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NELAP Certification Number: CA00046

DoD-ELAP Certification Number 4064.01

State Certification Number:

March 05, 2023

Watson Tanji
AECOM Honolulu
1001 Bishop Street, Suite 1600
Honolulu, HI 96813

RE: Red Hill AFFF Assessment Sampling
23B0184

Enclosed are the results of analyses for samples received by our laboratory on 2/24/2023. If you have any questions concerning this report, please feel free to contact me.

I certify that this data package is in compliance with the terms and conditions of the contract, both technically and for completeness. These test results meet all requirements of NELAC and DoD QSM. Release of the hard copy has been authorized by the Laboratory Manager or designee, as verified by the following signature.

Sincerely,

Paula McCartney For Karen Volpendesta
Project Manager

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Project: Red Hill AFFF Assessment Sampling
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Project Manager: Watson Tanji

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Data Validatable Report

Analysis Case Narrative

EPA 1633: Manual integrations were performed for this method in accordance with APPL's SOP. Chromatograms after manual integration are enclosed for specific samples and analytes. Abbreviated flags for technical justification are listed on the chromatogram.

Some extracted internal standards recovered outside of control limits in some samples; these samples were diluted and recovered in control, unless stated otherwise.

The analyte 6:2FTS recovered above the upper control limit in the SC00861-LCV1.

The analyte 9CI-PF3ONS recovered above the upper control limit in the SC00861-CCV1.

Samples in this Report

Lab ID	Sample	Matrix	Date Sampled	Date Received
23B0184-01	AF-RHMW03-WGN01LF-2302W3	Water	02/23/2023 12:15	02/24/2023
23B0184-02	AF-RHMW02-WGN01LF-2302W3	Water	02/23/2023 10:50	02/24/2023

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Containers Received

Lab ID	Container Type	Count	Preservation Check
23B0184-01	500mL P	2	
23B0184-02	500mL P	2	

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Project: Red Hill AFFF Assessment Sampling
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Reported: 03/05/2023 12:44

Sample Results

Sample: AF-RHMW03-WGN01LF-2302W3
23B0184-01 (Water)

Per- and Polyfluoroalkyl Substances

Analyte	Result /Qual	LOQ	LOD	DL	Units	Date Analyzed	DF	Method	Prep Batch
PFBA	1.6	1.5	0.76	0.20	ng/L	03/02/23	1	EPA 1633	BCB0459
PFPEA	3.1	0.76	0.38	0.062	ng/L	03/02/23	1	EPA 1633	BCB0459
PFHXA	1.8	0.38	0.19	0.052	ng/L	03/02/23	1	EPA 1633	BCB0459
PFHPA	1.6	0.38	0.19	0.039	ng/L	03/02/23	1	EPA 1633	BCB0459
PFOA	0.25 J	0.38	0.19	0.14	ng/L	03/02/23	1	EPA 1633	BCB0459
PFNA	0.093 J	0.38	0.19	0.078	ng/L	03/02/23	1	EPA 1633	BCB0459
PFDA	0.13 J	0.38	0.19	0.096	ng/L	03/02/23	1	EPA 1633	BCB0459
PFUnA	0.19 U IR1,	0.38	0.19	0.15	ng/L	03/02/23	1	EPA 1633	BCB0459
PFDOA	0.19 U	0.38	0.19	0.11	ng/L	03/02/23	1	EPA 1633	BCB0459
PFTRDA	0.28 U	0.38	0.28	0.19	ng/L	03/02/23	1	EPA 1633	BCB0459
PFTEDA	0.19 U IR2,	0.38	0.19	0.19	ng/L	03/02/23	1	EPA 1633	BCB0459
PFBS	0.19 U	0.38	0.19	0.035	ng/L	03/02/23	1	EPA 1633	BCB0459
PFPEs	0.19 U	0.38	0.19	0.059	ng/L	03/02/23	1	EPA 1633	BCB0459
PFHXS	0.19 U	0.38	0.19	0.030	ng/L	03/02/23	1	EPA 1633	BCB0459
PFHPS	0.19 U	0.38	0.19	0.049	ng/L	03/02/23	1	EPA 1633	BCB0459
PFOS	0.075 J	0.38	0.19	0.060	ng/L	03/02/23	1	EPA 1633	BCB0459
PFNS	0.19 U	0.38	0.19	0.12	ng/L	03/02/23	1	EPA 1633	BCB0459
PFDS	0.19 U	0.38	0.19	0.14	ng/L	03/02/23	1	EPA 1633	BCB0459
PFDOS	0.19 U	0.38	0.19	0.12	ng/L	03/02/23	1	EPA 1633	BCB0459
4:2FTS	0.76 U	1.5	0.76	0.28	ng/L	03/02/23	1	EPA 1633	BCB0459
6:2FTS	20	1.5	0.76	0.30	ng/L	03/02/23	1	EPA 1633	BCB0459
8:2FTS	0.76 U	1.5	0.76	0.078	ng/L	03/02/23	1	EPA 1633	BCB0459
PFOSA	0.19 U	0.38	0.19	0.099	ng/L	03/02/23	1	EPA 1633	BCB0459
NMeFOSA	0.76 U	1.5	0.76	0.45	ng/L	03/02/23	1	EPA 1633	BCB0459
NEtFOSA	0.76 U	1.5	0.76	0.39	ng/L	03/02/23	1	EPA 1633	BCB0459
NMeFOSAA	0.19 U	0.38	0.19	0.10	ng/L	03/02/23	1	EPA 1633	BCB0459
NEtFOSAA	0.19 U	0.38	0.19	0.11	ng/L	03/02/23	1	EPA 1633	BCB0459
NMeFOSE	1.1 U	1.5	1.1	0.96	ng/L	03/02/23	1	EPA 1633	BCB0459
NEtFOSE	1.1 U	1.5	1.1	0.99	ng/L	03/02/23	1	EPA 1633	BCB0459
HFPO-DA	0.38 U	0.76	0.38	0.17	ng/L	03/02/23	1	EPA 1633	BCB0459
ADONA	0.38 U	0.76	0.38	0.12	ng/L	03/02/23	1	EPA 1633	BCB0459
PFEESA	0.38 U	0.76	0.38	0.10	ng/L	03/02/23	1	EPA 1633	BCB0459
PFMPA	0.38 U	0.76	0.38	0.051	ng/L	03/02/23	1	EPA 1633	BCB0459
PFMBA	0.38 U	0.76	0.38	0.086	ng/L	03/02/23	1	EPA 1633	BCB0459
NFDHA	0.38 U	0.76	0.38	0.29	ng/L	03/02/23	1	EPA 1633	BCB0459
9CL-PF3ONS	0.38 U	0.76	0.38	0.20	ng/L	03/02/23	1	EPA 1633	BCB0459
11CL-PF3OUDS	0.38 U	0.76	0.38	0.20	ng/L	03/02/23	1	EPA 1633	BCB0459
3:3FTCA	0.76 U	1.5	0.76	0.54	ng/L	03/02/23	1	EPA 1633	BCB0459
5:3FTCA	0.76 U	1.5	0.76	0.42	ng/L	03/02/23	1	EPA 1633	BCB0459
7:3FTCA	0.76 U	1.5	0.76	0.52	ng/L	03/02/23	1	EPA 1633	BCB0459
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Surrogate: 13C4-PFBA	93.0%		10-130			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C5-PFPEA	69.6%		35-150			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C5-PFHXA	90.2%		55-150			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C4-PFHPA	86.4%		55-150			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C8-PFOA	96.1%		60-140			03/02/23	1	EPA 1633	BCB0459

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Reported: 03/05/2023 12:44

Sample Results

(Continued)

Sample: AF-RHMW03-WGN01LF-2302W3 (Continued)

23B0184-01 (Water)

Per- and Polyfluoroalkyl Substances (Continued)

Analyte	Result /Qual	LOQ	LOD	DL	Units	Date Analyzed	DF	Method	Prep Batch
Surrogate: 13C9-PFNA	108%		55-140			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C6-PFDA	104%		50-140			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C7-PFUnA	100%		30-140			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C2-PFDOA	77.8%		10-150			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C2-PFTEDA	63.3%		10-130			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C3-PFBS	92.8%		55-150			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C3-PFHXS	92.3%		55-150			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C8-PFOS	92.3%		45-140			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C2-4:2FTS	186%		60-200			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C2-6:2FTS	152%		60-200			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C2-8:2FTS	211% S2		50-200			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C2-8:2FTS	141%		50-200			03/02/23	10	EPA 1633	BCB0459
Surrogate: 13C8-PFOA	70.6%		30-130			03/02/23	1	EPA 1633	BCB0459
Surrogate: D3-NMEFOA	53.0%		15-130			03/02/23	1	EPA 1633	BCB0459
Surrogate: D5-NETFOA	52.6%		10-130			03/02/23	1	EPA 1633	BCB0459
Surrogate: D3-NMEFOSAA	164%		45-200			03/02/23	1	EPA 1633	BCB0459
Surrogate: D5-NETFOSAA	147%		10-200			03/02/23	1	EPA 1633	BCB0459
Surrogate: D7-NMEFOSE	48.8%		10-150			03/02/23	1	EPA 1633	BCB0459
Surrogate: D9-NETFOSE	55.8%		10-150			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C3-HFPO-DA	96.5%		25-160			03/02/23	1	EPA 1633	BCB0459

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Project: Red Hill AFFF Assessment Sampling
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Sample Results (Continued)

Sample: AF-RHMW02-WGN01LF-2302W3
23B0184-02 (Water)

Per- and Polyfluoroalkyl Substances

Analyte	Result /Qual	LOQ	LOD	DL	Units	Date Analyzed	DF	Method	Prep Batch
PFBA	0.80 U	1.6	0.80	0.21	ng/L	03/02/23	1	EPA 1633	BCB0459
PFPEA	2.5	0.80	0.40	0.065	ng/L	03/02/23	1	EPA 1633	BCB0459
PFHXA	0.37 J IR2,	0.40	0.20	0.055	ng/L	03/02/23	1	EPA 1633	BCB0459
PFHPA	0.44 IR1	0.40	0.20	0.041	ng/L	03/02/23	1	EPA 1633	BCB0459
PFOA	0.20 U	0.40	0.20	0.15	ng/L	03/02/23	1	EPA 1633	BCB0459
PFNA	0.20 U	0.40	0.20	0.081	ng/L	03/02/23	1	EPA 1633	BCB0459
PFDA	0.20 U	0.40	0.20	0.10	ng/L	03/02/23	1	EPA 1633	BCB0459
PFUnA	0.20 U IR2,	0.40	0.20	0.16	ng/L	03/02/23	1	EPA 1633	BCB0459
PFDOA	0.20 U	0.40	0.20	0.11	ng/L	03/02/23	1	EPA 1633	BCB0459
PFTRDA	0.30 U	0.40	0.30	0.20	ng/L	03/02/23	1	EPA 1633	BCB0459
PFTEDA	0.20 U	0.40	0.20	0.20	ng/L	03/02/23	1	EPA 1633	BCB0459
PFBS	0.20 U	0.40	0.20	0.037	ng/L	03/02/23	1	EPA 1633	BCB0459
PFPEs	0.20 U	0.40	0.20	0.062	ng/L	03/02/23	1	EPA 1633	BCB0459
PFHXS	0.20 U	0.40	0.20	0.031	ng/L	03/02/23	1	EPA 1633	BCB0459
PFHPS	0.20 U	0.40	0.20	0.051	ng/L	03/02/23	1	EPA 1633	BCB0459
PFOS	0.20 U	0.40	0.20	0.063	ng/L	03/02/23	1	EPA 1633	BCB0459
PFNS	0.20 U	0.40	0.20	0.12	ng/L	03/02/23	1	EPA 1633	BCB0459
PFDS	0.20 U	0.40	0.20	0.15	ng/L	03/02/23	1	EPA 1633	BCB0459
PFDOS	0.20 U	0.40	0.20	0.12	ng/L	03/02/23	1	EPA 1633	BCB0459
4:2FTS	0.80 U	1.6	0.80	0.29	ng/L	03/02/23	1	EPA 1633	BCB0459
6:2FTS	4.3	1.6	0.80	0.31	ng/L	03/02/23	1	EPA 1633	BCB0459
8:2FTS	0.80 U	1.6	0.80	0.082	ng/L	03/02/23	1	EPA 1633	BCB0459
PFOSA	0.20 U	0.40	0.20	0.10	ng/L	03/02/23	1	EPA 1633	BCB0459
NMeFOSA	0.80 U	1.6	0.80	0.47	ng/L	03/02/23	1	EPA 1633	BCB0459
NEtFOSA	0.80 U	1.6	0.80	0.41	ng/L	03/02/23	1	EPA 1633	BCB0459
NMeFOSAA	0.20 U	0.40	0.20	0.11	ng/L	03/02/23	1	EPA 1633	BCB0459
NEtFOSAA	0.20 U	0.40	0.20	0.11	ng/L	03/02/23	1	EPA 1633	BCB0459
NMeFOSE	1.2 U	1.6	1.2	1.0	ng/L	03/02/23	1	EPA 1633	BCB0459
NEtFOSE	1.2 U	1.6	1.2	1.0	ng/L	03/02/23	1	EPA 1633	BCB0459
HFPO-DA	0.40 U	0.80	0.40	0.17	ng/L	03/02/23	1	EPA 1633	BCB0459
ADONA	0.40 U	0.80	0.40	0.12	ng/L	03/02/23	1	EPA 1633	BCB0459
PFEESA	0.40 U	0.80	0.40	0.11	ng/L	03/02/23	1	EPA 1633	BCB0459
PFMPA	0.40 U	0.80	0.40	0.054	ng/L	03/02/23	1	EPA 1633	BCB0459
PFMBA	0.40 U	0.80	0.40	0.090	ng/L	03/02/23	1	EPA 1633	BCB0459
NFDHA	0.40 U	0.80	0.40	0.30	ng/L	03/02/23	1	EPA 1633	BCB0459
9CL-PF3ONS	0.40 U	0.80	0.40	0.21	ng/L	03/02/23	1	EPA 1633	BCB0459
11CL-PF3OUDS	0.40 U	0.80	0.40	0.20	ng/L	03/02/23	1	EPA 1633	BCB0459
3:3FTCA	0.80 U	1.6	0.80	0.57	ng/L	03/02/23	1	EPA 1633	BCB0459
5:3FTCA	0.80 U	1.6	0.80	0.44	ng/L	03/02/23	1	EPA 1633	BCB0459
7:3FTCA	0.80 U	1.6	0.80	0.55	ng/L	03/02/23	1	EPA 1633	BCB0459
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Surrogate: 13C4-PFBA	80.2%		10-130			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C5-PFPEA	58.2%		35-150			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C5-PFHXA	82.0%		55-150			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C4-PFHPA	84.3%		55-150			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C8-PFOA	85.2%		60-140			03/02/23	1	EPA 1633	BCB0459

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Project Manager: Watson Tanji

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Sample Results

(Continued)

Sample: AF-RHMW02-WGN01LF-2302W3 (Continued)

23B0184-02 (Water)

Per- and Polyfluoroalkyl Substances (Continued)

Analyte	Result /Qual	LOQ	LOD	DL	Units	Date Analyzed	DF	Method	Prep Batch
Surrogate: 13C9-PFNA	93.6%		55-140			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C6-PFDA	98.9%		50-140			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C7-PFUnA	97.5%		30-140			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C2-PFDOA	90.6%		10-150			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C2-PFTEDA	61.9%		10-130			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C3-PFBS	81.4%		55-150			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C3-PFHXS	90.3%		55-150			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C8-PFOS	95.3%		45-140			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C2-4:2FTS	167%		60-200			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C2-6:2FTS	154%		60-200			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C2-8:2FTS	240% S2		50-200			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C2-8:2FTS	92.8%		50-200			03/02/23	10	EPA 1633	BCB0459
Surrogate: 13C8-PFOA	68.4%		30-130			03/02/23	1	EPA 1633	BCB0459
Surrogate: D3-NMEFOA	47.5%		15-130			03/02/23	1	EPA 1633	BCB0459
Surrogate: D5-NETFOA	43.8%		10-130			03/02/23	1	EPA 1633	BCB0459
Surrogate: D3-NMEFOSAA	154%		45-200			03/02/23	1	EPA 1633	BCB0459
Surrogate: D5-NETFOSAA	157%		10-200			03/02/23	1	EPA 1633	BCB0459
Surrogate: D7-NMEFOSE	49.4%		10-150			03/02/23	1	EPA 1633	BCB0459
Surrogate: D9-NETFOSE	65.1%		10-150			03/02/23	1	EPA 1633	BCB0459
Surrogate: 13C3-HFPO-DA	87.7%		25-160			03/02/23	1	EPA 1633	BCB0459

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Project: Red Hill AFFF Assessment Sampling
Project Number: Red Hill AFFF Assessment Sampling
Project Manager: Watson Tanji

Reported: 03/05/2023 12:44

Quality Control

Per- and Polyfluoroalkyl Substances

Analyte	Result/Qual	LOQ	LOD	MDL	Spike Level	Source Result	%REC	%REC Limits	RPD	RPD Limit
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Method: EPA 1633

Batch: BCB0459 - EPA 1633

Blank (BCB0459-BLK1)

Prepared: 02/28/23 08:24 Analyzed: 03/02/23 10:59

	ng/L			
PFBA	0.80 U	1.6	0.80	0.21
PFPEA	0.40 U	0.80	0.40	0.065
PFHXA	0.20 U	0.40	0.20	0.055
PFHPA	0.20 U	0.40	0.20	0.041
PFOA	0.20 U	0.40	0.20	0.15
PFNA	0.20 U	0.40	0.20	0.082
PFDA	0.20 U	0.40	0.20	0.10
PFUnA	0.20 U	0.40	0.20	0.16
PFDOA	0.20 U	0.40	0.20	0.11
PFTDA	0.30 U	0.40	0.30	0.20
PFTEDA	0.20 U	0.40	0.20	0.20
PFBS	0.20 U	0.40	0.20	0.037
PFPEs	0.20 U	0.40	0.20	0.063
PFHXS	0.20 U	0.40	0.20	0.032
PFHPS	0.20 U	0.40	0.20	0.051
PFOS	0.102 J MI2,	0.40	0.20	0.064
PFNS	0.20 U	0.40	0.20	0.12
PFDS	0.20 U	0.40	0.20	0.15
PFDOS	0.20 U	0.40	0.20	0.12
4:2FTS	0.80 U	1.6	0.80	0.29
6:2FTS	0.80 U	1.6	0.80	0.31
8:2FTS	0.80 U	1.6	0.80	0.082
PFOSA	0.20 U	0.40	0.20	0.10
NMeFOSA	0.80 U	1.6	0.80	0.47
NEtFOSA	0.80 U	1.6	0.80	0.41
NMeFOSAA	0.20 U	0.40	0.20	0.11
NEtFOSAA	0.20 U	0.40	0.20	0.11
NMeFOSE	1.2 U	1.6	1.2	1.0
NEtFOSE	1.2 U	1.6	1.2	1.0
HFPO-DA	0.40 U	0.80	0.40	0.17
ADONA	0.40 U	0.80	0.40	0.12
PFEESA	0.40 U	0.80	0.40	0.11
PFMPA	0.40 U	0.80	0.40	0.054
PFMBA	0.40 U	0.80	0.40	0.091
NFDHA	0.40 U	0.80	0.40	0.30
9CL-PF3ONS	0.40 U	0.80	0.40	0.21
11CL-PF3OUDS	0.40 U	0.80	0.40	0.21
3:3FTCA	0.80 U	1.6	0.80	0.57
5:3FTCA	0.80 U	1.6	0.80	0.44
7:3FTCA	0.80 U	1.6	0.80	0.55

Surrogates

13C4-PFBA	30.3	32.0	94.6	10-130
13C5-PFPEA	14.1	16.0	88.1	35-150
13C5-PFHXA	7.60	8.00	95.1	55-150

AECOM Honolulu
1001 Bishop Street, Suite 1600
Honolulu, HI 96813

Project: Red Hill AFFF Assessment Sampling
Project Number: Red Hill AFFF Assessment Sampling
Project Manager: Watson Tanji

Reported: 03/05/2023 12:44

Quality Control (Continued)

Per- and Polyfluoroalkyl Substances (Continued)

Analyte	Result/Qual	LOQ	LOD	MDL	Spike Level	Source Result	%REC	%REC Limits	RPD	RPD Limit
Blank (BCB0459-BLK1)					Prepared: 02/28/23 08:24 Analyzed: 03/02/23 10:59					
	ng/L									
Surrogates										
13C4-PFHPA	7.57				8.00		94.6	55-150		
13C8-PFOA	7.93				8.00		99.1	60-140		
13C9-PFNA	4.13				4.00		103	55-140		
13C6-PFDA	3.88				4.00		96.9	50-140		
13C7-PFUnA	3.85				4.00		96.3	30-140		
13C2-PFDOA	3.98				4.00		99.5	10-150		
13C2-PFTEDA	3.70				4.00		92.5	10-130		
13C3-PFBS	8.09				8.00		101	55-150		
13C3-PFHXS	7.21				8.00		90.1	55-150		
13C8-PFOS	7.55				8.00		94.4	45-140		
13C2-4:2FTS	17.1				16.0		107	60-200		
13C2-6:2FTS	14.7				16.0		92.2	60-200		
13C2-8:2FTS	17.1				16.0		107	50-200		
13C8-PFOSA	7.45				8.00		93.1	30-130		
D3-NMEFOSA	4.00				8.00		50.0	15-130		
D5-NETFOSA	3.78				8.00		47.2	10-130		
D3-NMEFOSAA	17.3				16.0		108	45-200		
D5-NETFOSAA	16.9				16.0		105	10-200		
D7-NMEFOSE	45.9				80.0		57.4	10-150		
D9-NETFOSE	61.4				80.0		76.7	10-150		
13C3-HFPO-DA	29.3				32.0		91.6	25-160		

LCS (BCB0459-BS1)

Prepared: 02/28/23 08:24 Analyzed: 03/02/23 11:12

	ng/L									
PFBA	15.6				16.0		97.8	58-148		
PFPEA	7.64				8.00		95.4	54-152		
PFHXA	3.61				4.00		90.2	55-152		
PFHPA	3.47				4.00		86.8	54-154		
PFOA	3.68				4.00		91.9	52-161		
PFNA	4.21				4.00		105	59-149		
PFDA	4.00				4.00		99.9	52-147		
PFUnA	3.57				4.00		89.2	48-159		
PFDOA	3.98				4.00		99.5	64-142		
PFTRDA	3.96				4.00		98.9	49-148		
PFTEDA	3.41				4.00		85.3	47-161		
PFBS	3.34				3.54		94.2	62-144		
PFPEs	3.99				3.76		106	59-151		
PFHXS	3.66				3.66		100	57-146		
PFHPS	3.51				3.82		91.8	55-152		
PFOS	3.67				3.72		98.7	58-149		
PFNS	3.90				3.84		101	52-148		
PFDS	3.76				3.86		97.4	51-147		
PFDOS	4.26				3.88		110	36-145		
4:2FTS	13.8				15.0		92.0	67-146		
6:2FTS	16.3				15.2		107	61-151		
8:2FTS	14.9				15.4		97.0	63-152		
PFOSA	4.48				4.00		112	61-148		

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Project: Red Hill AFFF Assessment Sampling
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Project Manager: Watson Tanji

Reported: 03/05/2023 12:44

Quality Control (Continued)

Per- and Polyfluoroalkyl Substances (Continued)

Analyte	Result/Qual	LOQ	LOD	MDL	Spike Level	Source Result	%REC	%REC Limits	RPD	RPD Limit
LCS (BCB0459-BS1)					Prepared: 02/28/23 08:24 Analyzed: 03/02/23 11:12					
	ng/L									
NMeFOSA	15.3				16.0		95.4	63-145		
NEtFOSA	16.9				16.0		106	65-139		
NMeFOSAA	4.00				4.00		100	58-144		
NEtFOSAA	4.05				4.00		101	59-146		
NMeFOSE	16.2				16.0		101	71-136		
NEtFOSE	15.5				16.0		97.2	69-137		
HFPO-DA	7.82				8.00		97.8	63-144		
ADONA	8.03				7.56		106	68-146		
PFEESA	6.83				7.12		95.9	56-151		
PFMPA	6.84				8.00		85.6	51-145		
PFMBA	8.10				8.00		101	55-148		
NFDHA	6.63				8.00		82.9	48-161		
9CL-PF3ONS	8.85				7.48		118	56-156		
11CL-PF3OUDS	8.54				7.56		113	46-156		
3:3FTCA	13.2				16.0		82.7	62-129		
5:3FTCA	11.3				16.0		70.6	63-134		
7:3FTCA	12.4				16.0		77.4	50-138		
Surrogates										
13C4-PFBA	30.6				32.0		95.6	10-130		
13C5-PFPEA	13.8				16.0		86.2	35-150		
13C5-PFHXA	7.78				8.00		97.3	55-150		
13C4-PFHFA	7.16				8.00		89.5	55-150		
13C8-PFOA	7.31				8.00		91.4	60-140		
13C9-PFNA	3.96				4.00		99.0	55-140		
13C6-PFDA	4.09				4.00		102	50-140		
13C7-PFUnA	4.03				4.00		101	30-140		
13C2-PFDOA	4.11				4.00		103	10-150		
13C2-PFTEDA	3.98				4.00		99.4	10-130		
13C3-PFBS	8.84				8.00		110	55-150		
13C3-PFHXS	7.84				8.00		97.9	55-150		
13C8-PFOS	7.65				8.00		95.6	45-140		
13C2-4:2FTS	19.3				16.0		121	60-200		
13C2-6:2FTS	16.2				16.0		102	60-200		
13C2-8:2FTS	20.3				16.0		127	50-200		
13C8-PFOSA	6.78				8.00		84.8	30-130		
D3-NMEFOSA	4.34				8.00		54.2	15-130		
D5-NETFOSA	4.21				8.00		52.6	10-130		
D3-NMEFOSAA	17.7				16.0		111	45-200		
D5-NETFOSAA	15.9				16.0		99.2	10-200		
D7-NMEFOSE	45.8				80.0		57.3	10-150		
D9-NETFOSE	62.9				80.0		78.6	10-150		
13C3-HFPO-DA	26.5				32.0		82.9	25-160		

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Project: Red Hill AFFF Assessment Sampling

Project Number: Red Hill AFFF Assessment Sampling

Project Manager: Watson Tanji

Reported: 03/05/2023 12:44

Quality Control (Continued)

Per- and Polyfluoroalkyl Substances (Continued)

Analyte	Result/Qual	LOQ	LOD	MDL	Spike Level	Source Result	%REC	%REC Limits	RPD	RPD Limit
MRL Check (BCB0459-MRL1)					Prepared: 02/28/23 08:24 Analyzed: 03/02/23 11:25					
	ng/L									
PFBA	1.59 J				1.60		99.5	44-157		
PFPEA	0.916				0.800		114	57-148		
PFHXA	0.462				0.400		116	62-149		
PFHPA	0.391 J				0.400		97.6	56-150		
PFOA	0.492				0.400		123	57-161		
PFNA	0.297 J IR2,				0.400		74.2	53-157		
PFDA	0.498 IR2				0.400		124	43-158		
PFUnA	0.386 J IR2,				0.400		96.4	50-155		
PFDOA	0.451				0.400		113	60-141		
PFTRDA	0.321 J IR2,				0.400		80.4	52-140		
PFTEDA	0.410 IR1				0.400		102	52-156		
PFBS	0.364 J				0.354		103	63-145		
PFPEs	0.380 J				0.376		101	58-144		
PFHXS	0.379 J				0.366		104	44-158		
PFHPS	0.349 J				0.382		91.4	51-150		
PFOS	0.439				0.372		118	43-162		
PFNS	0.379 J				0.384		98.6	46-151		
PFDS	0.381 J				0.386		98.6	50-144		
PFDOS	0.388 J IR2,				0.388		100	30-138		
4:2FTS	1.54 J				1.50		103	52-158		
6:2FTS	1.97				1.52		130	48-158		
8:2FTS	1.65				1.54		108	46-165		
PFOSA	0.482				0.400		121	47-163		
NMeFOSA	1.65				1.60		103	54-155		
NEtFOSA	1.77				1.60		111	49-156		
NMeFOSAA	0.416				0.400		104	32-160		
NEtFOSAA	0.452				0.400		113	51-154		
NMeFOSE	1.73				1.60		108	56-151		
NEtFOSE	1.72				1.60		107	60-147		
HFPO-DA	0.796 J				0.800		99.5	58-154		
ADONA	0.787 J				0.756		104	61-148		
PFEESA	0.792 J				0.712		111	56-144		
PFMPA	0.761 J				0.800		95.1	48-150		
PFMBA	0.908				0.800		114	49-154		
NFDHA	0.725 J				0.800		90.6	47-160		
9CL-PF3ONS	0.875				0.748		117	44-167		
11CL-PF3OUDS	0.766 J				0.756		101	36-158		
3:3FTCA	1.60				1.60		100	32-161		
5:3FTCA	1.72				1.60		107	39-156		
7:3FTCA	1.12 J				1.60		69.9	36-149		
Surrogates										
13C4-PFBA	30.3				32.0		94.6	10-130		
13C5-PFPEA	14.8				16.0		92.3	35-150		
13C5-PFHXA	7.88				8.00		98.5	55-150		
13C4-PFHPA	7.58				8.00		94.7	55-150		
13C8-PFOA	7.90				8.00		98.7	60-140		
13C9-PFNA	3.68				4.00		91.9	55-140		

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Project: Red Hill AFFF Assessment Sampling
Project Number: Red Hill AFFF Assessment Sampling
Project Manager: Watson Tanji

Reported: 03/05/2023 12:44

Quality Control (Continued)

Per- and Polyfluoroalkyl Substances (Continued)

Analyte	Result/Qual	LOQ	LOD	MDL	Spike Level	Source Result	%REC	%REC Limits	RPD	RPD Limit
MRL Check (BCB0459-MRL1)					Prepared: 02/28/23 08:24 Analyzed: 03/02/23 11:25					
	ng/L									
Surrogates										
13C6-PFDA	4.09				4.00		102	50-140		
13C7-PFUnA	4.26				4.00		106	30-140		
13C2-PFDOA	3.98				4.00		99.6	10-150		
13C2-PFTEDA	3.64				4.00		91.0	10-130		
13C3-PFBS	7.76				8.00		97.0	55-150		
13C3-PFHXS	7.94				8.00		99.3	55-150		
13C8-PFOS	7.83				8.00		97.9	45-140		
13C2-4:2FTS	15.5				16.0		97.1	60-200		
13C2-6:2FTS	15.0				16.0		93.9	60-200		
13C2-8:2FTS	18.5				16.0		116	50-200		
13C8-PFOSA	7.12				8.00		89.0	30-130		
D3-NMEFOSA	4.35				8.00		54.4	15-130		
D5-NETFOSA	4.57				8.00		57.2	10-130		
D3-NMEFOSAA	18.2				16.0		113	45-200		
D5-NETFOSAA	16.9				16.0		106	10-200		
D7-NMEFOSE	46.8				80.0		58.6	10-150		
D9-NETFOSE	67.3				80.0		84.1	10-150		
13C3-HFPO-DA	31.0				32.0		97.0	25-160		

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Project: Red Hill AFFF Assessment Sampling
Project Number: Red Hill AFFF Assessment Sampling
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Reported: 03/05/2023 12:44

Notes and Definitions

Item	Definition
CV2	Calibration verification recovered above the upper control limit
IR1	Ion ratio below the lower control limit
IR2	Ion ratio above the upper control limit
IS1	Internal standard recovered below the lower control limit
J	Estimated value
MI2	Manual integration, non-target peak interference
S2	Surrogate recovered above the upper control limit
U	Not detected
Dry	Sample results reported on a dry weight basis.
DL	Dilution Factor
LOD	Limit of Detection
LOQ	Limit of Quantitation
DL	Detection Limit
*	Value outside control limits
RPD	Relative Percent Difference
%REC	Percent Recovery
Source	Sample that was matrix spiked or duplicated.

**WORK ORDER****23B0184**

Printed: 03/05/2023 12:44 pm

Project: Red Hill AFFF Assessment Sampling
Project Number: Red Hill AFFF Assessment Sampling
Project Manager: Karen Volpendesta
PO Number: 60697810

Report To:

AECOM Honolulu
 Watson Tanji
 1001 Bishop Street, Suite 1600
 Honolulu, HI 96813
 Phone: (808) 954-4512
 Fax: (808) 523-8950

Invoice To:

AECOM Honolulu
 Watson Tanji
 1001 Bishop Street, Suite 1600
 Honolulu, HI 96813
 Phone: (808) 954-4512
 Fax: (808) 523-8950

Date Received: 02/24/2023 01:40 PM
 Date Due: 03/03/2023 (5.00 day TAT)

Logged In By: Megan Salata
 Received By: Megan Salata

Analysis	Comments
23B0184-01 AF-RHMW03-WGN01LF-2302W3 [Water] Sampled 2/23/2023 12:15:00PM 1633 NONE	"Report relevant surrogates"
23B0184-02 AF-RHMW02-WGN01LF-2302W3 [Water] Sampled 2/23/2023 10:50:00AM 1633 NONE	"Report relevant surrogates"

23B0184**Sample Receipt Log**

Default Cooler

Samples Received at: **0.4°C**

Custody Seals	Yes	Were all containers sealed in separate bags?	Yes
Containers Intact	Yes	Did all containers arrive in good condition?	Yes
COC/Labels Agree	Yes	Correct containers/preserv. for tests indicated?	Yes
Preservation Confirmed	No	Sufficient volume sent for tests requested?	Yes
Received On Ice	Yes	Were bubbles absent in volatile samples?	No
Was a chain of custody received?	Yes	Sufficient remaining holding time for analyses?	Yes
COCs complete/signed in the appropriate places?	Yes	pH of non-VOA preserved containers documented?	No
Sample labels complete? Sample ID, date/time, etc.	Yes	Unpreserved vials received for VOA analysis?	No
Did all container labels agree with COCs?	Yes	If "yes", are unpreserved VOA vials noted on ARF?	No



APPL, Inc.
908 N Temperance
Clovis, CA 93611
www.applinc.com

ELECTRONIC CHAIN OF CUSTODY RECORD

Phone: (559) 275-2175

Fax: (559) 275-4422

C.O.C. 2302W3AFAL02

PLEASE PRINT

AECOM

Phone: 000-334-4312 / 003-190-4024

Address: 1001 Bishop St ste1600

Honolulu, HI 96813

Watson Tanji / Katie Abbott

Email: watson.tanji@aecom.com/katie.abbott@aecom.com

PLEASE PRINT

Invoice to:

AECOM

Company Name: **AECOM** Phone: _____

Address: _____

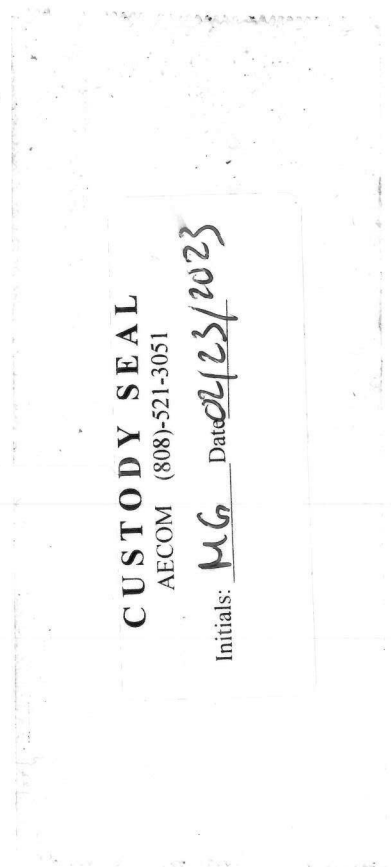
Fax: _____

Sheree Smith

Attn: Sheree Smith
Email: USAPimaging@aecom.com

[illegible]

Note: The first sampled date of the ARF will be used as the COC number unless indicated otherwise.



PFAS

SAMPLE DATA

FORM I

ANALYSIS DATA SHEET

AF-RHMW03-WGN01LF-2302W3

Laboratory:	APPL, LLC	Work Order:	23B0184
Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Matrix:	Water	Laboratory ID:	23B0184-01
		File ID:	S2023-03-02A (10)
Sampled:	02/23/23 12:15	Prepared:	02/28/23 08:33
		Analyzed:	03/02/23 11:38
Solids:		Preparation:	EPA 1633
		Dilution:	1
Initial/Final:	527.6 mL / 2 mL	Instrument:	Saphira
Batch:	BCB0459	Sequence:	SC00861
		Calibration:	2309009

COMPOUND	CONC. (ng/L)	LOQ	LOD	DL	Q
PFBA	1.6	1.5	0.76	0.20	
PFPEA	3.1	0.76	0.38	0.062	
PFHXA	1.8	0.38	0.19	0.052	
PFHPA	1.6	0.38	0.19	0.039	
PFOA	0.25 J	0.38	0.19	0.14	
PFNA	0.093 J	0.38	0.19	0.078	
PFDA	0.13 J	0.38	0.19	0.096	
PFUnA	0.19 U	0.38	0.19	0.15	IR1,
PFDOA	0.19 U	0.38	0.19	0.11	
PFTRDA	0.28 U	0.38	0.28	0.19	
PFTEDA	0.19 U	0.38	0.19	0.19	IR2,
PFBS	0.19 U	0.38	0.19	0.035	
PFPEs	0.19 U	0.38	0.19	0.059	
PFHXS	0.19 U	0.38	0.19	0.030	
PFHPS	0.19 U	0.38	0.19	0.049	
PFOS	0.075 J	0.38	0.19	0.060	
PFNS	0.19 U	0.38	0.19	0.12	
PFDS	0.19 U	0.38	0.19	0.14	
PFDOS	0.19 U	0.38	0.19	0.12	
4:2FTS	0.76 U	1.5	0.76	0.28	
6:2FTS	20	1.5	0.76	0.30	
8:2FTS	0.76 U	1.5	0.76	0.078	
PFOSA	0.19 U	0.38	0.19	0.099	
NMeFOSA	0.76 U	1.5	0.76	0.45	
NEtFOSA	0.76 U	1.5	0.76	0.39	
NMeFOSAA	0.19 U	0.38	0.19	0.10	
NEtFOSAA	0.19 U	0.38	0.19	0.11	
NMeFOSE	1.1 U	1.5	1.1	0.96	
NEtFOSE	1.1 U	1.5	1.1	0.99	
HFPO-DA	0.38 U	0.76	0.38	0.17	

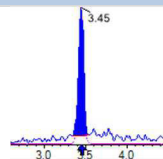
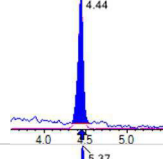
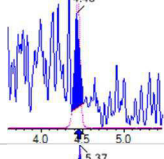
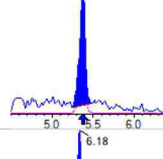
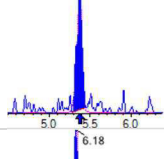
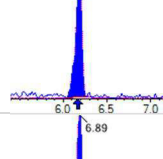
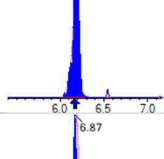
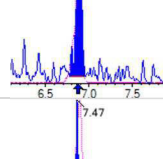
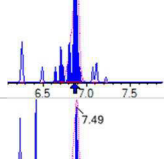
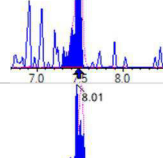
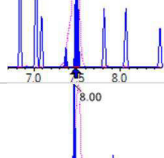
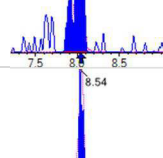
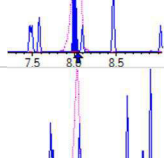
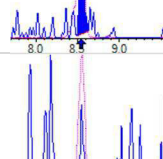
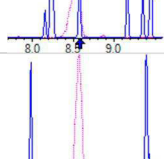
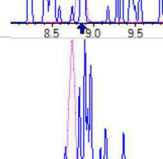
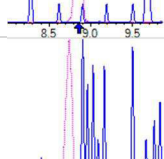
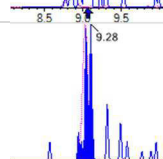
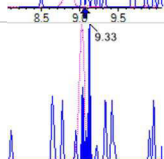
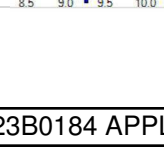
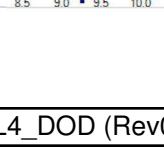
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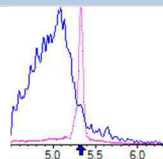
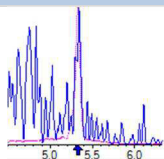
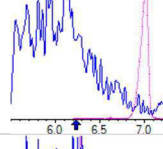
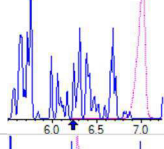
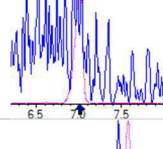
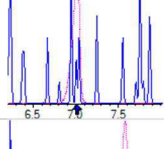
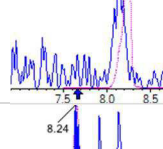
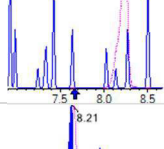
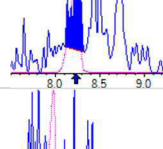
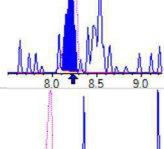
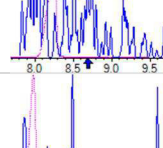
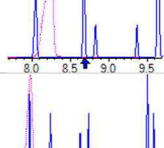
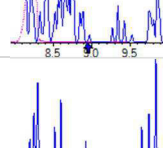
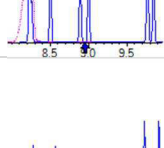
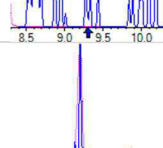
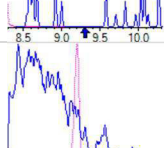
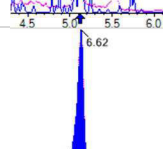
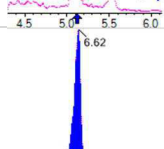
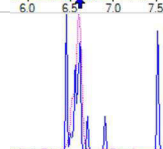
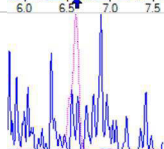
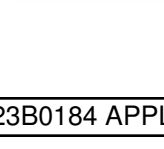
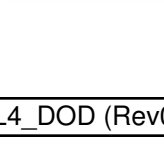
ANALYSIS DATA SHEET

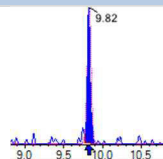
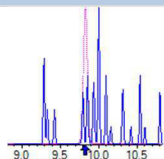
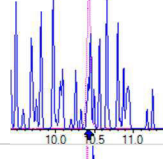
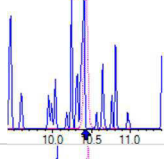
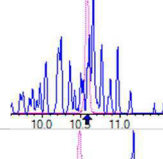
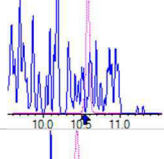
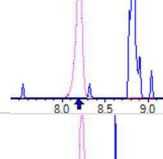
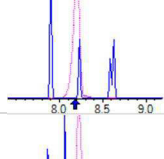
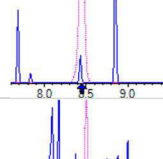
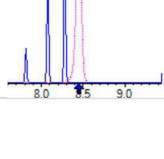
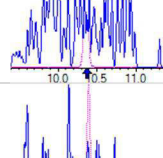
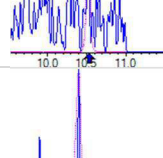
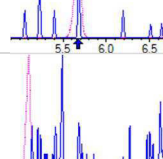
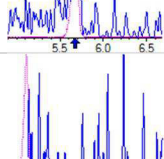
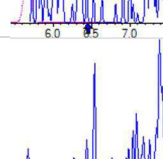
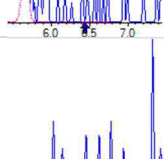
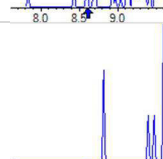
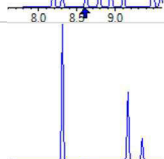
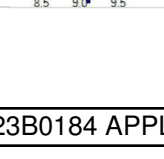
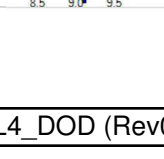
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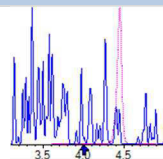
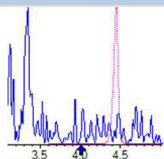
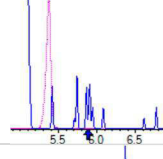
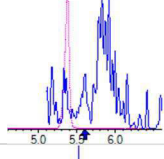
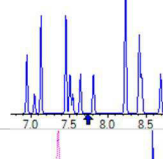
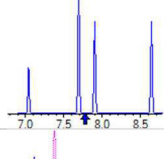
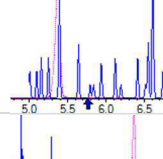
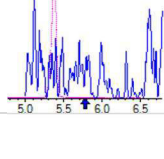
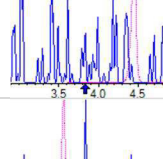
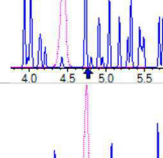
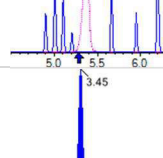
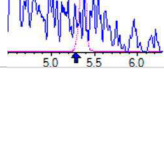
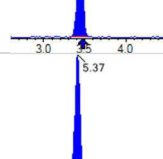
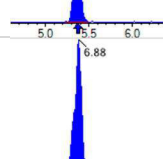
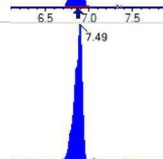
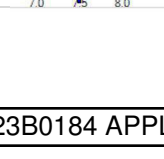
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Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Matrix:	Water	Laboratory ID:	23B0184-01
		File ID:	S2023-03-02A (10)
Sampled:	02/23/23 12:15	Prepared:	02/28/23 08:33
		Analyzed:	03/02/23 11:38
Solids:		Preparation:	EPA 1633
		Dilution:	1
Initial/Final:	527.6 mL / 2 mL	Instrument:	Saphira
Batch:	BCB0459	Sequence:	SC00861
		Calibration:	2309009

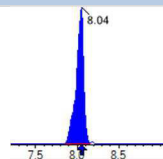
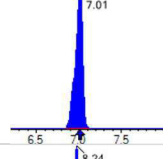
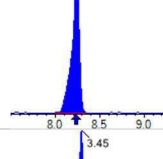
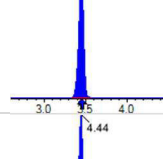
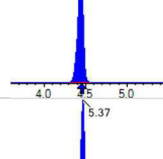
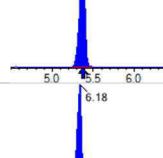
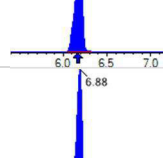
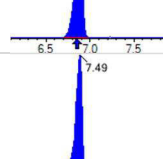
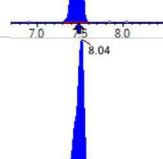
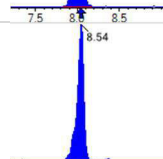
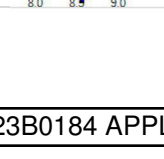
COMPOUND	CONC. (ng/L)	LOQ	LOD	DL	Q
ADONA	0.38 U	0.76	0.38	0.12	
PFEESA	0.38 U	0.76	0.38	0.10	
PFMPA	0.38 U	0.76	0.38	0.051	
PFMBA	0.38 U	0.76	0.38	0.086	
NFDHA	0.38 U	0.76	0.38	0.29	
9CL-PF3ONS	0.38 U	0.76	0.38	0.20	
11CL-PF3OUDS	0.38 U	0.76	0.38	0.20	
3:3FTCA	0.76 U	1.5	0.76	0.54	
5:3FTCA	0.76 U	1.5	0.76	0.42	
7:3FTCA	0.76 U	1.5	0.76	0.52	

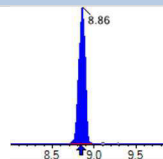
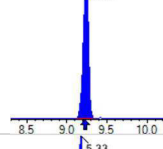
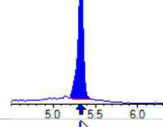
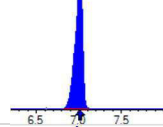
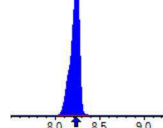
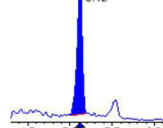
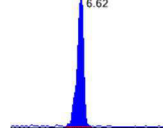
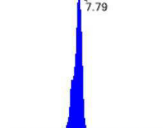
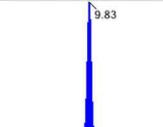
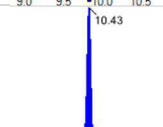
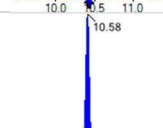
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT Ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) 44555	(3.45, 1.00) (0.00, N/A, 0.0)	55.3	N/A 0.0 0.0	0.4185	N/A			
PFPeA	(263.0 / 219.0) 237459 (263.0 / 69.0) 4528	(4.44, 1.00) (0.00, N/A, 0.6)	147.2 7.2	0.0191 160.0 162.1	0.8147	N/A			
PFHxA	(313.0 / 269.0) 236152 (313.0 / 119.0) 24088	(5.37, 1.00) (0.00, N/A, -0.1)	89.5 49.1	0.1020 109.0 98.7	0.4846	N/A			
PFHpA	(363.0 / 319.0) 171328 (363.0 / 169.0) 63772	(6.18, 1.00) (0.00, N/A, 0.0)	216.1 878040.4	0.3722 117.9 114.1	0.4113	N/A			
PFOA	(413.0 / 369.0) 39493 (413.0 / 169.0) 13545	(6.89, 1.00) (0.01, N/A, 1.2)	46.3 288603.1	0.3430 101.4 96.5	0.0670	N/A			
PFNA	(463.0 / 419.0) 13777 (463.0 / 169.0) 1931	(7.47, 1.00) (-0.02, N/A, -1.6)	62.8 80385.9	0.1402 59.9 60.5	0.0244	N/A			
PFDA	(513.0 / 469.0) 27370 (513.0 / 169.0) 2550	(8.01, 1.00) (-0.03, N/A, 0.9)	28.8 187.1	0.0932 75.2 68.8	0.0351	N/A			
PFUnA	(563.0 / 519.0) 11769 (563.0 / 169.0) N/A	(8.54, 1.00) (-0.01, N/A, #Value!)	35.3 N/A	N/A 0.0 0.0	0.0178	N/A			IR1,
PFDaA	(613.0 / 569.0) N/A (613.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFTTrDA	(663.0 / 619.0) N/A (663.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFTeDA	(713.0 / 669.0) 8802 (713.0 / 169.0) 3035	(9.28, 1.00) (0.04, N/A, -3.0)	46.9 135.3	0.3448 157.7 156.5	0.0209	N/A			IR2,

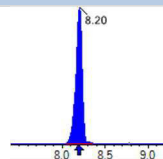
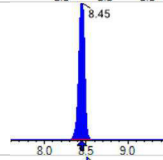
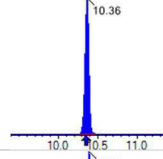
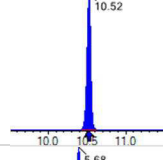
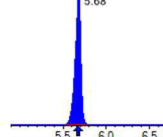
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) N/A (299.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFPeS	(349.0 / 80.0) N/A (349.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHxS	(399.0 / 80.0) N/A (399.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHpS	(449.0 / 80.0) N/A (449.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFOS	(499.0 / 80.0) 43105 (499.0 / 99.0) 13065	(8.24 , 1.00) (0.00 , N/A , 1.9)	14.6 22.4	0.3031 137.8 131.5	0.0198	N/A			
PFNS	(549.0 / 80.0) N/A (549.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDS	(599.0 / 80.0) N/A (599.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDoS	(699.0 / 80.0) N/A (699.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
4:2FTS	(327.0 / 307.0) N/A (327.0 / 81.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
6:2FTS	(427.0 / 407.0) 1280021 (427.0 / 81.0) 891962	(6.62 , 1.00) (0.00 , N/A , 0.0)	1299.9 384.8	0.6968 82.2 90.9	5.1662	N/A			
8:2FTS	(527.0 / 507.0) N/A (527.0 / 81.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) 15674 (498.0 / 478.0) N/A	(9.82 , 1.00) (-0.01 , N/A , #Value!)	97.1 N/A	N/A 0.0 0.0	0.0073	N/A			
NMeFOSA	(512.0 / 219.0) N/A (512.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NEtFOSA	(526.0 / 219.0) N/A (526.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NMeFOSAA	(570.0 / 419.0) N/A (570.0 / 483.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NEtFOSAA	(584.0 / 419.0) N/A (584.0 / 526.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NMeFOSE	(616.0 / 59.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
NEtFOSE	(630.0 / 59.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
HFPO-DA	(285.0 / 169.0) N/A (285.0 / 185.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
ADONA	(377.0 / 85.0) N/A (377.0 / 251.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
9Cl-Pf3ONS	(531.0 / 351.0) N/A (533.0 / 353.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
11Cl-Pf3OUDS	(631.0 / 451.0) N/A (633.0 / 453.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) N/A (241.0 / 117.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
5:3FTCA	(341.0 / 236.7) N/A (341.0 / 217.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
7:3FTCA	(441.0 / 317.0) N/A (441.0 / 337.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFEESA	(315.0 / 135.0) N/A (315.0 / 83.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFMPA	(229.0 / 85.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
PFMBA	(279.0 / 85.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
NFDHA	(295.0 / 201.0) N/A (295.0 / 85.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
13C3_PFBA_IIS	(216.0 / 172.0) 113798	(3.45, N/A) (N/A, -0.01, N/A)	582.3	N/A	0.3357 [1.0000]	33.6% { 54.3% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 426292	(5.37, N/A) (N/A, 0.00, N/A)	958.3	N/A	0.7167 [1.0000]	71.7% { 114.4% }			
13C4_PFOA_IIS	(417.0 / 372.0) 594675	(6.88, N/A) (N/A, 0.01, N/A)	1230.6	N/A	0.7311 [1.0000]	73.1% { 114.8% }			
13C5_PFNA_IIS	(468.0 / 423.0) 557290	(7.49, N/A) (N/A, 0.01, N/A)	634.5	N/A	0.7298 [1.0000]	73.0% { 101.2% }			

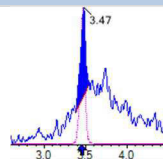
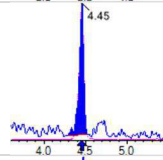
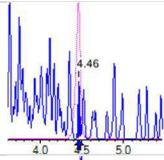
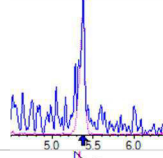
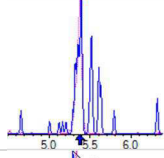
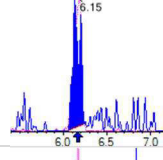
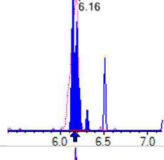
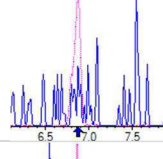
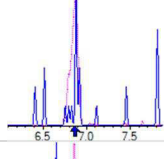
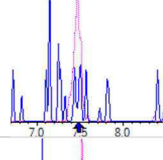
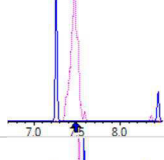
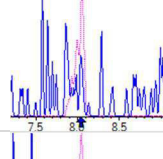
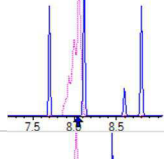
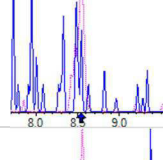
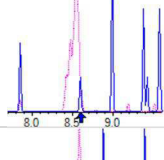
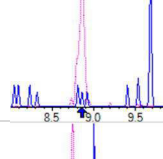
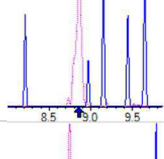
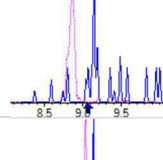
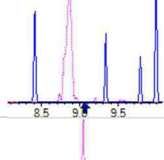
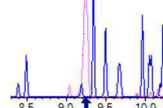
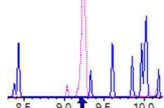
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 653387	(8.04 , N/A) (N/A , 0.01 , N/A)	664.8	N/A	0.9102 [1.0000]	91.0% { 136.5% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 894420	(7.01 , N/A) (N/A , 0.01 , N/A)	1377.7	N/A	0.7263 [1.0000]	72.6% { 101.1% }			
13C4_PFOS_IIS	(503.0 / 79.9) 1627797	(8.24 , N/A) (N/A , 0.01 , N/A)	626.6	N/A	0.9172 [1.0000]	91.7% { 126.2% }			
13C4_PFBA_EIS	(217.0 / 172.0) 1007193	(3.45 , N/A) (N/A , -0.01 , N/A)	2398.4	N/A	7.4432 [8.0000]	93.0% { 52.8% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 1323113	(4.44 , N/A) (N/A , -0.01 , N/A)	1986.7	N/A	2.7855 [4.0000]	69.6% { 75.5% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 1044419	(5.37 , N/A) (N/A , 0.00 , N/A)	1153.5	N/A	1.8032 [2.0000]	90.2% { 98.4% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 933032	(6.18 , N/A) (N/A , 0.02 , N/A)	2052.4	N/A	1.7272 [2.0000]	86.4% { 103.5% }			
13C8_PFOA_EIS	(421.0 / 376.0) 1223055	(6.88 , N/A) (N/A , 0.02 , N/A)	1038.5	N/A	1.9215 [2.0000]	96.1% { 115.2% }			
13C9_PFNA_EIS	(472.0 / 427.0) 597250	(7.49 , N/A) (N/A , 0.00 , N/A)	801.3	N/A	1.0754 [1.0000]	107.5% { 119.2% }			
13C6_PFDA_EIS	(519.0 / 474.0) 786107	(8.04 , N/A) (N/A , 0.01 , N/A)	825.1	N/A	1.0395 [1.0000]	103.9% { 125.7% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 750416	(8.54 , N/A) (N/A , 0.01 , N/A)	751.7	N/A	1.0016 [1.0000]	100.2% { 127.4% }			

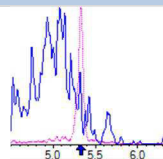
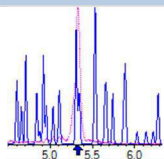
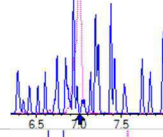
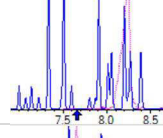
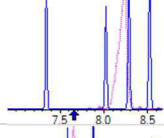
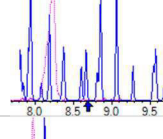
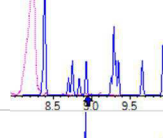
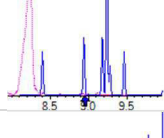
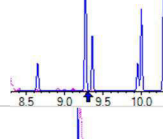
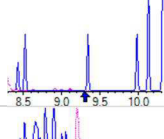
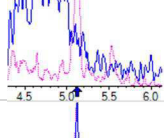
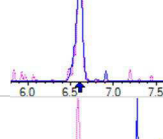
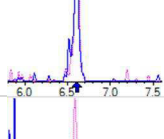
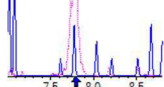
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 540547	(8.86 , N/A) (N/A , 0.01 , N/A)	702.9	N/A	0.7779 [1.0000]	77.8% { 120.7% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 457306	(9.24 , N/A) (N/A , 0.01 , N/A)	767.8	N/A	0.6331 [1.0000]	63.3% { 88.3% }			
13C3_PFBs_EIS	(302.0 / 80.0) 3353095	(5.33 , N/A) (N/A , 0.00 , N/A)	330.2	N/A	1.8552 [2.0000]	92.8% { 93.9% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 1743881	(7.01 , N/A) (N/A , 0.01 , N/A)	1036.7	N/A	1.8468 [2.0000]	92.3% { 99.5% }			
13C8_PFOS_EIS	(507.0 / 80.0) 3948803	(8.23 , N/A) (N/A , 0.01 , N/A)	1057.9	N/A	1.8465 [2.0000]	92.3% { 120.6% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 824481	(5.12 , N/A) (N/A , 0.00 , N/A)	142.8	N/A	7.4476 [4.0000]	186.2% { 204.5% }			S2,
13C2_6:2FTS_EIS	(429.0 / 81.0) 759688	(6.62 , N/A) (N/A , 0.02 , N/A)	371.7	N/A	6.0865 [4.0000]	152.2% { 178.0% }			S2,
13C2_8:2FTS_EIS	(529.0 / 81.0) 1113030	(7.79 , N/A) (N/A , 0.01 , N/A)	509.7	N/A	8.4327 [4.0000]	210.8% { 212.9% }			S2,
13C8_PFOA_EIS	(506.0 / 78.0) 5559063	(9.83 , N/A) (N/A , 0.01 , N/A)	2683.0	N/A	1.4112 [2.0000]	70.6% { 96.0% }			
D3_NMeFOA_EIS	(515.0 / 169.0) 896265	(10.43 , N/A) (N/A , 0.01 , N/A)	1679.0	N/A	1.0596 [2.0000]	53.0% { 67.5% }			
D5_NEtFOA_EIS	(531.0 / 169.0) 778901	(10.58 , N/A) (N/A , 0.01 , N/A)	2125.5	N/A	1.0516 [2.0000]	52.6% { 64.3% }			

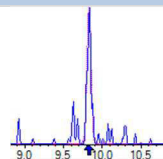
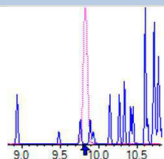
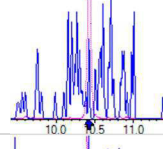
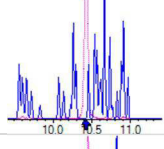
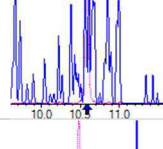
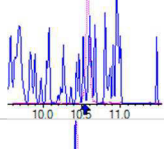
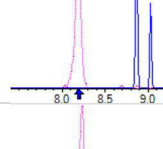
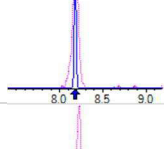
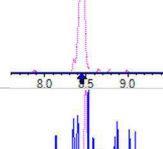
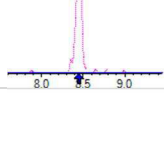
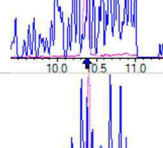
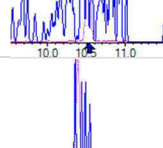
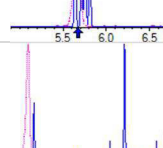
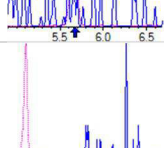
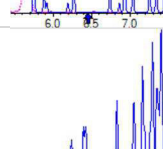
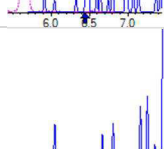
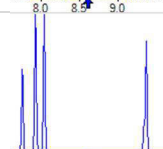
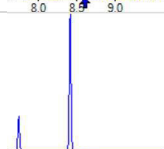
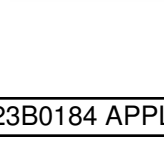
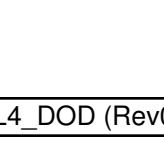
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT Ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1968287	(8.20 , N/A) (N/A , 0.01 , N/A)	1103.7	N/A	6.5581 [4.0000]	164.0% { 177.3% }			S2,
D5_EtFOSAA_EIS	(589.0 / 419.0) 1499321	(8.45 , N/A) (N/A , 0.01 , N/A)	3228.1	N/A	5.8664 [4.0000]	146.7% { 178.0% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 2811765	(10.36 , N/A) (N/A , 0.01 , N/A)	1950.6	N/A	9.7637 [20.0000]	48.8% { 65.7% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 3674642	(10.52 , N/A) (N/A , 0.01 , N/A)	1359.7	N/A	11.1550 [20.0000]	55.8% { 68.8% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 2810272	(5.68 , N/A) (N/A , 0.01 , N/A)	1925.2	N/A	7.7214 [8.0000]	96.5% { 101.0% }			

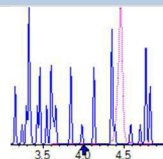
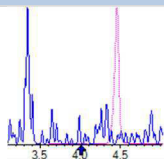
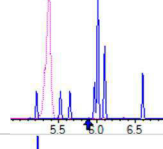
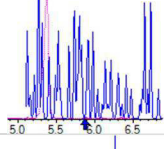
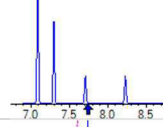
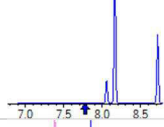
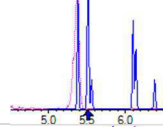
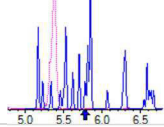
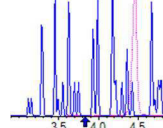
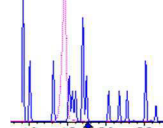
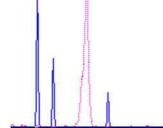
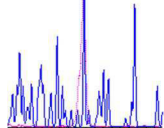
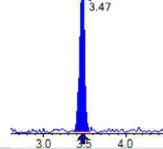
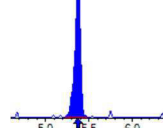
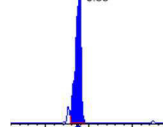
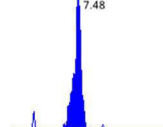
FORM I
ANALYSIS DATA SHEET
AF-RHMW03-WGN01LF-2302W3

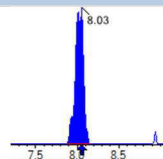
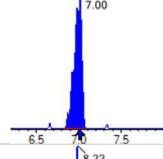
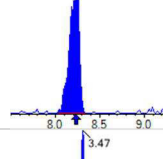
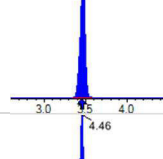
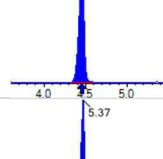
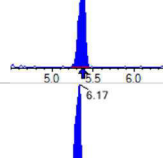
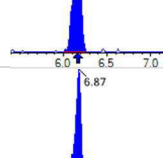
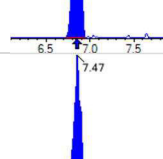
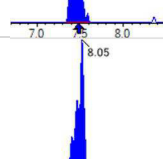
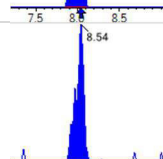
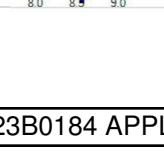
Laboratory:	APPL, LLC	Work Order:	23B0184		
Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling		
Matrix:	Water	Laboratory ID:	23B0184-01RE1	File ID:	S2023-03-02A (11)
Sampled:	02/23/23 12:15	Prepared:	02/28/23 08:33	Analyzed:	03/02/23 11:51
Solids:		Preparation:	EPA 1633	Dilution:	10
Initial/Final:	527.6 mL / 2 mL			Instrument:	Saphira
Batch:	BCB0459	Sequence:	SC00861	Calibration:	2309009

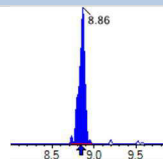
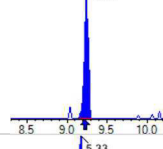
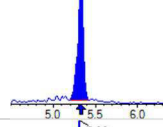
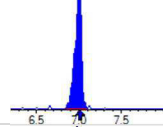
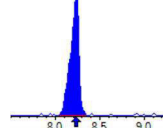
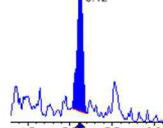
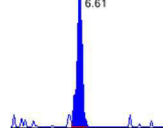
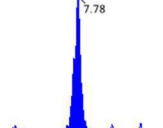
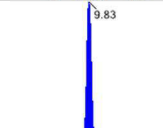
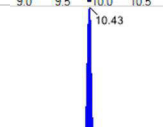
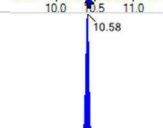
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) 8804	(3.47, 1.00) (0.01, N/A, 0.0)	20.0	N/A 0.0 0.0	0.0337	N/A			
PFPeA	(263.0 / 219.0) 39496 (263.0 / 69.0) 412	(4.45, 1.00) (0.00, N/A, -0.5)	80.6 4.5	0.0104 87.5 88.7	0.0897	N/A			
PFHxA	(313.0 / 269.0) N/A (313.0 / 119.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHpA	(363.0 / 319.0) 16988 (363.0 / 169.0) 5854	(6.15, 1.00) (-0.01, N/A, -0.3)	26.7 749.8	0.3446 109.2 105.7	0.0363	N/A			
PFOA	(413.0 / 369.0) N/A (413.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFNA	(463.0 / 419.0) N/A (463.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDA	(513.0 / 469.0) N/A (513.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFUnA	(563.0 / 519.0) N/A (563.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDaA	(613.0 / 569.0) N/A (613.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFTTrDA	(663.0 / 619.0) N/A (663.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFTeDA	(713.0 / 669.0) N/A (713.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

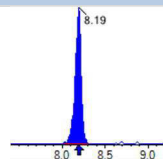
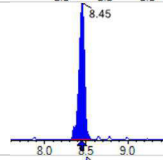
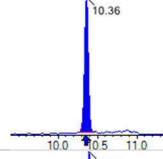
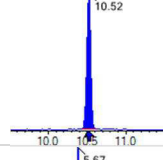
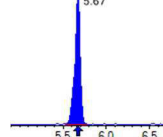
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) N/A (299.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFPeS	(349.0 / 80.0) N/A (349.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHxS	(399.0 / 80.0) N/A (399.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHpS	(449.0 / 80.0) N/A (449.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFOS	(499.0 / 80.0) N/A (499.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFNS	(549.0 / 80.0) N/A (549.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDS	(599.0 / 80.0) N/A (599.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDoS	(699.0 / 80.0) N/A (699.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
4:2FTS	(327.0 / 307.0) N/A (327.0 / 81.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
6:2FTS	(427.0 / 407.0) N/A (427.0 / 81.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
8:2FTS	(527.0 / 507.0) N/A (527.0 / 81.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) N/A (498.0 / 478.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NMeFOSA	(512.0 / 219.0) N/A (512.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NEtFOSA	(526.0 / 219.0) N/A (526.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NMeFOSAA	(570.0 / 419.0) N/A (570.0 / 483.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NEtFOSAA	(584.0 / 419.0) N/A (584.0 / 526.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NMeFOSE	(616.0 / 59.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
NEtFOSE	(630.0 / 59.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
HFPO-DA	(285.0 / 169.0) N/A (285.0 / 185.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
ADONA	(377.0 / 85.0) N/A (377.0 / 251.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
9Cl-Pf3ONS	(531.0 / 351.0) N/A (533.0 / 353.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
11Cl-Pf3OUDS	(631.0 / 451.0) N/A (633.0 / 453.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) N/A (241.0 / 117.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
5:3FTCA	(341.0 / 236.7) N/A (341.0 / 217.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
7:3FTCA	(441.0 / 317.0) N/A (441.0 / 337.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFEESA	(315.0 / 135.0) N/A (315.0 / 83.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFMPA	(229.0 / 85.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
PFMBA	(279.0 / 85.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
NFDHA	(295.0 / 201.0) N/A (295.0 / 85.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
13C3_PFBA_IIS	(216.0 / 172.0) 27631	(3.47, N/A) (N/A, 0.01, N/A)	158.1	N/A	0.8150 [1.0000]	81.5% { 13.2% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 47978	(5.37, N/A) (N/A, 0.00, N/A)	375.6	N/A	0.8067 [1.0000]	80.7% { 12.9% }			
13C4_PFOA_IIS	(417.0 / 372.0) 55325	(6.88, N/A) (N/A, 0.01, N/A)	6880.6	N/A	0.6801 [1.0000]	68.0% { 10.7% }			
13C5_PFNA_IIS	(468.0 / 423.0) 49370	(7.48, N/A) (N/A, 0.00, N/A)	522.7	N/A	0.6465 [1.0000]	64.7% { 9.0% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 61528	(8.03 , N/A) (N/A , 0.00 , N/A)	8616.1	N/A	0.8571 [1.0000]	85.7% { 12.9% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 81147	(7.00 , N/A) (N/A , 0.00 , N/A)	1371.4	N/A	0.6589 [1.0000]	65.9% { 9.2% }			
13C4_PFOS_IIS	(503.0 / 79.9) 154132	(8.22 , N/A) (N/A , -0.01 , N/A)	187.2	N/A	0.8684 [1.0000]	86.8% { 12.0% }			
13C4_PFBA_EIS	(217.0 / 172.0) 247052	(3.47 , N/A) (N/A , 0.01 , N/A)	1142.0	N/A	0.7519 [0.8000]	94.0% { 12.9% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 199867	(4.46 , N/A) (N/A , 0.01 , N/A)	698.0	N/A	0.3739 [0.4000]	93.5% { 11.4% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 127347	(5.37 , N/A) (N/A , 0.00 , N/A)	516.2	N/A	0.1953 [0.2000]	97.7% { 12.0% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 104918	(6.17 , N/A) (N/A , 0.01 , N/A)	555.9	N/A	0.1726 [0.2000]	86.3% { 11.6% }			
13C8_PFOA_EIS	(421.0 / 376.0) 114448	(6.87 , N/A) (N/A , 0.01 , N/A)	522.3	N/A	0.1933 [0.2000]	96.6% { 10.8% }			
13C9_PFNA_EIS	(472.0 / 427.0) 54093	(7.47 , N/A) (N/A , -0.01 , N/A)	7004.5	N/A	0.1099 [0.1000]	109.9% { 10.8% }			
13C6_PFDA_EIS	(519.0 / 474.0) 76339	(8.05 , N/A) (N/A , 0.02 , N/A)	49.6	N/A	0.1072 [0.1000]	107.2% { 12.2% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 57844	(8.54 , N/A) (N/A , 0.01 , N/A)	264.5	N/A	0.0820 [0.1000]	82.0% { 9.8% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (Δ RT-I[min], Δ RT-CV[min], Δ RT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDaA_EIS	(615.0 / 570.0) 59780	(8.86, N/A) (N/A, 0.01, N/A)	659.8	N/A	0.0914 [0.1000]	91.4% { 13.3% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 41291	(9.24, N/A) (N/A, 0.01, N/A)	866.4	N/A	0.0607 [0.1000]	60.7% { 8.0% }			
13C3_PFBs_EIS	(302.0 / 80.0) 401314	(5.33, N/A) (N/A, 0.00, N/A)	191.9	N/A	0.2447 [0.2000]	122.4% { 11.2% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 167562	(7.00, N/A) (N/A, 0.00, N/A)	586.3	N/A	0.1956 [0.2000]	97.8% { 9.6% }			
13C8_PFOS_EIS	(507.0 / 80.0) 408093	(8.23, N/A) (N/A, 0.00, N/A)	359.0	N/A	0.2015 [0.2000]	100.8% { 12.5% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 53646	(5.12, N/A) (N/A, 0.00, N/A)	34.9	N/A	0.5341 [0.4000]	133.5% { 13.3% }			
13C2_6:2FTS_EIS	(429.0 / 81.0) 51465	(6.61, N/A) (N/A, 0.00, N/A)	113.0	N/A	0.4545 [0.4000]	113.6% { 12.1% }			
13C2_8:2FTS_EIS	(529.0 / 81.0) 67379	(7.78, N/A) (N/A, 0.00, N/A)	178.5	N/A	0.5627 [0.4000]	140.7% { 12.9% }			
13C8_PFOsa_EIS	(506.0 / 78.0) 557374	(9.83, N/A) (N/A, 0.01, N/A)	1165.3	N/A	0.1494 [0.2000]	74.7% { 9.6% }			
D3_NMeFOsa_EIS	(515.0 / 169.0) 95966	(10.43, N/A) (N/A, 0.01, N/A)	408.5	N/A	0.1198 [0.2000]	59.9% { 7.2% }			
D5_NEtFOsa_EIS	(531.0 / 169.0) 86096	(10.58, N/A) (N/A, 0.01, N/A)	570.5	N/A	0.1228 [0.2000]	61.4% { 7.1% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 122964	(8.19 , N/A) (N/A , 0.00 , N/A)	643.3	N/A	0.4327 [0.4000]	108.2% { 11.1% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 99712	(8.45 , N/A) (N/A , 0.01 , N/A)	217926.1	N/A	0.4120 [0.4000]	103.0% { 11.8% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 352992	(10.36 , N/A) (N/A , 0.01 , N/A)	378.9	N/A	1.2945 [2.0000]	64.7% { 8.3% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 437798	(10.52 , N/A) (N/A , 0.01 , N/A)	584.2	N/A	1.4036 [2.0000]	70.2% { 8.2% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 315683	(5.67 , N/A) (N/A , 0.01 , N/A)	658.9	N/A	0.7707 [0.8000]	96.3% { 11.4% }			

FORM I

ANALYSIS DATA SHEET

AF-RHMW02-WGN01LF-2302W3

Laboratory:	APPL, LLC	Work Order:	23B0184
Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Matrix:	Water	Laboratory ID:	23B0184-02
		File ID:	S2023-03-02A (12)
Sampled:	02/23/23 10:50	Prepared:	02/28/23 08:33
		Analyzed:	03/02/23 12:03
Solids:		Preparation:	EPA 1633
		Dilution:	1
Initial/Final:	502.93 mL / 2 mL	Instrument:	Saphira
Batch:	BCB0459	Sequence:	SC00861
		Calibration:	2309009

COMPOUND	CONC. (ng/L)	LOQ	LOD	DL	Q
PFBA	0.80 U	1.6	0.80	0.21	
PFPEA	2.5	0.80	0.40	0.065	
PFHXA	0.37 J	0.40	0.20	0.055	IR2,
PFHPA	0.44	0.40	0.20	0.041	IR1
PFOA	0.20 U	0.40	0.20	0.15	
PFNA	0.20 U	0.40	0.20	0.081	
PFDA	0.20 U	0.40	0.20	0.10	
PFUnA	0.20 U	0.40	0.20	0.16	IR2,
PFDOA	0.20 U	0.40	0.20	0.11	
PFTRDA	0.30 U	0.40	0.30	0.20	
PFTEDA	0.20 U	0.40	0.20	0.20	
PFBS	0.20 U	0.40	0.20	0.037	
PFPEs	0.20 U	0.40	0.20	0.062	
PFHXS	0.20 U	0.40	0.20	0.031	
PFHPS	0.20 U	0.40	0.20	0.051	
PFOS	0.20 U	0.40	0.20	0.063	
PFNS	0.20 U	0.40	0.20	0.12	
PFDS	0.20 U	0.40	0.20	0.15	
PFDOS	0.20 U	0.40	0.20	0.12	
4:2FTS	0.80 U	1.6	0.80	0.29	
6:2FTS	4.3	1.6	0.80	0.31	
8:2FTS	0.80 U	1.6	0.80	0.082	
PFOSA	0.20 U	0.40	0.20	0.10	
NMeFOSA	0.80 U	1.6	0.80	0.47	
NEtFOSA	0.80 U	1.6	0.80	0.41	
NMeFOSAA	0.20 U	0.40	0.20	0.11	
NEtFOSAA	0.20 U	0.40	0.20	0.11	
NMeFOSE	1.2 U	1.6	1.2	1.0	
NEtFOSE	1.2 U	1.6	1.2	1.0	
HFPO-DA	0.40 U	0.80	0.40	0.17	

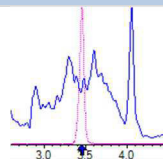
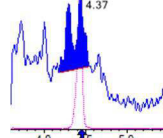
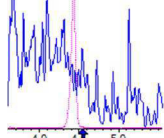
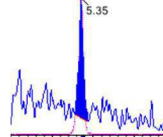
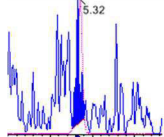
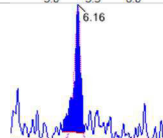
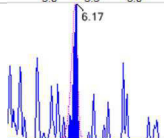
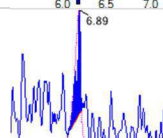
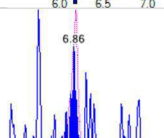
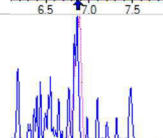
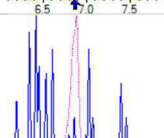
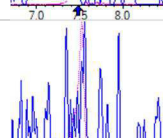
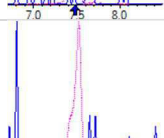
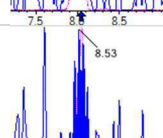
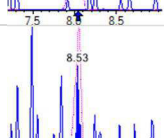
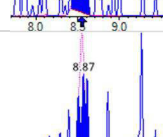
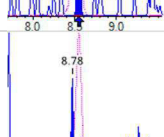
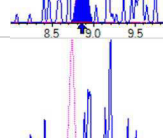
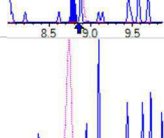
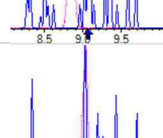
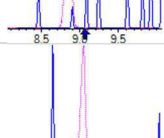
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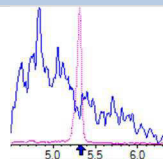
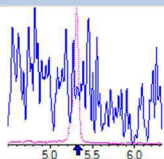
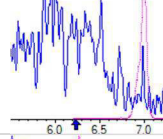
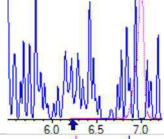
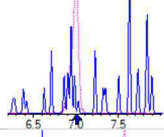
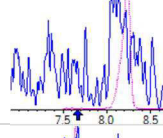
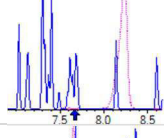
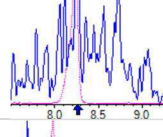
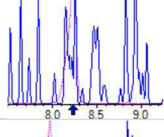
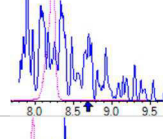
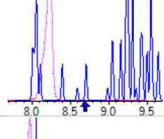
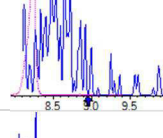
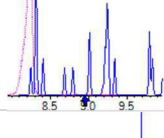
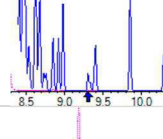
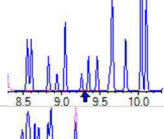
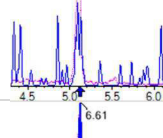
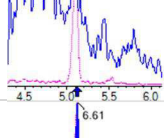
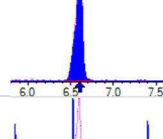
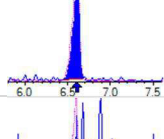
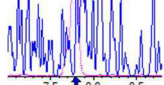
ANALYSIS DATA SHEET

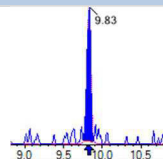
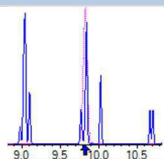
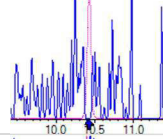
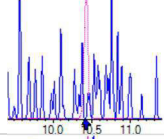
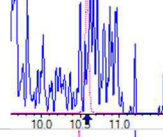
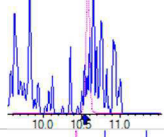
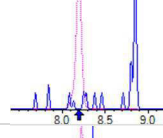
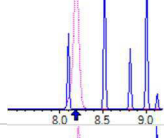
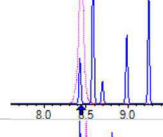
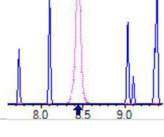
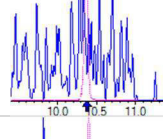
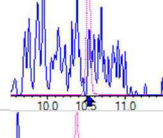
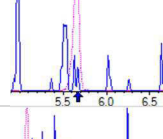
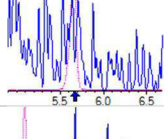
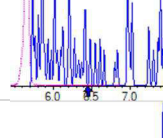
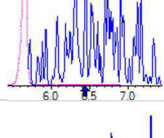
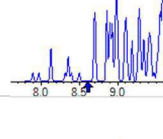
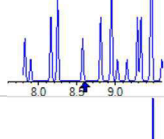
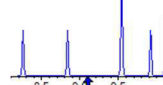
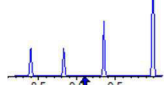
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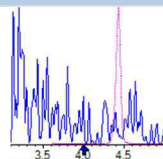
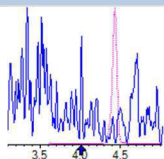
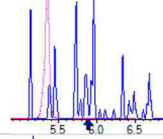
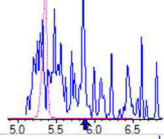
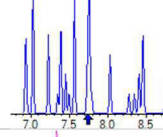
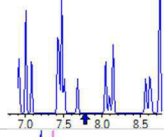
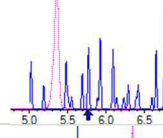
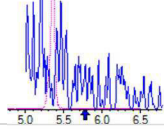
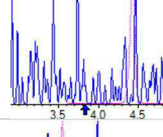
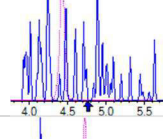
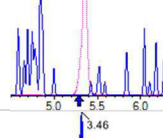
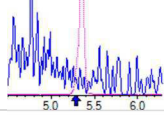
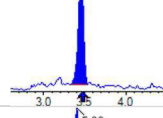
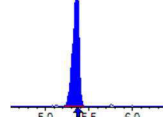
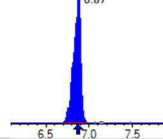
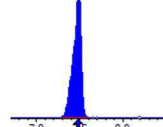
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Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Matrix:	Water	Laboratory ID:	23B0184-02
		File ID:	S2023-03-02A (12)
Sampled:	02/23/23 10:50	Prepared:	02/28/23 08:33
		Analyzed:	03/02/23 12:03
Solids:		Preparation:	EPA 1633
		Dilution:	1
Initial/Final:	502.93 mL / 2 mL	Instrument:	Saphira
Batch:	BCB0459	Sequence:	SC00861
		Calibration:	2309009

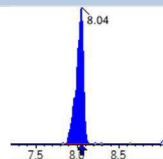
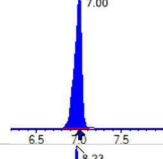
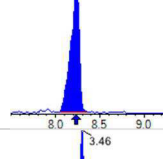
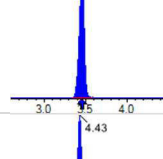
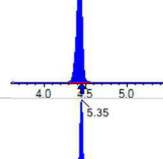
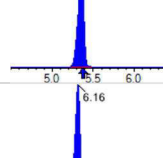
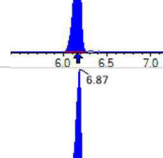
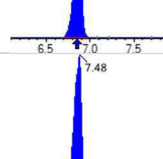
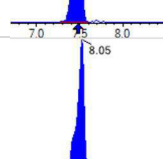
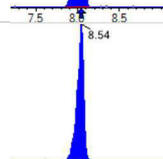
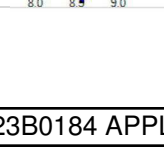
COMPOUND	CONC. (ng/L)	LOQ	LOD	DL	Q
ADONA	0.40 U	0.80	0.40	0.12	
PFEESA	0.40 U	0.80	0.40	0.11	
PFMPA	0.40 U	0.80	0.40	0.054	
PFMBA	0.40 U	0.80	0.40	0.090	
NFDHA	0.40 U	0.80	0.40	0.30	
9CL-PF3ONS	0.40 U	0.80	0.40	0.21	
11CL-PF3OUDS	0.40 U	0.80	0.40	0.20	
3:3FTCA	0.80 U	1.6	0.80	0.57	
5:3FTCA	0.80 U	1.6	0.80	0.44	
7:3FTCA	0.80 U	1.6	0.80	0.55	

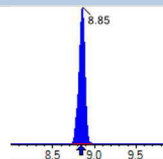
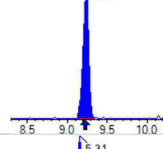
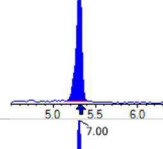
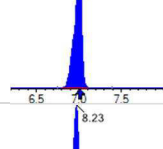
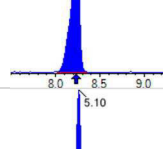
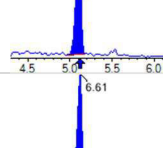
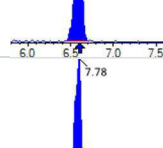
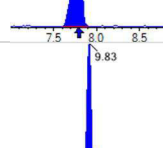
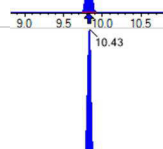
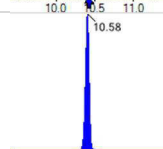
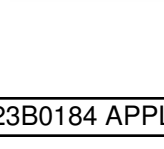
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
PFPeA	(263.0 / 219.0) 143246 (263.0 / 69.0) N/A	(4.37, 0.99) (-0.06, N/A, #Value!)	19.3 N/A	N/A 0.0 0.0	0.6337	N/A			
PFHxA	(313.0 / 269.0) 38046 (313.0 / 119.0) 5487	(5.35, 1.00) (0.00, N/A, 1.7)	30.8 16.1	0.1442 154.1 139.6	0.0925	N/A			IR2,
PFHpA	(363.0 / 319.0) 41645 (363.0 / 169.0) 5960	(6.16, 1.00) (0.00, N/A, -0.6)	36.8 47.7	0.1431 45.3 43.9	0.1104	N/A			IR1,
PFOA	(413.0 / 369.0) 10174 (413.0 / 169.0) 4461	(6.89, 1.00) (0.02, N/A, 1.9)	11.3 109.4	0.4384 129.6 123.4	0.0175	N/A			
PFNA	(463.0 / 419.0) N/A (463.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDA	(513.0 / 469.0) N/A (513.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFUnA	(563.0 / 519.0) 10444 (563.0 / 169.0) 2904	(8.53, 1.00) (-0.01, N/A, -0.2)	13.3 12.7	0.2780 258.3 190.4	0.0147	N/A			IR2,
PFDaA	(613.0 / 569.0) 9672 (613.0 / 169.0) 2167	(8.87, 1.00) (0.02, N/A, 5.9)	12.9 39.1	0.2241 141.0 118.7	0.0159	N/A			
PFTrDA	(663.0 / 619.0) N/A (663.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFTeDA	(713.0 / 669.0) N/A (713.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

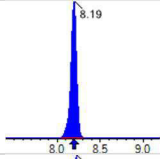
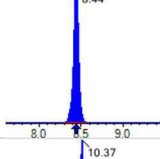
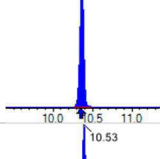
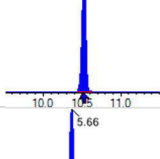
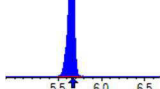
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) N/A (299.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFPeS	(349.0 / 80.0) N/A (349.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHxS	(399.0 / 80.0) N/A (399.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHpS	(449.0 / 80.0) N/A (449.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFOS	(499.0 / 80.0) N/A (499.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFNS	(549.0 / 80.0) N/A (549.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDS	(599.0 / 80.0) N/A (599.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDoS	(699.0 / 80.0) N/A (699.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
4:2FTS	(327.0 / 307.0) N/A (327.0 / 81.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
6:2FTS	(427.0 / 407.0) 283000 (427.0 / 81.0) 216030	(6.61 , 1.00) (0.00 , N/A , -0.2)	695.7 158.9	0.7634 90.1 99.6	1.0928	N/A			
8:2FTS	(527.0 / 507.0) N/A (527.0 / 81.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) 16155 (498.0 / 478.0) N/A	(9.83 , 1.00) (0.00 , N/A , #Value!)	58.6 N/A	N/A 0.0 0.0	0.0081	N/A			
NMeFOSA	(512.0 / 219.0) N/A (512.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NEtFOSA	(526.0 / 219.0) N/A (526.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NMeFOSAA	(570.0 / 419.0) N/A (570.0 / 483.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NEtFOSAA	(584.0 / 419.0) N/A (584.0 / 526.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NMeFOSE	(616.0 / 59.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
NEtFOSE	(630.0 / 59.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
HFPO-DA	(285.0 / 169.0) N/A (285.0 / 185.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
ADONA	(377.0 / 85.0) N/A (377.0 / 251.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
9Cl-Pf3ONS	(531.0 / 351.0) N/A (533.0 / 353.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
11Cl-PF3OUDS	(631.0 / 451.0) N/A (633.0 / 453.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) N/A (241.0 / 117.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
5:3FTCA	(341.0 / 236.7) N/A (341.0 / 217.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
7:3FTCA	(441.0 / 317.0) N/A (441.0 / 337.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFEESA	(315.0 / 135.0) N/A (315.0 / 83.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFMPA	(229.0 / 85.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
PFMBA	(279.0 / 85.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
NFDHA	(295.0 / 201.0) N/A (295.0 / 85.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
13C3_PFBA_IIS	(216.0 / 172.0) 91287	(3.46, N/A) (N/A, 0.00, N/A)	184.7	N/A	0.2693 [1.0000]	26.9% { 43.6% }			IS1,
13C2_PFHxA_IIS	(315.0 / 270.0) 395745	(5.36, N/A) (N/A, -0.01, N/A)	895.8	N/A	0.6654 [1.0000]	66.5% { 106.2% }			
13C4_PFOA_IIS	(417.0 / 372.0) 663067	(6.87, N/A) (N/A, 0.00, N/A)	756.7	N/A	0.8151 [1.0000]	81.5% { 127.9% }			
13C5_PFNA_IIS	(468.0 / 423.0) 686968	(7.48, N/A) (N/A, 0.01, N/A)	865.3	N/A	0.8996 [1.0000]	90.0% { 124.7% }			

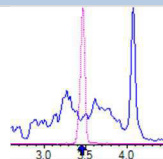
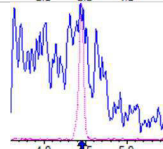
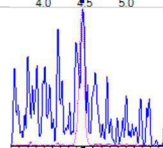
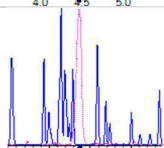
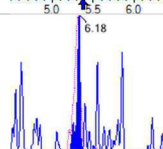
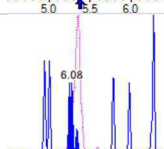
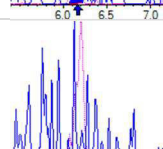
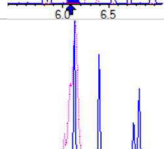
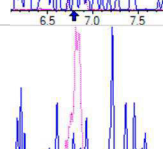
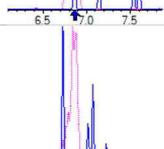
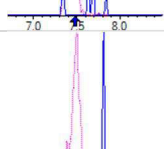
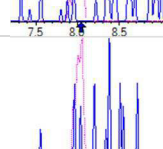
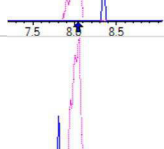
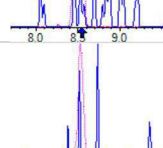
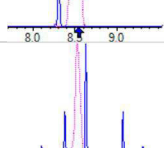
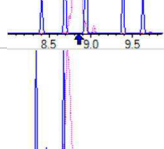
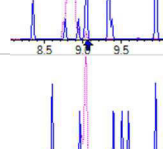
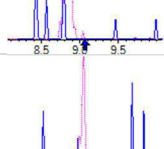
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT Ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 722703	(8.04 , N/A) (N/A , 0.01 , N/A)	451.6	N/A	1.0067 [1.0000]	100.7% { 151.0% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 920957	(7.00 , N/A) (N/A , 0.00 , N/A)	832.3	N/A	0.7478 [1.0000]	74.8% { 104.1% }			
13C4_PFOS_IIS	(503.0 / 79.9) 1558470	(8.23 , N/A) (N/A , 0.00 , N/A)	326.8	N/A	0.8781 [1.0000]	87.8% { 120.9% }			
13C4_PFBA_EIS	(217.0 / 172.0) 696316	(3.46 , N/A) (N/A , 0.00 , N/A)	1830.7	N/A	6.4148 [8.0000]	80.2% { 36.5% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 1026236	(4.43 , N/A) (N/A , -0.02 , N/A)	1088.0	N/A	2.3273 [4.0000]	58.2% { 58.5% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 881826	(5.35 , N/A) (N/A , -0.02 , N/A)	1289.8	N/A	1.6400 [2.0000]	82.0% { 83.1% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 845061	(6.16 , N/A) (N/A , 0.00 , N/A)	1400.4	N/A	1.6851 [2.0000]	84.3% { 93.7% }			
13C8_PFOA_EIS	(421.0 / 376.0) 1209781	(6.87 , N/A) (N/A , 0.01 , N/A)	1143.1	N/A	1.7046 [2.0000]	85.2% { 113.9% }			
13C9_PFNA_EIS	(472.0 / 427.0) 640940	(7.48 , N/A) (N/A , 0.00 , N/A)	592.8	N/A	0.9363 [1.0000]	93.6% { 127.9% }			
13C6_PFDA_EIS	(519.0 / 474.0) 827469	(8.05 , N/A) (N/A , 0.01 , N/A)	773.6	N/A	0.9892 [1.0000]	98.9% { 132.3% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 808085	(8.54 , N/A) (N/A , 0.01 , N/A)	1123.0	N/A	0.9751 [1.0000]	97.5% { 137.2% }			

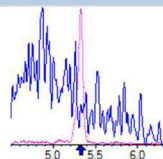
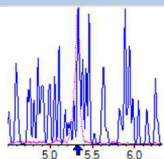
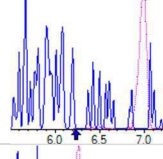
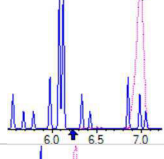
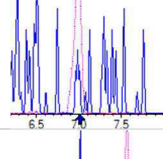
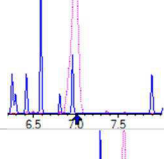
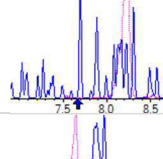
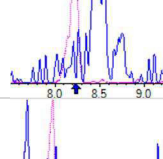
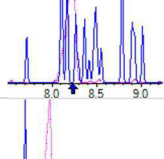
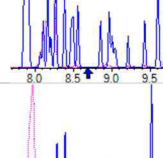
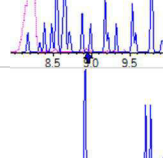
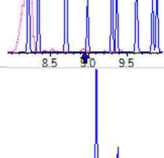
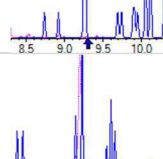
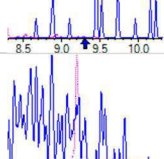
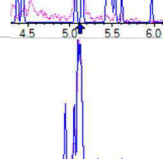
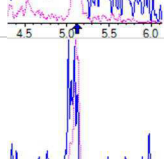
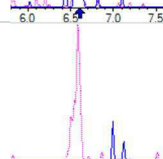
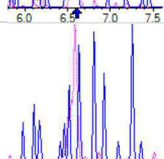
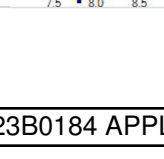
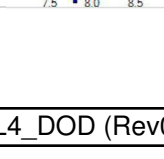
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT Ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDaA_EIS	(615.0 / 570.0) 696680	(8.85 , N/A) (N/A , 0.01 , N/A)	1249.7	N/A	0.9065 [1.0000]	90.6% { 155.6% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 494858	(9.24 , N/A) (N/A , 0.01 , N/A)	647.8	N/A	0.6194 [1.0000]	61.9% { 95.6% }			
13C3_PFBs_EIS	(302.0 / 80.0) 3031192	(5.31 , N/A) (N/A , -0.01 , N/A)	456.9	N/A	1.6288 [2.0000]	81.4% { 84.9% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 1755677	(7.00 , N/A) (N/A , 0.00 , N/A)	965.3	N/A	1.8057 [2.0000]	90.3% { 100.2% }			
13C8_PFOS_EIS	(507.0 / 80.0) 3901011	(8.23 , N/A) (N/A , 0.00 , N/A)	856.0	N/A	1.9053 [2.0000]	95.3% { 119.1% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 761263	(5.10 , N/A) (N/A , -0.02 , N/A)	223.5	N/A	6.6784 [4.0000]	167.0% { 188.9% }			S2,
13C2_6:2FTS_EIS	(429.0 / 81.0) 794033	(6.61 , N/A) (N/A , 0.00 , N/A)	444.7	N/A	6.1784 [4.0000]	154.5% { 186.1% }			S2,
13C2_8:2FTS_EIS	(529.0 / 81.0) 1304483	(7.78 , N/A) (N/A , 0.00 , N/A)	618.6	N/A	9.5985 [4.0000]	240.0% { 249.5% }			S2,
13C8_PFOSA_EIS	(506.0 / 78.0) 5158739	(9.83 , N/A) (N/A , 0.01 , N/A)	2310.4	N/A	1.3678 [2.0000]	68.4% { 89.1% }			
D3_NMeFOSA_EIS	(515.0 / 169.0) 768897	(10.43 , N/A) (N/A , 0.01 , N/A)	1598.0	N/A	0.9494 [2.0000]	47.5% { 57.9% }			
D5_NEtFOSA_EIS	(531.0 / 169.0) 620654	(10.58 , N/A) (N/A , 0.01 , N/A)	2045.3	N/A	0.8753 [2.0000]	43.8% { 51.2% }			

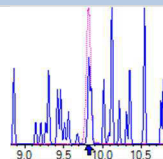
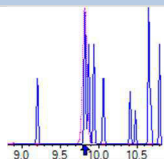
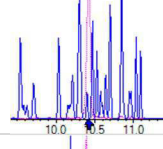
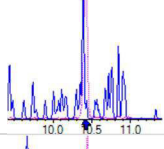
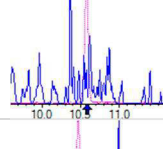
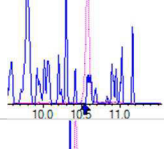
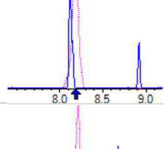
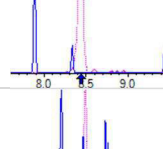
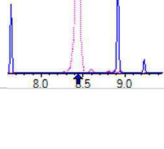
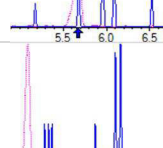
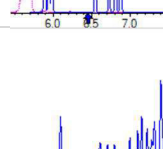
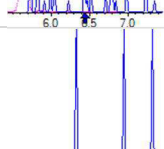
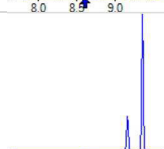
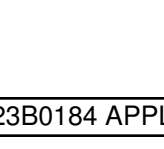
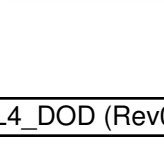
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1773724	(8.19 , N/A) (N/A , 0.00 , N/A)	1900.6	N/A	6.1727 [4.0000]	154.3% { 159.8% }			S2,
D5_EtFOSAA_EIS	(589.0 / 419.0) 1540850	(8.44 , N/A) (N/A , 0.00 , N/A)	75687.5	N/A	6.2971 [4.0000]	157.4% { 182.9% }			S2,
D7_NMeFOSE_EIS	(623.0 / 58.9) 2723371	(10.37 , N/A) (N/A , 0.01 , N/A)	1562.7	N/A	9.8775 [20.0000]	49.4% { 63.7% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 4104924	(10.53 , N/A) (N/A , 0.01 , N/A)	1215.5	N/A	13.0156 [20.0000]	65.1% { 76.9% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 2370664	(5.66 , N/A) (N/A , -0.01 , N/A)	1602.4	N/A	7.0163 [8.0000]	87.7% { 85.2% }			

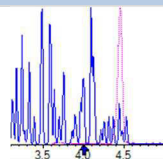
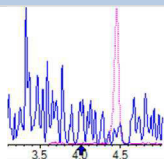
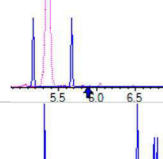
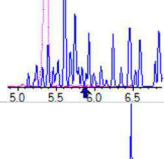
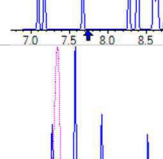
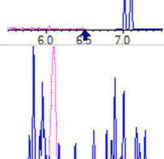
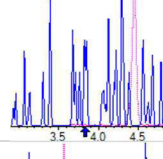

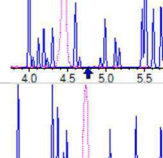
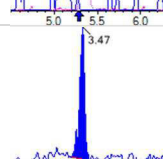
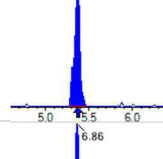

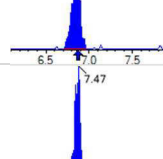
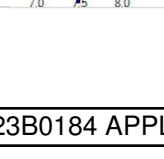
FORM I
ANALYSIS DATA SHEET
AF-RHMW02-WGN01LF-2302W3

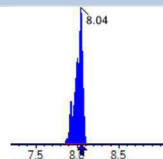
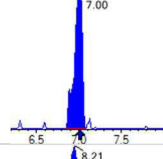
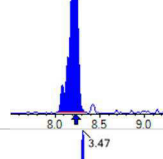
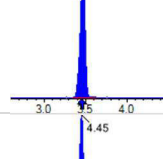
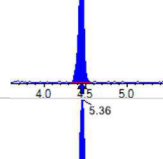
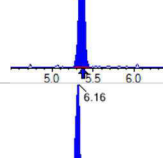
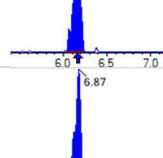
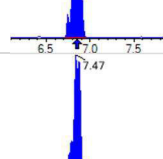
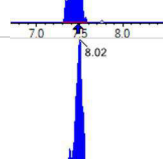
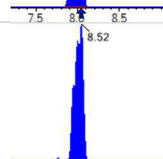
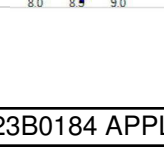
Laboratory:	APPL, LLC	Work Order:	23B0184		
Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling		
Matrix:	Water	Laboratory ID:	23B0184-02RE1	File ID:	S2023-03-02A (13)
Sampled:	02/23/23 10:50	Prepared:	02/28/23 08:33	Analyzed:	03/02/23 12:16
Solids:		Preparation:	EPA 1633	Dilution:	10
Initial/Final:	502.93 mL / 2 mL			Instrument:	Saphira
Batch:	BCB0459	Sequence:	SC00861	Calibration:	2309009

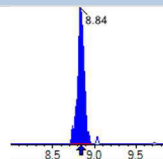
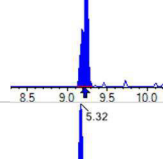
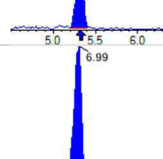
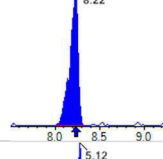
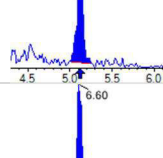
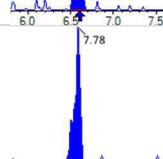
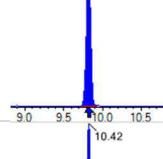
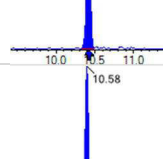
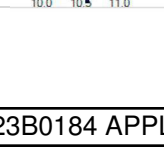
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
PFPeA	(263.0 / 219.0) N/A (263.0 / 69.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHxA	(313.0 / 269.0) N/A (313.0 / 119.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHpA	(363.0 / 319.0) 5572 (363.0 / 169.0) 1084	(6.18, 1.00) (0.02, N/A, 5.6)	41.9 310.6	0.1946 61.6 59.7	0.0110	N/A			
PFOA	(413.0 / 369.0) N/A (413.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFNA	(463.0 / 419.0) N/A (463.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDA	(513.0 / 469.0) N/A (513.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFUnA	(563.0 / 519.0) N/A (563.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDaA	(613.0 / 569.0) N/A (613.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFTrDA	(663.0 / 619.0) N/A (663.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFTeDA	(713.0 / 669.0) N/A (713.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

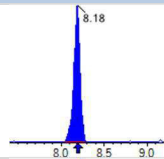
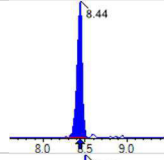
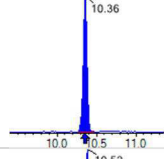
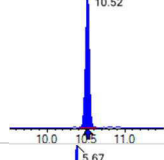
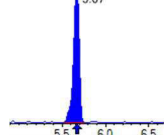
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) N/A (299.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFPeS	(349.0 / 80.0) N/A (349.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHxS	(399.0 / 80.0) N/A (399.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHpS	(449.0 / 80.0) N/A (449.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFOS	(499.0 / 80.0) N/A (499.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFNS	(549.0 / 80.0) N/A (549.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDS	(599.0 / 80.0) N/A (599.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDoS	(699.0 / 80.0) N/A (699.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
4:2FTS	(327.0 / 307.0) N/A (327.0 / 81.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
6:2FTS	(427.0 / 407.0) N/A (427.0 / 81.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
8:2FTS	(527.0 / 507.0) N/A (527.0 / 81.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) N/A (498.0 / 478.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NMeFOSA	(512.0 / 219.0) N/A (512.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NEtFOSA	(526.0 / 219.0) N/A (526.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NMeFOSAA	(570.0 / 419.0) N/A (570.0 / 483.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NEtFOSAA	(584.0 / 419.0) N/A (584.0 / 526.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NMeFOSE	(616.0 / 59.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
NEtFOSE	(630.0 / 59.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
HFPO-DA	(285.0 / 169.0) N/A (285.0 / 185.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
ADONA	(377.0 / 85.0) N/A (377.0 / 251.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
9Cl-Pf3ONS	(531.0 / 351.0) N/A (533.0 / 353.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
11Cl-Pf3OUDS	(631.0 / 451.0) N/A (633.0 / 453.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) N/A (241.0 / 117.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
5:3FTCA	(341.0 / 236.7) N/A (341.0 / 217.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
7:3FTCA	(441.0 / 317.0) N/A (441.0 / 337.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFEESA	(315.0 / 135.0) N/A (315.0 / 83.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFMPA	(229.0 / 85.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
PFMBA	(279.0 / 85.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
NFDHA	(295.0 / 201.0) N/A (295.0 / 85.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
13C3_PFBA_IIS	(216.0 / 172.0) 26332	(3.47, N/A) (N/A, 0.01, N/A)	87.4	N/A	0.7767 [1.0000]	77.7% { 12.6% }			
13C2_PFHx_A_IIS	(315.0 / 270.0) 54410	(5.37, N/A) (N/A, 0.00, N/A)	816.2	N/A	0.9148 [1.0000]	91.5% { 14.6% }			
13C4_PFOA_IIS	(417.0 / 372.0) 67278	(6.86, N/A) (N/A, -0.01, N/A)	432.1	N/A	0.8271 [1.0000]	82.7% { 13.0% }			
13C5_PFNA_IIS	(468.0 / 423.0) 62958	(7.47, N/A) (N/A, 0.00, N/A)	14911991.2	N/A	0.8245 [1.0000]	82.4% { 11.4% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 62809	(8.04 , N/A) (N/A , 0.00 , N/A)	72.3	N/A	0.8749 [1.0000]	87.5% { 13.1% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 107058	(7.00 , N/A) (N/A , 0.00 , N/A)	241.9	N/A	0.8693 [1.0000]	86.9% { 12.1% }			
13C4_PFOS_IIS	(503.0 / 79.9) 179391	(8.21 , N/A) (N/A , -0.01 , N/A)	177.2	N/A	1.0107 [1.0000]	101.1% { 13.9% }			
13C4_PFBA_EIS	(217.0 / 172.0) 206582	(3.47 , N/A) (N/A , 0.01 , N/A)	932.8	N/A	0.6598 [0.8000]	82.5% { 10.8% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 224235	(4.45 , N/A) (N/A , 0.00 , N/A)	565.0	N/A	0.3699 [0.4000]	92.5% { 12.8% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 139022	(5.36 , N/A) (N/A , -0.01 , N/A)	426.9	N/A	0.1880 [0.2000]	94.0% { 13.1% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 113533	(6.16 , N/A) (N/A , 0.00 , N/A)	1149.2	N/A	0.1647 [0.2000]	82.3% { 12.6% }			
13C8_PFOA_EIS	(421.0 / 376.0) 125487	(6.87 , N/A) (N/A , 0.00 , N/A)	2079.0	N/A	0.1743 [0.2000]	87.1% { 11.8% }			
13C9_PFNA_EIS	(472.0 / 427.0) 62596	(7.47 , N/A) (N/A , -0.01 , N/A)	8302.0	N/A	0.0998 [0.1000]	99.8% { 12.5% }			
13C6_PFDA_EIS	(519.0 / 474.0) 62015	(8.02 , N/A) (N/A , -0.01 , N/A)	67.1	N/A	0.0853 [0.1000]	85.3% { 9.9% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 87733	(8.52 , N/A) (N/A , -0.01 , N/A)	89416.7	N/A	0.1218 [0.1000]	121.8% { 14.9% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 65476	(8.84 , N/A) (N/A , -0.01 , N/A)	653.4	N/A	0.0980 [0.1000]	98.0% { 14.6% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 56559	(9.24 , N/A) (N/A , 0.01 , N/A)	326.4	N/A	0.0815 [0.1000]	81.5% { 10.9% }			
13C3_PFBs_EIS	(302.0 / 80.0) 424652	(5.32 , N/A) (N/A , 0.00 , N/A)	269.0	N/A	0.1963 [0.2000]	98.1% { 11.9% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 172077	(6.99 , N/A) (N/A , -0.01 , N/A)	601.5	N/A	0.1522 [0.2000]	76.1% { 9.8% }			
13C8_PFOS_EIS	(507.0 / 80.0) 403371	(8.22 , N/A) (N/A , 0.00 , N/A)	305.5	N/A	0.1712 [0.2000]	85.6% { 12.3% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 60601	(5.12 , N/A) (N/A , 0.00 , N/A)	61.7	N/A	0.4573 [0.4000]	114.3% { 15.0% }			
13C2_6:2FTS_EIS	(429.0 / 81.0) 54911	(6.60 , N/A) (N/A , 0.00 , N/A)	157.8	N/A	0.3675 [0.4000]	91.9% { 12.9% }			
13C2_8:2FTS_EIS	(529.0 / 81.0) 58650	(7.78 , N/A) (N/A , 0.00 , N/A)	293.0	N/A	0.3712 [0.4000]	92.8% { 11.2% }			
13C8_PFOsa_EIS	(506.0 / 78.0) 554028	(9.82 , N/A) (N/A , 0.00 , N/A)	1158.9	N/A	0.1276 [0.2000]	63.8% { 9.6% }			
D3_NMeFOsa_EIS	(515.0 / 169.0) 82032	(10.42 , N/A) (N/A , 0.00 , N/A)	397.3	N/A	0.0880 [0.2000]	44.0% { 6.2% }			
D5_NEtFOsa_EIS	(531.0 / 169.0) 71753	(10.58 , N/A) (N/A , 0.00 , N/A)	571.3	N/A	0.0879 [0.2000]	44.0% { 5.9% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 128673	(8.18 , N/A) (N/A , -0.01 , N/A)	701.3	N/A	0.3890 [0.4000]	97.3% { 11.6% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 127116	(8.44 , N/A) (N/A , 0.00 , N/A)	1489420.2	N/A	0.4513 [0.4000]	112.8% { 15.1% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 394512	(10.36 , N/A) (N/A , 0.00 , N/A)	647.8	N/A	1.2431 [2.0000]	62.2% { 9.2% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 475141	(10.52 , N/A) (N/A , 0.00 , N/A)	755.7	N/A	1.3088 [2.0000]	65.4% { 8.9% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 349384	(5.67 , N/A) (N/A , 0.00 , N/A)	622.3	N/A	0.7521 [0.8000]	94.0% { 12.6% }			

QUALITY CONTROL

SURROGATE SUMMARY SHEET

EPA 1633

Client: AECOM
 Work Order: 23B0184
 Project: Red Hill AFFF Assessment Sampling

Surrogate Compound	Spike level	% Recovery	Recovery Limits	Q
AF-RHMW03-WGN01LF-2302W3 (23B0184-01) ng/L				
		Lab File ID: S2023-03-02A (10)		Analyzed: 03/02/23 11:38
13C4-PFBA	30.3	93.0	10 - 130	
13C5-PFPEA	15.2	69.6	35 - 150	
13C5-PFHXA	7.58	90.2	55 - 150	
13C4-PFHFA	7.58	86.4	55 - 150	
13C8-PFOA	7.58	96.1	60 - 140	
13C9-PFNA	3.79	108	55 - 140	
13C6-PFDA	3.79	104	50 - 140	
13C7-PFUnA	3.79	100	30 - 140	
13C2-PFDOA	3.79	77.8	10 - 150	
13C2-PFTEDA	3.79	63.3	10 - 130	
13C3-PFBS	7.58	92.8	55 - 150	
13C3-PFHXS	7.58	92.3	55 - 150	
13C8-PFOS	7.58	92.3	45 - 140	
13C2-4:2FTS	15.2	186	60 - 200	
13C2-6:2FTS	15.2	152	60 - 200	
13C2-8:2FTS	15.2	211	50 - 200	*
13C8-PFOSA	7.58	70.6	30 - 130	
D3-NMEFOSA	7.58	53.0	15 - 130	
D5-NETFOSA	7.58	52.6	10 - 130	
D3-NMEFOSAA	15.2	164	45 - 200	
D5-NETFOSAA	15.2	147	10 - 200	
D7-NMEFOSE	75.8	48.8	10 - 150	
D9-NETFOSE	75.8	55.8	10 - 150	
13C3-HFPO-DA	30.3	96.5	25 - 160	
AF-RHMW03-WGN01LF-2302W3 (23B0184-01RE1) ng/L				
		Lab File ID: S2023-03-02A (11)		Analyzed: 03/02/23 11:51
13C2-8:2FTS	15.2	141	50 - 200	

SURROGATE SUMMARY SHEET

EPA 1633

Client: AECOM
 Work Order: 23B0184
 Project: Red Hill AFFF Assessment Sampling

Surrogate Compound	Spike level	% Recovery	Recovery Limits	Q
AF-RHMW02-WGN01LF-2302W3 (23B0184-02) ng/L				
		Lab File ID: S2023-03-02A (12)		Analyzed: 03/02/23 12:03
13C4-PFBA	31.8	80.2	10 - 130	
13C5-PFPEA	15.9	58.2	35 - 150	
13C5-PFHXA	7.95	82.0	55 - 150	
13C4-PFHFA	7.95	84.3	55 - 150	
13C8-PFOA	7.95	85.2	60 - 140	
13C9-PFNA	3.98	93.6	55 - 140	
13C6-PFDA	3.98	98.9	50 - 140	
13C7-PFUnA	3.98	97.5	30 - 140	
13C2-PFDOA	3.98	90.6	10 - 150	
13C2-PFTEDA	3.98	61.9	10 - 130	
13C3-PFBS	7.95	81.4	55 - 150	
13C3-PFHXS	7.95	90.3	55 - 150	
13C8-PFOS	7.95	95.3	45 - 140	
13C2-4:2FTS	15.9	167	60 - 200	
13C2-6:2FTS	15.9	154	60 - 200	
13C2-8:2FTS	15.9	240	50 - 200	*
13C8-PFOSA	7.95	68.4	30 - 130	
D3-NMEFOSA	7.95	47.5	15 - 130	
D5-NETFOSA	7.95	43.8	10 - 130	
D3-NMEFOSAA	15.9	154	45 - 200	
D5-NETFOSAA	15.9	157	10 - 200	
D7-NMEFOSE	79.5	49.4	10 - 150	
D9-NETFOSSE	79.5	65.1	10 - 150	
13C3-HFPO-DA	31.8	87.7	25 - 160	
AF-RHMW02-WGN01LF-2302W3 (23B0184-02RE1) ng/L				
		Lab File ID: S2023-03-02A (13)		Analyzed: 03/02/23 12:16
13C2-8:2FTS	15.9	92.8	50 - 200	

SURROGATE SUMMARY SHEET

EPA 1633

Client: AECOM
 Work Order: 23B0184
 Project: Red Hill AFFF Assessment Sampling

Surrogate Compound	Spike level	% Recovery	Recovery Limits	Q
Blank (BCB0459-BLK1) ng/L		Lab File ID: S2023-03-02A (7)		Analyzed: 03/02/23 10:59
13C4-PFBA	32.0	94.6	10 - 130	
13C5-PFPEA	16.0	88.1	35 - 150	
13C5-PFHXA	8.00	95.1	55 - 150	
13C4-PFHFA	8.00	94.6	55 - 150	
13C8-PFOA	8.00	99.1	60 - 140	
13C9-PFNA	4.00	103	55 - 140	
13C6-PFDA	4.00	96.9	50 - 140	
13C7-PFUnA	4.00	96.3	30 - 140	
13C2-PFDOA	4.00	99.5	10 - 150	
13C2-PFTEDA	4.00	92.5	10 - 130	
13C3-PFBS	8.00	101	55 - 150	
13C3-PFHXS	8.00	90.1	55 - 150	
13C8-PFOS	8.00	94.4	45 - 140	
13C2-4:2FTS	16.0	107	60 - 200	
13C2-6:2FTS	16.0	92.2	60 - 200	
13C2-8:2FTS	16.0	107	50 - 200	
13C8-PFOSA	8.00	93.1	30 - 130	
D3-NMEFOSA	8.00	50.0	15 - 130	
D5-NETFOSA	8.00	47.2	10 - 130	
D3-NMEFOSAA	16.0	108	45 - 200	
D5-NETFOSAA	16.0	105	10 - 200	
D7-NMEFOSE	80.0	57.4	10 - 150	
D9-NETFOSSE	80.0	76.7	10 - 150	
13C3-HFPO-DA	32.0	91.6	25 - 160	

SURROGATE SUMMARY SHEET

EPA 1633

Client: AECOM
 Work Order: 23B0184
 Project: Red Hill AFFF Assessment Sampling

Surrogate Compound	Spike level	% Recovery	Recovery Limits	Q
LCS (BCB0459-BS1) . ng/L		Lab File ID: S2023-03-02A (8)		Analyzed: 03/02/23 11:12
13C4-PFBA	32.0	95.6	10 - 130	
13C5-PFPEA	16.0	86.2	35 - 150	
13C5-PFHXA	8.00	97.3	55 - 150	
13C4-PFHFA	8.00	89.5	55 - 150	
13C8-PFOA	8.00	91.4	60 - 140	
13C9-PFNA	4.00	99.0	55 - 140	
13C6-PFDA	4.00	102	50 - 140	
13C7-PFUnA	4.00	101	30 - 140	
13C2-PFDOA	4.00	103	10 - 150	
13C2-PFTEDA	4.00	99.4	10 - 130	
13C3-PFBS	8.00	110	55 - 150	
13C3-PFHXS	8.00	97.9	55 - 150	
13C8-PFOS	8.00	95.6	45 - 140	
13C2-4:2FTS	16.0	121	60 - 200	
13C2-6:2FTS	16.0	102	60 - 200	
13C2-8:2FTS	16.0	127	50 - 200	
13C8-PFOSA	8.00	84.8	30 - 130	
D3-NMEFOSA	8.00	54.2	15 - 130	
D5-NETFOSA	8.00	52.6	10 - 130	
D3-NMEFOSAA	16.0	111	45 - 200	
D5-NETFOSAA	16.0	99.2	10 - 200	
D7-NMEFOSE	80.0	57.3	10 - 150	
D9-NETFOSSE	80.0	78.6	10 - 150	
13C3-HFPO-DA	32.0	82.9	25 - 160	

SURROGATE SUMMARY SHEET

EPA 1633

Client: AECOM
 Work Order: 23B0184
 Project: Red Hill AFFF Assessment Sampling

Surrogate Compound	Spike level	% Recovery	Recovery Limits	Q
MRL Check (BCB0459-MRL1) . ng/L		Lab File ID: S2023-03-02A (9)		Analyzed: 03/02/23 11:25
13C4-PFBA	32.0	94.6	10 - 130	
13C5-PFPEA	16.0	92.3	35 - 150	
13C5-PFHXA	8.00	98.5	55 - 150	
13C4-PFHFA	8.00	94.7	55 - 150	
13C8-PFOA	8.00	98.7	60 - 140	
13C9-PFNA	4.00	91.9	55 - 140	
13C6-PFDA	4.00	102	50 - 140	
13C7-PFUnA	4.00	106	30 - 140	
13C2-PFDOA	4.00	99.6	10 - 150	
13C2-PFTEDA	4.00	91.0	10 - 130	
13C3-PFBS	8.00	97.0	55 - 150	
13C3-PFHXS	8.00	99.3	55 - 150	
13C8-PFOS	8.00	97.9	45 - 140	
13C2-4:2FTS	16.0	97.1	60 - 200	
13C2-6:2FTS	16.0	93.9	60 - 200	
13C2-8:2FTS	16.0	116	50 - 200	
13C8-PFOSA	8.00	89.0	30 - 130	
D3-NMEFOSA	8.00	54.4	15 - 130	
D5-NETFOSA	8.00	57.2	10 - 130	
D3-NMEFOSAA	16.0	113	45 - 200	
D5-NETFOSAA	16.0	106	10 - 200	
D7-NMEFOSE	80.0	58.6	10 - 150	
D9-NETFOSSE	80.0	84.1	10 - 150	
13C3-HFPO-DA	32.0	97.0	25 - 160	

METHOD BLANK SUMMARY

EPA 1633

Laboratory: APPL, LLC Work Order: 23B0184
 Client: AECOM Project: Red Hill AFFF Assessment Sampling
 Blank ID: BCB0459-BLK1 Batch: BCB0459 Prepared: 02/28/2023 08:24

Client Sample ID	Laboratory Sample ID	Lab File ID	Time Analyzed
LCS	BCB0459-BS1	S2023-03-02A (8)	11:12
MRL Check	BCB0459-MRL1	S2023-03-02A (9)	11:25
AF-RHMW03-WGN01LF-2302W3	23B0184-01	S2023-03-02A (10)	11:38
Df 10	23B0184-01RE1	S2023-03-02A (11)	11:51
AF-RHMW02-WGN01LF-2302W3	23B0184-02	S2023-03-02A (12)	12:03
Df 10	23B0184-02RE1	S2023-03-02A (13)	12:16

ANALYSIS DATA SHEET

Blank

Laboratory:	APPL, LLC	Work Order:	23B0184
Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Matrix:	Water	Laboratory ID:	BCB0459-BLK1
		File ID:	S2023-03-02A (7)
Sampled:		Prepared:	02/28/23 08:24
		Analyzed:	03/02/23 10:59
Solids:		Preparation:	EPA 1633
		Dilution:	1
Batch:	BCB0459	Sequence:	SC00861
		Calibration:	2309009
Column:	1	Instrument:	Saphira

COMPOUND	CONC. (ng/L)	LOQ	LOD	DL	Q
PFBA	0.80 U	1.6	0.80	0.21	U
PFPEA	0.40 U	0.80	0.40	0.065	U
PFHXA	0.20 U	0.40	0.20	0.055	U
PFHPA	0.20 U	0.40	0.20	0.041	U
PFOA	0.20 U	0.40	0.20	0.15	U
PFNA	0.20 U	0.40	0.20	0.082	U
PFDA	0.20 U	0.40	0.20	0.10	U
PFUnA	0.20 U	0.40	0.20	0.16	U
PFDOA	0.20 U	0.40	0.20	0.11	U
PFTRDA	0.30 U	0.40	0.30	0.20	U
PFTEDA	0.20 U	0.40	0.20	0.20	U
PFBS	0.20 U	0.40	0.20	0.037	U
PFPEs	0.20 U	0.40	0.20	0.063	U
PFHXS	0.20 U	0.40	0.20	0.032	U
PFHPS	0.20 U	0.40	0.20	0.051	U
PFOS	0.102 J	0.40	0.20	0.064	MI2, J
PFNS	0.20 U	0.40	0.20	0.12	U
PFDS	0.20 U	0.40	0.20	0.15	U
PFDOS	0.20 U	0.40	0.20	0.12	U
4:2FTS	0.80 U	1.6	0.80	0.29	U
6:2FTS	0.80 U	1.6	0.80	0.31	U
8:2FTS	0.80 U	1.6	0.80	0.082	U
PFOSA	0.20 U	0.40	0.20	0.10	U
NMeFOSA	0.80 U	1.6	0.80	0.47	U
NEtFOSA	0.80 U	1.6	0.80	0.41	U
NMeFOSAA	0.20 U	0.40	0.20	0.11	U
NEtFOSAA	0.20 U	0.40	0.20	0.11	U
NMeFOSE	1.2 U	1.6	1.2	1.0	U
NEtFOSE	1.2 U	1.6	1.2	1.0	U
HFPO-DA	0.40 U	0.80	0.40	0.17	U

ANALYSIS DATA SHEET

Blank

Laboratory:	APPL, LLC	Work Order:	23B0184
Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Matrix:	Water	Laboratory ID:	BCB0459-BLK1
		File ID:	S2023-03-02A (7)
Sampled:		Prepared:	02/28/23 08:24
		Analyzed:	03/02/23 10:59
Solids:		Preparation:	EPA 1633
		Dilution:	1
Batch:	BCB0459	Sequence:	SC00861
		Calibration:	2309009
Column:	1	Instrument:	Saphira

COMPOUND	CONC. (ng/L)	LOQ	LOD	DL	Q
ADONA	0.40 U	0.80	0.40	0.12	U
PFEESA	0.40 U	0.80	0.40	0.11	U
PFMPA	0.40 U	0.80	0.40	0.054	U
PFMBA	0.40 U	0.80	0.40	0.091	U
NFDHA	0.40 U	0.80	0.40	0.30	U
9CL-PF3ONS	0.40 U	0.80	0.40	0.21	U
11CL-PF3OUDS	0.40 U	0.80	0.40	0.21	U
3:3FTCA	0.80 U	1.6	0.80	0.57	U
5:3FTCA	0.80 U	1.6	0.80	0.44	U
7:3FTCA	0.80 U	1.6	0.80	0.55	U

LCS / LCS DUPLICATE RECOVERY

EPA 1633

Laboratory: APPL, LLC

Work Order: 23B0184

Client: AECOM

Project: Red Hill AFFF Assessment Sampling

Matrix: Water

Preparation: EPA 1633

Batch: BCB0459

Laboratory ID: BCB0459-BS1

Column:

ANALYTE	SPIKE ADDED (ng/L)	LCS CONCENTRATION (ng/L)	LCS % REC.	QC LIMITS REC.
PFBA	16.0	15.6	97.8	58 - 148
PFPEA	8.00	7.64	95.4	54 - 152
PFHXA	4.00	3.61	90.2	55 - 152
PFHPA	4.00	3.47	86.8	54 - 154
PFOA	4.00	3.68	91.9	52 - 161
PFNA	4.00	4.21	105	59 - 149
PFDA	4.00	4.00	99.9	52 - 147
PFUnA	4.00	3.57	89.2	48 - 159
PFDOA	4.00	3.98	99.5	64 - 142
PFTRDA	4.00	3.96	98.9	49 - 148
PFTEDA	4.00	3.41	85.3	47 - 161
PFBS	3.54	3.34	94.2	62 - 144
PFPEs	3.76	3.99	106	59 - 151
PFHXS	3.66	3.66	100	57 - 146
PFHPS	3.82	3.51	91.8	55 - 152
PFOS	3.72	3.67	98.7	58 - 149
PFNS	3.84	3.90	101	52 - 148
PFDS	3.86	3.76	97.4	51 - 147
PFDOS	3.88	4.26	110	36 - 145
4:2FTS	15.0	13.8	92.0	67 - 146
6:2FTS	15.2	16.3	107	61 - 151
8:2FTS	15.4	14.9	97.0	63 - 152
PFOSA	4.00	4.48	112	61 - 148
NMeFOSA	16.0	15.3	95.4	63 - 145
NEtFOSA	16.0	16.9	106	65 - 139
NMeFOSAA	4.00	4.00	100	58 - 144
NEtFOSAA	4.00	4.05	101	59 - 146
NMeFOSE	16.0	16.2	101	71 - 136
NEtFOSE	16.0	15.5	97.2	69 - 137
HFPO-DA	8.00	7.82	97.8	63 - 144
ADONA	7.56	8.03	106	68 - 146
PFEESA	7.12	6.83	95.9	56 - 151
PFMPA	8.00	6.84	85.6	51 - 145
PFMBA	8.00	8.10	101	55 - 148

LCS / LCS DUPLICATE RECOVERY

EPA 1633

Laboratory: APPL, LLC

Work Order: 23B0184

Client: AECOM

Project: Red Hill AFFF Assessment Sampling

Matrix: Water

Preparation: EPA 1633

Batch: BCB0459

Laboratory ID: BCB0459-BS1

Column:

ANALYTE	SPIKE ADDED (ng/L)	LCS CONCENTRATION (ng/L)	LCS % REC.	QC LIMITS REC.
NFDHA	8.00	6.63	82.9	48 - 161
9CL-PF3ONS	7.48	8.85	118	56 - 156
11CL-PF3OUDS	7.56	8.54	113	46 - 156
3:3FTCA	16.0	13.2	82.7	62 - 129
5:3FTCA	16.0	11.3	70.6	63 - 134
7:3FTCA	16.0	12.4	77.4	50 - 138

CALIBRATION SUMMARY

Analyte	(Q1 / Q3)	Internal Standard	Multiplier	AcidFactor	Function	Qualifier
PFBA	(213.0 / 169.0)	13C4_PFBA_EIS	4.0000	1.0000	$y = 0.42284 x$ (std. dev. = 0.03761) (weighting: None)	%RSE=8.9
PFPeA	(263.0 / 219.0)	13C5_PFPeA_EIS	2.0000	1.0000	$y = 0.44057 x$ (std. dev. = 0.04807) (weighting: None)	%RSE=10.9
PFHxA	(313.0 / 269.0)	13C5_PFHxA_EIS	1.0000	1.0000	$y = 0.46662 x$ (std. dev. = 0.02871) (weighting: None)	%RSE=6.2
PFHpA	(363.0 / 319.0)	13C4_PFHpA_EIS	1.0000	1.0000	$y = 0.44650 x$ (std. dev. = 0.02803) (weighting: None)	%RSE=6.3
PFOA	(413.0 / 369.0)	13C8_PFOA_EIS	1.0000	1.0000	$y = 0.48177 x$ (std. dev. = 0.04560) (weighting: None)	%RSE=9.5
PFNA	(463.0 / 419.0)	13C9_PFNA_EIS	1.0000	1.0000	$y = 0.94459 x$ (std. dev. = 0.08740) (weighting: None)	%RSE=9.3
PFDA	(513.0 / 469.0)	13C6_PFDA_EIS	1.0000	1.0000	$y = 0.99147 x$ (std. dev. = 0.10952) (weighting: None)	%RSE=11.0
PFUnA	(563.0 / 519.0)	13C7_PFUnA_EIS	1.0000	1.0000	$y = 0.87937 x$ (std. dev. = 0.09881) (weighting: None)	%RSE=11.2
PFDaA	(613.0 / 569.0)	13C2_PFDaA_EIS	1.0000	1.0000	$y = 0.87244 x$ (std. dev. = 0.08918) (weighting: None)	%RSE=10.2
PFTTrDA	(663.0 / 619.0)	13C2_PFDaA_EIS	1.0000	1.0000	$y = 0.83953 x$ (std. dev. = 0.08426) (weighting: None)	%RSE=10.0
PFTeDA	(713.0 / 669.0)	13C2_PFTeDA_EIS	1.0000	1.0000	$y = 0.91950 x$ (std. dev. = 0.07289) (weighting: None)	%RSE=7.9
PFBS	(299.0 / 80.0)	13C3_PFBS_EIS	1.0000	0.8847	$y = 0.25153 x$ (std. dev. = 0.02274) (weighting: None)	%RSE=9.0
PFPeS	(349.0 / 80.0)	13C3_PFHxS_EIS	1.0000	0.9384	$y = 0.84236 x$ (std. dev. = 0.10508) (weighting: None)	%RSE=12.5
PFHxS	(399.0 / 80.0)	13C3_PFHxS_EIS	1.0000	0.9110	$y = 0.71750 x$ (std. dev. = 0.07265) (weighting: None)	%RSE=10.1
PFHpS	(449.0 / 80.0)	13C8_PFOS_EIS	1.0000	0.9514	$y = 0.42835 x$ (std. dev. = 0.03867) (weighting: None)	%RSE=9.0
PFOS	(499.0 / 80.0)	13C8_PFOS_EIS	1.0000	0.9275	$y = 0.51179 x$ (std. dev. = 0.04829) (weighting: None)	%RSE=9.4
PFNS	(549.0 / 80.0)	13C8_PFOS_EIS	1.0000	0.9599	$y = 0.53002 x$ (std. dev. = 0.07450) (weighting: None)	%RSE=14.1
PFDS	(599.0 / 80.0)	13C8_PFOS_EIS	1.0000	0.9631	$y = 0.59101 x$ (std. dev. = 0.06075) (weighting: None)	%RSE=10.3
PFDoS	(699.0 / 80.0)	13C8_PFOS_EIS	1.0000	0.9696	$y = 0.27466 x$ (std. dev. = 0.03210) (weighting: None)	%RSE=11.7
4:2FTS	(327.0 / 307.0)	13C2_4:2FTS_EIS	4.0000	0.9345	$y = 2.63771 x$ (std. dev. = 0.41713) (weighting: None)	%RSE=15.8
6:2FTS	(427.0 / 407.0)	13C2_6:2FTS_EIS	4.0000	0.9490	$y = 1.23810 x$ (std. dev. = 0.13799) (weighting: None)	%RSE=11.1
8:2FTS	(527.0 / 507.0)	13C2_8:2FTS_EIS	4.0000	0.9583	$y = 1.07578 x$ (std. dev. = 0.07559) (weighting: None)	%RSE=7.0
PFOSA	(498.0 / 78.0)	13C8_PFOSA_EIS	1.0000	1.0000	$y = 0.38855 x$ (std. dev. = 0.04463) (weighting: None)	%RSE=11.5
NMeFOSA	(512.0 / 219.0)	D3_NMeFOSA_EIS	4.0000	1.0000	$y = 1.71767 x$ (std. dev. = 0.15099) (weighting: None)	%RSE=8.8
NEtFOSA	(526.0 / 219.0)	D5_NEtFOSA_EIS	4.0000	1.0000	$y = 1.94294 x$ (std. dev. = 0.24785) (weighting: None)	%RSE=12.8
NMeFOSAA	(570.0 / 419.0)	D3_MeFOSAA_EIS	1.0000	1.0000	$y = 0.21367 x$ (std. dev. = 0.01792) (weighting: None)	%RSE=8.4
NEtFOSAA	(584.0 / 419.0)	D5_EtFOSAA_EIS	1.0000	1.0000	$y = 0.21793 x$ (std. dev. = 0.01891) (weighting: None)	%RSE=8.7
NMeFOSE	(616.0 / 59.0)	D7_NMeFOSE_EIS	4.0000	1.0000	$y = 0.20956 x$ (std. dev. = 0.01758) (weighting: None)	%RSE=8.4
NEtFOSE	(630.0 / 59.0)	D9_NEtFOSE_EIS	4.0000	1.0000	$y = 0.20209 x$ (std. dev. = 0.01915) (weighting: None)	%RSE=9.5
HFPO-DA	(285.0 / 169.0)	13C3_HFPODA_EIS	2.0000	1.0000	$y = 0.20496 x$ (std. dev. = 0.01868) (weighting: None)	%RSE=9.1
ADONA	(377.0 / 85.0)	13C3_HFPODA_EIS	2.0000	0.9427	$y = 0.71708 x$ (std. dev. = 0.11283) (weighting: None)	%RSE=15.7
9Cl-Pf3ONS	(531.0 / 351.0)	13C3_HFPODA_EIS	2.0000	0.9333	$y = 1.66491 x$ (std. dev. = 0.23461) (weighting: None)	%RSE=14.1
11Cl-Pf3OUDS	(631.0 / 451.0)	13C3_HFPODA_EIS	2.0000	0.9432	$y = 0.82728 x$ (std. dev. = 0.09326) (weighting: None)	%RSE=11.3
3:3FTCA	(241.0 / 177.0)	13C5_PFPeA_EIS	4.0000	1.0000	$y = 0.02602 x$ (std. dev. = 0.00252) (weighting: None)	%RSE=9.7
5:3FTCA	(341.0 / 236.7)	13C5_PFHxA_EIS	4.0000	1.0000	$y = 0.26681 x$ (std. dev. = 0.02199) (weighting: None)	%RSE=8.2
7:3FTCA	(441.0 / 317.0)	13C5_PFHxA_EIS	4.0000	1.0000	$y = 0.47968 x$ (std. dev. = 0.06527) (weighting: None)	%RSE=13.6
PFEESA	(315.0 / 135.0)	13C5_PFHxA_EIS	2.0000	0.8925	$y = 1.03498 x$ (std. dev. = 0.13837) (weighting: None)	%RSE=13.4
PFMPA	(229.0 / 85.0)	13C5_PFPeA_EIS	2.0000	1.0000	$y = 0.11264 x$ (std. dev. = 0.01032) (weighting: None)	%RSE=9.2
PFMBA	(279.0 / 85.0)	13C5_PFPeA_EIS	2.0000	1.0000	$y = 0.38591 x$ (std. dev. = 0.04577) (weighting: None)	%RSE=11.9
NFDHA	(295.0 / 201.0)	13C5_PFHxA_EIS	2.0000	1.0000	$y = 0.51906 x$ (std. dev. = 0.04487) (weighting: None)	%RSE=8.6
13C3_PFBA_IIS	(216.0 / 172.0)	13C3_PFBA_IIS	1.0000	1.0000	$y = 339023.8558 x$	%RSD=16.5
13C2_PFHxA_IIS	(315.0 / 270.0)	13C2_PFHxA_IIS	1.0000	1.0000	$y = 594763.6140 x$	%RSD=7.6
13C4_PFOA_IIS	(417.0 / 372.0)	13C4_PFOA_IIS	1.0000	1.0000	$y = 813434.6289 x$	%RSD=7.5
13C5_PFNA_IIS	(468.0 / 423.0)	13C5_PFNA_IIS	1.0000	1.0000	$y = 763620.2538 x$	%RSD=7.6

Analyte	(Q1 / Q3)	Internal Standard	Multiplier	AcidFactor	Function	Qualifier
13C2_PFDA_IIS	(515.0 / 470.1)	13C2_PFDA_IIS	1.0000	1.0000	$y = 717883.6124 x$	%RSD=5.2
18O2_PFHxS_IIS	(403.0 / 83.9)	18O2_PFHxS_IIS	1.0000	1.0000	$y = 1231554.2114 x$	%RSD=5.7
13C4_PFOS_IIS	(503.0 / 79.9)	13C4_PFOS_IIS	1.0000	1.0000	$y = 1774838.7334 x$	%RSD=7.8
13C4_PFBA_EIS	(217.0 / 172.0)	13C3_PFBA_IIS	8.0000	1.0000	$y = 9.5127 x$	%RSD=7.3
13C5_PFPeA_EIS	(268.0 / 223.0)	13C2_PFHxA_IIS	4.0000	1.0000	$y = 4.4570 x$	%RSD=8.2
13C5_PFHxA_EIS	(318.0 / 273.0)	13C2_PFHxA_IIS	2.0000	1.0000	$y = 2.7175 x$	%RSD=5.1
13C4_PFHpA_EIS	(367.0 / 322.0)	13C2_PFHxA_IIS	2.0000	1.0000	$y = 2.5344 x$	%RSD=4.4
13C8_PFOA_EIS	(421.0 / 376.0)	13C4_PFOA_IIS	2.0000	1.0000	$y = 2.1407 x$	%RSD=5.7
13C9_PFNA_EIS	(472.0 / 427.0)	13C5_PFNA_IIS	1.0000	1.0000	$y = 0.9965 x$	%RSD=7.1
13C6_PFDA_EIS	(519.0 / 474.0)	13C2_PFDA_IIS	1.0000	1.0000	$y = 1.1574 x$	%RSD=10.3
13C7_PFUnA_EIS	(570.0 / 525.0)	13C2_PFDA_IIS	1.0000	1.0000	$y = 1.1467 x$	%RSD=11.6
13C2_PFDaA_EIS	(615.0 / 570.0)	13C2_PFDA_IIS	1.0000	1.0000	$y = 1.0635 x$	%RSD=10.6
13C2_PFTeDA_EIS	(715.0 / 670.0)	13C2_PFDA_IIS	1.0000	1.0000	$y = 1.1055 x$	%RSD=12.9
13C3_PFBS_EIS	(302.0 / 80.0)	18O2_PFHxS_IIS	2.0000	1.0000	$y = 4.0416 x$	%RSD=12.5
13C3_PFHxS_EIS	(402.0 / 80.0)	18O2_PFHxS_IIS	2.0000	1.0000	$y = 2.1115 x$	%RSD=6.7
13C8_PFOS_EIS	(507.0 / 80.0)	13C4_PFOS_IIS	2.0000	1.0000	$y = 2.6275 x$	%RSD=7.0
13C2_4:2FTS_EIS	(329.0 / 81.0)	18O2_PFHxS_IIS	4.0000	1.0000	$y = 0.4951 x$	%RSD=10.4
13C2_6:2FTS_EIS	(429.0 / 81.0)	18O2_PFHxS_IIS	4.0000	1.0000	$y = 0.5582 x$	%RSD=7.8
13C2_8:2FTS_EIS	(529.0 / 81.0)	18O2_PFHxS_IIS	4.0000	1.0000	$y = 0.5903 x$	%RSD=8.3
13C8_PFOA_EIS	(506.0 / 78.0)	13C4_PFOS_IIS	2.0000	1.0000	$y = 4.8400 x$	%RSD=11.3
D3_NMeFOSA_EIS	(515.0 / 169.0)	13C4_PFOS_IIS	2.0000	1.0000	$y = 1.0393 x$	%RSD=8.6
D5_NEtFOSA_EIS	(531.0 / 169.0)	13C4_PFOS_IIS	2.0000	1.0000	$y = 0.9100 x$	%RSD=12.3
D3_MeFOSAA_EIS	(573.0 / 419.0)	13C4_PFOS_IIS	4.0000	1.0000	$y = 0.7375 x$	%RSD=7.1
D5_EtFOSAA_EIS	(589.0 / 419.0)	13C4_PFOS_IIS	4.0000	1.0000	$y = 0.6280 x$	%RSD=9.0
D7_NMeFOSE_EIS	(623.0 / 58.9)	13C4_PFOS_IIS	20.0000	1.0000	$y = 3.5383 x$	%RSD=9.7
D9_NEtFOSE_EIS	(639.0 / 58.9)	13C4_PFOS_IIS	20.0000	1.0000	$y = 4.0474 x$	%RSD=13.2
13C3_HFPODA_EIS	(287.0 / 169.0)	13C2_PFHxA_IIS	8.0000	1.0000	$y = 6.8302 x$	%RSD=4.0

x =Concentration Analyte

$$y = \text{Area Ratio} = \frac{\text{Area Analyte}}{\text{Area Internal Standard}}$$

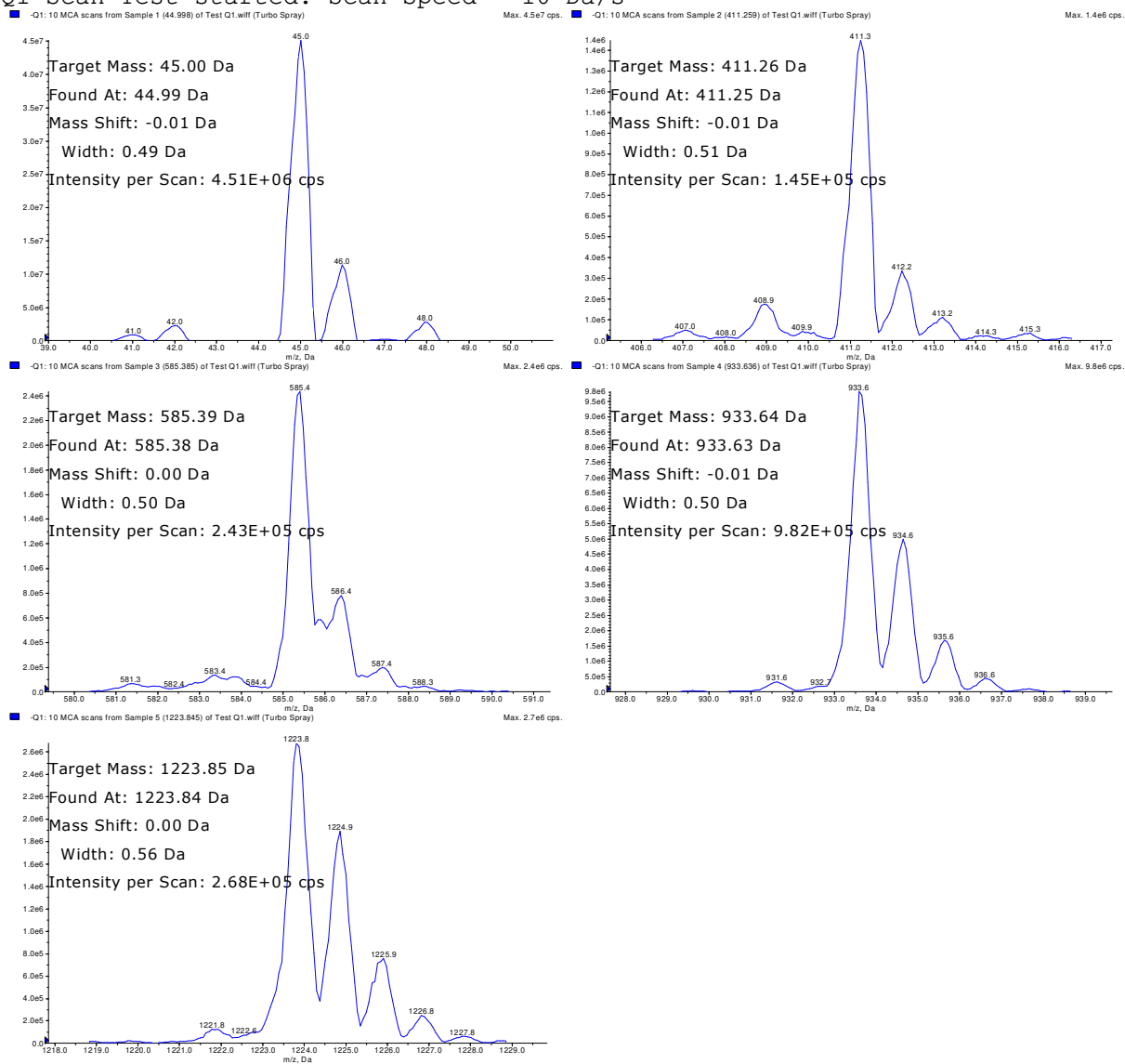
$$\text{Acid Factor} = \frac{\text{Molecular weight Acid}}{\text{Molecular weight Salt}}$$

$$\text{Multiplier} = \frac{\text{Concentration of Analyte}}{\text{Concentration of PFOA}} \text{ in curve standard mix}$$

$$\text{Result} \left(\frac{\text{ng}}{\text{mL}} \right) = x * \text{Multiplier} * \text{Acid Factor}$$

