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NELAP Certification Number: CA00046

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State Certification Number:

February 16, 2023

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

RE: Red Hill AFFF Assessment Sampling  
23B0091

Enclosed are the results of analyses for samples received by our laboratory on 2/11/2023. 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,

Greg Salata For Gregory Salata  
Project Manager

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Project: Red Hill AFFF Assessment Sampling  
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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 3:3FTcA recovered above the upper control limit in the SC00636-LCV1.

## Samples in this Report

Lab ID	Sample	Matrix	Date Sampled	Date Received
23B0091-01	AF-RHWW17-WGN01LF-2302W1	Water	02/10/2023 11:00	02/11/2023
23B0091-02	AF-RHWW17D-WGN01LF-2302W1	Water	02/10/2023 12:15	02/11/2023
23B0091-03	AF-RHWW17D-WQFB01-2302W1	Water	02/10/2023 11:30	02/11/2023
23B0091-04	AF-RHWW225401-WGN01B-2302W1	Water	02/10/2023 08:55	02/11/2023



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### Containers Received

Lab ID	Container Type	Count	Preservation Check
23B0091-01	500mL P	2	
23B0091-02	500mL P	2	
23B0091-03	500mL P	2	
23B0091-04	500mL P	2	

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## Sample Results

**Sample: AF-RHMW17-WGN01LF-2302W1**  
**23B0091-01 (Water)**

### Per- and Polyfluoroalkyl Substances

Analyte	Result /Qual	LOQ	LOD	DL	Units	Date Analyzed	DF	Method	Prep Batch
PFBA	2.1	1.4	0.71	0.18	ng/L	02/14/23	1	EPA 1633	BCB0232
PFPEA	5.3	0.71	0.35	0.057	ng/L	02/14/23	1	EPA 1633	BCB0232
PFHXA	2.5	0.35	0.18	0.048	ng/L	02/14/23	1	EPA 1633	BCB0232
PFHPA	0.44	0.35	0.18	0.036	ng/L	02/14/23	1	EPA 1633	BCB0232
PFOA	0.27 J	0.35	0.18	0.13	ng/L	02/14/23	1	EPA 1633	BCB0232
PFNA	0.18 U	0.35	0.18	0.072	ng/L	02/14/23	1	EPA 1633	BCB0232
PFDA	0.18 U	0.35	0.18	0.089	ng/L	02/14/23	1	EPA 1633	BCB0232
PFUnA	0.18 U	0.35	0.18	0.14	ng/L	02/14/23	1	EPA 1633	BCB0232
PFDOA	0.18 U	0.35	0.18	0.099	ng/L	02/14/23	1	EPA 1633	BCB0232
PFTRDA	0.26 U	0.35	0.26	0.18	ng/L	02/14/23	1	EPA 1633	BCB0232
PFTEDA	0.18 U IR2,	0.35	0.18	0.17	ng/L	02/14/23	1	EPA 1633	BCB0232
PFBS	0.22 J	0.35	0.18	0.032	ng/L	02/14/23	1	EPA 1633	BCB0232
PFPEs	0.18 U	0.35	0.18	0.055	ng/L	02/14/23	1	EPA 1633	BCB0232
PFHXS	0.18 U	0.35	0.18	0.028	ng/L	02/14/23	1	EPA 1633	BCB0232
PFHPS	0.18 U	0.35	0.18	0.045	ng/L	02/14/23	1	EPA 1633	BCB0232
PFOS	0.19 J MI5,	0.35	0.18	0.056	ng/L	02/14/23	1	EPA 1633	BCB0232
PFNS	0.18 U	0.35	0.18	0.11	ng/L	02/14/23	1	EPA 1633	BCB0232
PFDS	0.18 U	0.35	0.18	0.13	ng/L	02/14/23	1	EPA 1633	BCB0232
PFDOS	0.18 U	0.35	0.18	0.11	ng/L	02/14/23	1	EPA 1633	BCB0232
4:2FTS	0.71 U	1.4	0.71	0.26	ng/L	02/14/23	1	EPA 1633	BCB0232
6:2FTS	0.49 J	1.4	0.71	0.28	ng/L	02/14/23	1	EPA 1633	BCB0232
8:2FTS	0.71 U	1.4	0.71	0.073	ng/L	02/14/23	1	EPA 1633	BCB0232
PFOSA	0.18 U	0.35	0.18	0.092	ng/L	02/14/23	1	EPA 1633	BCB0232
NMeFOSA	0.71 U	1.4	0.71	0.42	ng/L	02/14/23	1	EPA 1633	BCB0232
NEtFOSA	0.71 U	1.4	0.71	0.36	ng/L	02/14/23	1	EPA 1633	BCB0232
NMeFOSAA	0.18 U	0.35	0.18	0.093	ng/L	02/14/23	1	EPA 1633	BCB0232
NEtFOSAA	0.18 U	0.35	0.18	0.10	ng/L	02/14/23	1	EPA 1633	BCB0232
NMeFOSE	1.1 U	1.4	1.1	0.89	ng/L	02/14/23	1	EPA 1633	BCB0232
NEtFOSE	1.1 U	1.4	1.1	0.92	ng/L	02/14/23	1	EPA 1633	BCB0232
HFPO-DA	0.35 U	0.71	0.35	0.15	ng/L	02/14/23	1	EPA 1633	BCB0232
ADONA	0.35 U	0.71	0.35	0.11	ng/L	02/14/23	1	EPA 1633	BCB0232
PFEESA	0.35 U	0.71	0.35	0.096	ng/L	02/14/23	1	EPA 1633	BCB0232
PFMPA	0.35 U	0.71	0.35	0.048	ng/L	02/14/23	1	EPA 1633	BCB0232
PFMBA	0.35 U	0.71	0.35	0.080	ng/L	02/14/23	1	EPA 1633	BCB0232
NFDHA	0.35 U	0.71	0.35	0.27	ng/L	02/14/23	1	EPA 1633	BCB0232
9CL-PF3ONS	0.35 U	0.71	0.35	0.18	ng/L	02/14/23	1	EPA 1633	BCB0232
11CL-PF3OUDS	0.35 U	0.71	0.35	0.18	ng/L	02/14/23	1	EPA 1633	BCB0232
3:3FTCA	0.71 U	1.4	0.71	0.51	ng/L	02/14/23	1	EPA 1633	BCB0232
5:3FTCA	0.71 U	1.4	0.71	0.39	ng/L	02/14/23	1	EPA 1633	BCB0232
7:3FTCA	0.71 U	1.4	0.71	0.49	ng/L	02/14/23	1	EPA 1633	BCB0232
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Surrogate: 13C4-PFBA	83.0%		10-130			02/14/23	1	EPA 1633	BCB0232
Surrogate: 13C5-PFPEA	91.0%		35-150			02/14/23	1	EPA 1633	BCB0232
Surrogate: 13C5-PFHXA	86.3%		55-150			02/14/23	1	EPA 1633	BCB0232
Surrogate: 13C4-PFHPA	81.7%		55-150			02/14/23	1	EPA 1633	BCB0232
Surrogate: 13C8-PFOA	82.7%		60-140			02/14/23	1	EPA 1633	BCB0232

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## Sample Results

(Continued)

**Sample: AF-RHMW17-WGN01LF-2302W1 (Continued)**

**23B0091-01 (Water)**

### Per- and Polyfluoroalkyl Substances (Continued)

Analyte	Result /Qual	LOQ	LOD	DL	Units	Date Analyzed	DF	Method	Prep Batch
Surrogate: 13C9-PFNA	81.7%		55-140			02/14/23	1	EPA 1633	BCB0232
Surrogate: 13C6-PFDA	84.3%		50-140			02/14/23	1	EPA 1633	BCB0232
Surrogate: 13C7-PFUnA	86.7%		30-140			02/14/23	1	EPA 1633	BCB0232
Surrogate: 13C2-PFDOA	91.2%		10-150			02/14/23	1	EPA 1633	BCB0232
Surrogate: 13C2-PFTEDA	80.2%		10-130			02/14/23	1	EPA 1633	BCB0232
Surrogate: 13C3-PFBS	84.9%		55-150			02/14/23	1	EPA 1633	BCB0232
Surrogate: 13C3-PFHXS	82.9%		55-150			02/14/23	1	EPA 1633	BCB0232
Surrogate: 13C8-PFOS	84.5%		45-140			02/14/23	1	EPA 1633	BCB0232
Surrogate: 13C2-4:2FTS	182%		60-200			02/14/23	1	EPA 1633	BCB0232
Surrogate: 13C2-6:2FTS	67.8%		60-200			02/14/23	1	EPA 1633	BCB0232
Surrogate: 13C2-8:2FTS	69.1%		50-200			02/14/23	1	EPA 1633	BCB0232
Surrogate: 13C8-PFOSA	77.8%		30-130			02/14/23	1	EPA 1633	BCB0232
Surrogate: D3-NMEFOSA	63.6%		15-130			02/14/23	1	EPA 1633	BCB0232
Surrogate: D5-NETFOSA	59.9%		10-130			02/14/23	1	EPA 1633	BCB0232
Surrogate: D3-NMEFOSAA	72.9%		45-200			02/14/23	1	EPA 1633	BCB0232
Surrogate: D5-NETFOSAA	81.4%		10-200			02/14/23	1	EPA 1633	BCB0232
Surrogate: D7-NMEFOSE	52.6%		10-150			02/14/23	1	EPA 1633	BCB0232
Surrogate: D9-NETFOSE	70.8%		10-150			02/14/23	1	EPA 1633	BCB0232
Surrogate: 13C3-HFPO-DA	90.1%		25-160			02/14/23	1	EPA 1633	BCB0232

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## Sample Results (Continued)

**Sample: AF-RHMW17D-WGN01LF-2302W1  
23B0091-02 (Water)**

### Per- and Polyfluoroalkyl Substances

Analyte	Result /Qual	LOQ	LOD	DL	Units	Date Analyzed	DF	Method	Prep Batch
PFBA	0.60 J	1.4	0.70	0.18	ng/L	02/15/23	1	EPA 1633	BCB0232
PFPEA	0.13 J	0.70	0.35	0.057	ng/L	02/15/23	1	EPA 1633	BCB0232
PFHXA	0.27 J IR2,	0.35	0.18	0.048	ng/L	02/15/23	1	EPA 1633	BCB0232
PFHPA	0.045 J IR2,	0.35	0.18	0.036	ng/L	02/15/23	1	EPA 1633	BCB0232
PFOA	0.18 U	0.35	0.18	0.13	ng/L	02/15/23	1	EPA 1633	BCB0232
PFNA	0.18 U	0.35	0.18	0.072	ng/L	02/15/23	1	EPA 1633	BCB0232
PFDA	0.18 U	0.35	0.18	0.089	ng/L	02/15/23	1	EPA 1633	BCB0232
PFUnA	0.18 U	0.35	0.18	0.14	ng/L	02/15/23	1	EPA 1633	BCB0232
PFDOA	0.18 U	0.35	0.18	0.098	ng/L	02/15/23	1	EPA 1633	BCB0232
PFTRDA	0.26 U	0.35	0.26	0.18	ng/L	02/15/23	1	EPA 1633	BCB0232
PFTEDA	0.18 U	0.35	0.18	0.17	ng/L	02/15/23	1	EPA 1633	BCB0232
PFBS	0.081 J	0.35	0.18	0.032	ng/L	02/15/23	1	EPA 1633	BCB0232
PFPEs	0.18 U	0.35	0.18	0.055	ng/L	02/15/23	1	EPA 1633	BCB0232
PFHXS	0.052 J	0.35	0.18	0.028	ng/L	02/15/23	1	EPA 1633	BCB0232
PFHPS	0.18 U	0.35	0.18	0.045	ng/L	02/15/23	1	EPA 1633	BCB0232
PFOS	0.12 J IR2,	0.35	0.18	0.056	ng/L	02/15/23	1	EPA 1633	BCB0232
PFNS	0.17 J IR1,	0.35	0.18	0.11	ng/L	02/15/23	1	EPA 1633	BCB0232
PFDS	0.18 U	0.35	0.18	0.13	ng/L	02/15/23	1	EPA 1633	BCB0232
PFDOS	0.18 U	0.35	0.18	0.11	ng/L	02/15/23	1	EPA 1633	BCB0232
4:2FTS	0.70 U	1.4	0.70	0.26	ng/L	02/15/23	1	EPA 1633	BCB0232
6:2FTS	6.1	1.4	0.70	0.28	ng/L	02/15/23	1	EPA 1633	BCB0232
8:2FTS	0.70 U	1.4	0.70	0.072	ng/L	02/15/23	1	EPA 1633	BCB0232
PFOSA	0.18 U	0.35	0.18	0.091	ng/L	02/15/23	1	EPA 1633	BCB0232
NMeFOSA	0.70 U	1.4	0.70	0.42	ng/L	02/15/23	1	EPA 1633	BCB0232
NEtFOSA	0.70 U	1.4	0.70	0.36	ng/L	02/15/23	1	EPA 1633	BCB0232
NMeFOSAA	0.18 U	0.35	0.18	0.093	ng/L	02/15/23	1	EPA 1633	BCB0232
NEtFOSAA	0.18 U	0.35	0.18	0.10	ng/L	02/15/23	1	EPA 1633	BCB0232
NMeFOSE	1.1 U	1.4	1.1	0.89	ng/L	02/15/23	1	EPA 1633	BCB0232
NEtFOSE	1.1 U	1.4	1.1	0.92	ng/L	02/15/23	1	EPA 1633	BCB0232
HFPO-DA	0.35 U	0.70	0.35	0.15	ng/L	02/15/23	1	EPA 1633	BCB0232
ADONA	0.35 U	0.70	0.35	0.11	ng/L	02/15/23	1	EPA 1633	BCB0232
PFEESA	0.35 U	0.70	0.35	0.096	ng/L	02/15/23	1	EPA 1633	BCB0232
PFMPA	0.35 U	0.70	0.35	0.047	ng/L	02/15/23	1	EPA 1633	BCB0232
PFMBA	0.35 U	0.70	0.35	0.080	ng/L	02/15/23	1	EPA 1633	BCB0232
NFDHA	0.35 U	0.70	0.35	0.26	ng/L	02/15/23	1	EPA 1633	BCB0232
9CL-PF3ONS	0.35 U	0.70	0.35	0.18	ng/L	02/15/23	1	EPA 1633	BCB0232
11CL-PF3OUDS	0.35 U	0.70	0.35	0.18	ng/L	02/15/23	1	EPA 1633	BCB0232
3:3FTCA	0.70 U	1.4	0.70	0.50	ng/L	02/15/23	1	EPA 1633	BCB0232
5:3FTCA	0.70 U	1.4	0.70	0.39	ng/L	02/15/23	1	EPA 1633	BCB0232
7:3FTCA	0.70 U	1.4	0.70	0.49	ng/L	02/15/23	1	EPA 1633	BCB0232
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Surrogate: 13C4-PFBA	81.8%		10-130			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C5-PFPEA	63.6%		35-150			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C5-PFHXA	76.2%		55-150			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C4-PFHPA	74.6%		55-150			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C8-PFOA	90.3%		60-140			02/15/23	1	EPA 1633	BCB0232

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Project Manager: Watson Tanji

Reported: 02/16/2023 15:45

## Sample Results

(Continued)

**Sample: AF-RHMW17D-WGN01LF-2302W1 (Continued)**

**23B0091-02 (Water)**

### Per- and Polyfluoroalkyl Substances (Continued)

Analyte	Result /Qual	LOQ	LOD	DL	Units	Date Analyzed	DF	Method	Prep Batch
Surrogate: 13C9-PFNA	85.9%		55-140			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C6-PFDA	90.4%		50-140			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C7-PFUnA	152% S2		30-140			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C7-PFUnA	71.8%		30-140			02/15/23	10	EPA 1633	BCB0232
Surrogate: 13C2-PFDOA	126%		10-150			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C2-PFTEDA	129%		10-130			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C3-PFBS	80.9%		55-150			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C3-PFHXS	73.5%		55-150			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C8-PFOS	78.2%		45-140			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C2-4:2FTS	190%		60-200			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C2-6:2FTS	165%		60-200			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C2-8:2FTS	110%		50-200			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C8-PFOA	67.0%		30-130			02/15/23	1	EPA 1633	BCB0232
Surrogate: D3-NMEFOA	52.1%		15-130			02/15/23	1	EPA 1633	BCB0232
Surrogate: D5-NETFOA	51.2%		10-130			02/15/23	1	EPA 1633	BCB0232
Surrogate: D3-NMEFOSAA	63.8%		45-200			02/15/23	1	EPA 1633	BCB0232
Surrogate: D5-NETFOSAA	99.3%		10-200			02/15/23	1	EPA 1633	BCB0232
Surrogate: D7-NMEFOSE	41.3%		10-150			02/15/23	1	EPA 1633	BCB0232
Surrogate: D9-NETFOSE	59.1%		10-150			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C3-HFPO-DA	79.1%		25-160			02/15/23	1	EPA 1633	BCB0232

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: 02/16/2023 15:45

**Sample Results****(Continued)**

**Sample: AF-RHMW17D-WQFB01-2302W1**  
**23B0091-03 (Water)**

**Per- and Polyfluoroalkyl Substances**

Analyte	Result /Qual	LOQ	LOD	DL	Units	Date Analyzed	DF	Method	Prep Batch
PFBA	0.72 U	1.4	0.72	0.19	ng/L	02/15/23	1	EPA 1633	BCB0232
PFPEA	0.36 U	0.72	0.36	0.058	ng/L	02/15/23	1	EPA 1633	BCB0232
PFHXA	0.18 U	0.36	0.18	0.049	ng/L	02/15/23	1	EPA 1633	BCB0232
PFHPA	0.18 U	0.36	0.18	0.037	ng/L	02/15/23	1	EPA 1633	BCB0232
PFOA	0.18 U	0.36	0.18	0.14	ng/L	02/15/23	1	EPA 1633	BCB0232
PFNA	0.18 U	0.36	0.18	0.073	ng/L	02/15/23	1	EPA 1633	BCB0232
PFDA	0.18 U	0.36	0.18	0.091	ng/L	02/15/23	1	EPA 1633	BCB0232
PFUnA	0.18 U	0.36	0.18	0.14	ng/L	02/15/23	1	EPA 1633	BCB0232
PFDOA	0.18 U	0.36	0.18	0.10	ng/L	02/15/23	1	EPA 1633	BCB0232
PFTRDA	0.27 U	0.36	0.27	0.18	ng/L	02/15/23	1	EPA 1633	BCB0232
PFTEDA	0.18 U	0.36	0.18	0.18	ng/L	02/15/23	1	EPA 1633	BCB0232
PFBS	0.18 U	0.36	0.18	0.033	ng/L	02/15/23	1	EPA 1633	BCB0232
PFPEs	0.18 U	0.36	0.18	0.056	ng/L	02/15/23	1	EPA 1633	BCB0232
PFHXS	0.18 U	0.36	0.18	0.028	ng/L	02/15/23	1	EPA 1633	BCB0232
PFHPS	0.18 U	0.36	0.18	0.046	ng/L	02/15/23	1	EPA 1633	BCB0232
PFOS	0.17 J M15,	0.36	0.18	0.057	ng/L	02/15/23	1	EPA 1633	BCB0232
PFNS	0.18 U	0.36	0.18	0.11	ng/L	02/15/23	1	EPA 1633	BCB0232
PFDS	0.18 U	0.36	0.18	0.14	ng/L	02/15/23	1	EPA 1633	BCB0232
PFDOS	0.18 U	0.36	0.18	0.11	ng/L	02/15/23	1	EPA 1633	BCB0232
4:2FTS	0.72 U	1.4	0.72	0.26	ng/L	02/15/23	1	EPA 1633	BCB0232
6:2FTS	0.72 U	1.4	0.72	0.28	ng/L	02/15/23	1	EPA 1633	BCB0232
8:2FTS	0.72 U	1.4	0.72	0.074	ng/L	02/15/23	1	EPA 1633	BCB0232
PFOSA	0.18 U	0.36	0.18	0.093	ng/L	02/15/23	1	EPA 1633	BCB0232
NMeFOSA	0.72 U	1.4	0.72	0.42	ng/L	02/15/23	1	EPA 1633	BCB0232
NEtFOSA	0.72 U	1.4	0.72	0.37	ng/L	02/15/23	1	EPA 1633	BCB0232
NMeFOSAA	0.18 U	0.36	0.18	0.095	ng/L	02/15/23	1	EPA 1633	BCB0232
NEtFOSAA	0.18 U	0.36	0.18	0.10	ng/L	02/15/23	1	EPA 1633	BCB0232
NMeFOSE	1.1 U	1.4	1.1	0.91	ng/L	02/15/23	1	EPA 1633	BCB0232
NEtFOSE	1.1 U	1.4	1.1	0.94	ng/L	02/15/23	1	EPA 1633	BCB0232
HFPO-DA	0.36 U	0.72	0.36	0.16	ng/L	02/15/23	1	EPA 1633	BCB0232
ADONA	0.36 U	0.72	0.36	0.11	ng/L	02/15/23	1	EPA 1633	BCB0232
PFEESA	0.36 U	0.72	0.36	0.098	ng/L	02/15/23	1	EPA 1633	BCB0232
PFMPA	0.36 U	0.72	0.36	0.048	ng/L	02/15/23	1	EPA 1633	BCB0232
PFMBA	0.36 U	0.72	0.36	0.081	ng/L	02/15/23	1	EPA 1633	BCB0232
NFDHA	0.36 U	0.72	0.36	0.27	ng/L	02/15/23	1	EPA 1633	BCB0232
9CL-PF3ONS	0.36 U	0.72	0.36	0.19	ng/L	02/15/23	1	EPA 1633	BCB0232
11CL-PF3OUDS	0.36 U	0.72	0.36	0.19	ng/L	02/15/23	1	EPA 1633	BCB0232
3:3FTCA	0.72 U	1.4	0.72	0.52	ng/L	02/15/23	1	EPA 1633	BCB0232
5:3FTCA	0.72 U	1.4	0.72	0.40	ng/L	02/15/23	1	EPA 1633	BCB0232
7:3FTCA	0.72 U	1.4	0.72	0.50	ng/L	02/15/23	1	EPA 1633	BCB0232
<hr/>									
Surrogate: 13C4-PFBA	86.7%		10-130			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C5-PFPEA	95.5%		35-150			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C5-PFHXA	88.7%		55-150			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C4-PFHPA	88.4%		55-150			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C8-PFOA	88.2%		60-140			02/15/23	1	EPA 1633	BCB0232

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

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## Sample Results

(Continued)

**Sample: AF-RHMW17D-WQFB01-2302W1 (Continued)**

**23B0091-03 (Water)**

### Per- and Polyfluoroalkyl Substances (Continued)

Analyte	Result /Qual	LOQ	LOD	DL	Units	Date Analyzed	DF	Method	Prep Batch
Surrogate: 13C9-PFNA	89.7%		55-140			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C6-PFDA	79.7%		50-140			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C7-PFUnA	78.6%		30-140			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C2-PFDOA	93.2%		10-150			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C2-PFTEDA	90.7%		10-130			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C3-PFBS	89.6%		55-150			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C3-PFHXS	88.1%		55-150			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C8-PFOS	83.8%		45-140			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C2-4:2FTS	178%		60-200			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C2-6:2FTS	80.7%		60-200			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C2-8:2FTS	70.5%		50-200			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C8-PFOA	72.7%		30-130			02/15/23	1	EPA 1633	BCB0232
Surrogate: D3-NMEFOA	54.7%		15-130			02/15/23	1	EPA 1633	BCB0232
Surrogate: D5-NETFOA	52.2%		10-130			02/15/23	1	EPA 1633	BCB0232
Surrogate: D3-NMEFOA	69.1%		45-200			02/15/23	1	EPA 1633	BCB0232
Surrogate: D5-NETFOA	69.7%		10-200			02/15/23	1	EPA 1633	BCB0232
Surrogate: D7-NMEFOE	45.2%		10-150			02/15/23	1	EPA 1633	BCB0232
Surrogate: D9-NETFOE	65.8%		10-150			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C3-HFPO-DA	89.8%		25-160			02/15/23	1	EPA 1633	BCB0232

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## Sample Results

(Continued)

**Sample: AF-RHMW225401-WGN01B-2302W1**  
**23B0091-04 (Water)**

### Per- and Polyfluoroalkyl Substances

Analyte	Result /Qual	LOQ	LOD	DL	Units	Date Analyzed	DF	Method	Prep Batch
PFBA	0.60 J	1.5	0.76	0.20	ng/L	02/15/23	1	EPA 1633	BCB0232
PFPEA	1.3	0.76	0.38	0.062	ng/L	02/15/23	1	EPA 1633	BCB0232
PFHXA	1.0	0.38	0.19	0.052	ng/L	02/15/23	1	EPA 1633	BCB0232
PFHPA	0.83	0.38	0.19	0.039	ng/L	02/15/23	1	EPA 1633	BCB0232
PFOA	1.0	0.38	0.19	0.14	ng/L	02/15/23	1	EPA 1633	BCB0232
PFNA	0.17 J IR1,	0.38	0.19	0.078	ng/L	02/15/23	1	EPA 1633	BCB0232
PFDA	0.19 U	0.38	0.19	0.096	ng/L	02/15/23	1	EPA 1633	BCB0232
PFUnA	0.19 U	0.38	0.19	0.15	ng/L	02/15/23	1	EPA 1633	BCB0232
PFDOA	0.19 U	0.38	0.19	0.11	ng/L	02/15/23	1	EPA 1633	BCB0232
PFTRDA	0.28 U	0.38	0.28	0.19	ng/L	02/15/23	1	EPA 1633	BCB0232
PFTEDA	0.19 U	0.38	0.19	0.19	ng/L	02/15/23	1	EPA 1633	BCB0232
PFBS	0.76	0.38	0.19	0.035	ng/L	02/15/23	1	EPA 1633	BCB0232
PFPEs	0.16 J	0.38	0.19	0.059	ng/L	02/15/23	1	EPA 1633	BCB0232
PFHXS	1.2	0.38	0.19	0.030	ng/L	02/15/23	1	EPA 1633	BCB0232
PFHPS	0.19 U	0.38	0.19	0.049	ng/L	02/15/23	1	EPA 1633	BCB0232
PFOS	1.1	0.38	0.19	0.060	ng/L	02/15/23	1	EPA 1633	BCB0232
PFNS	0.19 U	0.38	0.19	0.12	ng/L	02/15/23	1	EPA 1633	BCB0232
PFDS	0.19 U	0.38	0.19	0.14	ng/L	02/15/23	1	EPA 1633	BCB0232
PFDOS	0.19 U	0.38	0.19	0.12	ng/L	02/15/23	1	EPA 1633	BCB0232
4:2FTS	0.76 U	1.5	0.76	0.28	ng/L	02/15/23	1	EPA 1633	BCB0232
6:2FTS	0.76 U	1.5	0.76	0.30	ng/L	02/15/23	1	EPA 1633	BCB0232
8:2FTS	0.76 U	1.5	0.76	0.078	ng/L	02/15/23	1	EPA 1633	BCB0232
PFOSA	0.19 U	0.38	0.19	0.099	ng/L	02/15/23	1	EPA 1633	BCB0232
NMeFOSA	0.76 U	1.5	0.76	0.45	ng/L	02/15/23	1	EPA 1633	BCB0232
NEtFOSA	0.76 U	1.5	0.76	0.39	ng/L	02/15/23	1	EPA 1633	BCB0232
NMeFOSAA	0.19 U	0.38	0.19	0.10	ng/L	02/15/23	1	EPA 1633	BCB0232
NEtFOSAA	0.19 U	0.38	0.19	0.11	ng/L	02/15/23	1	EPA 1633	BCB0232
NMeFOSE	1.1 U	1.5	1.1	0.96	ng/L	02/15/23	1	EPA 1633	BCB0232
NEtFOSE	1.1 U	1.5	1.1	0.99	ng/L	02/15/23	1	EPA 1633	BCB0232
HFPO-DA	0.38 U	0.76	0.38	0.17	ng/L	02/15/23	1	EPA 1633	BCB0232
ADONA	0.38 U	0.76	0.38	0.12	ng/L	02/15/23	1	EPA 1633	BCB0232
PFEESA	0.38 U	0.76	0.38	0.10	ng/L	02/15/23	1	EPA 1633	BCB0232
PFMPA	0.38 U	0.76	0.38	0.051	ng/L	02/15/23	1	EPA 1633	BCB0232
PFMBA	0.38 U	0.76	0.38	0.086	ng/L	02/15/23	1	EPA 1633	BCB0232
NFDHA	0.38 U	0.76	0.38	0.29	ng/L	02/15/23	1	EPA 1633	BCB0232
9CL-PF3ONS	0.38 U	0.76	0.38	0.20	ng/L	02/15/23	1	EPA 1633	BCB0232
11CL-PF3OUDS	0.38 U	0.76	0.38	0.20	ng/L	02/15/23	1	EPA 1633	BCB0232
3:3FTCA	0.76 U	1.5	0.76	0.54	ng/L	02/15/23	1	EPA 1633	BCB0232
5:3FTCA	0.76 U	1.5	0.76	0.42	ng/L	02/15/23	1	EPA 1633	BCB0232
7:3FTCA	0.76 U	1.5	0.76	0.52	ng/L	02/15/23	1	EPA 1633	BCB0232
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Surrogate: 13C4-PFBA	84.7%		10-130			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C5-PFPEA	96.5%		35-150			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C5-PFHXA	91.5%		55-150			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C4-PFHPA	90.0%		55-150			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C8-PFOA	91.8%		60-140			02/15/23	1	EPA 1633	BCB0232



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Project Manager: Watson Tanji

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## Sample Results

(Continued)

**Sample: AF-RHMW225401-WGN01B-2302W1 (Continued)**

**23B0091-04 (Water)**

### Per- and Polyfluoroalkyl Substances (Continued)

Analyte	Result /Qual	LOQ	LOD	DL	Units	Date Analyzed	DF	Method	Prep Batch
Surrogate: 13C9-PFNA	94.9%		55-140			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C6-PFDA	87.1%		50-140			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C7-PFUnA	97.5%		30-140			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C2-PFDOA	95.2%		10-150			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C2-PFTEDA	82.0%		10-130			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C3-PFBS	96.3%		55-150			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C3-PFHXS	89.7%		55-150			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C8-PFOS	88.0%		45-140			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C2-4:2FTS	190%		60-200			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C2-6:2FTS	82.8%		60-200			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C2-8:2FTS	96.5%		50-200			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C8-PFOA	72.5%		30-130			02/15/23	1	EPA 1633	BCB0232
Surrogate: D3-NMEFOA	47.9%		15-130			02/15/23	1	EPA 1633	BCB0232
Surrogate: D5-NETFOA	47.7%		10-130			02/15/23	1	EPA 1633	BCB0232
Surrogate: D3-NMEFOA	90.3%		45-200			02/15/23	1	EPA 1633	BCB0232
Surrogate: D5-NETFOA	119%		10-200			02/15/23	1	EPA 1633	BCB0232
Surrogate: D7-NMEFOE	42.9%		10-150			02/15/23	1	EPA 1633	BCB0232
Surrogate: D9-NETFOE	59.4%		10-150			02/15/23	1	EPA 1633	BCB0232
Surrogate: 13C3-HFPO-DA	90.3%		25-160			02/15/23	1	EPA 1633	BCB0232

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## 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: BCB0232 - EPA 1633

#### Blank (BCB0232-BLK1)

Prepared: 02/13/23 07:19 Analyzed: 02/14/23 20:59

	ng/L			
PFBA	0.80 U	1.6	0.80	0.21
PFPEA	0.40 U	0.80	0.40	0.065
PFHXA	0.0870 J	0.40	0.20	0.055
PFHPA	0.20 U	0.40	0.20	0.041
PFOA	0.20 U IR2,	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	0.112 J MI5,	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 IR2,	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	25.7	32.0	80.4	10-130
13C5-PFPEA	13.4	16.0	83.5	35-150
13C5-PFHXA	5.93	8.00	74.1	55-150

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## Quality Control (Continued)

### Per- and Polyfluoroalkyl Substances (Continued)

Analyte	Result/Qual	LOQ	LOD	MDL	Spike Level	Source Result	%REC	%REC Limits	RPD	RPD Limit
<b>Blank (BCB0232-BLK1)</b>					Prepared: 02/13/23 07:19 Analyzed: 02/14/23 20:59					
	ng/L									
<b>Surrogates</b>										
13C4-PFHPA	6.25				8.00		78.1	55-150		
13C8-PFOA	5.97				8.00		74.7	60-140		
13C9-PFNA	2.73				4.00		68.3	55-140		
13C6-PFDA	3.01				4.00		75.3	50-140		
13C7-PFUnA	3.04				4.00		75.9	30-140		
13C2-PFDOA	3.62				4.00		90.6	10-150		
13C2-PFTEDA	3.29				4.00		82.4	10-130		
13C3-PFBS	5.99				8.00		74.9	55-150		
13C3-PFHXS	6.80				8.00		85.0	55-150		
13C8-PFOS	5.95				8.00		74.4	45-140		
13C2-4:2FTS	29.0				16.0		181	60-200		
13C2-6:2FTS	10.9				16.0		68.3	60-200		
13C2-8:2FTS	11.2				16.0		70.3	50-200		
13C8-PFOSA	6.05				8.00		75.7	30-130		
D3-NMEFOSA	4.00				8.00		50.0	15-130		
D5-NETFOSA	4.25				8.00		53.1	10-130		
D3-NMEFOSAA	10.6				16.0		66.2	45-200		
D5-NETFOSAA	10.5				16.0		65.7	10-200		
D7-NMEFOSE	38.7				80.0		48.3	10-150		
D9-NETFOSE	60.9				80.0		76.1	10-150		
13C3-HFPO-DA	26.7				32.0		83.4	25-160		

### LCS (BCB0232-BS1)

Prepared: 02/13/23 07:19 Analyzed: 02/14/23 21:12

	ng/L									
PFBA	16.7				16.0		105	58-148		
PFPEA	8.53				8.00		107	54-152		
PFHXA	4.36				4.00		109	55-152		
PFHPA	4.01				4.00		100	54-154		
PFOA	4.30				4.00		108	52-161		
PFNA	4.02				4.00		100	59-149		
PFDA	4.62				4.00		116	52-147		
PFUnA	4.11				4.00		103	48-159		
PFDOA	3.92				4.00		98.1	64-142		
PFTRDA	4.08				4.00		102	49-148		
PFTEDA	4.34				4.00		108	47-161		
PFBS	3.92				3.54		111	62-144		
PFPEs	3.94				3.76		105	59-151		
PFHXS	3.70				3.66		101	57-146		
PFHPS	4.10				3.82		107	55-152		
PFOS	3.80				3.72		102	58-149		
PFNS	4.06				3.84		106	52-148		
PFDS	3.98				3.86		103	51-147		
PFDOS	3.84				3.88		98.9	36-145		
4:2FTS	16.0				15.0		107	67-146		
6:2FTS	18.0				15.2		119	61-151		
8:2FTS	17.2				15.4		112	63-152		
PFOSA	4.63				4.00		116	61-148		

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

Reported: 02/16/2023 15:45

## Quality Control (Continued)

### Per- and Polyfluoroalkyl Substances (Continued)

Analyte	Result/Qual	LOQ	LOD	MDL	Spike Level	Source Result	%REC	%REC Limits	RPD	RPD Limit
<b>LCS (BCB0232-BS1)</b>					Prepared: 02/13/23 07:19 Analyzed: 02/14/23 21:12					
	ng/L									
NMeFOSA	16.1				16.0		101	63-145		
NEtFOSA	18.2				16.0		113	65-139		
NMeFOSAA	4.43				4.00		111	58-144		
NEtFOSAA	4.08				4.00		102	59-146		
NMeFOSE	18.5				16.0		116	71-136		
NEtFOSE	16.1				16.0		101	69-137		
HFPO-DA	8.21				8.00		103	63-144		
ADONA	7.95				7.56		105	68-146		
PFEESA	8.03				7.12		113	56-151		
PFMPA	8.03				8.00		100	51-145		
PFMBA	7.76				8.00		96.9	55-148		
NFDHA	8.99				8.00		112	48-161		
9CL-PF3ONS	7.63				7.48		102	56-156		
11CL-PF3OUDS	7.70				7.56		102	46-156		
3:3FTCA	16.9				16.0		106	62-129		
5:3FTCA	17.6				16.0		110	63-134		
7:3FTCA	16.0				16.0		99.8	50-138		
<b>Surrogates</b>										
13C4-PFBA	29.0				32.0		90.6	10-130		
13C5-PFPEA	14.8				16.0		92.4	35-150		
13C5-PFHXA	6.82				8.00		85.3	55-150		
13C4-PFHFA	7.15				8.00		89.4	55-150		
13C8-PFOA	7.12				8.00		89.1	60-140		
13C9-PFNA	3.43				4.00		85.8	55-140		
13C6-PFDA	3.17				4.00		79.2	50-140		
13C7-PFUnA	3.34				4.00		83.4	30-140		
13C2-PFDOA	3.35				4.00		83.8	10-150		
13C2-PFTEDA	3.04				4.00		76.1	10-130		
13C3-PFBS	6.76				8.00		84.5	55-150		
13C3-PFHXS	6.76				8.00		84.5	55-150		
13C8-PFOS	7.12				8.00		89.0	45-140		
13C2-4:2FTS	29.1				16.0		182	60-200		
13C2-6:2FTS	11.7				16.0		72.8	60-200		
13C2-8:2FTS	11.3				16.0		70.8	50-200		
13C8-PFOSA	6.81				8.00		85.1	30-130		
D3-NMEFOSA	4.68				8.00		58.6	15-130		
D5-NETFOSA	4.92				8.00		61.5	10-130		
D3-NMEFOSAA	12.1				16.0		75.6	45-200		
D5-NETFOSAA	12.4				16.0		77.8	10-200		
D7-NMEFOSE	41.7				80.0		52.1	10-150		
D9-NETFOSE	65.5				80.0		81.8	10-150		
13C3-HFPO-DA	29.0				32.0		90.8	25-160		

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: 02/16/2023 15:45

## Quality Control (Continued)

### Per- and Polyfluoroalkyl Substances (Continued)

Analyte	Result/Qual	LOQ	LOD	MDL	Spike Level	Source Result	%REC	%REC Limits	RPD	RPD Limit
<b>MRL Check (BCB0232-MRL1)</b>					Prepared: 02/13/23 07:19 Analyzed: 02/14/23 21:25					
	ng/L									
PFBA	1.89				1.60		118	44-157		
PFPEA	1.01				0.800		126	57-148		
PFHXA	0.544				0.400		136	62-149		
PFHPA	0.599				0.400		150	56-150		
PFOA	0.515 IR2				0.400		129	57-161		
PFNA	0.510				0.400		128	53-157		
PFDA	0.446				0.400		112	43-158		
PFUnA	0.495				0.400		124	50-155		
PFDOA	0.475 IR2				0.400		119	60-141		
PFTRDA	0.498				0.400		124	52-140		
PFTEDA	0.546				0.400		137	52-156		
PFBS	0.385 J				0.354		109	63-145		
PFPEs	0.477				0.376		127	58-144		
PFHXS	0.505				0.366		138	44-158		
PFHPS	0.419				0.382		110	51-150		
PFOS	0.526				0.372		141	43-162		
PFNS	0.430				0.384		112	46-151		
PFDS	0.455				0.386		118	50-144		
PFDOS	0.418				0.388		108	30-138		
4:2FTS	1.90				1.50		127	52-158		
6:2FTS	1.84				1.52		121	48-158		
8:2FTS	2.25				1.54		147	46-165		
PFOSA	0.542				0.400		136	47-163		
NMeFOSA	2.01				1.60		126	54-155		
NEtFOSA	1.89				1.60		118	49-156		
NMeFOSAA	0.495				0.400		124	32-160		
NEtFOSAA	0.488				0.400		122	51-154		
NMeFOSE	1.97				1.60		123	56-151		
NEtFOSE	1.77				1.60		111	60-147		
HFPO-DA	0.940				0.800		118	58-154		
ADONA	0.915				0.756		121	61-148		
PFEESA	0.810				0.712		114	56-144		
PFMPA	0.996				0.800		125	48-150		
PFMBA	0.903				0.800		113	49-154		
NFDHA	0.858				0.800		107	47-160		
9CL-PF3ONS	0.966				0.748		129	44-167		
11CL-PF3OUDS	0.935				0.756		124	36-158		
3:3FTCA	1.90				1.60		119	32-161		
5:3FTCA	1.66				1.60		104	39-156		
7:3FTCA	1.48 J				1.60		92.8	36-149		
<b>Surrogates</b>										
13C4-PFBA	28.6				32.0		89.4	10-130		
13C5-PFPEA	15.6				16.0		97.7	35-150		
13C5-PFHXA	7.70				8.00		96.3	55-150		
13C4-PFHPA	7.26				8.00		90.8	55-150		
13C8-PFOA	6.72				8.00		84.0	60-140		
13C9-PFNA	3.50				4.00		87.4	55-140		

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

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

Reported: 02/16/2023 15:45

## Quality Control (Continued)

### Per- and Polyfluoroalkyl Substances (Continued)

Analyte	Result/Qual	LOQ	LOD	MDL	Spike Level	Source Result	%REC	%REC Limits	RPD	RPD Limit
<b>MRL Check (BCB0232-MRL1)</b>					Prepared: 02/13/23 07:19 Analyzed: 02/14/23 21:25					
	ng/L									
<b>Surrogates</b>										
13C6-PFDA	3.59				4.00		89.8	50-140		
13C7-PFUnA	3.51				4.00		87.7	30-140		
13C2-PFDOA	3.47				4.00		86.8	10-150		
13C2-PFTEDA	3.52				4.00		88.1	10-130		
13C3-PFBS	6.83				8.00		85.4	55-150		
13C3-PFHXS	6.57				8.00		82.2	55-150		
13C8-PFOS	7.35				8.00		91.8	45-140		
13C2-4:2FTS	28.4				16.0		177	60-200		
13C2-6:2FTS	11.2				16.0		70.0	60-200		
13C2-8:2FTS	11.2				16.0		70.1	50-200		
13C8-PFOSA	7.12				8.00		89.0	30-130		
D3-NMEFOSA	5.07				8.00		63.4	15-130		
D5-NETFOSA	5.48				8.00		68.5	10-130		
D3-NMEFOSAA	13.4				16.0		83.5	45-200		
D5-NETFOSAA	14.1				16.0		88.3	10-200		
D7-NMEFOSE	43.3				80.0		54.1	10-150		
D9-NETFOSE	68.4				80.0		85.5	10-150		
13C3-HFPO-DA	30.9				32.0		96.5	25-160		

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: 02/16/2023 15:45

## 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
J	Estimated value
MI2	Manual integration, non-target peak interference
MI5	Manual integration, whole peak was not integrated
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****23B0091**

Printed: 02/16/2023 3:45 pm

**Project:** Red Hill AFFF Assessment Sampling  
**Project Number:** Red Hill AFFF Assessment Sampling  
**Project Manager:** Gregory Salata  
**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: 02/11/2023 11:05 AM  
 Date Due: 02/20/2023 (5.00 day TAT)

Logged In By: Megan Horne  
 Received By: Megan Horne

Analysis	Comments
<b>23B0091-01 AF-RHMW17-WGN01LF-2302W1 [Water] Sampled 2/10/2023 11:00:00AM</b>	
1633 NONE	"Report relevant surrogates"
<b>23B0091-02 AF-RHMW17D-WGN01LF-2302W1 [Water] Sampled 2/10/2023 12:15:00PM</b>	
1633 NONE	"Report relevant surrogates"
<b>23B0091-03 AF-RHMW17D-WQFB01-2302W1 [Water] Sampled 2/10/2023 11:30:00AM</b>	
1633 NONE	"Report relevant surrogates"
<b>23B0091-04 AF-RHMW225401-WGN01B-2302W1 [Water] Sampled 2/10/2023 8:55:00AM</b>	
1633 NONE	"Report relevant surrogates"

**23B0091****Sample Receipt Log**

Default Cooler

Samples Received at: **1.0°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

coc@appliance.com C.O.C. 2302W1AFAL10

Report to: PLEASE PRINT

AECOM

Phone: \_\_\_\_\_

Address: 1001 Bishop St ste1600

Honolulu, HI 96813

Watson Tanji / Katie Abbott

Email: [watson.tanji@aecom.com](mailto:watson.tanji@aecom.com)/[katie.abbott@aecom.com](mailto:katie.abbott@aecom.com)

Invoice to: PLEASE PRINT

Company Name: **AECOM**

Phone: \_\_\_\_\_

Address:

Fax: \_\_\_\_\_

Sheree Smith

E-mail: [USAPimaging@aecom.com](mailto:USAPimaging@aecom.com)

[illegible]

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

PLEASE PRINT

808-954-4512 / 303-796-4624

Phone: \_\_\_\_\_

Address: 1001 Bishop St ste1600

Honolulu, HI 96813

Watson Tanji / Katie Abbott

Email: [watson.tanji@aecom.com/katie.abbott@aecom.com](mailto:watson.tanji@aecom.com/katie.abbott@aecom.com)

**Invoice to:**

Company Name: **AECOM**

Phone:

Address:

Fax: \_\_\_\_\_

Sheree Smith

Attn: **Sheree Smith**

Email: [USAPimaging@aecom.com](mailto:USAPimaging@aecom.com)

[illegible]

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

**ELECTRONIC CHAIN OF CUSTODY RECORD**  
Phone: (559) 275-2175  
Fax: (559) 275-4422  
coc@applinc.com C.O.C. 2302W1AFAL013

[illegible]

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

**CUSTODY SEAL**  
AECOM (808)-521-3051  
Initials: JMM Date 2/10/23

PFAS

# SAMPLE DATA

# FORM I

## ANALYSIS DATA SHEET

### AF-RHMW17-WGN01LF-2302W1

Laboratory:	APPL, LLC	Work Order:	23B0091
Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Matrix:	Water	Laboratory ID:	23B0091-01
		File ID:	S2023-02-14D (20)
Sampled:	02/10/23 11:00	Prepared:	02/13/23 07:19
		Analyzed:	02/14/23 23:47
Solids:		Preparation:	EPA 1633
		Dilution:	1
Initial/Final:	566.32 mL / 2 mL	Instrument:	Saphira
Batch:	BCB0232	Sequence:	SC00636
		Calibration:	2307007

COMPOUND	CONC. (ng/L)	LOQ	LOD	DL	Q
PFBA	2.1	1.4	0.71	0.18	
PFPEA	5.3	0.71	0.35	0.057	
PFHXA	2.5	0.35	0.18	0.048	
PFHPA	0.44	0.35	0.18	0.036	
PFOA	0.27 J	0.35	0.18	0.13	
PFNA	0.18 U	0.35	0.18	0.072	
PFDA	0.18 U	0.35	0.18	0.089	
PFUnA	0.18 U	0.35	0.18	0.14	
PFDOA	0.18 U	0.35	0.18	0.099	
PFTRDA	0.26 U	0.35	0.26	0.18	
PFTEDA	0.18 U	0.35	0.18	0.17	IR2,
PFBS	0.22 J	0.35	0.18	0.032	
PFPEs	0.18 U	0.35	0.18	0.055	
PFHXS	0.18 U	0.35	0.18	0.028	
PFHPS	0.18 U	0.35	0.18	0.045	
PFOS	0.19 J	0.35	0.18	0.056	MI5,
PFNS	0.18 U	0.35	0.18	0.11	
PFDS	0.18 U	0.35	0.18	0.13	
PFDOS	0.18 U	0.35	0.18	0.11	
4:2FTS	0.71 U	1.4	0.71	0.26	
6:2FTS	0.49 J	1.4	0.71	0.28	
8:2FTS	0.71 U	1.4	0.71	0.073	
PFOSA	0.18 U	0.35	0.18	0.092	
NMeFOSA	0.71 U	1.4	0.71	0.42	
NEtFOSA	0.71 U	1.4	0.71	0.36	
NMeFOSAA	0.18 U	0.35	0.18	0.093	
NEtFOSAA	0.18 U	0.35	0.18	0.10	
NMeFOSE	1.1 U	1.4	1.1	0.89	
NEtFOSE	1.1 U	1.4	1.1	0.92	
HFPO-DA	0.35 U	0.71	0.35	0.15	

# FORM I

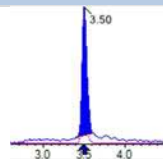
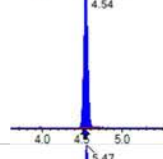
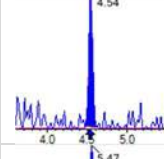
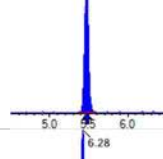
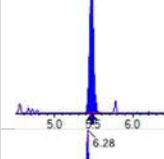
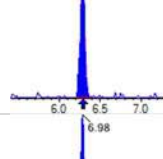
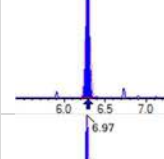
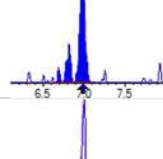
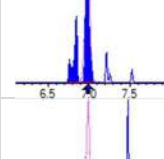
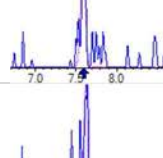
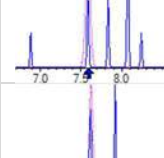
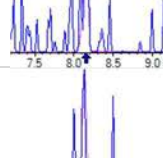
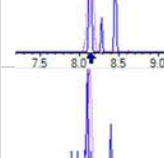
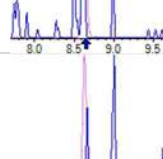
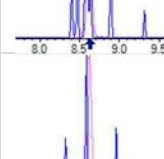
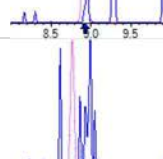
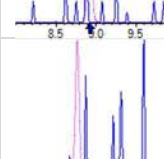
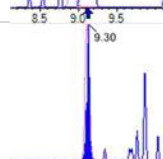
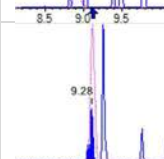
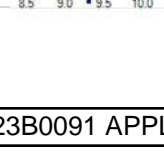
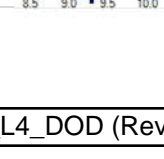
## ANALYSIS DATA SHEET

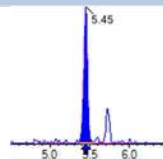
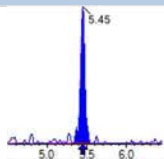
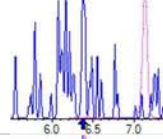
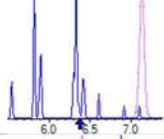
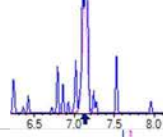
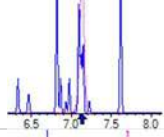
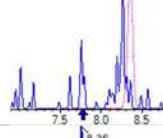
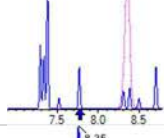
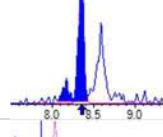
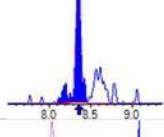
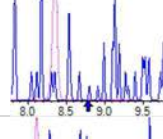
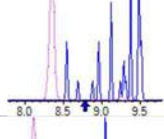
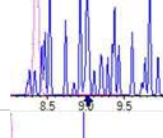
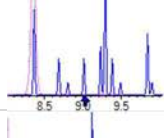
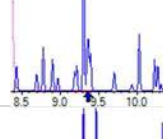
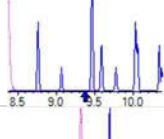
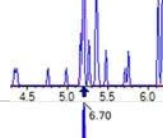
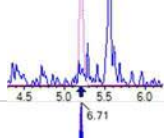
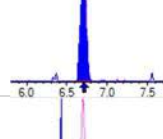
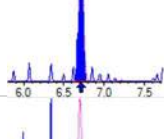
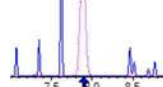
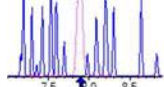
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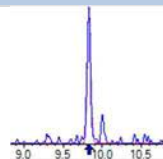
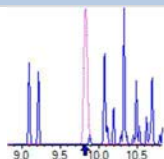
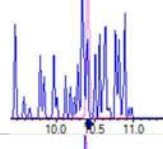
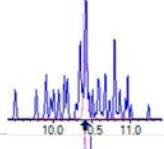
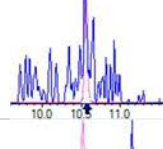
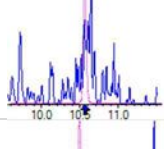
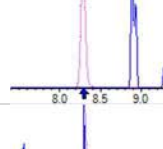
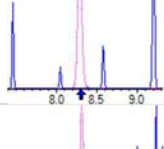
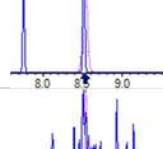
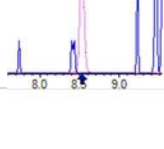
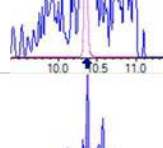
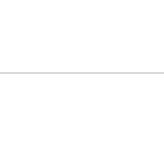
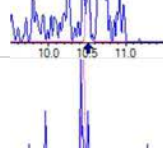
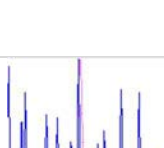
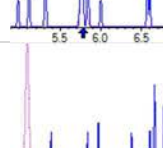
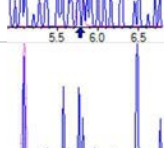
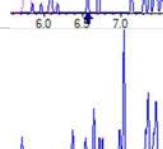
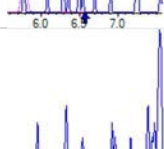
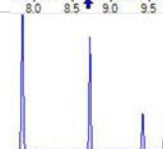
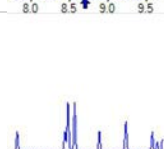
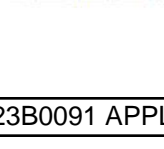
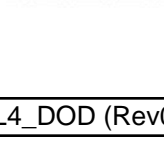
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Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Matrix:	Water	Laboratory ID:	23B0091-01
		File ID:	S2023-02-14D (20)
Sampled:	02/10/23 11:00	Prepared:	02/13/23 07:19
		Analyzed:	02/14/23 23:47
Solids:		Preparation:	EPA 1633
		Dilution:	1
Initial/Final:	566.32 mL / 2 mL	Instrument:	Saphira
Batch:	BCB0232	Sequence:	SC00636
		Calibration:	2307007

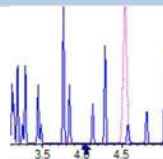
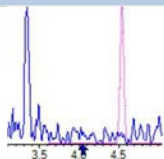
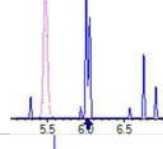
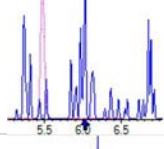
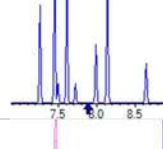
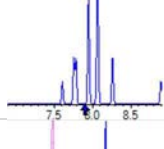
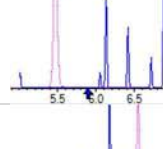
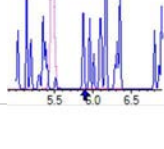
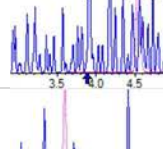
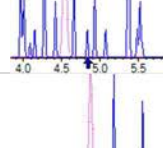
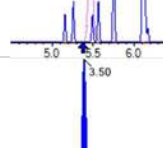
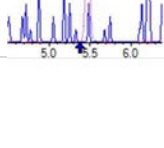
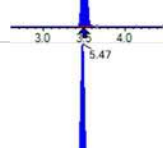
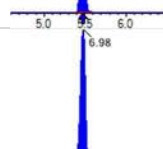
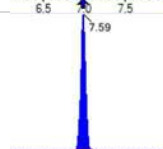

COMPOUND	CONC. (ng/L)	LOQ	LOD	DL	Q
ADONA	0.35 U	0.71	0.35	0.11	
PFEESA	0.35 U	0.71	0.35	0.096	
PFMPA	0.35 U	0.71	0.35	0.048	
PFMBA	0.35 U	0.71	0.35	0.080	
NFDHA	0.35 U	0.71	0.35	0.27	
9CL-PF3ONS	0.35 U	0.71	0.35	0.18	
11CL-PF3OUDS	0.35 U	0.71	0.35	0.18	
3:3FTCA	0.71 U	1.4	0.71	0.51	
5:3FTCA	0.71 U	1.4	0.71	0.39	
7:3FTCA	0.71 U	1.4	0.71	0.49	

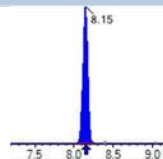
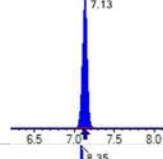
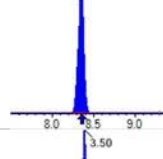
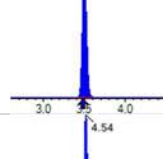
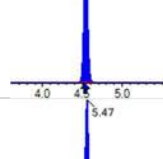
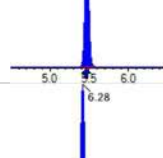
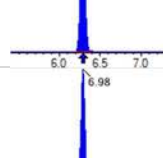
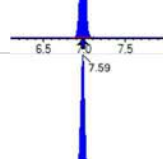
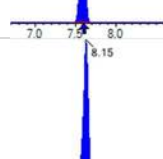
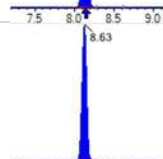
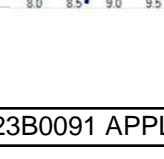


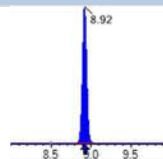
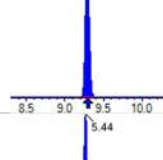
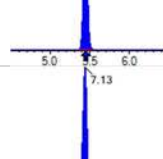
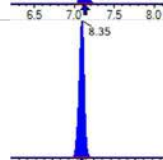
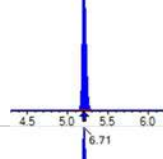
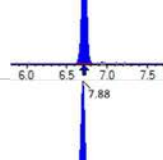
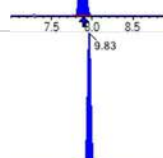
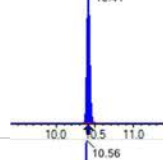
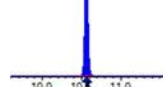
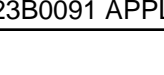
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	( 213.0 / 169.0 ) 150786	( 3.50 , 1.00 ) ( 0.00 , N/A , 0.0 )	95.0	N/A 0.0 0.0	0.6076	N/A			
PFPeA	( 263.0 / 219.0 ) 745047 ( 263.0 / 69.0 ) 7609	( 4.54 , 1.00 ) ( 0.00 , N/A , 0.1 )	1284.7 30.5	0.0102 95.4 92.7	1.4904	N/A			
PFHxA	( 313.0 / 269.0 ) 457210 ( 313.0 / 119.0 ) 47304	( 5.47 , 1.00 ) ( 0.00 , N/A , 0.0 )	1020.2 894.3	0.1035 103.9 108.7	0.7213	N/A			
PFHpA	( 363.0 / 319.0 ) 62846 ( 363.0 / 169.0 ) 23932	( 6.28 , 1.00 ) ( 0.00 , N/A , 0.1 )	1557.0 66890.5	0.3808 118.8 119.1	0.1258	N/A			
PFOA	( 413.0 / 369.0 ) 49309 ( 413.0 / 169.0 ) 18316	( 6.98 , 1.00 ) ( 0.00 , N/A , 0.4 )	2174.5 24169.9	0.3714 111.0 102.8	0.0777	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 ) 6601 ( 713.0 / 169.0 ) 2141	( 9.30 , 1.00 ) ( 0.02 , N/A , 1.7 )	92.1 471.6	0.3244 157.0 156.9	0.0139	N/A			IR2,