Tune 2021-11-23 Q1 NEG @ 10Da/s

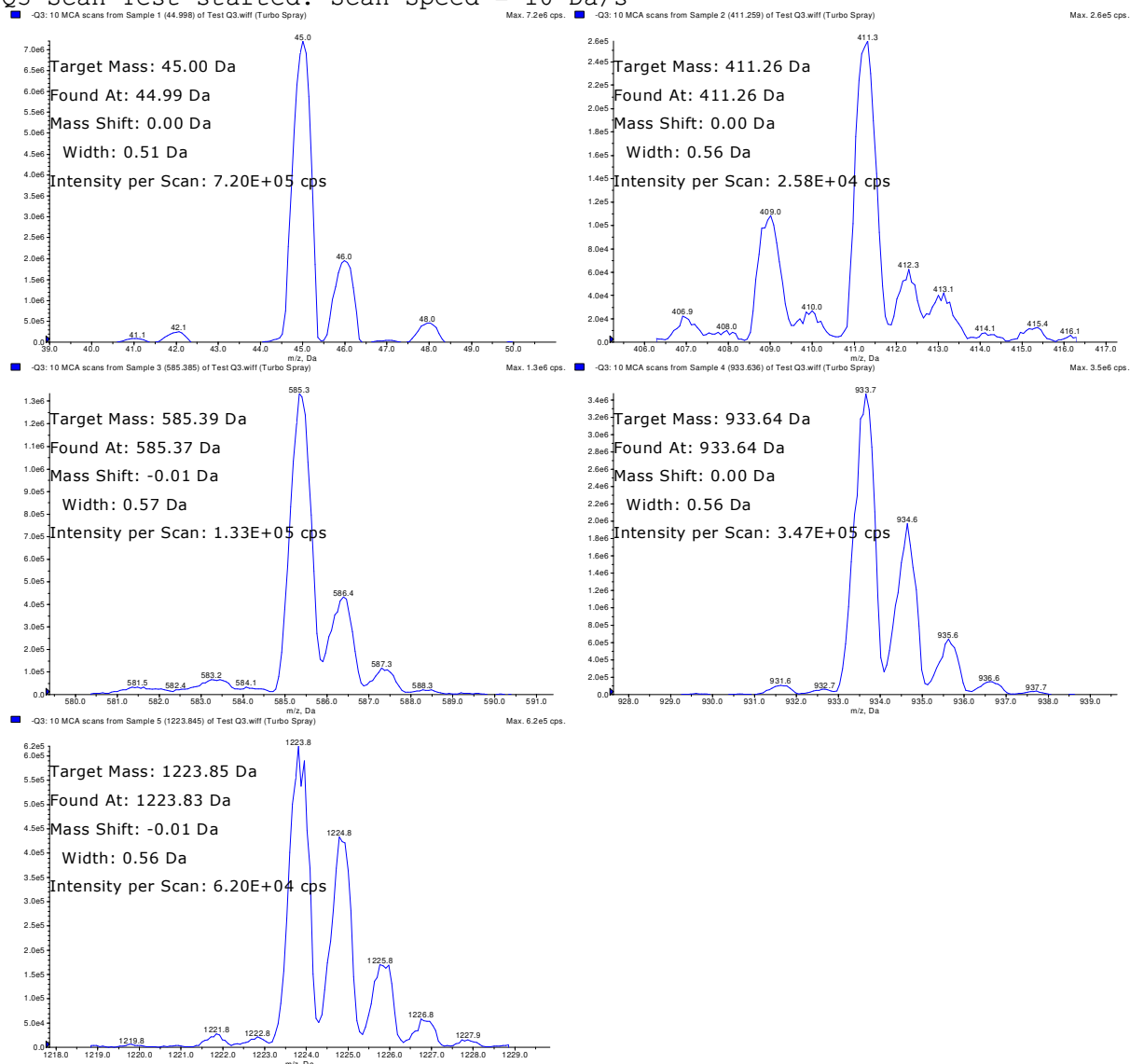
Q1 Scan Test started. Scan Speed = 10 Da/s



Target Mass	Found At	Delta	Width	Intensity	0.4<Width<0.6
45.00	44.99	-0.01	0.49	4.51E+06	PASS
411.26	411.25	-0.01	0.51	1.45E+05	PASS
585.39	585.38	0.00	0.50	2.43E+05	PASS
933.64	933.63	-0.01	0.50	9.82E+05	PASS
1223.85	1223.84	0.00	0.56	2.68E+05	PASS

Tune 2021-11-23 Q3 NEG @ 10Da/s

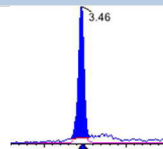
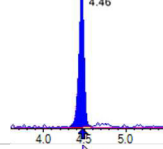
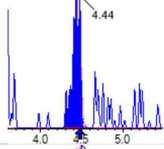
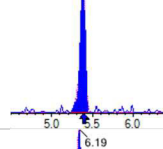
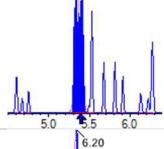
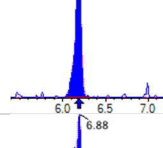
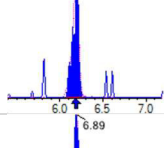
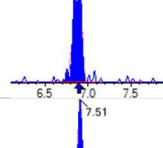
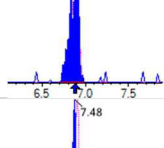
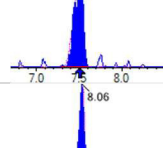
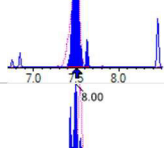
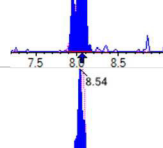
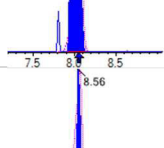
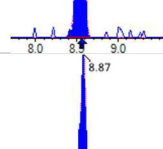
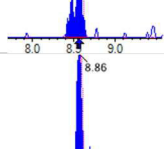
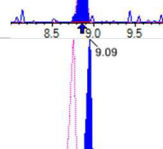
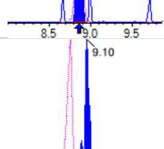
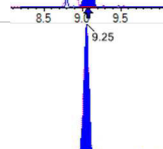
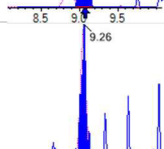
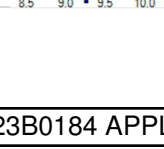
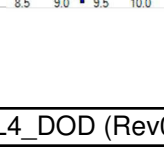
Q3 Scan Test started. Scan Speed = 10 Da/s

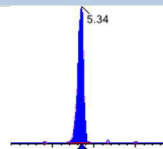
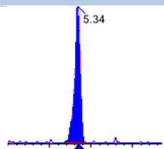
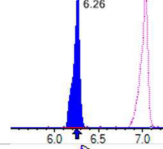
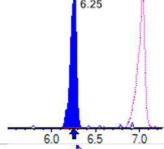
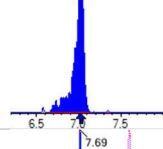
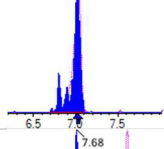
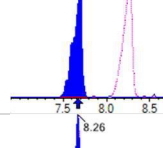
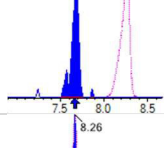
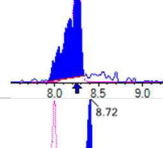
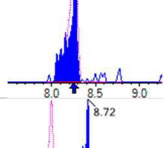
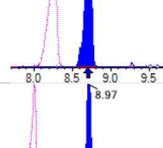
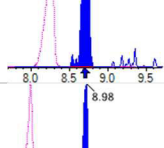
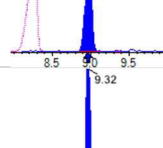
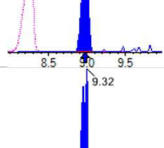
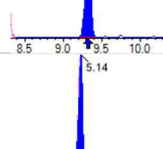
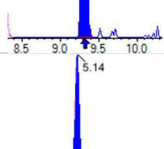
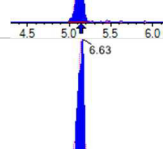
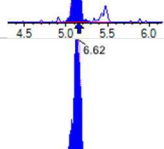
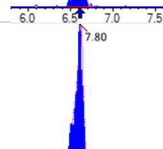
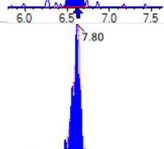
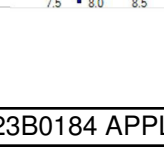
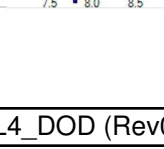


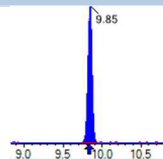
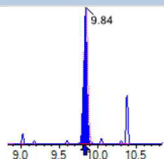
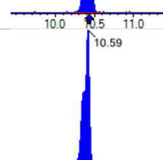
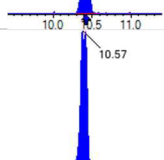
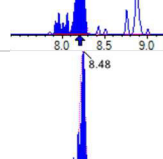
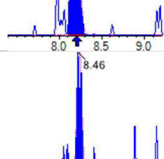
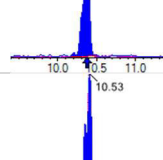
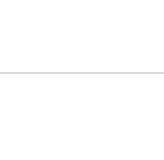
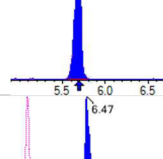
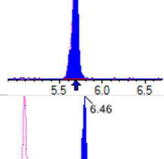
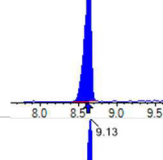
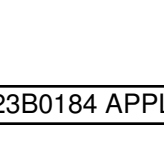
Target Mass	Found At	Delta	Width	Intensity	0.4<Width<0.6
45.00	44.99	0.00	0.51	7.20E+05	PASS
411.26	411.26	0.00	0.56	2.58E+04	PASS
585.39	585.37	-0.01	0.57	1.33E+05	PASS
933.64	933.64	0.00	0.56	3.47E+05	PASS
1223.85	1223.83	-0.01	0.56	6.20E+04	PASS

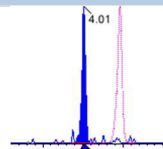
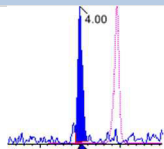
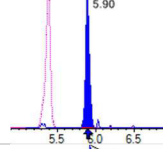
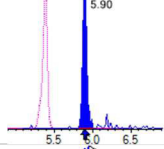
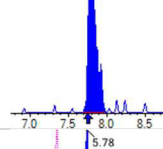
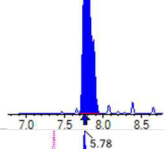
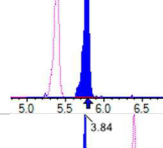
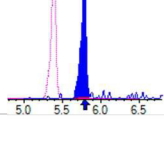
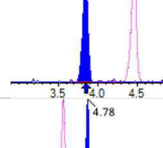
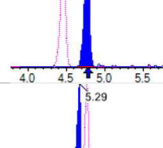
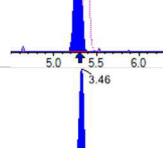
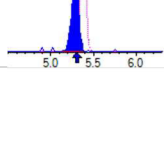
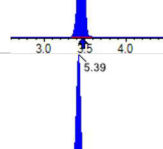
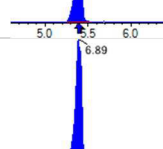
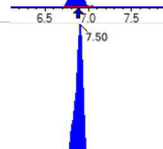
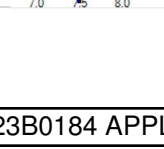
EPA 1633

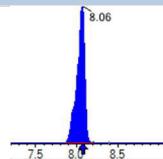
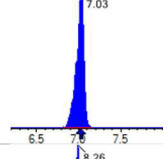
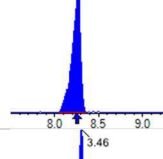
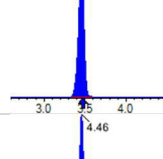
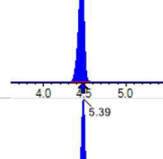
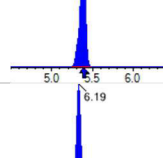
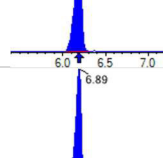
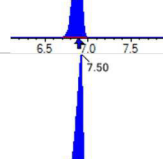
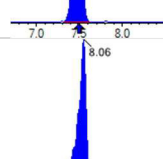
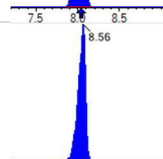
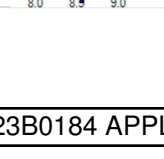
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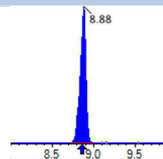
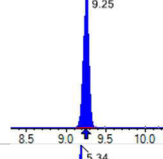
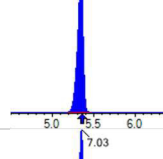
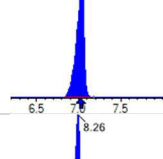
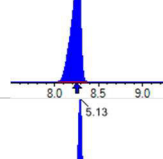
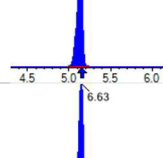
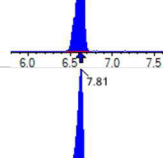
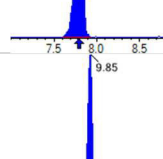
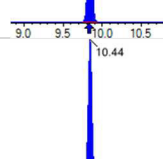
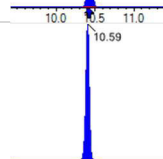
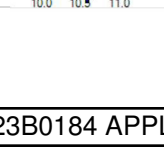
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-Imin, ΔRT-CVmin, ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration True ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) 159187	(3.46, 1.00) (0.00, N/A, 0.0)	82.9	N/A 0.0 0.0	0.4193 [0.4000]	104.8%			
PFPeA	(263.0 / 219.0) 136655 (263.0 / 69.0) 3570	(4.46, 1.00) (0.00, N/A, 1.2)	278.7 339.0	0.0261 219.2 219.2	0.2242 [0.2000]	112.1%			
PFHxA	(313.0 / 269.0) 78998 (313.0 / 119.0) 8528	(5.38, 1.00) (0.00, N/A, 0.9)	368.0 4622.7	0.1079 115.4 115.4	0.1037 [0.1000]	103.7%			
PFHpA	(363.0 / 319.0) 78992 (363.0 / 169.0) 18800	(6.19, 1.00) (0.00, N/A, -0.6)	1838.3 291.6	0.2380 75.4 75.4	0.1110 [0.1000]	111.0%			
PFOA	(413.0 / 369.0) 101357 (413.0 / 169.0) 41752	(6.88, 1.00) (-0.01, N/A, -0.3)	630.4 550231.2	0.4119 121.8 121.8	0.1196 [0.1000]	119.6%			
PFNA	(463.0 / 419.0) 83364 (463.0 / 169.0) 11049	(7.51, 1.00) (0.01, N/A, 1.9)	6507.3 1653.9	0.1325 56.6 56.6	0.1172 [0.1000]	117.2%			
PFDA	(513.0 / 469.0) 102743 (513.0 / 169.0) 8244	(8.06, 1.00) (-0.01, N/A, 3.2)	136.6 156320.6	0.0802 64.7 64.7	0.1147 [0.1000]	114.7%			
PFUnA	(563.0 / 519.0) 90007 (563.0 / 169.0) 16176	(8.54, 1.00) (-0.02, N/A, -0.8)	96.0 114.7	0.1797 167.0 167.0	0.1225 [0.1000]	122.5%			IR2,
PFDoA	(613.0 / 569.0) 66395 (613.0 / 169.0) 10579	(8.87, 1.00) (0.00, N/A, 1.0)	131.6 124.0	0.1593 100.3 100.3	0.0872 [0.1000]	87.2%			
PFTTrDA	(663.0 / 619.0) 80259 (663.0 / 169.0) 22186	(9.09, 1.02) (N/A, 0.02, -0.5)	592.8 515.8	0.2764 101.8 101.8	0.1095 [0.1000]	109.5%			
PFTeDA	(713.0 / 669.0) 85816 (713.0 / 169.0) 12133	(9.25, 1.00) (0.00, N/A, -0.5)	175.8 40.7	0.1414 64.7 64.7	0.1107 [0.1000]	110.7%			

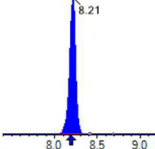
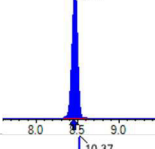
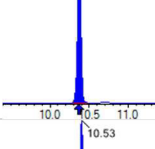
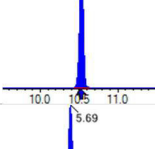
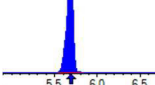
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-Imin, ΔRT-CVmin, ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration True ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) 135356 (299.0 / 99.0) 101238	(5.34, 1.00) (0.00, N/A, 0.1)	678.8 283.6	0.7479 118.9 118.9	0.0919 [0.0885]	103.9%			
PFPeS	(349.0 / 80.0) 252841 (349.0 / 99.0) 76134	(6.26, 0.89) (N/A, 0.01, 0.4)	370518.5 1638.4	0.3011 90.0 90.0	0.1054 [0.0938]	112.3%			
PFHxS	(399.0 / 80.0) 204921 (399.0 / 99.0) 64909	(7.03, 1.00) (0.00, N/A, 0.3)	551.2 2436.7	0.3168 99.0 99.0	0.0974 [0.0911]	106.9%			
PFHpS	(449.0 / 80.0) 213085 (449.0 / 99.0) 51597	(7.69, 0.93) (N/A, 0.03, 0.6)	1372.3 17063218.5	0.2421 89.2 89.2	0.0974 [0.0951]	102.4%			
PFOS	(499.0 / 80.0) 289776 (499.0 / 99.0) 54053	(8.26, 1.00) (0.00, N/A, -0.3)	77.6 65.7	0.1865 84.8 84.8	0.1081 [0.0927]	116.5%			
PFNS	(549.0 / 80.0) 298030 (549.0 / 99.0) 71535	(8.72, 1.06) (N/A, 0.02, -0.3)	2375659.5 202.7	0.2400 99.7 99.7	0.1111 [0.0960]	115.7%			
PFDS	(599.0 / 80.0) 302138 (599.0 / 99.0) 56240	(8.97, 1.09) (N/A, 0.01, -0.4)	385.1 297.5	0.1861 92.0 92.0	0.1013 [0.0963]	105.2%			
PFDoS	(699.0 / 80.0) 141394 (699.0 / 99.0) 41165	(9.32, 1.13) (N/A, 0.01, 0.2)	575.3 167.2	0.2911 149.0 149.0	0.1027 [0.0970]	105.9%			
4:2FTS	(327.0 / 307.0) 174639 (327.0 / 81.0) 126319	(5.14, 1.00) (0.00, N/A, 0.0)	619.3 215.3	0.7233 104.4 104.4	0.3678 [0.3738]	98.4%			
6:2FTS	(427.0 / 407.0) 82195 (427.0 / 81.0) 69148	(6.63, 1.00) (0.00, N/A, 0.5)	486.5 161.6	0.8413 99.3 99.3	0.3501 [0.3796]	92.2%			
8:2FTS	(527.0 / 507.0) 82323 (527.0 / 81.0) 57566	(7.80, 1.00) (-0.01, N/A, -0.1)	934.2 282.0	0.6993 85.7 85.7	0.4287 [0.3833]	111.8%			

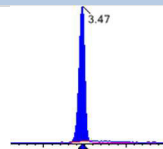
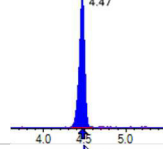
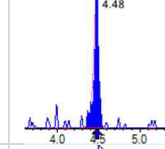
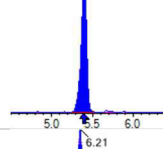
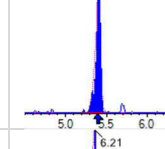
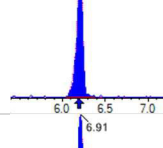
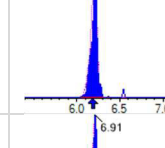
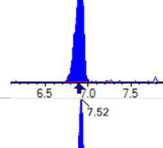
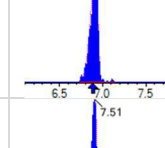
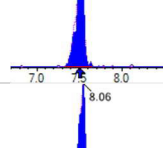
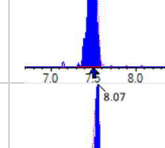
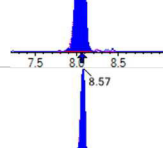
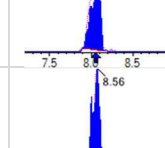
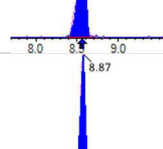
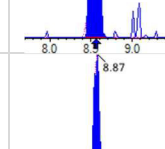
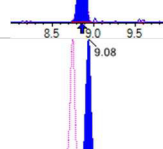
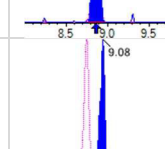
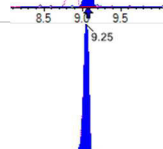
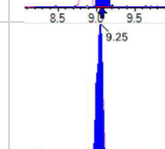
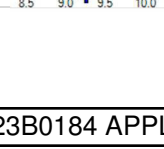
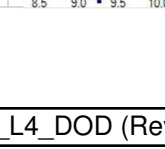
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-Imin, ΔRT-CVmin, ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration True ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) 408275 (498.0 / 478.0) 6632	(9.85, 1.00) (0.00, N/A, 0.4)	695.9 320.1	0.0162 82.7 82.7	0.1075 [0.1000]	107.5%			
NMeFOSA	(512.0 / 219.0) 316291 (512.0 / 169.0) 280188	(10.44, 1.00) (0.00, N/A, 0.8)	1451.0 1593.5	0.8859 107.2 107.2	0.4138 [0.4000]	103.5%			
NEIFOSA	(526.0 / 219.0) 340043 (526.0 / 169.0) 423688	(10.59, 1.00) (0.00, N/A, 1.0)	1539.8 1055.4	1.2460 98.4 98.4	0.4364 [0.4000]	109.1%			
NMeFOSAA	(570.0 / 419.0) 33321 (570.0 / 483.0) 17058	(8.22, 1.00) (0.01, N/A, 0.2)	269.8 73.6	0.5119 104.4 104.4	0.1164 [0.1000]	116.4%			
NEIFOSAA	(584.0 / 419.0) 28401 (584.0 / 526.0) 9020	(8.48, 1.00) (0.01, N/A, 0.8)	60179.4 330.5	0.3176 56.2 56.2	0.1100 [0.1000]	110.0%			
NMeFOSE	(616.0 / 59.0) 158489	(10.38, 1.00) (0.01, N/A, 0.0)	444.7	N/A 0.0 0.0	0.4432 [0.4000]	110.8%			
NEtFOSE	(630.0 / 59.0) 159842	(10.53, 1.00) (0.00, N/A, 0.0)	280.1	N/A 0.0 0.0	0.4426 [0.4000]	110.7%			
HFPO-DA	(285.0 / 169.0) 98966 (285.0 / 185.0) 204157	(5.69, 1.00) (0.00, N/A, -0.2)	695.0 375.7	2.0629 75.0 75.0	0.2247 [0.2000]	112.4%			
ADONA	(377.0 / 85.0) 350095 (377.0 / 251.0) 35281	(6.47, 1.14) (N/A, 0.00, 0.2)	426.5 135.0	0.1008 115.1 115.1	0.2142 [0.1885]	113.6%			
9CI-Pf3ONS	(531.0 / 351.0) 808793 (533.0 / 353.0) 215408	(8.63, 1.52) (N/A, 0.02, -0.2)	436.7 423.1	0.2663 90.4 90.4	0.2110 [0.1867]	113.0%			
11CI-PF3OUDS	(631.0 / 451.0) 401624 (633.0 / 453.0) 118778	(9.13, 1.61) (N/A, 0.02, 0.6)	1569.6 168942.3	0.2957 95.8 95.8	0.2131 [0.1886]	113.0%			

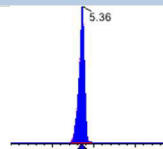
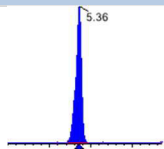
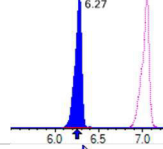
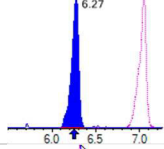
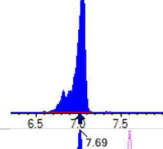
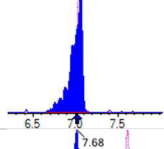
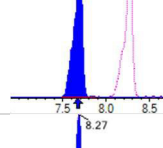
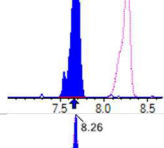
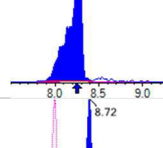
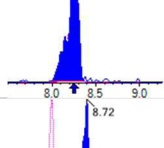
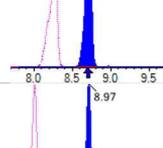
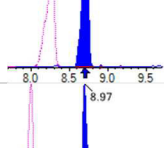
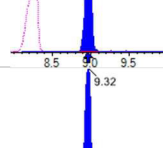
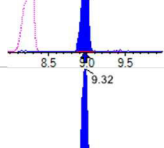
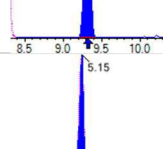
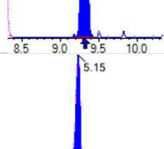
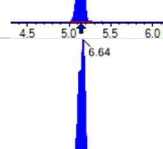
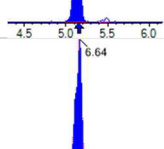
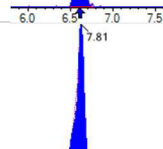
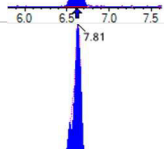
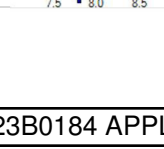
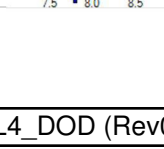
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-Imin, ΔRT-CVmin, ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration True ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) 6406 (241.0 / 117.0) 11620	(4.01, 0.90) (N/A, 0.00, 0.7)	141.0 65.8	1.8138 104.6 104.6	0.3558 [0.4000]	89.0%			
5:3FTCA	(341.0 / 236.7) 44851 (341.0 / 217.0) 93376	(5.90, 1.09) (N/A, 0.00, -0.2)	891.8 200.9	2.0819 123.8 123.8	0.4120 [0.4000]	103.0%			
7:3FTCA	(441.0 / 317.0) 97878 (441.0 / 337.0) 87848	(7.80, 1.45) (N/A, 0.01, 0.8)	121.2 179.6	0.8975 104.5 104.5	0.5001 [0.4000]	125.0%			
PFEESA	(315.0 / 135.0) 199103 (315.0 / 83.0) 61742	(5.78, 1.07) (N/A, 0.00, -0.3)	632.6 130.3	0.3101 110.2 110.2	0.2104 [0.1785]	117.9%			
PFMPA	(229.0 / 85.0) 34878	(3.84, 0.86) (N/A, -0.01, 0.0)	434.7	N/A 0.0 0.0	0.2238 [0.2000]	111.9%			
PFMBA	(279.0 / 85.0) 105375	(4.78, 1.07) (N/A, -0.01, 0.0)	463.3	N/A 0.0 0.0	0.1973 [0.2000]	98.7%			
NFDHA	(295.0 / 201.0) 95136 (295.0 / 85.0) 86110	(5.29, 0.98) (N/A, -0.01, 0.3)	1710.6 601.2	0.9051 86.7 86.7	0.2246 [0.2000]	112.3%			
13C3_PFBa_IIS	(216.0 / 172.0) 403828	(3.46, N/A) (N/A, -0.01, N/A)	1783.9	N/A	1.1912 [1.0000]	119.1% {122.5%}			
13C2_PFHxA_IIS	(315.0 / 270.0) 639133	(5.39, N/A) (N/A, -0.01, N/A)	1509.0	N/A	1.0746 [1.0000]	107.5% {115.0%}			
13C4_PFOA_IIS	(417.0 / 372.0) 834165	(6.89, N/A) (N/A, 0.01, N/A)	826.1	N/A	1.0255 [1.0000]	102.5% {111.2%}			
13C5_PFNA_IIS	(468.0 / 423.0) 783782	(7.50, N/A) (N/A, 0.01, N/A)	550.9	N/A	1.0264 [1.0000]	102.6% {102.9%}			

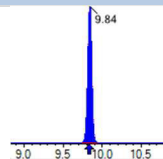
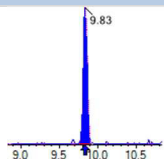
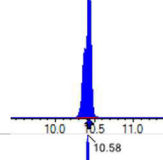
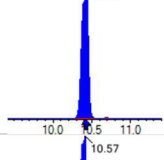
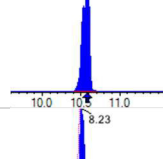
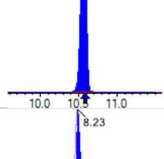
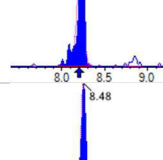
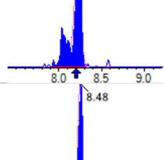
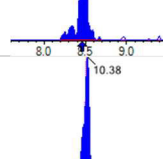
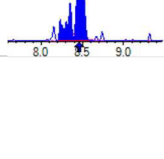
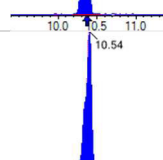
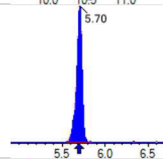
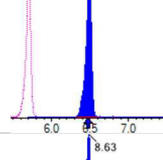
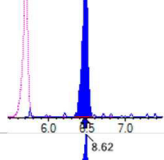
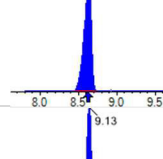
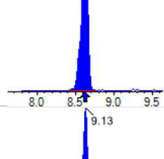
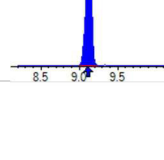
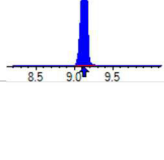
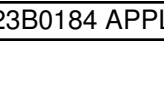
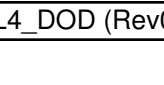
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-Imin, ΔRT-CVmin, ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration True ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 760809	(8.06, N/A) (N/A, 0.01, N/A)	862.0	N/A	1.0598 [1.0000]	106.0% {110.6%}			
18O2_PFHxS_IIS	(403.0 / 83.9) 1273266	(7.03, N/A) (N/A, 0.01, N/A)	1287.1	N/A	1.0339 [1.0000]	103.4% {106.0%}			
13C4_PFOS_IIS	(503.0 / 79.9) 1972538	(8.26, N/A) (N/A, 0.02, N/A)	1173.9	N/A	1.1114 [1.0000]	111.1% {113.9%}			
13C4_PFBA_EIS	(217.0 / 172.0) 3591649	(3.46, N/A) (N/A, -0.01, N/A)	2494.4	N/A	7.4797 [8.0000]	93.5% {122.6%}			
13C5_PFPeA_EIS	(268.0 / 223.0) 2767434	(4.46, N/A) (N/A, -0.01, N/A)	2035.0	N/A	3.8860 [4.0000]	97.2% {113.5%}			
13C5_PFHxA_EIS	(318.0 / 273.0) 1632151	(5.39, N/A) (N/A, -0.01, N/A)	1439.9	N/A	1.8795 [2.0000]	94.0% {106.3%}			
13C4_PFHpA_EIS	(367.0 / 322.0) 1594218	(6.19, N/A) (N/A, 0.00, N/A)	1543.0	N/A	1.9684 [2.0000]	98.4% {112.8%}			
13C8_PFOA_EIS	(421.0 / 376.0) 1758649	(6.89, N/A) (N/A, 0.01, N/A)	1993.9	N/A	1.9697 [2.0000]	98.5% {105.5%}			
13C9_PFNA_EIS	(472.0 / 427.0) 753093	(7.50, N/A) (N/A, 0.01, N/A)	1064.0	N/A	0.9642 [1.0000]	96.4% {106.2%}			
13C6_PFDA_EIS	(519.0 / 474.0) 903669	(8.06, N/A) (N/A, 0.03, N/A)	979.0	N/A	1.0262 [1.0000]	102.6% {123.0%}			
13C7_PFUnA_EIS	(570.0 / 525.0) 835427	(8.56, N/A) (N/A, 0.02, N/A)	1724.6	N/A	0.9576 [1.0000]	95.8% {111.0%}			

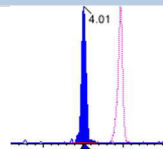
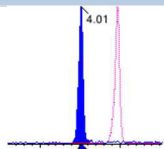
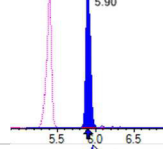
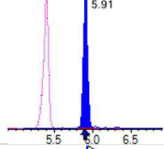
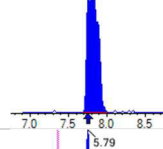
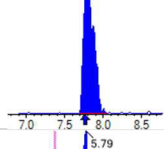
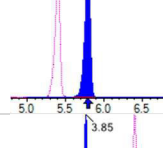
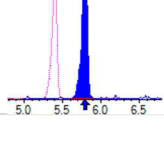
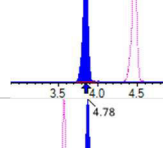
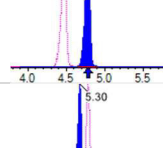
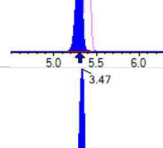
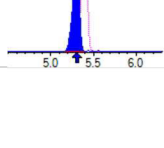
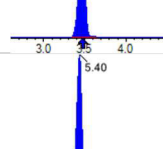
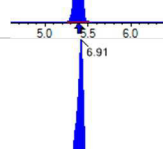
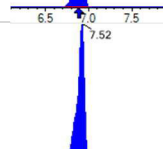
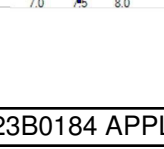
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-Imin, ΔRT-CVmin, ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration True ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 872979	(8.88, N/A) (N/A, 0.02, N/A)	918.6	N/A	1.0790 [1.0000]	107.9% {118.5%}			
13C2_PFTeDA_EIS	(715.0 / 670.0) 843285	(9.25, N/A) (N/A, 0.01, N/A)	918.5	N/A	1.0026 [1.0000]	100.3% {117.3%}			
13C3_PFBs_EIS	(302.0 / 80.0) 5181335	(5.34, N/A) (N/A, -0.01, N/A)	2161.9	N/A	2.0137 [2.0000]	100.7% {113.9%}			
13C3_PFHxS_EIS	(402.0 / 80.0) 2671902	(7.03, N/A) (N/A, 0.01, N/A)	1576.4	N/A	1.9876 [2.0000]	99.4% {112.8%}			
13C8_PFOS_EIS	(507.0 / 80.0) 4860196	(8.26, N/A) (N/A, 0.02, N/A)	1444.5	N/A	1.8755 [2.0000]	93.8% {113.8%}			
13C2_4:2FTS_EIS	(329.0 / 81.0) 672936	(5.13, N/A) (N/A, -0.01, N/A)	1458.1	N/A	4.2700 [4.0000]	106.8% {129.8%}			
13C2_6:2FTS_EIS	(429.0 / 81.0) 719951	(6.63, N/A) (N/A, 0.01, N/A)	1080.0	N/A	4.0519 [4.0000]	101.3% {121.9%}			
13C2_8:2FTS_EIS	(529.0 / 81.0) 684191	(7.81, N/A) (N/A, 0.02, N/A)	1141.8	N/A	3.6413 [4.0000]	91.0% {103.0%}			
13C8_PFOsa_EIS	(506.0 / 78.0) 9772490	(9.85, N/A) (N/A, 0.02, N/A)	2857.8	N/A	2.0472 [2.0000]	102.4% {125.4%}			
D3_NMeFOsa_EIS	(515.0 / 169.0) 1779886	(10.44, N/A) (N/A, 0.01, N/A)	2747.6	N/A	1.7364 [2.0000]	86.8% {103.5%}			
D5_NEtFOsa_EIS	(531.0 / 169.0) 1604328	(10.59, N/A) (N/A, 0.01, N/A)	3172.7	N/A	1.7875 [2.0000]	89.4% {102.6%}			

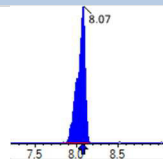
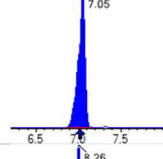
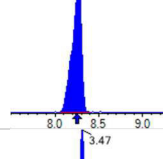
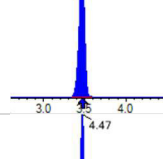
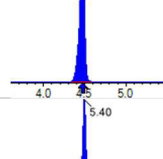
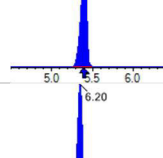
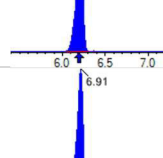
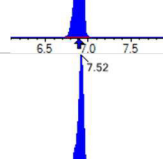
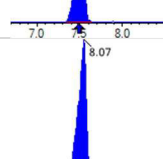
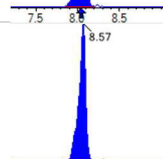
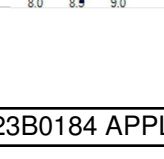
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-Imin, ΔRT-CVmin, ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration True ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1340165	(8.21 , N/A) (N/A , 0.02 , N/A)	998.5	N/A	3.6849 [4.0000]	92.1% { 104.2% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 1184365	(8.47 , N/A) (N/A , 0.01 , N/A)	6284.7	N/A	3.8242 [4.0000]	95.6% { 122.4% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 6825210	(10.37 , N/A) (N/A , 0.01 , N/A)	1607.3	N/A	19.5582 [20.0000]	97.8% { 112.8% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 7147806	(10.53 , N/A) (N/A , 0.01 , N/A)	1582.6	N/A	17.9062 [20.0000]	89.5% { 104.4% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 4297700	(5.69 , N/A) (N/A , 0.00 , N/A)	2513.5	N/A	7.8759 [8.0000]	98.4% { 114.9% }			

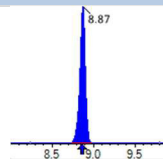
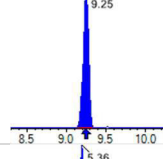
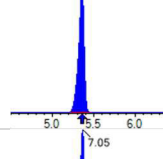
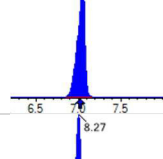
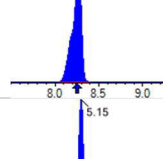
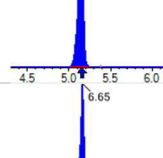
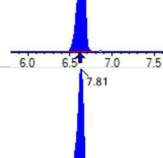
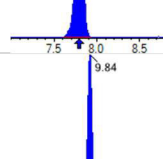
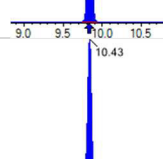
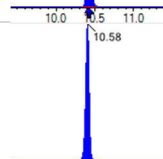
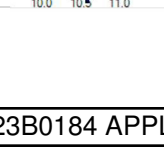
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) 755137	(3.47 , 1.00) (0.00 , N/A , 0.0)	128.6	N/A 0.0 0.0	1.7753 [2.0000]	88.8%			
PFPeA	(263.0 / 219.0) 599742 (263.0 / 69.0) 8058	(4.47 , 1.00) (0.00 , N/A , -0.4)	871.2 78.1	0.0134 112.7 112.7	0.8703 [1.0000]	87.0%			
PFHxA	(313.0 / 269.0) 406807 (313.0 / 119.0) 44060	(5.40 , 1.00) (0.00 , N/A , -0.5)	1470.0 164687.3	0.1083 115.8 115.8	0.4749 [0.5000]	95.0%			
PFHpA	(363.0 / 319.0) 371706 (363.0 / 169.0) 123076	(6.21 , 1.00) (0.00 , N/A , -0.4)	2643.1 48307.7	0.3311 104.9 104.9	0.4815 [0.5000]	96.3%			
PFOA	(413.0 / 369.0) 406570 (413.0 / 169.0) 138645	(6.91 , 1.00) (0.00 , N/A , -0.3)	1174.8 1146.7	0.3410 100.8 100.8	0.4368 [0.5000]	87.4%			
PFNA	(463.0 / 419.0) 365881 (463.0 / 169.0) 79262	(7.52 , 1.00) (0.00 , N/A , 0.8)	3933.7 1470.3	0.2166 92.5 92.5	0.4594 [0.5000]	91.9%			
PFDA	(513.0 / 469.0) 398042 (513.0 / 169.0) 45316	(8.06 , 1.00) (0.00 , N/A , -0.6)	333.7 53.8	0.1138 91.9 91.9	0.4433 [0.5000]	88.7%			
PFUnA	(563.0 / 519.0) 392695 (563.0 / 169.0) 44799	(8.57 , 1.00) (0.00 , N/A , 0.6)	521.6 109.5	0.1141 106.0 106.0	0.4731 [0.5000]	94.6%			
PFDoA	(613.0 / 569.0) 325040 (613.0 / 169.0) 64869	(8.87 , 1.00) (0.00 , N/A , 0.4)	441.0 1153.8	0.1996 125.6 125.6	0.4402 [0.5000]	88.0%			
PFTTrDA	(663.0 / 619.0) 367309 (663.0 / 169.0) 91442	(9.08 , 1.02) (N/A , 0.01 , -0.3)	614.8 534.0	0.2490 91.7 91.7	0.5169 [0.5000]	103.4%			
PFTeDA	(713.0 / 669.0) 422087 (713.0 / 169.0) 88136	(9.25 , 1.00) (0.00 , N/A , 0.3)	669.9 185.5	0.2088 95.5 95.5	0.5011 [0.5000]	100.2%			

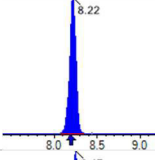
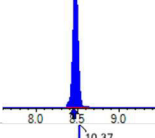
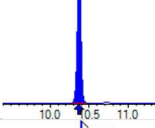
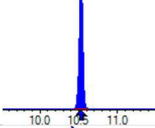
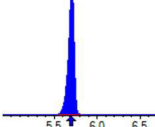
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) 718738 (299.0 / 99.0) 402042	(5.36, 1.00) (0.00, N/A, 0.0)	1415.0 677.2	0.5594 88.9 88.9	0.4229 [0.4424]	95.6%			
PFPeS	(349.0 / 80.0) 1150231 (349.0 / 99.0) 375229	(6.27, 0.89) (N/A, 0.02, 0.2)	1273024.4 2293.2	0.3262 97.5 97.5	0.4493 [0.4692]	95.8%			
PFHxS	(399.0 / 80.0) 951264 (399.0 / 99.0) 314947	(7.04, 1.00) (0.00, N/A, -0.5)	75741.3 718.5	0.3311 103.5 103.5	0.4235 [0.4555]	93.0%			
PFHpS	(449.0 / 80.0) 992727 (449.0 / 99.0) 265895	(7.69, 0.93) (N/A, 0.03, 0.5)	6833460.3 1532.1	0.2678 98.7 98.7	0.4297 [0.4757]	90.3%			
PFOS	(499.0 / 80.0) 1239344 (499.0 / 99.0) 254523	(8.27, 1.00) (0.00, N/A, 0.4)	192.7 172.4	0.2054 93.4 93.4	0.4377 [0.4637]	94.4%			
PFNS	(549.0 / 80.0) 1307203 (549.0 / 99.0) 313967	(8.72, 1.05) (N/A, 0.01, 0.1)	2179.6 70069509.6	0.2402 99.8 99.8	0.4613 [0.4799]	96.1%			
PFDS	(599.0 / 80.0) 1372070 (599.0 / 99.0) 303289	(8.97, 1.09) (N/A, 0.01, 0.2)	937.2 632.8	0.2210 109.2 109.2	0.4357 [0.4816]	90.5%			
PFDoS	(699.0 / 80.0) 610283 (699.0 / 99.0) 123735	(9.32, 1.13) (N/A, 0.01, 0.3)	978.6 307.0	0.2028 103.8 103.8	0.4198 [0.4848]	86.6%			
4:2FTS	(327.0 / 307.0) 860332 (327.0 / 81.0) 572842	(5.15, 1.00) (0.00, N/A, 0.1)	1361.7 672.6	0.6658 96.1 96.1	1.7863 [1.8691]	95.6%			
6:2FTS	(427.0 / 407.0) 413476 (427.0 / 81.0) 377891	(6.64, 1.00) (-0.01, N/A, -0.3)	873.2 587.0	0.9139 107.9 107.9	1.7091 [1.8981]	90.0%			
8:2FTS	(527.0 / 507.0) 387778 (527.0 / 81.0) 273625	(7.81, 1.00) (0.00, N/A, 0.0)	539.4 448.3	0.7056 86.5 86.5	1.8041 [1.9166]	94.1%			

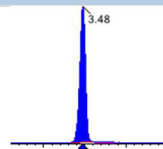
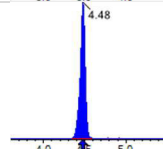
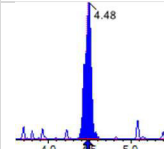
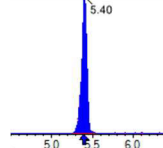
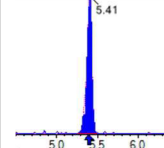
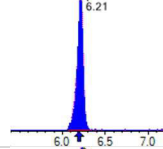
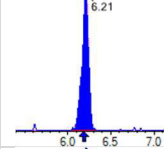
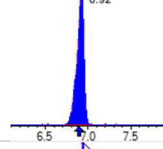
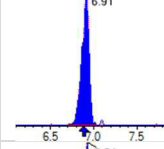
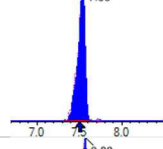
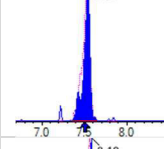
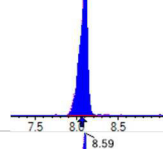
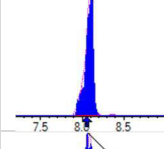
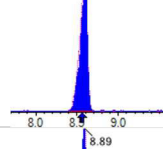
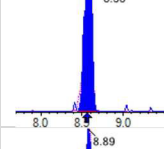
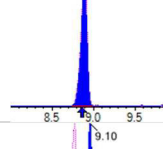
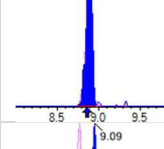
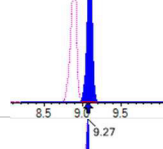
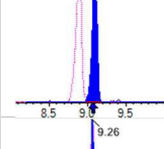
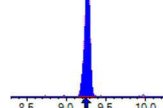
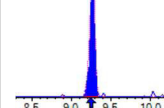
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) 1805841 (498.0 / 478.0) 40828	(9.84 , 1.00) (0.00 , N/A , 0.3)	2311.4 413.8	0.0226 115.1 115.1	0.4710 [0.5000]	94.2%			
NMeFOSA	(512.0 / 219.0) 1597506 (512.0 / 169.0) 1332810	(10.43 , 1.00) (0.00 , N/A , 1.1)	4495.4 3408.9	0.8343 101.0 101.0	1.8884 [2.0000]	94.4%			
NEIFOSA	(526.0 / 219.0) 1699037 (526.0 / 169.0) 2143008	(10.58 , 1.00) (-0.01 , N/A , 0.7)	4828.4 4504.2	1.2613 99.6 99.6	1.8685 [2.0000]	93.4%			
NMeFOSAA	(570.0 / 419.0) 132402 (570.0 / 483.0) 74102	(8.23 , 1.00) (0.02 , N/A , 0.4)	416.0 148.8	0.5597 114.1 114.1	0.4511 [0.5000]	90.2%			
NEIFOSAA	(584.0 / 419.0) 117619 (584.0 / 526.0) 67112	(8.48 , 1.00) (0.01 , N/A , -0.1)	11890.3 745.2	0.5706 101.0 101.0	0.4226 [0.5000]	84.5%			
NMeFOSE	(616.0 / 59.0) 679842	(10.38 , 1.00) (0.01 , N/A , 0.0)	966.5	N/A 0.0 0.0	1.8027 [2.0000]	90.1%			
NEtFOSE	(630.0 / 59.0) 761835	(10.54 , 1.00) (0.01 , N/A , 0.0)	950.5	N/A 0.0 0.0	1.7651 [2.0000]	88.3%			
HFPO-DA	(285.0 / 169.0) 420161 (285.0 / 185.0) 1079380	(5.70 , 1.00) (0.00 , N/A , 0.0)	1488.6 1172.5	2.5690 93.4 93.4	0.8951 [1.0000]	89.5%			
ADONA	(377.0 / 85.0) 1498766 (377.0 / 251.0) 136684	(6.49 , 1.14) (N/A , 0.03 , 0.7)	1154.5 195.9	0.0912 104.2 104.2	0.8603 [0.9427]	91.3%			
9CI-Pf3ONS	(531.0 / 351.0) 3523077 (533.0 / 353.0) 1048371	(8.63 , 1.51) (N/A , 0.02 , 0.3)	1108.0 609.3	0.2976 101.0 101.0	0.8623 [0.9333]	92.4%			
11CI-PF3OUDS	(631.0 / 451.0) 1691673 (633.0 / 453.0) 602232	(9.13 , 1.60) (N/A , 0.01 , -0.1)	977.9 1870.1	0.3560 115.4 115.4	0.8422 [0.9432]	89.3%			

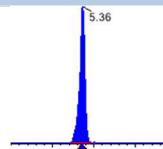
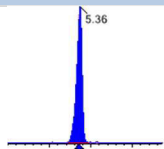
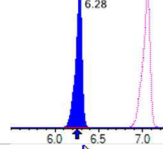
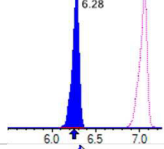
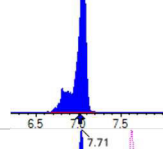
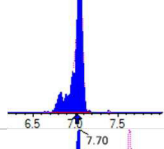
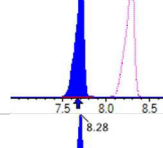
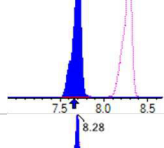
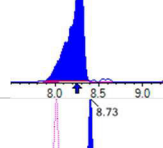
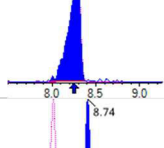
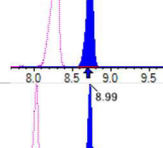
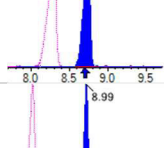
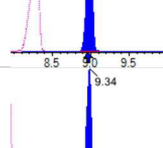
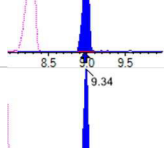
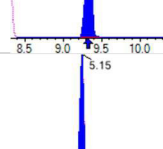
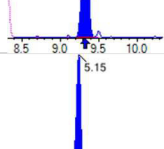
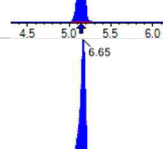
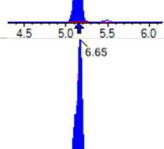
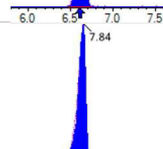
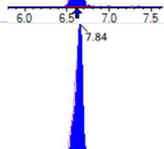
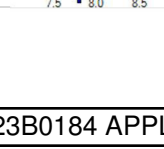
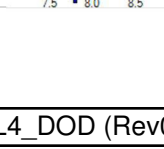
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) 38619 (241.0 / 117.0) 60683	(4.01 , 0.90) (N/A , 0.00 , -0.2)	520.0 314.2	1.5713 90.6 90.6	1.8976 [2.0000]	94.9%			
5:3FTCA	(341.0 / 236.7) 210131 (341.0 / 217.0) 377139	(5.90 , 1.09) (N/A , 0.01 , -0.3)	1291.8 727.5	1.7948 106.7 106.7	1.7160 [2.0000]	85.8%			
7:3FTCA	(441.0 / 317.0) 378938 (441.0 / 337.0) 344182	(7.81 , 1.45) (N/A , 0.02 , 0.4)	368.4 376.1	0.9083 105.8 105.8	1.7212 [2.0000]	86.1%			
PFEESA	(315.0 / 135.0) 867939 (315.0 / 83.0) 256094	(5.79 , 1.07) (N/A , 0.01 , -0.1)	1500.2 349.0	0.2951 104.8 104.8	0.8153 [0.8925]	91.4%			
PFMPA	(229.0 / 85.0) 152777	(3.85 , 0.86) (N/A , 0.00 , 0.0)	1068.2	N/A 0.0 0.0	0.8671 [1.0000]	86.7%			
PFMBA	(279.0 / 85.0) 537128	(4.78 , 1.07) (N/A , 0.00 , 0.0)	1545.8	N/A 0.0 0.0	0.8898 [1.0000]	89.0%			
NFDHA	(295.0 / 201.0) 436352 (295.0 / 85.0) 439776	(5.30 , 0.98) (N/A , 0.01 , 0.1)	981.4 1268.0	1.0078 96.5 96.5	0.9158 [1.0000]	91.6%			
13C3_PFBa_IIS	(216.0 / 172.0) 377842	(3.47 , N/A) (N/A , 0.00 , N/A)	1643.3	N/A	1.1145 [1.0000]	111.4% { 114.6% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 624524	(5.40 , N/A) (N/A , 0.00 , N/A)	1392.3	N/A	1.0500 [1.0000]	105.0% { 112.4% }			
13C4_PFOA_IIS	(417.0 / 372.0) 871802	(6.91 , N/A) (N/A , 0.03 , N/A)	1712.9	N/A	1.0718 [1.0000]	107.2% { 116.2% }			
13C5_PFNAl_IIS	(468.0 / 423.0) 831114	(7.52 , N/A) (N/A , 0.03 , N/A)	1511.9	N/A	1.0884 [1.0000]	108.8% { 109.2% }			

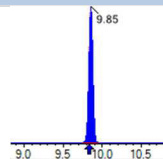
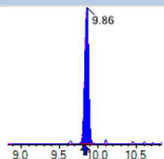
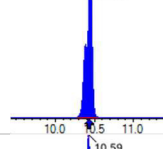
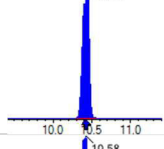
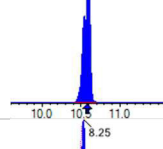
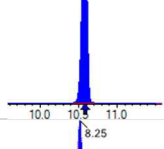
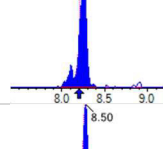
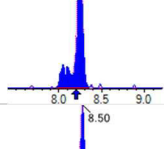
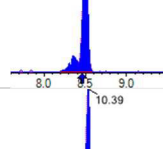
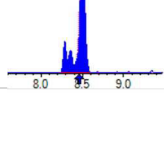
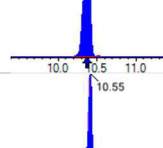
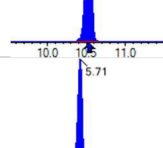
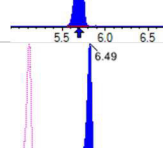
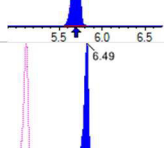
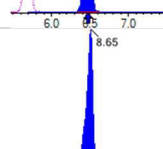
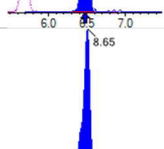
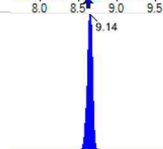
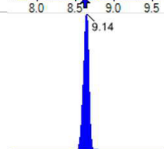
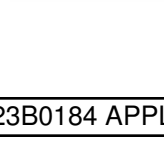
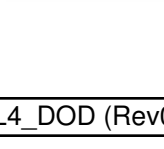
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 701666	(8.07 , N/A) (N/A , 0.02 , N/A)	721.9	N/A	0.9774 [1.0000]	97.7% { 102.0% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 1247541	(7.05 , N/A) (N/A , 0.03 , N/A)	1131.6	N/A	1.0130 [1.0000]	101.3% { 103.9% }			
13C4_PFOS_IIS	(503.0 / 79.9) 1747657	(8.26 , N/A) (N/A , 0.03 , N/A)	870.2	N/A	0.9847 [1.0000]	98.5% { 100.9% }			
13C4_PFBA_EIS	(217.0 / 172.0) 4023777	(3.47 , N/A) (N/A , 0.00 , N/A)	2704.2	N/A	8.9559 [8.0000]	111.9% { 137.4% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 3128479	(4.47 , N/A) (N/A , 0.00 , N/A)	2367.3	N/A	4.4958 [4.0000]	112.4% { 128.3% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 1835857	(5.40 , N/A) (N/A , 0.01 , N/A)	1541.1	N/A	2.1635 [2.0000]	108.2% { 119.6% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 1728938	(6.20 , N/A) (N/A , 0.02 , N/A)	1846.2	N/A	2.1846 [2.0000]	109.2% { 122.3% }			
13C8_PFOA_EIS	(421.0 / 376.0) 1932206	(6.91 , N/A) (N/A , 0.03 , N/A)	1467.1	N/A	2.0707 [2.0000]	103.5% { 116.0% }			
13C9_PFNA_EIS	(472.0 / 427.0) 843176	(7.52 , N/A) (N/A , 0.03 , N/A)	1032.4	N/A	1.0180 [1.0000]	101.8% { 118.9% }			
13C6_PFDA_EIS	(519.0 / 474.0) 905717	(8.07 , N/A) (N/A , 0.03 , N/A)	622.4	N/A	1.1152 [1.0000]	111.5% { 123.3% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 943818	(8.57 , N/A) (N/A , 0.02 , N/A)	1485.8	N/A	1.1731 [1.0000]	117.3% { 125.4% }			

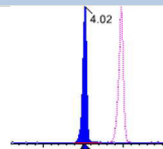
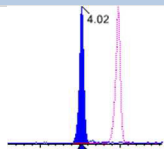
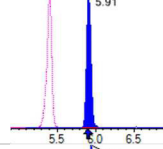
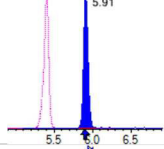
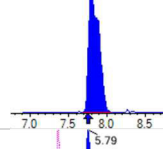
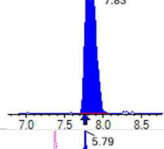
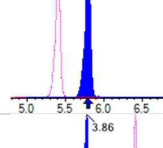
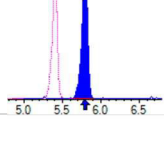
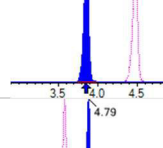
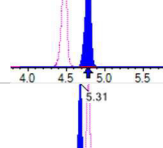
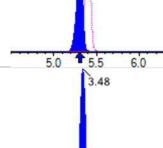
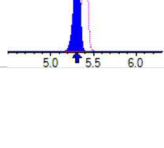
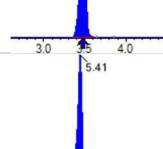
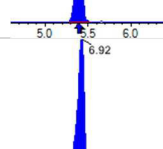
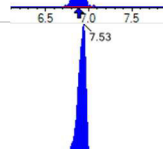
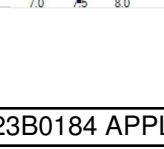
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 846417	(8.87, N/A) (N/A, 0.01, N/A)	1142.9	N/A	1.1343 [1.0000]	113.4% { 114.9% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 916114	(9.25, N/A) (N/A, 0.01, N/A)	1267.0	N/A	1.1810 [1.0000]	118.1% { 127.4% }			
13C3_PFBs_EIS	(302.0 / 80.0) 5977954	(5.36, N/A) (N/A, 0.01, N/A)	2326.7	N/A	2.3713 [2.0000]	118.6% { 131.4% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 2851985	(7.05, N/A) (N/A, 0.03, N/A)	1570.3	N/A	2.1654 [2.0000]	108.3% { 120.4% }			
13C8_PFOS_EIS	(507.0 / 80.0) 5131760	(8.27, N/A) (N/A, 0.03, N/A)	1312.8	N/A	2.2351 [2.0000]	111.8% { 120.1% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 682542	(5.15, N/A) (N/A, 0.00, N/A)	1100.1	N/A	4.4203 [4.0000]	110.5% { 131.6% }			
13C2_6:2FTS_EIS	(429.0 / 81.0) 741788	(6.65, N/A) (N/A, 0.03, N/A)	1147.7	N/A	4.2609 [4.0000]	106.5% { 125.6% }			
13C2_8:2FTS_EIS	(529.0 / 81.0) 765880	(7.81, N/A) (N/A, 0.02, N/A)	1009.0	N/A	4.1601 [4.0000]	104.0% { 115.3% }			
13C8_PFOsa_EIS	(506.0 / 78.0) 9868316	(9.84, N/A) (N/A, 0.01, N/A)	2159.0	N/A	2.3333 [2.0000]	116.7% { 126.7% }			
D3_NMeFOsa_EIS	(515.0 / 169.0) 1970012	(10.43, N/A) (N/A, 0.01, N/A)	2544.4	N/A	2.1692 [2.0000]	108.5% { 114.6% }			
D5_NEtFOsa_EIS	(531.0 / 169.0) 1871997	(10.58, N/A) (N/A, 0.01, N/A)	3281.6	N/A	2.3541 [2.0000]	117.7% { 119.7% }			

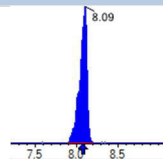
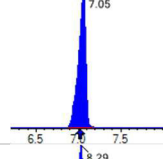
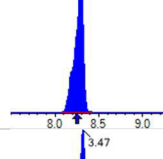
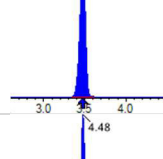
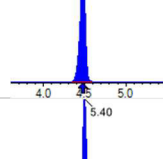
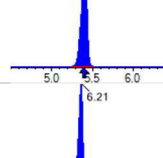
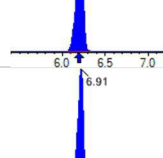
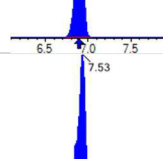
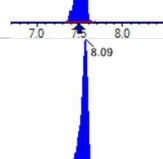
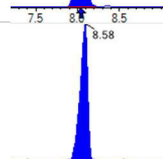
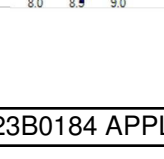
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1373660	(8.22 , N/A) (N/A , 0.02 , N/A)	1210.5	N/A	4.2630 [4.0000]	106.6% { 106.8% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 1277049	(8.47 , N/A) (N/A , 0.02 , N/A)	527735.4	N/A	4.6540 [4.0000]	116.4% { 132.0% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 7198350	(10.37 , N/A) (N/A , 0.01 , N/A)	1847.0	N/A	23.2817 [20.0000]	116.4% { 119.0% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 8543004	(10.53 , N/A) (N/A , 0.01 , N/A)	1631.4	N/A	24.1552 [20.0000]	120.8% { 124.7% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 4580346	(5.70 , N/A) (N/A , 0.01 , N/A)	2096.4	N/A	8.5902 [8.0000]	107.4% { 122.5% }			

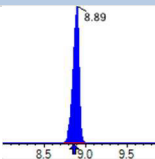
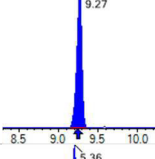
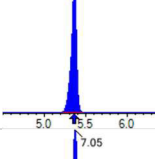
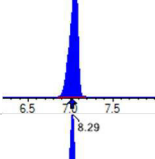
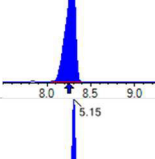
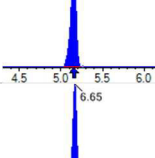
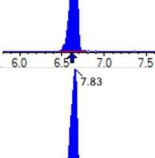
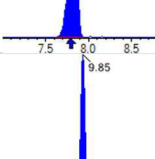
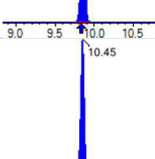
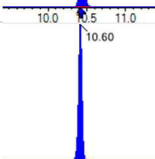
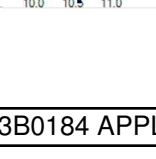
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) 1558824	(3.48 , 1.00) (0.00 , N/A , 0.0)	161.2	N/A 0.0 0.0	3.8585 [4.0000]	96.5%			
PFPeA	(263.0 / 219.0) 1265763 (263.0 / 69.0) 19161	(4.48 , 1.00) (0.00 , N/A , -0.3)	1452.9 127.2	0.0151 127.0 127.0	1.8650 [2.0000]	93.3%			
PFHxA	(313.0 / 269.0) 797424 (313.0 / 119.0) 86616	(5.40 , 1.00) (0.00 , N/A , -0.1)	2524.9 90779.8	0.1086 116.1 116.1	0.9514 [1.0000]	95.1%			
PFHpA	(363.0 / 319.0) 683494 (363.0 / 169.0) 206951	(6.21 , 1.00) (0.00 , N/A , 0.1)	4310.6 2135130.2	0.3028 95.9 95.9	0.9372 [1.0000]	93.7%			
PFOA	(413.0 / 369.0) 907671 (413.0 / 169.0) 315435	(6.92 , 1.00) (0.01 , N/A , 0.5)	1707.8 1430.0	0.3475 102.7 102.7	0.9602 [1.0000]	96.0%			
PFNA	(463.0 / 419.0) 707835 (463.0 / 169.0) 156984	(7.53 , 1.00) (0.00 , N/A , -0.3)	983.4 623.2	0.2218 94.7 94.7	0.8847 [1.0000]	88.5%			
PFDA	(513.0 / 469.0) 835540 (513.0 / 169.0) 108944	(8.09 , 1.00) (0.00 , N/A , -0.3)	550.1 61.4	0.1304 105.2 105.2	0.8794 [1.0000]	87.9%			
PFUnA	(563.0 / 519.0) 723977 (563.0 / 169.0) 90223	(8.59 , 1.00) (0.01 , N/A , 0.9)	609.2 421.9	0.1246 115.8 115.8	0.8854 [1.0000]	88.5%			
PFDoA	(613.0 / 569.0) 709027 (613.0 / 169.0) 103601	(8.89 , 1.00) (0.00 , N/A , -0.1)	956.4 376.4	0.1461 92.0 92.0	0.9490 [1.0000]	94.9%			
PFTTrDA	(663.0 / 619.0) 664752 (663.0 / 169.0) 170193	(9.10 , 1.02) (N/A , 0.03 , 0.1)	1282.6 488.6	0.2560 94.3 94.3	0.9246 [1.0000]	92.5%			
PFTeDA	(713.0 / 669.0) 776147 (713.0 / 169.0) 167772	(9.27 , 1.00) (0.00 , N/A , 0.1)	813.6 506.4	0.2162 98.9 98.9	0.9069 [1.0000]	90.7%			

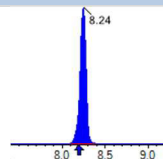
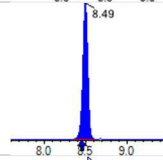
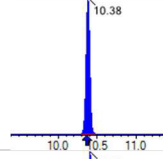
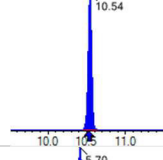
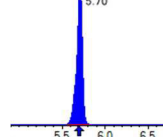
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) 1329800 (299.0 / 99.0) 875047	(5.36 , 1.00) (0.00 , N/A , -0.1)	1248.8 996.2	0.6580 104.6 104.6	0.7969 [0.8847]	90.1%			
PFPeS	(349.0 / 80.0) 2524286 (349.0 / 99.0) 821758	(6.28 , 0.89) (N/A , 0.03 , 0.1)	3988.1 18015.6	0.3255 97.3 97.3	0.9592 [0.9384]	102.2%			
PFHxS	(399.0 / 80.0) 1890138 (399.0 / 99.0) 597730	(7.05 , 1.00) (0.00 , N/A , 0.2)	120033.2 862.6	0.3162 98.8 98.8	0.8186 [0.9110]	89.9%			
PFHpS	(449.0 / 80.0) 2149516 (449.0 / 99.0) 597105	(7.71 , 0.93) (N/A , 0.05 , 0.2)	3207.3 22273.5	0.2778 102.3 102.3	0.9164 [0.9514]	96.3%			
PFOS	(499.0 / 80.0) 2507487 (499.0 / 99.0) 480040	(8.28 , 1.00) (0.00 , N/A , 0.1)	207.8 247.5	0.1914 87.0 87.0	0.8722 [0.9275]	94.0%			
PFNS	(549.0 / 80.0) 2661423 (549.0 / 99.0) 643064	(8.73 , 1.05) (N/A , 0.03 , -0.2)	852381.6 2828.0	0.2416 100.4 100.4	0.9251 [0.9599]	96.4%			
PFDS	(599.0 / 80.0) 2782413 (599.0 / 99.0) 634734	(8.99 , 1.08) (N/A , 0.02 , -0.1)	1562.1 908.1	0.2281 112.7 112.7	0.8703 [0.9631]	90.4%			
PFDoS	(699.0 / 80.0) 1327304 (699.0 / 99.0) 291014	(9.34 , 1.13) (N/A , 0.02 , 0.1)	1709.6 562.8	0.2193 112.2 112.2	0.8993 [0.9696]	92.8%			
4:2FTS	(327.0 / 307.0) 1697206 (327.0 / 81.0) 1180095	(5.15 , 1.00) (0.00 , N/A , -0.1)	1651.6 1084.2	0.6953 100.3 100.3	3.5442 [3.7381]	94.8%			
6:2FTS	(427.0 / 407.0) 842761 (427.0 / 81.0) 765235	(6.65 , 1.00) (0.00 , N/A , -0.1)	1665.8 851.8	0.9080 107.2 107.2	3.5474 [3.7962]	93.4%			
8:2FTS	(527.0 / 507.0) 778543 (527.0 / 81.0) 696144	(7.84 , 1.00) (0.00 , N/A , -0.2)	886.8 716.2	0.8942 109.6 109.6	3.8731 [3.8332]	101.0%			

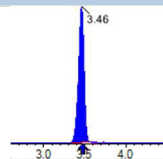
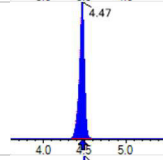
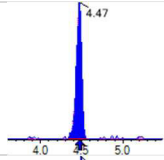
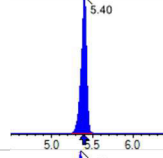
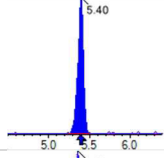
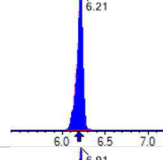
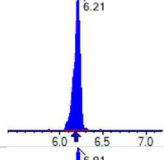
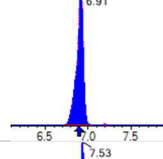
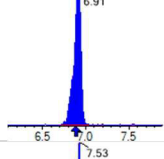
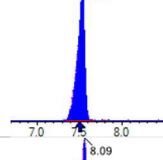
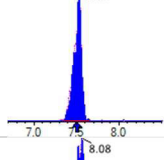
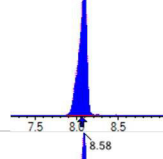
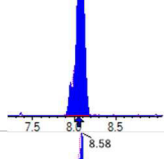
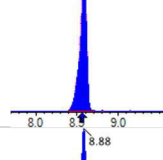
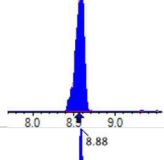
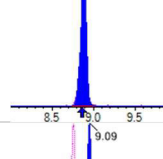
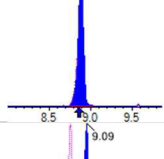
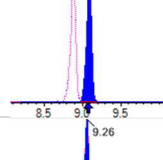
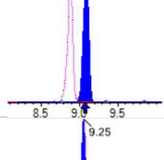
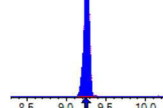
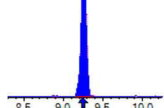
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) 3798776 (498.0 / 478.0) 69856	(9.85 , 1.00) (0.00 , N/A , -0.1)	2092.8 616.5	0.0184 93.6 93.6	0.9625 [1.0000]	96.2%			
NMeFOSA	(512.0 / 219.0) 3475090 (512.0 / 169.0) 2925163	(10.44 , 1.00) (0.00 , N/A , 0.9)	4208.6 3191.4	0.8418 101.9 101.9	3.9505 [4.0000]	98.8%			
NEiFOSA	(526.0 / 219.0) 3722150 (526.0 / 169.0) 4891746	(10.59 , 1.00) (-0.01 , N/A , 0.6)	6839.5 5197.2	1.3142 103.8 103.8	3.9325 [4.0000]	98.3%			
NMeFOSAA	(570.0 / 419.0) 288907 (570.0 / 483.0) 128440	(8.25 , 1.00) (0.01 , N/A , 0.3)	752.7 191.0	0.4446 90.7 90.7	0.9383 [1.0000]	93.8%			
NEiFOSAA	(584.0 / 419.0) 256591 (584.0 / 526.0) 136627	(8.50 , 1.00) (0.01 , N/A , -0.2)	4407.7 18005.9	0.5325 94.2 94.2	0.9990 [1.0000]	99.9%			
NMeFOSE	(616.0 / 59.0) 1414029	(10.39 , 1.00) (0.01 , N/A , 0.0)	1714.0	N/A 0.0 0.0	3.7753 [4.0000]	94.4%			
NEiFOSE	(630.0 / 59.0) 1663007	(10.55 , 1.00) (0.01 , N/A , 0.0)	926.7	N/A 0.0 0.0	3.7921 [4.0000]	94.8%			
HFPO-DA	(285.0 / 169.0) 907050 (285.0 / 185.0) 2341753	(5.71 , 1.00) (0.00 , N/A , 0.0)	1322.8 2111.7	2.5817 93.9 93.9	2.0087 [2.0000]	100.4%			
ADONA	(377.0 / 85.0) 3238705 (377.0 / 251.0) 299576	(6.49 , 1.14) (N/A , 0.03 , 0.0)	1602.5 745.8	0.0925 105.7 105.7	1.9325 [1.8854]	102.5%			
9CI-Pf3ONS	(531.0 / 351.0) 7337674 (533.0 / 353.0) 2229765	(8.65 , 1.52) (N/A , 0.04 , 0.1)	1610.2 1545.7	0.3039 103.2 103.2	1.8668 [1.8665]	100.0%			
11CI-PF3OUDS	(631.0 / 451.0) 3561286 (633.0 / 453.0) 1094280	(9.14 , 1.60) (N/A , 0.03 , 0.0)	1394.4 1681.1	0.3073 99.6 99.6	1.8429 [1.8864]	97.7%			

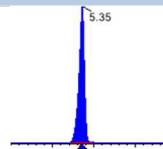
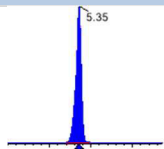
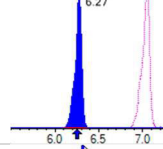
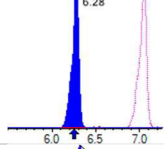
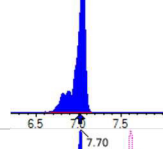
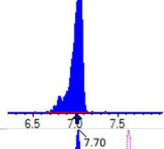
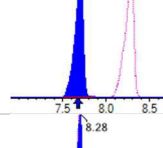
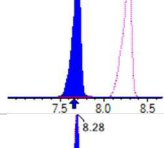
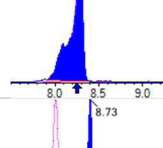
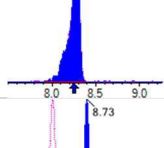
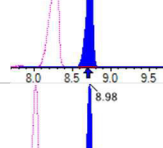
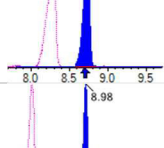
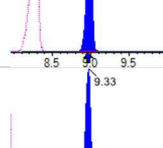
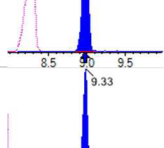
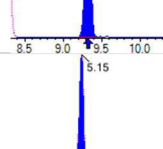
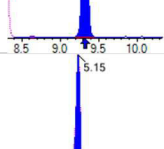
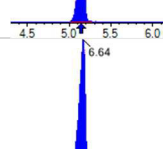
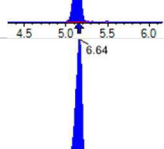
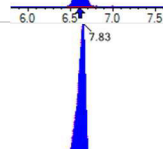
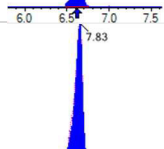
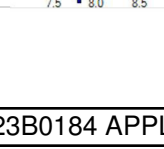
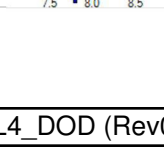
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) 74731 (241.0 / 117.0) 116652	(4.02, 0.90) (N/A, 0.01, 0.1)	589.2 482.5	1.5610 90.0 90.0	3.7286 [4.0000]	93.2%			
5:3FTCA	(341.0 / 236.7) 434752 (341.0 / 217.0) 717769	(5.91, 1.09) (N/A, 0.01, 0.0)	1085.3 866.2	1.6510 98.2 98.2	3.6285 [4.0000]	90.7%			
7:3FTCA	(441.0 / 317.0) 734732 (441.0 / 337.0) 609932	(7.82, 1.45) (N/A, 0.03, -0.6)	428.1 400.8	0.8301 96.7 96.7	3.4110 [4.0000]	85.3%			
PFEESA	(315.0 / 135.0) 1732259 (315.0 / 83.0) 518677	(5.79, 1.07) (N/A, 0.01, 0.0)	2155.7 702.4	0.2994 106.4 106.4	1.6632 [1.7849]	93.2%			
PFMPA	(229.0 / 85.0) 322638	(3.86, 0.86) (N/A, 0.01, 0.0)	2102.3	N/A 0.0 0.0	1.8593 [2.0000]	93.0%			
PFMBA	(279.0 / 85.0) 1037277	(4.79, 1.07) (N/A, 0.01, 0.0)	1702.1	N/A 0.0 0.0	1.7448 [2.0000]	87.2%			
NFDHA	(295.0 / 201.0) 898951 (295.0 / 85.0) 950784	(5.31, 0.98) (N/A, 0.01, -0.1)	1585.9 1690.4	1.0577 101.3 101.3	1.9283 [2.0000]	96.4%			
13C3_PFBa_IIS	(216.0 / 172.0) 382087	(3.48, N/A) (N/A, 0.01, N/A)	1420.6	N/A	1.1270 [1.0000]	112.7% { 115.9% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 625588	(5.41, N/A) (N/A, 0.01, N/A)	1047.7	N/A	1.0518 [1.0000]	105.2% { 112.6% }			
13C4_PFOA_IIS	(417.0 / 372.0) 841224	(6.92, N/A) (N/A, 0.03, N/A)	759.7	N/A	1.0342 [1.0000]	103.4% { 112.1% }			
13C5_PFNA_IIS	(468.0 / 423.0) 822059	(7.53, N/A) (N/A, 0.04, N/A)	1047.7	N/A	1.0765 [1.0000]	107.7% { 108.0% }			

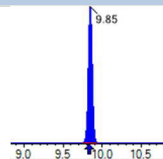
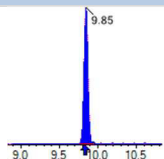
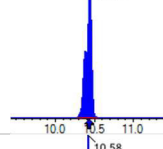
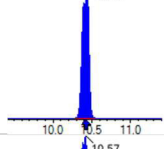
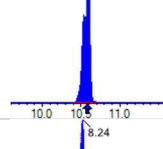
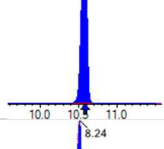
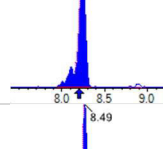
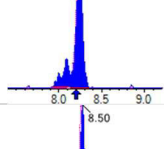
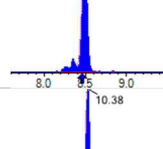
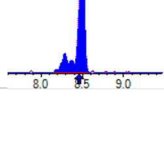
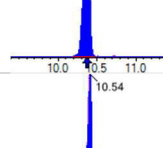
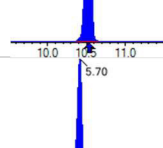
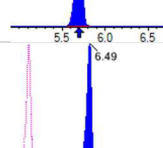
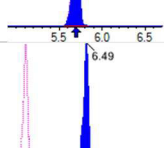
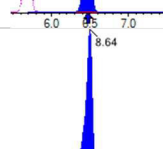
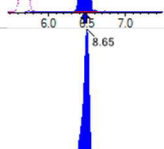
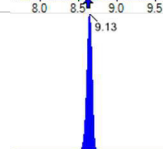
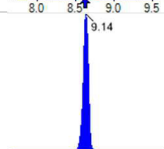
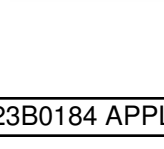
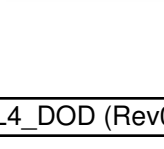
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 720343	(8.09 , N/A) (N/A , 0.04 , N/A)	779.3	N/A	1.0034 [1.0000]	100.3% { 104.7% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 1239843	(7.05 , N/A) (N/A , 0.04 , N/A)	1034.1	N/A	1.0067 [1.0000]	100.7% { 103.2% }			
13C4_PFOS_IIS	(503.0 / 79.9) 1851148	(8.29 , N/A) (N/A , 0.05 , N/A)	1210.5	N/A	1.0430 [1.0000]	104.3% { 106.9% }			
13C4_PFBA_EIS	(217.0 / 172.0) 3821781	(3.47 , N/A) (N/A , 0.01 , N/A)	3332.5	N/A	8.4118 [8.0000]	105.1% { 130.5% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 3080953	(4.48 , N/A) (N/A , 0.01 , N/A)	2005.0	N/A	4.4199 [4.0000]	110.5% { 126.4% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 1796236	(5.40 , N/A) (N/A , 0.01 , N/A)	2427.6	N/A	2.1132 [2.0000]	105.7% { 117.0% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 1633410	(6.21 , N/A) (N/A , 0.03 , N/A)	1499.8	N/A	2.0604 [2.0000]	103.0% { 115.6% }			
13C8_PFOA_EIS	(421.0 / 376.0) 1962058	(6.91 , N/A) (N/A , 0.03 , N/A)	1527.7	N/A	2.1791 [2.0000]	109.0% { 117.7% }			
13C9_PFNA_EIS	(472.0 / 427.0) 846974	(7.53 , N/A) (N/A , 0.04 , N/A)	1129.1	N/A	1.0339 [1.0000]	103.4% { 119.5% }			
13C6_PFDA_EIS	(519.0 / 474.0) 958247	(8.09 , N/A) (N/A , 0.05 , N/A)	762.0	N/A	1.1493 [1.0000]	114.9% { 130.5% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 929895	(8.58 , N/A) (N/A , 0.04 , N/A)	1111.9	N/A	1.1258 [1.0000]	112.6% { 123.5% }			

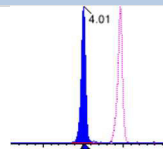
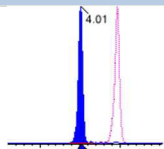
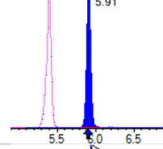
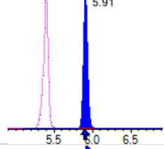
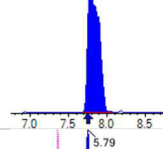
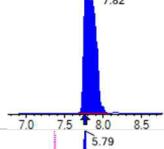
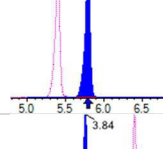
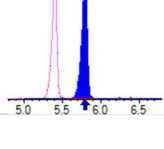
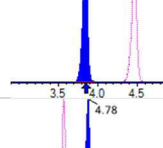
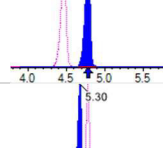
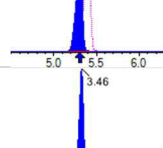
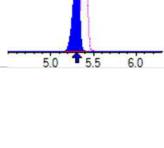
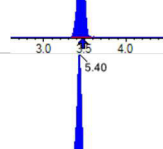
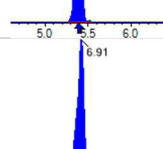
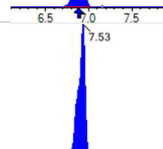
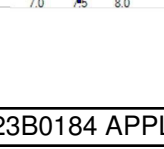
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 856390	(8.89 , N/A) (N/A , 0.03 , N/A)	865.0	N/A	1.1179 [1.0000]	111.8% { 116.2% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 930714	(9.27 , N/A) (N/A , 0.03 , N/A)	1015.0	N/A	1.1687 [1.0000]	116.9% { 129.4% }			
13C3_PFBs_EIS	(302.0 / 80.0) 5869688	(5.36 , N/A) (N/A , 0.01 , N/A)	2633.8	N/A	2.3428 [2.0000]	117.1% { 129.0% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 2931644	(7.05 , N/A) (N/A , 0.03 , N/A)	1945.4	N/A	2.2397 [2.0000]	112.0% { 123.8% }			
13C8_PFOS_EIS	(507.0 / 80.0) 5210104	(8.29 , N/A) (N/A , 0.05 , N/A)	976.1	N/A	2.1424 [2.0000]	107.1% { 121.9% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 678647	(5.15 , N/A) (N/A , 0.01 , N/A)	1162.9	N/A	4.4224 [4.0000]	110.6% { 130.9% }			
13C2_6:2FTS_EIS	(429.0 / 81.0) 728417	(6.65 , N/A) (N/A , 0.03 , N/A)	682.3	N/A	4.2101 [4.0000]	105.3% { 123.3% }			
13C2_8:2FTS_EIS	(529.0 / 81.0) 716231	(7.83 , N/A) (N/A , 0.04 , N/A)	958.0	N/A	3.9146 [4.0000]	97.9% { 107.9% }			
13C8_PFOsa_EIS	(506.0 / 78.0) 10158077	(9.85 , N/A) (N/A , 0.02 , N/A)	2799.0	N/A	2.2675 [2.0000]	113.4% { 130.4% }			
D3_NMeFOsa_EIS	(515.0 / 169.0) 2048481	(10.45 , N/A) (N/A , 0.02 , N/A)	2394.4	N/A	2.1295 [2.0000]	106.5% { 119.1% }			
D5_NEtFOsa_EIS	(531.0 / 169.0) 1948623	(10.60 , N/A) (N/A , 0.02 , N/A)	3224.4	N/A	2.3135 [2.0000]	115.7% { 124.6% }			

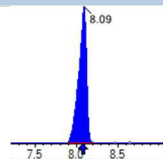
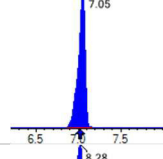
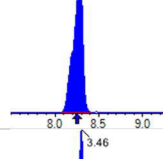
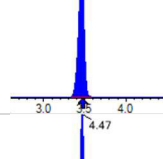
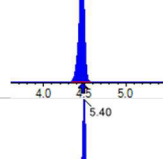
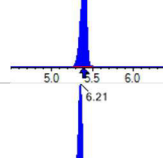
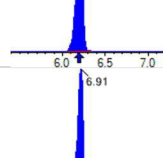
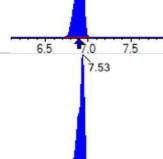
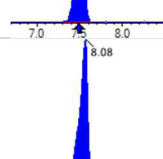
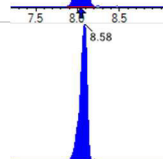
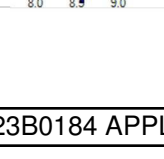
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1441118	(8.24 , N/A) (N/A , 0.05 , N/A)	1589.4	N/A	4.2223 [4.0000]	105.6% { 112.0% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 1178562	(8.49 , N/A) (N/A , 0.04 , N/A)	2950.5	N/A	4.0550 [4.0000]	101.4% { 121.8% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 7149088	(10.38 , N/A) (N/A , 0.02 , N/A)	2911.4	N/A	21.8297 [20.0000]	109.1% { 118.1% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 8680083	(10.54 , N/A) (N/A , 0.02 , N/A)	1249.1	N/A	23.1707 [20.0000]	115.9% { 126.7% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 4406473	(5.70 , N/A) (N/A , 0.02 , N/A)	2180.8	N/A	8.2501 [8.0000]	103.1% { 117.9% }			

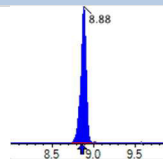
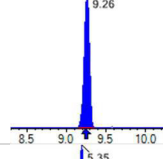
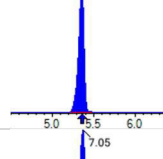
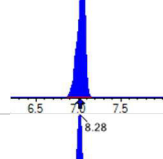
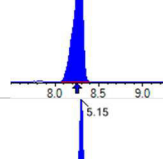
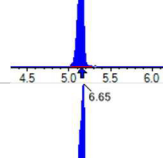
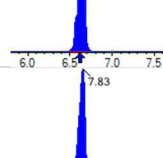
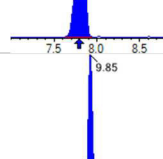
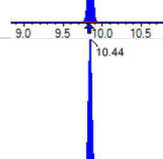
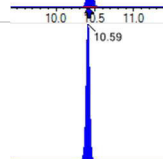
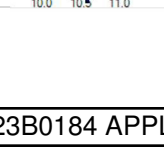
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) 2978159	(3.46 , 1.00) (0.00 , N/A , 0.0)	175.0	N/A 0.0 0.0	7.8088 [8.0000]	97.6%			
PFPeA	(263.0 / 219.0) 2437014 (263.0 / 69.0) 31666	(4.47 , 1.00) (0.00 , N/A , 0.0)	2356.6 276.8	0.0130 109.0 109.0	3.7735 [4.0000]	94.3%			
PFHxA	(313.0 / 269.0) 1573614 (313.0 / 119.0) 174994	(5.40 , 1.00) (0.00 , N/A , 0.2)	3255.3 49463.5	0.1112 118.8 118.8	1.9604 [2.0000]	98.0%			
PFHpA	(363.0 / 319.0) 1362121 (363.0 / 169.0) 445296	(6.21 , 1.00) (0.00 , N/A , 0.0)	2703.6 3563.0	0.3269 103.6 103.6	1.9223 [2.0000]	96.1%			
PFOA	(413.0 / 369.0) 1733709 (413.0 / 169.0) 569454	(6.91 , 1.00) (0.00 , N/A , 0.1)	3176.7 21911.3	0.3285 97.1 97.1	1.9559 [2.0000]	97.8%			
PFNA	(463.0 / 419.0) 1365813 (463.0 / 169.0) 344762	(7.53 , 1.00) (0.00 , N/A , -0.1)	15852.2 151029.7	0.2524 107.8 107.8	1.9786 [2.0000]	98.9%			
PFDA	(513.0 / 469.0) 1646083 (513.0 / 169.0) 189871	(8.09 , 1.00) (0.00 , N/A , 0.5)	981.7 659.9	0.1153 93.1 93.1	2.0731 [2.0000]	103.7%			
PFUnA	(563.0 / 519.0) 1447666 (563.0 / 169.0) 185027	(8.58 , 1.00) (0.00 , N/A , 0.2)	971.6 1467.0	0.1278 118.8 118.8	1.7969 [2.0000]	89.8%			
PFDoA	(613.0 / 569.0) 1433036 (613.0 / 169.0) 265342	(8.88 , 1.00) (0.00 , N/A , -0.1)	1619.1 1187.3	0.1852 116.5 116.5	2.1009 [2.0000]	105.0%			
PFTTrDA	(663.0 / 619.0) 1364367 (663.0 / 169.0) 377483	(9.09 , 1.02) (N/A , 0.02 , 0.0)	1093.2 972.3	0.2767 101.9 101.9	2.0787 [2.0000]	103.9%			
PFTeDA	(713.0 / 669.0) 1489305 (713.0 / 169.0) 313061	(9.26 , 1.00) (0.00 , N/A , 0.3)	1109.4 999.2	0.2102 96.1 96.1	1.8286 [2.0000]	91.4%			

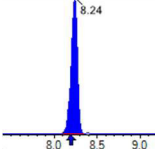
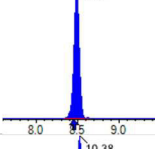
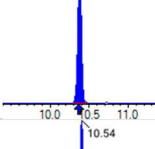
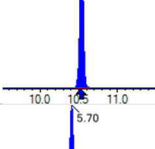
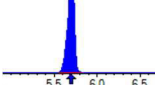
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) 2793737 (299.0 / 99.0) 1698297	(5.35 , 1.00) (0.00 , N/A , 0.0)	1854.8 1652.6	0.6079 96.6 96.6	1.7650 [1.7695]	99.7%			
PFPeS	(349.0 / 80.0) 4739230 (349.0 / 99.0) 1555828	(6.27 , 0.89) (N/A , 0.02 , 0.0)	52180.7 8853757.1	0.3283 98.1 98.1	1.8521 [1.8768]	98.7%			
PFHxS	(399.0 / 80.0) 3714721 (399.0 / 99.0) 1275949	(7.05 , 1.00) (0.00 , N/A , 0.3)	4966.7 857.5	0.3435 107.3 107.3	1.6546 [1.8220]	90.8%			
PFHpS	(449.0 / 80.0) 4164441 (449.0 / 99.0) 1203176	(7.70 , 0.93) (N/A , 0.04 , 0.0)	1250763.1 2433.7	0.2889 106.4 106.4	1.9076 [1.9028]	100.2%			
PFOS	(499.0 / 80.0) 4805372 (499.0 / 99.0) 1022399	(8.28 , 1.00) (0.00 , N/A , 0.3)	288.4 428.6	0.2128 96.7 96.7	1.7960 [1.8550]	96.8%			
PFNS	(549.0 / 80.0) 5359766 (549.0 / 99.0) 1220427	(8.73 , 1.05) (N/A , 0.03 , 0.2)	1071105.7 102953.8	0.2277 94.6 94.6	2.0018 [1.9198]	104.3%			
PFDS	(599.0 / 80.0) 6045855 (599.0 / 99.0) 1237926	(8.98 , 1.08) (N/A , 0.02 , 0.1)	1547.9 1053.5	0.2048 101.2 101.2	2.0318 [1.9262]	105.5%			
PFDoS	(699.0 / 80.0) 2614022 (699.0 / 99.0) 499622	(9.33 , 1.13) (N/A , 0.01 , -0.1)	944.0 873.7	0.1911 97.8 97.8	1.9030 [1.9391]	98.1%			
4:2FTS	(327.0 / 307.0) 3532565 (327.0 / 81.0) 2357912	(5.15 , 1.00) (0.00 , N/A , 0.0)	2192.5 1237.0	0.6675 96.3 96.3	7.7615 [7.4762]	103.8%			
6:2FTS	(427.0 / 407.0) 1902823 (427.0 / 81.0) 1462000	(6.64 , 1.00) (0.00 , N/A , 0.1)	1490.4 1254.7	0.7683 90.7 90.7	7.4774 [7.5923]	98.5%			
8:2FTS	(527.0 / 507.0) 1550550 (527.0 / 81.0) 1290969	(7.83 , 1.00) (0.00 , N/A , 0.1)	940.8 1574.4	0.8326 102.1 102.1	7.5141 [7.6663]	98.0%			

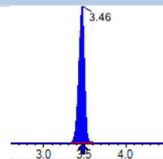
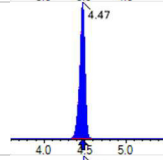
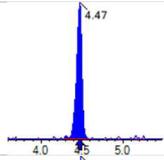
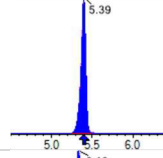
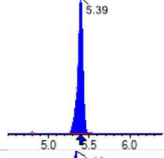
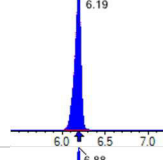
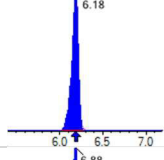
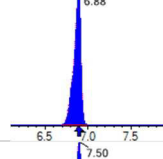
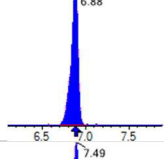
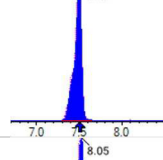
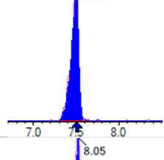
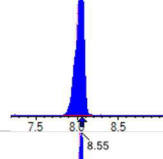
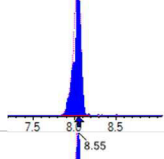
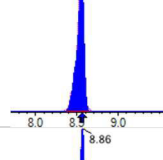
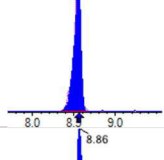
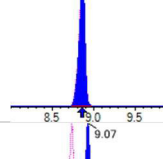
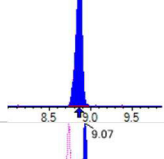
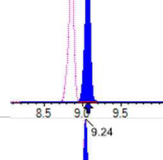
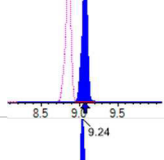
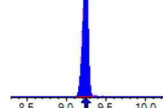
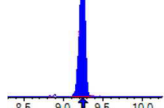
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) 7841826 (498.0 / 478.0) 139402	(9.85 , 1.00) (0.00 , N/A , 0.0)	3025.3 759.9	0.0178 90.5 90.5	2.0951 [2.0000]	104.8%			
NMeFOSA	(512.0 / 219.0) 7029796 (512.0 / 169.0) 5792823	(10.44 , 1.00) (0.00 , N/A , 1.1)	4347.7 7352.5	0.8240 99.7 99.7	8.1995 [8.0000]	102.5%			
NEIFOSA	(526.0 / 219.0) 7386563 (526.0 / 169.0) 9559463	(10.58 , 1.00) (-0.01 , N/A , 0.6)	12814.8 6845.3	1.2942 102.2 102.2	8.2050 [8.0000]	102.6%			
NMeFOSAA	(570.0 / 419.0) 609780 (570.0 / 483.0) 301348	(8.24 , 1.00) (0.00 , N/A , 0.0)	1023.7 225.8	0.4942 100.8 100.8	2.0743 [2.0000]	103.7%			
NEIFOSAA	(584.0 / 419.0) 476378 (584.0 / 526.0) 296111	(8.49 , 1.00) (0.01 , N/A , -0.3)	3396.7 926.5	0.6216 110.0 110.0	1.8606 [2.0000]	93.0%			
NMeFOSE	(616.0 / 59.0) 2838778	(10.38 , 1.00) (0.01 , N/A , 0.0)	2391.1	N/A 0.0 0.0	7.7446 [8.0000]	96.8%			
NEtFOSE	(630.0 / 59.0) 3278313	(10.54 , 1.00) (0.01 , N/A , 0.0)	1102.3	N/A 0.0 0.0	7.9731 [8.0000]	99.7%			
HFPO-DA	(285.0 / 169.0) 1639480 (285.0 / 185.0) 4591397	(5.70 , 1.00) (0.00 , N/A , -0.2)	1532.1 2560.1	2.8005 101.9 101.9	3.6719 [4.0000]	91.8%			
ADONA	(377.0 / 85.0) 6564313 (377.0 / 251.0) 559733	(6.49 , 1.14) (N/A , 0.03 , 0.0)	2563.4 884.9	0.0853 97.4 97.4	3.9614 [3.7708]	105.1%			
9Cl-Pf3ONS	(531.0 / 351.0) 15459546 (533.0 / 353.0) 4606512	(8.64 , 1.52) (N/A , 0.03 , -0.1)	1602.6 1289.7	0.2980 101.2 101.2	3.9779 [3.7330]	106.6%			
11Cl-Pf3OUDS	(631.0 / 451.0) 7357657 (633.0 / 453.0) 2245051	(9.13 , 1.60) (N/A , 0.02 , -0.1)	1410.8 1419.1	0.3051 98.9 98.9	3.8507 [3.7728]	102.1%			

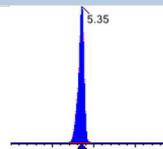
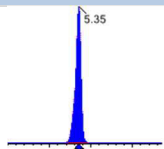
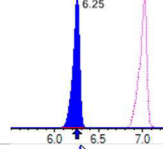
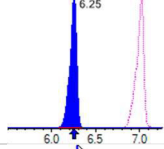
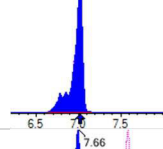
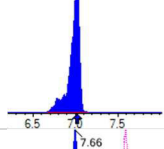
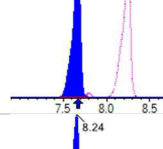
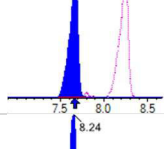
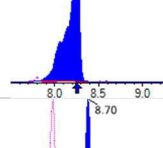
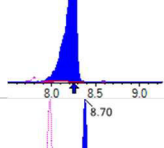
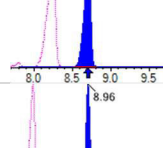
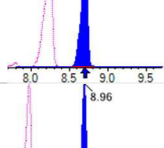
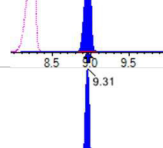
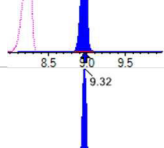
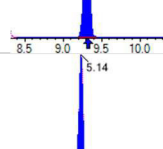
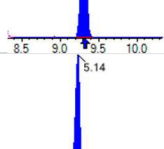
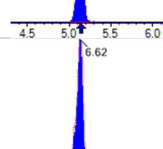
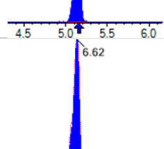
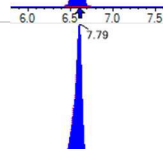
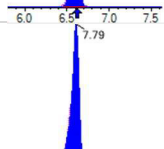
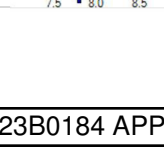
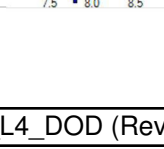
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) 146366 (241.0 / 117.0) 226482	(4.01 , 0.90) (N/A , -0.01 , -0.1)	764.7 683.5	1.5474 89.2 89.2	7.6742 [8.0000]	95.9%			
5:3FTCA	(341.0 / 236.7) 885987 (341.0 / 217.0) 1493057	(5.91 , 1.10) (N/A , 0.01 , 0.1)	1357.5 1658.6	1.6852 100.2 100.2	7.7213 [8.0000]	96.5%			
7:3FTCA	(441.0 / 317.0) 1487973 (441.0 / 337.0) 1357096	(7.82 , 1.45) (N/A , 0.03 , 0.1)	591.5 527.7	0.9120 106.2 106.2	7.2130 [8.0000]	90.2%			
PFEESA	(315.0 / 135.0) 3649655 (315.0 / 83.0) 989296	(5.79 , 1.07) (N/A , 0.01 , -0.1)	2483.2 1061.1	0.2711 96.3 96.3	3.6589 [3.5698]	102.5%			
PFMPA	(229.0 / 85.0) 605828	(3.84 , 0.86) (N/A , -0.01 , 0.0)	2138.3	N/A 0.0 0.0	3.6689 [4.0000]	91.7%			
PFMBA	(279.0 / 85.0) 2144549	(4.78 , 1.07) (N/A , 0.00 , 0.0)	1856.2	N/A 0.0 0.0	3.7910 [4.0000]	94.8%			
NFDHA	(295.0 / 201.0) 1692427 (295.0 / 85.0) 1830348	(5.30 , 0.98) (N/A , 0.01 , 0.0)	1825.5 1722.6	1.0815 103.6 103.6	3.7908 [4.0000]	94.8%			
13C3_PFBA_IIS	(216.0 / 172.0) 360206	(3.46 , N/A) (N/A , -0.01 , N/A)	1491.7	N/A	1.0625 [1.0000]	106.2% { 109.2% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 645560	(5.40 , N/A) (N/A , 0.00 , N/A)	1241.4	N/A	1.0854 [1.0000]	108.5% { 116.1% }			
13C4_PFOA_IIS	(417.0 / 372.0) 879294	(6.91 , N/A) (N/A , 0.03 , N/A)	1152.8	N/A	1.0810 [1.0000]	108.1% { 117.2% }			
13C5_PFNA_IIS	(468.0 / 423.0) 784728	(7.53 , N/A) (N/A , 0.04 , N/A)	596.7	N/A	1.0276 [1.0000]	102.8% { 103.1% }			

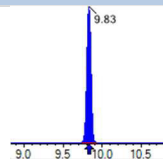
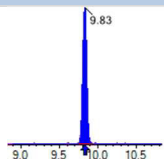
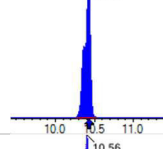
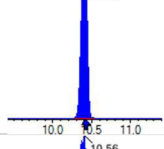
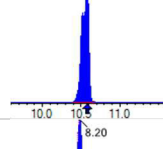
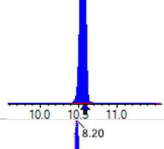
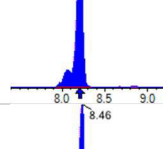
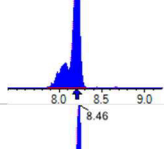
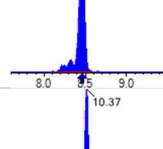
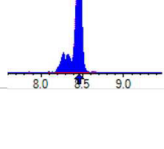
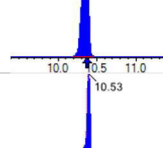
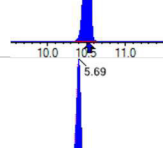
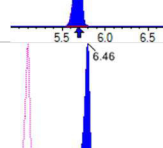
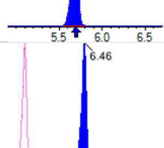
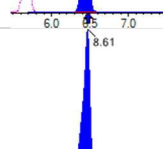
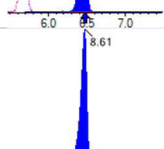
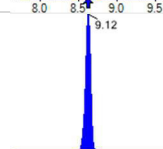
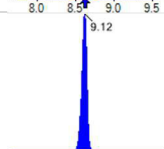
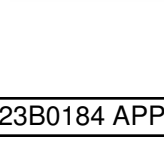
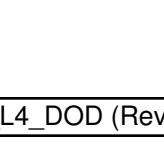
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 760726	(8.09 , N/A) (N/A , 0.03 , N/A)	938.9	N/A	1.0597 [1.0000]	106.0% { 110.5% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 1361278	(7.05 , N/A) (N/A , 0.03 , N/A)	1775.2	N/A	1.1053 [1.0000]	110.5% { 113.4% }			
13C4_PFOS_IIS	(503.0 / 79.9) 1884590	(8.28 , N/A) (N/A , 0.04 , N/A)	913.5	N/A	1.0618 [1.0000]	106.2% { 108.8% }			
13C4_PFBA_EIS	(217.0 / 172.0) 3607843	(3.46 , N/A) (N/A , 0.00 , N/A)	3262.4	N/A	8.4233 [8.0000]	105.3% { 123.2% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 2931788	(4.47 , N/A) (N/A , 0.00 , N/A)	1988.0	N/A	4.0758 [4.0000]	101.9% { 120.2% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 1720245	(5.40 , N/A) (N/A , 0.01 , N/A)	1781.8	N/A	1.9612 [2.0000]	98.1% { 112.1% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 1587008	(6.21 , N/A) (N/A , 0.02 , N/A)	1672.0	N/A	1.9400 [2.0000]	97.0% { 112.3% }			
13C8_PFOA_EIS	(421.0 / 376.0) 1839894	(6.91 , N/A) (N/A , 0.03 , N/A)	1579.8	N/A	1.9550 [2.0000]	97.7% { 110.4% }			
13C9_PFNA_EIS	(472.0 / 427.0) 730784	(7.53 , N/A) (N/A , 0.04 , N/A)	1059.9	N/A	0.9345 [1.0000]	93.5% { 103.1% }			
13C6_PFDA_EIS	(519.0 / 474.0) 800834	(8.08 , N/A) (N/A , 0.05 , N/A)	964.0	N/A	0.9095 [1.0000]	91.0% { 109.0% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 916176	(8.58 , N/A) (N/A , 0.04 , N/A)	1045.9	N/A	1.0503 [1.0000]	105.0% { 121.7% }			

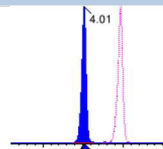
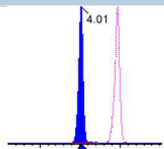
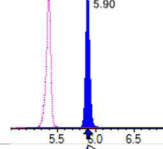
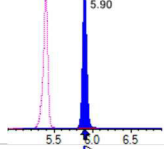
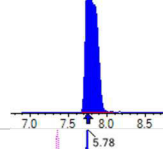
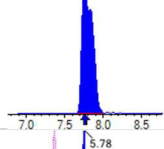
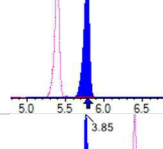
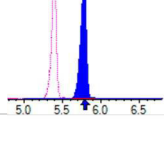
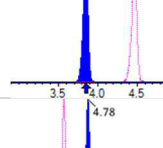
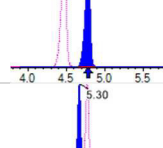
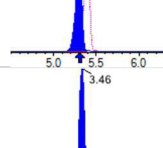
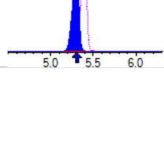
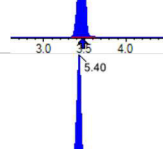
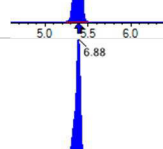
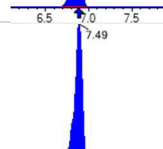
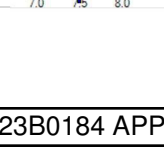
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 781819	(8.88 , N/A) (N/A , 0.02 , N/A)	961.0	N/A	0.9664 [1.0000]	96.6% { 106.1% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 885744	(9.26 , N/A) (N/A , 0.02 , N/A)	1144.9	N/A	1.0532 [1.0000]	105.3% { 123.2% }			
13C3_PFBs_EIS	(302.0 / 80.0) 5567594	(5.35 , N/A) (N/A , 0.00 , N/A)	2826.2	N/A	2.0240 [2.0000]	101.2% { 122.4% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 2850517	(7.05 , N/A) (N/A , 0.03 , N/A)	1193.5	N/A	1.9834 [2.0000]	99.2% { 120.4% }			
13C8_PFOS_EIS	(507.0 / 80.0) 4848965	(8.28 , N/A) (N/A , 0.04 , N/A)	1086.3	N/A	1.9585 [2.0000]	97.9% { 113.5% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 645015	(5.15 , N/A) (N/A , 0.00 , N/A)	1234.8	N/A	3.8283 [4.0000]	95.7% { 124.4% }			
13C2_6:2FTS_EIS	(429.0 / 81.0) 780256	(6.65 , N/A) (N/A , 0.03 , N/A)	1113.3	N/A	4.1074 [4.0000]	102.7% { 132.1% }			
13C2_8:2FTS_EIS	(529.0 / 81.0) 735263	(7.83 , N/A) (N/A , 0.04 , N/A)	856.5	N/A	3.6601 [4.0000]	91.5% { 110.7% }			
13C8_PFOsa_EIS	(506.0 / 78.0) 9632878	(9.85 , N/A) (N/A , 0.02 , N/A)	2860.5	N/A	2.1121 [2.0000]	105.6% { 123.6% }			
D3_NMeFOSA_EIS	(515.0 / 169.0) 1996541	(10.44 , N/A) (N/A , 0.02 , N/A)	2457.1	N/A	2.0387 [2.0000]	101.9% { 116.1% }			
D5_NEtFOSA_EIS	(531.0 / 169.0) 1853372	(10.59 , N/A) (N/A , 0.02 , N/A)	2454.9	N/A	2.1614 [2.0000]	108.1% { 118.5% }			

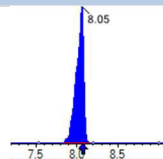
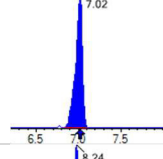
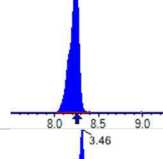
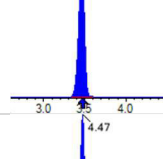
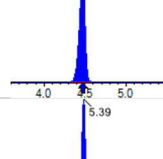
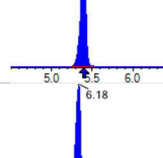
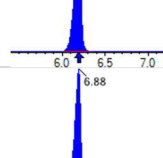
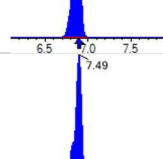
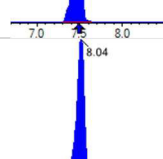
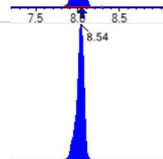
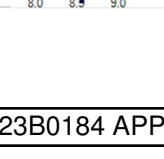
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1375836	(8.24 , N/A) (N/A , 0.04 , N/A)	1203.7	N/A	3.9595 [4.0000]	99.0% { 106.9% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 1174828	(8.49 , N/A) (N/A , 0.03 , N/A)	11028.6	N/A	3.9704 [4.0000]	99.3% { 121.4% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 6996507	(10.38 , N/A) (N/A , 0.02 , N/A)	1537.8	N/A	20.9847 [20.0000]	104.9% { 115.6% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 8138307	(10.54 , N/A) (N/A , 0.02 , N/A)	1520.9	N/A	21.3390 [20.0000]	106.7% { 118.8% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 4356913	(5.70 , N/A) (N/A , 0.01 , N/A)	1993.4	N/A	7.9049 [8.0000]	98.8% { 116.5% }			

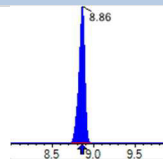
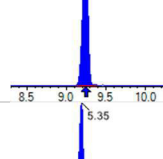
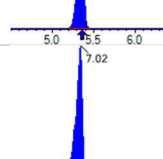
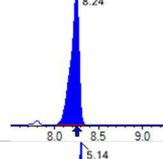
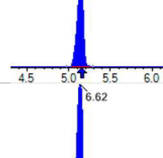
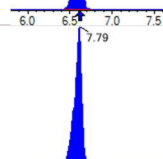
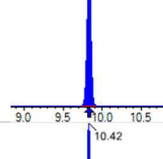
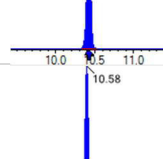
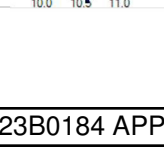
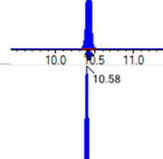
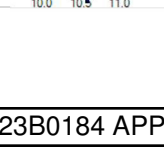
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) 6887982	(3.46 , 1.00) (0.00 , N/A , 0.0)	178.6	N/A 0.0 0.0	22.2419 [20.0000]	111.2%			
PFPeA	(263.0 / 219.0) 5837257 (263.0 / 69.0) 69565	(4.47 , 1.00) (0.00 , N/A , 0.1)	3551.0 286.4	0.0119 100.0 100.0	10.8683 [10.0000]	108.7%			
PFHxA	(313.0 / 269.0) 3859604 (313.0 / 119.0) 361138	(5.39 , 1.00) (0.00 , N/A , 0.0)	6341.4 2092.4	0.0936 100.0 100.0	5.3888 [5.0000]	107.8%			
PFHpA	(363.0 / 319.0) 3082886 (363.0 / 169.0) 973188	(6.19 , 1.00) (0.00 , N/A , 0.1)	4376.3 8634.7	0.3157 100.0 100.0	4.8857 [5.0000]	97.7%			
PFOA	(413.0 / 369.0) 4117111 (413.0 / 169.0) 1392975	(6.88 , 1.00) (0.00 , N/A , 0.3)	4757.1 8828.2	0.3383 100.0 100.0	5.1282 [5.0000]	102.6%			
PFNA	(463.0 / 419.0) 3488826 (463.0 / 169.0) 816729	(7.50 , 1.00) (0.00 , N/A , 0.2)	5185.8 1445.2	0.2341 100.0 100.0	5.2092 [5.0000]	104.2%			
PFDA	(513.0 / 469.0) 3981270 (513.0 / 169.0) 493389	(8.05 , 1.00) (0.01 , N/A , 0.0)	1319.6 442.8	0.1239 100.0 100.0	5.4676 [5.0000]	109.4%			
PFUnA	(563.0 / 519.0) 3556271 (563.0 / 169.0) 382747	(8.55 , 1.00) (0.01 , N/A , 0.0)	1457.7 903.2	0.1076 100.0 100.0	5.3732 [5.0000]	107.5%			
PFDoA	(613.0 / 569.0) 3274097 (613.0 / 169.0) 520170	(8.86 , 1.00) (0.00 , N/A , 0.0)	1539.4 998.3	0.1589 100.0 100.0	5.0927 [5.0000]	101.9%			
PFTTrDA	(663.0 / 619.0) 3198043 (663.0 / 169.0) 868288	(9.07 , 1.02) (N/A , 0.00 , -0.2)	2124.5 1303.1	0.2715 100.0 100.0	5.1695 [5.0000]	103.4%			
PFTeDA	(713.0 / 669.0) 3435322 (713.0 / 169.0) 751087	(9.24 , 1.00) (0.00 , N/A , 0.1)	1573.9 912.2	0.2186 100.0 100.0	5.1954 [5.0000]	103.9%			

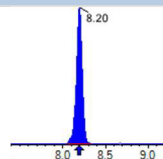
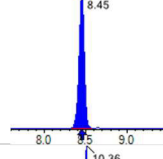
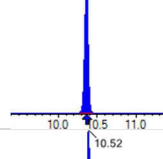
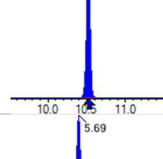
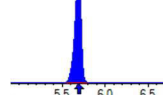
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) 6392257 (299.0 / 99.0) 4020950	(5.35 , 1.00) (0.00 , N/A , 0.0)	2259.7 2512.7	0.6290 100.0 100.0	4.9413 [4.4237]	111.7%			
PFPeS	(349.0 / 80.0) 11287040 (349.0 / 99.0) 3776028	(6.25 , 0.89) (N/A , 0.00 , 0.0)	1836879.2 278573.8	0.3345 100.0 100.0	5.3098 [4.6919]	113.2%			
PFHxS	(399.0 / 80.0) 9576642 (399.0 / 99.0) 3064686	(7.02 , 1.00) (0.00 , N/A , 0.2)	6710.1 2057.8	0.3200 100.0 100.0	5.1348 [4.5549]	112.7%			
PFHpS	(449.0 / 80.0) 10099121 (449.0 / 99.0) 2741856	(7.66 , 0.93) (N/A , 0.00 , 0.0)	27574.1 3023.4	0.2715 100.0 100.0	5.2502 [4.7570]	110.4%			
PFOS	(499.0 / 80.0) 11716008 (499.0 / 99.0) 2576734	(8.24 , 1.00) (0.00 , N/A , 0.1)	375.2 422.1	0.2199 100.0 100.0	4.9695 [4.6375]	107.2%			
PFNS	(549.0 / 80.0) 12692834 (549.0 / 99.0) 3054286	(8.70 , 1.06) (N/A , 0.00 , 0.0)	9516.6 33690.4	0.2406 100.0 100.0	5.3803 [4.7994]	112.1%			
PFDS	(599.0 / 80.0) 14131472 (599.0 / 99.0) 2860277	(8.96 , 1.09) (N/A , 0.00 , 0.1)	1888.5 1480.2	0.2024 100.0 100.0	5.3899 [4.8155]	111.9%			
PFDoS	(699.0 / 80.0) 7043385 (699.0 / 99.0) 1376106	(9.31 , 1.13) (N/A , 0.00 , 0.0)	1854.6 1342.4	0.1954 100.0 100.0	5.8193 [4.8478]	120.0%			
4:2FTS	(327.0 / 307.0) 7926427 (327.0 / 81.0) 5492761	(5.14 , 1.00) (0.00 , N/A , 0.0)	1931.3 1573.3	0.6930 100.0 100.0	21.6652 [18.6906]	115.9%			
6:2FTS	(427.0 / 407.0) 4266687 (427.0 / 81.0) 3615409	(6.62 , 1.00) (0.00 , N/A , 0.1)	1844.7 1832.6	0.8474 100.0 100.0	22.1428 [18.9808]	116.7%			
8:2FTS	(527.0 / 507.0) 3797855 (527.0 / 81.0) 3098407	(7.79 , 1.00) (0.00 , N/A , 0.1)	1632.5 1548.3	0.8158 100.0 100.0	20.3773 [19.1658]	106.3%			

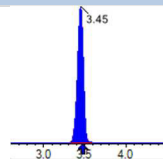
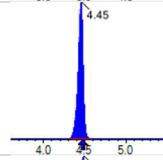
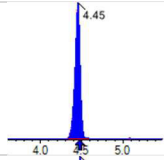
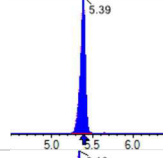
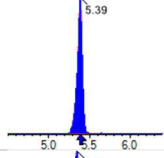
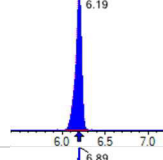
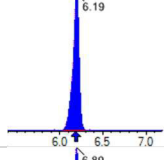
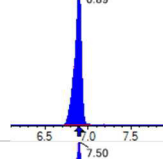
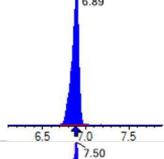
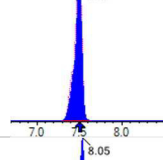
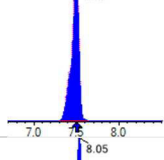
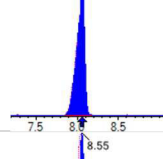
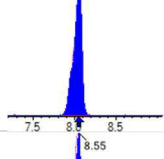
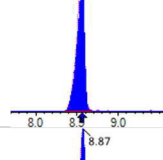
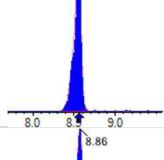
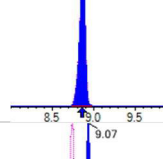
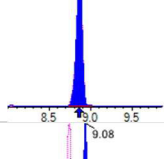
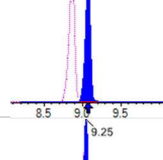
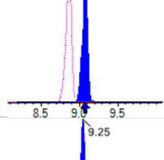
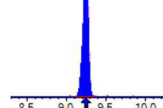
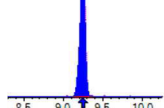
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) 17171365 (498.0 / 478.0) 337395	(9.83 , 1.00) (0.00 , N/A , 0.0)	2359.1 872.8	0.0196 100.0 100.0	5.6725 [5.0000]	113.5%			
NMeFOSA	(512.0 / 219.0) 16368076 (512.0 / 169.0) 13521783	(10.42 , 1.00) (0.00 , N/A , 1.1)	3723.2 3362.7	0.8261 100.0 100.0	22.1703 [20.0000]	110.9%			
NEIFOSA	(526.0 / 219.0) 17256788 (526.0 / 169.0) 21852773	(10.56 , 1.00) (-0.01 , N/A , 0.5)	15298.3 13721.8	1.2663 100.0 100.0	22.7188 [20.0000]	113.6%			
NMeFOSAA	(570.0 / 419.0) 1340653 (570.0 / 483.0) 657372	(8.20 , 1.00) (0.01 , N/A , -0.1)	1581.8 516.7	0.4903 100.0 100.0	4.8771 [5.0000]	97.5%			
NEIFOSAA	(584.0 / 419.0) 1136513 (584.0 / 526.0) 642114	(8.46 , 1.00) (0.01 , N/A , 0.0)	1923.2 1563.0	0.5650 100.0 100.0	5.3891 [5.0000]	107.8%			
NMeFOSE	(616.0 / 59.0) 6713767	(10.37 , 1.00) (0.01 , N/A , 0.0)	2748.1	N/A 0.0 0.0	21.1765 [20.0000]	105.9%			
NEIFOSE	(630.0 / 59.0) 7617213	(10.53 , 1.00) (0.01 , N/A , 0.0)	1231.6	N/A 0.0 0.0	22.0141 [20.0000]	110.1%			
HFPO-DA	(285.0 / 169.0) 4044639 (285.0 / 185.0) 11118904	(5.69 , 1.00) (0.00 , N/A , 0.1)	1829.2 2567.4	2.7490 100.0 100.0	10.5559 [10.0000]	105.6%			
ADONA	(377.0 / 85.0) 15597637 (377.0 / 251.0) 1365342	(6.46 , 1.14) (N/A , 0.00 , 0.1)	2420.5 1069.0	0.0875 100.0 100.0	10.9683 [9.4270]	116.4%			
9CI-PF3ONS	(531.0 / 351.0) 34989056 (533.0 / 353.0) 10306414	(8.61 , 1.51) (N/A , 0.00 , 0.0)	1896.9 1498.9	0.2946 100.0 100.0	10.4911 [9.3325]	112.4%			
11CI-PF3OUDS	(631.0 / 451.0) 17423179 (633.0 / 453.0) 5377206	(9.12 , 1.60) (N/A , 0.00 , -0.1)	2250.0 1533.2	0.3086 100.0 100.0	10.6258 [9.4321]	112.7%			

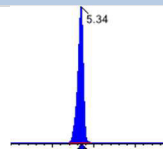
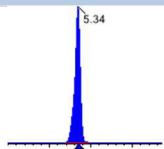
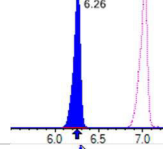
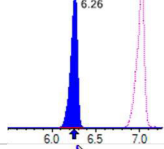
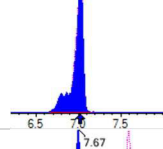
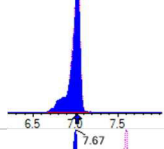
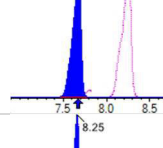
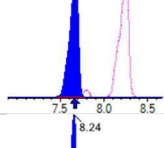
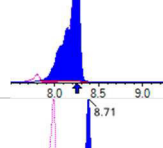
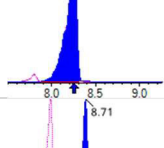
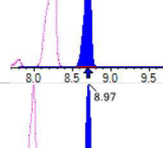
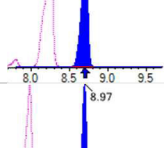
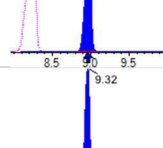
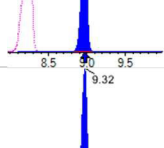
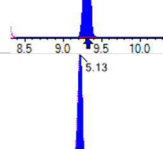
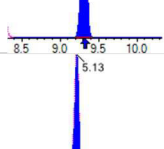
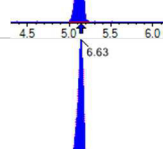
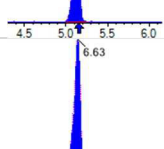
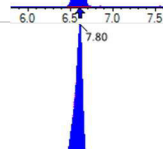
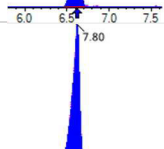
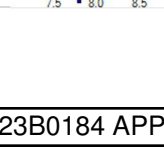
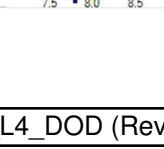
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) 304704 (241.0 / 117.0) 528426	(4.01 , 0.90) (N/A , 0.00 , 0.1)	1404.8 1425.4	1.7342 100.0 100.0	19.2104 [20.0000]	96.1%			
5:3FTCA	(341.0 / 236.7) 2127364 (341.0 / 217.0) 3577273	(5.90 , 1.09) (N/A , 0.00 , 0.0)	1973.1 1825.3	1.6816 100.0 100.0	20.7782 [20.0000]	103.9%			
7:3FTCA	(441.0 / 317.0) 3713939 (441.0 / 337.0) 3189506	(7.79 , 1.44) (N/A , 0.00 , 0.3)	919.4 703.9	0.8588 100.0 100.0	20.1770 [20.0000]	100.9%			
PFEESA	(315.0 / 135.0) 8678601 (315.0 / 83.0) 2443146	(5.78 , 1.07) (N/A , 0.00 , 0.1)	2157.1 1650.3	0.2815 100.0 100.0	9.7510 [8.9246]	109.3%			
PFMPA	(229.0 / 85.0) 1436993	(3.85 , 0.86) (N/A , 0.00 , 0.0)	2923.7	N/A 0.0 0.0	10.4644 [10.0000]	104.6%			
PFMBA	(279.0 / 85.0) 5400487	(4.78 , 1.07) (N/A , 0.00 , 0.0)	2597.0	N/A 0.0 0.0	11.4793 [10.0000]	114.8%			
NFDHA	(295.0 / 201.0) 4240196 (295.0 / 85.0) 4426346	(5.30 , 0.98) (N/A , 0.00 , 0.0)	1829.1 2739.3	1.0439 100.0 100.0	10.6441 [10.0000]	106.4%			
13C3_PFBa_IIS	(216.0 / 172.0) 329787	(3.46 , N/A) (N/A , 0.00 , N/A)	1620.4	N/A	0.9728 [1.0000]	97.3% { 100.0% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 555825	(5.40 , N/A) (N/A , 0.00 , N/A)	1867.9	N/A	0.9345 [1.0000]	93.5% { 100.0% }			
13C4_PFOA_IIS	(417.0 / 372.0) 750225	(6.88 , N/A) (N/A , 0.00 , N/A)	1252.2	N/A	0.9223 [1.0000]	92.2% { 100.0% }			
13C5_PFNAl_IIS	(468.0 / 423.0) 761426	(7.49 , N/A) (N/A , 0.00 , N/A)	990.9	N/A	0.9971 [1.0000]	99.7% { 100.0% }			

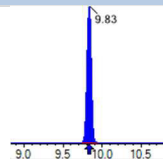
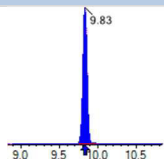
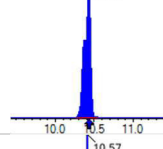
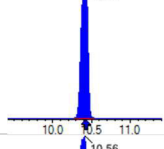
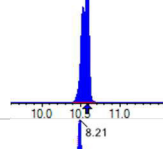
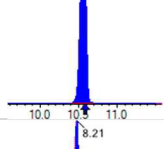
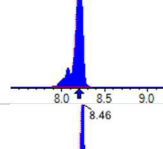
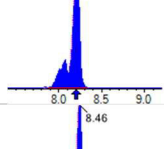
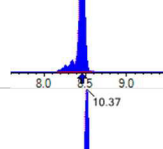
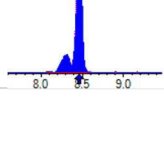
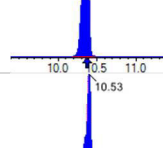
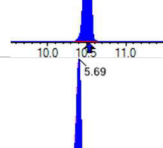
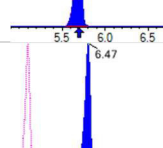
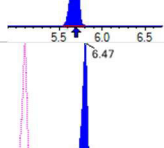
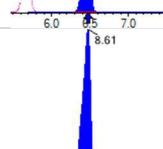
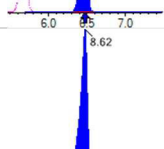
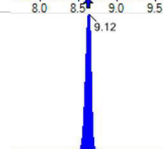
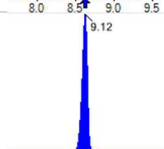
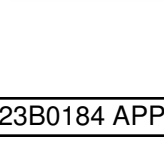
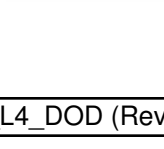
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 688199	(8.05 , N/A) (N/A , 0.00 , N/A)	716.6	N/A	0.9586 [1.0000]	95.9% { 100.0% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 1200902	(7.02 , N/A) (N/A , 0.00 , N/A)	1243.1	N/A	0.9751 [1.0000]	97.5% { 100.0% }			
13C4_PFOS_IIS	(503.0 / 79.9) 1732397	(8.24 , N/A) (N/A , 0.00 , N/A)	683.1	N/A	0.9761 [1.0000]	97.6% { 100.0% }			
13C4_PFBA_EIS	(217.0 / 172.0) 2929578	(3.46 , N/A) (N/A , 0.00 , N/A)	2587.4	N/A	7.4706 [8.0000]	93.4% { 100.0% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 2438177	(4.47 , N/A) (N/A , 0.00 , N/A)	2124.3	N/A	3.9368 [4.0000]	98.4% { 100.0% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 1534930	(5.39 , N/A) (N/A , 0.00 , N/A)	1516.3	N/A	2.0324 [2.0000]	101.6% { 100.0% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 1413227	(6.18 , N/A) (N/A , 0.00 , N/A)	1356.3	N/A	2.0064 [2.0000]	100.3% { 100.0% }			
13C8_PFOA_EIS	(421.0 / 376.0) 1666410	(6.88 , N/A) (N/A , 0.00 , N/A)	1500.9	N/A	2.0753 [2.0000]	103.8% { 100.0% }			
13C9_PFNA_EIS	(472.0 / 427.0) 709037	(7.49 , N/A) (N/A , 0.00 , N/A)	994.0	N/A	0.9344 [1.0000]	93.4% { 100.0% }			
13C6_PFDA_EIS	(519.0 / 474.0) 734422	(8.04 , N/A) (N/A , 0.00 , N/A)	933.8	N/A	0.9220 [1.0000]	92.2% { 100.0% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 752654	(8.54 , N/A) (N/A , 0.00 , N/A)	938.2	N/A	0.9538 [1.0000]	95.4% { 100.0% }			

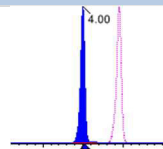
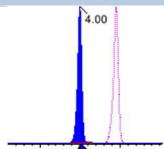
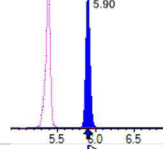
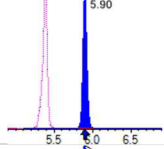
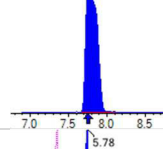
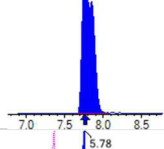
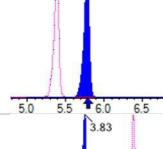
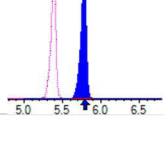
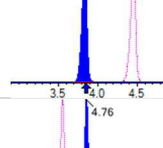
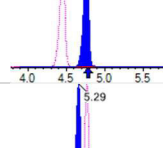
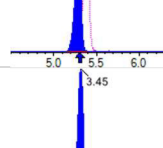
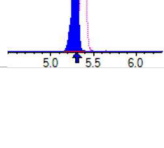
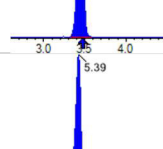
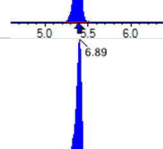
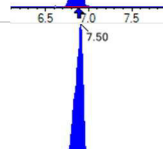
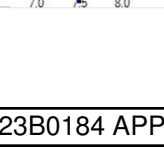
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 736893	(8.86 , N/A) (N/A , 0.00 , N/A)	1186.1	N/A	1.0069 [1.0000]	100.7% { 100.0% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 719115	(9.24 , N/A) (N/A , 0.00 , N/A)	1077.8	N/A	0.9452 [1.0000]	94.5% { 100.0% }			
13C3_PFBs_EIS	(302.0 / 80.0) 4550353	(5.35 , N/A) (N/A , 0.00 , N/A)	1731.8	N/A	1.8751 [2.0000]	93.8% { 100.0% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 2368002	(7.02 , N/A) (N/A , 0.00 , N/A)	1688.6	N/A	1.8677 [2.0000]	93.4% { 100.0% }			
13C8_PFOS_EIS	(507.0 / 80.0) 4272491	(8.24 , N/A) (N/A , 0.00 , N/A)	736.0	N/A	1.8772 [2.0000]	93.9% { 100.0% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 518489	(5.14 , N/A) (N/A , 0.00 , N/A)	990.0	N/A	3.4883 [4.0000]	87.2% { 100.0% }			
13C2_6:2FTS_EIS	(429.0 / 81.0) 590812	(6.62 , N/A) (N/A , 0.00 , N/A)	948.0	N/A	3.5255 [4.0000]	88.1% { 100.0% }			
13C2_8:2FTS_EIS	(529.0 / 81.0) 664084	(7.79 , N/A) (N/A , 0.00 , N/A)	712.1	N/A	3.7473 [4.0000]	93.7% { 100.0% }			
13C8_PFOsa_EIS	(506.0 / 78.0) 7790712	(9.83 , N/A) (N/A , 0.00 , N/A)	2362.7	N/A	1.8583 [2.0000]	92.9% { 100.0% }			
D3_NMeFOsa_EIS	(515.0 / 169.0) 1719281	(10.42 , N/A) (N/A , 0.00 , N/A)	2324.2	N/A	1.9098 [2.0000]	95.5% { 100.0% }			
D5_NEtFOsa_EIS	(531.0 / 169.0) 1563780	(10.58 , N/A) (N/A , 0.00 , N/A)	2719.7	N/A	1.9839 [2.0000]	99.2% { 100.0% }			

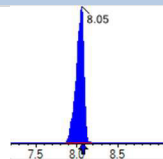
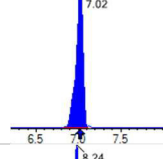
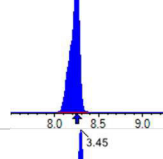
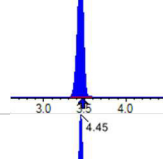
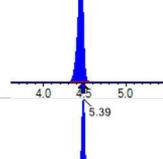
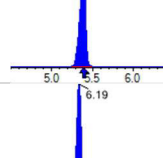
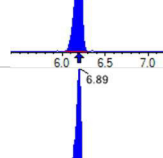
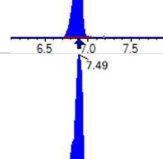
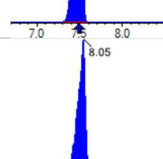
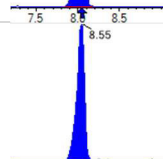
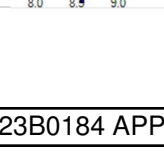
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1286524	(8.20 , N/A) (N/A , 0.00 , N/A)	1608.8	N/A	4.0277 [4.0000]	100.7% { 100.0% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 967676	(8.45 , N/A) (N/A , 0.00 , N/A)	3585.1	N/A	3.5576 [4.0000]	88.9% { 100.0% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 6051457	(10.36 , N/A) (N/A , 0.00 , N/A)	1774.7	N/A	19.7447 [20.0000]	98.7% { 100.0% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 6848678	(10.52 , N/A) (N/A , 0.00 , N/A)	1242.9	N/A	19.5351 [20.0000]	97.7% { 100.0% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 3738967	(5.69 , N/A) (N/A , 0.00 , N/A)	2143.5	N/A	7.8789 [8.0000]	98.5% { 100.0% }			

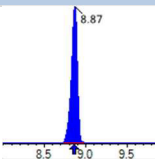
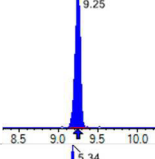
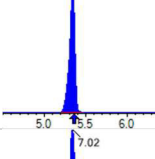
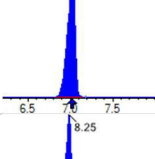
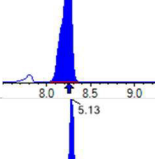
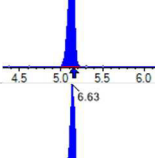
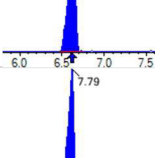
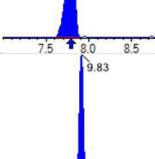
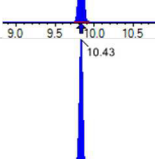
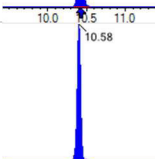
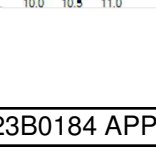
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) 13756191	(3.45 , 1.00) (0.00 , N/A , 0.0)	212.4	N/A 0.0 0.0	45.1688 [40.0000]	112.9%			
PFPeA	(263.0 / 219.0) 12096908 (263.0 / 69.0) 141038	(4.45 , 1.00) (0.00 , N/A , 0.1)	4551.8 1132.1	0.0117 97.8 97.8	23.3254 [20.0000]	116.6%			
PFHxA	(313.0 / 269.0) 7628712 (313.0 / 119.0) 762945	(5.39 , 1.00) (0.00 , N/A , 0.1)	5235.5 698940.0	0.1000 106.9 106.9	10.9129 [10.0000]	109.1%			
PFHpA	(363.0 / 319.0) 6730357 (363.0 / 169.0) 2062280	(6.19 , 1.00) (0.00 , N/A , 0.0)	8528.4 12388.9	0.3064 97.1 97.1	10.5540 [10.0000]	105.5%			
PFOA	(413.0 / 369.0) 8093602 (413.0 / 169.0) 2824786	(6.89 , 1.00) (0.00 , N/A , 0.0)	5483.0 129898.8	0.3490 103.2 103.2	10.4658 [10.0000]	104.7%			
PFNA	(463.0 / 419.0) 7308938 (463.0 / 169.0) 1544360	(7.50 , 1.00) (0.00 , N/A , 0.0)	8599.7 1209929.3	0.2113 90.3 90.3	10.7310 [10.0000]	107.3%			
PFDA	(513.0 / 469.0) 8220287 (513.0 / 169.0) 936600	(8.05 , 1.00) (0.00 , N/A , 0.1)	1821.6 1089.9	0.1139 91.9 91.9	11.1683 [10.0000]	111.7%			
PFUnA	(563.0 / 519.0) 6725530 (563.0 / 169.0) 838964	(8.55 , 1.00) (0.01 , N/A , 0.0)	1685.8 632.7	0.1247 115.9 115.9	10.4366 [10.0000]	104.4%			
PFDoA	(613.0 / 569.0) 7041213 (613.0 / 169.0) 1097210	(8.87 , 1.00) (0.00 , N/A , 0.1)	1642.2 999.2	0.1558 98.1 98.1	11.8918 [10.0000]	118.9%			
PFTTrDA	(663.0 / 619.0) 6352079 (663.0 / 169.0) 1408071	(9.07 , 1.02) (N/A , 0.00 , -0.3)	1643.1 1357.6	0.2217 81.6 81.6	11.1486 [10.0000]	111.5%			
PFTeDA	(713.0 / 669.0) 6953839 (713.0 / 169.0) 1458805	(9.25 , 1.00) (0.00 , N/A , 0.1)	1868.8 1375.3	0.2098 96.0 96.0	10.9787 [10.0000]	109.8%			

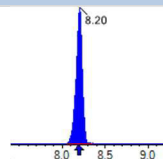
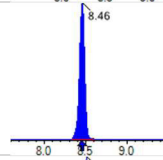
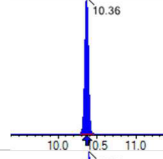
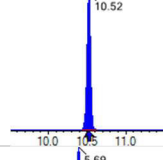
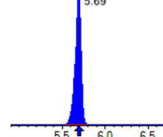
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) 12977840 (299.0 / 99.0) 7939442	(5.34 , 1.00) (0.00 , N/A , 0.0)	2308.0 2277.1	0.6118 97.3 97.3	9.9966 [8.8473]	113.0%			
PFPeS	(349.0 / 80.0) 21980088 (349.0 / 99.0) 7657852	(6.26 , 0.89) (N/A , 0.01 , -0.1)	12552.9 74884.5	0.3484 104.1 104.1	10.3002 [9.3838]	109.8%			
PFHxS	(399.0 / 80.0) 19582487 (399.0 / 99.0) 6332147	(7.02 , 1.00) (0.00 , N/A , 0.1)	7421.0 28788.2	0.3234 101.0 101.0	10.4590 [9.1098]	114.8%			
PFHpS	(449.0 / 80.0) 21196242 (449.0 / 99.0) 5607040	(7.67 , 0.93) (N/A , 0.00 , 0.1)	17960.6 109693.1	0.2645 97.4 97.4	10.8116 [9.5141]	113.6%			
PFOS	(499.0 / 80.0) 24116485 (499.0 / 99.0) 4922727	(8.25 , 1.00) (0.00 , N/A , 0.0)	364.6 548.8	0.2041 92.8 92.8	10.0367 [9.2749]	108.2%			
PFNS	(549.0 / 80.0) 25332042 (549.0 / 99.0) 5570113	(8.71 , 1.06) (N/A , 0.00 , 0.1)	6278.0 21597.1	0.2199 91.4 91.4	10.5356 [9.5989]	109.8%			
PFDS	(599.0 / 80.0) 28143793 (599.0 / 99.0) 5958051	(8.97 , 1.09) (N/A , 0.00 , 0.1)	2480.6 1769.0	0.2117 104.6 104.6	10.5322 [9.6311]	109.4%			
PFDoS	(699.0 / 80.0) 13398721 (699.0 / 99.0) 2532090	(9.32 , 1.13) (N/A , 0.00 , -0.1)	2343.1 1388.9	0.1890 96.7 96.7	10.8616 [9.6956]	112.0%			
4:2FTS	(327.0 / 307.0) 16267687 (327.0 / 81.0) 10789759	(5.13 , 1.00) (0.00 , N/A , -0.1)	2385.1 2040.9	0.6633 95.7 95.7	46.9279 [37.3811]	125.5%			
6:2FTS	(427.0 / 407.0) 8704318 (427.0 / 81.0) 7312928	(6.63 , 1.00) (0.00 , N/A , 0.0)	2162.2 2275.0	0.8401 99.1 99.1	44.6757 [37.9617]	117.7%			
8:2FTS	(527.0 / 507.0) 7908197 (527.0 / 81.0) 6505276	(7.80 , 1.00) (0.00 , N/A , -0.2)	1673.9 1834.2	0.8226 100.8 100.8	39.4573 [38.3315]	102.9%			

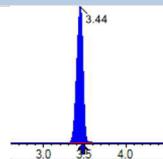
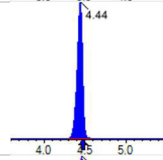
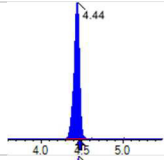
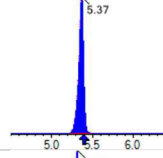
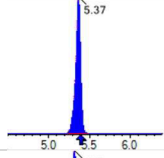
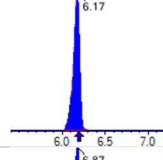
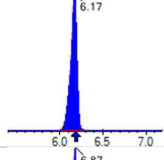
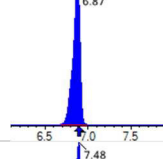
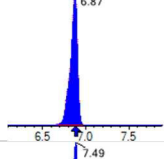
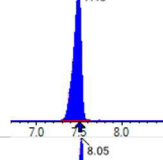
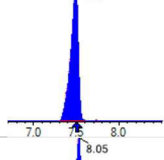
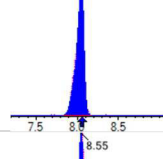
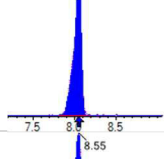
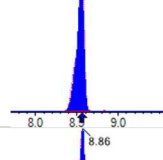
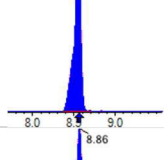
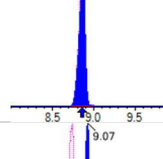
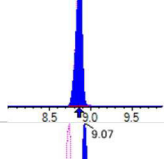
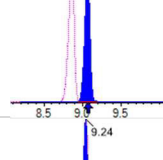
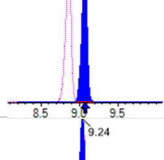
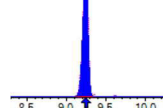
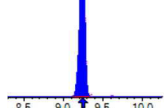
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) 31312377 (498.0 / 478.0) 678872	(9.83 , 1.00) (0.00 , N/A , 0.0)	2478.3 1594.4	0.0217 110.3 110.3	10.5200 [10.0000]	105.2%			
NMeFOSA	(512.0 / 219.0) 30750368 (512.0 / 169.0) 25658212	(10.42 , 1.00) (0.00 , N/A , 0.8)	2777.8 4757.7	0.8344 101.0 101.0	42.4322 [40.0000]	106.1%			
NEIFOSA	(526.0 / 219.0) 32433415 (526.0 / 169.0) 38954221	(10.57 , 1.00) (-0.01 , N/A , 0.4)	10684.8 10309.0	1.2011 94.8 94.8	45.2092 [40.0000]	113.0%			
NMeFOSAA	(570.0 / 419.0) 2604111 (570.0 / 483.0) 1318093	(8.21 , 1.00) (0.01 , N/A , 0.0)	2086.3 472.3	0.5062 103.2 103.2	10.3978 [10.0000]	104.0%			
NEIFOSAA	(584.0 / 419.0) 2386967 (584.0 / 526.0) 1238724	(8.46 , 1.00) (0.01 , N/A , 0.0)	2092.6 1743.6	0.5190 91.9 91.9	10.8287 [10.0000]	108.3%			
NMeFOSE	(616.0 / 59.0) 13284521	(10.37 , 1.00) (0.01 , N/A , 0.0)	3543.7	N/A 0.0 0.0	44.5128 [40.0000]	111.3%			
NEtFOSE	(630.0 / 59.0) 15130991	(10.53 , 1.00) (0.01 , N/A , 0.0)	1041.8	N/A 0.0 0.0	43.9064 [40.0000]	109.8%			
HFPO-DA	(285.0 / 169.0) 8522632 (285.0 / 185.0) 21710472	(5.69 , 1.00) (0.00 , N/A , 0.1)	2322.0 2558.3	2.5474 92.7 92.7	22.2268 [20.0000]	111.1%			
ADONA	(377.0 / 85.0) 30190474 (377.0 / 251.0) 2788031	(6.47 , 1.14) (N/A , 0.01 , -0.1)	2326.7 1959.1	0.0923 105.5 105.5	21.2148 [18.8540]	112.5%			
9CI-Pf3ONS	(531.0 / 351.0) 64220531 (533.0 / 353.0) 21002770	(8.61 , 1.51) (N/A , 0.00 , -0.2)	1841.9 2062.2	0.3270 111.0 111.0	19.2419 [18.6651]	103.1%			
11CI-PF3OUDS	(631.0 / 451.0) 31798587 (633.0 / 453.0) 10440648	(9.12 , 1.60) (N/A , 0.00 , -0.2)	1618.6 1530.7	0.3283 106.4 106.4	19.3788 [18.8642]	102.7%			

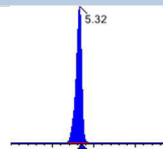
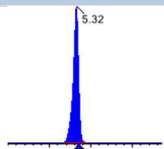
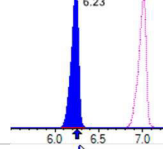
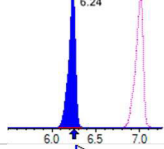
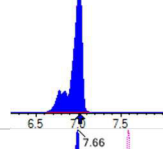
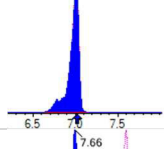
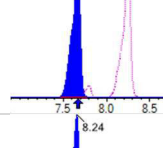
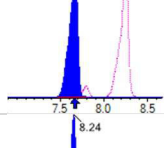
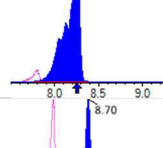
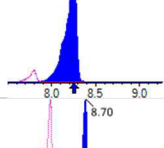
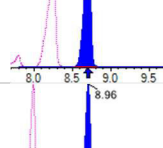
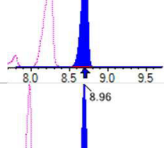
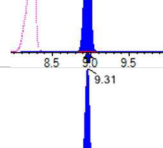
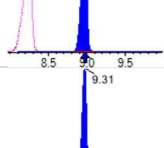
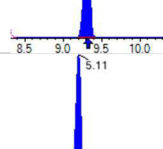
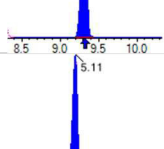
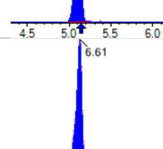
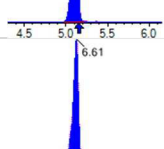
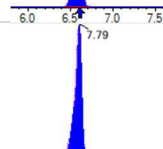
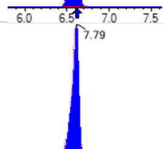
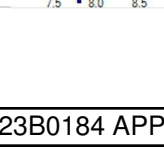
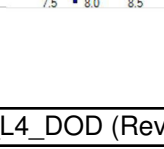
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) 681185 (241.0 / 117.0) 1074158	(4.00 , 0.90) (N/A , -0.02 , 0.0)	1390.7 1557.7	1.5769 90.9 90.9	44.4762 [40.0000]	111.2%			
5:3FTCA	(341.0 / 236.7) 4336443 (341.0 / 217.0) 7350647	(5.90 , 1.10) (N/A , 0.00 , 0.0)	2329.2 1923.9	1.6951 100.8 100.8	43.3947 [40.0000]	108.5%			
7:3FTCA	(441.0 / 317.0) 7430997 (441.0 / 337.0) 6401756	(7.79 , 1.45) (N/A , 0.00 , -0.1)	847.0 747.9	0.8615 100.3 100.3	41.3624 [40.0000]	103.4%			
PFEESA	(315.0 / 135.0) 17691646 (315.0 / 83.0) 4686976	(5.78 , 1.07) (N/A , 0.00 , 0.0)	2594.2 2429.5	0.2649 94.1 94.1	20.3659 [17.8492]	114.1%			
PFMPA	(229.0 / 85.0) 2952170	(3.83 , 0.86) (N/A , -0.02 , 0.0)	2852.7	N/A 0.0 0.0	22.2640 [20.0000]	111.3%			
PFMBA	(279.0 / 85.0) 10963289	(4.76 , 1.07) (N/A , -0.02 , 0.0)	2433.9	N/A 0.0 0.0	24.1339 [20.0000]	120.7%			
NFDHA	(295.0 / 201.0) 8574949 (295.0 / 85.0) 8583239	(5.29 , 0.98) (N/A , -0.01 , -0.1)	2419.3 2688.3	1.0010 95.9 95.9	22.0541 [20.0000]	110.3%			
13C3_PFBA_IIS	(216.0 / 172.0) 334286	(3.45 , N/A) (N/A , -0.01 , N/A)	1399.2	N/A	0.9860 [1.0000]	98.6% { 101.4% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 579108	(5.39 , N/A) (N/A , -0.01 , N/A)	1279.8	N/A	0.9737 [1.0000]	97.4% { 104.2% }			
13C4_PFOA_IIS	(417.0 / 372.0) 835085	(6.89 , N/A) (N/A , 0.01 , N/A)	950.6	N/A	1.0266 [1.0000]	102.7% { 111.3% }			
13C5_PFNA_IIS	(468.0 / 423.0) 770134	(7.50 , N/A) (N/A , 0.01 , N/A)	751.5	N/A	1.0085 [1.0000]	100.9% { 101.1% }			