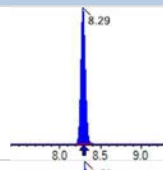
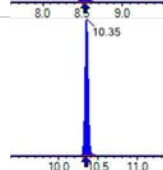
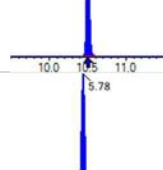


Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	( 299.0 / 80.0 ) 76818 ( 299.0 / 99.0 ) 51689	( 5.45 , 1.00 ) ( 0.00 , N/A , 0.0 )	141.6 109.9	0.6729 104.1 108.8	0.0635	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 ) 123559 ( 499.0 / 99.0 ) 38788	( 8.36 , 1.00 ) ( 0.00 , N/A , 0.2 )	76.6 117.1	0.3139 141.8 148.1	0.0546	N/A			MI5 DG 2023-02-15
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 ) 18411 ( 427.0 / 81.0 ) 17663	( 6.70 , 1.00 ) ( -0.01 , N/A , -0.5 )	7993.4 86.1	0.9594 115.9 124.2	0.1377	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 ) N/A	( N/A , N/A ) ( N/A , N/A , N/A )	N/A	N/A 0.0 0.0	0.0000	N/A			
NEiFOSE	( 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 ) 303297	( 3.50 , N/A ) ( N/A , 0.00 , N/A )	1301.7	N/A	1.3541 [ 1.0000 ]	135.4% { 138.2% }			
13C2_PFHxA_IIS	( 315.0 / 270.0 ) 610047	( 5.47 , N/A ) ( N/A , 0.00 , N/A )	2443.8	N/A	1.2085 [ 1.0000 ]	120.8% { 126.9% }			
13C4_PFOA_IIS	( 417.0 / 372.0 ) 761645	( 6.98 , N/A ) ( N/A , -0.01 , N/A )	3883.7	N/A	1.2018 [ 1.0000 ]	120.2% { 124.8% }			
13C5_PFNA_IIS	( 468.0 / 423.0 ) 729264	( 7.59 , N/A ) ( N/A , 0.00 , N/A )	2463.1	N/A	1.2069 [ 1.0000 ]	120.7% { 129.3% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) ( $\Delta$ RT-[min], $\Delta$ RT- CV[min], $\Delta$ 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 ) 693382	( 8.15, N/A ) ( N/A, 0.00, N/A )	1623.2	N/A	1.1983 [ 1.0000 ]	119.8% { 129.6% }			
18O2_PFHxS_IIS	( 403.0 / 83.9 ) 1094985	( 7.13, N/A ) ( N/A, 0.00, N/A )	4080.0	N/A	1.2786 [ 1.0000 ]	127.9% { 130.0% }			
13C4_PFOS_IIS	( 503.0 / 79.9 ) 1850620	( 8.35, N/A ) ( N/A, 0.00, N/A )	1594.3	N/A	1.2293 [ 1.0000 ]	122.9% { 138.0% }			
13C4_PFBA_EIS	( 217.0 / 172.0 ) 2396744	( 3.50, N/A ) ( N/A, 0.00, N/A )	6379.5	N/A	6.6434 [ 8.0000 ]	83.0% { 104.3% }			
13C5_PFPeA_EIS	( 268.0 / 223.0 ) 2430362	( 4.54, N/A ) ( N/A, 0.00, N/A )	4070.7	N/A	3.6411 [ 4.0000 ]	91.0% { 105.4% }			
13C5_PFHxA_EIS	( 318.0 / 273.0 ) 1442085	( 5.47, N/A ) ( N/A, 0.00, N/A )	2966.7	N/A	1.7258 [ 2.0000 ]	86.3% { 101.0% }			
13C4_PFHpA_EIS	( 367.0 / 322.0 ) 1183417	( 6.28, N/A ) ( N/A, -0.01, N/A )	4310.9	N/A	1.6346 [ 2.0000 ]	81.7% { 100.4% }			
13C8_PFOA_EIS	( 421.0 / 376.0 ) 1426121	( 6.98, N/A ) ( N/A, 0.00, N/A )	2526.8	N/A	1.6533 [ 2.0000 ]	82.7% { 100.4% }			
13C9_PFNA_EIS	( 472.0 / 427.0 ) 631651	( 7.59, N/A ) ( N/A, 0.00, N/A )	1600.2	N/A	0.8175 [ 1.0000 ]	81.7% { 103.8% }			
13C6_PFDA_EIS	( 519.0 / 474.0 ) 737180	( 8.15, N/A ) ( N/A, 0.00, N/A )	1082.7	N/A	0.8432 [ 1.0000 ]	84.3% { 98.9% }			
13C7_PFUnA_EIS	( 570.0 / 525.0 ) 753656	( 8.63, N/A ) ( N/A, 0.00, N/A )	1256.7	N/A	0.8669 [ 1.0000 ]	86.7% { 113.1% }			

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 ) 679973	( 8.92 , N/A ) ( N/A , 0.00 , N/A )	3742.5	N/A	0.9120 [ 1.0000 ]	91.2% { 103.3% }			
13C2_PFTeDA_EIS	( 715.0 / 670.0 ) 548053	( 9.29 , N/A ) ( N/A , 0.00 , N/A )	1562.3	N/A	0.8022 [ 1.0000 ]	80.2% { 93.6% }			
13C3_PFBs_EIS	( 302.0 / 80.0 ) 3893288	( 5.44 , N/A ) ( N/A , 0.00 , N/A )	3915.1	N/A	1.6989 [ 2.0000 ]	84.9% { 102.7% }			
13C3_PFHxS_EIS	( 402.0 / 80.0 ) 1973009	( 7.13 , N/A ) ( N/A , 0.00 , N/A )	1864.8	N/A	1.6575 [ 2.0000 ]	82.9% { 100.4% }			
13C8_PFOS_EIS	( 507.0 / 80.0 ) 4291630	( 8.35 , N/A ) ( N/A , 0.00 , N/A )	1641.6	N/A	1.6892 [ 2.0000 ]	84.5% { 101.3% }			
13C2_4:2FTS_EIS	( 329.0 / 81.0 ) 919440	( 5.20 , N/A ) ( N/A , 0.00 , N/A )	1800.1	N/A	7.2685 [ 4.0000 ]	181.7% { 264.8% }			S2,
13C2_6:2FTS_EIS	( 429.0 / 81.0 ) 376828	( 6.71 , N/A ) ( N/A , -0.01 , N/A )	609.4	N/A	2.7138 [ 4.0000 ]	67.8% { 90.6% }			
13C2_8:2FTS_EIS	( 529.0 / 81.0 ) 542384	( 7.88 , N/A ) ( N/A , -0.01 , N/A )	1232.4	N/A	2.7648 [ 4.0000 ]	69.1% { 89.9% }			
13C8_PFOSA_EIS	( 506.0 / 78.0 ) 7336212	( 9.83 , N/A ) ( N/A , 0.00 , N/A )	3915.5	N/A	1.5556 [ 2.0000 ]	77.8% { 88.3% }			
D3_NMeFOSA_EIS	( 515.0 / 169.0 ) 1413226	( 10.41 , N/A ) ( N/A , 0.00 , N/A )	2501.3	N/A	1.2719 [ 2.0000 ]	63.6% { 73.0% }			
D5_NEtFOSA_EIS	( 531.0 / 169.0 ) 1190414	( 10.56 , N/A ) ( N/A , 0.00 , N/A )	2565.2	N/A	1.1973 [ 2.0000 ]	59.9% { 66.7% }			

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 ) 1198772	( 8.29 , N/A ) ( N/A , 0.00 , N/A )	2133.4	N/A	2.9157 [ 4.0000 ]	72.9% { 90.5% }			
D5_EtFOSAA_EIS	( 589.0 / 419.0 ) 1096548	( 8.53 , N/A ) ( N/A , 0.00 , N/A )	13503.9	N/A	3.2572 [ 4.0000 ]	81.4% { 103.9% }			
D7_NMeFOSE_EIS	( 623.0 / 58.9 ) 4264171	( 10.35 , N/A ) ( N/A , 0.00 , N/A )	1804.9	N/A	10.5268 [ 20.0000 ]	52.6% { 57.4% }			
D9_NEtFOSE_EIS	( 639.0 / 58.9 ) 7345636	( 10.50 , N/A ) ( N/A , 0.00 , N/A )	2155.8	N/A	14.1601 [ 20.0000 ]	70.8% { 78.0% }			
13C3_HFPODA_EIS	( 287.0 / 169.0 ) 3668778	( 5.78 , N/A ) ( N/A , 0.00 , N/A )	3341.0	N/A	7.2110 [ 8.0000 ]	90.1% { 102.3% }			

# FORM I

## ANALYSIS DATA SHEET

AF-RHMW17D-WGN01LF-2302W1

Laboratory:	APPL, LLC	Work Order:	23B0091
Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Matrix:	Water	Laboratory ID:	23B0091-02
		File ID:	S2023-02-14D (22)
Sampled:	02/10/23 12:15	Prepared:	02/13/23 07:19
		Analyzed:	02/15/23 00:13
Solids:		Preparation:	EPA 1633
		Dilution:	1
Initial/Final:	569.47 mL / 2 mL	Instrument:	Saphira
Batch:	BCB0232	Sequence:	SC00636
		Calibration:	2307007

COMPOUND	CONC. (ng/L)	LOQ	LOD	DL	Q
PFBA	0.60 J	1.4	0.70	0.18	
PFPEA	0.13 J	0.70	0.35	0.057	
PFHXA	0.27 J	0.35	0.18	0.048	IR2,
PFHPA	0.045 J	0.35	0.18	0.036	IR2,
PFOA	0.18 U	0.35	0.18	0.13	
PFNA	0.18 U	0.35	0.18	0.072	
PFDA	0.18 U	0.35	0.18	0.089	
PFUnA	0.18 U	0.35	0.18	0.14	
PFDOA	0.18 U	0.35	0.18	0.098	
PFTRDA	0.26 U	0.35	0.26	0.18	
PFTEDA	0.18 U	0.35	0.18	0.17	
PFBS	0.081 J	0.35	0.18	0.032	
PFPEs	0.18 U	0.35	0.18	0.055	
PFHXS	0.052 J	0.35	0.18	0.028	
PFHPS	0.18 U	0.35	0.18	0.045	
PFOS	0.12 J	0.35	0.18	0.056	IR2,
PFNS	0.17 J	0.35	0.18	0.11	IR1,
PFDS	0.18 U	0.35	0.18	0.13	
PFDOS	0.18 U	0.35	0.18	0.11	
4:2FTS	0.70 U	1.4	0.70	0.26	
6:2FTS	6.1	1.4	0.70	0.28	
8:2FTS	0.70 U	1.4	0.70	0.072	
PFOSA	0.18 U	0.35	0.18	0.091	
NMeFOSA	0.70 U	1.4	0.70	0.42	
NEtFOSA	0.70 U	1.4	0.70	0.36	
NMeFOSAA	0.18 U	0.35	0.18	0.093	
NEtFOSAA	0.18 U	0.35	0.18	0.10	
NMeFOSE	1.1 U	1.4	1.1	0.89	
NEtFOSE	1.1 U	1.4	1.1	0.92	
HFPO-DA	0.35 U	0.70	0.35	0.15	



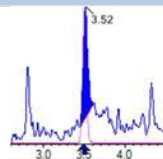
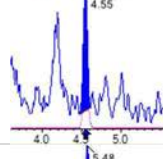
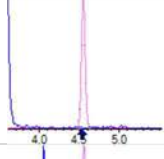
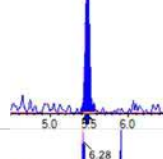
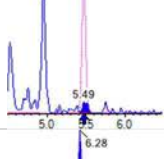
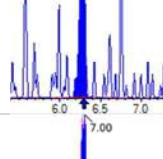
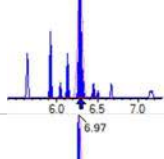
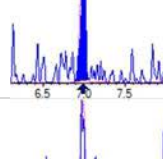
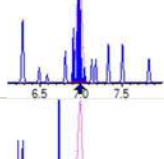
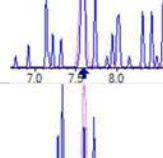
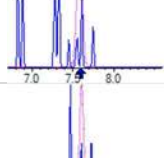
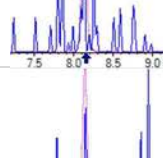
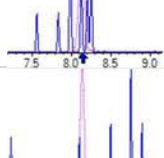
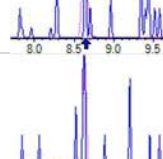
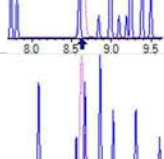
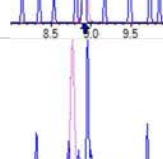
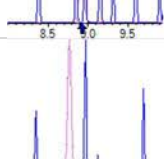
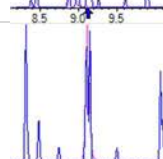
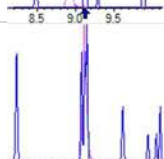
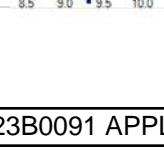
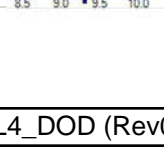
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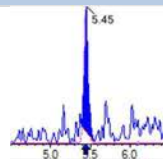
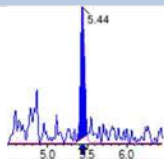
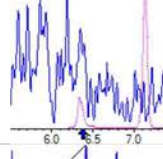
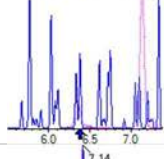
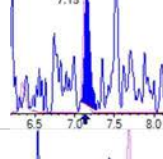
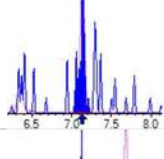
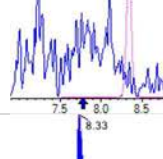
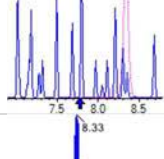
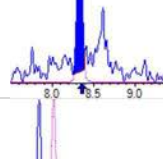
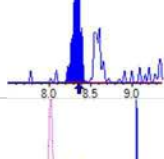
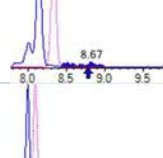
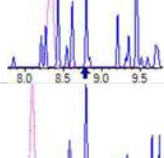
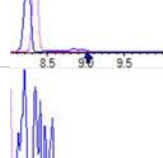
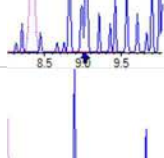
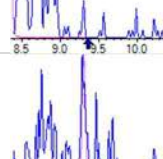
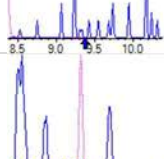
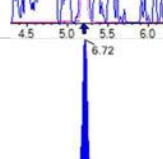
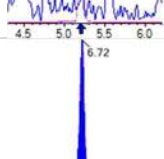
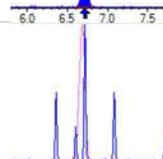
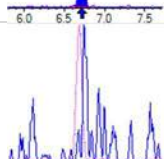
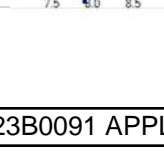
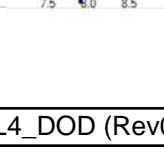
## ANALYSIS DATA SHEET

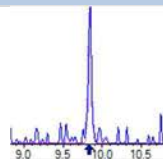
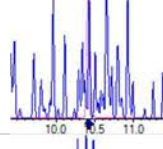
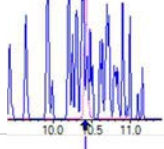
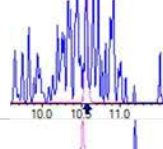
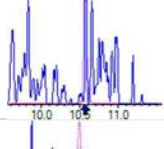
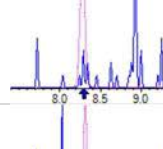
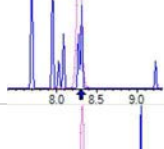
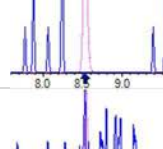
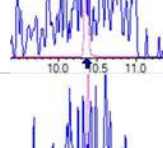
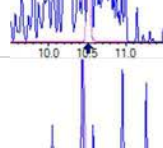
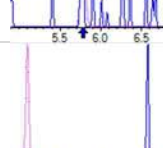
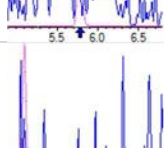
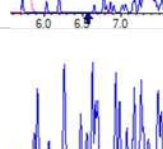
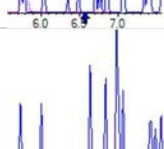
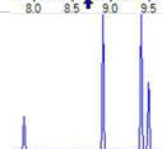
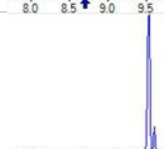
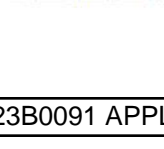
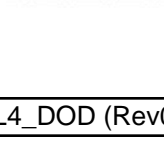
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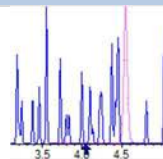
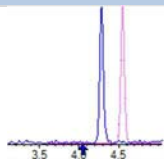
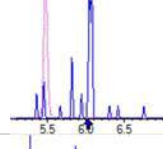
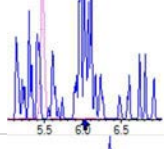
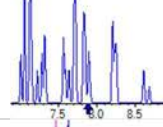
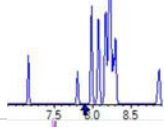
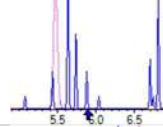
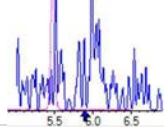
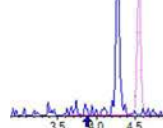
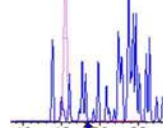
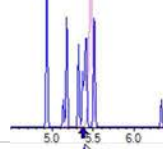
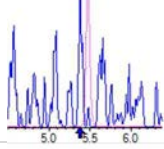
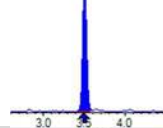
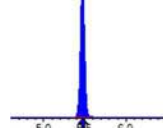
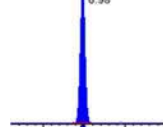
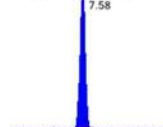
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Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Matrix:	Water	Laboratory ID:	23B0091-02
		File ID:	S2023-02-14D (22)
Sampled:	02/10/23 12:15	Prepared:	02/13/23 07:19
		Analyzed:	02/15/23 00:13
Solids:		Preparation:	EPA 1633
		Dilution:	1
Initial/Final:	569.47 mL / 2 mL	Instrument:	Saphira
Batch:	BCB0232	Sequence:	SC00636
		Calibration:	2307007

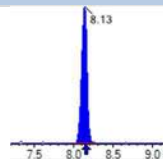
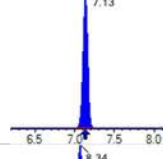
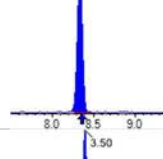
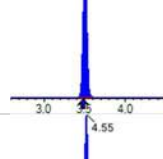
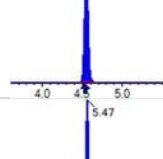
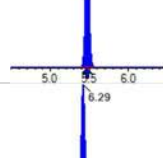
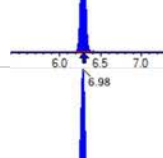
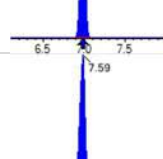
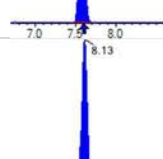
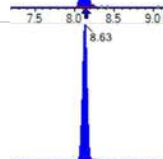
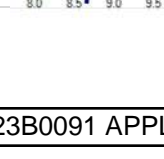
COMPOUND	CONC. (ng/L)	LOQ	LOD	DL	Q
ADONA	0.35 U	0.70	0.35	0.11	
PFEESA	0.35 U	0.70	0.35	0.096	
PFMPA	0.35 U	0.70	0.35	0.047	
PFMBA	0.35 U	0.70	0.35	0.080	
NFDHA	0.35 U	0.70	0.35	0.26	
9CL-PF3ONS	0.35 U	0.70	0.35	0.18	
11CL-PF3OUDS	0.35 U	0.70	0.35	0.18	
3:3FTCA	0.70 U	1.4	0.70	0.50	
5:3FTCA	0.70 U	1.4	0.70	0.39	
7:3FTCA	0.70 U	1.4	0.70	0.49	

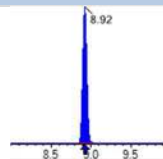
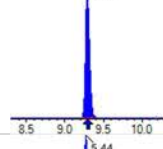
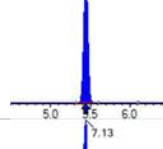
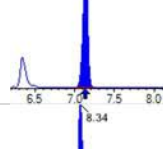
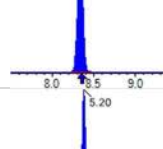
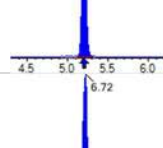
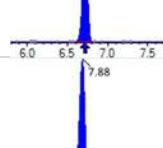
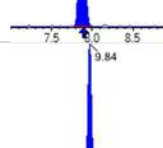
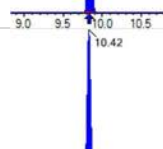
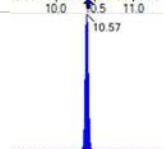

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	( 213.0 / 169.0 ) 24025	( 3.52, 1.00 ) ( 0.01, N/A, 0.0 )	22.8	N/A 0.0 0.0	0.1698	N/A			
PFPeA	( 263.0 / 219.0 ) 13917 ( 263.0 / 69.0 ) N/A	( 4.55, 1.00 ) ( 0.00, N/A, #Value! )	24.8 N/A	N/A 0.0 0.0	0.0375	N/A			
PFHxA	( 313.0 / 269.0 ) 45112 ( 313.0 / 119.0 ) 7149	( 5.48, 1.00 ) ( 0.01, N/A, -0.5 )	112.9 10.6	0.1585 159.1 166.5	0.0760	N/A			IR2,
PFHpA	( 363.0 / 319.0 ) 6218 ( 363.0 / 169.0 ) 7655	( 6.28, 1.00 ) ( -0.01, N/A, -0.1 )	37.0 3318878.0	1.2311 384.0 385.0	0.0128	N/A			IR2,
PFOA	( 413.0 / 369.0 ) 16617 ( 413.0 / 169.0 ) 6684	( 7.00, 1.00 ) ( 0.01, N/A, 1.5 )	41.8 7370.3	0.4022 120.2 111.3	0.0238	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 ) 30239 ( 299.0 / 99.0 ) 24898	( 5.45 , 1.00 ) ( 0.01 , N/A , 0.7 )	26.4 29.3	0.8234 127.3 133.1	0.0231	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 ) 22596 ( 399.0 / 99.0 ) 8888	( 7.15 , 1.00 ) ( 0.02 , N/A , 0.8 )	18.9 7014.0	0.3933 121.1 120.7	0.0148	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 ) 73474 ( 499.0 / 99.0 ) 25441	( 8.33 , 1.00 ) ( 0.00 , N/A , -0.1 )	35.8 73.7	0.3463 156.5 163.3	0.0333	N/A			IR2,
PFNS	( 549.0 / 80.0 ) 118235 ( 549.0 / 99.0 ) N/A	( 8.67 , 1.04 ) ( N/A , -0.11 , #Value! )	20.6 N/A	N/A 0.0 0.0	0.0476	N/A			IR1,
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 ) 642678 ( 427.0 / 81.0 ) 494592	( 6.72 , 1.00 ) ( 0.00 , N/A , -0.1 )	1308.5 642.2	0.7696 93.0 99.6	1.7432	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			

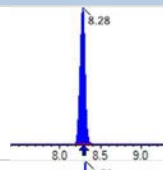
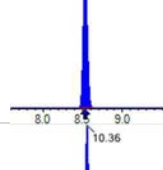
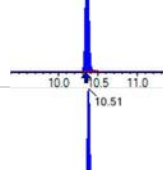
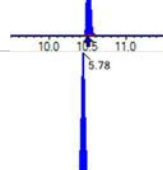

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	( 498.0 / 78.0 ) N/A ( 498.0 / 478.0 ) N/A	( N/A , N/A ) ( N/A , 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 ) N/A	( N/A , N/A ) ( N/A , N/A , N/A )	N/A	N/A 0.0 0.0	0.0000	N/A			
NEiFOSE	( 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 ) 175515	( 3.50 , N/A ) ( N/A , 0.00 , N/A )	670.4	N/A	0.7836 [ 1.0000 ]	78.4% { 80.0% }			
13C2_PFHxA_IIS	( 315.0 / 270.0 ) 647640	( 5.47 , N/A ) ( N/A , 0.00 , N/A )	2628.7	N/A	1.2830 [ 1.0000 ]	128.3% { 134.8% }			
13C4_PFOA_IIS	( 417.0 / 372.0 ) 766471	( 6.98 , N/A ) ( N/A , -0.01 , N/A )	1156.2	N/A	1.2094 [ 1.0000 ]	120.9% { 125.6% }			
13C5_PFNA_IIS	( 468.0 / 423.0 ) 600707	( 7.58 , N/A ) ( N/A , -0.01 , N/A )	1636.7	N/A	0.9941 [ 1.0000 ]	99.4% { 106.5% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) ( $\Delta$ RT-I[min], $\Delta$ RT-CV[min], $\Delta$ 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 ) 433817	( 8.13 , N/A ) ( N/A , -0.01 , N/A )	1338.3	N/A	0.7497 [ 1.0000 ]	75.0% { 81.1% }			
18O2_PFHxS_IIS	( 403.0 / 83.9 ) 1241954	( 7.13 , N/A ) ( N/A , 0.00 , N/A )	1485.7	N/A	1.4503 [ 1.0000 ]	145.0% { 147.5% }			
13C4_PFOS_IIS	( 503.0 / 79.9 ) 1951122	( 8.34 , N/A ) ( N/A , -0.02 , N/A )	645.7	N/A	1.2961 [ 1.0000 ]	129.6% { 145.5% }			
13C4_PFBA_EIS	( 217.0 / 172.0 ) 1366449	( 3.50 , N/A ) ( N/A , 0.01 , N/A )	4414.8	N/A	6.5451 [ 8.0000 ]	81.8% { 59.5% }			
13C5_PFPeA_EIS	( 268.0 / 223.0 ) 1803518	( 4.55 , N/A ) ( N/A , 0.01 , N/A )	2218.8	N/A	2.5452 [ 4.0000 ]	63.6% { 78.3% }			
13C5_PFHxA_EIS	( 318.0 / 273.0 ) 1351134	( 5.47 , N/A ) ( N/A , 0.00 , N/A )	2159.5	N/A	1.5231 [ 2.0000 ]	76.2% { 94.6% }			
13C4_PFHpA_EIS	( 367.0 / 322.0 ) 1147520	( 6.29 , N/A ) ( N/A , 0.00 , N/A )	1067.9	N/A	1.4930 [ 2.0000 ]	74.6% { 97.4% }			
13C8_PFOA_EIS	( 421.0 / 376.0 ) 1568350	( 6.98 , N/A ) ( N/A , 0.00 , N/A )	1879.2	N/A	1.8067 [ 2.0000 ]	90.3% { 110.4% }			
13C9_PFNA_EIS	( 472.0 / 427.0 ) 546819	( 7.59 , N/A ) ( N/A , 0.00 , N/A )	1655.4	N/A	0.8591 [ 1.0000 ]	85.9% { 89.8% }			
13C6_PFDA_EIS	( 519.0 / 474.0 ) 494565	( 8.13 , N/A ) ( N/A , -0.01 , N/A )	705.8	N/A	0.9042 [ 1.0000 ]	90.4% { 66.3% }			
13C7_PFUnA_EIS	( 570.0 / 525.0 ) 826806	( 8.63 , N/A ) ( N/A , 0.00 , N/A )	1273.4	N/A	1.5201 [ 1.0000 ]	152.0% { 124.1% }			S2,

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 ) 585909	( 8.92, N/A ) ( N/A, 0.00, N/A )	1192.5	N/A	1.2560 [ 1.0000 ]	125.6% { 89.0% }			
13C2_PFTeDA_EIS	( 715.0 / 670.0 ) 549953	( 9.29, N/A ) ( N/A, 0.00, N/A )	1182.1	N/A	1.2866 [ 1.0000 ]	128.7% { 93.9% }			
13C3_PFBs_EIS	( 302.0 / 80.0 ) 4207576	( 5.44, N/A ) ( N/A, 0.00, N/A )	1249.1	N/A	1.6187 [ 2.0000 ]	80.9% { 110.9% }			
13C3_PFHxS_EIS	( 402.0 / 80.0 ) 1983919	( 7.13, N/A ) ( N/A, 0.00, N/A )	342.9	N/A	1.4694 [ 2.0000 ]	73.5% { 100.9% }			
13C8_PFOS_EIS	( 507.0 / 80.0 ) 4189834	( 8.34, N/A ) ( N/A, -0.02, N/A )	1256.4	N/A	1.5642 [ 2.0000 ]	78.2% { 98.9% }			
13C2_4:2FTS_EIS	( 329.0 / 81.0 ) 1089157	( 5.20, N/A ) ( N/A, -0.01, N/A )	813.7	N/A	7.5913 [ 4.0000 ]	189.8% { 313.6% }			S2,
13C2_6:2FTS_EIS	( 429.0 / 81.0 ) 1039339	( 6.72, N/A ) ( N/A, 0.00, N/A )	620.7	N/A	6.5993 [ 4.0000 ]	165.0% { 249.8% }			S2,
13C2_8:2FTS_EIS	( 529.0 / 81.0 ) 981420	( 7.88, N/A ) ( N/A, -0.01, N/A )	723.4	N/A	4.4108 [ 4.0000 ]	110.3% { 162.7% }			
13C8_PFOsa_EIS	( 506.0 / 78.0 ) 6667246	( 9.84, N/A ) ( N/A, 0.01, N/A )	2634.0	N/A	1.3409 [ 2.0000 ]	67.0% { 80.3% }			
D3_NMeFOsa_EIS	( 515.0 / 169.0 ) 1220987	( 10.42, N/A ) ( N/A, 0.00, N/A )	2204.9	N/A	1.0423 [ 2.0000 ]	52.1% { 63.1% }			
D5_NEtFOsa_EIS	( 531.0 / 169.0 ) 1072476	( 10.57, N/A ) ( N/A, 0.01, N/A )	2767.3	N/A	1.0231 [ 2.0000 ]	51.2% { 60.1% }			

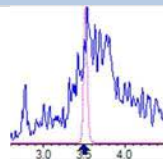
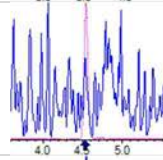
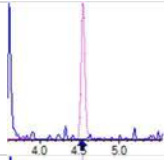
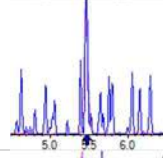
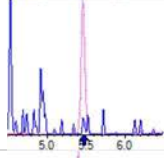
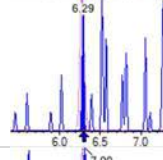
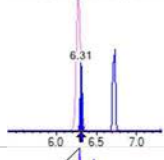
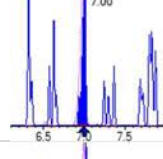
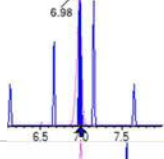
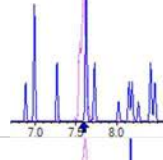
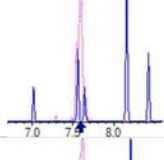
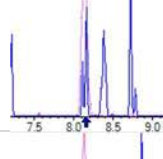
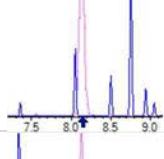
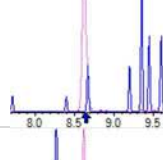
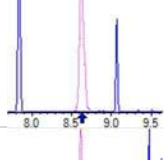
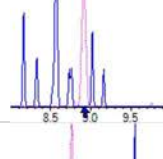
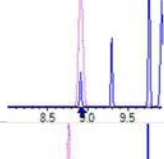
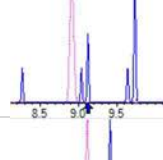
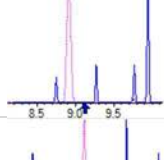
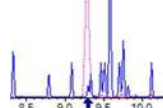
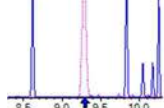


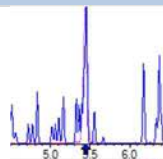
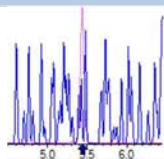
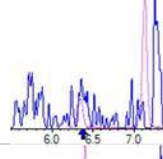
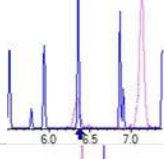
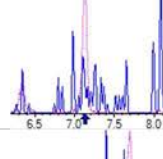
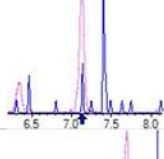
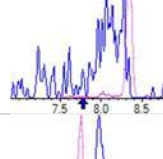
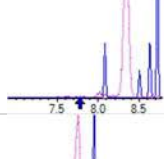
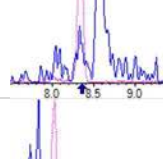
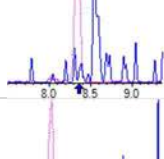
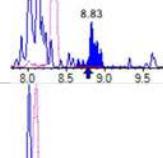
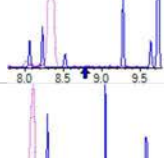
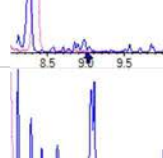
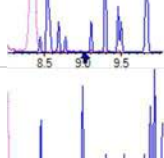
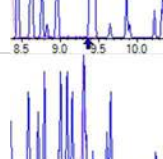
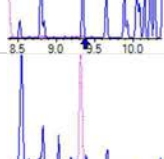
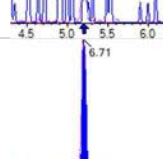
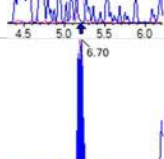
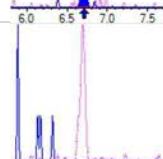
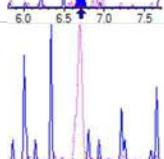
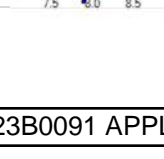
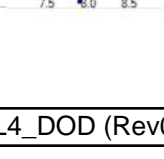
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	( 573.0 / 419.0 ) 1106094	( 8.28 , N/A ) ( N/A , -0.01 , N/A )	1469.8	N/A	2.5517 [ 4.0000 ]	63.8% { 83.5% }			
D5_EtFOSAA_EIS	( 589.0 / 419.0 ) 1410510	( 8.53 , N/A ) ( N/A , 0.00 , N/A )	20863.1	N/A	3.9740 [ 4.0000 ]	99.3% { 133.7% }			
D7_NMeFOSE_EIS	( 623.0 / 58.9 ) 3529153	( 10.36 , N/A ) ( N/A , 0.01 , N/A )	1940.0	N/A	8.2635 [ 20.0000 ]	41.3% { 47.5% }			
D9_NEtFOSE_EIS	( 639.0 / 58.9 ) 6468596	( 10.51 , N/A ) ( N/A , 0.01 , N/A )	2188.0	N/A	11.8271 [ 20.0000 ]	59.1% { 68.7% }			
13C3_HFPODA_EIS	( 287.0 / 169.0 ) 3417394	( 5.78 , N/A ) ( N/A , 0.00 , N/A )	3009.8	N/A	6.3271 [ 8.0000 ]	79.1% { 95.3% }			

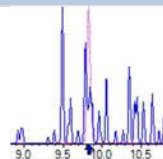
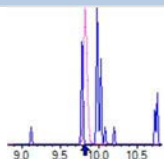
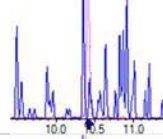
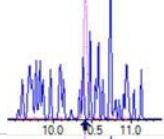
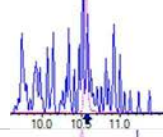
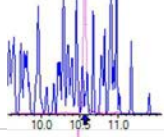
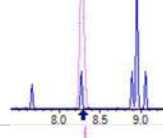
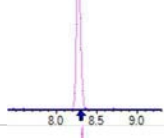
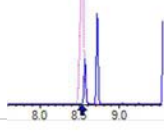
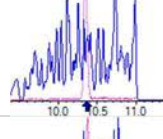
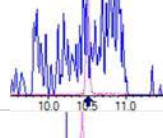
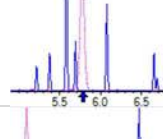
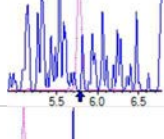
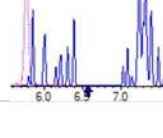
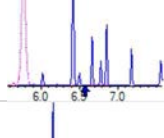
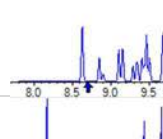
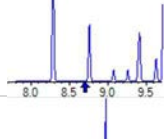
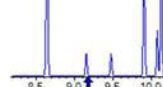



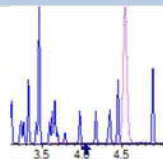
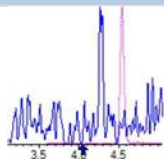
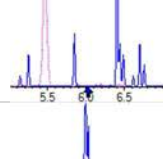
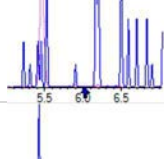
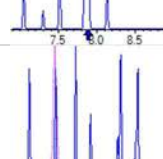
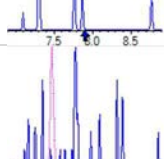
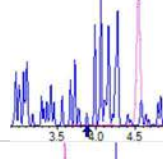

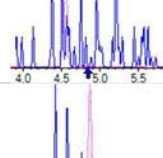
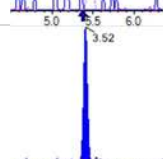
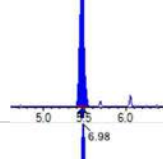

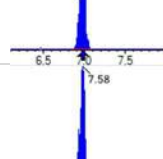
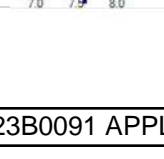
**FORM I**  
**ANALYSIS DATA SHEET**  
**AF-RHMW17D-WGN01LF-2302W1**

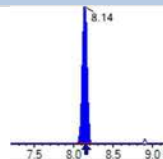
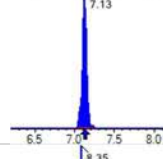
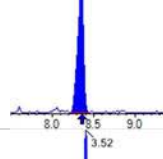
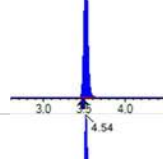
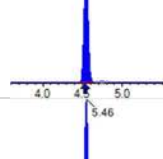
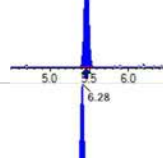
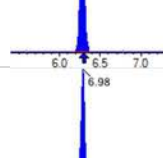
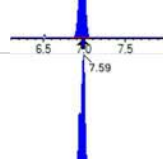
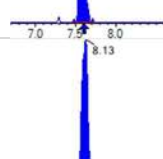
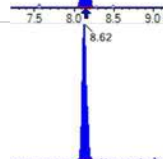
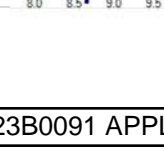
Laboratory:	APPL, LLC	Work Order:	23B0091		
Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling		
Matrix:	Water	Laboratory ID:	23B0091-02RE1	File ID:	S2023-02-14D (23)
Sampled:	02/10/23 12:15	Prepared:	02/13/23 07:19	Analyzed:	02/15/23 00:26
Solids:		Preparation:	EPA 1633	Dilution:	10
Initial/Final:	569.47 mL / 2 mL			Instrument:	Saphira
Batch:	BCB0232	Sequence:	SC00636	Calibration:	2307007

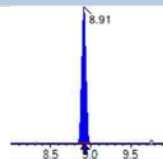
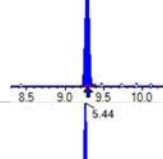
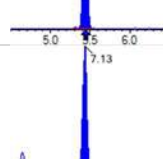
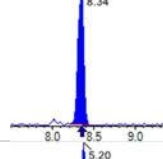
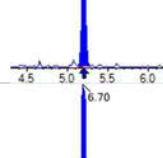
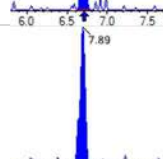
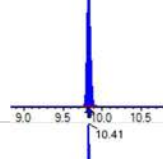
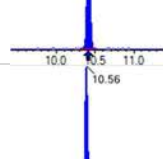
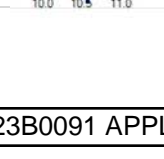
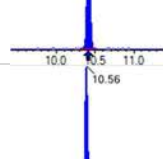
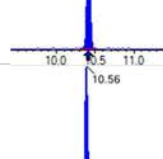
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	( 213.0 / 169.0 ) N/A	( N/A , N/A ) ( N/A , N/A , N/A )	N/A	N/A 0.0 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 ) 1190 ( 363.0 / 169.0 ) 493	( 6.29 , 1.00 ) ( 0.01 , N/A , -1.6 )	865.7 407.2	0.4146 129.3 129.7	0.0022	N/A			
PFOA	( 413.0 / 369.0 ) 3706 ( 413.0 / 169.0 ) 1678	( 7.00 , 1.00 ) ( 0.02 , N/A , 1.0 )	5603.2 80549.7	0.4526 135.3 125.3	0.0060	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			

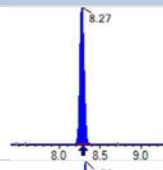
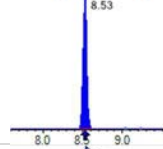
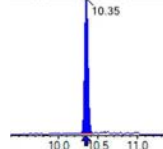
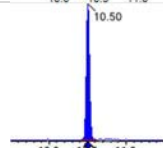
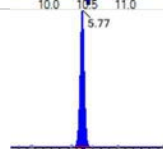
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	( 299.0 / 80.0 ) N/A ( 299.0 / 99.0 ) N/A	( N/A , N/A ) ( N/A , N/A , 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 ) 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 ) 15454 ( 549.0 / 99.0 ) N/A	( 8.83 , 1.06 ) ( N/A , 0.06 , #Value! )	70.8 N/A	N/A 0.0 0.0	0.0062	N/A			IR1,
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 ) 33509 ( 427.0 / 81.0 ) 14889	( 6.71 , 1.00 ) ( 0.00 , N/A , 0.7 )	1070.8 243.0	0.4443 53.7 57.5	0.2899	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 ) N/A	( N/A , N/A ) ( N/A , N/A , N/A )	N/A	N/A 0.0 0.0	0.0000	N/A			
NEiFOSE	( 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 ) 30788	( 3.52 , N/A ) ( N/A , 0.02 , N/A )	178.4	N/A	1.3745 [ 1.0000 ]	137.5% { 14.0% }			
13C2_PFHxA_IIS	( 315.0 / 270.0 ) 62060	( 5.47 , N/A ) ( N/A , -0.01 , N/A )	801.5	N/A	1.2294 [ 1.0000 ]	122.9% { 12.9% }			
13C4_PFOA_IIS	( 417.0 / 372.0 ) 73422	( 6.98 , N/A ) ( N/A , -0.01 , N/A )	34298689.2	N/A	1.1585 [ 1.0000 ]	115.8% { 12.0% }			
13C5_PFNA_IIS	( 468.0 / 423.0 ) 82086	( 7.58 , N/A ) ( N/A , -0.01 , N/A )	4582.8	N/A	1.3584 [ 1.0000 ]	135.8% { 14.6% }			

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 ) 87706	( 8.14, N/A ) ( N/A, -0.01, N/A )	24791741.1	N/A	1.5157 [ 1.0000 ]	151.6% { 16.4% }			
18O2_PFHxS_IIS	( 403.0 / 83.9 ) 114686	( 7.13, N/A ) ( N/A, -0.01, N/A )	208186.5	N/A	1.3392 [ 1.0000 ]	133.9% { 13.6% }			
13C4_PFOS_IIS	( 503.0 / 79.9 ) 207621	( 8.35, N/A ) ( N/A, -0.01, N/A )	215.3	N/A	1.3792 [ 1.0000 ]	137.9% { 15.5% }			
13C4_PFBA_EIS	( 217.0 / 172.0 ) 249290	( 3.52, N/A ) ( N/A, 0.02, N/A )	1603.0	N/A	0.6807 [ 0.8000 ]	85.1% { 10.9% }			
13C5_PFPeA_EIS	( 268.0 / 223.0 ) 246199	( 4.54, N/A ) ( N/A, 0.00, N/A )	828.8	N/A	0.3626 [ 0.4000 ]	90.6% { 10.7% }			
13C5_PFHxA_EIS	( 318.0 / 273.0 ) 143605	( 5.46, N/A ) ( N/A, -0.01, N/A )	604.0	N/A	0.1689 [ 0.2000 ]	84.5% { 10.1% }			
13C4_PFHpA_EIS	( 367.0 / 322.0 ) 129417	( 6.28, N/A ) ( N/A, -0.01, N/A )	2034.4	N/A	0.1757 [ 0.2000 ]	87.9% { 11.0% }			
13C8_PFOA_EIS	( 421.0 / 376.0 ) 139203	( 6.98, N/A ) ( N/A, 0.00, N/A )	19493.0	N/A	0.1674 [ 0.2000 ]	83.7% { 9.8% }			
13C9_PFNA_EIS	( 472.0 / 427.0 ) 69816	( 7.59, N/A ) ( N/A, 0.00, N/A )	45818.9	N/A	0.0803 [ 0.1000 ]	80.3% { 11.5% }			
13C6_PFDA_EIS	( 519.0 / 474.0 ) 92369	( 8.13, N/A ) ( N/A, -0.01, N/A )	941.6	N/A	0.0835 [ 0.1000 ]	83.5% { 12.4% }			
13C7_PFUnA_EIS	( 570.0 / 525.0 ) 78924	( 8.62, N/A ) ( N/A, -0.01, N/A )	995.2	N/A	0.0718 [ 0.1000 ]	71.8% { 11.8% }			

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 ) 70495	( 8.91 , N/A ) ( N/A , -0.01 , N/A )	113227.1	N/A	0.0747 [ 0.1000 ]	74.7% { 10.7% }			
13C2_PFTeDA_EIS	( 715.0 / 670.0 ) 64762	( 9.29 , N/A ) ( N/A , 0.00 , N/A )	571.0	N/A	0.0749 [ 0.1000 ]	74.9% { 11.1% }			
13C3_PFBs_EIS	( 302.0 / 80.0 ) 409334	( 5.44 , N/A ) ( N/A , -0.01 , N/A )	749.9	N/A	0.1705 [ 0.2000 ]	85.3% { 10.8% }			
13C3_PFHxS_EIS	( 402.0 / 80.0 ) 201384	( 7.13 , N/A ) ( N/A , -0.01 , N/A )	214.0	N/A	0.1615 [ 0.2000 ]	80.8% { 10.2% }			
13C8_PFOS_EIS	( 507.0 / 80.0 ) 424068	( 8.34 , N/A ) ( N/A , -0.01 , N/A )	282.6	N/A	0.1488 [ 0.2000 ]	74.4% { 10.0% }			
13C2_4:2FTS_EIS	( 329.0 / 81.0 ) 75192	( 5.20 , N/A ) ( N/A , -0.01 , N/A )	246.6	N/A	0.5675 [ 0.4000 ]	141.9% { 21.7% }			
13C2_6:2FTS_EIS	( 429.0 / 81.0 ) 32583	( 6.70 , N/A ) ( N/A , -0.01 , N/A )	148.4	N/A	0.2240 [ 0.4000 ]	56.0% { 7.8% }			
13C2_8:2FTS_EIS	( 529.0 / 81.0 ) 70887	( 7.89 , N/A ) ( N/A , 0.00 , N/A )	188.4	N/A	0.3450 [ 0.4000 ]	86.2% { 11.8% }			
13C8_PFOSA_EIS	( 506.0 / 78.0 ) 669689	( 9.83 , N/A ) ( N/A , -0.01 , N/A )	1852.7	N/A	0.1266 [ 0.2000 ]	63.3% { 8.1% }			
D3_NMeFOSA_EIS	( 515.0 / 169.0 ) 139739	( 10.41 , N/A ) ( N/A , 0.00 , N/A )	671.5	N/A	0.1121 [ 0.2000 ]	56.0% { 7.2% }			
D5_NEtFOSA_EIS	( 531.0 / 169.0 ) 114383	( 10.56 , N/A ) ( N/A , 0.00 , N/A )	809.3	N/A	0.1025 [ 0.2000 ]	51.3% { 6.4% }			

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 ) 209086	( 8.27 , N/A ) ( N/A , -0.01 , N/A )	901.8	N/A	0.4533 [ 0.4000 ]	113.3% { 15.8% }			
D5_EtFOSAA_EIS	( 589.0 / 419.0 ) 112026	( 8.53 , N/A ) ( N/A , 0.00 , N/A )	4842646.8	N/A	0.2966 [ 0.4000 ]	74.2% { 10.6% }			
D7_NMeFOSE_EIS	( 623.0 / 58.9 ) 458190	( 10.35 , N/A ) ( N/A , 0.00 , N/A )	399.1	N/A	1.0082 [ 2.0000 ]	50.4% { 6.2% }			
D9_NEtFOSE_EIS	( 639.0 / 58.9 ) 729184	( 10.50 , N/A ) ( N/A , 0.00 , N/A )	937.2	N/A	1.2529 [ 2.0000 ]	62.6% { 7.7% }			
13C3_HFPODA_EIS	( 287.0 / 169.0 ) 321442	( 5.77 , N/A ) ( N/A , -0.01 , N/A )	845.4	N/A	0.6211 [ 0.8000 ]	77.6% { 9.0% }			



# FORM I

## ANALYSIS DATA SHEET

### AF-RHMW17D-WQFB01-2302W1

Laboratory:	APPL, LLC	Work Order:	23B0091
Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Matrix:	Water	Laboratory ID:	23B0091-03
		File ID:	S2023-02-14D (24)
Sampled:	02/10/23 11:30	Prepared:	02/13/23 07:19
		Analyzed:	02/15/23 00:39
Solids:		Preparation:	EPA 1633
		Dilution:	1
Initial/Final:	556.9 mL / 2 mL	Instrument:	Saphira
Batch:	BCB0232	Sequence:	SC00636
		Calibration:	2307007

COMPOUND	CONC. (ng/L)	LOQ	LOD	DL	Q
PFBA	0.72 U	1.4	0.72	0.19	
PFPEA	0.36 U	0.72	0.36	0.058	
PFHXA	0.18 U	0.36	0.18	0.049	
PFHPA	0.18 U	0.36	0.18	0.037	
PFOA	0.18 U	0.36	0.18	0.14	
PFNA	0.18 U	0.36	0.18	0.073	
PFDA	0.18 U	0.36	0.18	0.091	
PFUnA	0.18 U	0.36	0.18	0.14	
PFDOA	0.18 U	0.36	0.18	0.10	
PFTRDA	0.27 U	0.36	0.27	0.18	
PFTEDA	0.18 U	0.36	0.18	0.18	
PFBS	0.18 U	0.36	0.18	0.033	
PFPEs	0.18 U	0.36	0.18	0.056	
PFHXS	0.18 U	0.36	0.18	0.028	
PFHPS	0.18 U	0.36	0.18	0.046	
PFOS	0.17 J	0.36	0.18	0.057	MI5,
PFNS	0.18 U	0.36	0.18	0.11	
PFDS	0.18 U	0.36	0.18	0.14	
PFDOS	0.18 U	0.36	0.18	0.11	
4:2FTS	0.72 U	1.4	0.72	0.26	
6:2FTS	0.72 U	1.4	0.72	0.28	
8:2FTS	0.72 U	1.4	0.72	0.074	
PFOSA	0.18 U	0.36	0.18	0.093	
NMeFOSA	0.72 U	1.4	0.72	0.42	
NEtFOSA	0.72 U	1.4	0.72	0.37	
NMeFOSAA	0.18 U	0.36	0.18	0.095	
NEtFOSAA	0.18 U	0.36	0.18	0.10	
NMeFOSE	1.1 U	1.4	1.1	0.91	
NEtFOSE	1.1 U	1.4	1.1	0.94	
HFPO-DA	0.36 U	0.72	0.36	0.16	

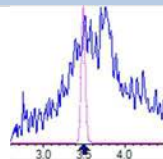
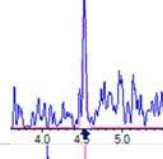
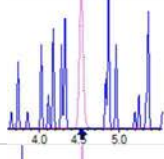
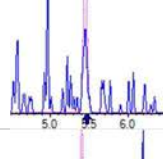
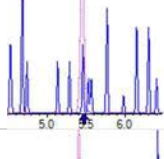
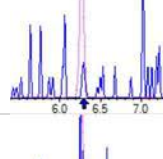
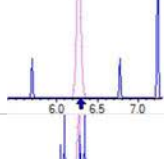
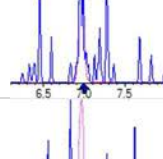
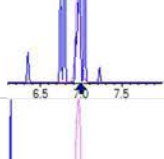
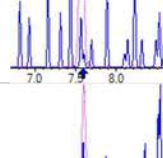
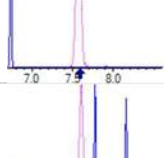
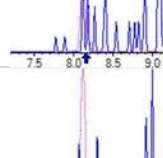
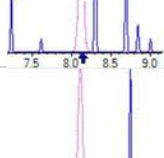
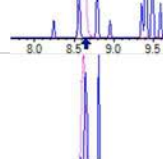
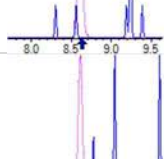
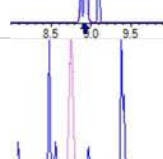
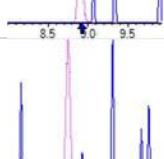
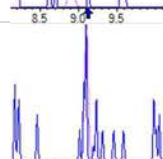
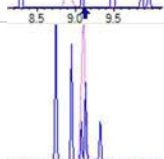
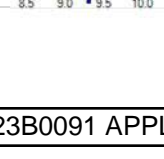
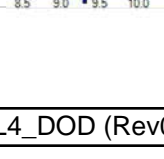
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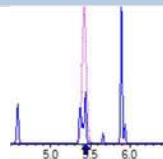
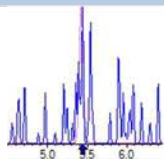
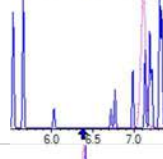
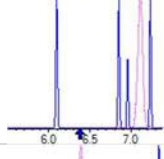
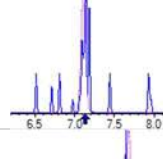
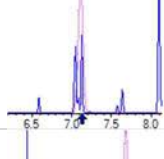
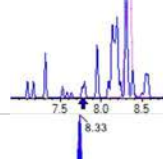
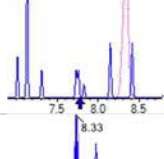
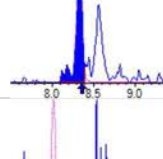
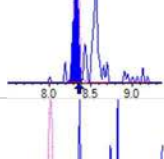
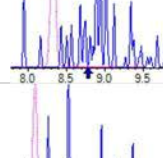
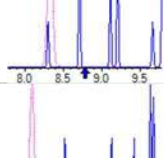
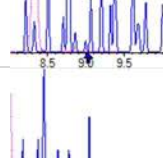
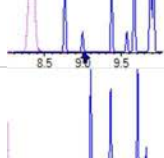
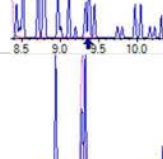
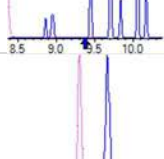
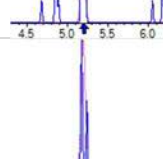
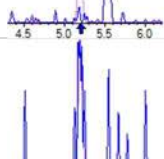
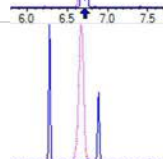
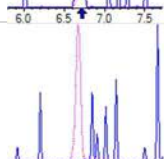
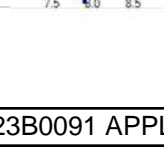
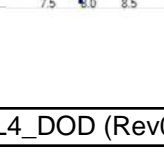
## ANALYSIS DATA SHEET

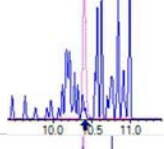
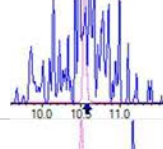
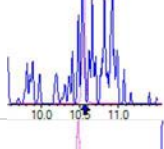
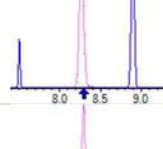
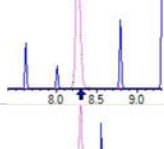
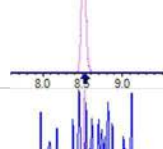
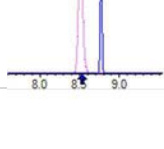
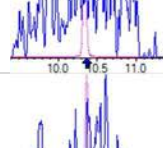
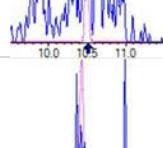
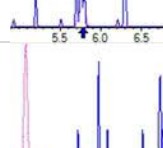
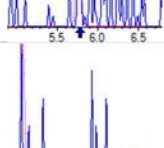
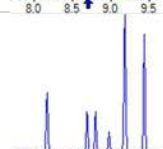
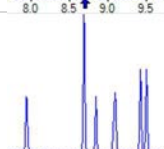
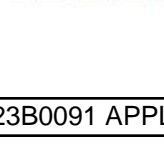
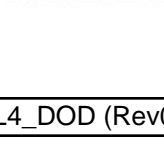
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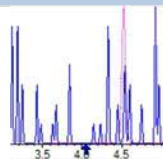
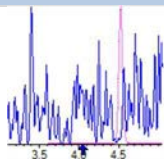
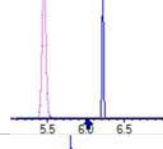
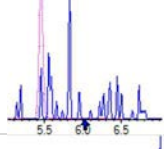
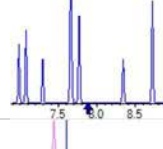
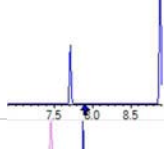
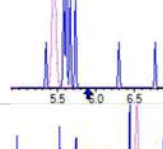
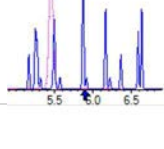
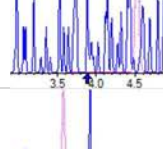
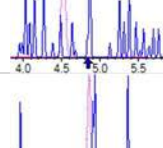
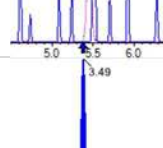
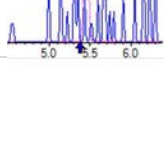
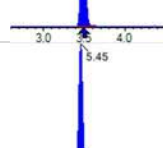
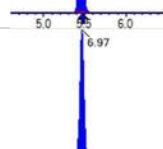
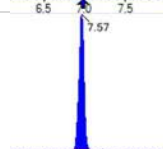

Laboratory:	APPL, LLC	Work Order:	23B0091
Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Matrix:	Water	Laboratory ID:	23B0091-03
		File ID:	S2023-02-14D (24)
Sampled:	02/10/23 11:30	Prepared:	02/13/23 07:19
		Analyzed:	02/15/23 00:39
Solids:		Preparation:	EPA 1633
		Dilution:	1
Initial/Final:	556.9 mL / 2 mL	Instrument:	Saphira
Batch:	BCB0232	Sequence:	SC00636
		Calibration:	2307007