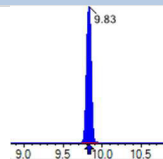
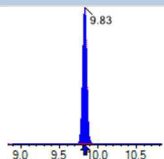
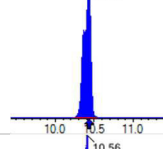
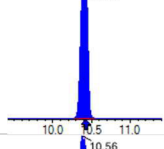
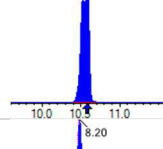
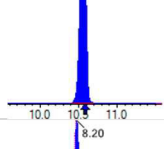
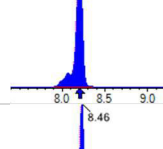
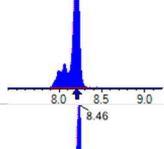
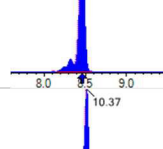
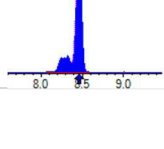
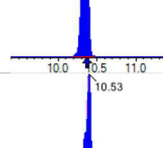
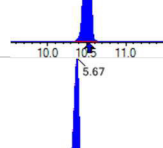
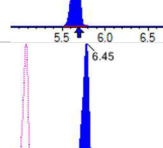
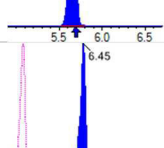
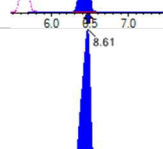
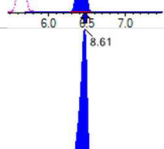
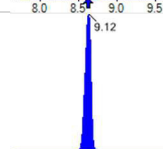
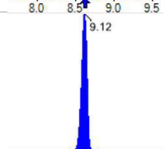
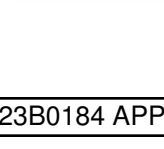
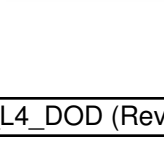
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 754411	(8.05 , N/A) (N/A , 0.00 , N/A)	1794.3	N/A	1.0509 [1.0000]	105.1% { 109.6% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 1188785	(7.02 , N/A) (N/A , 0.00 , N/A)	1465.1	N/A	0.9653 [1.0000]	96.5% { 99.0% }			
13C4_PFOS_IIS	(503.0 / 79.9) 1798446	(8.24 , N/A) (N/A , 0.01 , N/A)	1218.7	N/A	1.0133 [1.0000]	101.3% { 103.8% }			
13C4_PFBA_EIS	(217.0 / 172.0) 2881013	(3.45 , N/A) (N/A , -0.01 , N/A)	2734.6	N/A	7.2479 [8.0000]	90.6% { 98.3% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 2354297	(4.45 , N/A) (N/A , -0.02 , N/A)	2212.4	N/A	3.6486 [4.0000]	91.2% { 96.6% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 1498135	(5.39 , N/A) (N/A , -0.01 , N/A)	1965.1	N/A	1.9040 [2.0000]	95.2% { 97.6% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 1428247	(6.19 , N/A) (N/A , 0.01 , N/A)	1318.0	N/A	1.9462 [2.0000]	97.3% { 101.1% }			
13C8_PFOA_EIS	(421.0 / 376.0) 1605188	(6.89 , N/A) (N/A , 0.01 , N/A)	1288.6	N/A	1.7959 [2.0000]	89.8% { 96.3% }			
13C9_PFNA_EIS	(472.0 / 427.0) 721057	(7.49 , N/A) (N/A , 0.00 , N/A)	1444.7	N/A	0.9395 [1.0000]	94.0% { 101.7% }			
13C6_PFDA_EIS	(519.0 / 474.0) 742371	(8.05 , N/A) (N/A , 0.02 , N/A)	951.4	N/A	0.8502 [1.0000]	85.0% { 101.1% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 732822	(8.55 , N/A) (N/A , 0.00 , N/A)	893.1	N/A	0.8471 [1.0000]	84.7% { 97.4% }			

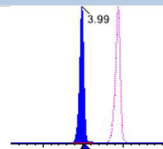
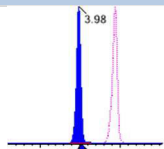
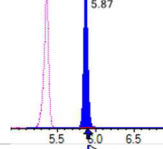
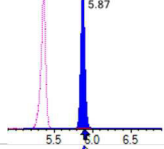
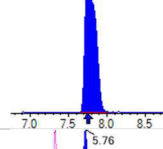
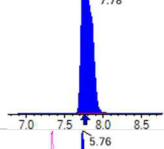
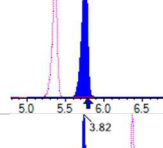
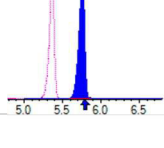
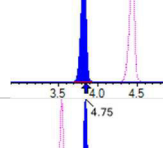
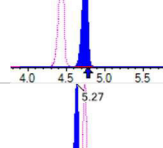
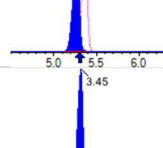
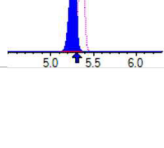
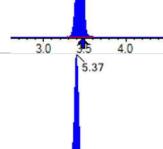
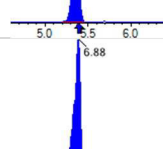
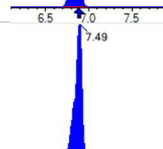
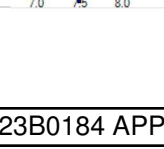
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 678675	(8.87, N/A) (N/A, 0.01, N/A)	2772.2	N/A	0.8459 [1.0000]	84.6% { 92.1% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 688843	(9.25, N/A) (N/A, 0.01, N/A)	739.3	N/A	0.8259 [1.0000]	82.6% { 95.8% }			
13C3_PFBs_EIS	(302.0 / 80.0) 4566480	(5.34, N/A) (N/A, -0.01, N/A)	1958.9	N/A	1.9009 [2.0000]	95.0% { 100.4% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 2377218	(7.02, N/A) (N/A, 0.00, N/A)	1309.9	N/A	1.8941 [2.0000]	94.7% { 100.4% }			
13C8_PFOS_EIS	(507.0 / 80.0) 4354524	(8.25, N/A) (N/A, 0.00, N/A)	434.0	N/A	1.8430 [2.0000]	92.2% { 101.9% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 491270	(5.13, N/A) (N/A, -0.02, N/A)	885.7	N/A	3.3388 [4.0000]	83.5% { 94.8% }			
13C2_6:2FTS_EIS	(429.0 / 81.0) 597386	(6.63, N/A) (N/A, 0.01, N/A)	975.8	N/A	3.6010 [4.0000]	90.0% { 101.1% }			
13C2_8:2FTS_EIS	(529.0 / 81.0) 714138	(7.79, N/A) (N/A, 0.01, N/A)	1008.7	N/A	4.0708 [4.0000]	101.8% { 107.5% }			
13C8_PFOsa_EIS	(506.0 / 78.0) 7660367	(9.83, N/A) (N/A, 0.01, N/A)	2237.9	N/A	1.7601 [2.0000]	88.0% { 98.3% }			
D3_NMeFOsa_EIS	(515.0 / 169.0) 1687624	(10.43, N/A) (N/A, 0.00, N/A)	2364.0	N/A	1.8058 [2.0000]	90.3% { 98.2% }			
D5_NEtFOsa_EIS	(531.0 / 169.0) 1476954	(10.58, N/A) (N/A, 0.00, N/A)	2489.8	N/A	1.8049 [2.0000]	90.2% { 94.4% }			

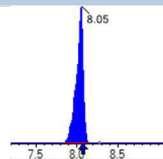
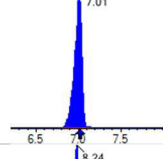
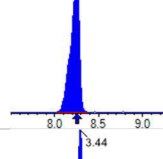
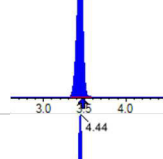
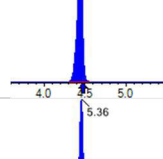
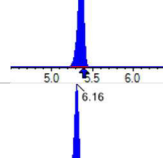
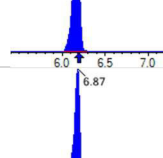
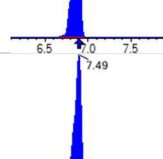
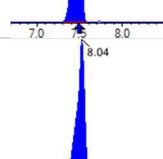
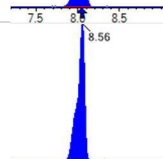
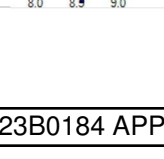
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1172136	(8.20 , N/A) (N/A , 0.01 , N/A)	1163.1	N/A	3.5348 [4.0000]	88.4% { 91.1% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 1011453	(8.46 , N/A) (N/A , 0.00 , N/A)	57422.4	N/A	3.5820 [4.0000]	89.6% { 104.5% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 5696500	(10.36 , N/A) (N/A , 0.00 , N/A)	1990.5	N/A	17.9039 [20.0000]	89.5% { 94.1% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 6821043	(10.52 , N/A) (N/A , 0.00 , N/A)	1437.2	N/A	18.7417 [20.0000]	93.7% { 99.6% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 3741666	(5.69 , N/A) (N/A , 0.00 , N/A)	2136.5	N/A	7.5676 [8.0000]	94.6% { 100.1% }			

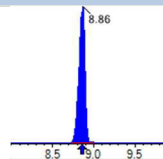
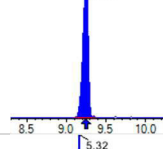
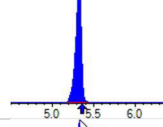
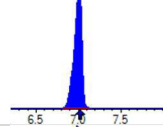
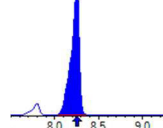
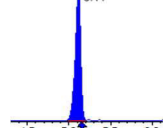
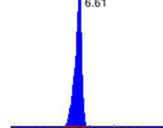
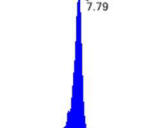
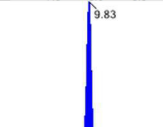
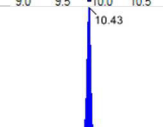

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) 23383373	(3.44 , 1.00) (0.00 , N/A , 0.0)	178.6	N/A 0.0 0.0	78.0490 [80.0000]	97.6%			
PFPeA	(263.0 / 219.0) 21458957 (263.0 / 69.0) 247776	(4.44 , 1.00) (0.00 , N/A , 0.0)	5369.8 1044.0	0.0115 96.9 96.9	38.7834 [40.0000]	97.0%			
PFHxA	(313.0 / 269.0) 14427222 (313.0 / 119.0) 1437857	(5.37 , 1.00) (0.00 , N/A , 0.0)	6552.7 208683.0	0.0997 106.5 106.5	19.6900 [20.0000]	98.4%			
PFHpA	(363.0 / 319.0) 12828245 (363.0 / 169.0) 3914525	(6.17 , 1.00) (0.00 , N/A , 0.0)	8203.2 30700.4	0.3051 96.7 96.7	20.9762 [20.0000]	104.9%			
PFOA	(413.0 / 369.0) 15358487 (413.0 / 169.0) 5405302	(6.87 , 1.00) (0.00 , N/A , -0.1)	6552.7 8960.8	0.3519 104.0 104.0	19.3192 [20.0000]	96.6%			
PFNA	(463.0 / 419.0) 13722178 (463.0 / 169.0) 3139510	(7.48 , 1.00) (0.00 , N/A , -0.1)	6782.1 997608.1	0.2288 97.7 97.7	19.0472 [20.0000]	95.2%			
PFDA	(513.0 / 469.0) 15164091 (513.0 / 169.0) 1754409	(8.05 , 1.00) (0.00 , N/A , 0.0)	1977.5 791.4	0.1157 93.4 93.4	18.3770 [20.0000]	91.9%			
PFUnA	(563.0 / 519.0) 13518716 (563.0 / 169.0) 1506565	(8.55 , 1.00) (-0.01 , N/A , 0.0)	1803.3 987.4	0.1114 103.5 103.5	18.7830 [20.0000]	93.9%			
PFDoA	(613.0 / 569.0) 12908177 (613.0 / 169.0) 2110662	(8.86 , 1.00) (0.00 , N/A , 0.1)	1612.7 1268.2	0.1635 102.9 102.9	20.6337 [20.0000]	103.2%			
PFTTrDA	(663.0 / 619.0) 11417572 (663.0 / 169.0) 2735060	(9.07 , 1.02) (N/A , 0.00 , -0.1)	2153.2 1270.0	0.2395 88.2 88.2	18.9665 [20.0000]	94.8%			
PFTeDA	(713.0 / 669.0) 13577078 (713.0 / 169.0) 2681370	(9.24 , 1.00) (0.00 , N/A , 0.1)	2625.5 1433.6	0.1975 90.3 90.3	20.1334 [20.0000]	100.7%			

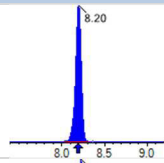
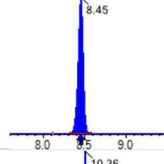
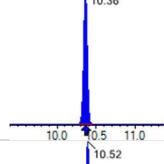
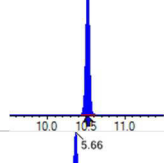
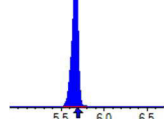
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) 22103106 (299.0 / 99.0) 14346921	(5.32 , 1.00) (0.00 , N/A , 0.0)	2399.5 2490.8	0.6491 103.2 103.2	17.2096 [17.6947]	97.3%			
PFPeS	(349.0 / 80.0) 39527279 (349.0 / 99.0) 14957759	(6.23 , 0.89) (N/A , -0.02 , -0.1)	12745.1 16575.6	0.3784 113.1 113.1	17.3040 [18.7676]	92.2%			
PFHxS	(399.0 / 80.0) 36183370 (399.0 / 99.0) 12034566	(7.01 , 1.00) (0.00 , N/A , 0.0)	8088.8 13392.5	0.3326 103.9 103.9	18.0536 [18.2197]	99.1%			
PFHpS	(449.0 / 80.0) 39494451 (449.0 / 99.0) 11202590	(7.66 , 0.93) (N/A , 0.00 , 0.1)	12534.6 46835.5	0.2836 104.5 104.5	18.9442 [19.0281]	99.6%			
PFOS	(499.0 / 80.0) 44333658 (499.0 / 99.0) 9725121	(8.24 , 1.00) (0.00 , N/A , -0.1)	349.7 622.1	0.2194 99.7 99.7	17.3509 [18.5499]	93.5%			
PFNS	(549.0 / 80.0) 46250056 (549.0 / 99.0) 11082689	(8.70 , 1.06) (N/A , 0.00 , -0.1)	8064.5 20553.9	0.2396 99.6 99.6	18.0889 [19.1977]	94.2%			
PFDS	(599.0 / 80.0) 47732243 (599.0 / 99.0) 10914827	(8.96 , 1.09) (N/A , 0.00 , 0.0)	2994.9 1968.7	0.2287 113.0 113.0	16.7981 [19.2621]	87.2%			
PFDoS	(699.0 / 80.0) 24262994 (699.0 / 99.0) 4721414	(9.31 , 1.13) (N/A , 0.00 , 0.0)	2020.0 2409.4	0.1946 99.6 99.6	18.4964 [19.3913]	95.4%			
4:2FTS	(327.0 / 307.0) 29767148 (327.0 / 81.0) 20604226	(5.11 , 1.00) (0.00 , N/A , 0.0)	1811.1 2139.1	0.6922 99.9 99.9	69.5480 [74.7622]	93.0%			
6:2FTS	(427.0 / 407.0) 16320654 (427.0 / 81.0) 14608467	(6.61 , 1.00) (0.00 , N/A , 0.1)	2530.3 2769.3	0.8951 105.6 105.6	75.9834 [75.9234]	100.1%			
8:2FTS	(527.0 / 507.0) 15535552 (527.0 / 81.0) 12858497	(7.79 , 1.00) (0.01 , N/A , 0.0)	2045.7 2402.4	0.8277 101.5 101.5	73.2925 [76.6631]	95.6%			

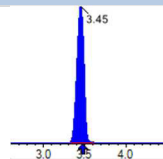
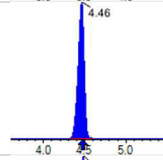
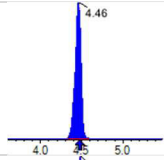
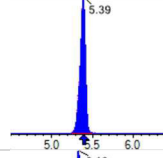
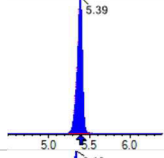
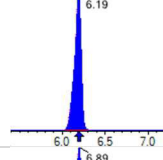
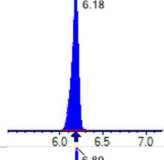
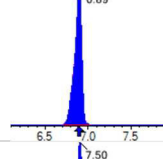
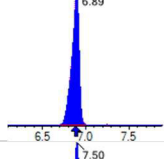
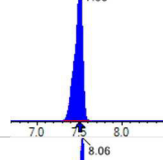
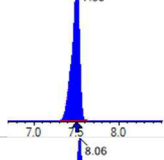
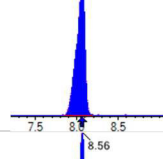
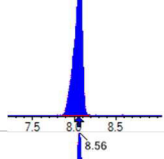
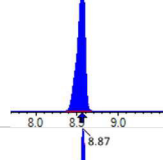
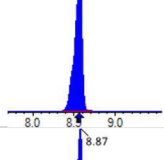
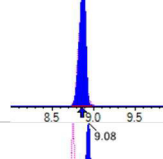
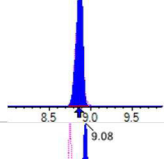
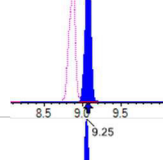
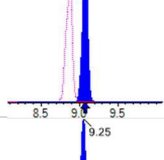
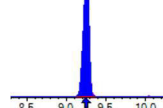
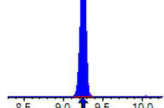
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) 47309564 (498.0 / 478.0) 1208162	(9.83 , 1.00) (0.00 , N/A , 0.0)	2587.2 1996.0	0.0255 130.0 130.0	15.7269 [20.0000]	78.6%			
NMeFOSA	(512.0 / 219.0) 50633693 (512.0 / 169.0) 42887648	(10.42 , 1.00) (-0.01 , N/A , 0.7)	2632.2 3591.1	0.8470 102.5 102.5	67.1487 [80.0000]	83.9%			
NEIFOSA	(526.0 / 219.0) 52831618 (526.0 / 169.0) 59879381	(10.56 , 1.00) (-0.01 , N/A , 0.2)	16358.7 16873.8	1.1334 89.5 89.5	75.9802 [80.0000]	95.0%			
NMeFOSAA	(570.0 / 419.0) 5372240 (570.0 / 483.0) 2423733	(8.20 , 1.00) (0.01 , N/A , -0.1)	2518.8 570.4	0.4512 92.0 92.0	20.3244 [20.0000]	101.6%			
NEIFOSAA	(584.0 / 419.0) 4636112 (584.0 / 526.0) 2673056	(8.46 , 1.00) (0.01 , N/A , -0.1)	2447.4 1722.5	0.5766 102.1 102.1	19.4219 [20.0000]	97.1%			
NMeFOSE	(616.0 / 59.0) 23903925	(10.37 , 1.00) (0.00 , N/A , 0.0)	3144.6	N/A 0.0 0.0	79.4621 [80.0000]	99.3%			
NEtFOSE	(630.0 / 59.0) 25464529	(10.53 , 1.00) (0.00 , N/A , 0.0)	997.0	N/A 0.0 0.0	79.2966 [80.0000]	99.1%			
HFPO-DA	(285.0 / 169.0) 16096251 (285.0 / 185.0) 37305083	(5.67 , 1.00) (0.00 , N/A , 0.0)	2168.1 2710.8	2.3176 84.3 84.3	39.5969 [40.0000]	99.0%			
ADONA	(377.0 / 85.0) 50291844 (377.0 / 251.0) 5091270	(6.45 , 1.14) (N/A , -0.02 , -0.2)	2013.6 1620.8	0.1012 115.7 115.7	33.3350 [37.7080]	88.4%			
9CI-Pf3ONS	(531.0 / 351.0) 95741997 (533.0 / 353.0) 37415789	(8.61 , 1.52) (N/A , -0.01 , -0.4)	1281.7 1930.6	0.3908 132.7 132.7	27.0589 [37.3302]	72.5%			
11CI-PF3OUDS	(631.0 / 451.0) 54217062 (633.0 / 453.0) 18287065	(9.12 , 1.61) (N/A , 0.00 , 0.0)	2334.8 2763.7	0.3373 109.3 109.3	31.1666 [37.7283]	82.6%			

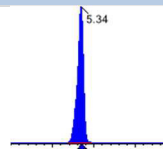
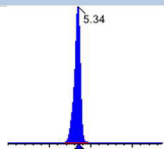
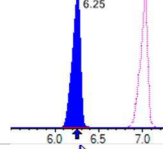
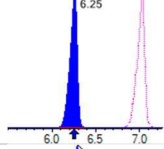
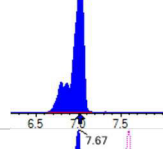
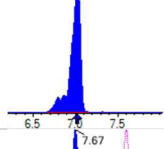
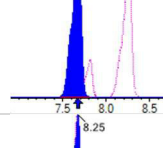
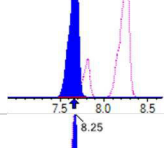
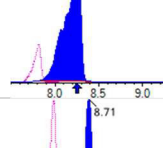
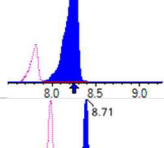
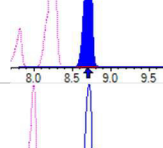
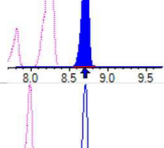
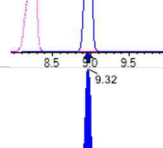
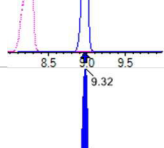
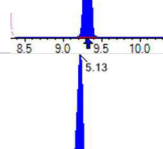
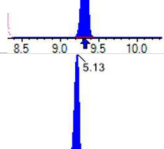
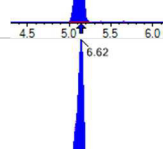
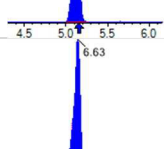
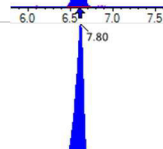
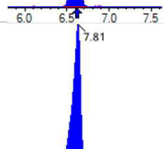
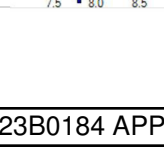
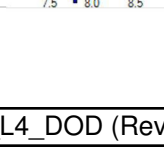
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) 1338965 (241.0 / 117.0) 2105387	(3.99, 0.90) (N/A, -0.03, 0.1)	1795.4 1619.8	1.5724 90.7 90.7	81.9434 [80.0000]	102.4%			
5:3FTCA	(341.0 / 236.7) 8659309 (341.0 / 217.0) 14561045	(5.87, 1.10) (N/A, -0.03, -0.1)	1842.3 2512.8	1.6815 100.0 100.0	82.6723 [80.0000]	103.3%			
7:3FTCA	(441.0 / 317.0) 14652173 (441.0 / 337.0) 12772655	(7.78, 1.45) (N/A, -0.01, 0.0)	1090.3 1120.6	0.8717 101.5 101.5	77.8099 [80.0000]	97.3%			
PFEESA	(315.0 / 135.0) 30437306 (315.0 / 83.0) 8443144	(5.76, 1.07) (N/A, -0.02, -0.1)	2175.8 2474.4	0.2774 98.5 98.5	33.4284 [35.6984]	93.6%			
PFMPA	(229.0 / 85.0) 5727862	(3.82, 0.86) (N/A, -0.03, 0.0)	2828.0	N/A 0.0 0.0	40.4890 [40.0000]	101.2%			
PFMBA	(279.0 / 85.0) 19369586	(4.75, 1.07) (N/A, -0.03, 0.0)	2266.2	N/A 0.0 0.0	39.9658 [40.0000]	99.9%			
NFDHA	(295.0 / 201.0) 16110694 (295.0 / 85.0) 16234898	(5.27, 0.98) (N/A, -0.03, 0.0)	2400.1 2111.0	1.0077 96.5 96.5	39.5318 [40.0000]	98.8%			
13C3_PFBA_IIS	(216.0 / 172.0) 292693	(3.45, N/A) (N/A, -0.02, N/A)	1130.6	N/A	0.8633 [1.0000]	86.3% { 88.8% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 568868	(5.37, N/A) (N/A, -0.03, N/A)	1276.6	N/A	0.9565 [1.0000]	95.6% { 102.3% }			
13C4_PFOA_IIS	(417.0 / 372.0) 793430	(6.88, N/A) (N/A, 0.00, N/A)	1116.4	N/A	0.9754 [1.0000]	97.5% { 105.8% }			
13C5_PFNA_IIS	(468.0 / 423.0) 673170	(7.49, N/A) (N/A, 0.00, N/A)	868.6	N/A	0.8816 [1.0000]	88.2% { 88.4% }			

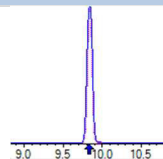
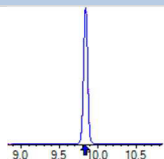
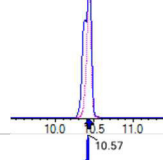
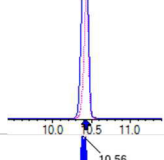
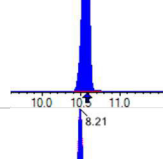
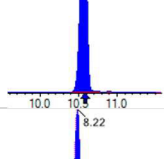
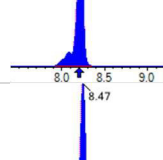
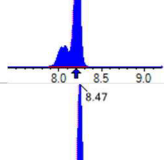
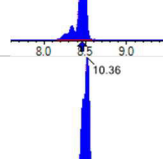
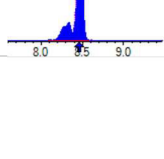
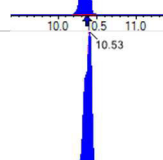
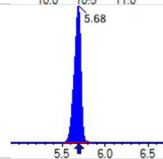
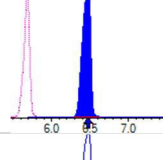
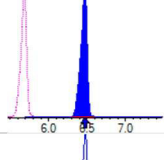
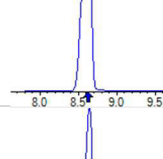
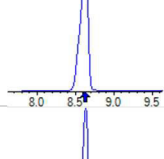
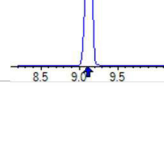
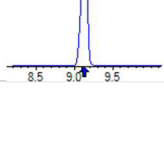
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 694088	(8.05 , N/A) (N/A , 0.00 , N/A)	849.4	N/A	0.9669 [1.0000]	96.7% { 100.9% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 1223798	(7.01 , N/A) (N/A , -0.01 , N/A)	1187.3	N/A	0.9937 [1.0000]	99.4% { 101.9% }			
13C4_PFOS_IIS	(503.0 / 79.9) 1695346	(8.24 , N/A) (N/A , 0.00 , N/A)	1052.4	N/A	0.9552 [1.0000]	95.5% { 97.9% }			
13C4_PFBA_EIS	(217.0 / 172.0) 2834169	(3.44 , N/A) (N/A , -0.02 , N/A)	3055.5	N/A	8.1433 [8.0000]	101.8% { 96.7% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 2511768	(4.44 , N/A) (N/A , -0.03 , N/A)	2146.6	N/A	3.9627 [4.0000]	99.1% { 103.0% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 1570281	(5.36 , N/A) (N/A , -0.03 , N/A)	1862.1	N/A	2.0316 [2.0000]	101.6% { 102.3% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 1369692	(6.16 , N/A) (N/A , -0.02 , N/A)	1255.9	N/A	1.9000 [2.0000]	95.0% { 96.9% }			
13C8_PFOA_EIS	(421.0 / 376.0) 1650128	(6.87 , N/A) (N/A , -0.01 , N/A)	1020.2	N/A	1.9431 [2.0000]	97.2% { 99.0% }			
13C9_PFNA_EIS	(472.0 / 427.0) 762691	(7.49 , N/A) (N/A , 0.00 , N/A)	1175.4	N/A	1.1369 [1.0000]	113.7% { 107.6% }			
13C6_PFDA_EIS	(519.0 / 474.0) 832267	(8.04 , N/A) (N/A , 0.01 , N/A)	741.2	N/A	1.0360 [1.0000]	103.6% { 113.3% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 818464	(8.56 , N/A) (N/A , 0.01 , N/A)	1062.4	N/A	1.0284 [1.0000]	102.8% { 108.7% }			

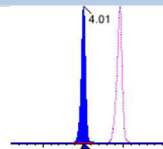
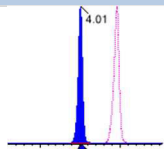
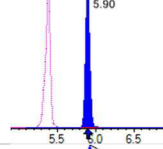
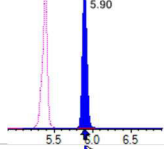
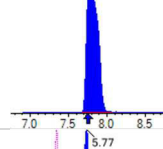
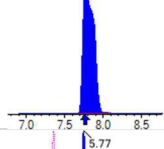
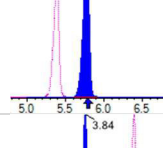
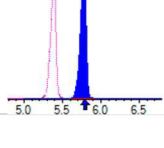
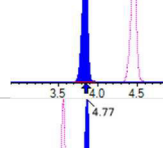
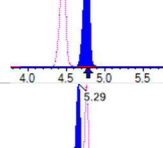
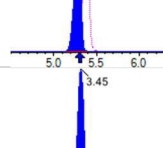
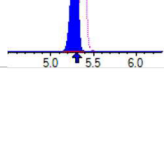
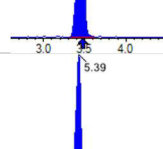
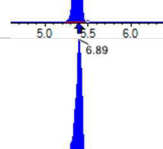
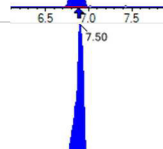
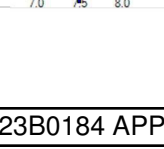
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 717052	(8.86 , N/A) (N/A , 0.00 , N/A)	1216.8	N/A	0.9714 [1.0000]	97.1% { 97.3% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 733391	(9.24 , N/A) (N/A , 0.00 , N/A)	686.9	N/A	0.9558 [1.0000]	95.6% { 102.0% }			
13C3_PFBs_EIS	(302.0 / 80.0) 4517665	(5.32 , N/A) (N/A , -0.03 , N/A)	1965.7	N/A	1.8268 [2.0000]	91.3% { 99.3% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 2544685	(7.01 , N/A) (N/A , -0.01 , N/A)	1461.3	N/A	1.9695 [2.0000]	98.5% { 107.5% }			
13C8_PFOS_EIS	(507.0 / 80.0) 4630519	(8.24 , N/A) (N/A , 0.00 , N/A)	383.3	N/A	2.0790 [2.0000]	104.0% { 108.4% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 606565	(5.11 , N/A) (N/A , -0.03 , N/A)	960.9	N/A	4.0045 [4.0000]	100.1% { 117.0% }			
13C2_6:2FTS_EIS	(429.0 / 81.0) 658583	(6.61 , N/A) (N/A , -0.01 , N/A)	1220.7	N/A	3.8563 [4.0000]	96.4% { 111.5% }			
13C2_8:2FTS_EIS	(529.0 / 81.0) 755264	(7.79 , N/A) (N/A , 0.00 , N/A)	941.1	N/A	4.1821 [4.0000]	104.6% { 113.7% }			
13C8_PFOsa_EIS	(506.0 / 78.0) 7742013	(9.83 , N/A) (N/A , 0.00 , N/A)	2144.5	N/A	1.8870 [2.0000]	94.4% { 99.4% }			
D3_NMeFOsa_EIS	(515.0 / 169.0) 1755992	(10.43 , N/A) (N/A , 0.00 , N/A)	1880.3	N/A	1.9932 [2.0000]	99.7% { 102.1% }			
D5_NEtFOsa_EIS	(531.0 / 169.0) 1431512	(10.58 , N/A) (N/A , 0.00 , N/A)	2700.7	N/A	1.8557 [2.0000]	92.8% { 91.5% }			

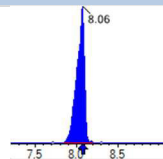
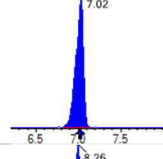
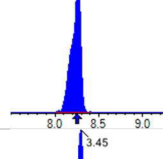
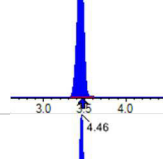
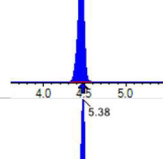
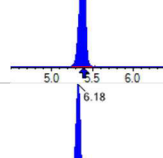
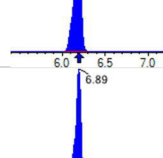
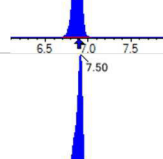
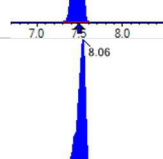
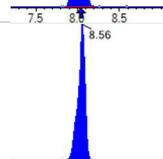
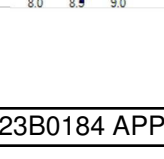
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1237081	(8.20 , N/A) (N/A , 0.00 , N/A)	1318.4	N/A	3.9576 [4.0000]	98.9% { 96.2% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 1095311	(8.45 , N/A) (N/A , 0.00 , N/A)	2552.9	N/A	4.1149 [4.0000]	102.9% { 113.2% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 5741908	(10.36 , N/A) (N/A , 0.00 , N/A)	2203.4	N/A	19.1441 [20.0000]	95.7% { 94.9% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 6356128	(10.52 , N/A) (N/A , 0.00 , N/A)	1331.0	N/A	18.5264 [20.0000]	92.6% { 92.8% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 3966715	(5.66 , N/A) (N/A , -0.02 , N/A)	1806.1	N/A	8.1672 [8.0000]	102.1% { 106.1% }			

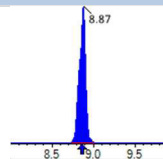
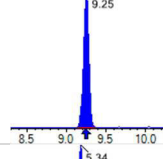
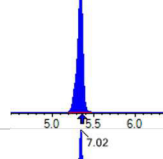
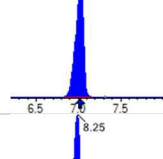
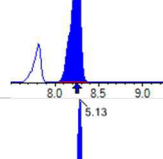
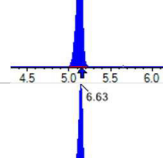
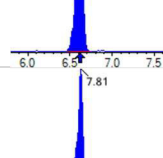
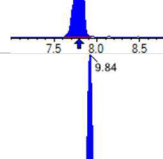
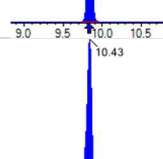
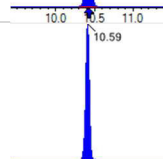
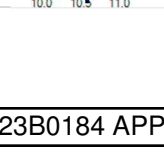
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) 41499707	(3.45 , 1.00) (0.00 , N/A , 0.0)	170.0	N/A 0.0 0.0	181.2987 [200.0000]	90.6%			
PFPeA	(263.0 / 219.0) 41492703 (263.0 / 69.0) 527610	(4.46 , 1.00) (0.00 , N/A , 0.0)	6500.0 1537.0	0.0127 106.7 106.7	91.0354 [100.0000]	91.0%			
PFHxA	(313.0 / 269.0) 29254814 (313.0 / 119.0) 3076533	(5.39 , 1.00) (0.00 , N/A , 0.0)	9581.4 34631.2	0.1052 112.4 112.4	46.3885 [50.0000]	92.8%			
PFHpA	(363.0 / 319.0) 27768317 (363.0 / 169.0) 8460728	(6.19 , 1.00) (0.00 , N/A , 0.1)	9861.5 8602.1	0.3047 96.5 96.5	47.3795 [50.0000]	94.8%			
PFOA	(413.0 / 369.0) 34736184 (413.0 / 169.0) 11436701	(6.89 , 1.00) (0.00 , N/A , 0.0)	8150.1 5447.4	0.3292 97.3 97.3	47.6927 [50.0000]	95.4%			
PFNA	(463.0 / 419.0) 32291769 (463.0 / 169.0) 7173063	(7.50 , 1.00) (0.00 , N/A , 0.1)	8163.1 19862.0	0.2221 94.9 94.9	48.4001 [50.0000]	96.8%			
PFDA	(513.0 / 469.0) 34749046 (513.0 / 169.0) 4007853	(8.06 , 1.00) (0.00 , N/A , 0.0)	1684.8 1564.4	0.1153 93.1 93.1	46.0761 [50.0000]	92.2%			
PFUnA	(563.0 / 519.0) 28504604 (563.0 / 169.0) 3359511	(8.56 , 1.00) (0.00 , N/A , 0.0)	1694.2 1225.1	0.1179 109.5 109.5	49.3643 [50.0000]	98.7%			
PFDoA	(613.0 / 569.0) 27250289 (613.0 / 169.0) 4441585	(8.87 , 1.00) (0.00 , N/A , -0.1)	2228.5 2230.0	0.1630 102.6 102.6	50.4535 [50.0000]	100.9%			
PFTTrDA	(663.0 / 619.0) 21050713 (663.0 / 169.0) 5464375	(9.08 , 1.02) (N/A , 0.01 , 0.0)	2417.7 2040.2	0.2596 95.6 95.6	40.5032 [50.0000]	81.0%			
PFTeDA	(713.0 / 669.0) 27069891 (713.0 / 169.0) 5799254	(9.25 , 1.00) (0.00 , N/A , 0.1)	2317.5 1573.0	0.2142 98.0 98.0	46.3137 [50.0000]	92.6%			

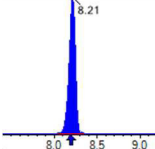
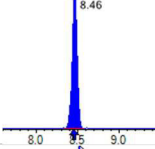
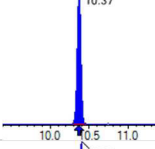
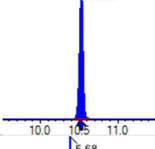
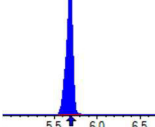
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) 41464583 (299.0 / 99.0) 27138490	(5.34 , 1.00) (0.00 , N/A , 0.1)	1925.6 2029.2	0.6545 104.0 104.0	39.2671 [44.2367]	88.8%			
PFPeS	(349.0 / 80.0) 71313453 (349.0 / 99.0) 31482926	(6.25 , 0.89) (N/A , 0.00 , -0.1)	10868.4 16838.6	0.4415 132.0 132.0	35.5932 [46.9191]	75.9%			
PFHxS	(399.0 / 80.0) 74332419 (399.0 / 99.0) 26768666	(7.02 , 1.00) (-0.01 , N/A , -0.1)	6106.9 13960.2	0.3601 112.5 112.5	42.2843 [45.5491]	92.8%			
PFHpS	(449.0 / 80.0) 74019843 (449.0 / 99.0) 24078257	(7.67 , 0.93) (N/A , 0.00 , -0.1)	8041.9 15002.9	0.3253 119.8 119.8	41.4790 [47.5703]	87.2%			
PFOS	(499.0 / 80.0) 90637354 (499.0 / 99.0) 21761174	(8.25 , 1.00) (0.00 , N/A , -0.4)	378.0 760.2	0.2401 109.2 109.2	41.4414 [46.3746]	89.4%			
PFNS	(549.0 / 80.0) 75048845 (549.0 / 99.0) 23194081	(8.71 , 1.06) (N/A , 0.01 , -0.2)	5666.8 10381.3	0.3091 128.4 128.4	34.2912 [47.9943]	71.4%			
PFDS	(599.0 / 80.0) N/A (599.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000 [48.1553]	N/A%			QC,
PFDoS	(699.0 / 80.0) 48525572 (699.0 / 99.0) 11072448	(9.32 , 1.13) (N/A , 0.01 , 0.0)	2124.1 2078.2	0.2282 116.8 116.8	43.2169 [48.4781]	89.1%			
4:2FTS	(327.0 / 307.0) 56217314 (327.0 / 81.0) 40288613	(5.13 , 1.00) (0.00 , N/A , 0.0)	2443.0 2574.8	0.7167 103.4 103.4	136.3114 [186.9055]	72.9%			
6:2FTS	(427.0 / 407.0) 38685986 (427.0 / 81.0) 32273693	(6.62 , 1.00) (0.00 , N/A , -0.1)	2477.0 2863.1	0.8342 98.5 98.5	173.4594 [189.8085]	91.4%			
8:2FTS	(527.0 / 507.0) 36938977 (527.0 / 81.0) 30134311	(7.80 , 1.00) (-0.01 , N/A , -0.2)	1880.3 2581.3	0.8158 100.0 100.0	172.6946 [191.6577]	90.1%			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) N/A (498.0 / 478.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000 [50.0000]	N/A%			QC,
NMeFOSA	(512.0 / 219.0) N/A (512.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000 [200.0000]	N/A%			QC,
NEIFOSA	(526.0 / 219.0) 87451751 (526.0 / 169.0) 93054225	(10.57 , 1.00) (-0.02 , N/A , 0.1)	13013.4 15860.2	1.0641 84.0 84.0	150.0336 [200.0000]	75.0%			
NMeFOSAA	(570.0 / 419.0) 12051539 (570.0 / 483.0) 5674644	(8.21 , 1.00) (0.00 , N/A , -0.1)	3220.1 665.3	0.4709 96.0 96.0	46.3670 [50.0000]	92.7%			
NEIFOSAA	(584.0 / 419.0) 10932768 (584.0 / 526.0) 6296395	(8.47 , 1.00) (0.01 , N/A , 0.0)	3458.5 2035.7	0.5759 101.9 101.9	49.6670 [50.0000]	99.3%			
NMeFOSE	(616.0 / 59.0) 45091490	(10.36 , 1.00) (-0.01 , N/A , 0.0)	3257.4	N/A 0.0 0.0	182.7506 [200.0000]	91.4%			
NEtFOSE	(630.0 / 59.0) 45200872	(10.53 , 1.00) (0.00 , N/A , 0.0)	1109.6	N/A 0.0 0.0	175.3347 [200.0000]	87.7%			
HFPO-DA	(285.0 / 169.0) 31842579 (285.0 / 185.0) 67618370	(5.68 , 1.00) (0.00 , N/A , 0.2)	2063.3 2680.9	2.1235 77.2 77.2	90.2171 [100.0000]	90.2%			
ADONA	(377.0 / 85.0) 86827157 (377.0 / 251.0) 11259712	(6.46 , 1.14) (N/A , 0.00 , -0.4)	1997.8 2166.4	0.1297 148.1 148.1	66.2830 [94.2700]	70.3%			
9CI-Pf3ONS	(531.0 / 351.0) N/A (533.0 / 353.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000 [93.3254]	N/A%			QC,
11CI-PF3OUDS	(631.0 / 451.0) N/A (633.0 / 453.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000 [94.3208]	N/A%			QC,

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) 3159042 (241.0 / 117.0) 5310509	(4.01 , 0.90) (N/A , -0.01 , 0.1)	1859.8 1969.3	1.6811 96.9 96.9	234.6937 [200.0000]	117.3%			
5:3FTCA	(341.0 / 236.7) 19519906 (341.0 / 217.0) 31848733	(5.90 , 1.10) (N/A , 0.00 , 0.0)	2417.2 2188.2	1.6316 97.0 97.0	216.5232 [200.0000]	108.3%			
7:3FTCA	(441.0 / 317.0) 36282186 (441.0 / 337.0) 31617035	(7.80 , 1.45) (N/A , -0.01 , 0.0)	1152.0 1207.2	0.8714 101.5 101.5	223.8599 [200.0000]	111.9%			
PFEESA	(315.0 / 135.0) 54624424 (315.0 / 83.0) 18021083	(5.77 , 1.07) (N/A , -0.01 , -0.1)	1702.1 2192.1	0.3299 117.2 117.2	69.7022 [89.2459]	78.1%			
PFMPA	(229.0 / 85.0) 11599193	(3.84 , 0.86) (N/A , -0.01 , 0.0)	2815.3	N/A 0.0 0.0	99.5344 [100.0000]	99.5%			
PFMBA	(279.0 / 85.0) 37911087	(4.77 , 1.07) (N/A , -0.01 , 0.0)	2255.9	N/A 0.0 0.0	94.9588 [100.0000]	95.0%			
NFDHA	(295.0 / 201.0) 31356860 (295.0 / 85.0) 31765052	(5.29 , 0.98) (N/A , -0.01 , 0.1)	2016.4 2162.5	1.0130 97.0 97.0	89.3954 [100.0000]	89.4%			
13C3_PFBA_IIS	(216.0 / 172.0) 231462	(3.45 , N/A) (N/A , -0.01 , N/A)	811.3	N/A	0.6827 [1.0000]	68.3% { 70.2% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 519504	(5.39 , N/A) (N/A , -0.01 , N/A)	1055.0	N/A	0.8735 [1.0000]	87.3% { 93.5% }			
13C4_PFOA_IIS	(417.0 / 372.0) 702253	(6.89 , N/A) (N/A , 0.01 , N/A)	1327.8	N/A	0.8633 [1.0000]	86.3% { 93.6% }			
13C5_PFNA_IIS	(468.0 / 423.0) 682549	(7.50 , N/A) (N/A , 0.01 , N/A)	766.9	N/A	0.8938 [1.0000]	89.4% { 89.6% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 662827	(8.06 , N/A) (N/A , 0.01 , N/A)	821.6	N/A	0.9233 [1.0000]	92.3% { 96.3% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 1117021	(7.02 , N/A) (N/A , 0.01 , N/A)	1053.0	N/A	0.9070 [1.0000]	90.7% { 93.0% }			
13C4_PFOS_IIS	(503.0 / 79.9) 1516588	(8.26 , N/A) (N/A , 0.02 , N/A)	1218.2	N/A	0.8545 [1.0000]	85.4% { 87.5% }			
13C4_PFBA_EIS	(217.0 / 172.0) 2165391	(3.45 , N/A) (N/A , -0.01 , N/A)	2598.1	N/A	7.8676 [8.0000]	98.3% { 73.9% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 2069084	(4.46 , N/A) (N/A , -0.01 , N/A)	1685.2	N/A	3.5744 [4.0000]	89.4% { 84.9% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 1351535	(5.38 , N/A) (N/A , -0.01 , N/A)	1418.5	N/A	1.9147 [2.0000]	95.7% { 88.1% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 1312625	(6.18 , N/A) (N/A , 0.00 , N/A)	2301.2	N/A	1.9939 [2.0000]	99.7% { 92.9% }			
13C8_PFOA_EIS	(421.0 / 376.0) 1511777	(6.89 , N/A) (N/A , 0.01 , N/A)	1768.5	N/A	2.0113 [2.0000]	100.6% { 90.7% }			
13C9_PFNA_EIS	(472.0 / 427.0) 706321	(7.50 , N/A) (N/A , 0.01 , N/A)	1268.4	N/A	1.0384 [1.0000]	103.8% { 99.6% }			
13C6_PFDA_EIS	(519.0 / 474.0) 760656	(8.06 , N/A) (N/A , 0.02 , N/A)	788.6	N/A	0.9915 [1.0000]	99.1% { 103.6% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 656647	(8.56 , N/A) (N/A , 0.01 , N/A)	1572.0	N/A	0.8640 [1.0000]	86.4% { 87.2% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 619074	(8.87, N/A) (N/A, 0.01, N/A)	2194.8	N/A	0.8782 [1.0000]	87.8% { 84.0% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 635660	(9.25, N/A) (N/A, 0.01, N/A)	930.2	N/A	0.8675 [1.0000]	86.7% { 88.4% }			
13C3_PFBs_EIS	(302.0 / 80.0) 3714321	(5.34, N/A) (N/A, -0.01, N/A)	1514.1	N/A	1.6455 [2.0000]	82.3% { 81.6% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 2231971	(7.02, N/A) (N/A, 0.01, N/A)	1384.5	N/A	1.8926 [2.0000]	94.6% { 94.3% }			
13C8_PFOS_EIS	(507.0 / 80.0) 3963601	(8.25, N/A) (N/A, 0.01, N/A)	242.4	N/A	1.9893 [2.0000]	99.5% { 92.8% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 584471	(5.13, N/A) (N/A, -0.01, N/A)	1097.7	N/A	4.2275 [4.0000]	105.7% { 112.7% }			
13C2_6:2FTS_EIS	(429.0 / 81.0) 683829	(6.63, N/A) (N/A, 0.01, N/A)	865.7	N/A	4.3869 [4.0000]	109.7% { 115.7% }			
13C2_8:2FTS_EIS	(529.0 / 81.0) 762145	(7.81, N/A) (N/A, 0.02, N/A)	672.8	N/A	4.6236 [4.0000]	115.6% { 114.8% }			
13C8_PFOsa_EIS	(506.0 / 78.0) 6365767	(9.84, N/A) (N/A, 0.01, N/A)	2313.5	N/A	1.7345 [2.0000]	86.7% { 81.7% }			
D3_NMeFOsa_EIS	(515.0 / 169.0) 1747358	(10.43, N/A) (N/A, 0.01, N/A)	1665.9	N/A	2.2172 [2.0000]	110.9% { 101.6% }			
D5_NEtFOsa_EIS	(531.0 / 169.0) 1200000	(10.59, N/A) (N/A, 0.01, N/A)	2164.9	N/A	1.7390 [2.0000]	86.9% { 76.7% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1216452	(8.21 , N/A) (N/A , 0.02 , N/A)	1189.8	N/A	4.3503 [4.0000]	108.8% { 94.6% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 1010035	(8.46 , N/A) (N/A , 0.01 , N/A)	3137.2	N/A	4.2418 [4.0000]	106.0% { 104.4% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 4709587	(10.37 , N/A) (N/A , 0.01 , N/A)	1590.2	N/A	17.5531 [20.0000]	87.8% { 77.8% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 5102587	(10.53 , N/A) (N/A , 0.01 , N/A)	1348.2	N/A	16.6257 [20.0000]	83.1% { 74.5% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 3444188	(5.68 , N/A) (N/A , -0.01 , N/A)	1774.4	N/A	7.7652 [8.0000]	97.1% { 92.1% }			

SECOND-SOURCE CALIBRATION VERIFICATION**EPA 1633****Laboratory:** APPL, LLC**SDG:****Client:** AECOM**Project:** Red Hill AFFF Assessment Sampling**Calibration:** 2309009**Laboratory ID:** SC00840-SCV1**Sequence:** SC00840**Standard ID:** 23B0089

ANALYTE	EXPECTED (ng/mL)	FOUND (ng/mL)	% DRIFT	QC LIMIT
PFBA	8.00	7.55	-5.6	30.00
PFPEA	4.00	3.74	-6.6	30.00
PFHXA	2.00	1.88	-5.8	30.00
PFHPA	2.00	1.74	-12.9	30.00
PFOA	2.00	1.83	-8.7	30.00
PFNA	2.00	1.89	-5.5	30.00
PFDA	2.00	1.79	-10.7	30.00
PFUnA	2.00	1.79	-10.7	30.00
PFDOA	2.00	1.80	-10.0	30.00
PFTRDA	2.00	1.84	-7.8	30.00
PFTEDA	2.00	1.67	-16.3	30.00
PFBS	1.77	1.65	-6.7	30.00
PFPEs	1.88	1.93	2.7	30.00
PFHXS	1.83	1.74	-5.2	30.00
PFHPS	1.91	1.77	-7.5	30.00
PFOS	1.86	1.64	-11.9	30.00
PFNS	1.92	1.77	-7.8	30.00
PFDS	1.93	1.74	-9.7	30.00
PFDOS	1.94	1.71	-11.7	30.00
4:2FTS	7.50	7.00	-6.7	30.00
6:2FTS	7.60	6.50	-14.4	30.00
8:2FTS	7.68	7.87	2.5	30.00
PFOSA	2.00	2.04	1.9	30.00
NMeFOSA	8.00	7.44	-7.0	30.00
NEtFOSA	8.00	6.78	-15.3	30.00
NMeFOSAA	2.00	1.91	-4.6	30.00
NEtFOSAA	2.00	1.89	-5.4	30.00

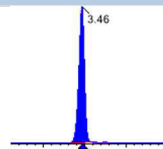
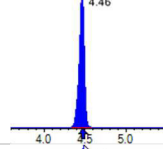
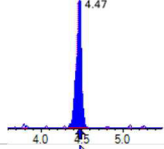
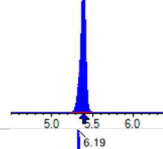
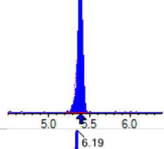
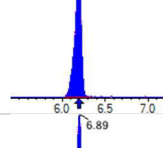
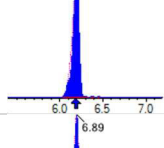
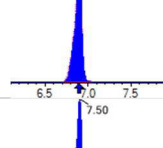
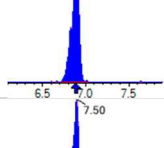
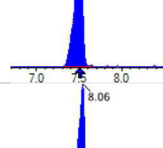
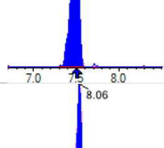
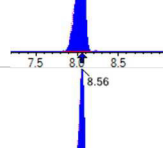
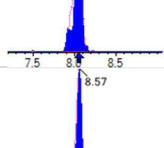
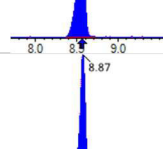
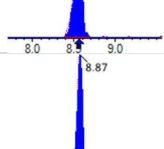
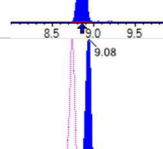
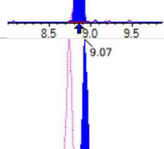
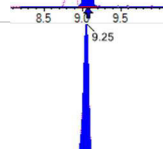
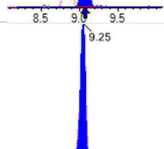
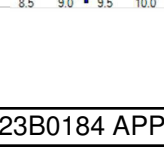
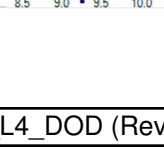
SECOND-SOURCE CALIBRATION VERIFICATION**EPA 1633****Laboratory:** APPL, LLC**SDG:****Client:** AECOM**Project:** Red Hill AFFF Assessment Sampling**Calibration:** 2309009**Laboratory ID:** SC00840-SCV1**Sequence:** SC00840**Standard ID:** 23B0089

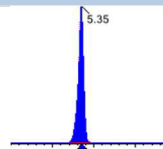
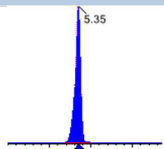
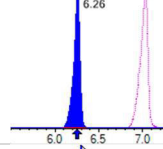
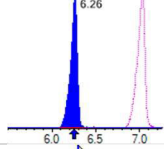
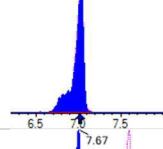
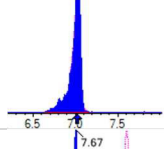
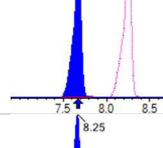
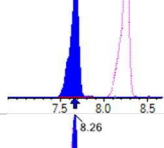
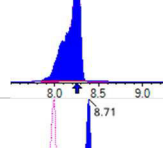
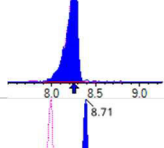
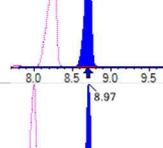
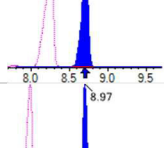
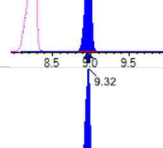
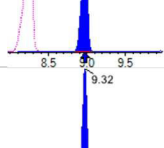
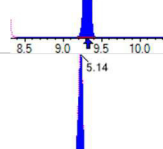
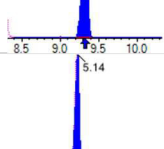
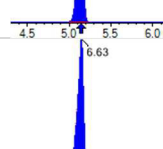
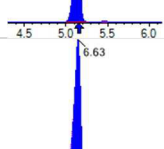
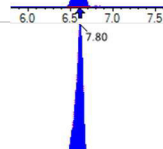
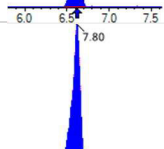
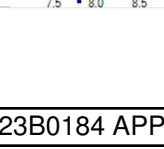
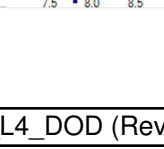
NMeFOSE	8.00	7.81	-2.4	30.00
NEtFOSE	8.00	7.88	-1.6	30.00
HFPO-DA	4.00	3.72	-6.9	30.00
ADONA	3.78	3.86	2.1	30.00
PFEESA	3.56	3.22	-9.5	30.00
PFMPA	4.00	3.52	-12.0	30.00
PFMBA	4.00	3.63	-9.3	30.00
NFDHA	4.00	3.74	-6.5	30.00
9CL-PF3ONS	3.74	3.86	3.1	30.00
11CL-PF3OUDS	3.78	3.61	-4.6	30.00
3:3FTCA	8.00	7.09	-11.4	30.00
5:3FTCA	8.00	6.99	-12.6	30.00
7:3FTCA	8.00	6.76	-15.5	30.00
13C4-PFBA	8.00	8.22	2.8	30.00
13C5-PFPEA	4.00	4.35	8.8	30.00
13C5-PFHXA	2.00	2.18	9.2	30.00
13C4-PFHFA	2.00	2.09	4.6	30.00
13C8-PFOA	2.00	2.07	3.4	30.00
13C9-PFNA	1.00	1.01	1.5	30.00
13C6-PFDA	1.00	1.06	6.3	30.00
13C7-PFUnA	1.00	1.11	10.8	30.00
13C2-PFDOA	1.00	1.08	7.6	30.00
13C2-PFTEDA	1.00	1.04	4.4	30.00
13C3-PFBS	2.00	2.04	2.0	30.00
13C3-PFHXS	2.00	1.96	-2.2	30.00
13C8-PFOS	2.00	2.19	9.5	30.00
13C2-4:2FTS	4.00	3.71	-7.2	30.00
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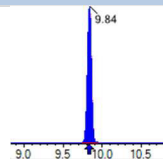
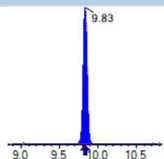
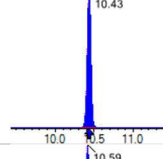
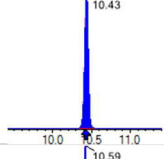
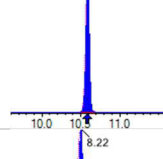
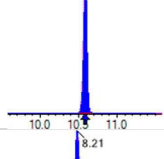
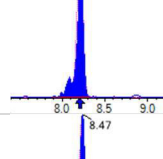
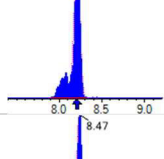
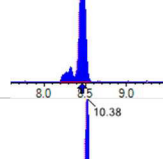
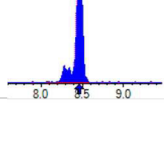
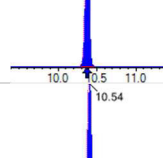
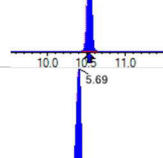
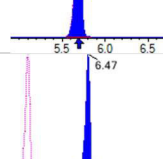
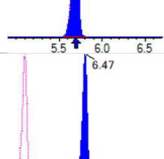
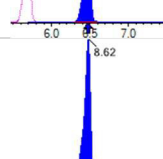
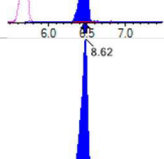
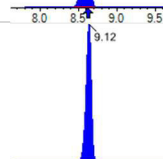
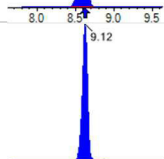
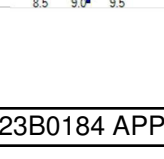
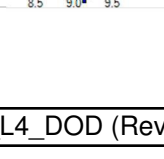
SECOND-SOURCE CALIBRATION VERIFICATION**EPA 1633****Laboratory:** APPL, LLC**SDG:****Client:** AECOM**Project:** Red Hill AFFF Assessment Sampling**Calibration:** 2309009**Laboratory ID:** SC00840-SCV1**Sequence:** SC00840**Standard ID:** 23B0089

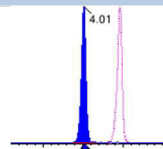
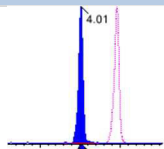
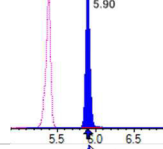
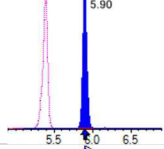
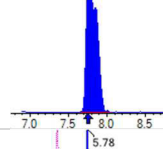
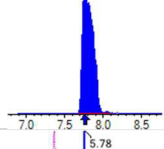
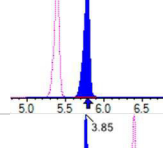
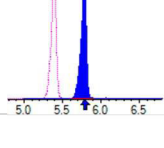
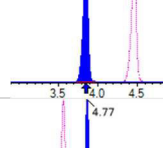
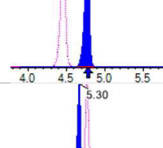
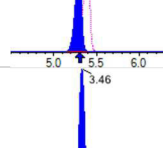
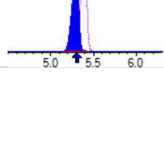
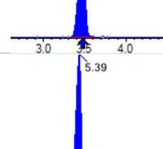
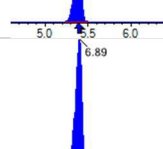
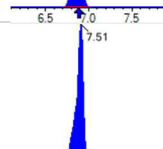
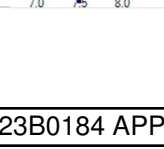
13C8-PFOSA	2.00	2.08	4.0	30.00
D3-NMEFOSA	2.00	2.16	7.9	30.00
D5-NETFOSA	2.00	2.24	12.1	30.00
D3-NMEFOSAA	4.00	4.43	10.7	30.00
D5-NETFOSAA	4.00	4.26	6.5	30.00
D7-NMEFOSE	20.0	22.4	12.2	30.00
D9-NETFOSSE	20.0	21.6	8.1	30.00
13C3-HFPO-DA	8.00	8.61	7.6	30.00

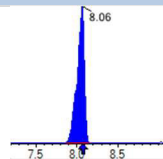
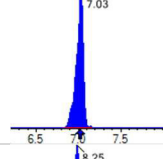
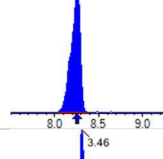
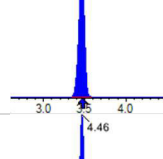
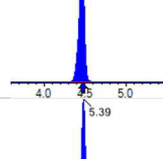
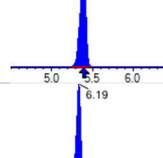
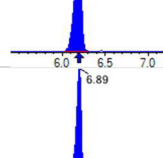
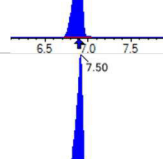
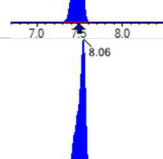
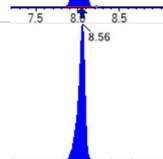
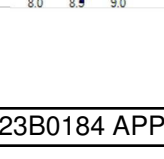
* Values outside of QC limits

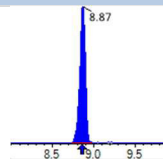
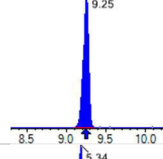
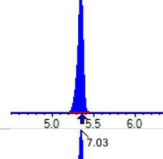
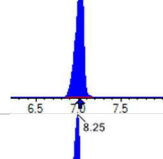
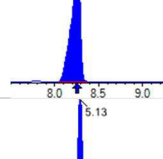
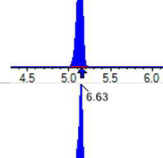
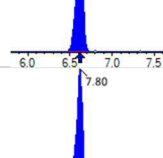
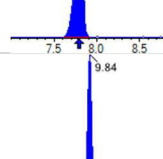
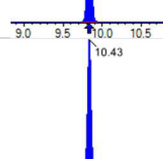
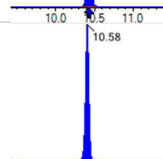
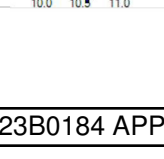
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) 2817870	(3.46 , 1.00) (0.00 , N/A , 0.0)	181.9	N/A 0.0 0.0	7.5535 [8.0000]	94.4%			
PFPeA	(263.0 / 219.0) 2335135 (263.0 / 69.0) 31036	(4.46 , 1.00) (0.00 , N/A , -0.3)	1911.5 382.7	0.0133 111.5 111.5	3.7379 [4.0000]	93.4%			
PFHxA	(313.0 / 269.0) 1526807 (313.0 / 119.0) 165903	(5.39 , 1.00) (0.00 , N/A , 0.1)	3392.4 200916.5	0.1087 116.1 116.1	1.8850 [2.0000]	94.2%			
PFHpA	(363.0 / 319.0) 1206369 (363.0 / 169.0) 359839	(6.19 , 1.00) (0.00 , N/A , -0.5)	10032.2 3701.9	0.2983 94.5 94.5	1.7427 [2.0000]	87.1%			
PFOA	(413.0 / 369.0) 1638255 (413.0 / 169.0) 536507	(6.89 , 1.00) (0.00 , N/A , 0.1)	2799.8 3657323.0	0.3275 96.8 96.8	1.8258 [2.0000]	91.3%			
PFNA	(463.0 / 419.0) 1410247 (463.0 / 169.0) 297200	(7.50 , 1.00) (0.00 , N/A , 0.1)	1981.8 910.0	0.2107 90.0 90.0	1.8895 [2.0000]	94.5%			
PFDA	(513.0 / 469.0) 1598327 (513.0 / 169.0) 194832	(8.06 , 1.00) (0.00 , N/A , -0.1)	1078.6 1072.3	0.1219 98.4 98.4	1.7866 [2.0000]	89.3%			
PFUnA	(563.0 / 519.0) 1463692 (563.0 / 169.0) 196601	(8.56 , 1.00) (0.00 , N/A , -0.4)	753.4 582.6	0.1343 124.8 124.8	1.7853 [2.0000]	89.3%			
PFDoA	(613.0 / 569.0) 1318905 (613.0 / 169.0) 210879	(8.87 , 1.00) (0.00 , N/A , -0.1)	1912.1 972.6	0.1599 100.6 100.6	1.8004 [2.0000]	90.0%			
PFTTrDA	(663.0 / 619.0) 1300053 (663.0 / 169.0) 359088	(9.08 , 1.02) (N/A , 0.01 , 0.3)	1205.8 628.4	0.2762 101.7 101.7	1.8443 [2.0000]	92.2%			
PFTeDA	(713.0 / 669.0) 1303207 (713.0 / 169.0) 311126	(9.25 , 1.00) (0.00 , N/A , 0.1)	1353.5 652.0	0.2387 109.2 109.2	1.6742 [2.0000]	83.7%			

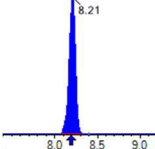
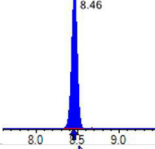
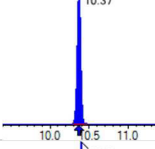
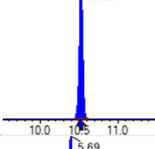
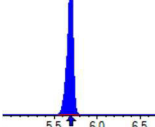
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) 2478161 (299.0 / 99.0) 1537615	(5.35 , 1.00) (0.00 , N/A , -0.1)	1939.1 1818.8	0.6205 98.6 98.6	1.6516 [1.7695]	93.3%			
PFPeS	(349.0 / 80.0) 4587368 (349.0 / 99.0) 1500852	(6.26 , 0.89) (N/A , 0.00 , -0.1)	2771577.9 711664.9	0.3272 97.8 97.8	1.9312 [1.8768]	102.9%			
PFHxS	(399.0 / 80.0) 3616887 (399.0 / 99.0) 1165097	(7.03 , 1.00) (0.00 , N/A , 0.0)	20772.6 1849.0	0.3221 100.7 100.7	1.7355 [1.8220]	95.3%			
PFHpS	(449.0 / 80.0) 4058678 (449.0 / 99.0) 1126402	(7.67 , 0.93) (N/A , 0.01 , -0.1)	143656.3 2081.2	0.2775 102.2 102.2	1.7669 [1.9028]	92.9%			
PFOS	(499.0 / 80.0) 4615403 (499.0 / 99.0) 978647	(8.25 , 1.00) (0.00 , N/A , -0.3)	331.4 311.6	0.2120 96.4 96.4	1.6394 [1.8550]	88.4%			
PFNS	(549.0 / 80.0) 4988466 (549.0 / 99.0) 1190660	(8.71 , 1.06) (N/A , 0.01 , 0.0)	9124.3 932984.3	0.2387 99.2 99.2	1.7708 [1.9198]	92.2%			
PFDS	(599.0 / 80.0) 5454478 (599.0 / 99.0) 1158548	(8.97 , 1.09) (N/A , 0.01 , -0.1)	1979.6 821.9	0.2124 104.9 104.9	1.7422 [1.9262]	90.4%			
PFDoS	(699.0 / 80.0) 2475315 (699.0 / 99.0) 526689	(9.32 , 1.13) (N/A , 0.01 , 0.0)	1521.1 1424.2	0.2128 108.9 108.9	1.7127 [1.9391]	88.3%			
4:2FTS	(327.0 / 307.0) 2907985 (327.0 / 81.0) 1965804	(5.14 , 1.00) (0.00 , N/A , 0.0)	2226.5 1411.0	0.6760 97.6 97.6	7.0011 [7.4762]	93.6%			
6:2FTS	(427.0 / 407.0) 1552065 (427.0 / 81.0) 1371714	(6.63 , 1.00) (0.00 , N/A , -0.1)	1418.0 1901.4	0.8838 104.3 104.3	6.5032 [7.5923]	85.7%			
8:2FTS	(527.0 / 507.0) 1662385 (527.0 / 81.0) 1252728	(7.80 , 1.00) (0.00 , N/A , 0.2)	1275.3 1311.9	0.7536 92.4 92.4	7.8691 [7.6663]	102.6%			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) 7066592 (498.0 / 478.0) 144068	(9.84 , 1.00) (0.00 , N/A , 0.1)	2727.2 1649.4	0.0204 103.8 103.8	2.0375 [2.0000]	101.9%			
NMeFOSA	(512.0 / 219.0) 6348828 (512.0 / 169.0) 4438785	(10.43 , 1.00) (0.00 , N/A , 0.0)	7203.0 6421.7	0.6992 84.6 84.6	7.4372 [8.0000]	93.0%			
NEIFOSA	(526.0 / 219.0) 5955989 (526.0 / 169.0) 6618343	(10.59 , 1.00) (0.01 , N/A , 0.0)	8894.1 8638.3	1.1112 87.8 87.8	6.7795 [8.0000]	84.7%			
NMeFOSAA	(570.0 / 419.0) 590048 (570.0 / 483.0) 273023	(8.22 , 1.00) (0.01 , N/A , 0.3)	1043.7 455.4	0.4627 94.4 94.4	1.9072 [2.0000]	95.4%			
NEIFOSAA	(584.0 / 419.0) 488870 (584.0 / 526.0) 289353	(8.47 , 1.00) (0.01 , N/A , -0.1)	1983.4 1109.1	0.5919 104.8 104.8	1.8919 [2.0000]	94.6%			
NMeFOSE	(616.0 / 59.0) 2878588	(10.38 , 1.00) (0.01 , N/A , 0.0)	3202.3	N/A 0.0 0.0	7.8064 [8.0000]	97.6%			
NEtFOSE	(630.0 / 59.0) 3087305	(10.54 , 1.00) (0.01 , N/A , 0.0)	2386.3	N/A 0.0 0.0	7.8753 [8.0000]	98.4%			
HFPO-DA	(285.0 / 169.0) 1639551 (285.0 / 185.0) 4211531	(5.69 , 1.00) (0.00 , N/A , -0.1)	1429.0 1885.9	2.5687 93.4 93.4	3.7229 [4.0000]	93.1%			
ADONA	(377.0 / 85.0) 6308450 (377.0 / 251.0) 598590	(6.47 , 1.14) (N/A , 0.01 , 0.0)	1940.2 941.8	0.0949 108.4 108.4	3.8596 [3.7708]	102.4%			
9CI-Pf3ONS	(531.0 / 351.0) 14787593 (533.0 / 353.0) 4359706	(8.62 , 1.52) (N/A , 0.01 , 0.0)	1980.7 1616.1	0.2948 100.1 100.1	3.8577 [3.7330]	103.3%			
11CI-PF3OUDS	(631.0 / 451.0) 6798708 (633.0 / 453.0) 2101242	(9.12 , 1.60) (N/A , 0.01 , 0.0)	1885.1 1241.0	0.3091 100.1 100.1	3.6075 [3.7728]	95.6%			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) 130823 (241.0 / 117.0) 204282	(4.01 , 0.90) (N/A , 0.00 , -0.1)	806.5 803.7	1.5615 90.0 90.0	7.0910 [8.0000]	88.6%			
5:3FTCA	(341.0 / 236.7) 809463 (341.0 / 217.0) 1316539	(5.90 , 1.09) (N/A , 0.00 , 0.0)	1308.1 1596.9	1.6264 96.7 96.7	6.9910 [8.0000]	87.4%			
7:3FTCA	(441.0 / 317.0) 1407442 (441.0 / 337.0) 1292691	(7.80 , 1.45) (N/A , 0.01 , 0.4)	531.4 654.7	0.9185 106.9 106.9	6.7612 [8.0000]	84.5%			
PFEESA	(315.0 / 135.0) 3241322 (315.0 / 83.0) 901031	(5.78 , 1.07) (N/A , 0.00 , 0.0)	2229.7 1038.2	0.2780 98.7 98.7	3.2203 [3.5698]	90.2%			
PFMPA	(229.0 / 85.0) 562290	(3.85 , 0.86) (N/A , 0.00 , 0.0)	1713.3	N/A 0.0 0.0	3.5204 [4.0000]	88.0%			
PFMBA	(279.0 / 85.0) 1984924	(4.77 , 1.07) (N/A , -0.01 , 0.0)	1394.5	N/A 0.0 0.0	3.6274 [4.0000]	90.7%			
NFDHA	(295.0 / 201.0) 1684250 (295.0 / 85.0) 1661347	(5.30 , 0.98) (N/A , 0.00 , 0.1)	1684.3 1820.0	0.9864 94.5 94.5	3.7385 [4.0000]	93.5%			
13C3_PFBa_IIS	(216.0 / 172.0) 361028	(3.46 , N/A) (N/A , 0.00 , N/A)	994.7	N/A	1.0649 [1.0000]	106.5% { 109.5% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 584798	(5.39 , N/A) (N/A , 0.00 , N/A)	1545.0	N/A	0.9832 [1.0000]	98.3% { 105.2% }			
13C4_PFOA_IIS	(417.0 / 372.0) 841453	(6.89 , N/A) (N/A , 0.01 , N/A)	1903.7	N/A	1.0344 [1.0000]	103.4% { 112.2% }			
13C5_PFNA_IIS	(468.0 / 423.0) 781350	(7.51 , N/A) (N/A , 0.02 , N/A)	1032.6	N/A	1.0232 [1.0000]	102.3% { 102.6% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 733505	(8.06 , N/A) (N/A , 0.01 , N/A)	1314.5	N/A	1.0218 [1.0000]	102.2% { 106.6% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 1280781	(7.03 , N/A) (N/A , 0.01 , N/A)	1417.7	N/A	1.0400 [1.0000]	104.0% { 106.7% }			
13C4_PFOS_IIS	(503.0 / 79.9) 1773012	(8.25 , N/A) (N/A , 0.01 , N/A)	1007.3	N/A	0.9990 [1.0000]	99.9% { 102.3% }			
13C4_PFBA_EIS	(217.0 / 172.0) 3529057	(3.46 , N/A) (N/A , 0.00 , N/A)	2821.1	N/A	8.2206 [8.0000]	102.8% { 120.5% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 2835943	(4.46 , N/A) (N/A , -0.01 , N/A)	1925.7	N/A	4.3522 [4.0000]	108.8% { 116.3% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 1735856	(5.39 , N/A) (N/A , 0.00 , N/A)	1455.4	N/A	2.1846 [2.0000]	109.2% { 113.1% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 1550387	(6.19 , N/A) (N/A , 0.00 , N/A)	1861.6	N/A	2.0921 [2.0000]	104.6% { 109.7% }			
13C8_PFOA_EIS	(421.0 / 376.0) 1862416	(6.89 , N/A) (N/A , 0.01 , N/A)	1601.0	N/A	2.0679 [2.0000]	103.4% { 111.8% }			
13C9_PFNA_EIS	(472.0 / 427.0) 790147	(7.50 , N/A) (N/A , 0.01 , N/A)	1916.9	N/A	1.0148 [1.0000]	101.5% { 111.4% }			
13C6_PFDA_EIS	(519.0 / 474.0) 902311	(8.06 , N/A) (N/A , 0.02 , N/A)	750.9	N/A	1.0628 [1.0000]	106.3% { 122.9% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 932343	(8.56 , N/A) (N/A , 0.01 , N/A)	1110.0	N/A	1.1085 [1.0000]	110.8% { 123.9% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 839650	(8.87 , N/A) (N/A , 0.01 , N/A)	737.4	N/A	1.0764 [1.0000]	107.6% { 113.9% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 846527	(9.25 , N/A) (N/A , 0.01 , N/A)	888.5	N/A	1.0439 [1.0000]	104.4% { 117.7% }			
13C3_PFBs_EIS	(302.0 / 80.0) 5277897	(5.34 , N/A) (N/A , -0.01 , N/A)	2357.4	N/A	2.0392 [2.0000]	102.0% { 116.0% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 2646125	(7.03 , N/A) (N/A , 0.01 , N/A)	1653.0	N/A	1.9569 [2.0000]	97.8% { 111.7% }			
13C8_PFOS_EIS	(507.0 / 80.0) 5101924	(8.25 , N/A) (N/A , 0.01 , N/A)	920.4	N/A	2.1903 [2.0000]	109.5% { 119.4% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 588640	(5.13 , N/A) (N/A , -0.01 , N/A)	1488.6	N/A	3.7132 [4.0000]	92.8% { 113.5% }			
13C2_6:2FTS_EIS	(429.0 / 81.0) 731765	(6.63 , N/A) (N/A , 0.01 , N/A)	1118.5	N/A	4.0942 [4.0000]	102.4% { 123.9% }			
13C2_8:2FTS_EIS	(529.0 / 81.0) 752730	(7.80 , N/A) (N/A , 0.01 , N/A)	977.5	N/A	3.9826 [4.0000]	99.6% { 113.3% }			
13C8_PFOsa_EIS	(506.0 / 78.0) 8926163	(9.84 , N/A) (N/A , 0.01 , N/A)	2503.2	N/A	2.0803 [2.0000]	104.0% { 114.6% }			
D3_NMeFOsa_EIS	(515.0 / 169.0) 1987938	(10.43 , N/A) (N/A , 0.01 , N/A)	2396.3	N/A	2.1577 [2.0000]	107.9% { 115.6% }			
D5_NEtFOsa_EIS	(531.0 / 169.0) 1808669	(10.58 , N/A) (N/A , 0.01 , N/A)	2456.6	N/A	2.2420 [2.0000]	112.1% { 115.7% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1447929	(8.21 , N/A) (N/A , 0.01 , N/A)	1062.9	N/A	4.4292 [4.0000]	110.7% { 112.5% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 1185674	(8.46 , N/A) (N/A , 0.01 , N/A)	6570.8	N/A	4.2592 [4.0000]	106.5% { 122.5% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 7038425	(10.37 , N/A) (N/A , 0.01 , N/A)	2368.7	N/A	22.4389 [20.0000]	112.2% { 116.3% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 7759341	(10.53 , N/A) (N/A , 0.01 , N/A)	1362.4	N/A	21.6257 [20.0000]	108.1% { 113.3% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 4297441	(5.69 , N/A) (N/A , 0.00 , N/A)	1853.2	N/A	8.6071 [8.0000]	107.6% { 114.9% }			

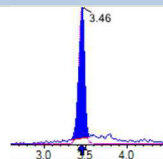
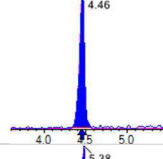
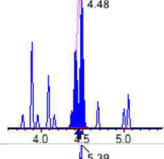
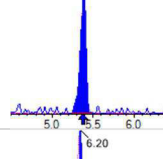
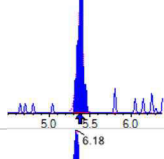
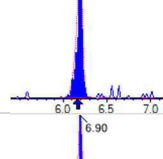
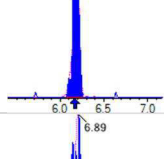
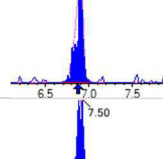
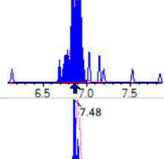
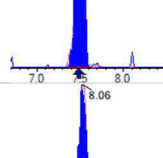
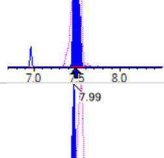
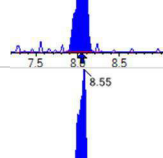
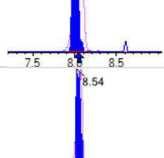
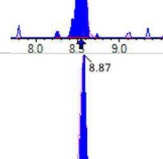
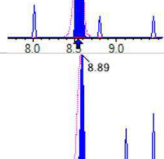
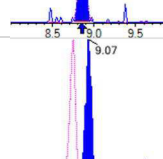
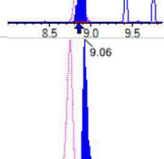
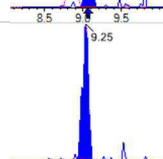
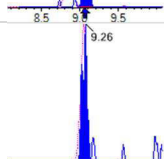
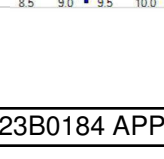
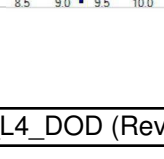
LOW-CONCENTRATION CALIBRATION VERIFICATION**EPA 1633****Laboratory:** APPL, LLC**SDG:****Client:** AECOM**Project:** Red Hill AFFF Assessment Sampling**Calibration:** 2309009**Laboratory ID:** SC00861-LCV1**Sequence:** SC00861**Standard ID:** 23B0080

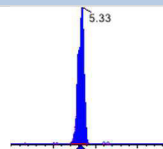
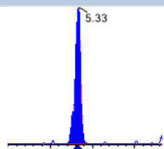
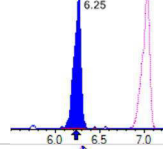
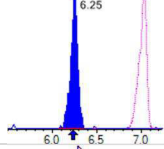
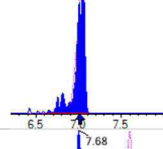
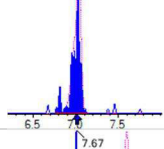
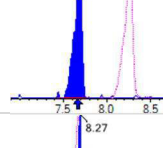
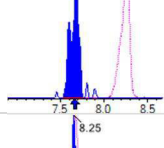
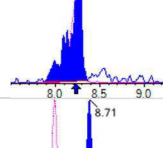
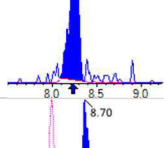
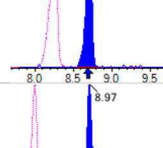
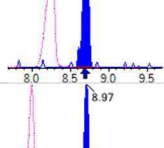
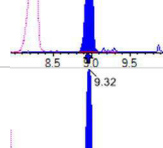
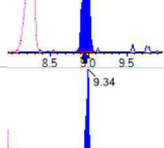
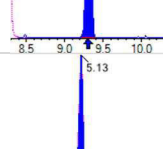
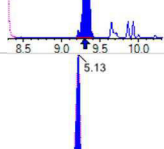
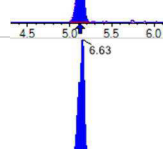
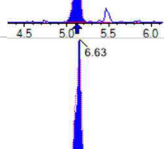
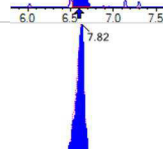
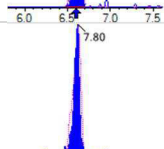
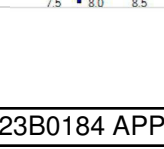
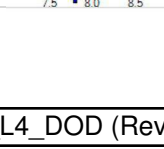
ANALYTE	EXPECTED (ng/mL)	FOUND (ng/mL)	% DRIFT	QC LIMIT
PFBA	0.400	0.432	8.0	30.00
PFPEA	0.200	0.219	9.7	30.00
PFHXA	0.100	0.103	2.9	30.00
PFHPA	0.100	0.117	16.9	30.00
PFOA	0.100	0.113	13.2	30.00
PFNA	0.100	0.105	4.9	30.00
PFDA	0.100	0.111	11.0	30.00
PFUnA	0.100	0.108	8.0	30.00
PFDOA	0.100	0.0884	-11.6	30.00
PFTRDA	0.100	0.100	0.3	30.00
PFTEDA	0.100	0.127	27.4	30.00
PFBS	0.0885	0.0976	10.3	30.00
PFPEs	0.0940	0.101	7.8	30.00
PFHXS	0.0915	0.0945	3.3	30.00
PFHPS	0.0955	0.0918	-3.9	30.00
PFOS	0.0930	0.111	18.9	30.00
PFNS	0.0960	0.0978	1.9	30.00
PFDS	0.0965	0.101	4.8	30.00
PFDOS	0.0970	0.117	20.6	30.00
4:2FTS	0.375	0.452	20.4	30.00
6:2FTS	0.380	0.531	39.9 *	30.00
8:2FTS	0.384	0.456	18.7	30.00
PFOSA	0.100	0.113	13.4	30.00
NMeFOSA	0.400	0.436	9.0	30.00
NEtFOSA	0.400	0.434	8.5	30.00
NMeFOSAA	0.100	0.123	22.7	30.00
NEtFOSAA	0.100	0.121	20.7	30.00
NMeFOSE	0.400	0.436	9.1	30.00

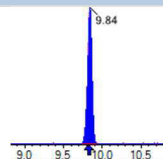
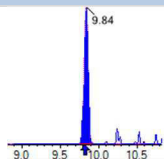
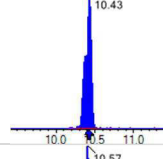
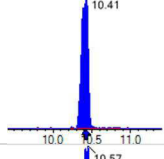
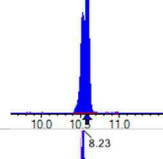
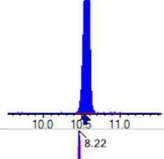
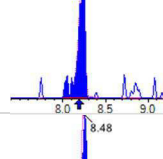
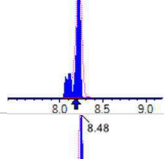
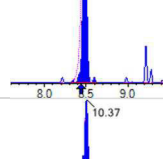
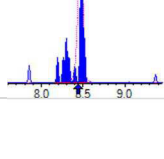
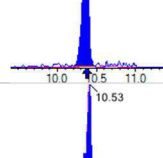
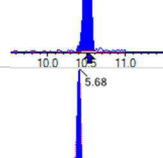
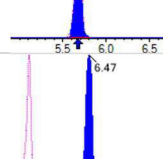
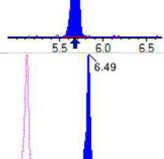
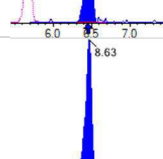
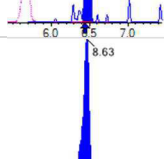
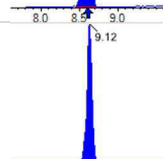
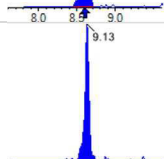
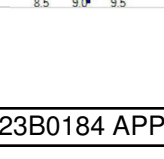
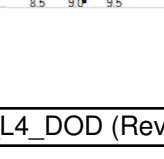
LOW-CONCENTRATION CALIBRATION VERIFICATION**EPA 1633****Laboratory:** APPL, LLC**SDG:****Client:** AECOM**Project:** Red Hill AFFF Assessment Sampling**Calibration:** 2309009**Laboratory ID:** SC00861-LCV1**Sequence:** SC00861**Standard ID:** 23B0080

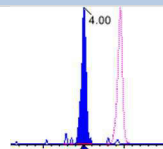
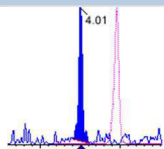
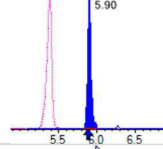
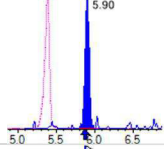
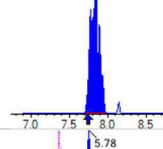
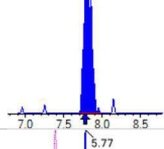
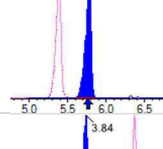
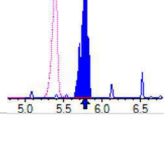
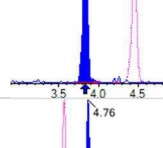
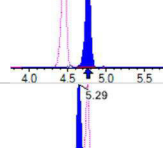
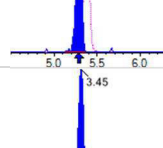
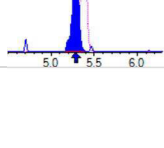
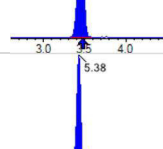
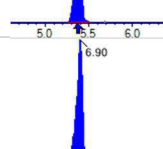
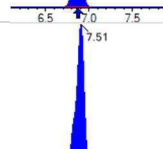
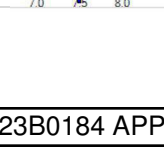
NEtFOSE	0.400	0.437	9.2	30.00
HFPO-DA	0.200	0.216	8.0	30.00
ADONA	0.189	0.208	9.9	30.00
PFEESA	0.178	0.192	7.8	30.00
PFMPA	0.200	0.189	-5.6	30.00
PFMBA	0.200	0.212	5.8	30.00
NFDHA	0.200	0.238	19.1	30.00
9CL-PF3ONS	0.187	0.238	27.1	30.00
11CL-PF3OUDS	0.189	0.217	15.0	30.00
3:3FTCA	0.400	0.388	-3.0	30.00
5:3FTCA	0.400	0.328	-17.9	30.00
7:3FTCA	0.400	0.400	-0.02	30.00

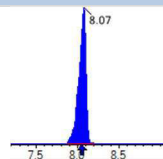
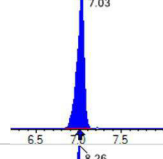
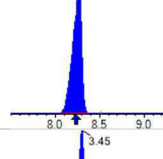
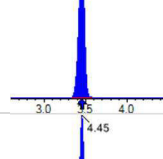
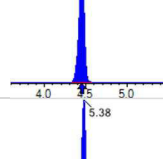
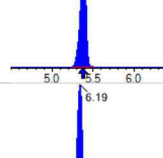
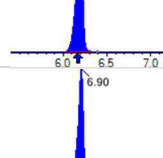
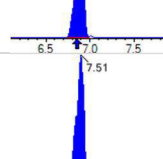
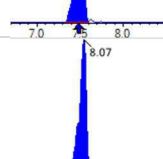
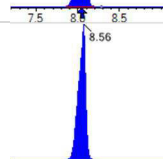
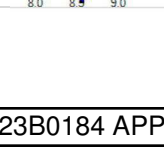
* Values outside of QC limits

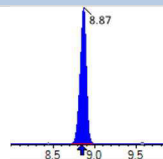
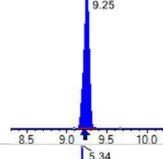
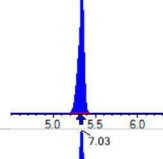
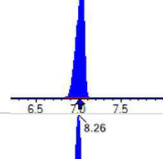
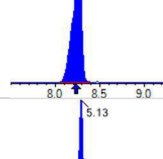
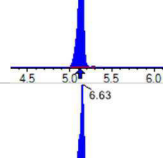
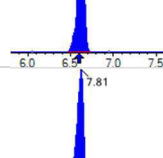
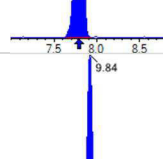
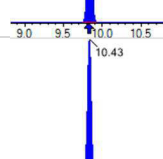
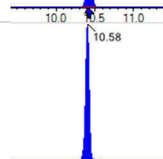
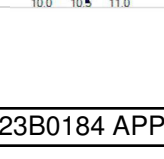
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT Ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) 100671	(3.46, 1.00) (0.00, N/A, 0.0)	63.1	N/A 0.0 0.0	0.4318 [0.4000]	108.0%			
PFPeA	(263.0 / 219.0) 94874 (263.0 / 69.0) 2577	(4.46, 1.00) (0.00, N/A, -1.3)	289.3 38.0	0.0272 227.9 230.9	0.2194 [0.2000]	109.7%			
PFHxA	(313.0 / 269.0) 57769 (313.0 / 119.0) 9878	(5.38, 1.00) (0.00, N/A, -0.3)	197.4 19763.6	0.1710 182.7 165.5	0.1029 [0.1000]	102.9%			IR2,
PFHpA	(363.0 / 319.0) 54365 (363.0 / 169.0) 15153	(6.20, 1.00) (0.01, N/A, 0.9)	6291.7 27088.4	0.2787 88.3 85.5	0.1169 [0.1000]	116.9%			
PFOA	(413.0 / 369.0) 67996 (413.0 / 169.0) 26021	(6.90, 1.00) (0.00, N/A, 0.9)	851.9 84.5	0.3827 113.1 107.7	0.1132 [0.1000]	113.2%			
PFNA	(463.0 / 419.0) 51070 (463.0 / 169.0) 16789	(7.50, 1.00) (-0.01, N/A, 1.4)	437.1 2333.3	0.3288 140.4 141.8	0.1049 [0.1000]	104.9%			
PFDA	(513.0 / 469.0) 73483 (513.0 / 169.0) 6247	(8.06, 1.00) (-0.01, N/A, 4.4)	100.7 179.9	0.0850 68.6 62.8	0.1110 [0.1000]	111.0%			
PFUnA	(563.0 / 519.0) 59842 (563.0 / 169.0) 7675	(8.55, 1.00) (-0.01, N/A, 0.8)	901951.5 244.4	0.1283 119.2 87.8	0.1080 [0.1000]	108.0%			
PFDoA	(613.0 / 569.0) 45797 (613.0 / 169.0) 6873	(8.87, 1.00) (0.01, N/A, -1.1)	133.1 8821.9	0.1501 94.5 79.5	0.0884 [0.1000]	88.4%			
PFTTrDA	(663.0 / 619.0) 49966 (663.0 / 169.0) 17744	(9.07, 1.02) (N/A, 0.02, 0.8)	136.8 24561.1	0.3551 130.8 124.8	0.1003 [0.1000]	100.3%			
PFTeDA	(713.0 / 669.0) 70105 (713.0 / 169.0) 13246	(9.25, 1.00) (0.00, N/A, -0.5)	72.9 182.4	0.1889 86.4 85.8	0.1274 [0.1000]	127.4%			

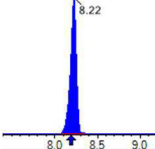
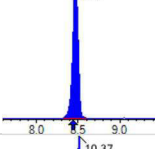
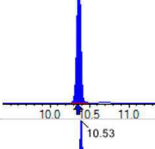
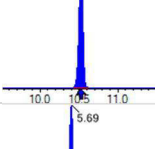
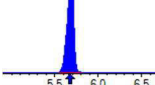
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) 114285 (299.0 / 99.0) 74655	(5.33 , 1.00) (-0.01 , N/A , 0.1)	541.1 334.2	0.6532 103.8 104.7	0.0976 [0.0885]	110.3%			
PFPeS	(349.0 / 80.0) 169183 (349.0 / 99.0) 64194	(6.25 , 0.89) (N/A , 0.03 , 0.3)	282453.0 173223.9	0.3794 113.4 113.3	0.1014 [0.0938]	108.0%			
PFHxS	(399.0 / 80.0) 138467 (399.0 / 99.0) 54566	(7.03 , 1.00) (0.00 , N/A , 0.5)	276.6 959.8	0.3941 123.1 114.1	0.0945 [0.0911]	103.8%			
PFHpS	(449.0 / 80.0) 155270 (449.0 / 99.0) 33892	(7.68 , 0.93) (N/A , 0.02 , 0.3)	70988.0 81.1	0.2183 80.4 78.0	0.0918 [0.0951]	96.5%			
PFOS	(499.0 / 80.0) 229188 (499.0 / 99.0) 50275	(8.27 , 1.00) (0.01 , N/A , 1.2)	58.0 55.5	0.2194 99.7 95.1	0.1106 [0.0927]	119.2%			
PFNS	(549.0 / 80.0) 202893 (549.0 / 99.0) 51396	(8.71 , 1.05) (N/A , 0.02 , 0.5)	1064.3 1100940.1	0.2533 105.3 109.9	0.0978 [0.0960]	101.9%			
PFDS	(599.0 / 80.0) 233103 (599.0 / 99.0) 50747	(8.97 , 1.09) (N/A , 0.02 , -0.1)	296.4 188.2	0.2177 107.6 102.0	0.1011 [0.0963]	105.0%			
PFDoS	(699.0 / 80.0) 124475 (699.0 / 99.0) 25878	(9.32 , 1.13) (N/A , 0.01 , -1.2)	2441.3 132.2	0.2079 106.4 99.6	0.1170 [0.0970]	120.6%			
4:2FTS	(327.0 / 307.0) 145925 (327.0 / 81.0) 90284	(5.13 , 1.00) (0.00 , N/A , 0.0)	636.1 316.6	0.6187 89.3 97.6	0.4515 [0.3738]	120.8%			
6:2FTS	(427.0 / 407.0) 80508 (427.0 / 81.0) 44678	(6.63 , 1.00) (0.00 , N/A , -0.2)	364.0 297.9	0.5549 65.5 72.4	0.5315 [0.3796]	140.0%			QC,
8:2FTS	(527.0 / 507.0) 68518 (527.0 / 81.0) 50620	(7.82 , 1.00) (0.01 , N/A , 0.9)	249.5 122.5	0.7388 90.6 92.6	0.4560 [0.3833]	119.0%			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) 289304 (498.0 / 478.0) 11384	(9.84 , 1.00) (0.00 , N/A , 0.0)	910.4 152.0	0.0394 200.3 193.0	0.1134 [0.1000]	113.4%			
NMeFOSA	(512.0 / 219.0) 269594 (512.0 / 169.0) 219956	(10.43 , 1.00) (0.00 , N/A , 1.0)	1132.2 806.8	0.8159 98.8 97.5	0.4361 [0.4000]	109.0%			
NEiFOSA	(526.0 / 219.0) 259947 (526.0 / 169.0) 340461	(10.57 , 1.00) (-0.01 , N/A , 0.2)	1657.9 1289.5	1.3097 103.4 102.2	0.4339 [0.4000]	108.5%			
NMeFOSAA	(570.0 / 419.0) 32321 (570.0 / 483.0) 5850	(8.23 , 1.00) (0.01 , N/A , 0.7)	637.4 396620.1	0.1810 36.9 40.9	0.1227 [0.1000]	122.7%			IR1,
NEiFOSAA	(584.0 / 419.0) 25711 (584.0 / 526.0) 8756	(8.48 , 1.00) (0.02 , N/A , 0.3)	39530.8 19060.0	0.3406 60.3 66.5	0.1207 [0.1000]	120.7%			
NMeFOSE	(616.0 / 59.0) 114149	(10.37 , 1.00) (0.00 , N/A , 0.0)	256.8	N/A 0.0 0.0	0.4364 [0.4000]	109.1%			
NEiFOSE	(630.0 / 59.0) 113151	(10.53 , 1.00) (0.01 , N/A , 0.0)	227.3	N/A 0.0 0.0	0.4370 [0.4000]	109.2%			
HFPO-DA	(285.0 / 169.0) 66795 (285.0 / 185.0) 176165	(5.68 , 1.00) (0.00 , N/A , -0.5)	168.7 402.7	2.6374 95.9 87.2	0.2160 [0.2000]	108.0%			
ADONA	(377.0 / 85.0) 238432 (377.0 / 251.0) 18151	(6.47 , 1.14) (N/A , 0.02 , -1.3)	355.5 121.5	0.0761 87.0 87.6	0.2077 [0.1885]	110.2%			
9CI-Pf3ONS	(531.0 / 351.0) 639863 (533.0 / 353.0) 200544	(8.63 , 1.52) (N/A , 0.03 , 0.1)	480.9 270.6	0.3134 106.4 100.7	0.2377 [0.1867]	127.4%			
11CI-PF3OUDS	(631.0 / 451.0) 287641 (633.0 / 453.0) 81203	(9.12 , 1.60) (N/A , 0.02 , -0.4)	2286.9 312.9	0.2823 91.5 91.2	0.2173 [0.1886]	115.2%			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) 4955 (241.0 / 117.0) 7572	(4.00, 0.90) (N/A, 0.00, -0.1)	266.7 54.6	1.5282 88.1 92.9	0.3880 [0.4000]	97.0%			
5:3FTCA	(341.0 / 236.7) 26371 (341.0 / 217.0) 41846	(5.90, 1.10) (N/A, 0.02, -0.1)	108895.3 134.0	1.5868 94.4 87.3	0.3284 [0.4000]	82.1%			
7:3FTCA	(441.0 / 317.0) 57727 (441.0 / 337.0) 43166	(7.82, 1.45) (N/A, 0.05, 1.0)	26312.1 275.6	0.7478 87.1 89.1	0.3999 [0.4000]	100.0%			
PFEESA	(315.0 / 135.0) 133935 (315.0 / 83.0) 40665	(5.78, 1.07) (N/A, 0.02, 0.1)	1480.0 211.7	0.3036 107.9 115.9	0.1919 [0.1785]	107.5%			
PFMPA	(229.0 / 85.0) 20875	(3.84, 0.86) (N/A, 0.00, 0.0)	245.9	N/A 0.0 0.0	0.1888 [0.2000]	94.4%			
PFMBA	(279.0 / 85.0) 80122	(4.76, 1.07) (N/A, 0.00, 0.0)	869.1	N/A 0.0 0.0	0.2115 [0.2000]	105.8%			
NFDHA	(295.0 / 201.0) 74437 (295.0 / 85.0) 73719	(5.29, 0.98) (N/A, 0.01, 0.0)	1474.6 328.5	0.9904 94.9 95.5	0.2383 [0.2000]	119.1%			
13C3_PFBA_IIS	(216.0 / 172.0) 254290	(3.45, N/A) (N/A, 0.00, N/A)	1252.9	N/A	0.7501 [1.0000]	75.0% { 121.4% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 465742	(5.38, N/A) (N/A, 0.01, N/A)	1348.1	N/A	0.7831 [1.0000]	78.3% { 125.0% }			
13C4_PFOA_IIS	(417.0 / 372.0) 591053	(6.90, N/A) (N/A, 0.03, N/A)	1296.5	N/A	0.7266 [1.0000]	72.7% { 114.1% }			
13C5_PFNA_IIS	(468.0 / 423.0) 604190	(7.51, N/A) (N/A, 0.03, N/A)	1233.9	N/A	0.7912 [1.0000]	79.1% { 109.7% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 581695	(8.07, N/A) (N/A, 0.04, N/A)	1129.3	N/A	0.8103 [1.0000]	81.0% { 121.5% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 956611	(7.03, N/A) (N/A, 0.03, N/A)	1489.0	N/A	0.7768 [1.0000]	77.7% { 108.1% }			
13C4_PFOS_IIS	(503.0 / 79.9) 1578124	(8.26, N/A) (N/A, 0.03, N/A)	928.0	N/A	0.8892 [1.0000]	88.9% { 122.4% }			
13C4_PFBA_EIS	(217.0 / 172.0) 2205354	(3.45, N/A) (N/A, 0.00, N/A)	3552.4	N/A	7.2935 [8.0000]	91.2% { 115.5% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 1963177	(4.45, N/A) (N/A, 0.00, N/A)	2292.7	N/A	3.7830 [4.0000]	94.6% { 112.0% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 1203726	(5.38, N/A) (N/A, 0.02, N/A)	949.0	N/A	1.9022 [2.0000]	95.1% { 113.4% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 1041690	(6.19, N/A) (N/A, 0.03, N/A)	1863.7	N/A	1.7650 [2.0000]	88.2% { 115.5% }			
13C8_PFOA_EIS	(421.0 / 376.0) 1247152	(6.90, N/A) (N/A, 0.03, N/A)	1434.7	N/A	1.9714 [2.0000]	98.6% { 117.5% }			
13C9_PFNA_EIS	(472.0 / 427.0) 515464	(7.51, N/A) (N/A, 0.02, N/A)	765.3	N/A	0.8561 [1.0000]	85.6% { 102.9% }			
13C6_PFDA_EIS	(519.0 / 474.0) 667496	(8.07, N/A) (N/A, 0.04, N/A)	902.0	N/A	0.9914 [1.0000]	99.1% { 106.8% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 630172	(8.56, N/A) (N/A, 0.03, N/A)	1226.1	N/A	0.9448 [1.0000]	94.5% { 107.0% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT Ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 593589	(8.87 , N/A) (N/A , 0.02 , N/A)	1125.5	N/A	0.9595 [1.0000]	96.0% { 132.5% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 598655	(9.25 , N/A) (N/A , 0.02 , N/A)	940.0	N/A	0.9309 [1.0000]	93.1% { 115.6% }			
13C3_PFBs_EIS	(302.0 / 80.0) 4119752	(5.34 , N/A) (N/A , 0.02 , N/A)	2217.7	N/A	2.1312 [2.0000]	106.6% { 115.4% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 1859452	(7.03 , N/A) (N/A , 0.03 , N/A)	1328.8	N/A	1.8411 [2.0000]	92.1% { 106.1% }			
13C8_PFOS_EIS	(507.0 / 80.0) 3756911	(8.26 , N/A) (N/A , 0.03 , N/A)	1189.7	N/A	1.8121 [2.0000]	90.6% { 114.7% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 458003	(5.13 , N/A) (N/A , 0.01 , N/A)	1020.7	N/A	3.8682 [4.0000]	96.7% { 113.6% }			
13C2_6:2FTS_EIS	(429.0 / 81.0) 464450	(6.63 , N/A) (N/A , 0.03 , N/A)	1840.7	N/A	3.4792 [4.0000]	87.0% { 108.8% }			
13C2_8:2FTS_EIS	(529.0 / 81.0) 535432	(7.81 , N/A) (N/A , 0.03 , N/A)	824.6	N/A	3.7929 [4.0000]	94.8% { 102.4% }			
13C8_PFOsa_EIS	(506.0 / 78.0) 6567165	(9.84 , N/A) (N/A , 0.02 , N/A)	2055.4	N/A	1.7196 [2.0000]	86.0% { 113.4% }			
D3_NMeFOsa_EIS	(515.0 / 169.0) 1439491	(10.43 , N/A) (N/A , 0.01 , N/A)	2001.6	N/A	1.7553 [2.0000]	87.8% { 108.4% }			
D5_NEtFOsa_EIS	(531.0 / 169.0) 1233365	(10.58 , N/A) (N/A , 0.01 , N/A)	2189.0	N/A	1.7176 [2.0000]	85.9% { 101.8% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1232516	(8.22 , N/A) (N/A , 0.03 , N/A)	1519.4	N/A	4.2358 [4.0000]	105.9% { 111.0% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 977798	(8.47 , N/A) (N/A , 0.03 , N/A)	3085.9	N/A	3.9463 [4.0000]	98.7% { 116.1% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 4992539	(10.37 , N/A) (N/A , 0.01 , N/A)	1165.1	N/A	17.8821 [20.0000]	89.4% { 116.7% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 5124957	(10.53 , N/A) (N/A , 0.01 , N/A)	1563.0	N/A	16.0474 [20.0000]	80.2% { 96.0% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 3017733	(5.69 , N/A) (N/A , 0.02 , N/A)	1897.1	N/A	7.5891 [8.0000]	94.9% { 108.5% }			

INITIAL AND CONTINUING CALIBRATION CHECK

EPA 1633

Laboratory: APPL, LLC

Work Order: 23B0184

Client: AECOM

Project: Red Hill AFFF Assessment Sampling

Instrument ID: Saphira

Calibration: 2309009

Standard ID: 23B0084

Sequence: SC00861

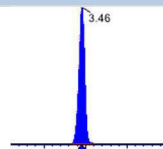
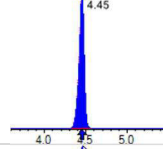
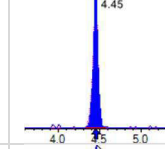
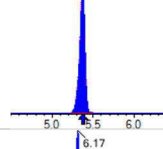
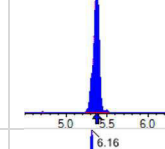
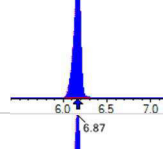
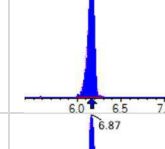
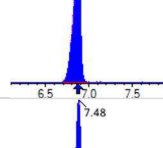
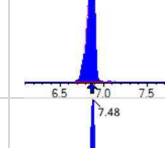
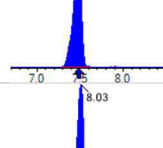
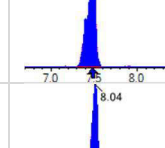
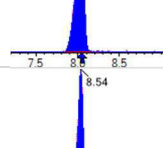
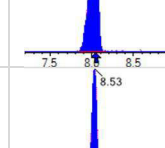
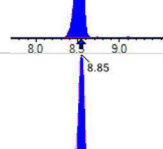
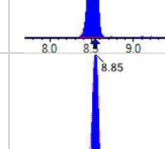
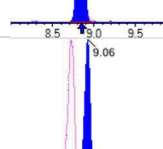
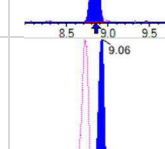
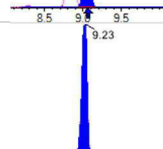
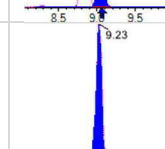
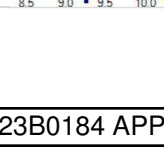
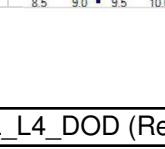
Lab Sample ID	Analyte	True	Found	%R	Units	Control Limit
SC00861-CCV1	PFBA	20.0	22.5	113	ng/mL	+/- 30.00%
	PFPEA	10.0	10.8	108	ng/mL	+/- 30.00%
	PFHXA	5.00	5.50	110	ng/mL	+/- 30.00%
	PFHPA	5.00	5.60	112	ng/mL	+/- 30.00%
	PFOA	5.00	5.42	108	ng/mL	+/- 30.00%
	PFNA	5.00	5.58	112	ng/mL	+/- 30.00%
	PFDA	5.00	4.95	99.0	ng/mL	+/- 30.00%
	PFUnA	5.00	5.34	107	ng/mL	+/- 30.00%
	PFDOA	5.00	5.79	116	ng/mL	+/- 30.00%
	PFTRDA	5.00	6.26	125	ng/mL	+/- 30.00%
	PFTEDA	5.00	5.07	101	ng/mL	+/- 30.00%
	PFBS	4.42	4.91	111	ng/mL	+/- 30.00%
	PFPEs	4.70	5.48	117	ng/mL	+/- 30.00%
	PFHXS	4.58	4.97	108	ng/mL	+/- 30.00%
	PFHPS	4.78	5.15	108	ng/mL	+/- 30.00%
	PFOS	4.65	5.29	114	ng/mL	+/- 30.00%
	PFNS	4.80	5.78	120	ng/mL	+/- 30.00%
	PFDS	4.82	5.52	115	ng/mL	+/- 30.00%
	PFDOS	4.85	5.54	114	ng/mL	+/- 30.00%
	4:2FTS	18.8	21.9	116	ng/mL	+/- 30.00%
	6:2FTS	19.0	24.2	127	ng/mL	+/- 30.00%
	8:2FTS	19.2	22.3	116	ng/mL	+/- 30.00%
	PFOSA	5.00	5.68	114	ng/mL	+/- 30.00%
	NMeFOSA	20.0	22.1	110	ng/mL	+/- 30.00%
	NEtFOSA	20.0	22.9	115	ng/mL	+/- 30.00%
	NMeFOSAA	5.00	5.17	103	ng/mL	+/- 30.00%
	NEtFOSAA	5.00	5.85	117	ng/mL	+/- 30.00%
	NMeFOSE	20.0	22.6	113	ng/mL	+/- 30.00%
	NEtFOSE	20.0	22.4	112	ng/mL	+/- 30.00%
	HFPO-DA	10.0	10.4	104	ng/mL	+/- 30.00%

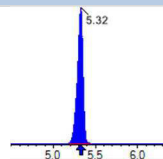
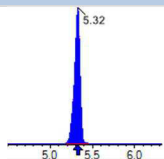
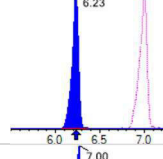
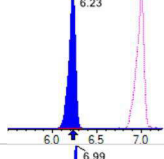
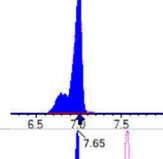
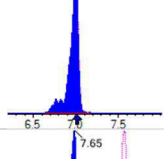
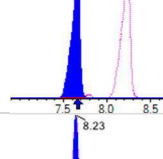
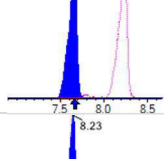
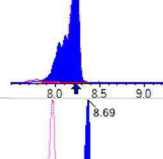
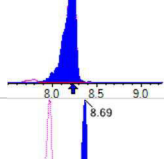
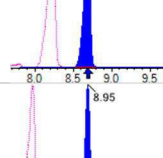
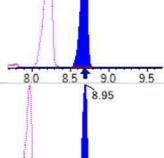
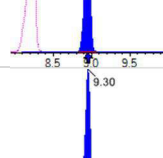
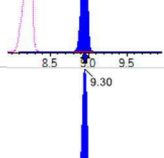
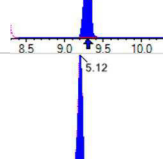
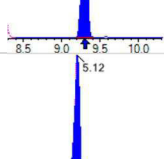
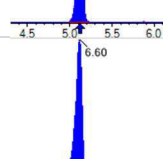
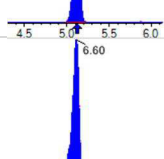
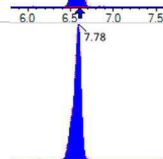
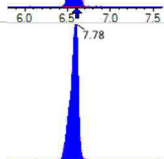
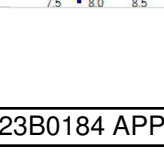
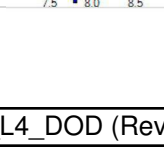
INITIAL AND CONTINUING CALIBRATION CHECK

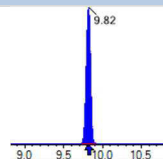
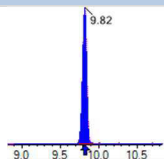
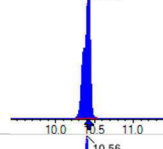
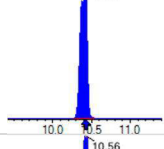
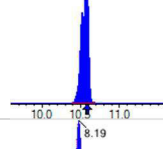
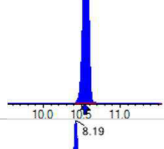
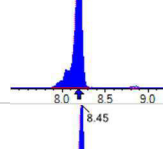
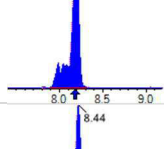
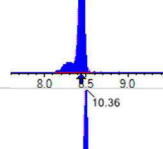
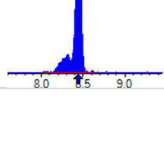
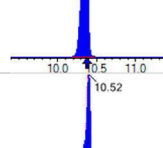
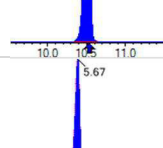
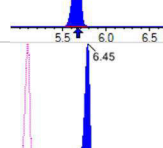
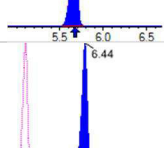
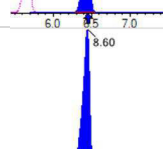
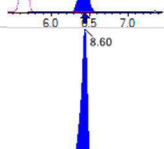
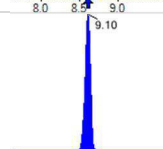
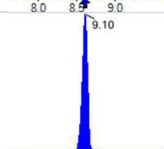
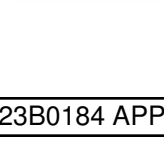
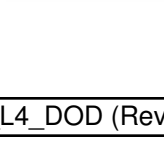
EPA 1633

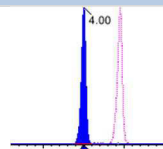
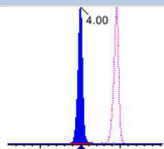
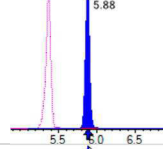
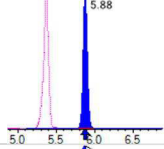
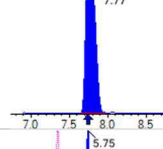
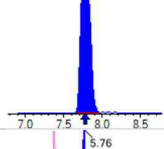
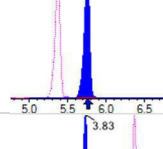
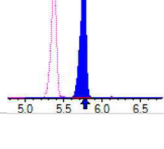
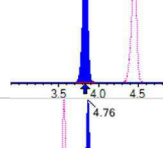
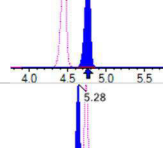
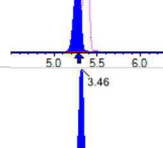
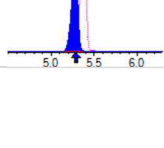
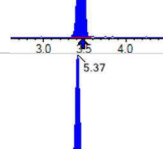
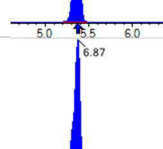
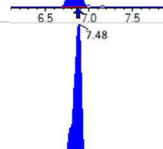
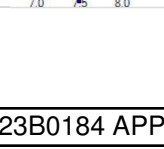
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Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Instrument ID:	Saphira	Calibration:	2309009
Standard ID:	23B0084	Sequence:	SC00861

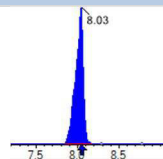
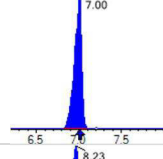
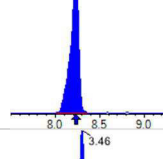
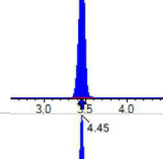
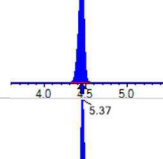
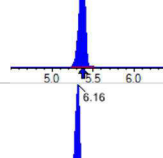
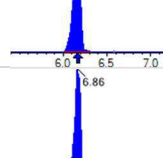
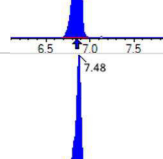
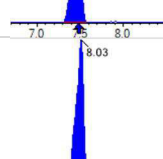
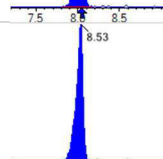
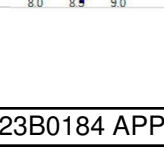
Lab Sample ID	Analyte	True	Found	%R	Units	Control Limit
SC00861-CCV1	ADONA	9.45	10.6	112	ng/mL	+/- 30.00%
	PFEESA	8.90	11.0	124	ng/mL	+/- 30.00%
	PFMPA	10.0	9.60	96.0	ng/mL	+/- 30.00%
	PFMBA	10.0	11.2	112	ng/mL	+/- 30.00%
	NFDHA	10.0	11.4	114	ng/mL	+/- 30.00%
	9CL-PF3ONS	9.35	12.3	132	ng/mL	+/- 30.00%
	11CL-PF3OUDS	9.45	10.9	115	ng/mL	+/- 30.00%
	3:3FTCA	20.0	18.5	92.6	ng/mL	+/- 30.00%
	5:3FTCA	20.0	20.7	103	ng/mL	+/- 30.00%
	7:3FTCA	20.0	20.7	103	ng/mL	+/- 30.00%

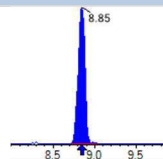
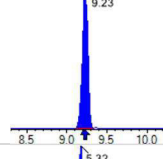
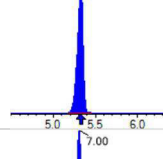
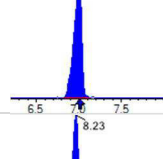
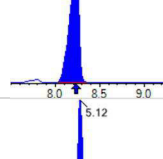
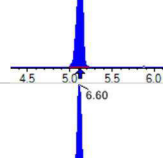
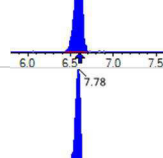
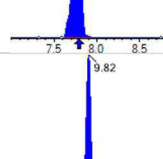
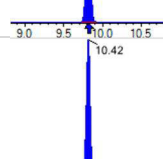
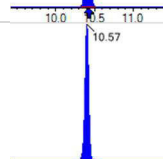
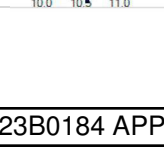
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) 4548912	(3.46, 1.00) (0.00, N/A, 0.0)	208.3	N/A 0.0 0.0	22.5424 [20.0000]	112.7%			
PFPeA	(263.0 / 219.0) 4171462 (263.0 / 69.0) 49078	(4.45, 1.00) (0.00, N/A, 0.2)	3220.6 380.3	0.0118 98.7 100.0	10.8039 [10.0000]	108.0%			
PFHxA	(313.0 / 269.0) 2724551 (313.0 / 119.0) 281468	(5.37, 1.00) (0.00, N/A, -0.1)	3989.3 3424.1	0.1033 110.4 100.0	5.4996 [5.0000]	110.0%			
PFHpA	(363.0 / 319.0) 2255747 (363.0 / 169.0) 735622	(6.17, 1.00) (0.00, N/A, 0.1)	16281.3 137494385.7	0.3261 103.3 100.0	5.6024 [5.0000]	112.0%			
PFOA	(413.0 / 369.0) 2771757 (413.0 / 169.0) 985121	(6.87, 1.00) (0.00, N/A, 0.0)	3943.8 1173.4	0.3554 105.0 100.0	5.4183 [5.0000]	108.4%			
PFNA	(463.0 / 419.0) 2638786 (463.0 / 169.0) 611641	(7.48, 1.00) (0.00, N/A, -0.1)	10890.1 49595.9	0.2318 99.0 100.0	5.5758 [5.0000]	111.5%			
PFDA	(513.0 / 469.0) 3068143 (513.0 / 169.0) 415499	(8.03, 1.00) (0.00, N/A, -0.2)	1317.6 1526.8	0.1354 109.3 100.0	4.9491 [5.0000]	99.0%			
PFUnA	(563.0 / 519.0) 2766764 (563.0 / 169.0) 404022	(8.54, 1.00) (0.00, N/A, 0.3)	1548.1 949.1	0.1460 135.7 100.0	5.3429 [5.0000]	106.9%			
PFDaA	(613.0 / 569.0) 2260446 (613.0 / 169.0) 426823	(8.85, 1.00) (0.00, N/A, 0.1)	1451.3 1077.3	0.1888 118.9 100.0	5.7854 [5.0000]	115.7%			
PFTrDA	(663.0 / 619.0) 2351876 (663.0 / 169.0) 669400	(9.06, 1.02) (N/A, 0.00, 0.1)	1796.5 1619.1	0.2846 104.8 100.0	6.2554 [5.0000]	125.1%			
PFTeDA	(713.0 / 669.0) 2415113 (713.0 / 169.0) 532104	(9.23, 1.00) (0.00, N/A, 0.2)	1356.2 654.2	0.2203 100.8 100.0	5.0736 [5.0000]	101.5%			

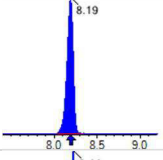
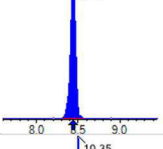
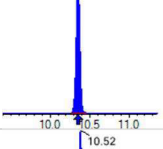
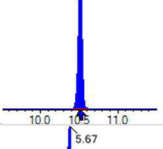
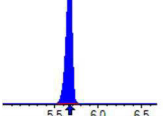
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) 4984630 (299.0 / 99.0) 3110636	(5.32 , 1.00) (0.00 , N/A , 0.0)	1778.8 1758.3	0.6240 99.2 100.0	4.9124 [4.4237]	111.0%			
PFPeS	(349.0 / 80.0) 8612457 (349.0 / 99.0) 2884067	(6.23 , 0.89) (N/A , 0.00 , 0.1)	41104.0 4439.0	0.3349 100.1 100.0	5.4768 [4.6919]	116.7%			
PFHxS	(399.0 / 80.0) 6852016 (399.0 / 99.0) 2367419	(7.00 , 1.00) (0.00 , N/A , 0.2)	11291.4 2083.3	0.3455 108.0 100.0	4.9662 [4.5549]	109.0%			
PFHpS	(449.0 / 80.0) 7590497 (449.0 / 99.0) 2123112	(7.65 , 0.93) (N/A , 0.00 , 0.1)	50496.7 3016.1	0.2797 103.0 100.0	5.1491 [4.7570]	108.2%			
PFOS	(499.0 / 80.0) 9553173 (499.0 / 99.0) 2202834	(8.23 , 1.00) (0.00 , N/A , 0.1)	299.4 392.4	0.2306 104.8 100.0	5.2875 [4.6375]	114.0%			
PFNS	(549.0 / 80.0) 10450261 (549.0 / 99.0) 2409220	(8.69 , 1.06) (N/A , 0.00 , 0.1)	10383.1 11083469.0	0.2305 95.8 100.0	5.7802 [4.7994]	120.4%			
PFDS	(599.0 / 80.0) 11100224 (599.0 / 99.0) 2370067	(8.95 , 1.09) (N/A , 0.00 , -0.1)	1784.5 1757.3	0.2135 105.5 100.0	5.5246 [4.8155]	114.7%			
PFDoS	(699.0 / 80.0) 5137682 (699.0 / 99.0) 1072259	(9.30 , 1.13) (N/A , 0.00 , 0.2)	1766.5 1437.5	0.2087 106.8 100.0	5.5390 [4.8478]	114.3%			
4:2FTS	(327.0 / 307.0) 6219957 (327.0 / 81.0) 3943918	(5.12 , 1.00) (0.00 , N/A , 0.0)	2180.7 1555.7	0.6341 91.5 100.0	21.8677 [18.6906]	117.0%			
6:2FTS	(427.0 / 407.0) 3361286 (427.0 / 81.0) 2575585	(6.60 , 1.00) (0.00 , N/A , 0.1)	1909.6 1418.8	0.7662 90.4 100.0	24.1525 [18.9808]	127.2%			
8:2FTS	(527.0 / 507.0) 3275071 (527.0 / 81.0) 2612661	(7.78 , 1.00) (0.00 , N/A , -0.1)	1916.4 1181.4	0.7977 97.8 100.0	22.3210 [19.1658]	116.5%			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) 12772746 (498.0 / 478.0) 260462	(9.82 , 1.00) (0.00 , N/A , 0.0)	3423.9 760.0	0.0204 103.8 100.0	5.6767 [5.0000]	113.5%			
NMeFOSA	(512.0 / 219.0) 12603940 (512.0 / 169.0) 10546339	(10.42 , 1.00) (0.00 , N/A , 1.0)	3383.3 5584.0	0.8367 101.3 100.0	22.0936 [20.0000]	110.5%			
NEIFOSA	(526.0 / 219.0) 13485514 (526.0 / 169.0) 17277295	(10.56 , 1.00) (-0.01 , N/A , 0.5)	16311.1 9846.1	1.2812 101.2 100.0	22.9146 [20.0000]	114.6%			
NMeFOSAA	(570.0 / 419.0) 1226521 (570.0 / 483.0) 543336	(8.19 , 1.00) (0.01 , N/A , 0.2)	1987.6 393.9	0.4430 90.3 100.0	5.1701 [5.0000]	103.4%			
NEIFOSAA	(584.0 / 419.0) 1073808 (584.0 / 526.0) 550098	(8.45 , 1.00) (0.01 , N/A , 0.2)	2530.1 1335.5	0.5123 90.7 100.0	5.8487 [5.0000]	117.0%			
NMeFOSE	(616.0 / 59.0) 5053303	(10.36 , 1.00) (0.01 , N/A , 0.0)	2353.4	N/A 0.0 0.0	22.5531 [20.0000]	112.8%			
NEtFOSE	(630.0 / 59.0) 6033767	(10.52 , 1.00) (0.01 , N/A , 0.0)	994.5	N/A 0.0 0.0	22.3649 [20.0000]	111.8%			
HFPO-DA	(285.0 / 169.0) 2971470 (285.0 / 185.0) 8990396	(5.67 , 1.00) (0.00 , N/A , 0.0)	1730.0 2332.2	3.0256 110.1 100.0	10.4253 [10.0000]	104.3%			
ADONA	(377.0 / 85.0) 11199590 (377.0 / 251.0) 973772	(6.45 , 1.14) (N/A , 0.00 , 0.4)	2110.5 1353.7	0.0869 99.3 100.0	10.5873 [9.4270]	112.3%			
9CI-Pf3ONS	(531.0 / 351.0) 30585757 (533.0 / 353.0) 9520967	(8.60 , 1.52) (N/A , 0.00 , -0.1)	71.2 2017.0	0.3113 105.7 100.0	12.3284 [9.3325]	132.1%			QC,
11CI-PF3OUDS	(631.0 / 451.0) 13270951 (633.0 / 453.0) 4108271	(9.10 , 1.61) (N/A , 0.00 , -0.1)	1748.0 1283.2	0.3096 100.3 100.0	10.8802 [9.4321]	115.4%			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) 211146 (241.0 / 117.0) 347177	(4.00 , 0.90) (N/A , 0.00 , 0.0)	1005.5 900.9	1.6442 94.8 100.0	18.5175 [20.0000]	92.6%			
5:3FTCA	(341.0 / 236.7) 1462752 (341.0 / 217.0) 2657445	(5.88 , 1.09) (N/A , 0.00 , 0.1)	1681.5 2137.1	1.8167 108.0 100.0	20.6549 [20.0000]	103.3%			
7:3FTCA	(441.0 / 317.0) 2634925 (441.0 / 337.0) 2212115	(7.77 , 1.45) (N/A , 0.00 , -0.3)	874.5 659.7	0.8395 97.8 100.0	20.6955 [20.0000]	103.5%			
PFEESA	(315.0 / 135.0) 6799730 (315.0 / 83.0) 1780668	(5.75 , 1.07) (N/A , 0.00 , -0.1)	2088.2 1947.1	0.2619 93.0 100.0	11.0453 [8.9246]	123.8%			
PFMPA	(229.0 / 85.0) 947665	(3.83 , 0.86) (N/A , 0.00 , 0.0)	2696.3	N/A 0.0 0.0	9.5996 [10.0000]	96.0%			
PFMBA	(279.0 / 85.0) 3799259	(4.76 , 1.07) (N/A , 0.00 , 0.0)	2135.3	N/A 0.0 0.0	11.2336 [10.0000]	112.3%			
NFDHA	(295.0 / 201.0) 3135249 (295.0 / 85.0) 3251821	(5.28 , 0.98) (N/A , 0.00 , -0.1)	1911.5 2232.5	1.0372 99.4 100.0	11.3784 [10.0000]	113.8%			
13C3_PFBA_IIS	(216.0 / 172.0) 209442	(3.46 , N/A) (N/A , 0.00 , N/A)	1075.8	N/A	0.6178 [1.0000]	61.8% { 100.0% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 372636	(5.37 , N/A) (N/A , 0.00 , N/A)	1397.3	N/A	0.6265 [1.0000]	62.7% { 100.0% }			
13C4_PFOA_IIS	(417.0 / 372.0) 518235	(6.87 , N/A) (N/A , 0.00 , N/A)	625.5	N/A	0.6371 [1.0000]	63.7% { 100.0% }			
13C5_PFNA_IIS	(468.0 / 423.0) 550835	(7.48 , N/A) (N/A , 0.00 , N/A)	983.2	N/A	0.7213 [1.0000]	72.1% { 100.0% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 478656	(8.03 , N/A) (N/A , 0.00 , N/A)	899.2	N/A	0.6668 [1.0000]	66.7% { 100.0% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 884962	(7.00 , N/A) (N/A , 0.00 , N/A)	1491.5	N/A	0.7186 [1.0000]	71.9% { 100.0% }			
13C4_PFOS_IIS	(503.0 / 79.9) 1289518	(8.23 , N/A) (N/A , 0.00 , N/A)	781.9	N/A	0.7266 [1.0000]	72.7% { 100.0% }			
13C4_PFBA_EIS	(217.0 / 172.0) 1908944	(3.46 , N/A) (N/A , 0.00 , N/A)	2831.4	N/A	7.6650 [8.0000]	95.8% { 100.0% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 1752774	(4.45 , N/A) (N/A , 0.00 , N/A)	1768.6	N/A	4.2214 [4.0000]	105.5% { 100.0% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 1061699	(5.37 , N/A) (N/A , 0.00 , N/A)	1659.1	N/A	2.0969 [2.0000]	104.8% { 100.0% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 901769	(6.16 , N/A) (N/A , 0.00 , N/A)	1509.1	N/A	1.9097 [2.0000]	95.5% { 100.0% }			
13C8_PFOA_EIS	(421.0 / 376.0) 1061814	(6.86 , N/A) (N/A , 0.00 , N/A)	1069.6	N/A	1.9143 [2.0000]	95.7% { 100.0% }			
13C9_PFNA_EIS	(472.0 / 427.0) 501022	(7.48 , N/A) (N/A , 0.00 , N/A)	919.1	N/A	0.9127 [1.0000]	91.3% { 100.0% }			
13C6_PFDA_EIS	(519.0 / 474.0) 625268	(8.03 , N/A) (N/A , 0.00 , N/A)	494.8	N/A	1.1286 [1.0000]	112.9% { 100.0% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 588877	(8.53 , N/A) (N/A , 0.00 , N/A)	1086.4	N/A	1.0729 [1.0000]	107.3% { 100.0% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 447839	(8.85 , N/A) (N/A , 0.00 , N/A)	1107.6	N/A	0.8798 [1.0000]	88.0% { 100.0% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 517685	(9.23 , N/A) (N/A , 0.00 , N/A)	1625.1	N/A	0.9783 [1.0000]	97.8% { 100.0% }			
13C3_PFBs_EIS	(302.0 / 80.0) 3569176	(5.32 , N/A) (N/A , 0.00 , N/A)	2282.7	N/A	1.9958 [2.0000]	99.8% { 100.0% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 1751796	(7.00 , N/A) (N/A , 0.00 , N/A)	1143.8	N/A	1.8750 [2.0000]	93.7% { 100.0% }			
13C8_PFOS_EIS	(507.0 / 80.0) 3274255	(8.23 , N/A) (N/A , 0.00 , N/A)	502.5	N/A	1.9327 [2.0000]	96.6% { 100.0% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 403096	(5.12 , N/A) (N/A , 0.00 , N/A)	1084.3	N/A	3.6801 [4.0000]	92.0% { 100.0% }			
13C2_6:2FTS_EIS	(429.0 / 81.0) 426712	(6.60 , N/A) (N/A , 0.00 , N/A)	888.2	N/A	3.4553 [4.0000]	86.4% { 100.0% }			
13C2_8:2FTS_EIS	(529.0 / 81.0) 522805	(7.78 , N/A) (N/A , 0.00 , N/A)	670.6	N/A	4.0033 [4.0000]	100.1% { 100.0% }			
13C8_PFOsa_EIS	(506.0 / 78.0) 5790807	(9.82 , N/A) (N/A , 0.00 , N/A)	2932.9	N/A	1.8556 [2.0000]	92.8% { 100.0% }			
D3_NMeFOsa_EIS	(515.0 / 169.0) 1328496	(10.42 , N/A) (N/A , 0.00 , N/A)	1750.4	N/A	1.9826 [2.0000]	99.1% { 100.0% }			
D5_NEtFOsa_EIS	(531.0 / 169.0) 1211594	(10.57 , N/A) (N/A , 0.00 , N/A)	2587.7	N/A	2.0650 [2.0000]	103.2% { 100.0% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1110285	(8.19 , N/A) (N/A , 0.00 , N/A)	1093.4	N/A	4.6698 [4.0000]	116.7% { 100.0% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 842448	(8.44 , N/A) (N/A , 0.00 , N/A)	7866.6	N/A	4.1610 [4.0000]	104.0% { 100.0% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 4276766	(10.35 , N/A) (N/A , 0.00 , N/A)	1601.6	N/A	18.7467 [20.0000]	93.7% { 100.0% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 5339898	(10.52 , N/A) (N/A , 0.00 , N/A)	1521.1	N/A	20.4627 [20.0000]	102.3% { 100.0% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 2781318	(5.67 , N/A) (N/A , 0.00 , N/A)	9231.6	N/A	8.7422 [8.0000]	109.3% { 100.0% }			

INITIAL AND CONTINUING CALIBRATION CHECK

EPA 1633

Laboratory: APPL, LLC

Work Order: 23B0184

Client: AECOM

Project: Red Hill AFFF Assessment Sampling

Instrument ID: Saphira

Calibration: 2309009

Standard ID: 23B0084

Sequence: SC00861

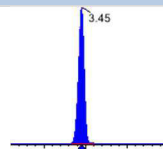
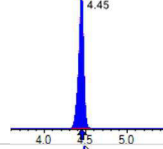
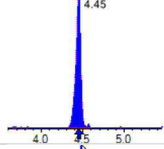
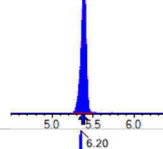
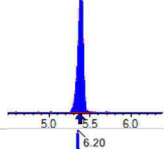
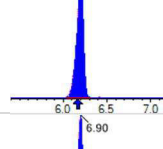
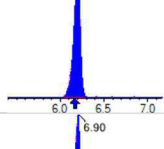
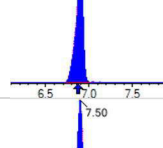
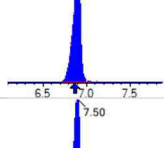
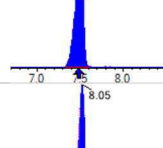
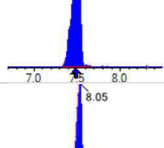
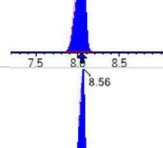
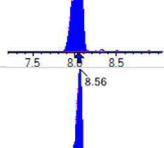
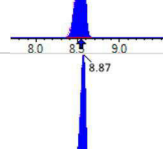
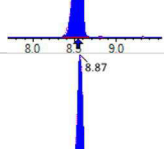
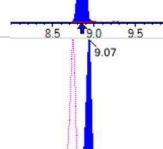
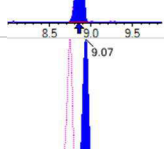
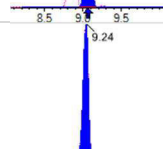
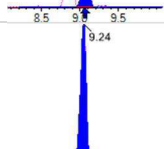
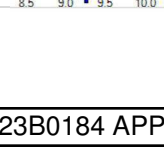
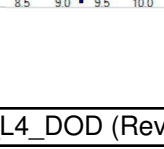
Lab Sample ID	Analyte	True	Found	%R	Units	Control Limit
SC00861-CCV2	PFBA	20.0	22.0	110	ng/mL	+/- 30.00%
	PFPEA	10.0	10.7	107	ng/mL	+/- 30.00%
	PFHXA	5.00	5.51	110	ng/mL	+/- 30.00%
	PFHPA	5.00	5.29	106	ng/mL	+/- 30.00%
	PFOA	5.00	5.18	104	ng/mL	+/- 30.00%
	PFNA	5.00	5.23	105	ng/mL	+/- 30.00%
	PFDA	5.00	5.35	107	ng/mL	+/- 30.00%
	PFUnA	5.00	5.21	104	ng/mL	+/- 30.00%
	PFDOA	5.00	5.23	105	ng/mL	+/- 30.00%
	PFTRDA	5.00	5.46	109	ng/mL	+/- 30.00%
	PFTEDA	5.00	5.03	101	ng/mL	+/- 30.00%
	PFBS	4.42	5.21	118	ng/mL	+/- 30.00%
	PFPEs	4.70	5.37	114	ng/mL	+/- 30.00%
	PFHXS	4.58	5.02	110	ng/mL	+/- 30.00%
	PFHPS	4.78	5.50	115	ng/mL	+/- 30.00%
	PFOS	4.65	4.96	107	ng/mL	+/- 30.00%
	PFNS	4.80	6.02	125	ng/mL	+/- 30.00%
	PFDS	4.82	5.90	122	ng/mL	+/- 30.00%
	PFDOS	4.85	5.83	120	ng/mL	+/- 30.00%
	4:2FTS	18.8	20.6	109	ng/mL	+/- 30.00%
	6:2FTS	19.0	22.6	119	ng/mL	+/- 30.00%
	8:2FTS	19.2	19.8	103	ng/mL	+/- 30.00%
	PFOSA	5.00	5.90	118	ng/mL	+/- 30.00%
	NMeFOSA	20.0	22.3	111	ng/mL	+/- 30.00%
	NEtFOSA	20.0	23.7	118	ng/mL	+/- 30.00%
	NMeFOSAA	5.00	5.72	114	ng/mL	+/- 30.00%
	NEtFOSAA	5.00	5.49	110	ng/mL	+/- 30.00%
	NMeFOSE	20.0	21.5	107	ng/mL	+/- 30.00%
	NEtFOSE	20.0	22.9	114	ng/mL	+/- 30.00%
	HFPO-DA	10.0	10.7	107	ng/mL	+/- 30.00%

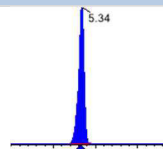
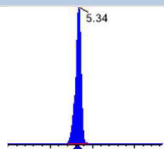
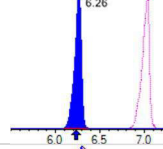
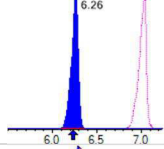
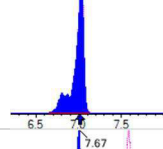
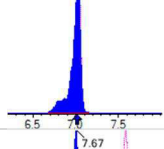
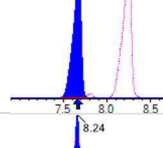
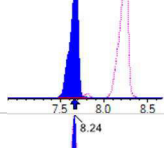
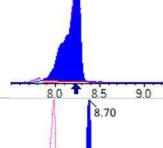
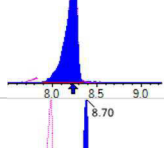
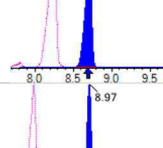
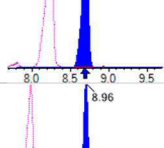
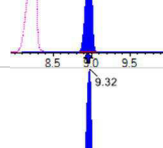
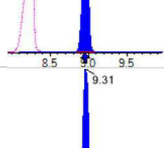
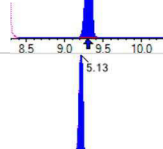
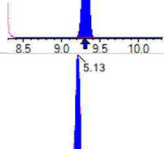
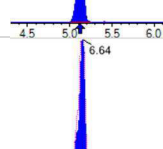
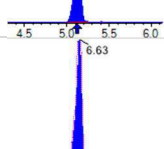
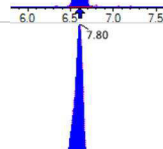
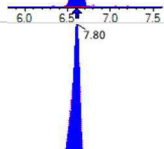
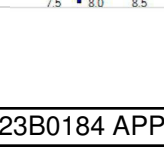
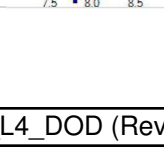
INITIAL AND CONTINUING CALIBRATION CHECK

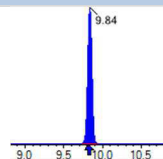
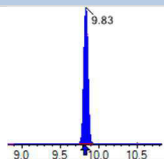
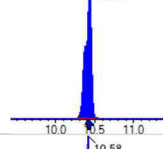
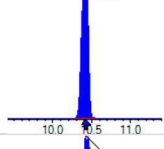
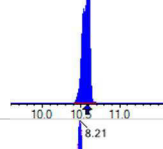
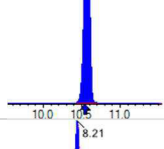
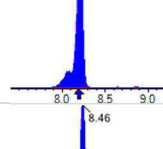
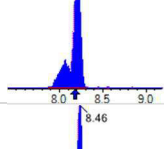
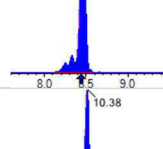
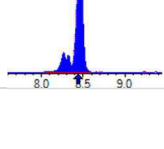
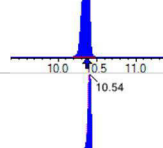
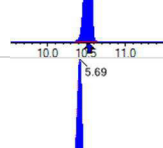
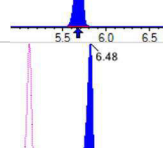
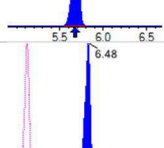
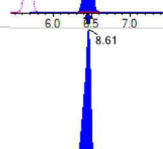
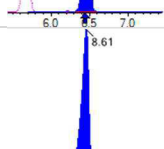
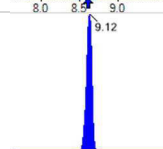
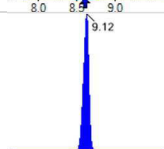
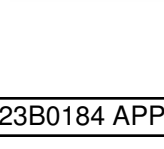
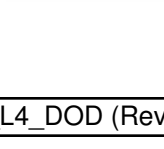
EPA 1633

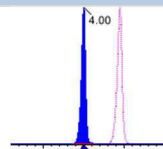
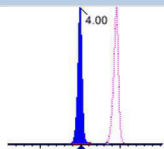
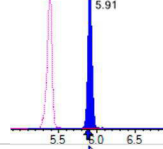
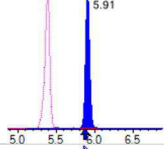
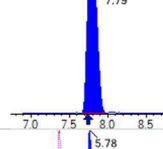
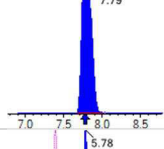
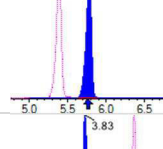
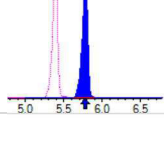
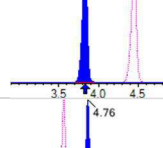
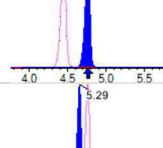
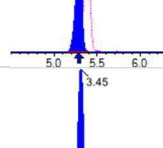
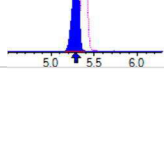
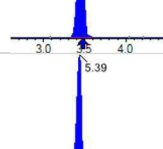
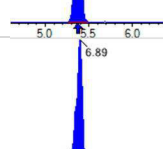
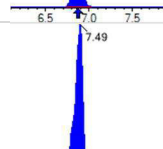
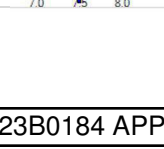
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Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Instrument ID:	Saphira	Calibration:	2309009
Standard ID:	23B0084	Sequence:	SC00861

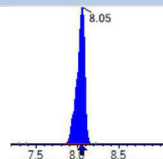
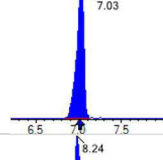
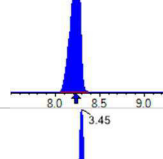
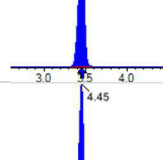
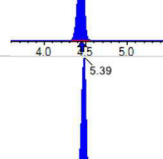
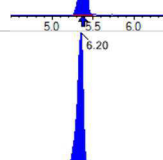
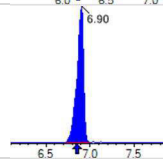
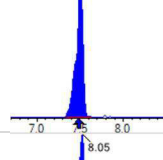
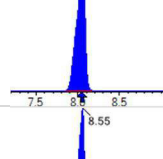
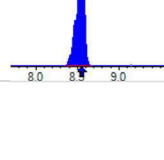
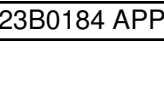
Lab Sample ID	Analyte	True	Found	%R	Units	Control Limit
SC00861-CCV2	ADONA	9.45	9.99	106	ng/mL	+/- 30.00%
	PFEESA	8.90	10.7	120	ng/mL	+/- 30.00%
	PFMPA	10.0	9.74	97.4	ng/mL	+/- 30.00%
	PFMBA	10.0	11.5	115	ng/mL	+/- 30.00%
	NFDHA	10.0	10.7	107	ng/mL	+/- 30.00%
	9CL-PF3ONS	9.35	11.5	123	ng/mL	+/- 30.00%
	11CL-PF3OUDS	9.45	10.8	114	ng/mL	+/- 30.00%
	3:3FTCA	20.0	19.1	95.7	ng/mL	+/- 30.00%
	5:3FTCA	20.0	19.7	98.4	ng/mL	+/- 30.00%
	7:3FTCA	20.0	19.3	96.7	ng/mL	+/- 30.00%

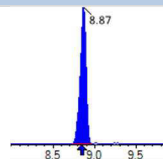
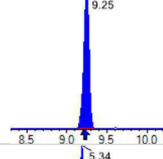
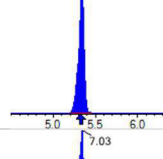
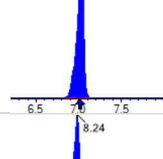
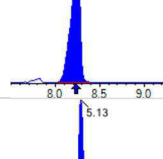
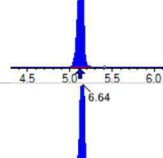
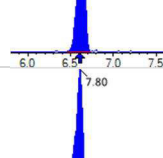
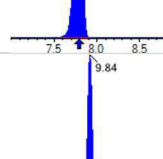
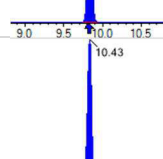
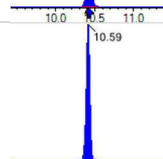
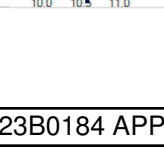
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT Ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) 5338181	(3.45, 1.00) (0.00, N/A, 0.0)	231.0	N/A 0.0 0.0	22.0489 [20.0000]	110.2%			
PFPeA	(263.0 / 219.0) 4859048 (263.0 / 69.0) 62008	(4.45, 1.00) (0.00, N/A, -0.2)	3157.5 561.1	0.0128 107.1 108.5	10.7496 [10.0000]	107.5%			
PFHxA	(313.0 / 269.0) 3227920 (313.0 / 119.0) 344151	(5.39, 1.00) (0.00, N/A, 0.0)	3868.6 32981.4	0.1066 113.9 103.2	5.5050 [5.0000]	110.1%			
PFHpA	(363.0 / 319.0) 2556902 (363.0 / 169.0) 784728	(6.20, 1.00) (0.00, N/A, 0.2)	4966.9 1542.6	0.3069 97.2 94.1	5.2943 [5.0000]	105.9%			
PFOA	(413.0 / 369.0) 3328107 (413.0 / 169.0) 1155013	(6.90, 1.00) (0.00, N/A, 0.2)	9550.7 12805241.6	0.3470 102.6 97.6	5.1776 [5.0000]	103.6%			
PFNA	(463.0 / 419.0) 2958383 (463.0 / 169.0) 683788	(7.50, 1.00) (-0.01, N/A, -0.1)	4820.1 2391.8	0.2311 98.7 99.7	5.2282 [5.0000]	104.6%			
PFDA	(513.0 / 469.0) 3419688 (513.0 / 169.0) 372120	(8.05, 1.00) (0.00, N/A, -0.2)	1568.8 613.3	0.1088 87.8 80.4	5.3459 [5.0000]	106.9%			
PFUnA	(563.0 / 519.0) 3017945 (563.0 / 169.0) 424660	(8.56, 1.00) (0.00, N/A, -0.1)	1419.7 695.5	0.1407 130.7 96.4	5.2086 [5.0000]	104.2%			
PFDaA	(613.0 / 569.0) 2968842 (613.0 / 169.0) 483803	(8.87, 1.00) (0.00, N/A, 0.1)	1834.6 1933.5	0.1630 102.6 86.3	5.2326 [5.0000]	104.7%			
PFTrDA	(663.0 / 619.0) 2983226 (663.0 / 169.0) 717279	(9.07, 1.02) (N/A, 0.02, 0.2)	1309.1 1139.2	0.2404 88.6 84.5	5.4641 [5.0000]	109.3%			
PFTeDA	(713.0 / 669.0) 3014745 (713.0 / 169.0) 628527	(9.24, 1.00) (0.00, N/A, 0.1)	1815.7 1142.8	0.2085 95.4 94.6	5.0295 [5.0000]	100.6%			