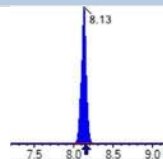
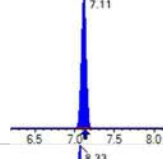
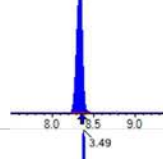
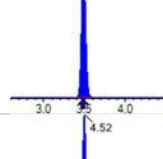
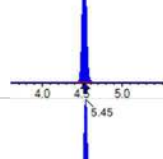
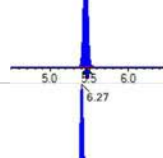
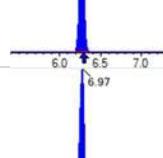
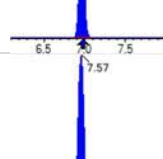
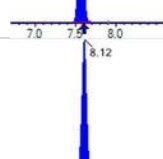
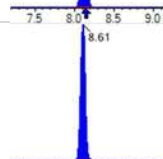
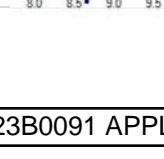
COMPOUND	CONC. (ng/L)	LOQ	LOD	DL	Q
ADONA	0.36 U	0.72	0.36	0.11	
PFEESA	0.36 U	0.72	0.36	0.098	
PFMPA	0.36 U	0.72	0.36	0.048	
PFMBA	0.36 U	0.72	0.36	0.081	
NFDHA	0.36 U	0.72	0.36	0.27	
9CL-PF3ONS	0.36 U	0.72	0.36	0.19	
11CL-PF3OUDS	0.36 U	0.72	0.36	0.19	
3:3FTCA	0.72 U	1.4	0.72	0.52	
5:3FTCA	0.72 U	1.4	0.72	0.40	
7:3FTCA	0.72 U	1.4	0.72	0.50	

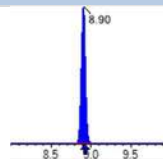
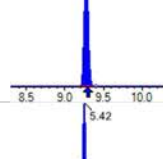
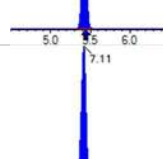
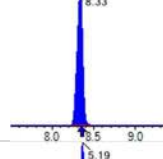
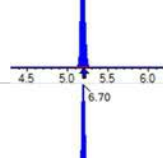
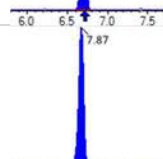
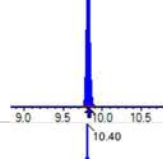
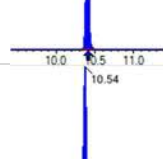
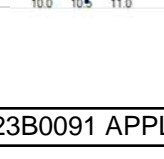
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	( 213.0 / 169.0 ) N/A	( N/A , N/A ) ( N/A , N/A , N/A )	N/A	N/A 0.0 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 ) 113179 ( 499.0 / 99.0 ) 22773	( 8.33 , 1.00 ) ( 0.00 , N/A , 0.3 )	91.7 105.7	0.2012 90.9 94.9	0.0476	N/A			MI5 DG 2023-02-15
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			

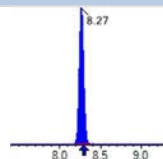
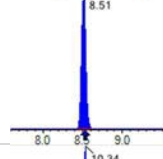
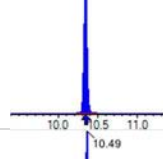
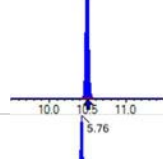
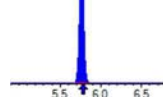
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	( 498.0 / 78.0 ) N/A ( 498.0 / 478.0 ) N/A	( N/A , N/A ) ( N/A , 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 ) N/A	( N/A , N/A ) ( N/A , N/A , N/A )	N/A	N/A 0.0 0.0	0.0000	N/A			
NEiFOSE	( 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 ) 297640	( 3.49 , N/A ) ( N/A , -0.01 , N/A )	1348.0	N/A	1.3288 [ 1.0000 ]	132.9% { 135.6% }			
13C2_PFHxA_IIS	( 315.0 / 270.0 ) 583935	( 5.45 , N/A ) ( N/A , -0.02 , N/A )	1731.4	N/A	1.1568 [ 1.0000 ]	115.7% { 121.5% }			
13C4_PFOA_IIS	( 417.0 / 372.0 ) 730818	( 6.97 , N/A ) ( N/A , -0.02 , N/A )	34039.5	N/A	1.1531 [ 1.0000 ]	115.3% { 119.7% }			
13C5_PFNA_IIS	( 468.0 / 423.0 ) 667133	( 7.57 , N/A ) ( N/A , -0.02 , N/A )	1303.8	N/A	1.1040 [ 1.0000 ]	110.4% { 118.3% }			

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
13C2_PFDA_IIS	( 515.0 / 470.1 ) 728704	( 8.13 , N/A ) ( N/A , -0.02 , N/A )	303.4	N/A	1.2593 [ 1.0000 ]	125.9% { 136.2% }			
18O2_PFHxS_IIS	( 403.0 / 83.9 ) 1057329	( 7.11 , N/A ) ( N/A , -0.02 , N/A )	2296.6	N/A	1.2347 [ 1.0000 ]	123.5% { 125.6% }			
13C4_PFOS_IIS	( 503.0 / 79.9 ) 1959790	( 8.33 , N/A ) ( N/A , -0.02 , N/A )	1457.9	N/A	1.3018 [ 1.0000 ]	130.2% { 146.1% }			
13C4_PFBA_EIS	( 217.0 / 172.0 ) 2456217	( 3.49 , N/A ) ( N/A , -0.01 , N/A )	5693.9	N/A	6.9377 [ 8.0000 ]	86.7% { 106.9% }			
13C5_PFPeA_EIS	( 268.0 / 223.0 ) 2441137	( 4.52 , N/A ) ( N/A , -0.02 , N/A )	3740.4	N/A	3.8208 [ 4.0000 ]	95.5% { 105.9% }			
13C5_PFHxA_EIS	( 318.0 / 273.0 ) 1418282	( 5.45 , N/A ) ( N/A , -0.02 , N/A )	3023.1	N/A	1.7732 [ 2.0000 ]	88.7% { 99.3% }			
13C4_PFHpA_EIS	( 367.0 / 322.0 ) 1225509	( 6.27 , N/A ) ( N/A , -0.02 , N/A )	2836.9	N/A	1.7684 [ 2.0000 ]	88.4% { 104.0% }			
13C8_PFOA_EIS	( 421.0 / 376.0 ) 1460484	( 6.97 , N/A ) ( N/A , -0.02 , N/A )	1347.9	N/A	1.7646 [ 2.0000 ]	88.2% { 102.8% }			
13C9_PFNA_EIS	( 472.0 / 427.0 ) 634157	( 7.57 , N/A ) ( N/A , -0.02 , N/A )	1263.0	N/A	0.8972 [ 1.0000 ]	89.7% { 104.2% }			
13C6_PFDA_EIS	( 519.0 / 474.0 ) 732702	( 8.12 , N/A ) ( N/A , -0.02 , N/A )	3762.3	N/A	0.7975 [ 1.0000 ]	79.7% { 98.3% }			
13C7_PFUnA_EIS	( 570.0 / 525.0 ) 718205	( 8.61 , N/A ) ( N/A , -0.02 , N/A )	6348.5	N/A	0.7861 [ 1.0000 ]	78.6% { 107.8% }			

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 ) 730421	( 8.90 , N/A ) ( N/A , -0.01 , N/A )	3033.9	N/A	0.9321 [ 1.0000 ]	93.2% { 111.0% }			
13C2_PFTeDA_EIS	( 715.0 / 670.0 ) 651427	( 9.28 , N/A ) ( N/A , -0.01 , N/A )	1478.8	N/A	0.9073 [ 1.0000 ]	90.7% { 111.2% }			
13C3_PFBs_EIS	( 302.0 / 80.0 ) 3963429	( 5.42 , N/A ) ( N/A , -0.02 , N/A )	4154.9	N/A	1.7911 [ 2.0000 ]	89.6% { 104.5% }			
13C3_PFHxS_EIS	( 402.0 / 80.0 ) 2025389	( 7.11 , N/A ) ( N/A , -0.02 , N/A )	1804.3	N/A	1.7621 [ 2.0000 ]	88.1% { 103.0% }			
13C8_PFOS_EIS	( 507.0 / 80.0 ) 4508729	( 8.33 , N/A ) ( N/A , -0.02 , N/A )	1813.6	N/A	1.6758 [ 2.0000 ]	83.8% { 106.4% }			
13C2_4:2FTS_EIS	( 329.0 / 81.0 ) 871156	( 5.19 , N/A ) ( N/A , -0.02 , N/A )	2786.6	N/A	7.1321 [ 4.0000 ]	178.3% { 250.9% }			S2,
13C2_6:2FTS_EIS	( 429.0 / 81.0 ) 433054	( 6.70 , N/A ) ( N/A , -0.02 , N/A )	1703.8	N/A	3.2298 [ 4.0000 ]	80.7% { 104.1% }			
13C2_8:2FTS_EIS	( 529.0 / 81.0 ) 534249	( 7.87 , N/A ) ( N/A , -0.02 , N/A )	1413.5	N/A	2.8203 [ 4.0000 ]	70.5% { 88.6% }			
13C8_PFOsa_EIS	( 506.0 / 78.0 ) 7259414	( 9.82 , N/A ) ( N/A , -0.01 , N/A )	2757.7	N/A	1.4535 [ 2.0000 ]	72.7% { 87.4% }			
D3_NMeFOsa_EIS	( 515.0 / 169.0 ) 1286443	( 10.40 , N/A ) ( N/A , -0.02 , N/A )	2189.4	N/A	1.0933 [ 2.0000 ]	54.7% { 66.5% }			
D5_NEtFOsa_EIS	( 531.0 / 169.0 ) 1098308	( 10.54 , N/A ) ( N/A , -0.02 , N/A )	2594.1	N/A	1.0431 [ 2.0000 ]	52.2% { 61.6% }			



Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) ( $\Delta$ RT-I[min], $\Delta$ RT- CV[min], $\Delta$ 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
D3_MeFOSAA_EIS	( 573.0 / 419.0 ) 1203370	( 8.27 , N/A ) ( N/A , -0.02 , N/A )	1396.8	N/A	2.7638 [ 4.0000 ]	69.1% { 90.9% }			
D5_EtFOSAA_EIS	( 589.0 / 419.0 ) 993867	( 8.51 , N/A ) ( N/A , -0.02 , N/A )	8058.6	N/A	2.7877 [ 4.0000 ]	69.7% { 94.2% }			
D7_NMeFOSE_EIS	( 623.0 / 58.9 ) 3876289	( 10.34 , N/A ) ( N/A , -0.01 , N/A )	1228.6	N/A	9.0362 [ 20.0000 ]	45.2% { 52.2% }			
D9_NEtFOSE_EIS	( 639.0 / 58.9 ) 7229919	( 10.49 , N/A ) ( N/A , -0.01 , N/A )	2343.6	N/A	13.1607 [ 20.0000 ]	65.8% { 76.7% }			
13C3_HFPODA_EIS	( 287.0 / 169.0 ) 3499731	( 5.76 , N/A ) ( N/A , -0.02 , N/A )	3540.7	N/A	7.1864 [ 8.0000 ]	89.8% { 97.6% }			

# FORM I

## ANALYSIS DATA SHEET

AF-RHMW225401-WGN01B-2302W1

Laboratory:	APPL, LLC	Work Order:	23B0091
Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Matrix:	Water	Laboratory ID:	23B0091-04
		File ID:	S2023-02-14D (26)
Sampled:	02/10/23 08:55	Prepared:	02/13/23 07:19
		Analyzed:	02/15/23 01:04
Solids:		Preparation:	EPA 1633
		Dilution:	1
Initial/Final:	527.55 mL / 2 mL	Instrument:	Saphira
Batch:	BCB0232	Sequence:	SC00636
		Calibration:	2307007

COMPOUND	CONC. (ng/L)	LOQ	LOD	DL	Q
PFBA	0.60 J	1.5	0.76	0.20	
PFPEA	1.3	0.76	0.38	0.062	
PFHXA	1.0	0.38	0.19	0.052	
PFHPA	0.83	0.38	0.19	0.039	
PFOA	1.0	0.38	0.19	0.14	
PFNA	0.17 J	0.38	0.19	0.078	IR1,
PFDA	0.19 U	0.38	0.19	0.096	
PFUnA	0.19 U	0.38	0.19	0.15	
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	
PFBS	0.76	0.38	0.19	0.035	
PFPEs	0.16 J	0.38	0.19	0.059	
PFHXS	1.2	0.38	0.19	0.030	
PFHPS	0.19 U	0.38	0.19	0.049	
PFOS	1.1	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	0.76 U	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	

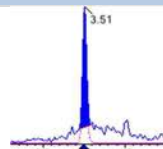
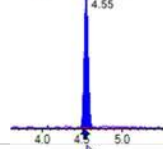
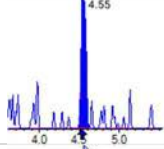
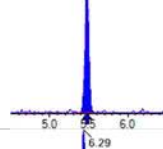
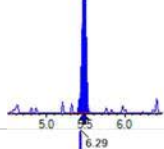
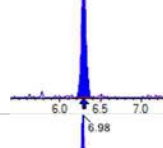
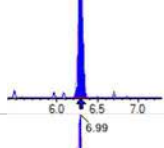
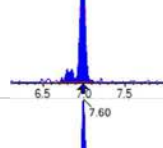
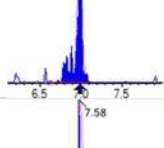
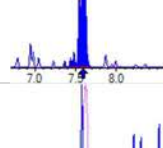
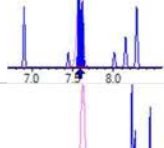
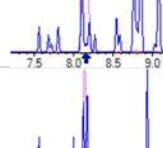
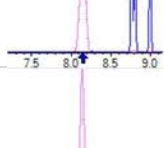
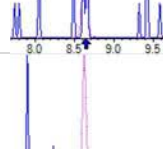
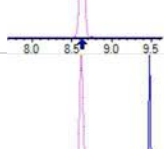
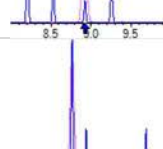
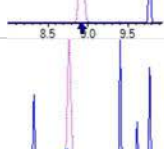
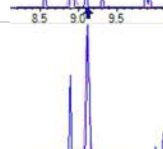
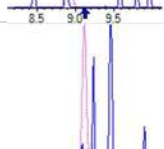
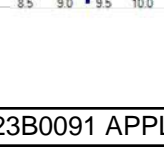
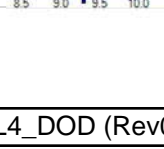
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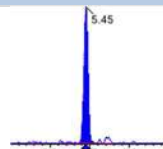
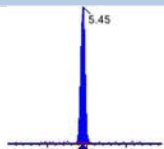
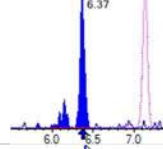
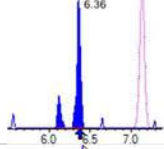
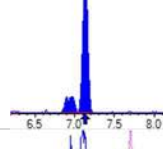
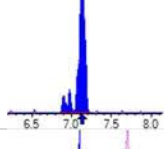
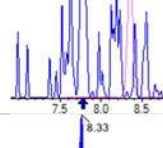
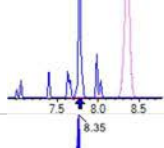
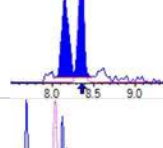
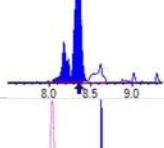
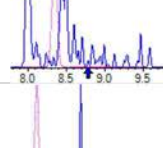
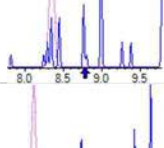
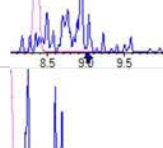
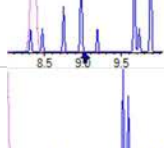
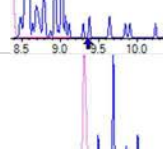
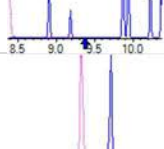
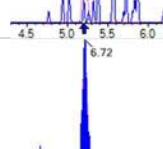
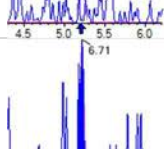
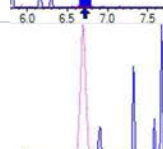
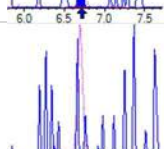

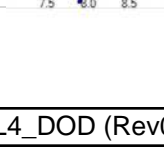
## ANALYSIS DATA SHEET

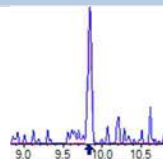
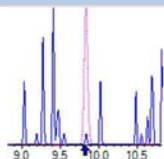
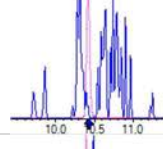
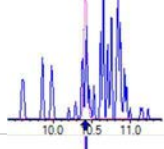
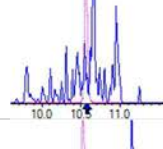
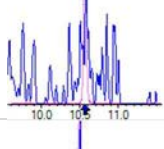
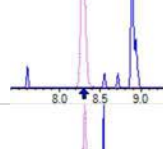
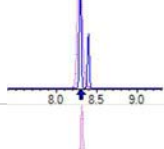
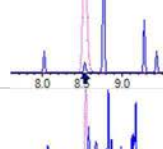
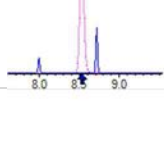
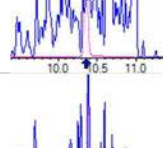
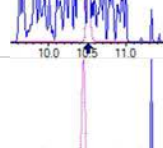
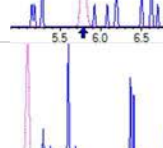
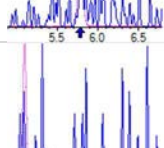
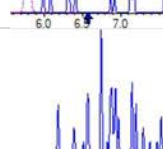
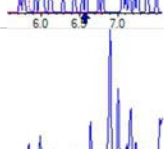
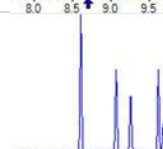
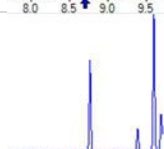
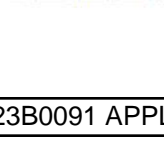
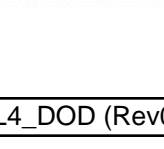
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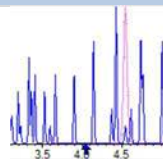
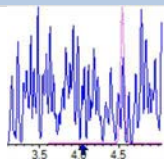
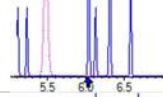
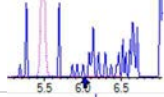
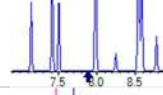
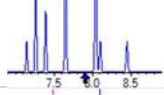
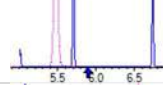
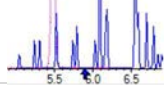
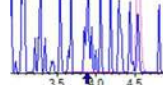
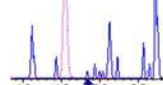
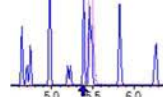
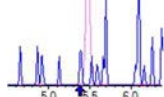
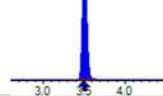
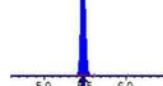
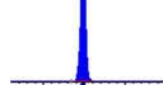
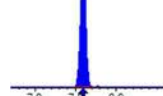
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Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Matrix:	Water	Laboratory ID:	23B0091-04
		File ID:	S2023-02-14D (26)
Sampled:	02/10/23 08:55	Prepared:	02/13/23 07:19
		Analyzed:	02/15/23 01:04
Solids:		Preparation:	EPA 1633
		Dilution:	1
Initial/Final:	527.55 mL / 2 mL	Instrument:	Saphira
Batch:	BCB0232	Sequence:	SC00636
		Calibration:	2307007

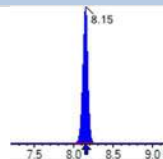
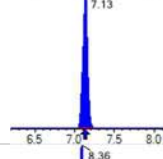
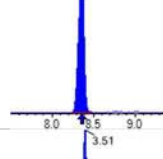
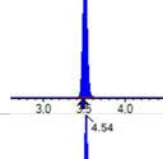
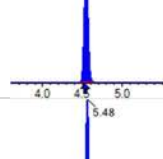
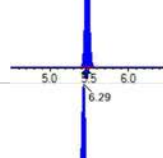
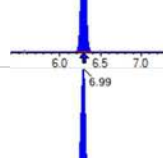
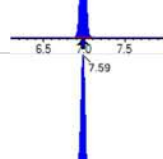
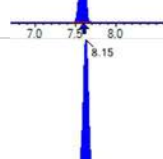
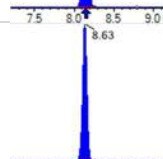
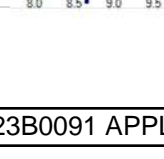
COMPOUND	CONC. (ng/L)	LOQ	LOD	DL	Q
ADONA	0.38 U	0.76	0.38	0.12	
PFEESA	0.38 U	0.76	0.38	0.10	
PFMPA	0.38 U	0.76	0.38	0.051	
PFMBA	0.38 U	0.76	0.38	0.086	
NFDHA	0.38 U	0.76	0.38	0.29	
9CL-PF3ONS	0.38 U	0.76	0.38	0.20	
11CL-PF3OUDS	0.38 U	0.76	0.38	0.20	
3:3FTCA	0.76 U	1.5	0.76	0.54	
5:3FTCA	0.76 U	1.5	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-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 ) 40576	( 3.51 , 1.00 ) ( 0.00 , N/A , 0.0 )	53.4	N/A 0.0 0.0	0.1578	N/A			
PFPeA	( 263.0 / 219.0 ) 181220 ( 263.0 / 69.0 ) 3237	( 4.55 , 1.00 ) ( 0.00 , N/A , -0.2 )	467.0 29.4	0.0179 166.8 162.1	0.3522	N/A			
PFHxA	( 313.0 / 269.0 ) 175493 ( 313.0 / 119.0 ) 22074	( 5.48 , 1.00 ) ( 0.00 , N/A , -0.1 )	445.3 4118.6	0.1258 126.3 132.2	0.2687	N/A			
PFHpA	( 363.0 / 319.0 ) 116486 ( 363.0 / 169.0 ) 35687	( 6.29 , 1.00 ) ( 0.00 , N/A , 0.2 )	573.8 1413.0	0.3064 95.6 95.8	0.2178	N/A			
PFOA	( 413.0 / 369.0 ) 180144 ( 413.0 / 169.0 ) 60264	( 6.98 , 1.00 ) ( 0.00 , N/A , -0.3 )	435.6 4111.9	0.3345 100.0 92.6	0.2757	N/A			
PFNA	( 463.0 / 419.0 ) 26611 ( 463.0 / 169.0 ) 2888	( 7.60 , 1.00 ) ( 0.00 , N/A , 1.1 )	2495.5 1191.4	0.1085 48.5 49.1	0.0458	N/A			IR1,
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			

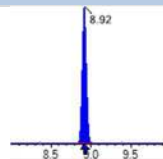
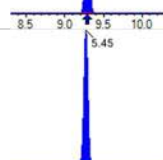
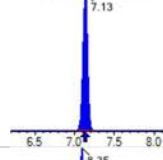
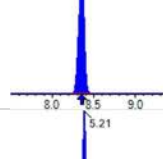
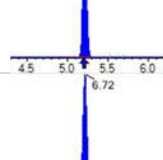
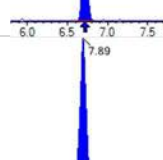
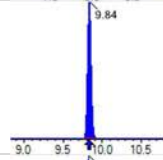
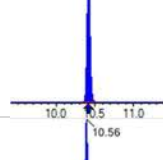
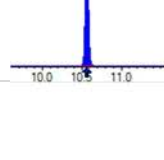
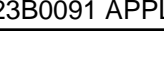
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	( 299.0 / 80.0 ) 250224 ( 299.0 / 99.0 ) 131433	( 5.45 , 1.00 ) ( 0.00 , N/A , 0.2 )	308.2 402.8	0.5253 81.2 84.9	0.1997	N/A			
PFPeS	( 349.0 / 80.0 ) 80181 ( 349.0 / 99.0 ) 16925	( 6.37 , 0.89 ) ( N/A , -0.01 , 0.3 )	11770.7 4521.8	0.2111 59.1 61.3	0.0426	N/A			
PFHxS	( 399.0 / 80.0 ) 473103 ( 399.0 / 99.0 ) 149135	( 7.13 , 1.00 ) ( 0.00 , N/A , 0.0 )	6258.1 2109.0	0.3152 97.1 96.8	0.3152	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 ) 735295 ( 499.0 / 99.0 ) 143824	( 8.33 , 1.00 ) ( -0.03 , N/A , -1.6 )	79.7 212.6	0.1956 88.4 92.3	0.2786	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 ) 7152 ( 427.0 / 81.0 ) 3877	( 6.72 , 1.00 ) ( 0.00 , N/A , 0.3 )	11642.7 30.6	0.5421 65.5 70.2	0.0480	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			

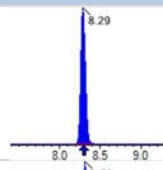
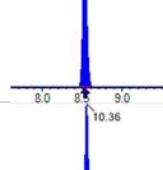
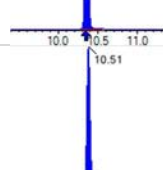
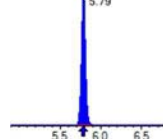

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) ( $\Delta$ RT-I[min], $\Delta$ RT-CV[min], $\Delta$ RT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	( 498.0 / 78.0 ) N/A ( 498.0 / 478.0 ) N/A	( 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 ) N/A	( N/A , N/A ) ( N/A , N/A , N/A )	N/A	N/A 0.0 0.0	0.0000	N/A			
NEiFOSE	( 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 ) 308013	( 3.51 , N/A ) ( N/A , 0.01 , N/A )	1408.4	N/A	1.3751 [ 1.0000 ]	137.5% { 140.4% }			
13C2_PFHxA_IIS	( 315.0 / 270.0 ) 592652	( 5.48 , N/A ) ( N/A , 0.00 , N/A )	1544.7	N/A	1.1740 [ 1.0000 ]	117.4% { 123.3% }			
13C4_PFOA_IIS	( 417.0 / 372.0 ) 705559	( 6.99 , N/A ) ( N/A , 0.00 , N/A )	1597.1	N/A	1.1133 [ 1.0000 ]	111.3% { 115.6% }			
13C5_PFNAl_IIS	( 468.0 / 423.0 ) 653697	( 7.59 , N/A ) ( N/A , 0.00 , N/A )	7650.5	N/A	1.0818 [ 1.0000 ]	108.2% { 115.9% }			

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_IIS	( 515.0 / 470.1 ) 774426	( 8.15 , N/A ) ( N/A , 0.00 , N/A )	2647.0	N/A	1.3384 [ 1.0000 ]	133.8% { 144.7% }			
18O2_PFHxS_IIS	( 403.0 / 83.9 ) 1000660	( 7.13 , N/A ) ( N/A , 0.00 , N/A )	5868.6	N/A	1.1685 [ 1.0000 ]	116.8% { 118.8% }			
13C4_PFOS_IIS	( 503.0 / 79.9 ) 2072436	( 8.36 , N/A ) ( N/A , 0.00 , N/A )	952.5	N/A	1.3767 [ 1.0000 ]	137.7% { 154.5% }			
13C4_PFBA_EIS	( 217.0 / 172.0 ) 2483715	( 3.51 , N/A ) ( N/A , 0.01 , N/A )	6145.9	N/A	6.7791 [ 8.0000 ]	84.7% { 108.1% }			
13C5_PFPeA_EIS	( 268.0 / 223.0 ) 2501775	( 4.54 , N/A ) ( N/A , 0.00 , N/A )	2905.1	N/A	3.8581 [ 4.0000 ]	96.5% { 108.5% }			
13C5_PFHxA_EIS	( 318.0 / 273.0 ) 1485717	( 5.48 , N/A ) ( N/A , 0.00 , N/A )	2954.7	N/A	1.8302 [ 2.0000 ]	91.5% { 104.1% }			
13C4_PFHpA_EIS	( 367.0 / 322.0 ) 1266447	( 6.29 , N/A ) ( N/A , 0.00 , N/A )	2007.7	N/A	1.8006 [ 2.0000 ]	90.0% { 107.5% }			
13C8_PFOA_EIS	( 421.0 / 376.0 ) 1467632	( 6.99 , N/A ) ( N/A , 0.00 , N/A )	1390.5	N/A	1.8367 [ 2.0000 ]	91.8% { 103.3% }			
13C9_PFNA_EIS	( 472.0 / 427.0 ) 657313	( 7.59 , N/A ) ( N/A , 0.00 , N/A )	1837.3	N/A	0.9490 [ 1.0000 ]	94.9% { 108.0% }			
13C6_PFDA_EIS	( 519.0 / 474.0 ) 850878	( 8.15 , N/A ) ( N/A , 0.00 , N/A )	1546.6	N/A	0.8714 [ 1.0000 ]	87.1% { 114.1% }			
13C7_PFUnA_EIS	( 570.0 / 525.0 ) 947170	( 8.63 , N/A ) ( N/A , 0.00 , N/A )	1516.0	N/A	0.9755 [ 1.0000 ]	97.5% { 142.2% }			



Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) ( $\Delta$ RT-I[min], $\Delta$ RT-CV[min], $\Delta$ 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_EIS	( 615.0 / 570.0 ) 793120	( 8.92, N/A ) ( N/A, 0.00, N/A )	2366.9	N/A	0.9524 [ 1.0000 ]	95.2% { 120.5% }			
13C2_PFTeDA_EIS	( 715.0 / 670.0 ) 625573	( 9.29, N/A ) ( N/A, 0.00, N/A )	11883.9	N/A	0.8198 [ 1.0000 ]	82.0% { 106.8% }			
13C3_PFBs_EIS	( 302.0 / 80.0 ) 4032385	( 5.45, N/A ) ( N/A, 0.00, N/A )	3050.2	N/A	1.9254 [ 2.0000 ]	96.3% { 106.3% }			
13C3_PFHxS_EIS	( 402.0 / 80.0 ) 1952312	( 7.13, N/A ) ( N/A, 0.00, N/A )	1642.9	N/A	1.7947 [ 2.0000 ]	89.7% { 99.3% }			
13C8_PFOS_EIS	( 507.0 / 80.0 ) 5005565	( 8.35, N/A ) ( N/A, 0.00, N/A )	1638.6	N/A	1.7593 [ 2.0000 ]	88.0% { 118.1% }			
13C2_4:2FTS_EIS	( 329.0 / 81.0 ) 880169	( 5.21, N/A ) ( N/A, 0.00, N/A )	1567.3	N/A	7.6140 [ 4.0000 ]	190.3% { 253.4% }			S2,
13C2_6:2FTS_EIS	( 429.0 / 81.0 ) 420156	( 6.72, N/A ) ( N/A, 0.00, N/A )	760.1	N/A	3.3111 [ 4.0000 ]	82.8% { 101.0% }			
13C2_8:2FTS_EIS	( 529.0 / 81.0 ) 692188	( 7.89, N/A ) ( N/A, 0.00, N/A )	799.8	N/A	3.8610 [ 4.0000 ]	96.5% { 114.7% }			
13C8_PFOsa_EIS	( 506.0 / 78.0 ) 7654815	( 9.84, N/A ) ( N/A, 0.00, N/A )	4530.1	N/A	1.4494 [ 2.0000 ]	72.5% { 92.1% }			
D3_NMeFOsa_EIS	( 515.0 / 169.0 ) 1191968	( 10.42, N/A ) ( N/A, 0.00, N/A )	2075.0	N/A	0.9579 [ 2.0000 ]	47.9% { 61.6% }			
D5_NEtFOsa_EIS	( 531.0 / 169.0 ) 1061267	( 10.56, N/A ) ( N/A, 0.00, N/A )	3052.8	N/A	0.9531 [ 2.0000 ]	47.7% { 59.5% }			

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 ) 1663563	( 8.29 , N/A ) ( N/A , 0.00 , N/A )	3470.3	N/A	3.6131 [ 4.0000 ]	90.3% { 125.6% }			
D5_EtFOSAA_EIS	( 589.0 / 419.0 ) 1792074	( 8.53 , N/A ) ( N/A , 0.00 , N/A )	627829.8	N/A	4.7534 [ 4.0000 ]	118.8% { 169.8% }			
D7_NMeFOSE_EIS	( 623.0 / 58.9 ) 3888993	( 10.36 , N/A ) ( N/A , 0.00 , N/A )	1403.2	N/A	8.5730 [ 20.0000 ]	42.9% { 52.4% }			
D9_NEtFOSE_EIS	( 639.0 / 58.9 ) 6903037	( 10.51 , N/A ) ( N/A , 0.00 , N/A )	1890.0	N/A	11.8826 [ 20.0000 ]	59.4% { 73.3% }			
13C3_HFPODA_EIS	( 287.0 / 169.0 ) 3572249	( 5.79 , N/A ) ( N/A , 0.00 , N/A )	3773.8	N/A	7.2274 [ 8.0000 ]	90.3% { 99.6% }			

# QUALITY CONTROL

# SURROGATE SUMMARY SHEET

EPA 1633

Client: AECOM  
 Work Order: 23B0091  
 Project: Red Hill AFFF Assessment Sampling

Surrogate Compound	Spike level	% Recovery	Recovery Limits	Q
AF-RHMW17-WGN01LF-2302W1 (23B0091-01) ng/L		Lab File ID: S2023-02-14D (20)		Analyzed: 02/14/23 23:47
13C4-PFBA	28.3	83.0	10 - 130	
13C5-PFPEA	14.1	91.0	35 - 150	
13C5-PFHXA	7.06	86.3	55 - 150	
13C4-PFHFA	7.06	81.7	55 - 150	
13C8-PFOA	7.06	82.7	60 - 140	
13C9-PFNA	3.53	81.7	55 - 140	
13C6-PFDA	3.53	84.3	50 - 140	
13C7-PFUnA	3.53	86.7	30 - 140	
13C2-PFDOA	3.53	91.2	10 - 150	
13C2-PFTEDA	3.53	80.2	10 - 130	
13C3-PFBS	7.06	84.9	55 - 150	
13C3-PFHXS	7.06	82.9	55 - 150	
13C8-PFOS	7.06	84.5	45 - 140	
13C2-4:2FTS	14.1	182	60 - 200	
13C2-6:2FTS	14.1	67.8	60 - 200	
13C2-8:2FTS	14.1	69.1	50 - 200	
13C8-PFOSA	7.06	77.8	30 - 130	
D3-NMEFOSA	7.06	63.6	15 - 130	
D5-NETFOSA	7.06	59.9	10 - 130	
D3-NMEFOSAA	14.1	72.9	45 - 200	
D5-NETFOSAA	14.1	81.4	10 - 200	
D7-NMEFOSE	70.6	52.6	10 - 150	
D9-NETFOSSE	70.6	70.8	10 - 150	
13C3-HFPO-DA	28.3	90.1	25 - 160	

## SURROGATE SUMMARY SHEET

EPA 1633

Client: AECOM  
 Work Order: 23B0091  
 Project: Red Hill AFFF Assessment Sampling

Surrogate Compound	Spike level	% Recovery	Recovery Limits	Q
<b>AF-RHMW17D-WGN01LF-2302W1 (23B0091-02 ) ng/L</b>		Lab File ID: S2023-02-14D (22)		Analyzed: 02/15/23 00:13
13C4-PFBA	28.1	81.8	10 - 130	
13C5-PFPEA	14.0	63.6	35 - 150	
13C5-PFHXA	7.02	76.2	55 - 150	
13C4-PFHFA	7.02	74.6	55 - 150	
13C8-PFOA	7.02	90.3	60 - 140	
13C9-PFNA	3.51	85.9	55 - 140	
13C6-PFDA	3.51	90.4	50 - 140	
13C7-PFUnA	3.51	152	30 - 140	*
13C2-PFDOA	3.51	126	10 - 150	
13C2-PFTEDA	3.51	129	10 - 130	
13C3-PFBS	7.02	80.9	55 - 150	
13C3-PFHXS	7.02	73.5	55 - 150	
13C8-PFOS	7.02	78.2	45 - 140	
13C2-4:2FTS	14.0	190	60 - 200	
13C2-6:2FTS	14.0	165	60 - 200	
13C2-8:2FTS	14.0	110	50 - 200	
13C8-PFOSA	7.02	67.0	30 - 130	
D3-NMEFOSA	7.02	52.1	15 - 130	
D5-NETFOSA	7.02	51.2	10 - 130	
D3-NMEFOSAA	14.0	63.8	45 - 200	
D5-NETFOSAA	14.0	99.3	10 - 200	
D7-NMEFOSE	70.2	41.3	10 - 150	
D9-NETFOSSE	70.2	59.1	10 - 150	
13C3-HFPO-DA	28.1	79.1	25 - 160	
<b>AF-RHMW17D-WGN01LF-2302W1 (23B0091-02RE1 ) ng/L</b>		Lab File ID: S2023-02-14D (23)		Analyzed: 02/15/23 00:26
13C7-PFUnA	3.51	71.8	30 - 140	

# SURROGATE SUMMARY SHEET

EPA 1633

Client: AECOM  
 Work Order: 23B0091  
 Project: Red Hill AFFF Assessment Sampling

Surrogate Compound	Spike level	% Recovery	Recovery Limits	Q
AF-RHMW17D-WQFB01-2302W1 (23B0091-03) . ng/L		Lab File ID: S2023-02-14D (24)		Analyzed: 02/15/23 00:39
13C4-PFBA	28.7	86.7	10 - 130	
13C5-PFPEA	14.4	95.5	35 - 150	
13C5-PFHXA	7.18	88.7	55 - 150	
13C4-PFHXA	7.18	88.4	55 - 150	
13C8-PFOA	7.18	88.2	60 - 140	
13C9-PFNA	3.59	89.7	55 - 140	
13C6-PFDA	3.59	79.7	50 - 140	
13C7-PFUnA	3.59	78.6	30 - 140	
13C2-PFDOA	3.59	93.2	10 - 150	
13C2-PFTEDA	3.59	90.7	10 - 130	
13C3-PFBS	7.18	89.6	55 - 150	
13C3-PFHXS	7.18	88.1	55 - 150	
13C8-PFOS	7.18	83.8	45 - 140	
13C2-4:2FTS	14.4	178	60 - 200	
13C2-6:2FTS	14.4	80.7	60 - 200	
13C2-8:2FTS	14.4	70.5	50 - 200	
13C8-PFOSA	7.18	72.7	30 - 130	
D3-NMEFOSA	7.18	54.7	15 - 130	
D5-NETFOSA	7.18	52.2	10 - 130	
D3-NMEFOSAA	14.4	69.1	45 - 200	
D5-NETFOSAA	14.4	69.7	10 - 200	
D7-NMEFOSE	71.8	45.2	10 - 150	
D9-NETFOSE	71.8	65.8	10 - 150	
13C3-HFPO-DA	28.7	89.8	25 - 160	

# SURROGATE SUMMARY SHEET

EPA 1633

Client: AECOM  
 Work Order: 23B0091  
 Project: Red Hill AFFF Assessment Sampling

Surrogate Compound	Spike level	% Recovery	Recovery Limits	Q
<b>AF-RHMW225401-WGN01B-2302W1 (23B0091-04) . ng/L</b>		Lab File ID: S2023-02-14D (26)		Analyzed: 02/15/23 01:04
13C4-PFBA	30.3	84.7	10 - 130	
13C5-PFPEA	15.2	96.5	35 - 150	
13C5-PFHXA	7.58	91.5	55 - 150	
13C4-PFHFA	7.58	90.0	55 - 150	
13C8-PFOA	7.58	91.8	60 - 140	
13C9-PFNA	3.79	94.9	55 - 140	
13C6-PFDA	3.79	87.1	50 - 140	
13C7-PFUnA	3.79	97.5	30 - 140	
13C2-PFDOA	3.79	95.2	10 - 150	
13C2-PFTEDA	3.79	82.0	10 - 130	
13C3-PFBS	7.58	96.3	55 - 150	
13C3-PFHXS	7.58	89.7	55 - 150	
13C8-PFOS	7.58	88.0	45 - 140	
13C2-4:2FTS	15.2	190	60 - 200	
13C2-6:2FTS	15.2	82.8	60 - 200	
13C2-8:2FTS	15.2	96.5	50 - 200	
13C8-PFOSA	7.58	72.5	30 - 130	
D3-NMEFOSA	7.58	47.9	15 - 130	
D5-NETFOSA	7.58	47.7	10 - 130	
D3-NMEFOSAA	15.2	90.3	45 - 200	
D5-NETFOSAA	15.2	119	10 - 200	
D7-NMEFOSE	75.8	42.9	10 - 150	
D9-NETFOSE	75.8	59.4	10 - 150	
13C3-HFPO-DA	30.3	90.3	25 - 160	

# SURROGATE SUMMARY SHEET

EPA 1633

Client: AECOM  
 Work Order: 23B0091  
 Project: Red Hill AFFF Assessment Sampling

Surrogate Compound	Spike level	% Recovery	Recovery Limits	Q
Blank (BCB0232-BLK1) ng/L		Lab File ID: S2023-02-14D (7)		Analyzed: 02/14/23 20:59
13C4-PFBA	32.0	80.4	10 - 130	
13C5-PFPEA	16.0	83.5	35 - 150	
13C5-PFHXA	8.00	74.1	55 - 150	
13C4-PFHFA	8.00	78.1	55 - 150	
13C8-PFOA	8.00	74.7	60 - 140	
13C9-PFNA	4.00	68.3	55 - 140	
13C6-PFDA	4.00	75.3	50 - 140	
13C7-PFUnA	4.00	75.9	30 - 140	
13C2-PFDOA	4.00	90.6	10 - 150	
13C2-PFTEDA	4.00	82.4	10 - 130	
13C3-PFBS	8.00	74.9	55 - 150	
13C3-PFHXS	8.00	85.0	55 - 150	
13C8-PFOS	8.00	74.4	45 - 140	
13C2-4:2FTS	16.0	181	60 - 200	
13C2-6:2FTS	16.0	68.3	60 - 200	
13C2-8:2FTS	16.0	70.3	50 - 200	
13C8-PFOSA	8.00	75.7	30 - 130	
D3-NMEFOSA	8.00	50.0	15 - 130	
D5-NETFOSA	8.00	53.1	10 - 130	
D3-NMEFOSAA	16.0	66.2	45 - 200	
D5-NETFOSAA	16.0	65.7	10 - 200	
D7-NMEFOSE	80.0	48.3	10 - 150	
D9-NETFOSE	80.0	76.1	10 - 150	
13C3-HFPO-DA	32.0	83.4	25 - 160	



# SURROGATE SUMMARY SHEET

EPA 1633

Client: AECOM  
 Work Order: 23B0091  
 Project: Red Hill AFFF Assessment Sampling

Surrogate Compound	Spike level	% Recovery	Recovery Limits	Q
LCS (BCB0232-BS1) . ng/L		Lab File ID: S2023-02-14D (8)		Analyzed: 02/14/23 21:12
13C4-PFBA	32.0	90.6	10 - 130	
13C5-PFPEA	16.0	92.4	35 - 150	
13C5-PFHXA	8.00	85.3	55 - 150	
13C4-PFHFA	8.00	89.4	55 - 150	
13C8-PFOA	8.00	89.1	60 - 140	
13C9-PFNA	4.00	85.8	55 - 140	
13C6-PFDA	4.00	79.2	50 - 140	
13C7-PFUnA	4.00	83.4	30 - 140	
13C2-PFDOA	4.00	83.8	10 - 150	
13C2-PFTEDA	4.00	76.1	10 - 130	
13C3-PFBS	8.00	84.5	55 - 150	
13C3-PFHXS	8.00	84.5	55 - 150	
13C8-PFOS	8.00	89.0	45 - 140	
13C2-4:2FTS	16.0	182	60 - 200	
13C2-6:2FTS	16.0	72.8	60 - 200	
13C2-8:2FTS	16.0	70.8	50 - 200	
13C8-PFOSA	8.00	85.1	30 - 130	
D3-NMEFOSA	8.00	58.6	15 - 130	
D5-NETFOSA	8.00	61.5	10 - 130	
D3-NMEFOSAA	16.0	75.6	45 - 200	
D5-NETFOSAA	16.0	77.8	10 - 200	
D7-NMEFOSE	80.0	52.1	10 - 150	
D9-NETFOSE	80.0	81.8	10 - 150	
13C3-HFPO-DA	32.0	90.8	25 - 160	

# SURROGATE SUMMARY SHEET

EPA 1633

Client: AECOM  
 Work Order: 23B0091  
 Project: Red Hill AFFF Assessment Sampling

Surrogate Compound	Spike level	% Recovery	Recovery Limits	Q
<b>MRL Check (BCB0232-MRL1) . ng/L</b>		Lab File ID: S2023-02-14D (9)		Analyzed: 02/14/23 21:25
13C4-PFBA	32.0	89.4	10 - 130	
13C5-PFPEA	16.0	97.7	35 - 150	
13C5-PFHXA	8.00	96.3	55 - 150	
13C4-PFHFA	8.00	90.8	55 - 150	
13C8-PFOA	8.00	84.0	60 - 140	
13C9-PFNA	4.00	87.4	55 - 140	
13C6-PFDA	4.00	89.8	50 - 140	
13C7-PFUnA	4.00	87.7	30 - 140	
13C2-PFDOA	4.00	86.8	10 - 150	
13C2-PFTEDA	4.00	88.1	10 - 130	
13C3-PFBS	8.00	85.4	55 - 150	
13C3-PFHXS	8.00	82.2	55 - 150	
13C8-PFOS	8.00	91.8	45 - 140	
13C2-4:2FTS	16.0	177	60 - 200	
13C2-6:2FTS	16.0	70.0	60 - 200	
13C2-8:2FTS	16.0	70.1	50 - 200	
13C8-PFOSA	8.00	89.0	30 - 130	
D3-NMEFOSA	8.00	63.4	15 - 130	
D5-NETFOSA	8.00	68.5	10 - 130	
D3-NMEFOSAA	16.0	83.5	45 - 200	
D5-NETFOSAA	16.0	88.3	10 - 200	
D7-NMEFOSE	80.0	54.1	10 - 150	
D9-NETFOSE	80.0	85.5	10 - 150	
13C3-HFPO-DA	32.0	96.5	25 - 160	

# METHOD BLANK SUMMARY

## EPA 1633

Laboratory:	APPL, LLC	Work Order:	23B0091
Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Blank ID:	BCB0232-BLK1	Batch:	BCB0232
		Prepared:	02/13/2023 07:19

Client Sample ID	Laboratory Sample ID	Lab File ID	Time Analyzed
LCS	BCB0232-BS1	S2023-02-14D (8)	21:12
MRL Check	BCB0232-MRL1	S2023-02-14D (9)	21:25
AF-RHMW17-WGN01LF-2302W1	23B0091-01	S2023-02-14D (20)	23:47
AF-RHMW17D-WGN01LF-2302W1	23B0091-02	S2023-02-14D (22)	00:13
DF10	23B0091-02RE1	S2023-02-14D (23)	00:26
AF-RHMW17D-WQFB01-2302W1	23B0091-03	S2023-02-14D (24)	00:39
AF-RHMW225401-WGN01B-2302W1	23B0091-04	S2023-02-14D (26)	01:04

# ANALYSIS DATA SHEET

Blank

Laboratory:	APPL, LLC	Work Order:	23B0091
Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Matrix:	Water	Laboratory ID:	BCB0232-BLK1
Sampled:		File ID:	S2023-02-14D (7)
		Prepared:	02/13/23 07:19
		Analyzed:	02/14/23 20:59
Solids:		Preparation:	EPA 1633
		Dilution:	1
Batch:	BCB0232	Sequence:	SC00636
		Calibration:	2307007
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	0.0870 J	0.40	0.20	0.055	J
PFHPA	0.20 U	0.40	0.20	0.041	U
PFOA	0.20 U	0.40	0.20	0.15	IR2, 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	0.112 J	0.40	0.20	0.064	MI5, 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:	23B0091
Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Matrix:	Water	Laboratory ID:	BCB0232-BLK1
		File ID:	S2023-02-14D (7)
Sampled:		Prepared:	02/13/23 07:19
		Analyzed:	02/14/23 20:59
Solids:		Preparation:	EPA 1633
		Dilution:	1
Batch:	BCB0232	Sequence:	SC00636
		Calibration:	2307007
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	IR2, 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: 23B0091

Client: AECOM

Project: Red Hill AFFF Assessment Sampling

Matrix: Water

Preparation: EPA 1633

Batch: BCB0232

Laboratory ID: BCB0232-BS1

Column:

ANALYTE	SPIKE ADDED (ng/L)	LCS CONCENTRATION (ng/L)	LCS % REC.	QC LIMITS REC.
PFBA	16.0	16.7	105	58 - 148
PFPEA	8.00	8.53	107	54 - 152
PFHXA	4.00	4.36	109	55 - 152
PFHPA	4.00	4.01	100	54 - 154
PFOA	4.00	4.30	108	52 - 161
PFNA	4.00	4.02	100	59 - 149
PFDA	4.00	4.62	116	52 - 147
PFUnA	4.00	4.11	103	48 - 159
PFDOA	4.00	3.92	98.1	64 - 142
PFTRDA	4.00	4.08	102	49 - 148
PFTEDA	4.00	4.34	108	47 - 161
PFBS	3.54	3.92	111	62 - 144
PFPEs	3.76	3.94	105	59 - 151
PFHXS	3.66	3.70	101	57 - 146
PFHPS	3.82	4.10	107	55 - 152
PFOS	3.72	3.80	102	58 - 149
PFNS	3.84	4.06	106	52 - 148
PFDS	3.86	3.98	103	51 - 147
PFDOS	3.88	3.84	98.9	36 - 145
4:2FTS	15.0	16.0	107	67 - 146
6:2FTS	15.2	18.0	119	61 - 151
8:2FTS	15.4	17.2	112	63 - 152
PFOSA	4.00	4.63	116	61 - 148
NMeFOSA	16.0	16.1	101	63 - 145
NEtFOSA	16.0	18.2	113	65 - 139
NMeFOSAA	4.00	4.43	111	58 - 144
NEtFOSAA	4.00	4.08	102	59 - 146
NMeFOSE	16.0	18.5	116	71 - 136
NEtFOSE	16.0	16.1	101	69 - 137
HFPO-DA	8.00	8.21	103	63 - 144
ADONA	7.56	7.95	105	68 - 146
PFEESA	7.12	8.03	113	56 - 151
PFMPA	8.00	8.03	100	51 - 145
PFMBA	8.00	7.76	96.9	55 - 148

**LCS / LCS DUPLICATE RECOVERY**

EPA 1633

Laboratory: APPL, LLC

Work Order: 23B0091

Client: AECOM

Project: Red Hill AFFF Assessment Sampling

Matrix: Water

Preparation: EPA 1633

Batch: BCB0232

Laboratory ID: BCB0232-BS1

Column:

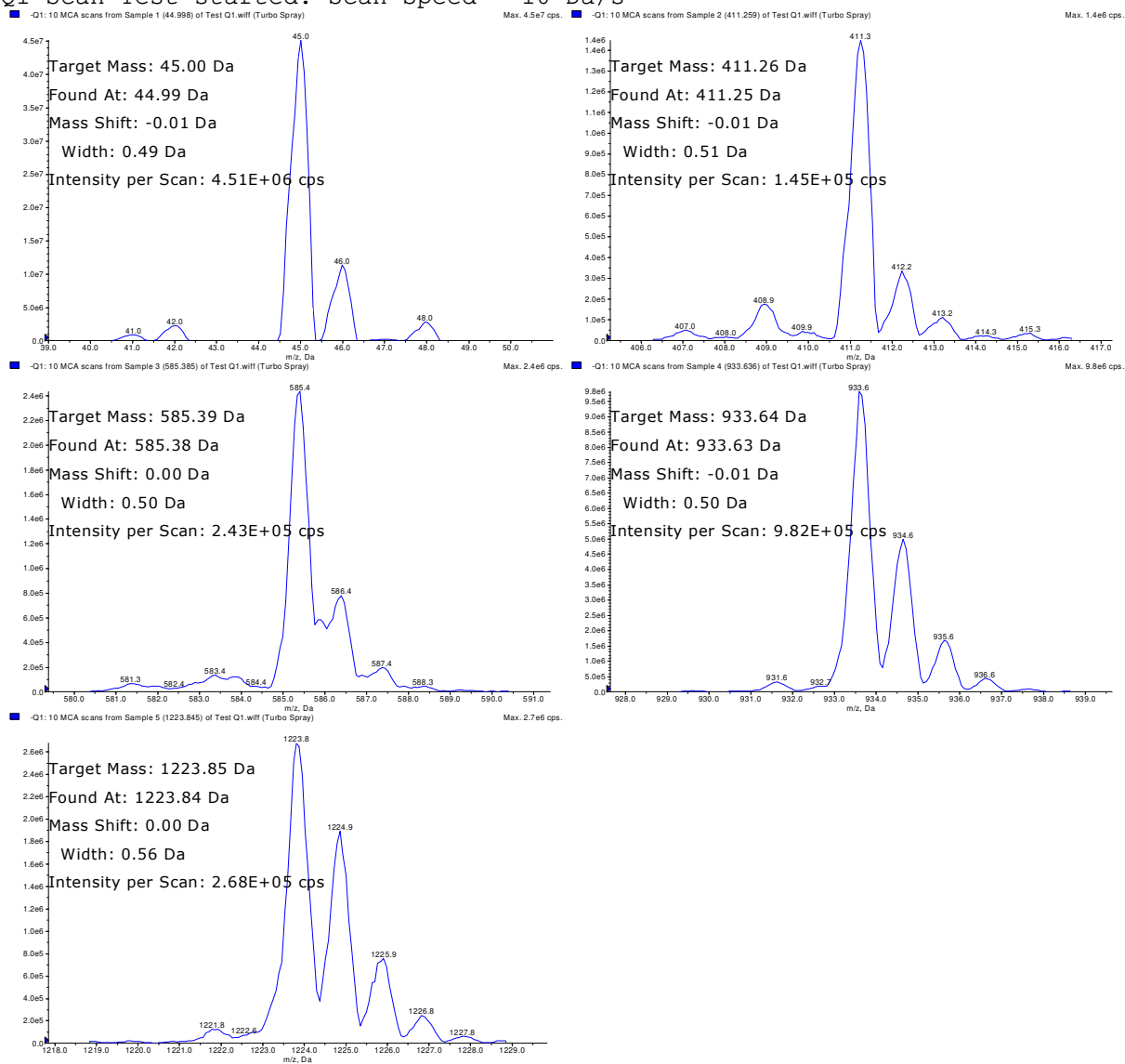
<b>ANALYTE</b>	<b>SPIKE ADDED (ng/L)</b>	<b>LCS CONCENTRATION (ng/L)</b>	<b>LCS % REC.</b>	<b>QC LIMITS REC.</b>
NFDHA	8.00	8.99	112	48 - 161
9CL-PF3ONS	7.48	7.63	102	56 - 156
11CL-PF3OUDS	7.56	7.70	102	46 - 156
3:3FTCA	16.0	16.9	106	62 - 129
5:3FTCA	16.0	17.6	110	63 - 134
7:3FTCA	16.0	16.0	99.8	50 - 138

# CALIBRATION SUMMARY



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

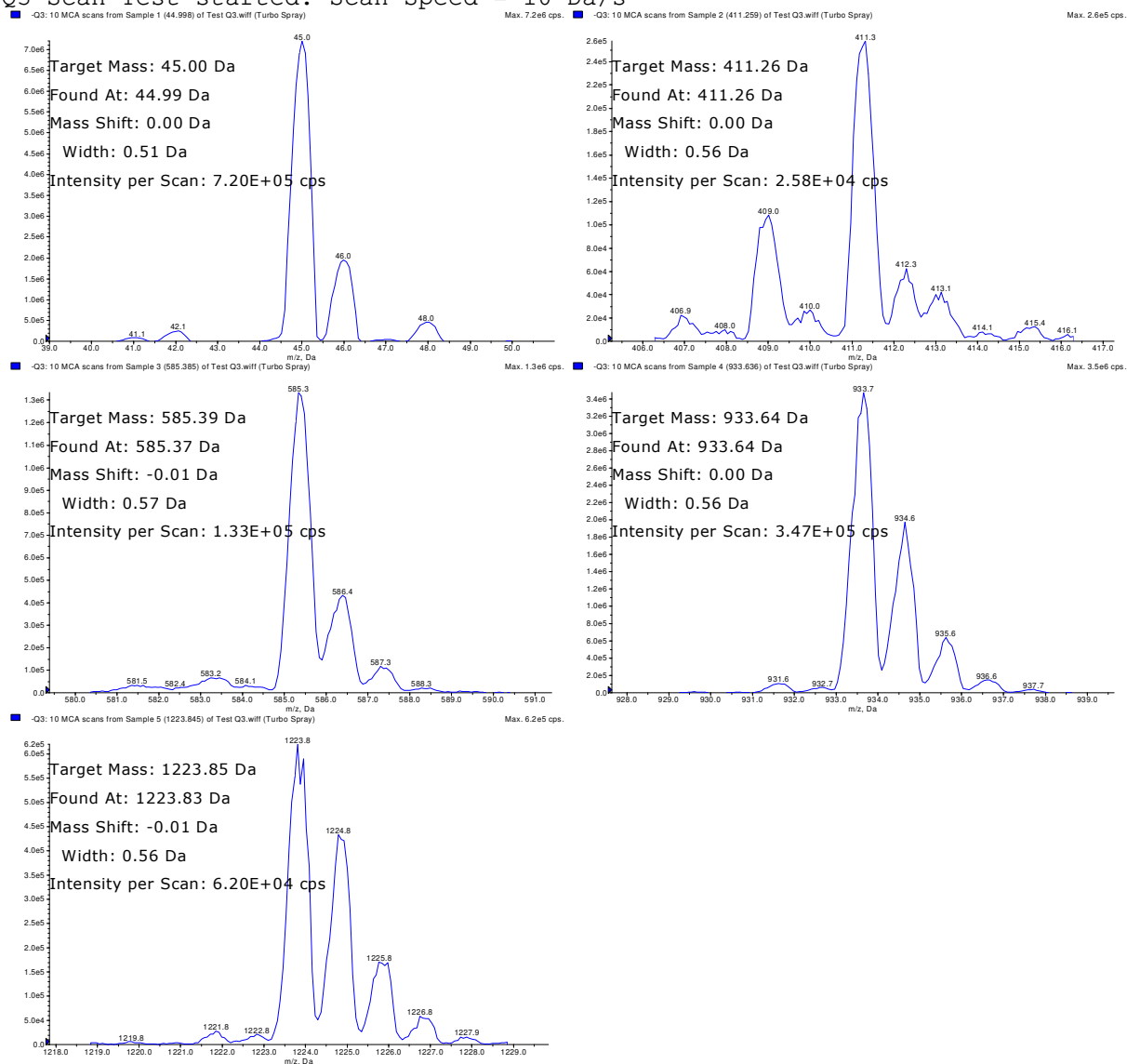
Q1 Scan Test started. Scan Speed = 10 Da/s