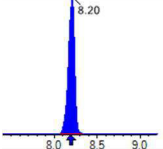
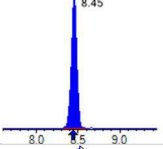
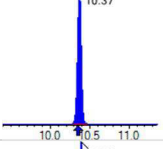
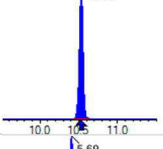
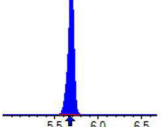
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) 5989097 (299.0 / 99.0) 3383337	(5.34 , 1.00) (0.00 , N/A , -0.1)	1949.4 1801.8	0.5649 89.8 90.5	5.2106 [4.4237]	117.8%			
PFPeS	(349.0 / 80.0) 9367886 (349.0 / 99.0) 3248376	(6.26 , 0.89) (N/A , 0.03 , 0.1)	254060.8 2106.4	0.3468 103.7 103.5	5.3692 [4.6919]	114.4%			
PFHxS	(399.0 / 80.0) 7678644 (399.0 / 99.0) 2482964	(7.03 , 1.00) (0.00 , N/A , 0.3)	5870.4 2200.2	0.3234 101.0 93.6	5.0160 [4.5549]	110.1%			
PFHpS	(449.0 / 80.0) 8897440 (449.0 / 99.0) 2355595	(7.67 , 0.93) (N/A , 0.01 , -0.2)	12644.5 2930.8	0.2647 97.5 94.7	5.5028 [4.7570]	115.7%			
PFOS	(499.0 / 80.0) 9824564 (499.0 / 99.0) 2295569	(8.24 , 1.00) (0.00 , N/A , 0.1)	359.9 536.6	0.2337 106.2 101.3	4.9577 [4.6375]	106.9%			
PFNS	(549.0 / 80.0) 11943385 (549.0 / 99.0) 2628800	(8.70 , 1.06) (N/A , 0.02 , 0.0)	15920.0 2083.5	0.2201 91.5 95.5	6.0229 [4.7994]	125.5%			
PFDS	(599.0 / 80.0) 13005400 (599.0 / 99.0) 2604283	(8.97 , 1.09) (N/A , 0.02 , 0.2)	2615.7 2836.8	0.2002 98.9 93.8	5.9013 [4.8155]	122.5%			
PFDoS	(699.0 / 80.0) 5926277 (699.0 / 99.0) 1133962	(9.32 , 1.13) (N/A , 0.01 , 0.1)	2154.7 1331.4	0.1913 97.9 91.7	5.8251 [4.8478]	120.2%			
4:2FTS	(327.0 / 307.0) 6666871 (327.0 / 81.0) 4644336	(5.13 , 1.00) (0.00 , N/A , 0.1)	1928.1 1795.8	0.6966 100.5 109.9	20.5620 [18.6906]	110.0%			
6:2FTS	(427.0 / 407.0) 3784321 (427.0 / 81.0) 3025908	(6.64 , 1.00) (0.00 , N/A , 0.2)	2680.0 1946.1	0.7996 94.4 104.4	22.5967 [18.9808]	119.1%			
8:2FTS	(527.0 / 507.0) 3654347 (527.0 / 81.0) 2972570	(7.80 , 1.00) (0.00 , N/A , -0.2)	1358.5 1503.0	0.8134 99.7 102.0	19.8235 [19.1658]	103.4%			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) 14533847 (498.0 / 478.0) 288708	(9.84 , 1.00) (0.00 , N/A , 0.1)	2529.5 1526.4	0.0199 101.1 97.4	5.8991 [5.0000]	118.0%			
NMeFOSA	(512.0 / 219.0) 14293665 (512.0 / 169.0) 11737586	(10.43 , 1.00) (0.00 , N/A , 1.0)	4225.0 4407.4	0.8212 99.4 98.1	22.2831 [20.0000]	111.4%			
NEIFOSA	(526.0 / 219.0) 15002764 (526.0 / 169.0) 19194601	(10.58 , 1.00) (-0.01 , N/A , 0.5)	15151.3 12967.1	1.2794 101.0 99.9	23.6636 [20.0000]	118.3%			
NMeFOSAA	(570.0 / 419.0) 1465592 (570.0 / 483.0) 635675	(8.21 , 1.00) (0.00 , N/A , -0.1)	1417.1 337.7	0.4337 88.5 97.9	5.7228 [5.0000]	114.5%			
NEIFOSAA	(584.0 / 419.0) 1091457 (584.0 / 526.0) 639646	(8.46 , 1.00) (0.01 , N/A , 0.0)	2331.4 980.6	0.5860 103.7 114.4	5.4871 [5.0000]	109.7%			
NMeFOSE	(616.0 / 59.0) 5733715	(10.38 , 1.00) (0.01 , N/A , 0.0)	2901.4	N/A 0.0 0.0	21.4885 [20.0000]	107.4%			
NEtFOSE	(630.0 / 59.0) 6887733	(10.54 , 1.00) (0.01 , N/A , 0.0)	1113.5	N/A 0.0 0.0	22.8720 [20.0000]	114.4%			
HFPO-DA	(285.0 / 169.0) 3518236 (285.0 / 185.0) 9642054	(5.69 , 1.00) (0.00 , N/A , 0.0)	2459.3 2224.1	2.7406 99.7 90.6	10.7133 [10.0000]	107.1%			
ADONA	(377.0 / 85.0) 12180658 (377.0 / 251.0) 1100483	(6.48 , 1.14) (N/A , 0.04 , 0.0)	2165.8 1246.9	0.0903 103.2 103.9	9.9939 [9.4270]	106.0%			
9CI-Pf3ONS	(531.0 / 351.0) 32913427 (533.0 / 353.0) 9617076	(8.61 , 1.51) (N/A , 0.02 , 0.0)	1752.8 1965.0	0.2922 99.2 93.9	11.5144 [9.3325]	123.4%			
11CI-PF3OUDS	(631.0 / 451.0) 15163742 (633.0 / 453.0) 4508832	(9.12 , 1.60) (N/A , 0.02 , 0.0)	1689.5 1576.3	0.2973 96.3 96.1	10.7900 [9.4321]	114.4%			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) 255558 (241.0 / 117.0) 415309	(4.00 , 0.90) (N/A , 0.00 , 0.2)	1806.8 1306.5	1.6251 93.7 98.8	19.1442 [20.0000]	95.7%			
5:3FTCA	(341.0 / 236.7) 1649342 (341.0 / 217.0) 2939910	(5.91 , 1.10) (N/A , 0.03 , 0.1)	1615.6 1869.1	1.7825 106.0 98.1	19.6771 [20.0000]	98.4%			
7:3FTCA	(441.0 / 317.0) 2914140 (441.0 / 337.0) 2511414	(7.79 , 1.45) (N/A , 0.02 , 0.3)	693.6 876.5	0.8618 100.4 102.7	19.3383 [20.0000]	96.7%			
PFEESA	(315.0 / 135.0) 7791259 (315.0 / 83.0) 2050264	(5.78 , 1.07) (N/A , 0.02 , 0.1)	3024.2 1910.1	0.2631 93.5 100.5	10.6928 [8.9246]	119.8%			
PFMPA	(229.0 / 85.0) 1126028	(3.83 , 0.86) (N/A , -0.01 , 0.0)	3160.6	N/A 0.0 0.0	9.7431 [10.0000]	97.4%			
PFMBA	(279.0 / 85.0) 4551250	(4.76 , 1.07) (N/A , 0.00 , 0.0)	2257.5	N/A 0.0 0.0	11.4948 [10.0000]	114.9%			
NFDHA	(295.0 / 201.0) 3475412 (295.0 / 85.0) 3828321	(5.29 , 0.98) (N/A , 0.01 , 0.0)	2136.0 2171.5	1.1015 105.5 106.2	10.6565 [10.0000]	106.6%			
13C3_PFBA_IIS	(216.0 / 172.0) 259282	(3.45 , N/A) (N/A , -0.01 , N/A)	913.1	N/A	0.7648 [1.0000]	76.5% { 123.8% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 471912	(5.39 , N/A) (N/A , 0.02 , N/A)	1506.8	N/A	0.7934 [1.0000]	79.3% { 126.6% }			
13C4_PFOA_IIS	(417.0 / 372.0) 631418	(6.89 , N/A) (N/A , 0.03 , N/A)	1259.9	N/A	0.7762 [1.0000]	77.6% { 121.8% }			
13C5_PFNA_IIS	(468.0 / 423.0) 584226	(7.49 , N/A) (N/A , 0.02 , N/A)	1393.6	N/A	0.7651 [1.0000]	76.5% { 106.1% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (Δ RT-I[min], Δ RT-CV[min], Δ RT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 594370	(8.05 , N/A) (N/A , 0.02 , N/A)	852.0	N/A	0.8279 [1.0000]	82.8% { 124.2% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 887629	(7.03 , N/A) (N/A , 0.03 , N/A)	879.3	N/A	0.7207 [1.0000]	72.1% { 100.3% }			
13C4_PFOS_IIS	(503.0 / 79.9) 1440719	(8.24 , N/A) (N/A , 0.01 , N/A)	882.6	N/A	0.8117 [1.0000]	81.2% { 111.7% }			
13C4_PFBA_EIS	(217.0 / 172.0) 2290296	(3.45 , N/A) (N/A , -0.01 , N/A)	3081.7	N/A	7.4286 [8.0000]	92.9% { 120.0% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 2051993	(4.45 , N/A) (N/A , 0.00 , N/A)	2046.6	N/A	3.9024 [4.0000]	97.6% { 117.1% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 1256616	(5.39 , N/A) (N/A , 0.02 , N/A)	1553.8	N/A	1.9598 [2.0000]	98.0% { 118.4% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 1081646	(6.20 , N/A) (N/A , 0.03 , N/A)	1219.9	N/A	1.8087 [2.0000]	90.4% { 119.9% }			
13C8_PFOA_EIS	(421.0 / 376.0) 1334218	(6.90 , N/A) (N/A , 0.03 , N/A)	950.8	N/A	1.9742 [2.0000]	98.7% { 125.7% }			
13C9_PFNA_EIS	(472.0 / 427.0) 599048	(7.50 , N/A) (N/A , 0.02 , N/A)	947.7	N/A	1.0289 [1.0000]	102.9% { 119.6% }			
13C6_PFDA_EIS	(519.0 / 474.0) 645193	(8.05 , N/A) (N/A , 0.01 , N/A)	1434.8	N/A	0.9379 [1.0000]	93.8% { 103.2% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 658904	(8.55 , N/A) (N/A , 0.02 , N/A)	1292.5	N/A	0.9668 [1.0000]	96.7% { 111.9% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT Ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 650323	(8.87 , N/A) (N/A , 0.02 , N/A)	896.2	N/A	1.0288 [1.0000]	102.9% { 145.2% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 651888	(9.25 , N/A) (N/A , 0.02 , N/A)	1349.0	N/A	0.9921 [1.0000]	99.2% { 125.9% }			
13C3_PFBs_EIS	(302.0 / 80.0) 4043026	(5.34 , N/A) (N/A , 0.01 , N/A)	2696.2	N/A	2.2540 [2.0000]	112.7% { 113.3% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 1943649	(7.03 , N/A) (N/A , 0.03 , N/A)	1243.0	N/A	2.0741 [2.0000]	103.7% { 111.0% }			
13C8_PFOS_EIS	(507.0 / 80.0) 3591314	(8.24 , N/A) (N/A , 0.01 , N/A)	464.3	N/A	1.8974 [2.0000]	94.9% { 109.7% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 459497	(5.13 , N/A) (N/A , 0.01 , N/A)	1133.6	N/A	4.1824 [4.0000]	104.6% { 114.0% }			
13C2_6:2FTS_EIS	(429.0 / 81.0) 513493	(6.64 , N/A) (N/A , 0.03 , N/A)	619.8	N/A	4.1455 [4.0000]	103.6% { 120.3% }			
13C2_8:2FTS_EIS	(529.0 / 81.0) 656842	(7.80 , N/A) (N/A , 0.02 , N/A)	1544.8	N/A	5.0145 [4.0000]	125.4% { 125.6% }			
13C8_PFOA_EIS	(506.0 / 78.0) 6340752	(9.84 , N/A) (N/A , 0.02 , N/A)	2678.1	N/A	1.8186 [2.0000]	90.9% { 109.5% }			
D3_NMeFOA_EIS	(515.0 / 169.0) 1493786	(10.43 , N/A) (N/A , 0.02 , N/A)	2808.1	N/A	1.9953 [2.0000]	99.8% { 112.4% }			
D5_NEtFOA_EIS	(531.0 / 169.0) 1305243	(10.59 , N/A) (N/A , 0.01 , N/A)	2664.0	N/A	1.9911 [2.0000]	99.6% { 107.7% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1198585	(8.20 , N/A) (N/A , 0.02 , N/A)	1042.2	N/A	4.5121 [4.0000]	112.8% { 108.0% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 912719	(8.45 , N/A) (N/A , 0.01 , N/A)	18948.3	N/A	4.0349 [4.0000]	100.9% { 108.3% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 5093047	(10.37 , N/A) (N/A , 0.02 , N/A)	1959.2	N/A	19.9819 [20.0000]	99.9% { 119.1% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 5960514	(10.53 , N/A) (N/A , 0.01 , N/A)	1338.5	N/A	20.4438 [20.0000]	102.2% { 111.6% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 3204565	(5.69 , N/A) (N/A , 0.02 , N/A)	1711.6	N/A	7.9536 [8.0000]	99.4% { 115.2% }			

INITIAL AND CONTINUING CALIBRATION CHECK

EPA 1633

Laboratory: APPL, LLC

Work Order: 23B0184

Client: AECOM

Project: Red Hill AFFF Assessment Sampling

Instrument ID: Saphira

Calibration: 2309009

Standard ID: 23B0084

Sequence: SC00861

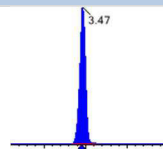
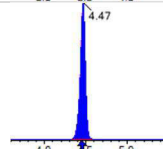
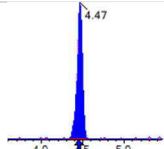
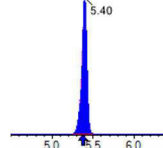
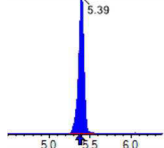
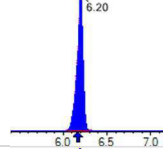
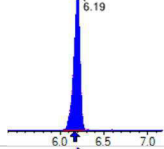
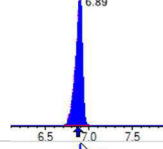
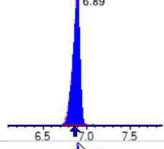
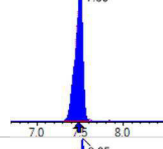
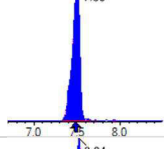
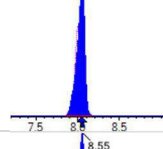
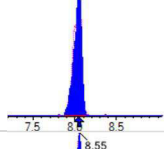
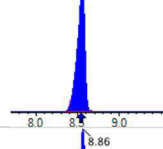
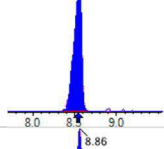
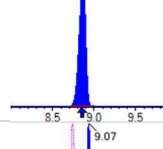
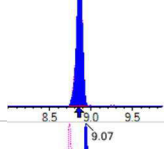
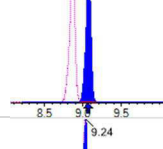
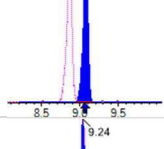
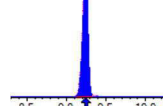
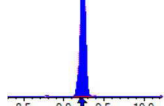
Lab Sample ID	Analyte	True	Found	%R	Units	Control Limit
SC00861-CCV3	PFBA	20.0	21.8	109	ng/mL	+/- 30.00%
	PFPEA	10.0	11.2	112	ng/mL	+/- 30.00%
	PFHXA	5.00	5.61	112	ng/mL	+/- 30.00%
	PFHPA	5.00	5.55	111	ng/mL	+/- 30.00%
	PFOA	5.00	5.55	111	ng/mL	+/- 30.00%
	PFNA	5.00	5.26	105	ng/mL	+/- 30.00%
	PFDA	5.00	4.99	99.9	ng/mL	+/- 30.00%
	PFUnA	5.00	4.93	98.5	ng/mL	+/- 30.00%
	PFDOA	5.00	5.47	109	ng/mL	+/- 30.00%
	PFTRDA	5.00	5.39	108	ng/mL	+/- 30.00%
	PFTEDA	5.00	5.09	102	ng/mL	+/- 30.00%
	PFBS	4.42	5.17	117	ng/mL	+/- 30.00%
	PFPEs	4.70	5.96	127	ng/mL	+/- 30.00%
	PFHXS	4.58	4.99	109	ng/mL	+/- 30.00%
	PFHPS	4.78	5.41	113	ng/mL	+/- 30.00%
	PFOS	4.65	5.04	108	ng/mL	+/- 30.00%
	PFNS	4.80	5.95	124	ng/mL	+/- 30.00%
	PFDS	4.82	5.84	121	ng/mL	+/- 30.00%
	PFDOS	4.85	5.83	120	ng/mL	+/- 30.00%
	4:2FTS	18.8	23.2	123	ng/mL	+/- 30.00%
	6:2FTS	19.0	23.6	124	ng/mL	+/- 30.00%
	8:2FTS	19.2	20.9	109	ng/mL	+/- 30.00%
	PFOSA	5.00	5.73	115	ng/mL	+/- 30.00%
	NMeFOSA	20.0	22.1	111	ng/mL	+/- 30.00%
	NEtFOSA	20.0	23.2	116	ng/mL	+/- 30.00%
	NMeFOSAA	5.00	5.20	104	ng/mL	+/- 30.00%
	NEtFOSAA	5.00	5.66	113	ng/mL	+/- 30.00%
	NMeFOSE	20.0	22.3	111	ng/mL	+/- 30.00%
	NEtFOSE	20.0	22.4	112	ng/mL	+/- 30.00%
	HFPO-DA	10.0	11.0	110	ng/mL	+/- 30.00%

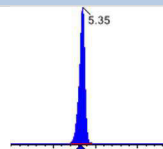
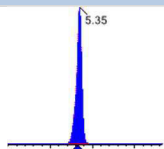
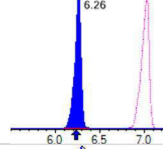
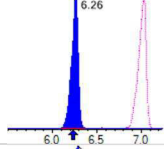
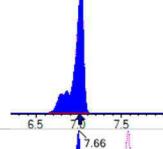
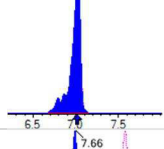
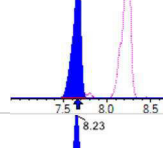
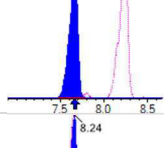
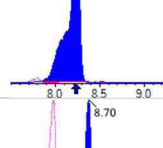
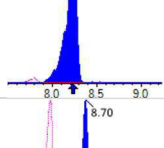
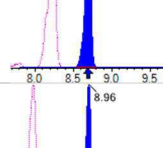
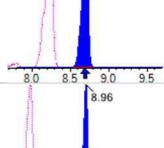
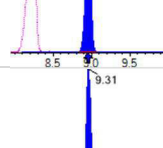
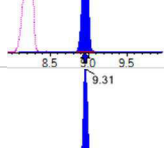
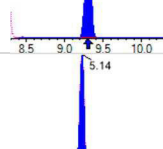
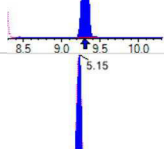
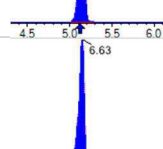
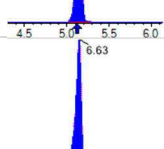
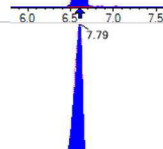
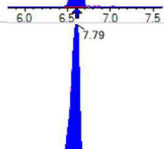
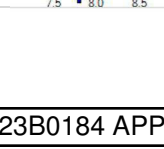
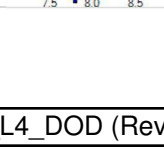
INITIAL AND CONTINUING CALIBRATION CHECK

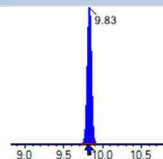
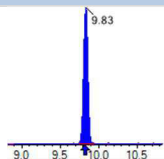
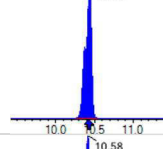
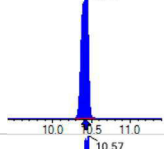
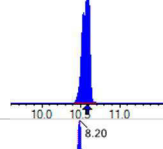
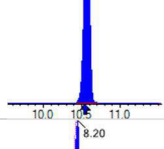
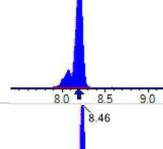
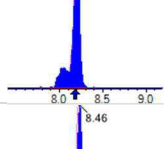
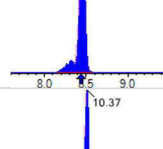
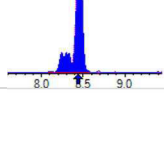
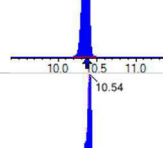
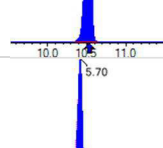
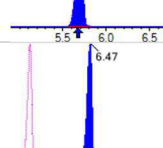
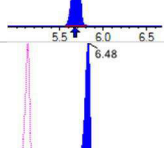
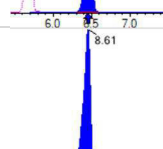
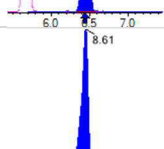
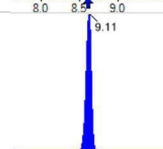
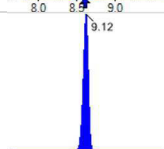
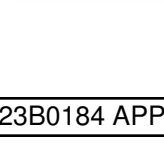
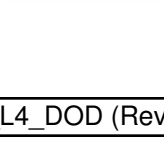
EPA 1633

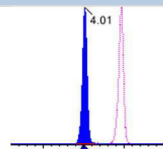
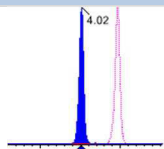
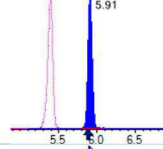
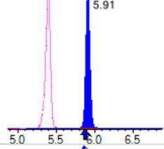
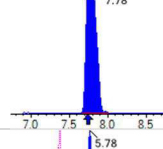
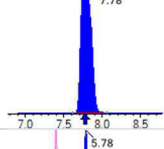
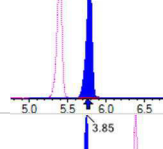
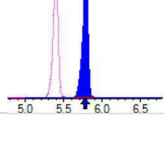
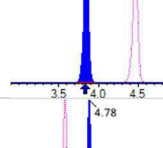
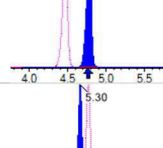
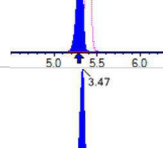
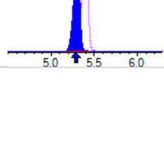
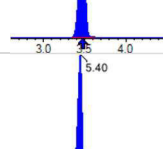
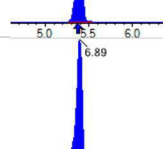
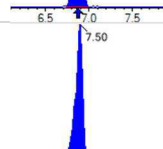
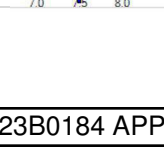
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Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Instrument ID:	Saphira	Calibration:	2309009
Standard ID:	23B0084	Sequence:	SC00861

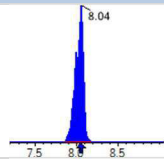
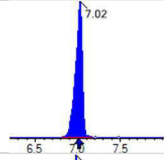
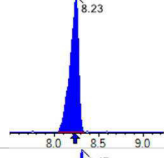
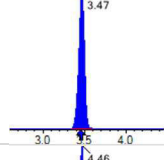
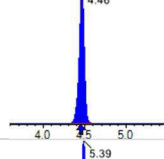
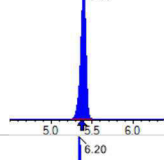
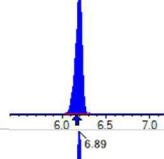
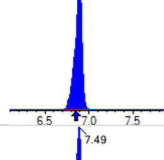
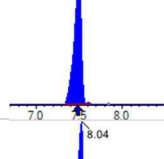
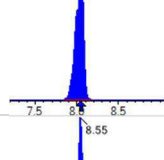
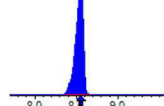
Lab Sample ID	Analyte	True	Found	%R	Units	Control Limit
SC00861-CCV3	ADONA	9.45	10.9	115	ng/mL	+/- 30.00%
	PFEESA	8.90	10.6	119	ng/mL	+/- 30.00%
	PFMPA	10.0	10.0	100	ng/mL	+/- 30.00%
	PFMBA	10.0	11.5	115	ng/mL	+/- 30.00%
	NFDHA	10.0	11.4	114	ng/mL	+/- 30.00%
	9CL-PF3ONS	9.35	12.1	130	ng/mL	+/- 30.00%
	11CL-PF3OUDS	9.45	11.7	123	ng/mL	+/- 30.00%
	3:3FTCA	20.0	19.1	95.7	ng/mL	+/- 30.00%
	5:3FTCA	20.0	19.6	97.8	ng/mL	+/- 30.00%
	7:3FTCA	20.0	19.7	98.7	ng/mL	+/- 30.00%

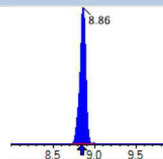
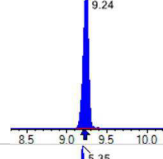
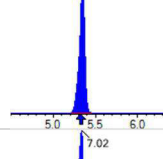
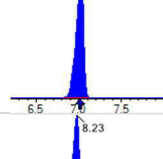
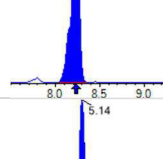
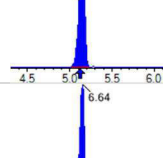
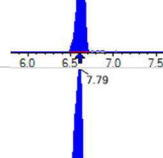
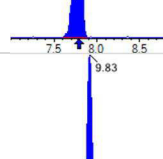
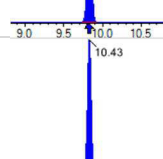
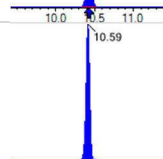
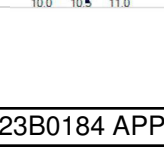
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT Ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) 5385910	(3.47, 1.00) (0.00, N/A, 0.0)	208.3	N/A 0.0 0.0	21.8119 [20.0000]	109.1%			
PFPeA	(263.0 / 219.0) 4888979 (263.0 / 69.0) 63781	(4.47, 1.00) (0.00, N/A, -0.1)	3485.6 377.3	0.0130 109.5 110.9	11.1574 [10.0000]	111.6%			
PFHxA	(313.0 / 269.0) 3220598 (313.0 / 119.0) 338608	(5.40, 1.00) (0.00, N/A, 0.1)	4977.1 2817.5	0.1051 112.4 101.8	5.6146 [5.0000]	112.3%			
PFHpA	(363.0 / 319.0) 2659586 (363.0 / 169.0) 811961	(6.20, 1.00) (0.00, N/A, 0.2)	5410.4 2364.0	0.3053 96.7 93.6	5.5543 [5.0000]	111.1%			
PFOA	(413.0 / 369.0) 3339357 (413.0 / 169.0) 1123065	(6.89, 1.00) (0.00, N/A, 0.1)	4556.2 87023.2	0.3363 99.4 94.6	5.5465 [5.0000]	110.9%			
PFNA	(463.0 / 419.0) 2947203 (463.0 / 169.0) 716993	(7.50, 1.00) (0.00, N/A, 0.1)	4457.9 800.0	0.2433 103.9 105.0	5.2598 [5.0000]	105.2%			
PFDA	(513.0 / 469.0) 3489148 (513.0 / 169.0) 405029	(8.05, 1.00) (0.01, N/A, 0.1)	1531.5 1122.6	0.1161 93.7 85.7	4.9930 [5.0000]	99.9%			
PFUnA	(563.0 / 519.0) 3070372 (563.0 / 169.0) 381470	(8.55, 1.00) (0.00, N/A, 0.2)	1690.3 705.0	0.1242 115.4 85.1	4.9254 [5.0000]	98.5%			
PFDoA	(613.0 / 569.0) 3025365 (613.0 / 169.0) 507186	(8.86, 1.00) (0.00, N/A, 0.1)	1415.9 1260.4	0.1676 105.5 88.8	5.4730 [5.0000]	109.5%			
PFTTrDA	(663.0 / 619.0) 2869547 (663.0 / 169.0) 699518	(9.07, 1.02) (N/A, 0.01, -0.2)	1751.9 916.1	0.2438 89.8 85.6	5.3947 [5.0000]	107.9%			
PFTeDA	(713.0 / 669.0) 2965307 (713.0 / 169.0) 674136	(9.24, 1.00) (0.00, N/A, 0.3)	2258.4 975.7	0.2273 104.0 103.2	5.0859 [5.0000]	101.7%			

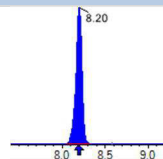
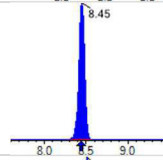
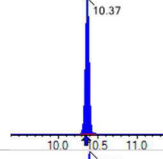
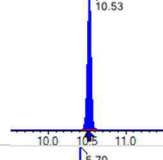
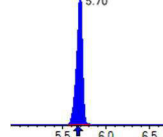
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) 5987621 (299.0 / 99.0) 3623340	(5.35 , 1.00) (0.00 , N/A , -0.1)	2227.8 2381.9	0.6051 96.2 97.0	5.1686 [4.4237]	116.8%			
PFPeS	(349.0 / 80.0) 10009015 (349.0 / 99.0) 3317972	(6.26 , 0.89) (N/A , 0.03 , 0.0)	249408.4 257124.5	0.3315 99.1 99.0	5.9614 [4.6919]	127.1%			
PFHxS	(399.0 / 80.0) 7353392 (399.0 / 99.0) 2496418	(7.02 , 1.00) (0.00 , N/A , 0.2)	5957.5 1522.5	0.3395 106.1 98.3	4.9917 [4.5549]	109.6%			
PFHpS	(449.0 / 80.0) 9032004 (449.0 / 99.0) 2463480	(7.66 , 0.93) (N/A , 0.01 , 0.3)	12243.9 5218.0	0.2728 100.5 97.5	5.4141 [4.7570]	113.8%			
PFOS	(499.0 / 80.0) 10297032 (499.0 / 99.0) 2405152	(8.23 , 1.00) (0.00 , N/A , -0.1)	419.8 469.0	0.2336 106.2 101.3	5.0362 [4.6375]	108.6%			
PFNS	(549.0 / 80.0) 12167193 (549.0 / 99.0) 2756629	(8.70 , 1.06) (N/A , 0.01 , 0.0)	38526.6 1243779.9	0.2266 94.2 98.3	5.9469 [4.7994]	123.9%			
PFDS	(599.0 / 80.0) 13274607 (599.0 / 99.0) 2734009	(8.96 , 1.09) (N/A , 0.01 , -0.1)	2100.0 1287.6	0.2060 101.8 96.5	5.8381 [4.8155]	121.2%			
PFDoS	(699.0 / 80.0) 6115918 (699.0 / 99.0) 1189735	(9.31 , 1.13) (N/A , 0.01 , 0.1)	1980.1 1348.6	0.1945 99.6 93.2	5.8265 [4.8478]	120.2%			
4:2FTS	(327.0 / 307.0) 7103675 (327.0 / 81.0) 4615285	(5.14 , 1.00) (0.00 , N/A , -0.2)	2160.6 1551.6	0.6497 93.8 102.5	23.1640 [18.6906]	123.9%			
6:2FTS	(427.0 / 407.0) 3743441 (427.0 / 81.0) 2895279	(6.63 , 1.00) (0.00 , N/A , 0.1)	2203.4 1472.4	0.7734 91.3 100.9	23.5846 [18.9808]	124.3%			
8:2FTS	(527.0 / 507.0) 3662986 (527.0 / 81.0) 2945706	(7.79 , 1.00) (0.00 , N/A , 0.2)	1113.0 1202.4	0.8042 98.6 100.8	20.9387 [19.1658]	109.3%			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) 15133365 (498.0 / 478.0) 310934	(9.83 , 1.00) (0.00 , N/A , -0.1)	2270.3 915.2	0.0205 104.6 100.8	5.7275 [5.0000]	114.5%			
NMeFOSA	(512.0 / 219.0) 14609009 (512.0 / 169.0) 12343375	(10.43 , 1.00) (0.00 , N/A , 1.0)	3087.2 4669.6	0.8449 102.3 101.0	22.1146 [20.0000]	110.6%			
NEIFOSA	(526.0 / 219.0) 15416905 (526.0 / 169.0) 19537072	(10.58 , 1.00) (-0.01 , N/A , 0.5)	16763.6 10125.9	1.2672 100.1 98.9	23.2279 [20.0000]	116.1%			
NMeFOSAA	(570.0 / 419.0) 1456834 (570.0 / 483.0) 709621	(8.20 , 1.00) (0.00 , N/A , -0.1)	1241.7 565.9	0.4871 99.3 110.0	5.1960 [5.0000]	103.9%			
NEIFOSAA	(584.0 / 419.0) 1196990 (584.0 / 526.0) 673466	(8.46 , 1.00) (0.01 , N/A , 0.0)	3220.3 1275.4	0.5626 99.6 109.8	5.6583 [5.0000]	113.2%			
NMeFOSE	(616.0 / 59.0) 6048063	(10.37 , 1.00) (0.01 , N/A , 0.0)	2737.0	N/A 0.0 0.0	22.2525 [20.0000]	111.3%			
NEtFOSE	(630.0 / 59.0) 7048496	(10.54 , 1.00) (0.01 , N/A , 0.0)	984.9	N/A 0.0 0.0	22.3501 [20.0000]	111.8%			
HFPO-DA	(285.0 / 169.0) 3493750 (285.0 / 185.0) 10013485	(5.70 , 1.00) (0.00 , N/A , 0.0)	1818.3 2423.4	2.8661 104.3 94.7	11.0062 [10.0000]	110.1%			
ADONA	(377.0 / 85.0) 12834142 (377.0 / 251.0) 1138685	(6.47 , 1.14) (N/A , 0.03 , -0.2)	2150.4 1041.8	0.0887 101.4 102.0	10.8938 [9.4270]	115.6%			
9CI-Pf3ONS	(531.0 / 351.0) 33472612 (533.0 / 353.0) 10461709	(8.61 , 1.51) (N/A , 0.01 , 0.0)	1812.8 1850.3	0.3125 106.1 100.4	12.1146 [9.3325]	129.8%			
11CI-PF3OUDS	(631.0 / 451.0) 15852254 (633.0 / 453.0) 4777699	(9.11 , 1.60) (N/A , 0.01 , -0.1)	2994.5 2467.2	0.3014 97.7 97.4	11.6696 [9.4321]	123.7%			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) 247760 (241.0 / 117.0) 402136	(4.01, 0.90) (N/A, 0.01, -0.1)	1304.6 1537.5	1.6231 93.6 98.7	19.1463 [20.0000]	95.7%			
5:3FTCA	(341.0 / 236.7) 1604009 (341.0 / 217.0) 2873653	(5.91, 1.10) (N/A, 0.03, -0.1)	2397.6 2144.5	1.7915 106.5 98.6	19.5615 [20.0000]	97.8%			
7:3FTCA	(441.0 / 317.0) 2911284 (441.0 / 337.0) 2529915	(7.78, 1.44) (N/A, 0.01, -0.2)	954.7 907.3	0.8690 101.2 103.5	19.7486 [20.0000]	98.7%			
PFEESA	(315.0 / 135.0) 7522293 (315.0 / 83.0) 1994785	(5.78, 1.07) (N/A, 0.03, 0.0)	2385.7 1456.8	0.2652 94.2 101.3	10.5531 [8.9246]	118.2%			
PFMPA	(229.0 / 85.0) 1123834	(3.85, 0.86) (N/A, 0.01, 0.0)	3777.1	N/A 0.0 0.0	10.0312 [10.0000]	100.3%			
PFMBA	(279.0 / 85.0) 4426525	(4.78, 1.07) (N/A, 0.01, 0.0)	2199.5	N/A 0.0 0.0	11.5329 [10.0000]	115.3%			
NFDHA	(295.0 / 201.0) 3621281 (295.0 / 85.0) 3781252	(5.30, 0.98) (N/A, 0.02, 0.1)	2369.5 1979.0	1.0442 100.0 100.7	11.3504 [10.0000]	113.5%			
13C3_PFBa_IIS	(216.0 / 172.0) 268140	(3.47, N/A) (N/A, 0.01, N/A)	1205.9	N/A	0.7909 [1.0000]	79.1% { 128.0% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 524905	(5.40, N/A) (N/A, 0.03, N/A)	982.8	N/A	0.8825 [1.0000]	88.3% { 140.9% }			
13C4_PFOA_IIS	(417.0 / 372.0) 625721	(6.89, N/A) (N/A, 0.02, N/A)	1025.5	N/A	0.7692 [1.0000]	76.9% { 120.7% }			
13C5_PFNAl_IIS	(468.0 / 423.0) 555435	(7.50, N/A) (N/A, 0.02, N/A)	808.6	N/A	0.7274 [1.0000]	72.7% { 100.8% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 563220	(8.04 , N/A) (N/A , 0.01 , N/A)	4123.3	N/A	0.7846 [1.0000]	78.5% { 117.7% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 930174	(7.02 , N/A) (N/A , 0.02 , N/A)	661.9	N/A	0.7553 [1.0000]	75.5% { 105.1% }			
13C4_PFOS_IIS	(503.0 / 79.9) 1524597	(8.23 , N/A) (N/A , 0.01 , N/A)	853.1	N/A	0.8590 [1.0000]	85.9% { 118.2% }			
13C4_PFBA_EIS	(217.0 / 172.0) 2335882	(3.47 , N/A) (N/A , 0.01 , N/A)	2821.5	N/A	7.3261 [8.0000]	91.6% { 122.4% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 1989165	(4.46 , N/A) (N/A , 0.02 , N/A)	2700.9	N/A	3.4010 [4.0000]	85.0% { 113.5% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 1229303	(5.39 , N/A) (N/A , 0.03 , N/A)	2356.0	N/A	1.7236 [2.0000]	86.2% { 115.8% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 1072429	(6.20 , N/A) (N/A , 0.03 , N/A)	980.3	N/A	1.6123 [2.0000]	80.6% { 118.9% }			
13C8_PFOA_EIS	(421.0 / 376.0) 1249686	(6.89 , N/A) (N/A , 0.03 , N/A)	1361.9	N/A	1.8660 [2.0000]	93.3% { 117.7% }			
13C9_PFNA_EIS	(472.0 / 427.0) 593192	(7.49 , N/A) (N/A , 0.01 , N/A)	1039.5	N/A	1.0717 [1.0000]	107.2% { 118.4% }			
13C6_PFDA_EIS	(519.0 / 474.0) 704823	(8.04 , N/A) (N/A , 0.01 , N/A)	1139.4	N/A	1.0812 [1.0000]	108.1% { 112.7% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 708894	(8.55 , N/A) (N/A , 0.01 , N/A)	1472.6	N/A	1.0977 [1.0000]	109.8% { 120.4% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT Ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 633598	(8.86 , N/A) (N/A , 0.02 , N/A)	1586.7	N/A	1.0578 [1.0000]	105.8% { 141.5% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 634092	(9.24 , N/A) (N/A , 0.01 , N/A)	888.2	N/A	1.0184 [1.0000]	101.8% { 122.5% }			
13C3_PFBs_EIS	(302.0 / 80.0) 4074883	(5.35 , N/A) (N/A , 0.02 , N/A)	2444.1	N/A	2.1679 [2.0000]	108.4% { 114.2% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 1870380	(7.02 , N/A) (N/A , 0.02 , N/A)	1365.6	N/A	1.9046 [2.0000]	95.2% { 106.8% }			
13C8_PFOS_EIS	(507.0 / 80.0) 3705322	(8.23 , N/A) (N/A , 0.01 , N/A)	496.0	N/A	1.8499 [2.0000]	92.5% { 113.2% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 434605	(5.14 , N/A) (N/A , 0.02 , N/A)	1243.3	N/A	3.7749 [4.0000]	94.4% { 107.8% }			
13C2_6:2FTS_EIS	(429.0 / 81.0) 486669	(6.64 , N/A) (N/A , 0.03 , N/A)	614.9	N/A	3.7492 [4.0000]	93.7% { 114.1% }			
13C2_8:2FTS_EIS	(529.0 / 81.0) 623330	(7.79 , N/A) (N/A , 0.01 , N/A)	1282.4	N/A	4.5410 [4.0000]	113.5% { 119.2% }			
13C8_PFOsa_EIS	(506.0 / 78.0) 6800198	(9.83 , N/A) (N/A , 0.01 , N/A)	2345.0	N/A	1.8431 [2.0000]	92.2% { 117.4% }			
D3_NMeFOSA_EIS	(515.0 / 169.0) 1538374	(10.43 , N/A) (N/A , 0.01 , N/A)	2048.3	N/A	1.9418 [2.0000]	97.1% { 115.8% }			
D5_NeFOSA_EIS	(531.0 / 169.0) 1366435	(10.59 , N/A) (N/A , 0.01 , N/A)	3656.1	N/A	1.9698 [2.0000]	98.5% { 112.8% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1312203	(8.20 , N/A) (N/A , 0.01 , N/A)	1044.4	N/A	4.6680 [4.0000]	116.7% { 118.2% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 970686	(8.45 , N/A) (N/A , 0.01 , N/A)	3642.8	N/A	4.0551 [4.0000]	101.4% { 115.2% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 5187812	(10.37 , N/A) (N/A , 0.01 , N/A)	1872.1	N/A	19.2339 [20.0000]	96.2% { 121.3% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 6242065	(10.53 , N/A) (N/A , 0.01 , N/A)	1319.1	N/A	20.2316 [20.0000]	101.2% { 116.9% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 3097562	(5.70 , N/A) (N/A , 0.03 , N/A)	1741.3	N/A	6.9118 [8.0000]	86.4% { 111.4% }			

INITIAL AND CONTINUING CALIBRATION CHECK

EPA 1633

Laboratory: APPL, LLC

Work Order: 23B0184

Client: AECOM

Project: Red Hill AFFF Assessment Sampling

Instrument ID: Saphira

Calibration: 2309009

Standard ID: 23B0084

Sequence: SC00861

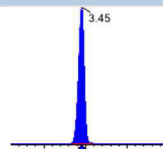
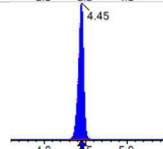
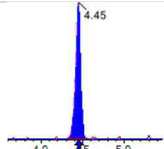
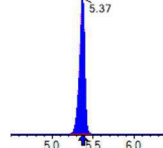
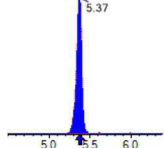
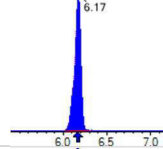
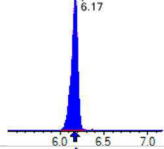
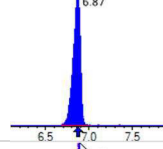
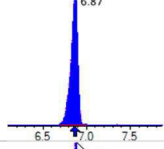
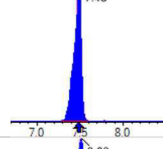
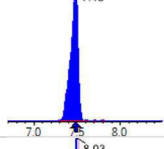
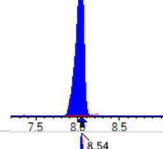
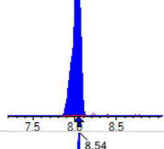
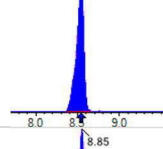
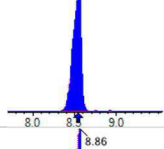
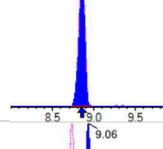
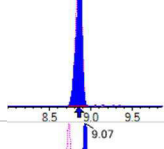
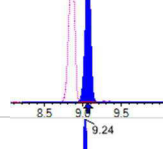
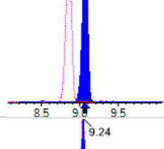
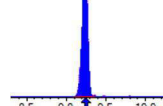
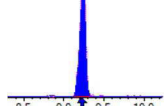
Lab Sample ID	Analyte	True	Found	%R	Units	Control Limit
SC00861-CCV4	PFBA	20.0	22.6	113	ng/mL	+/- 30.00%
	PFPEA	10.0	10.9	109	ng/mL	+/- 30.00%
	PFHXA	5.00	5.45	109	ng/mL	+/- 30.00%
	PFHPA	5.00	5.19	104	ng/mL	+/- 30.00%
	PFOA	5.00	5.38	108	ng/mL	+/- 30.00%
	PFNA	5.00	5.01	100	ng/mL	+/- 30.00%
	PFDA	5.00	5.26	105	ng/mL	+/- 30.00%
	PFUnA	5.00	5.32	106	ng/mL	+/- 30.00%
	PFDOA	5.00	5.54	111	ng/mL	+/- 30.00%
	PFTRDA	5.00	5.57	111	ng/mL	+/- 30.00%
	PFTEDA	5.00	5.07	101	ng/mL	+/- 30.00%
	PFBS	4.42	4.94	112	ng/mL	+/- 30.00%
	PFPEs	4.70	5.93	126	ng/mL	+/- 30.00%
	PFHXS	4.58	5.39	118	ng/mL	+/- 30.00%
	PFHPS	4.78	5.00	105	ng/mL	+/- 30.00%
	PFOS	4.65	4.99	107	ng/mL	+/- 30.00%
	PFNS	4.80	5.98	125	ng/mL	+/- 30.00%
	PFDS	4.82	5.65	117	ng/mL	+/- 30.00%
	PFDOS	4.85	5.72	118	ng/mL	+/- 30.00%
	4:2FTS	18.8	21.8	116	ng/mL	+/- 30.00%
	6:2FTS	19.0	24.4	129	ng/mL	+/- 30.00%
	8:2FTS	19.2	20.0	104	ng/mL	+/- 30.00%
	PFOSA	5.00	5.74	115	ng/mL	+/- 30.00%
	NMeFOSA	20.0	22.3	111	ng/mL	+/- 30.00%
	NEtFOSA	20.0	22.9	114	ng/mL	+/- 30.00%
	NMeFOSAA	5.00	5.13	103	ng/mL	+/- 30.00%
	NEtFOSAA	5.00	5.38	108	ng/mL	+/- 30.00%
	NMeFOSE	20.0	21.9	109	ng/mL	+/- 30.00%
	NEtFOSE	20.0	22.4	112	ng/mL	+/- 30.00%
	HFPO-DA	10.0	11.0	110	ng/mL	+/- 30.00%

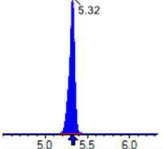
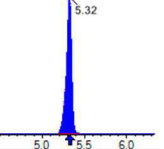
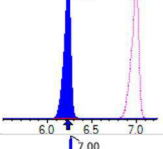
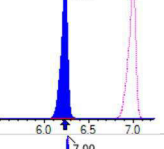
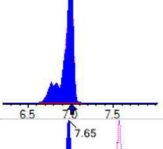
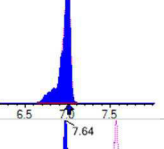
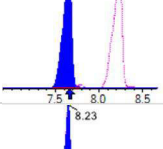
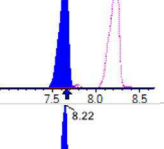
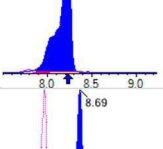
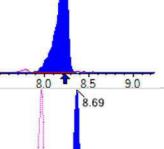
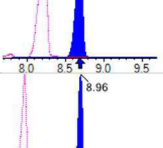
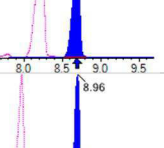
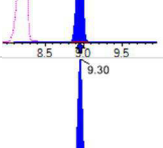
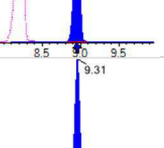
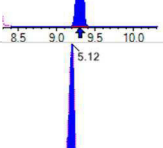
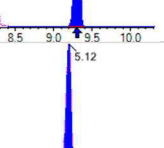
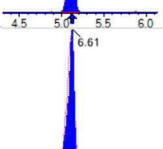
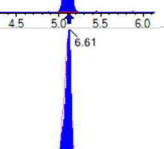
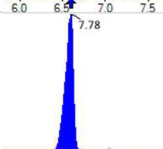
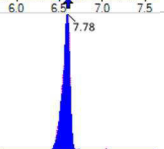

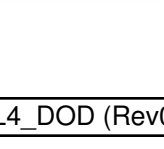
INITIAL AND CONTINUING CALIBRATION CHECK

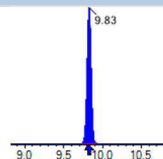
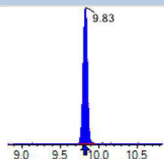
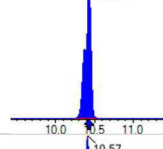
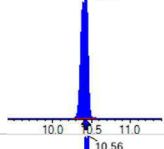
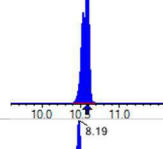
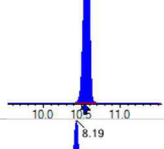
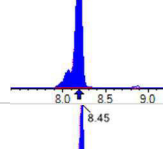
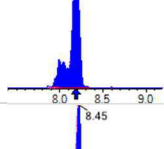
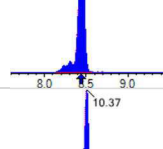
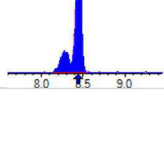
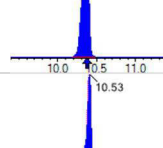
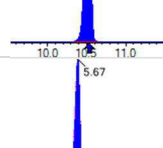
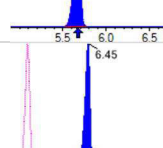
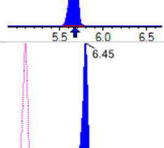
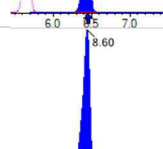
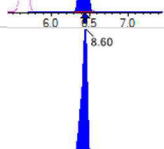
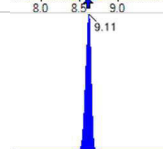
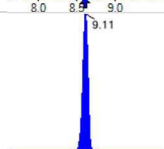
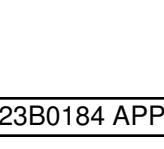
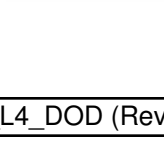
EPA 1633

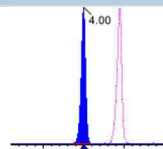
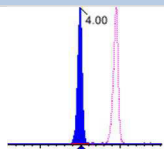
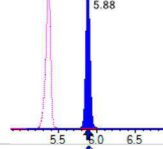
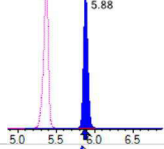
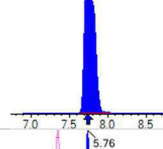
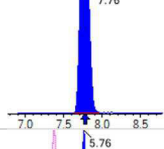
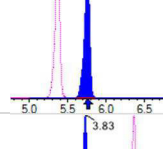
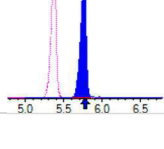
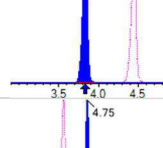
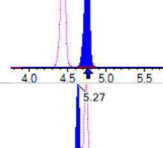
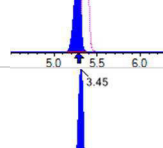
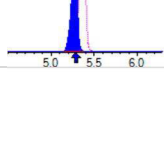
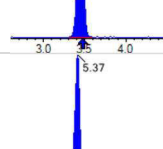
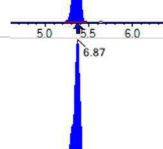
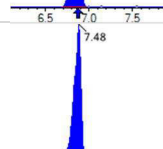
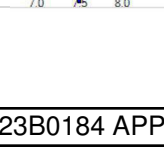
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Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Instrument ID:	Saphira	Calibration:	2309009
Standard ID:	23B0084	Sequence:	SC00861

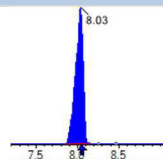
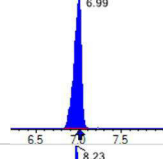
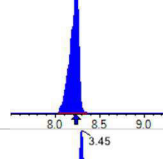
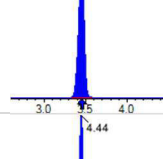
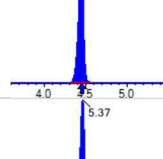
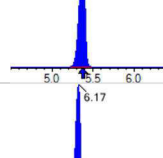
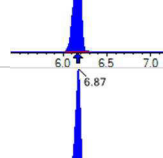
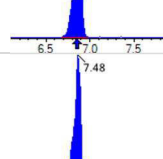
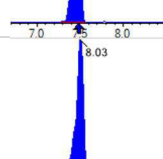
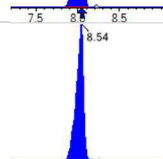
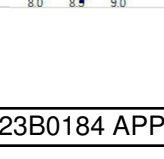
Lab Sample ID	Analyte	True	Found	%R	Units	Control Limit
SC00861-CCV4	ADONA	9.45	11.1	118	ng/mL	+/- 30.00%
	PFEESA	8.90	10.4	117	ng/mL	+/- 30.00%
	PFMPA	10.0	9.98	99.8	ng/mL	+/- 30.00%
	PFMBA	10.0	11.9	119	ng/mL	+/- 30.00%
	NFDHA	10.0	11.3	113	ng/mL	+/- 30.00%
	9CL-PF3ONS	9.35	12.1	130	ng/mL	+/- 30.00%
	11CL-PF3OUDS	9.45	11.1	118	ng/mL	+/- 30.00%
	3:3FTCA	20.0	19.4	96.9	ng/mL	+/- 30.00%
	5:3FTCA	20.0	19.0	94.9	ng/mL	+/- 30.00%
	7:3FTCA	20.0	19.6	97.8	ng/mL	+/- 30.00%

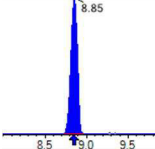
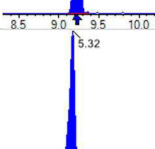
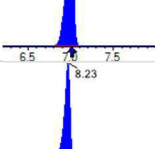
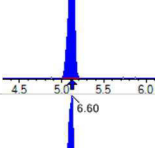
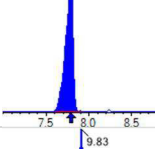
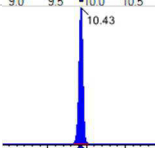
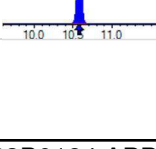
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT Ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) 5495975	(3.45, 1.00) (0.00, N/A, 0.0)	188.5	N/A 0.0 0.0	22.5736 [20.0000]	112.9%			
PFPeA	(263.0 / 219.0) 4817856 (263.0 / 69.0) 57835	(4.45, 1.00) (0.00, N/A, -0.1)	3292.8 533.9	0.0120 100.7 102.0	10.9250 [10.0000]	109.2%			
PFHxA	(313.0 / 269.0) 3160015 (313.0 / 119.0) 339199	(5.37, 1.00) (0.00, N/A, 0.1)	3793.8 8948239.1	0.1073 114.7 103.9	5.4453 [5.0000]	108.9%			
PFHpA	(363.0 / 319.0) 2567670 (363.0 / 169.0) 882578	(6.17, 1.00) (0.00, N/A, 0.1)	21766.2 2368.8	0.3437 108.9 105.4	5.1880 [5.0000]	103.8%			
PFOA	(413.0 / 369.0) 3211057 (413.0 / 169.0) 1082764	(6.87, 1.00) (0.00, N/A, -0.1)	6159.9 1281.3	0.3372 99.7 94.9	5.3838 [5.0000]	107.7%			
PFNA	(463.0 / 419.0) 2882055 (463.0 / 169.0) 683504	(7.48, 1.00) (0.00, N/A, 0.2)	9557.2 4860125.8	0.2372 101.3 102.3	5.0125 [5.0000]	100.2%			
PFDA	(513.0 / 469.0) 3464415 (513.0 / 169.0) 370919	(8.03, 1.00) (0.00, N/A, 0.0)	1188.1 988.2	0.1071 86.4 79.1	5.2625 [5.0000]	105.2%			
PFUnA	(563.0 / 519.0) 3140575 (563.0 / 169.0) 404619	(8.54, 1.00) (0.00, N/A, 0.0)	1250.8 1016.7	0.1288 119.7 88.2	5.3197 [5.0000]	106.4%			
PFDoA	(613.0 / 569.0) 2850779 (613.0 / 169.0) 530137	(8.85, 1.00) (0.00, N/A, -0.3)	1310.2 1056.1	0.1860 117.0 98.5	5.5429 [5.0000]	110.9%			
PFTTrDA	(663.0 / 619.0) 2758853 (663.0 / 169.0) 726697	(9.06, 1.02) (N/A, 0.01, -0.1)	1411.5 1001.0	0.2634 97.0 92.5	5.5744 [5.0000]	111.5%			
PFTeDA	(713.0 / 669.0) 2770969 (713.0 / 169.0) 609192	(9.24, 1.00) (0.00, N/A, -0.2)	2419.0 815.3	0.2198 100.6 99.8	5.0700 [5.0000]	101.4%			

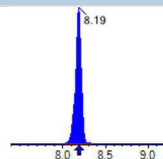
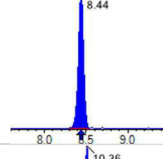
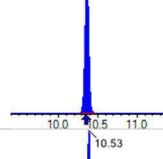
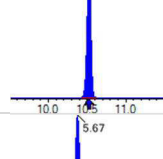
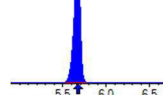
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) 5821062 (299.0 / 99.0) 3547103	(5.32 , 1.00) (0.00 , N/A , 0.0)	2282.1 2532.1	0.6094 96.9 97.6	4.9413 [4.4237]	111.7%			
PFPeS	(349.0 / 80.0) 9673917 (349.0 / 99.0) 3358853	(6.23 , 0.89) (N/A , 0.00 , 0.0)	32024.2 128055.8	0.3472 103.8 103.7	5.9278 [4.6919]	126.3%			
PFHxS	(399.0 / 80.0) 7718896 (399.0 / 99.0) 2455933	(7.00 , 1.00) (0.00 , N/A , 0.0)	6673.9 1751.4	0.3182 99.4 92.1	5.3908 [4.5549]	118.4%			
PFHpS	(449.0 / 80.0) 8663025 (449.0 / 99.0) 2504072	(7.65 , 0.93) (N/A , -0.01 , 0.2)	16391.7 186125.0	0.2891 106.5 103.3	5.0010 [4.7570]	105.1%			
PFOS	(499.0 / 80.0) 10587978 (499.0 / 99.0) 2277640	(8.23 , 1.00) (0.00 , N/A , 0.2)	393.7 508.9	0.2151 97.8 93.3	4.9871 [4.6375]	107.5%			
PFNS	(549.0 / 80.0) 12705823 (549.0 / 99.0) 2762341	(8.69 , 1.06) (N/A , 0.00 , 0.1)	32069.9 1682537.6	0.2174 90.3 94.3	5.9807 [4.7994]	124.6%			
PFDS	(599.0 / 80.0) 13347356 (599.0 / 99.0) 2687461	(8.96 , 1.09) (N/A , 0.01 , 0.1)	2401.9 1773.7	0.2013 99.5 94.3	5.6532 [4.8155]	117.4%			
PFDoS	(699.0 / 80.0) 6236153 (699.0 / 99.0) 1214730	(9.30 , 1.13) (N/A , 0.00 , -0.2)	2452.5 1375.9	0.1948 99.7 93.3	5.7215 [4.8478]	118.0%			
4:2FTS	(327.0 / 307.0) 6893698 (327.0 / 81.0) 4692872	(5.12 , 1.00) (0.00 , N/A , 0.2)	2360.9 1711.7	0.6807 98.2 107.4	21.7718 [18.6906]	116.5%			
6:2FTS	(427.0 / 407.0) 3592335 (427.0 / 81.0) 3044265	(6.61 , 1.00) (0.01 , N/A , 0.1)	1359.4 1723.0	0.8474 100.0 110.6	24.4488 [18.9808]	128.8%			
8:2FTS	(527.0 / 507.0) 3513879 (527.0 / 81.0) 2841665	(7.78 , 1.00) (0.00 , N/A , 0.0)	2185.7 1717.3	0.8087 99.1 101.4	20.0238 [19.1658]	104.5%			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) 15526995 (498.0 / 478.0) 313898	(9.83 , 1.00) (0.00 , N/A , 0.0)	3645.5 978.0	0.0202 102.9 99.1	5.7396 [5.0000]	114.8%			
NMeFOSA	(512.0 / 219.0) 14237667 (512.0 / 169.0) 11898638	(10.43 , 1.00) (0.00 , N/A , 1.0)	3979.9 6344.7	0.8357 101.2 99.9	22.2774 [20.0000]	111.4%			
NEIFOSA	(526.0 / 219.0) 15536639 (526.0 / 169.0) 19789072	(10.57 , 1.00) (-0.01 , N/A , 0.5)	13762.0 15403.8	1.2737 100.6 99.4	22.8584 [20.0000]	114.3%			
NMeFOSAA	(570.0 / 419.0) 1372293 (570.0 / 483.0) 626986	(8.19 , 1.00) (0.00 , N/A , 0.1)	1536.4 391.4	0.4569 93.2 103.1	5.1283 [5.0000]	102.6%			
NEIFOSAA	(584.0 / 419.0) 1236420 (584.0 / 526.0) 718046	(8.45 , 1.00) (0.01 , N/A , 0.0)	3064.5 1624.8	0.5807 102.8 113.4	5.3767 [5.0000]	107.5%			
NMeFOSE	(616.0 / 59.0) 6197344	(10.37 , 1.00) (0.01 , N/A , 0.0)	3028.1	N/A 0.0 0.0	21.8638 [20.0000]	109.3%			
NEtFOSE	(630.0 / 59.0) 7496584	(10.53 , 1.00) (0.01 , N/A , 0.0)	1130.0	N/A 0.0 0.0	22.3714 [20.0000]	111.9%			
HFPO-DA	(285.0 / 169.0) 3514895 (285.0 / 185.0) 9786018	(5.67 , 1.00) (0.00 , N/A , 0.0)	2040.8 2349.1	2.7842 101.3 92.0	11.0375 [10.0000]	110.4%			
ADONA	(377.0 / 85.0) 13148845 (377.0 / 251.0) 1125005	(6.45 , 1.14) (N/A , 0.00 , 0.0)	2272.3 1375.9	0.0856 97.7 98.4	11.1253 [9.4270]	118.0%			
9CI-Pf3ONS	(531.0 / 351.0) 33649260 (533.0 / 353.0) 10616742	(8.60 , 1.52) (N/A , 0.00 , -0.2)	1835.2 1998.8	0.3155 107.1 101.4	12.1396 [9.3325]	130.1%			QC,
11CI-PF3OUDS	(631.0 / 451.0) 15190575 (633.0 / 453.0) 4756996	(9.11 , 1.61) (N/A , 0.01 , 0.1)	1759.1 2589.8	0.3132 101.5 101.2	11.1468 [9.4321]	118.2%			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) 252517 (241.0 / 117.0) 385892	(4.00, 0.90) (N/A, 0.00, 0.1)	1253.2 1037.3	1.5282 88.1 92.9	19.3894 [20.0000]	96.9%			
5:3FTCA	(341.0 / 236.7) 1574440 (341.0 / 217.0) 2760527	(5.88, 1.10) (N/A, 0.00, 0.0)	2670.3 2019.2	1.7533 104.3 96.5	18.9791 [20.0000]	94.9%			
7:3FTCA	(441.0 / 317.0) 2918398 (441.0 / 337.0) 2561802	(7.76, 1.45) (N/A, -0.01, -0.3)	862.1 895.5	0.8778 102.2 104.6	19.5681 [20.0000]	97.8%			
PFEESA	(315.0 / 135.0) 7500399 (315.0 / 83.0) 1944360	(5.76, 1.07) (N/A, 0.00, -0.1)	2331.2 1393.9	0.2592 92.1 99.0	10.4008 [8.9246]	116.5%			
PFMPA	(229.0 / 85.0) 1124886	(3.83, 0.86) (N/A, 0.00, 0.0)	4137.3	N/A 0.0 0.0	9.9766 [10.0000]	99.8%			
PFMBA	(279.0 / 85.0) 4589021	(4.75, 1.07) (N/A, -0.01, 0.0)	2547.5	N/A 0.0 0.0	11.8800 [10.0000]	118.8%			
NFDHA	(295.0 / 201.0) 3658756 (295.0 / 85.0) 3775034	(5.27, 0.98) (N/A, 0.00, 0.1)	1887.0 2115.0	1.0318 98.8 99.5	11.3354 [10.0000]	113.4%			
13C3_PFBa_IIS	(216.0 / 172.0) 247053	(3.45, N/A) (N/A, -0.01, N/A)	902.7	N/A	0.7287 [1.0000]	72.9% { 118.0% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 488231	(5.37, N/A) (N/A, 0.00, N/A)	1034.2	N/A	0.8209 [1.0000]	82.1% { 131.0% }			
13C4_PFOA_IIS	(417.0 / 372.0) 604761	(6.87, N/A) (N/A, 0.00, N/A)	814.8	N/A	0.7435 [1.0000]	74.3% { 116.7% }			
13C5_PFNAl_IIS	(468.0 / 423.0) 618985	(7.48, N/A) (N/A, 0.00, N/A)	1087.3	N/A	0.8106 [1.0000]	81.1% { 112.4% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 593602	(8.03 , N/A) (N/A , 0.00 , N/A)	859.8	N/A	0.8269 [1.0000]	82.7% { 124.0% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 949244	(6.99 , N/A) (N/A , 0.00 , N/A)	1081.5	N/A	0.7708 [1.0000]	77.1% { 107.3% }			
13C4_PFOS_IIS	(503.0 / 79.9) 1528447	(8.23 , N/A) (N/A , 0.00 , N/A)	786.1	N/A	0.8612 [1.0000]	86.1% { 118.5% }			
13C4_PFBA_EIS	(217.0 / 172.0) 2303187	(3.45 , N/A) (N/A , -0.01 , N/A)	3079.9	N/A	7.8401 [8.0000]	98.0% { 120.7% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 2001932	(4.44 , N/A) (N/A , -0.01 , N/A)	2095.7	N/A	3.6800 [4.0000]	92.0% { 114.2% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 1243670	(5.37 , N/A) (N/A , 0.00 , N/A)	1596.4	N/A	1.8748 [2.0000]	93.7% { 117.1% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 1108453	(6.17 , N/A) (N/A , 0.01 , N/A)	1274.3	N/A	1.7916 [2.0000]	89.6% { 122.9% }			
13C8_PFOA_EIS	(421.0 / 376.0) 1237996	(6.87 , N/A) (N/A , 0.00 , N/A)	1361.8	N/A	1.9126 [2.0000]	95.6% { 116.6% }			
13C9_PFNA_EIS	(472.0 / 427.0) 608706	(7.48 , N/A) (N/A , -0.01 , N/A)	1113.4	N/A	0.9868 [1.0000]	98.7% { 121.5% }			
13C6_PFDA_EIS	(519.0 / 474.0) 663988	(8.03 , N/A) (N/A , 0.00 , N/A)	1011.8	N/A	0.9664 [1.0000]	96.6% { 106.2% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 671355	(8.54 , N/A) (N/A , 0.00 , N/A)	1128.8	N/A	0.9863 [1.0000]	98.6% { 114.0% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 589511	(8.85 , N/A) (N/A , 0.01 , N/A)	1124.5	N/A	0.9338 [1.0000]	93.4% { 131.6% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 594389	(9.24 , N/A) (N/A , 0.01 , N/A)	713.1	N/A	0.9058 [1.0000]	90.6% { 114.8% }			
13C3_PFBs_EIS	(302.0 / 80.0) 4143709	(5.32 , N/A) (N/A , 0.00 , N/A)	2058.3	N/A	2.1602 [2.0000]	108.0% { 116.1% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 1818000	(7.00 , N/A) (N/A , 0.00 , N/A)	991.4	N/A	1.8141 [2.0000]	90.7% { 103.8% }			
13C8_PFOS_EIS	(507.0 / 80.0) 3847519	(8.23 , N/A) (N/A , 0.00 , N/A)	593.4	N/A	1.9161 [2.0000]	95.8% { 117.5% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 448728	(5.12 , N/A) (N/A , 0.00 , N/A)	843.1	N/A	3.8193 [4.0000]	95.5% { 111.3% }			
13C2_6:2FTS_EIS	(429.0 / 81.0) 450516	(6.60 , N/A) (N/A , 0.00 , N/A)	1307.5	N/A	3.4010 [4.0000]	85.0% { 105.6% }			
13C2_8:2FTS_EIS	(529.0 / 81.0) 625278	(7.78 , N/A) (N/A , 0.00 , N/A)	926.4	N/A	4.4637 [4.0000]	111.6% { 119.6% }			
13C8_PFOsa_EIS	(506.0 / 78.0) 6962278	(9.83 , N/A) (N/A , 0.01 , N/A)	2923.1	N/A	1.8823 [2.0000]	94.1% { 120.2% }			
D3_NMeFOsa_EIS	(515.0 / 169.0) 1488313	(10.43 , N/A) (N/A , 0.01 , N/A)	2638.5	N/A	1.8739 [2.0000]	93.7% { 112.0% }			
D5_NEtFOsa_EIS	(531.0 / 169.0) 1399307	(10.58 , N/A) (N/A , 0.01 , N/A)	2373.5	N/A	2.0121 [2.0000]	100.6% { 115.5% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1252372	(8.19 , N/A) (N/A , 0.00 , N/A)	1220.3	N/A	4.4440 [4.0000]	111.1% { 112.8% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 1055171	(8.44 , N/A) (N/A , 0.00 , N/A)	18412.0	N/A	4.3970 [4.0000]	109.9% { 125.3% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 5410360	(10.36 , N/A) (N/A , 0.01 , N/A)	2015.6	N/A	20.0085 [20.0000]	100.0% { 126.5% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 6632572	(10.53 , N/A) (N/A , 0.01 , N/A)	1177.6	N/A	21.4431 [20.0000]	107.2% { 124.2% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 3107490	(5.67 , N/A) (N/A , 0.00 , N/A)	1834.2	N/A	7.4548 [8.0000]	93.2% { 111.7% }			

INITIAL AND CONTINUING CALIBRATION CHECK

EPA 1633

Laboratory: APPL, LLC

Work Order: 23B0184

Client: AECOM

Project: Red Hill AFFF Assessment Sampling

Instrument ID: Saphira

Calibration: 2309009

Standard ID: 23B0084

Sequence: SC00861

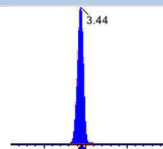
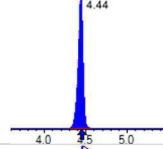
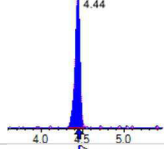
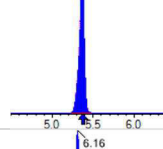
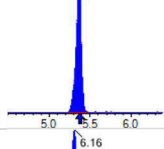
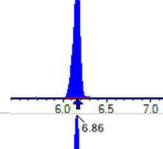
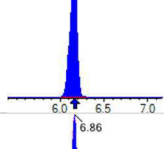
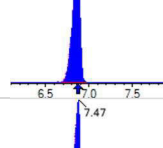
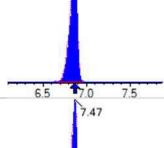
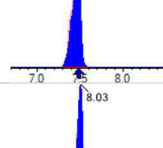
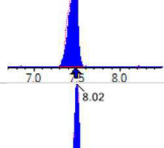
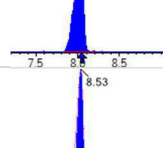
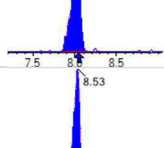
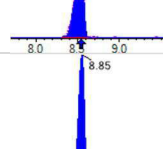
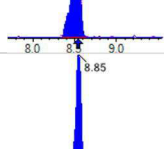
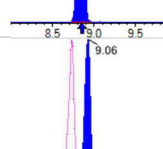
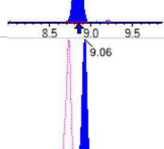
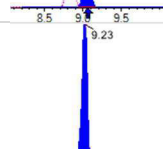
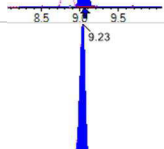
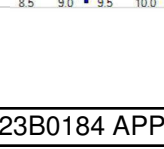
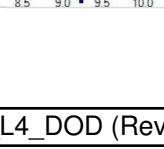
Lab Sample ID	Analyte	True	Found	%R	Units	Control Limit
SC00861-CCV5	PFBA	20.0	22.1	110	ng/mL	+/- 30.00%
	PFPEA	10.0	10.7	107	ng/mL	+/- 30.00%
	PFHXA	5.00	5.40	108	ng/mL	+/- 30.00%
	PFHPA	5.00	4.75	95.0	ng/mL	+/- 30.00%
	PFOA	5.00	5.40	108	ng/mL	+/- 30.00%
	PFNA	5.00	5.00	100	ng/mL	+/- 30.00%
	PFDA	5.00	4.83	96.5	ng/mL	+/- 30.00%
	PFUnA	5.00	5.03	101	ng/mL	+/- 30.00%
	PFDOA	5.00	5.23	105	ng/mL	+/- 30.00%
	PFTRDA	5.00	5.61	112	ng/mL	+/- 30.00%
	PFTEDA	5.00	5.27	105	ng/mL	+/- 30.00%
	PFBS	4.42	5.02	114	ng/mL	+/- 30.00%
	PFPEs	4.70	5.69	121	ng/mL	+/- 30.00%
	PFHXS	4.58	5.03	110	ng/mL	+/- 30.00%
	PFHPS	4.78	5.09	107	ng/mL	+/- 30.00%
	PFOS	4.65	5.12	110	ng/mL	+/- 30.00%
	PFNS	4.80	6.07	126	ng/mL	+/- 30.00%
	PFDS	4.82	5.89	122	ng/mL	+/- 30.00%
	PFDOS	4.85	6.03	124	ng/mL	+/- 30.00%
	4:2FTS	18.8	22.0	117	ng/mL	+/- 30.00%
	6:2FTS	19.0	24.0	126	ng/mL	+/- 30.00%
	8:2FTS	19.2	20.5	107	ng/mL	+/- 30.00%
	PFOSA	5.00	5.74	115	ng/mL	+/- 30.00%
	NMeFOSA	20.0	22.0	110	ng/mL	+/- 30.00%
	NEtFOSA	20.0	23.1	116	ng/mL	+/- 30.00%
	NMeFOSAA	5.00	5.17	103	ng/mL	+/- 30.00%
	NEtFOSAA	5.00	5.28	106	ng/mL	+/- 30.00%
	NMeFOSE	20.0	21.9	110	ng/mL	+/- 30.00%
	NEtFOSE	20.0	22.5	112	ng/mL	+/- 30.00%
	HFPO-DA	10.0	10.3	103	ng/mL	+/- 30.00%

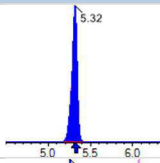
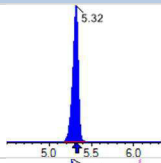
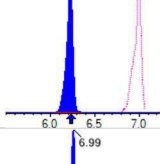
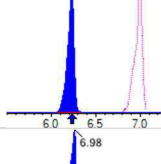
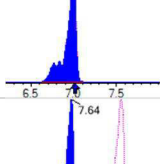
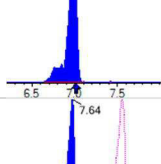
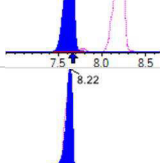
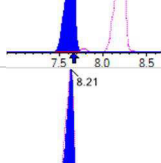
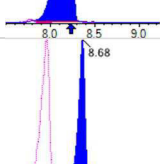
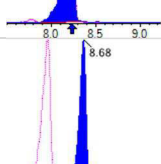
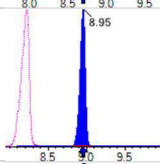
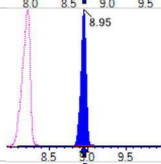
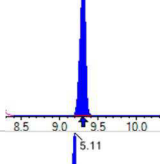
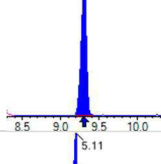
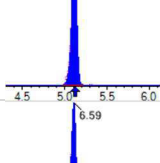
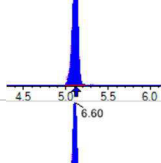
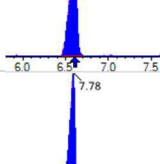
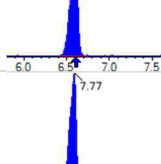
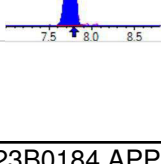
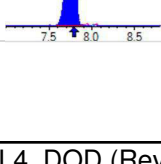


INITIAL AND CONTINUING CALIBRATION CHECK

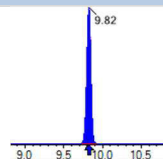
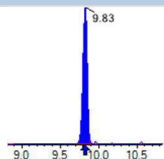
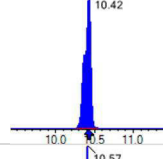
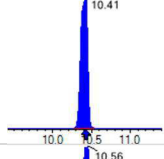
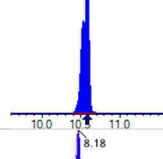
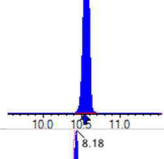
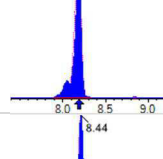
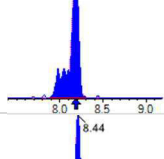
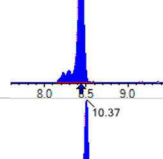
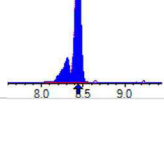
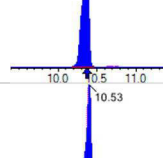
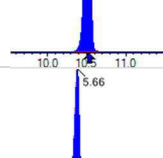
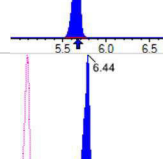
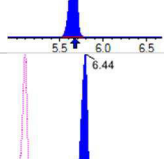
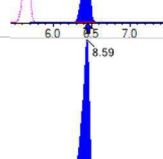
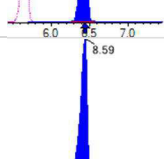
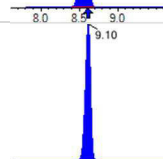
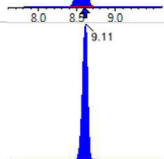
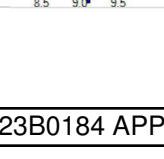
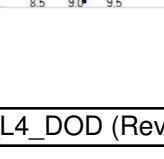
EPA 1633

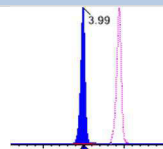
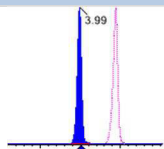
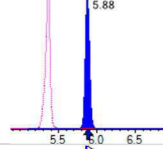
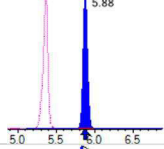
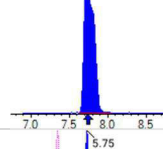
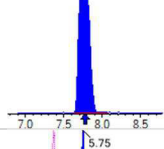
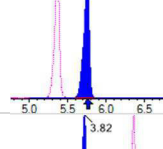
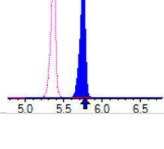
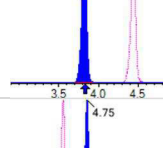
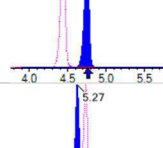
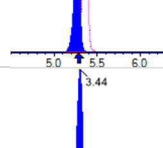
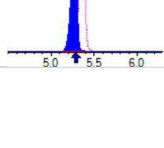
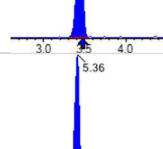
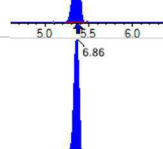
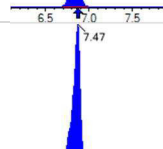
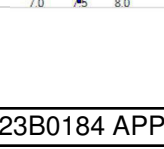
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Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Instrument ID:	Saphira	Calibration:	2309009
Standard ID:	23B0084	Sequence:	SC00861

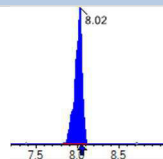
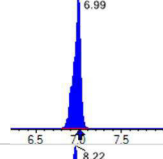
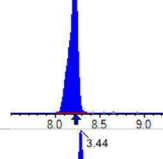
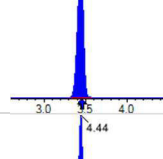
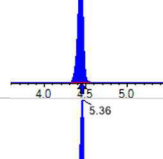
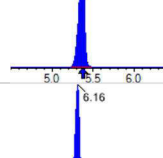
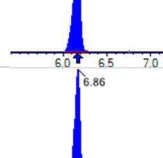
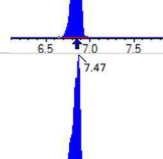
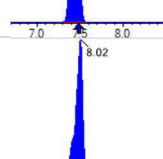
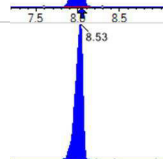
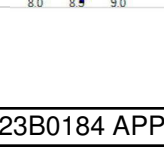
Lab Sample ID	Analyte	True	Found	%R	Units	Control Limit
SC00861-CCV5	ADONA	9.45	10.2	108	ng/mL	+/- 30.00%
	PFEESA	8.90	10.4	117	ng/mL	+/- 30.00%
	PFMPA	10.0	9.49	94.9	ng/mL	+/- 30.00%
	PFMBA	10.0	11.6	116	ng/mL	+/- 30.00%
	NFDHA	10.0	11.1	111	ng/mL	+/- 30.00%
	9CL-PF3ONS	9.35	12.1	130	ng/mL	+/- 30.00%
	11CL-PF3OUDS	9.45	11.9	126	ng/mL	+/- 30.00%
	3:3FTCA	20.0	18.6	93.1	ng/mL	+/- 30.00%
	5:3FTCA	20.0	19.7	98.4	ng/mL	+/- 30.00%
	7:3FTCA	20.0	19.8	99.2	ng/mL	+/- 30.00%

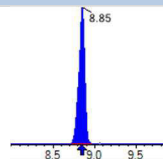
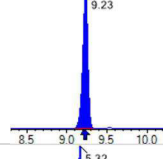
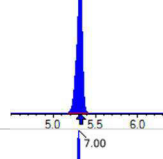
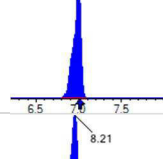
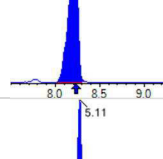
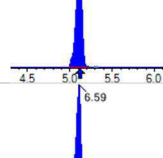
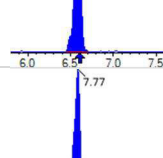
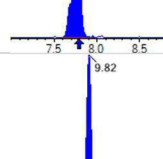
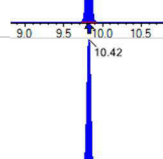
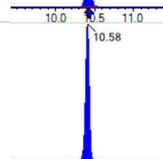
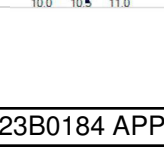
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) 4935596	(3.44, 1.00) (0.00, N/A, 0.0)	156.1	N/A 0.0 0.0	22.0556 [20.0000]	110.3%			
PFPeA	(263.0 / 219.0) 4491002 (263.0 / 69.0) 57281	(4.44, 1.00) (0.00, N/A, 0.1)	3226.2 335.4	0.0128 107.0 108.4	10.6747 [10.0000]	106.7%			
PFHxA	(313.0 / 269.0) 2919085 (313.0 / 119.0) 333877	(5.36, 1.00) (0.00, N/A, 0.1)	6700.5 1426.8	0.1144 122.2 110.7	5.4037 [5.0000]	108.1%			
PFHpA	(363.0 / 319.0) 2282872 (363.0 / 169.0) 710735	(6.16, 1.00) (0.00, N/A, 0.2)	6890.3 185888974.4	0.3113 98.6 95.5	4.7480 [5.0000]	95.0%			
PFOA	(413.0 / 369.0) 2967458 (413.0 / 169.0) 1011823	(6.86, 1.00) (0.00, N/A, -0.2)	5431.7 1820.4	0.3410 100.8 95.9	5.4021 [5.0000]	108.0%			
PFNA	(463.0 / 419.0) 2789770 (463.0 / 169.0) 611059	(7.47, 1.00) (0.00, N/A, 0.0)	25826.5 485381.2	0.2190 93.6 94.5	5.0029 [5.0000]	100.1%			
PFDA	(513.0 / 469.0) 3294848 (513.0 / 169.0) 358545	(8.03, 1.00) (0.00, N/A, 0.2)	1192.1 646.6	0.1088 87.8 80.4	4.8266 [5.0000]	96.5%			
PFUnA	(563.0 / 519.0) 2904653 (563.0 / 169.0) 396640	(8.53, 1.00) (0.00, N/A, 0.4)	1222.7 664.6	0.1366 126.9 93.5	5.0253 [5.0000]	100.5%			
PFDaA	(613.0 / 569.0) 2661420 (613.0 / 169.0) 419941	(8.85, 1.00) (0.00, N/A, -0.1)	1708.6 1255.8	0.1578 99.3 83.6	5.2304 [5.0000]	104.6%			
PFTTrDA	(663.0 / 619.0) 2744926 (663.0 / 169.0) 726201	(9.06, 1.02) (N/A, 0.00, 0.1)	1765.7 686.0	0.2646 97.4 93.0	5.6061 [5.0000]	112.1%			
PFTeDA	(713.0 / 669.0) 2805596 (713.0 / 169.0) 544631	(9.23, 1.00) (0.00, N/A, -0.1)	1432.6 725.1	0.1941 88.8 88.1	5.2728 [5.0000]	105.5%			

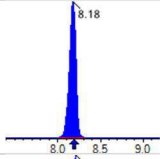
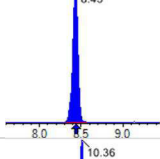
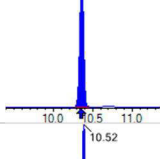
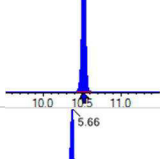
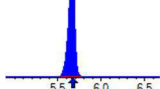
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) 5605382 (299.0 / 99.0) 3459254	(5.32 , 1.00) (0.00 , N/A , 0.0)	2213.5 1940.4	0.6171 98.1 98.9	5.0205 [4.4237]	113.5%			
PFPeS	(349.0 / 80.0) 9074016 (349.0 / 99.0) 3246315	(6.22 , 0.89) (N/A , -0.01 , -0.2)	24588.4 4168.3	0.3578 106.9 106.8	5.6898 [4.6919]	121.3%			
PFHxS	(399.0 / 80.0) 7042747 (399.0 / 99.0) 2426202	(6.99 , 1.00) (0.00 , N/A , 0.5)	5690.8 1464.8	0.3445 107.6 99.7	5.0332 [4.5549]	110.5%			
PFHpS	(449.0 / 80.0) 8158066 (449.0 / 99.0) 2301163	(7.64 , 0.93) (N/A , -0.01 , 0.0)	22096680.9 3944.1	0.2821 103.9 100.8	5.0918 [4.7570]	107.0%			
PFOS	(499.0 / 80.0) 10056378 (499.0 / 99.0) 2281211	(8.22 , 1.00) (0.01 , N/A , 0.1)	362.8 527.2	0.2268 103.1 98.4	5.1212 [4.6375]	110.4%			
PFNS	(549.0 / 80.0) 11928749 (549.0 / 99.0) 2750862	(8.68 , 1.06) (N/A , 0.00 , 0.0)	7454.0 674789.1	0.2306 95.8 100.0	6.0706 [4.7994]	126.5%			
PFDS	(599.0 / 80.0) 12852190 (599.0 / 99.0) 2703730	(8.95 , 1.09) (N/A , 0.00 , 0.1)	1858.7 1662.1	0.2104 103.9 98.5	5.8853 [4.8155]	122.2%			
PFDoS	(699.0 / 80.0) 6080519 (699.0 / 99.0) 1206736	(9.30 , 1.13) (N/A , 0.00 , 0.1)	2610.6 1108.5	0.1985 101.6 95.1	6.0315 [4.8478]	124.4%			
4:2FTS	(327.0 / 307.0) 6238851 (327.0 / 81.0) 4067350	(5.11 , 1.00) (0.00 , N/A , -0.1)	2230.8 1874.0	0.6519 94.1 102.8	21.9814 [18.6906]	117.6%			
6:2FTS	(427.0 / 407.0) 3388329 (427.0 / 81.0) 2786006	(6.59 , 1.00) (0.00 , N/A , -0.1)	1774.6 1292.7	0.8222 97.0 107.3	23.9723 [18.9808]	126.3%			
8:2FTS	(527.0 / 507.0) 3404295 (527.0 / 81.0) 2680230	(7.78 , 1.00) (0.00 , N/A , 0.3)	1786.5 1069.4	0.7873 96.5 98.7	20.4558 [19.1658]	106.7%			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) 14647011 (498.0 / 478.0) 297287	(9.82 , 1.00) (0.00 , N/A , -0.1)	2629.6 897.7	0.0203 103.3 99.5	5.7373 [5.0000]	114.7%			
NMeFOSA	(512.0 / 219.0) 14055323 (512.0 / 169.0) 11957694	(10.42 , 1.00) (0.00 , N/A , 1.0)	3606.0 5886.9	0.8508 103.0 101.7	21.9914 [20.0000]	110.0%			
NEIFOSA	(526.0 / 219.0) 15152977 (526.0 / 169.0) 19765378	(10.57 , 1.00) (-0.01 , N/A , 0.5)	9781.8 11687.2	1.3044 103.0 101.8	23.1325 [20.0000]	115.7%			
NMeFOSAA	(570.0 / 419.0) 1299423 (570.0 / 483.0) 608566	(8.18 , 1.00) (0.01 , N/A , 0.0)	1981.3 291.4	0.4683 95.5 105.7	5.1700 [5.0000]	103.4%			
NEIFOSAA	(584.0 / 419.0) 1077351 (584.0 / 526.0) 594377	(8.44 , 1.00) (0.01 , N/A , 0.1)	2213.0 911.0	0.5517 97.6 107.7	5.2835 [5.0000]	105.7%			
NMeFOSE	(616.0 / 59.0) 6076464	(10.37 , 1.00) (0.01 , N/A , 0.0)	2419.1	N/A 0.0 0.0	21.9078 [20.0000]	109.5%			
NEtFOSE	(630.0 / 59.0) 7153951	(10.53 , 1.00) (0.01 , N/A , 0.0)	1053.1	N/A 0.0 0.0	22.4868 [20.0000]	112.4%			
HFPO-DA	(285.0 / 169.0) 3150476 (285.0 / 185.0) 9418556	(5.66 , 1.00) (0.00 , N/A , 0.1)	2191.8 2763.5	2.9896 108.7 98.8	10.2559 [10.0000]	102.6%			
ADONA	(377.0 / 85.0) 11597064 (377.0 / 251.0) 1122149	(6.44 , 1.14) (N/A , 0.00 , 0.3)	2015.3 2058.5	0.0968 110.5 111.3	10.1721 [9.4270]	107.9%			
9CI-Pf3ONS	(531.0 / 351.0) 32455404 (533.0 / 353.0) 9852655	(8.59 , 1.52) (N/A , 0.00 , 0.0)	1714.0 1615.6	0.3036 103.1 97.5	12.1382 [9.3325]	130.1%			QC,
11CI-PF3OUDS	(631.0 / 451.0) 15647809 (633.0 / 453.0) 4653958	(9.10 , 1.61) (N/A , 0.00 , -0.1)	1981.7 1355.8	0.2974 96.4 96.1	11.9033 [9.4321]	126.2%			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) 231348 (241.0 / 117.0) 390405	(3.99, 0.90) (N/A, -0.01, 0.1)	1104.8 1242.7	1.6875 97.3 102.6	18.6203 [20.0000]	93.1%			
5:3FTCA	(341.0 / 236.7) 1520130 (341.0 / 217.0) 2645060	(5.88, 1.10) (N/A, 0.00, 0.0)	1456.9 2126.1	1.7400 103.5 95.8	19.6851 [20.0000]	98.4%			
7:3FTCA	(441.0 / 317.0) 2753025 (441.0 / 337.0) 2375590	(7.75, 1.45) (N/A, -0.02, -0.3)	688.2 505.3	0.8629 100.5 102.8	19.8301 [20.0000]	99.2%			
PFEESA	(315.0 / 135.0) 7010959 (315.0 / 83.0) 1897015	(5.75, 1.07) (N/A, 0.00, 0.2)	2210.8 2044.6	0.2706 96.1 103.3	10.4440 [8.9246]	117.0%			
PFMPA	(229.0 / 85.0) 1020786	(3.82, 0.86) (N/A, -0.01, 0.0)	2937.6	N/A 0.0 0.0	9.4897 [10.0000]	94.9%			
PFMBA	(279.0 / 85.0) 4290759	(4.75, 1.07) (N/A, -0.01, 0.0)	2302.2	N/A 0.0 0.0	11.6433 [10.0000]	116.4%			
NFDHA	(295.0 / 201.0) 3337791 (295.0 / 85.0) 3507307	(5.27, 0.98) (N/A, -0.01, 0.0)	2165.6 2000.0	1.0508 100.7 101.3	11.1089 [10.0000]	111.1%			
13C3_PFBa_IIS	(216.0 / 172.0) 228127	(3.44, N/A) (N/A, -0.02, N/A)	1069.8	N/A	0.6729 [1.0000]	67.3% { 108.9% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 454508	(5.36, N/A) (N/A, -0.01, N/A)	1283.8	N/A	0.7642 [1.0000]	76.4% { 122.0% }			
13C4_PFOA_IIS	(417.0 / 372.0) 588230	(6.86, N/A) (N/A, -0.01, N/A)	1305.5	N/A	0.7231 [1.0000]	72.3% { 113.5% }			
13C5_PFNAl_IIS	(468.0 / 423.0) 523274	(7.47, N/A) (N/A, -0.01, N/A)	932.4	N/A	0.6853 [1.0000]	68.5% { 95.0% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 540343	(8.02 , N/A) (N/A , -0.01 , N/A)	1221.3	N/A	0.7527 [1.0000]	75.3% { 112.9% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 828295	(6.99 , N/A) (N/A , -0.01 , N/A)	1967.0	N/A	0.6726 [1.0000]	67.3% { 93.6% }			
13C4_PFOS_IIS	(503.0 / 79.9) 1356701	(8.22 , N/A) (N/A , -0.01 , N/A)	761.2	N/A	0.7644 [1.0000]	76.4% { 105.2% }			
13C4_PFBA_EIS	(217.0 / 172.0) 2116930	(3.44 , N/A) (N/A , -0.02 , N/A)	2748.3	N/A	7.8039 [8.0000]	97.5% { 110.9% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 1909869	(4.44 , N/A) (N/A , -0.01 , N/A)	1972.9	N/A	3.7712 [4.0000]	94.3% { 109.0% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 1157700	(5.36 , N/A) (N/A , -0.01 , N/A)	1745.1	N/A	1.8746 [2.0000]	93.7% { 109.0% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 1076847	(6.16 , N/A) (N/A , 0.00 , N/A)	2063.9	N/A	1.8697 [2.0000]	93.5% { 119.4% }			
13C8_PFOA_EIS	(421.0 / 376.0) 1140200	(6.86 , N/A) (N/A , 0.00 , N/A)	1074.4	N/A	1.8110 [2.0000]	90.5% { 107.4% }			
13C9_PFNA_EIS	(472.0 / 427.0) 590342	(7.47 , N/A) (N/A , -0.02 , N/A)	956.2	N/A	1.1321 [1.0000]	113.2% { 117.8% }			
13C6_PFDA_EIS	(519.0 / 474.0) 688520	(8.02 , N/A) (N/A , -0.01 , N/A)	923.1	N/A	1.1009 [1.0000]	110.1% { 110.1% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 657297	(8.53 , N/A) (N/A , -0.01 , N/A)	602.1	N/A	1.0609 [1.0000]	106.1% { 111.6% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT Ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 583226	(8.85 , N/A) (N/A , 0.00 , N/A)	1482.3	N/A	1.0149 [1.0000]	101.5% { 130.2% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 578670	(9.23 , N/A) (N/A , 0.00 , N/A)	814.6	N/A	0.9687 [1.0000]	96.9% { 111.8% }			
13C3_PFBs_EIS	(302.0 / 80.0) 3927226	(5.32 , N/A) (N/A , -0.01 , N/A)	2041.9	N/A	2.3463 [2.0000]	117.3% { 110.0% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 1776595	(7.00 , N/A) (N/A , 0.00 , N/A)	997.9	N/A	2.0316 [2.0000]	101.6% { 101.4% }			
13C8_PFOS_EIS	(507.0 / 80.0) 3558688	(8.21 , N/A) (N/A , -0.01 , N/A)	564.3	N/A	1.9966 [2.0000]	99.8% { 108.7% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 402229	(5.11 , N/A) (N/A , -0.01 , N/A)	1577.3	N/A	3.9234 [4.0000]	98.1% { 99.8% }			
13C2_6:2FTS_EIS	(429.0 / 81.0) 433378	(6.59 , N/A) (N/A , -0.01 , N/A)	756.4	N/A	3.7494 [4.0000]	93.7% { 101.6% }			
13C2_8:2FTS_EIS	(529.0 / 81.0) 592984	(7.77 , N/A) (N/A , -0.01 , N/A)	525.1	N/A	4.8513 [4.0000]	121.3% { 113.4% }			
13C8_PFOsa_EIS	(506.0 / 78.0) 6570419	(9.82 , N/A) (N/A , 0.00 , N/A)	3069.3	N/A	2.0012 [2.0000]	100.1% { 113.5% }			
D3_NMeFOsa_EIS	(515.0 / 169.0) 1488360	(10.42 , N/A) (N/A , 0.01 , N/A)	1964.6	N/A	2.1111 [2.0000]	105.6% { 112.0% }			
D5_NEtFOsa_EIS	(531.0 / 169.0) 1348582	(10.58 , N/A) (N/A , 0.01 , N/A)	3019.3	N/A	2.1846 [2.0000]	109.2% { 111.3% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT Ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1176311	(8.18 , N/A) (N/A , -0.01 , N/A)	852.5	N/A	4.7025 [4.0000]	117.6% { 105.9% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 935646	(8.43 , N/A) (N/A , -0.01 , N/A)	20814.8	N/A	4.3924 [4.0000]	109.8% { 111.1% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 5294188	(10.36 , N/A) (N/A , 0.01 , N/A)	1336.1	N/A	22.0573 [20.0000]	110.3% { 123.8% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 6296944	(10.52 , N/A) (N/A , 0.01 , N/A)	1573.1	N/A	22.9352 [20.0000]	114.7% { 117.9% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 2997586	(5.66 , N/A) (N/A , 0.00 , N/A)	1536.8	N/A	7.7248 [8.0000]	96.6% { 107.8% }			

ANALYSIS SEQUENCE BLANKS

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 Client: AECOM
 Sequence: SC00840
 Calibration: 2309009

SDG:
 Project: Red Hill AFFF Assessment Sam
 Instrument: Saphira

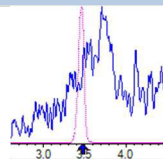
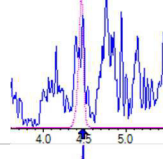
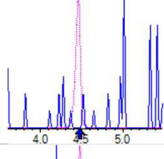
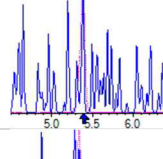
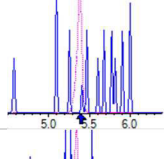
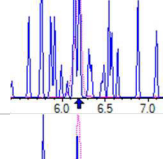
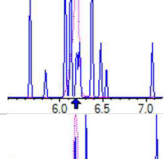
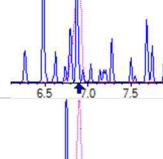
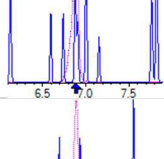
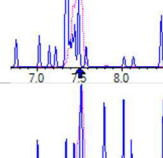
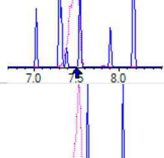
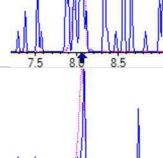
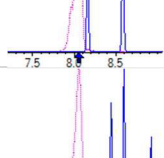
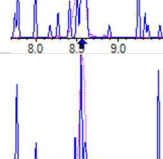
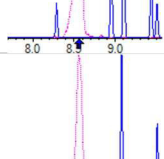
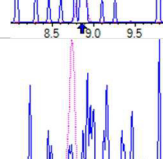
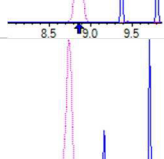
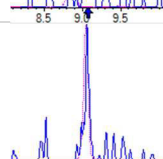
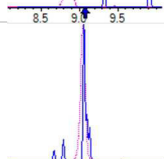
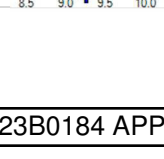
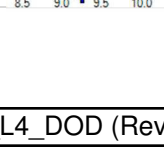
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	PFHXA	0.00	ng/mL	0.10	U
	PFHPA	0.00	ng/mL	0.10	U
	PFOA	0.00	ng/mL	0.10	U
	PFNA	0.00	ng/mL	0.10	U
	PFDA	0.00	ng/mL	0.10	U
	PFUnA	0.00	ng/mL	0.10	U
	PFDOA	0.00	ng/mL	0.10	U
	PFTRDA	0.00	ng/mL	0.10	U
	PFTEDA	0.00	ng/mL	0.10	U
	PFBS	0.00	ng/mL	0.10	U
	PFPEs	0.00	ng/mL	0.10	U
	PFHXS	0.00	ng/mL	0.10	U
	PFHPS	0.00	ng/mL	0.10	U
	PFOS	0.00	ng/mL	0.10	U
	PFNS	0.00	ng/mL	0.10	U
	PFDS	0.00	ng/mL	0.10	U
	PFDOS	0.00	ng/mL	0.10	U
	4:2FTS	0.00	ng/mL	0.40	U
	6:2FTS	0.00	ng/mL	0.40	U
	8:2FTS	0.00	ng/mL	0.40	U
	PFOSA	0.00	ng/mL	0.10	U
	NMeFOSA	0.00	ng/mL	0.40	U
	NEtFOSA	0.00	ng/mL	0.40	U
	NMeFOSAA	0.00	ng/mL	0.10	U
	NEtFOSAA	0.00	ng/mL	0.10	U
	NMeFOSE	0.101	ng/mL	0.40	U
	NEtFOSE	0.111	ng/mL	0.40	U
	HFPO-DA	0.00	ng/mL	0.20	U
	ADONA	0.00	ng/mL	0.20	U
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	PFMPA	0.00	ng/mL	0.20	U

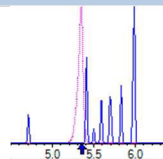
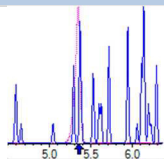
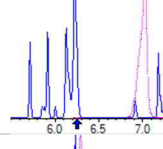
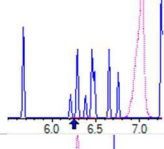
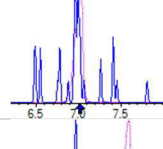
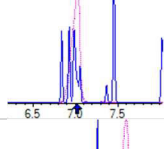
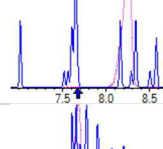
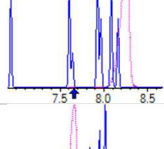
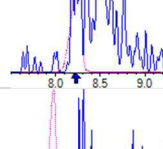
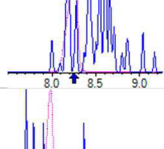
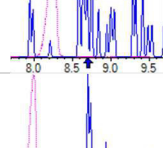
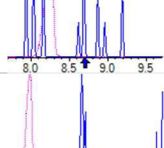
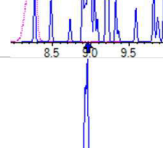
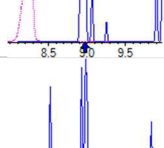
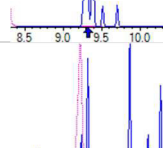
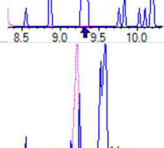
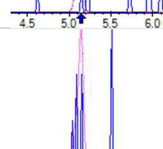
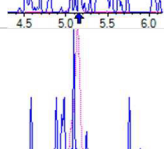
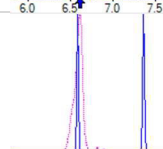
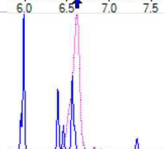
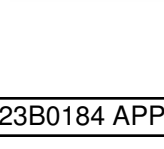
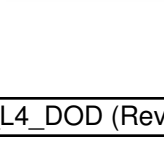
ANALYSIS SEQUENCE BLANKS

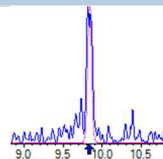
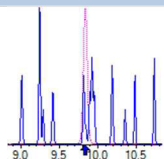
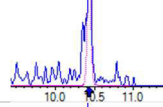
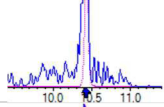
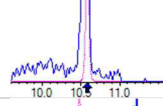
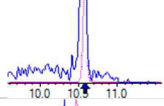
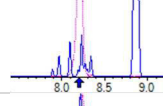
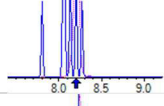
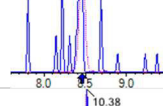
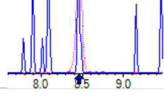
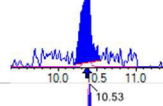
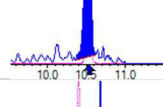
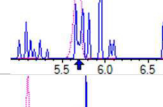
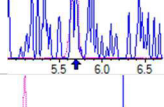
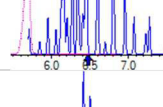
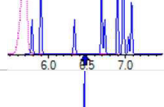
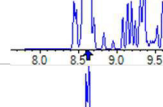
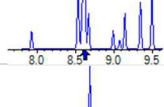
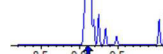
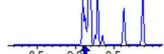
Laboratory: APPL, LLC
 Client: AECOM
 Sequence: SC00840
 Calibration: 2309009

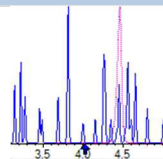
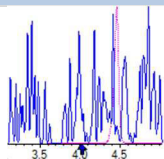
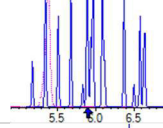
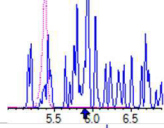
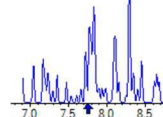
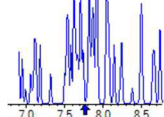
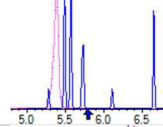
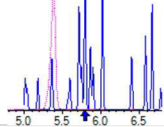
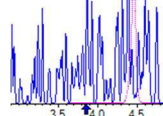
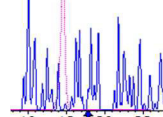
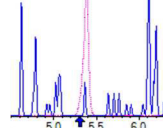
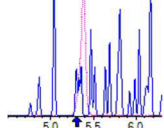
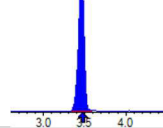
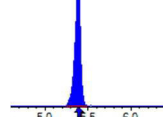
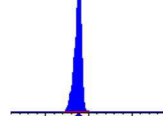

SDG:
 Project: Red Hill AFFF Assessment Sam
 Instrument: Saphira

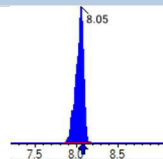
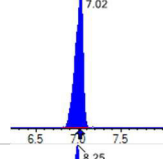
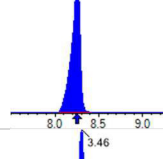
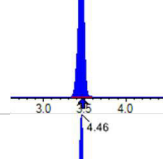
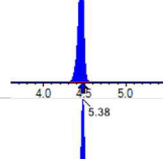
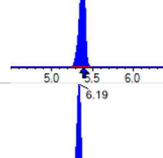
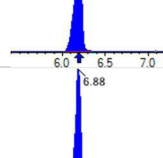
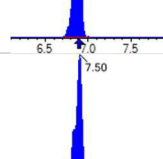
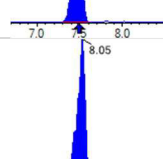
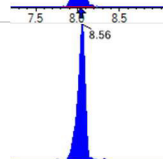
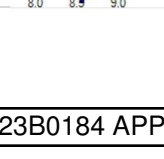
Lab Sample ID	Analyte	Found	Units	RL	C
SC00840-ICB1	PFMBA	0.00	ng/mL	0.20	U
	NFDHA	0.00	ng/mL	0.20	U
	9CL-PF3ONS	0.00	ng/mL	0.20	U
	11CL-PF3OUDS	0.00	ng/mL	0.20	U
	3:3FTCA	0.00	ng/mL	0.40	U
	5:3FTCA	0.00	ng/mL	0.40	U
	7:3FTCA	0.00	ng/mL	0.40	U
	13C4-PFBA	6.83	ng/mL		
	13C5-PFPEA	3.93	ng/mL		
	13C5-PFHXA	1.88	ng/mL		
	13C4-PFHPA	1.95	ng/mL		
	13C8-PFOA	1.61	ng/mL		
	13C9-PFNA	0.909	ng/mL		
	13C6-PFDA	0.869	ng/mL		
	13C7-PFUnA	0.862	ng/mL		
	13C2-PFDOA	1.04	ng/mL		
	13C2-PFTEDA	0.906	ng/mL		
	13C3-PFBS	1.82	ng/mL		
	13C3-PFHXS	1.70	ng/mL		
	13C8-PFOS	1.93	ng/mL		
	13C2-4:2FTS	3.28	ng/mL		
	13C2-6:2FTS	3.18	ng/mL		
	13C2-8:2FTS	3.42	ng/mL		
	13C8-PFOSA	1.86	ng/mL		
	D3-NMEFOSA	1.87	ng/mL		
	D5-NETFOSA	2.05	ng/mL		
	D3-NMEFOSAA	4.07	ng/mL		
	D5-NETFOSAA	4.08	ng/mL		
	D7-NMEFOSE	19.9	ng/mL		
	D9-NETFOSSE	20.0	ng/mL		
	13C3-HFPO-DA	7.79	ng/mL		

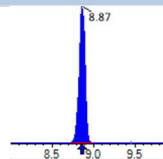
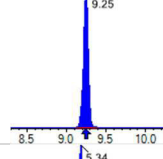
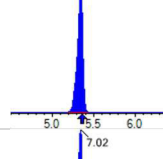
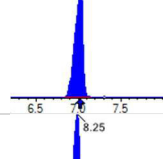
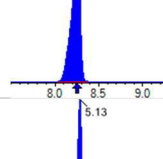
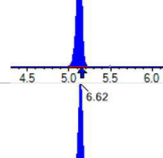
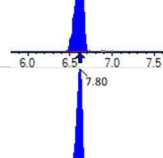
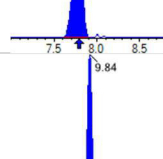
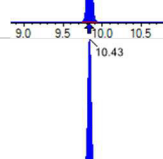
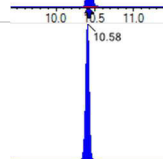
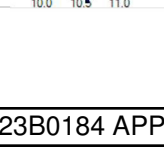
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
PFPeA	(263.0 / 219.0) N/A (263.0 / 69.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHxA	(313.0 / 269.0) N/A (313.0 / 119.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHpA	(363.0 / 319.0) N/A (363.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFOA	(413.0 / 369.0) N/A (413.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFNA	(463.0 / 419.0) N/A (463.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDA	(513.0 / 469.0) N/A (513.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFUnA	(563.0 / 519.0) N/A (563.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDaA	(613.0 / 569.0) N/A (613.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFTTrDA	(663.0 / 619.0) N/A (663.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFTeDA	(713.0 / 669.0) N/A (713.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

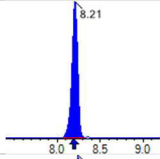
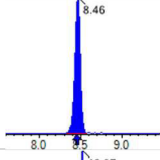
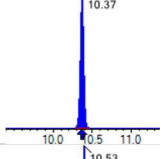
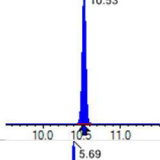
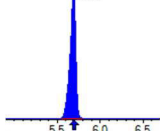
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) N/A (299.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFPeS	(349.0 / 80.0) N/A (349.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHxS	(399.0 / 80.0) N/A (399.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHpS	(449.0 / 80.0) N/A (449.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFOS	(499.0 / 80.0) N/A (499.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFNS	(549.0 / 80.0) N/A (549.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDS	(599.0 / 80.0) N/A (599.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDoS	(699.0 / 80.0) N/A (699.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
4:2FTS	(327.0 / 307.0) N/A (327.0 / 81.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
6:2FTS	(427.0 / 407.0) N/A (427.0 / 81.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
8:2FTS	(527.0 / 507.0) N/A (527.0 / 81.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) N/A (498.0 / 478.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NMeFOSA	(512.0 / 219.0) N/A (512.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NEiFOSA	(526.0 / 219.0) N/A (526.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NMeFOSAA	(570.0 / 419.0) N/A (570.0 / 483.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NEiFOSAA	(584.0 / 419.0) N/A (584.0 / 526.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NMeFOSE	(616.0 / 59.0) 30935	(10.38 , 1.00) (0.01 , N/A , 0.0)	57.0	N/A 0.0 0.0	0.1012	N/A			
NEiFOSE	(630.0 / 59.0) 37534	(10.53 , 1.00) (0.00 , N/A , 0.0)	73.6	N/A 0.0 0.0	0.1107	N/A			
HFPO-DA	(285.0 / 169.0) N/A (285.0 / 185.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
ADONA	(377.0 / 85.0) N/A (377.0 / 251.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
9CI-Pf3ONS	(531.0 / 351.0) N/A (533.0 / 353.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
11CI-PF3OUDS	(631.0 / 451.0) N/A (633.0 / 453.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) N/A (241.0 / 117.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
5:3FTCA	(341.0 / 236.7) N/A (341.0 / 217.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
7:3FTCA	(441.0 / 317.0) N/A (441.0 / 337.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFEESA	(315.0 / 135.0) N/A (315.0 / 83.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFMPA	(229.0 / 85.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
PFMBA	(279.0 / 85.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
NFDHA	(295.0 / 201.0) N/A (295.0 / 85.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
13C3_PFBA_IIS	(216.0 / 172.0) 308902	(3.46 , N/A) (N/A , -0.01 , N/A)	1070.8	N/A	0.9112 [1.0000]	91.1% { 93.7% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 492240	(5.38 , N/A) (N/A , -0.01 , N/A)	5902.7	N/A	0.8276 [1.0000]	82.8% { 88.6% }			
13C4_PFOA_IIS	(417.0 / 372.0) 779414	(6.89 , N/A) (N/A , 0.00 , N/A)	1766.6	N/A	0.9582 [1.0000]	95.8% { 103.9% }			
13C5_PFNA_IIS	(468.0 / 423.0) 701819	(7.50 , N/A) (N/A , 0.01 , N/A)	659.7	N/A	0.9191 [1.0000]	91.9% { 92.2% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 662877	(8.05 , N/A) (N/A , 0.00 , N/A)	1116.5	N/A	0.9234 [1.0000]	92.3% { 96.3% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 1153284	(7.02 , N/A) (N/A , 0.00 , N/A)	1358.4	N/A	0.9364 [1.0000]	93.6% { 96.0% }			
13C4_PFOS_IIS	(503.0 / 79.9) 1655832	(8.25 , N/A) (N/A , 0.01 , N/A)	894.9	N/A	0.9329 [1.0000]	93.3% { 95.6% }			
13C4_PFBA_EIS	(217.0 / 172.0) 2507442	(3.46 , N/A) (N/A , 0.00 , N/A)	2830.1	N/A	6.8265 [8.0000]	85.3% { 85.6% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 2156125	(4.46 , N/A) (N/A , -0.01 , N/A)	1996.4	N/A	3.9311 [4.0000]	98.3% { 88.4% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 1260382	(5.38 , N/A) (N/A , -0.01 , N/A)	1789.2	N/A	1.8845 [2.0000]	94.2% { 82.1% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 1216516	(6.19 , N/A) (N/A , 0.01 , N/A)	1409.3	N/A	1.9503 [2.0000]	97.5% { 86.1% }			
13C8_PFOA_EIS	(421.0 / 376.0) 1343118	(6.88 , N/A) (N/A , 0.00 , N/A)	1415.4	N/A	1.6100 [2.0000]	80.5% { 80.6% }			
13C9_PFNA_EIS	(472.0 / 427.0) 635603	(7.50 , N/A) (N/A , 0.01 , N/A)	879.4	N/A	0.9088 [1.0000]	90.9% { 89.6% }			
13C6_PFDA_EIS	(519.0 / 474.0) 667077	(8.05 , N/A) (N/A , 0.02 , N/A)	877.9	N/A	0.8695 [1.0000]	86.9% { 90.8% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 655229	(8.56 , N/A) (N/A , 0.01 , N/A)	630.8	N/A	0.8620 [1.0000]	86.2% { 87.1% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 736171	(8.87, N/A) (N/A, 0.01, N/A)	1616.9	N/A	1.0443 [1.0000]	104.4% { 99.9% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 663664	(9.25, N/A) (N/A, 0.01, N/A)	941.8	N/A	0.9056 [1.0000]	90.6% { 92.3% }			
13C3_PFBs_EIS	(302.0 / 80.0) 4248266	(5.34, N/A) (N/A, -0.01, N/A)	3325.1	N/A	1.8229 [2.0000]	91.1% { 93.4% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 2066130	(7.02, N/A) (N/A, 0.00, N/A)	1133.7	N/A	1.6969 [2.0000]	84.8% { 87.3% }			
13C8_PFOS_EIS	(507.0 / 80.0) 4203475	(8.25, N/A) (N/A, 0.01, N/A)	1574.8	N/A	1.9323 [2.0000]	96.6% { 98.4% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 468716	(5.13, N/A) (N/A, -0.01, N/A)	1403.3	N/A	3.2836 [4.0000]	82.1% { 90.4% }			
13C2_6:2FTS_EIS	(429.0 / 81.0) 511696	(6.62, N/A) (N/A, 0.01, N/A)	812.2	N/A	3.1794 [4.0000]	79.5% { 86.6% }			
13C2_8:2FTS_EIS	(529.0 / 81.0) 582261	(7.80, N/A) (N/A, 0.01, N/A)	687.3	N/A	3.4212 [4.0000]	85.5% { 87.7% }			
13C8_PFOsa_EIS	(506.0 / 78.0) 7457193	(9.84, N/A) (N/A, 0.01, N/A)	2452.0	N/A	1.8610 [2.0000]	93.0% { 95.7% }			
D3_NMeFOsa_EIS	(515.0 / 169.0) 1606033	(10.43, N/A) (N/A, 0.01, N/A)	2085.7	N/A	1.8665 [2.0000]	93.3% { 93.4% }			
D5_NEtFOsa_EIS	(531.0 / 169.0) 1547178	(10.58, N/A) (N/A, 0.01, N/A)	2981.0	N/A	2.0536 [2.0000]	102.7% { 98.9% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1241849	(8.21 , N/A) (N/A , 0.01 , N/A)	1058.0	N/A	4.0676 [4.0000]	101.7% { 96.5% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 1059492	(8.46 , N/A) (N/A , 0.01 , N/A)	4007.6	N/A	4.0753 [4.0000]	101.9% { 109.5% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 5837509	(10.37 , N/A) (N/A , 0.01 , N/A)	2444.0	N/A	19.9273 [20.0000]	99.6% { 96.5% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 6713177	(10.53 , N/A) (N/A , 0.01 , N/A)	1443.3	N/A	20.0340 [20.0000]	100.2% { 98.0% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 3273875	(5.69 , N/A) (N/A , 0.00 , N/A)	2225.9	N/A	7.7900 [8.0000]	97.4% { 87.6% }			

ANALYSIS SEQUENCE BLANKS

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 Client: AECOM
 Sequence: SC00861
 Calibration: 2309009

SDG:
 Project: Red Hill AFFF Assessment Sam
 Instrument: Saphira

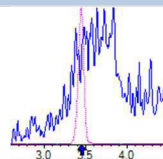
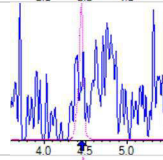
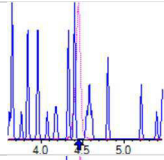
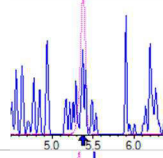
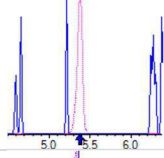
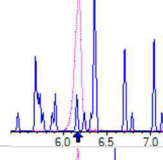
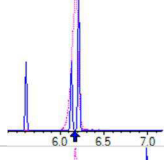
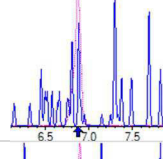
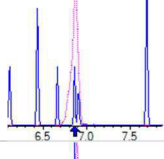
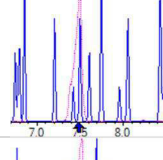
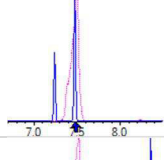
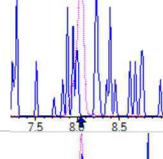
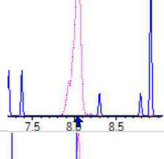
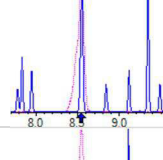
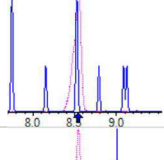
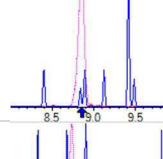
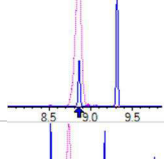
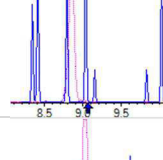
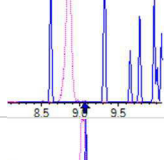
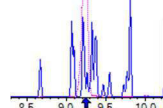
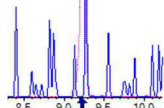
Lab Sample ID	Analyte	Found	Units	RL	C
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	PFHXA	0.00	ng/mL	0.10	U
	PFHPA	0.00	ng/mL	0.10	U
	PFOA	0.00	ng/mL	0.10	U
	PFNA	0.00	ng/mL	0.10	U
	PFDA	0.00	ng/mL	0.10	U
	PFUnA	0.00	ng/mL	0.10	U
	PFDOA	0.00	ng/mL	0.10	U
	PFTRDA	0.00	ng/mL	0.10	U
	PFTEDA	0.00	ng/mL	0.10	U
	PFBS	0.00	ng/mL	0.10	U
	PFPEs	0.00	ng/mL	0.10	U
	PFHXS	0.00	ng/mL	0.10	U
	PFHPS	0.00	ng/mL	0.10	U
	PFOS	0.00	ng/mL	0.10	U
	PFNS	0.00	ng/mL	0.10	U
	PFDS	0.00	ng/mL	0.10	U
	PFDOS	0.00	ng/mL	0.10	U
	4:2FTS	0.00	ng/mL	0.40	U
	6:2FTS	0.00	ng/mL	0.40	U
	8:2FTS	0.00	ng/mL	0.40	U
	PFOSA	0.00	ng/mL	0.10	U
	NMeFOSA	0.00	ng/mL	0.40	U
	NEtFOSA	0.00	ng/mL	0.40	U
	NMeFOSAA	0.00	ng/mL	0.10	U
	NEtFOSAA	0.00	ng/mL	0.10	U
	NMeFOSE	0.00	ng/mL	0.40	U
	NEtFOSE	0.00	ng/mL	0.40	U
	HFPO-DA	0.00	ng/mL	0.20	U
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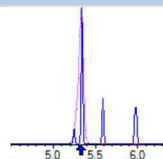
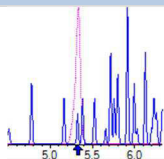
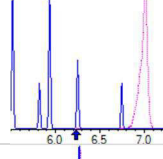
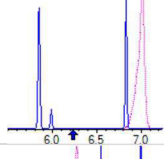
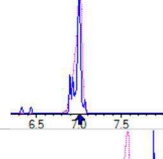
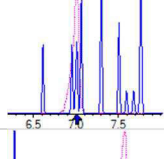
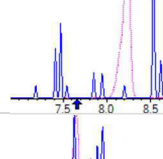
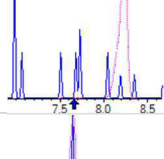
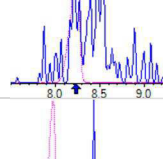
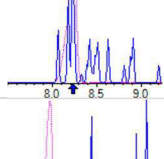
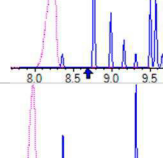
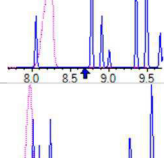
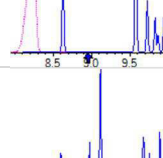
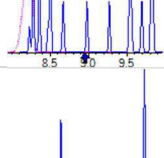
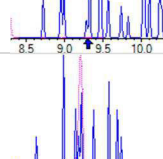
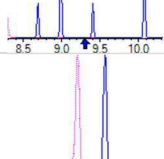
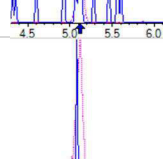
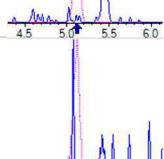
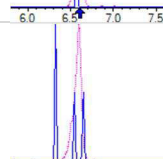
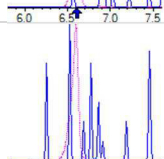
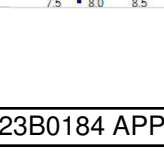
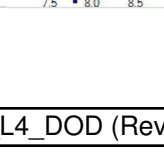
ANALYSIS SEQUENCE BLANKS

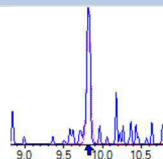
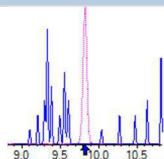
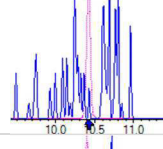
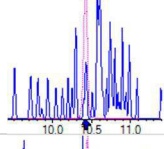
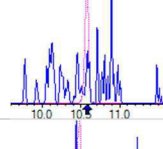
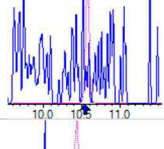
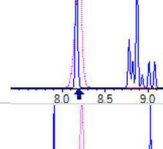
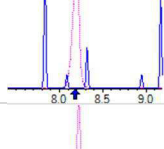
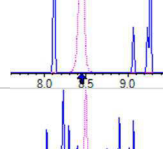
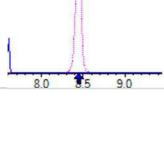
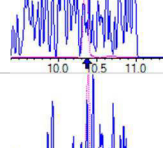
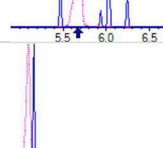
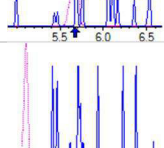
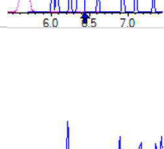
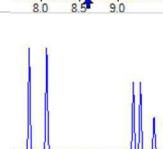
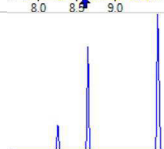
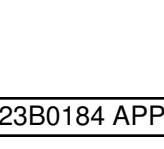
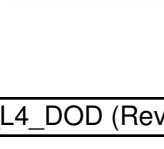
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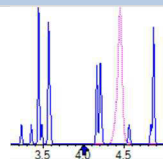
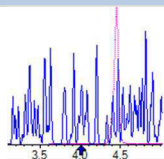
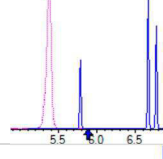
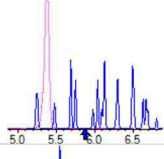
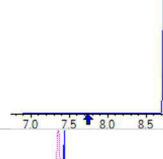
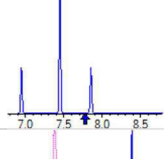
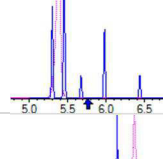
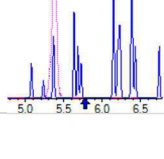
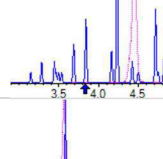
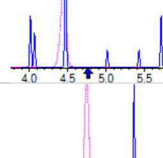
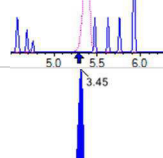
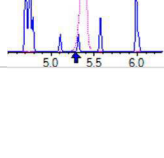
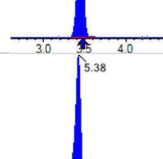
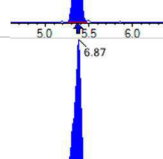
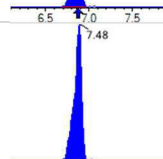
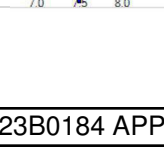
SDG:
 Project: Red Hill AFFF Assessment Sam
 Instrument: Saphira

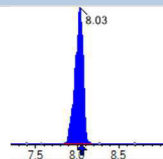
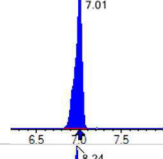
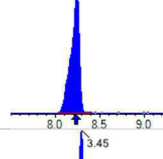
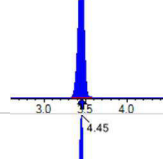
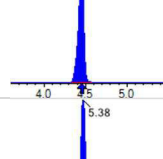
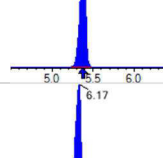
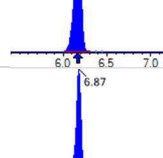
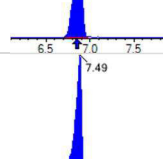
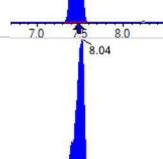
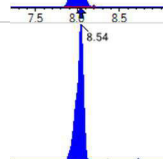
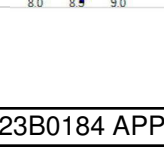
Lab Sample ID	Analyte	Found	Units	RL	C
SC00861-CCB1	PFMBA	0.00	ng/mL	0.20	U
	NFDHA	0.00	ng/mL	0.20	U
	9CL-PF3ONS	0.00	ng/mL	0.20	U
	11CL-PF3OUDS	0.00	ng/mL	0.20	U
	3:3FTCA	0.00	ng/mL	0.40	U
	5:3FTCA	0.00	ng/mL	0.40	U
	7:3FTCA	0.00	ng/mL	0.40	U
	13C4-PFBA	7.06	ng/mL		
	13C5-PFPEA	3.75	ng/mL		
	13C5-PFHXA	1.88	ng/mL		
	13C4-PFHPA	1.87	ng/mL		
	13C8-PFOA	1.70	ng/mL		
	13C9-PFNA	0.847	ng/mL		
	13C6-PFDA	0.960	ng/mL		
	13C7-PFUnA	0.904	ng/mL		
	13C2-PFDOA	0.910	ng/mL		
	13C2-PFTEDA	0.858	ng/mL		
	13C3-PFBS	2.21	ng/mL		
	13C3-PFHXS	1.89	ng/mL		
	13C8-PFOS	1.77	ng/mL		
	13C2-4:2FTS	4.35	ng/mL		
	13C2-6:2FTS	3.51	ng/mL		
	13C2-8:2FTS	4.38	ng/mL		
	13C8-PFOSA	1.64	ng/mL		
	D3-NMEFOSA	1.61	ng/mL		
	D5-NETFOSA	1.68	ng/mL		
	D3-NMEFOSAA	3.89	ng/mL		
	D5-NETFOSAA	3.96	ng/mL		
	D7-NMEFOSE	15.8	ng/mL		
	D9-NETFOSSE	17.6	ng/mL		
	13C3-HFPO-DA	7.71	ng/mL		

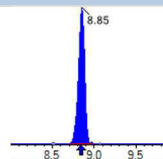
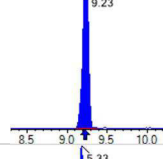
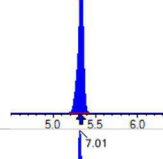
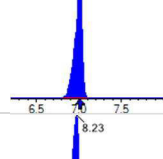
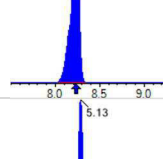
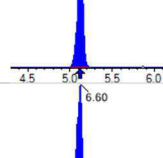
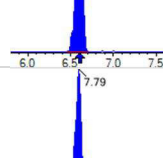
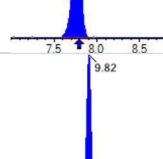
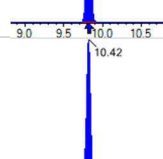
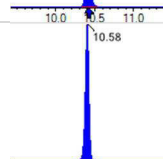
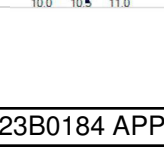
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-Imin, ΔRT-CVmin, ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration True ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
PFPeA	(263.0 / 219.0) N/A (263.0 / 69.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHxA	(313.0 / 269.0) N/A (313.0 / 119.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHpA	(363.0 / 319.0) N/A (363.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFOA	(413.0 / 369.0) N/A (413.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFNA	(463.0 / 419.0) N/A (463.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDA	(513.0 / 469.0) N/A (513.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFUnA	(563.0 / 519.0) N/A (563.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDaA	(613.0 / 569.0) N/A (613.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFTTrDA	(663.0 / 619.0) N/A (663.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFTeDA	(713.0 / 669.0) N/A (713.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

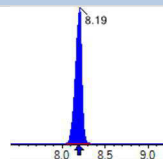
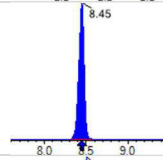
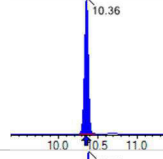
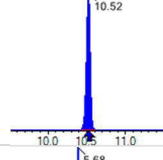
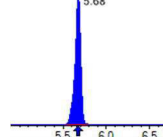
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-Imin, ΔRT-CVmin, ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration True ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) N/A (299.0 / 99.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFPeS	(349.0 / 80.0) N/A (349.0 / 99.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHxS	(399.0 / 80.0) N/A (399.0 / 99.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHpS	(449.0 / 80.0) N/A (449.0 / 99.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFOS	(499.0 / 80.0) N/A (499.0 / 99.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFNS	(549.0 / 80.0) N/A (549.0 / 99.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDS	(599.0 / 80.0) N/A (599.0 / 99.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDoS	(699.0 / 80.0) N/A (699.0 / 99.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
4:2FTS	(327.0 / 307.0) N/A (327.0 / 81.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
6:2FTS	(427.0 / 407.0) N/A (427.0 / 81.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
8:2FTS	(527.0 / 507.0) N/A (527.0 / 81.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-Imin, ΔRT-CVmin, ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration True ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) N/A (498.0 / 478.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NMeFOSA	(512.0 / 219.0) N/A (512.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NEtFOSA	(526.0 / 219.0) N/A (526.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NMeFOSAA	(570.0 / 419.0) N/A (570.0 / 483.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NEtFOSAA	(584.0 / 419.0) N/A (584.0 / 526.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NMeFOSE	(616.0 / 59.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
NEtFOSE	(630.0 / 59.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
HFPO-DA	(285.0 / 169.0) N/A (285.0 / 185.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
ADONA	(377.0 / 85.0) N/A (377.0 / 251.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
9CI-Pf3ONS	(531.0 / 351.0) N/A (533.0 / 353.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
11CI-PF3OUDS	(631.0 / 451.0) N/A (633.0 / 453.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-Imin, ΔRT-CVmin, ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration True ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) N/A (241.0 / 117.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
5:3FTCA	(341.0 / 236.7) N/A (341.0 / 217.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
7:3FTCA	(441.0 / 317.0) N/A (441.0 / 337.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFEESA	(315.0 / 135.0) N/A (315.0 / 83.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFMPA	(229.0 / 85.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
PFMBA	(279.0 / 85.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
NFDHA	(295.0 / 201.0) N/A (295.0 / 85.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
13C3_PFBA_IIS	(216.0 / 172.0) 259382	(3.45, N/A) (N/A, -0.01, N/A)	972.1	N/A	0.7651 [1.0000]	76.5% { 123.8% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 462225	(5.38, N/A) (N/A, 0.01, N/A)	1362.4	N/A	0.7772 [1.0000]	77.7% { 124.0% }			
13C4_PFOA_IIS	(417.0 / 372.0) 643344	(6.87, N/A) (N/A, 0.01, N/A)	1038.6	N/A	0.7909 [1.0000]	79.1% { 124.1% }			
13C5_PFNA_IIS	(468.0 / 423.0) 652768	(7.48, N/A) (N/A, 0.01, N/A)	1078.9	N/A	0.8548 [1.0000]	85.5% { 118.5% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-Imin, ΔRT-CVmin, ΔRT ion(s))	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration True ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 613138	(8.03, N/A) (N/A, 0.00, N/A)	686.9	N/A	0.8541 [1.0000]	85.4% {128.1%}			
18O2_PFHxS_IIS	(403.0 / 83.9) 913123	(7.01, N/A) (N/A, 0.01, N/A)	1035.8	N/A	0.7414 [1.0000]	74.1% {103.2%}			
13C4_PFOS_IIS	(503.0 / 79.9) 1583184	(8.24, N/A) (N/A, 0.01, N/A)	664.9	N/A	0.8920 [1.0000]	89.2% {122.8%}			
13C4_PFBA_EIS	(217.0 / 172.0) 2178436	(3.45, N/A) (N/A, -0.01, N/A)	3048.6	N/A	7.0630 [8.0000]	88.3% {114.1%}			
13C5_PFPeA_EIS	(268.0 / 223.0) 1930465	(4.45, N/A) (N/A, 0.00, N/A)	2461.0	N/A	3.7482 [4.0000]	93.7% {110.1%}			
13C5_PFHxA_EIS	(318.0 / 273.0) 1179641	(5.38, N/A) (N/A, 0.01, N/A)	1951.8	N/A	1.8783 [2.0000]	93.9% {111.1%}			
13C4_PFHpA_EIS	(367.0 / 322.0) 1098257	(6.17, N/A) (N/A, 0.01, N/A)	1120.3	N/A	1.8750 [2.0000]	93.7% {121.8%}			
13C8_PFOA_EIS	(421.0 / 376.0) 1173007	(6.87, N/A) (N/A, 0.01, N/A)	1272.7	N/A	1.7035 [2.0000]	85.2% {110.5%}			
13C9_PFNA_EIS	(472.0 / 427.0) 550952	(7.49, N/A) (N/A, 0.00, N/A)	1349.8	N/A	0.8470 [1.0000]	84.7% {110.0%}			
13C6_PFDA_EIS	(519.0 / 474.0) 681534	(8.04, N/A) (N/A, 0.01, N/A)	3806.1	N/A	0.9604 [1.0000]	96.0% {109.0%}			
13C7_PFUnA_EIS	(570.0 / 525.0) 635750	(8.54, N/A) (N/A, 0.01, N/A)	869.8	N/A	0.9043 [1.0000]	90.4% {108.0%}			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-Imin, ΔRT-CVmin, ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration True ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 593160	(8.85, N/A) (N/A, 0.00, N/A)	992.6	N/A	0.9097 [1.0000]	91.0% {132.4%}			
13C2_PFTeDA_EIS	(715.0 / 670.0) 581770	(9.23, N/A) (N/A, 0.00, N/A)	583.5	N/A	0.8583 [1.0000]	85.8% {112.4%}			
13C3_PFBs_EIS	(302.0 / 80.0) 4080851	(5.33, N/A) (N/A, 0.01, N/A)	2221.5	N/A	2.2116 [2.0000]	110.6% {114.3%}			
13C3_PFHxS_EIS	(402.0 / 80.0) 1818502	(7.01, N/A) (N/A, 0.01, N/A)	1750.0	N/A	1.8863 [2.0000]	94.3% {103.8%}			
13C8_PFOS_EIS	(507.0 / 80.0) 3678813	(8.23, N/A) (N/A, 0.00, N/A)	1198.6	N/A	1.7687 [2.0000]	88.4% {112.4%}			
13C2_4:2FTS_EIS	(329.0 / 81.0) 491947	(5.13, N/A) (N/A, 0.01, N/A)	1049.8	N/A	4.3528 [4.0000]	108.8% {122.0%}			
13C2_6:2FTS_EIS	(429.0 / 81.0) 447376	(6.60, N/A) (N/A, 0.00, N/A)	1404.0	N/A	3.5109 [4.0000]	87.8% {104.8%}			
13C2_8:2FTS_EIS	(529.0 / 81.0) 590569	(7.79, N/A) (N/A, 0.01, N/A)	966.7	N/A	4.3827 [4.0000]	109.6% {113.0%}			
13C8_PFOA_EIS	(506.0 / 78.0) 6300558	(9.82, N/A) (N/A, 0.00, N/A)	3290.9	N/A	1.6445 [2.0000]	82.2% {108.8%}			
D3_NMeFOA_EIS	(515.0 / 169.0) 1328065	(10.42, N/A) (N/A, 0.00, N/A)	2179.8	N/A	1.6143 [2.0000]	80.7% {100.0%}			
D5_NEtFOA_EIS	(531.0 / 169.0) 1212487	(10.58, N/A) (N/A, 0.00, N/A)	2193.9	N/A	1.6832 [2.0000]	84.2% {100.1%}			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-Imin, ΔRT-CVmin, ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration True ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1135653	(8.19 , N/A) (N/A , 0.01 , N/A)	1026.7	N/A	3.8905 [4.0000]	97.3% { 102.3% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 984152	(8.45 , N/A) (N/A , 0.01 , N/A)	10204.2	N/A	3.9592 [4.0000]	99.0% { 116.8% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 4411738	(10.36 , N/A) (N/A , 0.00 , N/A)	1251.5	N/A	15.7513 [20.0000]	78.8% { 103.2% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 5629685	(10.52 , N/A) (N/A , 0.00 , N/A)	1255.8	N/A	17.5715 [20.0000]	87.9% { 105.4% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 3043789	(5.68 , N/A) (N/A , 0.01 , N/A)	2420.8	N/A	7.7128 [8.0000]	96.4% { 109.4% }			

ANALYSIS SEQUENCE BLANKS

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 Client: AECOM
 Sequence: SC00861
 Calibration: 2309009

SDG:
 Project: Red Hill AFFF Assessment Sam
 Instrument: Saphira

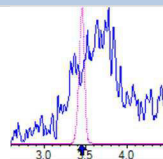
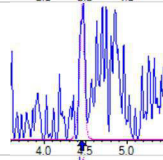
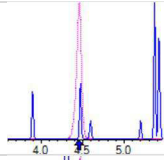
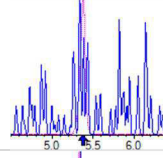
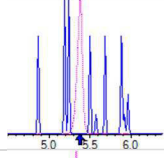
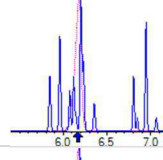
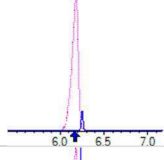
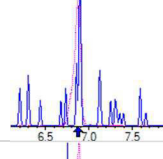
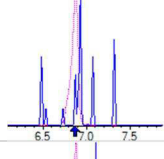
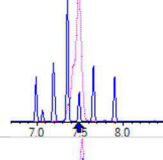
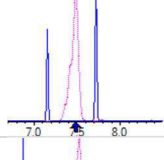
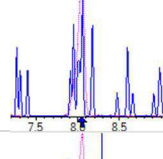
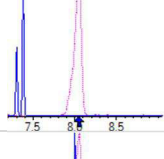
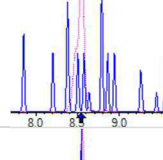
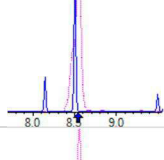
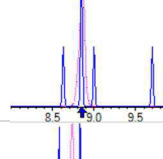
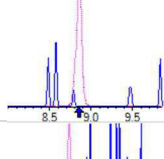
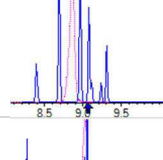
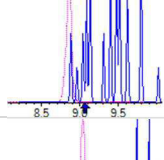
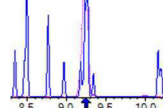
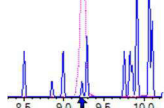
Lab Sample ID	Analyte	Found	Units	RL	C
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	PFPEA	0.00	ng/mL	0.20	U
	PFHXA	0.00	ng/mL	0.10	U
	PFHPA	0.00	ng/mL	0.10	U
	PFOA	0.00	ng/mL	0.10	U
	PFNA	0.00	ng/mL	0.10	U
	PFDA	0.00	ng/mL	0.10	U
	PFUnA	0.00	ng/mL	0.10	U
	PFDOA	0.00	ng/mL	0.10	U
	PFTRDA	0.00	ng/mL	0.10	U
	PFTEDA	0.00	ng/mL	0.10	U
	PFBS	0.00	ng/mL	0.10	U
	PFPEs	0.00	ng/mL	0.10	U
	PFHXS	0.00	ng/mL	0.10	U
	PFHPS	0.00	ng/mL	0.10	U
	PFOS	0.00	ng/mL	0.10	U
	PFNS	0.00	ng/mL	0.10	U
	PFDS	0.00	ng/mL	0.10	U
	PFDOS	0.00	ng/mL	0.10	U
	4:2FTS	0.00	ng/mL	0.40	U
	6:2FTS	0.00	ng/mL	0.40	U
	8:2FTS	0.00	ng/mL	0.40	U
	PFOSA	0.00	ng/mL	0.10	U
	NMeFOSA	0.00	ng/mL	0.40	U
	NEtFOSA	0.00	ng/mL	0.40	U
	NMeFOSAA	0.00	ng/mL	0.10	U
	NEtFOSAA	0.00	ng/mL	0.10	U
	NMeFOSE	0.00	ng/mL	0.40	U
	NEtFOSE	0.00	ng/mL	0.40	U
	HFPO-DA	0.00	ng/mL	0.20	U
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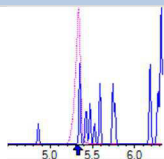
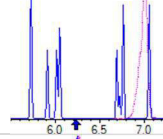
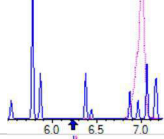
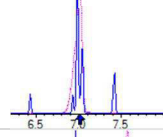
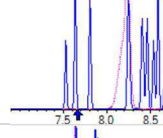
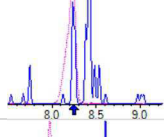
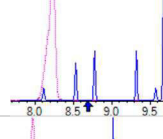
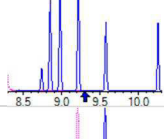
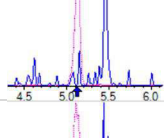
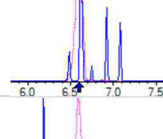
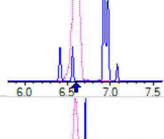
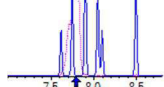
ANALYSIS SEQUENCE BLANKS

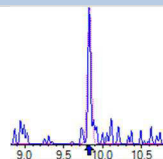
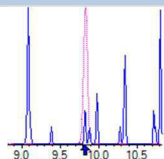
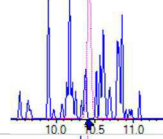
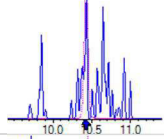
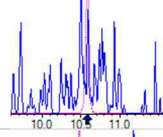
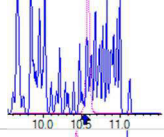
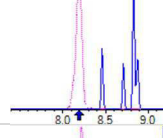
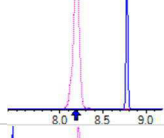
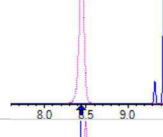
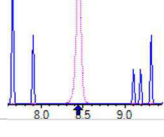
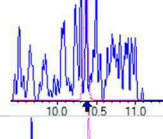
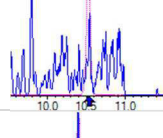
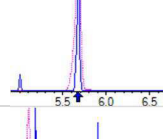
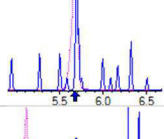
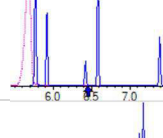
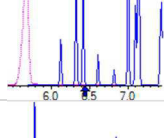
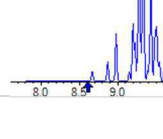
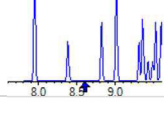
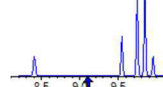
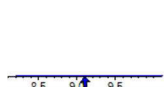
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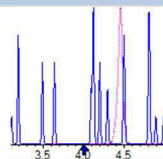
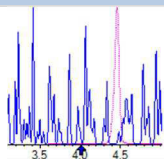
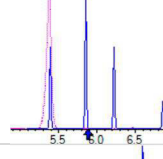
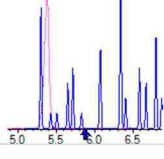
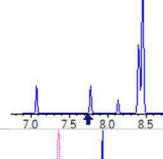
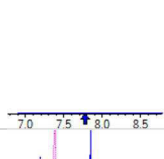
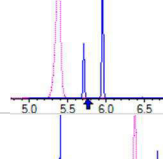
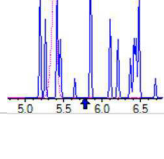
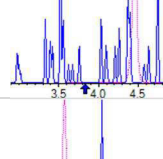
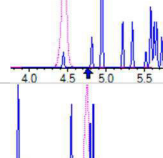
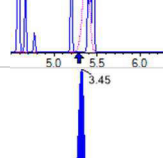
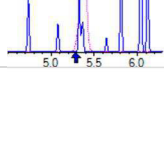
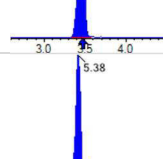
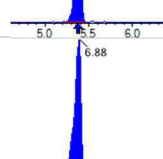
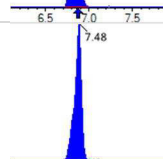
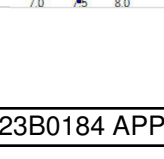
SDG:
 Project: Red Hill AFFF Assessment Sam
 Instrument: Saphira

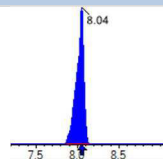
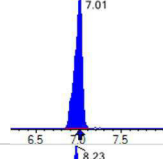
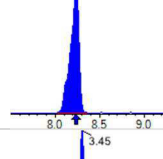
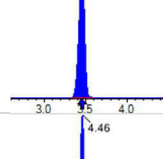
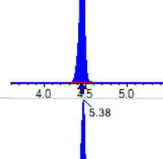
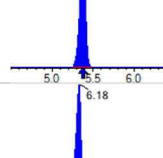
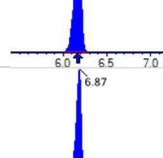
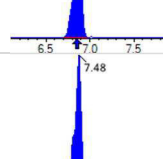
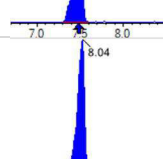
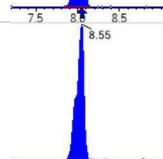
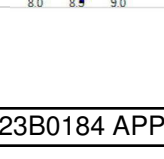
Lab Sample ID	Analyte	Found	Units	RL	C
SC00861-CCB2	PFMBA	0.00	ng/mL	0.20	U
	NFDHA	0.00	ng/mL	0.20	U
	9CL-PF3ONS	0.00	ng/mL	0.20	U
	11CL-PF3OUDS	0.00	ng/mL	0.20	U
	3:3FTCA	0.00	ng/mL	0.40	U
	5:3FTCA	0.00	ng/mL	0.40	U
	7:3FTCA	0.00	ng/mL	0.40	U
	13C4-PFBA	6.92	ng/mL		
	13C5-PFPEA	3.83	ng/mL		
	13C5-PFHXA	1.91	ng/mL		
	13C4-PFHPA	1.89	ng/mL		
	13C8-PFOA	1.71	ng/mL		
	13C9-PFNA	0.904	ng/mL		
	13C6-PFDA	0.844	ng/mL		
	13C7-PFUnA	0.852	ng/mL		
	13C2-PFDOA	0.809	ng/mL		
	13C2-PFTEDA	0.824	ng/mL		
	13C3-PFBS	2.01	ng/mL		
	13C3-PFHXS	1.78	ng/mL		
	13C8-PFOS	1.85	ng/mL		
	13C2-4:2FTS	3.61	ng/mL		
	13C2-6:2FTS	3.27	ng/mL		
	13C2-8:2FTS	3.87	ng/mL		
	13C8-PFOSA	1.81	ng/mL		
	D3-NMEFOSA	1.89	ng/mL		
	D5-NETFOSA	1.99	ng/mL		
	D3-NMEFOSAA	4.42	ng/mL		
	D5-NETFOSAA	4.17	ng/mL		
	D7-NMEFOSE	18.1	ng/mL		
	D9-NETFOSSE	20.5	ng/mL		
	13C3-HFPO-DA	7.95	ng/mL		

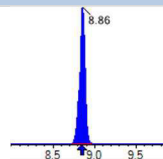
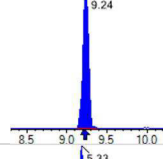
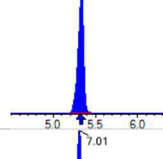
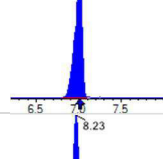
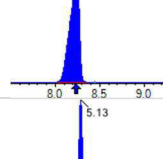
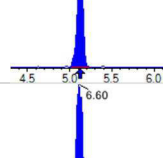
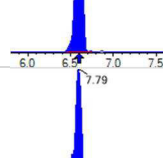
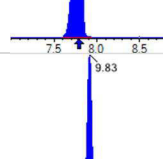
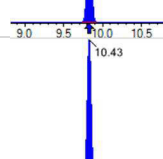
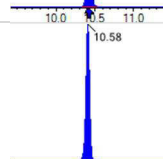
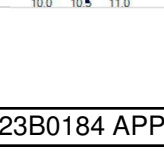
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
PFPeA	(263.0 / 219.0) N/A (263.0 / 69.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHxA	(313.0 / 269.0) N/A (313.0 / 119.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHpA	(363.0 / 319.0) N/A (363.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFOA	(413.0 / 369.0) N/A (413.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFNA	(463.0 / 419.0) N/A (463.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDA	(513.0 / 469.0) N/A (513.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFUnA	(563.0 / 519.0) N/A (563.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDaA	(613.0 / 569.0) N/A (613.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFTrDA	(663.0 / 619.0) N/A (663.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFTeDA	(713.0 / 669.0) N/A (713.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

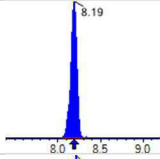
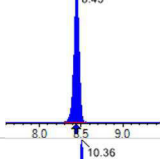
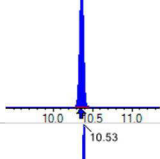
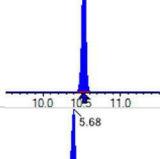
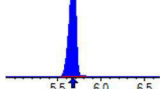
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) N/A (299.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFPeS	(349.0 / 80.0) N/A (349.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHxS	(399.0 / 80.0) N/A (399.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHpS	(449.0 / 80.0) N/A (449.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFOS	(499.0 / 80.0) N/A (499.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFNS	(549.0 / 80.0) N/A (549.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDS	(599.0 / 80.0) N/A (599.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDoS	(699.0 / 80.0) N/A (699.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
4:2FTS	(327.0 / 307.0) N/A (327.0 / 81.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
6:2FTS	(427.0 / 407.0) N/A (427.0 / 81.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
8:2FTS	(527.0 / 507.0) N/A (527.0 / 81.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) N/A (498.0 / 478.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NMeFOSA	(512.0 / 219.0) N/A (512.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NEtFOSA	(526.0 / 219.0) N/A (526.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NMeFOSAA	(570.0 / 419.0) N/A (570.0 / 483.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NEtFOSAA	(584.0 / 419.0) N/A (584.0 / 526.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NMeFOSE	(616.0 / 59.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
NEtFOSE	(630.0 / 59.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
HFPO-DA	(285.0 / 169.0) N/A (285.0 / 185.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
ADONA	(377.0 / 85.0) N/A (377.0 / 251.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
9Cl-Pf3ONS	(531.0 / 351.0) N/A (533.0 / 353.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
11Cl-PF3OUDS	(631.0 / 451.0) N/A (633.0 / 453.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) N/A (241.0 / 117.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
5:3FTCA	(341.0 / 236.7) N/A (341.0 / 217.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
7:3FTCA	(441.0 / 317.0) N/A (441.0 / 337.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFEESA	(315.0 / 135.0) N/A (315.0 / 83.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFMPA	(229.0 / 85.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
PFMBA	(279.0 / 85.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
NFDHA	(295.0 / 201.0) N/A (295.0 / 85.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
13C3_PFBA_IIS	(216.0 / 172.0) 259745	(3.45, N/A) (N/A, 0.00, N/A)	1171.7	N/A	0.7662 [1.0000]	76.6% { 124.0% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 437032	(5.38, N/A) (N/A, 0.01, N/A)	752.9	N/A	0.7348 [1.0000]	73.5% { 117.3% }			
13C4_PFOA_IIS	(417.0 / 372.0) 644563	(6.88, N/A) (N/A, 0.01, N/A)	976.5	N/A	0.7924 [1.0000]	79.2% { 124.4% }			
13C5_PFNA_IIS	(468.0 / 423.0) 564549	(7.48, N/A) (N/A, 0.01, N/A)	880.7	N/A	0.7393 [1.0000]	73.9% { 102.5% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 611748	(8.04 , N/A) (N/A , 0.01 , N/A)	2205.2	N/A	0.8522 [1.0000]	85.2% { 127.8% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 929086	(7.01 , N/A) (N/A , 0.01 , N/A)	736.9	N/A	0.7544 [1.0000]	75.4% { 105.0% }			
13C4_PFOS_IIS	(503.0 / 79.9) 1463564	(8.23 , N/A) (N/A , 0.00 , N/A)	938.9	N/A	0.8246 [1.0000]	82.5% { 113.5% }			
13C4_PFBA_EIS	(217.0 / 172.0) 2137466	(3.45 , N/A) (N/A , 0.00 , N/A)	3604.0	N/A	6.9205 [8.0000]	86.5% { 112.0% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 1863981	(4.46 , N/A) (N/A , 0.01 , N/A)	2047.9	N/A	3.8278 [4.0000]	95.7% { 106.3% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 1136918	(5.38 , N/A) (N/A , 0.01 , N/A)	1142.2	N/A	1.9146 [2.0000]	95.7% { 107.1% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 1045891	(6.18 , N/A) (N/A , 0.01 , N/A)	1580.0	N/A	1.8885 [2.0000]	94.4% { 116.0% }			
13C8_PFOA_EIS	(421.0 / 376.0) 1180044	(6.87 , N/A) (N/A , 0.01 , N/A)	1257.4	N/A	1.7105 [2.0000]	85.5% { 111.1% }			
13C9_PFNA_EIS	(472.0 / 427.0) 508331	(7.48 , N/A) (N/A , 0.00 , N/A)	749.0	N/A	0.9036 [1.0000]	90.4% { 101.5% }			
13C6_PFDA_EIS	(519.0 / 474.0) 597263	(8.04 , N/A) (N/A , 0.01 , N/A)	719.7	N/A	0.8435 [1.0000]	84.4% { 95.5% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 597806	(8.55 , N/A) (N/A , 0.01 , N/A)	478.8	N/A	0.8522 [1.0000]	85.2% { 101.5% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 526216	(8.86 , N/A) (N/A , 0.01 , N/A)	1471.3	N/A	0.8088 [1.0000]	80.9% { 117.5% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 557015	(9.24 , N/A) (N/A , 0.01 , N/A)	933.1	N/A	0.8236 [1.0000]	82.4% { 107.6% }			
13C3_PFBs_EIS	(302.0 / 80.0) 3773097	(5.33 , N/A) (N/A , 0.01 , N/A)	2261.2	N/A	2.0097 [2.0000]	100.5% { 105.7% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 1749183	(7.01 , N/A) (N/A , 0.01 , N/A)	1101.5	N/A	1.7833 [2.0000]	89.2% { 99.9% }			
13C8_PFOS_EIS	(507.0 / 80.0) 3552760	(8.23 , N/A) (N/A , 0.01 , N/A)	1437.7	N/A	1.8477 [2.0000]	92.4% { 108.5% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 415489	(5.13 , N/A) (N/A , 0.01 , N/A)	1082.5	N/A	3.6131 [4.0000]	90.3% { 103.1% }			
13C2_6:2FTS_EIS	(429.0 / 81.0) 424394	(6.60 , N/A) (N/A , 0.00 , N/A)	997.7	N/A	3.2733 [4.0000]	81.8% { 99.5% }			
13C2_8:2FTS_EIS	(529.0 / 81.0) 531238	(7.79 , N/A) (N/A , 0.01 , N/A)	1058.6	N/A	3.8747 [4.0000]	96.9% { 101.6% }			
13C8_PFOsa_EIS	(506.0 / 78.0) 6423740	(9.83 , N/A) (N/A , 0.01 , N/A)	2441.2	N/A	1.8137 [2.0000]	90.7% { 110.9% }			
D3_NMeFOsa_EIS	(515.0 / 169.0) 1436714	(10.43 , N/A) (N/A , 0.01 , N/A)	2311.7	N/A	1.8891 [2.0000]	94.5% { 108.1% }			
D5_NEtFOsa_EIS	(531.0 / 169.0) 1322935	(10.58 , N/A) (N/A , 0.01 , N/A)	2278.4	N/A	1.9866 [2.0000]	99.3% { 109.2% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT Ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1194041	(8.19 , N/A) (N/A , 0.00 , N/A)	1385.6	N/A	4.4248 [4.0000]	110.6% { 107.5% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 958344	(8.45 , N/A) (N/A , 0.01 , N/A)	78120.9	N/A	4.1705 [4.0000]	104.3% { 113.8% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 4688097	(10.36 , N/A) (N/A , 0.01 , N/A)	2301.0	N/A	18.1060 [20.0000]	90.5% { 109.6% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 6079488	(10.53 , N/A) (N/A , 0.01 , N/A)	1080.0	N/A	20.5264 [20.0000]	102.6% { 113.9% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 2968182	(5.68 , N/A) (N/A , 0.01 , N/A)	1590.9	N/A	7.9548 [8.0000]	99.4% { 106.7% }			

ANALYSIS SEQUENCE BLANKS

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 Client: AECOM
 Sequence: SC00861
 Calibration: 2309009

SDG:
 Project: Red Hill AFFF Assessment Sam
 Instrument: Saphira

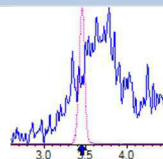
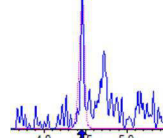
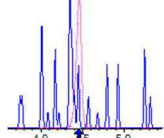
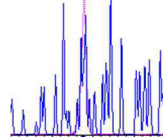
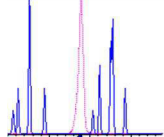
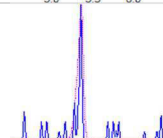
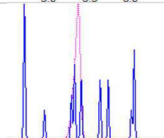
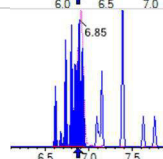
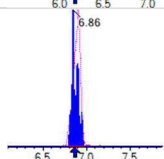
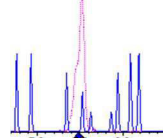
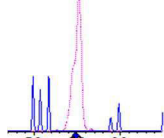
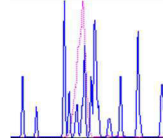
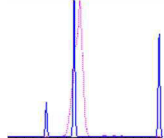
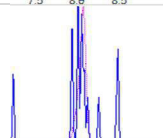
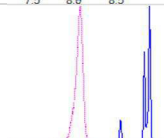
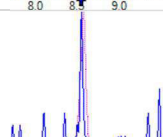
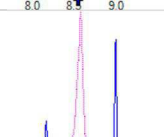
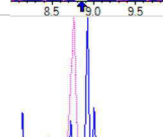
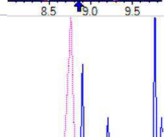
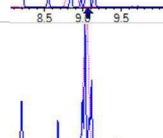
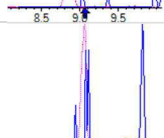
Lab Sample ID	Analyte	Found	Units	RL	C
SC00861-CCB3	PFBA	0.00	ng/mL	0.40	U
	PFPEA	0.00	ng/mL	0.20	U
	PFHXA	0.00	ng/mL	0.10	U
	PFHPA	0.00	ng/mL	0.10	U
	PFOA	0.0134	ng/mL	0.10	U
	PFNA	0.00	ng/mL	0.10	U
	PFDA	0.00	ng/mL	0.10	U
	PFUnA	0.00	ng/mL	0.10	U
	PFDOA	0.00	ng/mL	0.10	U
	PFTRDA	0.00	ng/mL	0.10	U
	PFTEDA	0.00	ng/mL	0.10	U
	PFBS	0.00	ng/mL	0.10	U
	PFPEs	0.00	ng/mL	0.10	U
	PFHXS	0.00	ng/mL	0.10	U
	PFHPS	0.00	ng/mL	0.10	U
	PFOS	0.00	ng/mL	0.10	U
	PFNS	0.000672	ng/mL	0.10	U
	PFDS	0.000841	ng/mL	0.10	U
	PFDOS	0.00109	ng/mL	0.10	U
	4:2FTS	0.00443	ng/mL	0.40	U
	6:2FTS	0.00810	ng/mL	0.40	U
	8:2FTS	0.00326	ng/mL	0.40	U
	PFOSA	0.00435	ng/mL	0.10	U
	NMeFOSA	0.00329	ng/mL	0.40	U
	NEtFOSA	0.00629	ng/mL	0.40	U
	NMeFOSAA	0.00824	ng/mL	0.10	U
	NEtFOSAA	0.00	ng/mL	0.10	U
	NMeFOSE	0.0144	ng/mL	0.40	U
	NEtFOSE	0.00614	ng/mL	0.40	U
	HFPO-DA	0.00300	ng/mL	0.20	U
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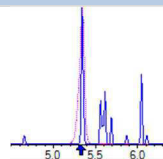
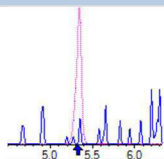
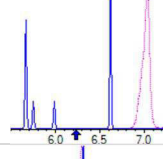
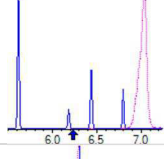
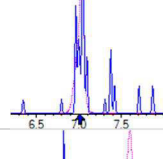
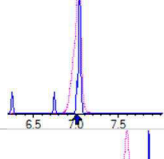
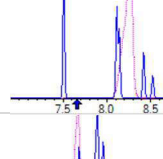
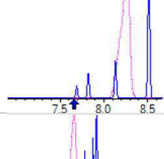
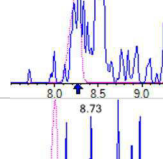
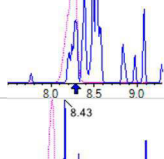
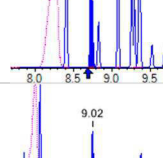
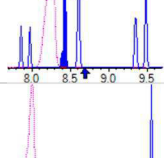
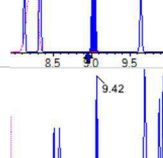
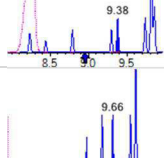
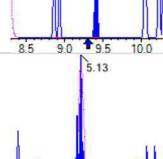
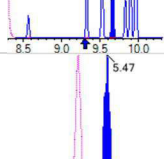
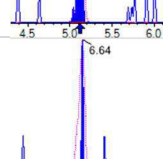
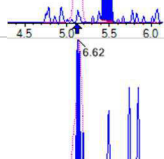
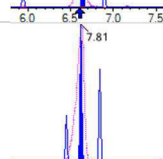
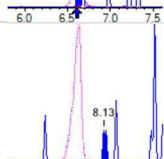
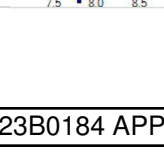
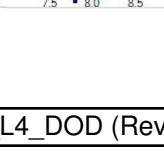
ANALYSIS SEQUENCE BLANKS

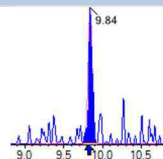
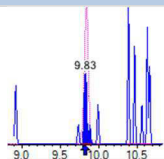
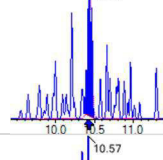
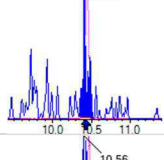
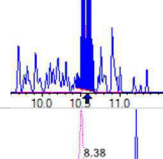
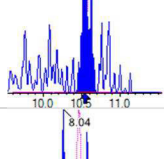
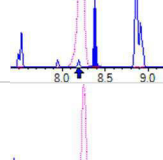
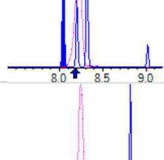
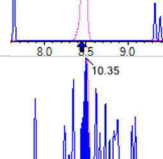
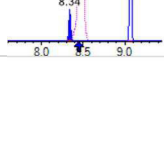
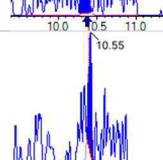
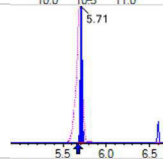
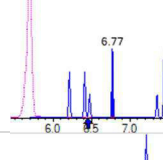
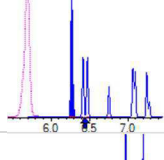
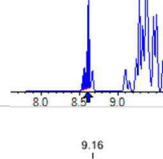
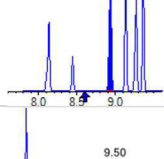
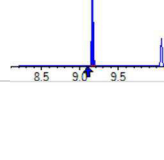
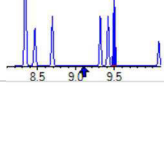
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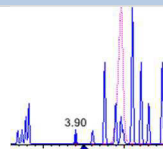
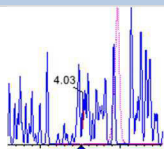
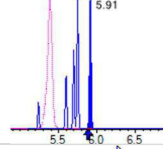
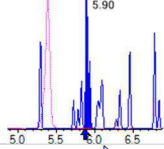
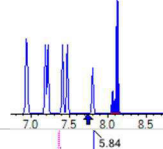
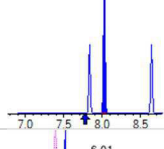
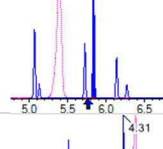
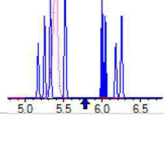
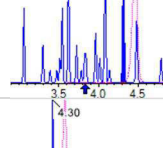
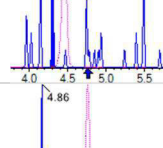
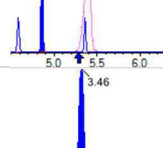
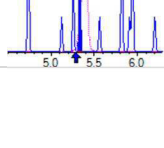
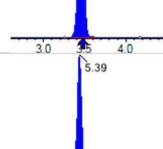
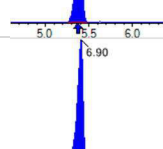
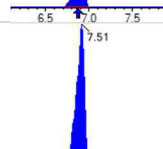
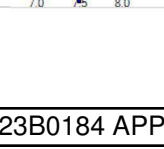
SDG:
 Project: Red Hill AFFF Assessment Sam
 Instrument: Saphira

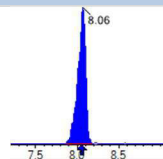
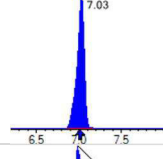
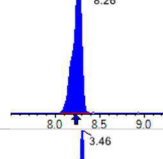
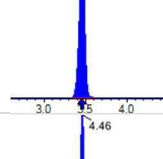
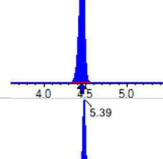
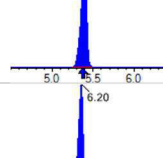
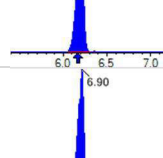
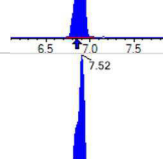
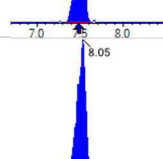
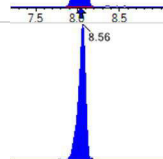
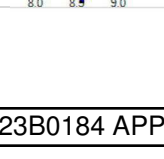
Lab Sample ID	Analyte	Found	Units	RL	C
SC00861-CCB3	PFMBA	0.00117	ng/mL	0.20	U
	NFDHA	0.00133	ng/mL	0.20	U
	9CL-PF3ONS	0.000886	ng/mL	0.20	U
	11CL-PF3OUDS	0.000409	ng/mL	0.20	U
	3:3FTCA	0.00197	ng/mL	0.40	U
	5:3FTCA	0.00991	ng/mL	0.40	U
	7:3FTCA	0.00927	ng/mL	0.40	U
	13C4-PFBA	7.01	ng/mL		
	13C5-PFPEA	3.65	ng/mL		
	13C5-PFHXA	1.83	ng/mL		
	13C4-PFHPA	1.75	ng/mL		
	13C8-PFOA	1.61	ng/mL		
	13C9-PFNA	0.980	ng/mL		
	13C6-PFDA	0.974	ng/mL		
	13C7-PFUnA	1.04	ng/mL		
	13C2-PFDOA	1.01	ng/mL		
	13C2-PFTEDA	0.992	ng/mL		
	13C3-PFBS	2.01	ng/mL		
	13C3-PFHXS	1.74	ng/mL		
	13C8-PFOS	1.77	ng/mL		
	13C2-4:2FTS	3.90	ng/mL		
	13C2-6:2FTS	3.33	ng/mL		
	13C2-8:2FTS	4.21	ng/mL		
	13C8-PFOSA	1.75	ng/mL		
	D3-NMEFOSA	1.77	ng/mL		
	D5-NETFOSA	2.07	ng/mL		
	D3-NMEFOSAA	4.30	ng/mL		
	D5-NETFOSAA	3.90	ng/mL		
	D7-NMEFOSE	18.8	ng/mL		
	D9-NETFOSSE	20.5	ng/mL		
	13C3-HFPO-DA	7.49	ng/mL		

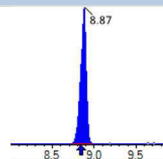
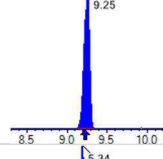
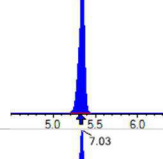
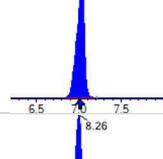
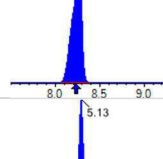
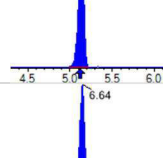
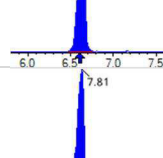
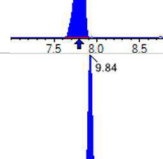
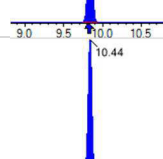
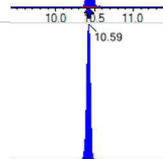
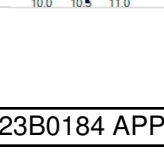
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
PFPeA	(263.0 / 219.0) N/A (263.0 / 69.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHxA	(313.0 / 269.0) N/A (313.0 / 119.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHpA	(363.0 / 319.0) N/A (363.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFOA	(413.0 / 369.0) 8073 (413.0 / 169.0) 1666	(6.85, 0.99) (-0.05, N/A, -0.7)	48745.2 116298.4	0.2063 61.0 58.1	0.0134	N/A			
PFNA	(463.0 / 419.0) N/A (463.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDA	(513.0 / 469.0) N/A (513.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFUnA	(563.0 / 519.0) N/A (563.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDaA	(613.0 / 569.0) N/A (613.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFTTrDA	(663.0 / 619.0) N/A (663.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFTeDA	(713.0 / 669.0) N/A (713.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

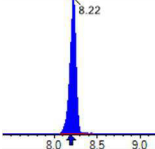
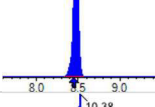
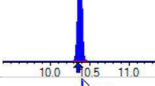
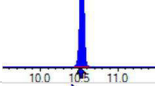
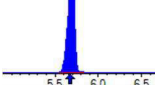
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) N/A (299.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFPeS	(349.0 / 80.0) N/A (349.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHxS	(399.0 / 80.0) N/A (399.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHpS	(449.0 / 80.0) N/A (449.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFOS	(499.0 / 80.0) N/A (499.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFNS	(549.0 / 80.0) 1377 (549.0 / 99.0) 2288	(8.73 , 1.06) (N/A , 0.05 , 18.1)	122.1 1021.2	1.6620 690.7 720.9	0.0007	N/A			IR2,
PFDS	(599.0 / 80.0) 1913 (599.0 / 99.0) 538	(9.02 , 1.09) (N/A , 0.07 , -21.5)	390.1 15.4	0.2812 138.9 131.7	0.0008	N/A			
PFDoS	(699.0 / 80.0) 1142 (699.0 / 99.0) 831	(9.42 , 1.14) (N/A , 0.11 , -14.8)	38.7 44.9	0.7275 372.4 348.6	0.0011	N/A			IR2,
4:2FTS	(327.0 / 307.0) 1521 (327.0 / 81.0) 12079	(5.13 , 1.00) (-0.01 , N/A , -20.8)	56.2 60.3	7.9411 1145.9 1252.4	0.0044	N/A			IR2,
6:2FTS	(427.0 / 407.0) 1237 (427.0 / 81.0) 1857	(6.64 , 1.00) (0.00 , N/A , 1.0)	973.9 5258.9	1.5019 177.2 196.0	0.0081	N/A			IR2,
8:2FTS	(527.0 / 507.0) 573 (527.0 / 81.0) 779	(7.81 , 1.00) (0.00 , N/A , -18.9)	381.0 31.4	1.3588 166.6 170.3	0.0033	N/A			IR2,

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) 11466 (498.0 / 478.0) 617	(9.84, 1.00) (0.00, N/A, 0.9)	30.0 17.1	0.0538 273.9 264.0	0.0043	N/A			
NMeFOSA	(512.0 / 219.0) 2078 (512.0 / 169.0) 2619	(10.44, 1.00) (0.00, N/A, 1.8)	24.0 36.6	1.2603 152.6 150.6	0.0033	N/A			IR2,
NEiFOSA	(526.0 / 219.0) 4588 (526.0 / 169.0) 6481	(10.57, 1.00) (-0.02, N/A, 1.0)	31.9 22.0	1.4127 111.6 110.3	0.0063	N/A			
NMeFOSAA	(570.0 / 419.0) 2233 (570.0 / 483.0) 1196	(8.38, 1.02) (0.16, N/A, 20.1)	19333.8 3277.6	0.5354 109.2 120.9	0.0082	N/A			RT,
NEiFOSAA	(584.0 / 419.0) N/A (584.0 / 526.0) 606	(N/A, N/A) (N/A, N/A, N/A)	N/A 796323.3	N/A 0.0 0.0	0.0000	N/A			
NMeFOSE	(616.0 / 59.0) 4020	(10.35, 1.00) (-0.02, N/A, 0.0)	15.6	N/A 0.0 0.0	0.0144	N/A			
NEiFOSE	(630.0 / 59.0) 2058	(10.55, 1.00) (0.01, N/A, 0.0)	10.3	N/A 0.0 0.0	0.0061	N/A			
HFPO-DA	(285.0 / 169.0) 972 (285.0 / 185.0) 3489	(5.71, 1.00) (0.02, N/A, 0.9)	178341.9 56.8	3.5881 130.5 118.6	0.0030	N/A			
ADONA	(377.0 / 85.0) 483 (377.0 / 251.0) 1903	(6.77, 1.19) (N/A, 0.33, 30.1)	106.3 86.7	3.9403 4501.4 4531.8	0.0004	N/A			IR2,
9CI-Pf3ONS	(531.0 / 351.0) 2499 (533.0 / 353.0) 1776	(8.62, 1.51) (N/A, 0.02, -18.9)	19.9 26.0	0.7106 241.2 228.3	0.0009	N/A			IR2,
11CI-PF3OUDS	(631.0 / 451.0) 568 (633.0 / 453.0) 1227	(9.16, 1.61) (N/A, 0.06, -20.0)	190.7 70.0	2.1625 700.7 698.6	0.0004	N/A			IR2,

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) 26 (241.0 / 117.0) 57	(3.90, 0.87) (N/A, -0.10, -7.8)	3.0 2.4	2.2220 128.1 135.1	0.0020	N/A			
5:3FTCA	(341.0 / 236.7) 812 (341.0 / 217.0) 3531	(5.91, 1.10) (N/A, 0.04, 1.0)	5775.6 24.2	4.3470 258.5 239.3	0.0099	N/A			IR2,
7:3FTCA	(441.0 / 317.0) 1365 (441.0 / 337.0) 398	(8.12, 1.51) (N/A, 0.35, 5.9)	70.4 452.1	0.2917 34.0 34.8	0.0093	N/A			IR1,
PFEESA	(315.0 / 135.0) 1333 (315.0 / 83.0) 908	(5.84, 1.08) (N/A, 0.08, -10.2)	875.9 58.2	0.6809 241.9 260.0	0.0019	N/A			IR2,
PFMPA	(229.0 / 85.0) 240	(4.31, 0.97) (N/A, 0.48, 0.0)	20.5	N/A 0.0 0.0	0.0021	N/A			RT,
PFMBA	(279.0 / 85.0) 453	(4.30, 0.96) (N/A, -0.46, 0.0)	24.7	N/A 0.0 0.0	0.0012	N/A			RT,
NFDHA	(295.0 / 201.0) 425 (295.0 / 85.0) 308	(4.86, 0.90) (N/A, -0.42, -28.5)	179627.8 55.0	0.7248 69.4 69.9	0.0013	N/A			RT,
13C3_PFBa_IIS	(216.0 / 172.0) 279071	(3.46, N/A) (N/A, 0.00, N/A)	985.7	N/A	0.8232 [1.0000]	82.3% { 133.2% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 494581	(5.39, N/A) (N/A, 0.02, N/A)	1479.1	N/A	0.8316 [1.0000]	83.2% { 132.7% }			
13C4_PFOA_IIS	(417.0 / 372.0) 727004	(6.90, N/A) (N/A, 0.03, N/A)	1620.2	N/A	0.8937 [1.0000]	89.4% { 140.3% }			
13C5_PFNAl_IIS	(468.0 / 423.0) 565724	(7.51, N/A) (N/A, 0.03, N/A)	972.4	N/A	0.7408 [1.0000]	74.1% { 102.7% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 589012	(8.06 , N/A) (N/A , 0.03 , N/A)	688.5	N/A	0.8205 [1.0000]	82.0% { 123.1% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 1009080	(7.03 , N/A) (N/A , 0.03 , N/A)	1360.7	N/A	0.8194 [1.0000]	81.9% { 114.0% }			
13C4_PFOS_IIS	(503.0 / 79.9) 1598763	(8.26 , N/A) (N/A , 0.03 , N/A)	800.0	N/A	0.9008 [1.0000]	90.1% { 124.0% }			
13C4_PFBA_EIS	(217.0 / 172.0) 2325825	(3.46 , N/A) (N/A , 0.00 , N/A)	2888.7	N/A	7.0089 [8.0000]	87.6% { 121.8% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 2011939	(4.46 , N/A) (N/A , 0.01 , N/A)	2170.1	N/A	3.6509 [4.0000]	91.3% { 114.8% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 1228285	(5.39 , N/A) (N/A , 0.02 , N/A)	1730.6	N/A	1.8278 [2.0000]	91.4% { 115.7% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 1095522	(6.20 , N/A) (N/A , 0.04 , N/A)	1742.2	N/A	1.7480 [2.0000]	87.4% { 121.5% }			
13C8_PFOA_EIS	(421.0 / 376.0) 1254978	(6.90 , N/A) (N/A , 0.04 , N/A)	1420.7	N/A	1.6128 [2.0000]	80.6% { 118.2% }			
13C9_PFNA_EIS	(472.0 / 427.0) 552448	(7.52 , N/A) (N/A , 0.04 , N/A)	1061.9	N/A	0.9799 [1.0000]	98.0% { 110.3% }			
13C6_PFDA_EIS	(519.0 / 474.0) 664303	(8.05 , N/A) (N/A , 0.02 , N/A)	592.2	N/A	0.9744 [1.0000]	97.4% { 106.2% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 705645	(8.56 , N/A) (N/A , 0.03 , N/A)	859.5	N/A	1.0448 [1.0000]	104.5% { 119.8% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 629710	(8.87, N/A) (N/A, 0.03, N/A)	1339.1	N/A	1.0053 [1.0000]	100.5% { 140.6% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 645668	(9.25, N/A) (N/A, 0.02, N/A)	1454.7	N/A	0.9916 [1.0000]	99.2% { 124.7% }			
13C3_PFBs_EIS	(302.0 / 80.0) 4093059	(5.34, N/A) (N/A, 0.02, N/A)	1692.2	N/A	2.0073 [2.0000]	100.4% { 114.7% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 1858431	(7.03, N/A) (N/A, 0.03, N/A)	991.8	N/A	1.7444 [2.0000]	87.2% { 106.1% }			
13C8_PFOS_EIS	(507.0 / 80.0) 3708917	(8.26, N/A) (N/A, 0.03, N/A)	1234.7	N/A	1.7658 [2.0000]	88.3% { 113.3% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 486615	(5.13, N/A) (N/A, 0.01, N/A)	1037.5	N/A	3.8962 [4.0000]	97.4% { 120.7% }			
13C2_6:2FTS_EIS	(429.0 / 81.0) 468445	(6.64, N/A) (N/A, 0.03, N/A)	1008.0	N/A	3.3267 [4.0000]	83.2% { 109.8% }			
13C2_8:2FTS_EIS	(529.0 / 81.0) 626346	(7.81, N/A) (N/A, 0.03, N/A)	1277.9	N/A	4.2062 [4.0000]	105.2% { 119.8% }			
13C8_PFOsa_EIS	(506.0 / 78.0) 6787909	(9.84, N/A) (N/A, 0.02, N/A)	2785.6	N/A	1.7544 [2.0000]	87.7% { 117.2% }			
D3_NMeFOSA_EIS	(515.0 / 169.0) 1468681	(10.44, N/A) (N/A, 0.02, N/A)	1764.9	N/A	1.7678 [2.0000]	88.4% { 110.6% }			
D5_NEtFOSA_EIS	(531.0 / 169.0) 1502392	(10.59, N/A) (N/A, 0.02, N/A)	2870.1	N/A	2.0653 [2.0000]	103.3% { 124.0% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1268350	(8.22 , N/A) (N/A , 0.03 , N/A)	1269.8	N/A	4.3027 [4.0000]	107.6% { 114.2% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 979331	(8.47 , N/A) (N/A , 0.03 , N/A)	11787581.6	N/A	3.9014 [4.0000]	97.5% { 116.2% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 5317743	(10.38 , N/A) (N/A , 0.02 , N/A)	2313.6	N/A	18.8010 [20.0000]	94.0% { 124.3% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 6636993	(10.54 , N/A) (N/A , 0.02 , N/A)	1399.6	N/A	20.5137 [20.0000]	102.6% { 124.3% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 3161126	(5.69 , N/A) (N/A , 0.03 , N/A)	1887.9	N/A	7.4862 [8.0000]	93.6% { 113.7% }			

ANALYSIS SEQUENCE BLANKS

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 Client: AECOM
 Sequence: SC00861
 Calibration: 2309009

SDG:
 Project: Red Hill AFFF Assessment Sam
 Instrument: Saphira

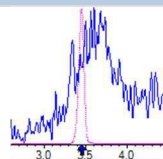
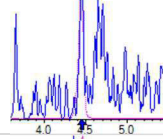
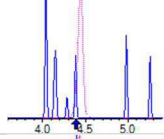
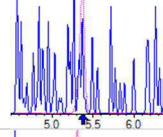
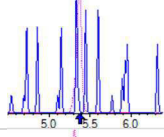
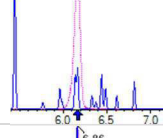
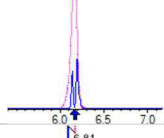
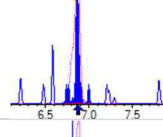
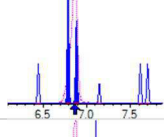
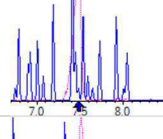
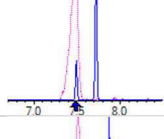
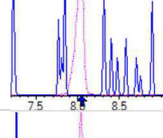
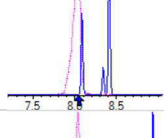
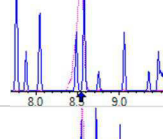
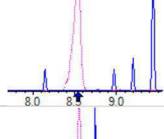
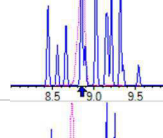
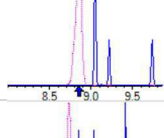
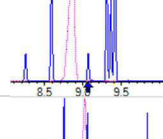
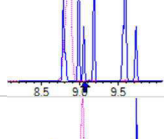
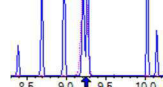
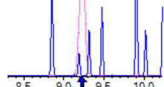
Lab Sample ID	Analyte	Found	Units	RL	C
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	PFPEA	0.00	ng/mL	0.20	U
	PFHXA	0.00	ng/mL	0.10	U
	PFHPA	0.00	ng/mL	0.10	U
	PFOA	0.0129	ng/mL	0.10	U
	PFNA	0.00	ng/mL	0.10	U
	PFDA	0.00	ng/mL	0.10	U
	PFUnA	0.00	ng/mL	0.10	U
	PFDOA	0.00	ng/mL	0.10	U
	PFTRDA	0.00	ng/mL	0.10	U
	PFTEDA	0.00	ng/mL	0.10	U
	PFBS	0.00	ng/mL	0.10	U
	PFPEs	0.00	ng/mL	0.10	U
	PFHXS	0.00	ng/mL	0.10	U
	PFHPS	0.00	ng/mL	0.10	U
	PFOS	0.00	ng/mL	0.10	U
	PFNS	0.000866	ng/mL	0.10	U
	PFDS	0.000952	ng/mL	0.10	U
	PFDOS	0.00110	ng/mL	0.10	U
	4:2FTS	0.00111	ng/mL	0.40	U
	6:2FTS	0.00661	ng/mL	0.40	U
	8:2FTS	0.0335	ng/mL	0.40	U
	PFOSA	0.00386	ng/mL	0.10	U
	NMeFOSA	0.00712	ng/mL	0.40	U
	NEtFOSA	0.00486	ng/mL	0.40	U
	NMeFOSAA	0.00307	ng/mL	0.10	U
	NEtFOSAA	0.00634	ng/mL	0.10	U
	NMeFOSE	0.0118	ng/mL	0.40	U
	NEtFOSE	0.00933	ng/mL	0.40	U
	HFPO-DA	0.00228	ng/mL	0.20	U
	ADONA	0.00	ng/mL	0.20	U
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	PFMPA	0.00204	ng/mL	0.20	U

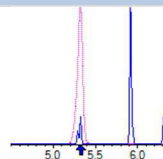
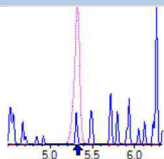
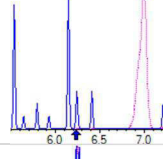
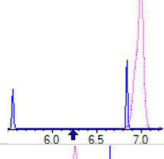
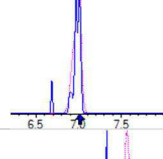
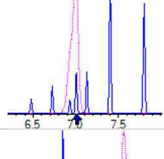
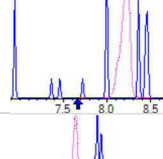
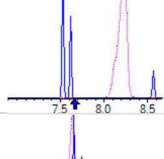
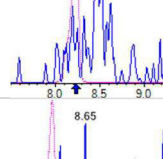
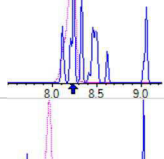
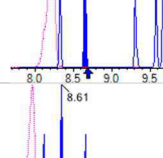
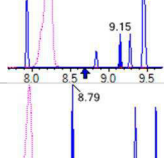
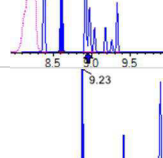
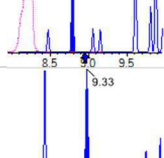
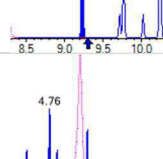
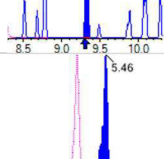
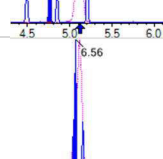
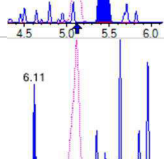
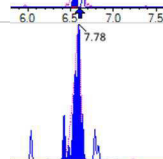
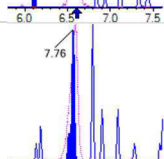
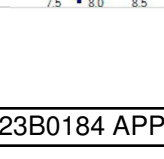
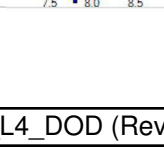
ANALYSIS SEQUENCE BLANKS

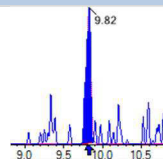
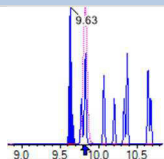
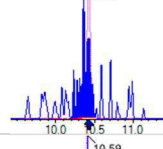
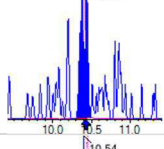
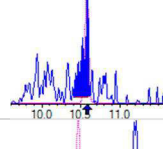
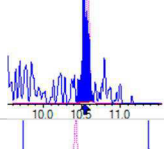
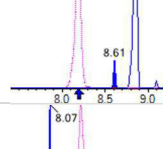
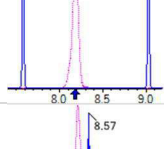
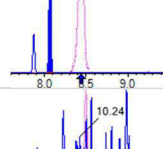
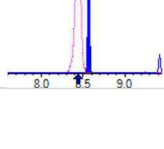
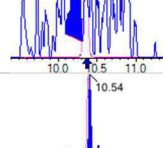
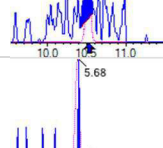
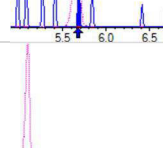
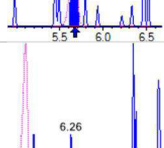
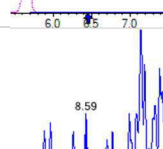
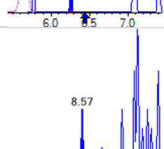
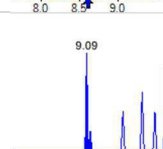
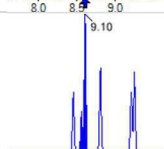
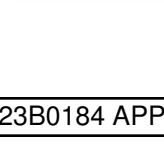
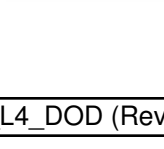
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 Calibration: 2309009

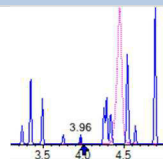
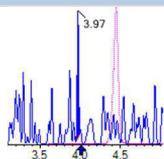
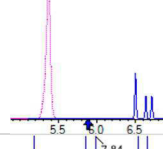
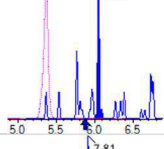
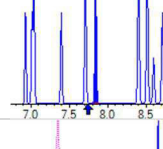
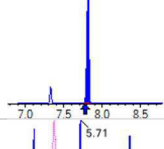
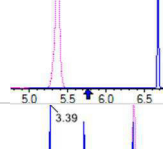
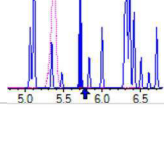
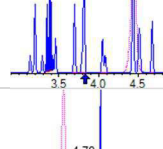
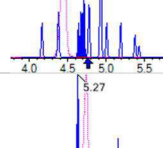
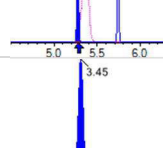
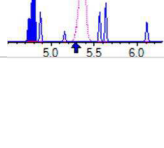
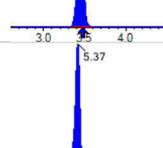
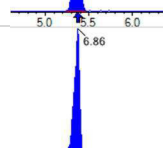
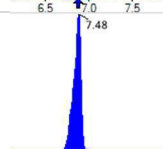
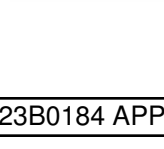
SDG:
 Project: Red Hill AFFF Assessment Sam
 Instrument: Saphira

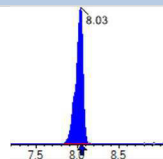
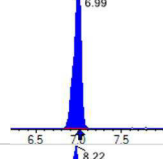
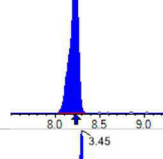
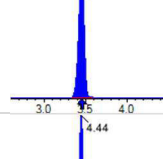
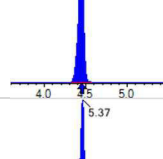
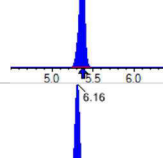
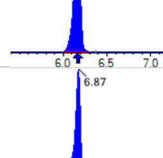
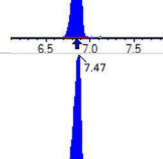
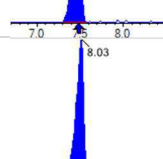
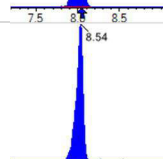
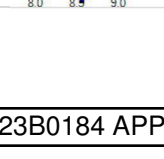
Lab Sample ID	Analyte	Found	Units	RL	C
SC00861-CCB4	PFMBA	0.00187	ng/mL	0.20	U
	NFDHA	0.00529	ng/mL	0.20	U
	9CL-PF3ONS	0.000460	ng/mL	0.20	U
	11CL-PF3OUDS	0.000853	ng/mL	0.20	U
	3:3FTCA	0.00204	ng/mL	0.40	U
	5:3FTCA	0.00	ng/mL	0.40	U
	7:3FTCA	0.00376	ng/mL	0.40	U
	13C4-PFBA	6.87	ng/mL		
	13C5-PFPEA	3.35	ng/mL		
	13C5-PFHXA	1.77	ng/mL		
	13C4-PFHPA	1.64	ng/mL		
	13C8-PFOA	1.79	ng/mL		
	13C9-PFNA	0.832	ng/mL		
	13C6-PFDA	0.927	ng/mL		
	13C7-PFUnA	0.917	ng/mL		
	13C2-PFDOA	0.909	ng/mL		
	13C2-PFTEDA	0.923	ng/mL		
	13C3-PFBS	2.04	ng/mL		
	13C3-PFHXS	1.84	ng/mL		
	13C8-PFOS	1.76	ng/mL		
	13C2-4:2FTS	3.56	ng/mL		
	13C2-6:2FTS	3.69	ng/mL		
	13C2-8:2FTS	4.09	ng/mL		
	13C8-PFOSA	1.64	ng/mL		
	D3-NMEFOSA	1.80	ng/mL		
	D5-NETFOSA	2.01	ng/mL		
	D3-NMEFOSAA	4.06	ng/mL		
	D5-NETFOSAA	3.82	ng/mL		
	D7-NMEFOSE	18.5	ng/mL		
	D9-NETFOSSE	20.1	ng/mL		
	13C3-HFPO-DA	6.73	ng/mL		

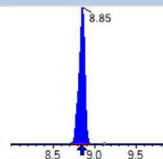
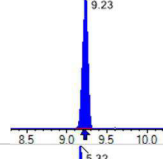
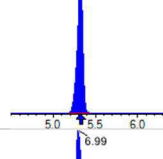
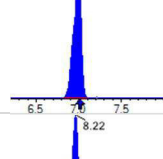
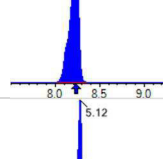
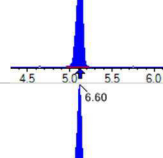
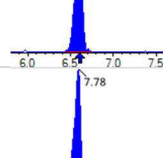
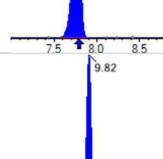
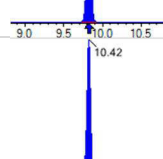
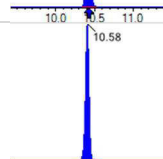
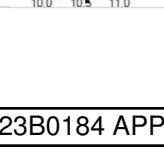
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
PFPeA	(263.0 / 219.0) N/A (263.0 / 69.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHxA	(313.0 / 269.0) N/A (313.0 / 119.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHpA	(363.0 / 319.0) N/A (363.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFOA	(413.0 / 369.0) 7961 (413.0 / 169.0) 1988	(6.86, 1.00) (0.00, N/A, 3.5)	251399.0 5331.5	0.2497 73.8 70.3	0.0129	N/A			
PFNA	(463.0 / 419.0) N/A (463.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDA	(513.0 / 469.0) N/A (513.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFUnA	(563.0 / 519.0) N/A (563.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDaA	(613.0 / 569.0) N/A (613.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFTrDA	(663.0 / 619.0) N/A (663.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFTeDA	(713.0 / 669.0) N/A (713.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

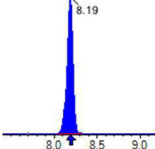
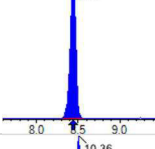
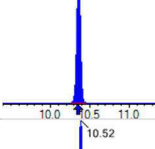
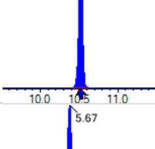
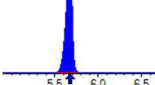
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) N/A (299.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFPeS	(349.0 / 80.0) N/A (349.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHxS	(399.0 / 80.0) N/A (399.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHpS	(449.0 / 80.0) N/A (449.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFOS	(499.0 / 80.0) N/A (499.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFNS	(549.0 / 80.0) 1867 (549.0 / 99.0) 379	(8.65 , 1.05) (N/A , -0.03 , -29.8)	5234.0 44975.6	0.2028 84.3 88.0	0.0009	N/A			
PFDS	(599.0 / 80.0) 2282 (599.0 / 99.0) 1189	(8.61 , 1.05) (N/A , -0.34 , -10.9)	40.0 72.7	0.5208 257.3 243.9	0.0010	N/A			IR2,
PFDoS	(699.0 / 80.0) 1215 (699.0 / 99.0) 2951	(9.23 , 1.12) (N/A , -0.07 , -5.8)	240.8 34.0	2.4295 1243.5 1164.1	0.0011	N/A			IR2,
4:2FTS	(327.0 / 307.0) 343 (327.0 / 81.0) 13220	(4.76 , 0.93) (-0.36 , N/A , -42.0)	520.8 60.8	38.5782 5567.1 6084.2	0.0011	N/A			RT, IR2,
6:2FTS	(427.0 / 407.0) 1105 (427.0 / 81.0) 731	(6.56 , 0.99) (-0.05 , N/A , 26.9)	6582.0 448.1	0.6618 78.1 86.4	0.0066	N/A			
8:2FTS	(527.0 / 507.0) 5661 (527.0 / 81.0) 2641	(7.78 , 1.00) (0.00 , N/A , 1.3)	162.3 36.5	0.4664 57.2 58.5	0.0335	N/A			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) 10062 (498.0 / 478.0) 674	(9.82 , 1.00) (0.00 , N/A , 11.5)	29.7 32.0	0.0670 340.8 328.4	0.0039	N/A			
NMeFOSA	(512.0 / 219.0) 4814 (512.0 / 169.0) 2862	(10.37 , 0.99) (-0.05 , N/A , -1.9)	42.3 20.1	0.5945 72.0 71.1	0.0071	N/A			
NEiFOSA	(526.0 / 219.0) 3640 (526.0 / 169.0) 4632	(10.59 , 1.00) (0.01 , N/A , 3.0)	34.6 28.9	1.2727 100.5 99.3	0.0049	N/A			
NMeFOSAA	(570.0 / 419.0) 830 (570.0 / 483.0) N/A	(8.61 , 1.05) (0.42 , N/A , #Value!)	312.1 N/A	N/A 0.0 0.0	0.0031	N/A			RT,IR1,
NEiFOSAA	(584.0 / 419.0) 1396 (584.0 / 526.0) 1452	(8.07 , 0.96) (-0.37 , N/A , -30.1)	8045.6 295.0	1.0404 184.1 203.1	0.0063	N/A			RT,IR2,
NMeFOSE	(616.0 / 59.0) 3401	(10.24 , 0.99) (-0.11 , N/A , 0.0)	6.4	N/A 0.0 0.0	0.0118	N/A			RT,
NEiFOSE	(630.0 / 59.0) 3227	(10.54 , 1.00) (0.02 , N/A , 0.0)	16.1	N/A 0.0 0.0	0.0093	N/A			
HFPO-DA	(285.0 / 169.0) 710 (285.0 / 185.0) 3907	(5.68 , 1.00) (0.01 , N/A , 1.0)	46.7 47.0	5.5055 200.3 182.0	0.0023	N/A			IR2,
ADONA	(377.0 / 85.0) N/A (377.0 / 251.0) 381	(N/A , N/A) (N/A , N/A , N/A)	N/A 3548.1	N/A 0.0 0.0	0.0000	N/A			
9CI-Pf3ONS	(531.0 / 351.0) 1244 (533.0 / 353.0) 622	(8.59 , 1.52) (N/A , -0.01 , 1.2)	6.1 13.3	0.5001 169.8 160.7	0.0005	N/A			IR2,
11CI-PF3OUDS	(631.0 / 451.0) 1135 (633.0 / 453.0) 1703	(9.09 , 1.60) (N/A , -0.02 , -1.1)	305.9 625.6	1.4995 485.9 484.4	0.0009	N/A			IR2,

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) 26 (241.0 / 117.0) 449	(3.96, 0.89) (N/A, -0.04, -0.5)	4.7 14.2	17.1352 988.1 1042.1	0.0020	N/A			IR2,
5:3FTCA	(341.0 / 236.7) N/A (341.0 / 217.0) 2679	(N/A, N/A) (N/A, N/A, N/A)	N/A 36.0	N/A 0.0 0.0	0.0000	N/A			
7:3FTCA	(441.0 / 317.0) 574 (441.0 / 337.0) 1907	(7.84, 1.46) (N/A, 0.07, 1.8)	15.8 15726.5	3.3230 386.9 395.8	0.0038	N/A			IR2,
PFEESA	(315.0 / 135.0) N/A (315.0 / 83.0) 1397	(N/A, N/A) (N/A, N/A, N/A)	N/A 21.1	N/A 0.0 0.0	0.0000	N/A			
PFMPA	(229.0 / 85.0) 226	(3.39, 0.76) (N/A, -0.44, 0.0)	24.6	N/A 0.0 0.0	0.0020	N/A			RT,
PFMBA	(279.0 / 85.0) 712	(4.70, 1.06) (N/A, -0.06, 0.0)	21.2	N/A 0.0 0.0	0.0019	N/A			
NFDHA	(295.0 / 201.0) 1745 (295.0 / 85.0) 2652	(5.27, 0.98) (N/A, 0.00, 28.8)	79170.2 246.8	1.5195 145.6 146.5	0.0053	N/A			
13C3_PFBA_IIS	(216.0 / 172.0) 286011	(3.45, N/A) (N/A, -0.01, N/A)	1257.8	N/A	0.8436 [1.0000]	84.4% { 136.6% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 527602	(5.37, N/A) (N/A, 0.00, N/A)	951.3	N/A	0.8871 [1.0000]	88.7% { 141.6% }			
13C4_PFOA_IIS	(417.0 / 372.0) 666354	(6.86, N/A) (N/A, 0.00, N/A)	1294.1	N/A	0.8192 [1.0000]	81.9% { 128.6% }			
13C5_PFNA_IIS	(468.0 / 423.0) 657623	(7.48, N/A) (N/A, 0.00, N/A)	1248.1	N/A	0.8612 [1.0000]	86.1% { 119.4% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 627947	(8.03 , N/A) (N/A , 0.00 , N/A)	770.1	N/A	0.8747 [1.0000]	87.5% { 131.2% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 995626	(6.99 , N/A) (N/A , -0.01 , N/A)	871.6	N/A	0.8084 [1.0000]	80.8% { 112.5% }			
13C4_PFOS_IIS	(503.0 / 79.9) 1685321	(8.22 , N/A) (N/A , 0.00 , N/A)	685.0	N/A	0.9496 [1.0000]	95.0% { 130.7% }			
13C4_PFBA_EIS	(217.0 / 172.0) 2336909	(3.45 , N/A) (N/A , 0.00 , N/A)	3207.0	N/A	6.8714 [8.0000]	85.9% { 122.4% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 1970472	(4.44 , N/A) (N/A , -0.01 , N/A)	2125.8	N/A	3.3518 [4.0000]	83.8% { 112.4% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 1272267	(5.37 , N/A) (N/A , 0.00 , N/A)	1950.6	N/A	1.7747 [2.0000]	88.7% { 119.8% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 1097319	(6.16 , N/A) (N/A , 0.00 , N/A)	1796.5	N/A	1.6413 [2.0000]	82.1% { 121.7% }			
13C8_PFOA_EIS	(421.0 / 376.0) 1279792	(6.87 , N/A) (N/A , 0.00 , N/A)	1193.4	N/A	1.7944 [2.0000]	89.7% { 120.5% }			
13C9_PFNA_EIS	(472.0 / 427.0) 545298	(7.47 , N/A) (N/A , -0.01 , N/A)	669.5	N/A	0.8321 [1.0000]	83.2% { 108.8% }			
13C6_PFDA_EIS	(519.0 / 474.0) 674007	(8.03 , N/A) (N/A , 0.00 , N/A)	1089.6	N/A	0.9274 [1.0000]	92.7% { 107.8% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 660364	(8.54 , N/A) (N/A , 0.00 , N/A)	1154.0	N/A	0.9171 [1.0000]	91.7% { 112.1% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT Ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 607126	(8.85 , N/A) (N/A , 0.01 , N/A)	1327.1	N/A	0.9091 [1.0000]	90.9% { 135.6% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 641032	(9.23 , N/A) (N/A , 0.00 , N/A)	1012.2	N/A	0.9234 [1.0000]	92.3% { 123.8% }			
13C3_PFBs_EIS	(302.0 / 80.0) 4102577	(5.32 , N/A) (N/A , 0.00 , N/A)	2313.3	N/A	2.0391 [2.0000]	102.0% { 114.9% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 1939311	(6.99 , N/A) (N/A , -0.01 , N/A)	1218.4	N/A	1.8450 [2.0000]	92.2% { 110.7% }			
13C8_PFOS_EIS	(507.0 / 80.0) 3904558	(8.22 , N/A) (N/A , 0.00 , N/A)	1274.1	N/A	1.7635 [2.0000]	88.2% { 119.3% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 438179	(5.12 , N/A) (N/A , 0.00 , N/A)	1089.2	N/A	3.5558 [4.0000]	88.9% { 108.7% }			
13C2_6:2FTS_EIS	(429.0 / 81.0) 512500	(6.60 , N/A) (N/A , 0.00 , N/A)	988.9	N/A	3.6887 [4.0000]	92.2% { 120.1% }			
13C2_8:2FTS_EIS	(529.0 / 81.0) 601605	(7.78 , N/A) (N/A , 0.00 , N/A)	961.9	N/A	4.0947 [4.0000]	102.4% { 115.1% }			
13C8_PFOSA_EIS	(506.0 / 78.0) 6706537	(9.82 , N/A) (N/A , 0.00 , N/A)	3245.8	N/A	1.6444 [2.0000]	82.2% { 115.8% }			
D3_NMeFOSA_EIS	(515.0 / 169.0) 1574786	(10.42 , N/A) (N/A , 0.01 , N/A)	2169.6	N/A	1.7982 [2.0000]	89.9% { 118.5% }			
D5_NEtFOSA_EIS	(531.0 / 169.0) 1540606	(10.58 , N/A) (N/A , 0.00 , N/A)	3125.5	N/A	2.0091 [2.0000]	100.5% { 127.2% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1262600	(8.19 , N/A) (N/A , 0.00 , N/A)	1279.8	N/A	4.0632 [4.0000]	101.6% { 113.7% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 1010247	(8.44 , N/A) (N/A , 0.00 , N/A)	1961.2	N/A	3.8179 [4.0000]	95.4% { 119.9% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 5501063	(10.36 , N/A) (N/A , 0.01 , N/A)	2007.6	N/A	18.4502 [20.0000]	92.3% { 128.6% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 6848751	(10.52 , N/A) (N/A , 0.01 , N/A)	1698.9	N/A	20.0810 [20.0000]	100.4% { 128.3% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 3033472	(5.67 , N/A) (N/A , 0.00 , N/A)	1498.9	N/A	6.7342 [8.0000]	84.2% { 109.1% }			

ANALYSIS SEQUENCE BLANKS

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 Client: AECOM
 Sequence: SC00861
 Calibration: 2309009

SDG:
 Project: Red Hill AFFF Assessment Sam
 Instrument: Saphira

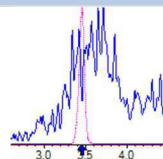
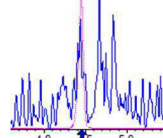
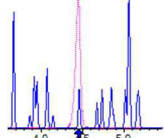
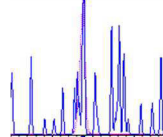
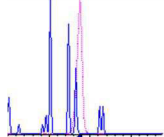
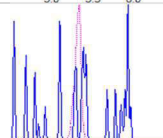
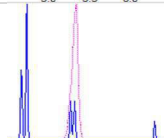
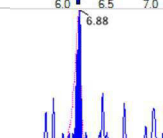
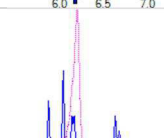
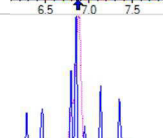
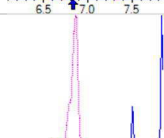
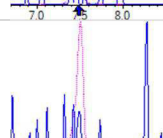
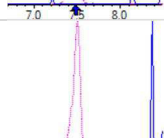
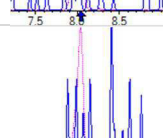
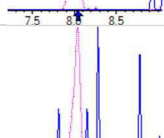
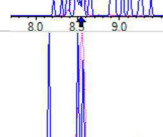
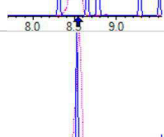
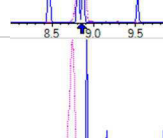
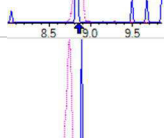
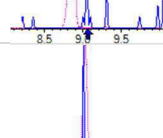
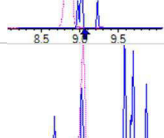
Lab Sample ID	Analyte	Found	Units	RL	C
SC00861-CCB5	PFBA	0.00	ng/mL	0.40	U
	PFPEA	0.00	ng/mL	0.20	U
	PFHXA	0.00	ng/mL	0.10	U
	PFHPA	0.00	ng/mL	0.10	U
	PFOA	0.0141	ng/mL	0.10	U
	PFNA	0.00	ng/mL	0.10	U
	PFDA	0.00	ng/mL	0.10	U
	PFUnA	0.00	ng/mL	0.10	U
	PFDOA	0.00	ng/mL	0.10	U
	PFTRDA	0.00	ng/mL	0.10	U
	PFTEDA	0.00	ng/mL	0.10	U
	PFBS	0.00	ng/mL	0.10	U
	PFPEs	0.00	ng/mL	0.10	U
	PFHXS	0.00	ng/mL	0.10	U
	PFHPS	0.00	ng/mL	0.10	U
	PFOS	0.00	ng/mL	0.10	U
	PFNS	0.000898	ng/mL	0.10	U
	PFDS	0.000585	ng/mL	0.10	U
	PFDOS	0.00200	ng/mL	0.10	U
	4:2FTS	0.00318	ng/mL	0.40	U
	6:2FTS	0.00	ng/mL	0.40	U
	8:2FTS	0.00476	ng/mL	0.40	U
	PFOSA	0.00376	ng/mL	0.10	U
	NMeFOSA	0.00475	ng/mL	0.40	U
	NEtFOSA	0.00558	ng/mL	0.40	U
	NMeFOSAA	0.00829	ng/mL	0.10	U
	NEtFOSAA	0.000846	ng/mL	0.10	U
	NMeFOSE	0.0138	ng/mL	0.40	U
	NEtFOSE	0.0174	ng/mL	0.40	U
	HFPO-DA	0.00231	ng/mL	0.20	U
	ADONA	0.000522	ng/mL	0.20	U
	PFEESA	0.000738	ng/mL	0.20	U
	PFMPA	0.00262	ng/mL	0.20	U

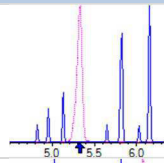
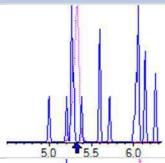
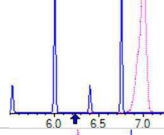
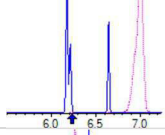
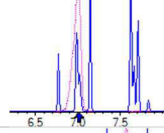
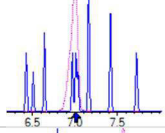
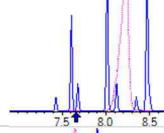
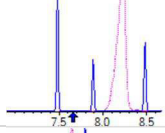
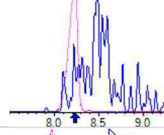
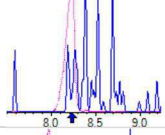
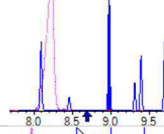
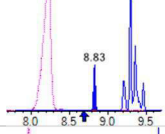
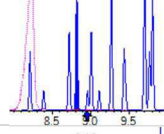
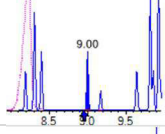
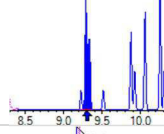
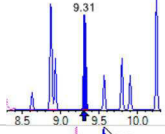
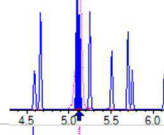
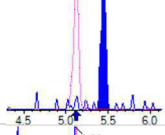
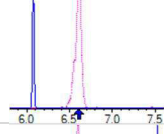
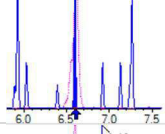
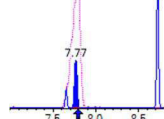
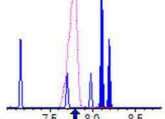
ANALYSIS SEQUENCE BLANKS

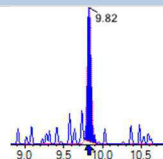
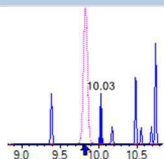
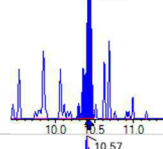
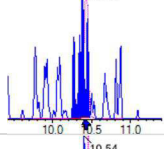
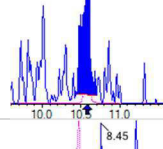
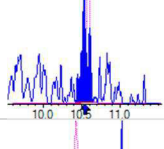
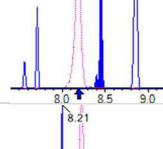
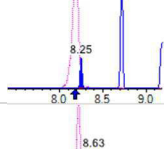
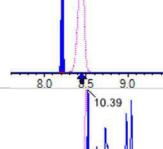
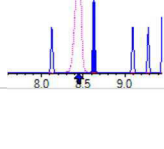
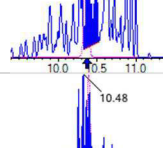
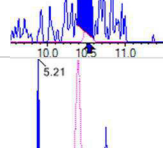
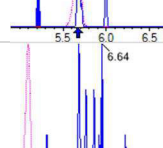
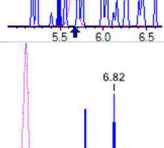
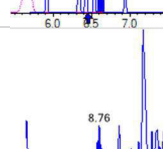
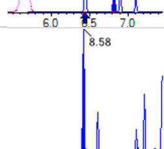
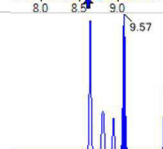
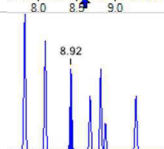
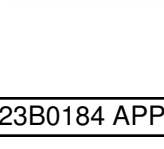
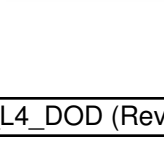
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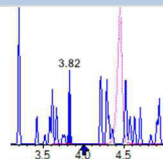
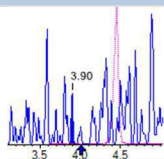
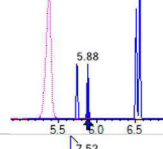
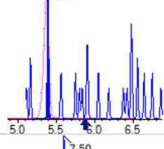
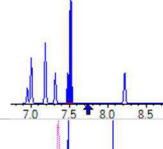
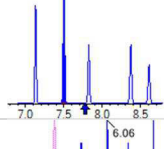
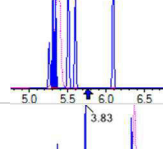
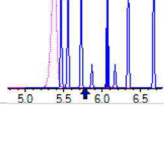
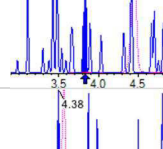
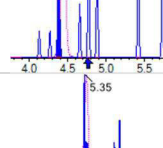
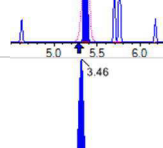
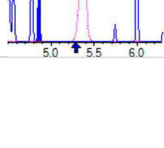
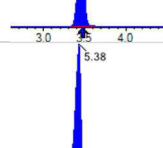
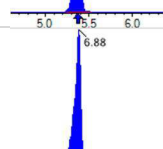
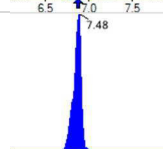
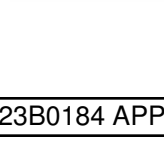
SDG:
 Project: Red Hill AFFF Assessment Sam
 Instrument: Saphira

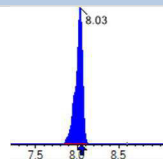
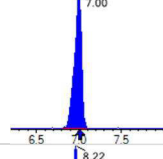
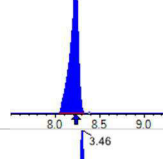
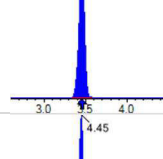
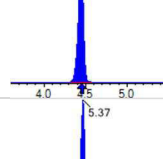
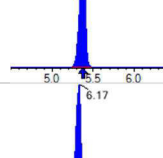
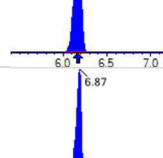
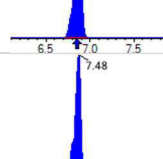
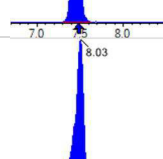
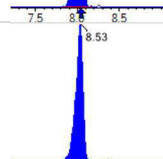
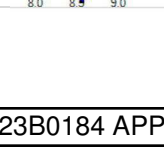
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	NFDHA	0.00303	ng/mL	0.20	U
	9CL-PF3ONS	0.00101	ng/mL	0.20	U
	11CL-PF3OUDS	0.000769	ng/mL	0.20	U
	3:3FTCA	0.0152	ng/mL	0.40	U
	5:3FTCA	0.00352	ng/mL	0.40	U
	7:3FTCA	0.0139	ng/mL	0.40	U
	13C4-PFBA	7.02	ng/mL		
	13C5-PFPEA	3.84	ng/mL		
	13C5-PFHXA	1.82	ng/mL		
	13C4-PFHPA	1.76	ng/mL		
	13C8-PFOA	1.91	ng/mL		
	13C9-PFNA	0.891	ng/mL		
	13C6-PFDA	0.890	ng/mL		
	13C7-PFUnA	1.03	ng/mL		
	13C2-PFDOA	0.962	ng/mL		
	13C2-PFTEDA	0.869	ng/mL		
	13C3-PFBS	2.11	ng/mL		
	13C3-PFHXS	1.92	ng/mL		
	13C8-PFOS	1.79	ng/mL		
	13C2-4:2FTS	4.20	ng/mL		
	13C2-6:2FTS	3.56	ng/mL		
	13C2-8:2FTS	4.03	ng/mL		
	13C8-PFOSA	1.72	ng/mL		
	D3-NMEFOSA	1.78	ng/mL		
	D5-NETFOSA	1.96	ng/mL		
	D3-NMEFOSAA	4.12	ng/mL		
	D5-NETFOSAA	3.96	ng/mL		
	D7-NMEFOSE	18.4	ng/mL		
	D9-NETFOSE	20.5	ng/mL		
	13C3-HFPO-DA	7.49	ng/mL		

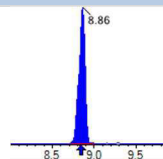
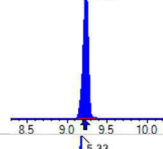
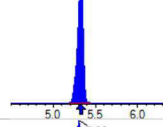
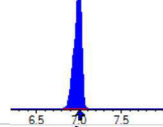
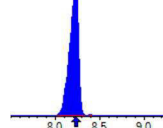
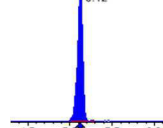
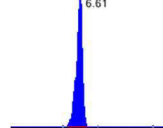
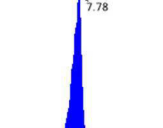
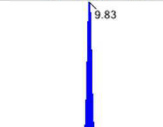
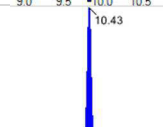
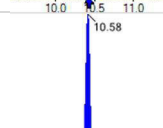
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
PFPeA	(263.0 / 219.0) N/A (263.0 / 69.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHxA	(313.0 / 269.0) N/A (313.0 / 119.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHpA	(363.0 / 319.0) N/A (363.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFOA	(413.0 / 369.0) 8666 (413.0 / 169.0) N/A	(6.88 , 1.00) (0.01 , N/A , #Value!)	970.5 N/A	N/A 0.0 0.0	0.0141	N/A			IR1,
PFNA	(463.0 / 419.0) N/A (463.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDA	(513.0 / 469.0) N/A (513.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFUnA	(563.0 / 519.0) N/A (563.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDaA	(613.0 / 569.0) N/A (613.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFTTrDA	(663.0 / 619.0) N/A (663.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFTeDA	(713.0 / 669.0) N/A (713.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

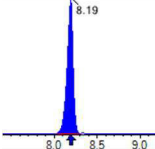
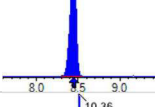
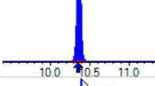
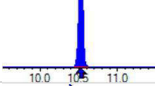
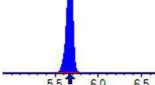
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) N/A (299.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFPeS	(349.0 / 80.0) N/A (349.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHxS	(399.0 / 80.0) N/A (399.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHpS	(449.0 / 80.0) N/A (449.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFOS	(499.0 / 80.0) N/A (499.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFNS	(549.0 / 80.0) 1971 (549.0 / 99.0) 1020	(8.98 , 1.09) (N/A , 0.29 , 9.0)	7507.6 51303.6	0.5176 215.1 224.5	0.0009	N/A			IR2,
PFDS	(599.0 / 80.0) 1426 (599.0 / 99.0) 587	(8.82 , 1.07) (N/A , -0.13 , -10.8)	16.3 13.7	0.4117 203.4 192.8	0.0006	N/A			IR2,
PFDoS	(699.0 / 80.0) 2250 (699.0 / 99.0) 1563	(9.30 , 1.13) (N/A , 0.00 , -0.9)	32.6 52.2	0.6946 355.5 332.8	0.0020	N/A			IR2,
4:2FTS	(327.0 / 307.0) 1107 (327.0 / 81.0) 16783	(5.11 , 1.00) (-0.02 , N/A , -20.5)	70.3 72.6	15.1550 2187.0 2390.1	0.0032	N/A			IR2,
6:2FTS	(427.0 / 407.0) N/A (427.0 / 81.0) 944	(N/A , N/A) (N/A , N/A , N/A)	N/A 123.1	N/A 0.0 0.0	0.0000	N/A			
8:2FTS	(527.0 / 507.0) 755 (527.0 / 81.0) 1176	(7.77 , 1.00) (-0.01 , N/A , -20.3)	107.9 892.8	1.5574 190.9 195.2	0.0048	N/A			IR2,

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) 10324 (498.0 / 478.0) 253	(9.82 , 1.00) (-0.01 , N/A , -12.3)	40.3 40.6	0.0245 124.6 120.1	0.0038	N/A			
NMeFOSA	(512.0 / 219.0) 3188 (512.0 / 169.0) 2933	(10.43 , 1.00) (0.00 , N/A , 2.8)	68.3 38.7	0.9199 111.4 109.9	0.0047	N/A			
NEiFOSA	(526.0 / 219.0) 4089 (526.0 / 169.0) 5784	(10.57 , 1.00) (-0.01 , N/A , 2.1)	27.3 27.3	1.4146 111.7 110.4	0.0056	N/A			
NMeFOSAA	(570.0 / 419.0) 2279 (570.0 / 483.0) 582	(8.45 , 1.03) (0.26 , N/A , 12.1)	143.7 2662.8	0.2554 52.1 57.7	0.0083	N/A			RT,
NEiFOSAA	(584.0 / 419.0) 194 (584.0 / 526.0) 415	(8.21 , 0.97) (-0.23 , N/A , -25.0)	3545.6 469.5	2.1389 378.6 417.5	0.0008	N/A			RT,IR2,
NMeFOSE	(616.0 / 59.0) 3974	(10.39 , 1.00) (0.03 , N/A , 0.0)	17.0	N/A 0.0 0.0	0.0138	N/A			
NEiFOSE	(630.0 / 59.0) 6182	(10.48 , 1.00) (-0.04 , N/A , 0.0)	14.6	N/A 0.0 0.0	0.0174	N/A			
HFPO-DA	(285.0 / 169.0) 723 (285.0 / 185.0) 1271	(5.21 , 0.92) (-0.46 , N/A , -16.5)	2463695.7 16.2	1.7582 64.0 58.1	0.0023	N/A			RT,
ADONA	(377.0 / 85.0) 605 (377.0 / 251.0) 965	(6.64 , 1.17) (N/A , 0.19 , -11.1)	72.1 433.4	1.5956 1822.8 1835.2	0.0005	N/A			IR2,
9CI-Pf3ONS	(531.0 / 351.0) 2753 (533.0 / 353.0) 1659	(8.76 , 1.54) (N/A , 0.16 , 10.4)	8.0 33.7	0.6027 204.6 193.6	0.0010	N/A			IR2,
11CI-PF3OUDS	(631.0 / 451.0) 1028 (633.0 / 453.0) 592	(9.57 , 1.69) (N/A , 0.47 , 39.1)	459.6 44.9	0.5759 186.6 186.0	0.0008	N/A			RT,IR2,

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) 201 (241.0 / 117.0) 176	(3.82, 0.86) (N/A, -0.18, -4.3)	11.6 4.9	0.8730 50.3 53.1	0.0152	N/A			
5:3FTCA	(341.0 / 236.7) 276 (341.0 / 217.0) 1183	(5.88, 1.09) (N/A, 0.00, 29.4)	2327.8 17.6	4.2871 255.0 236.0	0.0035	N/A			IR2,
7:3FTCA	(441.0 / 317.0) 1956 (441.0 / 337.0) 1241	(7.52, 1.40) (N/A, -0.26, 1.1)	128.8 807.2	0.6347 73.9 75.6	0.0139	N/A			
PFEESA	(315.0 / 135.0) 503 (315.0 / 83.0) 778	(5.32, 0.99) (N/A, -0.43, -44.5)	1726.5 57.3	1.5460 549.2 590.4	0.0007	N/A			RT, IR2,
PFMPA	(229.0 / 85.0) 300	(3.83, 0.86) (N/A, 0.00, 0.0)	16.9	N/A 0.0 0.0	0.0026	N/A			
PFMBA	(279.0 / 85.0) 351	(4.38, 0.98) (N/A, -0.38, 0.0)	27.3	N/A 0.0 0.0	0.0009	N/A			
NFDHA	(295.0 / 201.0) 925 (295.0 / 85.0) 637	(5.35, 1.00) (N/A, 0.07, 29.5)	532.2 119.3	0.6891 66.0 66.4	0.0030	N/A			
13C3_PFBA_IIS	(216.0 / 172.0) 271124	(3.46, N/A) (N/A, 0.00, N/A)	1232.5	N/A	0.7997 [1.0000]	80.0% { 129.5% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 476808	(5.38, N/A) (N/A, 0.01, N/A)	1521.9	N/A	0.8017 [1.0000]	80.2% { 128.0% }			
13C4_PFOA_IIS	(417.0 / 372.0) 626290	(6.88, N/A) (N/A, 0.01, N/A)	884.7	N/A	0.7699 [1.0000]	77.0% { 120.9% }			
13C5_PFNA_IIS	(468.0 / 423.0) 637959	(7.48, N/A) (N/A, 0.00, N/A)	1453.6	N/A	0.8354 [1.0000]	83.5% { 115.8% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 622825	(8.03 , N/A) (N/A , 0.00 , N/A)	735.1	N/A	0.8676 [1.0000]	86.8% { 130.1% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 950571	(7.00 , N/A) (N/A , 0.00 , N/A)	891.2	N/A	0.7718 [1.0000]	77.2% { 107.4% }			
13C4_PFOS_IIS	(503.0 / 79.9) 1693167	(8.22 , N/A) (N/A , 0.00 , N/A)	1009.9	N/A	0.9540 [1.0000]	95.4% { 131.3% }			
13C4_PFBA_EIS	(217.0 / 172.0) 2263007	(3.46 , N/A) (N/A , 0.00 , N/A)	3134.3	N/A	7.0194 [8.0000]	87.7% { 118.5% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 2039446	(4.45 , N/A) (N/A , 0.00 , N/A)	2319.4	N/A	3.8387 [4.0000]	96.0% { 116.4% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 1176624	(5.37 , N/A) (N/A , 0.00 , N/A)	1662.0	N/A	1.8162 [2.0000]	90.8% { 110.8% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 1065958	(6.17 , N/A) (N/A , 0.01 , N/A)	1737.5	N/A	1.7642 [2.0000]	88.2% { 118.2% }			
13C8_PFOA_EIS	(421.0 / 376.0) 1280075	(6.87 , N/A) (N/A , 0.01 , N/A)	1685.0	N/A	1.9096 [2.0000]	95.5% { 120.6% }			
13C9_PFNA_EIS	(472.0 / 427.0) 566263	(7.48 , N/A) (N/A , -0.01 , N/A)	1016.0	N/A	0.8907 [1.0000]	89.1% { 113.0% }			
13C6_PFDA_EIS	(519.0 / 474.0) 641428	(8.03 , N/A) (N/A , 0.00 , N/A)	660.0	N/A	0.8898 [1.0000]	89.0% { 102.6% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 732422	(8.53 , N/A) (N/A , 0.00 , N/A)	768.7	N/A	1.0256 [1.0000]	102.6% { 124.4% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT Ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 637162	(8.86 , N/A) (N/A , 0.01 , N/A)	1191.5	N/A	0.9620 [1.0000]	96.2% { 142.3% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 598531	(9.24 , N/A) (N/A , 0.01 , N/A)	866.3	N/A	0.8693 [1.0000]	86.9% { 115.6% }			
13C3_PFBs_EIS	(302.0 / 80.0) 4048621	(5.33 , N/A) (N/A , 0.00 , N/A)	2793.2	N/A	2.1077 [2.0000]	105.4% { 113.4% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 1926802	(7.00 , N/A) (N/A , 0.00 , N/A)	1149.4	N/A	1.9199 [2.0000]	96.0% { 110.0% }			
13C8_PFOS_EIS	(507.0 / 80.0) 3974757	(8.22 , N/A) (N/A , -0.01 , N/A)	1267.0	N/A	1.7869 [2.0000]	89.3% { 121.4% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 494220	(5.12 , N/A) (N/A , 0.00 , N/A)	1045.6	N/A	4.2006 [4.0000]	105.0% { 122.6% }			
13C2_6:2FTS_EIS	(429.0 / 81.0) 471899	(6.61 , N/A) (N/A , 0.01 , N/A)	1060.7	N/A	3.5575 [4.0000]	88.9% { 110.6% }			
13C2_8:2FTS_EIS	(529.0 / 81.0) 564908	(7.78 , N/A) (N/A , 0.00 , N/A)	919.1	N/A	4.0271 [4.0000]	100.7% { 108.1% }			
13C8_PFOsa_EIS	(506.0 / 78.0) 7057789	(9.83 , N/A) (N/A , 0.01 , N/A)	2150.6	N/A	1.7225 [2.0000]	86.1% { 121.9% }			
D3_NMeFOsa_EIS	(515.0 / 169.0) 1563351	(10.43 , N/A) (N/A , 0.01 , N/A)	2187.5	N/A	1.7768 [2.0000]	88.8% { 117.7% }			
D5_NEtFOsa_EIS	(531.0 / 169.0) 1509189	(10.58 , N/A) (N/A , 0.01 , N/A)	2643.0	N/A	1.9590 [2.0000]	97.9% { 124.6% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT Ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1286867	(8.19 , N/A) (N/A , 0.00 , N/A)	1222.3	N/A	4.1221 [4.0000]	103.1% { 115.9% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 1052288	(8.44 , N/A) (N/A , 0.00 , N/A)	15525.0	N/A	3.9584 [4.0000]	99.0% { 124.9% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 5507522	(10.36 , N/A) (N/A , 0.01 , N/A)	2155.9	N/A	18.3863 [20.0000]	91.9% { 128.8% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 7039175	(10.53 , N/A) (N/A , 0.01 , N/A)	1667.6	N/A	20.5437 [20.0000]	102.7% { 131.8% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 3048704	(5.67 , N/A) (N/A , 0.01 , N/A)	1694.5	N/A	7.4890 [8.0000]	93.6% { 109.6% }			

ANALYSIS SEQUENCE BLANKS

Laboratory: APPL, LLC
 Client: AECOM
 Sequence: SC00861
 Calibration: 2309009

SDG:
 Project: Red Hill AFFF Assessment Sam
 Instrument: Saphira

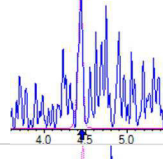
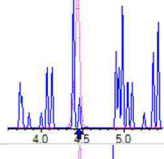
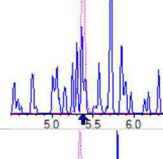
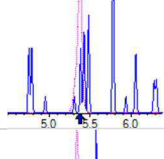
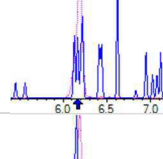
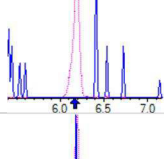
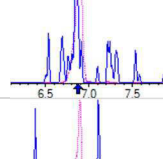
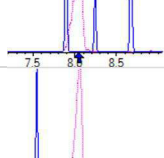
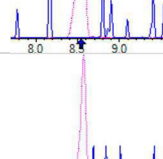
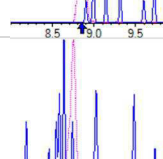
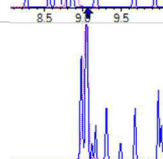
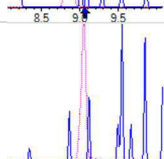
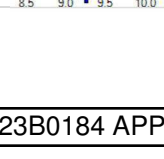
Lab Sample ID	Analyte	Found	Units	RL	C
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	PFPEA	0.00	ng/mL	0.20	U
	PFHXA	0.00	ng/mL	0.10	U
	PFHPA	0.00	ng/mL	0.10	U
	PFOA	0.00	ng/mL	0.10	U
	PFNA	0.00	ng/mL	0.10	U
	PFDA	0.00	ng/mL	0.10	U
	PFUnA	0.00	ng/mL	0.10	U
	PFDOA	0.00	ng/mL	0.10	U
	PFTRDA	0.00	ng/mL	0.10	U
	PFTEDA	0.00	ng/mL	0.10	U
	PFBS	0.00	ng/mL	0.10	U
	PFPEs	0.00	ng/mL	0.10	U
	PFHXS	0.00	ng/mL	0.10	U
	PFHPS	0.00	ng/mL	0.10	U
	PFOS	0.00	ng/mL	0.10	U
	PFNS	0.000617	ng/mL	0.10	U
	PFDS	0.00110	ng/mL	0.10	U
	PFDOS	0.00439	ng/mL	0.10	U
	4:2FTS	0.00377	ng/mL	0.40	U
	6:2FTS	0.00569	ng/mL	0.40	U
	8:2FTS	0.00914	ng/mL	0.40	U
	PFOSA	0.00401	ng/mL	0.10	U
	NMeFOSA	0.00502	ng/mL	0.40	U
	NEtFOSA	0.00532	ng/mL	0.40	U
	NMeFOSAA	0.00239	ng/mL	0.10	U
	NEtFOSAA	0.00	ng/mL	0.10	U
	NMeFOSE	0.0139	ng/mL	0.40	U
	NEtFOSE	0.0149	ng/mL	0.40	U
	HFPO-DA	0.00240	ng/mL	0.20	U
	ADONA	0.00274	ng/mL	0.20	U
	PFEESA	0.000907	ng/mL	0.20	U
	PFMPA	0.00387	ng/mL	0.20	U

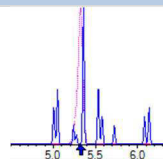
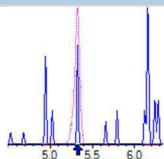
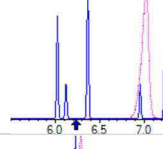
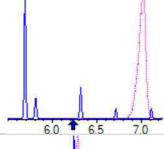
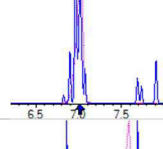
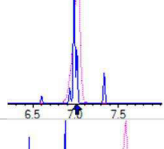
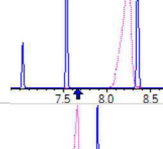
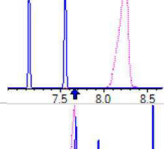
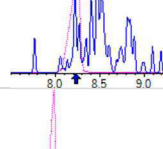
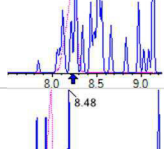
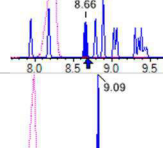
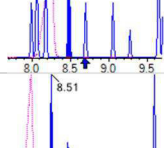
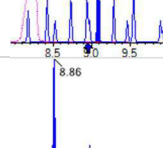
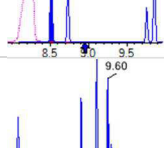
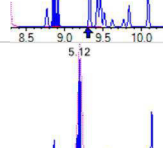
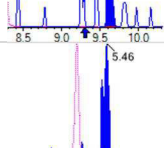
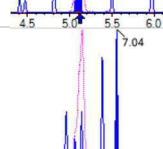
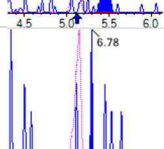
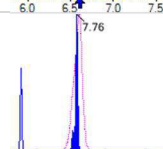
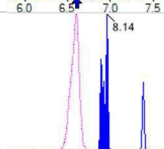
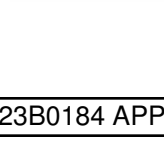
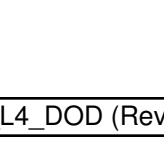
ANALYSIS SEQUENCE BLANKS

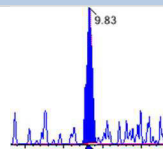
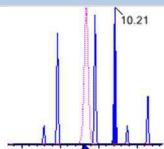
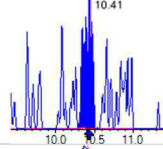
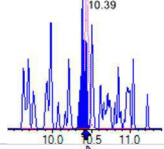
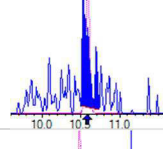
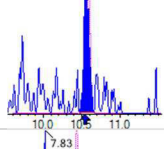
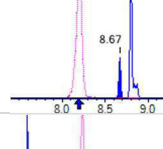
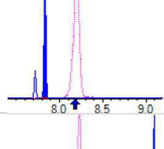
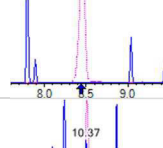
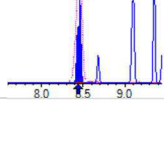
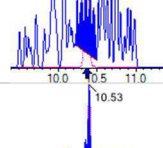
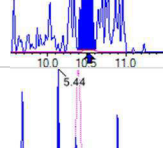
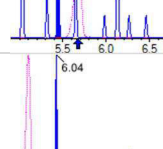
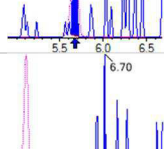
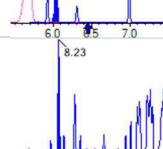
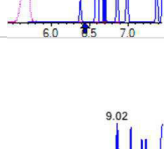
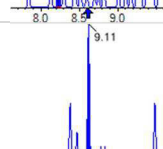
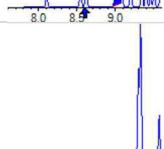
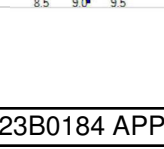
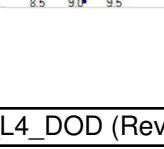
Laboratory: APPL, LLC
 Client: AECOM
 Sequence: SC00861
 Calibration: 2309009

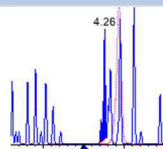
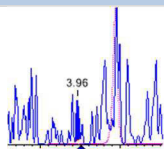
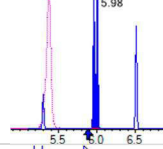
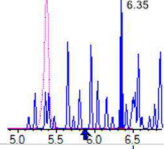
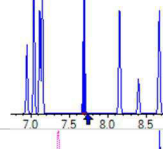
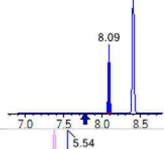
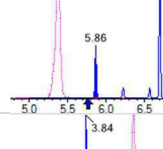
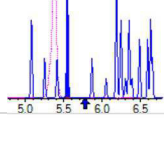
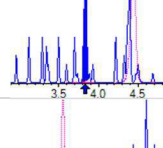
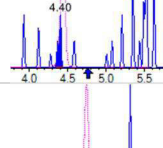
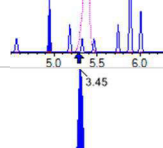
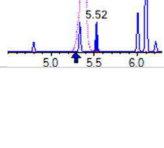
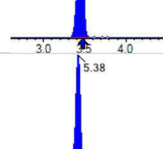
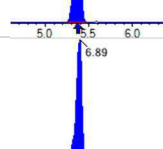
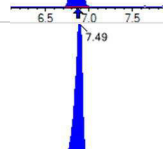
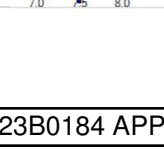
SDG:
 Project: Red Hill AFFF Assessment Sam
 Instrument: Saphira

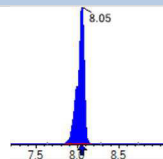
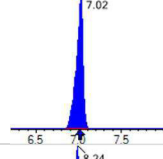
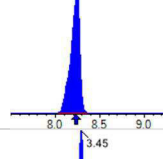
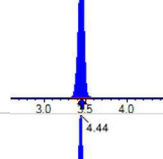
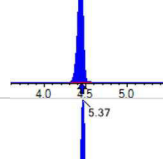
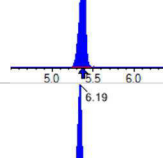
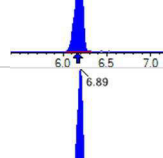
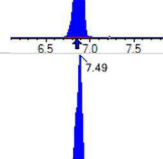
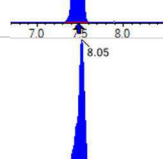
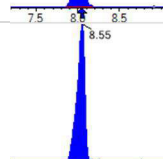
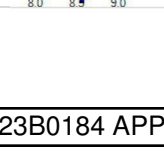
Lab Sample ID	Analyte	Found	Units	RL	C
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	NFDHA	0.00129	ng/mL	0.20	U
	9CL-PF3ONS	0.000700	ng/mL	0.20	U
	11CL-PF3OUDS	0.00154	ng/mL	0.20	U
	3:3FTCA	0.0211	ng/mL	0.40	U
	5:3FTCA	0.0127	ng/mL	0.40	U
	7:3FTCA	0.00491	ng/mL	0.40	U
	13C4-PFBA	7.09	ng/mL		
	13C5-PFPEA	3.77	ng/mL		
	13C5-PFHXA	1.89	ng/mL		
	13C4-PFHPA	1.85	ng/mL		
	13C8-PFOA	1.76	ng/mL		
	13C9-PFNA	0.952	ng/mL		
	13C6-PFDA	0.881	ng/mL		
	13C7-PFUnA	0.879	ng/mL		
	13C2-PFDOA	0.816	ng/mL		
	13C2-PFTEDA	0.868	ng/mL		
	13C3-PFBS	1.95	ng/mL		
	13C3-PFHXS	1.69	ng/mL		
	13C8-PFOS	1.68	ng/mL		
	13C2-4:2FTS	3.48	ng/mL		
	13C2-6:2FTS	3.12	ng/mL		
	13C2-8:2FTS	3.83	ng/mL		
	13C8-PFOSA	1.81	ng/mL		
	D3-NMEFOSA	1.89	ng/mL		
	D5-NETFOSA	2.02	ng/mL		
	D3-NMEFOSAA	4.00	ng/mL		
	D5-NETFOSAA	3.73	ng/mL		
	D7-NMEFOSE	19.7	ng/mL		
	D9-NETFOSSE	21.6	ng/mL		
	13C3-HFPO-DA	7.34	ng/mL		

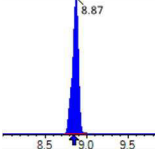
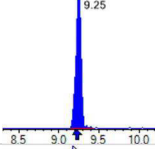
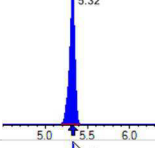
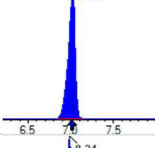
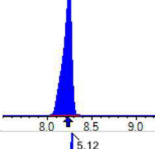
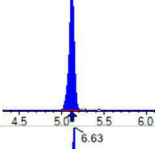
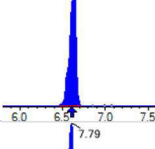
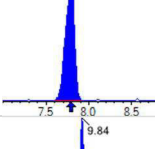
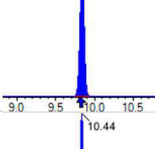
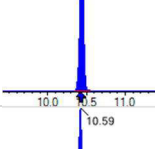
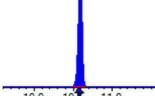
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
PFPeA	(263.0 / 219.0) N/A (263.0 / 69.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHxA	(313.0 / 269.0) N/A (313.0 / 119.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHpA	(363.0 / 319.0) N/A (363.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFOA	(413.0 / 369.0) N/A (413.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFNA	(463.0 / 419.0) N/A (463.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDA	(513.0 / 469.0) N/A (513.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFUnA	(563.0 / 519.0) N/A (563.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDaA	(613.0 / 569.0) N/A (613.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFTTrDA	(663.0 / 619.0) N/A (663.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFTeDA	(713.0 / 669.0) N/A (713.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

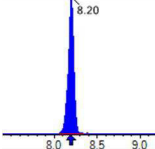
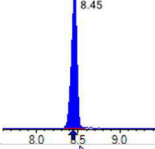
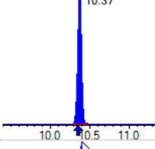
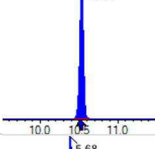
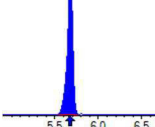
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) N/A (299.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFPeS	(349.0 / 80.0) N/A (349.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHxS	(399.0 / 80.0) N/A (399.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHpS	(449.0 / 80.0) N/A (449.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFOS	(499.0 / 80.0) N/A (499.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFNS	(549.0 / 80.0) 1244 (549.0 / 99.0) 1037	(8.66 , 1.05) (N/A , -0.03 , 10.6)	925787.9 98.8	0.8338 346.5 361.7	0.0006	N/A			IR2,
PFDS	(599.0 / 80.0) 2460 (599.0 / 99.0) 829	(9.09 , 1.10) (N/A , 0.14 , 34.2)	39.8 5095.4	0.3371 166.5 157.9	0.0011	N/A			IR2,
PFDoS	(699.0 / 80.0) 4541 (699.0 / 99.0) 2067	(8.86 , 1.08) (N/A , -0.44 , -44.3)	42.7 26.5	0.4553 233.0 218.1	0.0044	N/A			RT,IR2,
4:2FTS	(327.0 / 307.0) 1192 (327.0 / 81.0) 14012	(5.12 , 1.00) (0.00 , N/A , -20.7)	153.7 45.9	11.7590 1696.9 1854.5	0.0038	N/A			IR2,
6:2FTS	(427.0 / 407.0) 843 (427.0 / 81.0) 827	(7.04 , 1.06) (0.41 , N/A , 15.5)	3366.3 83.1	0.9805 115.7 128.0	0.0057	N/A			RT,
8:2FTS	(527.0 / 507.0) 1509 (527.0 / 81.0) 2552	(7.76 , 1.00) (-0.02 , N/A , -22.5)	642.0 107339.9	1.6912 207.3 212.0	0.0091	N/A			IR2,

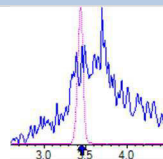
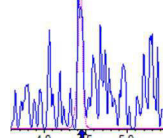
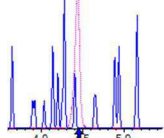
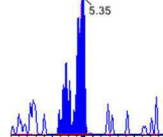
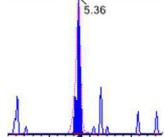
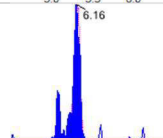
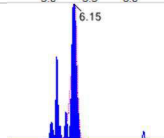
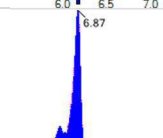
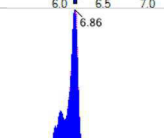
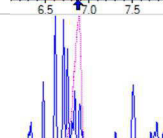
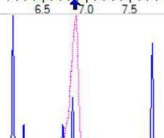
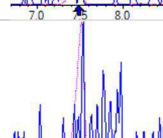
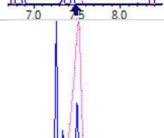
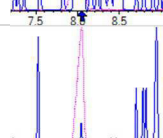
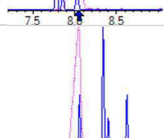
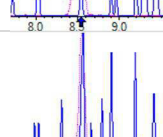
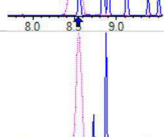
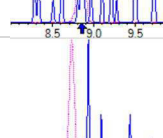
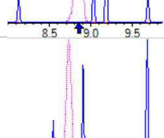
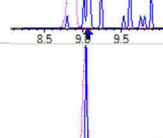
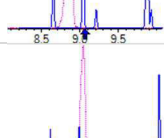
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) 11263 (498.0 / 478.0) 705	(9.83 , 1.00) (-0.01 , N/A , -22.7)	29.0 137.0	0.0626 318.5 306.9	0.0040	N/A			
NMeFOSA	(512.0 / 219.0) 3491 (512.0 / 169.0) 1852	(10.41 , 1.00) (-0.03 , N/A , 1.3)	17.4 22.3	0.5306 64.2 63.4	0.0050	N/A			
NEiFOSA	(526.0 / 219.0) 3929 (526.0 / 169.0) 4614	(10.54 , 1.00) (-0.05 , N/A , -1.0)	29.6 27.8	1.1744 92.7 91.7	0.0053	N/A			
NMeFOSAA	(570.0 / 419.0) 622 (570.0 / 483.0) 1914	(8.67 , 1.06) (0.47 , N/A , 50.2)	1394.3 8277.6	3.0760 627.3 694.4	0.0024	N/A			RT, IR2,
NEiFOSAA	(584.0 / 419.0) N/A (584.0 / 526.0) 1037	(N/A , N/A) (N/A , N/A , N/A)	N/A 80.7	N/A 0.0 0.0	0.0000	N/A			
NMeFOSE	(616.0 / 59.0) 4187	(10.37 , 1.00) (-0.01 , N/A , 0.0)	7.7	N/A 0.0 0.0	0.0139	N/A			
NEiFOSE	(630.0 / 59.0) 5443	(10.53 , 1.00) (-0.01 , N/A , 0.0)	14.3	N/A 0.0 0.0	0.0149	N/A			
HFPO-DA	(285.0 / 169.0) 759 (285.0 / 185.0) 2204	(5.44 , 0.96) (-0.23 , N/A , -13.8)	92.9 23.7	2.9034 105.6 96.0	0.0024	N/A			RT,
ADONA	(377.0 / 85.0) 3220 (377.0 / 251.0) 934	(6.04 , 1.06) (N/A , -0.41 , -39.5)	1924.3 72.7	0.2902 331.5 333.8	0.0027	N/A			RT, IR2,
9CI-Pf3ONS	(531.0 / 351.0) 1930 (533.0 / 353.0) 1214	(8.23 , 1.45) (N/A , -0.37 , -47.5)	10.0 13.7	0.6292 213.6 202.1	0.0007	N/A			IR2,
11CI-PF3OUDS	(631.0 / 451.0) 2080 (633.0 / 453.0) N/A	(9.11 , 1.60) (N/A , 0.01 , #Value!)	246.2 N/A	N/A 0.0 0.0	0.0015	N/A			IR1,

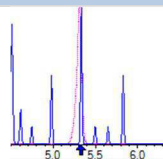
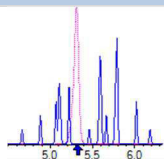
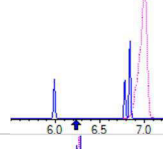
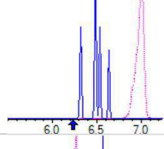
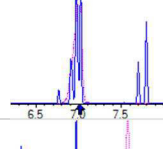
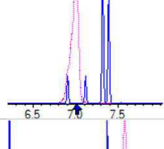
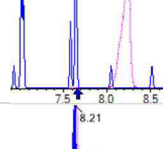
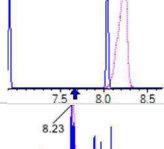
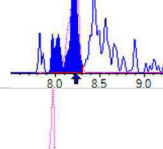
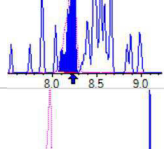
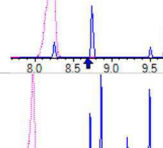
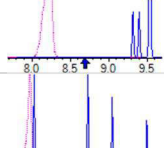
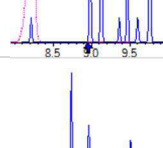
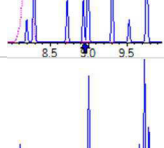
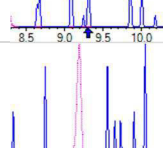
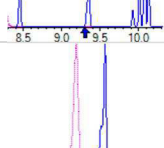
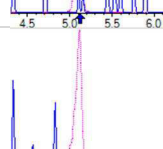
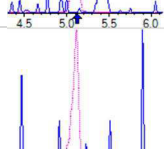
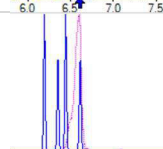
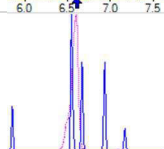
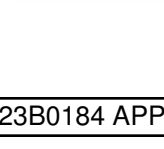
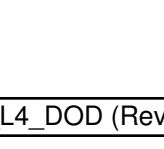
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) 284 (241.0 / 117.0) 313	(4.26, 0.96) (N/A, 0.26, 17.6)	16.4 4.1	1.1033 63.6 67.1	0.0211	N/A			
5:3FTCA	(341.0 / 236.7) 1075 (341.0 / 217.0) 1835	(5.98, 1.11) (N/A, 0.11, -21.8)	5666.3 14.2	1.7067 101.5 93.9	0.0127	N/A			
7:3FTCA	(441.0 / 317.0) 746 (441.0 / 337.0) 200	(7.69, 1.43) (N/A, -0.08, -23.7)	34.3 3001.3	0.2679 31.2 31.9	0.0049	N/A			IR1,
PFEESA	(315.0 / 135.0) 667 (315.0 / 83.0) 954	(5.86, 1.09) (N/A, 0.11, 19.2)	6136.6 18.4	1.4310 508.3 546.5	0.0009	N/A			IR2,
PFMPA	(229.0 / 85.0) 451	(3.84, 0.87) (N/A, 0.01, 0.0)	28.8	N/A 0.0 0.0	0.0039	N/A			
PFMBA	(279.0 / 85.0) 351	(4.40, 0.99) (N/A, -0.36, 0.0)	14.8	N/A 0.0 0.0	0.0009	N/A			
NFDHA	(295.0 / 201.0) 423 (295.0 / 85.0) 334	(4.94, 0.92) (N/A, -0.33, -34.8)	58.0 22.8	0.7894 75.6 76.1	0.0013	N/A			
13C3_PFBA_IIS	(216.0 / 172.0) 273955	(3.45, N/A) (N/A, -0.01, N/A)	1282.9	N/A	0.8081 [1.0000]	80.8% { 130.8% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 492894	(5.38, N/A) (N/A, 0.01, N/A)	1611.4	N/A	0.8287 [1.0000]	82.9% { 132.3% }			
13C4_PFOA_IIS	(417.0 / 372.0) 673502	(6.89, N/A) (N/A, 0.02, N/A)	910.4	N/A	0.8280 [1.0000]	82.8% { 130.0% }			
13C5_PFNA_IIS	(468.0 / 423.0) 613770	(7.49, N/A) (N/A, 0.01, N/A)	531.4	N/A	0.8038 [1.0000]	80.4% { 111.4% }			

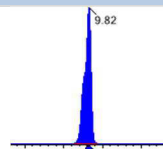
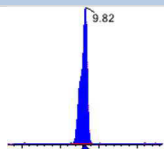
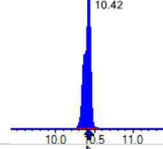
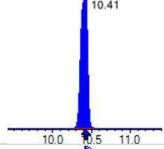
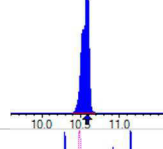
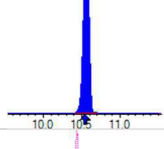
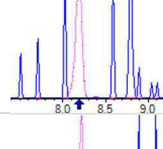
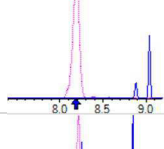
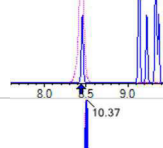
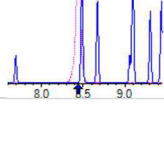
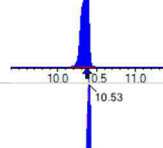
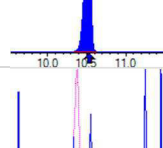
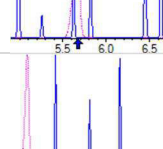
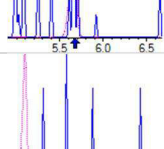
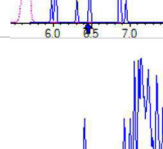
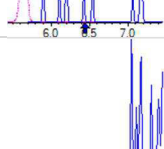
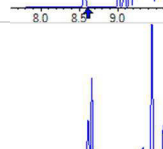
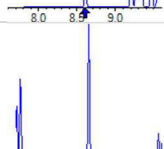
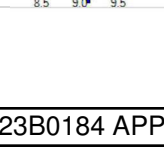
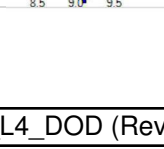
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 659574	(8.05 , N/A) (N/A , 0.02 , N/A)	796.1	N/A	0.9188 [1.0000]	91.9% { 137.8% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 1041601	(7.02 , N/A) (N/A , 0.02 , N/A)	1401.1	N/A	0.8458 [1.0000]	84.6% { 117.7% }			
13C4_PFOS_IIS	(503.0 / 79.9) 1650591	(8.24 , N/A) (N/A , 0.01 , N/A)	1308.0	N/A	0.9300 [1.0000]	93.0% { 128.0% }			
13C4_PFBA_EIS	(217.0 / 172.0) 2308899	(3.45 , N/A) (N/A , -0.01 , N/A)	3317.6	N/A	7.0878 [8.0000]	88.6% { 121.0% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 2070748	(4.44 , N/A) (N/A , -0.01 , N/A)	2121.2	N/A	3.7704 [4.0000]	94.3% { 118.1% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 1266820	(5.37 , N/A) (N/A , 0.01 , N/A)	1857.4	N/A	1.8916 [2.0000]	94.6% { 119.3% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 1155411	(6.19 , N/A) (N/A , 0.03 , N/A)	904.0	N/A	1.8498 [2.0000]	92.5% { 128.1% }			
13C8_PFOA_EIS	(421.0 / 376.0) 1269012	(6.89 , N/A) (N/A , 0.03 , N/A)	922.5	N/A	1.7604 [2.0000]	88.0% { 119.5% }			
13C9_PFNA_EIS	(472.0 / 427.0) 581982	(7.49 , N/A) (N/A , 0.01 , N/A)	793.1	N/A	0.9515 [1.0000]	95.2% { 116.2% }			
13C6_PFDA_EIS	(519.0 / 474.0) 672453	(8.05 , N/A) (N/A , 0.01 , N/A)	954.2	N/A	0.8809 [1.0000]	88.1% { 107.5% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 664815	(8.55 , N/A) (N/A , 0.02 , N/A)	877.9	N/A	0.8790 [1.0000]	87.9% { 112.9% }			

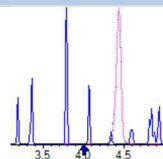
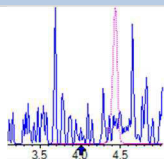
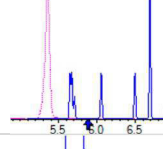
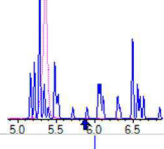
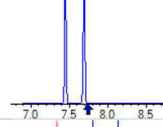
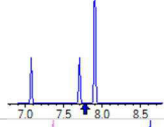
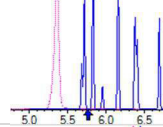
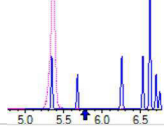
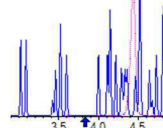
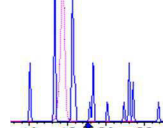
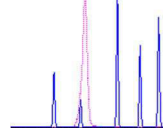
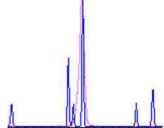
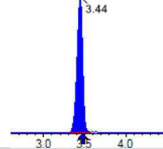
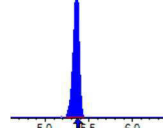
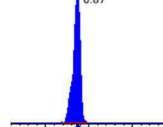
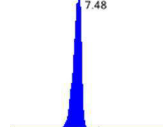
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 572606	(8.87, N/A) (N/A, 0.02, N/A)	4491.7	N/A	0.8163 [1.0000]	81.6% { 127.9% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 633220	(9.25, N/A) (N/A, 0.02, N/A)	826.6	N/A	0.8684 [1.0000]	86.8% { 122.3% }			
13C3_PFBs_EIS	(302.0 / 80.0) 4098406	(5.32, N/A) (N/A, 0.00, N/A)	2075.0	N/A	1.9471 [2.0000]	97.4% { 114.8% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 1856685	(7.02, N/A) (N/A, 0.02, N/A)	1618.0	N/A	1.6884 [2.0000]	84.4% { 106.0% }			
13C8_PFOS_EIS	(507.0 / 80.0) 3653316	(8.24, N/A) (N/A, 0.01, N/A)	1310.6	N/A	1.6847 [2.0000]	84.2% { 111.6% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 448004	(5.12, N/A) (N/A, 0.00, N/A)	894.5	N/A	3.4750 [4.0000]	86.9% { 111.1% }			
13C2_6:2FTS_EIS	(429.0 / 81.0) 453928	(6.63, N/A) (N/A, 0.02, N/A)	952.0	N/A	3.1229 [4.0000]	78.1% { 106.4% }			
13C2_8:2FTS_EIS	(529.0 / 81.0) 587937	(7.79, N/A) (N/A, 0.01, N/A)	992.6	N/A	3.8250 [4.0000]	95.6% { 112.5% }			
13C8_PFOsa_EIS	(506.0 / 78.0) 7229775	(9.84, N/A) (N/A, 0.02, N/A)	2462.4	N/A	1.8100 [2.0000]	90.5% { 124.8% }			
D3_NMeFOsa_EIS	(515.0 / 169.0) 1620569	(10.44, N/A) (N/A, 0.02, N/A)	2639.3	N/A	1.8894 [2.0000]	94.5% { 122.0% }			
D5_NEtFOsa_EIS	(531.0 / 169.0) 1519563	(10.59, N/A) (N/A, 0.02, N/A)	2390.8	N/A	2.0233 [2.0000]	101.2% { 125.4% }			

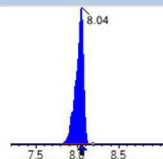
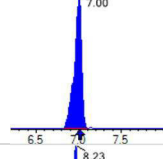
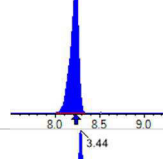
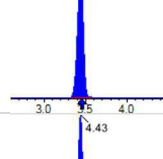
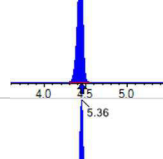
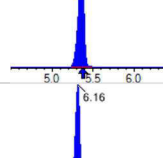
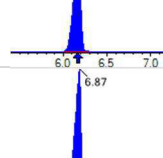
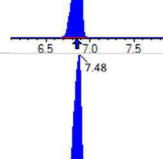
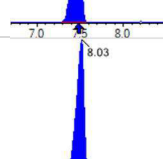
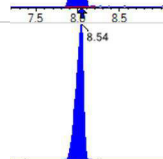
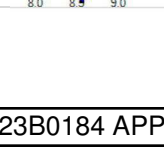
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1217850	(8.20 , N/A) (N/A , 0.01 , N/A)	1597.3	N/A	4.0017 [4.0000]	100.0% { 109.7% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 967838	(8.45 , N/A) (N/A , 0.02 , N/A)	3390.9	N/A	3.7346 [4.0000]	93.4% { 114.9% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 5749889	(10.37 , N/A) (N/A , 0.02 , N/A)	2214.5	N/A	19.6905 [20.0000]	98.5% { 134.4% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 7208966	(10.53 , N/A) (N/A , 0.02 , N/A)	1407.6	N/A	21.5819 [20.0000]	107.9% { 135.0% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 3089588	(5.68 , N/A) (N/A , 0.01 , N/A)	2067.3	N/A	7.3418 [8.0000]	91.8% { 111.1% }			

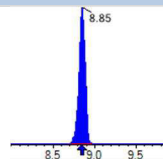
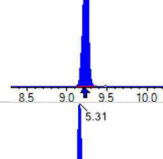
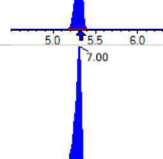
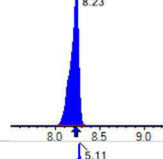
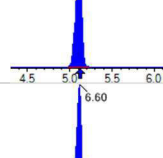
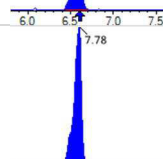
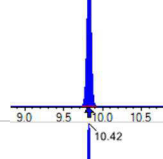
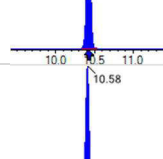
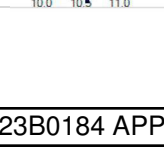
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
PFPeA	(263.0 / 219.0) N/A (263.0 / 69.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHxA	(313.0 / 269.0) 20914 (313.0 / 119.0) 3390	(5.35, 1.00) (0.00, N/A, -0.3)	1534.0 157445.5	0.1621 173.2 156.9	0.0374	N/A			IR2,
PFHpA	(363.0 / 319.0) 46446 (363.0 / 169.0) 16480	(6.16, 1.00) (-0.01, N/A, 0.6)	79552.3 193667.5	0.3548 112.4 108.8	0.0948	N/A			
PFOA	(413.0 / 369.0) 5366062 (413.0 / 169.0) 2000926	(6.87, 1.00) (0.00, N/A, 0.2)	2457.6 1886.2	0.3729 110.2 104.9	9.9885	N/A			
PFNA	(463.0 / 419.0) N/A (463.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDA	(513.0 / 469.0) N/A (513.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFUnA	(563.0 / 519.0) N/A (563.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDoA	(613.0 / 569.0) N/A (613.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFTTrDA	(663.0 / 619.0) N/A (663.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFTeDA	(713.0 / 669.0) N/A (713.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

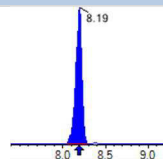
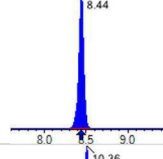
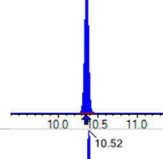
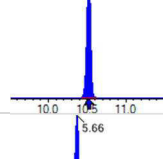
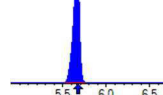
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) N/A (299.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFPeS	(349.0 / 80.0) N/A (349.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHxS	(399.0 / 80.0) N/A (399.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHpS	(449.0 / 80.0) N/A (449.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFOS	(499.0 / 80.0) 46786 (499.0 / 99.0) 8486	(8.21 , 1.00) (-0.02 , N/A , -0.9)	25.0 47.3	0.1814 82.5 78.7	0.0221	N/A			M12 ABK 3/2/23
PFNS	(549.0 / 80.0) N/A (549.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDS	(599.0 / 80.0) N/A (599.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDoS	(699.0 / 80.0) N/A (699.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
4:2FTS	(327.0 / 307.0) N/A (327.0 / 81.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
6:2FTS	(427.0 / 407.0) N/A (427.0 / 81.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
8:2FTS	(527.0 / 507.0) N/A (527.0 / 81.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

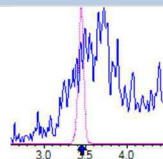
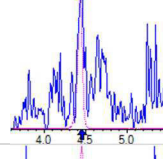
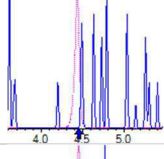
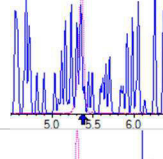
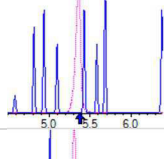
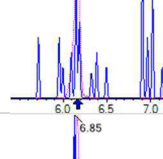
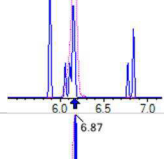
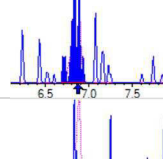
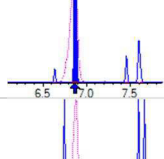
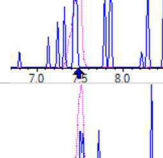
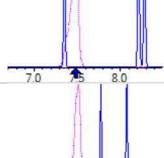
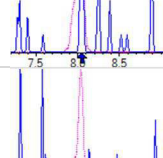
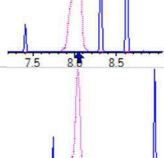
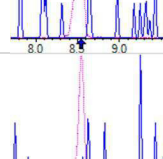
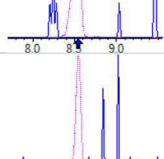
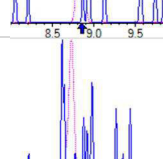
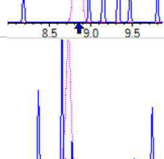
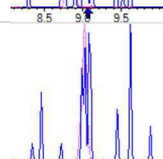
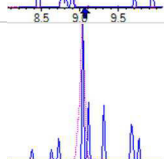
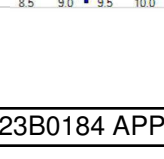
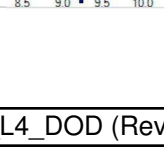
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) 29095021 (498.0 / 478.0) 632261	(9.82 , 1.00) (-0.01 , N/A , -0.1)	1871.3 807.9	0.0217 110.6 106.6	11.5260	N/A			
NMeFOSA	(512.0 / 219.0) 6324076 (512.0 / 169.0) 5392083	(10.42 , 1.00) (0.00 , N/A , 0.9)	2965.5 4025.4	0.8526 103.2 101.9	10.2094	N/A			
NEIFOSA	(526.0 / 219.0) 6923536 (526.0 / 169.0) 8947469	(10.57 , 1.00) (-0.01 , N/A , 0.7)	6904.9 5515.4	1.2923 102.1 100.9	10.4126	N/A			
NMeFOSAA	(570.0 / 419.0) N/A (570.0 / 483.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NEIFOSAA	(584.0 / 419.0) N/A (584.0 / 526.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NMeFOSE	(616.0 / 59.0) 2956878	(10.37 , 1.00) (0.01 , N/A , 0.0)	1828.6	N/A 0.0 0.0	10.8288	N/A			
NEtFOSE	(630.0 / 59.0) 3574961	(10.53 , 1.00) (0.01 , N/A , 0.0)	1253.6	N/A 0.0 0.0	11.4811	N/A			
HFPO-DA	(285.0 / 169.0) N/A (285.0 / 185.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
ADONA	(377.0 / 85.0) N/A (377.0 / 251.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
9CI-Pf3ONS	(531.0 / 351.0) N/A (533.0 / 353.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
11CI-PF3OUDS	(631.0 / 451.0) N/A (633.0 / 453.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

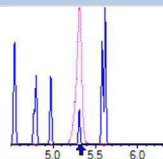
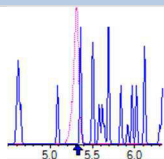
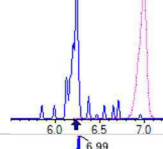
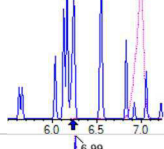
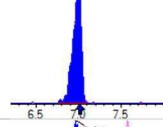
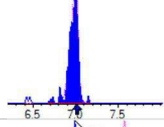
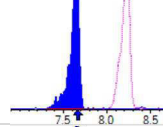
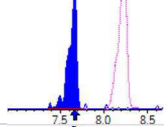
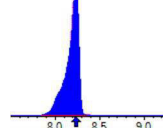
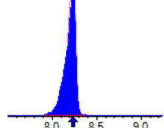
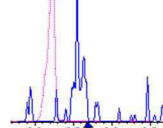
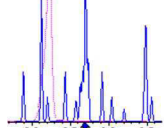
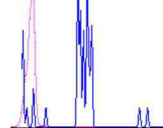
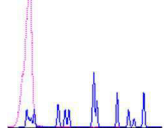
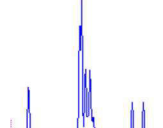
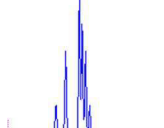
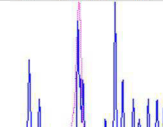
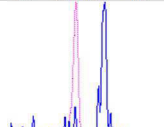
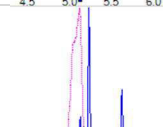
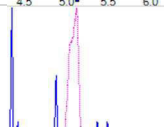
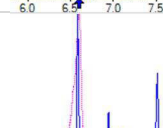
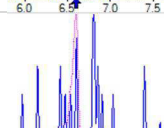
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) N/A (241.0 / 117.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
5:3FTCA	(341.0 / 236.7) N/A (341.0 / 217.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
7:3FTCA	(441.0 / 317.0) N/A (441.0 / 337.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFEESA	(315.0 / 135.0) N/A (315.0 / 83.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFMPA	(229.0 / 85.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
PFMBA	(279.0 / 85.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
NFDHA	(295.0 / 201.0) N/A (295.0 / 85.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
13C3_PFBA_IIS	(216.0 / 172.0) 263189	(3.44, N/A) (N/A, -0.02, N/A)	1161.8	N/A	0.7763 [1.0000]	77.6% { 125.7% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 486148	(5.36, N/A) (N/A, -0.01, N/A)	1346.3	N/A	0.8174 [1.0000]	81.7% { 130.5% }			
13C4_PFOA_IIS	(417.0 / 372.0) 626127	(6.87, N/A) (N/A, 0.00, N/A)	1098.7	N/A	0.7697 [1.0000]	77.0% { 120.8% }			
13C5_PFNA_IIS	(468.0 / 423.0) 603593	(7.48, N/A) (N/A, 0.00, N/A)	762.4	N/A	0.7904 [1.0000]	79.0% { 109.6% }			

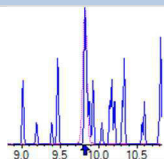
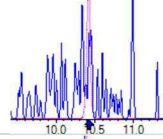
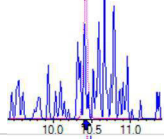
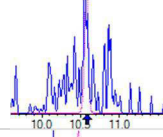
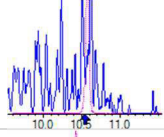
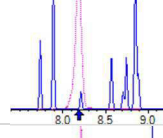
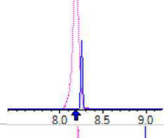
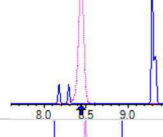
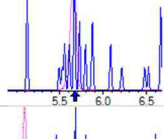
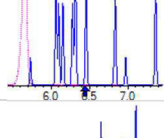
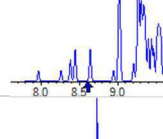
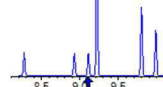
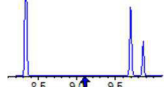
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 630670	(8.04 , N/A) (N/A , 0.01 , N/A)	856.0	N/A	0.8785 [1.0000]	87.9% { 131.8% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 948459	(7.00 , N/A) (N/A , 0.00 , N/A)	785.1	N/A	0.7701 [1.0000]	77.0% { 107.2% }			
13C4_PFOS_IIS	(503.0 / 79.9) 1559281	(8.23 , N/A) (N/A , 0.00 , N/A)	1014.0	N/A	0.8785 [1.0000]	87.9% { 120.9% }			
13C4_PFBA_EIS	(217.0 / 172.0) 2173559	(3.44 , N/A) (N/A , -0.02 , N/A)	3128.9	N/A	6.9453 [8.0000]	86.8% { 113.9% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 1915580	(4.43 , N/A) (N/A , -0.01 , N/A)	1869.9	N/A	3.5363 [4.0000]	88.4% { 109.3% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 1199495	(5.36 , N/A) (N/A , -0.01 , N/A)	1374.2	N/A	1.8159 [2.0000]	90.8% { 113.0% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 1097070	(6.16 , N/A) (N/A , 0.00 , N/A)	1467.4	N/A	1.7808 [2.0000]	89.0% { 121.7% }			
13C8_PFOA_EIS	(421.0 / 376.0) 1115098	(6.87 , N/A) (N/A , 0.00 , N/A)	1263.3	N/A	1.6639 [2.0000]	83.2% { 105.0% }			
13C9_PFNA_EIS	(472.0 / 427.0) 595094	(7.48 , N/A) (N/A , -0.01 , N/A)	950.3	N/A	0.9894 [1.0000]	98.9% { 118.8% }			
13C6_PFDA_EIS	(519.0 / 474.0) 680474	(8.03 , N/A) (N/A , 0.00 , N/A)	593.0	N/A	0.9322 [1.0000]	93.2% { 108.8% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 710403	(8.54 , N/A) (N/A , 0.00 , N/A)	876.2	N/A	0.9823 [1.0000]	98.2% { 120.6% }			

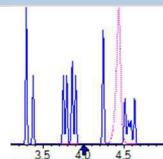
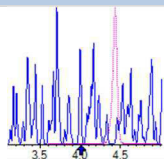
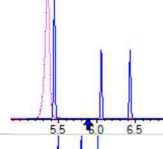
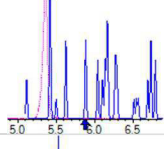
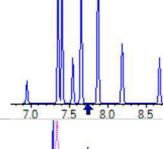
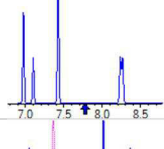
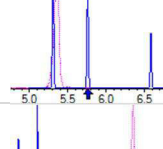
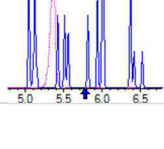
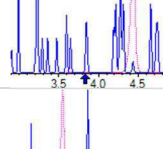
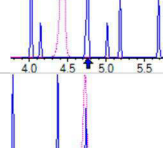
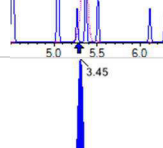
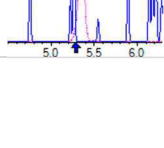
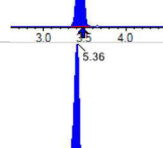
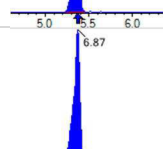
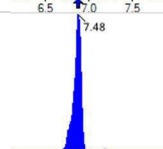
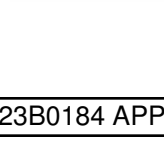
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 565637	(8.85, N/A) (N/A, 0.01, N/A)	26578.7	N/A	0.8434 [1.0000]	84.3% { 126.3% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 547140	(9.24, N/A) (N/A, 0.01, N/A)	1673.6	N/A	0.7848 [1.0000]	78.5% { 105.7% }			
13C3_PFBs_EIS	(302.0 / 80.0) 4091234	(5.31, N/A) (N/A, -0.01, N/A)	2987.0	N/A	2.1346 [2.0000]	106.7% { 114.6% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 1906909	(7.00, N/A) (N/A, 0.00, N/A)	1063.9	N/A	1.9044 [2.0000]	95.2% { 108.9% }			
13C8_PFOS_EIS	(507.0 / 80.0) 3830150	(8.23, N/A) (N/A, 0.01, N/A)	1266.9	N/A	1.8697 [2.0000]	93.5% { 117.0% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 443131	(5.11, N/A) (N/A, -0.01, N/A)	1286.3	N/A	3.7748 [4.0000]	94.4% { 109.9% }			
13C2_6:2FTS_EIS	(429.0 / 81.0) 429055	(6.60, N/A) (N/A, 0.00, N/A)	932.7	N/A	3.2417 [4.0000]	81.0% { 100.5% }			
13C2_8:2FTS_EIS	(529.0 / 81.0) 574387	(7.78, N/A) (N/A, 0.00, N/A)	1223.6	N/A	4.1038 [4.0000]	102.6% { 109.9% }			
13C8_PFOsa_EIS	(506.0 / 78.0) 6496638	(9.83, N/A) (N/A, 0.01, N/A)	3022.4	N/A	1.7217 [2.0000]	86.1% { 112.2% }			
D3_NMeFOSA_EIS	(515.0 / 169.0) 1442511	(10.42, N/A) (N/A, 0.01, N/A)	1625.6	N/A	1.7803 [2.0000]	89.0% { 108.6% }			
D5_NEtFOSA_EIS	(531.0 / 169.0) 1368894	(10.58, N/A) (N/A, 0.01, N/A)	2903.4	N/A	1.9294 [2.0000]	96.5% { 113.0% }			

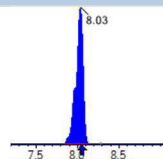
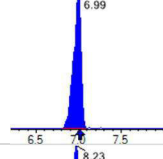
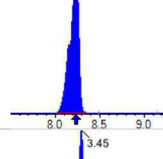
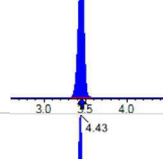
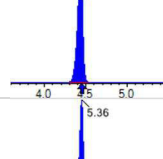
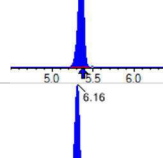
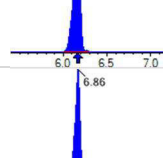
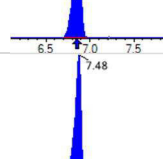
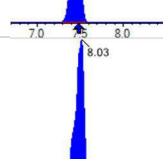
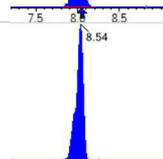
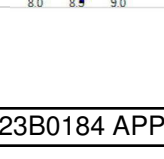
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1161076	(8.19 , N/A) (N/A , 0.00 , N/A)	1324.6	N/A	4.0385 [4.0000]	101.0% { 104.6% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 976183	(8.44 , N/A) (N/A , 0.00 , N/A)	10709.4	N/A	3.9874 [4.0000]	99.7% { 115.9% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 5211932	(10.36 , N/A) (N/A , 0.01 , N/A)	2135.5	N/A	18.8935 [20.0000]	94.5% { 121.9% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 6163085	(10.52 , N/A) (N/A , 0.01 , N/A)	1572.6	N/A	19.5313 [20.0000]	97.7% { 115.4% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 3030560	(5.66 , N/A) (N/A , -0.01 , N/A)	2385.7	N/A	7.3014 [8.0000]	91.3% { 109.0% }			

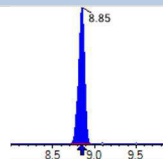
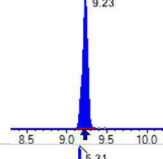
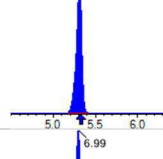
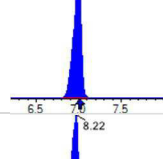
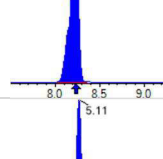
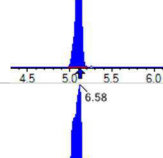
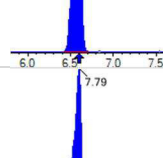
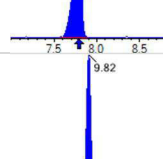
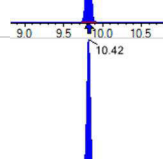
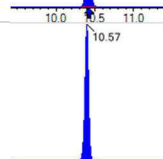
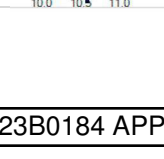
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
PFPeA	(263.0 / 219.0) N/A (263.0 / 69.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHxA	(313.0 / 269.0) N/A (313.0 / 119.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHpA	(363.0 / 319.0) N/A (363.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFOA	(413.0 / 369.0) 8431 (413.0 / 169.0) 2814	(6.85, 1.00) (-0.01, N/A, -1.6)	227.3 3001202.7	0.3338 98.7 93.9	0.0127	N/A			
PFNA	(463.0 / 419.0) N/A (463.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDA	(513.0 / 469.0) N/A (513.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFUnA	(563.0 / 519.0) N/A (563.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDaA	(613.0 / 569.0) N/A (613.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFTTrDA	(663.0 / 619.0) N/A (663.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFTeDA	(713.0 / 669.0) N/A (713.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

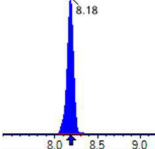
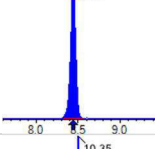
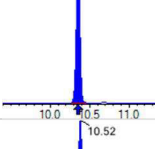
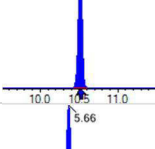
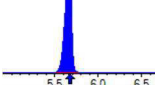
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) N/A (299.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFPeS	(349.0 / 80.0) N/A (349.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHxS	(399.0 / 80.0) 214395 (399.0 / 99.0) 77892	(6.99 , 1.00) (0.00 , N/A , 0.2)	1079.9 10619.5	0.3633 113.5 105.2	0.1334	N/A			
PFHpS	(449.0 / 80.0) 426335 (449.0 / 99.0) 130495	(7.64 , 0.93) (N/A , -0.01 , -1.1)	21642.7 421.0	0.3061 112.7 109.4	0.2294	N/A			
PFOS	(499.0 / 80.0) 20019087 (499.0 / 99.0) 4705539	(8.23 , 1.00) (0.00 , N/A , -0.1)	623.6 739.0	0.2351 106.9 101.9	8.7895	N/A			
PFNS	(549.0 / 80.0) N/A (549.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDS	(599.0 / 80.0) N/A (599.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDoS	(699.0 / 80.0) N/A (699.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
4:2FTS	(327.0 / 307.0) N/A (327.0 / 81.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
6:2FTS	(427.0 / 407.0) N/A (427.0 / 81.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
8:2FTS	(527.0 / 507.0) N/A (527.0 / 81.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) N/A (498.0 / 478.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NMeFOSA	(512.0 / 219.0) N/A (512.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NEtFOSA	(526.0 / 219.0) N/A (526.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NMeFOSAA	(570.0 / 419.0) N/A (570.0 / 483.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NEtFOSAA	(584.0 / 419.0) N/A (584.0 / 526.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NMeFOSE	(616.0 / 59.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
NEtFOSE	(630.0 / 59.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
HFPO-DA	(285.0 / 169.0) N/A (285.0 / 185.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
ADONA	(377.0 / 85.0) N/A (377.0 / 251.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
9CI-Pf3ONS	(531.0 / 351.0) N/A (533.0 / 353.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
11CI-PF3OUDS	(631.0 / 451.0) N/A (633.0 / 453.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) N/A (241.0 / 117.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
5:3FTCA	(341.0 / 236.7) N/A (341.0 / 217.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
7:3FTCA	(441.0 / 317.0) N/A (441.0 / 337.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFEESA	(315.0 / 135.0) N/A (315.0 / 83.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFMPA	(229.0 / 85.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
PFMBA	(279.0 / 85.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
NFDHA	(295.0 / 201.0) N/A (295.0 / 85.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
13C3_PFBA_IIS	(216.0 / 172.0) 295055	(3.45, N/A) (N/A, -0.01, N/A)	1465.6	N/A	0.8703 [1.0000]	87.0% { 140.9% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 564167	(5.36, N/A) (N/A, -0.01, N/A)	1551.3	N/A	0.9486 [1.0000]	94.9% { 151.4% }			
13C4_PFOA_IIS	(417.0 / 372.0) 704732	(6.87, N/A) (N/A, 0.00, N/A)	804.7	N/A	0.8664 [1.0000]	86.6% { 136.0% }			
13C5_PFNA_IIS	(468.0 / 423.0) 666345	(7.48, N/A) (N/A, 0.00, N/A)	1108.7	N/A	0.8726 [1.0000]	87.3% { 121.0% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 701833	(8.03 , N/A) (N/A , 0.00 , N/A)	848.9	N/A	0.9776 [1.0000]	97.8% { 146.6% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 1009154	(6.99 , N/A) (N/A , -0.01 , N/A)	1089.4	N/A	0.8194 [1.0000]	81.9% { 114.0% }			
13C4_PFOS_IIS	(503.0 / 79.9) 1726780	(8.23 , N/A) (N/A , 0.00 , N/A)	812.2	N/A	0.9729 [1.0000]	97.3% { 133.9% }			
13C4_PFBA_EIS	(217.0 / 172.0) 2566806	(3.45 , N/A) (N/A , -0.01 , N/A)	2894.7	N/A	7.3160 [8.0000]	91.5% { 134.5% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 2304373	(4.43 , N/A) (N/A , -0.02 , N/A)	2252.3	N/A	3.6658 [4.0000]	91.6% { 131.5% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 1454945	(5.36 , N/A) (N/A , -0.01 , N/A)	1858.2	N/A	1.8980 [2.0000]	94.9% { 137.0% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 1212237	(6.16 , N/A) (N/A , 0.00 , N/A)	1984.8	N/A	1.6956 [2.0000]	84.8% { 134.4% }			
13C8_PFOA_EIS	(421.0 / 376.0) 1381070	(6.86 , N/A) (N/A , 0.00 , N/A)	1762.5	N/A	1.8309 [2.0000]	91.5% { 130.1% }			
13C9_PFNA_EIS	(472.0 / 427.0) 685873	(7.48 , N/A) (N/A , 0.00 , N/A)	1436.8	N/A	1.0329 [1.0000]	103.3% { 136.9% }			
13C6_PFDA_EIS	(519.0 / 474.0) 786423	(8.03 , N/A) (N/A , 0.00 , N/A)	996.8	N/A	0.9681 [1.0000]	96.8% { 125.8% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 699586	(8.54 , N/A) (N/A , 0.00 , N/A)	744.1	N/A	0.8693 [1.0000]	86.9% { 118.8% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 677538	(8.85 , N/A) (N/A , 0.00 , N/A)	1193.6	N/A	0.9078 [1.0000]	90.8% { 151.3% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 695264	(9.23 , N/A) (N/A , 0.00 , N/A)	1034.0	N/A	0.8961 [1.0000]	89.6% { 134.3% }			
13C3_PFBs_EIS	(302.0 / 80.0) 4701475	(5.31 , N/A) (N/A , -0.01 , N/A)	2715.5	N/A	2.3055 [2.0000]	115.3% { 131.7% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 2040400	(6.99 , N/A) (N/A , -0.01 , N/A)	1667.4	N/A	1.9151 [2.0000]	95.8% { 116.5% }			
13C8_PFOS_EIS	(507.0 / 80.0) 4127601	(8.22 , N/A) (N/A , 0.00 , N/A)	1285.8	N/A	1.8195 [2.0000]	91.0% { 126.1% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 567942	(5.11 , N/A) (N/A , -0.02 , N/A)	1384.5	N/A	4.5470 [4.0000]	113.7% { 140.9% }			
13C2_6:2FTS_EIS	(429.0 / 81.0) 656162	(6.58 , N/A) (N/A , -0.02 , N/A)	962.4	N/A	4.6594 [4.0000]	116.5% { 153.8% }			
13C2_8:2FTS_EIS	(529.0 / 81.0) 713212	(7.79 , N/A) (N/A , 0.01 , N/A)	942.1	N/A	4.7892 [4.0000]	119.7% { 136.4% }			
13C8_PFOsa_EIS	(506.0 / 78.0) 7798317	(9.82 , N/A) (N/A , 0.00 , N/A)	2781.8	N/A	1.8661 [2.0000]	93.3% { 134.7% }			
D3_NMeFOsa_EIS	(515.0 / 169.0) 1652666	(10.42 , N/A) (N/A , 0.00 , N/A)	2297.5	N/A	1.8418 [2.0000]	92.1% { 124.4% }			
D5_NEtFOsa_EIS	(531.0 / 169.0) 1534805	(10.57 , N/A) (N/A , 0.00 , N/A)	3252.4	N/A	1.9534 [2.0000]	97.7% { 126.7% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT Ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1257828	(8.18 , N/A) (N/A , 0.00 , N/A)	1016.1	N/A	3.9507 [4.0000]	98.8% { 113.3% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 1170891	(8.44 , N/A) (N/A , 0.00 , N/A)	5546.7	N/A	4.3188 [4.0000]	108.0% { 139.0% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 5857253	(10.35 , N/A) (N/A , 0.00 , N/A)	1520.2	N/A	19.1732 [20.0000]	95.9% { 137.0% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 7186480	(10.52 , N/A) (N/A , 0.00 , N/A)	1322.4	N/A	20.5653 [20.0000]	102.8% { 134.6% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 3645600	(5.66 , N/A) (N/A , -0.01 , N/A)	2204.8	N/A	7.5686 [8.0000]	94.6% { 131.1% }			

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QUALITY CONTROL RAW DATA

ANALYSIS DATA SHEET

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Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Matrix:	Water	Laboratory ID:	BCB0459-BLK1
		File ID:	S2023-03-02A (7)
Sampled:		Prepared:	02/28/23 08:24
		Analyzed:	03/02/23 10:59
Solids:		Preparation:	EPA 1633
		Dilution:	1
Batch:	BCB0459	Sequence:	SC00861
		Calibration:	2309009
Column:	1	Instrument:	Saphira

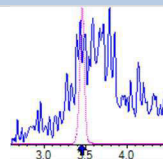
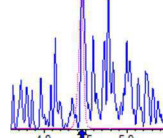
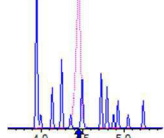
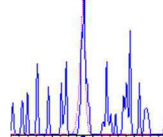
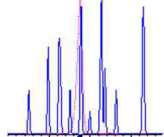
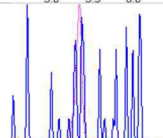
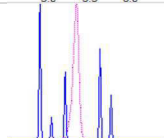
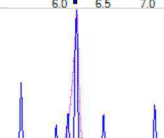
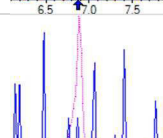
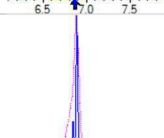
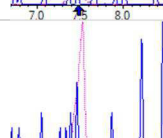
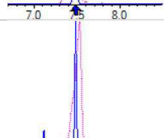
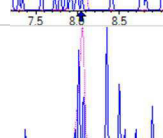
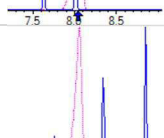
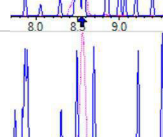
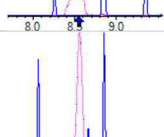
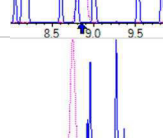
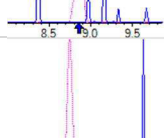
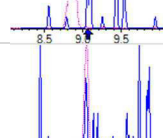
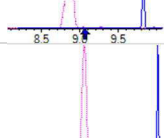
COMPOUND	CONC. (ng/L)	LOQ	LOD	DL	Q
PFBA	0.80 U	1.6	0.80	0.21	U
PFPEA	0.40 U	0.80	0.40	0.065	U
PFHXA	0.20 U	0.40	0.20	0.055	U
PFHPA	0.20 U	0.40	0.20	0.041	U
PFOA	0.20 U	0.40	0.20	0.15	U
PFNA	0.20 U	0.40	0.20	0.082	U
PFDA	0.20 U	0.40	0.20	0.10	U
PFUnA	0.20 U	0.40	0.20	0.16	U
PFDOA	0.20 U	0.40	0.20	0.11	U
PFTRDA	0.30 U	0.40	0.30	0.20	U
PFTEDA	0.20 U	0.40	0.20	0.20	U
PFBS	0.20 U	0.40	0.20	0.037	U
PFPEs	0.20 U	0.40	0.20	0.063	U
PFHXS	0.20 U	0.40	0.20	0.032	U
PFHPS	0.20 U	0.40	0.20	0.051	U
PFOS	0.102 J	0.40	0.20	0.064	MI2, J
PFNS	0.20 U	0.40	0.20	0.12	U
PFDS	0.20 U	0.40	0.20	0.15	U
PFDOS	0.20 U	0.40	0.20	0.12	U
4:2FTS	0.80 U	1.6	0.80	0.29	U
6:2FTS	0.80 U	1.6	0.80	0.31	U
8:2FTS	0.80 U	1.6	0.80	0.082	U
PFOSA	0.20 U	0.40	0.20	0.10	U
NMeFOSA	0.80 U	1.6	0.80	0.47	U
NEtFOSA	0.80 U	1.6	0.80	0.41	U
NMeFOSAA	0.20 U	0.40	0.20	0.11	U
NEtFOSAA	0.20 U	0.40	0.20	0.11	U
NMeFOSE	1.2 U	1.6	1.2	1.0	U
NEtFOSE	1.2 U	1.6	1.2	1.0	U
HFPO-DA	0.40 U	0.80	0.40	0.17	U

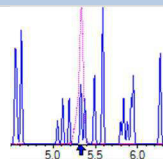
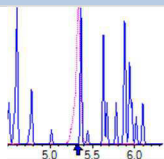
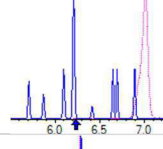
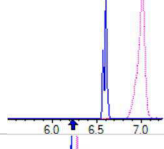
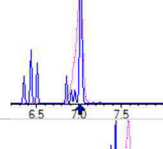
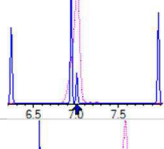
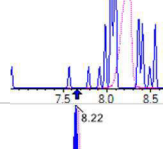
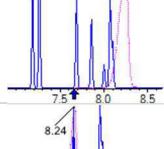
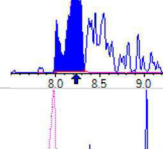
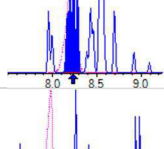
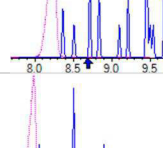
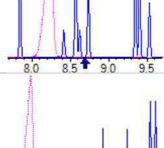
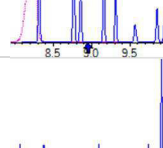
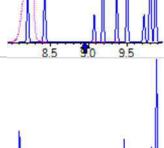
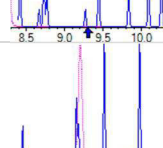
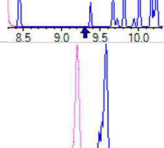
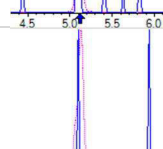
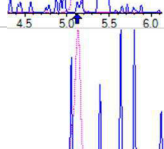
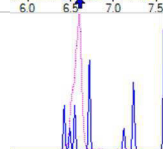
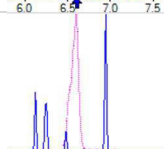
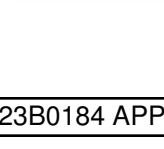
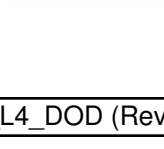
ANALYSIS DATA SHEET

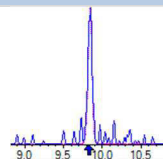
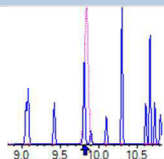
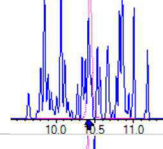
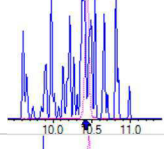
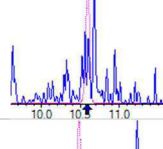
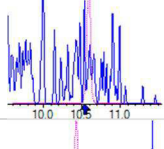
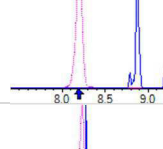
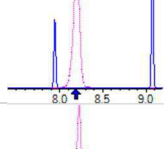
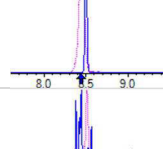
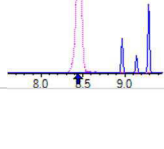
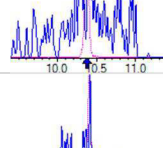
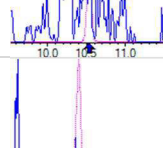
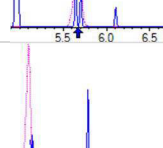
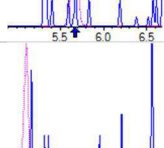
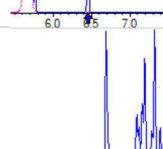
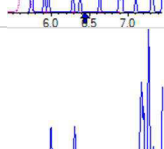