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

Analyte	( Q1 / Q3 )	Internal Standard	Multiplier	AcidFactor	Function	Qualifier
PFBA	( 213.0 / 169.0 )	13C4_PFBA_EIS	4.0000	1.0000	$y = 0.41417 x$ (std. dev. = 0.03642) (weighting: None)	%RSE=8.8
PFPeA	( 263.0 / 219.0 )	13C5_PFPeA_EIS	2.0000	1.0000	$y = 0.41137 x$ (std. dev. = 0.03924) (weighting: None)	%RSE=9.5
PFHxA	( 313.0 / 269.0 )	13C5_PFHxA_EIS	1.0000	1.0000	$y = 0.43957 x$ (std. dev. = 0.04840) (weighting: None)	%RSE=11.0
PFHpA	( 363.0 / 319.0 )	13C4_PFHpA_EIS	1.0000	1.0000	$y = 0.42221 x$ (std. dev. = 0.04136) (weighting: None)	%RSE=9.8
PFOA	( 413.0 / 369.0 )	13C8_PFOA_EIS	1.0000	1.0000	$y = 0.44519 x$ (std. dev. = 0.03337) (weighting: None)	%RSE=7.5
PFNA	( 463.0 / 419.0 )	13C9_PFNA_EIS	1.0000	1.0000	$y = 0.88477 x$ (std. dev. = 0.05485) (weighting: None)	%RSE=6.2
PFDA	( 513.0 / 469.0 )	13C6_PFDA_EIS	1.0000	1.0000	$y = 0.87964 x$ (std. dev. = 0.08735) (weighting: None)	%RSE=9.9
PFUnA	( 563.0 / 519.0 )	13C7_PFUnA_EIS	1.0000	1.0000	$y = 0.83084 x$ (std. dev. = 0.10366) (weighting: None)	%RSE=12.5
PFDaA	( 613.0 / 569.0 )	13C2_PFDaA_EIS	1.0000	1.0000	$y = 0.87937 x$ (std. dev. = 0.06795) (weighting: None)	%RSE=7.7
PFTTrDA	( 663.0 / 619.0 )	13C2_PFDaA_EIS	1.0000	1.0000	$y = 0.79619 x$ (std. dev. = 0.07839) (weighting: None)	%RSE=9.8
PFTeDA	( 713.0 / 669.0 )	13C2_PFTeDA_EIS	1.0000	1.0000	$y = 0.86535 x$ (std. dev. = 0.04299) (weighting: None)	%RSE=5.0
PFBS	( 299.0 / 80.0 )	13C3_PFBS_EIS	1.0000	0.8847	$y = 0.27491 x$ (std. dev. = 0.02287) (weighting: None)	%RSE=8.3
PFPeS	( 349.0 / 80.0 )	13C3_PFHxS_EIS	1.0000	0.9384	$y = 0.90486 x$ (std. dev. = 0.09986) (weighting: None)	%RSE=11.0
PFHxS	( 399.0 / 80.0 )	13C3_PFHxS_EIS	1.0000	0.9110	$y = 0.70041 x$ (std. dev. = 0.05642) (weighting: None)	%RSE=8.1
PFHpS	( 449.0 / 80.0 )	13C8_PFOS_EIS	1.0000	0.9514	$y = 0.36246 x$ (std. dev. = 0.04648) (weighting: None)	%RSE=12.8
PFOS	( 499.0 / 80.0 )	13C8_PFOS_EIS	1.0000	0.9275	$y = 0.48901 x$ (std. dev. = 0.04989) (weighting: None)	%RSE=10.2
PFNS	( 549.0 / 80.0 )	13C8_PFOS_EIS	1.0000	0.9599	$y = 0.56877 x$ (std. dev. = 0.05568) (weighting: None)	%RSE=9.8
PFDS	( 599.0 / 80.0 )	13C8_PFOS_EIS	1.0000	0.9631	$y = 0.62077 x$ (std. dev. = 0.05855) (weighting: None)	%RSE=9.4
PFDoS	( 699.0 / 80.0 )	13C8_PFOS_EIS	1.0000	0.9696	$y = 0.39858 x$ (std. dev. = 0.05576) (weighting: None)	%RSE=14.0
4:2FTS	( 327.0 / 307.0 )	13C2_4:2FTS_EIS	4.0000	0.9345	$y = 2.55237 x$ (std. dev. = 0.26579) (weighting: None)	%RSE=10.4
6:2FTS	( 427.0 / 407.0 )	13C2_6:2FTS_EIS	4.0000	0.9490	$y = 1.34660 x$ (std. dev. = 0.14334) (weighting: None)	%RSE=10.6
8:2FTS	( 527.0 / 507.0 )	13C2_8:2FTS_EIS	4.0000	0.9583	$y = 1.04382 x$ (std. dev. = 0.12314) (weighting: None)	%RSE=11.8
PFOSA	( 498.0 / 78.0 )	13C8_PFOSA_EIS	1.0000	1.0000	$y = 0.39991 x$ (std. dev. = 0.04179) (weighting: None)	%RSE=10.4
NMeFOSA	( 512.0 / 219.0 )	D3_NMeFOSA_EIS	4.0000	1.0000	$y = -0.01499 x^2 + 1.74034 x - 0.02935$ (r = 0.99892) (weighting: 1 / x)	%RSE=7.9
NEtFOSA	( 526.0 / 219.0 )	D5_NEtFOSA_EIS	4.0000	1.0000	$y = 1.86847 x$ (std. dev. = 0.19576) (weighting: None)	%RSE=10.5
NMeFOSAA	( 570.0 / 419.0 )	D3_MeFOSAA_EIS	1.0000	1.0000	$y = 0.19619 x$ (std. dev. = 0.01798) (weighting: None)	%RSE=9.2
NEtFOSAA	( 584.0 / 419.0 )	D5_EtFOSAA_EIS	1.0000	1.0000	$y = 0.20998 x$ (std. dev. = 0.01387) (weighting: None)	%RSE=6.6
NMeFOSE	( 616.0 / 59.0 )	D7_NMeFOSE_EIS	4.0000	1.0000	$y = 0.19995 x$ (std. dev. = 0.01319) (weighting: None)	%RSE=6.6
NEtFOSE	( 630.0 / 59.0 )	D9_NEtFOSE_EIS	4.0000	1.0000	$y = 0.19033 x$ (std. dev. = 0.01474) (weighting: None)	%RSE=7.7
HFPO-DA	( 285.0 / 169.0 )	13C3_HFPODA_EIS	2.0000	1.0000	$y = 0.19770 x$ (std. dev. = 0.01992) (weighting: None)	%RSE=10.1
ADONA	( 377.0 / 85.0 )	13C3_HFPODA_EIS	2.0000	0.9427	$y = 0.64094 x$ (std. dev. = 0.09352) (weighting: None)	%RSE=14.6
9Cl-Pf3ONS	( 531.0 / 351.0 )	13C3_HFPODA_EIS	2.0000	0.9333	$y = -0.04473 x^2 + 2.10957 x - 0.05721$ (r = 0.99774) (weighting: 1 / x)	%RSE=10.3
11Cl-Pf3OUDS	( 631.0 / 451.0 )	13C3_HFPODA_EIS	2.0000	0.9432	$y = 1.04469 x$ (std. dev. = 0.11673) (weighting: None)	%RSE=11.2
3:3FTCA	( 241.0 / 177.0 )	13C5_PFPeA_EIS	4.0000	1.0000	$y = 1.86535e-4 x^2 + 0.01946 x - 4.63779e-5$ (r = 0.99934) (weighting: 1 / x)	%RSE=7.1
5:3FTCA	( 341.0 / 236.7 )	13C5_PFHxA_EIS	4.0000	1.0000	$y = 0.24044 x$ (std. dev. = 0.03492) (weighting: None)	%RSE=14.5
7:3FTCA	( 441.0 / 317.0 )	13C5_PFHxA_EIS	4.0000	1.0000	$y = 0.44417 x$ (std. dev. = 0.05674) (weighting: None)	%RSE=12.8
PFEESA	( 315.0 / 135.0 )	13C5_PFHxA_EIS	2.0000	0.8925	$y = 0.94250 x$ (std. dev. = 0.11701) (weighting: None)	%RSE=12.4
PFMPA	( 229.0 / 85.0 )	13C5_PFPeA_EIS	2.0000	1.0000	$y = 0.08985 x$ (std. dev. = 0.01015) (weighting: None)	%RSE=11.3
PFMBA	( 279.0 / 85.0 )	13C5_PFPeA_EIS	2.0000	1.0000	$y = 0.37681 x$ (std. dev. = 0.03401) (weighting: None)	%RSE=9.0
NFDHA	( 295.0 / 201.0 )	13C5_PFHxA_EIS	2.0000	1.0000	$y = 0.48796 x$ (std. dev. = 0.05376) (weighting: None)	%RSE=11.0
13C3_PFBA_IIS	( 216.0 / 172.0 )	13C3_PFBA_IIS	1.0000	1.0000	$y = 223990.8344 x$	%RSD=15.7
13C2_PFHxA_IIS	( 315.0 / 270.0 )	13C2_PFHxA_IIS	1.0000	1.0000	$y = 504804.2832 x$	%RSD=8.3
13C4_PFOA_IIS	( 417.0 / 372.0 )	13C4_PFOA_IIS	1.0000	1.0000	$y = 633776.3615 x$	%RSD=10.0
13C5_PFNA_IIS	( 468.0 / 423.0 )	13C5_PFNA_IIS	1.0000	1.0000	$y = 604267.6924 x$	%RSD=7.7

Analyte	( Q1 / Q3 )	Internal Standard	Multiplier	AcidFactor	Function	Qualifier
13C2_PFDA_IIS	( 515.0 / 470.1 )	13C2_PFDA_IIS	1.0000	1.0000	y = 578638.3434 x	%RSD=6.0
18O2_PFHxS_IIS	( 403.0 / 83.9 )	18O2_PFHxS_IIS	1.0000	1.0000	y = 856363.7448 x	%RSD=5.6
13C4_PFOS_IIS	( 503.0 / 79.9 )	13C4_PFOS_IIS	1.0000	1.0000	y = 1505389.7322 x	%RSD=6.7
13C4_PFBA_EIS	( 217.0 / 172.0 )	13C3_PFBA_IIS	8.0000	1.0000	y = 9.5160 x	%RSD=6.0
13C5_PFPeA_EIS	( 268.0 / 223.0 )	13C2_PFHxA_IIS	4.0000	1.0000	y = 4.3765 x	%RSD=12.0
13C5_PFHxA_EIS	( 318.0 / 273.0 )	13C2_PFHxA_IIS	2.0000	1.0000	y = 2.7395 x	%RSD=10.4
13C4_PFHpA_EIS	( 367.0 / 322.0 )	13C2_PFHxA_IIS	2.0000	1.0000	y = 2.3736 x	%RSD=9.3
13C8_PFOA_EIS	( 421.0 / 376.0 )	13C4_PFOA_IIS	2.0000	1.0000	y = 2.2651 x	%RSD=10.4
13C9_PFNA_EIS	( 472.0 / 427.0 )	13C5_PFNA_IIS	1.0000	1.0000	y = 1.0595 x	%RSD=8.0
13C6_PFDA_EIS	( 519.0 / 474.0 )	13C2_PFDA_IIS	1.0000	1.0000	y = 1.2608 x	%RSD=10.3
13C7_PFUnA_EIS	( 570.0 / 525.0 )	13C2_PFDA_IIS	1.0000	1.0000	y = 1.2538 x	%RSD=12.9
13C2_PFDaA_EIS	( 615.0 / 570.0 )	13C2_PFDA_IIS	1.0000	1.0000	y = 1.0753 x	%RSD=12.1
13C2_PFTeDA_EIS	( 715.0 / 670.0 )	13C2_PFDA_IIS	1.0000	1.0000	y = 0.9853 x	%RSD=8.1
13C3_PFBS_EIS	( 302.0 / 80.0 )	18O2_PFHxS_IIS	2.0000	1.0000	y = 4.1858 x	%RSD=8.8
13C3_PFHxS_EIS	( 402.0 / 80.0 )	18O2_PFHxS_IIS	2.0000	1.0000	y = 2.1742 x	%RSD=6.1
13C8_PFOS_EIS	( 507.0 / 80.0 )	13C4_PFOS_IIS	2.0000	1.0000	y = 2.7457 x	%RSD=8.1
13C2_4:2FTS_EIS	( 329.0 / 81.0 )	18O2_PFHxS_IIS	4.0000	1.0000	y = 0.4621 x	%RSD=6.6
13C2_6:2FTS_EIS	( 429.0 / 81.0 )	18O2_PFHxS_IIS	4.0000	1.0000	y = 0.5072 x	%RSD=8.2
13C2_8:2FTS_EIS	( 529.0 / 81.0 )	18O2_PFHxS_IIS	4.0000	1.0000	y = 0.7166 x	%RSD=12.2
13C8_PFOSA_EIS	( 506.0 / 78.0 )	13C4_PFOS_IIS	2.0000	1.0000	y = 5.0967 x	%RSD=12.6
D3_NMeFOSA_EIS	( 515.0 / 169.0 )	13C4_PFOS_IIS	2.0000	1.0000	y = 1.2008 x	%RSD=7.5
D5_NEtFOSA_EIS	( 531.0 / 169.0 )	13C4_PFOS_IIS	2.0000	1.0000	y = 1.0745 x	%RSD=15.8
D3_MeFOSAA_EIS	( 573.0 / 419.0 )	13C4_PFOS_IIS	4.0000	1.0000	y = 0.8887 x	%RSD=7.4
D5_EtFOSAA_EIS	( 589.0 / 419.0 )	13C4_PFOS_IIS	4.0000	1.0000	y = 0.7277 x	%RSD=6.9
D7_NMeFOSE_EIS	( 623.0 / 58.9 )	13C4_PFOS_IIS	20.0000	1.0000	y = 4.3778 x	%RSD=11.4
D9_NEtFOSE_EIS	( 639.0 / 58.9 )	13C4_PFOS_IIS	20.0000	1.0000	y = 5.6063 x	%RSD=15.4
13C3_HFPODA_EIS	( 287.0 / 169.0 )	13C2_PFHxA_IIS	8.0000	1.0000	y = 6.6719 x	%RSD=9.1

x=Concentration Analyte

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

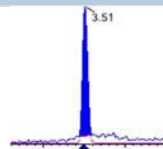
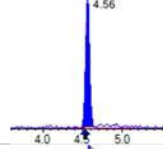
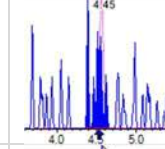
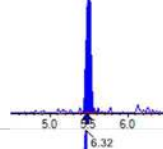
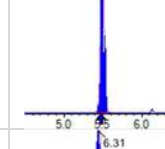
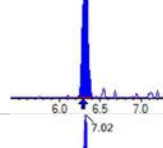
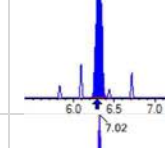
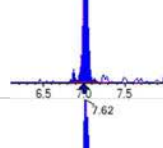
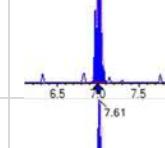
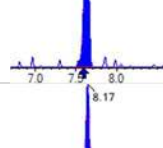
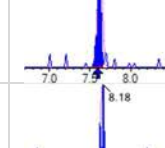
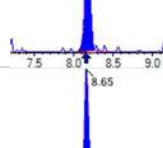
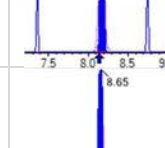
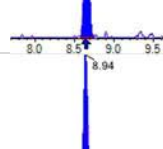
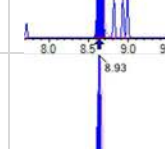
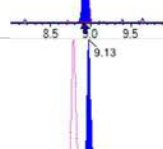
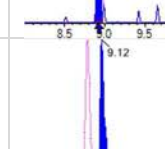
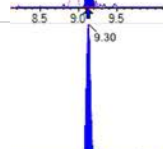
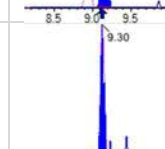
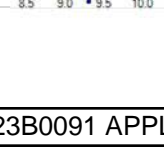
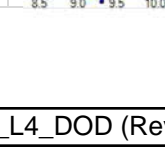
$$\text{Acid Factor} = \frac{\text{Molecular weight Acid}}{\text{Molecular weight Salt}}$$

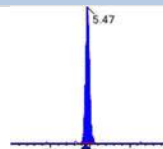
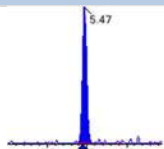
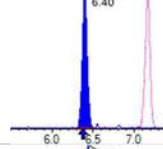
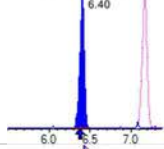
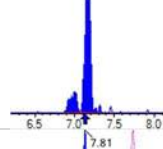
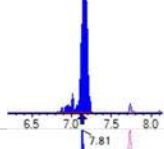
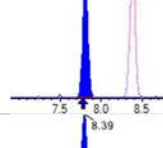
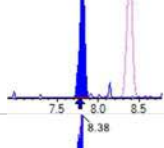
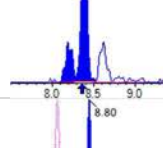
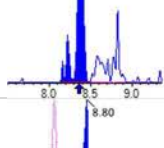
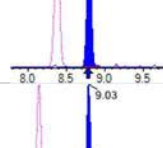
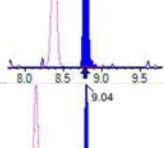
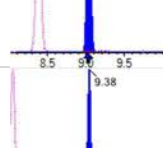
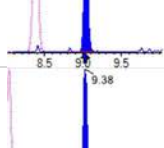
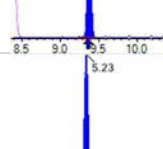
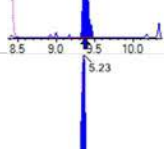
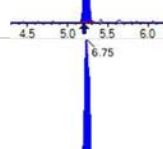
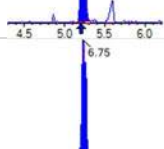
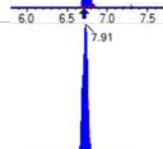
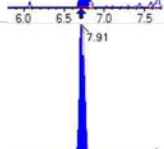


$$\text{Multiplier} = \frac{\text{Concentration of Analyte}}{\text{Concentration of PFOA}} \text{ in curve standard mix}$$

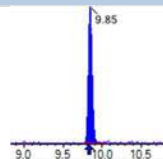
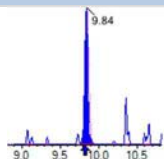
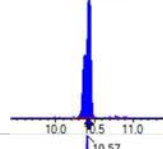
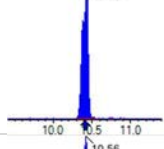
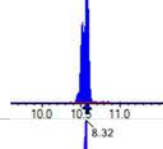
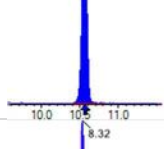
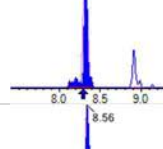
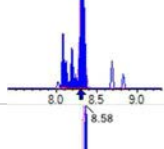
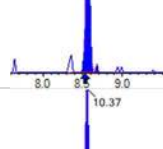
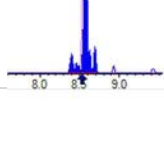
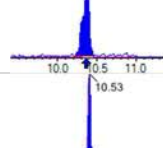
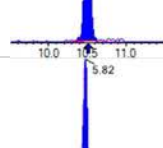
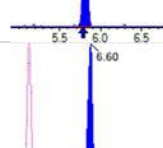
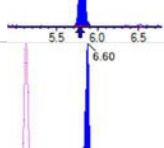
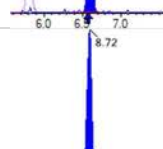
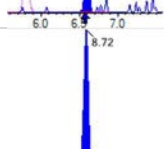
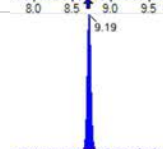
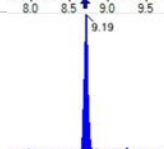
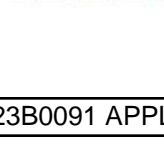
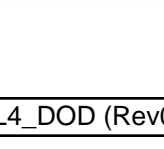
$$\text{Result} \left( \frac{\text{ng}}{\text{mL}} \right) = x * \text{Multiplier} * \text{Acid Factor}$$

# EPA 1633

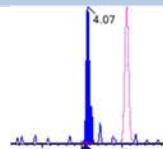
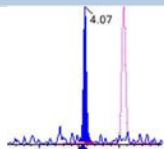
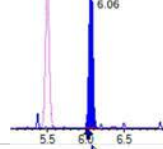
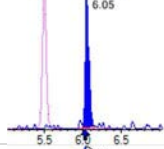
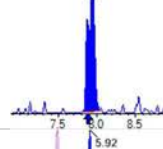
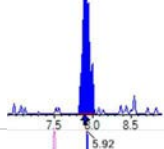
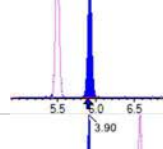
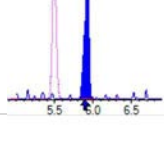
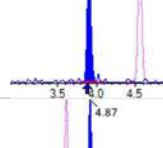
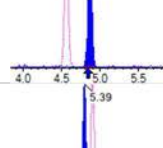
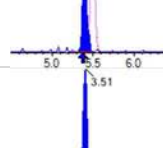
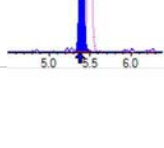
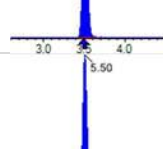
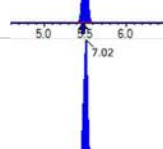
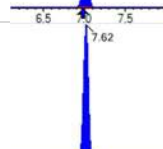

Initial Calibration: SC00647

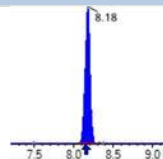
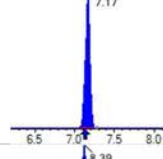
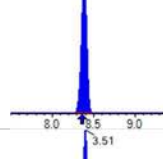
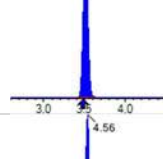
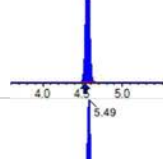
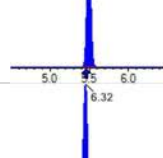
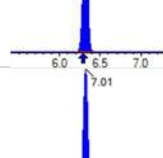
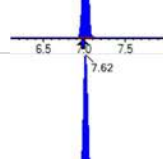
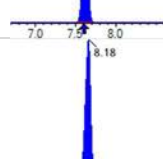
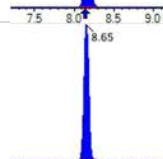
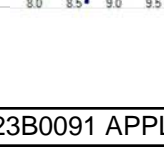
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-Imin, ΔRT-CVmin, ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration True ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	(213.0 / 169.0) 89209	(3.51, 1.00) (0.00, N/A, 0.0)	76.9	N/A 0.0 0.0	0.3517 [0.4000]	87.9%			
PFPeA	(263.0 / 219.0) 89024 (263.0 / 69.0) 1335	(4.56, 1.00) (0.00, N/A, 6.9)	310.9 5964.0	0.0150 140.0 140.0	0.1743 [0.2000]	87.2%			
PFHxA	(313.0 / 269.0) 59236 (313.0 / 119.0) 7304	(5.50, 1.00) (0.00, N/A, 0.5)	372.7 42522.7	0.1233 123.8 123.8	0.0903 [0.1000]	90.3%			
PFHpA	(363.0 / 319.0) 51122 (363.0 / 169.0) 17342	(6.32, 1.00) (0.00, N/A, 0.3)	55898.0 812.6	0.3392 105.8 105.8	0.0951 [0.1000]	95.1%			
PFOA	(413.0 / 369.0) 70061 (413.0 / 169.0) 21539	(7.02, 1.00) (0.00, N/A, 0.1)	656.9 1399.5	0.3074 91.9 91.9	0.1047 [0.1000]	104.7%			
PFNA	(463.0 / 419.0) 54818 (463.0 / 169.0) 11781	(7.62, 1.00) (0.00, N/A, 0.7)	5330.4 644.4	0.2149 96.0 96.0	0.0937 [0.1000]	93.7%			
PFDA	(513.0 / 469.0) 56800 (513.0 / 169.0) 4296	(8.17, 1.00) (0.00, N/A, -0.7)	157.8 6379.1	0.0756 62.2 62.2	0.0887 [0.1000]	88.7%			
PFUnA	(563.0 / 519.0) 60153 (563.0 / 169.0) 5180	(8.65, 1.00) (0.00, N/A, 0.4)	229.8 793.8	0.0861 82.1 82.1	0.0873 [0.1000]	87.3%			
PFDaA	(613.0 / 569.0) 63649 (613.0 / 169.0) 8537	(8.94, 1.00) (0.00, N/A, 0.4)	581.7 121.0	0.1341 82.0 82.0	0.0963 [0.1000]	96.3%			
PFTTrDA	(663.0 / 619.0) 62304 (663.0 / 169.0) 19614	(9.13, 1.02) (N/A, 0.01, 0.6)	434.9 11588627.4	0.3148 131.7 131.7	0.1041 [0.1000]	104.1%			
PFTeDA	(713.0 / 669.0) 49918 (713.0 / 169.0) 13191	(9.30, 1.00) (0.00, N/A, 0.1)	460.4 598.4	0.2643 127.9 127.9	0.0964 [0.1000]	96.4%			

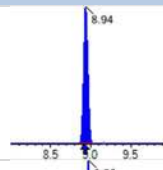
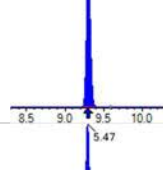
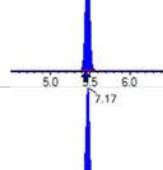
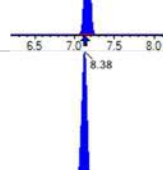
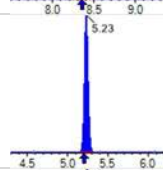
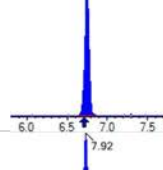
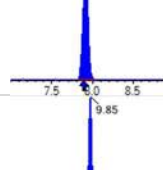
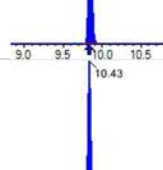
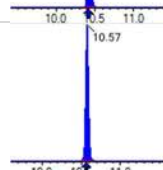
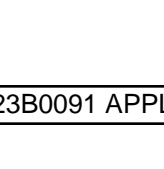
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-Imin, ΔRT-CVmin, ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration True ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) 104864 (299.0 / 99.0) 61681	(5.47, 1.00) (0.00, N/A, 0.0)	937.2 202.1	0.5882 91.0 91.0	0.0900 [0.0885]	101.8%			
PFPeS	(349.0 / 80.0) 165326 (349.0 / 99.0) 57545	(6.40, 0.89) (N/A, 0.03, -0.4)	1188.9 210907.1	0.3481 97.5 97.5	0.0881 [0.0938]	93.9%			
PFHxS	(399.0 / 80.0) 142609 (399.0 / 99.0) 51178	(7.17, 1.00) (0.00, N/A, 0.2)	2284.2 5030.8	0.3589 110.5 110.5	0.0953 [0.0911]	104.6%			
PFHpS	(449.0 / 80.0) 143003 (449.0 / 99.0) 48860	(7.81, 0.93) (N/A, 0.03, -0.1)	1479.4 20139600.5	0.3417 125.5 125.5	0.0859 [0.0951]	90.3%			
PFOS	(499.0 / 80.0) 240900 (499.0 / 99.0) 55675	(8.39, 1.00) (0.00, N/A, 0.4)	144.5 95.1	0.2311 104.4 104.4	0.1046 [0.0927]	112.7%			
PFNS	(549.0 / 80.0) 267881 (549.0 / 99.0) 72412	(8.80, 1.05) (N/A, 0.02, 0.1)	2678.9 730.3	0.2703 123.3 123.3	0.1035 [0.0960]	107.8%			
PFDS	(599.0 / 80.0) 266169 (599.0 / 99.0) 60916	(9.03, 1.08) (N/A, 0.01, -0.5)	821.7 424.5	0.2289 105.0 105.0	0.0945 [0.0963]	98.1%			
PFDoS	(699.0 / 80.0) 164788 (699.0 / 99.0) 41858	(9.38, 1.12) (N/A, 0.02, 0.2)	625.6 256.9	0.2540 122.0 122.0	0.0917 [0.0970]	94.6%			
4:2FTS	(327.0 / 307.0) 100994 (327.0 / 81.0) 64806	(5.23, 1.00) (0.00, N/A, -0.2)	587.5 260.2	0.6417 95.1 95.1	0.3649 [0.3738]	97.6%			
6:2FTS	(427.0 / 407.0) 52063 (427.0 / 81.0) 50004	(6.75, 1.00) (0.00, N/A, 0.0)	1057.7 255.3	0.9604 116.0 116.0	0.3227 [0.3796]	85.0%			
8:2FTS	(527.0 / 507.0) 54092 (527.0 / 81.0) 51796	(7.91, 1.00) (0.00, N/A, 0.5)	22137206.7 343.8	0.9576 115.2 115.2	0.2983 [0.3833]	77.8%			

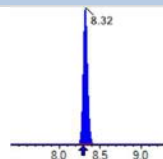
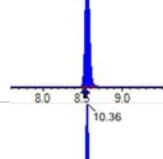
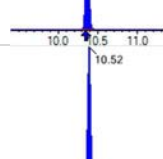
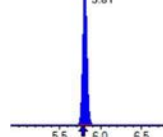

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-Imin, ΔRT-CVmin, ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration True ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) 326059 (498.0 / 478.0) 6395	(9.85, 1.00) (0.00, N/A, 0.5)	435.1 92.9	0.0196 95.5 95.5	0.0981 [0.1000]	98.1%			
NMeFOSA	(512.0 / 219.0) 311700 (512.0 / 169.0) 251496	(10.43, 1.00) (0.00, N/A, 0.8)	910.7 988.1	0.8069 98.5 98.5	0.4355 [0.4000]	108.9%			
NEiFOSA	(526.0 / 219.0) 325495 (526.0 / 169.0) 402906	(10.57, 1.00) (0.00, N/A, 0.8)	1489.9 1292.0	1.2378 97.5 97.5	0.3950 [0.4000]	98.7%			
NMeFOSAA	(570.0 / 419.0) 31281 (570.0 / 483.0) 12615	(8.32, 1.00) (0.01, N/A, 0.0)	81874.8 66.4	0.4033 85.0 85.0	0.1066 [0.1000]	106.6%			
NEiFOSAA	(584.0 / 419.0) 23406 (584.0 / 526.0) 18199	(8.56, 1.00) (0.01, N/A, -0.8)	455204.0 175.6	0.7775 152.4 152.4	0.0995 [0.1000]	99.5%			IR2,
NMeFOSE	(616.0 / 59.0) 145674	(10.37, 1.00) (0.01, N/A, 0.0)	321.7	N/A 0.0 0.0	0.4009 [0.4000]	100.2%			
NEiFOSE	(630.0 / 59.0) 156006	(10.53, 1.00) (0.01, N/A, 0.0)	279.4	N/A 0.0 0.0	0.4008 [0.4000]	100.2%			
HFPO-DA	(285.0 / 169.0) 66037 (285.0 / 185.0) 177079	(5.82, 1.00) (0.01, N/A, 0.3)	708.1 472.4	2.6815 97.4 97.4	0.1832 [0.2000]	91.6%			
ADONA	(377.0 / 85.0) 224483 (377.0 / 251.0) 22053	(6.60, 1.14) (N/A, 0.03, 0.0)	373.1 122.9	0.0982 111.4 111.4	0.1811 [0.1885]	96.0%			
9CI-PF3ONS	(531.0 / 351.0) 629773 (533.0 / 353.0) 194028	(8.72, 1.50) (N/A, 0.03, 0.2)	557.7 584.4	0.3081 88.5 88.5	0.2039 [0.1867]	109.2%			
11CI-PF3OUDS	(631.0 / 451.0) 362238 (633.0 / 453.0) 128996	(9.19, 1.58) (N/A, 0.02, -0.1)	784.7 5104.8	0.3561 98.4 98.4	0.1793 [0.1886]	95.1%			

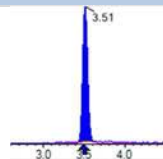
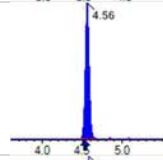
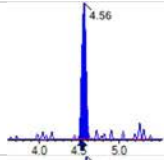
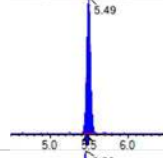
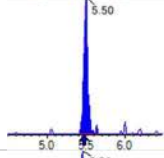
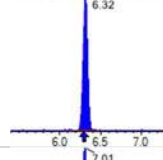
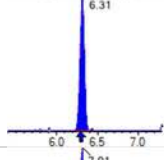
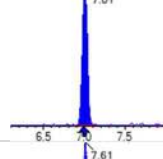
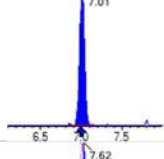
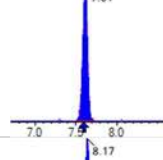
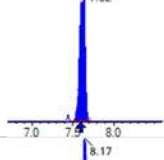
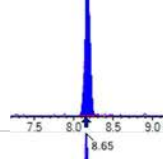
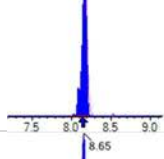
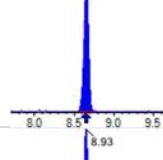
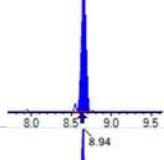
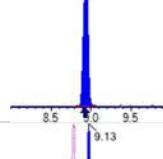
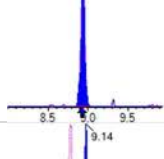
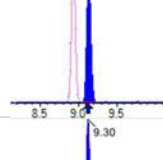
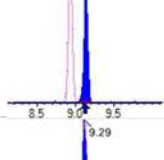
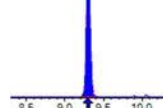
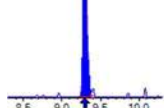


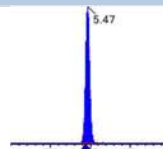
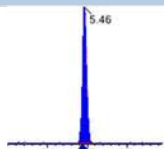
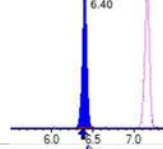
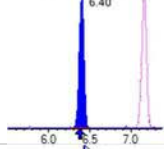
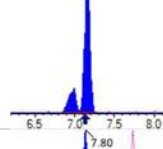
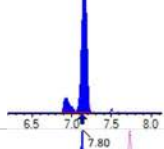
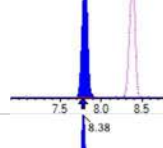
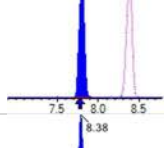
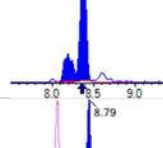
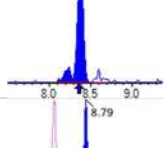
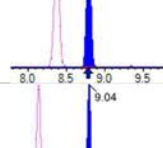
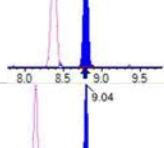
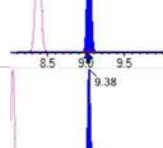
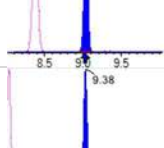
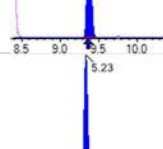
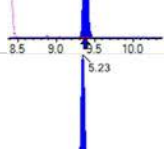
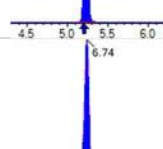
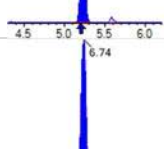
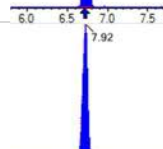
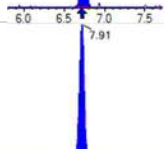


Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-Imin, ΔRT-CVmin, ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration True ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	(241.0 / 177.0) 4574 (241.0 / 117.0) 6911	(4.07, 0.89) (N/A, 0.03, -0.1)	137.7 66.2	1.5109 95.7 95.7	0.3879 [0.4000]	97.0%			
5:3FTCA	(341.0 / 236.7) 36025 (341.0 / 217.0) 50605	(6.06, 1.10) (N/A, 0.04, 0.9)	289.4 119.0	1.4047 82.4 82.4	0.4014 [0.4000]	100.3%			
7:3FTCA	(441.0 / 317.0) 64735 (441.0 / 337.0) 54368	(7.93, 1.44) (N/A, 0.05, 0.7)	93.1 99.2	0.8399 97.6 97.6	0.3904 [0.4000]	97.6%			
PFEESA	(315.0 / 135.0) 130937 (315.0 / 83.0) 37471	(5.92, 1.08) (N/A, 0.03, -0.2)	767.3 153.5	0.2862 112.0 112.0	0.1661 [0.1785]	93.0%			
PFMPA	(229.0 / 85.0) 18495	(3.90, 0.86) (N/A, 0.02, 0.0)	183.4	N/A 0.0 0.0	0.1658 [0.2000]	82.9%			
PFMBA	(279.0 / 85.0) 92630	(4.87, 1.07) (N/A, 0.03, 0.0)	415.5	N/A 0.0 0.0	0.1980 [0.2000]	99.0%			
NFDHA	(295.0 / 201.0) 67723 (295.0 / 85.0) 57064	(5.39, 0.98) (N/A, 0.02, -0.5)	301.0 266.7	0.8426 84.2 84.2	0.1859 [0.2000]	93.0%			
13C3_PFBA_IIS	(216.0 / 172.0) 254010	(3.51, N/A) (N/A, 0.02, N/A)	1726.7	N/A	1.1340 [1.0000]	113.4% {112.7%}			
13C2_PFHxA_IIS	(315.0 / 270.0) 577490	(5.50, N/A) (N/A, 0.03, N/A)	3009.2	N/A	1.1440 [1.0000]	114.4% {120.3%}			
13C4_PFOA_IIS	(417.0 / 372.0) 690977	(7.02, N/A) (N/A, 0.04, N/A)	1738.7	N/A	1.0903 [1.0000]	109.0% {116.4%}			
13C5_PFNA_IIS	(468.0 / 423.0) 636163	(7.62, N/A) (N/A, 0.03, N/A)	4733.4	N/A	1.0528 [1.0000]	105.3% {107.3%}			

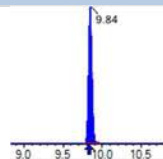
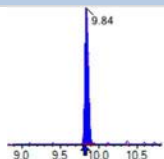
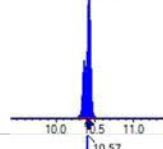
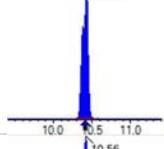
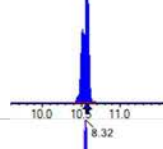
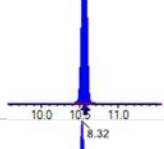
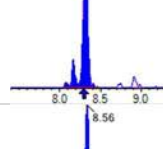
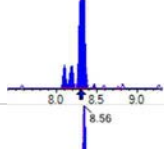
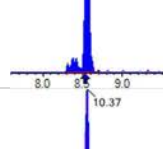
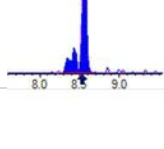
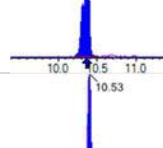
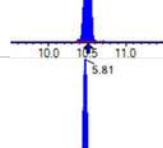
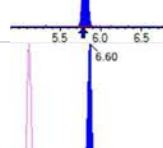
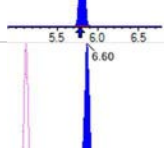
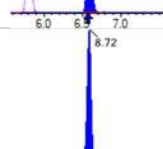
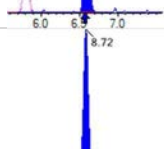
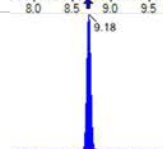
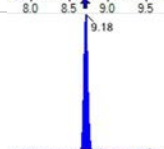
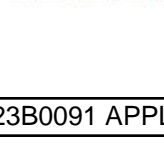
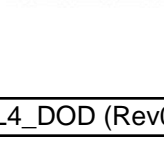
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-Imin, ΔRT-CVmin, ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration True ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	(515.0 / 470.1) 593318	(8.18, N/A) (N/A, 0.03, N/A)	1103.0	N/A	1.0254 [1.0000]	102.5% {109.0%}			
18O2_PFHxS_IIS	(403.0 / 83.9) 870629	(7.17, N/A) (N/A, 0.04, N/A)	3746.4	N/A	1.0167 [1.0000]	101.7% {103.1%}			
13C4_PFOS_IIS	(503.0 / 79.9) 1609689	(8.39, N/A) (N/A, 0.03, N/A)	1109.3	N/A	1.0693 [1.0000]	106.9% {106.5%}			
13C4_PFBA_EIS	(217.0 / 172.0) 2449883	(3.51, N/A) (N/A, 0.02, N/A)	4854.2	N/A	8.1084 [8.0000]	101.4% {104.9%}			
13C5_PFPeA_EIS	(268.0 / 223.0) 2482651	(4.56, N/A) (N/A, 0.03, N/A)	4201.2	N/A	3.9292 [4.0000]	98.2% {102.6%}			
13C5_PFHxA_EIS	(318.0 / 273.0) 1493141	(5.49, N/A) (N/A, 0.03, N/A)	4148.5	N/A	1.8876 [2.0000]	94.4% {97.3%}			
13C4_PFHpA_EIS	(367.0 / 322.0) 1272791	(6.32, N/A) (N/A, 0.03, N/A)	2795.4	N/A	1.8571 [2.0000]	92.9% {100.7%}			
13C8_PFOA_EIS	(421.0 / 376.0) 1503687	(7.01, N/A) (N/A, 0.03, N/A)	2224.2	N/A	1.9215 [2.0000]	96.1% {97.6%}			
13C9_PFNA_EIS	(472.0 / 427.0) 661416	(7.62, N/A) (N/A, 0.03, N/A)	1339.6	N/A	0.9813 [1.0000]	98.1% {94.1%}			
13C6_PFDA_EIS	(519.0 / 474.0) 728048	(8.18, N/A) (N/A, 0.03, N/A)	2745.6	N/A	0.9732 [1.0000]	97.3% {93.7%}			
13C7_PFUnA_EIS	(570.0 / 525.0) 829502	(8.65, N/A) (N/A, 0.02, N/A)	2125.0	N/A	1.1151 [1.0000]	111.5% {107.5%}			

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_EIS	( 615.0 / 570.0 ) 751841	( 8.94 , N/A ) ( N/A , 0.02 , N/A )	1472.4	N/A	1.1784 [ 1.0000 ]	117.8% { 114.5% }			
13C2_PFTeDA_EIS	( 715.0 / 670.0 ) 598593	( 9.30 , N/A ) ( N/A , 0.01 , N/A )	1534.2	N/A	1.0239 [ 1.0000 ]	102.4% { 98.1% }			
13C3_PFBs_EIS	( 302.0 / 80.0 ) 3748603	( 5.47 , N/A ) ( N/A , 0.03 , N/A )	3158.7	N/A	2.0572 [ 2.0000 ]	102.9% { 94.9% }			
13C3_PFHxS_EIS	( 402.0 / 80.0 ) 1946389	( 7.17 , N/A ) ( N/A , 0.03 , N/A )	2265.9	N/A	2.0565 [ 2.0000 ]	102.8% { 97.8% }			
13C8_PFOS_EIS	( 507.0 / 80.0 ) 4369903	( 8.38 , N/A ) ( N/A , 0.03 , N/A )	2552.8	N/A	1.9775 [ 2.0000 ]	98.9% { 98.9% }			
13C2_4:2FTS_EIS	( 329.0 / 81.0 ) 405361	( 5.23 , N/A ) ( N/A , 0.03 , N/A )	1641.7	N/A	4.0303 [ 4.0000 ]	100.8% { 99.0% }			
13C2_6:2FTS_EIS	( 429.0 / 81.0 ) 454861	( 6.75 , N/A ) ( N/A , 0.04 , N/A )	1114.4	N/A	4.1199 [ 4.0000 ]	103.0% { 99.3% }			
13C2_8:2FTS_EIS	( 529.0 / 81.0 ) 665899	( 7.92 , N/A ) ( N/A , 0.03 , N/A )	1555.4	N/A	4.2691 [ 4.0000 ]	106.7% { 110.1% }			
13C8_PFOsa_EIS	( 506.0 / 78.0 ) 8308651	( 9.85 , N/A ) ( N/A , 0.02 , N/A )	3294.2	N/A	2.0255 [ 2.0000 ]	101.3% { 96.9% }			
D3_NMeFOsa_EIS	( 515.0 / 169.0 ) 1948846	( 10.43 , N/A ) ( N/A , 0.02 , N/A )	3119.7	N/A	2.0165 [ 2.0000 ]	100.8% { 106.8% }			
D5_NEtFOsa_EIS	( 531.0 / 169.0 ) 1764295	( 10.57 , N/A ) ( N/A , 0.02 , N/A )	2916.1	N/A	2.0400 [ 2.0000 ]	102.0% { 102.4% }			

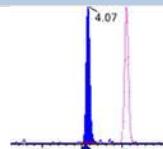
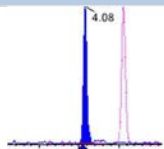
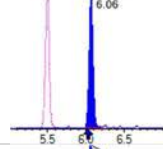
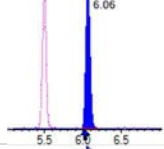
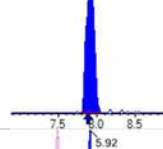
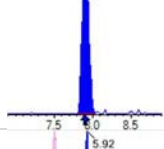
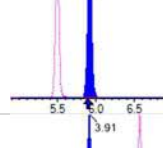
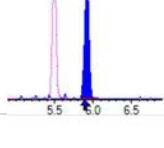
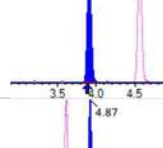
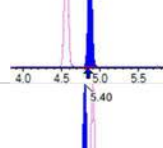
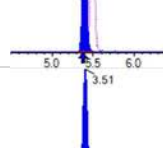
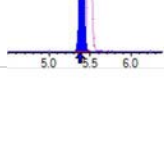
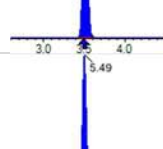
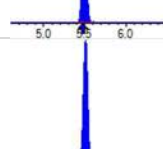
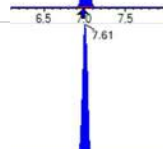

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-Imin, ΔRT-CVmin, ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration True ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	(573.0 / 419.0) 1495876	(8.32, N/A) (N/A, 0.03, N/A)	2110.6	N/A	4.1829 [4.0000]	104.6% {100.5%}			
D5_EtFOSAA_EIS	(589.0 / 419.0) 1120624	(8.56, N/A) (N/A, 0.02, N/A)	4180.6	N/A	3.8269 [4.0000]	95.7% {99.9%}			
D7_NMeFOSE_EIS	(623.0 / 58.9) 7269308	(10.36, N/A) (N/A, 0.02, N/A)	1974.3	N/A	20.6314 [20.0000]	103.2% {101.1%}			
D9_NEtFOSE_EIS	(639.0 / 58.9) 8179426	(10.52, N/A) (N/A, 0.02, N/A)	2057.8	N/A	18.1274 [20.0000]	90.6% {85.9%}			
13C3_HFPODA_EIS	(287.0 / 169.0) 3647056	(5.81, N/A) (N/A, 0.03, N/A)	3400.0	N/A	7.5725 [8.0000]	94.7% {99.6%}			

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
PFBA	( 213.0 / 169.0 ) 451149	( 3.51 , 1.00 ) ( 0.00 , N/A , 0.0 )	144.9	N/A 0.0 0.0	1.8493 [ 2.0000 ]	92.5%			
PFPeA	( 263.0 / 219.0 ) 452555 ( 263.0 / 69.0 ) 8313	( 4.56 , 1.00 ) ( 0.00 , N/A , 0.3 )	1325.9 88.5	0.0184 171.5 171.5	0.9113 [ 1.0000 ]	91.1%			
PFHxA	( 313.0 / 269.0 ) 289264 ( 313.0 / 119.0 ) 25240	( 5.49 , 1.00 ) ( 0.00 , N/A , -0.2 )	2436.9 1164772.0	0.0873 87.6 87.6	0.4579 [ 0.5000 ]	91.6%			
PFHpA	( 363.0 / 319.0 ) 250921 ( 363.0 / 169.0 ) 65829	( 6.32 , 1.00 ) ( 0.00 , N/A , 0.1 )	3944.0 8977.6	0.2624 81.8 81.8	0.4549 [ 0.5000 ]	91.0%			
PFOA	( 413.0 / 369.0 ) 313034 ( 413.0 / 169.0 ) 106346	( 7.01 , 1.00 ) ( 0.00 , N/A , -0.1 )	1450.9 9317.1	0.3397 101.5 101.5	0.4538 [ 0.5000 ]	90.8%			
PFNA	( 463.0 / 419.0 ) 284469 ( 463.0 / 169.0 ) 62891	( 7.61 , 1.00 ) ( 0.00 , N/A , -0.2 )	31533.0 17199442.2	0.2211 98.8 98.8	0.5039 [ 0.5000 ]	100.8%			
PFDA	( 513.0 / 469.0 ) 303672 ( 513.0 / 169.0 ) 41031	( 8.17 , 1.00 ) ( 0.00 , N/A , 0.3 )	433.1 1465214.7	0.1351 111.1 111.1	0.4463 [ 0.5000 ]	89.3%			
PFUnA	( 563.0 / 519.0 ) 289025 ( 563.0 / 169.0 ) 39284	( 8.65 , 1.00 ) ( 0.00 , N/A , 0.0 )	915.6 1903603.1	0.1359 129.6 129.6	0.4511 [ 0.5000 ]	90.2%			
PFDaA	( 613.0 / 569.0 ) 291794 ( 613.0 / 169.0 ) 48414	( 8.93 , 1.00 ) ( 0.00 , N/A , -0.1 )	1118.5 762.6	0.1659 101.4 101.4	0.5240 [ 0.5000 ]	104.8%			
PFTTrDA	( 663.0 / 619.0 ) 246586 ( 663.0 / 169.0 ) 80494	( 9.13 , 1.02 ) ( N/A , 0.01 , -0.2 )	2744.7 100483.4	0.3264 136.6 136.6	0.4891 [ 0.5000 ]	97.8%			
PFTeDA	( 713.0 / 669.0 ) 240170 ( 713.0 / 169.0 ) 59327	( 9.30 , 1.00 ) ( 0.00 , N/A , 0.4 )	675.8 329.7	0.2470 119.5 119.5	0.5061 [ 0.5000 ]	101.2%			

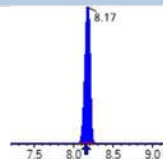
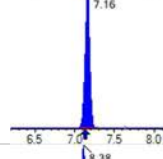
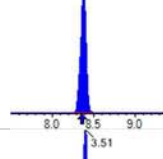
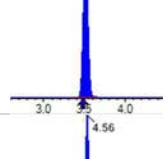
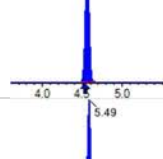
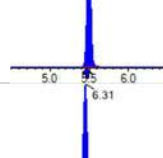
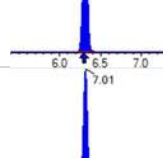
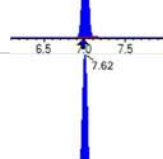
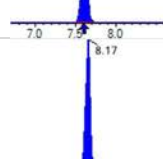
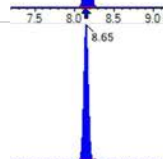
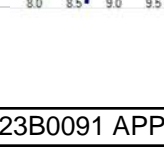
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	( 299.0 / 80.0 ) 488992 ( 299.0 / 99.0 ) 318173	( 5.47 , 1.00 ) ( 0.00 , N/A , 0.1 )	1533.7 797.1	0.6507 100.6 100.6	0.4102 [ 0.4424 ]	92.7%			
PFPeS	( 349.0 / 80.0 ) 884263 ( 349.0 / 99.0 ) 328050	( 6.40 , 0.89 ) ( N/A , 0.03 , 0.0 )	76134.6 12284.9	0.3710 103.9 103.9	0.4660 [ 0.4692 ]	99.3%			
PFHxS	( 399.0 / 80.0 ) 613514 ( 399.0 / 99.0 ) 203107	( 7.16 , 1.00 ) ( 0.00 , N/A , 0.2 )	45927.5 249.2	0.3311 101.9 101.9	0.4055 [ 0.4555 ]	89.0%			
PFHpS	( 449.0 / 80.0 ) 737280 ( 449.0 / 99.0 ) 229850	( 7.80 , 0.93 ) ( N/A , 0.03 , 0.0 )	4303.0 425623.2	0.3118 114.5 114.5	0.4457 [ 0.4757 ]	93.7%			
PFOS	( 499.0 / 80.0 ) 1008273 ( 499.0 / 99.0 ) 218013	( 8.38 , 1.00 ) ( 0.00 , N/A , -0.2 )	174.8 455.8	0.2162 97.7 97.7	0.4405 [ 0.4637 ]	95.0%			
PFNS	( 549.0 / 80.0 ) 1109039 ( 549.0 / 99.0 ) 278597	( 8.79 , 1.05 ) ( N/A , 0.02 , 0.0 )	9198421.7 19330.3	0.2512 114.6 114.6	0.4311 [ 0.4799 ]	89.8%			
PFDS	( 599.0 / 80.0 ) 1241916 ( 599.0 / 99.0 ) 290512	( 9.04 , 1.08 ) ( N/A , 0.02 , 0.0 )	1447.0 1355.0	0.2339 107.3 107.3	0.4438 [ 0.4816 ]	92.2%			
PFDoS	( 699.0 / 80.0 ) 801641 ( 699.0 / 99.0 ) 188250	( 9.38 , 1.12 ) ( N/A , 0.01 , -0.1 )	1658.6 1272.5	0.2348 112.8 112.8	0.4491 [ 0.4848 ]	92.6%			
4:2FTS	( 327.0 / 307.0 ) 513694 ( 327.0 / 81.0 ) 373584	( 5.23 , 1.00 ) ( 0.00 , N/A , 0.1 )	2186.0 971.2	0.7272 107.8 107.8	1.9300 [ 1.8691 ]	103.3%			
6:2FTS	( 427.0 / 407.0 ) 304578 ( 427.0 / 81.0 ) 194493	( 6.74 , 1.00 ) ( 0.00 , N/A , 0.2 )	16204.4 543.4	0.6386 77.1 77.1	1.9163 [ 1.8981 ]	101.0%			
8:2FTS	( 527.0 / 507.0 ) 298908 ( 527.0 / 81.0 ) 263515	( 7.92 , 1.00 ) ( 0.00 , N/A , 0.4 )	1027.2 1232.9	0.8816 106.1 106.1	1.8930 [ 1.9166 ]	98.8%			

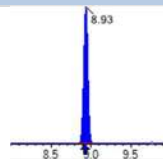
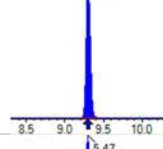
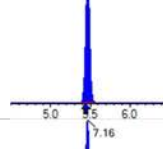
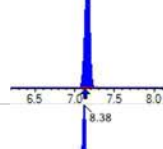
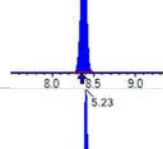
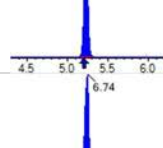
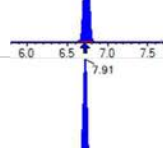
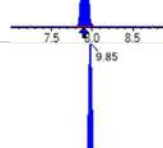
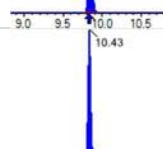
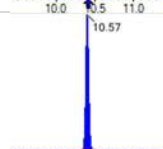
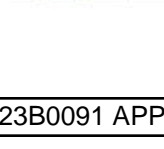
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	( 498.0 / 78.0 ) 1656779 ( 498.0 / 478.0 ) 33096	( 9.84 , 1.00 ) ( 0.00 , N/A , 0.2 )	1723.9 736.7	0.0200 97.3 97.3	0.4992 [ 0.5000 ]	99.8%			
NMeFOSA	( 512.0 / 219.0 ) 1474591 ( 512.0 / 169.0 ) 1275492	( 10.43 , 1.00 ) ( 0.00 , N/A , 0.7 )	2716.9 2319.6	0.8650 105.6 105.6	1.7977 [ 2.0000 ]	89.9%			
NEIFOSA	( 526.0 / 219.0 ) 1722014 ( 526.0 / 169.0 ) 2236482	( 10.57 , 1.00 ) ( 0.00 , N/A , 1.0 )	3690.3 4294.5	1.2988 102.3 102.3	1.9274 [ 2.0000 ]	96.4%			
NMeFOSAA	( 570.0 / 419.0 ) 108171 ( 570.0 / 483.0 ) 55304	( 8.32 , 1.00 ) ( 0.01 , N/A , 0.1 )	2008.1 6004.0	0.5113 107.7 107.7	0.4261 [ 0.5000 ]	85.2%			
NEIFOSAA	( 584.0 / 419.0 ) 98385 ( 584.0 / 526.0 ) 67969	( 8.56 , 1.00 ) ( 0.00 , N/A , -0.2 )	8363.3 1160.0	0.6908 135.4 135.4	0.4314 [ 0.5000 ]	86.3%			
NMeFOSE	( 616.0 / 59.0 ) 669858	( 10.37 , 1.00 ) ( 0.01 , N/A , 0.0 )	1001.5	N/A 0.0 0.0	1.8949 [ 2.0000 ]	94.7%			
NEtFOSE	( 630.0 / 59.0 ) 888349	( 10.53 , 1.00 ) ( 0.01 , N/A , 0.0 )	857.4	N/A 0.0 0.0	1.9103 [ 2.0000 ]	95.5%			
HFPO-DA	( 285.0 / 169.0 ) 321441 ( 285.0 / 185.0 ) 865673	( 5.81 , 1.00 ) ( 0.00 , N/A , 0.0 )	1278.5 1204.7	2.6931 97.8 97.8	0.9265 [ 1.0000 ]	92.6%			
ADONA	( 377.0 / 85.0 ) 1107868 ( 377.0 / 251.0 ) 118038	( 6.60 , 1.14 ) ( N/A , 0.03 , -0.3 )	1437.6 728.1	0.1065 120.8 120.8	0.9285 [ 0.9427 ]	98.5%			
9CI-Pf3ONS	( 531.0 / 351.0 ) 3303252 ( 533.0 / 353.0 ) 984795	( 8.72 , 1.50 ) ( N/A , 0.02 , 0.1 )	2197.8 1257.6	0.2981 85.6 85.6	0.8924 [ 0.9333 ]	95.6%			
11CI-PF3OUDS	( 631.0 / 451.0 ) 1784838 ( 633.0 / 453.0 ) 612728	( 9.18 , 1.58 ) ( N/A , 0.01 , 0.0 )	1962.5 1840.4	0.3433 94.9 94.9	0.9182 [ 0.9432 ]	97.4%			

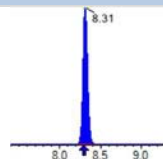
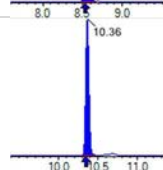
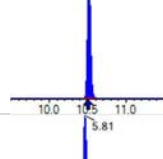