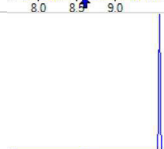
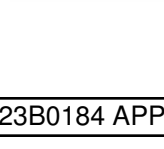
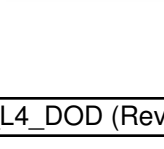
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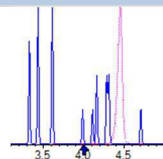
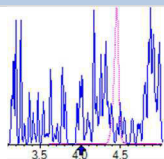
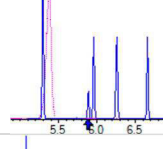
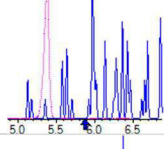
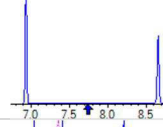
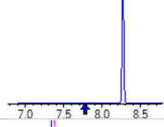
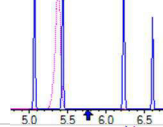
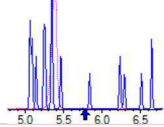
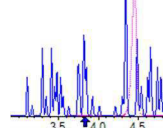
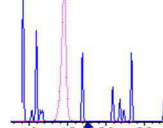
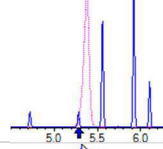
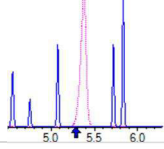
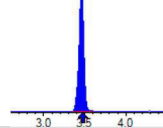
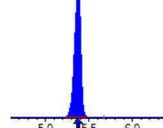
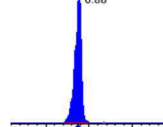
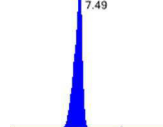
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Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Matrix:	Water	Laboratory ID:	BCB0459-BLK1
Sampled:		Prepared:	02/28/23 08:24
Solids:		Preparation:	EPA 1633
Batch:	BCB0459	Sequence:	SC00861
Column:	1	Calibration:	2309009
		Instrument:	Saphira

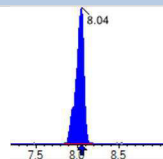
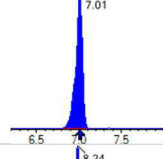
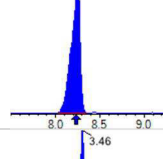
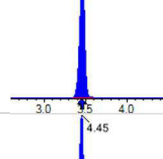
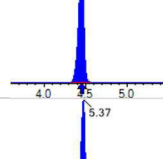
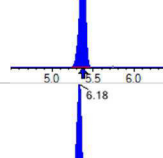
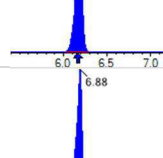
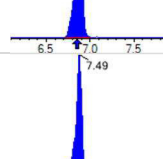
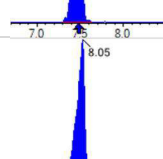
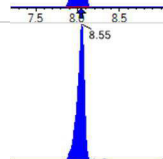
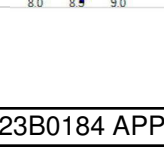
COMPOUND	CONC. (ng/L)	LOQ	LOD	DL	Q
ADONA	0.40 U	0.80	0.40	0.12	U
PFEESA	0.40 U	0.80	0.40	0.11	U
PFMPA	0.40 U	0.80	0.40	0.054	U
PFMBA	0.40 U	0.80	0.40	0.091	U
NFDHA	0.40 U	0.80	0.40	0.30	U
9CL-PF3ONS	0.40 U	0.80	0.40	0.21	U
11CL-PF3OUDS	0.40 U	0.80	0.40	0.21	U
3:3FTCA	0.80 U	1.6	0.80	0.57	U
5:3FTCA	0.80 U	1.6	0.80	0.44	U
7:3FTCA	0.80 U	1.6	0.80	0.55	U

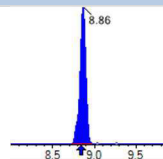
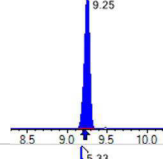
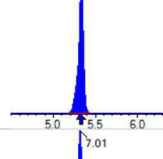
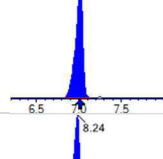
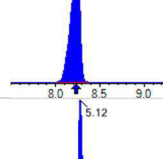
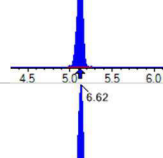
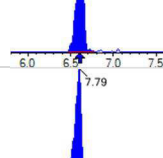
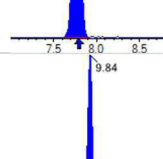
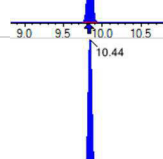
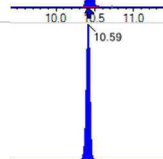
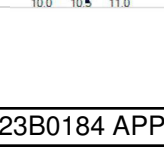
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
PFPeA	(263.0 / 219.0) N/A (263.0 / 69.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHxA	(313.0 / 269.0) N/A (313.0 / 119.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHpA	(363.0 / 319.0) N/A (363.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFOA	(413.0 / 369.0) N/A (413.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFNA	(463.0 / 419.0) N/A (463.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDA	(513.0 / 469.0) N/A (513.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFUnA	(563.0 / 519.0) N/A (563.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDaA	(613.0 / 569.0) N/A (613.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFTTrDA	(663.0 / 619.0) N/A (663.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFTeDA	(713.0 / 669.0) N/A (713.0 / 169.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

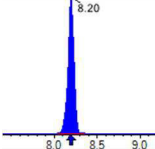
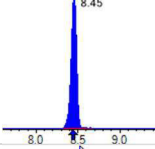
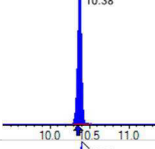
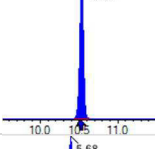
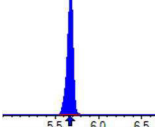
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) N/A (299.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFPeS	(349.0 / 80.0) N/A (349.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHxS	(399.0 / 80.0) N/A (399.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFHpS	(449.0 / 80.0) N/A (449.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFOS	(499.0 / 80.0) 54021 (499.0 / 99.0) 7797	(8.22 , 1.00) (-0.01 , N/A , -1.2)	20.6 23.1	0.1443 65.6 62.6	0.0256	N/A			M12 ABK 3/2/23
PFNS	(549.0 / 80.0) N/A (549.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDS	(599.0 / 80.0) N/A (599.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFDoS	(699.0 / 80.0) N/A (699.0 / 99.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
4:2FTS	(327.0 / 307.0) N/A (327.0 / 81.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
6:2FTS	(427.0 / 407.0) N/A (427.0 / 81.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
8:2FTS	(527.0 / 507.0) N/A (527.0 / 81.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) N/A (498.0 / 478.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NMeFOSA	(512.0 / 219.0) N/A (512.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NEtFOSA	(526.0 / 219.0) N/A (526.0 / 169.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NMeFOSAA	(570.0 / 419.0) N/A (570.0 / 483.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NEtFOSAA	(584.0 / 419.0) N/A (584.0 / 526.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NMeFOSE	(616.0 / 59.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
NEtFOSE	(630.0 / 59.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
HFPO-DA	(285.0 / 169.0) N/A (285.0 / 185.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
ADONA	(377.0 / 85.0) N/A (377.0 / 251.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
9Cl-Pf3ONS	(531.0 / 351.0) N/A (533.0 / 353.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
11Cl-Pf3OUDS	(631.0 / 451.0) N/A (633.0 / 453.0) N/A	(N/A , N/A) (N/A , N/A , N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) N/A (241.0 / 117.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
5:3FTCA	(341.0 / 236.7) N/A (341.0 / 217.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
7:3FTCA	(441.0 / 317.0) N/A (441.0 / 337.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFEESA	(315.0 / 135.0) N/A (315.0 / 83.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
PFMPA	(229.0 / 85.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
PFMBA	(279.0 / 85.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
NFDHA	(295.0 / 201.0) N/A (295.0 / 85.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
13C3_PFBA_IIS	(216.0 / 172.0) 247271	(3.46, N/A) (N/A, 0.00, N/A)	1269.4	N/A	0.7294 [1.0000]	72.9% { 118.1% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 494238	(5.37, N/A) (N/A, 0.00, N/A)	1424.1	N/A	0.8310 [1.0000]	83.1% { 132.6% }			
13C4_PFOA_IIS	(417.0 / 372.0) 589925	(6.88, N/A) (N/A, 0.01, N/A)	1569.5	N/A	0.7252 [1.0000]	72.5% { 113.8% }			
13C5_PFNA_IIS	(468.0 / 423.0) 551453	(7.49, N/A) (N/A, 0.01, N/A)	1296.0	N/A	0.7222 [1.0000]	72.2% { 100.1% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 616942	(8.04 , N/A) (N/A , 0.01 , N/A)	715.6	N/A	0.8594 [1.0000]	85.9% { 128.9% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 952370	(7.01 , N/A) (N/A , 0.01 , N/A)	939.7	N/A	0.7733 [1.0000]	77.3% { 107.6% }			
13C4_PFOS_IIS	(503.0 / 79.9) 1543777	(8.24 , N/A) (N/A , 0.01 , N/A)	830.0	N/A	0.8698 [1.0000]	87.0% { 119.7% }			
13C4_PFBA_EIS	(217.0 / 172.0) 2226192	(3.46 , N/A) (N/A , 0.00 , N/A)	3240.9	N/A	7.5714 [8.0000]	94.6% { 116.6% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 1940736	(4.45 , N/A) (N/A , 0.00 , N/A)	2099.7	N/A	3.5241 [4.0000]	88.1% { 110.7% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 1276722	(5.37 , N/A) (N/A , 0.01 , N/A)	1243.1	N/A	1.9012 [2.0000]	95.1% { 120.3% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 1185333	(6.18 , N/A) (N/A , 0.02 , N/A)	1901.8	N/A	1.8926 [2.0000]	94.6% { 131.4% }			
13C8_PFOA_EIS	(421.0 / 376.0) 1251228	(6.88 , N/A) (N/A , 0.02 , N/A)	1705.7	N/A	1.9816 [2.0000]	99.1% { 117.8% }			
13C9_PFNA_EIS	(472.0 / 427.0) 567184	(7.49 , N/A) (N/A , 0.01 , N/A)	848.8	N/A	1.0321 [1.0000]	103.2% { 113.2% }			
13C6_PFDA_EIS	(519.0 / 474.0) 691807	(8.05 , N/A) (N/A , 0.02 , N/A)	3221.4	N/A	0.9688 [1.0000]	96.9% { 110.6% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 681370	(8.55 , N/A) (N/A , 0.01 , N/A)	660.5	N/A	0.9632 [1.0000]	96.3% { 115.7% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 652904	(8.86 , N/A) (N/A , 0.02 , N/A)	722.8	N/A	0.9951 [1.0000]	99.5% { 145.8% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 630786	(9.25 , N/A) (N/A , 0.02 , N/A)	1365.4	N/A	0.9249 [1.0000]	92.5% { 121.8% }			
13C3_PFBs_EIS	(302.0 / 80.0) 3894546	(5.33 , N/A) (N/A , 0.01 , N/A)	2449.0	N/A	2.0236 [2.0000]	101.2% { 109.1% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 1812755	(7.01 , N/A) (N/A , 0.01 , N/A)	1165.2	N/A	1.8029 [2.0000]	90.1% { 103.5% }			
13C8_PFOS_EIS	(507.0 / 80.0) 3829127	(8.24 , N/A) (N/A , 0.01 , N/A)	1706.6	N/A	1.8880 [2.0000]	94.4% { 116.9% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 503143	(5.12 , N/A) (N/A , 0.00 , N/A)	1000.6	N/A	4.2684 [4.0000]	106.7% { 124.8% }			
13C2_6:2FTS_EIS	(429.0 / 81.0) 489907	(6.62 , N/A) (N/A , 0.01 , N/A)	488.6	N/A	3.6862 [4.0000]	92.2% { 114.8% }			
13C2_8:2FTS_EIS	(529.0 / 81.0) 599733	(7.79 , N/A) (N/A , 0.01 , N/A)	656.4	N/A	4.2673 [4.0000]	106.7% { 114.7% }			
13C8_PFOsa_EIS	(506.0 / 78.0) 6958315	(9.84 , N/A) (N/A , 0.02 , N/A)	2841.8	N/A	1.8625 [2.0000]	93.1% { 120.2% }			
D3_NMeFOSA_EIS	(515.0 / 169.0) 801873	(10.44 , N/A) (N/A , 0.02 , N/A)	1749.2	N/A	0.9996 [2.0000]	50.0% { 60.4% }			
D5_NEtFOSA_EIS	(531.0 / 169.0) 663663	(10.59 , N/A) (N/A , 0.02 , N/A)	1991.8	N/A	0.9448 [2.0000]	47.2% { 54.8% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1230320	(8.20 , N/A) (N/A , 0.01 , N/A)	1124.5	N/A	4.3224 [4.0000]	108.1% { 110.8% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 1021315	(8.45 , N/A) (N/A , 0.01 , N/A)	2570.8	N/A	4.2136 [4.0000]	105.3% { 121.2% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 3133414	(10.38 , N/A) (N/A , 0.02 , N/A)	1446.1	N/A	11.4728 [20.0000]	57.4% { 73.3% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 4793099	(10.54 , N/A) (N/A , 0.02 , N/A)	1318.5	N/A	15.3422 [20.0000]	76.7% { 89.8% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 3091820	(5.68 , N/A) (N/A , 0.01 , N/A)	2348.0	N/A	7.3271 [8.0000]	91.6% { 111.2% }			

ANALYSIS DATA SHEET

LCS

Laboratory:	APPL, LLC	Work Order:	23B0184
Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Matrix:	Water	Laboratory ID:	BCB0459-BS1
		File ID:	S2023-03-02A (8)
Sampled:		Prepared:	02/28/23 08:24
		Analyzed:	03/02/23 11:12
Solids:		Preparation:	EPA 1633
		Dilution:	1
Batch:	BCB0459	Sequence:	SC00861
		Calibration:	2309009
Column:	1	Instrument:	Saphira

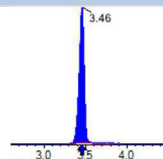
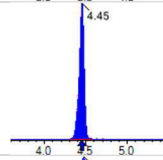
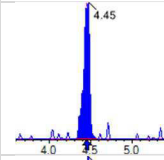
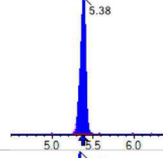
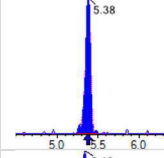
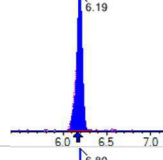
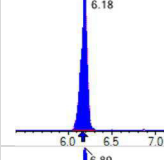
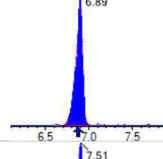
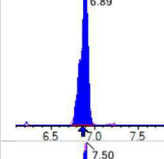
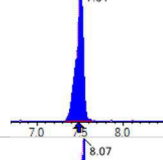
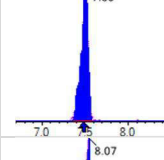
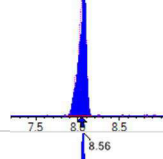
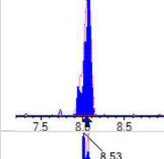
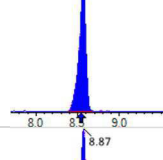
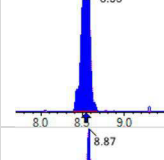
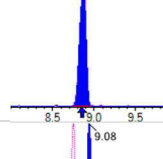
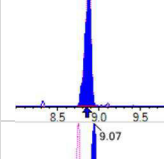
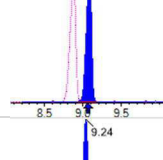
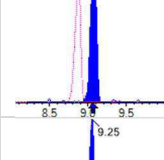
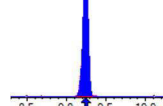
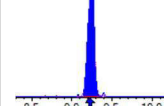
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PFPEA	7.64	0.80	0.065	
PFHXA	3.61	0.40	0.055	
PFHPA	3.47	0.40	0.041	
PFOA	3.68	0.40	0.15	
PFNA	4.21	0.40	0.082	
PFDA	4.00	0.40	0.10	
PFUnA	3.57	0.40	0.16	
PFDOA	3.98	0.40	0.11	
PFTRDA	3.96	0.40	0.20	
PFTEDA	3.41	0.40	0.20	
PFBS	3.34	0.40	0.037	
PFPEs	3.99	0.40	0.063	
PFHXS	3.66	0.40	0.032	
PFHPS	3.51	0.40	0.051	
PFOS	3.67	0.40	0.064	
PFNS	3.90	0.40	0.12	
PFDS	3.76	0.40	0.15	
PFDOS	4.26	0.40	0.12	
4:2FTS	13.8	1.6	0.29	
6:2FTS	16.3	1.6	0.31	
8:2FTS	14.9	1.6	0.082	
PFOSA	4.48	0.40	0.10	
NMeFOSA	15.3	1.6	0.47	
NEtFOSA	16.9	1.6	0.41	
NMeFOSAA	4.00	0.40	0.11	
NEtFOSAA	4.05	0.40	0.11	
NMeFOSE	16.2	1.6	1.0	
NEtFOSE	15.5	1.6	1.0	
HFPO-DA	7.82	0.80	0.17	

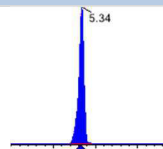
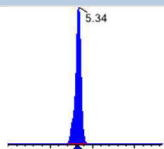
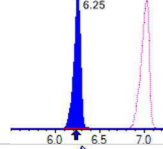
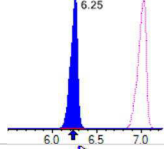
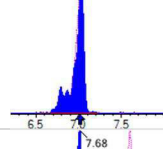
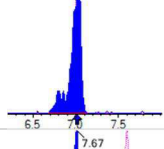
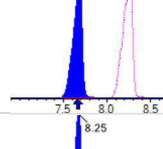
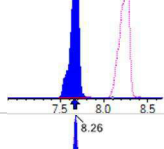
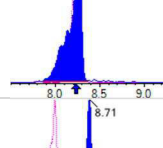
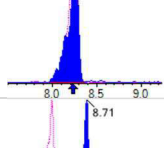
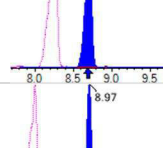
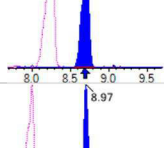
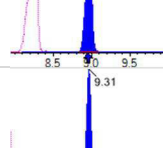
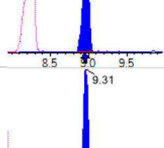
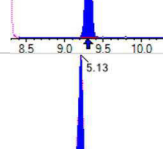
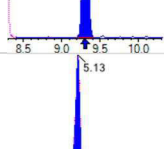
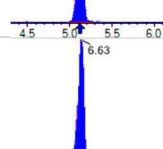
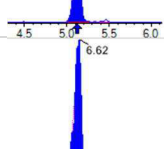
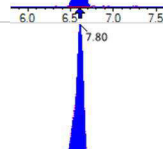
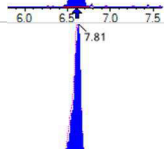
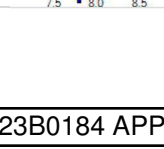
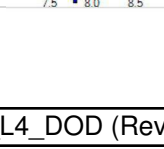
ANALYSIS DATA SHEET

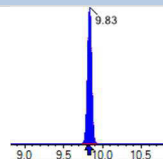
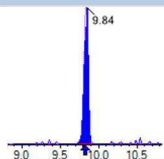
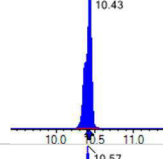
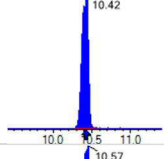
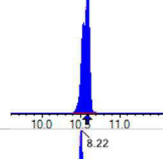
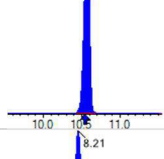
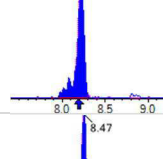
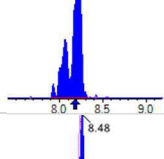
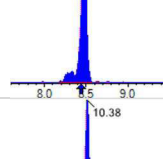
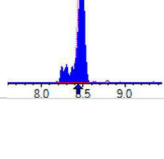
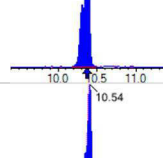
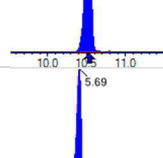
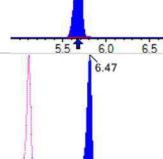
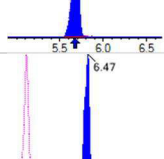
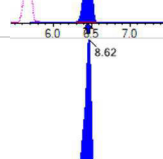
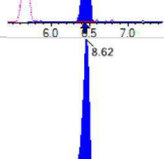
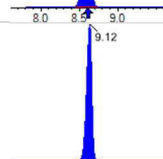
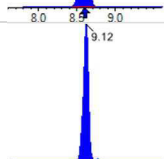
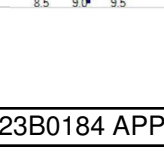
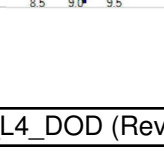
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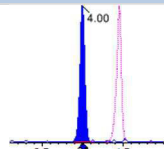
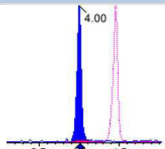
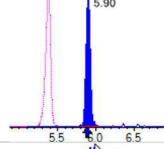
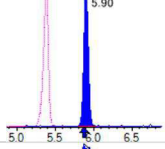
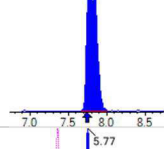
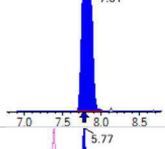
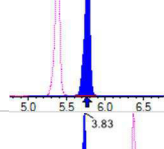
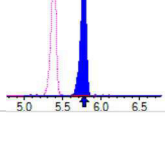
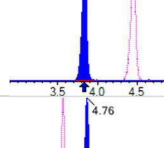
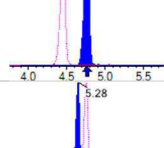
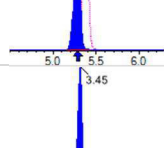
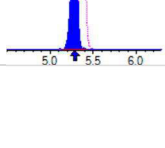
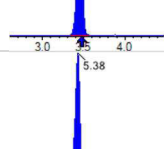
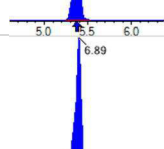
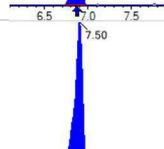
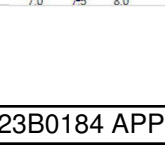
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Matrix:	Water	Laboratory ID:	BCB0459-BS1
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Sampled:		Prepared:	02/28/23 08:24
		Analyzed:	03/02/23 11:12
Solids:		Preparation:	EPA 1633
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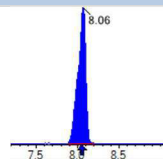
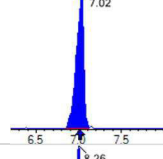
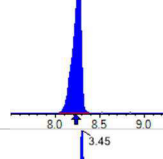
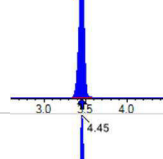
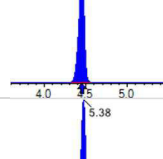
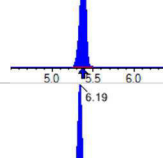
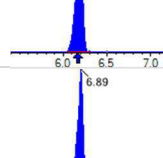
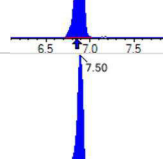
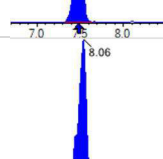
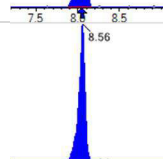
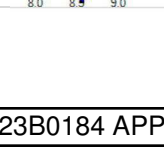
COMPOUND	CONC. (ng/L)	LOQ	DL	Q
ADONA	8.03	0.80	0.12	
PFEESA	6.83	0.80	0.11	
PFMPA	6.84	0.80	0.054	
PFMBA	8.10	0.80	0.091	
NFDHA	6.63	0.80	0.30	
9CL-PF3ONS	8.85	0.80	0.21	
11CL-PF3OUDS	8.54	0.80	0.21	
3:3FTCA	13.2	1.6	0.57	
5:3FTCA	11.3	1.6	0.44	
7:3FTCA	12.4	1.6	0.55	

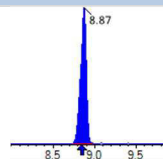
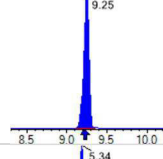
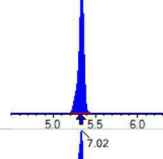
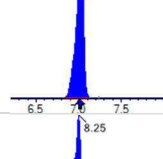
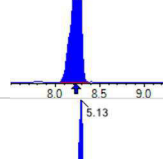
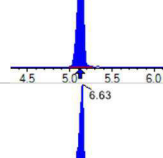
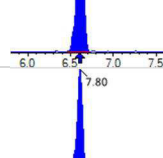
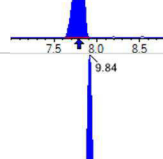
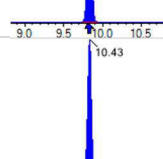
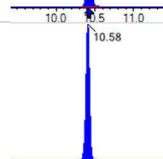
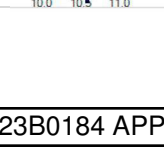
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT Ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) 920526	(3.46 , 1.00) (0.00 , N/A , 0.0)	140.1	N/A 0.0 0.0	3.9119 [4.0000]	97.8%			
PFPeA	(263.0 / 219.0) 816971 (263.0 / 69.0) 13372	(4.45 , 1.00) (0.00 , N/A , -0.1)	1543.0 116.9	0.0164 137.3 139.1	1.9090 [2.0000]	95.4%			
PFHxA	(313.0 / 269.0) 562358 (313.0 / 119.0) 65176	(5.38 , 1.00) (0.00 , N/A , 0.2)	1588.6 11634.9	0.1159 123.9 112.2	0.9018 [1.0000]	90.2%			
PFHpA	(363.0 / 319.0) 444492 (363.0 / 169.0) 187739	(6.19 , 1.00) (0.00 , N/A , 0.2)	3089.5 3162890.4	0.4224 133.8 129.5	0.8681 [1.0000]	86.8%			
PFOA	(413.0 / 369.0) 526288 (413.0 / 169.0) 185476	(6.89 , 1.00) (0.00 , N/A , 0.0)	2284.8 17379.3	0.3524 104.2 99.2	0.9191 [1.0000]	91.9%			
PFNA	(463.0 / 419.0) 567626 (463.0 / 169.0) 106895	(7.51 , 1.00) (0.01 , N/A , 0.3)	8803.0 1168.6	0.1883 80.4 81.2	1.0524 [1.0000]	105.2%			
PFDA	(513.0 / 469.0) 650510 (513.0 / 169.0) 87331	(8.07 , 1.00) (0.01 , N/A , 0.1)	684.1 530.4	0.1342 108.3 99.1	0.9989 [1.0000]	99.9%			
PFUnA	(563.0 / 519.0) 502934 (563.0 / 169.0) 68651	(8.56 , 1.00) (0.00 , N/A , 1.7)	850.1 665.3	0.1365 126.8 93.5	0.8915 [1.0000]	89.2%			
PFDaA	(613.0 / 569.0) 526444 (613.0 / 169.0) 101779	(8.87 , 1.00) (-0.01 , N/A , -0.3)	911.0 1137.3	0.1933 121.7 102.4	0.9949 [1.0000]	99.5%			
PFTTrDA	(663.0 / 619.0) 503552 (663.0 / 169.0) 125332	(9.08 , 1.02) (N/A , 0.02 , 0.4)	868.4 459.9	0.2489 91.7 87.4	0.9889 [1.0000]	98.9%			
PFTeDA	(713.0 / 669.0) 478001 (713.0 / 169.0) 121605	(9.24 , 1.00) (0.00 , N/A , -0.1)	1048.7 641.5	0.2544 116.4 115.5	0.8527 [1.0000]	85.3%			

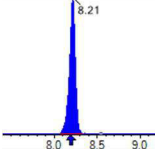
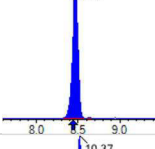
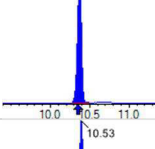
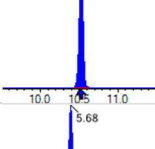
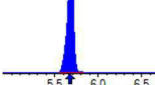
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) 903617 (299.0 / 99.0) 564492	(5.34 , 1.00) (0.00 , N/A , 0.1)	1362.1 1599.5	0.6247 99.3 100.1	0.8338 [0.8847]	94.2%			
PFPeS	(349.0 / 80.0) 1583169 (349.0 / 99.0) 589628	(6.25 , 0.89) (N/A , 0.02 , 0.0)	86354.6 10369.6	0.3724 111.3 111.2	0.9986 [0.9384]	106.4%			
PFHxS	(399.0 / 80.0) 1274216 (399.0 / 99.0) 427127	(7.03 , 1.00) (0.01 , N/A , 0.6)	3790.4 1090.2	0.3352 104.7 97.0	0.9160 [0.9110]	100.6%			
PFHpS	(449.0 / 80.0) 1445055 (449.0 / 99.0) 361440	(7.68 , 0.93) (N/A , 0.02 , 0.1)	616188.5 29417.4	0.2501 92.1 89.4	0.8769 [0.9514]	92.2%			
PFOS	(499.0 / 80.0) 1854222 (499.0 / 99.0) 446555	(8.25 , 1.00) (0.01 , N/A , -0.1)	208.0 302.5	0.2408 109.5 104.4	0.9181 [0.9275]	99.0%			
PFNS	(549.0 / 80.0) 1969080 (549.0 / 99.0) 499906	(8.71 , 1.06) (N/A , 0.02 , -0.1)	11308.0 5543588.9	0.2539 105.5 110.1	0.9743 [0.9599]	101.5%			
PFDS	(599.0 / 80.0) 2112208 (599.0 / 99.0) 441900	(8.97 , 1.09) (N/A , 0.02 , 0.1)	1371.9 1181.1	0.2092 103.4 98.0	0.9404 [0.9631]	97.6%			
PFDoS	(699.0 / 80.0) 1103364 (699.0 / 99.0) 248315	(9.31 , 1.13) (N/A , 0.01 , 0.0)	1088.6 656.4	0.2251 115.2 107.8	1.0641 [0.9696]	109.8%			
4:2FTS	(327.0 / 307.0) 1243634 (327.0 / 81.0) 825355	(5.13 , 1.00) (0.00 , N/A , 0.0)	1450.4 855.8	0.6637 95.8 104.7	3.4499 [3.7381]	92.3%			
6:2FTS	(427.0 / 407.0) 641861 (427.0 / 81.0) 473060	(6.63 , 1.00) (0.00 , N/A , 0.1)	867.9 584.6	0.7370 87.0 96.2	4.0654 [3.7962]	107.1%			
8:2FTS	(527.0 / 507.0) 670180 (527.0 / 81.0) 541519	(7.80 , 1.00) (0.00 , N/A , -0.5)	850.6 654.1	0.8080 99.0 101.3	3.7247 [3.8332]	97.2%			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) 2599515 (498.0 / 478.0) 57353	(9.83 , 1.00) (0.00 , N/A , -0.5)	2282.1 270.2	0.0221 112.3 108.2	1.1198 [1.0000]	112.0%			
NMeFOSA	(512.0 / 219.0) 1344166 (512.0 / 169.0) 1133100	(10.43 , 1.00) (0.00 , N/A , 1.0)	2708.8 2020.8	0.8430 102.0 100.7	3.8146 [4.0000]	95.4%			
NEIFOSA	(526.0 / 219.0) 1433002 (526.0 / 169.0) 1893488	(10.57 , 1.00) (-0.01 , N/A , 0.4)	3533.2 3375.7	1.3213 104.3 103.1	4.2338 [4.0000]	105.8%			
NMeFOSAA	(570.0 / 419.0) 253805 (570.0 / 483.0) 111347	(8.22 , 1.00) (0.01 , N/A , 0.5)	1034.7 126.8	0.4387 89.5 99.0	1.0006 [1.0000]	100.1%			
NEIFOSAA	(584.0 / 419.0) 200447 (584.0 / 526.0) 107757	(8.47 , 1.00) (0.01 , N/A , -0.6)	13483.5 172003.7	0.5376 95.1 104.9	1.0132 [1.0000]	101.3%			
NMeFOSE	(616.0 / 59.0) 626051	(10.38 , 1.00) (0.01 , N/A , 0.0)	899.1	N/A 0.0 0.0	4.0463 [4.0000]	101.2%			
NEtFOSE	(630.0 / 59.0) 910402	(10.54 , 1.00) (0.01 , N/A , 0.0)	784.3	N/A 0.0 0.0	3.8870 [4.0000]	97.2%			
HFPO-DA	(285.0 / 169.0) 573861 (285.0 / 185.0) 1640403	(5.69 , 1.00) (0.00 , N/A , 0.0)	1299.3 1533.9	2.8585 104.0 94.5	1.9550 [2.0000]	97.8%			
ADONA	(377.0 / 85.0) 2185748 (377.0 / 251.0) 210581	(6.47 , 1.14) (N/A , 0.03 , 0.1)	1582.9 590.4	0.0963 110.1 110.8	2.0064 [1.8854]	106.4%			
9CI-Pf3ONS	(531.0 / 351.0) 5653435 (533.0 / 353.0) 1729830	(8.62 , 1.52) (N/A , 0.02 , 0.0)	1707.9 1187.5	0.3060 103.9 98.3	2.2127 [1.8665]	118.5%			
11CI-PF3OUDS	(631.0 / 451.0) 2682263 (633.0 / 453.0) 839432	(9.12 , 1.61) (N/A , 0.02 , 0.1)	2178.7 855.8	0.3130 101.4 101.1	2.1353 [1.8864]	113.2%			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) 41789 (241.0 / 117.0) 67267	(4.00 , 0.90) (N/A , 0.00 , -0.2)	485.7 357.6	1.6097 92.8 97.9	3.3065 [4.0000]	82.7%			
5:3FTCA	(341.0 / 236.7) 251738 (341.0 / 217.0) 437579	(5.90 , 1.10) (N/A , 0.02 , -0.1)	761.3 639.9	1.7382 103.4 95.7	2.8239 [4.0000]	70.6%			
7:3FTCA	(441.0 / 317.0) 496491 (441.0 / 337.0) 447225	(7.81 , 1.45) (N/A , 0.03 , -0.2)	461.3 875.5	0.9008 104.9 107.3	3.0980 [4.0000]	77.4%			
PFEESA	(315.0 / 135.0) 1323075 (315.0 / 83.0) 370994	(5.77 , 1.07) (N/A , 0.02 , 0.0)	2132.8 822.3	0.2804 99.6 107.1	1.7074 [1.7849]	95.7%			
PFMPA	(229.0 / 85.0) 187240	(3.83 , 0.86) (N/A , 0.00 , 0.0)	1070.2	N/A 0.0 0.0	1.7112 [2.0000]	85.6%			
PFMBA	(279.0 / 85.0) 758712	(4.76 , 1.07) (N/A , 0.00 , 0.0)	1518.8	N/A 0.0 0.0	2.0240 [2.0000]	101.2%			
NFDHA	(295.0 / 201.0) 575058 (295.0 / 85.0) 640325	(5.28 , 0.98) (N/A , 0.01 , 0.1)	1650.2 1182.9	1.1135 106.7 107.4	1.6580 [2.0000]	82.9%			
13C3_PFBA_IIS	(216.0 / 172.0) 244669	(3.45 , N/A) (N/A , 0.00 , N/A)	1156.6	N/A	0.7217 [1.0000]	72.2% { 116.8% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 505590	(5.38 , N/A) (N/A , 0.01 , N/A)	1570.3	N/A	0.8501 [1.0000]	85.0% { 135.7% }			
13C4_PFOA_IIS	(417.0 / 372.0) 607388	(6.89 , N/A) (N/A , 0.03 , N/A)	926.5	N/A	0.7467 [1.0000]	74.7% { 117.2% }			
13C5_PFNA_IIS	(468.0 / 423.0) 578860	(7.50 , N/A) (N/A , 0.03 , N/A)	765.9	N/A	0.7580 [1.0000]	75.8% { 105.1% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 554686	(8.06 , N/A) (N/A , 0.03 , N/A)	713.5	N/A	0.7727 [1.0000]	77.3% { 115.9% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 853978	(7.02 , N/A) (N/A , 0.02 , N/A)	804.0	N/A	0.6934 [1.0000]	69.3% { 96.5% }			
13C4_PFOS_IIS	(503.0 / 79.9) 1456400	(8.26 , N/A) (N/A , 0.03 , N/A)	1001.5	N/A	0.8206 [1.0000]	82.1% { 112.9% }			
13C4_PFBA_EIS	(217.0 / 172.0) 2226034	(3.45 , N/A) (N/A , 0.00 , N/A)	3508.1	N/A	7.6514 [8.0000]	95.6% { 116.6% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 1942773	(4.45 , N/A) (N/A , 0.00 , N/A)	1959.6	N/A	3.4486 [4.0000]	86.2% { 110.8% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 1336433	(5.38 , N/A) (N/A , 0.01 , N/A)	1634.1	N/A	1.9454 [2.0000]	97.3% { 125.9% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 1146760	(6.19 , N/A) (N/A , 0.03 , N/A)	1577.0	N/A	1.7899 [2.0000]	89.5% { 127.2% }			
13C8_PFOA_EIS	(421.0 / 376.0) 1188587	(6.89 , N/A) (N/A , 0.03 , N/A)	1000.2	N/A	1.8283 [2.0000]	91.4% { 111.9% }			
13C9_PFNA_EIS	(472.0 / 427.0) 570976	(7.50 , N/A) (N/A , 0.02 , N/A)	1103.3	N/A	0.9898 [1.0000]	99.0% { 114.0% }			
13C6_PFDA_EIS	(519.0 / 474.0) 656844	(8.06 , N/A) (N/A , 0.03 , N/A)	750.5	N/A	1.0231 [1.0000]	102.3% { 105.0% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 641512	(8.56 , N/A) (N/A , 0.03 , N/A)	1018.5	N/A	1.0086 [1.0000]	100.9% { 108.9% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT Ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 606532	(8.87 , N/A) (N/A , 0.02 , N/A)	971.6	N/A	1.0282 [1.0000]	102.8% { 135.4% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 609654	(9.25 , N/A) (N/A , 0.02 , N/A)	1564.3	N/A	0.9942 [1.0000]	99.4% { 117.8% }			
13C3_PFBs_EIS	(302.0 / 80.0) 3811894	(5.34 , N/A) (N/A , 0.01 , N/A)	2106.5	N/A	2.2089 [2.0000]	110.4% { 106.8% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 1766111	(7.02 , N/A) (N/A , 0.02 , N/A)	1113.7	N/A	1.9589 [2.0000]	97.9% { 100.8% }			
13C8_PFOS_EIS	(507.0 / 80.0) 3660212	(8.25 , N/A) (N/A , 0.02 , N/A)	767.1	N/A	1.9130 [2.0000]	95.6% { 111.8% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 510869	(5.13 , N/A) (N/A , 0.01 , N/A)	1026.5	N/A	4.8333 [4.0000]	120.8% { 126.7% }			
13C2_6:2FTS_EIS	(429.0 / 81.0) 484088	(6.63 , N/A) (N/A , 0.02 , N/A)	526.3	N/A	4.0621 [4.0000]	101.6% { 113.4% }			
13C2_8:2FTS_EIS	(529.0 / 81.0) 641108	(7.80 , N/A) (N/A , 0.02 , N/A)	734.0	N/A	5.0873 [4.0000]	127.2% { 122.6% }			
13C8_PFOsa_EIS	(506.0 / 78.0) 5974624	(9.84 , N/A) (N/A , 0.01 , N/A)	2616.3	N/A	1.6952 [2.0000]	84.8% { 103.2% }			
D3_NMeFOSA_EIS	(515.0 / 169.0) 820589	(10.43 , N/A) (N/A , 0.01 , N/A)	1894.5	N/A	1.0843 [2.0000]	54.2% { 61.8% }			
D5_NEtFOSA_EIS	(531.0 / 169.0) 696809	(10.58 , N/A) (N/A , 0.01 , N/A)	1888.3	N/A	1.0515 [2.0000]	52.6% { 57.5% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT Ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1187171	(8.21 , N/A) (N/A , 0.03 , N/A)	1273.1	N/A	4.4210 [4.0000]	110.5% { 106.9% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 907796	(8.47 , N/A) (N/A , 0.03 , N/A)	2224.1	N/A	3.9700 [4.0000]	99.2% { 107.8% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 2953244	(10.37 , N/A) (N/A , 0.02 , N/A)	890.9	N/A	11.4619 [20.0000]	57.3% { 69.1% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 4635821	(10.53 , N/A) (N/A , 0.01 , N/A)	1189.8	N/A	15.7290 [20.0000]	78.6% { 86.8% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 2864341	(5.68 , N/A) (N/A , 0.02 , N/A)	2292.7	N/A	6.6356 [8.0000]	82.9% { 103.0% }			

ANALYSIS DATA SHEET

MRL Check

Laboratory:	APPL, LLC	Work Order:	23B0184
Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Matrix:	Water	Laboratory ID:	BCB0459-MRL1
		File ID:	S2023-03-02A (9)
Sampled:		Prepared:	02/28/23 08:24
		Analyzed:	03/02/23 11:25
Solids:		Preparation:	EPA 1633
		Dilution:	1
Batch:	BCB0459	Sequence:	SC00861
		Calibration:	2309009
Column:	1	Instrument:	Saphira

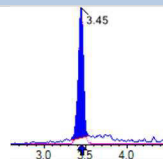
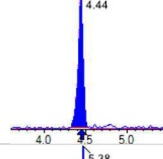
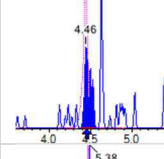
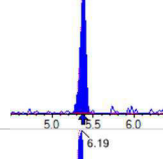
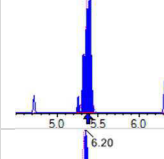
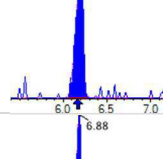
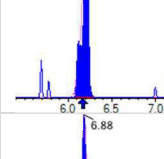
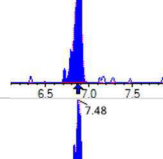
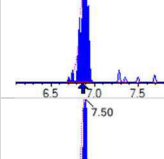
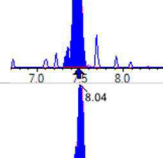
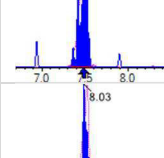
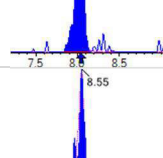
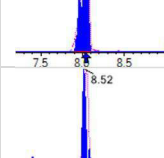
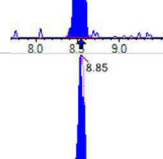
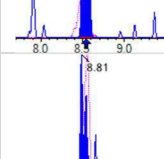
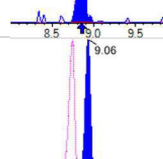
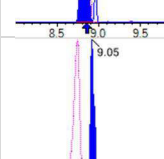
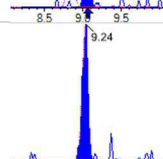
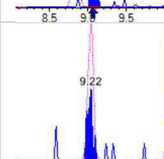
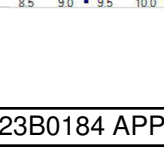
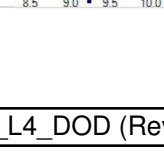
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PFPEA	0.916	0.80	0.065	
PFHXA	0.462	0.40	0.055	
PFHPA	0.391	0.40	0.041	J
PFOA	0.492	0.40	0.15	
PFNA	0.297	0.40	0.082	IR2, J
PFDA	0.498	0.40	0.10	IR2
PFUnA	0.386	0.40	0.16	IR2, J
PFDOA	0.451	0.40	0.11	
PFTRDA	0.321	0.40	0.20	IR2, J
PFTEDA	0.410	0.40	0.20	IR1
PFBS	0.364	0.40	0.037	J
PFPEs	0.380	0.40	0.063	J
PFHXS	0.379	0.40	0.032	J
PFHPS	0.349	0.40	0.051	J
PFOS	0.439	0.40	0.064	
PFNS	0.379	0.40	0.12	J
PFDS	0.381	0.40	0.15	J
PFDOS	0.388	0.40	0.12	IR2, J
4:2FTS	1.54	1.6	0.29	J
6:2FTS	1.97	1.6	0.31	
8:2FTS	1.65	1.6	0.082	
PFOSA	0.482	0.40	0.10	
NMeFOSA	1.65	1.6	0.47	
NEtFOSA	1.77	1.6	0.41	
NMeFOSAA	0.416	0.40	0.11	
NEtFOSAA	0.452	0.40	0.11	
NMeFOSE	1.73	1.6	1.0	
NEtFOSE	1.72	1.6	1.0	
HFPO-DA	0.796	0.80	0.17	J

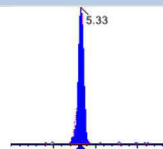
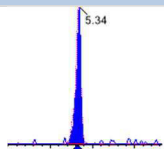
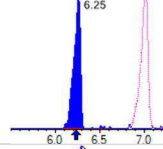
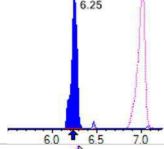
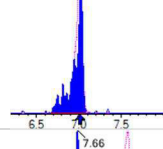
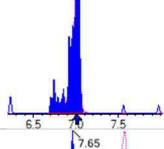
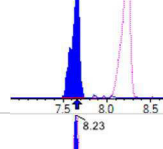
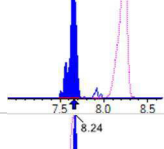
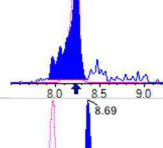
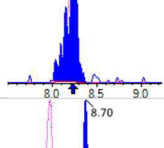
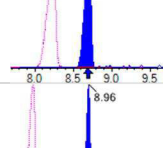
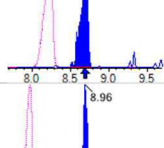
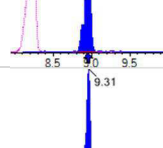
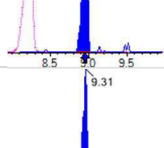
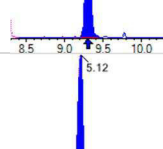
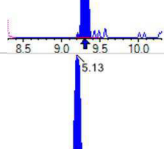
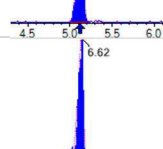
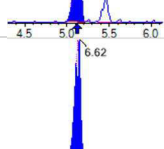
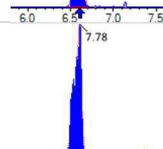
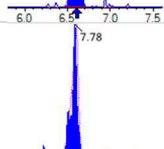
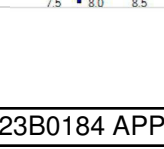
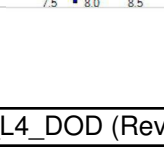
ANALYSIS DATA SHEET

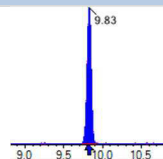
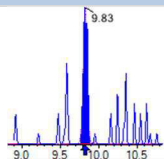
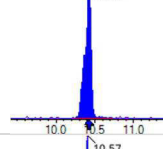
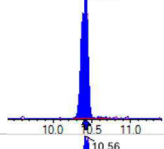
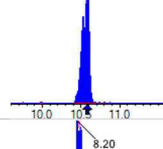
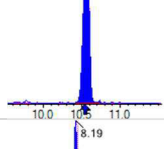
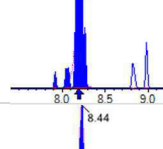
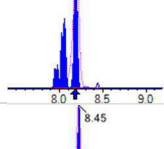
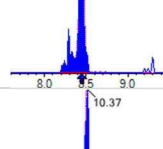
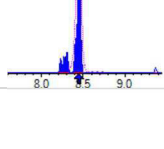
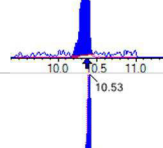
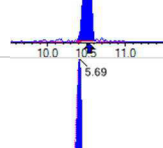
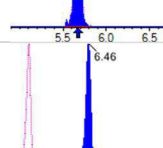
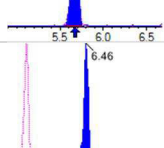
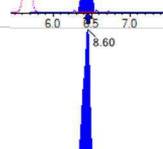
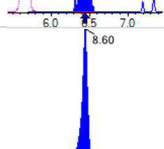
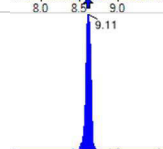
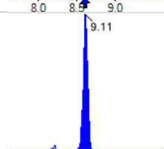
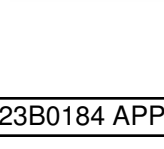
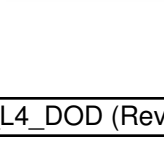
MRL Check

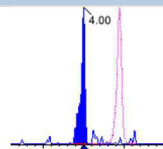
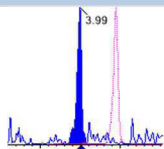
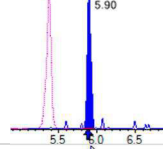
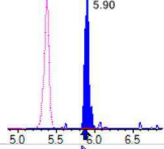
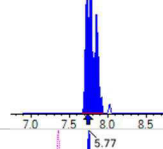
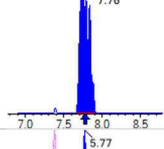
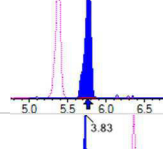
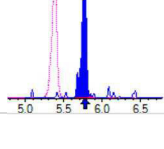
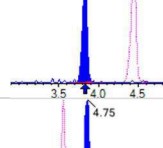
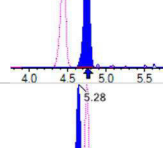
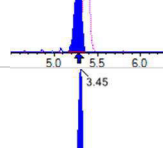
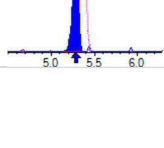
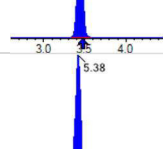
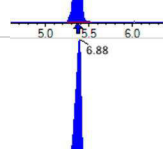
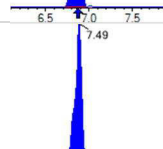
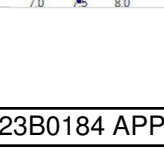
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Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Matrix:	Water	Laboratory ID:	BCB0459-MRL1
		File ID:	S2023-03-02A (9)
Sampled:		Prepared:	02/28/23 08:24
		Analyzed:	03/02/23 11:25
Solids:		Preparation:	EPA 1633
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Batch:	BCB0459	Sequence:	SC00861
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Column:	1	Instrument:	Saphira

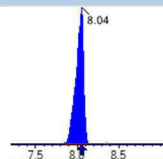
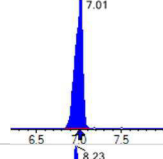
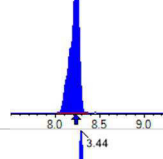
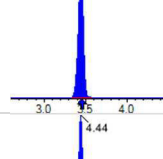
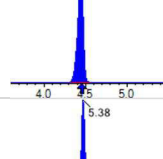
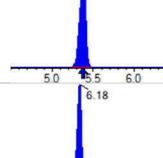
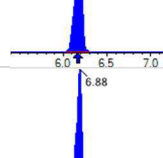
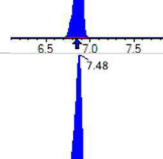
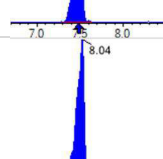
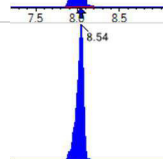
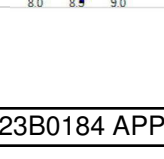
COMPOUND	CONC. (ng/L)	LOQ	DL	Q
ADONA	0.787	0.80	0.12	J
PFEESA	0.792	0.80	0.11	J
PFMPA	0.761	0.80	0.054	J
PFMBA	0.908	0.80	0.091	
NFDHA	0.725	0.80	0.30	J
9CL-PF3ONS	0.875	0.80	0.21	
11CL-PF3OUDS	0.766	0.80	0.21	J
3:3FTCA	1.60	1.6	0.57	
5:3FTCA	1.72	1.6	0.44	
7:3FTCA	1.12	1.6	0.55	J

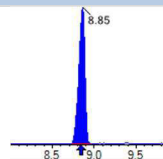
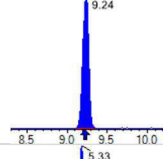
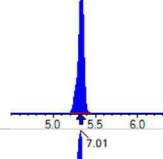
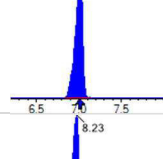
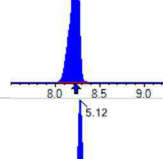
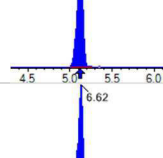
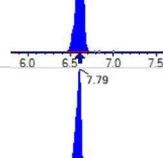
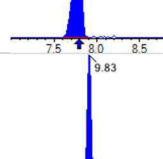
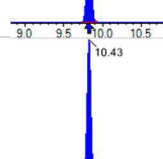
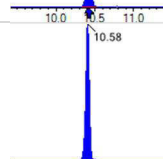
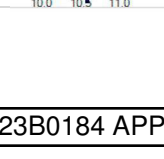
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) 91686	(3.45, 1.00) (0.00, N/A, 0.0)	83.3	N/A 0.0 0.0	0.3978 [0.4000]	99.5%			
PFPeA	(263.0 / 219.0) 94768 (263.0 / 69.0) 1036	(4.44, 1.00) (0.00, N/A, -0.9)	327.1 15.9	0.0109 91.7 92.9	0.2289 [0.2000]	114.5%			
PFHxA	(313.0 / 269.0) 65956 (313.0 / 119.0) 8609	(5.38, 1.00) (0.00, N/A, -0.3)	307.5 355178.8	0.1305 139.5 126.4	0.1156 [0.1000]	115.6%			
PFHpA	(363.0 / 319.0) 47796 (363.0 / 169.0) 15434	(6.19, 1.00) (0.01, N/A, -0.6)	1248.0 417.9	0.3229 102.3 99.0	0.0976 [0.1000]	97.6%			
PFOA	(413.0 / 369.0) 69878 (413.0 / 169.0) 34275	(6.88, 1.00) (0.00, N/A, -0.1)	13730.0 578.3	0.4905 145.0 138.0	0.1231 [0.1000]	123.1%			
PFNA	(463.0 / 419.0) 37383 (463.0 / 169.0) 14039	(7.48, 1.00) (0.00, N/A, -1.2)	28848.0 260650.5	0.3755 160.4 162.0	0.0742 [0.1000]	74.2%			IR2,
PFDA	(513.0 / 469.0) 79784 (513.0 / 169.0) 16322	(8.04, 1.00) (-0.01, N/A, 0.6)	85.2 10459841.5	0.2046 165.1 151.1	0.1244 [0.1000]	124.4%			IR2,
PFUnA	(563.0 / 519.0) 56613 (563.0 / 169.0) 11614	(8.55, 1.00) (0.01, N/A, 2.1)	123.1 146.7	0.2052 190.6 140.5	0.0964 [0.1000]	96.4%			IR2,
PFDaA	(613.0 / 569.0) 56974 (613.0 / 169.0) 7637	(8.85, 1.00) (-0.01, N/A, 2.4)	167.1 34.8	0.1340 84.4 71.0	0.1127 [0.1000]	112.7%			
PFTTrDA	(663.0 / 619.0) 39102 (663.0 / 169.0) 21380	(9.06, 1.02) (N/A, 0.00, 0.8)	108.8 418.8	0.5468 201.4 192.1	0.0804 [0.1000]	80.4%			IR2,
PFTeDA	(713.0 / 669.0) 51851 (713.0 / 169.0) 4916	(9.24, 1.00) (0.00, N/A, 1.0)	110.1 38.0	0.0948 43.4 43.0	0.1024 [0.1000]	102.4%			IR1,

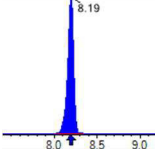
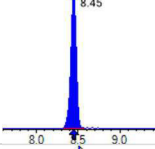
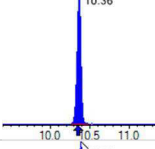
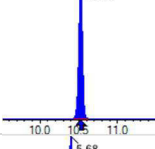
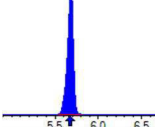
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) 88760 (299.0 / 99.0) 57379	(5.33, 1.00) (0.00, N/A, -0.7)	488.3 144.9	0.6464 102.8 103.6	0.0910 [0.0885]	102.8%			
PFPeS	(349.0 / 80.0) 156363 (349.0 / 99.0) 60274	(6.25, 0.89) (N/A, 0.02, 0.2)	239204.7 1329.4	0.3855 115.2 115.1	0.0949 [0.0938]	101.1%			
PFHxS	(399.0 / 80.0) 136912 (399.0 / 99.0) 46255	(7.02, 1.00) (0.01, N/A, 1.0)	424.8 28564.2	0.3378 105.6 97.8	0.0947 [0.0911]	104.0%			
PFHpS	(449.0 / 80.0) 146810 (449.0 / 99.0) 57468	(7.66, 0.93) (N/A, 0.01, 0.5)	897.7 106.0	0.3914 144.2 139.9	0.0873 [0.0951]	91.7%			
PFOS	(499.0 / 80.0) 226532 (499.0 / 99.0) 53810	(8.23, 1.00) (0.00, N/A, -0.7)	62.3 67.4	0.2375 108.0 103.0	0.1099 [0.0927]	118.5%			
PFNS	(549.0 / 80.0) 195247 (549.0 / 99.0) 49159	(8.69, 1.06) (N/A, 0.01, -0.3)	87376.1 749.7	0.2518 104.6 109.2	0.0946 [0.0960]	98.6%			
PFDS	(599.0 / 80.0) 218218 (599.0 / 99.0) 56950	(8.96, 1.09) (N/A, 0.01, -0.1)	373.8 606.0	0.2610 128.9 122.2	0.0952 [0.0963]	98.8%			
PFDoS	(699.0 / 80.0) 102702 (699.0 / 99.0) 31740	(9.31, 1.13) (N/A, 0.01, 0.2)	533.3 134.0	0.3091 158.2 148.1	0.0970 [0.0970]	100.1%			IR2,
4:2FTS	(327.0 / 307.0) 114248 (327.0 / 81.0) 91648	(5.12, 1.00) (0.00, N/A, 0.0)	466.2 261.1	0.8022 115.8 126.5	0.3848 [0.3738]	102.9%			
6:2FTS	(427.0 / 407.0) 73644 (427.0 / 81.0) 60259	(6.62, 1.00) (0.00, N/A, 0.1)	6548.0 302.3	0.8182 96.6 106.8	0.4922 [0.3796]	129.7%			
8:2FTS	(527.0 / 507.0) 69251 (527.0 / 81.0) 50280	(7.78, 1.00) (0.00, N/A, 0.3)	484.0 163.5	0.7261 89.0 91.0	0.4130 [0.3833]	107.7%			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) 293379 (498.0 / 478.0) 3016	(9.83 , 1.00) (0.00 , N/A , -0.2)	790.5 27.5	0.0103 52.3 50.4	0.1206 [0.1000]	120.6%			
NMeFOSA	(512.0 / 219.0) 145787 (512.0 / 169.0) 131781	(10.43 , 1.00) (0.00 , N/A , 0.9)	595.2 703.8	0.9039 109.4 108.0	0.4134 [0.4000]	103.3%			
NEIFOSA	(526.0 / 219.0) 162623 (526.0 / 169.0) 205405	(10.57 , 1.00) (-0.01 , N/A , 0.9)	939.2 652.9	1.2631 99.7 98.6	0.4430 [0.4000]	110.7%			
NMeFOSAA	(570.0 / 419.0) 27015 (570.0 / 483.0) 16626	(8.20 , 1.00) (0.00 , N/A , 0.6)	200127.7 13740.6	0.6155 125.5 138.9	0.1040 [0.1000]	104.0%			
NEIFOSAA	(584.0 / 419.0) 23772 (584.0 / 526.0) 11890	(8.44 , 1.00) (0.00 , N/A , -0.3)	130.5 4626.2	0.5002 88.5 97.6	0.1129 [0.1000]	112.9%			
NMeFOSE	(616.0 / 59.0) 68261	(10.37 , 1.00) (0.01 , N/A , 0.0)	130.5	N/A 0.0 0.0	0.4330 [0.4000]	108.3%			
NEtFOSE	(630.0 / 59.0) 107200	(10.53 , 1.00) (0.01 , N/A , 0.0)	230.2	N/A 0.0 0.0	0.4291 [0.4000]	107.3%			
HFPO-DA	(285.0 / 169.0) 61688 (285.0 / 185.0) 169076	(5.69 , 1.00) (0.01 , N/A , 0.3)	6403.7 1023.6	2.7408 99.7 90.6	0.1990 [0.2000]	99.5%			
ADONA	(377.0 / 85.0) 226334 (377.0 / 251.0) 25354	(6.46 , 1.14) (N/A , 0.01 , -0.1)	1012.8 176.6	0.1120 128.0 128.8	0.1967 [0.1885]	104.3%			
9Cl-Pf3ONS	(531.0 / 351.0) 590275 (533.0 / 353.0) 184727	(8.60 , 1.51) (N/A , 0.00 , -0.3)	348.7 361.1	0.3130 106.2 100.5	0.2188 [0.1867]	117.2%			
11Cl-PF3OUDS	(631.0 / 451.0) 254185 (633.0 / 453.0) 91563	(9.11 , 1.60) (N/A , 0.01 , -0.1)	663.7 3585.6	0.3602 116.7 116.4	0.1916 [0.1886]	101.6%			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) 4898 (241.0 / 117.0) 8827	(4.00 , 0.90) (N/A , 0.00 , 0.4)	142.3 52.6	1.8022 103.9 109.6	0.4006 [0.4000]	100.2%			
5:3FTCA	(341.0 / 236.7) 34995 (341.0 / 217.0) 43734	(5.90 , 1.10) (N/A , 0.02 , -0.1)	293.5 186.4	1.2497 74.3 68.8	0.4291 [0.4000]	107.3%			
7:3FTCA	(441.0 / 317.0) 40991 (441.0 / 337.0) 49859	(7.77 , 1.44) (N/A , 0.00 , 0.3)	724.4 36789.6	1.2163 141.6 144.9	0.2796 [0.4000]	69.9%			QC,
PFEESA	(315.0 / 135.0) 140320 (315.0 / 83.0) 38382	(5.77 , 1.07) (N/A , 0.02 , 0.3)	771.5 148.7	0.2735 97.2 104.5	0.1979 [0.1785]	110.9%			
PFMPA	(229.0 / 85.0) 20134	(3.83 , 0.86) (N/A , 0.00 , 0.0)	316.8	N/A 0.0 0.0	0.1902 [0.2000]	95.1%			
PFMBA	(279.0 / 85.0) 82321	(4.75 , 1.07) (N/A , -0.01 , 0.0)	357.4	N/A 0.0 0.0	0.2270 [0.2000]	113.5%			
NFDHA	(295.0 / 201.0) 57504 (295.0 / 85.0) 65515	(5.28 , 0.98) (N/A , 0.00 , -0.1)	1143.2 666.0	1.1393 109.1 109.8	0.1812 [0.2000]	90.6%			
13C3_PFBA_IIS	(216.0 / 172.0) 242191	(3.45 , N/A) (N/A , -0.01 , N/A)	1224.1	N/A	0.7144 [1.0000]	71.4% { 115.6% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 456742	(5.38 , N/A) (N/A , 0.01 , N/A)	1214.2	N/A	0.7679 [1.0000]	76.8% { 122.6% }			
13C4_PFOA_IIS	(417.0 / 372.0) 557666	(6.88 , N/A) (N/A , 0.01 , N/A)	947.2	N/A	0.6856 [1.0000]	68.6% { 107.6% }			
13C5_PFNA_IIS	(468.0 / 423.0) 581892	(7.49 , N/A) (N/A , 0.01 , N/A)	1369.8	N/A	0.7620 [1.0000]	76.2% { 105.6% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 547211	(8.04, N/A) (N/A, 0.01, N/A)	1051.2	N/A	0.7623 [1.0000]	76.2% { 114.3% }			
18O2_PFHxS_IIS	(403.0 / 83.9) 875338	(7.01, N/A) (N/A, 0.01, N/A)	1211.8	N/A	0.7108 [1.0000]	71.1% { 98.9% }			
13C4_PFOS_IIS	(503.0 / 79.9) 1452401	(8.23, N/A) (N/A, 0.01, N/A)	897.5	N/A	0.8183 [1.0000]	81.8% { 112.6% }			
13C4_PFBA_EIS	(217.0 / 172.0) 2180079	(3.44, N/A) (N/A, -0.01, N/A)	4281.9	N/A	7.5701 [8.0000]	94.6% { 114.2% }			
13C5_PFPeA_EIS	(268.0 / 223.0) 1879238	(4.44, N/A) (N/A, -0.01, N/A)	1919.9	N/A	3.6926 [4.0000]	92.3% { 107.2% }			
13C5_PFHxA_EIS	(318.0 / 273.0) 1222707	(5.38, N/A) (N/A, 0.01, N/A)	1437.7	N/A	1.9702 [2.0000]	98.5% { 115.2% }			
13C4_PFHpA_EIS	(367.0 / 322.0) 1096410	(6.18, N/A) (N/A, 0.02, N/A)	1819.2	N/A	1.8943 [2.0000]	94.7% { 121.6% }			
13C8_PFOA_EIS	(421.0 / 376.0) 1178173	(6.88, N/A) (N/A, 0.02, N/A)	1692.8	N/A	1.9739 [2.0000]	98.7% { 111.0% }			
13C9_PFNA_EIS	(472.0 / 427.0) 533109	(7.48, N/A) (N/A, 0.00, N/A)	903.4	N/A	0.9194 [1.0000]	91.9% { 106.4% }			
13C6_PFDA_EIS	(519.0 / 474.0) 646825	(8.04, N/A) (N/A, 0.01, N/A)	1440.6	N/A	1.0213 [1.0000]	102.1% { 103.4% }			
13C7_PFUnA_EIS	(570.0 / 525.0) 667821	(8.54, N/A) (N/A, 0.01, N/A)	30684.5	N/A	1.0643 [1.0000]	106.4% { 113.4% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT Ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	(615.0 / 570.0) 579521	(8.85 , N/A) (N/A , 0.01 , N/A)	685.9	N/A	0.9958 [1.0000]	99.6% { 129.4% }			
13C2_PFTeDA_EIS	(715.0 / 670.0) 550497	(9.24 , N/A) (N/A , 0.01 , N/A)	1007.6	N/A	0.9100 [1.0000]	91.0% { 106.3% }			
13C3_PFBs_EIS	(302.0 / 80.0) 3432474	(5.33 , N/A) (N/A , 0.01 , N/A)	3113.6	N/A	1.9405 [2.0000]	97.0% { 96.2% }			
13C3_PFHxS_EIS	(402.0 / 80.0) 1835540	(7.01 , N/A) (N/A , 0.01 , N/A)	1367.7	N/A	1.9862 [2.0000]	99.3% { 104.8% }			
13C8_PFOS_EIS	(507.0 / 80.0) 3736414	(8.23 , N/A) (N/A , 0.00 , N/A)	1429.7	N/A	1.9582 [2.0000]	97.9% { 114.1% }			
13C2_4:2FTS_EIS	(329.0 / 81.0) 420817	(5.12 , N/A) (N/A , 0.00 , N/A)	1288.7	N/A	3.8841 [4.0000]	97.1% { 104.4% }			
13C2_6:2FTS_EIS	(429.0 / 81.0) 458780	(6.62 , N/A) (N/A , 0.02 , N/A)	1163.1	N/A	3.7558 [4.0000]	93.9% { 107.5% }			
13C2_8:2FTS_EIS	(529.0 / 81.0) 597458	(7.79 , N/A) (N/A , 0.01 , N/A)	548.0	N/A	4.6252 [4.0000]	115.6% { 114.3% }			
13C8_PFOsa_EIS	(506.0 / 78.0) 6259920	(9.83 , N/A) (N/A , 0.01 , N/A)	3338.0	N/A	1.7810 [2.0000]	89.0% { 108.1% }			
D3_NMeFOSA_EIS	(515.0 / 169.0) 821331	(10.43 , N/A) (N/A , 0.01 , N/A)	1710.1	N/A	1.0882 [2.0000]	54.4% { 61.8% }			
D5_NEtFOSA_EIS	(531.0 / 169.0) 755770	(10.58 , N/A) (N/A , 0.01 , N/A)	1775.3	N/A	1.1436 [2.0000]	57.2% { 62.4% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [True] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1215548	(8.19 , N/A) (N/A , 0.01 , N/A)	1318.6	N/A	4.5392 [4.0000]	113.5% { 109.5% }			
D5_EtFOSAA_EIS	(589.0 / 419.0) 966077	(8.45 , N/A) (N/A , 0.01 , N/A)	90642.9	N/A	4.2365 [4.0000]	105.9% { 114.7% }			
D7_NMeFOSE_EIS	(623.0 / 58.9) 3008921	(10.36 , N/A) (N/A , 0.01 , N/A)	1257.7	N/A	11.7101 [20.0000]	58.6% { 70.4% }			
D9_NEtFOSE_EIS	(639.0 / 58.9) 4944456	(10.52 , N/A) (N/A , 0.01 , N/A)	1266.1	N/A	16.8224 [20.0000]	84.1% { 92.6% }			
13C3_HFPODA_EIS	(287.0 / 169.0) 3025088	(5.68 , N/A) (N/A , 0.02 , N/A)	2258.8	N/A	7.7575 [8.0000]	97.0% { 108.8% }			

PREPARATION BATCH SUMMARY

EPA 1633

Laboratory: APPL, LLC

Work Order: 23B0184

Client: AECOM

Project: Red Hill AFFF Assessment Sampling

Batch: BCB0459

Batch Matrix: Water

Preparation: EPA 1633

SAMPLE NAME	LAB SAMPLE ID	DATE PREPARED	INITIAL VOL./WEIGHT mL	FINAL VOL. mL
AF-RHMW03-WGN01LF-2302W3	23B0184-01	02/28/23 08:33	527.60	2.00
AF-RHMW03-WGN01LF-2302W3	23B0184-01RE1	02/28/23 08:33	527.60	2.00
AF-RHMW02-WGN01LF-2302W3	23B0184-02	02/28/23 08:33	502.93	2.00
AF-RHMW02-WGN01LF-2302W3	23B0184-02RE1	02/28/23 08:33	502.93	2.00
Blank	BCB0459-BLK1	02/28/23 08:24	500.00	2.00
LCS	BCB0459-BS1	02/28/23 08:24	500.00	2.00
MRL Check	BCB0459-MRL1	02/28/23 08:24	500.00	2.00

PREPARATION BENCH SHEET

Organics

BCB0459

Print Date/Time: 03/03/2023 5:57 pm

Matrix: Water

Prepared using: PFAS - EPA 1633

Analyses		Spiking Solution(s)				Surrogate Solution(s)			
1633		23B0578 PFAS - MIX 1633 10ng/mL				23B0575 MPFAC-HIF-ES 20.0ng/mL			
Lab Number	Sample and Source ID	Date Due	Extract by	Prepared	Initial (mL)	Final (mL)	ul Spike	ul Surrogate	Extraction Comments
23B0184-01	AF-RHWW03-WGN01LF-2302W3	03/03/2023	03/23/2023	2/28/2023 8:33:00AM	527.6	2		200	"Report relevant surrogates"
23B0184-01RE1	AF-RHWW03-WGN01LF-2302W3	03/03/2023	03/23/2023	2/28/2023 8:33:00AM	527.6	2		200	"Report relevant surrogates"
23B0184-02	AF-RHWW02-WGN01LF-2302W3	03/03/2023	03/23/2023	2/28/2023 8:33:00AM	502.93	2		200	"Report relevant surrogates"
23B0184-02RE1	AF-RHWW02-WGN01LF-2302W3	03/03/2023	03/23/2023	2/28/2023 8:33:00AM	502.93	2		200	"Report relevant surrogates"
23B0200-01	AF-RHWW17S-WGN01B-2302W3	03/07/2023	03/24/2023	2/28/2023 8:33:00AM	556.68	2		200	"Report relevant surrogates"
23B0200-01RE1	AF-RHWW17S-WGN01B-2302W3	03/07/2023	03/24/2023	2/28/2023 8:33:00AM	556.68	2		200	"Report relevant surrogates"
23B0200-02	AF-RHWW17D-WGN01LF-2302W3	03/07/2023	03/24/2023	2/28/2023 8:33:00AM	553.82	2		200	"Report relevant surrogates"
23B0200-02RE1	AF-RHWW17D-WGN01LF-2302W3	03/07/2023	03/24/2023	2/28/2023 8:33:00AM	553.82	2		200	"Report relevant surrogates"
23B0200-03	AF-RHWW17D-WQFB01-2302W3	03/07/2023	03/24/2023	2/28/2023 8:33:00AM	556.11	2		200	"Report relevant surrogates"
23B0200-03RE1	AF-RHWW17D-WQFB01-2302W3	03/07/2023	03/24/2023	2/28/2023 8:33:00AM	556.11	2		200	"Report relevant surrogates"
23B0200-04	AF-RHWW17-WGN01LF-2302W3	03/07/2023	03/24/2023	2/28/2023 8:33:00AM	563.12	2		200	"Report relevant surrogates"
23B0200-04RE1	AF-RHWW17-WGN01LF-2302W3	03/07/2023	03/24/2023	2/28/2023 8:33:00AM	563.12	2		200	"Report relevant surrogates"
BCB0459-BLK1	Blank			2/28/2023 8:24:00AM	500	2	0	200	
BCB0459-BS1	LCS			2/28/2023 8:24:00AM	500	2	200	200	
BCB0459-MRL1	MRL Check			2/28/2023 8:24:00AM	500	2	20	200	

PREPARATION BENCH SHEET

Organics

BCB0459

(Continued)

Matrix: Water

Analyses
1633

Spiking Solution(s)
23B0578 PFAS - MIX 1633 10ng/mL

Start Date/Time	
StopDate/Time	

Batch Comments:

Spiked by: PAF 2/28/23 3:05pm
Witness: DAG
Balance #: WB-2
Cartridge: Oasis
Concentration: 3/1/23 1:35pm - 3:35pm

Prepared using: PFAS - EPA 1633

Surrogate Solution(s)		
23B0575	MPFAC-HIF-ES 20.0ng/mL	
Reagents		
<u>Standard</u>	<u>Description</u>	<u>LotNum</u>
22C0296	Envi-carb	122395
23B0048	Am. Ac. preservative	*
23B0116	Reagent -0.3M Formic Acid	M13H051
23B0118	Reagent - 0.05MFA wash	x
23B0413	Reagent - 1.0% Ammonia Hydroxide	219481

Spiking Witnessed By	Preparation Reviewed By	Date

Extracts Received By	Date

INJECTION LOG - ANALYSIS SEQUENCE SUMMARY

EPA 1633

Laboratory: APPL, LLC

SDG:

Client: AECOM

Project: Red Hill AFFF Assessment Sampling

Sequence: SC00840

Instrument: Saphira

Calibration: 2309009

Sample Name	Lab Sample ID	Lab File ID	Analysis Date/Time
Cal Standard	SC00840-CAL1	S2023-02-28A (1)	02/28/23 15:52
Cal Standard	SC00840-CAL2	S2023-02-28A (2)	02/28/23 16:05
Cal Standard	SC00840-CAL3	S2023-02-28A (3)	02/28/23 16:18
Cal Standard	SC00840-CAL4	S2023-02-28A (4)	02/28/23 16:31
Cal Standard	SC00840-CAL5	S2023-02-28A (5)	02/28/23 16:44
Cal Standard	SC00840-CAL6	S2023-02-28A (6)	02/28/23 16:56
Cal Standard	SC00840-CAL7	S2023-02-28A (7)	02/28/23 17:09
Cal Standard	SC00840-CAL8	S2023-02-28A (8)	02/28/23 17:22
Initial Cal Blank	SC00840-ICB1	S2023-02-28A (9)	02/28/23 17:35
Secondary Cal Check	SC00840-SCV1	S2023-02-28A (10)	02/28/23 17:48

INJECTION LOG - ANALYSIS SEQUENCE SUMMARY

EPA 1633

Laboratory: APPL, LLC

SDG:

Client: AECOM

Project: Red Hill AFFF Assessment Sampling

Sequence: SC00861

Instrument: Saphira

Calibration: 2309009

Sample Name	Lab Sample ID	Lab File ID	Analysis Date/Time
Calibration Blank	SC00861-CCB1	S2023-03-02A (1)	03/02/23 09:42
Low Cal Check	SC00861-LCV1	S2023-03-02A (2)	03/02/23 09:55
Calibration Check	SC00861-CCV1	S2023-03-02A (3)	03/02/23 10:07
Performance Mix	SC00861-PEM1	S2023-03-02A (4)	03/02/23 10:20
Performance Mix	SC00861-PEM2	S2023-03-02A (5)	03/02/23 10:33
Calibration Blank	SC00861-CCB2	S2023-03-02A (6)	03/02/23 10:46
Blank	BCB0459-BLK1	S2023-03-02A (7)	03/02/23 10:59
LCS	BCB0459-BS1	S2023-03-02A (8)	03/02/23 11:12
MRL Check	BCB0459-MRL1	S2023-03-02A (9)	03/02/23 11:25
AF-RHMW03-WGN01LF-2302W3	23B0184-01	S2023-03-02A (10)	03/02/23 11:38
AF-RHMW03-WGN01LF-2302W3	23B0184-01RE1	S2023-03-02A (11)	03/02/23 11:51
AF-RHMW02-WGN01LF-2302W3	23B0184-02	S2023-03-02A (12)	03/02/23 12:03
AF-RHMW02-WGN01LF-2302W3	23B0184-02RE1	S2023-03-02A (13)	03/02/23 12:16
Calibration Check	SC00861-CCV2	S2023-03-02A (22)	03/02/23 14:12
Calibration Blank	SC00861-CCB3	S2023-03-02A (23)	03/02/23 14:25
Calibration Check	SC00861-CCV3	S2023-03-02A (40)	03/02/23 18:04
Calibration Blank	SC00861-CCB4	S2023-03-02A (41)	03/02/23 18:17
Calibration Check	SC00861-CCV4	S2023-03-02A (52)	03/02/23 20:39
Calibration Blank	SC00861-CCB5	S2023-03-02A (53)	03/02/23 20:52
Calibration Check	SC00861-CCV5	S2023-03-02A (55)	03/02/23 21:18
Calibration Blank	SC00861-CCB6	S2023-03-02A (56)	03/02/23 21:31



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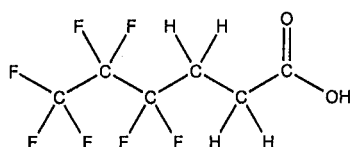
CERTIFICATE OF ANALYSIS DOCUMENTATION

PRODUCT CODE: FPrPA
COMPOUND: 3-Perfluoropropyl propanoic acid

LOT NUMBER: FPrPA1020

STRUCTURE:

CAS #: 356-02-5



MOLECULAR FORMULA: $C_8H_5F_7O_2$
CONCENTRATION: $50.0 \pm 2.5 \mu\text{g/mL}$
CHEMICAL PURITY: >98%
LAST TESTED: (mm/dd/yyyy) 11/12/2020
EXPIRY DATE: (mm/dd/yyyy) 11/12/2025
RECOMMENDED STORAGE: Refrigerate ampoule

MOLECULAR WEIGHT: 242.09
SOLVENT(S): Methanol

DOCUMENTATION/ DATA ATTACHED:

Figure 1: LC/MS Data (TIC and Mass Spectrum)
Figure 2: LC/MS/MS Data (Selected MRM Transitions)

ADDITIONAL INFORMATION:

- See page 2 for further details.
- Contains <1% of the unsaturated 3:3 telomer acid ($C_8H_3F_7O_2$) as an impurity determined by ^{19}F NMR.

FOR LABORATORY USE ONLY: NOT FOR HUMAN OR DRUG USE

Certified By: 
B.G. Chittim, General Manager

Date: 11/27/2020
(mm/dd/yyyy)

Wellington Laboratories Inc., 345 Southgate Dr. Guelph ON N1G 3M5 CANADA
519-822-2436 • Fax: 519-822-2849 • info@well-labs.com

INTENDED USE:

The products prepared by Wellington Laboratories Inc. are for laboratory use only. This certified reference material (CRM) was designed to be used as a standard for the identification and/or quantification of the specific chemical compound it contains.

HANDLING:

This product should only be used by qualified personnel familiar with its potential hazards and trained in the handling of hazardous chemicals. Due care should be exercised to prevent unnecessary human contact or ingestion. All procedures should be carried out in a well-functioning fume hood and suitable gloves, eye protection, and clothing should be worn at all times. Waste should be disposed of according to national and regional regulations. Safety Data Sheets (SDSs) are available upon request.

SYNTHESIS / CHARACTERIZATION:

Our products are synthesized using single-product unambiguous routes whenever possible. They are then characterized, and their structures and purities confirmed, using a combination of the most relevant techniques, such as NMR, GC/MS, LC/MS/MS, SFC/UV/MS/MS, x-ray crystallography, and melting point. Isotopic purities of mass-labelled compounds are also confirmed using HRGC/HRMS and/or LC/MS/MS.

HOMOGENEITY:

Prior to solution preparation, crystalline material is tested for homogeneity using a variety of techniques (as stated above) and its solubility in a given diluent is taken into consideration. Duplicate solutions of a new product are prepared from the same crystalline lot and, after the addition of an appropriate internal standard, they are compared by GC/MS, LC/MS/MS, and/or SFC/UV/MS/MS. The relative response factors of the analyte of interest in each solution are required to be <5% RSD. New solution lots of existing products are compared to older lots in the same manner, which further confirms the homogeneity of the crystalline material as well as the stability and homogeneity of the solutions in the storage containers. In order to maintain the integrity of the assigned value(s), and associated uncertainty, the dilution or injection of a subsample of this product should be performed using calibrated measuring equipment.

UNCERTAINTY:

The maximum combined relative standard uncertainty of our reference standard solutions is calculated using the following equation:

The combined relative standard uncertainty, $u_c(y)$, of a value y and the uncertainty of the independent parameters

x_1, x_2, \dots, x_n on which it depends is:

$$u_c(y(x_1, x_2, \dots, x_n)) = \sqrt{\sum_{i=1}^n u(y, x_i)^2}$$

where x is expressed as a relative standard uncertainty of the individual parameter.

The individual uncertainties taken into account include those associated with weights (calibration of the balance) and volumes (calibration of the volumetric glassware). An expanded maximum combined percent relative uncertainty of $\pm 5\%$ (calculated with a coverage factor of 2 and a level of confidence of 95%) is stated on the Certificate of Analysis for all of our products.

TRACEABILITY:

All reference standard solutions are traceable to specific crystalline lots. The microbalances used for solution preparation are regularly calibrated by an external ISO/IEC 17025 accredited laboratory. In addition, their calibration is verified prior to each weighing using calibrated external weights traceable to an ISO/IEC 17025 accredited laboratory. All volumetric glassware used is calibrated, of Class A tolerance, and traceable to an ISO/IEC 17025 accredited laboratory. For certain products, traceability to international interlaboratory studies has also been established.

EXPIRY DATE / PERIOD OF VALIDITY:

Ongoing stability studies of this product have demonstrated stability in its composition and concentration, until the specified expiry date, in the unopened ampoule. Monitoring for any degradation or change in concentration of the listed analyte(s) is performed on a routine basis.

LIMITED WARRANTY:

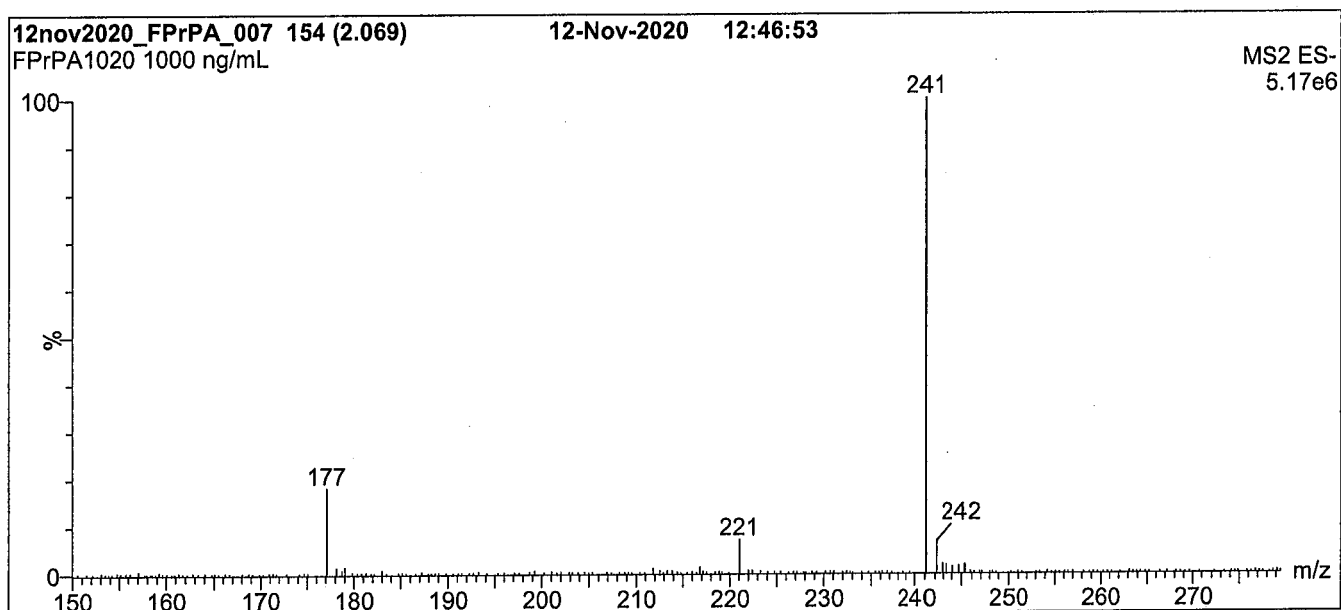
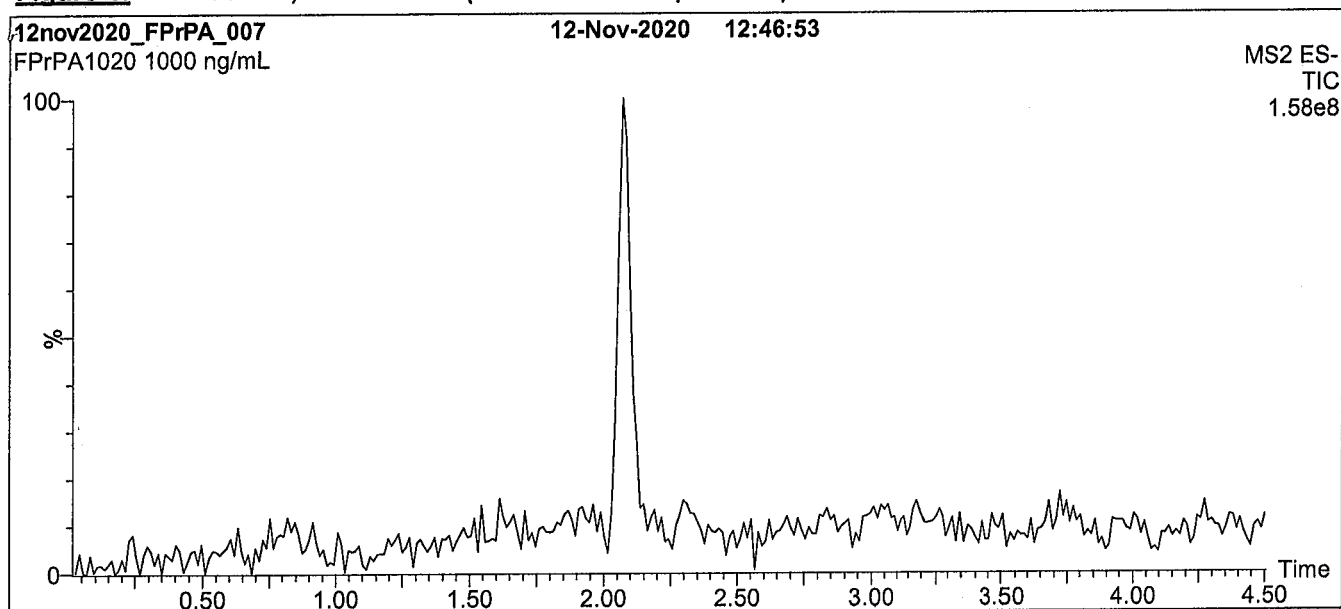
At the time of shipment, all products are warranted to be free of defects in material and workmanship and to conform to the stated technical and purity specifications.

QUALITY MANAGEMENT:

This product was produced using a Quality Management System registered to the latest versions of ISO 9001 by SAI Global, ISO/IEC 17025 by the Canadian Association for Laboratory Accreditation Inc. (CALA; A1226), and ISO 17034 by ANSI-ASQ National Accreditation Board (ANAB; AR-1523).



For additional information or assistance concerning this or any other products from Wellington Laboratories Inc., please visit our website at www.well-labs.com or contact us directly at info@well-labs.com

Figure 1: FPrPA; LC/MS Data (TIC and Mass Spectrum)**Conditions for Figure 1:**

Waters Acquity Ultra Performance LC
Waters Xevo TQ-S micro MS

Chromatographic Conditions:

Column: Acquity UPLC BEH Shield RP₁₈
1.7 μ m, 2.1 x 100 mm

Mobile phase: Gradient

Start: 60% H₂O / 40% (80:20 MeOH:ACN)
(both with 10 mM NH₄OAc buffer)
Ramp to 90% organic over 8 min and hold for 2 min
before returning to initial conditions in 0.75 min.
Time: 12 min

Flow: 300 μ L/min

MS Parameters:

Experiment: Full Scan (150 - 850 amu)

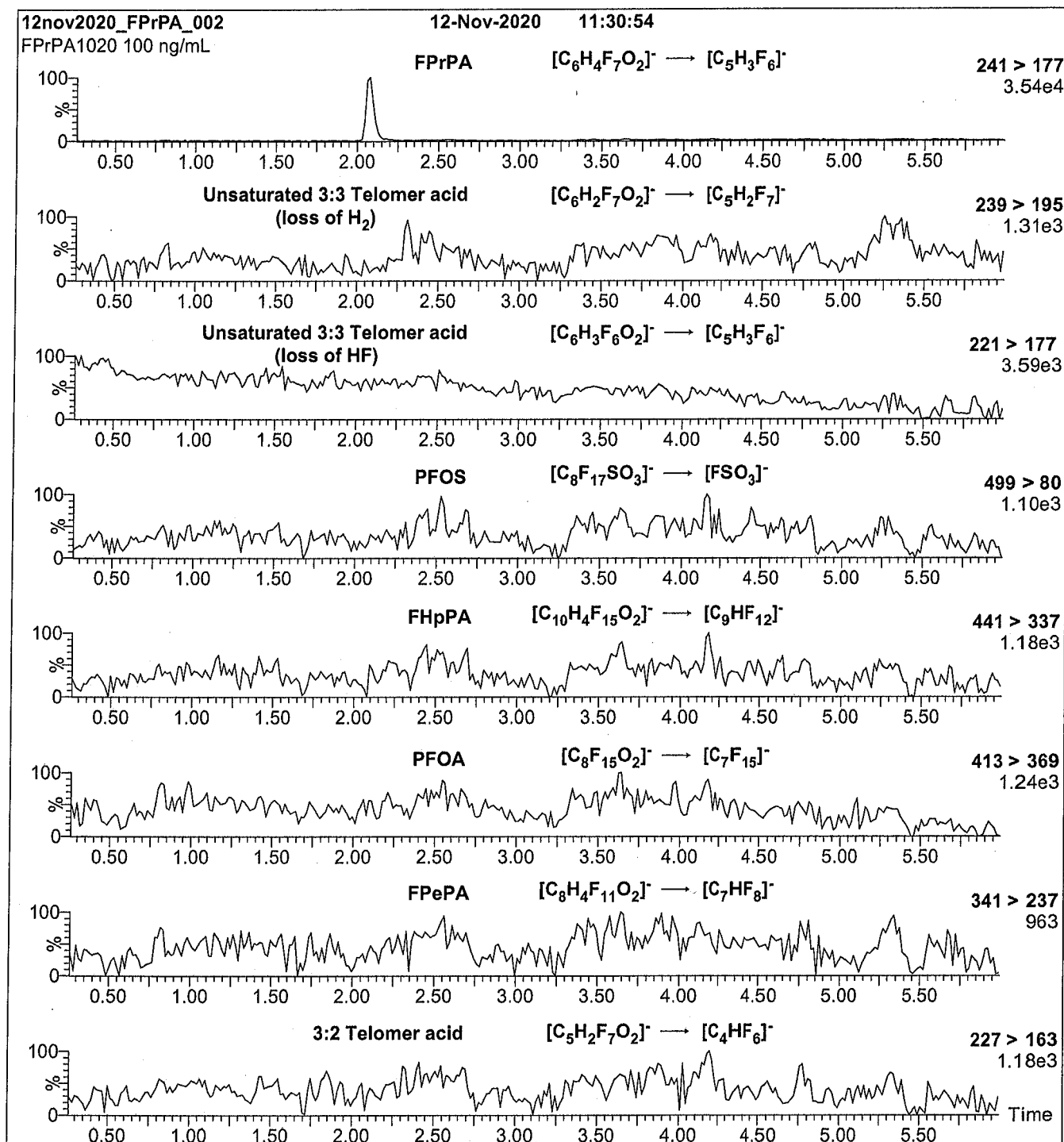
Source: Electrospray (negative)

Capillary Voltage (kV) = 0.50

Cone Voltage (V) = 18.50

Desolvation Temperature (°C) = 500

Desolvation Gas Flow (L/hr) = 1000

Figure 2: FPrPA; LC/MS/MS Data (Selected MRM Transitions)**Conditions for Figure 2:**

Injection: On-column (FPrPA)

Mobile phase: Same as Figure 1

Flow: 300 μ L/min**MS Parameters:**

Collision Gas (mbar) = 3.49e-3

Collision Energy (eV) = 10

Analytical Standard Record

21L0004

Description: PFAS - SAS 3:3FTA 50ug/mL
Standard Type: Analyte Spike
Solvent: MeOH
Final Volume (mLs): 1
Vials: 1
Comments: 3:3 FTCA 50.0ug/mL

Expires: 06/05/2022
Prepared: 12/07/2021
Prepared By: Hart Hedgpeth
Department: PFAS
Last Edit: 12/07/2021 16:03 by HGH

Analyte	Parent	CAS Number	Concentration	Units
3:3 FTA		113507-82-7	50	ug/mL



WELLINGTON LABORATORIES

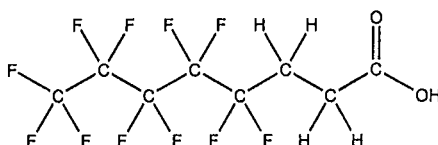
CERTIFICATE OF ANALYSIS DOCUMENTATION

PRODUCT CODE: FPePA
COMPOUND: 3-Perfluoropentyl propanoic acid

LOT NUMBER: FPePA1120

STRUCTURE:

CAS #: 914637-49-3



MOLECULAR FORMULA: $C_8H_5F_{11}O_2$
CONCENTRATION: $50.0 \pm 2.5 \mu\text{g/mL}$
CHEMICAL PURITY: >98%
LAST TESTED: (mm/dd/yyyy) 11/11/2020
EXPIRY DATE: (mm/dd/yyyy) 11/11/2025
RECOMMENDED STORAGE: Refrigerate ampoule

MOLECULAR WEIGHT: 342.11
SOLVENT(S): Methanol

DOCUMENTATION/ DATA ATTACHED:

Figure 1: LC/MS Data (TIC and Mass Spectrum)
Figure 2: LC/MS/MS Data (Selected MRM Transitions)

ADDITIONAL INFORMATION:

- See page 2 for further details.
- Contains <1% of the unsaturated 5:3 telomer acid ($C_8H_3F_{11}O_2$) as an impurity determined by ^{19}F NMR.

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Certified By: 
B.G. Chittim, General Manager

Date: 11/27/2020
(mm/dd/yyyy)

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where x is expressed as a relative standard uncertainty of the individual parameter.

The individual uncertainties taken into account include those associated with weights (calibration of the balance) and volumes (calibration of the volumetric glassware). An expanded maximum combined percent relative uncertainty of $\pm 5\%$ (calculated with a coverage factor of 2 and a level of confidence of 95%) is stated on the Certificate of Analysis for all of our products.

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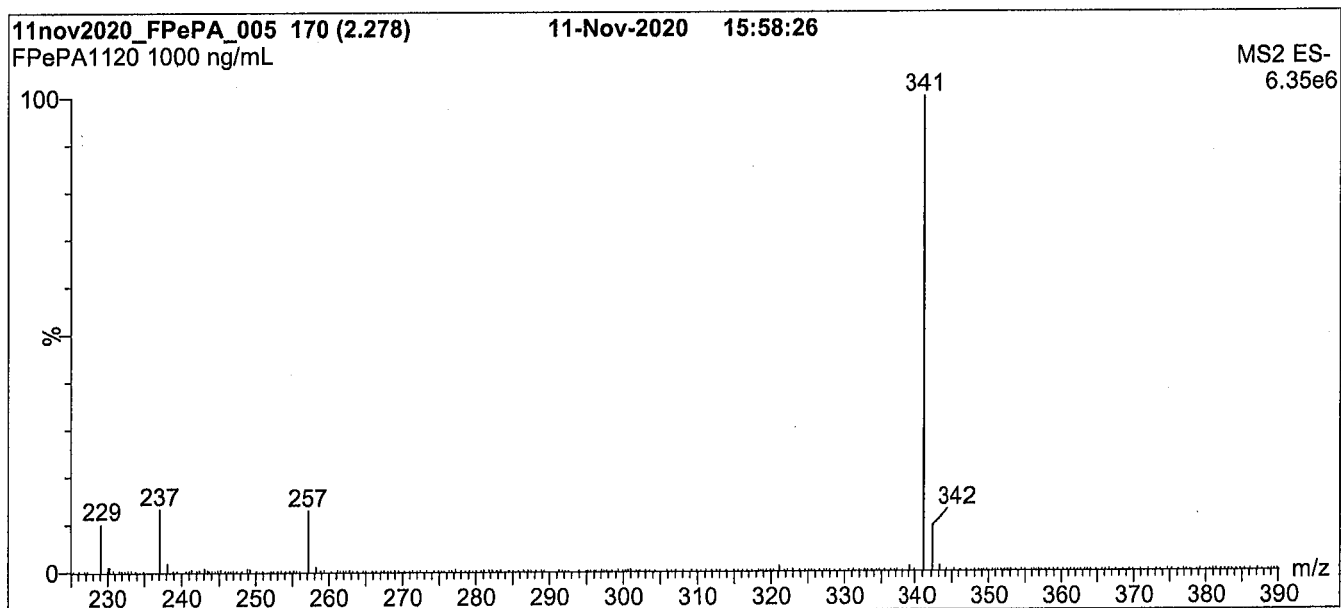
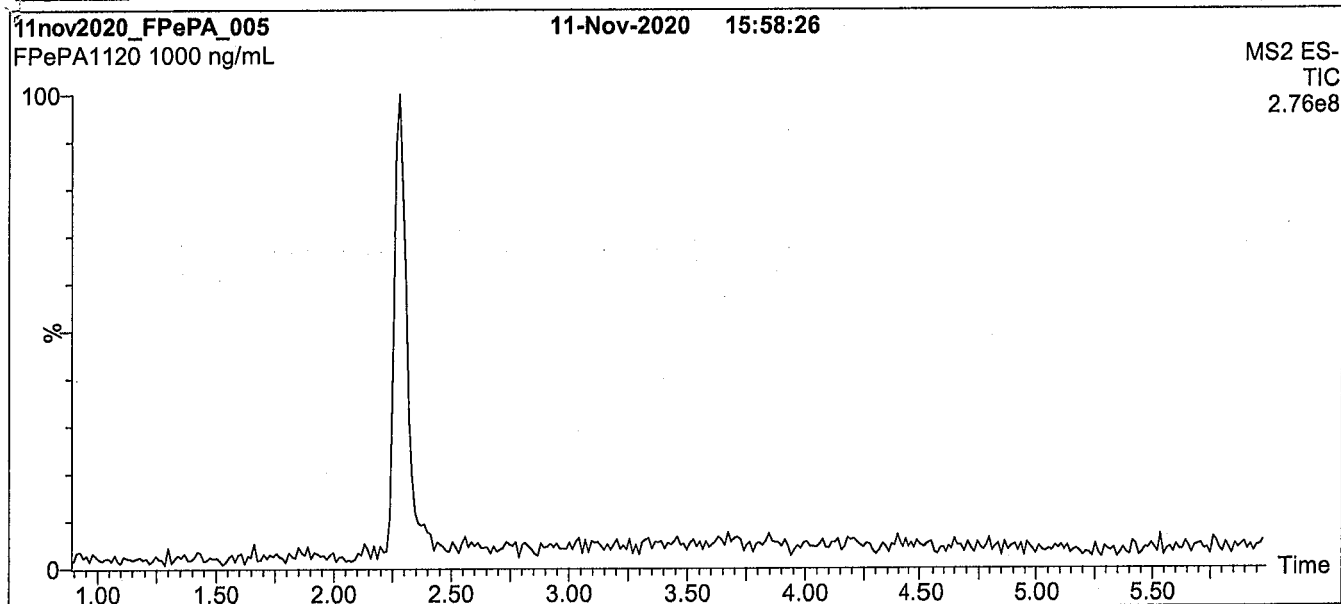
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Waters Xevo TQ-S micro MS

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1.7 μ m, 2.1 x 100 mm

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Start: 45% H₂O / 55% (80:20 MeOH:ACN)
(both with 10 mM NH₄OAc buffer)
Ramp to 90% organic over 8 min and hold for
2 min before returning to initial conditions in 0.75 min.
Time: 12 min

Flow: 300 μ L/min

MS Parameters:

Experiment: Full Scan (225 - 850 amu)

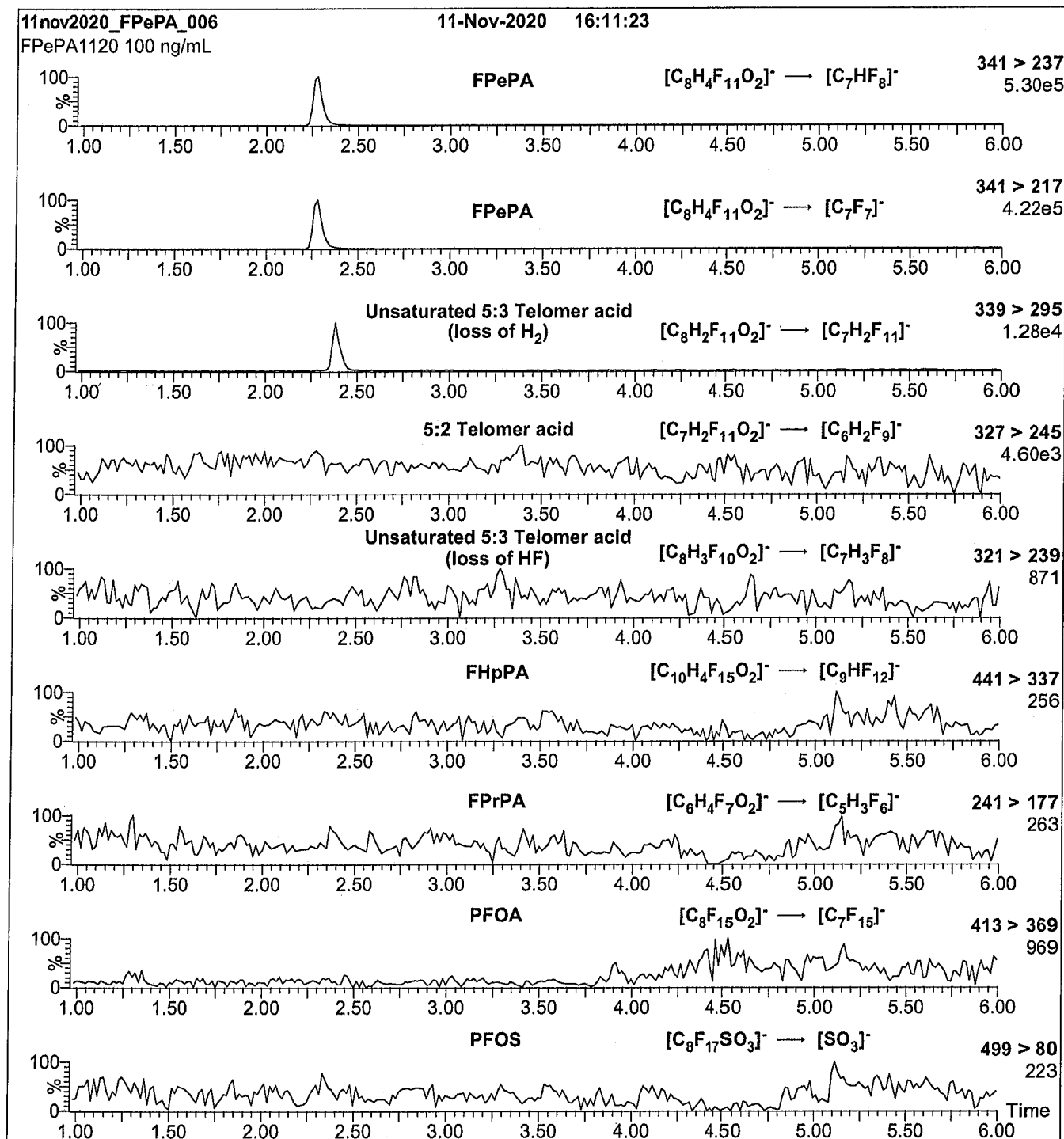
Source: Electrospray (negative)

Capillary Voltage (kV) = 0.50

Cone Voltage (V) = 18.50

Desolvation Temperature (°C) = 500

Desolvation Gas Flow (L/hr) = 1000

Figure 2: FPePA; LC/MS/MS Data (Selected MRM Transitions)**Conditions for Figure 2:**

Injection: On-column (FPePA)

Mobile phase: Same as Figure 1

Flow: 300 μ L/min**MS Parameters:**

Collision Gas (mbar) = 3.24e-3

Collision Energy (eV) = 10

Analytical Standard Record

21L0005

Description: PFAS - SAS 5:3FTA 50ug/mL
Standard Type: Analyte Spike
Solvent: MeOH
Final Volume (mLs): 1.2
Vials: 1
Comments: 5:3 FTCA 50.0ug/mL

Expires: 06/05/2022
Prepared: 12/07/2021
Prepared By: Hart Hedgpeth
Department: PFAS
Last Edit: 12/07/2021 16:03 by HGH

Analyte	Parent	CAS Number	Concentration	Units
5:3 FTA		914637-49-3	50	ug/mL



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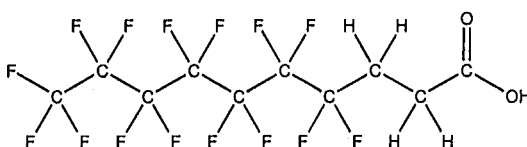
CERTIFICATE OF ANALYSIS DOCUMENTATION

PRODUCT CODE: FHpPA
COMPOUND: 3-Perfluoroheptyl propanoic acid

LOT NUMBER: FHpPA1020

STRUCTURE:

CAS #: 812-70-4



MOLECULAR FORMULA: $C_{10}H_{16}F_{15}O_2$
CONCENTRATION: $50.0 \pm 2.5 \mu\text{g/mL}$
CHEMICAL PURITY: >98%
LAST TESTED: (mm/dd/yyyy) 11/12/2020
EXPIRY DATE: (mm/dd/yyyy) 11/12/2025
RECOMMENDED STORAGE: Refrigerate ampoule

MOLECULAR WEIGHT: 442.12
SOLVENT(S): Methanol

DOCUMENTATION/ DATA ATTACHED:

Figure 1: LC/MS Data (TIC and Mass Spectrum)
Figure 2: LC/MS/MS Data (Selected MRM Transitions)

ADDITIONAL INFORMATION:

- See page 2 for further details.

FOR LABORATORY USE ONLY: NOT FOR HUMAN OR DRUG USE

Certified By: 
B.G. Chittim, General Manager

Date: 11/27/2020
(mm/dd/yyyy)

Wellington Laboratories Inc., 345 Southgate Dr. Guelph ON N1G 3M5 CANADA
519-822-2436 • Fax: 519-822-2849 • info@well-labs.com

INTENDED USE:

The products prepared by Wellington Laboratories Inc. are for laboratory use only. This certified reference material (CRM) was designed to be used as a standard for the identification and/or quantification of the specific chemical compound it contains.

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Prior to solution preparation, crystalline material is tested for homogeneity using a variety of techniques (as stated above) and its solubility in a given diluent is taken into consideration. Duplicate solutions of a new product are prepared from the same crystalline lot and, after the addition of an appropriate internal standard, they are compared by GC/MS, LC/MS/MS, and/or SFC/UV/MS/MS. The relative response factors of the analyte of interest in each solution are required to be <5% RSD. New solution lots of existing products are compared to older lots in the same manner, which further confirms the homogeneity of the crystalline material as well as the stability and homogeneity of the solutions in the storage containers. In order to maintain the integrity of the assigned value(s), and associated uncertainty, the dilution or injection of a subsample of this product should be performed using calibrated measuring equipment.

UNCERTAINTY:

The maximum combined relative standard uncertainty of our reference standard solutions is calculated using the following equation:

The combined relative standard uncertainty, $u_c(y)$, of a value y and the uncertainty of the independent parameters

x_1, x_2, \dots, x_n on which it depends is:

$$u_c(y(x_1, x_2, \dots, x_n)) = \sqrt{\sum_{i=1}^n u(y, x_i)^2}$$

where x is expressed as a relative standard uncertainty of the individual parameter.

The individual uncertainties taken into account include those associated with weights (calibration of the balance) and volumes (calibration of the volumetric glassware). An expanded maximum combined percent relative uncertainty of $\pm 5\%$ (calculated with a coverage factor of 2 and a level of confidence of 95%) is stated on the Certificate of Analysis for all of our products.

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LIMITED WARRANTY:

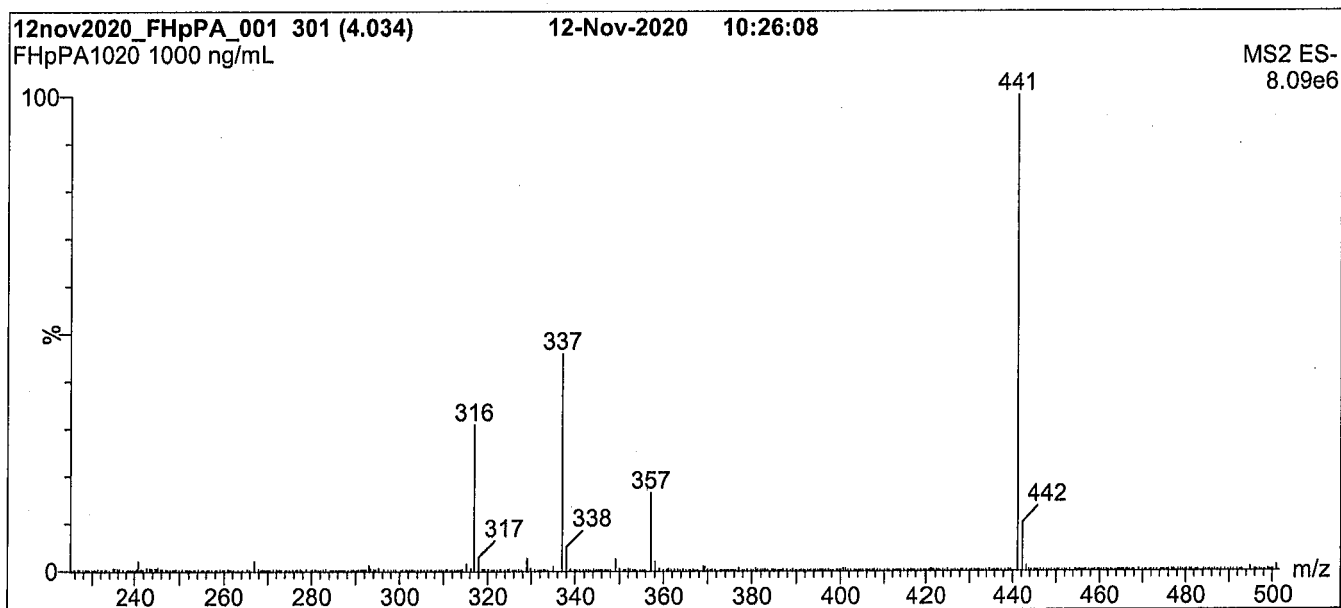
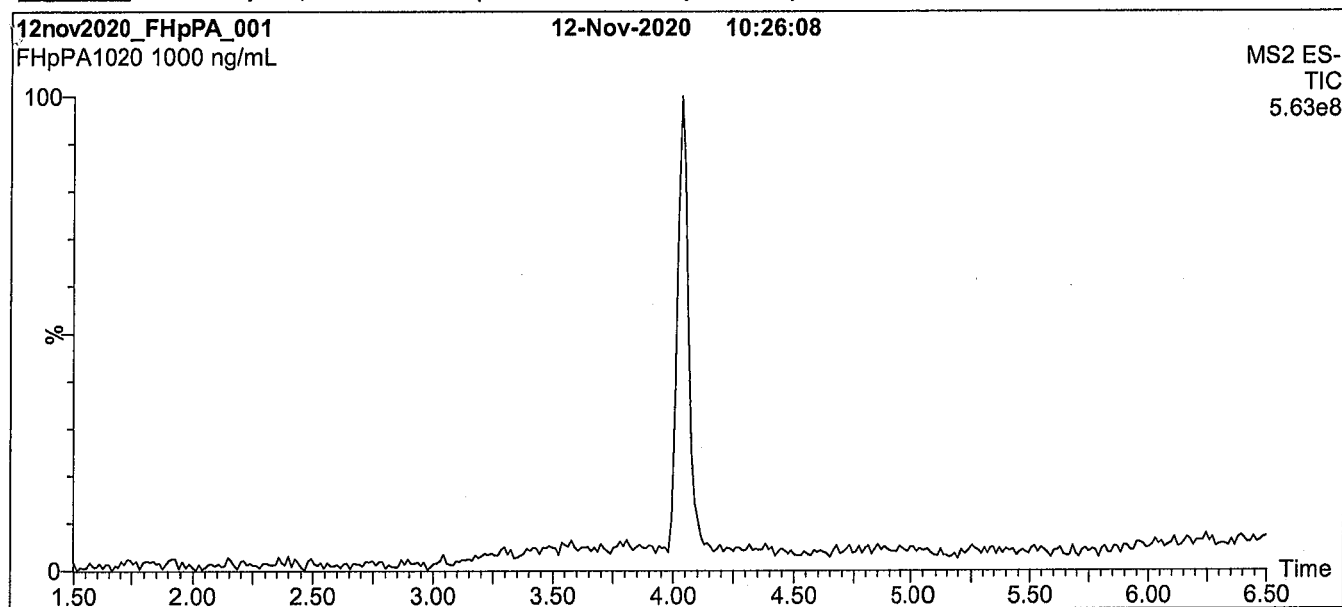
At the time of shipment, all products are warranted to be free of defects in material and workmanship and to conform to the stated technical and purity specifications.

QUALITY MANAGEMENT:

This product was produced using a Quality Management System registered to the latest versions of ISO 9001 by SAI Global, ISO/IEC 17025 by the Canadian Association for Laboratory Accreditation Inc. (CALA; A1226), and ISO 17034 by ANSI-ASQ National Accreditation Board (ANAB; AR-1523).



For additional information or assistance concerning this or any other products from Wellington Laboratories Inc., please visit our website at www.well-labs.com or contact us directly at info@well-labs.com

Figure 1: FHpPA; LC/MS Data (TIC and Mass Spectrum)**Conditions for Figure 1:**

Waters Acquity Ultra Performance LC
Waters Xevo TQ-S micro MS

Chromatographic Conditions:

Column: Acquity UPLC BEH Shield RP₁₈
1.7 μ m, 2.1 x 100 mm

Mobile phase: Gradient

Start: 45% H₂O / 55% (80:20 MeOH:ACN)
(both with 10 mM NH₄OAc buffer)

Ramp to 90% organic over 8 min and hold for
2 min before returning to initial conditions in 0.75 min.

Time: 12 min

Flow: 300 μ L/min

MS Parameters:

Experiment: Full Scan (225 - 850 amu)

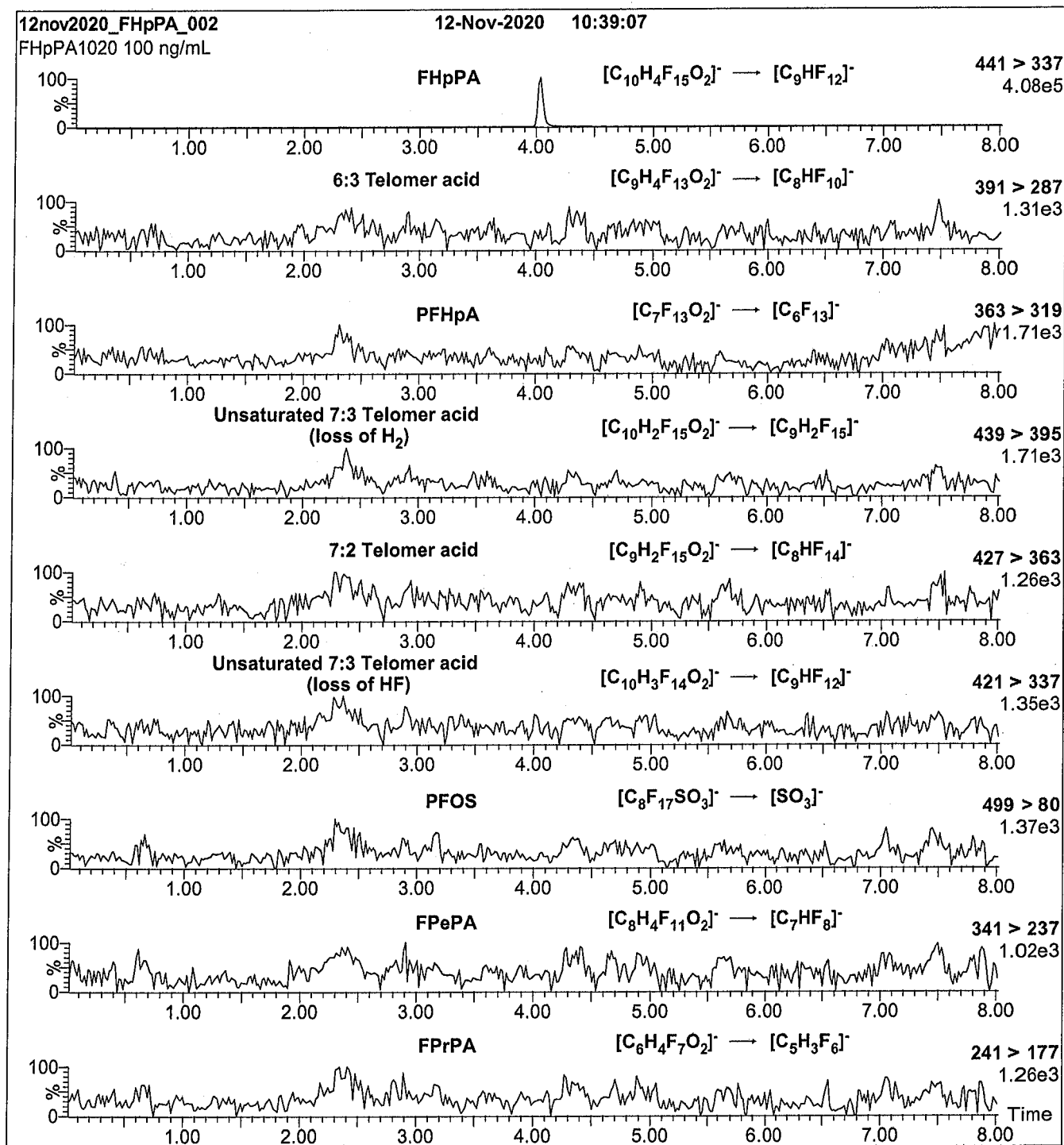
Source: Electrospray (negative)

Capillary Voltage (kV) = 0.50

Cone Voltage (V) = 28.50

Desolvation Temperature (°C) = 500

Desolvation Gas Flow (L/hr) = 1000

Figure 2: FHpPA; LC/MS/MS Data (Selected MRM Transitions)**Conditions for Figure 2:**

Injection: On-column (FHpPA)

Mobile phase: Same as Figure 1

Flow: 300 μ L/min**MS Parameters:**

Collision Gas (mbar) = 3.41e-3

Collision Energy (eV) = 8

Analytical Standard Record

21L0007

Description: PFAS - SAS 7:3FTA 50ug/mL
Standard Type: Analyte Spike
Solvent: MeOH
Final Volume (mLs): 1.2
Vials: 1
Comments: 7:3 FTCA 50.0ug/mL

Expires: 06/05/2022
Prepared: 12/07/2021
Prepared By: Hart Hedgpeth
Department: PFAS
Last Edit: 12/07/2021 16:16 by HGH

Analyte	Parent	CAS Number	Concentration	Units
7:3 FTA		812-70-4	50	ug/mL



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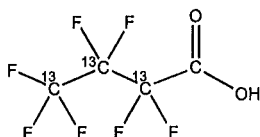
CERTIFICATE OF ANALYSIS DOCUMENTATION

PRODUCT CODE: M3PFBA
COMPOUND: Perfluoro-n-(2,3,4-¹³C₃)butanoic acid

LOT NUMBER: M3PFBA0721

STRUCTURE:

CAS #: Not available



MOLECULAR FORMULA: ¹³C₃¹²CHF₇O₂
CONCENTRATION: 50.0 ± 2.5 µg/mL

MOLECULAR WEIGHT: 217.02
SOLVENT(S): Methanol
Water (<1%)

CHEMICAL PURITY: >98%

ISOTOPIC PURITY: ≥99%¹³C
(2,3,4-¹³C₃)

LAST TESTED: (mm/dd/yyyy) 08/19/2021

EXPIRY DATE: (mm/dd/yyyy) 08/19/2026

RECOMMENDED STORAGE: Store ampoule in a cool, dark place

DOCUMENTATION/ DATA ATTACHED:

Figure 1: LC/MS Data (Full Scan and Mass Spectrum)


Figure 2: LC/MS/MS Data (Selected MRM Transitions)

ADDITIONAL INFORMATION:

- See page 2 for further details.
- Contains 4 mole eq. of NaOH to prevent conversion of the carboxylic acid to the methyl ester.
- Contains ~0.2% of perfluoro-n-(¹³C₃)propanoic acid and also contains ~1.0% of perfluoro-n-(1,2,3,4-¹³C₄)butanoic acid due to the naturally occurring isotopic abundance of ¹³C in the unlabelled carbon atom.

FOR LABORATORY USE ONLY: NOT FOR HUMAN OR DRUG USE

Certified By:


B.G. Chittim, General Manager

Date: 08/25/2021
(mm/dd/yyyy)

Wellington Laboratories Inc., 345 Southgate Dr. Guelph ON N1G 3M5 CANADA
519-822-2436 • Fax: 519-822-2849 • info@well-labs.com

INTENDED USE:

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SYNTHESIS / CHARACTERIZATION:

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UNCERTAINTY:

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x_1, x_2, \dots, x_n on which it depends is:

$$u_c(y(x_1, x_2, \dots, x_n)) = \sqrt{\sum_{i=1}^n u(y, x_i)^2}$$

where x is expressed as a relative standard uncertainty of the individual parameter.

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EXPIRY DATE / PERIOD OF VALIDITY:

Ongoing stability studies of this product have demonstrated stability in its composition and concentration, until the specified expiry date, in the unopened ampoule. Monitoring for any degradation or change in concentration of the listed analyte(s) is performed on a routine basis.

LIMITED WARRANTY:

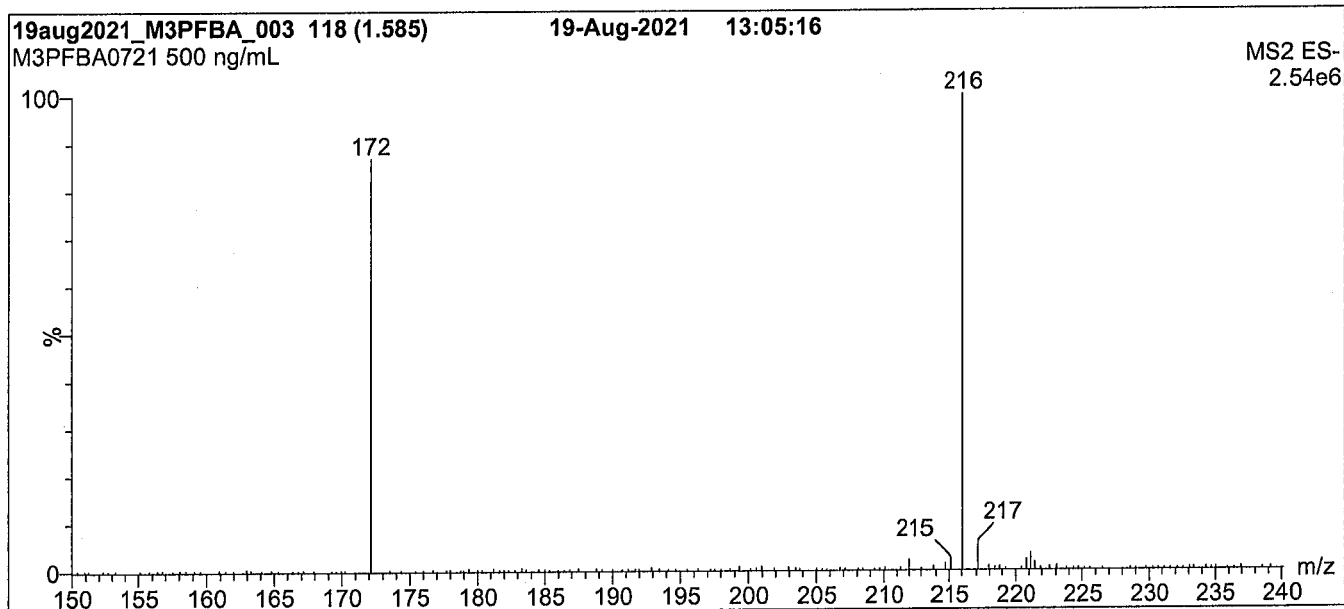
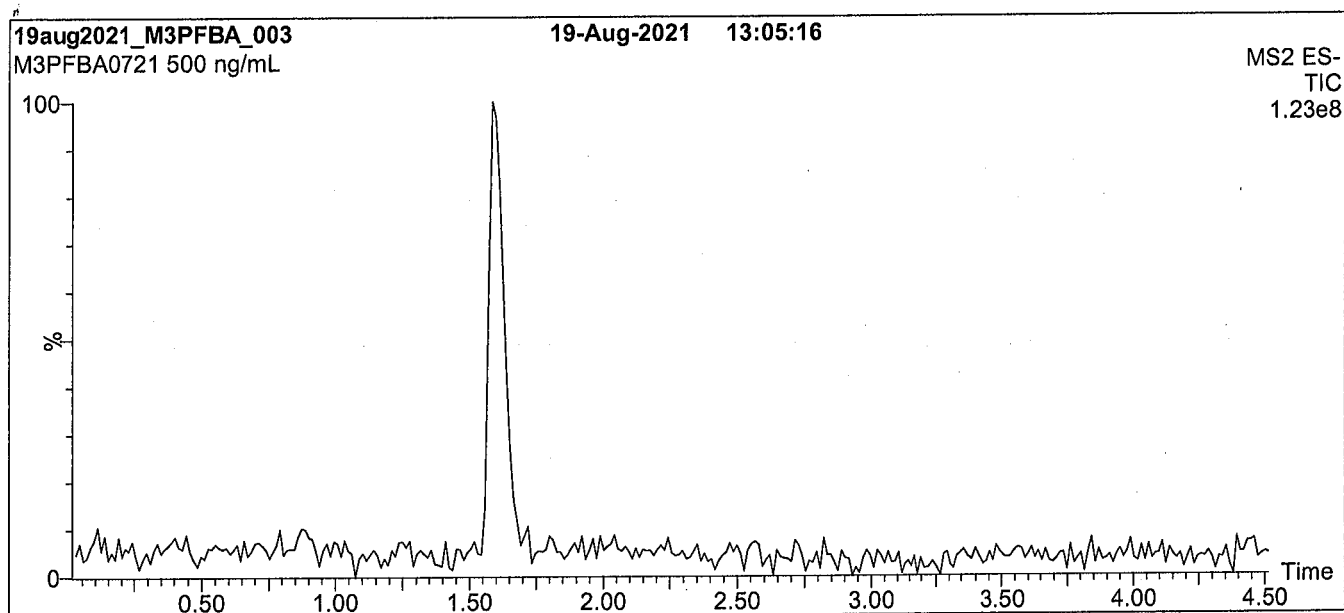
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Figure 1: M3PFBA; LC/MS Data (Full Scan and Mass Spectrum)**Conditions for Figure 1:**

Waters Acquity Ultra Performance LC
Waters Xevo TQ-S micro MS

Chromatographic Conditions:

Column: Acquity UPLC BEH Shield RP₁₈
1.7 μ m, 2.1 x 100 mm

Mobile phase: Gradient

Start: 60% H₂O / 40% (80:20 MeOH:ACN)
(both with 10 mM NH₄OAc buffer)
Ramp to 90% organic over 8 min and hold for
2 min before returning to initial conditions in 0.5 min.
Time: 12 min

Flow: 300 μ L/min

MS Parameters:

Experiment: Full Scan (150 - 850 amu)

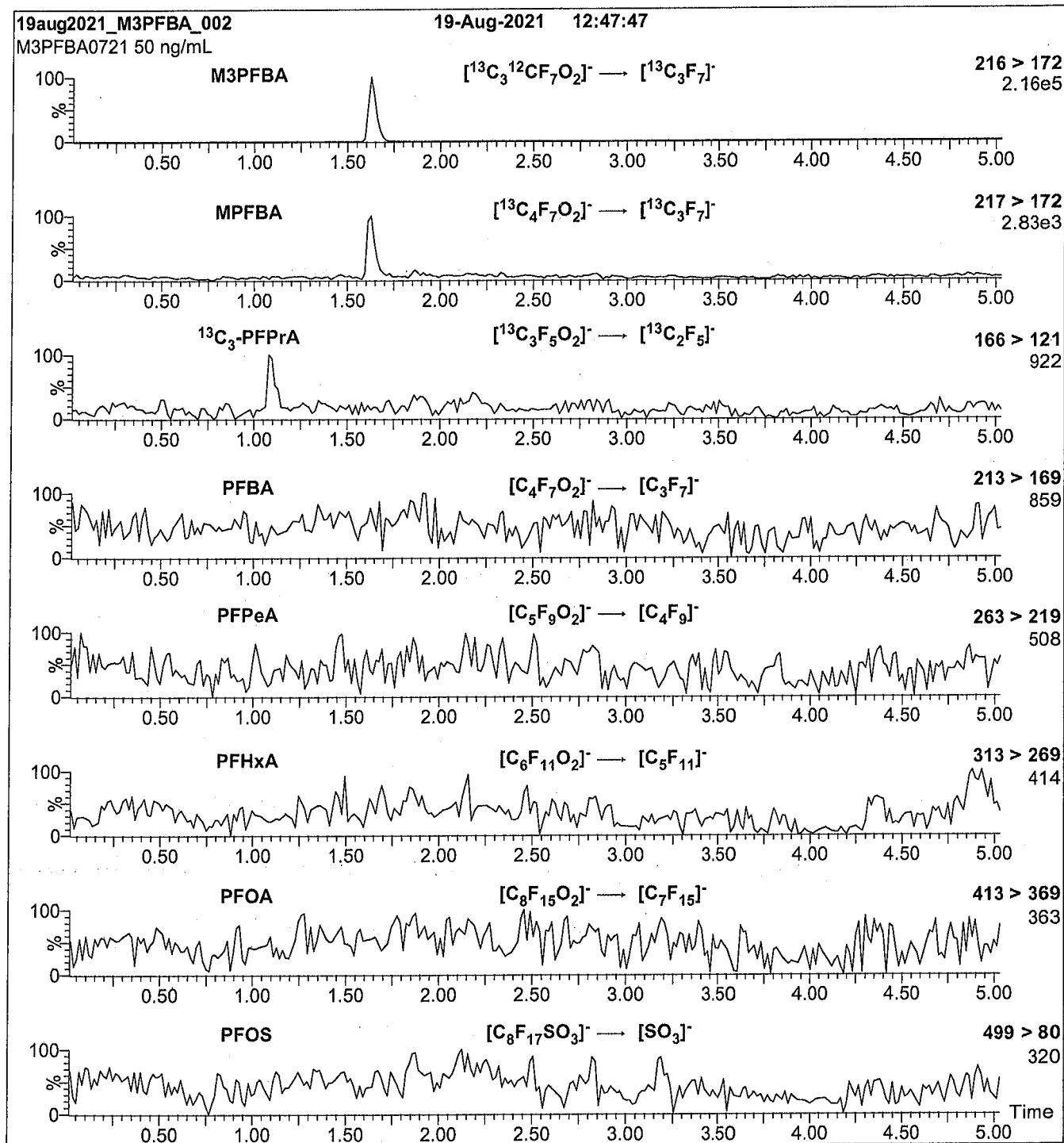
Source: Electrospray (negative)

Capillary Voltage (kV) = 2.00

Cone Voltage (V) = 10.00

Desolvation Temperature (°C) = 500

Desolvation Gas Flow (L/hr) = 1000

Figure 2: M3PFBA; LC/MS/MS Data (Selected MRM Transitions)**Conditions for Figure 2:**

Injection: On-column (M3PFBA)

Mobile phase: Same as Figure 1

Flow: 300 $\mu\text{L}/\text{min}$ **MS Parameters:**

Collision Gas (mbar) = 3.45e-3

Collision Energy (eV) = 8

Analytical Standard Record

22A0116

Description: PFAS - IIS M3PFBA 50ug/mL
Standard Type: Analyte Spike
Solvent: MeOH
Final Volume (mLs): 1.2
Vials: 1

Expires: 08/19/2026
Prepared: 08/19/2021
Prepared By: Dipti Gokal
Department: PFAS
Last Edit: 01/20/2022 15:48 by HGH

Analyte	Parent	CAS Number	Concentration	Units
13C3-PFBA		13C3-PFBA	50	ug/mL



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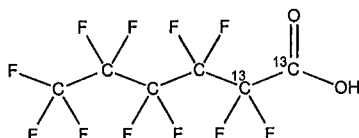
CERTIFICATE OF ANALYSIS DOCUMENTATION

PRODUCT CODE: MPFHxA
COMPOUND: Perfluoro-n-(1,2-¹³C₂)hexanoic acid

LOT NUMBER: MPFHxA0921

STRUCTURE:

CAS #: 960315-47-3



MOLECULAR FORMULA: ¹³C₂¹²C₄HF₁₁O₂
CONCENTRATION: 50.0 ± 2.5 µg/mL

MOLECULAR WEIGHT: 316.04
SOLVENT(S): Methanol
Water (<1%)

CHEMICAL PURITY: >98%

ISOTOPIC PURITY: ≥99% ¹³C
(1,2-¹³C₂)

LAST TESTED: (mm/dd/yyyy) 10/04/2021

EXPIRY DATE: (mm/dd/yyyy) 10/04/2026

RECOMMENDED STORAGE: Store ampoule in a cool, dark place

DOCUMENTATION/ DATA ATTACHED:

Figure 1: LC/MS Data (Full Scan and Mass Spectrum)

Figure 2: LC/MS/MS Data (Selected MRM Transitions)

ADDITIONAL INFORMATION:

- See page 2 for further details.
- Contains 4 mole eq. of NaOH to prevent conversion of the carboxylic acid to the methyl ester.

FOR LABORATORY USE ONLY: NOT FOR HUMAN OR DRUG USE

Certified By: 
B.G. Chittim, General Manager

Date: 10/22/2021
(mm/dd/yyyy)

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519-822-2436 • Fax: 519-822-2849 • info@well-labs.com

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where x is expressed as a relative standard uncertainty of the individual parameter.

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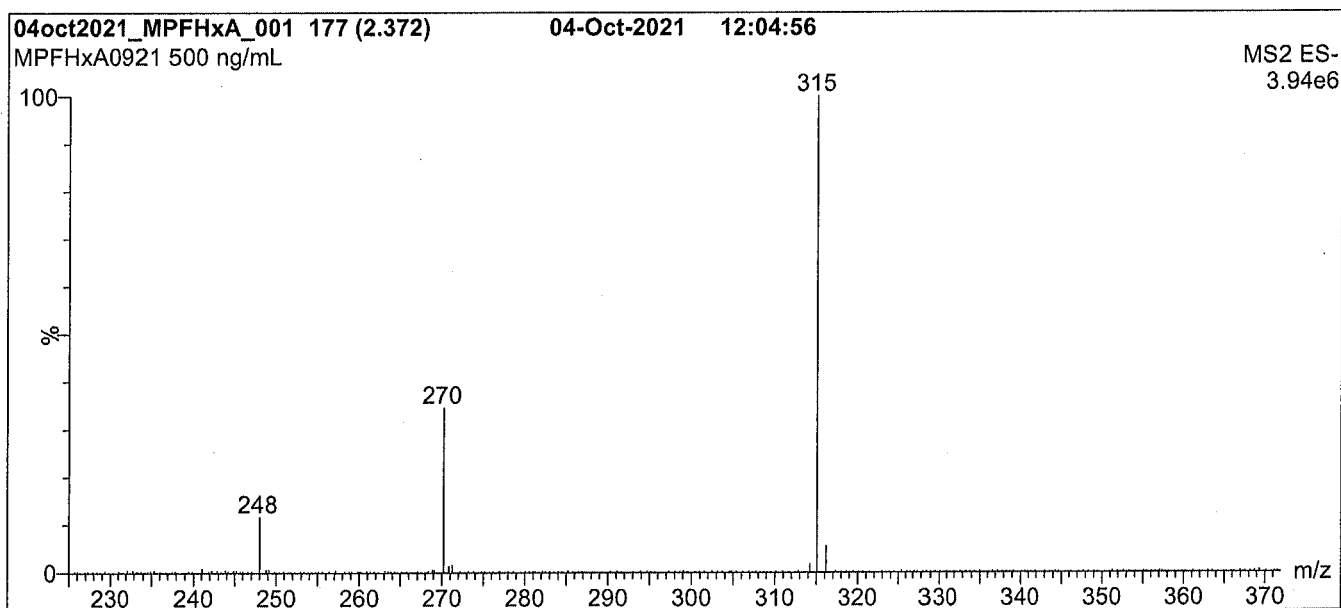
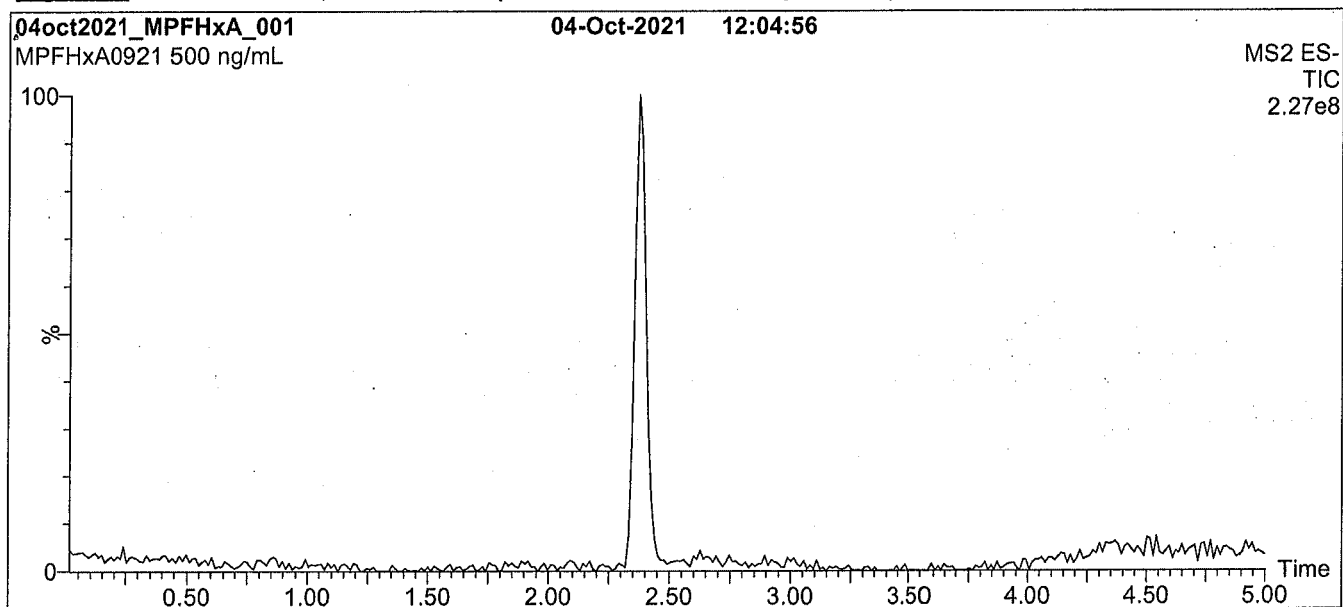
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Figure 1: MPFHxA; LC/MS Data (Full Scan and Mass Spectrum)**Conditions for Figure 1:**

Waters Acquity Ultra Performance LC
Waters Xevo TQ-S micro MS

Chromatographic Conditions:

Column: Acquity UPLC BEH Shield RP₁₈
1.7 μ m, 2.1 x 100 mm

Mobile phase: Gradient

Start: 50% H₂O / 50% (80:20 MeOH:ACN)
(both with 10 mM NH₄OAc buffer)
Ramp to 90% organic over 7 min and hold for
2 min before returning to initial conditions in 1 min.
Time: 12 min

Flow: 300 μ L/min

MS Parameters:

Experiment: Full Scan (225 - 850 amu)

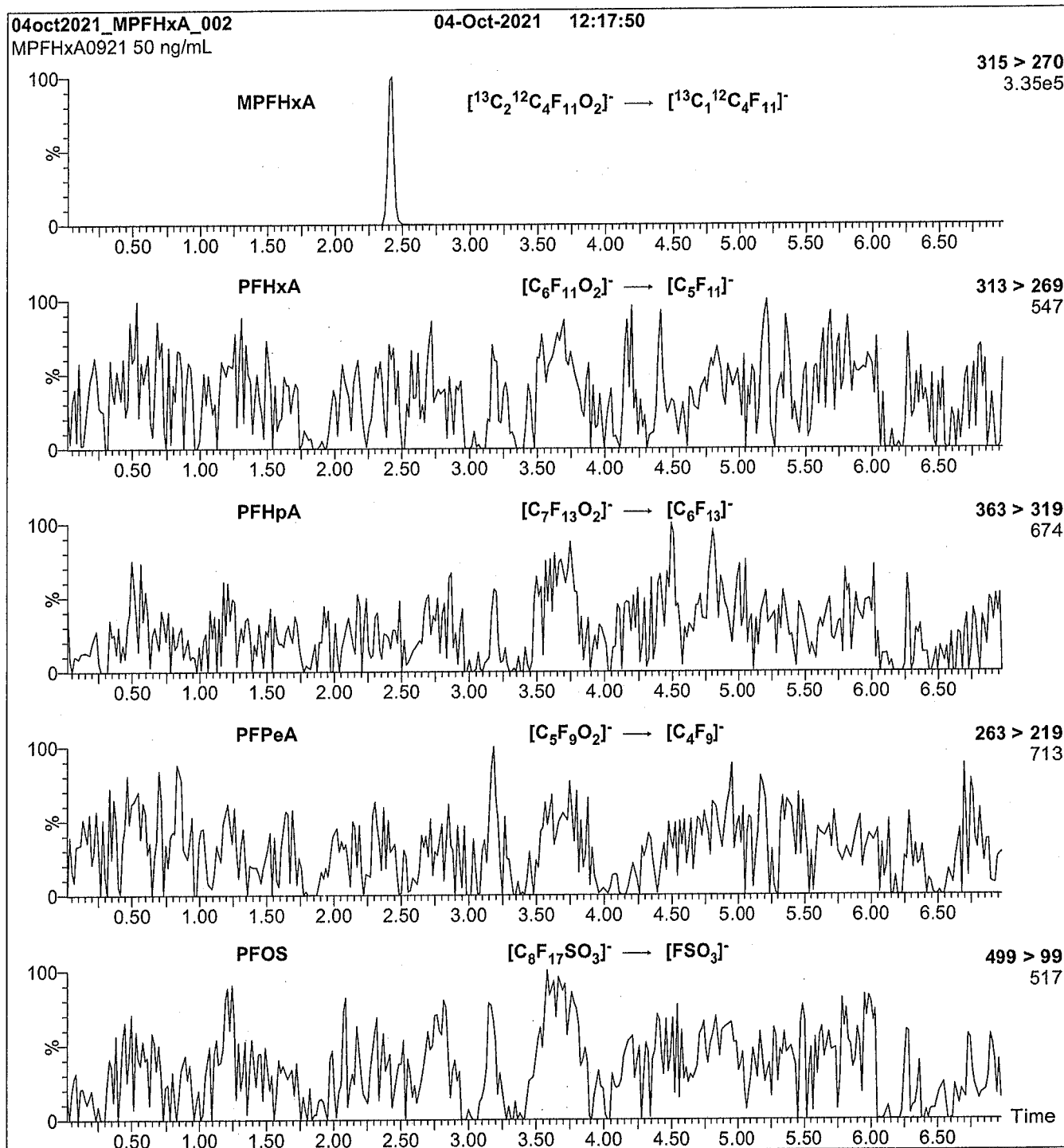
Source: Electrospray (negative)

Capillary Voltage (kV) = 2.00

Cone Voltage (V) = 10.00

Desolvation Temperature (°C) = 500

Desolvation Gas Flow (L/hr) = 1000

Figure 2: MPFHxA; LC/MS/MS Data (Selected MRM Transitions)**Conditions for Figure 2:**

Injection: On-column (MPFHxA)

Mobile phase: Same as Figure 1

Flow: 300 $\mu\text{L}/\text{min}$ **MS Parameters:**

Collision Gas (mbar) = 3.31e-3

Collision Energy (eV) = 8

Analytical Standard Record

22A0117

Description: PFAS - IIS MPFHxA 50ug/mL
Standard Type: Analyte Spike
Solvent: MeOH
Final Volume (mLs): 1.2
Vials: 1

Expires: 10/04/2026
Prepared: 10/04/2021
Prepared By: Dipti Gokal
Department: PFAS
Last Edit: 01/20/2022 15:48 by HGH

Analyte	Parent	CAS Number	Concentration	Units
13C2-PFHxA		13C2-PFHxA	50	ug/mL

Analytical Standard Record

22A0117

Description: PFAS - IIS MPFHxA 50ug/mL
Standard Type: Analyte Spike
Solvent: MeOH
Final Volume (mLs): 1.2
Vials: 1

Expires: 10/04/2026
Prepared: 10/04/2021
Prepared By: Dipti Gokal
Department: PFAS
Last Edit: 01/20/2022 15:48 by HGH

Analyte	Parent	CAS Number	Concentration	Units
13C2-PFHxA		13C2-PFHxA	50	ug/mL



WELLINGTON LABORATORIES

CERTIFICATE OF ANALYSIS DOCUMENTATION

PRODUCT CODE:

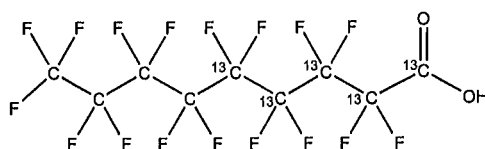
MPFNA

LOT NUMBER:

MPFNA1021

COMPOUND:Perfluoro-n-(1,2,3,4,5-¹³C₅)nonanoic acid**STRUCTURE:****CAS #:**

960315-49-5

**MOLECULAR FORMULA:**¹³C₅¹²C₄HF₁₇O₂**MOLECULAR WEIGHT:**

469.04

CONCENTRATION:

50.0 ± 2.5 µg/mL

SOLVENT(S):

Methanol

Water (<1%)

CHEMICAL PURITY:

>98%

ISOTOPIC PURITY:≥99% ¹³C(1,2,3,4,5-¹³C₅)**LAST TESTED:** (mm/dd/yyyy)

10/29/2021

EXPIRY DATE: (mm/dd/yyyy)

10/29/2026

RECOMMENDED STORAGE:

Store ampoule in a cool, dark place

DOCUMENTATION/ DATA ATTACHED:

Figure 1: LC/MS Data (Full Scan and Mass Spectrum)

Figure 2: LC/MS/MS Data (Selected MRM Transitions)

ADDITIONAL INFORMATION:

- See page 2 for further details.
- Contains 4 mole eq. of NaOH to prevent conversion of the carboxylic acid to the methyl ester.

FOR LABORATORY USE ONLY: NOT FOR HUMAN OR DRUG USE

Certified By:

B.G. Chittim, General Manager
Date: 11/01/2021
(mm/dd/yyyy)

Wellington Laboratories Inc., 345 Southgate Dr. Guelph ON N1G 3M5 CANADA
519-822-2436 • Fax: 519-822-2849 • info@well-labs.com

INTENDED USE:

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HANDLING:

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UNCERTAINTY:

The maximum combined relative standard uncertainty of our reference standard solutions is calculated using the following equation:

The combined relative standard uncertainty, $u_c(y)$, of a value y and the uncertainty of the independent parameters

x_1, x_2, \dots, x_n on which it depends is:

$$u_c(y(x_1, x_2, \dots, x_n)) = \sqrt{\sum_{i=1}^n u(y, x_i)^2}$$

where x is expressed as a relative standard uncertainty of the individual parameter.

The individual uncertainties taken into account include those associated with weights (calibration of the balance) and volumes (calibration of the volumetric glassware). An expanded maximum combined percent relative uncertainty of $\pm 5\%$ (calculated with a coverage factor of 2 and a level of confidence of 95%) is stated on the Certificate of Analysis for all of our products.