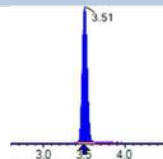
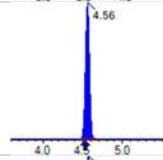
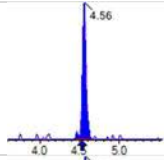
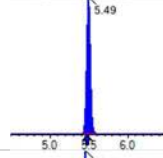
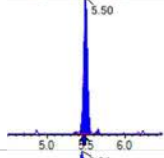
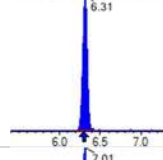
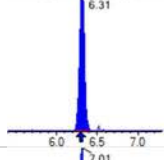
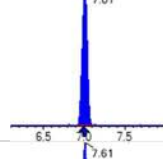
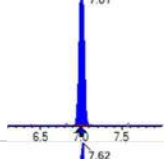
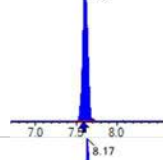
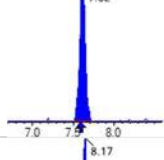
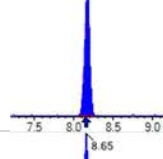
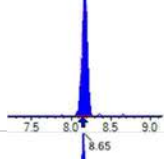
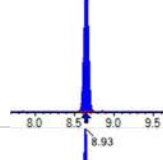
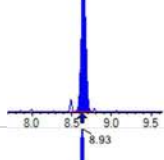
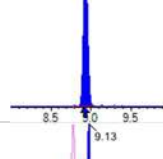
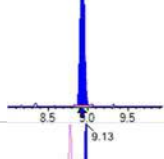
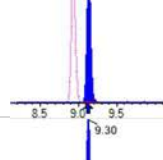
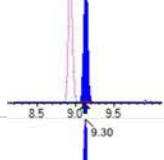
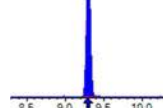
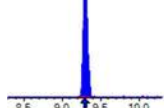
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	( 241.0 / 177.0 ) 22880 ( 241.0 / 117.0 ) 36417	( 4.07 , 0.89 ) ( N/A , 0.03 , -0.1 )	390.3 261.9	1.5917 100.9 100.9	1.9481 [ 2.0000 ]	97.4%			
5:3FTCA	( 341.0 / 236.7 ) 155711 ( 341.0 / 217.0 ) 279237	( 6.06 , 1.10 ) ( N/A , 0.04 , 0.1 )	515.7 785.8	1.7933 105.2 105.2	1.8024 [ 2.0000 ]	90.1%			
7:3FTCA	( 441.0 / 317.0 ) 303061 ( 441.0 / 337.0 ) 252225	( 7.91 , 1.44 ) ( N/A , 0.03 , 0.2 )	412.3 382.9	0.8323 96.7 96.7	1.8990 [ 2.0000 ]	94.9%			
PFEESA	( 315.0 / 135.0 ) 668291 ( 315.0 / 83.0 ) 182692	( 5.92 , 1.08 ) ( N/A , 0.03 , -0.1 )	2261.2 359.4	0.2734 107.0 107.0	0.8806 [ 0.8925 ]	98.7%			
PFMPA	( 229.0 / 85.0 ) 101708	( 3.91 , 0.86 ) ( N/A , 0.03 , 0.0 )	888.8	N/A 0.0 0.0	0.9376 [ 1.0000 ]	93.8%			
PFMBA	( 279.0 / 85.0 ) 405237	( 4.87 , 1.07 ) ( N/A , 0.03 , 0.0 )	1278.3	N/A 0.0 0.0	0.8908 [ 1.0000 ]	89.1%			
NFDHA	( 295.0 / 201.0 ) 315461 ( 295.0 / 85.0 ) 324582	( 5.40 , 0.98 ) ( N/A , 0.03 , 0.0 )	1056.6 916.7	1.0289 102.8 102.8	0.8996 [ 1.0000 ]	90.0%			
13C3_PFBA_IIS	( 216.0 / 172.0 ) 237086	( 3.51 , N/A ) ( N/A , 0.02 , N/A )	1428.1	N/A	1.0585 [ 1.0000 ]	105.8% { 105.2% }			
13C2_PFHxA_IIS	( 315.0 / 270.0 ) 474334	( 5.49 , N/A ) ( N/A , 0.03 , N/A )	5166.1	N/A	0.9396 [ 1.0000 ]	94.0% { 98.8% }			
13C4_PFOA_IIS	( 417.0 / 372.0 ) 597727	( 7.01 , N/A ) ( N/A , 0.03 , N/A )	1166.1	N/A	0.9431 [ 1.0000 ]	94.3% { 100.7% }			
13C5_PFNA_IIS	( 468.0 / 423.0 ) 600355	( 7.61 , N/A ) ( N/A , 0.03 , N/A )	4826.5	N/A	0.9935 [ 1.0000 ]	99.4% { 101.3% }			

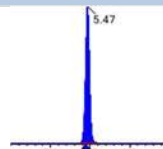
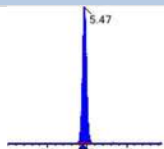
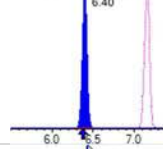
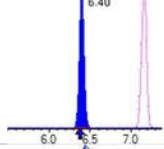
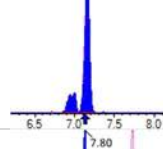
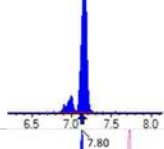
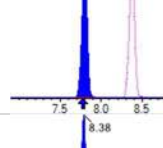
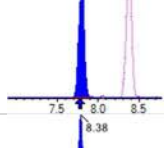
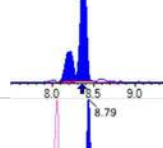
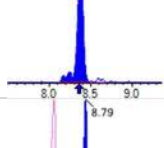
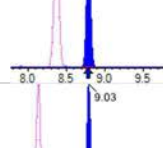
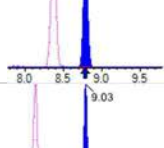
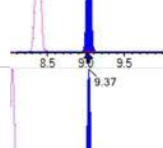
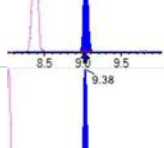
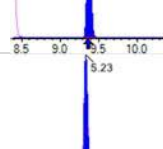
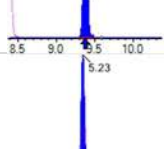
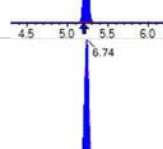
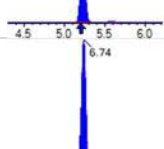
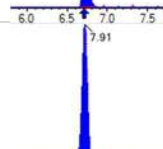
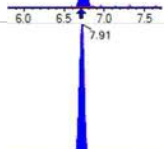
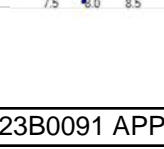
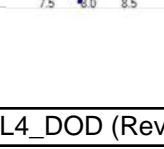


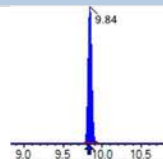
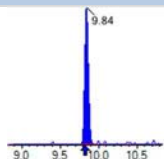
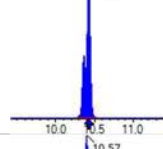
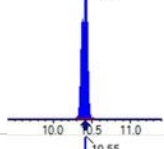
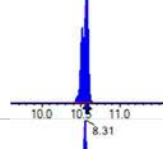
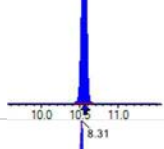
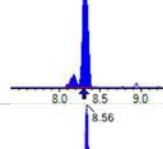
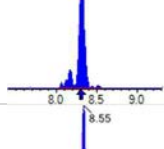
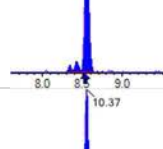
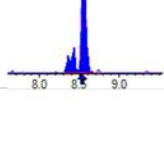
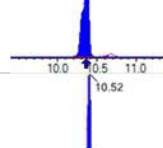
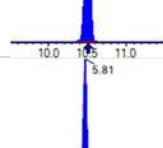
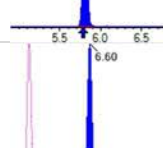
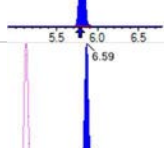
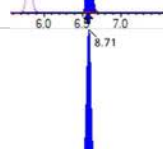
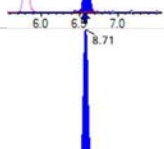
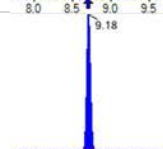
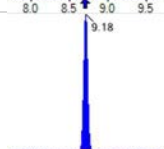
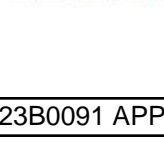
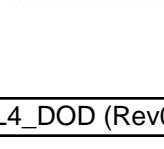
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	( 515.0 / 470.1 ) 537002	( 8.17 , N/A ) ( N/A , 0.02 , N/A )	10203.3	N/A	0.9280 [ 1.0000 ]	92.8% { 98.6% }			
18O2_PFHxS_IIS	( 403.0 / 83.9 ) 872778	( 7.16 , N/A ) ( N/A , 0.03 , N/A )	1067.1	N/A	1.0192 [ 1.0000 ]	101.9% { 103.4% }			
13C4_PFOS_IIS	( 503.0 / 79.9 ) 1485383	( 8.38 , N/A ) ( N/A , 0.02 , N/A )	1347.5	N/A	0.9867 [ 1.0000 ]	98.7% { 98.2% }			
13C4_PFBA_EIS	( 217.0 / 172.0 ) 2356057	( 3.51 , N/A ) ( N/A , 0.02 , N/A )	5406.3	N/A	8.3544 [ 8.0000 ]	104.4% { 100.9% }			
13C5_PFPeA_EIS	( 268.0 / 223.0 ) 2414394	( 4.56 , N/A ) ( N/A , 0.03 , N/A )	3420.2	N/A	4.6522 [ 4.0000 ]	116.3% { 99.8% }			
13C5_PFHxA_EIS	( 318.0 / 273.0 ) 1437224	( 5.49 , N/A ) ( N/A , 0.03 , N/A )	2433.0	N/A	2.2121 [ 2.0000 ]	110.6% { 93.7% }			
13C4_PFHpA_EIS	( 367.0 / 322.0 ) 1306373	( 6.31 , N/A ) ( N/A , 0.03 , N/A )	2350.6	N/A	2.3206 [ 2.0000 ]	116.0% { 103.4% }			
13C8_PFOA_EIS	( 421.0 / 376.0 ) 1549605	( 7.01 , N/A ) ( N/A , 0.03 , N/A )	3163.0	N/A	2.2891 [ 2.0000 ]	114.5% { 100.6% }			
13C9_PFNA_EIS	( 472.0 / 427.0 ) 638082	( 7.62 , N/A ) ( N/A , 0.02 , N/A )	2377.6	N/A	1.0031 [ 1.0000 ]	100.3% { 90.8% }			
13C6_PFDA_EIS	( 519.0 / 474.0 ) 773492	( 8.17 , N/A ) ( N/A , 0.03 , N/A )	2146.3	N/A	1.1424 [ 1.0000 ]	114.2% { 99.6% }			
13C7_PFUnA_EIS	( 570.0 / 525.0 ) 771164	( 8.65 , N/A ) ( N/A , 0.02 , N/A )	2481.9	N/A	1.1454 [ 1.0000 ]	114.5% { 100.0% }			

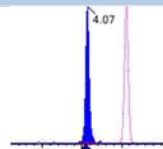
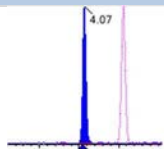
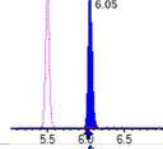
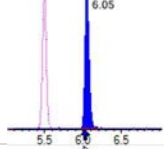
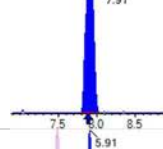
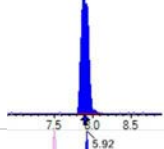
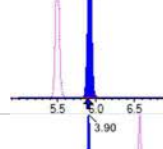
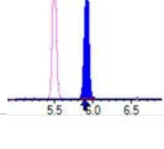
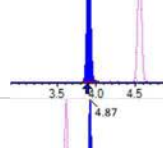
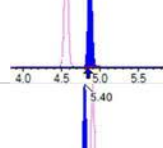
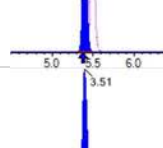
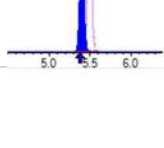
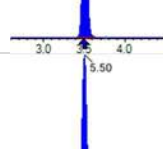
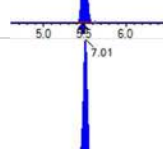
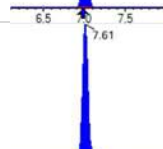

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	( 615.0 / 570.0 ) 633220	( 8.93 , N/A ) ( N/A , 0.02 , N/A )	1548.4	N/A	1.0966 [ 1.0000 ]	109.7% { 96.4% }			
13C2_PFTeDA_EIS	( 715.0 / 670.0 ) 548363	( 9.30 , N/A ) ( N/A , 0.01 , N/A )	1640.3	N/A	1.0363 [ 1.0000 ]	103.6% { 89.9% }			
13C3_PFBs_EIS	( 302.0 / 80.0 ) 3836135	( 5.47 , N/A ) ( N/A , 0.03 , N/A )	4222.9	N/A	2.1001 [ 2.0000 ]	105.0% { 97.1% }			
13C3_PFHxS_EIS	( 402.0 / 80.0 ) 1967780	( 7.16 , N/A ) ( N/A , 0.03 , N/A )	2175.8	N/A	2.0740 [ 2.0000 ]	103.7% { 98.9% }			
13C8_PFOS_EIS	( 507.0 / 80.0 ) 4341643	( 8.38 , N/A ) ( N/A , 0.03 , N/A )	2646.7	N/A	2.1291 [ 2.0000 ]	106.5% { 98.2% }			
13C2_4:2FTS_EIS	( 329.0 / 81.0 ) 389814	( 5.23 , N/A ) ( N/A , 0.03 , N/A )	2328.3	N/A	3.8662 [ 4.0000 ]	96.7% { 95.2% }			
13C2_6:2FTS_EIS	( 429.0 / 81.0 ) 448065	( 6.74 , N/A ) ( N/A , 0.03 , N/A )	1550.7	N/A	4.0484 [ 4.0000 ]	101.2% { 97.8% }			
13C2_8:2FTS_EIS	( 529.0 / 81.0 ) 579858	( 7.91 , N/A ) ( N/A , 0.03 , N/A )	1886.6	N/A	3.7084 [ 4.0000 ]	92.7% { 95.8% }			
13C8_PFOsa_EIS	( 506.0 / 78.0 ) 8299727	( 9.85 , N/A ) ( N/A , 0.02 , N/A )	2955.9	N/A	2.1926 [ 2.0000 ]	109.6% { 96.8% }			
D3_NMeFOsa_EIS	( 515.0 / 169.0 ) 1966659	( 10.43 , N/A ) ( N/A , 0.01 , N/A )	2317.1	N/A	2.2052 [ 2.0000 ]	110.3% { 107.8% }			
D5_NEtFOsa_EIS	( 531.0 / 169.0 ) 1912622	( 10.57 , N/A ) ( N/A , 0.01 , N/A )	3094.7	N/A	2.3966 [ 2.0000 ]	119.8% { 111.0% }			

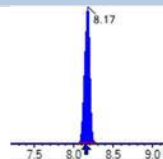
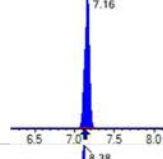
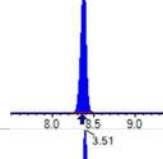
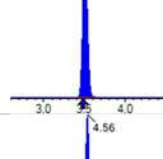
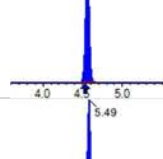
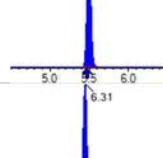
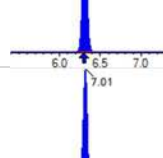
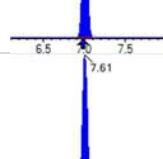
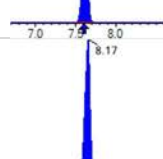
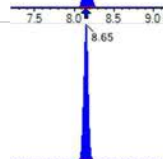
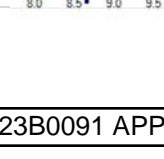
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	( 573.0 / 419.0 ) 1294019	( 8.31 , N/A ) ( N/A , 0.03 , N/A )	2109.7	N/A	3.9213 [ 4.0000 ]	98.0% { 86.9% }			
D5_EtFOSAA_EIS	( 589.0 / 419.0 ) 1086211	( 8.56 , N/A ) ( N/A , 0.02 , N/A )	9289.8	N/A	4.0198 [ 4.0000 ]	100.5% { 96.8% }			
D7_NMeFOSE_EIS	( 623.0 / 58.9 ) 7071882	( 10.36 , N/A ) ( N/A , 0.01 , N/A )	1362.0	N/A	21.7507 [ 20.0000 ]	108.8% { 98.4% }			
D9_NEtFOSE_EIS	( 639.0 / 58.9 ) 9773335	( 10.52 , N/A ) ( N/A , 0.01 , N/A )	2205.6	N/A	23.4725 [ 20.0000 ]	117.4% { 102.6% }			
13C3_HFPODA_EIS	( 287.0 / 169.0 ) 3509898	( 5.81 , N/A ) ( N/A , 0.03 , N/A )	3188.3	N/A	8.8726 [ 8.0000 ]	110.9% { 95.9% }			

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 ) 992985	( 3.51 , 1.00 ) ( 0.00 , N/A , 0.0 )	192.4	N/A 0.0 0.0	4.0401 [ 4.0000 ]	101.0%			
PFPeA	( 263.0 / 219.0 ) 995190 ( 263.0 / 69.0 ) 13200	( 4.56 , 1.00 ) ( 0.00 , N/A , 0.3 )	2220.7 225.4	0.0133 123.9 123.9	2.0489 [ 2.0000 ]	102.4%			
PFHxA	( 313.0 / 269.0 ) 632186 ( 313.0 / 119.0 ) 57415	( 5.49 , 1.00 ) ( 0.00 , N/A , -0.1 )	2410.2 171374.7	0.0908 91.2 91.2	0.9528 [ 1.0000 ]	95.3%			
PFHpA	( 363.0 / 319.0 ) 507737 ( 363.0 / 169.0 ) 157722	( 6.31 , 1.00 ) ( 0.00 , N/A , 0.0 )	26405.3 2125592.4	0.3106 96.9 96.9	0.9408 [ 1.0000 ]	94.1%			
PFOA	( 413.0 / 369.0 ) 622937 ( 413.0 / 169.0 ) 205185	( 7.01 , 1.00 ) ( 0.00 , N/A , 0.1 )	6482.6 6529.1	0.3294 98.4 98.4	0.9571 [ 1.0000 ]	95.7%			
PFNA	( 463.0 / 419.0 ) 570086 ( 463.0 / 169.0 ) 115433	( 7.61 , 1.00 ) ( 0.00 , N/A , -0.3 )	4299.3 26480.6	0.2025 90.5 90.5	0.9731 [ 1.0000 ]	97.3%			
PFDA	( 513.0 / 469.0 ) 667559 ( 513.0 / 169.0 ) 85346	( 8.17 , 1.00 ) ( 0.00 , N/A , -0.1 )	713.4 2307.5	0.1278 105.1 105.1	0.9768 [ 1.0000 ]	97.7%			
PFUnA	( 563.0 / 519.0 ) 656549 ( 563.0 / 169.0 ) 83205	( 8.65 , 1.00 ) ( 0.00 , N/A , 0.1 )	1652.5 448.0	0.1267 120.9 120.9	1.0212 [ 1.0000 ]	102.1%			
PFDoA	( 613.0 / 569.0 ) 556848 ( 613.0 / 169.0 ) 102128	( 8.93 , 1.00 ) ( 0.00 , N/A , 0.1 )	675.4 232.8	0.1834 112.1 112.1	0.9839 [ 1.0000 ]	98.4%			
PFTTrDA	( 663.0 / 619.0 ) 522444 ( 663.0 / 169.0 ) 151041	( 9.13 , 1.02 ) ( N/A , 0.01 , -0.1 )	1144.8 8191.0	0.2891 121.0 121.0	1.0195 [ 1.0000 ]	101.9%			
PFTeDA	( 713.0 / 669.0 ) 522743 ( 713.0 / 169.0 ) 134113	( 9.30 , 1.00 ) ( 0.00 , N/A , 0.0 )	1196.0 1042.9	0.2566 124.1 124.1	0.9899 [ 1.0000 ]	99.0%			

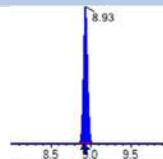
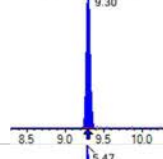
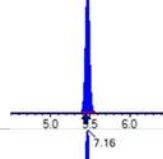
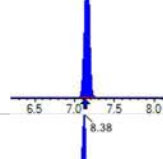
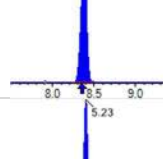
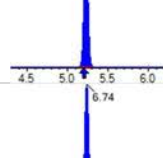
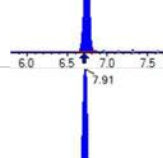
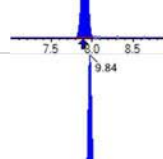
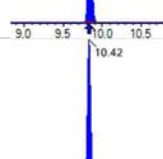
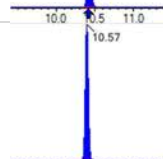
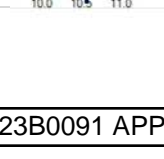
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	( 299.0 / 80.0 ) 1115761 ( 299.0 / 99.0 ) 660695	( 5.47 , 1.00 ) ( 0.00 , N/A , 0.1 )	3318.3 1288.5	0.5921 91.6 91.6	0.9169 [ 0.8847 ]	103.6%			
PFPeS	( 349.0 / 80.0 ) 1898329 ( 349.0 / 99.0 ) 641927	( 6.40 , 0.89 ) ( N/A , 0.03 , 0.1 )	28058.9 8629.1	0.3382 94.7 94.7	0.9722 [ 0.9384 ]	103.6%			
PFHxS	( 399.0 / 80.0 ) 1346938 ( 399.0 / 99.0 ) 485366	( 7.16 , 1.00 ) ( 0.00 , N/A , 0.0 )	7988.0 934.1	0.3603 111.0 111.0	0.8651 [ 0.9110 ]	95.0%			
PFHpS	( 449.0 / 80.0 ) 1649623 ( 449.0 / 99.0 ) 406605	( 7.80 , 0.93 ) ( N/A , 0.02 , 0.0 )	831232.4 8656.3	0.2465 90.5 90.5	0.9551 [ 0.9514 ]	100.4%			
PFOS	( 499.0 / 80.0 ) 2103149 ( 499.0 / 99.0 ) 460724	( 8.38 , 1.00 ) ( 0.00 , N/A , 0.0 )	307.7 265.9	0.2191 99.0 99.0	0.8798 [ 0.9275 ]	94.9%			
PFNS	( 549.0 / 80.0 ) 2497676 ( 549.0 / 99.0 ) 597897	( 8.79 , 1.05 ) ( N/A , 0.02 , 0.0 )	26410.0 39648.2	0.2394 109.2 109.2	0.9297 [ 0.9599 ]	96.9%			
PFDS	( 599.0 / 80.0 ) 2775816 ( 599.0 / 99.0 ) 601185	( 9.03 , 1.08 ) ( N/A , 0.01 , 0.1 )	2370.9 1333.7	0.2166 99.4 99.4	0.9499 [ 0.9631 ]	98.6%			
PFDoS	( 699.0 / 80.0 ) 1869304 ( 699.0 / 99.0 ) 376149	( 9.37 , 1.12 ) ( N/A , 0.01 , -0.1 )	1682.1 879.2	0.2012 96.7 96.7	1.0029 [ 0.9696 ]	103.4%			
4:2FTS	( 327.0 / 307.0 ) 1102979 ( 327.0 / 81.0 ) 700459	( 5.23 , 1.00 ) ( 0.00 , N/A , 0.0 )	2554.0 1188.0	0.6351 94.1 94.1	3.7263 [ 3.7381 ]	99.7%			
6:2FTS	( 427.0 / 407.0 ) 551597 ( 427.0 / 81.0 ) 491370	( 6.74 , 1.00 ) ( 0.00 , N/A , 0.0 )	4455.6 936.5	0.8908 107.6 107.6	3.5034 [ 3.7962 ]	92.3%			
8:2FTS	( 527.0 / 507.0 ) 648567 ( 527.0 / 81.0 ) 536762	( 7.91 , 1.00 ) ( 0.00 , N/A , -0.1 )	1501.8 2426.4	0.8276 99.6 99.6	4.3681 [ 3.8332 ]	114.0%			

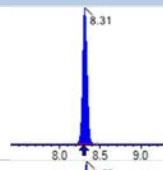




Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T. ) ( $\Delta$ RT-[min], $\Delta$ RT- CV[min], $\Delta$ RT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	( 498.0 / 78.0 ) 3648720 ( 498.0 / 478.0 ) 66647	( 9.84 , 1.00 ) ( 0.00 , N/A , 0.0 )	3061.1 348.5	0.0183 88.9 88.9	1.0638 [ 1.0000 ]	106.4%			
NMeFOSA	( 512.0 / 219.0 ) 3232329 ( 512.0 / 169.0 ) 2709837	( 10.42 , 1.00 ) ( 0.00 , N/A , 0.9 )	4031.3 4123.4	0.8384 102.3 102.3	3.8883 [ 4.0000 ]	97.2%			
NEIFOSA	( 526.0 / 219.0 ) 3650558 ( 526.0 / 169.0 ) 4758356	( 10.57 , 1.00 ) ( 0.00 , N/A , 1.0 )	6399.0 5545.6	1.3035 102.7 102.7	4.0961 [ 4.0000 ]	102.4%			
NMeFOSAA	( 570.0 / 419.0 ) 279159 ( 570.0 / 483.0 ) 118936	( 8.31 , 1.00 ) ( 0.00 , N/A , 0.0 )	4984.7 185484.6	0.4261 89.7 89.7	1.0227 [ 1.0000 ]	102.3%			
NEIFOSAA	( 584.0 / 419.0 ) 257131 ( 584.0 / 526.0 ) 141719	( 8.56 , 1.00 ) ( 0.00 , N/A , 0.1 )	2290.5 2169.6	0.5512 108.0 108.0	0.9982 [ 1.0000 ]	99.8%			
NMeFOSE	( 616.0 / 59.0 ) 1483945	( 10.37 , 1.00 ) ( 0.01 , N/A , 0.0 )	1605.1	N/A 0.0 0.0	3.9770 [ 4.0000 ]	99.4%			
NEIFOSE	( 630.0 / 59.0 ) 1945357	( 10.52 , 1.00 ) ( 0.01 , N/A , 0.0 )	1205.3	N/A 0.0 0.0	4.1721 [ 4.0000 ]	104.3%			
HFPO-DA	( 285.0 / 169.0 ) 702936 ( 285.0 / 185.0 ) 2029139	( 5.81 , 1.00 ) ( 0.00 , N/A , 0.1 )	1495.7 2469.1	2.8867 104.8 104.8	1.9094 [ 2.0000 ]	95.5%			
ADONA	( 377.0 / 85.0 ) 2546378 ( 377.0 / 251.0 ) 217566	( 6.60 , 1.14 ) ( N/A , 0.03 , 0.2 )	2490.7 407.3	0.0854 96.9 96.9	2.0112 [ 1.8854 ]	106.7%			
9CI-PF3ONS	( 531.0 / 351.0 ) 6975004 ( 533.0 / 353.0 ) 2111294	( 8.71 , 1.50 ) ( N/A , 0.02 , 0.1 )	2422.7 1747.7	0.3027 86.9 86.9	1.7421 [ 1.8665 ]	93.3%			
11CI-PF3OUDS	( 631.0 / 451.0 ) 3820731 ( 633.0 / 453.0 ) 1412412	( 9.18 , 1.58 ) ( N/A , 0.01 , -0.1 )	2015.7 2421.3	0.3697 102.2 102.2	1.8524 [ 1.8864 ]	98.2%			

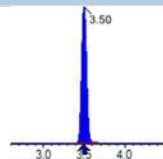
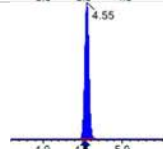
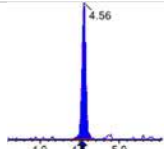
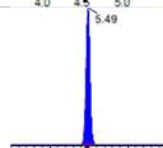
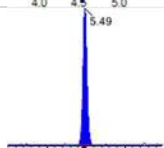
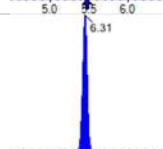
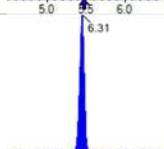
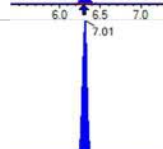
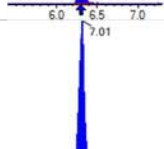
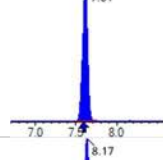
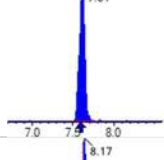
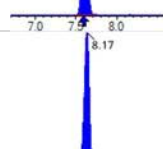
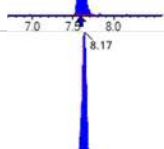
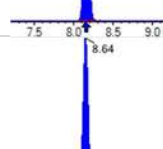
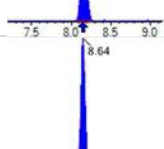
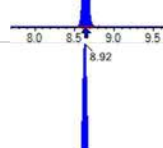
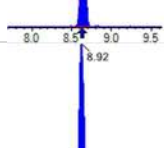
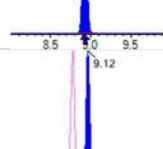
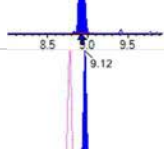
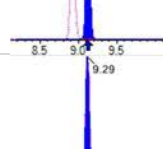
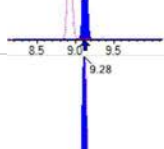
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) ( $\Delta$ RT-[min], $\Delta$ RT- CV[min], $\Delta$ RT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	( 241.0 / 177.0 ) 49606 ( 241.0 / 117.0 ) 73731	( 4.07 , 0.89 ) ( N/A , 0.02 , 0.1 )	645.5 531.6	1.4863 94.2 94.2	4.2828 [ 4.0000 ]	107.1%			
5:3FTCA	( 341.0 / 236.7 ) 307972 ( 341.0 / 217.0 ) 563339	( 6.05 , 1.10 ) ( N/A , 0.02 , 0.1 )	1195.2 949.5	1.8292 107.3 107.3	3.3942 [ 4.0000 ]	84.9%			
7:3FTCA	( 441.0 / 317.0 ) 606609 ( 441.0 / 337.0 ) 509153	( 7.91 , 1.44 ) ( N/A , 0.03 , 0.5 )	637.1 611.4	0.8393 97.5 97.5	3.6190 [ 4.0000 ]	90.5%			
PFEESA	( 315.0 / 135.0 ) 1415175 ( 315.0 / 83.0 ) 430101	( 5.91 , 1.08 ) ( N/A , 0.03 , -0.1 )	2671.2 1261.5	0.3039 119.0 119.0	1.7755 [ 1.7849 ]	99.5%			
PFMPA	( 229.0 / 85.0 ) 210517	( 3.90 , 0.86 ) ( N/A , 0.02 , 0.0 )	1653.4	N/A 0.0 0.0	1.9842 [ 2.0000 ]	99.2%			
PFMBA	( 279.0 / 85.0 ) 870406	( 4.87 , 1.07 ) ( N/A , 0.03 , 0.0 )	2115.8	N/A 0.0 0.0	1.9563 [ 2.0000 ]	97.8%			
NFDHA	( 295.0 / 201.0 ) 737616 ( 295.0 / 85.0 ) 776112	( 5.40 , 0.98 ) ( N/A , 0.03 , 0.0 )	2100.7 2541.0	1.0522 105.1 105.1	2.0028 [ 2.0000 ]	100.1%			
13C3_PFBA_IIS	( 216.0 / 172.0 ) 247463	( 3.51 , N/A ) ( N/A , 0.02 , N/A )	1430.4	N/A	1.1048 [ 1.0000 ]	110.5% { 109.8% }			
13C2_PFHxA_IIS	( 315.0 / 270.0 ) 545767	( 5.50 , N/A ) ( N/A , 0.03 , N/A )	7771.3	N/A	1.0811 [ 1.0000 ]	108.1% { 113.7% }			
13C4_PFOA_IIS	( 417.0 / 372.0 ) 680981	( 7.01 , N/A ) ( N/A , 0.03 , N/A )	2337.2	N/A	1.0745 [ 1.0000 ]	107.4% { 114.8% }			
13C5_PFNA_IIS	( 468.0 / 423.0 ) 635510	( 7.61 , N/A ) ( N/A , 0.02 , N/A )	1046.2	N/A	1.0517 [ 1.0000 ]	105.2% { 107.2% }			

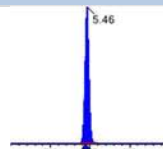
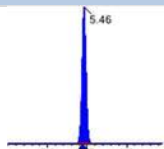
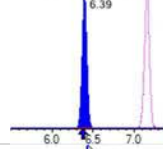
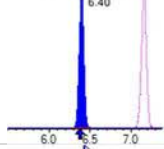
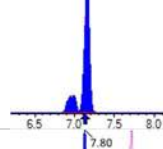
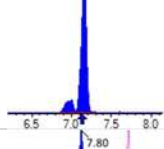
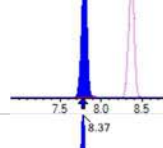
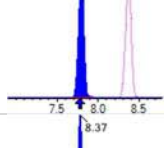
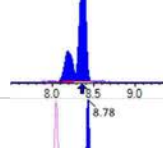
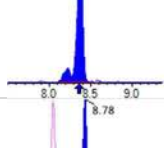
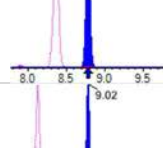
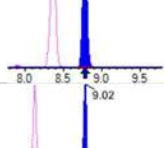
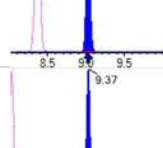
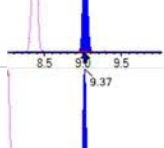
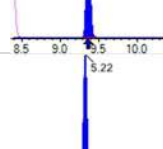
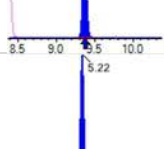
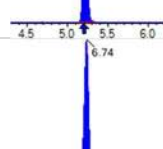
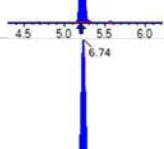
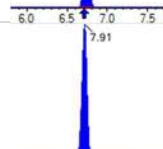
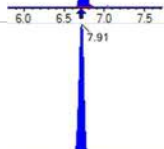
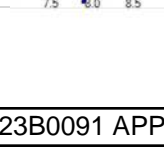
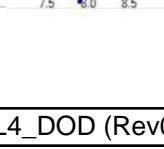
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	( 515.0 / 470.1 ) 586872	( 8.17, N/A ) ( N/A, 0.02, N/A )	1991.3	N/A	1.0142 [ 1.0000 ]	101.4% { 107.8% }			
18O2_PFHxS_IIS	( 403.0 / 83.9 ) 877200	( 7.16, N/A ) ( N/A, 0.03, N/A )	2151.0	N/A	1.0243 [ 1.0000 ]	102.4% { 103.9% }			
13C4_PFOS_IIS	( 503.0 / 79.9 ) 1527312	( 8.38, N/A ) ( N/A, 0.02, N/A )	1792.9	N/A	1.0146 [ 1.0000 ]	101.5% { 101.0% }			
13C4_PFBA_EIS	( 217.0 / 172.0 ) 2373686	( 3.51, N/A ) ( N/A, 0.02, N/A )	4602.1	N/A	8.0640 [ 8.0000 ]	100.8% { 101.7% }			
13C5_PFPeA_EIS	( 268.0 / 223.0 ) 2361531	( 4.56, N/A ) ( N/A, 0.03, N/A )	3090.6	N/A	3.9547 [ 4.0000 ]	98.9% { 97.6% }			
13C5_PFHxA_EIS	( 318.0 / 273.0 ) 1509504	( 5.49, N/A ) ( N/A, 0.03, N/A )	2234.0	N/A	2.0193 [ 2.0000 ]	101.0% { 98.4% }			
13C4_PFHpA_EIS	( 367.0 / 322.0 ) 1278281	( 6.31, N/A ) ( N/A, 0.03, N/A )	2293.3	N/A	1.9735 [ 2.0000 ]	98.7% { 101.2% }			
13C8_PFOA_EIS	( 421.0 / 376.0 ) 1461964	( 7.01, N/A ) ( N/A, 0.03, N/A )	2029.4	N/A	1.8956 [ 2.0000 ]	94.8% { 94.9% }			
13C9_PFNA_EIS	( 472.0 / 427.0 ) 662113	( 7.61, N/A ) ( N/A, 0.02, N/A )	2039.6	N/A	0.9833 [ 1.0000 ]	98.3% { 94.2% }			
13C6_PFDA_EIS	( 519.0 / 474.0 ) 776912	( 8.17, N/A ) ( N/A, 0.02, N/A )	9707.2	N/A	1.0499 [ 1.0000 ]	105.0% { 100.0% }			
13C7_PFUnA_EIS	( 570.0 / 525.0 ) 773791	( 8.65, N/A ) ( N/A, 0.01, N/A )	2173.1	N/A	1.0516 [ 1.0000 ]	105.2% { 100.3% }			

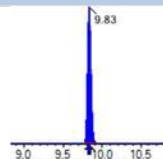
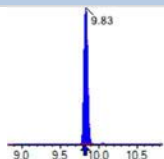
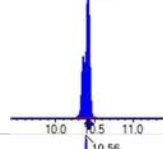
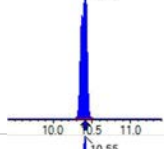
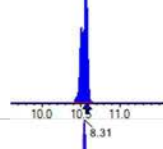
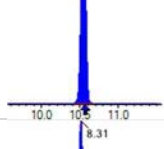
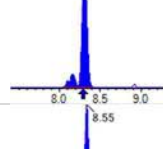
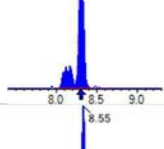
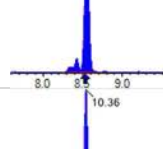
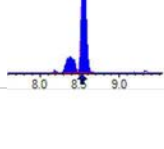
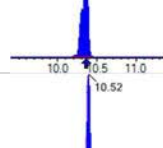
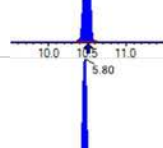
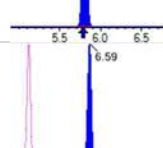
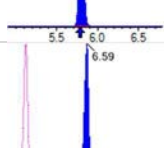
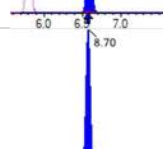
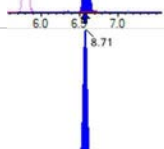
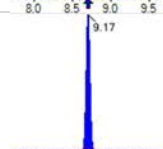
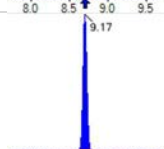
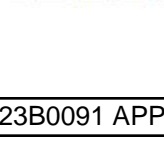
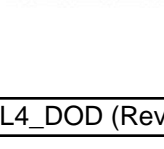


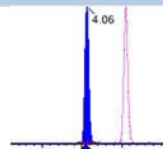
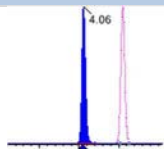
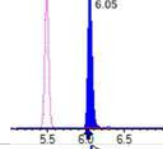
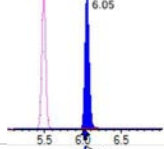
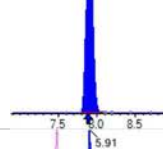
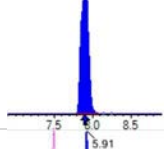
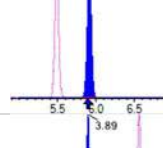
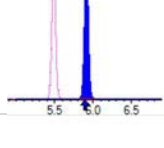
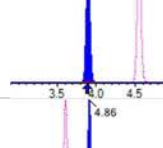
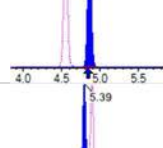
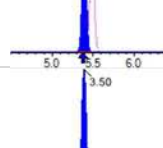
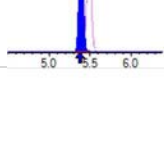
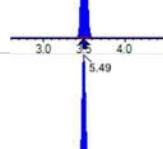
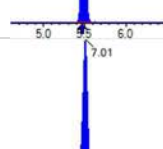
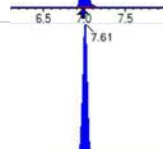

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	( 615.0 / 570.0 ) 643629	( 8.93 , N/A ) ( N/A , 0.01 , N/A )	2450.6	N/A	1.0199 [ 1.0000 ]	102.0% { 98.0% }			
13C2_PFTeDA_EIS	( 715.0 / 670.0 ) 610224	( 9.30 , N/A ) ( N/A , 0.01 , N/A )	802.3	N/A	1.0553 [ 1.0000 ]	105.5% { 100.0% }			
13C3_PFBs_EIS	( 302.0 / 80.0 ) 3916474	( 5.47 , N/A ) ( N/A , 0.03 , N/A )	3181.3	N/A	2.1333 [ 2.0000 ]	106.7% { 99.2% }			
13C3_PFHxS_EIS	( 402.0 / 80.0 ) 2025045	( 7.16 , N/A ) ( N/A , 0.02 , N/A )	1903.3	N/A	2.1236 [ 2.0000 ]	106.2% { 101.8% }			
13C8_PFOS_EIS	( 507.0 / 80.0 ) 4533818	( 8.38 , N/A ) ( N/A , 0.02 , N/A )	2062.8	N/A	2.1623 [ 2.0000 ]	108.1% { 102.6% }			
13C2_4:2FTS_EIS	( 329.0 / 81.0 ) 433513	( 5.23 , N/A ) ( N/A , 0.03 , N/A )	1584.2	N/A	4.2779 [ 4.0000 ]	106.9% { 105.9% }			
13C2_6:2FTS_EIS	( 429.0 / 81.0 ) 443849	( 6.74 , N/A ) ( N/A , 0.03 , N/A )	630.4	N/A	3.9901 [ 4.0000 ]	99.8% { 96.9% }			
13C2_8:2FTS_EIS	( 529.0 / 81.0 ) 545242	( 7.91 , N/A ) ( N/A , 0.03 , N/A )	949.4	N/A	3.4694 [ 4.0000 ]	86.7% { 90.1% }			
13C8_PFOsa_EIS	( 506.0 / 78.0 ) 8576988	( 9.84 , N/A ) ( N/A , 0.01 , N/A )	3982.4	N/A	2.2037 [ 2.0000 ]	110.2% { 100.1% }			
D3_NMeFOSA_EIS	( 515.0 / 169.0 ) 1961100	( 10.42 , N/A ) ( N/A , 0.01 , N/A )	2880.3	N/A	2.1386 [ 2.0000 ]	106.9% { 107.5% }			
D5_NEtFOSA_EIS	( 531.0 / 169.0 ) 1907916	( 10.57 , N/A ) ( N/A , 0.01 , N/A )	3172.4	N/A	2.3251 [ 2.0000 ]	116.3% { 110.7% }			

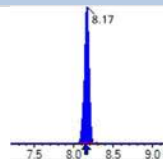
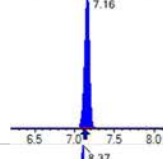
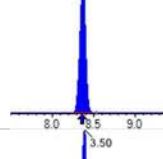
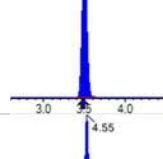
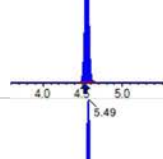
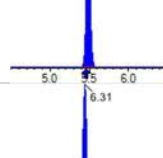
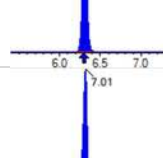
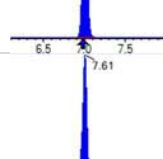
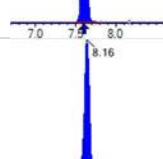
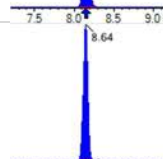
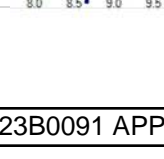
Analyte	( Q1 / Q3 ) Area Counts*min	R.T. ( R.T [min] , R.R.T. ) ( ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	( 573.0 / 419.0 ) 1391320	( 8.31 , N/A ) ( N/A , 0.02 , N/A )	1765.7	N/A	4.1004 [ 4.0000 ]	102.5% { 93.5% }			
D5_EtFOSAA_EIS	( 589.0 / 419.0 ) 1226806	( 8.55 , N/A ) ( N/A , 0.02 , N/A )	33894.6	N/A	4.4155 [ 4.0000 ]	110.4% { 109.4% }			
D7_NMeFOSE_EIS	( 623.0 / 58.9 ) 7464643	( 10.36 , N/A ) ( N/A , 0.01 , N/A )	1414.6	N/A	22.3284 [ 20.0000 ]	111.6% { 103.8% }			
D9_NEtFOSE_EIS	( 639.0 / 58.9 ) 9799462	( 10.51 , N/A ) ( N/A , 0.01 , N/A )	2582.9	N/A	22.8891 [ 20.0000 ]	114.4% { 102.9% }			
13C3_HFPODA_EIS	( 287.0 / 169.0 ) 3724342	( 5.81 , N/A ) ( N/A , 0.03 , N/A )	3728.1	N/A	8.1824 [ 8.0000 ]	102.3% { 101.7% }			

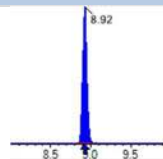
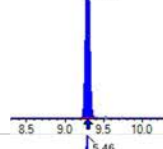
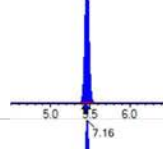
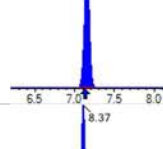
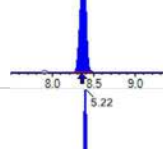
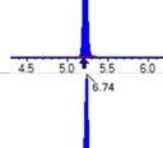
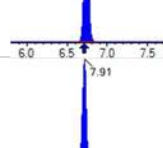
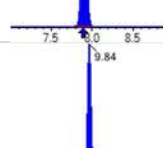
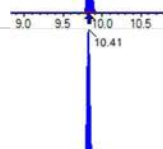
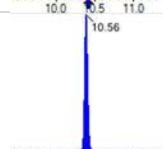
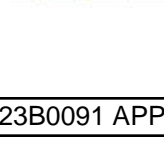
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) ( $\Delta$ RT-I[min], $\Delta$ RT- CV[min], $\Delta$ RT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	( 213.0 / 169.0 ) 2040417	( 3.50 , 1.00 ) ( 0.00 , N/A , 0.0 )	222.5	N/A 0.0 0.0	8.2532 [ 8.0000 ]	103.2%			
PFPeA	( 263.0 / 219.0 ) 2041470 ( 263.0 / 69.0 ) 23902	( 4.55 , 1.00 ) ( 0.00 , N/A , -0.2 )	2800.6 239.1	0.0117 109.3 109.3	4.0952 [ 4.0000 ]	102.4%			
PFHxA	( 313.0 / 269.0 ) 1336473 ( 313.0 / 119.0 ) 125042	( 5.49 , 1.00 ) ( 0.00 , N/A , 0.1 )	5541.8 2724595.8	0.0936 93.9 93.9	2.0232 [ 2.0000 ]	101.2%			
PFHpA	( 363.0 / 319.0 ) 1046730 ( 363.0 / 169.0 ) 341278	( 6.31 , 1.00 ) ( 0.00 , N/A , -0.1 )	16010.9 7246.9	0.3260 101.7 101.7	1.9309 [ 2.0000 ]	96.5%			
PFOA	( 413.0 / 369.0 ) 1332664 ( 413.0 / 169.0 ) 469250	( 7.01 , 1.00 ) ( 0.00 , N/A , 0.0 )	6220.3 2713549.8	0.3521 105.2 105.2	2.0866 [ 2.0000 ]	104.3%			
PFNA	( 463.0 / 419.0 ) 1137106 ( 463.0 / 169.0 ) 290595	( 7.61 , 1.00 ) ( 0.00 , N/A , 0.0 )	8852.4 4937.2	0.2556 114.2 114.2	2.0394 [ 2.0000 ]	102.0%			
PFDA	( 513.0 / 469.0 ) 1436390 ( 513.0 / 169.0 ) 173502	( 8.17 , 1.00 ) ( 0.00 , N/A , -0.2 )	1533.1 2906.4	0.1208 99.3 99.3	2.1390 [ 2.0000 ]	107.0%			
PFUnA	( 563.0 / 519.0 ) 1214746 ( 563.0 / 169.0 ) 160306	( 8.64 , 1.00 ) ( 0.00 , N/A , 0.1 )	1539.9 3496.0	0.1320 125.9 125.9	1.9413 [ 2.0000 ]	97.1%			
PFDaA	( 613.0 / 569.0 ) 1115680 ( 613.0 / 169.0 ) 230062	( 8.92 , 1.00 ) ( 0.00 , N/A , 0.2 )	1334.3 600.4	0.2062 126.1 126.1	2.0539 [ 2.0000 ]	102.7%			
PFTTrDA	( 663.0 / 619.0 ) 1066226 ( 663.0 / 169.0 ) 270733	( 9.12 , 1.02 ) ( N/A , 0.00 , -0.1 )	1920.4 3654.3	0.2539 106.2 106.2	2.1679 [ 2.0000 ]	108.4%			
PFTeDA	( 713.0 / 669.0 ) 1064969 ( 713.0 / 169.0 ) 204784	( 9.29 , 1.00 ) ( 0.00 , N/A , 0.2 )	1633.8 783.2	0.1923 93.0 93.0	2.0298 [ 2.0000 ]	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
PFBS	( 299.0 / 80.0 ) 2223132 ( 299.0 / 99.0 ) 1362361	( 5.46 , 1.00 ) ( 0.00 , N/A , 0.0 )	3112.7 2641.0	0.6128 94.8 94.8	1.8073 [ 1.7695 ]	102.1%			
PFPeS	( 349.0 / 80.0 ) 3780216 ( 349.0 / 99.0 ) 1256747	( 6.39 , 0.89 ) ( N/A , 0.02 , -0.1 )	1878642.0 330746896.9	0.3325 93.1 93.1	1.9706 [ 1.8768 ]	105.0%			
PFHxS	( 399.0 / 80.0 ) 2792019 ( 399.0 / 99.0 ) 984345	( 7.16 , 1.00 ) ( 0.00 , N/A , 0.0 )	139093.1 2342.3	0.3526 108.6 108.6	1.8254 [ 1.8220 ]	100.2%			
PFHpS	( 449.0 / 80.0 ) 3285266 ( 449.0 / 99.0 ) 948388	( 7.80 , 0.93 ) ( N/A , 0.02 , 0.1 )	15910125.3 18002.5	0.2887 106.0 106.0	1.9219 [ 1.9028 ]	101.0%			
PFOS	( 499.0 / 80.0 ) 4400886 ( 499.0 / 99.0 ) 965758	( 8.37 , 1.00 ) ( 0.00 , N/A , -0.1 )	484.0 429.4	0.2194 99.2 99.2	1.8603 [ 1.8550 ]	100.3%			
PFNS	( 549.0 / 80.0 ) 5077665 ( 549.0 / 99.0 ) 1142350	( 8.78 , 1.05 ) ( N/A , 0.01 , 0.1 )	12572.7 141808.6	0.2250 102.6 102.6	1.9098 [ 1.9198 ]	99.5%			
PFDS	( 599.0 / 80.0 ) 5627897 ( 599.0 / 99.0 ) 1220356	( 9.02 , 1.08 ) ( N/A , 0.00 , 0.0 )	3007.5 2088.2	0.2168 99.5 99.5	1.9460 [ 1.9262 ]	101.0%			
PFDoS	( 699.0 / 80.0 ) 3473138 ( 699.0 / 99.0 ) 799506	( 9.37 , 1.12 ) ( N/A , 0.00 , 0.0 )	3432.2 1574.3	0.2302 110.6 110.6	1.8829 [ 1.9391 ]	97.1%			
4:2FTS	( 327.0 / 307.0 ) 2147611 ( 327.0 / 81.0 ) 1465550	( 5.22 , 1.00 ) ( 0.00 , N/A , 0.0 )	4169.6 2224.4	0.6824 101.1 101.1	8.0095 [ 7.4762 ]	107.1%			
6:2FTS	( 427.0 / 407.0 ) 1217522 ( 427.0 / 81.0 ) 905821	( 6.74 , 1.00 ) ( 0.00 , N/A , 0.1 )	2807.5 1230.9	0.7440 89.9 89.9	8.2518 [ 7.5923 ]	108.7%			
8:2FTS	( 527.0 / 507.0 ) 1325429 ( 527.0 / 81.0 ) 1143451	( 7.91 , 1.00 ) ( 0.00 , N/A , 0.0 )	1768.2 1699.8	0.8627 103.8 103.8	8.0433 [ 7.6663 ]	104.9%			

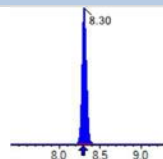
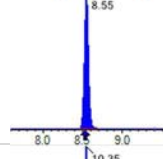
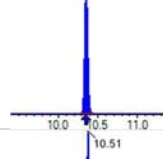
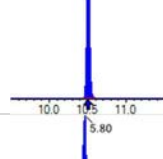

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	( 498.0 / 78.0 ) 7454165 ( 498.0 / 478.0 ) 153863	( 9.83 , 1.00 ) ( 0.00 , N/A , 0.1 )	3632.9 1289.8	0.0206 100.5 100.5	2.1877 [ 2.0000 ]	109.4%			
NMeFOSA	( 512.0 / 219.0 ) 6581539 ( 512.0 / 169.0 ) 5434702	( 10.42 , 1.00 ) ( 0.00 , N/A , 0.9 )	4400.8 4035.5	0.8257 100.8 100.8	8.0911 [ 8.0000 ]	101.1%			
NEIFOSA	( 526.0 / 219.0 ) 7369810 ( 526.0 / 169.0 ) 9562461	( 10.56 , 1.00 ) ( 0.00 , N/A , 0.9 )	10194.3 8766.4	1.2975 102.2 102.2	8.3766 [ 8.0000 ]	104.7%			
NMeFOSAA	( 570.0 / 419.0 ) 558224 ( 570.0 / 483.0 ) 249670	( 8.31 , 1.00 ) ( 0.00 , N/A , -0.1 )	3385.1 183.6	0.4473 94.2 94.2	2.1736 [ 2.0000 ]	108.7%			
NEIFOSAA	( 584.0 / 419.0 ) 468621 ( 584.0 / 526.0 ) 259589	( 8.55 , 1.00 ) ( 0.01 , N/A , 0.2 )	253752.3 16601.7	0.5539 108.6 108.6	1.9370 [ 2.0000 ]	96.9%			
NMeFOSE	( 616.0 / 59.0 ) 2915088	( 10.36 , 1.00 ) ( 0.01 , N/A , 0.0 )	2699.4	N/A 0.0 0.0	7.9620 [ 8.0000 ]	99.5%			
NEIFOSE	( 630.0 / 59.0 ) 3776358	( 10.52 , 1.00 ) ( 0.01 , N/A , 0.0 )	1374.9	N/A 0.0 0.0	8.1217 [ 8.0000 ]	101.5%			
HFPO-DA	( 285.0 / 169.0 ) 1383460 ( 285.0 / 185.0 ) 4010692	( 5.80 , 1.00 ) ( 0.00 , N/A , 0.0 )	2428.0 3386.2	2.8990 105.3 105.3	4.0649 [ 4.0000 ]	101.6%			
ADONA	( 377.0 / 85.0 ) 4962626 ( 377.0 / 251.0 ) 457532	( 6.59 , 1.14 ) ( N/A , 0.02 , -0.1 )	3737.6 918.9	0.0922 104.6 104.6	4.2400 [ 3.7708 ]	112.4%			
9CI-PF3ONS	( 531.0 / 351.0 ) 14406479 ( 533.0 / 353.0 ) 4641144	( 8.70 , 1.50 ) ( N/A , 0.01 , -0.1 )	3319.3 3063.0	0.3222 92.5 92.5	3.9280 [ 3.7330 ]	105.2%			
11CI-PF3OUDS	( 631.0 / 451.0 ) 8500924 ( 633.0 / 453.0 ) 2975725	( 9.17 , 1.58 ) ( N/A , 0.00 , 0.2 )	3323.4 2149.4	0.3500 96.7 96.7	4.4584 [ 3.7728 ]	118.2%			

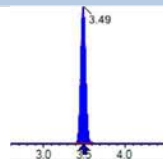
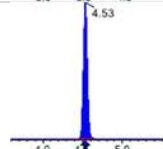
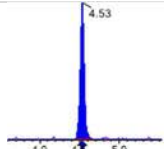
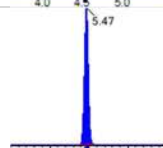
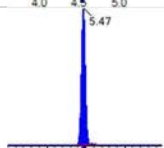
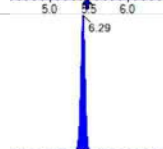
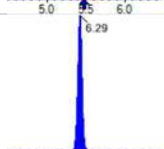
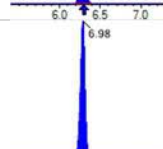
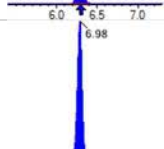
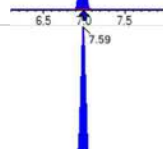
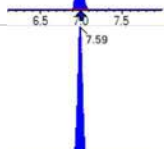
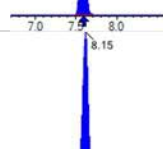
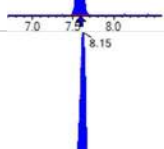
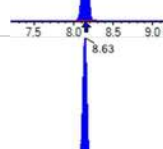
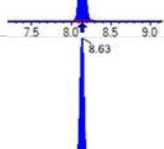
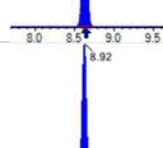
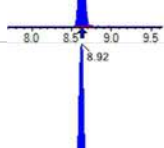
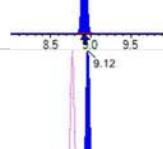
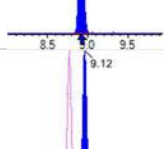
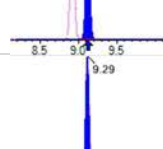
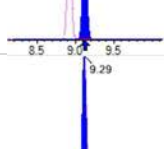
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T. ) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	( 241.0 / 177.0 ) 98166 ( 241.0 / 117.0 ) 149343	( 4.06 , 0.89 ) ( N/A , 0.02 , 0.1 )	980.2 901.4	1.5213 96.4 96.4	8.1739 [ 8.0000 ]	102.2%			
5:3FTCA	( 341.0 / 236.7 ) 659057 ( 341.0 / 217.0 ) 1151481	( 6.05 , 1.10 ) ( N/A , 0.02 , -0.1 )	1706.3 1760.2	1.7472 102.5 102.5	7.2959 [ 8.0000 ]	91.2%			
7:3FTCA	( 441.0 / 317.0 ) 1221396 ( 441.0 / 337.0 ) 1047529	( 7.90 , 1.44 ) ( N/A , 0.02 , 0.2 )	612.9 919.3	0.8576 99.7 99.7	7.3193 [ 8.0000 ]	91.5%			
PFEESA	( 315.0 / 135.0 ) 2916970 ( 315.0 / 83.0 ) 815959	( 5.91 , 1.08 ) ( N/A , 0.02 , -0.1 )	3510.3 2085.3	0.2797 109.5 109.5	3.6759 [ 3.5698 ]	103.0%			
PFMPA	( 229.0 / 85.0 ) 425241	( 3.89 , 0.86 ) ( N/A , 0.01 , 0.0 )	2450.5	N/A 0.0 0.0	3.9053 [ 4.0000 ]	97.6%			
PFMBA	( 279.0 / 85.0 ) 1832503	( 4.86 , 1.07 ) ( N/A , 0.02 , 0.0 )	3046.2	N/A 0.0 0.0	4.0131 [ 4.0000 ]	100.3%			
NFDHA	( 295.0 / 201.0 ) 1471114 ( 295.0 / 85.0 ) 1539319	( 5.39 , 0.98 ) ( N/A , 0.02 , 0.2 )	2648.0 2571.6	1.0464 104.5 104.5	4.0123 [ 4.0000 ]	100.3%			
13C3_PFBA_IIS	( 216.0 / 172.0 ) 258935	( 3.50 , N/A ) ( N/A , 0.01 , N/A )	1459.6	N/A	1.1560 [ 1.0000 ]	115.6% { 114.9% }			
13C2_PFHxA_IIS	( 315.0 / 270.0 ) 527362	( 5.49 , N/A ) ( N/A , 0.02 , N/A )	2491.9	N/A	1.0447 [ 1.0000 ]	104.5% { 109.8% }			
13C4_PFOA_IIS	( 417.0 / 372.0 ) 724530	( 7.01 , N/A ) ( N/A , 0.03 , N/A )	1417.5	N/A	1.1432 [ 1.0000 ]	114.3% { 122.1% }			
13C5_PFNA_IIS	( 468.0 / 423.0 ) 687055	( 7.61 , N/A ) ( N/A , 0.02 , N/A )	10320.7	N/A	1.1370 [ 1.0000 ]	113.7% { 115.9% }			

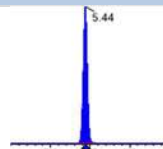
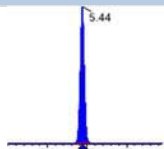
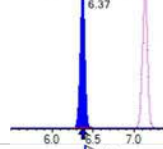
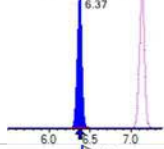
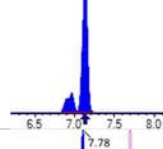
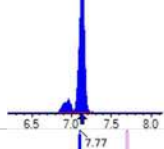
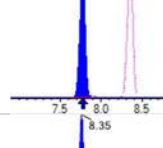
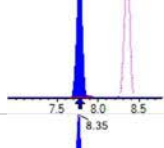
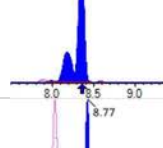
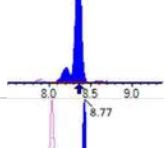
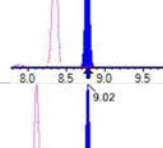
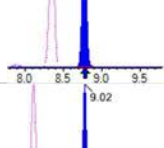
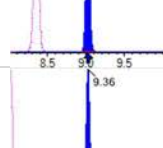
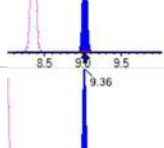
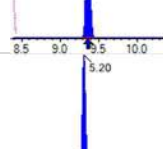
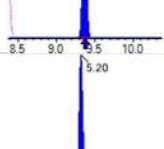
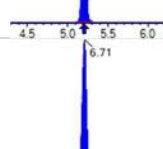
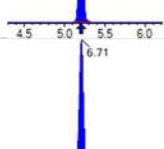
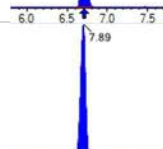
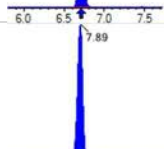
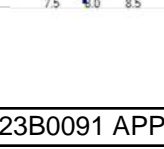
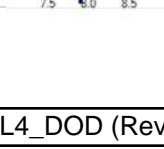
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	( 515.0 / 470.1 ) 642605	( 8.17, N/A ) ( N/A, 0.02, N/A )	3000.0	N/A	1.1105 [ 1.0000 ]	111.1% { 118.0% }			
18O2_PFHxS_IIS	( 403.0 / 83.9 ) 945086	( 7.16, N/A ) ( N/A, 0.03, N/A )	1162.6	N/A	1.1036 [ 1.0000 ]	110.4% { 111.9% }			
13C4_PFOS_IIS	( 503.0 / 79.9 ) 1662514	( 8.37, N/A ) ( N/A, 0.01, N/A )	1471.1	N/A	1.1044 [ 1.0000 ]	110.4% { 109.9% }			
13C4_PFBA_EIS	( 217.0 / 172.0 ) 2387665	( 3.50, N/A ) ( N/A, 0.01, N/A )	5160.5	N/A	7.7521 [ 8.0000 ]	96.9% { 102.3% }			
13C5_PFPeA_EIS	( 268.0 / 223.0 ) 2423641	( 4.55, N/A ) ( N/A, 0.02, N/A )	3349.2	N/A	4.2004 [ 4.0000 ]	105.0% { 100.2% }			
13C5_PFHxA_EIS	( 318.0 / 273.0 ) 1502798	( 5.49, N/A ) ( N/A, 0.02, N/A )	3626.2	N/A	2.0804 [ 2.0000 ]	104.0% { 97.9% }			
13C4_PFHpA_EIS	( 367.0 / 322.0 ) 1283933	( 6.31, N/A ) ( N/A, 0.02, N/A )	2445.9	N/A	2.0514 [ 2.0000 ]	102.6% { 101.6% }			
13C8_PFOA_EIS	( 421.0 / 376.0 ) 1434645	( 7.01, N/A ) ( N/A, 0.03, N/A )	2756.5	N/A	1.7484 [ 2.0000 ]	87.4% { 93.1% }			
13C9_PFNA_EIS	( 472.0 / 427.0 ) 630189	( 7.61, N/A ) ( N/A, 0.02, N/A )	1674.8	N/A	0.8657 [ 1.0000 ]	86.6% { 89.7% }			
13C6_PFDA_EIS	( 519.0 / 474.0 ) 763399	( 8.16, N/A ) ( N/A, 0.02, N/A )	2225.3	N/A	0.9422 [ 1.0000 ]	94.2% { 98.3% }			
13C7_PFUnA_EIS	( 570.0 / 525.0 ) 753151	( 8.64, N/A ) ( N/A, 0.01, N/A )	13028.6	N/A	0.9348 [ 1.0000 ]	93.5% { 97.6% }			

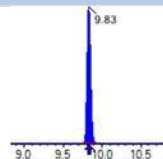
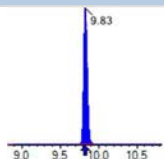
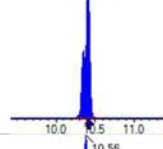
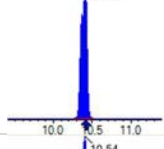
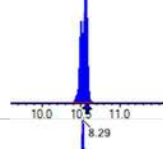
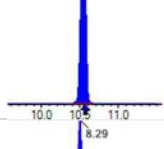
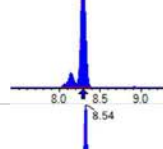
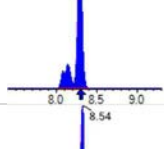
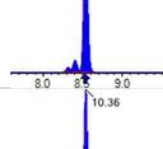
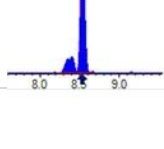
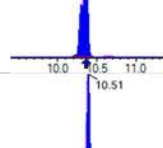
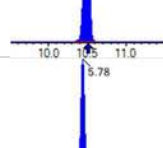
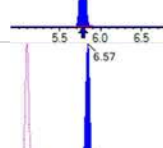
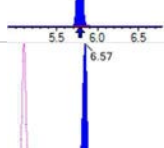
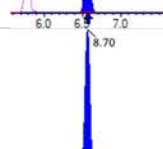
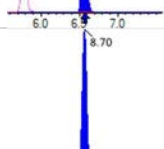
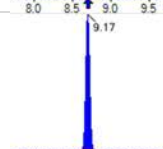
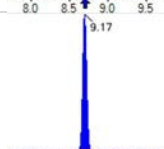

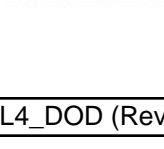
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) ( $\Delta$ RT-I[min], $\Delta$ RT- CV[min], $\Delta$ RT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	( 615.0 / 570.0 ) 617721	( 8.92, N/A ) ( N/A, 0.01, N/A )	2492.7	N/A	0.8939 [ 1.0000 ]	89.4% { 94.0% }			
13C2_PFTeDA_EIS	( 715.0 / 670.0 ) 606299	( 9.29, N/A ) ( N/A, 0.00, N/A )	1178.7	N/A	0.9575 [ 1.0000 ]	95.8% { 99.4% }			
13C3_PFBs_EIS	( 302.0 / 80.0 ) 3958718	( 5.46, N/A ) ( N/A, 0.02, N/A )	3417.7	N/A	2.0014 [ 2.0000 ]	100.1% { 100.2% }			
13C3_PFHxS_EIS	( 402.0 / 80.0 ) 1989400	( 7.16, N/A ) ( N/A, 0.02, N/A )	2679.8	N/A	1.9364 [ 2.0000 ]	96.8% { 100.0% }			
13C8_PFOS_EIS	( 507.0 / 80.0 ) 4487002	( 8.37, N/A ) ( N/A, 0.02, N/A )	1488.1	N/A	1.9659 [ 2.0000 ]	98.3% { 101.5% }			
13C2_4:2FTS_EIS	( 329.0 / 81.0 ) 392697	( 5.22, N/A ) ( N/A, 0.02, N/A )	1473.7	N/A	3.5968 [ 4.0000 ]	89.9% { 95.9% }			
13C2_6:2FTS_EIS	( 429.0 / 81.0 ) 415946	( 6.74, N/A ) ( N/A, 0.03, N/A )	1899.8	N/A	3.4707 [ 4.0000 ]	86.8% { 90.8% }			
13C2_8:2FTS_EIS	( 529.0 / 81.0 ) 605135	( 7.91, N/A ) ( N/A, 0.02, N/A )	1540.6	N/A	3.5739 [ 4.0000 ]	89.3% { 100.0% }			
13C8_PFOsa_EIS	( 506.0 / 78.0 ) 8520316	( 9.84, N/A ) ( N/A, 0.00, N/A )	3894.3	N/A	2.0111 [ 2.0000 ]	100.6% { 99.4% }			
D3_NMeFOsa_EIS	( 515.0 / 169.0 ) 1919031	( 10.41, N/A ) ( N/A, 0.00, N/A )	2947.3	N/A	1.9225 [ 2.0000 ]	96.1% { 105.2% }			
D5_NEtFOsa_EIS	( 531.0 / 169.0 ) 1883486	( 10.56, N/A ) ( N/A, 0.00, N/A )	3509.4	N/A	2.1087 [ 2.0000 ]	105.4% { 109.3% }			

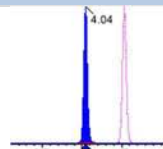
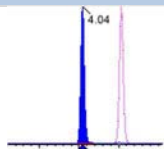
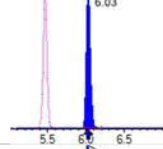
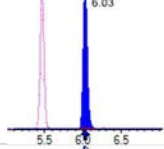
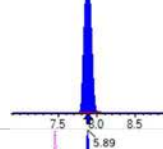
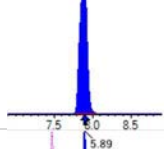
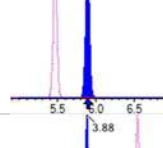
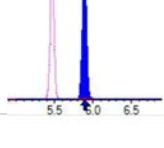
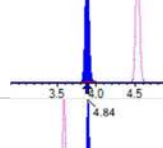
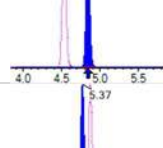
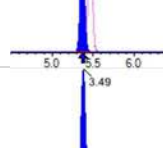
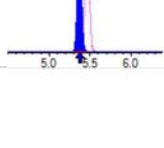
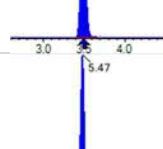
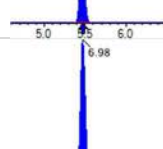
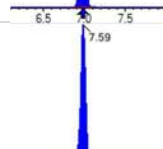



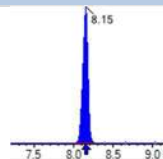
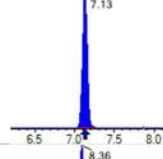
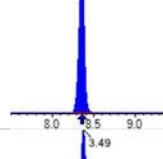
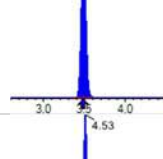
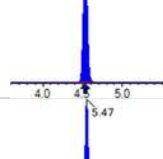
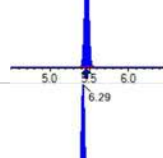
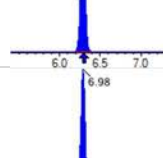
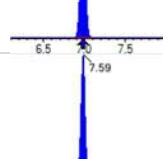
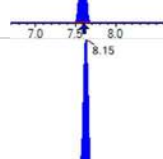
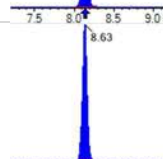
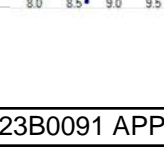
Analyte	( Q1 / Q3 ) Area Counts*min	R.T. ( R.T [min] , R.R.T. ) ( ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	( 573.0 / 419.0 ) 1309006	( 8.30 , N/A ) ( N/A , 0.02 , N/A )	1009.4	N/A	3.5440 [ 4.0000 ]	88.6% { 87.9% }			
D5_EtFOSAA_EIS	( 589.0 / 419.0 ) 1152185	( 8.55 , N/A ) ( N/A , 0.01 , N/A )	15689.4	N/A	3.8097 [ 4.0000 ]	95.2% { 102.7% }			
D7_NMeFOSE_EIS	( 623.0 / 58.9 ) 7324449	( 10.35 , N/A ) ( N/A , 0.00 , N/A )	2458.1	N/A	20.1274 [ 20.0000 ]	100.6% { 101.9% }			
D9_NEtFOSE_EIS	( 639.0 / 58.9 ) 9772079	( 10.51 , N/A ) ( N/A , 0.00 , N/A )	2034.7	N/A	20.9689 [ 20.0000 ]	104.8% { 102.6% }			
13C3_HFPODA_EIS	( 287.0 / 169.0 ) 3443023	( 5.80 , N/A ) ( N/A , 0.02 , N/A )	4078.3	N/A	7.8284 [ 8.0000 ]	97.9% { 94.0% }			

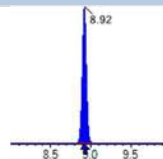
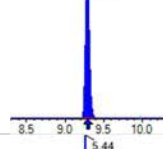
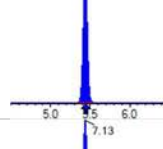
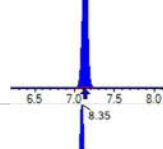
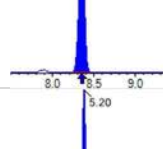
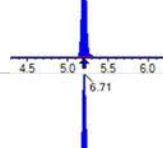
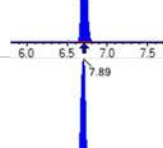
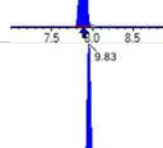
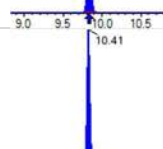
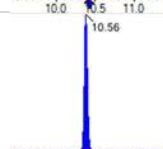

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	( 213.0 / 169.0 ) 4625962	( 3.49 , 1.00 ) ( 0.00 , N/A , 0.0 )	269.9	N/A 0.0 0.0	19.1353 [ 20.0000 ]	95.7%			
PFPeA	( 263.0 / 219.0 ) 4849452 ( 263.0 / 69.0 ) 51930	( 4.53 , 1.00 ) ( 0.00 , N/A , 0.1 )	6300.0 525.7	0.0107 100.0 100.0	9.7482 [ 10.0000 ]	97.5%			
PFHxA	( 313.0 / 269.0 ) 3231153 ( 313.0 / 119.0 ) 321815	( 5.47 , 1.00 ) ( 0.00 , N/A , 0.0 )	7539.0 569520.8	0.0996 100.0 100.0	4.7905 [ 5.0000 ]	95.8%			
PFHpA	( 363.0 / 319.0 ) 2374219 ( 363.0 / 169.0 ) 761184	( 6.29 , 1.00 ) ( 0.00 , N/A , 0.0 )	27517.9 4626.3	0.3206 100.0 100.0	4.4501 [ 5.0000 ]	89.0%			
PFOA	( 413.0 / 369.0 ) 3119950 ( 413.0 / 169.0 ) 1043952	( 6.98 , 1.00 ) ( 0.00 , N/A , 0.1 )	11212.8 2366.5	0.3346 100.0 100.0	4.5495 [ 5.0000 ]	91.0%			
PFNA	( 463.0 / 419.0 ) 2777308 ( 463.0 / 169.0 ) 621436	( 7.59 , 1.00 ) ( 0.00 , N/A , 0.1 )	143516.6 7999265.7	0.2238 100.0 100.0	4.4669 [ 5.0000 ]	89.3%			
PFDA	( 513.0 / 469.0 ) 3288150 ( 513.0 / 169.0 ) 399859	( 8.15 , 1.00 ) ( 0.00 , N/A , -0.2 )	2295.7 8100.9	0.1216 100.0 100.0	4.8121 [ 5.0000 ]	96.2%			
PFUnA	( 563.0 / 519.0 ) 2957188 ( 563.0 / 169.0 ) 310031	( 8.63 , 1.00 ) ( 0.00 , N/A , 0.2 )	2421.9 3060.8	0.1048 100.0 100.0	4.6144 [ 5.0000 ]	92.3%			
PFDoA	( 613.0 / 569.0 ) 2625478 ( 613.0 / 169.0 ) 429428	( 8.92 , 1.00 ) ( 0.00 , N/A , 0.0 )	2165.7 1615.7	0.1636 100.0 100.0	4.5454 [ 5.0000 ]	90.9%			
PFTTrDA	( 663.0 / 619.0 ) 2589019 ( 663.0 / 169.0 ) 618790	( 9.12 , 1.02 ) ( N/A , 0.00 , 0.1 )	2797.5 1841.7	0.2390 100.0 100.0	4.9505 [ 5.0000 ]	99.0%			
PFTeDA	( 713.0 / 669.0 ) 2457334 ( 713.0 / 169.0 ) 507818	( 9.29 , 1.00 ) ( 0.00 , N/A , 0.1 )	3479.2 625.2	0.2067 100.0 100.0	4.6552 [ 5.0000 ]	93.1%			

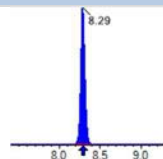
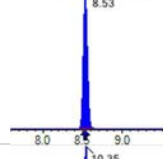
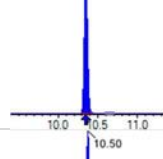
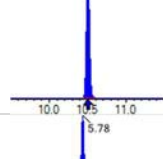
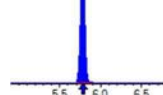
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	( 299.0 / 80.0 ) 5078337 ( 299.0 / 99.0 ) 3283985	( 5.44 , 1.00 ) ( 0.00 , N/A , 0.0 )	4191.7 3173.0	0.6467 100.0 100.0	4.1378 [ 4.4237 ]	93.5%			
PFPeS	( 349.0 / 80.0 ) 8825978 ( 349.0 / 99.0 ) 3150960	( 6.37 , 0.89 ) ( N/A , 0.00 , 0.0 )	52478.2 43714.8	0.3570 100.0 100.0	4.5997 [ 4.6919 ]	98.0%			
PFHxS	( 399.0 / 80.0 ) 6499323 ( 399.0 / 99.0 ) 2110618	( 7.13 , 1.00 ) ( 0.00 , N/A , 0.1 )	4671.8 15342595.9	0.3247 100.0 100.0	4.2481 [ 4.5549 ]	93.3%			
PFHpS	( 449.0 / 80.0 ) 7537575 ( 449.0 / 99.0 ) 2051876	( 7.78 , 0.93 ) ( N/A , 0.00 , 0.1 )	18633.8 4259.7	0.2722 100.0 100.0	4.4756 [ 4.7570 ]	94.1%			
PFOS	( 499.0 / 80.0 ) 10091750 ( 499.0 / 99.0 ) 2233382	( 8.35 , 1.00 ) ( 0.00 , N/A , 0.0 )	1680.0 1507.9	0.2213 100.0 100.0	4.3298 [ 4.6375 ]	93.4%			
PFNS	( 549.0 / 80.0 ) 12319974 ( 549.0 / 99.0 ) 2701138	( 8.77 , 1.05 ) ( N/A , 0.00 , 0.0 )	41212.2 30751.4	0.2192 100.0 100.0	4.7034 [ 4.7994 ]	98.0%			
PFDS	( 599.0 / 80.0 ) 13893059 ( 599.0 / 99.0 ) 3028577	( 9.02 , 1.08 ) ( N/A , 0.00 , 0.0 )	4249.5 2452.4	0.2180 100.0 100.0	4.8759 [ 4.8155 ]	101.3%			
PFDoS	( 699.0 / 80.0 ) 9237910 ( 699.0 / 99.0 ) 1922828	( 9.36 , 1.12 ) ( N/A , 0.00 , -0.1 )	3957.0 2185.3	0.2081 100.0 100.0	5.0833 [ 4.8478 ]	104.9%			
4:2FTS	( 327.0 / 307.0 ) 4991486 ( 327.0 / 81.0 ) 3368015	( 5.20 , 1.00 ) ( 0.00 , N/A , 0.0 )	4068.6 2907.8	0.6748 100.0 100.0	17.8604 [ 18.6906 ]	95.6%			
6:2FTS	( 427.0 / 407.0 ) 2761537 ( 427.0 / 81.0 ) 2285757	( 6.71 , 1.00 ) ( 0.00 , N/A , 0.0 )	1751.6 2382.2	0.8277 100.0 100.0	16.9949 [ 18.9808 ]	89.5%			
8:2FTS	( 527.0 / 507.0 ) 3235546 ( 527.0 / 81.0 ) 2689124	( 7.89 , 1.00 ) ( 0.00 , N/A , 0.0 )	2836.4 1954.5	0.8311 100.0 100.0	19.6365 [ 19.1658 ]	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
PFOSA	( 498.0 / 78.0 ) 16873776 ( 498.0 / 478.0 ) 346600	( 9.83 , 1.00 ) ( 0.00 , N/A , 0.1 )	3623.3 1365.4	0.0205 100.0 100.0	4.9230 [ 5.0000 ]	98.5%			
NMeFOSA	( 512.0 / 219.0 ) 15062148 ( 512.0 / 169.0 ) 12339105	( 10.41 , 1.00 ) ( 0.00 , N/A , 1.0 )	4316.3 4090.2	0.8192 100.0 100.0	19.8909 [ 20.0000 ]	99.5%			
NEIFOSA	( 526.0 / 219.0 ) 16221852 ( 526.0 / 169.0 ) 20591347	( 10.56 , 1.00 ) ( 0.00 , N/A , 0.8 )	12815.2 15235.1	1.2694 100.0 100.0	20.1464 [ 20.0000 ]	100.7%			
NMeFOSAA	( 570.0 / 419.0 ) 1293168 ( 570.0 / 483.0 ) 613882	( 8.29 , 1.00 ) ( 0.00 , N/A , -0.2 )	3254.9 276085.0	0.4747 100.0 100.0	4.4279 [ 5.0000 ]	88.6%			
NEIFOSAA	( 584.0 / 419.0 ) 1211099 ( 584.0 / 526.0 ) 617983	( 8.54 , 1.00 ) ( 0.01 , N/A , 0.1 )	1922.1 7076.5	0.5103 100.0 100.0	5.1422 [ 5.0000 ]	102.8%			
NMeFOSE	( 616.0 / 59.0 ) 6860263	( 10.36 , 1.00 ) ( 0.01 , N/A , 0.0 )	3314.5	N/A 0.0 0.0	19.0869 [ 20.0000 ]	95.4%			
NEIFOSE	( 630.0 / 59.0 ) 8750560	( 10.51 , 1.00 ) ( 0.01 , N/A , 0.0 )	1947.4	N/A 0.0 0.0	19.3134 [ 20.0000 ]	96.6%			
HFPO-DA	( 285.0 / 169.0 ) 3384631 ( 285.0 / 185.0 ) 9322395	( 5.78 , 1.00 ) ( 0.00 , N/A , 0.0 )	3354.6 4730.4	2.7543 100.0 100.0	9.3521 [ 10.0000 ]	93.5%			
ADONA	( 377.0 / 85.0 ) 11635291 ( 377.0 / 251.0 ) 1025920	( 6.57 , 1.14 ) ( N/A , 0.00 , -0.1 )	5483.9 2252.2	0.0882 100.0 100.0	9.3484 [ 9.4270 ]	99.2%			
9CI-Pf3ONS	( 531.0 / 351.0 ) 31179230 ( 533.0 / 353.0 ) 10857577	( 8.70 , 1.50 ) ( N/A , 0.00 , 0.0 )	3496.7 3130.1	0.3482 100.0 100.0	8.3839 [ 9.3325 ]	89.8%			
11CI-PF3OUDS	( 631.0 / 451.0 ) 18911923 ( 633.0 / 453.0 ) 6842706	( 9.17 , 1.59 ) ( N/A , 0.00 , 0.0 )	2197.8 3639.0	0.3618 100.0 100.0	9.3273 [ 9.4321 ]	98.9%			

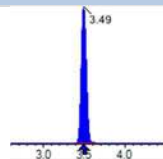
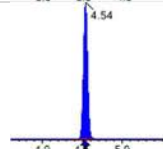
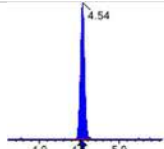
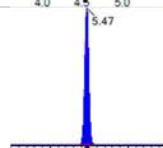
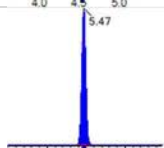
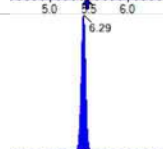
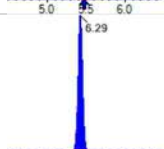
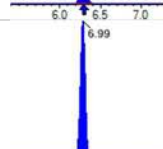
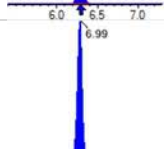
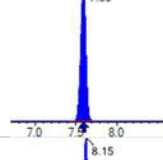
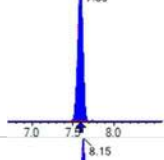
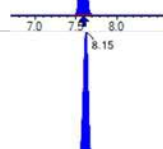
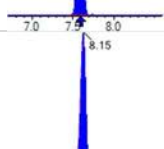
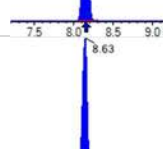
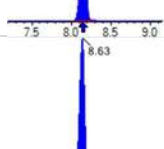
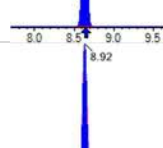
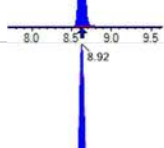
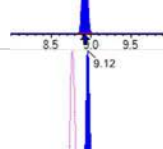
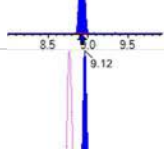
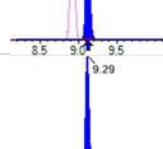
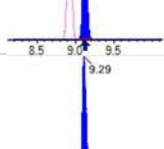
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	( 241.0 / 177.0 ) 221117 ( 241.0 / 117.0 ) 348920	( 4.04 , 0.89 ) ( N/A , 0.00 , 0.0 )	1710.7 1270.6	1.5780 100.0 100.0	18.0210 [ 20.0000 ]	90.1%			
5:3FTCA	( 341.0 / 236.7 ) 1559646 ( 341.0 / 217.0 ) 2659637	( 6.03 , 1.10 ) ( N/A , 0.00 , 0.0 )	1471.7 2247.5	1.7053 100.0 100.0	16.9094 [ 20.0000 ]	84.5%			
7:3FTCA	( 441.0 / 317.0 ) 2829193 ( 441.0 / 337.0 ) 2434614	( 7.88 , 1.44 ) ( N/A , 0.00 , -0.2 )	1104.9 1471.1	0.8605 100.0 100.0	16.6044 [ 20.0000 ]	83.0%			
PFEESA	( 315.0 / 135.0 ) 6867136 ( 315.0 / 83.0 ) 1753875	( 5.89 , 1.08 ) ( N/A , 0.00 , 0.1 )	4733.3 2624.5	0.2554 100.0 100.0	8.4754 [ 8.9246 ]	95.0%			
PFMPA	( 229.0 / 85.0 ) 1012996	( 3.88 , 0.86 ) ( N/A , 0.00 , 0.0 )	4456.4	N/A 0.0 0.0	9.3225 [ 10.0000 ]	93.2%			
PFMBA	( 279.0 / 85.0 ) 4261572	( 4.84 , 1.07 ) ( N/A , 0.00 , 0.0 )	4291.0	N/A 0.0 0.0	9.3520 [ 10.0000 ]	93.5%			
NFDHA	( 295.0 / 201.0 ) 3469986 ( 295.0 / 85.0 ) 3474472	( 5.37 , 0.98 ) ( N/A , 0.00 , 0.0 )	3832.4 3353.4	1.0013 100.0 100.0	9.2688 [ 10.0000 ]	92.7%			
13C3_PFBA_IIS	( 216.0 / 172.0 ) 225438	( 3.49 , N/A ) ( N/A , 0.00 , N/A )	1151.4	N/A	1.0065 [ 1.0000 ]	100.6% { 100.0% }			
13C2_PFHxA_IIS	( 315.0 / 270.0 ) 480171	( 5.47 , N/A ) ( N/A , 0.00 , N/A )	1976.3	N/A	0.9512 [ 1.0000 ]	95.1% { 100.0% }			
13C4_PFOA_IIS	( 417.0 / 372.0 ) 593445	( 6.98 , N/A ) ( N/A , 0.00 , N/A )	1655.7	N/A	0.9364 [ 1.0000 ]	93.6% { 100.0% }			
13C5_PFNA_IIS	( 468.0 / 423.0 ) 592832	( 7.59 , N/A ) ( N/A , 0.00 , N/A )	3326.2	N/A	0.9811 [ 1.0000 ]	98.1% { 100.0% }			

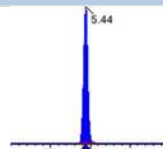
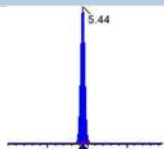
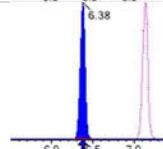
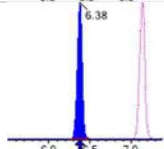
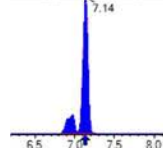
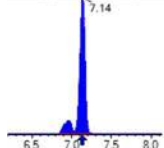
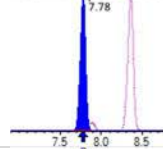
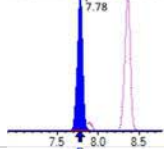
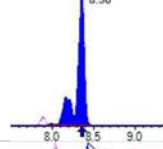
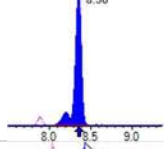
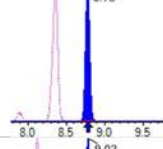
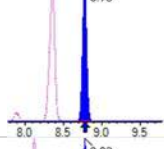
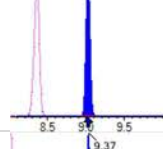
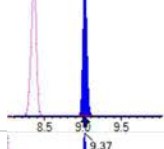
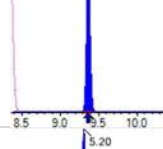
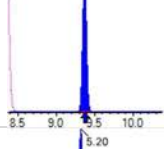
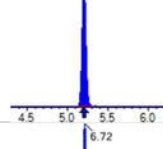
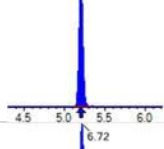
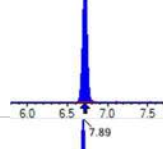
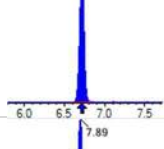
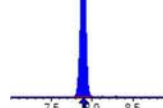
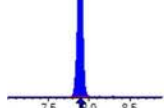
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	( 515.0 / 470.1 ) 544568	( 8.15 , N/A ) ( N/A , 0.00 , N/A )	905.1	N/A	0.9411 [ 1.0000 ]	94.1% { 100.0% }			
18O2_PFHxS_IIS	( 403.0 / 83.9 ) 844212	( 7.13 , N/A ) ( N/A , 0.00 , N/A )	1879.8	N/A	0.9858 [ 1.0000 ]	98.6% { 100.0% }			
13C4_PFOS_IIS	( 503.0 / 79.9 ) 1512102	( 8.36 , N/A ) ( N/A , 0.00 , N/A )	1058.0	N/A	1.0045 [ 1.0000 ]	100.4% { 100.0% }			
13C4_PFBA_EIS	( 217.0 / 172.0 ) 2334765	( 3.49 , N/A ) ( N/A , 0.00 , N/A )	4444.3	N/A	8.7067 [ 8.0000 ]	108.8% { 100.0% }			
13C5_PFPeA_EIS	( 268.0 / 223.0 ) 2418611	( 4.53 , N/A ) ( N/A , 0.00 , N/A )	3591.2	N/A	4.6036 [ 4.0000 ]	115.1% { 100.0% }			
13C5_PFHxA_EIS	( 318.0 / 273.0 ) 1534443	( 5.47 , N/A ) ( N/A , 0.00 , N/A )	3297.1	N/A	2.3330 [ 2.0000 ]	116.7% { 100.0% }			
13C4_PFHpA_EIS	( 367.0 / 322.0 ) 1263643	( 6.29 , N/A ) ( N/A , 0.00 , N/A )	2333.7	N/A	2.2174 [ 2.0000 ]	110.9% { 100.0% }			
13C8_PFOA_EIS	( 421.0 / 376.0 ) 1540426	( 6.98 , N/A ) ( N/A , 0.00 , N/A )	3544.1	N/A	2.2920 [ 2.0000 ]	114.6% { 100.0% }			
13C9_PFNA_EIS	( 472.0 / 427.0 ) 702736	( 7.59 , N/A ) ( N/A , 0.00 , N/A )	1753.9	N/A	1.1188 [ 1.0000 ]	111.9% { 100.0% }			
13C6_PFDA_EIS	( 519.0 / 474.0 ) 776808	( 8.15 , N/A ) ( N/A , 0.00 , N/A )	1570.3	N/A	1.1314 [ 1.0000 ]	113.1% { 100.0% }			
13C7_PFUnA_EIS	( 570.0 / 525.0 ) 771339	( 8.63 , N/A ) ( N/A , 0.00 , N/A )	4873.0	N/A	1.1297 [ 1.0000 ]	113.0% { 100.0% }			

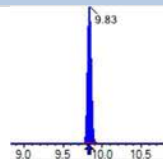
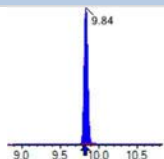
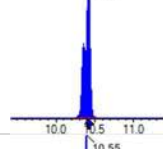
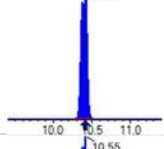
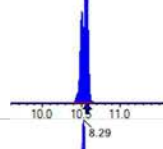
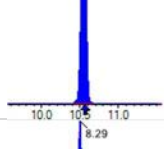
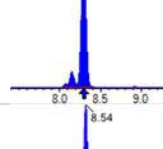
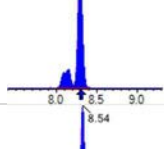
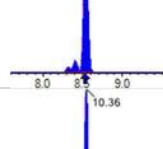
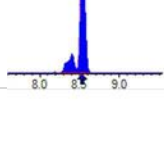
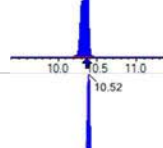
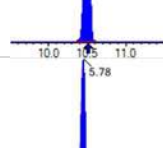
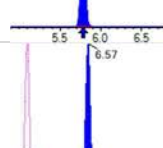
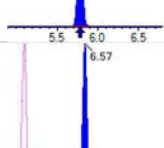
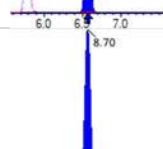
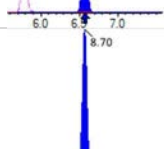
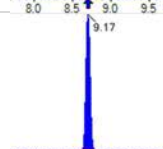
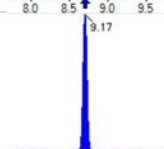
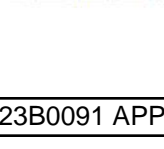
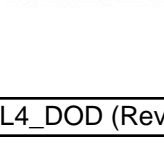
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	( 615.0 / 570.0 ) 656853	( 8.92 , N/A ) ( N/A , 0.00 , N/A )	1995.2	N/A	1.1217 [ 1.0000 ]	112.2% { 100.0% }			
13C2_PFTeDA_EIS	( 715.0 / 670.0 ) 610004	( 9.29 , N/A ) ( N/A , 0.00 , N/A )	1873.5	N/A	1.1368 [ 1.0000 ]	113.7% { 100.0% }			
13C3_PFBs_EIS	( 302.0 / 80.0 ) 3949791	( 5.44 , N/A ) ( N/A , 0.00 , N/A )	3731.4	N/A	2.2355 [ 2.0000 ]	111.8% { 100.0% }			
13C3_PFHxS_EIS	( 402.0 / 80.0 ) 1989895	( 7.13 , N/A ) ( N/A , 0.00 , N/A )	2011.6	N/A	2.1683 [ 2.0000 ]	108.4% { 100.0% }			
13C8_PFOS_EIS	( 507.0 / 80.0 ) 4420657	( 8.35 , N/A ) ( N/A , 0.00 , N/A )	1390.6	N/A	2.1295 [ 2.0000 ]	106.5% { 100.0% }			
13C2_4:2FTS_EIS	( 329.0 / 81.0 ) 409305	( 5.20 , N/A ) ( N/A , 0.00 , N/A )	1608.0	N/A	4.1969 [ 4.0000 ]	104.9% { 100.0% }			
13C2_6:2FTS_EIS	( 429.0 / 81.0 ) 458078	( 6.71 , N/A ) ( N/A , 0.00 , N/A )	1511.4	N/A	4.2789 [ 4.0000 ]	107.0% { 100.0% }			
13C2_8:2FTS_EIS	( 529.0 / 81.0 ) 605084	( 7.89 , N/A ) ( N/A , 0.00 , N/A )	1458.0	N/A	4.0006 [ 4.0000 ]	100.0% { 100.0% }			
13C8_PFOsa_EIS	( 506.0 / 78.0 ) 8570962	( 9.83 , N/A ) ( N/A , 0.00 , N/A )	2834.2	N/A	2.2243 [ 2.0000 ]	111.2% { 100.0% }			
D3_NMeFOsa_EIS	( 515.0 / 169.0 ) 1824778	( 10.41 , N/A ) ( N/A , 0.00 , N/A )	2895.5	N/A	2.0100 [ 2.0000 ]	100.5% { 100.0% }			
D5_NEtFOsa_EIS	( 531.0 / 169.0 ) 1723765	( 10.56 , N/A ) ( N/A , 0.00 , N/A )	3505.4	N/A	2.1218 [ 2.0000 ]	106.1% { 100.0% }			

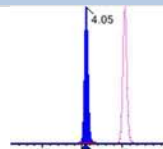
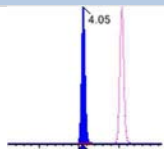
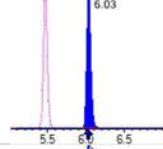
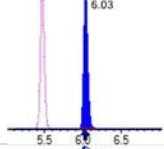
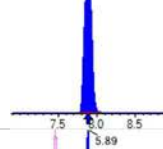
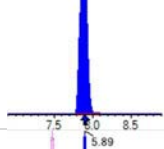
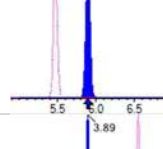
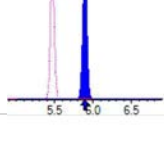
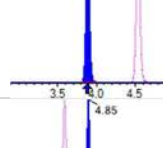
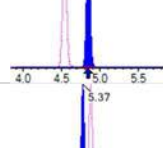
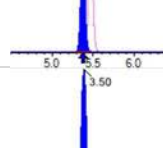
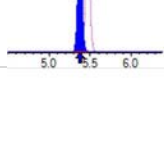
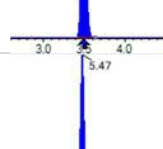
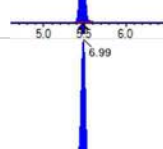
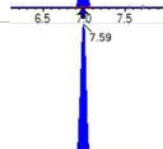

Analyte	( Q1 / Q3 ) Area Counts*min	R.T. ( R.T [min] , R.R.T. ) ( ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	( 573.0 / 419.0 ) 1488594	( 8.29 , N/A ) ( N/A , 0.00 , N/A )	1552.1	N/A	4.4312 [ 4.0000 ]	110.8% { 100.0% }			
D5_EtFOSAA_EIS	( 589.0 / 419.0 ) 1121658	( 8.53 , N/A ) ( N/A , 0.00 , N/A )	3086.8	N/A	4.0777 [ 4.0000 ]	101.9% { 100.0% }			
D7_NMeFOSE_EIS	( 623.0 / 58.9 ) 7190335	( 10.35 , N/A ) ( N/A , 0.00 , N/A )	1480.1	N/A	21.7243 [ 20.0000 ]	108.6% { 100.0% }			
D9_NEtFOSE_EIS	( 639.0 / 58.9 ) 9522239	( 10.50 , N/A ) ( N/A , 0.00 , N/A )	2110.5	N/A	22.4653 [ 20.0000 ]	112.3% { 100.0% }			
13C3_HFPODA_EIS	( 287.0 / 169.0 ) 3661243	( 5.78 , N/A ) ( N/A , 0.00 , N/A )	3134.5	N/A	9.1427 [ 8.0000 ]	114.3% { 100.0% }			

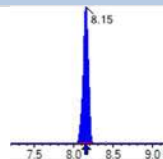
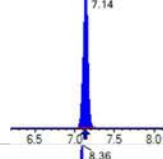
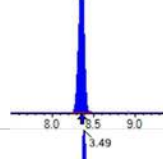
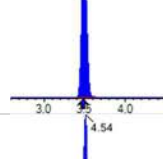
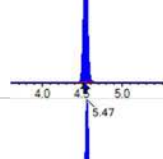
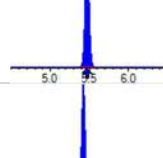
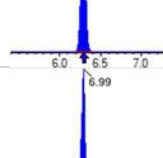
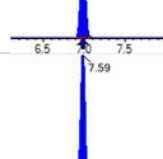
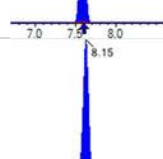
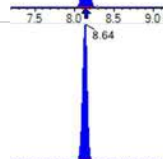
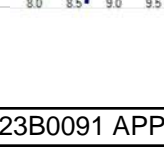


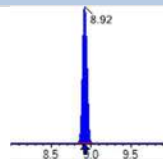
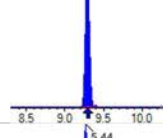
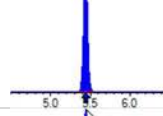
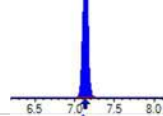
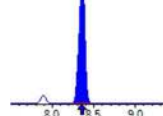
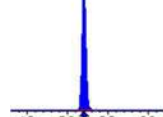
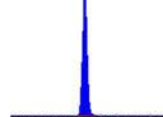
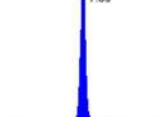
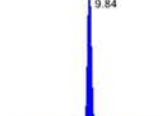
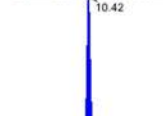
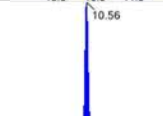
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	( 213.0 / 169.0 ) 9233467	( 3.49 , 1.00 ) ( 0.00 , N/A , 0.0 )	224.3	N/A 0.0 0.0	46.9444 [ 40.0000 ]	117.4%			
PFPeA	( 263.0 / 219.0 ) 9531366 ( 263.0 / 69.0 ) 126268	( 4.54 , 1.00 ) ( 0.00 , N/A , 0.0 )	6000.9 942.8	0.0132 123.7 123.7	23.8670 [ 20.0000 ]	119.3%			
PFHxA	( 313.0 / 269.0 ) 6393140 ( 313.0 / 119.0 ) 631426	( 5.47 , 1.00 ) ( 0.00 , N/A , 0.0 )	8330.5 54939.1	0.0988 99.2 99.2	12.4668 [ 10.0000 ]	124.7%			
PFHpA	( 363.0 / 319.0 ) 5097470 ( 363.0 / 169.0 ) 1520229	( 6.29 , 1.00 ) ( 0.00 , N/A , 0.1 )	8366.5 19589.4	0.2982 93.0 93.0	11.2965 [ 10.0000 ]	113.0%			
PFOA	( 413.0 / 369.0 ) 6305799 ( 413.0 / 169.0 ) 2127178	( 6.99 , 1.00 ) ( 0.00 , N/A , 0.1 )	9194.0 1722.9	0.3373 100.8 100.8	10.4049 [ 10.0000 ]	104.0%			
PFNA	( 463.0 / 419.0 ) 5831024 ( 463.0 / 169.0 ) 1289232	( 7.59 , 1.00 ) ( 0.00 , N/A , 0.0 )	7838.8 3394.3	0.2211 98.8 98.8	10.4692 [ 10.0000 ]	104.7%			
PFDA	( 513.0 / 469.0 ) 6609858 ( 513.0 / 169.0 ) 772819	( 8.15 , 1.00 ) ( 0.00 , N/A , -0.1 )	2492.5 1306.9	0.1169 96.1 96.1	11.9377 [ 10.0000 ]	119.4%			
PFUnA	( 563.0 / 519.0 ) 6120046 ( 563.0 / 169.0 ) 718714	( 8.63 , 1.00 ) ( 0.00 , N/A , 0.0 )	2434.2 1645.4	0.1174 112.0 112.0	12.5450 [ 10.0000 ]	125.4%			
PFDoA	( 613.0 / 569.0 ) 5503400 ( 613.0 / 169.0 ) 936186	( 8.92 , 1.00 ) ( 0.00 , N/A , 0.0 )	2937.9 2305.5	0.1701 104.0 104.0	11.5680 [ 10.0000 ]	115.7%			
PFTTrDA	( 663.0 / 619.0 ) 4864186 ( 663.0 / 169.0 ) 1198575	( 9.12 , 1.02 ) ( N/A , 0.00 , 0.0 )	2848.6 2088.0	0.2464 103.1 103.1	11.2925 [ 10.0000 ]	112.9%			
PFTeDA	( 713.0 / 669.0 ) 4740592 ( 713.0 / 169.0 ) 999348	( 9.29 , 1.00 ) ( 0.00 , N/A , 0.2 )	3599.3 1036.6	0.2108 102.0 102.0	10.7960 [ 10.0000 ]	108.0%			

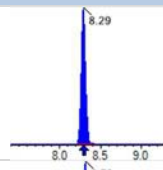
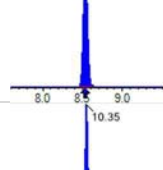
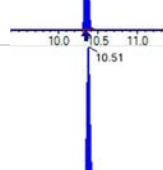
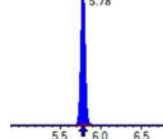

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	( 299.0 / 80.0 ) 10390212 ( 299.0 / 99.0 ) 6605887	( 5.44 , 1.00 ) ( 0.00 , N/A , 0.0 )	4357.0 4771.2	0.6358 98.3 98.3	10.2872 [ 8.8473 ]	116.3%			
PFPeS	( 349.0 / 80.0 ) 17873849 ( 349.0 / 99.0 ) 6341955	( 6.38 , 0.89 ) ( N/A , 0.00 , -0.1 )	64324.3 304115.8	0.3548 99.4 99.4	11.1269 [ 9.3838 ]	118.6%			
PFHxS	( 399.0 / 80.0 ) 13307133 ( 399.0 / 99.0 ) 4364301	( 7.14 , 1.00 ) ( 0.00 , N/A , 0.0 )	6437.0 4162.2	0.3280 101.0 101.0	10.3896 [ 9.1098 ]	114.0%			
PFHpS	( 449.0 / 80.0 ) 16488908 ( 449.0 / 99.0 ) 4352074	( 7.78 , 0.93 ) ( N/A , 0.00 , -0.1 )	14934.5 33644616.2	0.2639 97.0 97.0	12.2275 [ 9.5141 ]	128.5%			
PFOS	( 499.0 / 80.0 ) 20346672 ( 499.0 / 99.0 ) 4392227	( 8.36 , 1.00 ) ( 0.00 , N/A , 0.1 )	1739.1 2878.9	0.2159 97.5 97.5	10.9024 [ 9.2749 ]	117.5%			
PFNS	( 549.0 / 80.0 ) 23660442 ( 549.0 / 99.0 ) 5643402	( 8.78 , 1.05 ) ( N/A , 0.00 , 0.0 )	25366.7 18550.7	0.2385 108.8 108.8	11.2811 [ 9.5989 ]	117.5%			
PFDS	( 599.0 / 80.0 ) 26146019 ( 599.0 / 99.0 ) 6144955	( 9.02 , 1.08 ) ( N/A , 0.00 , 0.1 )	4520.9 3068.3	0.2350 107.8 107.8	11.4601 [ 9.6311 ]	119.0%			
PFDoS	( 699.0 / 80.0 ) 17995927 ( 699.0 / 99.0 ) 3844433	( 9.37 , 1.12 ) ( N/A , 0.00 , 0.0 )	3280.7 2642.6	0.2136 102.6 102.6	12.3673 [ 9.6956 ]	127.6%			
4:2FTS	( 327.0 / 307.0 ) 10683219 ( 327.0 / 81.0 ) 7205169	( 5.20 , 1.00 ) ( 0.00 , N/A , 0.0 )	6478.7 5055.2	0.6744 100.0 100.0	44.2283 [ 37.3811 ]	118.3%			
6:2FTS	( 427.0 / 407.0 ) 5942947 ( 427.0 / 81.0 ) 4683835	( 6.72 , 1.00 ) ( 0.00 , N/A , -0.1 )	3868.4 3031.6	0.7881 95.2 95.2	44.0754 [ 37.9617 ]	116.1%			
8:2FTS	( 527.0 / 507.0 ) 6511862 ( 527.0 / 81.0 ) 5633982	( 7.89 , 1.00 ) ( 0.00 , N/A , 0.0 )	2935.1 2349.4	0.8652 104.1 104.1	42.5374 [ 38.3315 ]	111.0%			

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 ) 28496764 ( 498.0 / 478.0 ) 688439	( 9.83 , 1.00 ) ( 0.00 , N/A , -0.1)	3755.5 1485.3	0.0242 117.6 117.6	10.8845 [ 10.0000 ]	108.8%			
NMeFOSA	( 512.0 / 219.0 ) 27427411 ( 512.0 / 169.0 ) 23151406	( 10.42 , 1.00 ) ( 0.00 , N/A , 1.0)	3933.1 5131.3	0.8441 103.0 103.0	43.4554 [ 40.0000 ]	108.6%			
NEIFOSA	( 526.0 / 219.0 ) 30152734 ( 526.0 / 169.0 ) 36496875	( 10.55 , 1.00 ) ( -0.01 , N/A , 0.5)	12529.4 13453.4	1.2104 95.4 95.4	47.2706 [ 40.0000 ]	118.2%			
NMeFOSAA	( 570.0 / 419.0 ) 2565724 ( 570.0 / 483.0 ) 1265795	( 8.29 , 1.00 ) ( 0.00 , N/A , 0.2)	2950.6 2120.3	0.4933 103.9 103.9	10.9996 [ 10.0000 ]	110.0%			
NEIFOSAA	( 584.0 / 419.0 ) 2217053 ( 584.0 / 526.0 ) 1367295	( 8.54 , 1.00 ) ( 0.00 , N/A , 0.0)	2737.6 1586.6	0.6167 120.9 120.9	10.9014 [ 10.0000 ]	109.0%			
NMeFOSE	( 616.0 / 59.0 ) 13103112	( 10.36 , 1.00 ) ( 0.01 , N/A , 0.0)	3716.4	N/A 0.0 0.0	45.6930 [ 40.0000 ]	114.2%			
NEiFOSE	( 630.0 / 59.0 ) 16273237	( 10.52 , 1.00 ) ( 0.01 , N/A , 0.0)	1832.3	N/A 0.0 0.0	45.8199 [ 40.0000 ]	114.5%			
HFPO-DA	( 285.0 / 169.0 ) 7145918 ( 285.0 / 185.0 ) 18638748	( 5.78 , 1.00 ) ( 0.00 , N/A , 0.1)	3242.8 4909.2	2.6083 94.7 94.7	24.4771 [ 20.0000 ]	122.4%			
ADONA	( 377.0 / 85.0 ) 22877224 ( 377.0 / 251.0 ) 2187719	( 6.57 , 1.14 ) ( N/A , 0.00 , 0.1)	4690.0 2291.0	0.0956 108.5 108.5	22.7861 [ 18.8540 ]	120.9%			
9CI-Pf3ONS	( 531.0 / 351.0 ) 52552445 ( 533.0 / 353.0 ) 20544699	( 8.70 , 1.50 ) ( N/A , 0.00 , 0.0)	3095.8 2832.2	0.3909 112.3 112.3	20.6280 [ 18.6651 ]	110.5%			
11CI-PF3OUDS	( 631.0 / 451.0 ) 33770888 ( 633.0 / 453.0 ) 12872268	( 9.17 , 1.59 ) ( N/A , 0.00 , 0.0)	2732.6 2758.3	0.3812 105.3 105.3	20.6476 [ 18.8642 ]	109.5%			

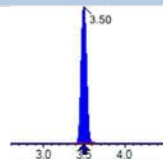
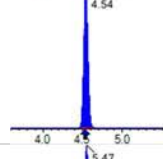
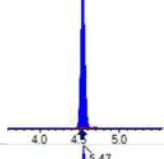
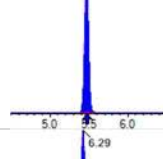
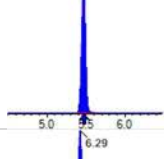
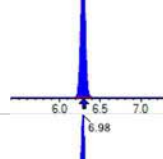
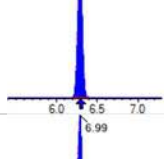
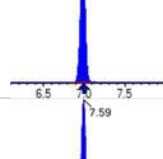
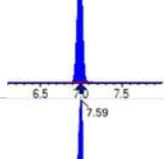
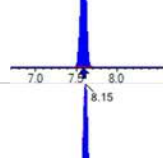
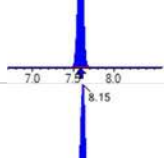
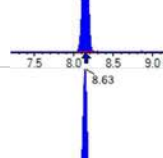
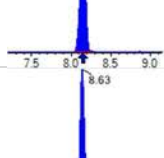
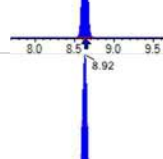
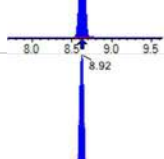
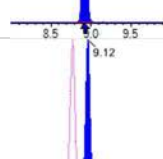
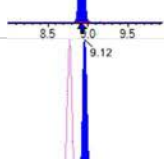
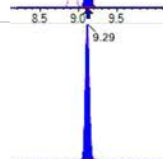
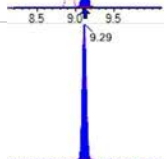
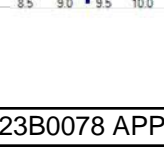
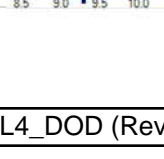
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
3:3FTCA	( 241.0 / 177.0 ) 453435 ( 241.0 / 117.0 ) 709343	( 4.05 , 0.89 ) ( N/A , 0.01 , 0.0 )	1696.2 1830.7	1.5644 99.1 99.1	43.4779 [ 40.0000 ]	108.7%			
5:3FTCA	( 341.0 / 236.7 ) 3202760 ( 341.0 / 217.0 ) 5838898	( 6.03 , 1.10 ) ( N/A , 0.01 , 0.0 )	1874.5 2496.4	1.8231 106.9 106.9	45.6716 [ 40.0000 ]	114.2%			
7:3FTCA	( 441.0 / 317.0 ) 6044367 ( 441.0 / 337.0 ) 5231397	( 7.88 , 1.44 ) ( N/A , 0.00 , 0.0 )	1248.9 1229.4	0.8655 100.6 100.6	46.6584 [ 40.0000 ]	116.6%			
PFEESA	( 315.0 / 135.0 ) 13856058 ( 315.0 / 83.0 ) 3611442	( 5.89 , 1.08 ) ( N/A , 0.00 , 0.1 )	4699.3 3584.0	0.2606 102.1 102.1	22.4929 [ 17.8492 ]	126.0%			
PFMPA	( 229.0 / 85.0 ) 2074408	( 3.89 , 0.86 ) ( N/A , 0.01 , 0.0 )	4384.2	N/A 0.0 0.0	23.7810 [ 20.0000 ]	118.9%			
PFMBA	( 279.0 / 85.0 ) 8783910	( 4.85 , 1.07 ) ( N/A , 0.00 , 0.0 )	5239.5	N/A 0.0 0.0	24.0123 [ 20.0000 ]	120.1%			
NFDHA	( 295.0 / 201.0 ) 7069388 ( 295.0 / 85.0 ) 7165355	( 5.37 , 0.98 ) ( N/A , 0.00 , 0.0 )	5626.6 4988.4	1.0136 101.2 101.2	24.8368 [ 20.0000 ]	124.2%			
13C3_PFBa_IIS	( 216.0 / 172.0 ) 224891	( 3.50 , N/A ) ( N/A , 0.01 , N/A )	1167.6	N/A	1.0040 [ 1.0000 ]	100.4% { 99.8% }			
13C2_PFHxA_IIS	( 315.0 / 270.0 ) 503477	( 5.47 , N/A ) ( N/A , 0.00 , N/A )	2210.5	N/A	0.9974 [ 1.0000 ]	99.7% { 104.9% }			
13C4_PFOA_IIS	( 417.0 / 372.0 ) 642862	( 6.99 , N/A ) ( N/A , 0.01 , N/A )	2214.2	N/A	1.0143 [ 1.0000 ]	101.4% { 108.3% }			
13C5_PFNAl_IIS	( 468.0 / 423.0 ) 572418	( 7.59 , N/A ) ( N/A , 0.00 , N/A )	11193.0	N/A	0.9473 [ 1.0000 ]	94.7% { 96.6% }			

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 ) 590795	( 8.15 , N/A ) ( N/A , 0.00 , N/A )	2371.5	N/A	1.0210 [ 1.0000 ]	102.1% { 108.5% }			
18O2_PFHxS_IIS	( 403.0 / 83.9 ) 837150	( 7.14 , N/A ) ( N/A , 0.01 , N/A )	1125.2	N/A	0.9776 [ 1.0000 ]	97.8% { 99.2% }			
13C4_PFOS_IIS	( 503.0 / 79.9 ) 1482150	( 8.36 , N/A ) ( N/A , 0.00 , N/A )	1109.8	N/A	0.9846 [ 1.0000 ]	98.5% { 98.0% }			
13C4_PFBA_EIS	( 217.0 / 172.0 ) 1899580	( 3.49 , N/A ) ( N/A , 0.01 , N/A )	4448.0	N/A	7.1011 [ 8.0000 ]	88.8% { 81.4% }			
13C5_PFPeA_EIS	( 268.0 / 223.0 ) 1941584	( 4.54 , N/A ) ( N/A , 0.01 , N/A )	3195.2	N/A	3.5246 [ 4.0000 ]	88.1% { 80.3% }			
13C5_PFHxA_EIS	( 318.0 / 273.0 ) 1166626	( 5.47 , N/A ) ( N/A , 0.00 , N/A )	3846.9	N/A	1.6917 [ 2.0000 ]	84.6% { 76.0% }			
13C4_PFHpA_EIS	( 367.0 / 322.0 ) 1068755	( 6.29 , N/A ) ( N/A , 0.00 , N/A )	2525.5	N/A	1.7886 [ 2.0000 ]	89.4% { 84.6% }			
13C8_PFOA_EIS	( 421.0 / 376.0 ) 1361303	( 6.99 , N/A ) ( N/A , 0.01 , N/A )	1818.4	N/A	1.8698 [ 2.0000 ]	93.5% { 88.4% }			
13C9_PFNA_EIS	( 472.0 / 427.0 ) 629508	( 7.59 , N/A ) ( N/A , 0.00 , N/A )	3985.1	N/A	1.0379 [ 1.0000 ]	103.8% { 89.6% }			
13C6_PFDA_EIS	( 519.0 / 474.0 ) 629459	( 8.15 , N/A ) ( N/A , 0.00 , N/A )	1155.8	N/A	0.8450 [ 1.0000 ]	84.5% { 81.0% }			
13C7_PFUnA_EIS	( 570.0 / 525.0 ) 587177	( 8.64 , N/A ) ( N/A , 0.00 , N/A )	3704.7	N/A	0.7927 [ 1.0000 ]	79.3% { 76.1% }			