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LIMITED WARRANTY:

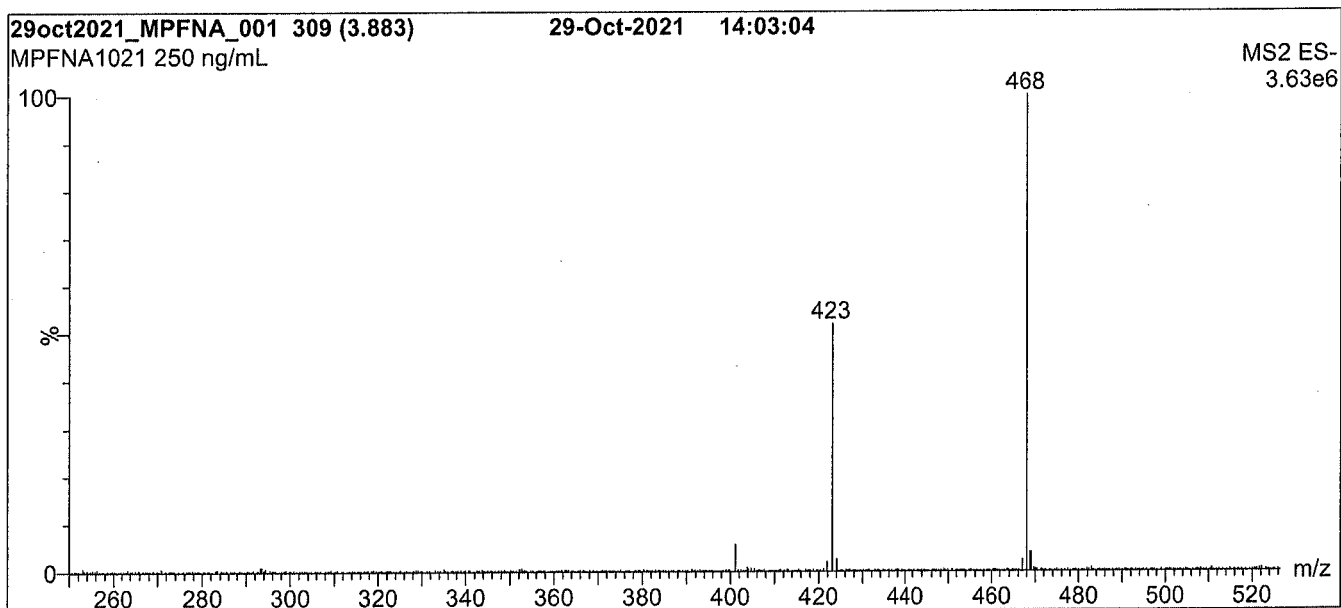
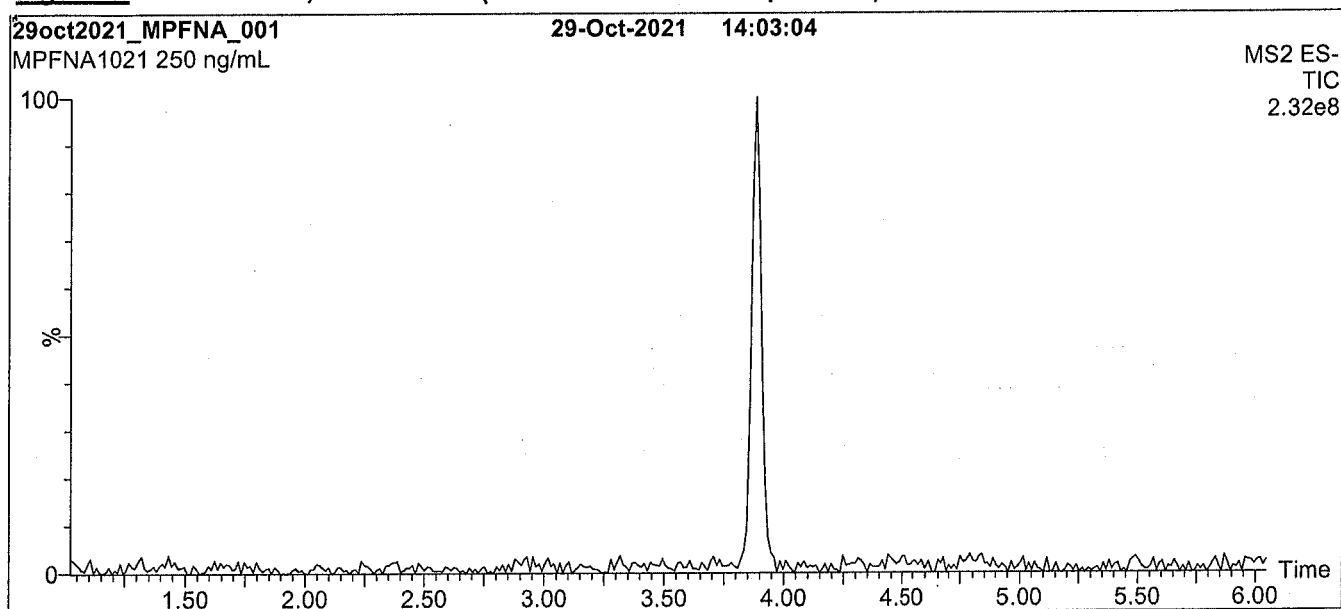
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QUALITY MANAGEMENT:

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Figure 1: MPFNA; LC/MS Data (Full Scan and Mass Spectrum)**Conditions for Figure 1:**

Waters Acquity Ultra Performance LC
Waters Xevo TQ-S micro MS

Chromatographic Conditions:

Column: Acquity UPLC BEH Shield RP₁₈
1.7 μ m, 2.1 x 100 mm

Mobile phase: Gradient

Start: 40% H₂O / 60% (80:20 MeOH:ACN)
(both with 10 mM NH₄OAc buffer)
Ramp to 90% organic over 8 min and hold for
2 min before returning to initial conditions in 0.75 min.
Time: 12 min

Flow: 300 μ L/min

MS Parameters:

Experiment: Full Scan (250 - 850 amu)

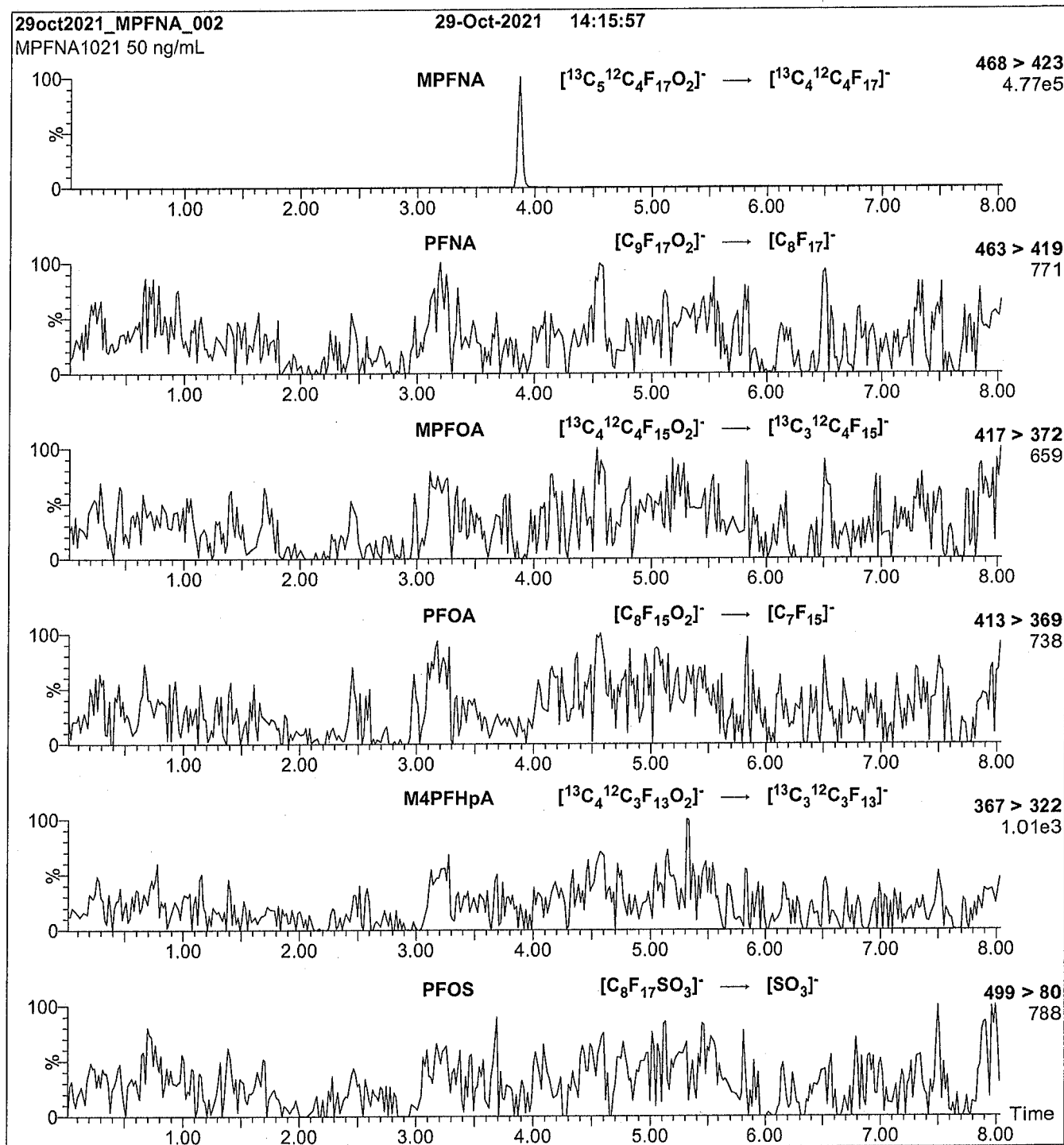
Source: Electrospray (negative)

Capillary Voltage (kV) = 2.00

Cone Voltage (V) = 10.00

Desolvation Temperature (°C) = 500

Desolvation Gas Flow (L/hr) = 1000

Figure 2: MPFNA; LC/MS/MS Data (Selected MRM Transitions)**Conditions for Figure 2:**

Injection: On-column (MPFNA)

Mobile phase: Same as Figure 1

Flow: 300 $\mu\text{L}/\text{min}$ **MS Parameters:**

Collision Gas (mbar) = 3.16e-3

Collision Energy (eV) = 10

Analytical Standard Record

22A0118

Description: PFAS - IIS MPFNA 50ug/mL
Standard Type: Analyte Spike
Solvent: MeOH
Final Volume (mLs): 1.2
Vials: 1

Expires: 10/29/2026
Prepared: 10/29/2021
Prepared By: Dipti Gokal
Department: PFAS
Last Edit: 01/20/2022 15:48 by HGH

Analyte	Parent	CAS Number	Concentration	Units
13C5-PFNA		13C5-PFNA	50	ug/mL

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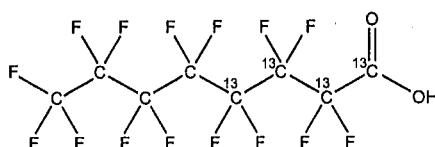
CERTIFICATE OF ANALYSIS DOCUMENTATION

PRODUCT CODE: MPFOA
COMPOUND: Perfluoro-n-(1,2,3,4-¹³C₄)octanoic acid

LOT NUMBER: MPFOA1121

STRUCTURE:

CAS #: 960315-48-4



MOLECULAR FORMULA: ¹³C₄¹²C₄HF₁₅O₂
CONCENTRATION: 50.0 ± 2.5 µg/mL

MOLECULAR WEIGHT: 418.04
SOLVENT(S): Methanol
Water (<1%)

CHEMICAL PURITY: >98%
LAST TESTED: (mm/dd/yyyy) 12/07/2021
EXPIRY DATE: (mm/dd/yyyy) 12/07/2026

ISOTOPIC PURITY: ≥99% ¹³C
(1,2,3,4-¹³C₄)

RECOMMENDED STORAGE: Store ampoule in a cool, dark place

DOCUMENTATION/ DATA ATTACHED:

Figure 1: LC/MS Data (Full Scan and Mass Spectrum)
Figure 2: LC/MS/MS Data (Selected MRM Transitions)

ADDITIONAL INFORMATION:

- See page 2 for further details.
- Contains 4 mole eq. of NaOH to prevent conversion of the carboxylic acid to the methyl ester.

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Certified By: 
B.G. Chittim, General Manager

Date: 12/20/2021
(mm/dd/yyyy)

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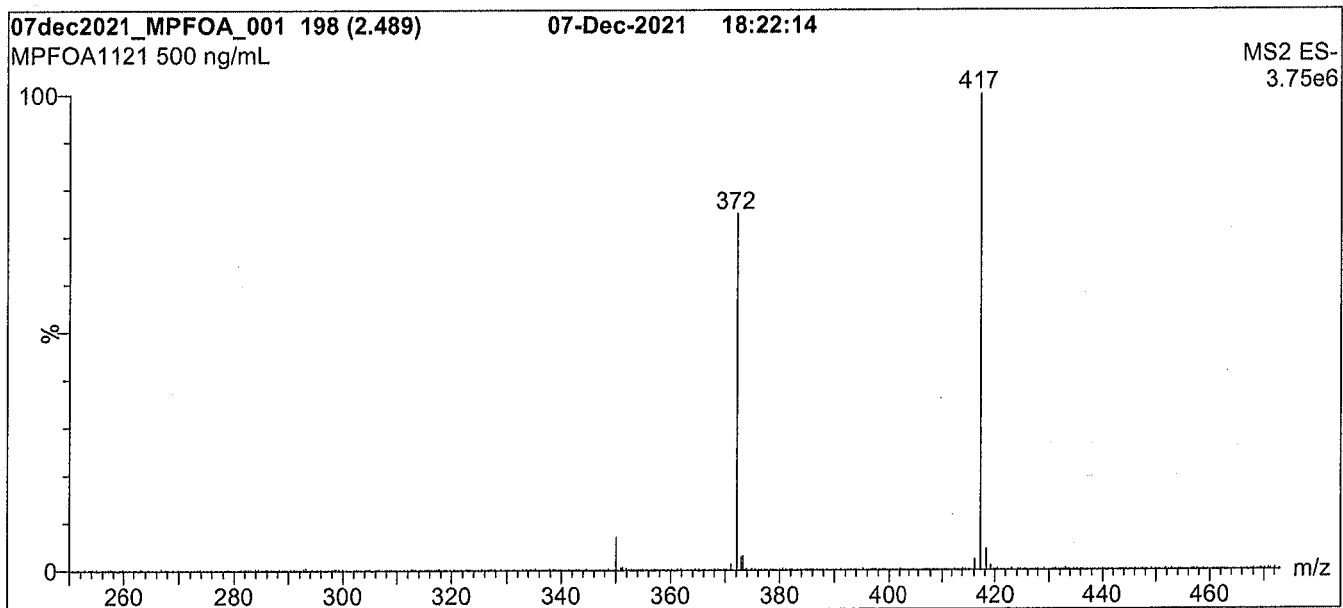
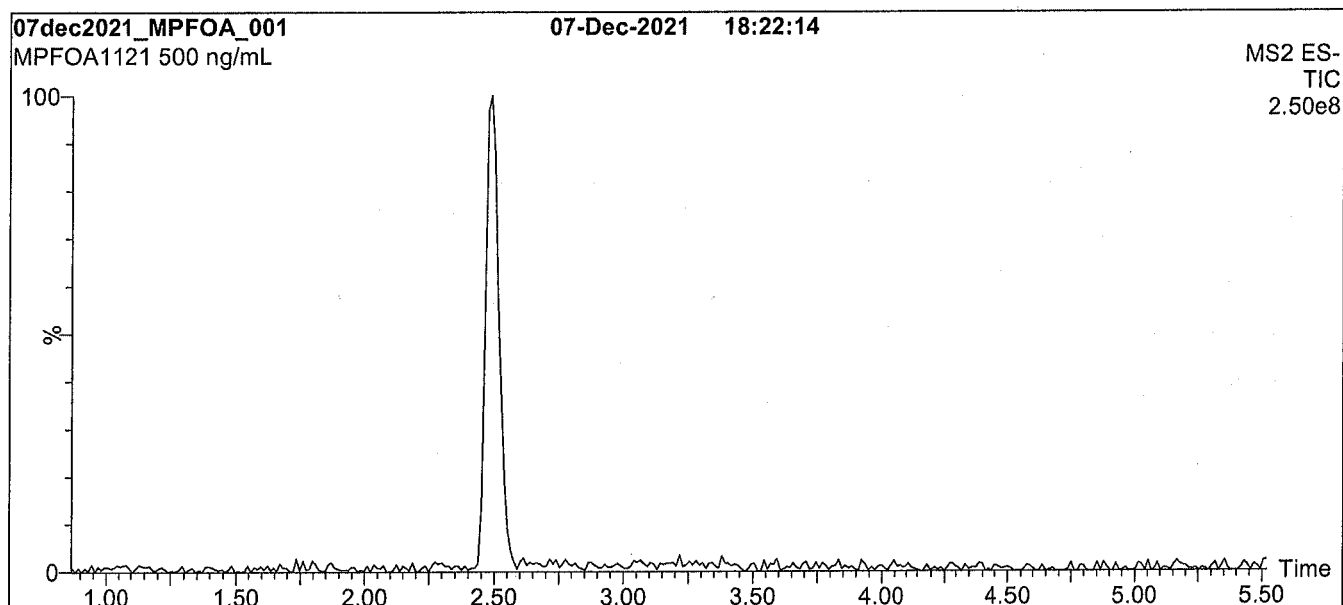
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Waters Xevo TQ-S micro MS

Chromatographic Conditions:

Column: Acquity UPLC BEH Shield RP₁₈
1.7 μ m, 2.1 x 100 mm

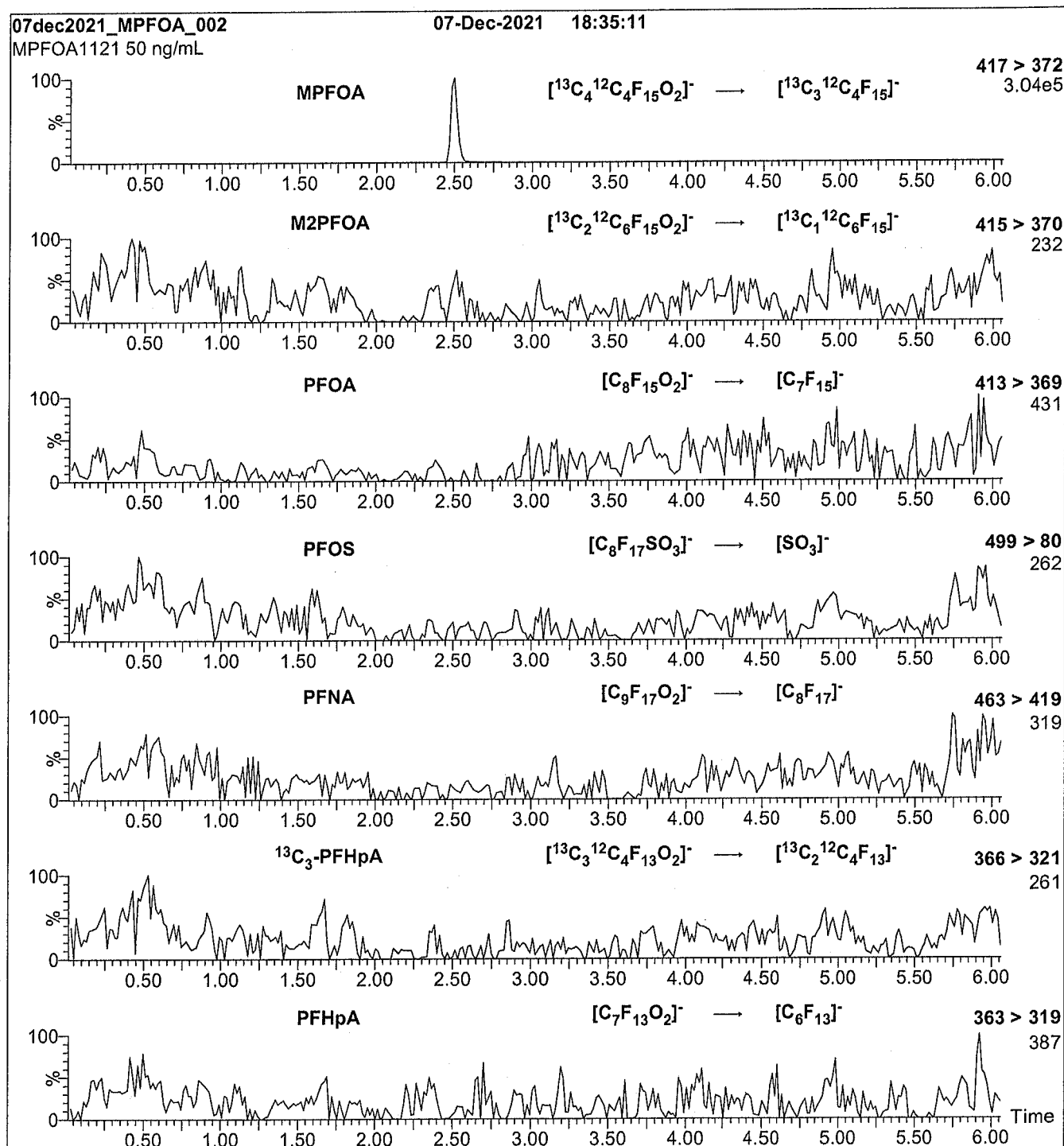
Mobile phase: Gradient
Start: 40% H₂O / 60% (80:20 MeOH:ACN)
(both with 10 mM NH₄OAc buffer)
Ramp to 90% organic over 8 min and hold for 2 min
before returning to initial conditions in 0.75 min.
Time: 12 min

Flow: 300 μ L/min

MS Parameters:

Experiment: Full Scan (250 - 850 amu)

Source: Electrospray (negative)
Capillary Voltage (kV) = 2.00
Cone Voltage (V) = 10.00
Desolvation Temperature (°C) = 500
Desolvation Gas Flow (L/hr) = 1000

Figure 2: MPFOA; LC/MS/MS Data (Selected MRM Transitions)**Conditions for Figure 2:**

Injection: On-column (MPFOA)

Mobile phase: Same as Figure 1

Flow: 300 $\mu\text{L}/\text{min}$ **MS Parameters:**

Collision Gas (mbar) = 3.39e-3

Collision Energy (eV) = 8

Analytical Standard Record

22A0119

Description:	PFAS - IIS MPFOA 50ug/mL	Expires:	12/07/2026
Standard Type:	Analyte Spike	Prepared:	12/07/2021
Solvent:	MeOH	Prepared By:	Dipti Gokal
Final Volume (mls):	1.2	Department:	PFAS
Vials:	1	Last Edit:	01/20/2022 15:48 by HGH

Analyte	Parent	CAS Number	Concentration	Units
13C4-PFOA		13C4-PFOA	50	ug/mL



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CERTIFICATE OF ANALYSIS DOCUMENTATION

PRODUCT CODE:

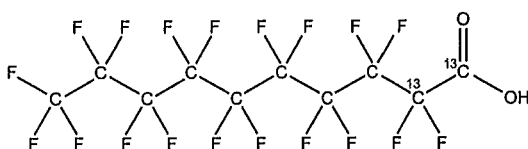
MPFDA

LOT NUMBER:

MPFDA1221

COMPOUND:Perfluoro-n-(1,2-¹³C₂)decanoic acid**STRUCTURE:****CAS #:**

960315-50-8

**MOLECULAR FORMULA:**¹³C₂¹²C₈H₁₉O₂**MOLECULAR WEIGHT:**

516.07

CONCENTRATION:

50.0 ± 2.5 µg/mL

SOLVENT(S):

Methanol

Water (<1%)

CHEMICAL PURITY:

>98%

ISOTOPIC PURITY:≥99% ¹³C**LAST TESTED:** (mm/dd/yyyy)

12/08/2021

(1,2-¹³C₂)**EXPIRY DATE:** (mm/dd/yyyy)

12/08/2026

RECOMMENDED STORAGE:

Store ampoule in a cool, dark place

DOCUMENTATION/ DATA ATTACHED:

Figure 1: LC/MS Data (Full Scan and Mass Spectrum)

Figure 2: LC/MS/MS Data (Selected MRM Transitions)

ADDITIONAL INFORMATION:

- See page 2 for further details.
- Contains 4 mole eq. of NaOH to prevent conversion of the carboxylic acid to the methyl ester.

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Certified By:

B.G. Chittim, General Manager

Date:

12/13/2021

(mm/dd/yyyy)

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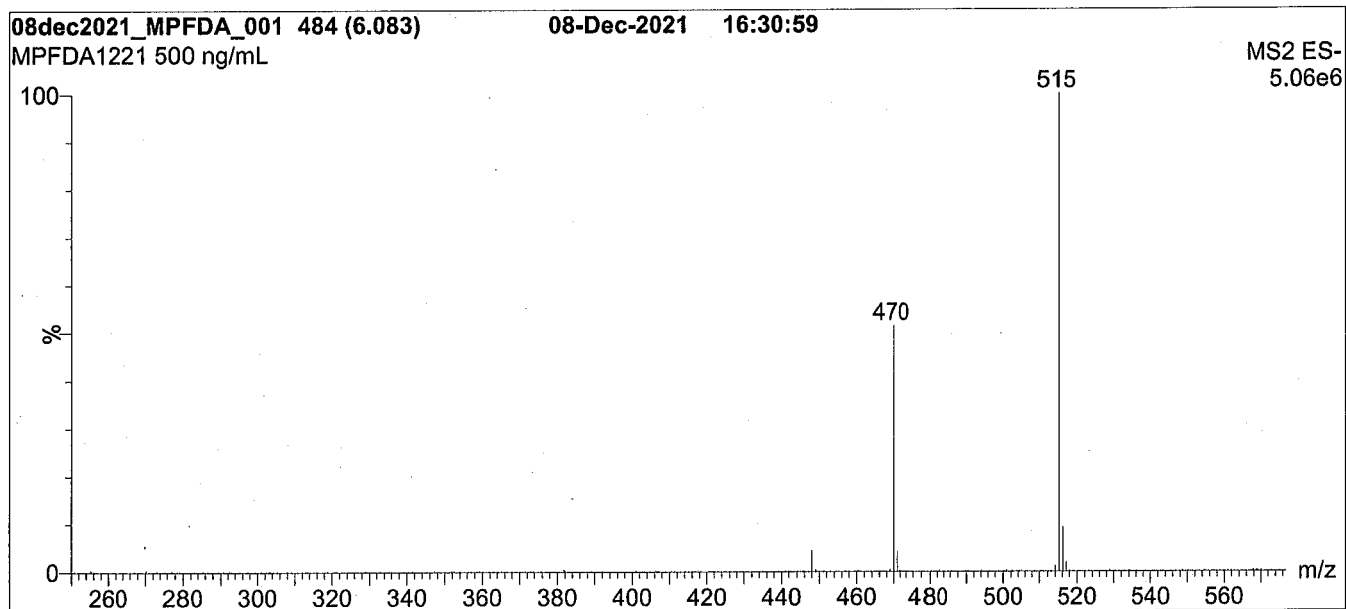
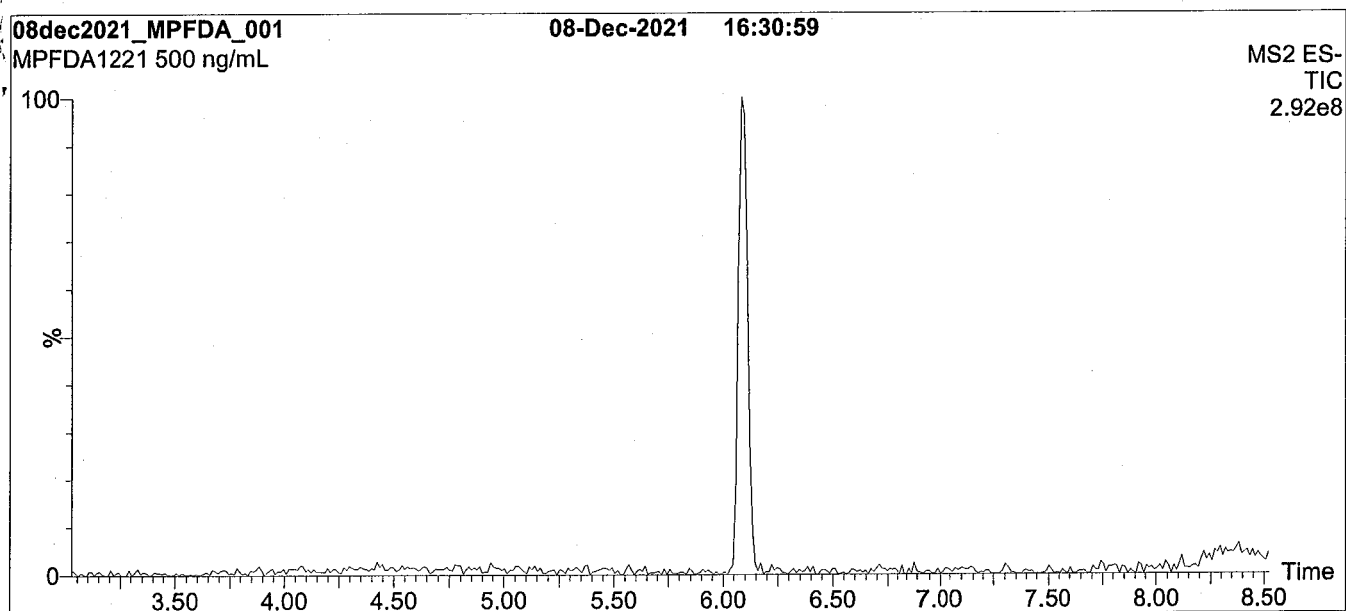
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Waters Acquity Ultra Performance LC
Waters Xevo TQ-S micro MS

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1.7 μ m, 2.1 x 100 mm

Mobile phase: Gradient

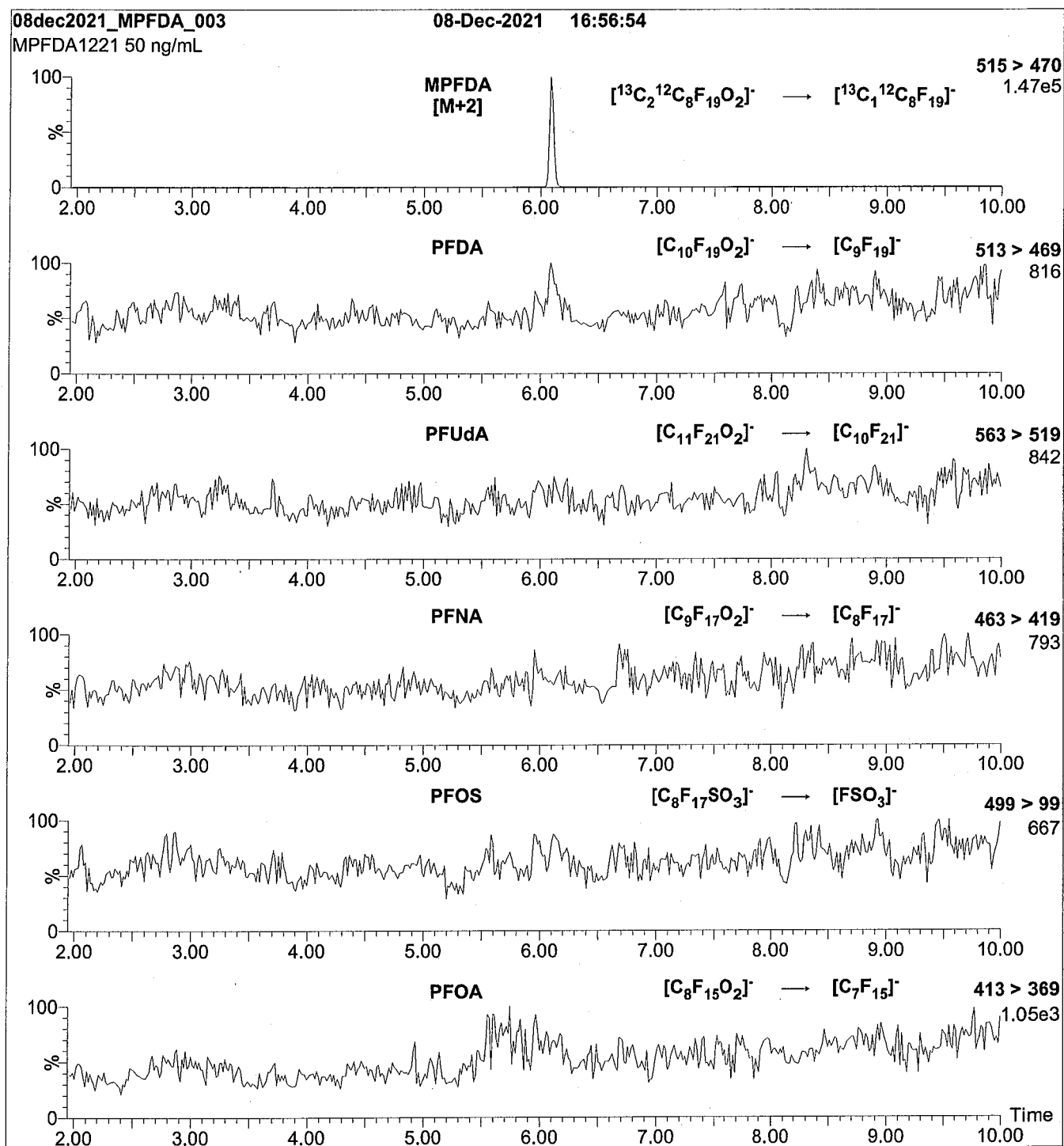
Start: 50% H₂O / 50% (80:20 MeOH:ACN)
(both with 10 mM NH₄OAc buffer)
Ramp to 90% organic over 9 min and hold for
1 min before returning to initial conditions in 0.75 min.
Time: 12 min

Flow: 300 μ L/min

MS Parameters:

Experiment: Full Scan (250 - 850 amu)

Source: Electrospray (negative)
Capillary Voltage (kV) = 2.00
Cone Voltage (V) = 10.00
Desolvation Temperature (°C) = 500
Desolvation Gas Flow (L/hr) = 1000

Figure 2: MPFDA; LC/MS/MS Data (Selected MRM Transitions)**Conditions for Figure 2:**

Injection: On-column (MPFDA)

Mobile phase: Same as Figure 1

Flow: 300 μ L/min**MS Parameters:**

Collision Gas (mbar) = 3.39e-3

Collision Energy (eV) = 10

Analytical Standard Record

22A0120

Description: PFAS - IIS MPFDA 50ug/mL
Standard Type: Analyte Spike
Solvent: MeOH
Final Volume (mLs): 1.2
Vials: 1

Expires: 12/08/2026
Prepared: 12/08/2021
Prepared By: Dipti Gokal
Department: PFAS
Last Edit: 01/20/2022 15:49 by HGH

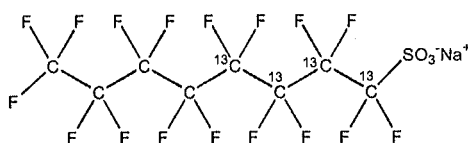
Analyte	Parent	CAS Number	Concentration	Units
13C2-PFDA		13C2-PFDA	50	ug/mL



WELLINGTON LABORATORIES

CERTIFICATE OF ANALYSIS DOCUMENTATION

PRODUCT CODE: MPFOS **LOT NUMBER:** MPFOS0821
COMPOUND: Sodium perfluoro-1-(1,2,3,4-¹³C₄)octanesulfonate
STRUCTURE: **CAS #:** 960315-53-1



MOLECULAR FORMULA: ¹³C₄¹²C₄F₁₇SO₃Na **MOLECULAR WEIGHT:** 526.08
CONCENTRATION: 50.0 ± 2.5 µg/mL (Na salt) **SOLVENT(S):** Methanol
 47.9 ± 2.4 µg/mL (MPFOS acid)
 47.8 ± 2.4 µg/mL (MPFOS anion)
CHEMICAL PURITY: >98% **ISOTOPIC PURITY:** ≥99% ¹³C
LAST TESTED: (mm/dd/yyyy) 08/18/2021 (1,2,3,4-¹³C₄)
EXPIRY DATE: (mm/dd/yyyy) 08/18/2026
RECOMMENDED STORAGE: Store ampoule in a cool, dark place

DOCUMENTATION/ DATA ATTACHED:

Figure 1: LC/MS Data (Full Scan and Mass Spectrum)
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ADDITIONAL INFORMATION:

- See page 2 for further details.
- Contains ~0.4% sodium perfluoro-1-(¹³C₃)heptanesulfonate.

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Certified By: 
 B.G. Chittim, General Manager

Date: 08/19/2021
 (mm/dd/yyyy)

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LIMITED WARRANTY:

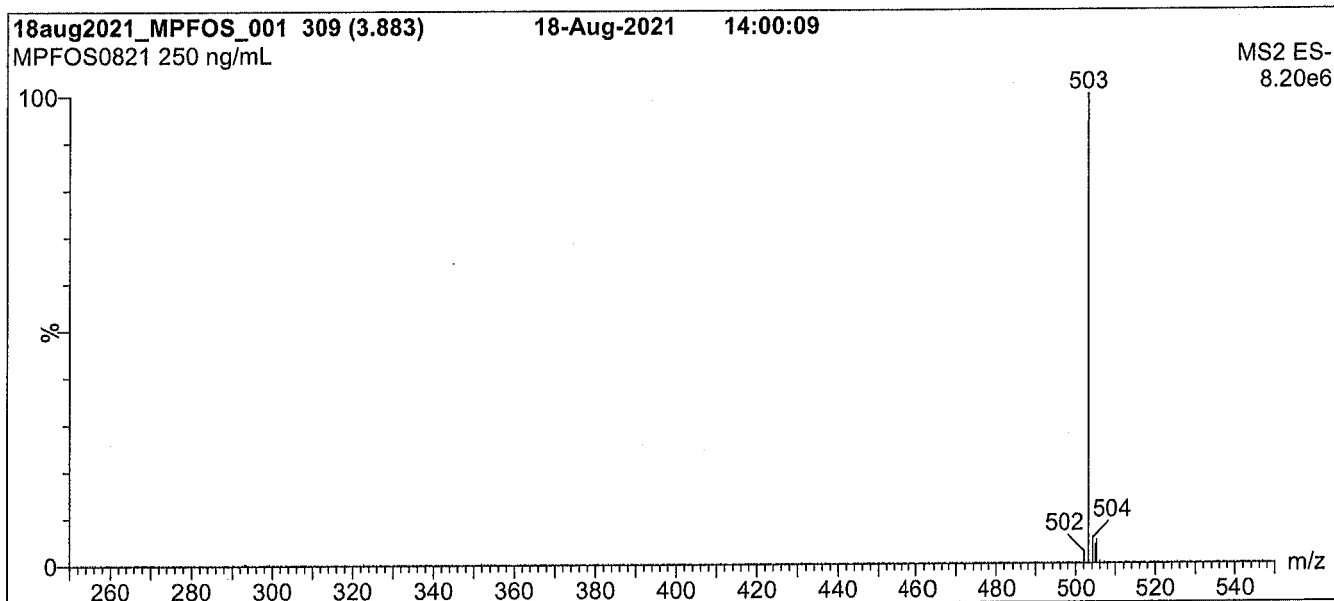
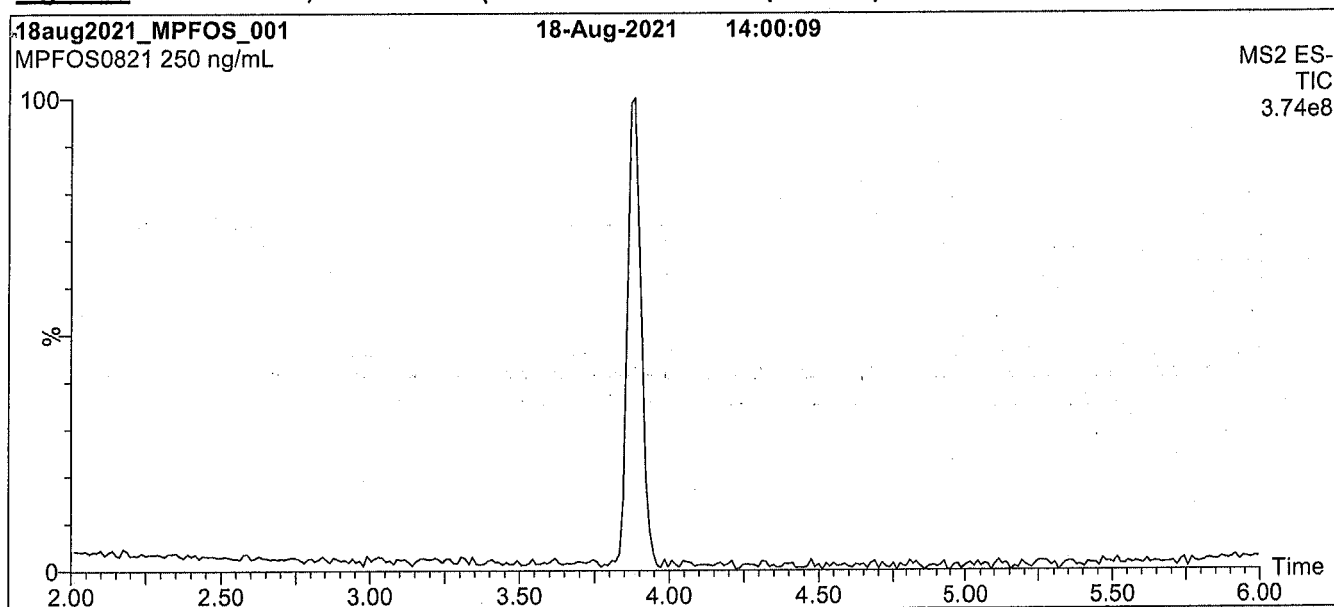
At the time of shipment, all products are warranted to be free of defects in material and workmanship and to conform to the stated technical and purity specifications.

QUALITY MANAGEMENT:

This product was produced using a Quality Management System registered to the latest versions of ISO 9001 by SAI Global, ISO/IEC 17025 by the Canadian Association for Laboratory Accreditation Inc. (CALA; A1226), and ISO 17034 by ANSI National Accreditation Board (ANAB; AR-1523).



For additional information or assistance concerning this or any other products from Wellington Laboratories Inc., please visit our website at www.well-labs.com or contact us directly at info@well-labs.com

Figure 1: MPFOS; LC/MS Data (Full Scan and Mass Spectrum)**Conditions for Figure 1:**

Waters Acquity Ultra Performance LC
Waters Xevo TQ-S micro MS

Chromatographic Conditions:

Column: Acquity UPLC BEH Shield RP₁₈
1.7 μ m, 2.1 x 100 mm

Mobile phase: Gradient

Start: 40% H₂O / 60% (80:20 MeOH:ACN)
(both with 10 mM NH₄OAc buffer)
Ramp to 90% organic over 8 min and hold for
2 min before returning to initial conditions in 0.75 min.
Time: 12 min

Flow: 300 μ L/min

MS Parameters:

Experiment: Full Scan (250 - 850 amu)

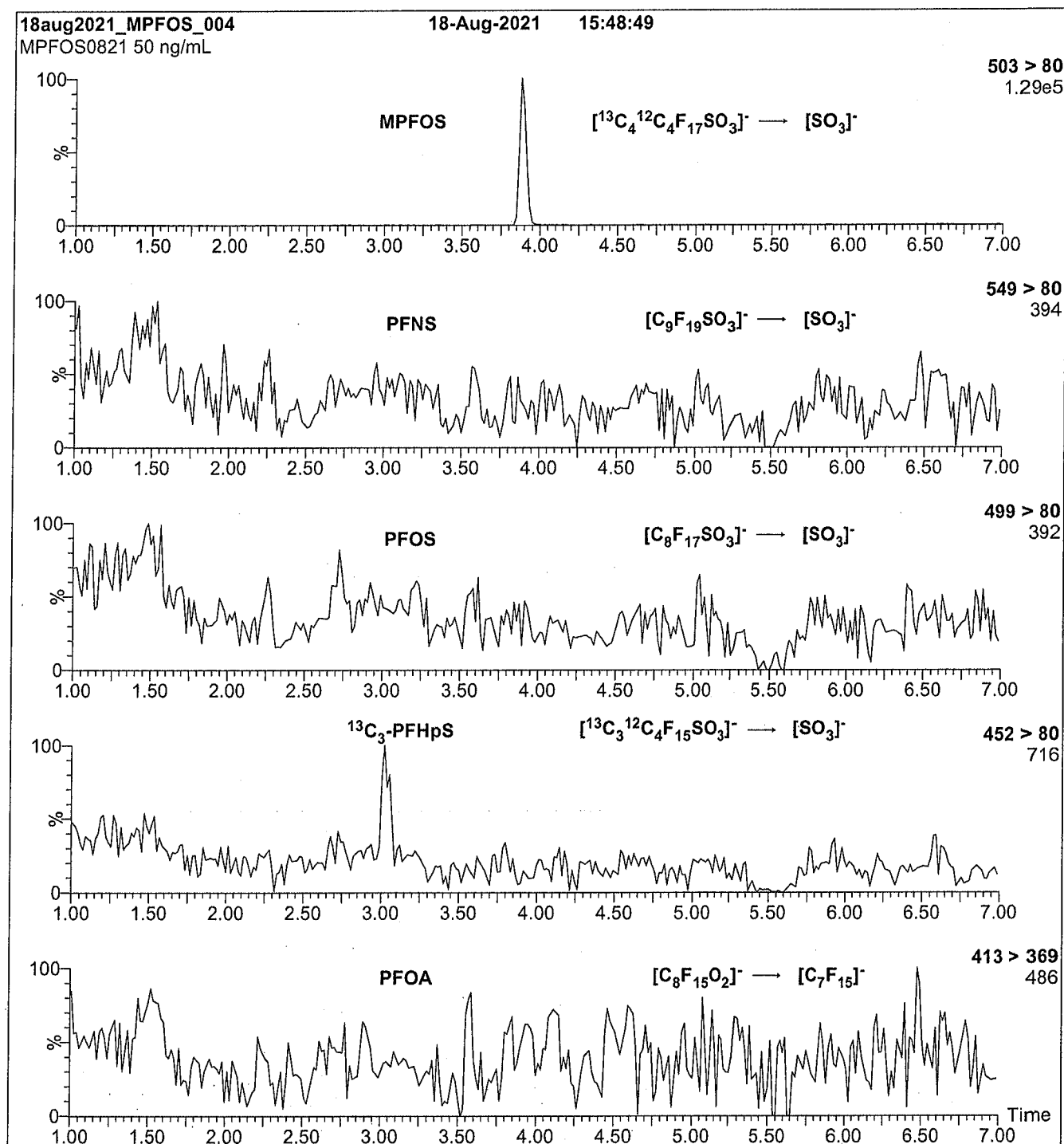
Source: Electrospray (negative)

Capillary Voltage (kV) = 2.00

Cone Voltage (V) = 10.00

Desolvation Temperature (°C) = 500

Desolvation Gas Flow (L/hr) = 1000

Figure 2: MPFOS; LC/MS/MS Data (Selected MRM Transitions)**Conditions for Figure 2:**

Injection: On-column (MPFOS)

Mobile phase: Same as Figure 1

Flow: 300 $\mu\text{L}/\text{min}$ **MS Parameters:**

Collision Gas (mbar) = 3.39e-3

Collision Energy (eV) = 42

Analytical Standard Record

22A0121

Description: PFAS - IIS MPFOS 50ug/mL
Standard Type: Analyte Spike
Solvent: MeOH
Final Volume (mLs): 1.2
Vials: 1

Expires: 08/18/2026
Prepared: 08/18/2021
Prepared By: Dipti Gokal
Department: PFAS
Last Edit: 01/20/2022 15:49 by HGH

Analyte	Parent	CAS Number	Concentration	Units
13C4-PFOS		13C4-PFOS	50	ug/mL



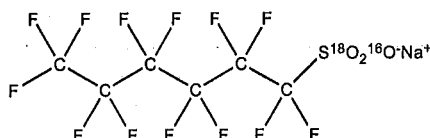
WELLINGTON LABORATORIES

CERTIFICATE OF ANALYSIS DOCUMENTATION

PRODUCT CODE: MPFHxS
COMPOUND: Sodium perfluoro-1-hexane(¹⁸O₂)sulfonate

LOT NUMBER: MPFHxS1021

STRUCTURE:



CAS #: 1585941-14-5

MOLECULAR FORMULA: C₆F₁₃S¹⁸O₂¹⁶ONa
CONCENTRATION: 50.0 ± 2.5 µg/mL (Na salt)
47.4 ± 2.4 µg/mL (MPFHxS acid)
47.3 ± 2.4 µg/mL (MPFHxS anion)

MOLECULAR WEIGHT: 426.10
SOLVENT(S): Methanol

CHEMICAL PURITY: >98%
LAST TESTED: (mm/dd/yyyy) 10/29/2021
EXPIRY DATE: (mm/dd/yyyy) 10/29/2026
RECOMMENDED STORAGE: Store ampoule in a cool, dark place

ISOTOPIC PURITY: >94% (¹⁸O₂)

DOCUMENTATION/ DATA ATTACHED:

Figure 1: LC/MS Data (Full Scan and Mass Spectrum)

Figure 2: LC/MS/MS Data (Selected MRM Transitions)

ADDITIONAL INFORMATION:

- See page 2 for further details.
- The response factor for MPFHxS (C₆F₁₃S¹⁸O₂¹⁶O) has been observed to be up to 10% lower than for PFHxS (C₆F₁₃S¹⁸O₃) when both compounds are injected together. This difference may vary between instruments.
- Contains ~0.6% of sodium perfluoro-1-octane(¹⁸O₂)sulfonate (¹⁸O₂-PFOS) and ~0.3% of sodium perfluoro-1-heptane(¹⁸O₂)sulfonate (¹⁸O₂-PFHpS).
- Due to the isotopic purity of the starting material (¹⁸O₂ >94%), MPFHxS contains ~0.3% of PFHxS. This value agrees with the theoretical percent relative abundance that is expected based on the stated isotopic purity.

FOR LABORATORY USE ONLY: NOT FOR HUMAN OR DRUG USE

Certified By: 
B.G. Chittim, General Manager

Date: 11/05/2021
(mm/dd/yyyy)

Wellington Laboratories Inc., 345 Southgate Dr. Guelph ON N1G 3M5 CANADA
519-822-2436 • Fax: 519-822-2849 • info@well-labs.com

INTENDED USE:

The products prepared by Wellington Laboratories Inc. are for laboratory use only. This certified reference material (CRM) was designed to be used as a standard for the identification and/or quantification of the specific chemical compound it contains.

HANDLING:

This product should only be used by qualified personnel familiar with its potential hazards and trained in the handling of hazardous chemicals. Due care should be exercised to prevent unnecessary human contact or ingestion. All procedures should be carried out in a well-functioning fume hood and suitable gloves, eye protection, and clothing should be worn at all times. Waste should be disposed of according to national and regional regulations. Safety Data Sheets (SDSs) are available upon request.

SYNTHESIS / CHARACTERIZATION:

Our products are synthesized using single-product unambiguous routes whenever possible. They are then characterized, and their structures and purities confirmed, using a combination of the most relevant techniques, such as NMR, GC/MS, LC/MS/MS, SFC/UV/MS/MS, x-ray crystallography, and melting point. Isotopic purities of mass-labelled compounds are also confirmed using HRGC/HRMS and/or LC/MS/MS.

HOMOGENEITY:

Prior to solution preparation, crystalline material is tested for homogeneity using a variety of techniques (as stated above) and its solubility in a given diluent is taken into consideration. Duplicate solutions of a new product are prepared from the same crystalline lot and, after the addition of an appropriate internal standard, they are compared by GC/MS, LC/MS/MS, and/or SFC/UV/MS/MS. The relative response factors of the analyte of interest in each solution are required to be <5% RSD. New solution lots of existing products are compared to older lots in the same manner, which further confirms the homogeneity of the crystalline material as well as the stability and homogeneity of the solutions in the storage containers. In order to maintain the integrity of the assigned value(s), and associated uncertainty, the dilution or injection of a subsample of this product should be performed using calibrated measuring equipment.

UNCERTAINTY:

The maximum combined relative standard uncertainty of our reference standard solutions is calculated using the following equation:

The combined relative standard uncertainty, $u_c(y)$, of a value y and the uncertainty of the independent parameters

x_1, x_2, \dots, x_n on which it depends is:

$$u_c(y(x_1, x_2, \dots, x_n)) = \sqrt{\sum_{i=1}^n u(y, x_i)^2}$$

where x is expressed as a relative standard uncertainty of the individual parameter.

The individual uncertainties taken into account include those associated with weights (calibration of the balance) and volumes (calibration of the volumetric glassware). An expanded maximum combined percent relative uncertainty of $\pm 5\%$ (calculated with a coverage factor of 2 and a level of confidence of 95%) is stated on the Certificate of Analysis for all of our products.

TRACEABILITY:

All reference standard solutions are traceable to specific crystalline lots. The microbalances used for solution preparation are regularly calibrated by an external ISO/IEC 17025 accredited laboratory. In addition, their calibration is verified prior to each weighing using calibrated external weights traceable to an ISO/IEC 17025 accredited laboratory. All volumetric glassware used is calibrated, of Class A tolerance, and traceable to an ISO/IEC 17025 accredited laboratory. For certain products, traceability to international interlaboratory studies has also been established.

EXPIRY DATE / PERIOD OF VALIDITY:

Ongoing stability studies of this product have demonstrated stability in its composition and concentration, until the specified expiry date, in the unopened ampoule. Monitoring for any degradation or change in concentration of the listed analyte(s) is performed on a routine basis.

LIMITED WARRANTY:

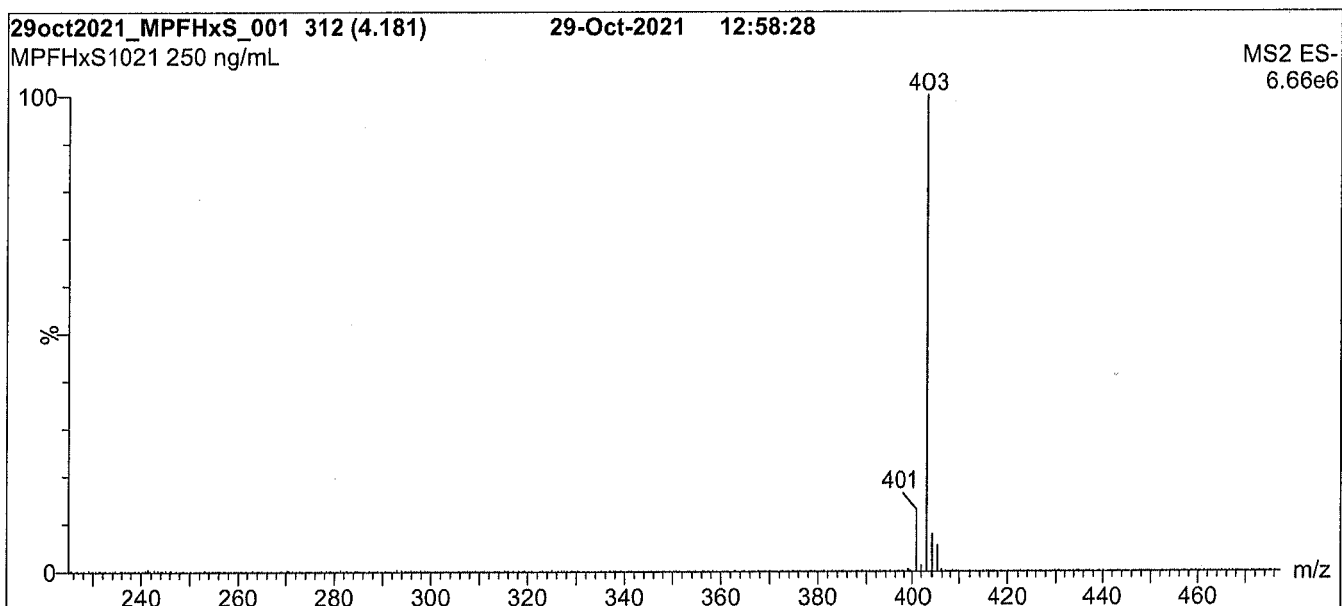
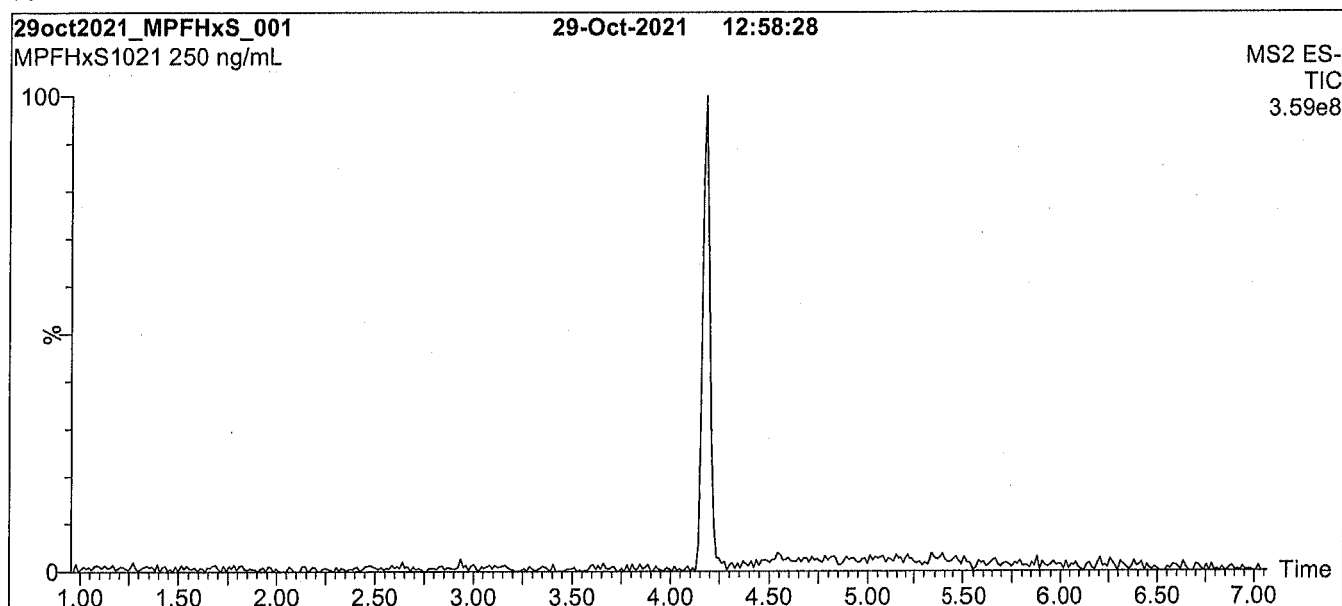
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Figure 1: MPFHxS; LC/MS Data (Full Scan and Mass Spectrum)**Conditions for Figure 1:**

Waters Acquity Ultra Performance LC
Waters Xevo TQ-S micro MS

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1.7 μ m, 2.1 x 100 mm

Mobile phase: Gradient

Start: 50% H₂O / 50% (80:20 MeOH:ACN)
(both with 10 mM NH₄OAc buffer)
Ramp to 90% organic over 9 min and hold for
1 min before returning to initial conditions in 0.75 min.
Time: 12 min

Flow: 300 μ L/min

MS Parameters:

Experiment: Full Scan (225 - 850 amu)

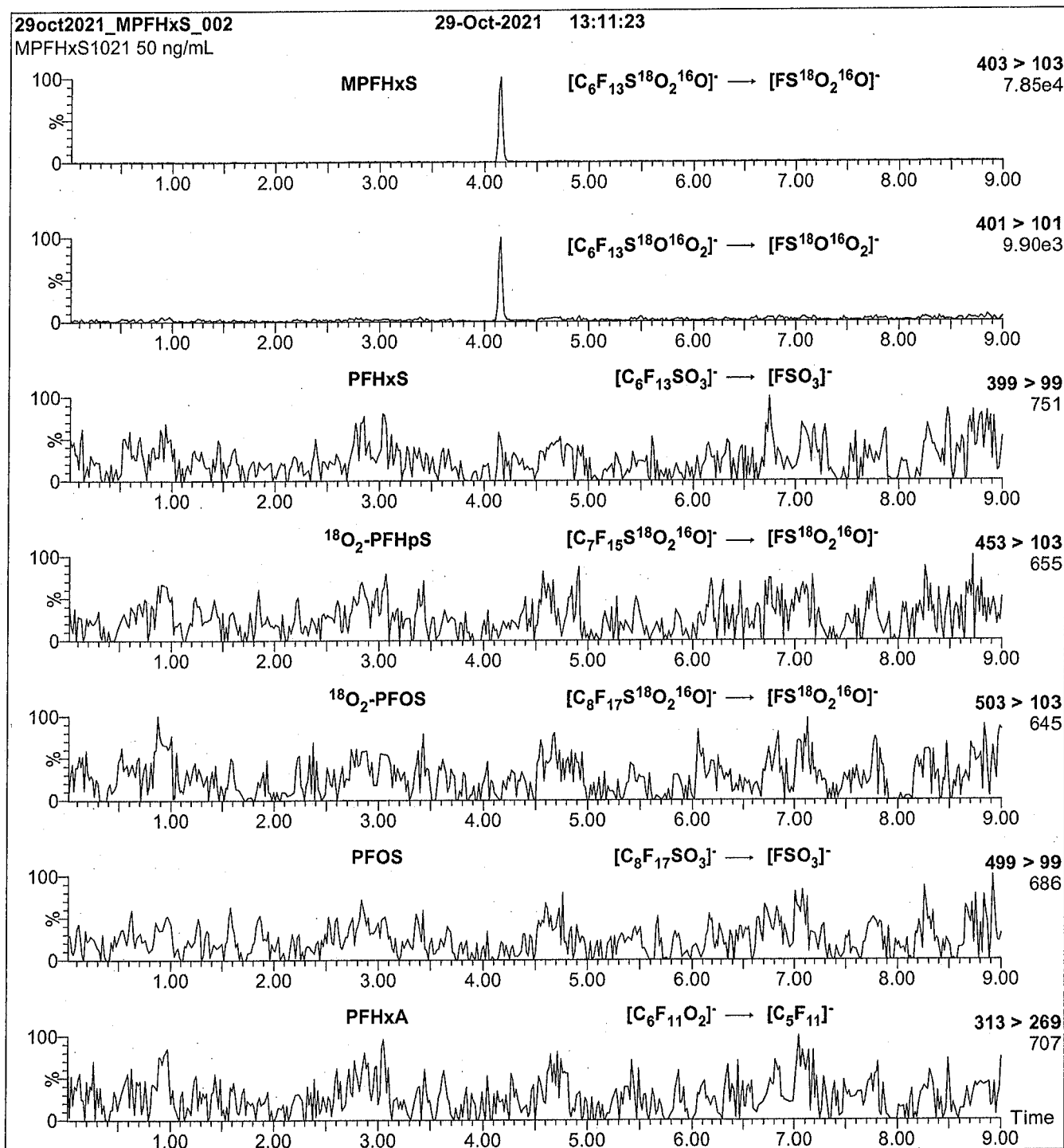
Source: Electrospray (negative)

Capillary Voltage (kV) = 2.00

Cone Voltage (V) = 10.00

Desolvation Temperature (°C) = 500

Desolvation Gas Flow (L/hr) = 1000

Figure 2: MPFHxS; LC/MS/MS Data (Selected MRM Transitions)**Conditions for Figure 2:**

Injection: On-column (MPFHxS)

Mobile phase: Same as Figure 1

Flow: 300 μ L/min**MS Parameters:**

Collision Gas (mbar) = 3.16e-3

Collision Energy (eV) = 32

Analytical Standard Record

22A0122

Description: PFAS - IIS MPFHxS 50ug/mL
Standard Type: Analyte Spike
Solvent: MeOH
Final Volume (mLs): 1.2
Vials: 1

Expires: 10/29/2026
Prepared: 10/29/2021
Prepared By: Dipti Gokal
Department: PFAS
Last Edit: 01/20/2022 15:49 by HGH

Analyte	Parent	CAS Number	Concentration	Units
18O2-PFHXS		1802-PFHXS	50	ug/mL

Analytical Standard Record

22A0122

Description: PFAS - IIS MPFHxS 50ug/mL
Standard Type: Analyte Spike
Solvent: MeOH
Final Volume (mLs): 1.2
Vials: 1

Expires: 10/29/2026
Prepared: 10/29/2021
Prepared By: Dipti Gokal
Department: PFAS
Last Edit: 01/20/2022 15:49 by HGH

Analyte	Parent	CAS Number	Concentration	Units
18O2-PFHXS		18O2-PFHXS	50	ug/mL



Certificate of Analysis

Taurodeoxycholic Acid, Sodium Salt - CAS 1180-95-6 - Calbiochem

Batch Number: 3761825
Material Number: 580221-5GM
Molecular Formula: $C_{26}H_{44}NO_6S \cdot Na$
Molecular Weight: 521.7
CAS Number: 1180-95-6

Quality Release Date: 05 OCT 2021
Recommended Retest Date: 30 SEP 2023

Analytical Data

Test	Tolerance	Result
Solubility:		H ₂ O (100 mg/ml)
Chloride:		<0.01%
Loss on drying:	≤5.0 %	0.1%
Color:		White
Form:		Powder
TLC:	≥95.0 %	≥95.00%
IR:		Conforms to reference
Optical rotation:	35.0 ° - 39.0 °	+36.38°
Water by Karl Fischer:	≤5.0 %	2.86%
Carbon:	≥0.00 %	57.78%
Hydrogen:	≥0.00 %	8.32 %
Nitrogen:	≥0.00 %	2.77 %

Storage and Handling: +15°C to +30°C

This lot conforms to specifications established by EMD Millipore Corporation for this product.

Issued by **Jamie Thomas**

This document has been electronically produced and is valid without a signature

Quality Control/ Assurance Signature

05 OCT 2021

Date

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Technical Support NA +1-800-221-1975 | email: www.millipore.com/techservices | www.calbiochem.com

Technical Support All Other Countries - Contact Your Local Office

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Darmstadt, Germany

580221-5GM/09-MAY-2018/EA

Analytical Standard Record

22A0123

Description:	PFAS Taurodeoxycholic Acid, Sodium Salt	Expires:	09/30/2023
Standard Type:	Other	Prepared:	10/05/2021
Solvent:	n/a	Prepared By:	Dipti Gokal
Final Volume (mls):	1	Department:	PFAS
Vials:	1	Last Edit:	01/20/2022 15:54 by DAG

Analyte	Parent	CAS Number	Concentration	Units
TAURODEOXYCHOLIC ACID		516-50-7	1	ug/mL

Analytical Standard Record

22A0123

Description:	PFAS Taurodeoxycholic Acid, Sodium Salt	Expires:	09/30/2023
Standard Type:	Other	Prepared:	10/05/2021
Solvent:	n/a	Prepared By:	Dipti Gokal
Final Volume (mls):	1	Department:	PFAS
Vials:	1	Last Edit:	01/20/2022 15:54 by DAG

Analyte	Parent	CAS Number	Concentration	Units
TAURODEOXYCHOLIC ACID		516-50-7	1	ug/mL

Analytical Standard Record

22A0234

Description:	PFAS IIS 7C 5ug/mL	Expires:	01/20/2023
Standard Type:	Internal Standard	Prepared:	01/20/2022
Solvent:	MeOH/61252	Prepared By:	Dipti Gokal
Final Volume (mLs):	12	Department:	PFAS
Vials:	1	Last Edit:	01/20/2022 15:49 by HGH
Comments:	mpfna had more left over than others.		

Analyte	Parent	CAS Number	Concentration	Units
13C3-PFBA	22A0116	13C3-PFBA	5	ug/mL
13C2-PFHxA	22A0117	13C2-PFHxA	5	ug/mL
13C5-PFNA	22A0118	13C5-PFNA	5	ug/mL
13C4-PFOA	22A0119	13C4-PFOA	5	ug/mL
13C2-PFDA	22A0120	13C2-PFDA	5	ug/mL
13C4-PFOS	22A0121	13C4-PFOS	5	ug/mL
18O2-PFHXS	22A0122	18O2-PFHXS	5	ug/mL

Parent Standards used:

Standard	Description	Prepared	Prepared By	Lot Nbr	Expires	Last Edit	(mLs)
22A0116	PFAS - IIS M3PFBA 50ug/mL	08/19/2021	Wellington Laboratories	M3PFBA0721	08/19/2026	01/20/2022 15:48 by HGH	1.2
22A0117	PFAS - IIS MPFHxA 50ug/mL	10/04/2021	Wellington Laboratories	MPFHxA0921	10/04/2026	01/20/2022 15:48 by HGH	1.2
22A0118	PFAS - IIS MPFNA 50ug/mL	10/29/2021	Wellington Laboratories	MPFNA1021	10/29/2026	01/20/2022 15:48 by HGH	1.2
22A0119	PFAS - IIS MPFOA 50ug/mL	12/07/2021	Wellington Laboratories	MPFOA1121	12/07/2026	01/20/2022 15:48 by HGH	1.2
22A0120	PFAS - IIS MPFDA 50ug/mL	12/08/2021	Wellington Laboratories	MPFDA1221	12/08/2026	01/20/2022 15:49 by HGH	1.2
22A0121	PFAS - IIS MPFOS 50ug/mL	08/18/2021	Wellington Laboratories	MPFOS0821	08/18/2026	01/20/2022 15:49 by HGH	1.2
22A0122	PFAS - IIS MPFHxS 50ug/mL	10/29/2021	Wellington Laboratories	MPFHxS1021	10/29/2026	01/20/2022 15:49 by HGH	1.2

Analytical Standard Record

22A0234

Description:	PFAS IIS 7C 5ug/mL	Expires:	01/20/2023
Standard Type:	Internal Standard	Prepared:	01/20/2022
Solvent:	MeOH/61252	Prepared By:	Dipti Gokal
Final Volume (mLs):	12	Department:	PFAS
Vials:	1	Last Edit:	01/20/2022 15:49 by HGH
Comments:	mpfna had more left over than others.		

Analyte	Parent	CAS Number	Concentration	Units
13C3-PFBA	22A0116	13C3-PFBA	5	ug/mL
13C2-PFHxA	22A0117	13C2-PFHxA	5	ug/mL
13C5-PFNA	22A0118	13C5-PFNA	5	ug/mL
13C4-PFOA	22A0119	13C4-PFOA	5	ug/mL
13C2-PFDA	22A0120	13C2-PFDA	5	ug/mL
13C4-PFOS	22A0121	13C4-PFOS	5	ug/mL
18O2-PFHXS	22A0122	18O2-PFHXS	5	ug/mL

Parent Standards used:

Standard	Description	Prepared	Prepared By	Lot Nbr	Expires	Last Edit	(mLs)
22A0116	PFAS - IIS M3PFBA 50ug/mL	08/19/2021	Wellington Laboratories	M3PFBA0721	08/19/2026	01/20/2022 15:48 by HGH	1.2
22A0117	PFAS - IIS MPFHxA 50ug/mL	10/04/2021	Wellington Laboratories	MPFHxA0921	10/04/2026	01/20/2022 15:48 by HGH	1.2
22A0118	PFAS - IIS MPFNA 50ug/mL	10/29/2021	Wellington Laboratories	MPFNA1021	10/29/2026	01/20/2022 15:48 by HGH	1.2
22A0119	PFAS - IIS MPFOA 50ug/mL	12/07/2021	Wellington Laboratories	MPFOA1121	12/07/2026	01/20/2022 15:48 by HGH	1.2
22A0120	PFAS - IIS MPFDA 50ug/mL	12/08/2021	Wellington Laboratories	MPFDA1221	12/08/2026	01/20/2022 15:49 by HGH	1.2
22A0121	PFAS - IIS MPFOS 50ug/mL	08/18/2021	Wellington Laboratories	MPFOS0821	08/18/2026	01/20/2022 15:49 by HGH	1.2
22A0122	PFAS - IIS MPFHxS 50ug/mL	10/29/2021	Wellington Laboratories	MPFHxS1021	10/29/2026	01/20/2022 15:49 by HGH	1.2

Analytical Standard Record

22A0234

Description:	PFAS IIS 7C 5ug/mL	Expires:	01/20/2023
Standard Type:	Internal Standard	Prepared:	01/20/2022
Solvent:	MeOH/61252	Prepared By:	Dipti Gokal
Final Volume (mLs):	12	Department:	PFAS
Vials:	1	Last Edit:	01/20/2022 15:49 by HGH
Comments:	mpfna had more left over than others.		

Analyte	Parent	CAS Number	Concentration	Units
13C3-PFBA	22A0116	13C3-PFBA	5	ug/mL
13C2-PFHxA	22A0117	13C2-PFHxA	5	ug/mL
13C5-PFNA	22A0118	13C5-PFNA	5	ug/mL
13C4-PFOA	22A0119	13C4-PFOA	5	ug/mL
13C2-PFDA	22A0120	13C2-PFDA	5	ug/mL
13C4-PFOS	22A0121	13C4-PFOS	5	ug/mL
18O2-PFHXS	22A0122	18O2-PFHXS	5	ug/mL

Parent Standards used:

Standard	Description	Prepared	Prepared By	Lot Nbr	Expires	Last Edit	(mLs)
22A0116	PFAS - IIS M3PFBA 50ug/mL	08/19/2021	Wellington Laboratories	M3PFBA0721	08/19/2026	01/20/2022 15:48 by HGH	1.2
22A0117	PFAS - IIS MPFHxA 50ug/mL	10/04/2021	Wellington Laboratories	MPFHxA0921	10/04/2026	01/20/2022 15:48 by HGH	1.2
22A0118	PFAS - IIS MPFNA 50ug/mL	10/29/2021	Wellington Laboratories	MPFNA1021	10/29/2026	01/20/2022 15:48 by HGH	1.2
22A0119	PFAS - IIS MPFOA 50ug/mL	12/07/2021	Wellington Laboratories	MPFOA1121	12/07/2026	01/20/2022 15:48 by HGH	1.2
22A0120	PFAS - IIS MPFDA 50ug/mL	12/08/2021	Wellington Laboratories	MPFDA1221	12/08/2026	01/20/2022 15:49 by HGH	1.2
22A0121	PFAS - IIS MPFOS 50ug/mL	08/18/2021	Wellington Laboratories	MPFOS0821	08/18/2026	01/20/2022 15:49 by HGH	1.2
22A0122	PFAS - IIS MPFHxS 50ug/mL	10/29/2021	Wellington Laboratories	MPFHxS1021	10/29/2026	01/20/2022 15:49 by HGH	1.2



WELLINGTON LABORATORIES

CERTIFICATE OF ANALYSIS DOCUMENTATION

PRODUCT CODE:

N-MeFOSE-M

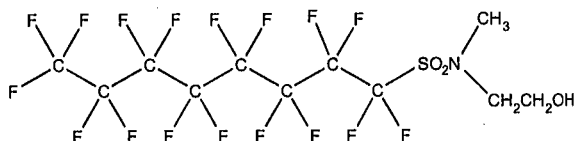
LOT NUMBER: NMeFOSE0921M**COMPOUND:**

2-(N-methylperfluoro-1-octanesulfonamido)-ethanol

22C0307

STRUCTURE:**CAS #:**

24448-09-7

**MOLECULAR FORMULA:** $C_{11}H_8F_{17}NO_3S$ **MOLECULAR WEIGHT:** 557.22**CONCENTRATION:** $50.0 \pm 2.5 \mu\text{g/mL}$ **SOLVENT(S):**

Methanol

CHEMICAL PURITY:

>98%

LAST TESTED: (mm/dd/yyyy)

09/22/2021 (HRGC/LRMS)

09/23/2021 (LC/MS)

EXPIRY DATE: (mm/dd/yyyy)

09/23/2026

RECOMMENDED STORAGE:

Store ampoule in a cool, dark place

DOCUMENTATION/ DATA ATTACHED:

Figure 1: HRGC/LRMS Data (Full Scan and Mass Spectrum)

Figure 2: LC/MS Data (Full Scan and Mass Spectrum)

Figure 3: LC/MS/MS Data (Selected MRM Transitions)

ADDITIONAL INFORMATION:

- See page 2 for further details.
- In order to see the molecular ion (adduct free), the LC mobile phase should be free of ammonium acetate buffer.

FOR LABORATORY USE ONLY: NOT FOR HUMAN OR DRUG USE

Certified By:

B.G. Chittim, General Manager

Date:

09/28/2021
(mm/dd/yyyy)

Wellington Laboratories Inc., 345 Southgate Dr. Guelph ON N1G 3M5 CANADA
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HANDLING:

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The combined relative standard uncertainty, $u_c(y)$, of a value y and the uncertainty of the independent parameters

x_1, x_2, \dots, x_n on which it depends is:

$$u_c(y(x_1, x_2, \dots, x_n)) = \sqrt{\sum_{i=1}^n u(y, x_i)^2}$$

where x is expressed as a relative standard uncertainty of the individual parameter.

The individual uncertainties taken into account include those associated with weights (calibration of the balance) and volumes (calibration of the volumetric glassware). An expanded maximum combined percent relative uncertainty of $\pm 5\%$ (calculated with a coverage factor of 2 and a level of confidence of 95%) is stated on the Certificate of Analysis for all of our products.

TRACEABILITY:

All reference standard solutions are traceable to specific crystalline lots. The microbalances used for solution preparation are regularly calibrated by an external ISO/IEC 17025 accredited laboratory. In addition, their calibration is verified prior to each weighing using calibrated external weights traceable to an ISO/IEC 17025 accredited laboratory. All volumetric glassware used is calibrated, of Class A tolerance, and traceable to an ISO/IEC 17025 accredited laboratory. For certain products, traceability to international interlaboratory studies has also been established.

EXPIRY DATE / PERIOD OF VALIDITY:

Ongoing stability studies of this product have demonstrated stability in its composition and concentration, until the specified expiry date, in the unopened ampoule. Monitoring for any degradation or change in concentration of the listed analyte(s) is performed on a routine basis.

LIMITED WARRANTY:

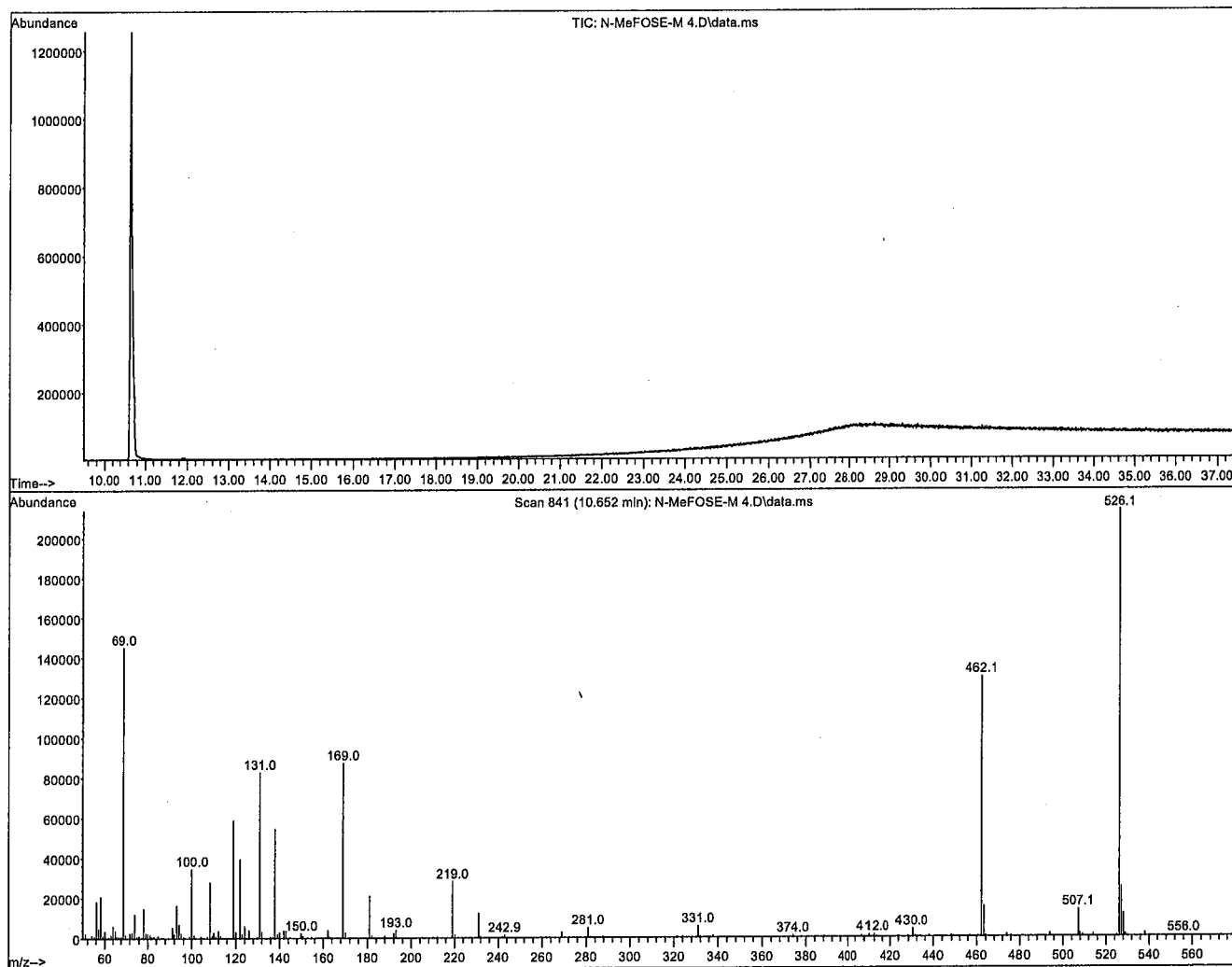
At the time of shipment, all products are warranted to be free of defects in material and workmanship and to conform to the stated technical and purity specifications.

QUALITY MANAGEMENT:

This product was produced using a Quality Management System registered to the latest versions of ISO 9001 by SAI Global, ISO/IEC 17025 by the Canadian Association for Laboratory Accreditation Inc. (CALA; A1226), and ISO 17034 by ANSI National Accreditation Board (ANAB; AR-1523).



For additional information or assistance concerning this or any other products from Wellington Laboratories Inc., please visit our website at www.well-labs.com or contact us directly at info@well-labs.com

Figure 1: N-MeFOSE-M; HRGC/LRMS Data (Full Scan and Mass Spectrum)**Conditions for Figure 1:**

Agilent 7890A HRGC
 Agilent 5975C MSD

Chromatographic Conditions:

Column: 30 m DB-5 (0.25 mm id, 0.25 μ m film thickness) Agilent J&W

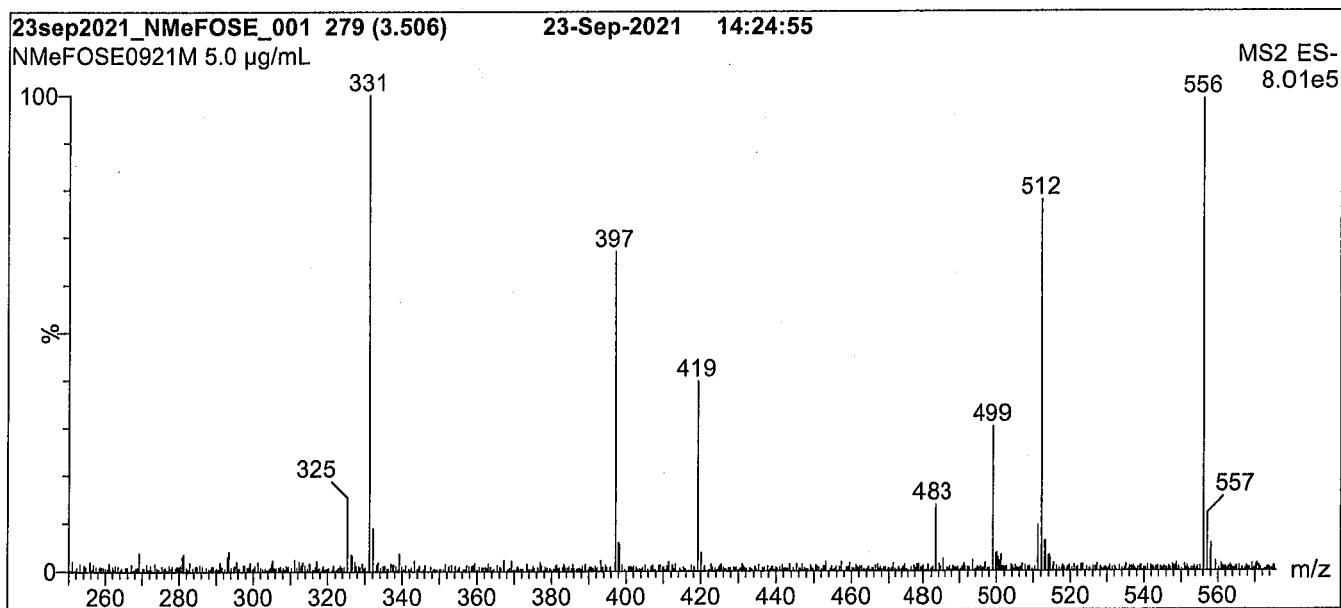
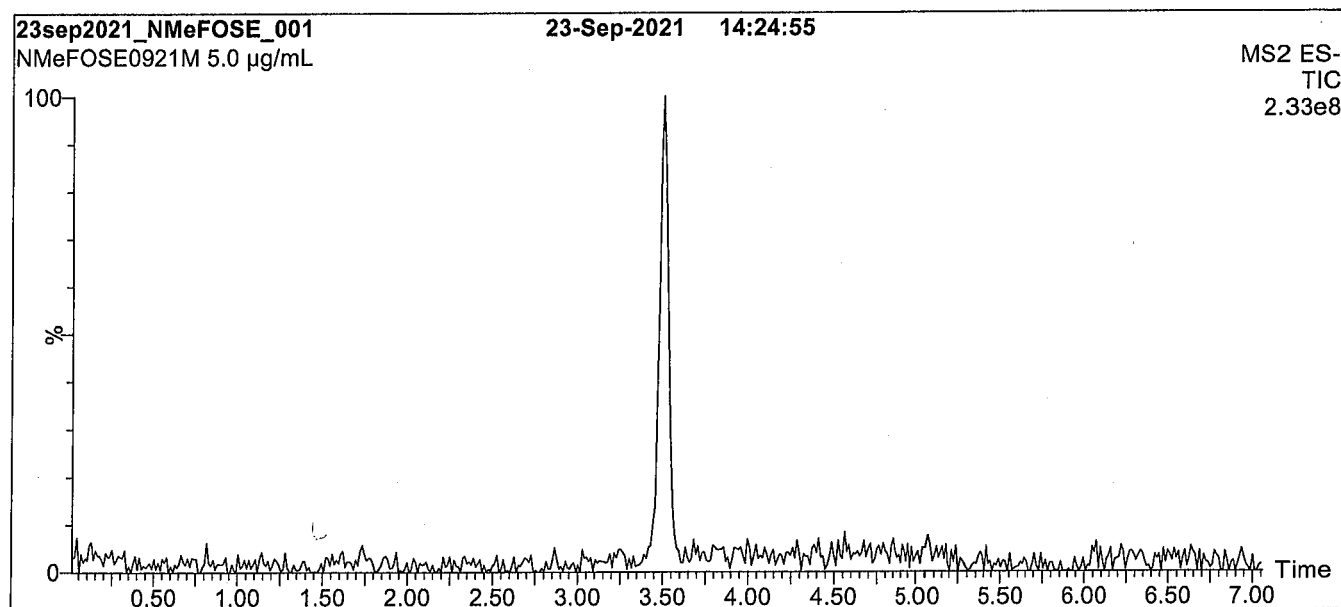
Flow: Constant at 1 mL/min

Injector: 250°C (Splitless Injection)

Oven: 100°C (5 min)
 10°C/min to 310°C
 310°C (10 min)

Ionization: EI+

Detector: 230°C
 Full Scan (50-1000 amu)

Figure 2: N-MeFOSE-M; LC/MS Data (Full Scan and Mass Spectrum)**Conditions for Figure 2:**

Waters Acquity Ultra Performance LC
Waters Xevo TQ-S micro MS

Chromatographic Conditions:

Column: Acquity UPLC BEH Shield RP₁₈
1.7 µm, 2.1 x 100 mm

Mobile phase: Gradient

Start: 30% H₂O / 70% MeOH

Ramp to 90% organic over 8 min and hold for
1.5 min before returning to initial conditions in 1 min.

Time: 12 min

Flow: 300 µL/min

MS Parameters:

Experiment: Full Scan (250 - 850 amu)

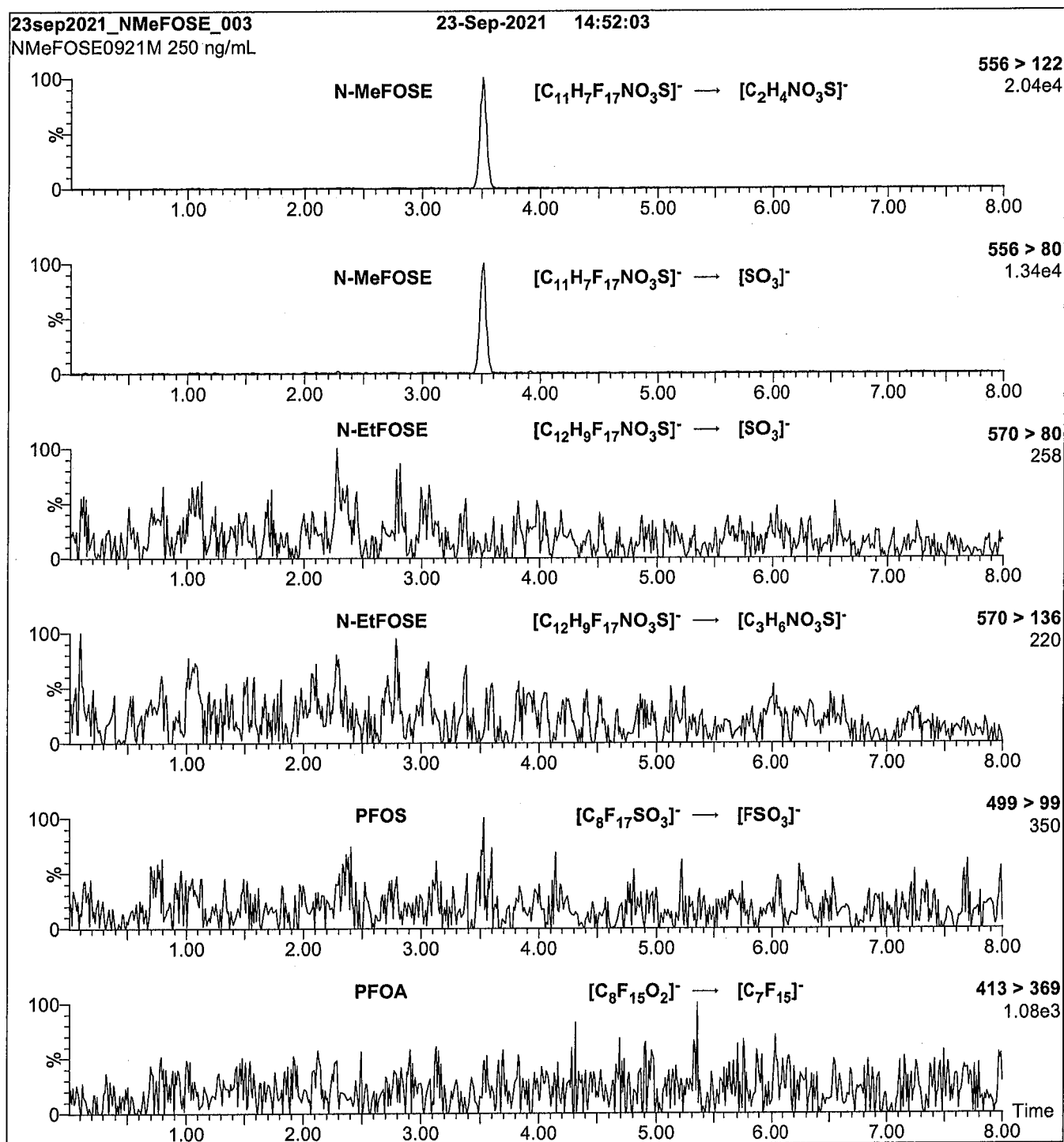
Source: Electrospray (negative)

Capillary Voltage (kV) = 2.00

Cone Voltage (V) = 65.00

Desolvation Temperature (°C) = 450

Desolvation Gas Flow (L/hr) = 1000

Figure 3: N-MeFOSE-M; LC/MS/MS Data (Selected MRM Transitions)**Conditions for Figure 3:**

Injection: On-column (N-MeFOSE-M)

Mobile phase: Same as Figure 2

Flow: 300 μ L/min**MS Parameters:**

Collision Gas (mbar) = 3.14e-3

Collision Energy (eV) = 36

g'

x

Analytical Standard Record

22C0307

Description:	PFAS - SAS N-MeFOSE 50ug/mL	Expires:	09/23/2026
Standard Type:	Analyte Spike	Prepared:	03/15/2022
Solvent:	Methanol	Prepared By:	Wellington Laboratories (Lot#: NMeFOSE0921M)
Final Volume (mLs):	1	Department:	PFAS
Vials:	1	Last Edit:	03/15/2022 15:59 by DAG

Analyte	Parent	CAS Number	Concentration	Units
NMeFOSE		24448-09-7	50	ug/mL



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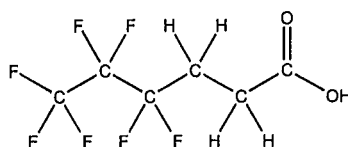
CERTIFICATE OF ANALYSIS DOCUMENTATION

PRODUCT CODE: FPrPA
COMPOUND: 3-Perfluoropropyl propanoic acid

LOT NUMBER: FPrPA0122
22C0308

STRUCTURE:

CAS #: 356-02-5



MOLECULAR FORMULA: $C_8H_5F_7O_2$
CONCENTRATION: $50.0 \pm 2.5 \mu\text{g/mL}$
CHEMICAL PURITY: >98%
LAST TESTED: (mm/dd/yyyy) 02/03/2022
EXPIRY DATE: (mm/dd/yyyy) 02/03/2027
RECOMMENDED STORAGE: Refrigerate ampoule

MOLECULAR WEIGHT: 242.09
SOLVENT(S): Methanol

DOCUMENTATION/ DATA ATTACHED:

Figure 1: LC/MS Data (Full Scan and Mass Spectrum)
Figure 2: LC/MS/MS Data (Selected MRM Transitions)

ADDITIONAL INFORMATION:

- See page 2 for further details.
- Contains <1% of the unsaturated 3:3 telomer acid ($C_8H_3F_7O_2$) as an impurity determined by ^{19}F NMR.

FOR LABORATORY USE ONLY: NOT FOR HUMAN OR DRUG USE

Certified By: 
B.G. Chittim, General Manager

Date: 02/04/2022
(mm/dd/yyyy)

Wellington Laboratories Inc., 345 Southgate Dr. Guelph ON N1G 3M5 CANADA
519-822-2436 • Fax: 519-822-2849 • info@well-labs.com

INTENDED USE:

The products prepared by Wellington Laboratories Inc. are for laboratory use only. This certified reference material (CRM) was designed to be used as a standard for the identification and/or quantification of the specific chemical compound it contains.

HANDLING:

This product should only be used by qualified personnel familiar with its potential hazards and trained in the handling of hazardous chemicals. Due care should be exercised to prevent unnecessary human contact or ingestion. All procedures should be carried out in a well-functioning fume hood and suitable gloves, eye protection, and clothing should be worn at all times. Waste should be disposed of according to national and regional regulations. Safety Data Sheets (SDSs) are available upon request.

SYNTHESIS / CHARACTERIZATION:

Our products are synthesized using single-product unambiguous routes whenever possible. They are then characterized, and their structures and purities confirmed, using a combination of the most relevant techniques, such as NMR, GC/MS, LC/MS/MS, SFC/UV/MS/MS, x-ray crystallography, and melting point. Isotopic purities of mass-labelled compounds are also confirmed using HRGC/HRMS and/or LC/MS/MS.

HOMOGENEITY:

Prior to solution preparation, crystalline material is tested for homogeneity using a variety of techniques (as stated above) and its solubility in a given diluent is taken into consideration. Duplicate solutions of a new product are prepared from the same crystalline lot and, after the addition of an appropriate internal standard, they are compared by GC/MS, LC/MS/MS, and/or SFC/UV/MS/MS. The relative response factors of the analyte of interest in each solution are required to be <5% RSD. New solution lots of existing products are compared to older lots in the same manner, which further confirms the homogeneity of the crystalline material as well as the stability and homogeneity of the solutions in the storage containers. In order to maintain the integrity of the assigned value(s), and associated uncertainty, the dilution or injection of a subsample of this product should be performed using calibrated measuring equipment.

UNCERTAINTY:

The maximum combined relative standard uncertainty of our reference standard solutions is calculated using the following equation:

The combined relative standard uncertainty, $u_c(y)$, of a value y and the uncertainty of the independent parameters

x_1, x_2, \dots, x_n on which it depends is:

$$u_c(y(x_1, x_2, \dots, x_n)) = \sqrt{\sum_{i=1}^n u(y, x_i)^2}$$

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TRACEABILITY:

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EXPIRY DATE / PERIOD OF VALIDITY:

Ongoing stability studies of this product have demonstrated stability in its composition and concentration, until the specified expiry date, in the unopened ampoule. Monitoring for any degradation or change in concentration of the listed analyte(s) is performed on a routine basis.

LIMITED WARRANTY:

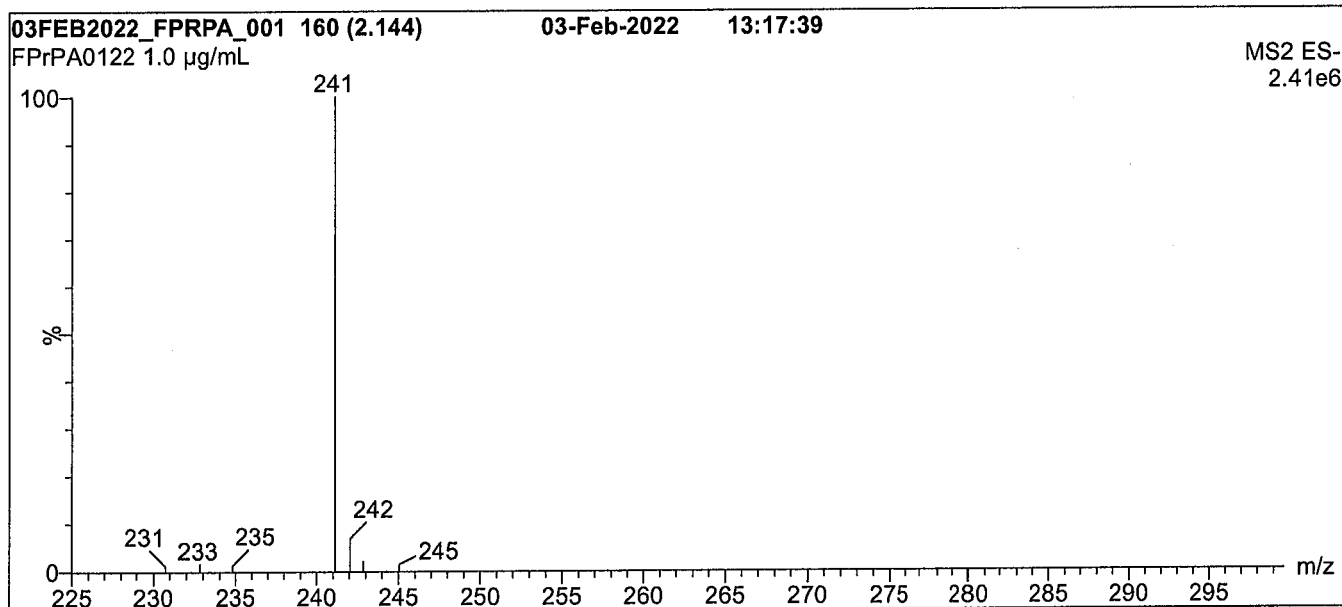
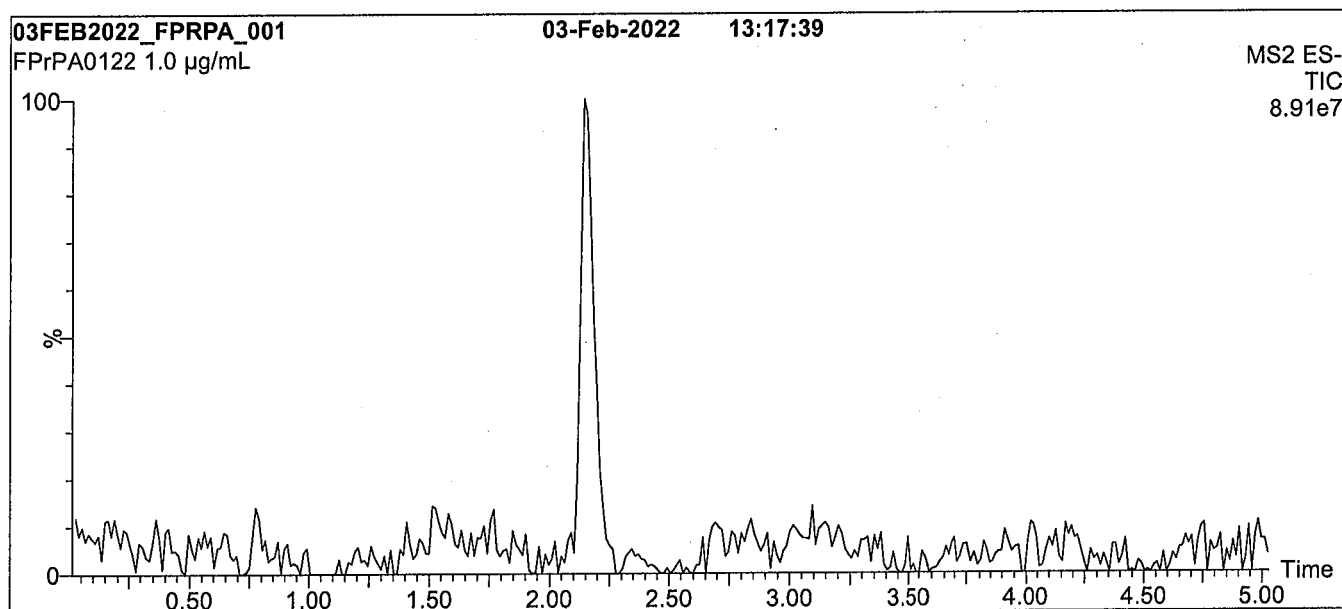
At the time of shipment, all products are warranted to be free of defects in material and workmanship and to conform to the stated technical and purity specifications.

QUALITY MANAGEMENT:

This product was produced using a Quality Management System registered to the latest versions of ISO 9001 by SAI Global, ISO/IEC 17025 by the Canadian Association for Laboratory Accreditation Inc. (CALA; A1226), and ISO 17034 by ANSI National Accreditation Board (ANAB; AR-1523).



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Figure 1: FPrPA; LC/MS Data (Full Scan and Mass Spectrum)**Conditions for Figure 1:**

Waters Acquity Ultra Performance LC
Waters Xevo TQ-S micro MS

Chromatographic Conditions:

Column: Acquity UPLC BEH Shield RP₁₈
1.7 µm, 2.1 x 100 mm

Mobile phase: Gradient

Start: 60% H₂O / 40% (80:20 MeOH:ACN)
(both with 10 mM NH₄OAc buffer)
Ramp to 90% organic over 8 min and hold for 2 min
before returning to initial conditions in 0.75 min.
Time: 12 min

Flow: 300 µL/min

MS Parameters:

Experiment: Full Scan (225 - 850 amu)

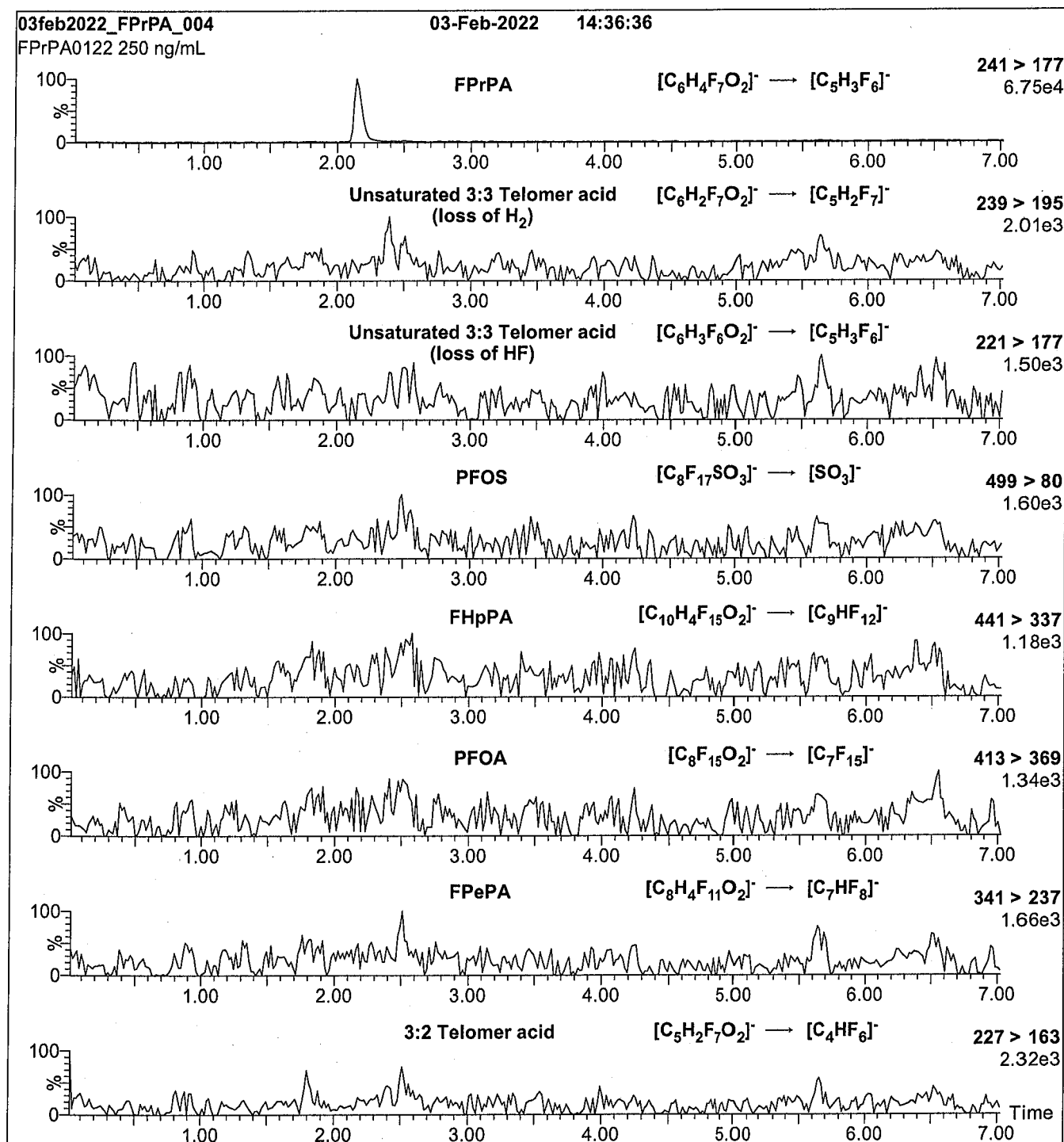
Source: Electrospray (negative)

Capillary Voltage (kV) = 2.00

Cone Voltage (V) = 10.00

Desolvation Temperature (°C) = 500

Desolvation Gas Flow (L/hr) = 1000

Figure 2: FPrPA; LC/MS/MS Data (Selected MRM Transitions)**Conditions for Figure 2:**

Injection: On-column (FPrPA)

Mobile phase: Same as Figure 1

Flow: 300 μ L/min**MS Parameters:**

Collision Gas (mbar) = 3.33e-3

Collision Energy (eV) = 10

Analytical Standard Record

22C0308

Description:	PFAS - SAS FPrPA 50ug/mL	Expires:	02/03/2027
Standard Type:	Analyte Spike	Prepared:	03/15/2022
Solvent:	Methanol	Prepared By:	Wellington Laboratories (Lot#: FPrPA0122)
Final Volume (mLs):	1	Department:	PFAS
Vials:	1	Last Edit:	03/15/2022 15:59 by DAG

Analyte	Parent	CAS Number	Concentration	Units
3:3FTCA		113507-82-7	50	ug/mL



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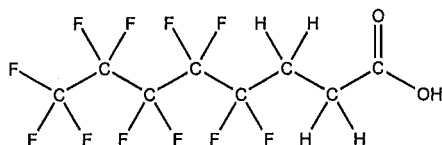
CERTIFICATE OF ANALYSIS DOCUMENTATION

PRODUCT CODE: FPePA
COMPOUND: 3-Perfluoropentyl propanoic acid

LOT NUMBER: FPePA1221
22C0309

STRUCTURE:

CAS #: 914637-49-3



MOLECULAR FORMULA: $C_8H_5F_{11}O_2$
CONCENTRATION: $50.0 \pm 2.5 \mu\text{g/mL}$
CHEMICAL PURITY: >98%
LAST TESTED: (mm/dd/yyyy) 01/05/2022
EXPIRY DATE: (mm/dd/yyyy) 01/05/2027
RECOMMENDED STORAGE: Refrigerate ampoule

MOLECULAR WEIGHT: 342.11
SOLVENT(S): Methanol

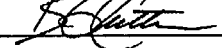
DOCUMENTATION/ DATA ATTACHED:

Figure 1: LC/MS Data (Full Scan and Mass Spectrum)
Figure 2: LC/MS/MS Data (Selected MRM Transitions)

ADDITIONAL INFORMATION:

- See page 2 for further details.
- Contains <0.5% of the unsaturated 5:3 telomer acid ($C_8H_3F_{11}O_2$) as an impurity determined by ^1H NMR.

FOR LABORATORY USE ONLY: NOT FOR HUMAN OR DRUG USE

Certified By: 
B.G. Chittim, General Manager

Date: 01/06/2022
(mm/dd/yyyy)

Wellington Laboratories Inc., 345 Southgate Dr. Guelph ON N1G 3M5 CANADA
519-822-2436 • Fax: 519-822-2849 • info@well-labs.com

INTENDED USE:

The products prepared by Wellington Laboratories Inc. are for laboratory use only. This certified reference material (CRM) was designed to be used as a standard for the identification and/or quantification of the specific chemical compound it contains.

HANDLING:

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HOMOGENEITY:

Prior to solution preparation, crystalline material is tested for homogeneity using a variety of techniques (as stated above) and its solubility in a given diluent is taken into consideration. Duplicate solutions of a new product are prepared from the same crystalline lot and, after the addition of an appropriate internal standard, they are compared by GC/MS, LC/MS/MS, and/or SFC/UV/MS/MS. The relative response factors of the analyte of interest in each solution are required to be <5% RSD. New solution lots of existing products are compared to older lots in the same manner, which further confirms the homogeneity of the crystalline material as well as the stability and homogeneity of the solutions in the storage containers. In order to maintain the integrity of the assigned value(s), and associated uncertainty, the dilution or injection of a subsample of this product should be performed using calibrated measuring equipment.

UNCERTAINTY:

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x_1, x_2, \dots, x_n on which it depends is:

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EXPIRY DATE / PERIOD OF VALIDITY:

Ongoing stability studies of this product have demonstrated stability in its composition and concentration, until the specified expiry date, in the unopened ampoule. Monitoring for any degradation or change in concentration of the listed analyte(s) is performed on a routine basis.

LIMITED WARRANTY:

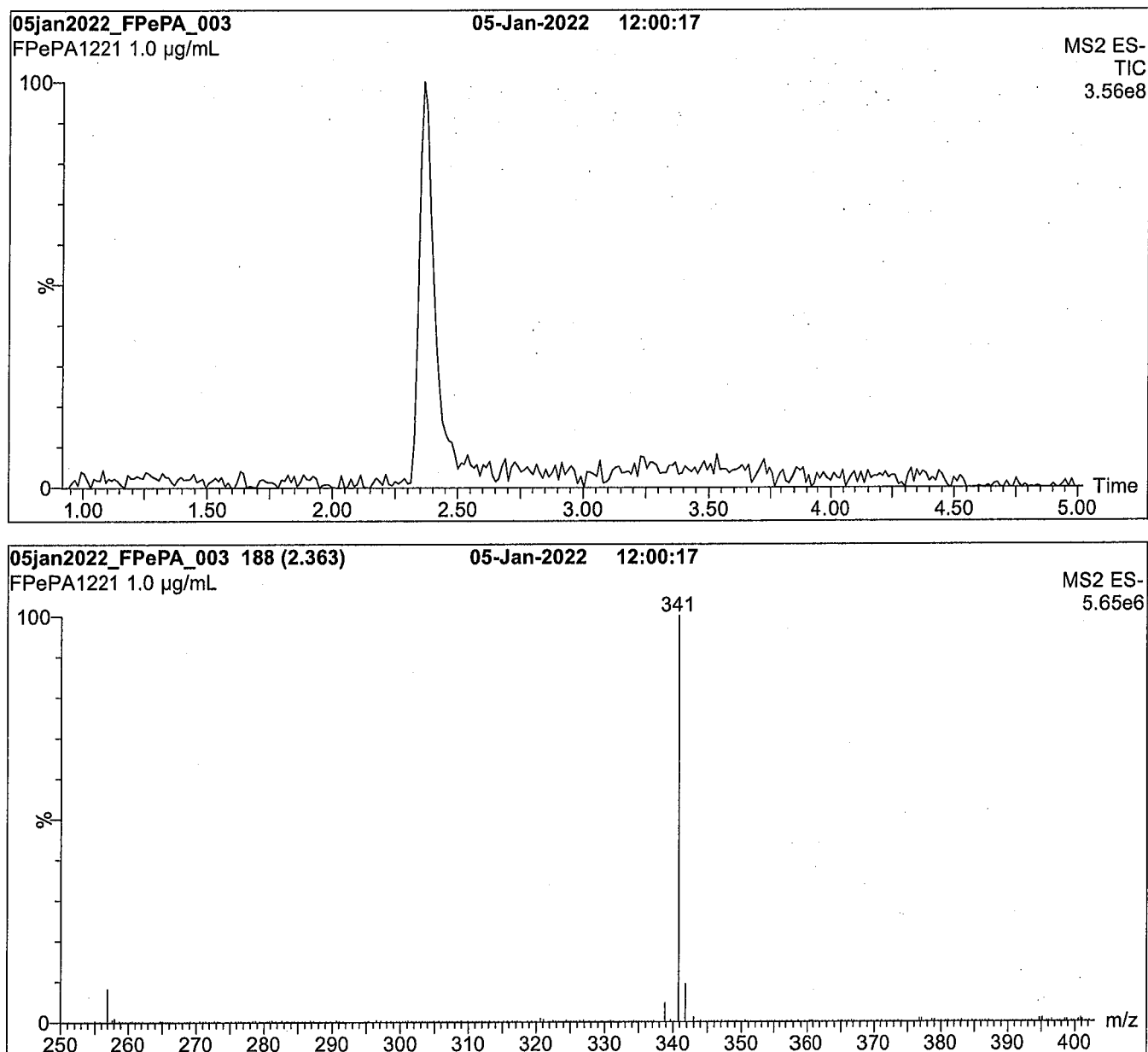
At the time of shipment, all products are warranted to be free of defects in material and workmanship and to conform to the stated technical and purity specifications.

QUALITY MANAGEMENT:

This product was produced using a Quality Management System registered to the latest versions of ISO 9001 by SAI Global, ISO/IEC 17025 by the Canadian Association for Laboratory Accreditation Inc. (CALA; A1226), and ISO 17034 by ANSI National Accreditation Board (ANAB; AR-1523).



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Figure 1: FPePA; LC/MS Data (Full Scan and Mass Spectrum)**Conditions for Figure 1:**

Waters Acquity Ultra Performance LC
Waters Xevo TQ-S micro MS

Chromatographic Conditions:

Column: Acquity UPLC BEH Shield RP₁₈
1.7 µm, 2.1 x 100 mm

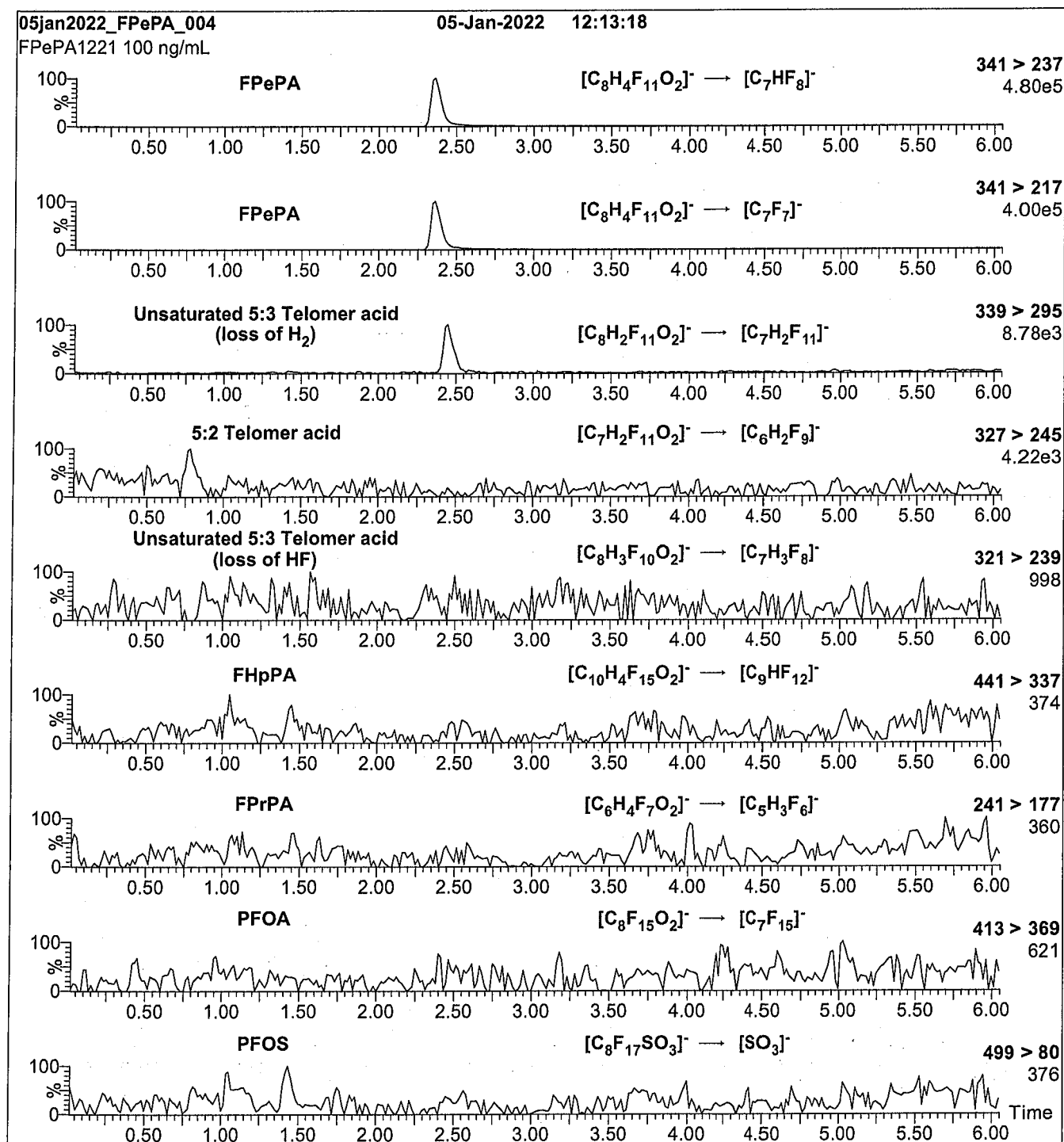
Mobile phase: Gradient
Start: 45% H₂O / 55% (80:20 MeOH:ACN)
(both with 10 mM NH₄OAc buffer)
Ramp to 90% organic over 7 min and hold for
3 min before returning to initial conditions in 0.75 min.
Time: 12 min

Flow: 300 µL/min

MS Parameters:

Experiment: Full Scan (250 - 850 amu)

Source: Electrospray (negative)
Capillary Voltage (kV) = 0.50
Cone Voltage (V) = 18.50
Desolvation Temperature (°C) = 500
Desolvation Gas Flow (L/hr) = 1000

Figure 2: FPePA; LC/MS/MS Data (Selected MRM Transitions)**Conditions for Figure 2:**

Injection: On-column (FPePA)

Mobile phase: Same as Figure 1

Flow: 300 μ L/min**MS Parameters:**

Collision Gas (mbar) = 3.09e-3

Collision Energy (eV) = 10

Analytical Standard Record

22C0309

Description:	PFAS - SAS FPePA 50ug/mL	Expires:	01/05/2027
Standard Type:	Analyte Spike	Prepared:	03/15/2022
Solvent:	Methanol	Prepared By:	Wellington Laboratories (Lot#:
Final Volume (mLs):	1	Department:	PFAS1221)
Vials:	1	Last Edit:	03/15/2022 15:59 by DAG

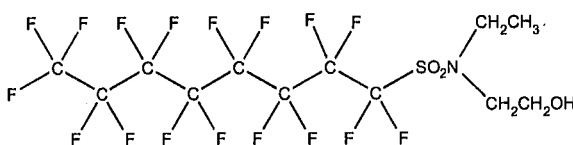
Analyte	Parent	CAS Number	Concentration	Units
5:3FTCA		914637-49-3	50	ug/mL



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CERTIFICATE OF ANALYSIS DOCUMENTATION

PRODUCT CODE: N-EtFOSE-M **LOT NUMBER:** NEtFOSE0921M
COMPOUND: 2-(N-ethylperfluoro-1-octanesulfonamido)-ethanol **22C0310**
STRUCTURE: **CAS #:** 1691-99-2



MOLECULAR FORMULA: $C_{12}H_{10}F_{17}NO_3S$ **MOLECULAR WEIGHT:** 571.25
CONCENTRATION: $50.0 \pm 2.5 \mu\text{g/mL}$ **SOLVENT(S):** Methanol
CHEMICAL PURITY: >98%
LAST TESTED: (mm/dd/yyyy) 09/22/2021 (HRGC/LRMS)
 09/23/2021 (LC/MS)
EXPIRY DATE: (mm/dd/yyyy) 09/23/2026
RECOMMENDED STORAGE: Store ampoule in a cool, dark place

DOCUMENTATION/ DATA ATTACHED:

Figure 1: HRGC/LRMS Data (Full Scan and Mass Spectrum)
 Figure 2: LC/MS Data (Full Scan and Mass Spectrum)
 Figure 3: LC/MS/MS Data (Selected MRM Transitions)

ADDITIONAL INFORMATION:

- See page 2 for further details.
- In order to see the molecular ion (adduct free), the LC mobile phase should be free of ammonium acetate buffer.

FOR LABORATORY USE ONLY: NOT FOR HUMAN OR DRUG USE

Certified By: 
 B.G. Chittim, General Manager

Date: 10/20/2021
 (mm/dd/yyyy)

Wellington Laboratories Inc., 345 Southgate Dr. Guelph ON N1G 3M5 CANADA
 519-822-2436 • Fax: 519-822-2849 • info@well-labs.com

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SYNTHESIS / CHARACTERIZATION:

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HOMOGENEITY:

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Ongoing stability studies of this product have demonstrated stability in its composition and concentration, until the specified expiry date, in the unopened ampoule. Monitoring for any degradation or change in concentration of the listed analyte(s) is performed on a routine basis.

LIMITED WARRANTY:

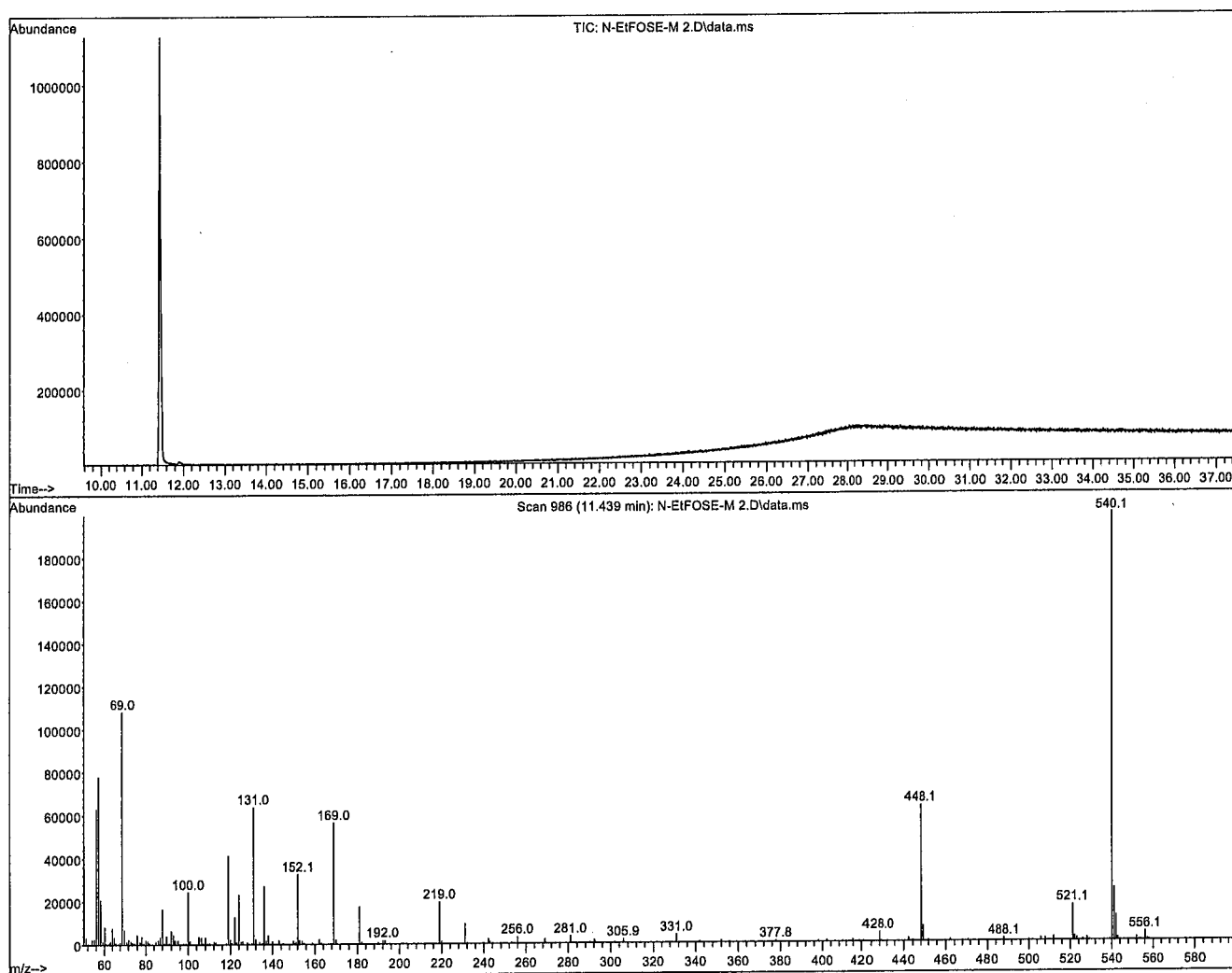
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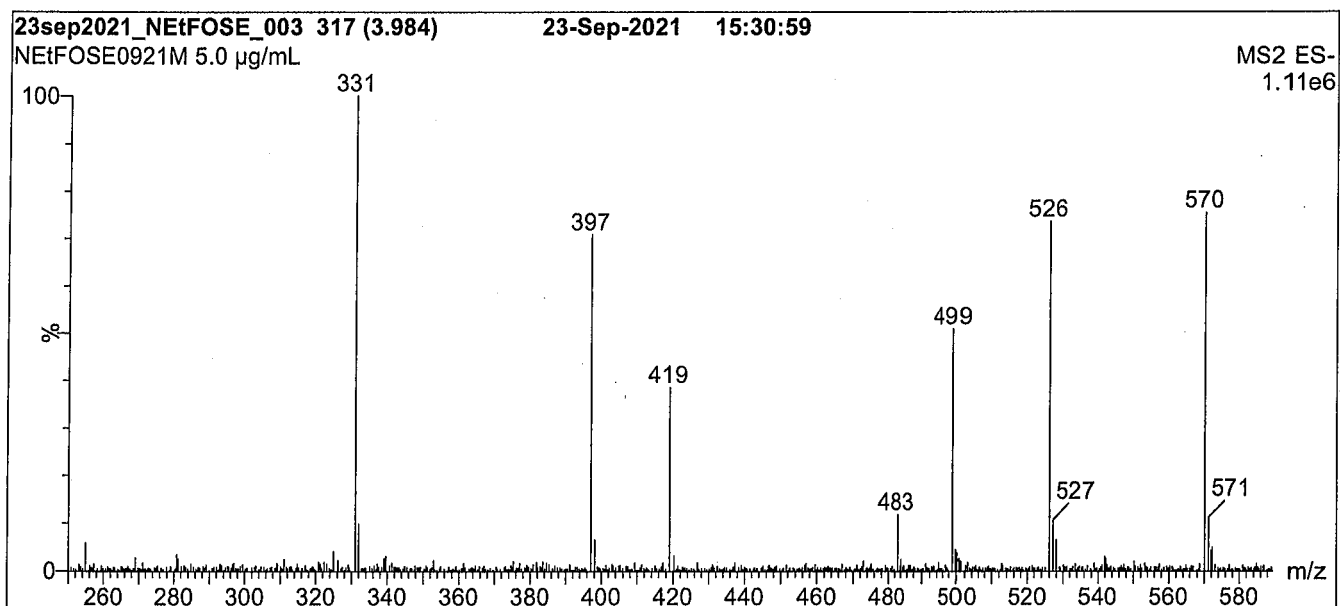
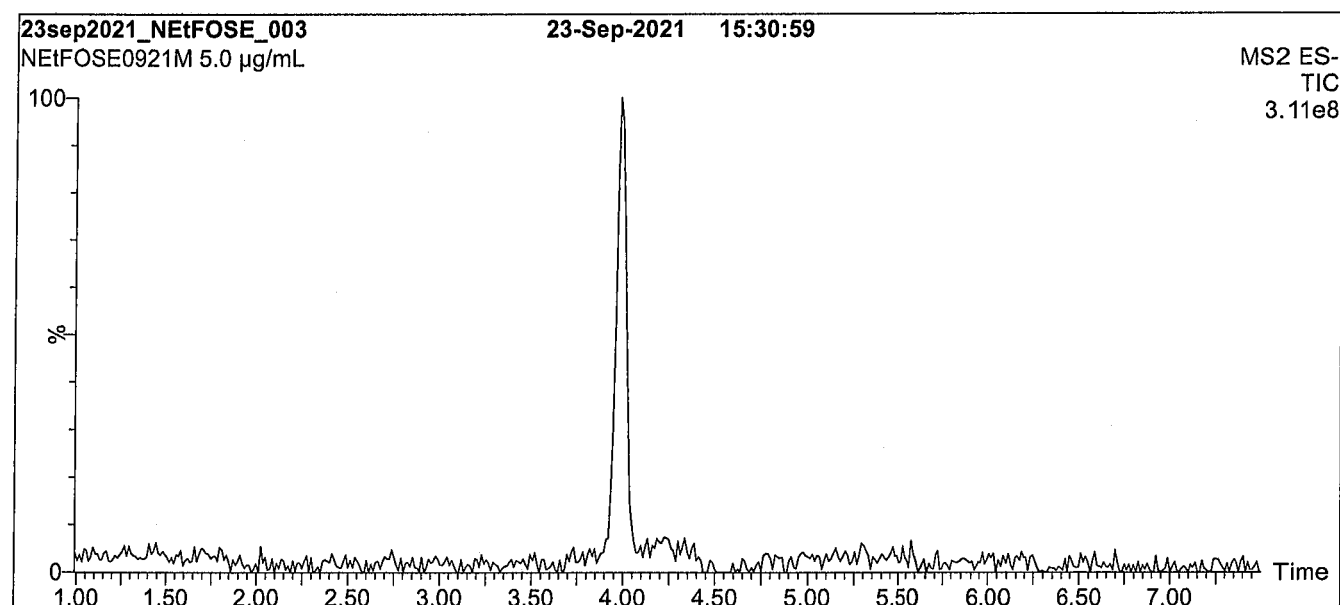
For additional information or assistance concerning this or any other products from Wellington Laboratories Inc., please visit our website at www.well-labs.com or contact us directly at info@well-labs.com

Figure 1: N-EtFOSE-M; HRGC/LRMS Data (Full Scan and Mass Spectrum)**Conditions for Figure 1:**

Agilent 7890A HRGC
Agilent 5975C MSD

Chromatographic Conditions:

Column: 30 m DB-5 (0.25 mm id, 0.25 μ m film thickness) Agilent J&W
Flow: Constant at 1 mL/min
Injector: 250°C (Splitless Injection)
Oven: 100°C (5 min)
10°C/min to 325°C
325°C (10 min)
Ionization: EI+
Detector: 230°C
Full Scan (50-1000 amu)

Figure 2: N-EtFOSE-M; LC/MS Data (Full Scan and Mass Spectrum)**Conditions for Figure 2:**

Waters Acquity Ultra Performance LC
Waters Xevo TQ-S micro MS

Chromatographic Conditions:

Column: Acquity UPLC BEH Shield RP₁₈
1.7 µm, 2.1 x 100 mm

Mobile phase: Gradient

Start: 30% H₂O / 70% MeOH

Ramp to 90% organic over 8 min and hold for
1.5 min before returning to initial conditions in 1 min.

Time: 12 min

Flow: 300 µL/min

MS Parameters:

Experiment: Full Scan (250 - 850 amu)

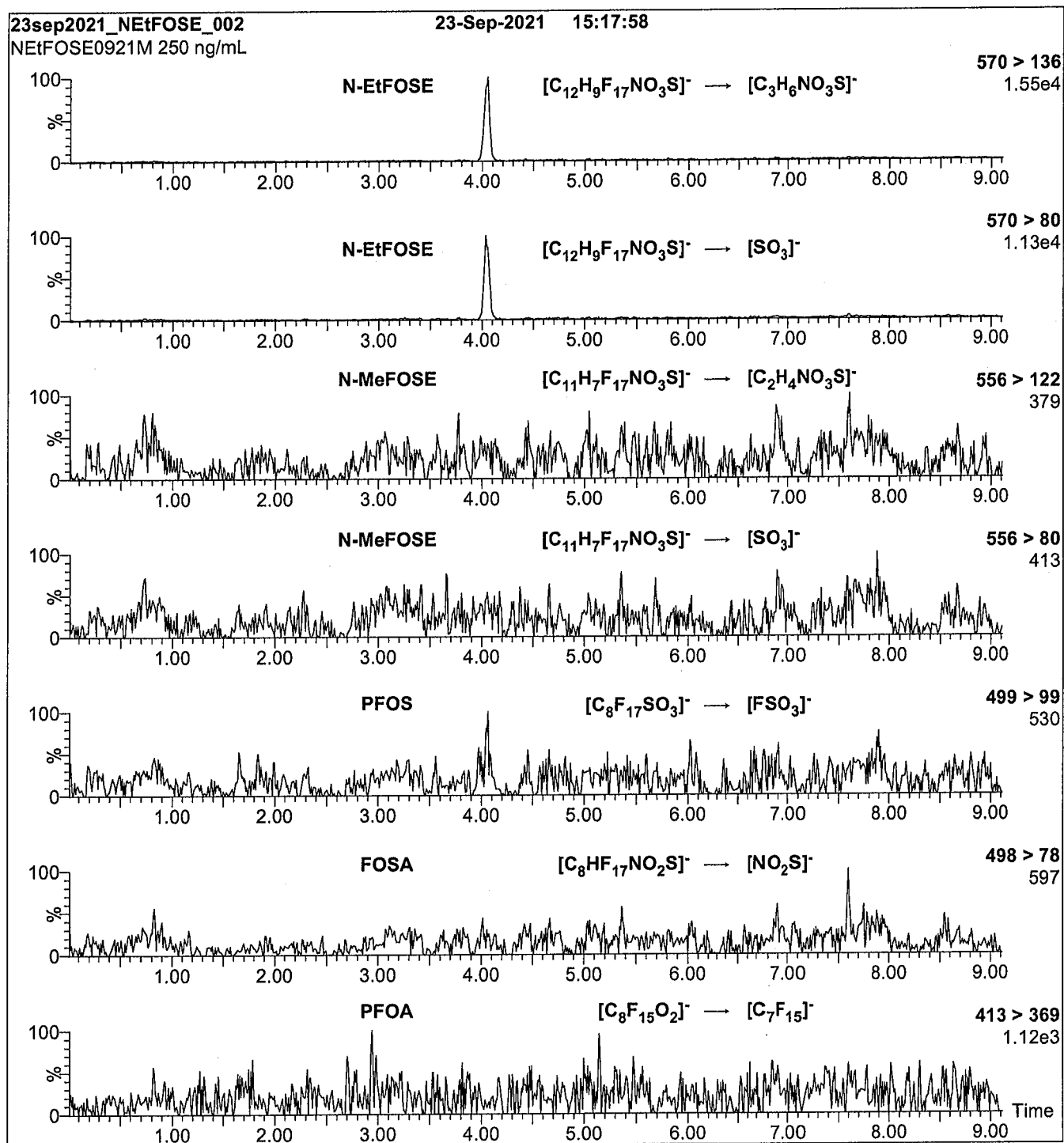
Source: Electrospray (negative)

Capillary Voltage (kV) = 2.00

Cone Voltage (V) = 65.00

Desolvation Temperature (°C) = 450

Desolvation Gas Flow (L/hr) = 1000

Figure 3: N-EtFOSE-M; LC/MS/MS Data (Selected MRM Transitions)**Conditions for Figure 3:**

Injection: On-column (N-EtFOSE-M)

Mobile phase: Same as Figure 2

Flow: 300 μ L/min**MS Parameters:**

Collision Gas (mbar) = 3.14e-3

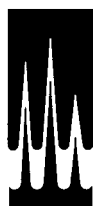
Collision Energy (eV) = 32

Analytical Standard Record

22C0310

Description:	PFAS - SAS NETFOSE 50ug/mL	Expires:	09/23/2026
Standard Type:	Analyte Spike	Prepared:	03/15/2022
Solvent:	Methanol	Prepared By:	Wellington Laboratories (Lot#:
Final Volume (mLs):	1	Department:	NETFOSE0921M)
Vials:	1	Last Edit:	03/15/2022 15:59 by DAG

Analyte	Parent	CAS Number	Concentration	Units
NETFOSE		1691-99-2	50	ug/mL



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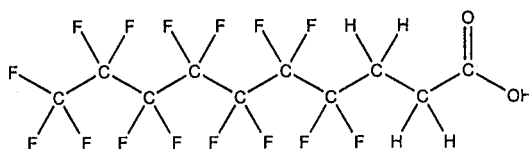
CERTIFICATE OF ANALYSIS DOCUMENTATION

PRODUCT CODE: FHpPA
COMPOUND: 3-Perfluoroheptyl propanoic acid

LOT NUMBER: FHpPA1020
22C0311

STRUCTURE:

CAS #: 812-70-4



MOLECULAR FORMULA: $C_{10}H_6F_{16}O_2$
CONCENTRATION: $50.0 \pm 2.5 \mu\text{g/mL}$
CHEMICAL PURITY: >98%
LAST TESTED: (mm/dd/yyyy) 11/12/2020
EXPIRY DATE: (mm/dd/yyyy) 11/12/2025
RECOMMENDED STORAGE: Refrigerate ampoule

MOLECULAR WEIGHT: 442.12
SOLVENT(S): Methanol

DOCUMENTATION/ DATA ATTACHED:

Figure 1: LC/MS Data (TIC and Mass Spectrum)
Figure 2: LC/MS/MS Data (Selected MRM Transitions)

ADDITIONAL INFORMATION:

- See page 2 for further details.

FOR LABORATORY USE ONLY: NOT FOR HUMAN OR DRUG USE

Certified By: 
B.G. Chittim, General Manager

Date: 11/27/2020
(mm/dd/yyyy)

Wellington Laboratories Inc., 345 Southgate Dr. Guelph ON N1G 3M5 CANADA
519-822-2436 • Fax: 519-822-2849 • info@well-labs.com

INTENDED USE:

The products prepared by Wellington Laboratories Inc. are for laboratory use only. This certified reference material (CRM) was designed to be used as a standard for the identification and/or quantification of the specific chemical compound it contains.

HANDLING:

This product should only be used by qualified personnel familiar with its potential hazards and trained in the handling of hazardous chemicals. Due care should be exercised to prevent unnecessary human contact or ingestion. All procedures should be carried out in a well-functioning fume hood and suitable gloves, eye protection, and clothing should be worn at all times. Waste should be disposed of according to national and regional regulations. Safety Data Sheets (SDSs) are available upon request.

SYNTHESIS / CHARACTERIZATION:

Our products are synthesized using single-product unambiguous routes whenever possible. They are then characterized, and their structures and purities confirmed, using a combination of the most relevant techniques, such as NMR, GC/MS, LC/MS/MS, SFC/UV/MS/MS, x-ray crystallography, and melting point. Isotopic purities of mass-labelled compounds are also confirmed using HRGC/HRMS and/or LC/MS/MS.

HOMOGENEITY:

Prior to solution preparation, crystalline material is tested for homogeneity using a variety of techniques (as stated above) and its solubility in a given diluent is taken into consideration. Duplicate solutions of a new product are prepared from the same crystalline lot and, after the addition of an appropriate internal standard, they are compared by GC/MS, LC/MS/MS, and/or SFC/UV/MS/MS. The relative response factors of the analyte of interest in each solution are required to be <5% RSD. New solution lots of existing products are compared to older lots in the same manner, which further confirms the homogeneity of the crystalline material as well as the stability and homogeneity of the solutions in the storage containers. In order to maintain the integrity of the assigned value(s), and associated uncertainty, the dilution or injection of a subsample of this product should be performed using calibrated measuring equipment.

UNCERTAINTY:

The maximum combined relative standard uncertainty of our reference standard solutions is calculated using the following equation:

The combined relative standard uncertainty, $u_c(y)$, of a value y and the uncertainty of the independent parameters

x_1, x_2, \dots, x_n on which it depends is:

$$u_c(y(x_1, x_2, \dots, x_n)) = \sqrt{\sum_{i=1}^n u(y, x_i)^2}$$

where x is expressed as a relative standard uncertainty of the individual parameter.

The individual uncertainties taken into account include those associated with weights (calibration of the balance) and volumes (calibration of the volumetric glassware). An expanded maximum combined percent relative uncertainty of $\pm 5\%$ (calculated with a coverage factor of 2 and a level of confidence of 95%) is stated on the Certificate of Analysis for all of our products.

TRACEABILITY:

All reference standard solutions are traceable to specific crystalline lots. The microbalances used for solution preparation are regularly calibrated by an external ISO/IEC 17025 accredited laboratory. In addition, their calibration is verified prior to each weighing using calibrated external weights traceable to an ISO/IEC 17025 accredited laboratory. All volumetric glassware used is calibrated, of Class A tolerance, and traceable to an ISO/IEC 17025 accredited laboratory. For certain products, traceability to international interlaboratory studies has also been established.

EXPIRY DATE / PERIOD OF VALIDITY:

Ongoing stability studies of this product have demonstrated stability in its composition and concentration, until the specified expiry date, in the unopened ampoule. Monitoring for any degradation or change in concentration of the listed analyte(s) is performed on a routine basis.

LIMITED WARRANTY:

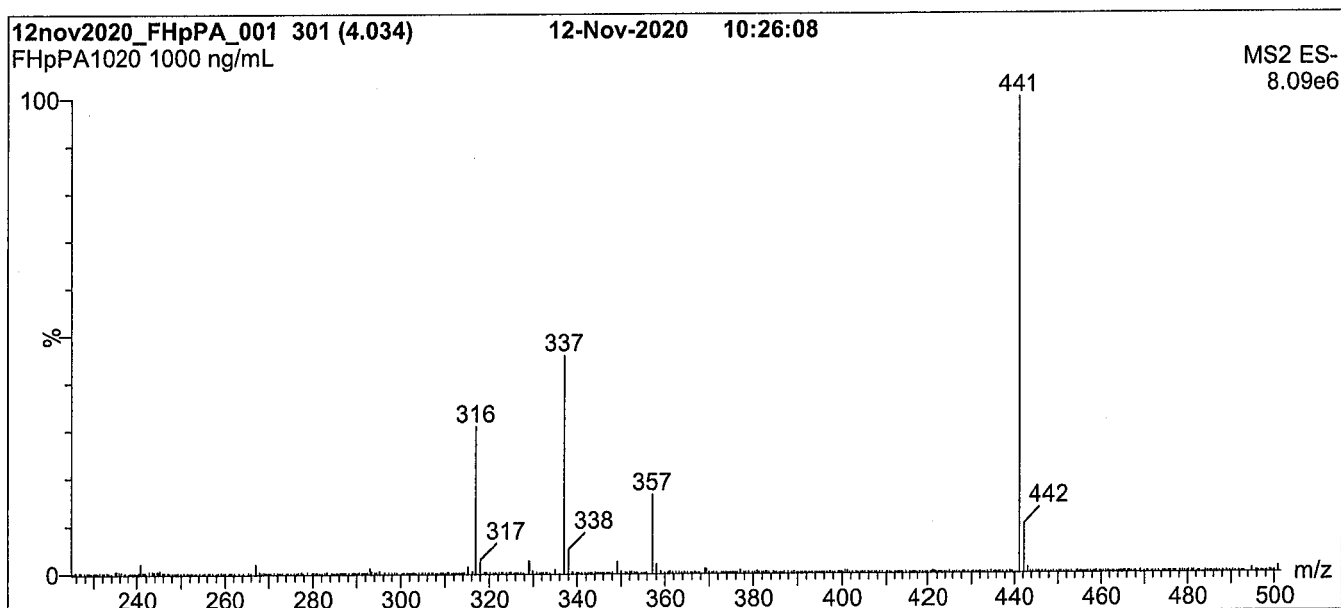
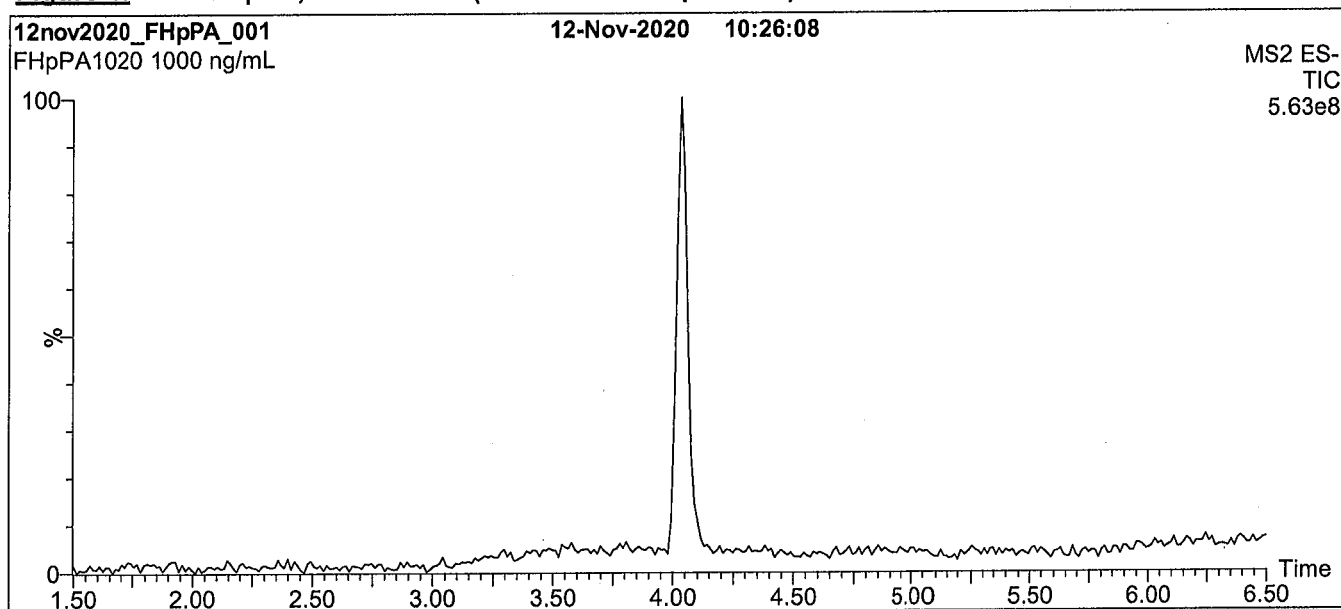
At the time of shipment, all products are warranted to be free of defects in material and workmanship and to conform to the stated technical and purity specifications.

QUALITY MANAGEMENT:

This product was produced using a Quality Management System registered to the latest versions of ISO 9001 by SAI Global, ISO/IEC 17025 by the Canadian Association for Laboratory Accreditation Inc. (CALA; A1226), and ISO 17034 by ANSI-ASQ National Accreditation Board (ANAB; AR-1523).



For additional information or assistance concerning this or any other products from Wellington Laboratories Inc., please visit our website at www.well-labs.com or contact us directly at info@well-labs.com

Figure 1: FHpPA; LC/MS Data (TIC and Mass Spectrum)**Conditions for Figure 1:**

Waters Acquity Ultra Performance LC
Waters Xevo TQ-S micro MS

Chromatographic Conditions:

Column: Acquity UPLC BEH Shield RP₁₈
1.7 μ m, 2.1 x 100 mm

Mobile phase: Gradient

Start: 45% H₂O / 55% (80:20 MeOH:ACN)
(both with 10 mM NH₄OAc buffer)
Ramp to 90% organic over 8 min and hold for
2 min before returning to initial conditions in 0.75 min.
Time: 12 min

Flow: 300 μ L/min

MS Parameters:

Experiment: Full Scan (225 - 850 amu)

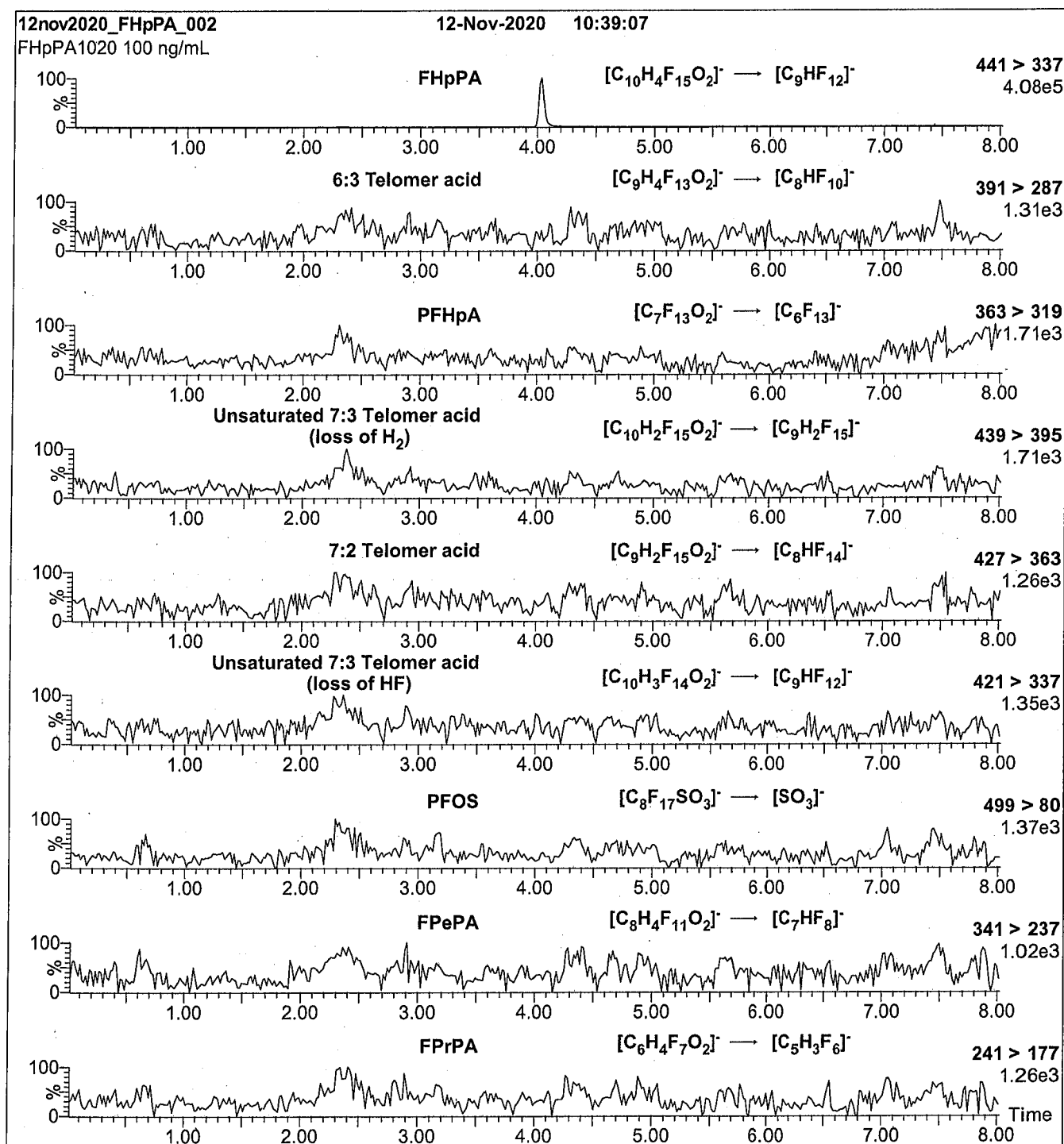
Source: Electrospray (negative)

Capillary Voltage (kV) = 0.50

Cone Voltage (V) = 28.50

Desolvation Temperature (°C) = 500

Desolvation Gas Flow (L/hr) = 1000

Figure 2: FHpPA; LC/MS/MS Data (Selected MRM Transitions)**Conditions for Figure 2:**

Injection: On-column (FHpPA)

Mobile phase: Same as Figure 1

Flow: 300 μ L/min**MS Parameters:**

Collision Gas (mbar) = 3.41e-3

Collision Energy (eV) = 8

Analytical Standard Record

22C0311

Description:	PFAS - SAS FHpPA 50ug/mL	Expires:	11/12/2025
Standard Type:	Analyte Spike	Prepared:	03/15/2022
Solvent:	Methanol	Prepared By:	Wellington Laboratories (Lot#: FHpPA1020)
Final Volume (mLs):	1	Department:	PFAS
Vials:	1	Last Edit:	03/15/2022 16:00 by DAG

Analyte	Parent	CAS Number	Concentration	Units
7:3FTCA		812-70-4	50	ug/mL

Analytical Standard Record

22C0311

Description:	PFAS - SAS FHpPA 50ug/mL	Expires:	11/12/2025
Standard Type:	Analyte Spike	Prepared:	03/15/2022
Solvent:	Methanol	Prepared By:	Wellington Laboratories (Lot#: FHpPA1020)
Final Volume (mLs):	1	Department:	PFAS
Vials:	1	Last Edit:	03/15/2022 16:00 by DAG

Analyte	Parent	CAS Number	Concentration	Units
7:3FTCA		812-70-4	50	ug/mL



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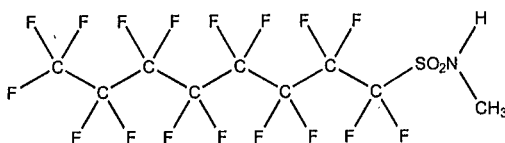
CERTIFICATE OF ANALYSIS DOCUMENTATION

PRODUCT CODE: N-MeFOSA-M
COMPOUND: N-methylperfluoro-1-octanesulfonamide

LOT NUMBER: NMeFOSA0721M
22C0312

STRUCTURE:

CAS #: 31506-32-8



MOLECULAR FORMULA: $C_9H_4F_{17}NO_2S$
CONCENTRATION: $50.0 \pm 2.5 \mu\text{g/mL}$
CHEMICAL PURITY: >98%
LAST TESTED: (mm/dd/yyyy) 08/03/2021
EXPIRY DATE: (mm/dd/yyyy) 08/03/2026
RECOMMENDED STORAGE: Store ampoule in a cool, dark place

MOLECULAR WEIGHT: 513.17
SOLVENT(S): Methanol

DOCUMENTATION/ DATA ATTACHED:

Figure 1: LC/MS Data (Full Scan and Mass Spectrum)
Figure 2: LC/MS/MS Data (Selected MRM Transitions)

ADDITIONAL INFORMATION:

- See page 2 for further details.