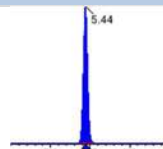
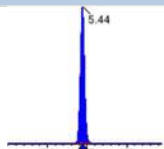
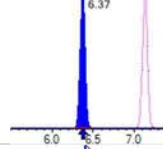
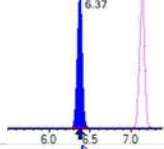
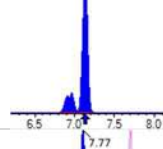
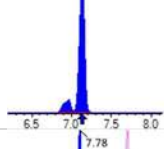
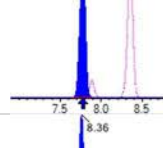
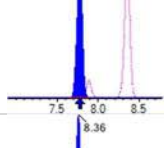
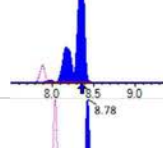
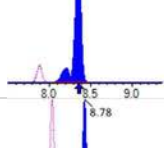
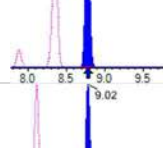
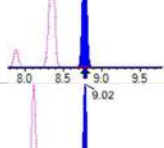
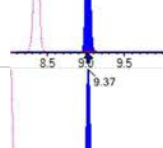
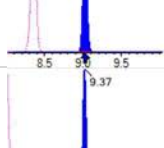
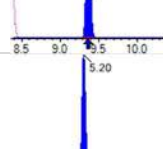
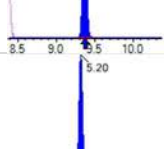
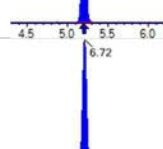
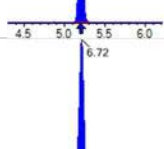
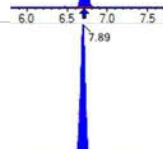
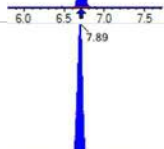

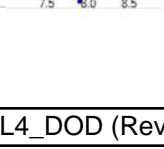
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDa_EIS	( 615.0 / 570.0 ) 541005	( 8.92 , N/A ) ( N/A , 0.00 , N/A )	53274.1	N/A	0.8516 [ 1.0000 ]	85.2% { 82.4% }			
13C2_PFTeDA_EIS	( 715.0 / 670.0 ) 507428	( 9.29 , N/A ) ( N/A , 0.00 , N/A )	1491.2	N/A	0.8717 [ 1.0000 ]	87.2% { 83.2% }			
13C3_PFBs_EIS	( 302.0 / 80.0 ) 3250497	( 5.44 , N/A ) ( N/A , 0.00 , N/A )	3936.3	N/A	1.8552 [ 2.0000 ]	92.8% { 82.3% }			
13C3_PFHxS_EIS	( 402.0 / 80.0 ) 1665870	( 7.14 , N/A ) ( N/A , 0.00 , N/A )	2186.2	N/A	1.8305 [ 2.0000 ]	91.5% { 83.7% }			
13C8_PFOS_EIS	( 507.0 / 80.0 ) 3539645	( 8.36 , N/A ) ( N/A , 0.01 , N/A )	1147.9	N/A	1.7396 [ 2.0000 ]	87.0% { 80.1% }			
13C2_4:2FTS_EIS	( 329.0 / 81.0 ) 353761	( 5.20 , N/A ) ( N/A , 0.00 , N/A )	1744.3	N/A	3.6579 [ 4.0000 ]	91.4% { 86.4% }			
13C2_6:2FTS_EIS	( 429.0 / 81.0 ) 380113	( 6.72 , N/A ) ( N/A , 0.01 , N/A )	1460.9	N/A	3.5806 [ 4.0000 ]	89.5% { 83.0% }			
13C2_8:2FTS_EIS	( 529.0 / 81.0 ) 562167	( 7.89 , N/A ) ( N/A , 0.00 , N/A )	1081.9	N/A	3.7482 [ 4.0000 ]	93.7% { 92.9% }			
13C8_PFOsa_EIS	( 506.0 / 78.0 ) 6546821	( 9.84 , N/A ) ( N/A , 0.00 , N/A )	2955.2	N/A	1.7333 [ 2.0000 ]	86.7% { 76.4% }			
D3_NMeFOSA_EIS	( 515.0 / 169.0 ) 1603154	( 10.42 , N/A ) ( N/A , 0.00 , N/A )	2177.9	N/A	1.8015 [ 2.0000 ]	90.1% { 87.9% }			
D5_NEtFOSA_EIS	( 531.0 / 169.0 ) 1365559	( 10.56 , N/A ) ( N/A , 0.00 , N/A )	2826.0	N/A	1.7149 [ 2.0000 ]	85.7% { 79.2% }			

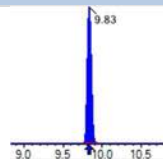
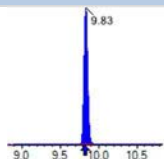
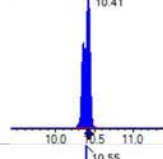
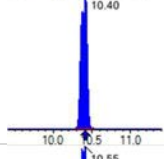
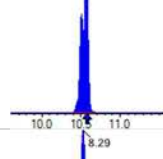
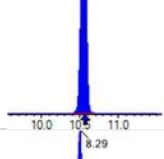
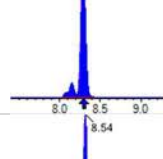
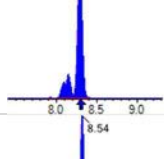
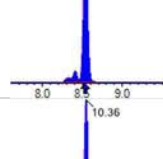
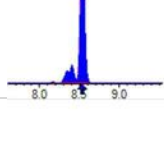
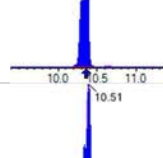
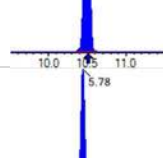
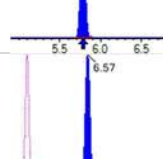
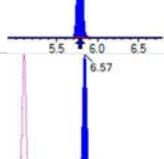
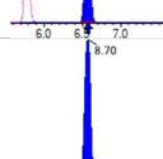
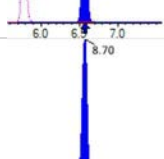
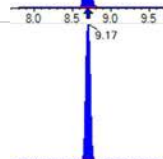
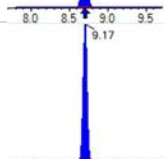
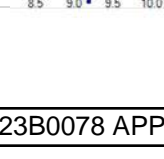
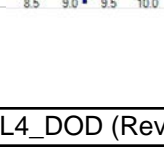
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	( 573.0 / 419.0 ) 1188918	( 8.29 , N/A ) ( N/A , 0.00 , N/A )	1548.6	N/A	3.6106 [ 4.0000 ]	90.3% { 79.9% }			
D5_EtFOSAA_EIS	( 589.0 / 419.0 ) 968558	( 8.53 , N/A ) ( N/A , 0.00 , N/A )	1321416.8	N/A	3.5922 [ 4.0000 ]	89.8% { 86.4% }			
D7_NMeFOSE_EIS	( 623.0 / 58.9 ) 5736778	( 10.35 , N/A ) ( N/A , 0.00 , N/A )	1892.8	N/A	17.6829 [ 20.0000 ]	88.4% { 79.8% }			
D9_NEtFOSE_EIS	( 639.0 / 58.9 ) 7464155	( 10.51 , N/A ) ( N/A , 0.00 , N/A )	2307.6	N/A	17.9656 [ 20.0000 ]	89.8% { 78.4% }			
13C3_HFPODA_EIS	( 287.0 / 169.0 ) 2953400	( 5.78 , N/A ) ( N/A , 0.00 , N/A )	2802.1	N/A	7.0337 [ 8.0000 ]	87.9% { 80.7% }			

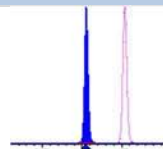
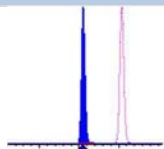
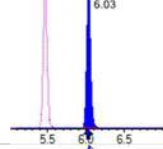
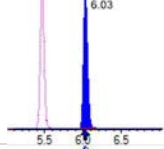
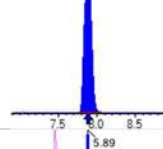
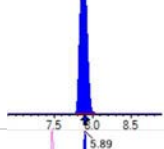
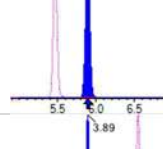
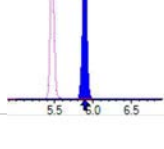
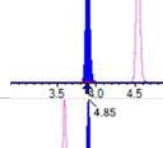
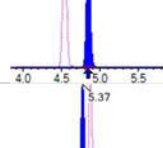
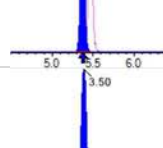
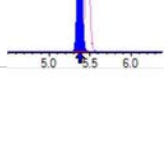
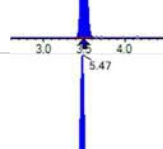
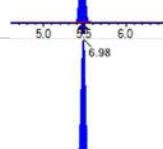
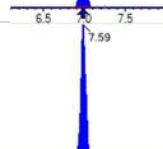



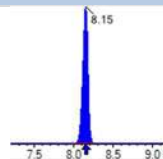
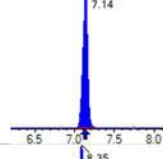
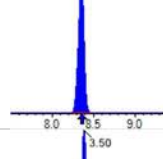
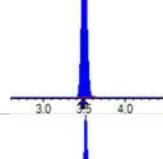
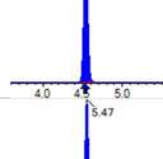
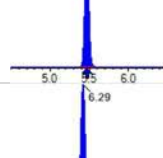
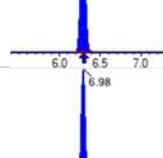
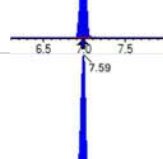
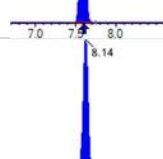
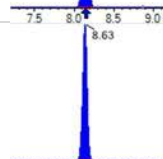
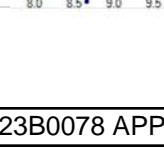
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	( 213.0 / 169.0 ) 15337689	( 3.50 , 1.00 ) ( 0.00 , N/A , 0.0 )	202.8	N/A 0.0 0.0	82.1737 [ 80.0000 ]	102.7%			
PFPeA	( 263.0 / 219.0 ) 16682791 ( 263.0 / 69.0 ) 214318	( 4.54 , 1.00 ) ( 0.00 , N/A , 0.1 )	7620.4 1415.7	0.0128 120.0 120.0	40.3999 [ 40.0000 ]	101.0%			
PFHxA	( 313.0 / 269.0 ) 11582071 ( 313.0 / 119.0 ) 1164126	( 5.47 , 1.00 ) ( 0.00 , N/A , 0.0 )	10132.7 83880.5	0.1005 100.9 100.9	20.9265 [ 20.0000 ]	104.6%			
PFHpA	( 363.0 / 319.0 ) 10236624 ( 363.0 / 169.0 ) 2996363	( 6.29 , 1.00 ) ( 0.00 , N/A , 0.0 )	9527.2 83284.9	0.2927 91.3 91.3	22.4112 [ 20.0000 ]	112.1%			
PFOA	( 413.0 / 369.0 ) 12773098 ( 413.0 / 169.0 ) 3940700	( 6.98 , 1.00 ) ( 0.00 , N/A , -0.1 )	9301.8 8448.1	0.3085 92.2 92.2	22.4084 [ 20.0000 ]	112.0%			
PFNA	( 463.0 / 419.0 ) 11486325 ( 463.0 / 169.0 ) 2596327	( 7.59 , 1.00 ) ( 0.00 , N/A , -0.1 )	9149.1 1094562.3	0.2260 101.0 101.0	20.8564 [ 20.0000 ]	104.3%			
PFDA	( 513.0 / 469.0 ) 13202559 ( 513.0 / 169.0 ) 1478374	( 8.15 , 1.00 ) ( 0.00 , N/A , 0.0 )	3679.6 1643.9	0.1120 92.1 92.1	20.3910 [ 20.0000 ]	102.0%			
PFUnA	( 563.0 / 519.0 ) 12098447 ( 563.0 / 169.0 ) 1401009	( 8.63 , 1.00 ) ( 0.00 , N/A , 0.2 )	3830.9 1876.5	0.1158 110.5 110.5	21.9467 [ 20.0000 ]	109.7%			
PFDaA	( 613.0 / 569.0 ) 9962922 ( 613.0 / 169.0 ) 1708344	( 8.92 , 1.00 ) ( 0.00 , N/A , 0.0 )	3464.3 2061.8	0.1715 104.8 104.8	19.4226 [ 20.0000 ]	97.1%			
PFTTrDA	( 663.0 / 619.0 ) 8892680 ( 663.0 / 169.0 ) 2243556	( 9.12 , 1.02 ) ( N/A , 0.00 , 0.1 )	3771.0 1929.7	0.2523 105.6 105.6	19.1473 [ 20.0000 ]	95.7%			
PFTeDA	( 713.0 / 669.0 ) 9909351 ( 713.0 / 169.0 ) 1996611	( 9.29 , 1.00 ) ( 0.00 , N/A , 0.0 )	4270.0 1672.9	0.2015 97.5 97.5	21.0189 [ 20.0000 ]	105.1%			

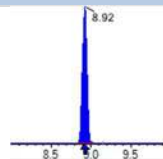
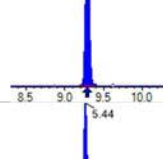
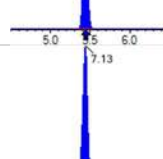
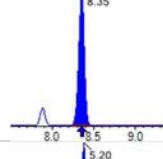
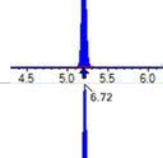
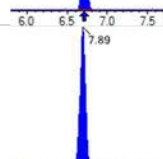
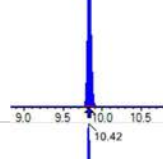
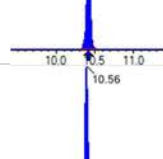
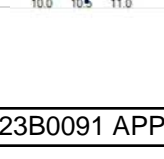
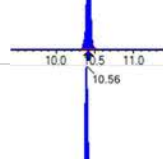
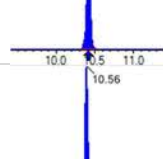


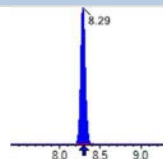
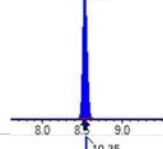
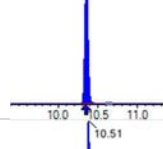
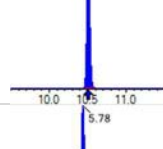

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	( 299.0 / 80.0 ) 18432508 ( 299.0 / 99.0 ) 11551282	( 5.44 , 1.00 ) ( 0.00 , N/A , 0.0 )	4327.3 4075.2	0.6267 96.9 96.9	17.7178 [ 17.6947 ]	100.1%			
PFPeS	( 349.0 / 80.0 ) 31713610 ( 349.0 / 99.0 ) 12209635	( 6.37 , 0.89 ) ( N/A , 0.00 , -0.1 )	19247.5 6160374.7	0.3850 107.8 107.8	19.1741 [ 18.7676 ]	102.2%			
PFHxS	( 399.0 / 80.0 ) 25528591 ( 399.0 / 99.0 ) 8369716	( 7.13 , 1.00 ) ( 0.00 , N/A , 0.0 )	7561.6 283033.5	0.3279 101.0 101.0	19.3577 [ 18.2197 ]	106.2%			
PFHpS	( 449.0 / 80.0 ) 29354713 ( 449.0 / 99.0 ) 8525062	( 7.77 , 0.93 ) ( N/A , 0.00 , 0.0 )	69900.5 28178.3	0.2904 106.7 106.7	19.8365 [ 19.0281 ]	104.2%			
PFOS	( 499.0 / 80.0 ) 37594232 ( 499.0 / 99.0 ) 8592679	( 8.36 , 1.00 ) ( 0.00 , N/A , 0.0 )	2042.7 3023.4	0.2286 103.3 103.3	18.3565 [ 18.5499 ]	99.0%			
PFNS	( 549.0 / 80.0 ) 40002905 ( 549.0 / 99.0 ) 10885578	( 8.78 , 1.05 ) ( N/A , 0.00 , 0.0 )	9789.3 69993.2	0.2721 124.1 124.1	17.3803 [ 19.1977 ]	90.5%			
PFDS	( 599.0 / 80.0 ) 43319573 ( 599.0 / 99.0 ) 11405776	( 9.02 , 1.08 ) ( N/A , 0.00 , 0.0 )	4704.2 4394.0	0.2633 120.8 120.8	17.3024 [ 19.2621 ]	89.8%			
PFDoS	( 699.0 / 80.0 ) 31495074 ( 699.0 / 99.0 ) 7427296	( 9.37 , 1.12 ) ( N/A , 0.00 , 0.0 )	3407.1 3335.6	0.2358 113.3 113.3	19.7235 [ 19.3913 ]	101.7%			
4:2FTS	( 327.0 / 307.0 ) 19385545 ( 327.0 / 81.0 ) 13393654	( 5.20 , 1.00 ) ( 0.00 , N/A , 0.0 )	4804.2 3388.8	0.6909 102.4 102.4	72.0914 [ 74.7622 ]	96.4%			
6:2FTS	( 427.0 / 407.0 ) 12377441 ( 427.0 / 81.0 ) 9446075	( 6.72 , 1.00 ) ( 0.00 , N/A , 0.0 )	3948.2 3770.7	0.7632 92.2 92.2	81.9573 [ 75.9234 ]	107.9%			
8:2FTS	( 527.0 / 507.0 ) 13494970 ( 527.0 / 81.0 ) 11411983	( 7.89 , 1.00 ) ( 0.00 , N/A , 0.0 )	3531.0 3647.8	0.8456 101.7 101.7	78.5620 [ 76.6631 ]	102.5%			

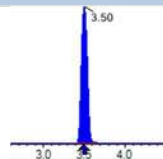
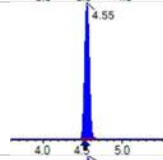
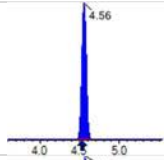
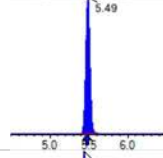
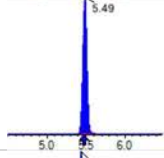
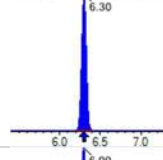
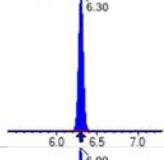
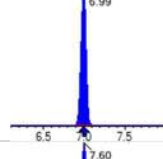
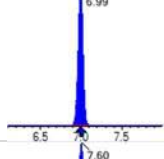
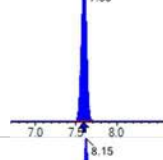
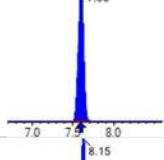
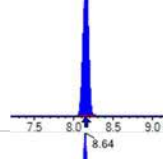
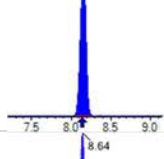
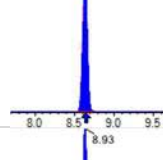
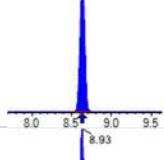
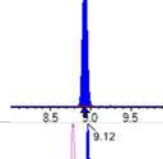
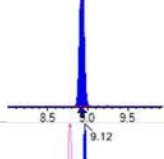
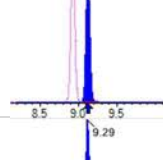
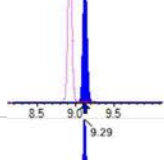
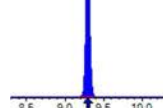
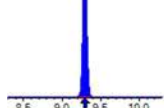
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) ( $\Delta$ RT-[min], $\Delta$ RT- CV[min], $\Delta$ RT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	( 498.0 / 78.0 ) 45374007 ( 498.0 / 478.0 ) 1211984	( 9.83 , 1.00 ) ( 0.00 , N/A , 0.1 )	3138.6 2816.6	0.0267 130.0 130.0	15.7940 [ 20.0000 ]	79.0%			
NMeFOSA	( 512.0 / 219.0 ) 46430830 ( 512.0 / 169.0 ) 38979424	( 10.41 , 1.00 ) ( -0.01 , N/A , 0.6 )	3026.9 3553.6	0.8395 102.5 102.5	75.0182 [ 80.0000 ]	93.8%			
NEIFOSA	( 526.0 / 219.0 ) 49087453 ( 526.0 / 169.0 ) 56621172	( 10.55 , 1.00 ) ( -0.01 , N/A , 0.3 )	16025.9 16309.8	1.1535 90.9 90.9	78.9558 [ 80.0000 ]	98.7%			
NMeFOSAA	( 570.0 / 419.0 ) 4928976 ( 570.0 / 483.0 ) 2272830	( 8.29 , 1.00 ) ( 0.00 , N/A , -0.1 )	3144.1 1671.5	0.4611 97.1 97.1	20.3987 [ 20.0000 ]	102.0%			
NEIFOSAA	( 584.0 / 419.0 ) 4563311 ( 584.0 / 526.0 ) 2631013	( 8.54 , 1.00 ) ( 0.01 , N/A , 0.1 )	2694.2 2137.1	0.5766 113.0 113.0	20.4343 [ 20.0000 ]	102.2%			
NMeFOSE	( 616.0 / 59.0 ) 24449401	( 10.36 , 1.00 ) ( 0.01 , N/A , 0.0 )	4286.5	N/A 0.0 0.0	82.4264 [ 80.0000 ]	103.0%			
NEtFOSE	( 630.0 / 59.0 ) 28256391	( 10.51 , 1.00 ) ( 0.01 , N/A , 0.0 )	1566.1	N/A 0.0 0.0	79.8411 [ 80.0000 ]	99.8%			
HFPO-DA	( 285.0 / 169.0 ) 13130220 ( 285.0 / 185.0 ) 32292921	( 5.78 , 1.00 ) ( 0.00 , N/A , 0.0 )	4252.9 4645.1	2.4594 89.3 89.3	41.6650 [ 40.0000 ]	104.2%			
ADONA	( 377.0 / 85.0 ) 38742745 ( 377.0 / 251.0 ) 3813720	( 6.57 , 1.14 ) ( N/A , 0.00 , 0.0 )	4573.0 2959.6	0.0984 111.6 111.6	35.7483 [ 37.7080 ]	94.8%			
9CI-Pf3ONS	( 531.0 / 351.0 ) 75971960 ( 533.0 / 353.0 ) 33582790	( 8.70 , 1.50 ) ( N/A , 0.00 , 0.0 )	2457.2 3506.4	0.4420 126.9 126.9	35.2512 [ 37.3302 ]	94.4%			
11CI-PF3OUDS	( 631.0 / 451.0 ) 55193805 ( 633.0 / 453.0 ) 22582441	( 9.17 , 1.59 ) ( N/A , 0.00 , -0.1 )	3979.8 3614.0	0.4091 113.1 113.1	31.2618 [ 37.7283 ]	82.9%			

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
3:3FTCA	( 241.0 / 177.0 ) 902077 ( 241.0 / 117.0 ) 1433216	( 4.05 , 0.89 ) ( N/A , 0.01 , 0.1 )	2153.7 2284.4	1.5888 100.7 100.7	77.8383 [ 80.0000 ]	97.3%			
5:3FTCA	( 341.0 / 236.7 ) 6878497 ( 341.0 / 217.0 ) 11289214	( 6.03 , 1.10 ) ( N/A , 0.01 , 0.1 )	3227.4 2937.3	1.6412 96.2 96.2	90.8834 [ 80.0000 ]	113.6%			
7:3FTCA	( 441.0 / 317.0 ) 12072417 ( 441.0 / 337.0 ) 10707762	( 7.88 , 1.44 ) ( N/A , 0.00 , 0.0 )	1547.1 2006.4	0.8870 103.1 103.1	86.3460 [ 80.0000 ]	107.9%			
PFEESA	( 315.0 / 135.0 ) 24328200 ( 315.0 / 83.0 ) 6999908	( 5.89 , 1.08 ) ( N/A , 0.00 , 0.0 )	5805.7 4209.3	0.2877 112.7 112.7	36.5918 [ 35.6984 ]	102.5%			
PFMPA	( 229.0 / 85.0 ) 3687422	( 3.89 , 0.86 ) ( N/A , 0.01 , 0.0 )	5387.3	N/A 0.0 0.0	40.8816 [ 40.0000 ]	102.2%			
PFMBA	( 279.0 / 85.0 ) 15245914	( 4.85 , 1.07 ) ( N/A , 0.00 , 0.0 )	6035.2	N/A 0.0 0.0	40.3060 [ 40.0000 ]	100.8%			
NFDHA	( 295.0 / 201.0 ) 12950881 ( 295.0 / 85.0 ) 12829443	( 5.37 , 0.98 ) ( N/A , 0.00 , 0.0 )	4942.1 5261.9	0.9906 98.9 98.9	42.1582 [ 40.0000 ]	105.4%			
13C3_PFBA_IIS	( 216.0 / 172.0 ) 185676	( 3.50 , N/A ) ( N/A , 0.01 , N/A )	986.1	N/A	0.8289 [ 1.0000 ]	82.9% { 82.4% }			
13C2_PFHxA_IIS	( 315.0 / 270.0 ) 474941	( 5.47 , N/A ) ( N/A , 0.00 , N/A )	6192.6	N/A	0.9408 [ 1.0000 ]	94.1% { 98.9% }			
13C4_PFOA_IIS	( 417.0 / 372.0 ) 608223	( 6.98 , N/A ) ( N/A , 0.01 , N/A )	3392.9	N/A	0.9597 [ 1.0000 ]	96.0% { 102.5% }			
13C5_PFNA_IIS	( 468.0 / 423.0 ) 546126	( 7.59 , N/A ) ( N/A , 0.00 , N/A )	43952.5	N/A	0.9038 [ 1.0000 ]	90.4% { 92.1% }			

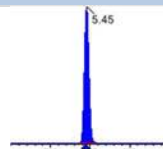
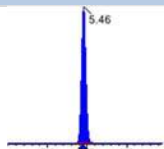
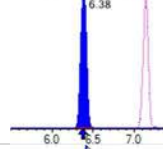
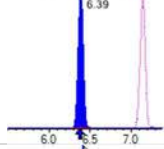
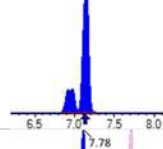
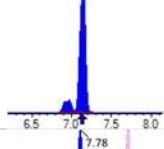
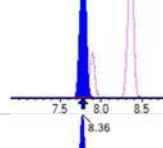
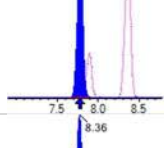
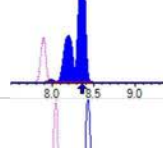
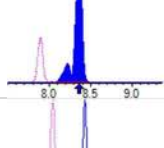
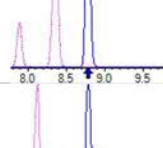
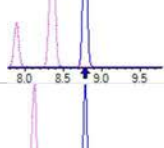
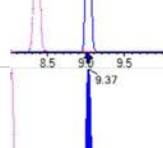
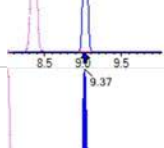
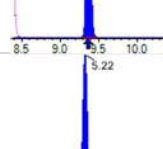
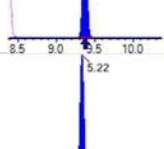
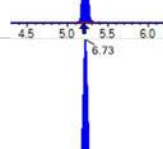
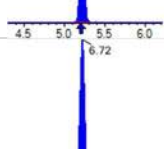
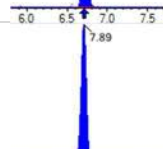
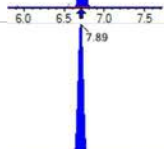

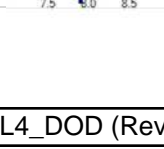
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
13C2_PFDA_IIS	( 515.0 / 470.1 ) 584998	( 8.15 , N/A ) ( N/A , 0.00 , N/A )	1234.2	N/A	1.0110 [ 1.0000 ]	101.1% { 107.4% }			
18O2_PFHxS_IIS	( 403.0 / 83.9 ) 824707	( 7.14 , N/A ) ( N/A , 0.01 , N/A )	1224.7	N/A	0.9630 [ 1.0000 ]	96.3% { 97.7% }			
13C4_PFOS_IIS	( 503.0 / 79.9 ) 1345342	( 8.35 , N/A ) ( N/A , 0.00 , N/A )	1649.3	N/A	0.8937 [ 1.0000 ]	89.4% { 89.0% }			
13C4_PFBA_EIS	( 217.0 / 172.0 ) 1802619	( 3.50 , N/A ) ( N/A , 0.01 , N/A )	4896.0	N/A	8.1618 [ 8.0000 ]	102.0% { 77.2% }			
13C5_PFPeA_EIS	( 268.0 / 223.0 ) 2007646	( 4.54 , N/A ) ( N/A , 0.01 , N/A )	3229.3	N/A	3.8635 [ 4.0000 ]	96.6% { 83.0% }			
13C5_PFHxA_EIS	( 318.0 / 273.0 ) 1259108	( 5.47 , N/A ) ( N/A , 0.00 , N/A )	2504.4	N/A	1.9355 [ 2.0000 ]	96.8% { 82.1% }			
13C4_PFHpA_EIS	( 367.0 / 322.0 ) 1081833	( 6.29 , N/A ) ( N/A , 0.00 , N/A )	5563.9	N/A	1.9193 [ 2.0000 ]	96.0% { 85.6% }			
13C8_PFOA_EIS	( 421.0 / 376.0 ) 1280381	( 6.98 , N/A ) ( N/A , 0.00 , N/A )	1843.3	N/A	1.8588 [ 2.0000 ]	92.9% { 83.1% }			
13C9_PFNA_EIS	( 472.0 / 427.0 ) 622463	( 7.59 , N/A ) ( N/A , 0.00 , N/A )	1881.4	N/A	1.0757 [ 1.0000 ]	107.6% { 88.6% }			
13C6_PFDA_EIS	( 519.0 / 474.0 ) 736065	( 8.14 , N/A ) ( N/A , 0.00 , N/A )	1519.2	N/A	0.9979 [ 1.0000 ]	99.8% { 94.8% }			
13C7_PFUnA_EIS	( 570.0 / 525.0 ) 663506	( 8.63 , N/A ) ( N/A , 0.00 , N/A )	2209.0	N/A	0.9046 [ 1.0000 ]	90.5% { 86.0% }			

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 ) 583320	( 8.92, N/A ) ( N/A, 0.00, N/A )	4662.7	N/A	0.9273 [ 1.0000 ]	92.7% { 88.8% }			
13C2_PFTeDA_EIS	( 715.0 / 670.0 ) 544806	( 9.29, N/A ) ( N/A, 0.00, N/A )	1065.6	N/A	0.9451 [ 1.0000 ]	94.5% { 89.3% }			
13C3_PFBs_EIS	( 302.0 / 80.0 ) 3348089	( 5.44, N/A ) ( N/A, 0.00, N/A )	2918.5	N/A	1.9398 [ 2.0000 ]	97.0% { 84.8% }			
13C3_PFHxS_EIS	( 402.0 / 80.0 ) 1715256	( 7.13, N/A ) ( N/A, 0.00, N/A )	1905.6	N/A	1.9132 [ 2.0000 ]	95.7% { 86.2% }			
13C8_PFOS_EIS	( 507.0 / 80.0 ) 3884369	( 8.35, N/A ) ( N/A, 0.00, N/A )	849.6	N/A	2.1031 [ 2.0000 ]	105.2% { 87.9% }			
13C2_4:2FTS_EIS	( 329.0 / 81.0 ) 393824	( 5.20, N/A ) ( N/A, 0.00, N/A )	1464.1	N/A	4.1337 [ 4.0000 ]	103.3% { 96.2% }			
13C2_6:2FTS_EIS	( 429.0 / 81.0 ) 425746	( 6.72, N/A ) ( N/A, 0.01, N/A )	1188.0	N/A	4.0710 [ 4.0000 ]	101.8% { 92.9% }			
13C2_8:2FTS_EIS	( 529.0 / 81.0 ) 630798	( 7.89, N/A ) ( N/A, 0.00, N/A )	1493.4	N/A	4.2693 [ 4.0000 ]	106.7% { 104.2% }			
13C8_PFOsa_EIS	( 506.0 / 78.0 ) 7183878	( 9.83, N/A ) ( N/A, 0.00, N/A )	3258.9	N/A	2.0954 [ 2.0000 ]	104.8% { 83.8% }			
D3_NMeFOsa_EIS	( 515.0 / 169.0 ) 1698408	( 10.42, N/A ) ( N/A, 0.00, N/A )	2308.8	N/A	2.1027 [ 2.0000 ]	105.1% { 93.1% }			
D5_NEtFOsa_EIS	( 531.0 / 169.0 ) 1330948	( 10.56, N/A ) ( N/A, 0.01, N/A )	3073.0	N/A	1.8414 [ 2.0000 ]	92.1% { 77.2% }			

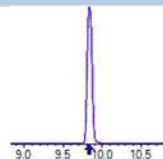
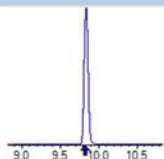
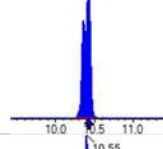
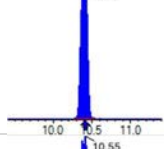
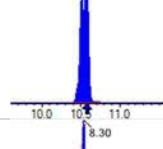
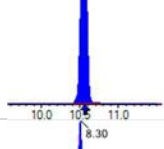
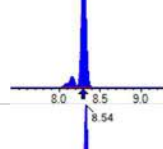
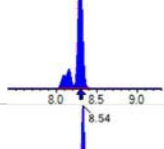
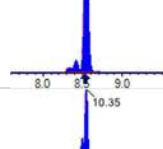
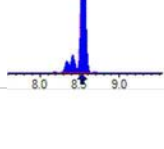
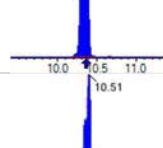
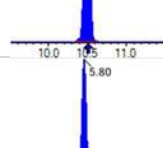
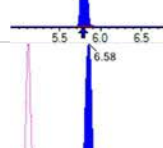
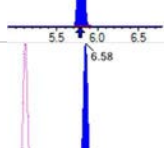
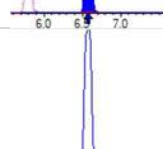
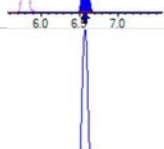
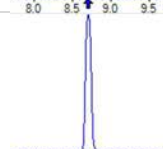
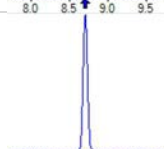
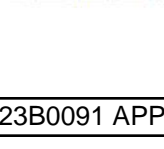
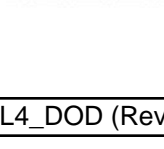
Analyte	( Q1 / Q3 ) Area Counts*min	R.T. ( R.T [min] , R.R.T. ) ( ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
D3_MeFOSAA_EIS	( 573.0 / 419.0 ) 1231612	( 8.29 , N/A ) ( N/A , 0.00 , N/A )	2039.8	N/A	4.1206 [ 4.0000 ]	103.0% { 82.7% }			
D5_EtFOSAA_EIS	( 589.0 / 419.0 ) 1063539	( 8.53 , N/A ) ( N/A , 0.00 , N/A )	7788.5	N/A	4.3456 [ 4.0000 ]	108.6% { 94.8% }			
D7_NMeFOSE_EIS	( 623.0 / 58.9 ) 5933968	( 10.35 , N/A ) ( N/A , 0.00 , N/A )	2294.5	N/A	20.1507 [ 20.0000 ]	100.8% { 82.5% }			
D9_NEtFOSE_EIS	( 639.0 / 58.9 ) 7437914	( 10.51 , N/A ) ( N/A , 0.00 , N/A )	2061.1	N/A	19.7230 [ 20.0000 ]	98.6% { 78.1% }			
13C3_HFPODA_EIS	( 287.0 / 169.0 ) 3188049	( 5.78 , N/A ) ( N/A , 0.00 , N/A )	2502.4	N/A	8.0487 [ 8.0000 ]	100.6% { 87.1% }			

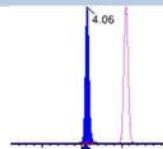
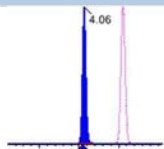
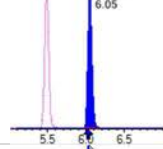
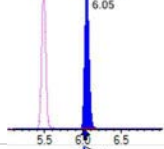
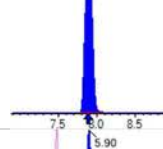
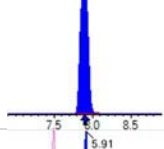
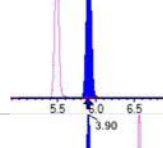
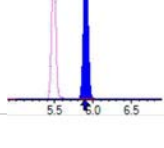
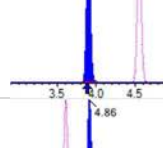
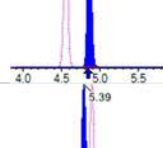
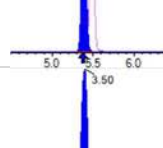
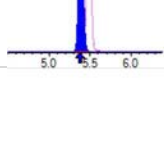
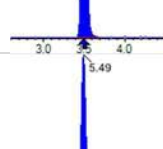
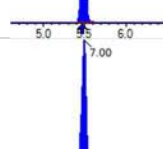
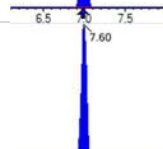

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	( 213.0 / 169.0 ) 30157574	( 3.50 , 1.00 ) ( 0.00 , N/A , 0.0 )	199.4	N/A 0.0 0.0	199.3851 [ 200.0000 ]	99.7%			
PFPeA	( 263.0 / 219.0 ) 33180312 ( 263.0 / 69.0 ) 437514	( 4.55 , 1.00 ) ( 0.00 , N/A , -0.1 )	7659.6 4196.6	0.0132 123.1 123.1	99.0620 [ 100.0000 ]	99.1%			
PFHxA	( 313.0 / 269.0 ) 24352789 ( 313.0 / 119.0 ) 2602017	( 5.49 , 1.00 ) ( 0.00 , N/A , 0.0 )	9698.3 258522.9	0.1068 107.3 107.3	48.3136 [ 50.0000 ]	96.6%			
PFHpA	( 363.0 / 319.0 ) 23305554 ( 363.0 / 169.0 ) 6794628	( 6.30 , 1.00 ) ( 0.00 , N/A , 0.1 )	13241.3 34181.2	0.2915 90.9 90.9	54.6205 [ 50.0000 ]	109.2%			
PFOA	( 413.0 / 369.0 ) 27748645 ( 413.0 / 169.0 ) 9269777	( 6.99 , 1.00 ) ( 0.00 , N/A , 0.0 )	9517.6 12954.3	0.3341 99.8 99.8	48.7358 [ 50.0000 ]	97.5%			
PFNA	( 463.0 / 419.0 ) 26643346 ( 463.0 / 169.0 ) 5700123	( 7.60 , 1.00 ) ( 0.00 , N/A , -0.1 )	10839.5 55499.9	0.2139 95.6 95.6	53.9772 [ 50.0000 ]	108.0%			
PFDA	( 513.0 / 469.0 ) 27897726 ( 513.0 / 169.0 ) 3358162	( 8.15 , 1.00 ) ( 0.00 , N/A , 0.0 )	3311.3 2578.7	0.1204 99.0 99.0	49.9184 [ 50.0000 ]	99.8%			
PFUnA	( 563.0 / 519.0 ) 25375749 ( 563.0 / 169.0 ) 3201701	( 8.64 , 1.00 ) ( 0.00 , N/A , 0.0 )	4015.5 2264.0	0.1262 120.3 120.3	47.9194 [ 50.0000 ]	95.8%			
PFDoA	( 613.0 / 569.0 ) 22253713 ( 613.0 / 169.0 ) 3793752	( 8.93 , 1.00 ) ( 0.00 , N/A , 0.0 )	3711.7 3062.5	0.1705 104.2 104.2	47.0721 [ 50.0000 ]	94.1%			
PFTTrDA	( 663.0 / 619.0 ) 17139294 ( 663.0 / 169.0 ) 4669255	( 9.12 , 1.02 ) ( N/A , 0.00 , 0.0 )	2774.9 2031.0	0.2724 114.0 114.0	40.0412 [ 50.0000 ]	80.1%			
PFTeDA	( 713.0 / 669.0 ) 21814245 ( 713.0 / 169.0 ) 4999446	( 9.29 , 1.00 ) ( 0.00 , N/A , 0.0 )	2027.5 1844.5	0.2292 110.9 110.9	47.8819 [ 50.0000 ]	95.8%			

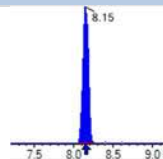
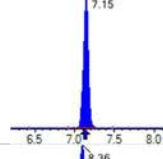
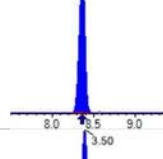
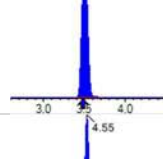
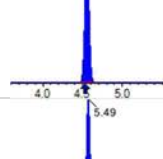
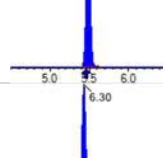
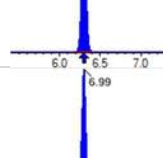
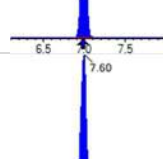
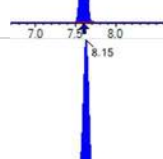
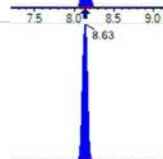
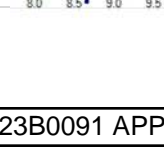


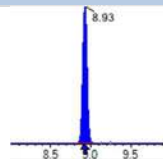
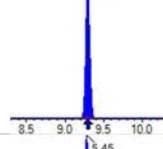
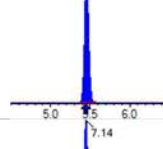
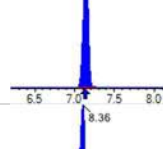
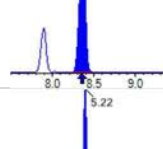
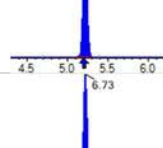
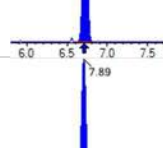
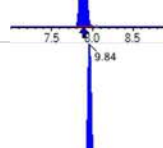
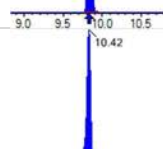
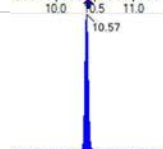
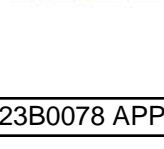
Analyte	(Q1 / Q3) Area Counts*min	R.T. ( R.T [min] , R.R.T. ) ( ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	( 299.0 / 80.0 ) 33764092 ( 299.0 / 99.0 ) 22999680	( 5.45 , 1.00 ) ( 0.00 , N/A , 0.0 )	4850.9 4549.7	0.6812 105.3 105.3	39.7222 [ 44.2367 ]	89.8%			
PFPeS	( 349.0 / 80.0 ) 57760543 ( 349.0 / 99.0 ) 25912545	( 6.38 , 0.89 ) ( N/A , 0.01 , -0.1 )	11585.7 28547.7	0.4486 125.7 125.7	37.2690 [ 46.9191 ]	79.4%			
PFHxS	( 399.0 / 80.0 ) 54967905 ( 399.0 / 99.0 ) 19970270	( 7.14 , 1.00 ) ( 0.00 , N/A , 0.1 )	7010.9 10956.1	0.3633 111.9 111.9	44.4819 [ 45.5491 ]	97.7%			
PFHpS	( 449.0 / 80.0 ) 55548911 ( 449.0 / 99.0 ) 18510688	( 7.78 , 0.93 ) ( N/A , 0.01 , 0.0 )	14572.4 49735.1	0.3332 122.4 122.4	41.7551 [ 47.5703 ]	87.8%			
PFOS	( 499.0 / 80.0 ) 74516074 ( 499.0 / 99.0 ) 18648612	( 8.36 , 1.00 ) ( 0.00 , N/A , -0.1 )	2183.8 3445.8	0.2503 113.1 113.1	40.4730 [ 46.3746 ]	87.3%			
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 [ 47.9943 ]	N/A%			QC,
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 ) 54329438 ( 699.0 / 99.0 ) 16259426	( 9.37 , 1.12 ) ( N/A , 0.01 , 0.1 )	2658.8 3615.0	0.2993 143.8 143.8	37.8463 [ 48.4781 ]	78.1%			
4:2FTS	( 327.0 / 307.0 ) 39943309 ( 327.0 / 81.0 ) 29603285	( 5.22 , 1.00 ) ( 0.00 , N/A , 0.0 )	5776.1 5249.2	0.7411 109.8 109.8	153.2757 [ 186.9055 ]	82.0%			
6:2FTS	( 427.0 / 407.0 ) 29385691 ( 427.0 / 81.0 ) 23406118	( 6.73 , 1.00 ) ( 0.00 , N/A , 0.0 )	3922.8 5055.0	0.7965 96.2 96.2	188.8167 [ 189.8085 ]	99.5%			
8:2FTS	( 527.0 / 507.0 ) 32033130 ( 527.0 / 81.0 ) 26056569	( 7.89 , 1.00 ) ( 0.00 , N/A , 0.1 )	2531.0 3355.4	0.8134 97.9 97.9	169.8673 [ 191.6577 ]	88.6%			

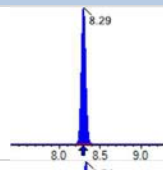
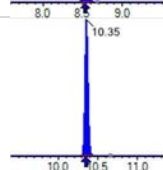
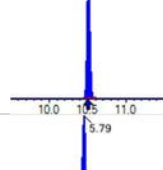
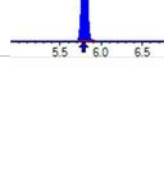



Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	( 498.0 / 78.0 ) N/A ( 498.0 / 478.0 ) N/A	( N/A , N/A ) ( N/A , 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 ) 76556778 ( 512.0 / 169.0 ) 65118925	( 10.41 , 1.00 ) ( -0.02 , N/A , 0.3 )	2387.5 3511.2	0.8506 103.8 103.8	206.1452 [ 200.0000 ]	103.1%			
NEIFOSA	( 526.0 / 219.0 ) 82865198 ( 526.0 / 169.0 ) 88002594	( 10.55 , 1.00 ) ( -0.02 , N/A , 0.1 )	16651.6 18744.9	1.0620 83.7 83.7	160.3500 [ 200.0000 ]	80.2%			
NMeFOSAA	( 570.0 / 419.0 ) 12224827 ( 570.0 / 483.0 ) 5469499	( 8.30 , 1.00 ) ( 0.00 , N/A , 0.1 )	4664.0 2463.8	0.4474 94.2 94.2	48.3498 [ 50.0000 ]	96.7%			
NEIFOSAA	( 584.0 / 419.0 ) 10977527 ( 584.0 / 526.0 ) 6240818	( 8.54 , 1.00 ) ( 0.01 , N/A , 0.0 )	2642.6 1856.2	0.5685 111.4 111.4	51.7785 [ 50.0000 ]	103.6%			
NMeFOSE	( 616.0 / 59.0 ) 45235489	( 10.35 , 1.00 ) ( 0.00 , N/A , 0.0 )	3201.0	N/A 0.0 0.0	186.7620 [ 200.0000 ]	93.4%			
NEtFOSE	( 630.0 / 59.0 ) 47659476	( 10.51 , 1.00 ) ( 0.00 , N/A , 0.0 )	1276.6	N/A 0.0 0.0	175.0607 [ 200.0000 ]	87.5%			
HFPO-DA	( 285.0 / 169.0 ) 27064622 ( 285.0 / 185.0 ) 56107744	( 5.80 , 1.00 ) ( 0.00 , N/A , 0.0 )	4176.1 4089.0	2.0731 75.3 75.3	98.6051 [ 100.0000 ]	98.6%			
ADONA	( 377.0 / 85.0 ) 63651267 ( 377.0 / 251.0 ) 8645557	( 6.58 , 1.14 ) ( N/A , 0.01 , 0.0 )	2896.3 3684.8	0.1358 154.0 154.0	67.4326 [ 94.2700 ]	71.5%			IR2,
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 ) 2348441 ( 241.0 / 117.0 ) 3695398	( 4.06 , 0.89 ) ( N/A , 0.02 , 0.0 )	2293.1 2774.8	1.5736 99.7 99.7	200.2861 [ 200.0000 ]	100.1%			
5:3FTCA	( 341.0 / 236.7 ) 16701638 ( 341.0 / 217.0 ) 27334931	( 6.05 , 1.10 ) ( N/A , 0.02 , 0.0 )	2530.4 3328.2	1.6367 96.0 96.0	242.3031 [ 200.0000 ]	121.2%			
7:3FTCA	( 441.0 / 317.0 ) 30019668 ( 441.0 / 337.0 ) 25999080	( 7.89 , 1.44 ) ( N/A , 0.01 , 0.0 )	1616.8 1863.7	0.8661 100.6 100.6	235.7560 [ 200.0000 ]	117.9%			
PFEESA	( 315.0 / 135.0 ) 44504868 ( 315.0 / 83.0 ) 14830352	( 5.90 , 1.08 ) ( N/A , 0.02 , 0.0 )	3473.2 4026.5	0.3332 130.5 130.5	73.5006 [ 89.2459 ]	82.4%			
PFMPA	( 229.0 / 85.0 ) 8205055	( 3.90 , 0.86 ) ( N/A , 0.02 , 0.0 )	5685.5	N/A 0.0 0.0	112.1506 [ 100.0000 ]	112.2%			
PFMBA	( 279.0 / 85.0 ) 30499882	( 4.86 , 1.07 ) ( N/A , 0.02 , 0.0 )	5844.0	N/A 0.0 0.0	99.4098 [ 100.0000 ]	99.4%			
NFDHA	( 295.0 / 201.0 ) 26402610 ( 295.0 / 85.0 ) 26363507	( 5.39 , 0.98 ) ( N/A , 0.02 , 0.0 )	3971.6 5150.6	0.9985 99.7 99.7	94.3710 [ 100.0000 ]	94.4%			
13C3_PFBA_IIS	( 216.0 / 172.0 ) 158427	( 3.50 , N/A ) ( N/A , 0.01 , N/A )	747.3	N/A	0.7073 [ 1.0000 ]	70.7% { 70.3% }			
13C2_PFHxA_IIS	( 315.0 / 270.0 ) 454892	( 5.49 , N/A ) ( N/A , 0.02 , N/A )	3086.9	N/A	0.9011 [ 1.0000 ]	90.1% { 94.7% }			
13C4_PFOA_IIS	( 417.0 / 372.0 ) 531465	( 7.00 , N/A ) ( N/A , 0.02 , N/A )	2829.3	N/A	0.8386 [ 1.0000 ]	83.9% { 89.6% }			
13C5_PFNA_IIS	( 468.0 / 423.0 ) 563682	( 7.60 , N/A ) ( N/A , 0.01 , N/A )	1482.9	N/A	0.9328 [ 1.0000 ]	93.3% { 95.1% }			

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 ) 548948	( 8.15 , N/A ) ( N/A , 0.00 , N/A )	1038.3	N/A	0.9487 [ 1.0000 ]	94.9% { 100.8% }			
18O2_PFHxS_IIS	( 403.0 / 83.9 ) 779148	( 7.15 , N/A ) ( N/A , 0.01 , N/A )	934.0	N/A	0.9098 [ 1.0000 ]	91.0% { 92.3% }			
13C4_PFOS_IIS	( 503.0 / 79.9 ) 1418626	( 8.36 , N/A ) ( N/A , 0.01 , N/A )	1316.0	N/A	0.9424 [ 1.0000 ]	94.2% { 93.8% }			
13C4_PFBA_EIS	( 217.0 / 172.0 ) 1460766	( 3.50 , N/A ) ( N/A , 0.01 , N/A )	3458.8	N/A	7.7515 [ 8.0000 ]	96.9% { 62.6% }			
13C5_PFPeA_EIS	( 268.0 / 223.0 ) 1628440	( 4.55 , N/A ) ( N/A , 0.02 , N/A )	2993.6	N/A	3.2719 [ 4.0000 ]	81.8% { 67.3% }			
13C5_PFHxA_EIS	( 318.0 / 273.0 ) 1146711	( 5.49 , N/A ) ( N/A , 0.02 , N/A )	3359.8	N/A	1.8404 [ 2.0000 ]	92.0% { 74.7% }			
13C4_PFHpA_EIS	( 367.0 / 322.0 ) 1010584	( 6.30 , N/A ) ( N/A , 0.01 , N/A )	1691.9	N/A	1.8719 [ 2.0000 ]	93.6% { 80.0% }			
13C8_PFOA_EIS	( 421.0 / 376.0 ) 1278931	( 6.99 , N/A ) ( N/A , 0.01 , N/A )	2254.9	N/A	2.1248 [ 2.0000 ]	106.2% { 83.0% }			
13C9_PFNA_EIS	( 472.0 / 427.0 ) 557891	( 7.60 , N/A ) ( N/A , 0.01 , N/A )	2610.8	N/A	0.9341 [ 1.0000 ]	93.4% { 79.4% }			
13C6_PFDA_EIS	( 519.0 / 474.0 ) 635337	( 8.15 , N/A ) ( N/A , 0.00 , N/A )	2044.6	N/A	0.9179 [ 1.0000 ]	91.8% { 81.8% }			
13C7_PFUnA_EIS	( 570.0 / 525.0 ) 637370	( 8.63 , N/A ) ( N/A , 0.00 , N/A )	1798.4	N/A	0.9261 [ 1.0000 ]	92.6% { 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
13C2_PFDa_EIS	( 615.0 / 570.0 ) 537611	( 8.93, N/A ) ( N/A, 0.01, N/A )	1771.9	N/A	0.9107 [ 1.0000 ]	91.1% { 81.8% }			
13C2_PFTeDA_EIS	( 715.0 / 670.0 ) 526472	( 9.29, N/A ) ( N/A, 0.00, N/A )	1321.1	N/A	0.9733 [ 1.0000 ]	97.3% { 86.3% }			
13C3_PFBs_EIS	( 302.0 / 80.0 ) 2735550	( 5.45, N/A ) ( N/A, 0.02, N/A )	3421.3	N/A	1.6775 [ 2.0000 ]	83.9% { 69.3% }			
13C3_PFHxS_EIS	( 402.0 / 80.0 ) 1607241	( 7.14, N/A ) ( N/A, 0.01, N/A )	2197.8	N/A	1.8976 [ 2.0000 ]	94.9% { 80.8% }			
13C8_PFOS_EIS	( 507.0 / 80.0 ) 3491992	( 8.36, N/A ) ( N/A, 0.01, N/A )	536.1	N/A	1.7930 [ 2.0000 ]	89.7% { 79.0% }			
13C2_4:2FTS_EIS	( 329.0 / 81.0 ) 381662	( 5.22, N/A ) ( N/A, 0.02, N/A )	1246.7	N/A	4.2402 [ 4.0000 ]	106.0% { 93.2% }			
13C2_6:2FTS_EIS	( 429.0 / 81.0 ) 438735	( 6.73, N/A ) ( N/A, 0.01, N/A )	1279.4	N/A	4.4405 [ 4.0000 ]	111.0% { 95.8% }			
13C2_8:2FTS_EIS	( 529.0 / 81.0 ) 692502	( 7.89, N/A ) ( N/A, 0.01, N/A )	2008.3	N/A	4.9610 [ 4.0000 ]	124.0% { 114.4% }			
13C8_PFOsa_EIS	( 506.0 / 78.0 ) 5474172	( 9.84, N/A ) ( N/A, 0.01, N/A )	2848.0	N/A	1.5142 [ 2.0000 ]	75.7% { 63.9% }			
D3_NMeFOsa_EIS	( 515.0 / 169.0 ) 1535713	( 10.42, N/A ) ( N/A, 0.01, N/A )	1409.7	N/A	1.8030 [ 2.0000 ]	90.2% { 84.2% }			
D5_NEtFOsa_EIS	( 531.0 / 169.0 ) 1106313	( 10.57, N/A ) ( N/A, 0.01, N/A )	2709.4	N/A	1.4515 [ 2.0000 ]	72.6% { 64.2% }			

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 ) 1288746	( 8.29 , N/A ) ( N/A , 0.01 , N/A )	2014.3	N/A	4.0890 [ 4.0000 ]	102.2% { 86.6% }			
D5_EtFOSAA_EIS	( 589.0 / 419.0 ) 1009687	( 8.54 , N/A ) ( N/A , 0.01 , N/A )	2733.1	N/A	3.9125 [ 4.0000 ]	97.8% { 90.0% }			
D7_NMeFOSE_EIS	( 623.0 / 58.9 ) 4845449	( 10.35 , N/A ) ( N/A , 0.01 , N/A )	1915.8	N/A	15.6043 [ 20.0000 ]	78.0% { 67.4% }			
D9_NEtFOSE_EIS	( 639.0 / 58.9 ) 5721656	( 10.51 , N/A ) ( N/A , 0.01 , N/A )	1896.2	N/A	14.3883 [ 20.0000 ]	71.9% { 60.1% }			
13C3_HFPODA_EIS	( 287.0 / 169.0 ) 2776685	( 5.79 , N/A ) ( N/A , 0.02 , N/A )	2495.0	N/A	7.3191 [ 8.0000 ]	91.5% { 75.8% }			

**SECOND-SOURCE CALIBRATION VERIFICATION****EPA 1633****Laboratory:** APPL, LLC**SDG:****Client:** AECOM**Project:** Red Hill AFFF Assessment Sampling**Calibration:** 2307007**Laboratory ID:** SC00647-SCV1**Sequence:** SC00647**Standard ID:** 23B0089

ANALYTE	EXPECTED (ng/mL)	FOUND (ng/mL)	% DRIFT	QC LIMIT
PFBA	8.00	7.72	-3.5	30.00
PFPEA	4.00	4.05	1.3	30.00
PFHXA	2.00	1.91	-4.5	30.00
PFHPA	2.00	2.00	-0.04	30.00
PFOA	2.00	1.96	-1.9	30.00
PFNA	2.00	2.05	2.3	30.00
PFDA	2.00	1.89	-5.7	30.00
PFUnA	2.00	1.99	-0.5	30.00
PFDOA	2.00	1.94	-3.1	30.00
PFTRDA	2.00	2.07	3.5	30.00
PFTEDA	2.00	1.79	-10.5	30.00
PFBS	1.77	1.71	-3.2	30.00
PFPEs	1.88	1.97	4.9	30.00
PFHXS	1.83	1.79	-2.1	30.00
PFHPS	1.91	1.86	-2.5	30.00
PFOS	1.86	1.76	-5.3	30.00
PFNS	1.92	1.88	-1.8	30.00
PFDS	1.93	1.96	1.4	30.00
PFDOS	1.94	2.02	3.9	30.00
4:2FTS	7.50	7.61	1.5	30.00
6:2FTS	7.60	7.11	-6.4	30.00
8:2FTS	7.68	7.91	3.1	30.00
PFOSA	2.00	2.06	2.8	30.00
NMeFOSA	8.00	7.51	-6.1	30.00
NEtFOSA	8.00	7.19	-10.2	30.00
NMeFOSAA	2.00	1.87	-6.4	30.00
NEtFOSAA	2.00	1.96	-2.2	30.00

**SECOND-SOURCE CALIBRATION VERIFICATION****EPA 1633****Laboratory:** APPL, LLC**SDG:****Client:** AECOM**Project:** Red Hill AFFF Assessment Sampling**Calibration:** 2307007**Laboratory ID:** SC00647-SCV1**Sequence:** SC00647**Standard ID:** 23B0089