FOR LABORATORY USE ONLY: NOT FOR HUMAN OR DRUG USE

Certified By: 
B.G. Chittim, General Manager

Date: 08/04/2021
(mm/dd/yyyy)

Wellington Laboratories Inc., 345 Southgate Dr. Guelph ON N1G 3M5 CANADA
519-822-2436 • Fax: 519-822-2849 • info@well-labs.com

INTENDED USE:

The products prepared by Wellington Laboratories Inc. are for laboratory use only. This certified reference material (CRM) was designed to be used as a standard for the identification and/or quantification of the specific chemical compound it contains.

HANDLING:

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Prior to solution preparation, crystalline material is tested for homogeneity using a variety of techniques (as stated above) and its solubility in a given diluent is taken into consideration. Duplicate solutions of a new product are prepared from the same crystalline lot and, after the addition of an appropriate internal standard, they are compared by GC/MS, LC/MS/MS, and/or SFC/UV/MS/MS. The relative response factors of the analyte of interest in each solution are required to be <5% RSD. New solution lots of existing products are compared to older lots in the same manner, which further confirms the homogeneity of the crystalline material as well as the stability and homogeneity of the solutions in the storage containers. In order to maintain the integrity of the assigned value(s), and associated uncertainty, the dilution or injection of a subsample of this product should be performed using calibrated measuring equipment.

UNCERTAINTY:

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x_1, x_2, \dots, x_n on which it depends is:

$$u_c(y(x_1, x_2, \dots, x_n)) = \sqrt{\sum_{i=1}^n u(y, x_i)^2}$$

where x is expressed as a relative standard uncertainty of the individual parameter.

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EXPIRY DATE / PERIOD OF VALIDITY:

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LIMITED WARRANTY:

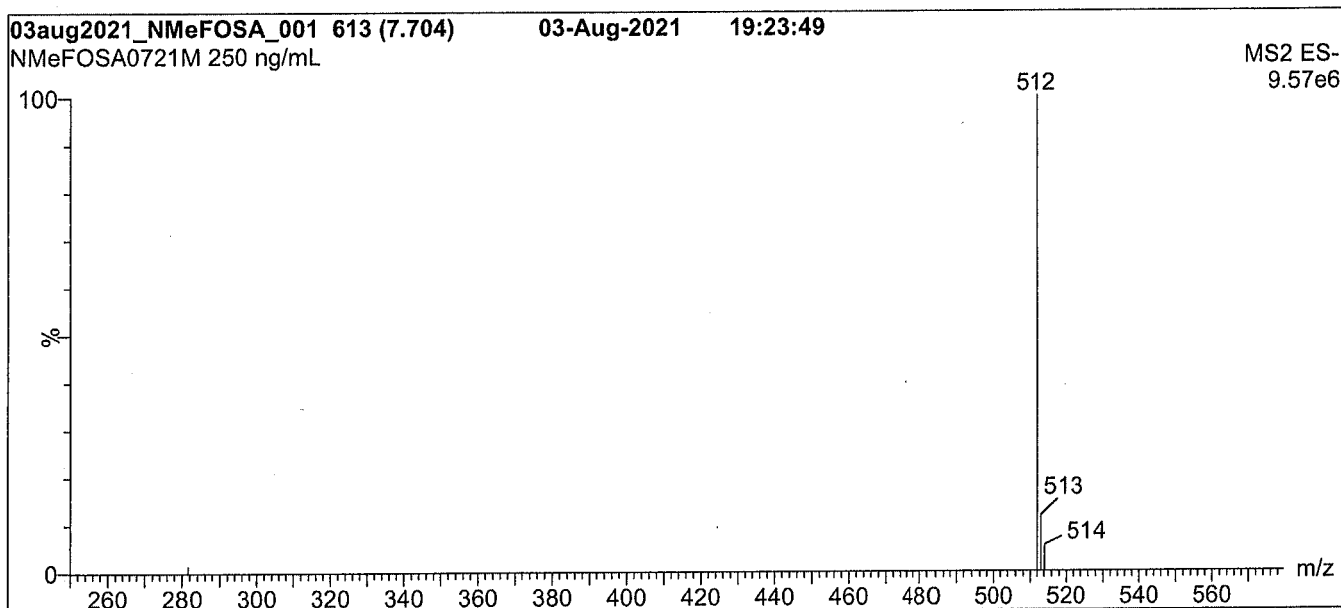
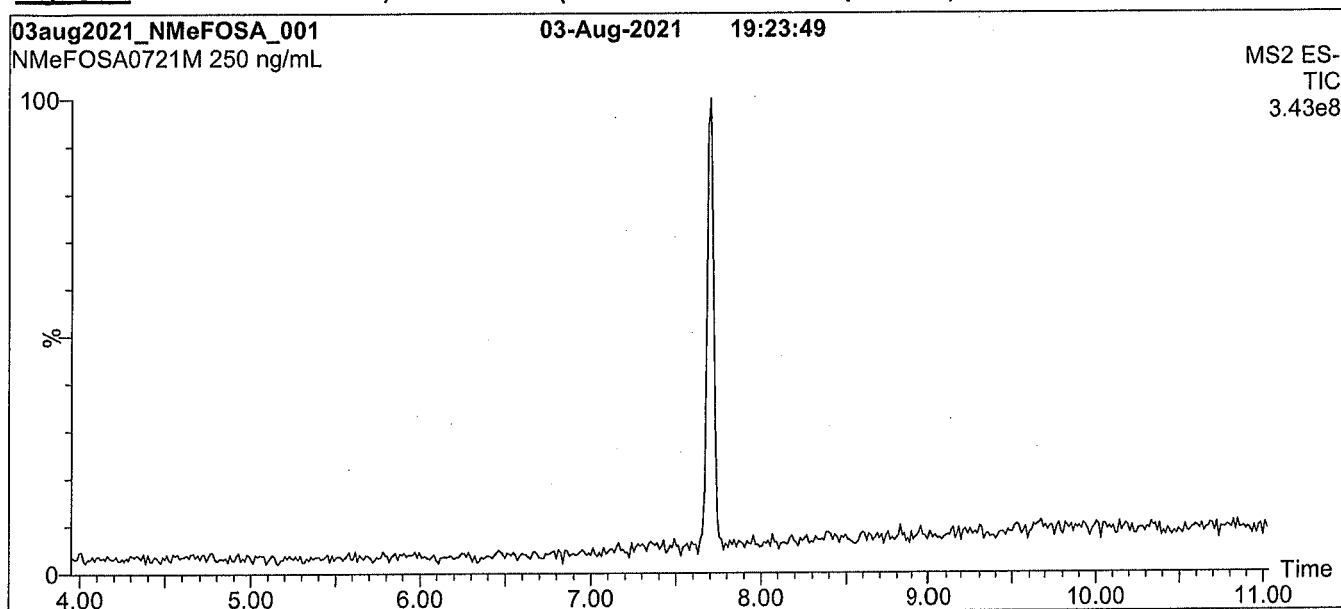
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For additional information or assistance concerning this or any other products from Wellington Laboratories Inc., please visit our website at www.well-labs.com or contact us directly at info@well-labs.com

Figure 1: N-MeFOSA-M; LC/MS Data (Full Scan and Mass Spectrum)**Conditions for Figure 1:**

Waters Acquity Ultra Performance LC
Waters Xevo TQ-S micro MS

Chromatographic Conditions:

Column: Acquity UPLC BEH Shield RP₁₈
1.7 μ m, 2.1 x 100 mm

Mobile phase: Gradient

Start: 40% H₂O / 60% (80:20 MeOH:ACN)
(both with 10 mM NH₄OAc buffer)
Ramp to 90% organic over 8 min and hold for
2 min before returning to initial conditions in 0.75 min.
Time: 12 min

Flow: 300 μ L/min

MS Parameters:

Experiment: Full Scan (250 - 850 amu)

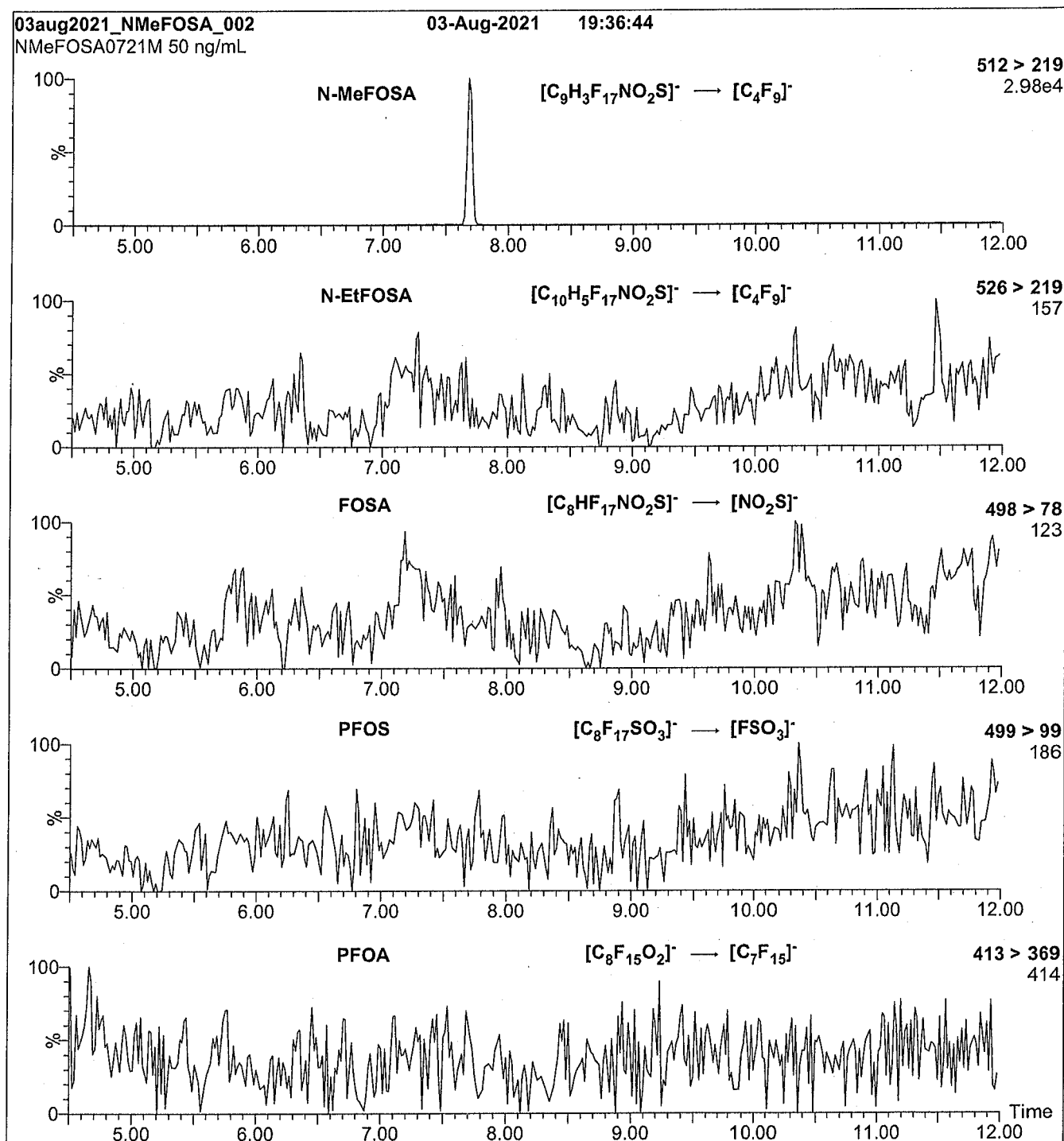
Source: Electrospray (negative)

Capillary Voltage (kV) = 1.00

Cone Voltage (V) = 44.00

Desolvation Temperature (°C) = 500

Desolvation Gas Flow (L/hr) = 1000

Figure 2: N-MeFOSA-M; LC/MS/MS Data (Selected MRM Transitions)**Conditions for Figure 2:**

Injection: On-column (N-MeFOSA-M)

Mobile phase: Same as Figure 1

Flow: 300 μ L/min**MS Parameters:**

Collision Gas (mbar) = 3.41e-3

Collision Energy (eV) = 24

Analytical Standard Record

22C0312

Description: PFAS - SAS NMeFOSA 50ug/mL
Standard Type: Analyte Spike
Solvent: Methanol
Final Volume (mLs): 1
Vials: 1

Expires: 08/03/2026
Prepared: 03/15/2022
Prepared By: Dipti Gokal
Department: PFAS
Last Edit: 03/15/2022 16:00 by DAG

Analyte	Parent	CAS Number	Concentration	Units
NMeFOSA		31506-32-8	50	ug/mL



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CERTIFICATE OF ANALYSIS DOCUMENTATION

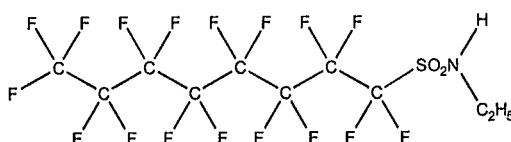
PRODUCT CODE: N-EtFOSA-M
COMPOUND: N-ethylperfluoro-1-octanesulfonamide

LOT NUMBER: NEtFOSA0821M

22C0313

STRUCTURE:

CAS #: 4151-50-2



MOLECULAR FORMULA: $C_{10}H_{17}F_{17}NO_2S$
CONCENTRATION: $50.0 \pm 2.5 \mu\text{g/mL}$
CHEMICAL PURITY: >98%

MOLECULAR WEIGHT: 527.20
SOLVENT(S): Methanol

LAST TESTED: (mm/dd/yyyy) 08/12/2021
EXPIRY DATE: (mm/dd/yyyy) 08/12/2026
RECOMMENDED STORAGE: Store ampoule in a cool, dark place

DOCUMENTATION/ DATA ATTACHED:

Figure 1: LC/MS Data (Full Scan and Mass Spectrum)
Figure 2: LC/MS/MS Data (Selected MRM Transitions)

ADDITIONAL INFORMATION:

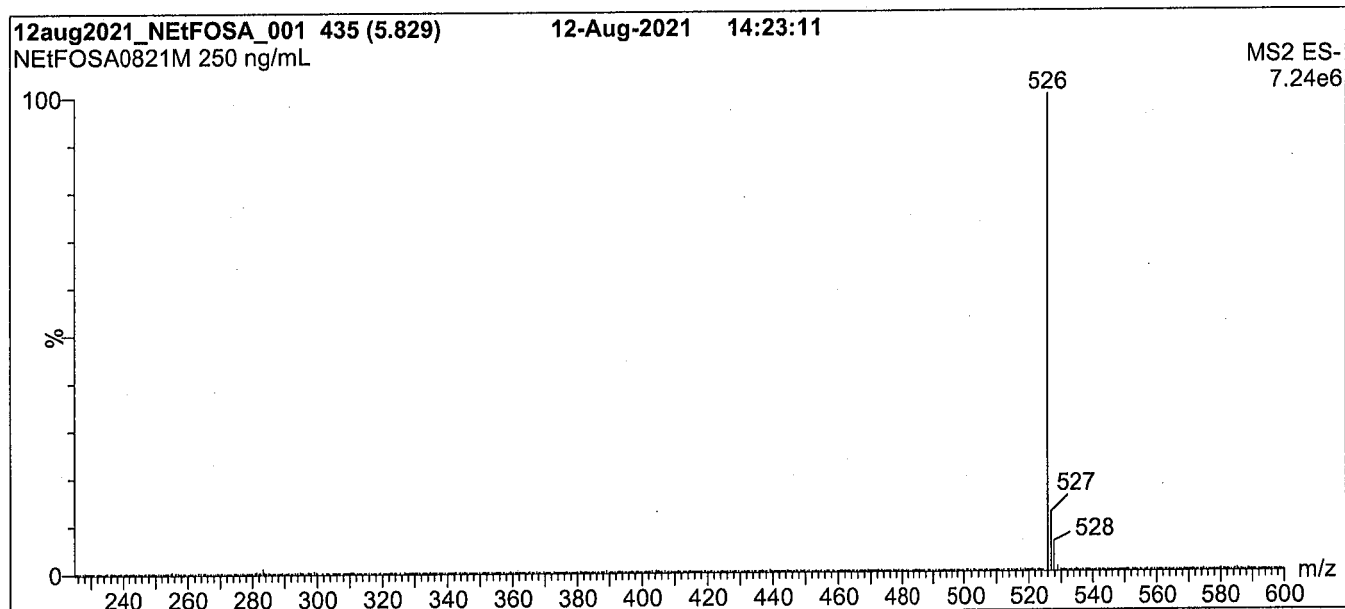
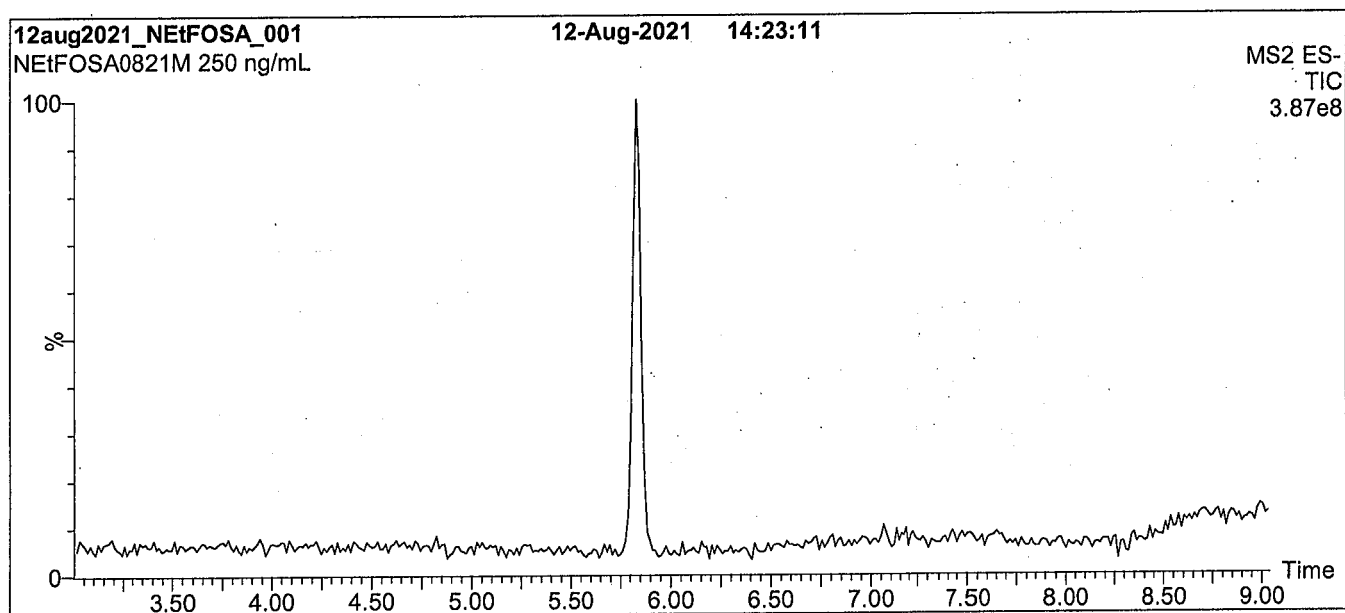
- See page 2 for further details.

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Certified By: 
B.G. Chittim, General Manager

Date: 08/16/2021
(mm/dd/yyyy)

Wellington Laboratories Inc., 345 Southgate Dr. Guelph ON N1G 3M5 CANADA
519-822-2436 • Fax: 519-822-2849 • info@well-labs.com

Figure 1: N-EtFOSA-M; LC/MS Data (Full Scan and Mass Spectrum)**Conditions for Figure 1:**

Waters Acquity Ultra Performance LC
Waters Xevo TQ-S micro MS

Chromatographic Conditions:

Column: Acquity UPLC BEH Shield RP₁₈
1.7 μ m, 2.1 x 100 mm

Mobile phase: Gradient

Start: 30% H₂O / 70% (80:20 MeOH:ACN)
(both with 10 mM NH₄OAc buffer)
Ramp to 90% organic over 8 min and hold for
2 min before returning to initial conditions in 0.75 min.
Time: 12 min

Flow: 300 μ L/min

MS Parameters:

Experiment: Full Scan (225 - 850 amu)

Source: Electrospray (negative)

Capillary Voltage (kV) = 1.00

Cone Voltage (V) = 44.00

Desolvation Temperature (°C) = 500

Desolvation Gas Flow (L/hr) = 1000

Analytical Standard Record

22C0313

Description:	PFAS - SAS NETFOSA 50ug/mL	Expires:	08/12/2026
Standard Type:	Other	Prepared:	03/15/2022
Solvent:	Methanol	Prepared By:	Wellington Laboratories (Lot#:
Final Volume (mL):	1	Department:	NETFOSA0821M)
Vials:	1	Last Edit:	08/17/2022 10:49 by LYA

Analyte	Parent	CAS Number	Concentration	Units
NETFOSA		4151-50-2	50	ug/mL

**WELLINGTON**
LABORATORIES**CERTIFICATE OF ANALYSIS**
DOCUMENTATION**PFAC-MXF** 22F0058**Native Replacement PFAS
Solution/Mixture**

PRODUCT CODE: PFAC-MXF
LOT NUMBER: PFACMXF0122
SOLVENT(S): Methanol / Water (<1%)
DATE PREPARED: (mm/dd/yyyy) 01/10/2022
LAST TESTED: (mm/dd/yyyy) 01/11/2022
EXPIRY DATE: (mm/dd/yyyy) 01/11/2025
RECOMMENDED STORAGE: Refrigerate ampoule

DESCRIPTION:

PFAC-MXF is a solution/mixture of sodium dodecafluoro-3H-4,8-dioxanonanoate (NaDONA), the major and minor components of F-53B (9Cl-PF3ONS and 11Cl-PF3OUdS), and GenX (HFPO-DA). The components and their concentrations are given in Table A.

The individual native components of this mixture all have chemical purities of >98%.

DOCUMENTATION/ DATA ATTACHED:

Table A: Components and Concentrations of the Solution/Mixture
Figure 1: LC/MS Data (SIR)
Figure 2: LC/MS/MS Data (Selected MRM Transitions)

ADDITIONAL INFORMATION:

- See page 2 for further details.
- Contains 4 mole eq. of NaOH to prevent conversion of the carboxylic acid to the methyl ester.

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519-822-2436 • Fax: 519-822-2849 • info@well-labs.com

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HANDLING:

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UNCERTAINTY:

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x_1, x_2, \dots, x_n on which it depends is:

$$u_c(y(x_1, x_2, \dots, x_n)) = \sqrt{\sum_{i=1}^n u(y, x_i)^2}$$

where x is expressed as a relative standard uncertainty of the individual parameter.

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EXPIRY DATE / PERIOD OF VALIDITY:

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QUALITY MANAGEMENT:

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For additional information or assistance concerning this or any other products from Wellington Laboratories Inc., please visit our website at www.well-labs.com or contact us directly at info@well-labs.com

Table A: PFAC-MXF; Components and Concentrations (ng/mL; \pm 5% in Methanol/Water (<1%))

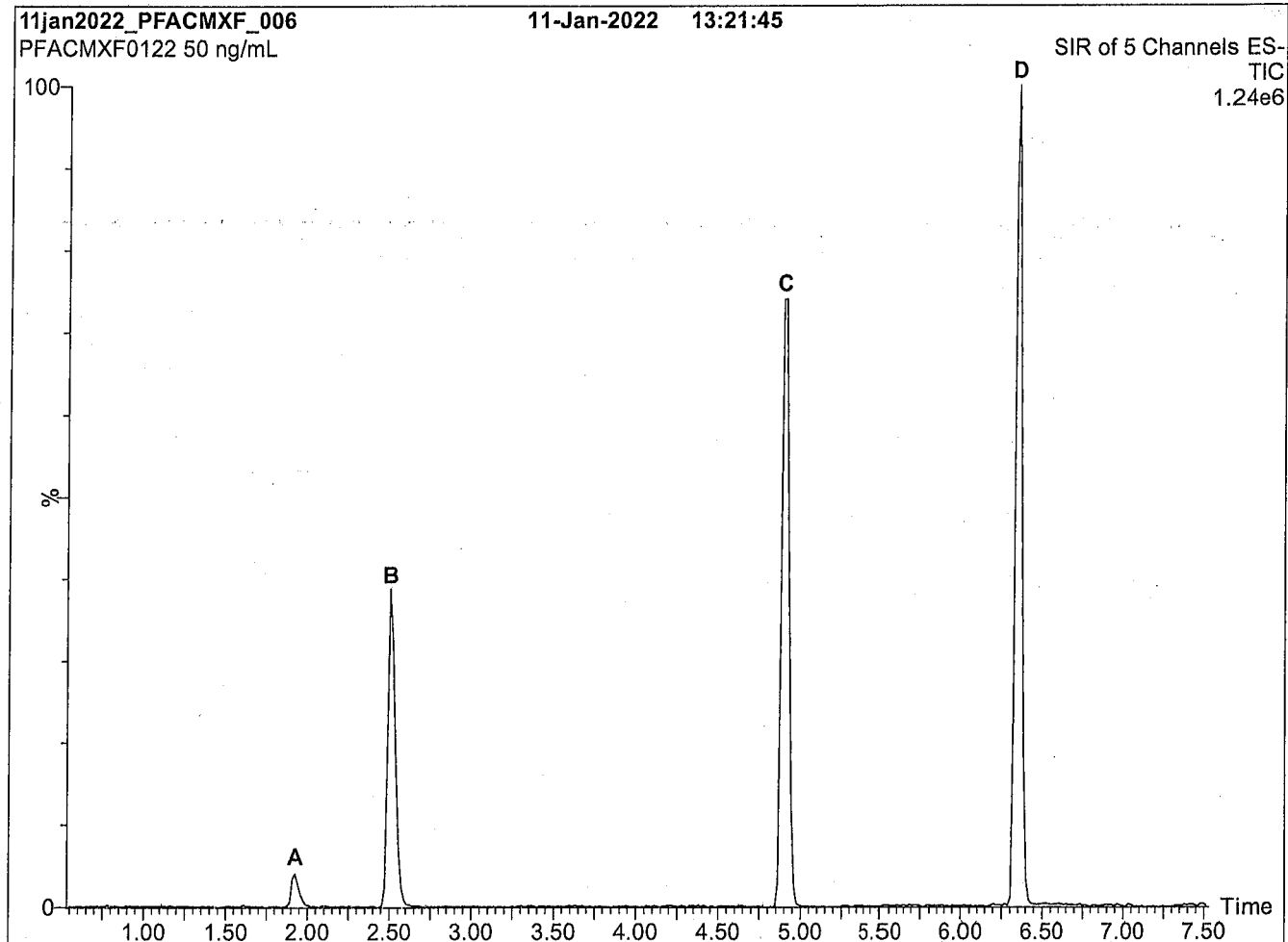
Compound	Acronym	Concentration* (ng/ml)		Peak Assignment in Figure 1
2,3,3,3-Tetrafluoro-2-(1,1,2,2,3,3,3-heptafluoropropoxy)-propanoic acid	HFPO-DA	2000		A
Compound	Acronym	Concentration* (ng/mL)		Peak Assignment in Figure 1
		as the salt	as the acid	
Sodium dodecafluoro-3H-4,8-dioxanonoate	NaDONA	2000	1890	B
Potassium 9-chlorohexadecafluoro-3-oxanonane-1-sulfonate	9CI-PF3ONS	2000	1870	C
Potassium 11-chloroeicosafluoro-3-oxaundecane-1-sulfonate	11CI-PF3OUdS	2000	1890	D

* Concentrations have been rounded to three significant figures.

Certified By: 

B.G. Chittim, General Manager

Date: 01/12/2022
(mm/dd/yyyy)

Figure 1: PFAC-MXF; LC/MS Data (SIR)**Conditions for Figure 1:**

Waters Acquity Ultra Performance LC
Waters Xevo TQ-S micro MS

Chromatographic Conditions:

Column: Acquity UPLC BEH Shield RP₁₈
1.7 μ m, 2.1 x 100 mm

Mobile phase: Gradient

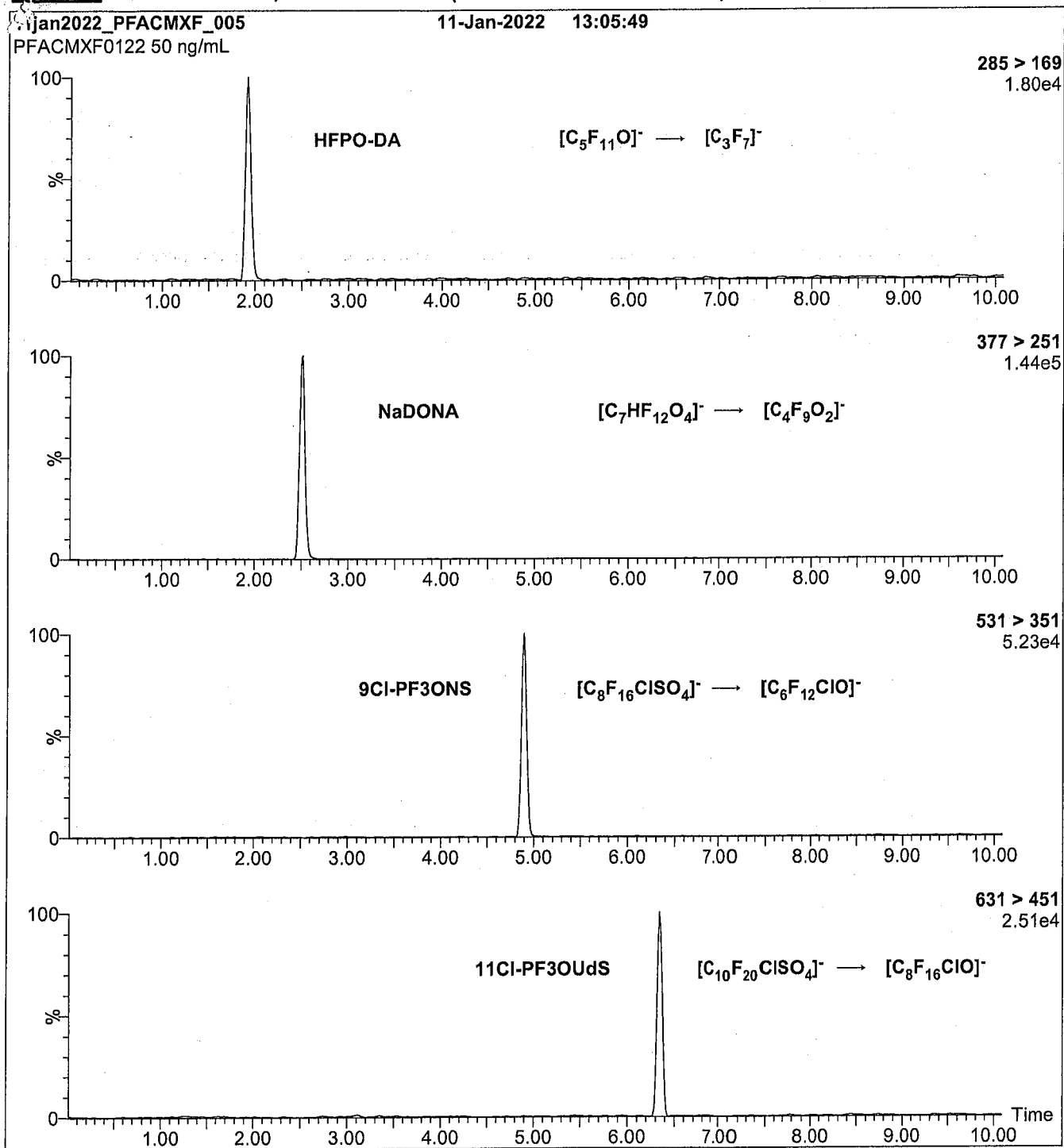
Start: 45% H₂O / 55% (80:20 MeOH:ACN)
(both with 10 mM NH₄OAc buffer)
Ramp to 90% organic over 8 min and hold for 2 min
before returning to initial conditions in 0.75 min.
Time: 12 min

Flow: 300 μ L/min

MS Parameters:

Experiment: SIR

Source: Electrospray (negative)
Capillary Voltage (kV) = 2.00
Cone Voltage (V) = variable (15-74)
Desolvation Temperature (°C) = 325
Desolvation Gas Flow (L/hr) = 1000

Figure 2: PFAC-MXF; LC/MS/MS Data (Selected MRM Transitions)**Conditions for Figure 2:**

Injection: On-column (PFAC-MXF)

Mobile phase: Same as Figure 1

Flow: 300 μ L/min**MS Parameters:**

Collision Gas (mbar) = 3.43e-3

Collision Energy (eV) = 6-60 (variable)

Analytical Standard Record

22F0058

Description: PFAS - MIX MXF 2ug/mL
Standard Type: Other
Solvent: MeOH
Final Volume (mLs): 1.2
Vials: 1

Expires: 01/11/2025
Prepared: 01/10/2022
Prepared By: Lizbeth Andres
Department: PFAS
Last Edit: 09/15/2022 09:32 by DAG

Analyte	Parent	CAS Number	Concentration	Units
11CL-PF3OUDS		763051-92-9	1.89	ug/mL
9CL-PF3ONS		756426-58-1	1.87	ug/mL
ADONA		919005-14-4	1.89	ug/mL
HFPO-DA		13252-13-6	2	ug/mL



WELLINGTON LABORATORIES

CERTIFICATE OF ANALYSIS DOCUMENTATION

PFAC-MXH 22F0059

Native Per- and Poly-fluoroalkyl Substance Solution/Mixture

PRODUCT CODE: PFAC-MXH
LOT NUMBER: PFACMXH0921
SOLVENT(S): Methanol / Isopropanol (2%) / Water (<1%)
DATE PREPARED: (mm/dd/yyyy) 09/09/2021
LAST TESTED: (mm/dd/yyyy) 09/14/2021
EXPIRY DATE: (mm/dd/yyyy) 09/14/2026
RECOMMENDED STORAGE: Refrigerate ampoule

DESCRIPTION:

PFAC-MXH is a solution/mixture of eleven native linear perfluoroalkylcarboxylic acids (C_4 - C_{14}), eight native perfluoroalkanesulfonates (C_4 , C_5 , C_7 , C_8 , C_{10} and C_{12} linear; C_6 and C_8 linear and branched), three native fluorotelomer sulfonates (4:2, 6:2, and 8:2), two native linear and branched perfluorooctanesulfonamidoacetic acids, and perfluoro-1-octanesulfonamide (FOSA). The components and their concentrations are given in Table A.

The individual components of this mixture all have chemical purities of >98%.

DOCUMENTATION/ DATA ATTACHED:

Table A: Components and Concentrations of the Solution/Mixture
 Table B: Isomeric Components and Percent Composition of br-NMeFOSAA
 Table C: Isomeric Components and Percent Composition of br-NEtFOSAA
 Table D: Isomeric Components and Percent Composition of PFHxSK
 Table E: Isomeric Components and Percent Composition of PFOSK
 Figure 1: LC/MS Data (SIR)
 Figure 2: LC/MS/MS Data (Selected MRM Transitions)

ADDITIONAL INFORMATION:

- See page 2 for further details.
- Contains 4 mole eq. of NaOH to prevent conversion of the carboxylic acids to their respective methyl esters.

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Wellington Laboratories Inc., 345 Southgate Dr. Guelph ON N1G 3M5 CANADA
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INTENDED USE:

The products prepared by Wellington Laboratories Inc. are for laboratory use only. This certified reference material (CRM) was designed to be used as a standard for the identification and/or quantification of the specific chemical compounds it contains.

HANDLING:

This product should only be used by qualified personnel familiar with its potential hazards and trained in the handling of hazardous chemicals. Due care should be exercised to prevent unnecessary human contact or ingestion. All procedures should be carried out in a well-functioning fume hood and suitable gloves, eye protection, and clothing should be worn at all times. Waste should be disposed of according to national and regional regulations. Safety Data Sheets (SDSs) are available upon request.

SYNTHESIS / CHARACTERIZATION:

Our products are synthesized using single-product unambiguous routes whenever possible. They are then characterized, and their structures and purities confirmed, using a combination of the most relevant techniques, such as NMR, GC/MS, LC/MS/MS, SFC/UV/MS/MS, x-ray crystallography, and melting point. Isotopic purities of mass-labelled compounds are also confirmed using HRGC/HRMS and/or LC/MS/MS.

HOMOGENEITY:

Prior to solution preparation, crystalline material is tested for homogeneity using a variety of techniques (as stated above) and its solubility in a given diluent is taken into consideration. Duplicate solutions of a new product are prepared from the same crystalline lot and, after the addition of an appropriate internal standard, they are compared by GC/MS, LC/MS/MS, and/or SFC/UV/MS/MS. The relative response factors of the analyte of interest in each solution are required to be <5% RSD. New solution lots of existing products, as well as mixtures and calibration solutions, are compared to older lots in a similar manner. This further confirms the homogeneity of the crystalline material as well as the stability and homogeneity of the solutions in the storage containers. In order to maintain the integrity of the assigned value(s), and associated uncertainty, the dilution or injection of a subsample of this product should be performed using calibrated measuring equipment.

UNCERTAINTY:

The maximum combined relative standard uncertainty of our reference standard solutions is calculated using the following equation:

The combined relative standard uncertainty, $u_c(y)$, of a value y and the uncertainty of the independent parameters

x_1, x_2, \dots, x_n on which it depends is:

$$u_c(y(x_1, x_2, \dots, x_n)) = \sqrt{\sum_{i=1}^n u(y, x_i)^2}$$

where x is expressed as a relative standard uncertainty of the individual parameter.

The individual uncertainties taken into account include those associated with weights (calibration of the balance) and volumes (calibration of the volumetric glassware). An expanded maximum combined percent relative uncertainty of $\pm 5\%$ (calculated with a coverage factor of 2 and a level of confidence of 95%) is stated on the Certificate of Analysis for all of our products.

TRACEABILITY:

All reference standard solutions are traceable to specific crystalline lots. The microbalances used for solution preparation are regularly calibrated by an external ISO/IEC 17025 accredited laboratory. In addition, their calibration is verified prior to each weighing using calibrated external weights traceable to an ISO/IEC 17025 accredited laboratory. All volumetric glassware used is calibrated, of Class A tolerance, and traceable to an ISO/IEC 17025 accredited laboratory. For certain products, traceability to international interlaboratory studies has also been established.

EXPIRY DATE / PERIOD OF VALIDITY:

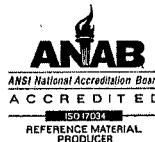
Ongoing stability studies of this product have demonstrated stability in its composition and concentration, until the specified expiry date, in the unopened ampoule. Monitoring for any degradation or change in concentration of the listed analyte(s) is performed on a routine basis.

LIMITED WARRANTY:

At the time of shipment, all products are warranted to be free of defects in material and workmanship and to conform to the stated technical and purity specifications.

QUALITY MANAGEMENT:

This product was produced using a Quality Management System registered to the latest versions of ISO 9001 by SAI Global, ISO/IEC 17025 by the Canadian Association for Laboratory Accreditation Inc. (CALA; A1226), and ISO 17034 by ANSI National Accreditation Board (ANAB; AR-1523).



For additional information or assistance concerning this or any other products from Wellington Laboratories Inc., please visit our website at www.well-labs.com or contact us directly at info@well-labs.com

Table A: PFAC-MXH; Components and Concentrations
($\mu\text{g/mL}$, $\pm 5\%$ in methanol / isopropanol (2%) / water (<1%))

Compound	Acronym	Concentration* (µg/mL)		Peak Assignment in Figure 1
Perfluoro-n-butanoic acid	PFBA	4.00		1
Perfluoro-n-pentanoic acid	PFPeA	2.00		2
Perfluoro-n-hexanoic acid	PFHxA	1.00		5
Perfluoro-n-heptanoic acid	PFHpA	1.00		7
Perfluoro-n-octanoic acid	PFOA	1.00		11
Perfluoro-n-nonanoic acid	PFNA	1.00		14
Perfluoro-n-decanoic acid	PFDA	1.00		18
Perfluoro-n-undecanoic acid	PFUdA	1.00		23
Perfluoro-n-dodecanoic acid	PFDoA	1.00		26
Perfluoro-n-tridecanoic acid	PFTrDA	1.00		27
Perfluoro-n-tetradecanoic acid	PFTeDA	1.00		29
Perfluoro-1-octanesulfonamide	FOSA	1.00		25
N-methylperfluorooctanesulfonamidoacetic acid ^a	N-MeFOSAA: linear isomer	0.760		20
	N-MeFOSAA: ∑ branched isomers	0.240		17
N-ethylperfluorooctanesulfonamidoacetic acid ^b	N-EtFOSAA: linear isomer	0.775		22
	N-EtFOSAA: ∑ branched isomers	0.225		21
Compound	Acronym	Concentration* (µg/mL)		Peak Assignment in Figure 1
		as the salt	as the acid	
Potassium perfluoro-1-butanesulfonate	L-PFBS	1.00	0.887	3
Sodium perfluoro-1-pentanesulfonate	L-PFPeS	1.00	0.941	6
Potassium perfluorohexanesulfonate ^c	PFHxSK: linear isomer	0.811	0.741	9
	PFHxSK: ∑ branched isomers	0.189	0.173	8
Sodium perfluoro-1-heptanesulfonate	L-PFHpS	1.00	0.953	12
Potassium perfluorooctanesulfonate ^d	PFOSK: linear isomer	0.788	0.732	15
	PFOSK: ∑ branched isomers	0.211	0.196	13
Sodium perfluoro-1-nonanesulfonate	L-PFNS	1.00	0.962	19
Sodium perfluoro-1-decanesulfonate	L-PFDS	1.00	0.965	24
Sodium perfluoro-1-dodecanesulfonate	L-PFDoS	1.00	0.970	28
Sodium 1H,1H,2H,2H-perfluorohexanesulfonate	4:2FTS	4.00	3.75	4
Sodium 1H,1H,2H,2H-perfluorooctanesulfonate	6:2FTS	4.00	3.80	10
Sodium 1H,1H,2H,2H-perfluorodecanesulfonate	8:2FTS	4.00	3.84	16

^a See Table B for percent composition of linear and branched N-MeFOSAA isomers.

^b See Table C for percent composition of linear and branched N-EtFOSAA isomers.

^c See Table D for percent composition of linear and branched PFHxSK isomers.

^d See Table E for percent composition of linear and branched PFOSK isomers.

* Concentrations have been rounded to three significant figures.

Certified By: 

B.G. Chittim, General Manager

Date: 09/23/2021

(mm/dd/yyyy)

Table B: br-NMeFOSAA; Isomeric Components and Percent Composition (by ^{19}F -NMR)*

Isomer	Compound	Structure	Percent Composition by ^{19}F -NMR	
1	N-methylperfluoro-1-octanesulfonamidoacetic acid	$\text{CF}_3(\text{CF}_2)_7\text{SO}_2\text{N}(\text{CH}_3)\text{CH}_2\text{CO}_2\text{H}$	76.0	76.0
2	N-methylperfluoro-3-methylheptanesulfonamidoacetic acid	$\text{CF}_3(\text{CF}_2)_3\text{CF}(\text{CF}_3)\text{SO}_2\text{N}(\text{CH}_3)\text{CH}_2\text{CO}_2\text{H}$	0.7	24.0
3	N-methylperfluoro-4-methylheptanesulfonamidoacetic acid	$\text{CF}_3(\text{CF}_2)_2\text{CF}(\text{CF}_3)\text{SO}_2\text{N}(\text{CH}_3)\text{CH}_2\text{CO}_2\text{H}$	2.0	
4	N-methylperfluoro-5-methylheptanesulfonamidoacetic acid	$\text{CF}_3\text{CF}_2\text{CF}(\text{CF}_3)\text{SO}_2\text{N}(\text{CH}_3)\text{CH}_2\text{CO}_2\text{H}$	6.0	
5	N-methylperfluoro-6-methylheptanesulfonamidoacetic acid	$\text{CF}_3\text{CF}(\text{CF}_3)\text{SO}_2\text{N}(\text{CH}_3)\text{CH}_2\text{CO}_2\text{H}$	14.0	
6	N-methylperfluoro-5,5-dimethylhexanesulfonamidoacetic acid	$\text{CF}_3\text{C}(\text{CF}_3)_2\text{SO}_2\text{N}(\text{CH}_3)\text{CH}_2\text{CO}_2\text{H}$	0.2	
7	Other Unidentified Isomers		1.1	

* Percent of total N-methylperfluorooctanesulfonamidoacetic acid isomers only.

Table C: br-NEtFOSAA; Isomeric Components and Percent Composition (by ^{19}F -NMR)*

Isomer	Compound	Structure	Percent Composition by ^{19}F -NMR	
1	N-ethylperfluoro-1-octanesulfonamidoacetic acid	$\text{CF}_3(\text{CF}_2)_7\text{SO}_2\text{NCH}_2\text{CO}_2\text{H}$ C_2H_5	77.5	77.5
2	N-ethylperfluoro-3-methylheptanesulfonamidoacetic acid	$\text{CF}_3(\text{CF}_2)_3\text{CF}(\text{CF}_2)_2\text{SO}_2\text{NCH}_2\text{CO}_2\text{H}$ CF_3 C_2H_5	2.3	22.5
3	N-ethylperfluoro-4-methylheptanesulfonamidoacetic acid	$\text{CF}_3(\text{CF}_2)_2\text{CF}(\text{CF}_2)_3\text{SO}_2\text{NCH}_2\text{CO}_2\text{H}$ CF_3 C_2H_5	2.2	
4	N-ethylperfluoro-5-methylheptanesulfonamidoacetic acid	$\text{CF}_3\text{CF}_2\text{CF}(\text{CF}_2)_4\text{SO}_2\text{NCH}_2\text{CO}_2\text{H}$ CF_3 C_2H_5	5.4	
5	N-ethylperfluoro-6-methylheptanesulfonamidoacetic acid	$\text{CF}_3\text{CF}(\text{CF}_2)_5\text{SO}_2\text{NCH}_2\text{CO}_2\text{H}$ CF_3 C_2H_5	10.4	
6	N-ethylperfluoro-5,5-dimethylhexanesulfonamidoacetic acid	CF_3 $\text{CF}_3\text{C}(\text{CF}_2)_4\text{SO}_2\text{NCH}_2\text{CO}_2\text{H}$ CF_3 C_2H_5	0.3	
7	N-ethylperfluoro-4,5-dimethylhexanesulfonamidoacetic acid	CF_3 $\text{CF}_3\text{CFCF}(\text{CF}_2)_3\text{SO}_2\text{NCH}_2\text{CO}_2\text{H}$ CF_3 C_2H_5	0.3	
8	N-ethylperfluoro-3,5-dimethylhexanesulfonamidoacetic acid	CF_3 $\text{CF}_3\text{CFCF}_2\text{CF}(\text{CF}_2)_2\text{SO}_2\text{NCH}_2\text{CO}_2\text{H}$ CF_3 C_2H_5	0.3	
9	Other Unidentified Isomers		1.3	

* Percent of total N-ethylperfluorooctanesulfonamidoacetic acid isomers only.

Table D: PFHxSK; Isomeric Components and Percent Composition (by ^{19}F -NMR)*

Isomer	Compound	Structure	Percent Composition by ^{19}F -NMR	
1	Potassium perfluoro-1-hexanesulfonate	$\text{CF}_3\text{CF}_2\text{CF}_2\text{CF}_2\text{CF}_2\text{CF}_2\text{SO}_3^-\text{K}^+$	81.1	81.1
2	Potassium 1-trifluoromethylperfluoropentanesulfonate**	$\begin{array}{c} \text{CF}_3\text{CF}_2\text{CF}_2\text{CF}_2\text{CF}(\text{SO}_3^-\text{K}^+)\text{CF}_3 \\ \\ \text{CF}_3 \end{array}$	2.9	18.9
3	Potassium 2-trifluoromethylperfluoropentanesulfonate	$\begin{array}{c} \text{CF}_3\text{CF}_2\text{CF}_2\text{CF}(\text{CF}_3)\text{CF}_2\text{SO}_3^-\text{K}^+ \\ \\ \text{CF}_3 \end{array}$	1.4	
4	Potassium 3-trifluoromethylperfluoropentanesulfonate	$\begin{array}{c} \text{CF}_3\text{CF}_2\text{CF}(\text{CF}_3)\text{CF}_2\text{CF}_2\text{SO}_3^-\text{K}^+ \\ \\ \text{CF}_3 \end{array}$	5.0	
5	Potassium 4-trifluoromethylperfluoropentanesulfonate	$\begin{array}{c} \text{CF}_3\text{CF}(\text{CF}_3)\text{CF}_2\text{CF}_2\text{CF}_2\text{SO}_3^-\text{K}^+ \\ \\ \text{CF}_3 \end{array}$	8.9	
6	Potassium 3,3-di(trifluoromethyl)perfluorobutanesulfonate	$\begin{array}{c} \text{CF}_3 \\ \\ \text{CF}_3\text{C}(\text{CF}_3)_2\text{CF}_2\text{SO}_3^-\text{K}^+ \\ \\ \text{CF}_3 \end{array}$	0.2	
7	Other Unidentified Isomers		0.5	

* Percent of total perfluorohexanesulfonate isomers only.

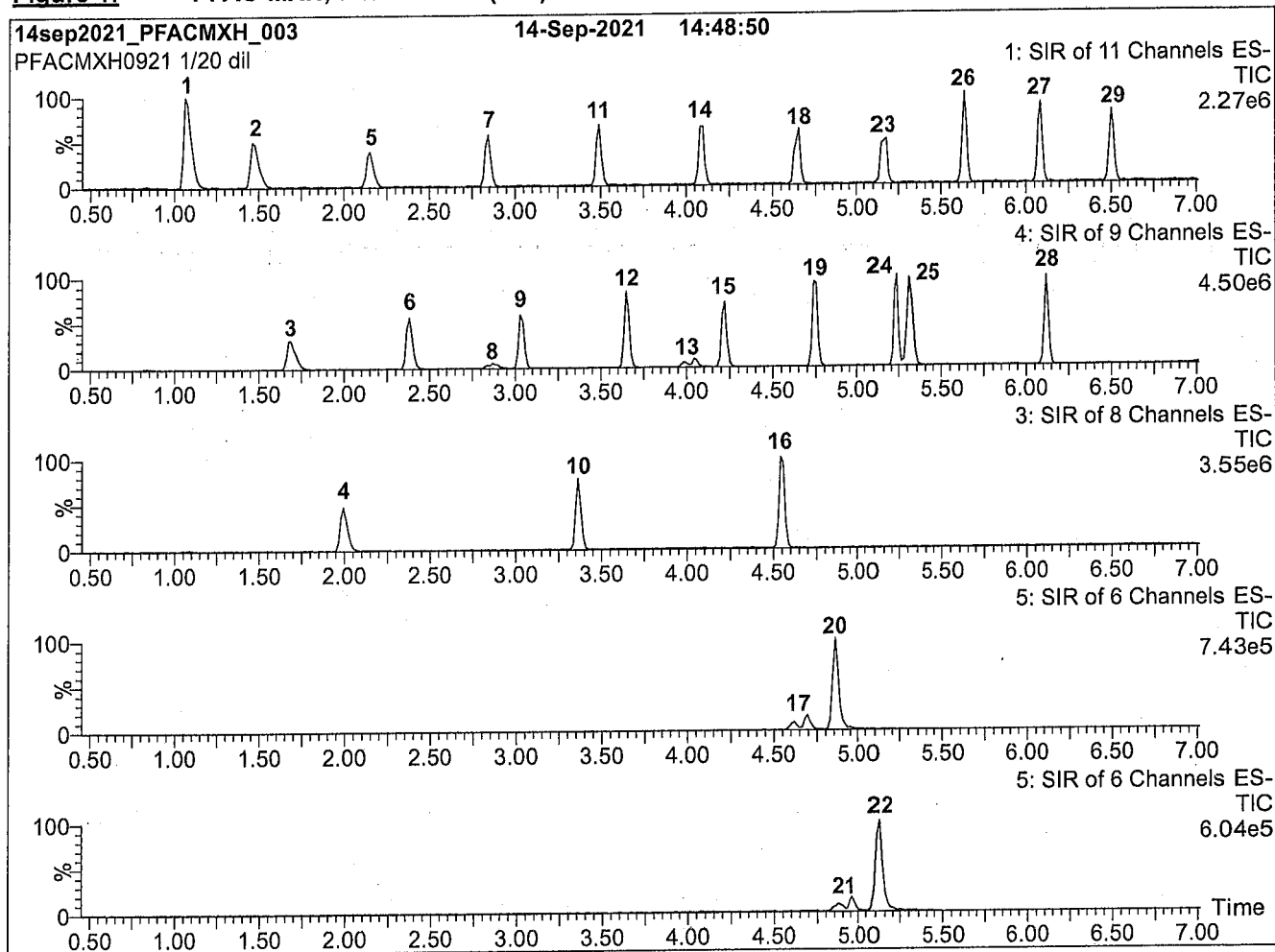
** Systematic Name: Potassium perfluorohexane-2-sulfonate.

Table E: PFOSK; Isomeric Components and Percent Composition (by ^{19}F -NMR)*

Isomer	Compound	Structure	Percent Composition by ^{19}F -NMR	
1	Potassium perfluoro-1-octanesulfonate	$\text{CF}_3\text{CF}_2\text{CF}_2\text{CF}_2\text{CF}_2\text{CF}_2\text{CF}_2\text{SO}_3^-\text{K}^+$	78.8	78.8
2	Potassium 1-trifluoromethylperfluoroheptanesulfonate**	$\text{CF}_3\text{CF}_2\text{CF}_2\text{CF}_2\text{CF}_2\text{CF}_2\text{C}(\text{SO}_3^-\text{K}^+)\text{CF}_3$	1.2	21.1
3	Potassium 2-trifluoromethylperfluoroheptanesulfonate	$\text{CF}_3\text{CF}_2\text{CF}_2\text{CF}_2\text{CF}_2\text{C}(\text{SO}_3^-\text{K}^+)\text{CF}_2\text{CF}_3$	0.6	
4	Potassium 3-trifluoromethylperfluoroheptanesulfonate	$\text{CF}_3\text{CF}_2\text{CF}_2\text{CF}_2\text{C}(\text{SO}_3^-\text{K}^+)\text{CF}_2\text{CF}_2\text{CF}_3$	1.9	
5	Potassium 4-trifluoromethylperfluoroheptanesulfonate	$\text{CF}_3\text{CF}_2\text{CF}_2\text{C}(\text{SO}_3^-\text{K}^+)\text{CF}_2\text{CF}_2\text{CF}_2\text{CF}_3$	2.2	
6	Potassium 5-trifluoromethylperfluoroheptanesulfonate	$\text{CF}_3\text{CF}_2\text{C}(\text{SO}_3^-\text{K}^+)\text{CF}_2\text{CF}_2\text{CF}_2\text{CF}_2\text{CF}_3$	4.5	
7	Potassium 6-trifluoromethylperfluoroheptanesulfonate	$\text{CF}_3\text{C}(\text{SO}_3^-\text{K}^+)\text{CF}_2\text{CF}_2\text{CF}_2\text{CF}_2\text{CF}_2\text{CF}_3$	10.0	
8	Potassium 5,5-di(trifluoromethyl)perfluorohexanesulfonate	$\text{CF}_3\text{C}(\text{CF}_3)_2\text{CF}_2\text{CF}_2\text{CF}_2\text{SO}_3^-\text{K}^+$	0.2	
9	Potassium 4,4-di(trifluoromethyl)perfluorohexanesulfonate	$\text{CF}_3\text{CF}_2\text{C}(\text{CF}_3)_2\text{CF}_2\text{CF}_2\text{SO}_3^-\text{K}^+$	0.03	
10	Potassium 4,5-di(trifluoromethyl)perfluorohexanesulfonate	$\text{CF}_3\text{C}(\text{CF}_3)\text{CF}_2\text{C}(\text{CF}_3)\text{CF}_2\text{SO}_3^-\text{K}^+$	0.4	
11	Potassium 3,5-di(trifluoromethyl)perfluorohexanesulfonate	$\text{CF}_3\text{C}(\text{CF}_3)\text{CF}_2\text{C}(\text{CF}_3)\text{CF}_2\text{SO}_3^-\text{K}^+$	0.07	

* Percent of total perfluorooctanesulfonate isomers only.

** Systematic Name: Potassium perfluorooctane-2-sulfonate.

Figure 1: PFAC-MXH; LC/MS Data (SIR)**Conditions for Figure 1:**

Waters Acquity Ultra Performance LC
Waters Xevo TQ-S micro MS

Chromatographic Conditions:

Column: Acquity UPLC BEH Shield RP₁₈
1.7 μ m, 2.1 x 100 mm

Mobile phase: Gradient

Start: 50% H₂O / 50% (80:20 MeOH:ACN)
(both with 10 mM NH₄OAc buffer)
Ramp to 90% organic over 9 min and hold for 2 min
before returning to initial conditions in 1 min.
Time: 15 min

Flow: 300 μ L/min

MS Parameters:

Experiment: SIR

Source: Electrospray (negative)
Capillary Voltage (kV) = 2.50
Cone Voltage (V) = variable (2-74)
Desolvation Temperature (°C) = 350
Desolvation Gas Flow (L/hr) = 1000

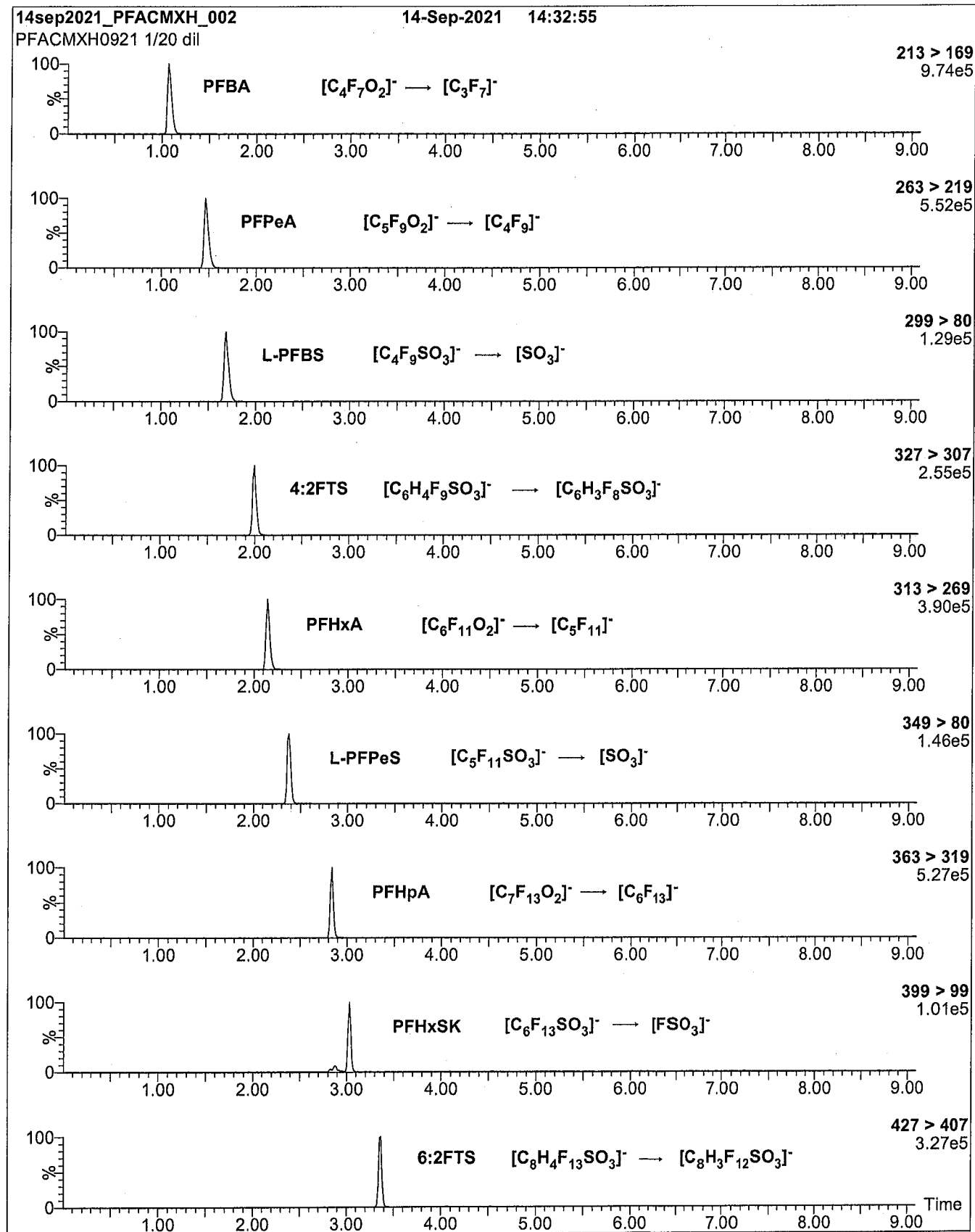
Figure 2: PFAC-MXH; LC/MS/MS Data (Selected MRM Transitions)

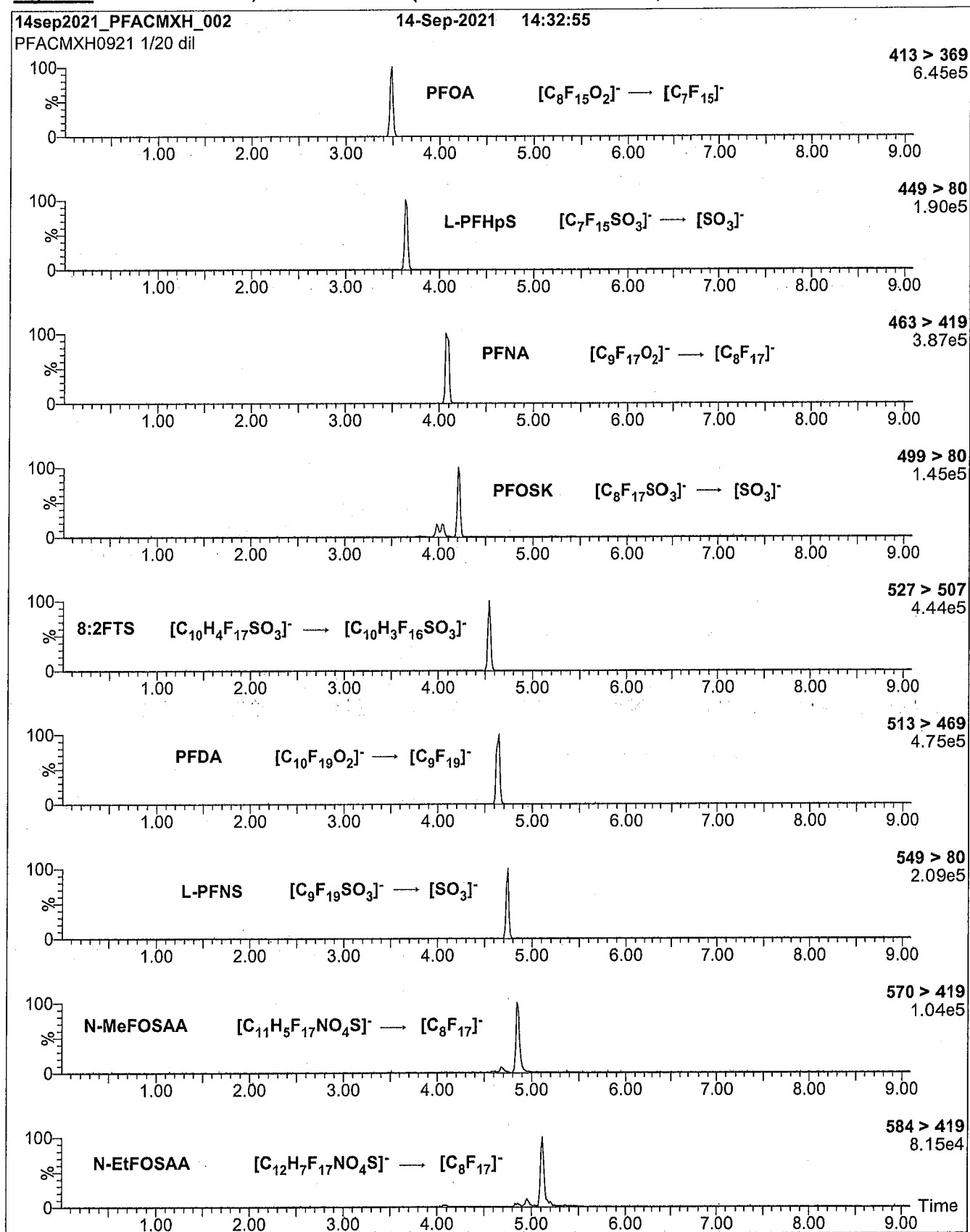
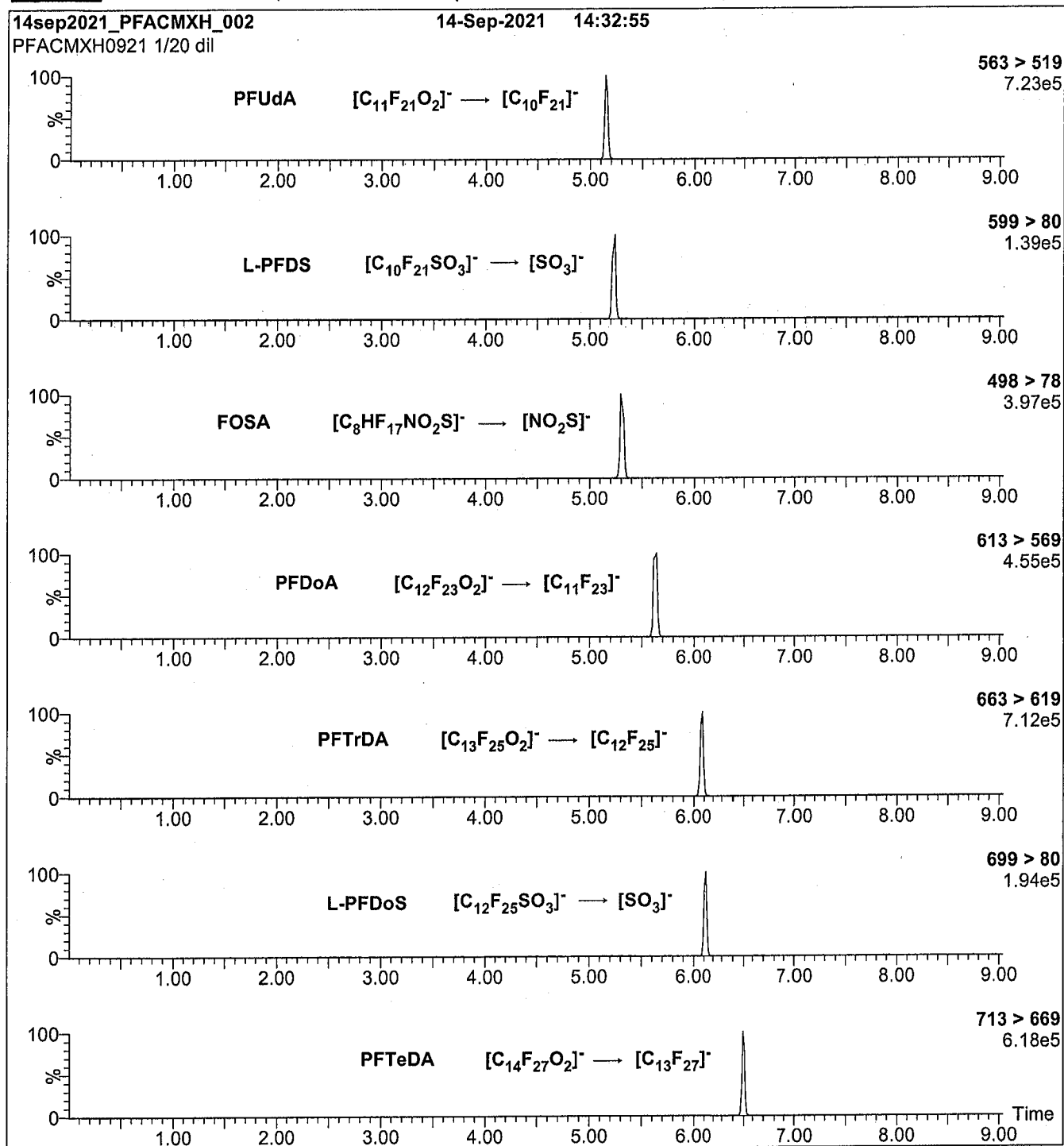
Figure 2: PFAC-MXH; LC/MS/MS Data (Selected MRM Transitions)

Figure 2: PFAC-MXH; LC/MS/MS Data (Selected MRM Transitions)**Conditions for Figure 2:**

Injection: On-column (PFAC-MXH)

Mobile phase: Same as Figure 1

Flow: 300 μ L/min**MS Parameters:**

Collision Gas (mbar) = 3.31e-3

Collision Energy (eV) = 6-60 (variable)

Analytical Standard Record

22F0059

Description: PFAS - MIX MXH 2ug/mL
 Standard Type: Other
 Solvent: MeOH
 Final Volume (mLs): 1.2
 Vials: 1

Expires: 09/14/2026
 Prepared: 09/09/2021
 Prepared By: Lizbeth Andres
 Department: PFAS
 Last Edit: 09/15/2022 09:33 by DAG

Analyte	Parent	CAS Number	Concentration	Units
4:2FTS		757124-72-4	3.75	ug/mL
6:2FTS		27619-97-2	3.8	ug/mL
8:2FTS		39108-34-4	3.84	ug/mL
NEtFOSAA		2991-50-6	1	ug/mL
NMeFOSAA		2355-31-9	1	ug/mL
PFBA		375-22-4	4	ug/mL
PFBS		375-73-5	0.887	ug/mL
PFDA		335-76-2	1	ug/mL
PFDOA		307-55-1	1	ug/mL
PFDOS		79780-39-5	0.97	ug/mL
PFDS		335-77-3	0.965	ug/mL
PFHPA		375-85-9	1	ug/mL
PFHPS		375-92-8	0.953	ug/mL
PFHXA		307-24-4	1	ug/mL
PFHXS		355-46-4	0.914	ug/mL
PFNA		375-95-1	1	ug/mL
PFNS		68259-12-1	0.962	ug/mL
PFOA		335-67-1	1	ug/mL
PFOS		1763-23-1	0.928	ug/mL
PFOSA		754-91-6	1	ug/mL
PFPEA		2706-90-3	2	ug/mL
PFPEs		630402-22-1	0.941	ug/mL
PFTEDA		376-06-7	1	ug/mL
PFTRDA		72629-94-8	1	ug/mL
PFUnA		2058-94-8	1	ug/mL

**WELLINGTON**
LABORATORIES**CERTIFICATE OF ANALYSIS**
DOCUMENTATION**PFAC-MXG** 22F0061**Native Perfluoroalkyl Ether Carboxylic
Acids and Sulfonate Solution/Mixture**

PRODUCT CODE: PFAC-MXG
LOT NUMBER: PFACMXG0222
SOLVENT(S): Methanol/Water (<1%)
DATE PREPARED: (mm/dd/yyyy) 02/07/2022
LAST TESTED: (mm/dd/yyyy) 02/22/2022
EXPIRY DATE: (mm/dd/yyyy) 02/22/2027
RECOMMENDED STORAGE: Store ampoule in a cool, dark place

DESCRIPTION:

PFAC-MXG is a solution/mixture of three native perfluoroalkyl ether carboxylic acids and a native perfluoroalkyl ether sulfonate. The components and their concentrations are given in Table A.

The individual components all have chemical purities of >98%.

DOCUMENTATION/ DATA ATTACHED:

Table A: Components and Concentrations of the Solution/Mixture
Figure 1: LC/MS Data (SIR)
Figure 2: LC/MS/MS Data (Selected MRM Transitions)

ADDITIONAL INFORMATION:

- See page 2 for further details.
- Contains 4 mole eq. of NaOH to prevent conversion of the carboxylic acids to their respective methyl esters.

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INTENDED USE:

The products prepared by Wellington Laboratories Inc. are for laboratory use only. This certified reference material (CRM) was designed to be used as a standard for the identification and/or quantification of the specific chemical compounds it contains.

HANDLING:

This product should only be used by qualified personnel familiar with its potential hazards and trained in the handling of hazardous chemicals. Due care should be exercised to prevent unnecessary human contact or ingestion. All procedures should be carried out in a well-functioning fume hood and suitable gloves, eye protection, and clothing should be worn at all times. Waste should be disposed of according to national and regional regulations. Safety Data Sheets (SDSs) are available upon request.

SYNTHESIS / CHARACTERIZATION:

Our products are synthesized using single-product unambiguous routes whenever possible. They are then characterized, and their structures and purities confirmed, using a combination of the most relevant techniques, such as NMR, GC/MS, LC/MS/MS, SFC/UV/MS/MS, x-ray crystallography, and melting point. Isotopic purities of mass-labelled compounds are also confirmed using HRGC/HRMS and/or LC/MS/MS.

HOMOGENEITY:

Prior to solution preparation, crystalline material is tested for homogeneity using a variety of techniques (as stated above) and its solubility in a given diluent is taken into consideration. Duplicate solutions of a new product are prepared from the same crystalline lot and, after the addition of an appropriate internal standard, they are compared by GC/MS, LC/MS/MS, and/or SFC/UV/MS/MS. The relative response factors of the analyte of interest in each solution are required to be <5% RSD. New solution lots of existing products, as well as mixtures and calibration solutions, are compared to older lots in a similar manner. This further confirms the homogeneity of the crystalline material as well as the stability and homogeneity of the solutions in the storage containers. In order to maintain the integrity of the assigned value(s), and associated uncertainty, the dilution or injection of a subsample of this product should be performed using calibrated measuring equipment.

UNCERTAINTY:

The maximum combined relative standard uncertainty of our reference standard solutions is calculated using the following equation:

The combined relative standard uncertainty, $u_c(y)$, of a value y and the uncertainty of the independent parameters

x_1, x_2, \dots, x_n on which it depends is:

$$u_c(y(x_1, x_2, \dots, x_n)) = \sqrt{\sum_{i=1}^n u(y, x_i)^2}$$

where x is expressed as a relative standard uncertainty of the individual parameter.

The individual uncertainties taken into account include those associated with weights (calibration of the balance) and volumes (calibration of the volumetric glassware). An expanded maximum combined percent relative uncertainty of $\pm 5\%$ (calculated with a coverage factor of 2 and a level of confidence of 95%) is stated on the Certificate of Analysis for all of our products.

TRACEABILITY:

All reference standard solutions are traceable to specific crystalline lots. The microbalances used for solution preparation are regularly calibrated by an external ISO/IEC 17025 accredited laboratory. In addition, their calibration is verified prior to each weighing using calibrated external weights traceable to an ISO/IEC 17025 accredited laboratory. All volumetric glassware used is calibrated, of Class A tolerance, and traceable to an ISO/IEC 17025 accredited laboratory. For certain products, traceability to international interlaboratory studies has also been established.

EXPIRY DATE / PERIOD OF VALIDITY:

Ongoing stability studies of this product have demonstrated stability in its composition and concentration, until the specified expiry date, in the unopened ampoule. Monitoring for any degradation or change in concentration of the listed analyte(s) is performed on a routine basis.

LIMITED WARRANTY:

At the time of shipment, all products are warranted to be free of defects in material and workmanship and to conform to the stated technical and purity specifications.

QUALITY MANAGEMENT:

This product was produced using a Quality Management System registered to the latest versions of ISO 9001 by SAI Global, ISO/IEC 17025 by the Canadian Association for Laboratory Accreditation Inc. (CALA; A1226), and ISO 17034 by ANSI National Accreditation Board (ANAB; AR-1523).



For additional information or assistance concerning this or any other products from Wellington Laboratories Inc., please visit our website at www.well-labs.com or contact us directly at info@well-labs.com

Table A: PFAC-MXG; Components and Concentrations (ng/mL; \pm 5% in methanol/water (<1%))

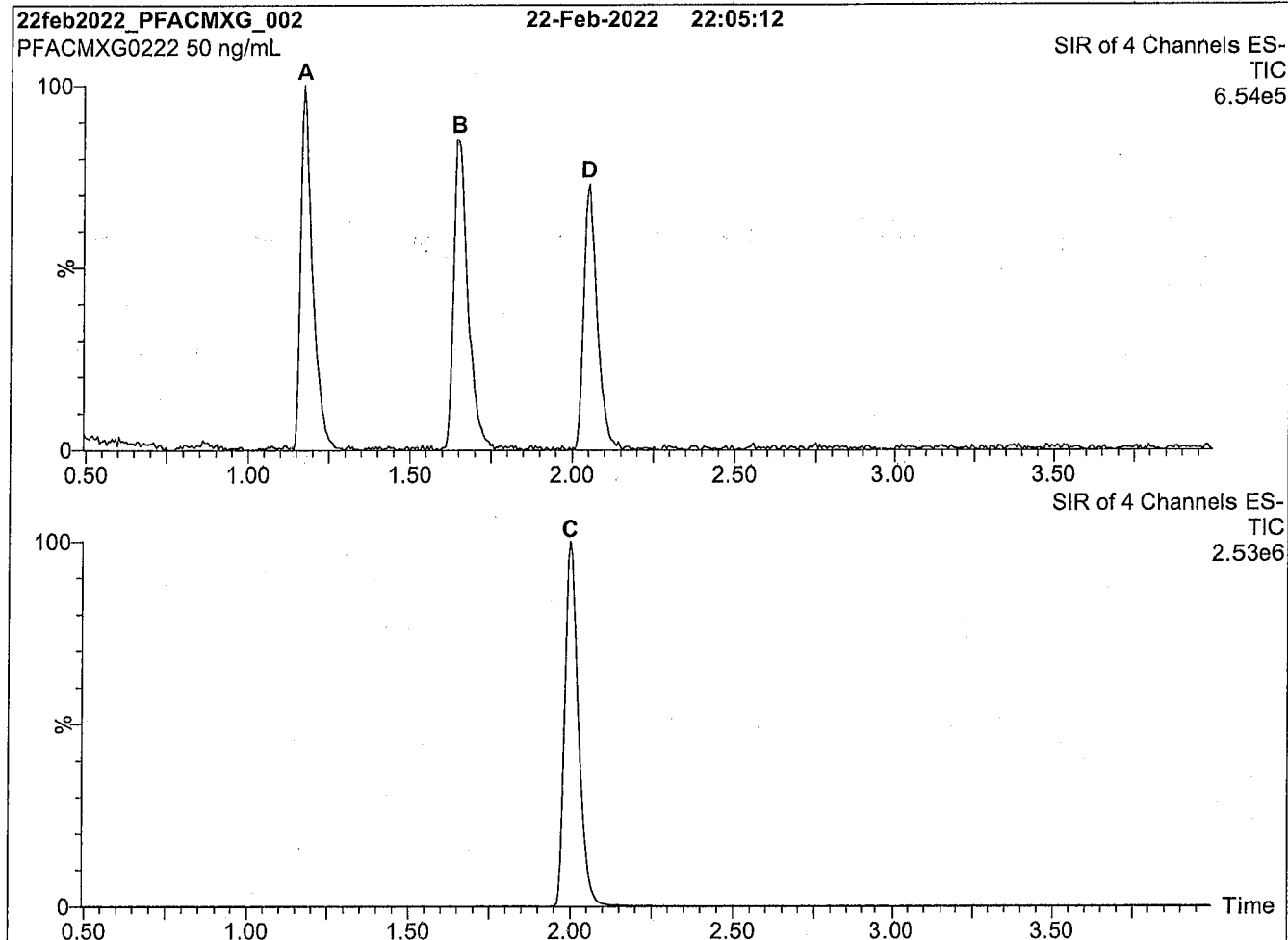
Compound	Acronym	Concentration (ng/mL)		Peak Assignment in Figure 1
Perfluoro-4-oxapentanoic acid	PF4OPeA	2000		A
Perfluoro-5-oxahexanoic acid	PF5OHxA	2000		B
Perfluoro-3,6-dioxaheptanoic acid	3,6-OPFHpA	2000		D
Compound	Acronym	Concentration* (ng/mL)		Peak Assignment in Figure 1
		as the salt	as the acid	
Potassium perfluoro(2-ethoxyethane)sulfonate	PFEESA	2000	1780	C

* Concentrations have been rounded to three significant figures.

Certified By: 

B.G. Chittim, General Manager

Date: 03/03/2022
(mm/dd/yyyy)

Figure 1: PFAC-MXG; LC/MS Data (SIR)**Conditions for Figure 1:**

Waters Acquity Ultra Performance LC
Waters Xevo TQ-S micro MS

Chromatographic Conditions:

Column: Acquity UPLC BEH Shield RP₁₈
1.7 μ m, 2.1 x 100 mm

Mobile phase: Gradient

Start: 50% H₂O / 50% (80:20 MeOH:ACN)
(both with 10 mM NH₄OAc buffer)
Ramp to 90% organic over 8 min and hold for 2 min
before returning to initial conditions in 0.75 min.
Time: 12 min

Flow: 300 μ L/min

MS Parameters:

Experiment: SIR

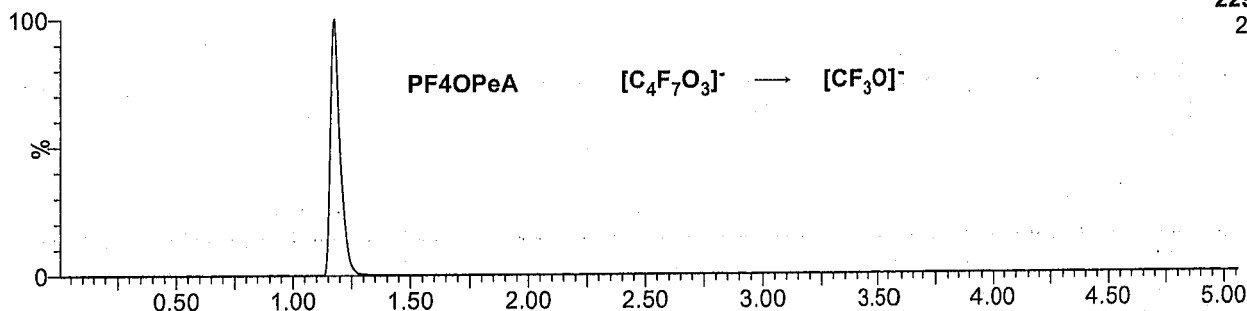
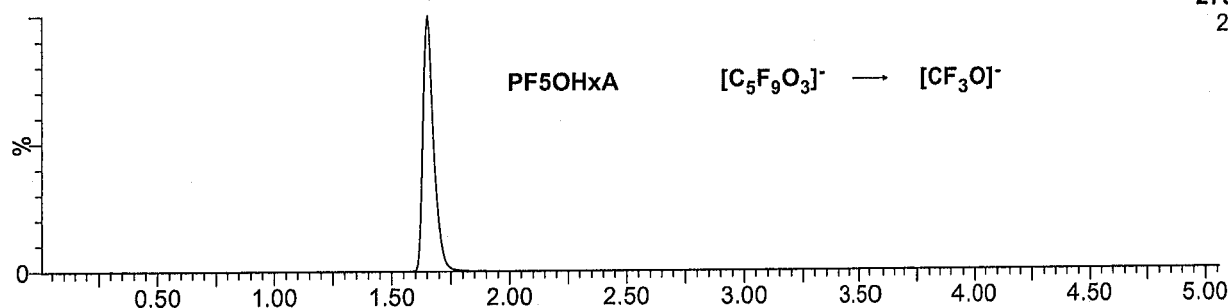
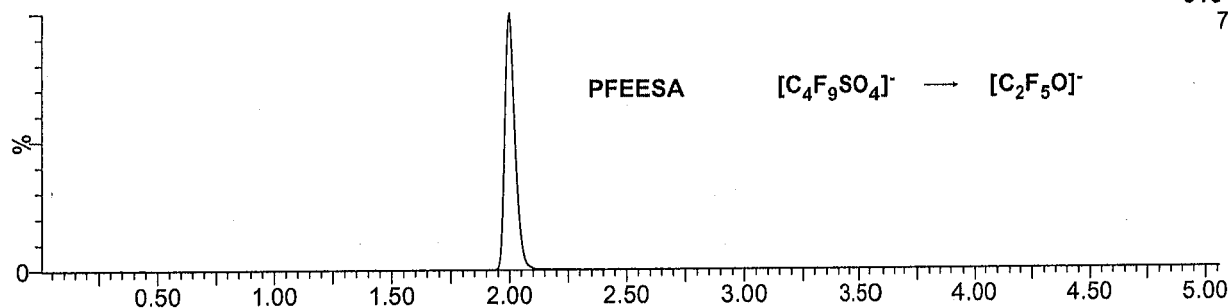
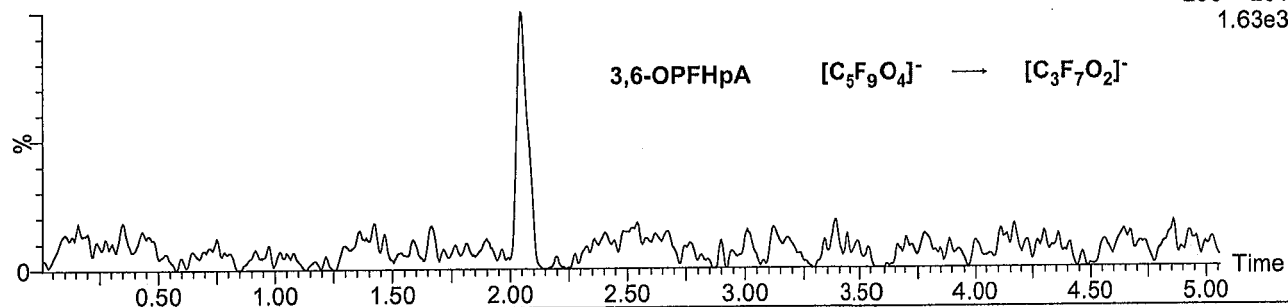
Source: Electrospray (negative)
Capillary Voltage (kV) = 1.00
Cone Voltage (V) = variable (15-35)
Desolvation Temperature ($^{\circ}$ C) = 500
Desolvation Gas Flow (L/hr) = 1000

Figure 2: PFAC-MXG; LC/MS/MS Data (Selected MRM Transitions)

22feb2022_PFACMXG_003

22-Feb-2022 22:18:09

PFACMXG0222 50 ng/mL

229 > 85
2.70e5279 > 85
2.89e5315 > 135
7.56e5295 > 201
1.63e3**Conditions for Figure 2:**

Injection: On-column (PFAC-MXG)

Mobile phase: Same as Figure 1

Flow: 300 μ L/min**MS Parameters:**

Collision Gas (mbar) = 3.33e-3

Collision Energy (eV) = 8-48 (variable)

Analytical Standard Record

22F0061

Description:	PFAS - MIX MXG 2ug/mL	Expires:	02/22/2027
Standard Type:	Other	Prepared:	02/07/2022
Solvent:	MeOH	Prepared By:	Lizbeth Andres
Final Volume (mLs):	1	Department:	PFAS
Vials:	1	Last Edit:	09/15/2022 09:34 by DAG
Comments:	contains NFHDA PFMBA PFMPA PFEESA @ 2ug/mL		

Analyte	Parent	CAS Number	Concentration	Units
NFDHA		151772-58-6	2	ug/mL
PFEESA		113507-82-7	1.78	ug/mL
PFMBA		863090-89-5	2	ug/mL
PFMPA		377-73-1	2	ug/mL

Analytical Standard Record

22F0445

Description: TDCA 1000ug/mL
Standard Type: Other
Solvent: 62097
Final Volume (mLs): 25
Vials: 1

Expires: 09/19/2023
Prepared: 01/12/2022
Prepared By: Dipti Gokal
Department: PFAS
Last Edit: 11/14/2022 14:31 by DAG

Analyte	Parent	CAS Number	Concentration	Units
TAURODEOXYCHOLIC ACID	22A0123	516-50-7	1000	ug/mL

Parent Standards used:

Standard	Description	Prepared	Prepared By	Lot Nbr	Expires	Last Edit	(mLs)
22A0123	PFAS Taurodeoxycholic Acid, Sodium Salt neat	10/09/2021	Calbiochem	3761825	09/30/2023	06/22/2022 13:01 by DAG	25000

Analytical Standard Record

22F0446

Description: TCDA 100ug/mL
Standard Type: Other
Solvent: 62097
Final Volume (mL): 3
Vials: 1

Expires: 09/19/2023
Prepared: 01/13/2022
Prepared By: Dipti Gokal
Department: PFAS
Last Edit: 06/22/2022 13:05 by DAG

Analyte	Parent	CAS Number	Concentration	Units
TAURODEOXYCHOLIC ACID	22F0445	516-50-7	100	ug/mL

Parent Standards used:

Standard	Description	Prepared	Prepared By	Lot Nbr	Expires	Last Edit	(mL)
22F0445	TCDA 1000ug/mL	01/12/2022	Calbiochem	3761825	09/19/2023	06/22/2022 13:05 by DAG	0.3

Analytical Standard Record

22I0153

Description: PFAS - MIX 1633 200ng/mL
 Standard Type: Analyte Spike
 Solvent: MeOH
 Final Volume (mLs): 6
 Vials: 1

Expires: 01/11/2025
 Prepared: 09/13/2022
 Prepared By: Dipti Gokal
 Department: PFAS
 Last Edit: 09/15/2022 09:34 by DAG

Analyte	Parent	CAS Number	Concentration	Units
NMeFOSE	22C0307	24448-09-7	0.8	ug/mL
3:3FTCA	22C0308	113507-82-7	0.8	ug/mL
5:3FTCA	22C0309	914637-49-3	0.8	ug/mL
NEtFOSE	22C0310	1691-99-2	0.8	ug/mL
7:3FTCA	22C0311	812-70-4	0.8	ug/mL
NMeFOSA	22C0312	31506-32-8	0.8	ug/mL
NEtFOSA	22C0313	4151-50-2	0.8	ug/mL
11CL-PF3OUDS	22F0058	763051-92-9	0.378	ug/mL
9CL-PF3ONS	22F0058	756426-58-1	0.374	ug/mL
ADONA	22F0058	919005-14-4	0.378	ug/mL
HFPO-DA	22F0058	13252-13-6	0.4	ug/mL
4:2FTS	22F0059	757124-72-4	0.75	ug/mL
6:2FTS	22F0059	27619-97-2	0.76	ug/mL
8:2FTS	22F0059	39108-34-4	0.768	ug/mL
NEtFOSAA	22F0059	2991-50-6	0.2	ug/mL
NMeFOSAA	22F0059	2355-31-9	0.2	ug/mL
PFBA	22F0059	375-22-4	0.8	ug/mL
PFBS	22F0059	375-73-5	0.177	ug/mL
PFDA	22F0059	335-76-2	0.2	ug/mL
PFDOA	22F0059	307-55-1	0.2	ug/mL
PFDOS	22F0059	79780-39-5	0.194	ug/mL
PFDS	22F0059	335-77-3	0.193	ug/mL
PFHPA	22F0059	375-85-9	0.2	ug/mL
PFHPS	22F0059	375-92-8	0.191	ug/mL
PFHXA	22F0059	307-24-4	0.2	ug/mL
PFHXS	22F0059	355-46-4	0.183	ug/mL
PFNA	22F0059	375-95-1	0.2	ug/mL
PFNS	22F0059	68259-12-1	0.192	ug/mL
PFOA	22F0059	335-67-1	0.2	ug/mL
PFOS	22F0059	1763-23-1	0.186	ug/mL
PFOSA	22F0059	754-91-6	0.2	ug/mL
PFPEA	22F0059	2706-90-3	0.4	ug/mL
PFPEs	22F0059	630402-22-1	0.188	ug/mL
PFTEDA	22F0059	376-06-7	0.2	ug/mL
PFTRDA	22F0059	72629-94-8	0.2	ug/mL
PFUnA	22F0059	2058-94-8	0.2	ug/mL
NFDHA	22F0061	151772-58-6	0.4	ug/mL
PFEESA	22F0061	113507-82-7	0.356	ug/mL
PFMBA	22F0061	863090-89-5	0.4	ug/mL
PFMPA	22F0061	377-73-1	0.4	ug/mL

Analytical Standard Record

22I0153

Parent Standards used:

Standard	Description	Prepared	Prepared By	Lot Nbr	Expires	Last Edit		(mls)
22C0307	PFAS - SAS N-MeFOSE 50ug/mL	03/15/2022	Wellington Laboratories	NMeFOSE0921M	09/23/2026	03/15/2022 15:59	by DAG	0.096
22C0308	PFAS - SAS FPrPA 50ug/mL	03/15/2022	Wellington Laboratories	FPrPA0122	02/03/2027	03/15/2022 15:59	by DAG	0.096
22C0309	PFAS - SAS FPePA 50ug/mL	03/15/2022	Wellington Laboratories	FPePA1221	01/05/2027	03/15/2022 15:59	by DAG	0.096
22C0310	PFAS - SAS NEtFOSE 50ug/mL	03/15/2022	Wellington Laboratories	NEtFOSE0921M	09/23/2026	03/15/2022 15:59	by DAG	0.096
22C0311	PFAS - SAS FHpPA 50ug/mL	03/15/2022	Wellington Laboratories	HHpPA1020	11/12/2025	03/15/2022 16:00	by DAG	0.096
22C0312	PFAS - SAS NMeFOSA 50ug/mL	03/15/2022	Wellington Laboratories	NMeFOSA0721M	08/03/2026	03/15/2022 16:00	by DAG	0.096
22C0313	PFAS - SAS NEtFOSA 50ug/mL	03/15/2022	Wellington Laboratories	NEtFOSA0821M	08/12/2026	08/17/2022 10:49	by LYA	0.096
22F0058	PFAS - MIX MXF 2ug/mL	01/10/2022	Wellington Laboratories	PFACMXF0122	01/11/2025	09/15/2022 09:32	by DAG	1.2
22F0059	PFAS - MIX MXH 2ug/mL	09/09/2021	Wellington Laboratories	PFACMXH0921	09/14/2026	09/15/2022 09:33	by DAG	1.2
22F0061	PFAS - MIX MXG 2ug/mL	02/07/2022	Wellington Laboratories	PFACMXG0222	02/22/2027	09/15/2022 09:34	by DAG	1.2

Analytical Standard Record

22J0297

Description:	T-PFOA	Expires:	01/27/2027
Standard Type:	Other	Prepared:	01/27/2022
Solvent:	MeOH	Prepared By:	Wellington Laboratories (Lot#:
Final Volume (mls):	1	Department:	PFOA0122)
Vials:	1	Last Edit:	10/18/2022 12:59 by HGH

Analyte	Parent	CAS Number	Concentration	Units
PFOA		335-67-1	50	ug/mL



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CERTIFICATE OF ANALYSIS DOCUMENTATION

PRODUCT CODE: T-PFOA **LOT NUMBER:** TPFOA0122
COMPOUND: Ammonium perfluorooctanoate (Technical Grade)
STRUCTURE: (see Table A) **CAS #:** 3825-26-1
 (for linear ammonium perfluorooctanoate)
MOLECULAR FORMULA: $C_8F_{15}O_2NH_4$
CONCENTRATION: $50.0 \pm 2.5 \mu\text{g/mL}$ (gravimetric)
CHEMICAL PURITY: Technical material
SOLVENT(S): Methanol/Water (<1%)
LAST TESTED: (mm/dd/yyyy) 01/27/2022
EXPIRY DATE: (mm/dd/yyyy) 01/27/2027
RECOMMENDED STORAGE: Store ampoule in a cool, dark place

DOCUMENTATION/ DATA ATTACHED:

Table A: Isomeric Components and Percent Composition
 Figure 1: LC/MS Data (Full Scan and Mass Spectrum)
 Figure 2: LC/MS Data (SIR)
 Figure 3: LC/MS/MS Data (Selected MRM Transitions)
 Figure 4: LC/MS Elution Profile of the Perfluorooctanoic Acid Isomers

ADDITIONAL INFORMATION:

- See page 2 for further details.
- This technical mixture is >97% ammonium perfluorooctanoate (branched and linear isomers). The remaining 3% consists of common impurities such as the perfluoroheptanoic and perfluorohexanoic acids.
- It is recommended that this solution be used as a *qualitative or semi-quantitative standard only*.
- Contains 4 mole eq. of NaOH to prevent conversion of any carboxylic acids to their corresponding methyl esters.
- The molecular weight of perfluoro-n-octanoic acid is 414.07 g/mol.

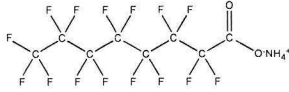
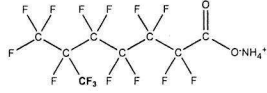
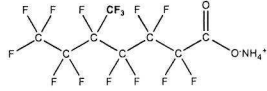
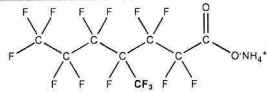
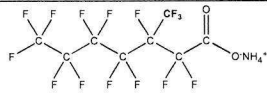
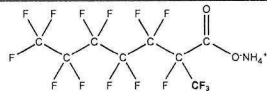
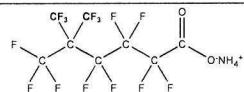
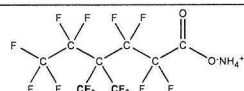
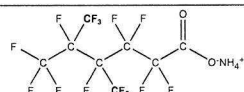
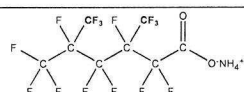
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Certified By: 
 B.G. Chittim, General Manager

Date: 01/27/2022
 (mm/dd/yyyy)

Wellington Laboratories Inc., 345 Southgate Dr. Guelph ON N1G 3M5 CANADA
 519-822-2436 • Fax: 519-822-2849 • info@well-labs.com

Table A: T-PFOA; Isomeric Components and Percent Composition (by ^{19}F -NMR)*

Isomer	Compound	Structure	Percent Composition by ^{19}F -NMR
1	Ammonium perfluoro-n-octanoate		79
2	Ammonium 6-trifluoromethylperfluoroheptanoate		9.0
3	Ammonium 5-trifluoromethylperfluoroheptanoate		4.5
4	Ammonium 4-trifluoromethylperfluoroheptanoate		4.0
5	Ammonium 3-trifluoromethylperfluoroheptanoate		3.0
6	Ammonium 2-trifluoromethylperfluoroheptanoate		0.50
7	Ammonium 5,5-bis(trifluoromethyl)perfluorohexanoate		
8	Ammonium 4,4-bis(trifluoromethyl)perfluorohexanoate		
9	Ammonium 4,5-bis(trifluoromethyl)perfluorohexanoate		
10	Ammonium 3,5-bis(trifluoromethyl)perfluorohexanoate		

* Percent Composition was determined by ^{19}F -NMR. The percentages displayed are of total ammonium perfluorooctanoate isomers only (isomers are labelled in Figure 4).

Analytical Standard Record

22J0298

Description:	br-FOSA	Expires:	10/07/2027
Standard Type:	Other	Prepared:	09/14/2022
Solvent:	Isopropanol	Prepared By:	Wellington Laboratories (Lot#:
Final Volume (mls):	1	Department:	br-FOSA0922)
Vials:	1	Last Edit:	10/18/2022 13:03 by HGH

Analyte	Parent	CAS Number	Concentration	Units
PFOSA		754-91-6	50	ug/mL



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CERTIFICATE OF ANALYSIS DOCUMENTATION

br-FOSA

Perfluorooctanesulfonamide Isomeric Mix

<u>PRODUCT CODE:</u>	br-FOSA
<u>LOT NUMBER:</u>	brFOSA0922
<u>CONCENTRATION:</u>	50.0 ± 2.5 µg/mL
<u>SOLVENT(S):</u>	Isopropanol
<u>DATE PREPARED:</u> (mm/dd/yyyy)	09/14/2022
<u>LAST TESTED:</u> (mm/dd/yyyy)	10/07/2022
<u>EXPIRY DATE:</u> (mm/dd/yyyy)	10/07/2027
<u>RECOMMENDED STORAGE:</u>	Refrigerate ampoule

DESCRIPTION:

The chemical purity has been determined to be ≥98% perfluorooctanesulfonamide (linear and branched isomers). The full name, structure, and percent composition for each of the identified isomeric components are given in Table A.

DOCUMENTATION/ DATA ATTACHED:

Table A: Isomeric Components and Percent Composition by ¹⁹F-NMR
 Figure 1: LC/MS Data (Full Scan and Mass Spectrum)
 Figure 2: LC/MS Data (SIR)
 Figure 3: LC/MS/MS Data (Selected MRM Transitions)

ADDITIONAL INFORMATION:

- See page 2 for further details.
- CAS #: 754-91-6 (for linear isomer).

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Table A: br-FOSA; Isomeric Components and Percent Composition (by ^{19}F -NMR)*

Isomer	Compound	Structure	Percent Composition by ^{19}F -NMR
1	Perfluoro-1-octanesulfonamide	$\text{CF}_3(\text{CF}_2)_7\text{SO}_2\text{NH}_2$	66.6
2	Perfluoro-1-methyl-1-heptanesulfonamide**	$\text{CF}_3(\text{CF}_2)_5\text{C}(\text{CF}_3)\text{SO}_2\text{NH}_2$	0.8
3	Perfluoro-2-methyl-1-heptanesulfonamide	$\text{CF}_3(\text{CF}_2)_4\text{C}(\text{CF}_3)\text{CF}_2\text{SO}_2\text{NH}_2$	0.3
4	Perfluoro-3-methyl-1-heptanesulfonamide	$\text{CF}_3(\text{CF}_2)_3\text{C}(\text{CF}_3)\text{CF}(\text{CF}_2)_2\text{SO}_2\text{NH}_2$	4.2
5	Perfluoro-4-methyl-1-heptanesulfonamide	$\text{CF}_3(\text{CF}_2)_2\text{C}(\text{CF}_3)\text{CF}(\text{CF}_2)_3\text{SO}_2\text{NH}_2$	3.5
6	Perfluoro-5-methyl-1-heptanesulfonamide	$\text{CF}_3\text{CF}_2\text{C}(\text{CF}_3)\text{CF}(\text{CF}_2)_4\text{SO}_2\text{NH}_2$	7.8
7	Perfluoro-6-methyl-1-heptanesulfonamide	$\text{CF}_3\text{C}(\text{CF}_3)\text{CF}(\text{CF}_2)_5\text{SO}_2\text{NH}_2$	16.8
8	Perfluoro-5,5-dimethyl-1-hexanesulfonamide	$\text{CF}_3\text{C}(\text{CF}_3)_2\text{CF}(\text{CF}_2)_4\text{SO}_2\text{NH}_2$	0.2

* Percent of total perfluorooctanesulfonamide isomers only.

** Systematic Name: Perfluoro-2-octanesulfonamide.

Certified By: 
B.G. Chittim, General Manager

Date: 11/15/2022
(mm/dd/yyyy)

Analytical Standard Record

22J0298

Description:	br-FOSA	Expires:	10/07/2027
Standard Type:	Other	Prepared:	09/14/2022
Solvent:	Isopropanol	Prepared By:	Wellington Laboratories (Lot#: br-FOSA0922)
Final Volume (mls):	1	Department:	PFAS
Vials:	1	Last Edit:	10/18/2022 13:03 by HGH

Analyte	Parent	CAS Number	Concentration	Units
PFOSA		754-91-6	50	ug/mL

Analytical Standard Record

22J0301

Description:	br-NMeFOSA	Expires:	08/23/2027
Standard Type:	Other	Prepared:	08/23/2022
Solvent:	MeOH	Prepared By:	Wellington Laboratories (Lot#:
Final Volume (mls):	1	Department:	br-NMeFOSA0822)
Vials:	1	Last Edit:	10/18/2022 13:37 by HGH

Analyte	Parent	CAS Number	Concentration	Units
NMeFOSA		31506-32-8	50	ug/mL



WELLINGTON LABORATORIES

CERTIFICATE OF ANALYSIS DOCUMENTATION

br-NMeFOSA

N-Methylperfluorooctanesulfonamide Isomeric Mix

<u>PRODUCT CODE:</u>	br-NMeFOSA
<u>LOT NUMBER:</u>	brNMeFOSA0822
<u>CONCENTRATION:</u>	50.0 ± 2.5 µg/mL
<u>SOLVENT(S):</u>	Methanol
<u>DATE PREPARED:</u> (mm/dd/yyyy)	08/18/2022
<u>LAST TESTED:</u> (mm/dd/yyyy)	08/23/2022
<u>EXPIRY DATE:</u> (mm/dd/yyyy)	08/23/2027
<u>RECOMMENDED STORAGE:</u>	Store ampoule in a cool, dark place

DESCRIPTION:

The chemical purity has been determined to be ≥98% N-methylperfluorooctanesulfonamide (linear and branched isomers). The full name, structure, and percent composition for each of the identified isomeric components are given in Table A.

DOCUMENTATION/ DATA ATTACHED:

Table A: Isomeric Components and Percent Composition by ¹⁹F-NMR
 Figure 1: LC/MS Data (Full Scan and Mass Spectrum)
 Figure 2: LC/MS Data (SIR)
 Figure 3: LC/MS/MS Data (Selected MRM Transitions)

ADDITIONAL INFORMATION:

- See page 2 for further details.
- CAS #: 31506-32-8 (for linear isomer).

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Table A: br-NMeFOSA; Isomeric Components and Percent Composition (by ^{19}F -NMR)*

Isomer	Compound	Structure	Percent Composition by ^{19}F -NMR
1	N-Methylperfluoro-1-octanesulfonamide	$\text{CF}_3(\text{CF}_2)_7\text{SO}_2\text{NHCH}_3$	72.3
2	N-Methylperfluoro-3-methyl-1-heptanesulfonamide	$\text{CF}_3(\text{CF}_2)_3\text{CF}(\text{CF}_2)_2\text{SO}_2\text{NHCH}_3$	2.1
3	N-Methylperfluoro-4-methyl-1-heptanesulfonamide	$\text{CF}_3(\text{CF}_2)_2\text{CF}(\text{CF}_2)_3\text{SO}_2\text{NHCH}_3$	2.6
4	N-Methylperfluoro-5-methyl-1-heptanesulfonamide	$\text{CF}_3\text{CF}_2\text{CF}(\text{CF}_2)_4\text{SO}_2\text{NHCH}_3$	6.7
5	N-Methylperfluoro-6-methyl-1-heptanesulfonamide	$\text{CF}_3\text{CF}(\text{CF}_2)_5\text{SO}_2\text{NHCH}_3$	16.2
6	N-Methylperfluoro-5,5-dimethyl-1-hexanesulfonamide	$\text{CF}_3\text{C}(\text{CF}_3)(\text{CF}_2)_4\text{SO}_2\text{NHCH}_3$	0.04

* Percent of total N-methylperfluorooctanesulfonamide isomers only.

Certified By: _____


B.G. Chittim, General Manager

Date: 11/15/2022
(mm/dd/yyyy)

Analytical Standard Record

22J0301

Description:	br-NMeFOSA	Expires:	08/23/2027
Standard Type:	Other	Prepared:	08/23/2022
Solvent:	MeOH	Prepared By:	Wellington Laboratories (Lot#: br-NMeFOSA0822)
Final Volume (mls):	1	Department:	PPAS
Vials:	1	Last Edit:	10/18/2022 13:37 by HGH

Analyte	Parent	CAS Number	Concentration	Units
NMeFOSA		31506-32-8	50	ug/mL

Analytical Standard Record

22J0302

Description:	br-NETfOSA	Expires:	10/07/2027
Standard Type:	Other	Prepared:	10/07/2022
Solvent:	MeOH	Prepared By:	Wellington Laboratories (Lot#: PFA5)
Final Volume (mls):	1	Department:	NETfOSA0922)
Vials:	1	Last Edit:	10/18/2022 13:38 by HGH

Analyte	Parent	CAS Number	Concentration	Units
NETfOSA		4151-50-2	50	ug/mL



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CERTIFICATE OF ANALYSIS DOCUMENTATION

br-NEtFOSA

N-Ethylperfluorooctanesulfonamide Isomeric Mix

<u>PRODUCT CODE:</u>	br-NEtFOSA
<u>LOT NUMBER:</u>	brNEtFOSA0922
<u>CONCENTRATION:</u>	50.0 ± 2.5 µg/mL
<u>SOLVENT(S):</u>	Methanol
<u>DATE PREPARED:</u> (mm/dd/yyyy)	08/23/2022
<u>LAST TESTED:</u> (mm/dd/yyyy)	10/07/2022
<u>EXPIRY DATE:</u> (mm/dd/yyyy)	10/07/2027
<u>RECOMMENDED STORAGE:</u>	Store ampoule in a cool, dark place

DESCRIPTION:

The chemical purity has been determined to be ≥98% N-ethylperfluorooctanesulfonamide (linear and branched isomers). The full name, structure, and percent composition for each of the identified isomeric components are given in Table A.

DOCUMENTATION/ DATA ATTACHED:

Table A: Isomeric Components and Percent Composition by ¹⁹F-NMR
 Figure 1: LC/MS Data (Full Scan and Mass Spectrum)
 Figure 2: LC/MS Data (SIR)
 Figure 3: LC/MS/MS Data (Selected MRM Transitions)

ADDITIONAL INFORMATION:

- See page 2 for further details.
- CAS #: 4151-50-2 (for linear isomer).

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Table A: br-NEtFOSA; Isomeric Components and Percent Composition (by ^{19}F -NMR)*

Isomer	Compound	Structure	Percent Composition by ^{19}F -NMR
1	N-Ethylperfluoro-1-octanesulfonamide	$\text{CF}_3(\text{CF}_2)_7\text{SO}_2\text{NHCH}_2\text{CH}_3$	73.8
2	N-Ethylperfluoro-1-methyl-1-heptanesulfonamide**	$\text{CF}_3(\text{CF}_2)_5\text{CF}(\text{SO}_2\text{NHCH}_2\text{CH}_3)\text{CF}_3$	0.1
3	N-Ethylperfluoro-3-methyl-1-heptanesulfonamide	$\text{CF}_3(\text{CF}_2)_3\text{CF}(\text{CF}_2)_2\text{SO}_2\text{NHCH}_2\text{CH}_3$	2.3
4	N-Ethylperfluoro-4-methyl-1-heptanesulfonamide	$\text{CF}_3(\text{CF}_2)_2\text{CF}(\text{CF}_2)_3\text{SO}_2\text{NHCH}_2\text{CH}_3$	2.6
5	N-Ethylperfluoro-5-methyl-1-heptanesulfonamide	$\text{CF}_3\text{CF}_2\text{CF}(\text{CF}_2)_4\text{SO}_2\text{NHCH}_2\text{CH}_3$	6.2
6	N-Ethylperfluoro-6-methyl-1-heptanesulfonamide	$\text{CF}_3\text{CF}(\text{CF}_2)_5\text{SO}_2\text{NHCH}_2\text{CH}_3$	14.8
7	N-Ethylperfluoro-5,5-dimethyl-1-hexanesulfonamide	$\text{CF}_3\text{C}(\text{CF}_3)(\text{CF}_2)_4\text{SO}_2\text{NHCH}_2\text{CH}_3$	0.2

* Percent of total N-ethylperfluorooctanesulfonamide isomers only.

** Systematic Name: N-Ethylperfluoro-2-octanesulfonamide.

Certified By: _____

B.G. Chittim, General Manager

Date: 11/15/2022

(mm/dd/yyyy)

Analytical Standard Record

22J0303

Description:	br-NMeFOSE	Expires:	10/07/2027
Standard Type:	Other	Prepared:	10/07/2022
Solvent:	MeOH	Prepared By:	Wellington Laboratories (Lot#:
Final Volume (mls):	1	Department:	br-NMeFOSE0922)
Vials:	1	Last Edit:	10/18/2022 13:41 by HGH

Analyte	Parent	CAS Number	Concentration	Units
NETFOSE		1691-99-2	50	ug/mL

**WELLINGTON
LABORATORIES****CERTIFICATE OF ANALYSIS
DOCUMENTATION****br-NMeFOSE****2-(N-Methylperfluorooctanesulfonamido)ethanol
Isomeric Mix**

<u>PRODUCT CODE:</u>	br-NMeFOSE
<u>LOT NUMBER:</u>	brNMeFOSE0922
<u>CONCENTRATION:</u>	50.0 ± 2.5 µg/mL
<u>SOLVENT(S):</u>	Methanol
<u>DATE PREPARED:</u> (mm/dd/yyyy)	09/02/2022
<u>LAST TESTED:</u> (mm/dd/yyyy)	09/07/2022 (HRGC/LRMS) 10/07/2022 (LC/MS)
<u>EXPIRY DATE:</u> (mm/dd/yyyy)	10/07/2027
<u>RECOMMENDED STORAGE:</u>	Store ampoule in a cool, dark place

DESCRIPTION:

The chemical purity has been determined to be ≥98% 2-(N-methylperfluorooctanesulfonamido)ethanol linear and branched isomers. The full name, structure, and percent composition for each of the isomeric components are given in Table A.

DOCUMENTATION/ DATA ATTACHED:

Table A: Isomeric Components and Percent Composition by ¹⁹F-NMR
Figure 1: HRGC/LRMS Data (Full Scan and Mass Spectrum)
Figure 2: LC/MS Data (Full Scan and Mass Spectrum)
Figure 3: LC/MS Data (SIR)
Figure 4: LC/MS/MS Data (Selected MRM Transitions)

ADDITIONAL INFORMATION:

- See page 2 for further details.
- CAS #: 24448-09-7 (for linear isomer).

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Analytical Standard Record

22J0304

Description:	br-NETFOSE	Expires:	10/07/2027
Standard Type:	Other	Prepared:	10/07/2022
Solvent:	MeOH	Prepared By:	Wellington Laboratories (Lot#: PNS)
Final Volume (mls):	1	Department:	br-NETFOSE1022)
Vials:	1	Last Edit:	10/18/2022 13:43 by HGH

Analyte	Parent	CAS Number	Concentration	Units
NETFOSE		1691-99-2	50	ug/mL



WELLINGTON LABORATORIES

CERTIFICATE OF ANALYSIS DOCUMENTATION

br-NEtFOSE

2-(N-Ethylperfluorooctanesulfonamido)ethanol Isomeric Mix

<u>PRODUCT CODE:</u>	br-NEtFOSE
<u>LOT NUMBER:</u>	brNEtFOSE1022
<u>CONCENTRATION:</u>	50.0 ± 2.5 µg/mL
<u>SOLVENT(S):</u>	Methanol
<u>DATE PREPARED:</u> (mm/dd/yyyy)	09/12/2022
<u>LAST TESTED:</u> (mm/dd/yyyy)	09/12/2022 (HRGC/LRMS) 10/07/2022 (LC/MS)
<u>EXPIRY DATE:</u> (mm/dd/yyyy)	10/07/2027
<u>RECOMMENDED STORAGE:</u>	Store ampoule in a cool, dark place

DESCRIPTION:

The chemical purity has been determined to be ≥98% 2-(N-ethylperfluorooctanesulfonamido)ethanol linear and branched isomers. The full name, structure, and percent composition for each of the isomeric components are given in Table A.

DOCUMENTATION/ DATA ATTACHED:

Table A: Isomeric Components and Percent Composition by ¹⁹F-NMR
 Figure 1: HRGC/LRMS Data (Full Scan and Mass Spectrum)
 Figure 2: LC/MS Data (Full Scan and Mass Spectrum)
 Figure 3: LC/MS Data (SIR)
 Figure 4: LC/MS/MS Data (Selected MRM Transitions)

ADDITIONAL INFORMATION:

- See page 2 for further details.
- CAS #: 1691-99-2 (for linear isomer).

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Table A: br-NEtFOSE; Isomeric Components and Percent Composition (by ^{19}F -NMR)*

Isomer	Compound	Structure	Percent Composition by ^{19}F -NMR
1	2-(N-Ethylperfluoro-1-octanesulfonamido)ethanol	$\text{CF}_3(\text{CF}_2)_7\text{SO}_2\text{NCH}_2\text{CH}_2\text{OH}$ CH_2CH_3	64.6
2	2-(N-Ethylperfluoro-2-methyl-1-heptanesulfonamido)ethanol	$\text{CF}_3(\text{CF}_2)_4\text{CFCF}_2\text{SO}_2\text{NCH}_2\text{CH}_2\text{OH}$ CF_3 CH_2CH_3	0.2
3	2-(N-Ethylperfluoro-3-methyl-1-heptanesulfonamido)ethanol	$\text{CF}_3(\text{CF}_2)_3\text{CF}(\text{CF}_2)_2\text{SO}_2\text{NCH}_2\text{CH}_2\text{OH}$ CF_3 CH_2CH_3	4.1
4	2-(N-Ethylperfluoro-4-methyl-1-heptanesulfonamido)ethanol	$\text{CF}_3(\text{CF}_2)_2\text{CF}(\text{CF}_2)_3\text{SO}_2\text{NCH}_2\text{CH}_2\text{OH}$ CF_3 CH_2CH_3	4.3
5	2-(N-Ethylperfluoro-5-methyl-1-heptanesulfonamido)ethanol	$\text{CF}_3\text{CF}_2\text{CF}(\text{CF}_2)_4\text{SO}_2\text{NCH}_2\text{CH}_2\text{OH}$ CF_3 CH_2CH_3	8.8
6	2-(N-Ethylperfluoro-6-methyl-1-heptanesulfonamido)ethanol	$\text{CF}_3\text{CF}(\text{CF}_2)_5\text{SO}_2\text{NCH}_2\text{CH}_2\text{OH}$ CF_3 CH_2CH_3	17.8
7	2-(N-Ethylperfluoro-5,5-dimethyl-1-hexanesulfonamido)ethanol	CF_3 $\text{CF}_3\text{C}(\text{CF}_2)_4\text{SO}_2\text{NCH}_2\text{CH}_2\text{OH}$ CF_3 CH_2CH_3	0.3

* Percent of total 2-(N-ethylperfluorooctanesulfonamido)ethanol isomers only.

Certified By: 
B.G. Chittim, General Manager

Date: 11/14/2022
(mm/dd/yyyy)

Analytical Standard Record

22J0420

Description: PFAS RES-MIX 1000ng/mL
 Standard Type: Other
 Solvent: MeOH
 Final Volume (mLs): 1
 Vials: 1

Expires: 04/24/2023
 Prepared: 10/26/2022
 Prepared By: Dipti Gokal
 Department: PFAS
 Last Edit: 10/26/2022 10:16 by HGH

Analyte	Parent	CAS Number	Concentration	Units
PFOA	22J0297	335-67-1	1	ug/mL
PFOSA	22J0298	754-91-6	1	ug/mL
NMeFOSA	22J0301	31506-32-8	1	ug/mL
NEtFOSA	22J0302	4151-50-2	1	ug/mL
NMeFOSE	22J0303	24448-09-7	1	ug/mL
NEtFOSE	22J0304	1691-99-2	1	ug/mL

Parent Standards used:

Standard	Description	Prepared	Prepared By	Lot Nbr	Expires	Last Edit	(mLs)
22J0297	T-PFOA	01/27/2022	Wellington Laboratories	TPFOA0122	01/27/2027	10/18/2022 12:59 by HGH	0.02
22J0298	br-FOSA	09/14/2022	Wellington Laboratories	br-FOSA0922	10/07/2027	10/18/2022 13:03 by HGH	0.02
22J0301	br-NMeFOSA	08/23/2022	Wellington Laboratories	beNMeFOSA0822	08/23/2027	10/18/2022 13:37 by HGH	0.02
22J0302	br-NEtFOSA	10/07/2022	Wellington Laboratories	beNEtFOSA0922	10/07/2027	10/18/2022 13:38 by HGH	0.02
22J0303	br-NMeFOSE	10/07/2022	Wellington Laboratories	beNMeFOSE0922	10/07/2027	10/26/2022 10:16 by HGH	0.02
22J0304	br-NEtFOSE	10/07/2022	Wellington Laboratories	beNEtFOSE1022	10/07/2027	10/18/2022 13:43 by HGH	0.02

Analytical Standard Record

22J0448

Description: PFAS - MIX 1633 20ng/mL
 Standard Type: Analyte Spike
 Solvent: MeOH
 Final Volume (mLs): 10
 Vials: 1

Expires: 04/25/2023
 Prepared: 10/27/2022
 Prepared By: Dipti Gokal
 Department: PFAS
 Last Edit: 10/27/2022 08:51 by DAG

Analyte	Parent	CAS Number	Concentration	Units
11CL-PF3OUDS	22I0153	763051-92-9	0.0378	ug/mL
3:3FTCA	22I0153	113507-82-7	0.08	ug/mL
4:2FTS	22I0153	757124-72-4	0.075	ug/mL
5:3FTCA	22I0153	914637-49-3	0.08	ug/mL
6:2FTS	22I0153	27619-97-2	0.076	ug/mL
7:3FTCA	22I0153	812-70-4	0.08	ug/mL
8:2FTS	22I0153	39108-34-4	0.0768	ug/mL
9CL-PF3ONS	22I0153	756426-58-1	0.0374	ug/mL
ADONA	22I0153	919005-14-4	0.0378	ug/mL
HFPO-DA	22I0153	13252-13-6	0.04	ug/mL
NETFOSA	22I0153	4151-50-2	0.08	ug/mL
NETFOSAA	22I0153	2991-50-6	0.02	ug/mL
NETFOSE	22I0153	1691-99-2	0.08	ug/mL
NFDHA	22I0153	151772-58-6	0.04	ug/mL
NMeFOSA	22I0153	31506-32-8	0.08	ug/mL
NMeFOSAA	22I0153	2355-31-9	0.02	ug/mL
NMeFOSE	22I0153	24448-09-7	0.08	ug/mL
PFBA	22I0153	375-22-4	0.08	ug/mL
PFBS	22I0153	375-73-5	0.0177	ug/mL
PFDA	22I0153	335-76-2	0.02	ug/mL
PFDOA	22I0153	307-55-1	0.02	ug/mL
PFDOS	22I0153	79780-39-5	0.0194	ug/mL
PFDS	22I0153	335-77-3	0.0193	ug/mL
PFEESA	22I0153	113507-82-7	0.0356	ug/mL
PFHPA	22I0153	375-85-9	0.02	ug/mL
PFHPS	22I0153	375-92-8	0.0191	ug/mL
PFHXA	22I0153	307-24-4	0.02	ug/mL
PFHXS	22I0153	355-46-4	0.0183	ug/mL
PFMBA	22I0153	863090-89-5	0.04	ug/mL
PFMPA	22I0153	377-73-1	0.04	ug/mL
PFNA	22I0153	375-95-1	0.02	ug/mL
PFNS	22I0153	68259-12-1	0.0192	ug/mL
PFOA	22I0153	335-67-1	0.02	ug/mL
PFOS	22I0153	1763-23-1	0.0186	ug/mL
PFOSA	22I0153	754-91-6	0.02	ug/mL
PFPEA	22I0153	2706-90-3	0.04	ug/mL
PFPEs	22I0153	630402-22-1	0.0188	ug/mL
PFTEDA	22I0153	376-06-7	0.02	ug/mL
PFTRDA	22I0153	72629-94-8	0.02	ug/mL
PFUnA	22I0153	2058-94-8	0.02	ug/mL

Analytical Standard Record

22J0448**Parent Standards used:**

Standard	Description	Prepared	Prepared By	Lot Nbr	Expires	Last Edit	(mls)
22I0153	PFAS - MIX 1633 200ng/mL	09/13/2022	In house	x	01/11/2025	09/15/2022 09:34 by DAG	1

Analytical Standard Record

22K0180

Description:	PFAS - MIX MXF 2 ug/mL	Expires:	01/11/2025
Standard Type:	Analyte Spike	Prepared:	01/10/2022
Solvent:	MeOH	Prepared By:	Wellington Laboratories (Lot#:
Final Volume (mLs):	1.2	Department:	PFAS MXF0122)
Vials:	1	Last Edit:	11/08/2022 13:30 by ABK

Analyte	Parent	CAS Number	Concentration	Units
11CL-PF3OUDS		763051-92-9	1.89	ug/mL
9CL-PF3ONS		756426-58-1	1.87	ug/mL
ADONA		919005-14-4	1.89	ug/mL
HFPO-DA		13252-13-6	2	ug/mL

Analytical Standard Record

22K0181

Description:	PFAS - MIX MXG 2 ug/mL	Expires:	02/22/2027
Standard Type:	Analyte Spike	Prepared:	02/07/2022
Solvent:	MeOH	Prepared By:	Wellington Laboratories (Lot#:
Final Volume (mLs):	1.2	Department:	PFAS MXG0222)
Vials:	1	Last Edit:	11/08/2022 13:31 by ABK

Analyte	Parent	CAS Number	Concentration	Units
NFDHA		151772-58-6	2	ug/mL
PFEESA		113507-82-7	1.78	ug/mL
PFMBA		863090-89-5	2	ug/mL
PFMPA		377-73-1	2	ug/mL

Analytical Standard Record

22K0182

Description:	PFAS - MIX MXH 1 ug/mL	Expires:	08/08/2027
Standard Type:	Analyte Spike	Prepared:	08/05/2022
Solvent:	MeOH	Prepared By:	Wellington Laboratories (Lot#:
Final Volume (mLs):	1.2	Department:	PFAS MXH0822)
Vials:	1	Last Edit:	11/08/2022 13:35 by ABK

Analyte	Parent	CAS Number	Concentration	Units
4:2FTS		757124-72-4	3.75	ug/mL
6:2FTS		27619-97-2	3.8	ug/mL
8:2FTS		39108-34-4	3.84	ug/mL
NETFOSAA		2991-50-6	1	ug/mL
NMeFOSAA		2355-31-9	1	ug/mL
PFBA		375-22-4	4	ug/mL
PFBS		375-73-5	0.887	ug/mL
PFDA		335-76-2	1	ug/mL
PFDOA		307-55-1	1	ug/mL
PFDOS		79780-39-5	0.97	ug/mL
PFDS		335-77-3	0.965	ug/mL
PFHPA		375-85-9	1	ug/mL
PFHPS		375-92-8	0.953	ug/mL
PFHXA		307-24-4	1	ug/mL
PFHXS		355-46-4	0.914	ug/mL
PFNA		375-95-1	1	ug/mL
PFNS		68259-12-1	0.962	ug/mL
PFOA		335-67-1	1	ug/mL
PFOS		1763-23-1	0.928	ug/mL
PFOSA		754-91-6	1	ug/mL
PFPEA		2706-90-3	2	ug/mL
PFPEs		630402-22-1	0.941	ug/mL
PFTEDA		376-06-7	1	ug/mL
PFTRDA		72629-94-8	1	ug/mL
PFUnA		2058-94-8	1	ug/mL

Analytical Standard Record

23A0022

Description: PFOS 0.4mg/ml
Standard Type: Other
Solvent: 62097
Final Volume (mls): 40
Vials: 1

Expires: 07/03/2023
Prepared: 01/04/2023
Prepared By: Dipti Gokal
Department: PFAS
Last Edit: 01/04/2023 11:14 by DAG

Analyte	Parent	CAS Number	Concentration	Units
PFOS	23A0024	1763-23-1	400	ug/mL

Parent Standards used:

Standard	Description	Prepared	Prepared By	Lot Nbr	Expires	Last Edit	(mls)
23A0024	PFOS 40%	01/04/2023	Sigma-Aldrich	0000100807	11/10/2023	01/04/2023 11:14 by DAG	0.04

Analytical Standard Record

23A0024

Description: PFOS 40%
Standard Type: Other
Solvent: methanol 77283
Final Volume (mls): 1
Vials: 1
Comments: neat-77283

Expires: 11/10/2023
Prepared: 01/04/2023
Prepared By: Dipti Gokal
Department: PFAS
Last Edit: 01/04/2023 11:14 by DAG

Analyte	Parent	CAS Number	Concentration	Units
PFOS		1763-23-1	400000	ug/g

Analytical Standard Record

23A0025

Description: PFOS 0.4ug/ml
Standard Type: Other
Solvent: 62097
Final Volume (mls): 40
Vials: 1

Expires: 07/03/2023
Prepared: 01/04/2023
Prepared By: Dipti Gokal
Department: PFAS
Last Edit: 01/04/2023 11:18 by DAG

Analyte	Parent	CAS Number	Concentration	Units
PFOS	23A0022	1763-23-1	0.4	ug/mL

Parent Standards used:

Standard	Description	Prepared	Prepared By	Lot Nbr	Expires	Last Edit	(mls)
23A0022	PFOS 0.4mg/ml	01/04/2023	In house	3761825	07/03/2023	01/04/2023 11:14 by DAG	0.04

Analytical Standard Record

23A0025

Description: PFOS 0.4ug/ml
Standard Type: Other
Solvent: 62097
Final Volume (mls): 40
Vials: 1

Expires: 07/03/2023
Prepared: 01/04/2023
Prepared By: Dipti Gokal
Department: PFAS
Last Edit: 01/04/2023 11:18 by DAG

Analyte	Parent	CAS Number	Concentration	Units
PFOS	23A0022	1763-23-1	0.4	ug/mL

Parent Standards used:

Standard	Description	Prepared	Prepared By	Lot Nbr	Expires	Last Edit	(mls)
23A0022	PFOS 0.4mg/ml	01/04/2023	In house	3761825	07/03/2023	01/04/2023 11:14 by DAG	0.04

Analytical Standard Record

23A0371

Description: PFAS - MIX 1633 200ng/mL
 Standard Type: Analyte Spike
 Solvent: MeOH 62244
 Final Volume (mLs): 5
 Vials: 1

Expires: 07/18/2023
 Prepared: 01/19/2023
 Prepared By: Dipti Gokal
 Department: PFAS
 Last Edit: 01/19/2023 14:21 by HGH

Analyte	Parent	CAS Number	Concentration	Units
3:3FTCA	21L0004	113507-82-7	0.8	ug/mL
5:3FTCA	21L0005	914637-49-3	0.8	ug/mL
7:3FTCA	21L0007	812-70-4	0.8	ug/mL
NMeFOSA	22J0301	31506-32-8	0.8	ug/mL
NEtFOSA	22J0302	4151-50-2	0.8	ug/mL
NMeFOSE	22J0303	24448-09-7	0.8	ug/mL
NEtFOSE	22J0304	1691-99-2	0.8	ug/mL
11CL-PF3OUDS	22K0180	763051-92-9	0.378	ug/mL
9CL-PF3ONS	22K0180	756426-58-1	0.374	ug/mL
ADONA	22K0180	919005-14-4	0.378	ug/mL
HFPO-DA	22K0180	13252-13-6	0.4	ug/mL
NFDHA	22K0181	151772-58-6	0.4	ug/mL
PFEESA	22K0181	113507-82-7	0.356	ug/mL
PFMBA	22K0181	863090-89-5	0.4	ug/mL
PFMPA	22K0181	377-73-1	0.4	ug/mL
4:2FTS	22K0182	757124-72-4	0.75	ug/mL
6:2FTS	22K0182	27619-97-2	0.76	ug/mL
8:2FTS	22K0182	39108-34-4	0.768	ug/mL
NEtFOSAA	22K0182	2991-50-6	0.2	ug/mL
NMeFOSAA	22K0182	2355-31-9	0.2	ug/mL
PFBA	22K0182	375-22-4	0.8	ug/mL
PFBS	22K0182	375-73-5	0.177	ug/mL
PFDA	22K0182	335-76-2	0.2	ug/mL
PFDOA	22K0182	307-55-1	0.2	ug/mL
PFDOS	22K0182	79780-39-5	0.194	ug/mL
PFDS	22K0182	335-77-3	0.193	ug/mL
PFHPA	22K0182	375-85-9	0.2	ug/mL
PFHPS	22K0182	375-92-8	0.191	ug/mL
PFHXA	22K0182	307-24-4	0.2	ug/mL
PFHXS	22K0182	355-46-4	0.183	ug/mL
PFNA	22K0182	375-95-1	0.2	ug/mL
PFNS	22K0182	68259-12-1	0.192	ug/mL
PFOA	22K0182	335-67-1	0.2	ug/mL
PFOS	22K0182	1763-23-1	0.186	ug/mL
PFOSA	22K0182	754-91-6	0.2	ug/mL
PFPEA	22K0182	2706-90-3	0.4	ug/mL
PFPEs	22K0182	630402-22-1	0.188	ug/mL
PFTEDA	22K0182	376-06-7	0.2	ug/mL
PFTRDA	22K0182	72629-94-8	0.2	ug/mL
PFUnA	22K0182	2058-94-8	0.2	ug/mL

Analytical Standard Record

23A0371

Parent Standards used:

Standard	Description	Prepared	Prepared By	Lot Nbr	Expires	Last Edit		(mls)
21L0004	PFAS - SAS 3:3FTA 50ug/mL	12/07/2021	Wellington Laboratories	FPrPA1020	11/12/2025	10/31/2022 14:39	by DAG	0.08
21L0005	PFAS - SAS 5:3FTA 50ug/mL	12/07/2021	Wellington Laboratories	FPePA1120	11/11/2025	10/31/2022 14:41	by DAG	0.08
21L0007	PFAS - SAS 7:3FTA 50ug/mL	12/07/2021	Wellington Laboratories	FHpPA1020	11/12/2025	10/31/2022 14:42	by DAG	0.08
22J0301	br-NMeFOSA	08/23/2022	Wellington Laboratories	beNMeFOSA0822	08/23/2027	10/18/2022 13:37	by HGH	0.08
22J0302	br-NEtFOSA	10/07/2022	Wellington Laboratories	beNEtFOSA0922	10/07/2027	10/18/2022 13:38	by HGH	0.08
22J0303	br-NMeFOSE	10/07/2022	Wellington Laboratories	beNMeFOSE0922	10/07/2027	10/26/2022 10:16	by HGH	0.08
22J0304	br-NEtFOSE	10/07/2022	Wellington Laboratories	beNEtFOSE1022	10/07/2027	10/18/2022 13:43	by HGH	0.08
22K0180	PFAS - MIX MXF 2 ug/mL	01/10/2022	Wellington Laboratories	PFACMXF0122	01/11/2025	11/08/2022 16:39	by DAG	1
22K0181	PFAS - MIX MXG 2 ug/mL	02/07/2022	Wellington Laboratories	PFACMXG0222	02/22/2027	11/08/2022 16:39	by DAG	1
22K0182	PFAS - MIX MXH 1 ug/mL	08/05/2022	Wellington Laboratories	PFACMXH0822	08/08/2027	11/08/2022 16:38	by DAG	1

Analytical Standard Record

23A0390

Description:	MPFAC-HIF-ES-EIS	Expires:	11/23/2025
Standard Type:	Other	Prepared:	10/28/2022
Solvent:	MeOH	Prepared By:	Wellington Laboratories (Lot#:
Final Volume (mls):	1.2	Department:	MPFAC-HIFES1022)
Vials:	1	Last Edit:	01/23/2023 15:26 by ABK
Lot Number:	MPFAC-HIFES1022		

Analyte	Parent	CAS Number	Concentration	Units
13C2-4:2FTS		13C2-4:2FTS	1	ug/mL
13C2-6:2FTS		13C2-6:2FTS	1	ug/mL
13C2-8:2FTS		13C2-8:2FTS	1	ug/mL
13C2-PFDOA		13C2-PFDOA	0.25	ug/mL
13C2-PFTEDA		13C2-PFTEDA	0.25	ug/mL
13C3-HFPO-DA		13C3-HFPO-DA	2	ug/mL
13C3-PFBS		13C3-PFBS	0.5	ug/mL
13C3-PFHXS		13C3-PFHXS	0.5	ug/mL
13C4-PFBA		13C4-PFBA	2	ug/mL
13C4-PFHHPA		13C4-PFHHPA	0.5	ug/mL
13C5-PFHXA		13C5-PFHXA	0.5	ug/mL
13C5-PFPEA		13C5-PFPEA	1	ug/mL
13C6-PFDA		13C6-PFDA	0.25	ug/mL
13C7-PFUDA		13C7-PFUDA	0.25	ug/mL
13C8-PFOA		13C8-PFOA	0.5	ug/mL
13C8-PFOS		13C8-PFOS	0.5	ug/mL
13C8-PFOSA		13C8-PFOSA	0.5	ug/mL
13C9-PFNA		13C9-PFNA	0.25	ug/mL
D3-NMEFOSA		D3-NMEFOSA	0.5	ug/mL
D3-NMEFOSAA		D3-NMEFOSAA	1	ug/mL
D5-NETFOSA		D5-NETFOSA	0.5	ug/mL
D5-NETFOSAA		D5-NETFOSAA	1	ug/mL
D7-NMEFOSE		D7-NMEFOSE	5	ug/mL
D9-NETFOSSE		D9-NETFOSSE	5	ug/mL



WELLINGTON LABORATORIES

CERTIFICATE OF ANALYSIS DOCUMENTATION

MPFAC-HIF-ES

**Mass-Labelled PFAS Extraction
Standard Solution/Mixture**

PRODUCT CODE: MPFAC-HIF-ES
LOT NUMBER: MPFACHIFES1022
SOLVENT(S): Methanol/Isopropanol (1%)/Water (<1%)
DATE PREPARED: (mm/dd/yyyy) 10/28/2022
LAST TESTED: (mm/dd/yyyy) 11/23/2022
EXPIRY DATE: (mm/dd/yyyy) 11/23/2025
RECOMMENDED STORAGE: Refrigerate ampoule

DESCRIPTION:

MPFAC-HIF-ES is a solution/mixture of ten mass-labelled (^{13}C) perfluoroalkylcarboxylic acids (C_4 - C_{12} , C_{14}), three mass-labelled (^{13}C) perfluoroalkanesulfonates (C_4 , C_6 , and C_8), three mass-labelled (one ^{13}C and two ^2H) perfluoro-1-octanesulfonamides, three mass-labelled (^{13}C) fluorotelomer sulfonates (4:2, 6:2, and 8:2), two mass-labelled (^2H) perfluorooctanesulfonamidoacetic acids, two mass-labelled (^2H) perfluorooctanesulfonamidoethanols, and mass-labelled (^{13}C) hexafluoropropylene oxide dimer acid (GenX, M3HFPO-DA). The components and their concentrations are given in Table A.

The individual ^{13}C -labelled components all have chemical purities >98% and isotopic purities of $\geq 99\%$. The individual ^2H -labelled components all have chemical purities >98% and isotopic purities of $\geq 98\%$.

DOCUMENTATION/ DATA ATTACHED:

Table A: Components and Concentrations of the Solution/Mixture
 Figure 1: LC/MS Data (SIR)
 Figure 2: LC/MS/MS Data (Selected MRM Transitions)

ADDITIONAL INFORMATION:

- See page 2 for further details.
- Contains 4 mole eq. of NaOH to prevent conversion of the carboxylic acids to their respective methyl esters.

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Table A: MPFAC-HIF-ES; Components and Concentrations
(ng/mL, \pm 5% in methanol/isopropanol (1%)/water (<1%))

Compound	Acronym	Concentration (ng/mL)		Peak Assignment in Figure 1
Perfluoro-n-(¹³ C ₄)butanoic acid	MPFBA	2000		1
Perfluoro-n-(¹³ C ₅)pentanoic acid	M5PFPeA	1000		2
Perfluoro-n-(1,2,3,4,6- ¹³ C ₅)hexanoic acid	M5PFHxA	500		5
Perfluoro-n-(1,2,3,4- ¹³ C ₄)heptanoic acid	M4PFHpA	500		7
Perfluoro-n-(¹³ C ₈)octanoic acid	M8PFOA	500		10
Perfluoro-n-(¹³ C ₉)nonanoic acid	M9PFNA	250		11
Perfluoro-n-(1,2,3,4,5,6- ¹³ C ₆)decanoic acid	M6PFDA	250		14
Perfluoro-n-(1,2,3,4,5,6,7- ¹³ C ₇)undecanoic acid	M7PFUdA	250		18
Perfluoro-n-(1,2- ¹³ C ₂)dodecanoic acid	MPFDoA	250		19
Perfluoro-n-(1,2- ¹³ C ₂)tetradecanoic acid	M2PFTeDA	250		22
Perfluoro-1-(¹³ C ₈)octanesulfonamide	M8FOSA	500		17
N-methyl-d ₃ -perfluoro-1-octanesulfonamide	d-N-MeFOSA	500		21
N-ethyl-d ₅ -perfluoro-1-octanesulfonamide	d-N-EtFOSA	500		24
N-methyl-d ₃ -perfluoro-1-octanesulfonamidoacetic acid	d3-N-MeFOSAA	1000		15
N-ethyl-d ₅ -perfluoro-1-octanesulfonamidoacetic acid	d5-N-EtFOSAA	1000		16
2-(N-methyl-d ₃ -perfluoro-1-octanesulfonamido)ethan-d ₄ -ol	d7-N-MeFOSE	5000		20
2-(N-ethyl-d ₅ -perfluoro-1-octanesulfonamido)ethan-d ₄ -ol	d9-N-EtFOSE	5000		23
2,3,3,3-Tetrafluoro-2-(1,1,2,2,3,3,3-heptafluoropropoxy)(¹³ C ₃)propanoic acid	M3HFPO-DA	2000		6
Compound	Acronym	Concentration* (ng/mL)		Peak Assignment in Figure 1
		as the salt	as the acid	
Sodium perfluoro-1-(2,3,4- ¹³ C ₃)butanesulfonate	M3PFBS	500	466	3
Sodium perfluoro-1-(1,2,3- ¹³ C ₃)hexanesulfonate	M3PFHxS	500	474	8
Sodium perfluoro-1-(¹³ C ₈)octanesulfonate	M8PFOS	500	479	12
Sodium 1H,1H,2H,2H-perfluoro-(1,2- ¹³ C ₂)hexanesulfonate	M2-4:2FTS	1000	938	4
Sodium 1H,1H,2H,2H-perfluoro-(1,2- ¹³ C ₂)octanesulfonate	M2-6:2FTS	1000	951	9
Sodium 1H,1H,2H,2H-perfluoro-(1,2- ¹³ C ₂)decanesulfonate	M2-8:2FTS	1000	960	13

* Concentrations have been rounded to three significant figures.

Certified By:


B.G. Chittim, General Manager

Date: 11/24/2022
(mm/dd/yyyy)

Analytical Standard Record

23B0165

Description:	MPFAC-HIF-ES-EIS	Expires:	08/06/2023
Standard Type:	Other	Prepared:	02/07/2023
Solvent:	MeOH	Prepared By:	Wellington Laboratories (Lot#:
Final Volume (mls):	1.2	Department:	MPFAC-HIFES1022)
Vials:	1	Last Edit:	02/07/2023 16:13 by ABK
Lot Number:	MPFAC-HIFES1022		

Analyte	Parent	CAS Number	Concentration	Units
13C2-4:2FTS		13C2-4:2FTS	1	ug/mL
13C2-6:2FTS		13C2-6:2FTS	1	ug/mL
13C2-8:2FTS		13C2-8:2FTS	1	ug/mL
13C2-PFDOA		13C2-PFDOA	0.25	ug/mL
13C2-PFTEDA		13C2-PFTEDA	0.25	ug/mL
13C3-HFPO-DA		13C3-HFPO-DA	2	ug/mL
13C3-PFBS		13C3-PFBS	0.5	ug/mL
13C3-PFHXS		13C3-PFHXS	0.5	ug/mL
13C4-PFBA		13C4-PFBA	2	ug/mL
13C4-PFHHPA		13C4-PFHHPA	0.5	ug/mL
13C5-PFHXA		13C5-PFHXA	0.5	ug/mL
13C5-PFPEA		13C5-PFPEA	1	ug/mL
13C6-PFDA		13C6-PFDA	0.25	ug/mL
13C7-PFUDA		13C7-PFUDA	0.25	ug/mL
13C8-PFOA		13C8-PFOA	0.5	ug/mL
13C8-PFOS		13C8-PFOS	0.5	ug/mL
13C8-PFOSA		13C8-PFOSA	0.5	ug/mL
13C9-PFNA		13C9-PFNA	0.25	ug/mL
D3-NMEFOSA		D3-NMEFOSA	0.5	ug/mL
D3-NMEFOSAA		D3-NMEFOSAA	1	ug/mL
D5-NETFOSA		D5-NETFOSA	0.5	ug/mL
D5-NETFOSAA		D5-NETFOSAA	1	ug/mL
D7-NMEFOSE		D7-NMEFOSE	5	ug/mL
D9-NETFOSSE		D9-NETFOSSE	5	ug/mL



WELLINGTON LABORATORIES

CERTIFICATE OF ANALYSIS DOCUMENTATION

MPFAC-HIF-ES

**Mass-Labelled PFAS Extraction
Standard Solution/Mixture**

PRODUCT CODE: MPFAC-HIF-ES
LOT NUMBER: MPFACHIFES1022
SOLVENT(S): Methanol/Isopropanol (1%)/Water (<1%)
DATE PREPARED: (mm/dd/yyyy) 10/28/2022
LAST TESTED: (mm/dd/yyyy) 11/23/2022
EXPIRY DATE: (mm/dd/yyyy) 11/23/2025
RECOMMENDED STORAGE: Refrigerate ampoule

DESCRIPTION:

MPFAC-HIF-ES is a solution/mixture of ten mass-labelled (^{13}C) perfluoroalkylcarboxylic acids (C_4 - C_{12} , C_{14}), three mass-labelled (^{13}C) perfluoroalkanesulfonates (C_4 , C_6 , and C_8), three mass-labelled (one ^{13}C and two ^2H) perfluoro-1-octanesulfonamides, three mass-labelled (^{13}C) fluorotelomer sulfonates (4:2, 6:2, and 8:2), two mass-labelled (^2H) perfluorooctanesulfonamidoacetic acids, two mass-labelled (^2H) perfluorooctanesulfonamidoethanols, and mass-labelled (^{13}C) hexafluoropropylene oxide dimer acid (GenX, M3HFPO-DA). The components and their concentrations are given in Table A.

The individual ^{13}C -labelled components all have chemical purities >98% and isotopic purities of $\geq 99\%$. The individual ^2H -labelled components all have chemical purities >98% and isotopic purities of $\geq 98\%$.

DOCUMENTATION/ DATA ATTACHED:

Table A: Components and Concentrations of the Solution/Mixture
 Figure 1: LC/MS Data (SIR)
 Figure 2: LC/MS/MS Data (Selected MRM Transitions)

ADDITIONAL INFORMATION:

- See page 2 for further details.
- Contains 4 mole eq. of NaOH to prevent conversion of the carboxylic acids to their respective methyl esters.

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Table A: MPFAC-HIF-ES; Components and Concentrations
(ng/mL, \pm 5% in methanol/isopropanol (1%)/water (<1%))

Compound	Acronym	Concentration (ng/mL)		Peak Assignment in Figure 1
Perfluoro-n-(¹³ C ₄)butanoic acid	MPFBA	2000		1
Perfluoro-n-(¹³ C ₅)pentanoic acid	M5PFPeA	1000		2
Perfluoro-n-(1,2,3,4,6- ¹³ C ₅)hexanoic acid	M5PFHxA	500		5
Perfluoro-n-(1,2,3,4- ¹³ C ₄)heptanoic acid	M4PFHpA	500		7
Perfluoro-n-(¹³ C ₈)octanoic acid	M8PFOA	500		10
Perfluoro-n-(¹³ C ₉)nonanoic acid	M9PFNA	250		11
Perfluoro-n-(1,2,3,4,5,6- ¹³ C ₆)decanoic acid	M6PFDA	250		14
Perfluoro-n-(1,2,3,4,5,6,7- ¹³ C ₇)undecanoic acid	M7PFUdA	250		18
Perfluoro-n-(1,2- ¹³ C ₂)dodecanoic acid	MPFDoA	250		19
Perfluoro-n-(1,2- ¹³ C ₂)tetradecanoic acid	M2PFTeDA	250		22
Perfluoro-1-(¹³ C ₈)octanesulfonamide	M8FOSA	500		17
N-methyl-d ₃ -perfluoro-1-octanesulfonamide	d-N-MeFOSA	500		21
N-ethyl-d ₅ -perfluoro-1-octanesulfonamide	d-N-EtFOSA	500		24
N-methyl-d ₃ -perfluoro-1-octanesulfonamidoacetic acid	d3-N-MeFOSAA	1000		15
N-ethyl-d ₅ -perfluoro-1-octanesulfonamidoacetic acid	d5-N-EtFOSAA	1000		16
2-(N-methyl-d ₃ -perfluoro-1-octanesulfonamido)ethan-d ₄ -ol	d7-N-MeFOSE	5000		20
2-(N-ethyl-d ₅ -perfluoro-1-octanesulfonamido)ethan-d ₄ -ol	d9-N-EtFOSE	5000		23
2,3,3,3-Tetrafluoro-2-(1,1,2,2,3,3,3-heptafluoropropoxy)(¹³ C ₃)propanoic acid	M3HFPO-DA	2000		6
Compound	Acronym	Concentration* (ng/mL)		Peak Assignment in Figure 1
		as the salt	as the acid	
Sodium perfluoro-1-(2,3,4- ¹³ C ₃)butanesulfonate	M3PFBS	500	466	3
Sodium perfluoro-1-(1,2,3- ¹³ C ₃)hexanesulfonate	M3PFHxS	500	474	8
Sodium perfluoro-1-(¹³ C ₈)octanesulfonate	M8PFOS	500	479	12
Sodium 1H,1H,2H,2H-perfluoro-(1,2- ¹³ C ₂)hexanesulfonate	M2-4:2FTS	1000	938	4
Sodium 1H,1H,2H,2H-perfluoro-(1,2- ¹³ C ₂)octanesulfonate	M2-6:2FTS	1000	951	9
Sodium 1H,1H,2H,2H-perfluoro-(1,2- ¹³ C ₂)decanesulfonate	M2-8:2FTS	1000	960	13

* Concentrations have been rounded to three significant figures.

Certified By:


B.G. Chittim, General Manager

Date: 11/24/2022
(mm/dd/yyyy)