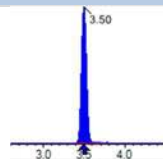
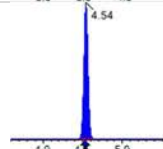
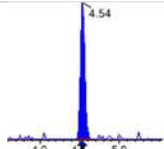
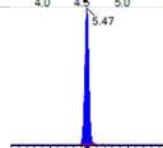
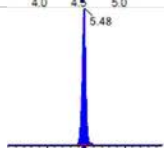
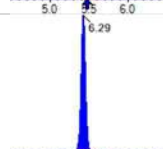
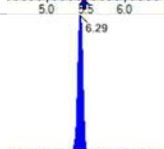
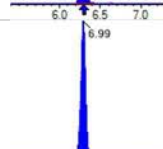
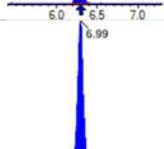
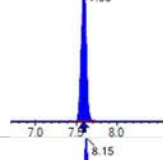
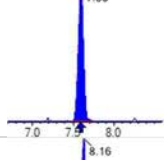
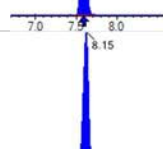
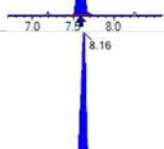
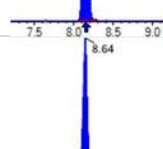
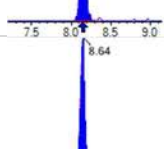
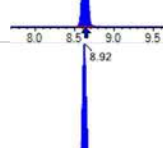
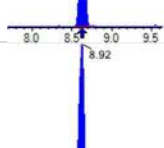
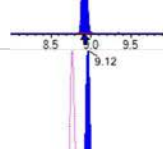
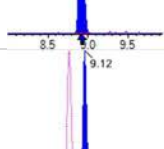
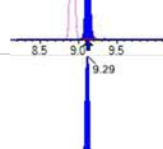
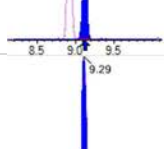
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NEtFOSE	8.00	8.29	3.6	30.00
HFPO-DA	4.00	3.67	-8.2	30.00
ADONA	3.78	4.12	9.1	30.00
PFEESA	3.56	3.63	1.9	30.00
PFMPA	4.00	4.17	4.3	30.00
PFMBA	4.00	3.96	-1.0	30.00
NFDHA	4.00	3.90	-2.4	30.00
9CL-PF3ONS	3.74	3.74	-0.03	30.00
11CL-PF3OUDS	3.78	4.31	13.9	30.00
3:3FTCA	8.00	8.22	2.7	30.00
5:3FTCA	8.00	7.53	-5.9	30.00
7:3FTCA	8.00	7.65	-4.4	30.00
13C4-PFBA	8.00	8.13	1.6	30.00
13C5-PFPEA	4.00	4.18	4.5	30.00
13C5-PFHXA	2.00	2.14	6.9	30.00
13C4-PFHFA	2.00	2.05	2.7	30.00
13C8-PFOA	2.00	1.91	-4.5	30.00
13C9-PFNA	1.00	1.05	5.5	30.00
13C6-PFDA	1.00	1.08	8.4	30.00
13C7-PFUnA	1.00	1.02	1.9	30.00
13C2-PFDOA	1.00	0.992	-0.8	30.00
13C2-PFTEDA	1.00	1.11	10.6	30.00
13C3-PFBS	2.00	2.17	8.5	30.00
13C3-PFHXS	2.00	2.09	4.6	30.00
13C8-PFOS	2.00	1.98	-0.8	30.00
13C2-4:2FTS	4.00	3.89	-2.9	30.00
13C2-6:2FTS	4.00	4.20	5.1	30.00
13C2-8:2FTS	4.00	3.89	-2.9	30.00

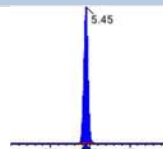
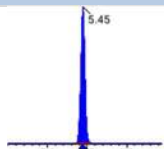
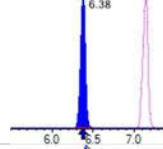
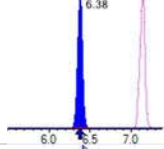
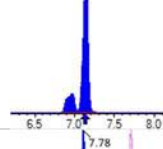
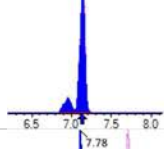
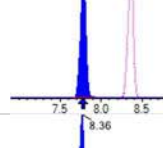
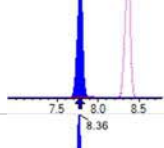
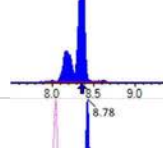
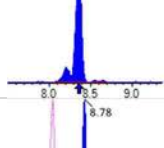
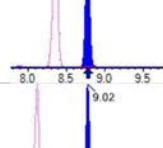
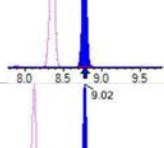
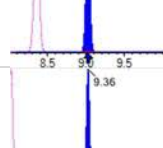
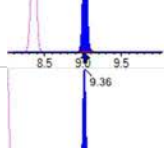
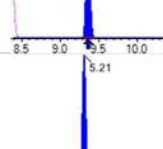
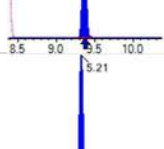
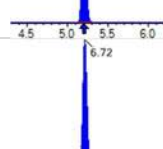
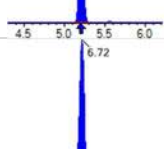
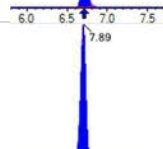
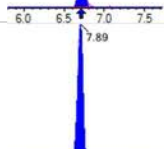
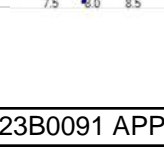
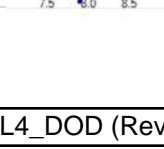
**SECOND-SOURCE CALIBRATION VERIFICATION****EPA 1633****Laboratory:** APPL, LLC**SDG:****Client:** AECOM**Project:** Red Hill AFFF Assessment Sampling**Calibration:** 2307007**Laboratory ID:** SC00647-SCV1**Sequence:** SC00647**Standard ID:** 23B0089

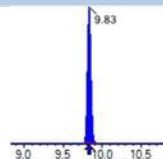
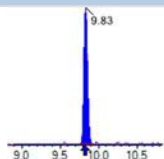
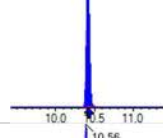
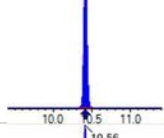
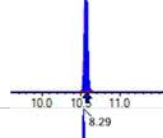
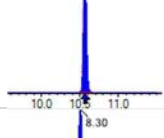
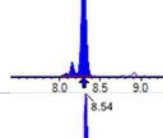
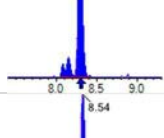
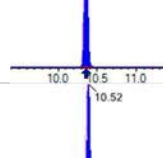
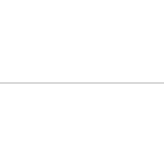
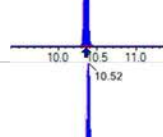
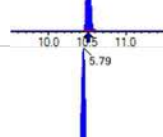
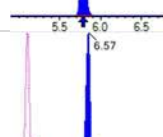
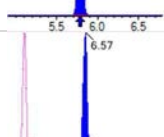
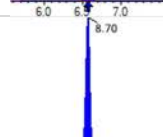
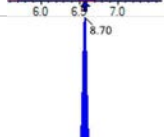
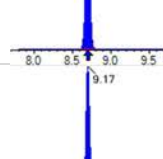
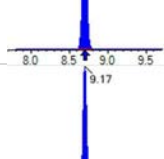
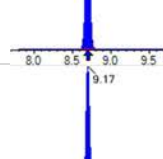
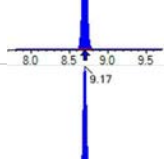
13C8-PFOSA	2.00	2.05	2.4	30.00
D3-NMEFOSA	2.00	2.02	1.1	30.00
D5-NETFOSA	2.00	2.07	3.3	30.00
D3-NMEFOSAA	4.00	3.99	-0.2	30.00
D5-NETFOSAA	4.00	3.83	-4.3	30.00
D7-NMEFOSE	20.0	20.3	1.7	30.00
D9-NETFOSSE	20.0	19.8	-1.2	30.00
13C3-HFPO-DA	8.00	8.77	9.6	30.00

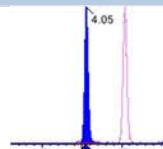
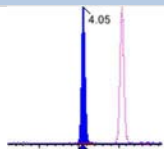
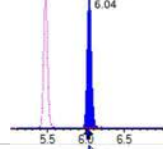
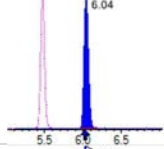
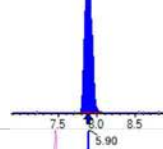
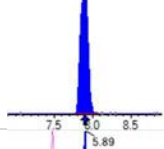
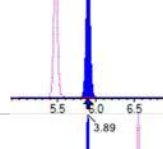
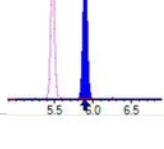
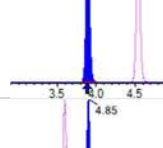
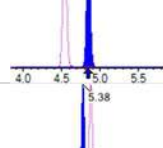
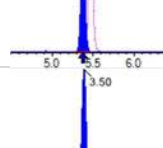
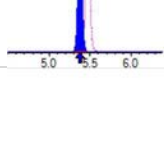
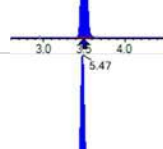
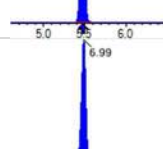
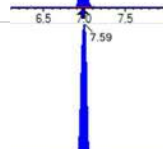

\* Values outside of QC limits

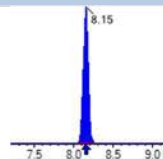
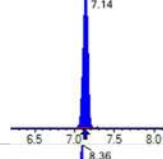
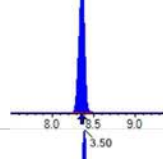
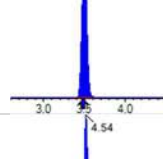
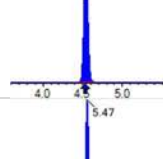
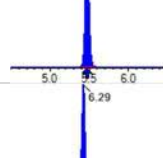
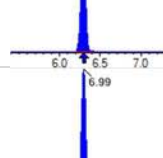
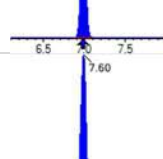
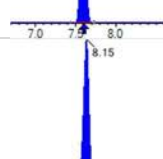
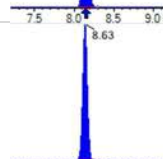
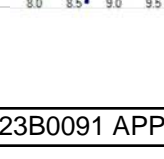


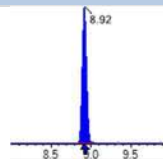
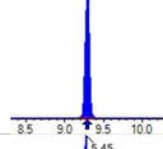
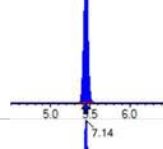
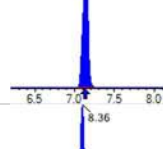
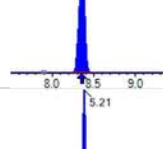
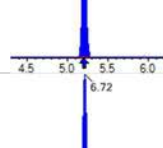
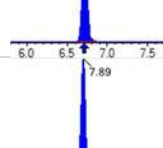
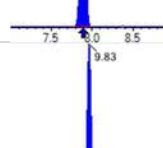
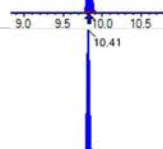
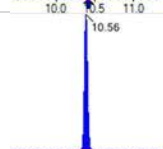
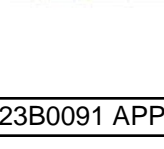
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) ( $\Delta$ RT-[min], $\Delta$ RT- CV[min], $\Delta$ RT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	( 213.0 / 169.0 ) 1855837	( 3.50 , 1.00 ) ( 0.00 , N/A , 0.0 )	214.2	N/A 0.0 0.0	7.7193 [ 8.0000 ]	96.5%			
PFPeA	( 263.0 / 219.0 ) 1816304 ( 263.0 / 69.0 ) 22222	( 4.54 , 1.00 ) ( 0.00 , N/A , 0.3 )	2797.4 227.3	0.0122 114.3 114.3	4.0506 [ 4.0000 ]	101.3%			
PFHxA	( 313.0 / 269.0 ) 1172574 ( 313.0 / 119.0 ) 109802	( 5.47 , 1.00 ) ( 0.00 , N/A , -0.2 )	5660.8 4014564.1	0.0936 94.0 94.0	1.9097 [ 2.0000 ]	95.5%			
PFHpA	( 363.0 / 319.0 ) 980875 ( 363.0 / 169.0 ) 294159	( 6.29 , 1.00 ) ( 0.00 , N/A , 0.0 )	43558.2 199114.1	0.2999 93.5 93.5	1.9991 [ 2.0000 ]	100.0%			
PFOA	( 413.0 / 369.0 ) 1269405 ( 413.0 / 169.0 ) 434013	( 6.99 , 1.00 ) ( 0.00 , N/A , -0.1 )	9343.6 2964.4	0.3419 102.2 102.2	1.9616 [ 2.0000 ]	98.1%			
PFNA	( 463.0 / 419.0 ) 1176960 ( 463.0 / 169.0 ) 257649	( 7.60 , 1.00 ) ( 0.00 , N/A , 0.0 )	10298.1 5708.0	0.2189 97.8 97.8	2.0460 [ 2.0000 ]	102.3%			
PFDA	( 513.0 / 469.0 ) 1273474 ( 513.0 / 169.0 ) 146381	( 8.15 , 1.00 ) ( 0.00 , N/A , -0.3 )	1241.8 651.1	0.1149 94.5 94.5	1.8850 [ 2.0000 ]	94.3%			
PFUnA	( 563.0 / 519.0 ) 1187758 ( 563.0 / 169.0 ) 148979	( 8.64 , 1.00 ) ( 0.00 , N/A , 0.0 )	1593.9 1026.2	0.1254 119.6 119.6	1.9902 [ 2.0000 ]	99.5%			
PFDoA	( 613.0 / 569.0 ) 1022182 ( 613.0 / 169.0 ) 171947	( 8.92 , 1.00 ) ( 0.00 , N/A , -0.4 )	1893.1 1840.4	0.1682 102.8 102.8	1.9381 [ 2.0000 ]	96.9%			
PFTTrDA	( 663.0 / 619.0 ) 988300 ( 663.0 / 169.0 ) 288733	( 9.12 , 1.02 ) ( N/A , 0.00 , 0.1 )	1735.7 884.1	0.2922 122.2 122.2	2.0696 [ 2.0000 ]	103.5%			
PFTeDA	( 713.0 / 669.0 ) 948401 ( 713.0 / 169.0 ) 229129	( 9.29 , 1.00 ) ( 0.00 , N/A , 0.3 )	1207.7 516.0	0.2416 116.9 116.9	1.7896 [ 2.0000 ]	89.5%			

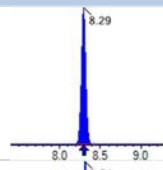
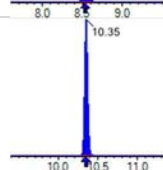
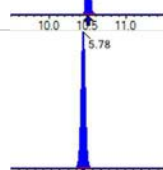


Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	( 299.0 / 80.0 ) 1944081 ( 299.0 / 99.0 ) 1240225	( 5.45 , 1.00 ) ( 0.00 , N/A , 0.0 )	3671.2 2990.3	0.6379 98.7 98.7	1.7133 [ 1.7695 ]	96.8%			
PFPeS	( 349.0 / 80.0 ) 3474653 ( 349.0 / 99.0 ) 1195565	( 6.38 , 0.89 ) ( N/A , 0.01 , 0.0 )	40358961.1 12054.4	0.3441 96.4 96.4	1.9714 [ 1.8768 ]	105.0%			
PFHxS	( 399.0 / 80.0 ) 2517471 ( 399.0 / 99.0 ) 811394	( 7.14 , 1.00 ) ( 0.00 , N/A , 0.0 )	63031.5 6250.9	0.3223 99.2 99.2	1.7914 [ 1.8220 ]	98.3%			
PFHpS	( 449.0 / 80.0 ) 3034087 ( 449.0 / 99.0 ) 825667	( 7.78 , 0.93 ) ( N/A , 0.01 , 0.1 )	22648185.7 36047.8	0.2721 100.0 100.0	1.8631 [ 1.9028 ]	97.9%			
PFOS	( 499.0 / 80.0 ) 3970613 ( 499.0 / 99.0 ) 875204	( 8.36 , 1.00 ) ( 0.00 , N/A , -0.1 )	1176.9 1242.3	0.2204 99.6 99.6	1.7618 [ 1.8550 ]	95.0%			
PFNS	( 549.0 / 80.0 ) 4773847 ( 549.0 / 99.0 ) 1126371	( 8.78 , 1.05 ) ( N/A , 0.00 , 0.1 )	16422.3 9408.7	0.2359 107.6 107.6	1.8848 [ 1.9198 ]	98.2%			
PFDS	( 599.0 / 80.0 ) 5389437 ( 599.0 / 99.0 ) 1155934	( 9.02 , 1.08 ) ( N/A , 0.00 , 0.0 )	3964.5 1730.1	0.2145 98.4 98.4	1.9561 [ 1.9262 ]	101.6%			
PFDoS	( 699.0 / 80.0 ) 3541779 ( 699.0 / 99.0 ) 730787	( 9.36 , 1.12 ) ( N/A , 0.00 , 0.0 )	1827.6 950.3	0.2063 99.1 99.1	2.0155 [ 1.9391 ]	103.9%			
4:2FTS	( 327.0 / 307.0 ) 1875591 ( 327.0 / 81.0 ) 1270793	( 5.21 , 1.00 ) ( 0.00 , N/A , -0.1 )	4384.0 2751.3	0.6775 100.4 100.4	7.6108 [ 7.4762 ]	101.8%			
6:2FTS	( 427.0 / 407.0 ) 1081545 ( 427.0 / 81.0 ) 826407	( 6.72 , 1.00 ) ( 0.00 , N/A , -0.1 )	2419.4 1671.5	0.7641 92.3 92.3	7.1118 [ 7.5923 ]	93.7%			
8:2FTS	( 527.0 / 507.0 ) 1206144 ( 527.0 / 81.0 ) 1085149	( 7.89 , 1.00 ) ( 0.00 , N/A , -0.2 )	1836.7 1318.7	0.8997 108.2 108.2	7.9144 [ 7.6663 ]	103.2%			

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 ) 6738795 ( 498.0 / 478.0 ) 139185	( 9.83 , 1.00 ) ( 0.00 , N/A , 0.1 )	3420.6 518.2	0.0207 100.6 100.6	2.0569 [ 2.0000 ]	102.8%			
NMeFOSA	( 512.0 / 219.0 ) 6068879 ( 512.0 / 169.0 ) 4266094	( 10.41 , 1.00 ) ( 0.00 , N/A , 0.0 )	8283.1 7296.2	0.7029 85.8 85.8	7.5097 [ 8.0000 ]	93.9%			
NEiFOSA	( 526.0 / 219.0 ) 5846000 ( 526.0 / 169.0 ) 6647247	( 10.56 , 1.00 ) ( 0.01 , N/A , 0.0 )	12368.8 8629.6	1.1371 89.6 89.6	7.1864 [ 8.0000 ]	89.8%			
NMeFOSAA	( 570.0 / 419.0 ) 511022 ( 570.0 / 483.0 ) 237635	( 8.29 , 1.00 ) ( 0.00 , N/A , -0.1 )	1494.6 1575.2	0.4650 98.0 98.0	1.8726 [ 2.0000 ]	93.6%			
NEiFOSAA	( 584.0 / 419.0 ) 448836 ( 584.0 / 526.0 ) 239853	( 8.54 , 1.00 ) ( 0.01 , N/A , 0.0 )	9002.5 186834.1	0.5344 104.7 104.7	1.9560 [ 2.0000 ]	97.8%			
NMeFOSE	( 616.0 / 59.0 ) 2858949	( 10.36 , 1.00 ) ( 0.01 , N/A , 0.0 )	2885.4	N/A 0.0 0.0	8.1903 [ 8.0000 ]	102.4%			
NEiFOSE	( 630.0 / 59.0 ) 3426435	( 10.52 , 1.00 ) ( 0.01 , N/A , 0.0 )	2679.0	N/A 0.0 0.0	8.2899 [ 8.0000 ]	103.6%			
HFPO-DA	( 285.0 / 169.0 ) 1266072 ( 285.0 / 185.0 ) 3773275	( 5.79 , 1.00 ) ( 0.00 , N/A , 0.0 )	2016.7 3174.2	2.9803 108.2 108.2	3.6730 [ 4.0000 ]	91.8%			
ADONA	( 377.0 / 85.0 ) 4888516 ( 377.0 / 251.0 ) 451545	( 6.57 , 1.14 ) ( N/A , 0.00 , 0.0 )	2673.1 1707.9	0.0924 104.8 104.8	4.1239 [ 3.7708 ]	109.4%			
9CI-Pf3ONS	( 531.0 / 351.0 ) 13910227 ( 533.0 / 353.0 ) 4138206	( 8.70 , 1.50 ) ( N/A , 0.00 , 0.1 )	3737.3 2578.6	0.2975 85.4 85.4	3.7389 [ 3.7330 ]	100.2%			
11CI-PF3OUDS	( 631.0 / 451.0 ) 8315091 ( 633.0 / 453.0 ) 2851144	( 9.17 , 1.59 ) ( N/A , 0.00 , 0.0 )	2700.8 2174.5	0.3429 94.8 94.8	4.3058 [ 3.7728 ]	114.1%			

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 ) 88786 ( 241.0 / 117.0 ) 140594	( 4.05 , 0.89 ) ( N/A , 0.01 , 0.1 )	767.3 708.5	1.5835 100.4 100.4	8.2179 [ 8.0000 ]	102.7%			
5:3FTCA	( 341.0 / 236.7 ) 631861 ( 341.0 / 217.0 ) 1085339	( 6.04 , 1.10 ) ( N/A , 0.01 , 0.1 )	1441.5 1128.7	1.7177 100.7 100.7	7.5252 [ 8.0000 ]	94.1%			
7:3FTCA	( 441.0 / 317.0 ) 1186511 ( 441.0 / 337.0 ) 1063466	( 7.89 , 1.44 ) ( N/A , 0.01 , -0.3 )	696.0 619.9	0.8963 104.2 104.2	7.6493 [ 8.0000 ]	95.6%			
PFEESA	( 315.0 / 135.0 ) 2674575 ( 315.0 / 83.0 ) 698671	( 5.90 , 1.08 ) ( N/A , 0.01 , 0.0 )	4876.6 1560.2	0.2612 102.3 102.3	3.6260 [ 3.5698 ]	101.6%			
PFMPA	( 229.0 / 85.0 ) 408761	( 3.89 , 0.86 ) ( N/A , 0.01 , 0.0 )	2687.6	N/A 0.0 0.0	4.1734 [ 4.0000 ]	104.3%			
PFMBA	( 279.0 / 85.0 ) 1626572	( 4.85 , 1.07 ) ( N/A , 0.01 , 0.0 )	2449.4	N/A 0.0 0.0	3.9601 [ 4.0000 ]	99.0%			
NFDHA	( 295.0 / 201.0 ) 1330784 ( 295.0 / 85.0 ) 1362923	( 5.38 , 0.98 ) ( N/A , 0.01 , 0.0 )	2155.6 2870.7	1.0242 102.3 102.3	3.9048 [ 4.0000 ]	97.6%			
13C3_PFBA_IIS	( 216.0 / 172.0 ) 240066	( 3.50 , N/A ) ( N/A , 0.01 , N/A )	1412.3	N/A	1.0718 [ 1.0000 ]	107.2% { 106.5% }			
13C2_PFHxA_IIS	( 315.0 / 270.0 ) 476905	( 5.47 , N/A ) ( N/A , 0.01 , N/A )	2780.9	N/A	0.9447 [ 1.0000 ]	94.5% { 99.3% }			
13C4_PFOA_IIS	( 417.0 / 372.0 ) 672267	( 6.99 , N/A ) ( N/A , 0.01 , N/A )	1232.7	N/A	1.0607 [ 1.0000 ]	106.1% { 113.3% }			
13C5_PFNA_IIS	( 468.0 / 423.0 ) 581759	( 7.59 , N/A ) ( N/A , 0.01 , N/A )	3145.9	N/A	0.9628 [ 1.0000 ]	96.3% { 98.1% }			

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 ) 562140	( 8.15 , N/A ) ( N/A , 0.00 , N/A )	25539.1	N/A	0.9715 [ 1.0000 ]	97.1% { 103.2% }			
18O2_PFHxS_IIS	( 403.0 / 83.9 ) 804048	( 7.14 , N/A ) ( N/A , 0.01 , N/A )	2686.8	N/A	0.9389 [ 1.0000 ]	93.9% { 95.2% }			
13C4_PFOS_IIS	( 503.0 / 79.9 ) 1569000	( 8.36 , N/A ) ( N/A , 0.01 , N/A )	1276.6	N/A	1.0423 [ 1.0000 ]	104.2% { 103.8% }			
13C4_PFBA_EIS	( 217.0 / 172.0 ) 2321888	( 3.50 , N/A ) ( N/A , 0.01 , N/A )	4695.7	N/A	8.1311 [ 8.0000 ]	101.6% { 99.4% }			
13C5_PFPeA_EIS	( 268.0 / 223.0 ) 2180078	( 4.54 , N/A ) ( N/A , 0.01 , N/A )	3494.2	N/A	4.1780 [ 4.0000 ]	104.5% { 90.1% }			
13C5_PFHxA_EIS	( 318.0 / 273.0 ) 1396881	( 5.47 , N/A ) ( N/A , 0.01 , N/A )	2597.4	N/A	2.1384 [ 2.0000 ]	106.9% { 91.0% }			
13C4_PFHpA_EIS	( 367.0 / 322.0 ) 1162104	( 6.29 , N/A ) ( N/A , 0.00 , N/A )	3086.1	N/A	2.0532 [ 2.0000 ]	102.7% { 92.0% }			
13C8_PFOA_EIS	( 421.0 / 376.0 ) 1453557	( 6.99 , N/A ) ( N/A , 0.01 , N/A )	6980.9	N/A	1.9092 [ 2.0000 ]	95.5% { 94.4% }			
13C9_PFNA_EIS	( 472.0 / 427.0 ) 650184	( 7.60 , N/A ) ( N/A , 0.00 , N/A )	4374.6	N/A	1.0548 [ 1.0000 ]	105.5% { 92.5% }			
13C6_PFDA_EIS	( 519.0 / 474.0 ) 768006	( 8.15 , N/A ) ( N/A , 0.01 , N/A )	1522.6	N/A	1.0836 [ 1.0000 ]	108.4% { 98.9% }			
13C7_PFUnA_EIS	( 570.0 / 525.0 ) 718332	( 8.63 , N/A ) ( N/A , 0.00 , N/A )	1490.8	N/A	1.0192 [ 1.0000 ]	101.9% { 93.1% }			

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 ) 599764	( 8.92 , N/A ) ( N/A , 0.00 , N/A )	1573.7	N/A	0.9922 [ 1.0000 ]	99.2% { 91.3% }			
13C2_PFTeDA_EIS	( 715.0 / 670.0 ) 612411	( 9.29 , N/A ) ( N/A , 0.00 , N/A )	2166.8	N/A	1.1056 [ 1.0000 ]	110.6% { 100.4% }			
13C3_PFBs_EIS	( 302.0 / 80.0 ) 3651854	( 5.45 , N/A ) ( N/A , 0.01 , N/A )	3505.6	N/A	2.1701 [ 2.0000 ]	108.5% { 92.5% }			
13C3_PFHxS_EIS	( 402.0 / 80.0 ) 1827789	( 7.14 , N/A ) ( N/A , 0.01 , N/A )	2735.6	N/A	2.0911 [ 2.0000 ]	104.6% { 91.9% }			
13C8_PFOS_EIS	( 507.0 / 80.0 ) 4274649	( 8.36 , N/A ) ( N/A , 0.01 , N/A )	1379.1	N/A	1.9845 [ 2.0000 ]	99.2% { 96.7% }			
13C2_4:2FTS_EIS	( 329.0 / 81.0 ) 360923	( 5.21 , N/A ) ( N/A , 0.01 , N/A )	1374.9	N/A	3.8857 [ 4.0000 ]	97.1% { 88.2% }			
13C2_6:2FTS_EIS	( 429.0 / 81.0 ) 428716	( 6.72 , N/A ) ( N/A , 0.01 , N/A )	1900.2	N/A	4.2047 [ 4.0000 ]	105.1% { 93.6% }			
13C2_8:2FTS_EIS	( 529.0 / 81.0 ) 559642	( 7.89 , N/A ) ( N/A , 0.00 , N/A )	1311.1	N/A	3.8850 [ 4.0000 ]	97.1% { 92.5% }			
13C8_PFOsa_EIS	( 506.0 / 78.0 ) 8192381	( 9.83 , N/A ) ( N/A , 0.00 , N/A )	3577.3	N/A	2.0489 [ 2.0000 ]	102.4% { 95.6% }			
D3_NMeFOSA_EIS	( 515.0 / 169.0 ) 1905355	( 10.41 , N/A ) ( N/A , 0.00 , N/A )	2669.3	N/A	2.0226 [ 2.0000 ]	101.1% { 104.4% }			
D5_NEtFOSA_EIS	( 531.0 / 169.0 ) 1741498	( 10.56 , N/A ) ( N/A , 0.00 , N/A )	3801.6	N/A	2.0659 [ 2.0000 ]	103.3% { 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
D3_MeFOSAA_EIS	( 573.0 / 419.0 ) 1390921	( 8.29 , N/A ) ( N/A , 0.01 , N/A )	2287.2	N/A	3.9903 [ 4.0000 ]	99.8% { 93.4% }			
D5_EtFOSAA_EIS	( 589.0 / 419.0 ) 1092823	( 8.54 , N/A ) ( N/A , 0.00 , N/A )	20352.5	N/A	3.8288 [ 4.0000 ]	95.7% { 97.4% }			
D7_NMeFOSE_EIS	( 623.0 / 58.9 ) 6983155	( 10.35 , N/A ) ( N/A , 0.00 , N/A )	2149.5	N/A	20.3332 [ 20.0000 ]	101.7% { 97.1% }			
D9_NEtFOSE_EIS	( 639.0 / 58.9 ) 8686678	( 10.50 , N/A ) ( N/A , 0.00 , N/A )	2029.5	N/A	19.7508 [ 20.0000 ]	98.8% { 91.2% }			
13C3_HFPODA_EIS	( 287.0 / 169.0 ) 3487091	( 5.78 , N/A ) ( N/A , 0.01 , N/A )	4321.4	N/A	8.7674 [ 8.0000 ]	109.6% { 95.2% }			

**LOW-CONCENTRATION CALIBRATION VERIFICATION****EPA 1633****Laboratory:** APPL, LLC**SDG:****Client:** AECOM**Project:** Red Hill AFFF Assessment Sampling**Calibration:** 2307007**Laboratory ID:** SC00636-LCV1**Sequence:** SC00636**Standard ID:** 23B0080

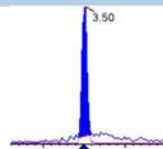
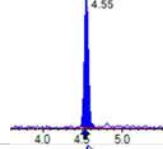
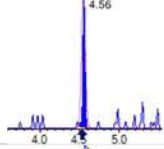
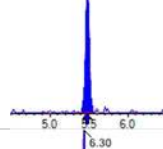
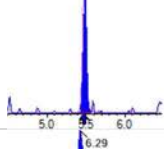
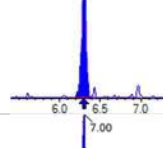
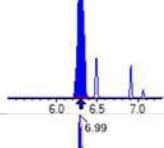
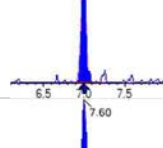
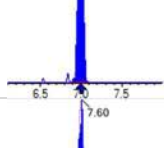
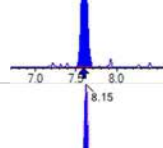
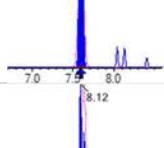
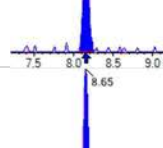
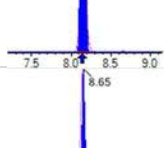
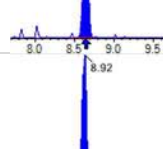
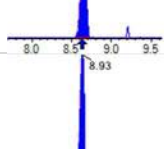
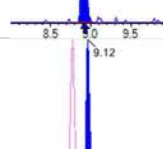
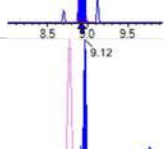
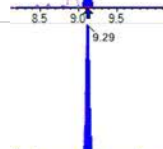
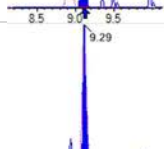
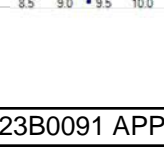
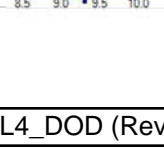
ANALYTE	EXPECTED (ng/mL)	FOUND (ng/mL)	% DRIFT	QC LIMIT
PFBA	0.400	0.353	-11.8	30.00
PFPEA	0.200	0.180	-9.9	30.00
PFHXA	0.100	0.0894	-10.6	30.00
PFHPA	0.100	0.0818	-18.2	30.00
PFOA	0.100	0.110	9.6	30.00
PFNA	0.100	0.0966	-3.4	30.00
PFDA	0.100	0.105	4.6	30.00
PFUnA	0.100	0.0806	-19.4	30.00
PFDOA	0.100	0.106	6.0	30.00
PFTRDA	0.100	0.119	19.3	30.00
PFTEDA	0.100	0.0850	-15.0	30.00
PFBS	0.0885	0.0801	-9.5	30.00
PFPEs	0.0940	0.0927	-1.4	30.00
PFHXS	0.0915	0.0876	-4.2	30.00
PFHPS	0.0955	0.0836	-12.5	30.00
PFOS	0.0930	0.0914	-1.8	30.00
PFNS	0.0960	0.0815	-15.1	30.00
PFDS	0.0965	0.0851	-11.8	30.00
PFDOS	0.0970	0.0846	-12.8	30.00
4:2FTS	0.375	0.376	0.3	30.00
6:2FTS	0.380	0.421	10.7	30.00
8:2FTS	0.384	0.377	-1.8	30.00
PFOSA	0.100	0.0987	-1.3	30.00
NMeFOSA	0.400	0.427	6.8	30.00
NEtFOSA	0.400	0.382	-4.6	30.00
NMeFOSAA	0.100	0.107	7.5	30.00
NEtFOSAA	0.100	0.0870	-13.0	30.00
NMeFOSE	0.400	0.344	-14.1	30.00

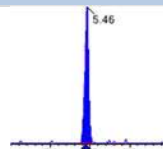
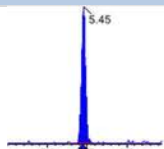
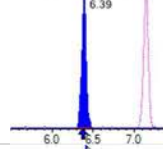
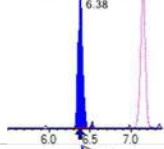
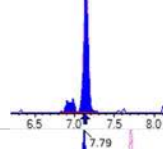
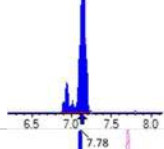
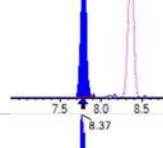
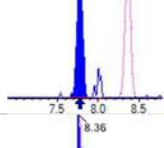
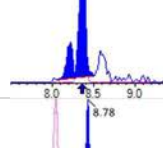
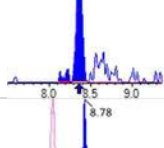
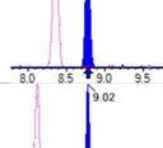
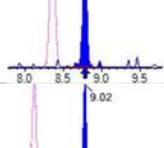
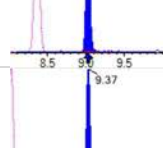
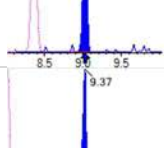
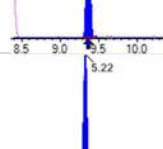
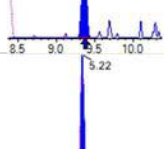
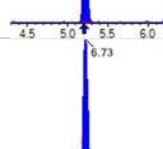
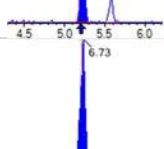
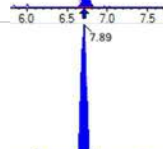
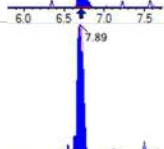

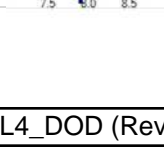


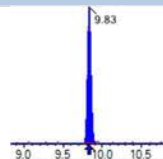
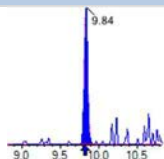
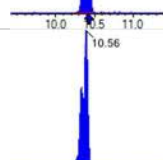
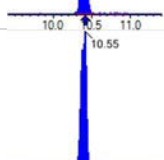
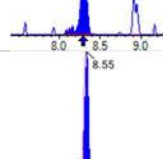
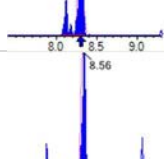
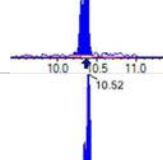
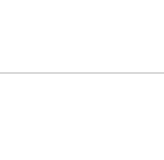
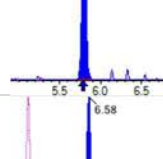
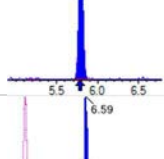
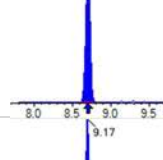
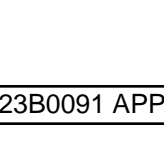
**LOW-CONCENTRATION CALIBRATION VERIFICATION****EPA 1633****Laboratory:** APPL, LLC**SDG:****Client:** AECOM**Project:** Red Hill AFFF Assessment Sampling**Calibration:** 2307007**Laboratory ID:** SC00636-LCV1**Sequence:** SC00636**Standard ID:** 23B0080

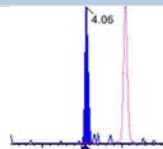
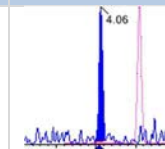
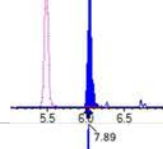
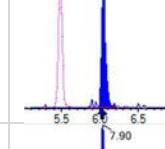
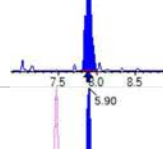
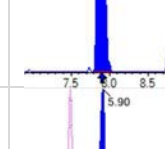
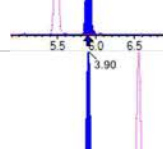
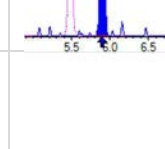
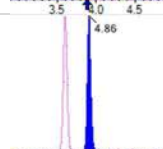
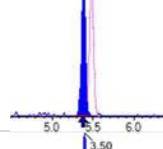
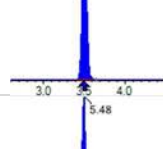
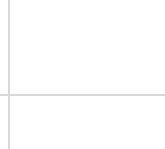
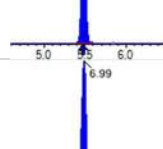
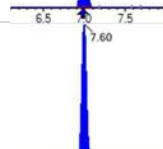
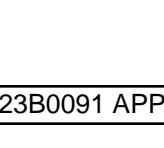
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HFPO-DA	0.200	0.191	-4.4	30.00
ADONA	0.189	0.203	7.3	30.00
PFEESA	0.178	0.171	-3.7	30.00
PFMPA	0.200	0.190	-4.8	30.00
PFMBA	0.200	0.181	-9.6	30.00
NFDHA	0.200	0.166	-16.9	30.00
9CL-PF3ONS	0.187	0.207	10.6	30.00
11CL-PF3OUDS	0.189	0.186	-1.7	30.00
3:3FTCA	0.400	0.592	47.9 *	30.00
5:3FTCA	0.400	0.381	-4.6	30.00
7:3FTCA	0.400	0.345	-13.8	30.00

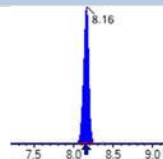
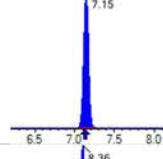
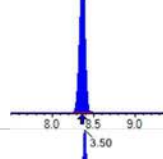
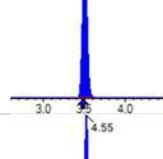
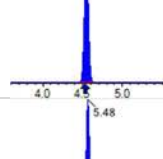
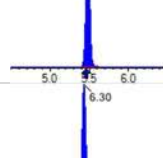
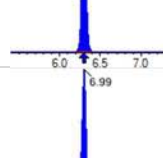
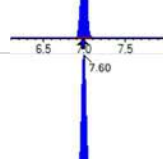
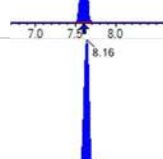
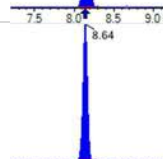
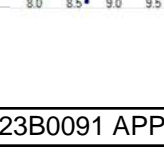
\* Values outside of QC limits

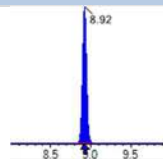
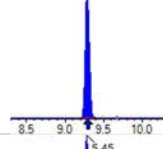
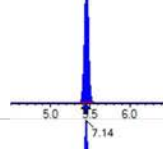
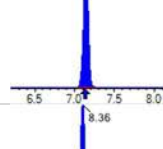
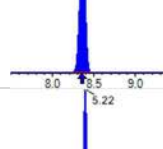
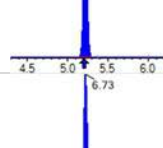
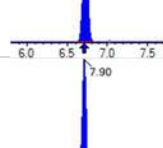
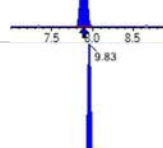
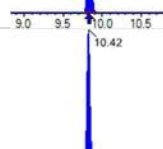
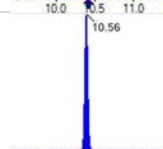
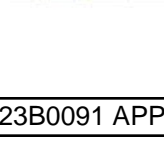
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	( 213.0 / 169.0 ) 89749	( 3.50 , 1.00 ) ( 0.00 , N/A , 0.0 )	83.1	N/A 0.0 0.0	0.3530 [ 0.4000 ]	88.2%			
PFPeA	( 263.0 / 219.0 ) 88501 ( 263.0 / 69.0 ) 1206	( 4.55 , 1.00 ) ( 0.00 , N/A , -0.5 )	315.4 80.9	0.0136 127.3 123.7	0.1802 [ 0.2000 ]	90.1%			
PFHxA	( 313.0 / 269.0 ) 57401 ( 313.0 / 119.0 ) 12709	( 5.48 , 1.00 ) ( 0.00 , N/A , -0.6 )	679.3 31051.0	0.2214 222.3 232.6	0.0894 [ 0.1000 ]	89.4%			IR2,
PFHpA	( 363.0 / 319.0 ) 44905 ( 363.0 / 169.0 ) 9505	( 6.30 , 1.00 ) ( 0.00 , N/A , 0.5 )	1576.7 212939.8	0.2117 66.0 66.2	0.0818 [ 0.1000 ]	81.8%			
PFOA	( 413.0 / 369.0 ) 72107 ( 413.0 / 169.0 ) 18807	( 7.00 , 1.00 ) ( 0.00 , N/A , 0.4 )	794.9 219.8	0.2608 77.9 72.2	0.1096 [ 0.1000 ]	109.6%			
PFNA	( 463.0 / 419.0 ) 56510 ( 463.0 / 169.0 ) 6018	( 7.60 , 1.00 ) ( 0.00 , N/A , 0.0 )	113020.1 485720.1	0.1065 47.6 48.2	0.0966 [ 0.1000 ]	96.6%			IR1,
PFDA	( 513.0 / 469.0 ) 66426 ( 513.0 / 169.0 ) 11070	( 8.15 , 1.00 ) ( 0.00 , N/A , 1.7 )	149.0 150981.0	0.1667 137.0 153.6	0.1046 [ 0.1000 ]	104.6%			IR2,
PFUnA	( 563.0 / 519.0 ) 48076 ( 563.0 / 169.0 ) 10153	( 8.65 , 1.00 ) ( 0.01 , N/A , 0.0 )	286.4 193.9	0.2112 201.4 180.6	0.0806 [ 0.1000 ]	80.6%			IR2,
PFDoA	( 613.0 / 569.0 ) 55813 ( 613.0 / 169.0 ) 5366	( 8.92 , 1.00 ) ( 0.00 , N/A , -0.8 )	333.9 236627.1	0.0961 58.8 57.3	0.1060 [ 0.1000 ]	106.0%			
PFTTrDA	( 663.0 / 619.0 ) 56883 ( 663.0 / 169.0 ) 9826	( 9.12 , 1.02 ) ( N/A , 0.00 , -0.1 )	1323.4 319.7	0.1727 72.3 64.2	0.1193 [ 0.1000 ]	119.3%			
PFTeDA	( 713.0 / 669.0 ) 44396 ( 713.0 / 169.0 ) 14795	( 9.29 , 1.00 ) ( 0.01 , N/A , 0.4 )	254.6 1204.7	0.3333 161.3 161.2	0.0850 [ 0.1000 ]	85.0%			IR2,

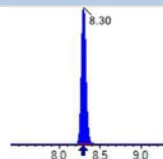
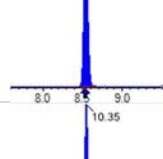
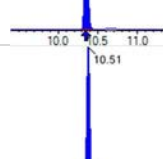
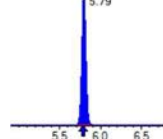

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	( 299.0 / 80.0 ) 99486 ( 299.0 / 99.0 ) 73734	( 5.46 , 1.00 ) ( 0.01 , N/A , 0.3 )	744.6 393.8	0.7411 114.6 119.8	0.0801 [ 0.0885 ]	90.5%			
PFPeS	( 349.0 / 80.0 ) 175663 ( 349.0 / 99.0 ) 66355	( 6.39 , 0.89 ) ( N/A , 0.01 , 0.6 )	38186.9 72060.5	0.3777 105.8 109.7	0.0927 [ 0.0938 ]	98.8%			
PFHxS	( 399.0 / 80.0 ) 132387 ( 399.0 / 99.0 ) 49757	( 7.14 , 1.00 ) ( 0.00 , N/A , 0.2 )	358.2 143.4	0.3758 115.7 115.4	0.0876 [ 0.0911 ]	96.2%			
PFHpS	( 449.0 / 80.0 ) 141021 ( 449.0 / 99.0 ) 35771	( 7.79 , 0.93 ) ( N/A , 0.01 , 0.7 )	934831.8 5992.8	0.2537 93.2 91.1	0.0836 [ 0.0951 ]	87.9%			
PFOS	( 499.0 / 80.0 ) 213307 ( 499.0 / 99.0 ) 42170	( 8.37 , 1.00 ) ( 0.00 , N/A , 0.4 )	106.2 99.8	0.1977 89.3 93.2	0.0914 [ 0.0927 ]	98.5%			MI2 DG 2023-02-15
PFNS	( 549.0 / 80.0 ) 213928 ( 549.0 / 99.0 ) 45264	( 8.78 , 1.05 ) ( N/A , 0.01 , 0.0 )	74758786.6 48165.3	0.2116 96.5 92.0	0.0815 [ 0.0960 ]	84.9%			
PFDS	( 599.0 / 80.0 ) 243008 ( 599.0 / 99.0 ) 53790	( 9.02 , 1.08 ) ( N/A , 0.00 , 0.0 )	524.9 268.0	0.2214 101.5 103.1	0.0851 [ 0.0963 ]	88.4%			
PFDoS	( 699.0 / 80.0 ) 154067 ( 699.0 / 99.0 ) 29623	( 9.37 , 1.12 ) ( N/A , 0.00 , -0.1 )	574.3 152.1	0.1923 92.4 89.6	0.0846 [ 0.0970 ]	87.3%			
4:2FTS	( 327.0 / 307.0 ) 94608 ( 327.0 / 81.0 ) 64089	( 5.22 , 1.00 ) ( 0.00 , N/A , 0.0 )	866.2 304.2	0.6774 100.4 98.8	0.3761 [ 0.3738 ]	100.6%			
6:2FTS	( 427.0 / 407.0 ) 66995 ( 427.0 / 81.0 ) 40797	( 6.73 , 1.00 ) ( 0.00 , N/A , -0.4 )	11419.5 454.9	0.6090 73.6 78.8	0.4205 [ 0.3796 ]	110.8%			
8:2FTS	( 527.0 / 507.0 ) 55251 ( 527.0 / 81.0 ) 40716	( 7.89 , 1.00 ) ( -0.01 , N/A , 0.3 )	2476.6 104.7	0.7369 88.7 95.2	0.3772 [ 0.3833 ]	98.4%			

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 ) 330920 ( 498.0 / 478.0 ) 8093	( 9.83 , 1.00 ) ( 0.00 , N/A , -0.3)	584.2 67.5	0.0245 119.1 115.7	0.0987 [ 0.1000 ]	98.7%			
NMeFOSA	( 512.0 / 219.0 ) 313558 ( 512.0 / 169.0 ) 261812	( 10.42 , 1.00 ) ( 0.00 , N/A , 0.9)	1477.7 1134.6	0.8350 101.9 102.0	0.4271 [ 0.4000 ]	106.8%			
NEIFOSA	( 526.0 / 219.0 ) 330588 ( 526.0 / 169.0 ) 409433	( 10.56 , 1.00 ) ( 0.00 , N/A , 0.8)	1826.7 1107.9	1.2385 97.6 99.8	0.3816 [ 0.4000 ]	95.4%			
NMeFOSAA	( 570.0 / 419.0 ) 26335 ( 570.0 / 483.0 ) 12069	( 8.30 , 1.00 ) ( 0.01 , N/A , 0.1)	1302.0 3948.6	0.4583 96.5 98.9	0.1075 [ 0.1000 ]	107.5%			
NEIFOSAA	( 584.0 / 419.0 ) 19685 ( 584.0 / 526.0 ) 8911	( 8.55 , 1.00 ) ( 0.01 , N/A , -0.4)	483.6 1067.2	0.4527 88.7 77.2	0.0870 [ 0.1000 ]	87.0%			
NMeFOSE	( 616.0 / 59.0 ) 125955	( 10.36 , 1.00 ) ( 0.01 , N/A , 0.0)	270.9	N/A 0.0 0.0	0.3436 [ 0.4000 ]	85.9%			
NEtFOSE	( 630.0 / 59.0 ) 147219	( 10.52 , 1.00 ) ( 0.01 , N/A , 0.0)	330.0	N/A 0.0 0.0	0.3780 [ 0.4000 ]	94.5%			
HFPO-DA	( 285.0 / 169.0 ) 67925 ( 285.0 / 185.0 ) 181166	( 5.79 , 1.00 ) ( 0.00 , N/A , -0.1)	347.1 461.6	2.6672 96.8 96.1	0.1912 [ 0.2000 ]	95.6%			
ADONA	( 377.0 / 85.0 ) 247859 ( 377.0 / 251.0 ) 39547	( 6.58 , 1.14 ) ( N/A , 0.01 , -0.2)	677.2 304.2	0.1596 181.0 170.9	0.2028 [ 0.1885 ]	107.6%			IR2,
9CI-Pf3ONS	( 531.0 / 351.0 ) 632769 ( 533.0 / 353.0 ) 224435	( 8.70 , 1.50 ) ( N/A , 0.01 , 0.2)	891.7 636.6	0.3547 101.9 107.3	0.2069 [ 0.1867 ]	110.8%			
11CI-Pf3OUDS	( 631.0 / 451.0 ) 369670 ( 633.0 / 453.0 ) 116593	( 9.17 , 1.58 ) ( N/A , 0.00 , 0.0)	1442.4 951.3	0.3154 87.2 88.5	0.1857 [ 0.1886 ]	98.4%			

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 ) 6773 ( 241.0 / 117.0 ) 7278	( 4.06 , 0.89 ) ( N/A , 0.00 , -0.3 )	24.7 41.6	1.0746 68.1 65.7	0.5916 [ 0.4000 ]	147.9%			QC,
5:3FTCA	( 341.0 / 236.7 ) 33493 ( 341.0 / 217.0 ) 57323	( 6.05 , 1.10 ) ( N/A , 0.01 , 0.3 )	1093.5 252.7	1.7115 100.4 106.3	0.3814 [ 0.4000 ]	95.4%			
7:3FTCA	( 441.0 / 317.0 ) 55915 ( 441.0 / 337.0 ) 53743	( 7.89 , 1.44 ) ( N/A , 0.01 , -0.6 )	235.1 656.6	0.9612 111.7 118.2	0.3447 [ 0.4000 ]	86.2%			
PFEESA	( 315.0 / 135.0 ) 132209 ( 315.0 / 83.0 ) 40181	( 5.90 , 1.08 ) ( N/A , 0.01 , 0.0 )	978.1 103.6	0.3039 119.0 110.3	0.1714 [ 0.1785 ]	96.0%			
PFMPA	( 229.0 / 85.0 ) 20431	( 3.90 , 0.86 ) ( N/A , 0.01 , 0.0 )	190.9	N/A 0.0 0.0	0.1904 [ 0.2000 ]	95.2%			
PFMBA	( 279.0 / 85.0 ) 81328	( 4.86 , 1.07 ) ( N/A , 0.01 , 0.0 )	269.9	N/A 0.0 0.0	0.1807 [ 0.2000 ]	90.4%			
NFDHA	( 295.0 / 201.0 ) 59262 ( 295.0 / 85.0 ) 71242	( 5.38 , 0.98 ) ( N/A , 0.01 , 0.0 )	276.7 323.1	1.2022 120.1 113.7	0.1663 [ 0.2000 ]	83.1%			
13C3_PFBa_IIS	( 216.0 / 172.0 ) 254197	( 3.50 , N/A ) ( N/A , 0.00 , N/A )	1338.7	N/A	1.1349 [ 1.0000 ]	113.5% { 115.8% }			
13C2_PFHxA_IIS	( 315.0 / 270.0 ) 491981	( 5.48 , N/A ) ( N/A , 0.01 , N/A )	1630.6	N/A	0.9746 [ 1.0000 ]	97.5% { 102.4% }			
13C4_PFOA_IIS	( 417.0 / 372.0 ) 668125	( 6.99 , N/A ) ( N/A , 0.01 , N/A )	1052.6	N/A	1.0542 [ 1.0000 ]	105.4% { 109.5% }			
13C5_PFNA_IIS	( 468.0 / 423.0 ) 592459	( 7.60 , N/A ) ( N/A , 0.01 , N/A )	1880.2	N/A	0.9805 [ 1.0000 ]	98.0% { 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 ) 630481	( 8.16 , N/A ) ( N/A , 0.01 , N/A )	2462.1	N/A	1.0896 [ 1.0000 ]	109.0% { 117.8% }			
18O2_PFHxS_IIS	( 403.0 / 83.9 ) 908827	( 7.15 , N/A ) ( N/A , 0.01 , N/A )	2139.2	N/A	1.0613 [ 1.0000 ]	106.1% { 107.9% }			
13C4_PFOS_IIS	( 503.0 / 79.9 ) 1612834	( 8.36 , N/A ) ( N/A , 0.01 , N/A )	1531.1	N/A	1.0714 [ 1.0000 ]	107.1% { 120.3% }			
13C4_PFBA_EIS	( 217.0 / 172.0 ) 2455524	( 3.50 , N/A ) ( N/A , 0.00 , N/A )	5171.7	N/A	8.1210 [ 8.0000 ]	101.5% { 106.9% }			
13C5_PFPeA_EIS	( 268.0 / 223.0 ) 2388199	( 4.55 , N/A ) ( N/A , 0.01 , N/A )	3718.0	N/A	4.4366 [ 4.0000 ]	110.9% { 103.6% }			
13C5_PFHxA_EIS	( 318.0 / 273.0 ) 1460798	( 5.48 , N/A ) ( N/A , 0.01 , N/A )	2255.9	N/A	2.1677 [ 2.0000 ]	108.4% { 102.3% }			
13C4_PFHpA_EIS	( 367.0 / 322.0 ) 1300427	( 6.30 , N/A ) ( N/A , 0.01 , N/A )	2254.7	N/A	2.2272 [ 2.0000 ]	111.4% { 110.4% }			
13C8_PFOA_EIS	( 421.0 / 376.0 ) 1477825	( 6.99 , N/A ) ( N/A , 0.01 , N/A )	2422.6	N/A	1.9531 [ 2.0000 ]	97.7% { 104.1% }			
13C9_PFNA_EIS	( 472.0 / 427.0 ) 661014	( 7.60 , N/A ) ( N/A , 0.01 , N/A )	1629.0	N/A	1.0530 [ 1.0000 ]	105.3% { 108.6% }			
13C6_PFDA_EIS	( 519.0 / 474.0 ) 721955	( 8.16 , N/A ) ( N/A , 0.01 , N/A )	1135.4	N/A	0.9082 [ 1.0000 ]	90.8% { 96.8% }			
13C7_PFUnA_EIS	( 570.0 / 525.0 ) 718253	( 8.64 , N/A ) ( N/A , 0.01 , N/A )	1468.1	N/A	0.9086 [ 1.0000 ]	90.9% { 107.8% }			

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 ) 598800	( 8.92 , N/A ) ( N/A , 0.00 , N/A )	2065.9	N/A	0.8832 [ 1.0000 ]	88.3% { 91.0% }			
13C2_PFTeDA_EIS	( 715.0 / 670.0 ) 603361	( 9.29 , N/A ) ( N/A , 0.00 , N/A )	1636.3	N/A	0.9712 [ 1.0000 ]	97.1% { 103.0% }			
13C3_PFBs_EIS	( 302.0 / 80.0 ) 3996709	( 5.45 , N/A ) ( N/A , 0.01 , N/A )	3613.4	N/A	2.1012 [ 2.0000 ]	105.1% { 105.4% }			
13C3_PFHxS_EIS	( 402.0 / 80.0 ) 1965140	( 7.14 , N/A ) ( N/A , 0.01 , N/A )	2949.5	N/A	1.9891 [ 2.0000 ]	99.5% { 100.0% }			
13C8_PFOS_EIS	( 507.0 / 80.0 ) 4428708	( 8.36 , N/A ) ( N/A , 0.01 , N/A )	2537.5	N/A	2.0002 [ 2.0000 ]	100.0% { 104.5% }			
13C2_4:2FTS_EIS	( 329.0 / 81.0 ) 368431	( 5.22 , N/A ) ( N/A , 0.01 , N/A )	1579.7	N/A	3.5092 [ 4.0000 ]	87.7% { 106.1% }			
13C2_6:2FTS_EIS	( 429.0 / 81.0 ) 449092	( 6.73 , N/A ) ( N/A , 0.01 , N/A )	1839.0	N/A	3.8967 [ 4.0000 ]	97.4% { 107.9% }			
13C2_8:2FTS_EIS	( 529.0 / 81.0 ) 537880	( 7.90 , N/A ) ( N/A , 0.01 , N/A )	1391.3	N/A	3.3035 [ 4.0000 ]	82.6% { 89.2% }			
13C8_PFOsa_EIS	( 506.0 / 78.0 ) 8386134	( 9.83 , N/A ) ( N/A , 0.00 , N/A )	3400.6	N/A	2.0404 [ 2.0000 ]	102.0% { 100.9% }			
D3_NMeFOsa_EIS	( 515.0 / 169.0 ) 2005925	( 10.42 , N/A ) ( N/A , 0.00 , N/A )	2844.4	N/A	2.0715 [ 2.0000 ]	103.6% { 103.6% }			
D5_NEtFOsa_EIS	( 531.0 / 169.0 ) 1854416	( 10.56 , N/A ) ( N/A , 0.00 , N/A )	2835.5	N/A	2.1401 [ 2.0000 ]	107.0% { 104.0% }			

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 ) 1249118	( 8.30 , N/A ) ( N/A , 0.01 , N/A )	1478.4	N/A	3.4861 [ 4.0000 ]	87.2% { 94.3% }			
D5_EtFOSAA_EIS	( 589.0 / 419.0 ) 1078073	( 8.54 , N/A ) ( N/A , 0.01 , N/A )	54306.2	N/A	3.6744 [ 4.0000 ]	91.9% { 102.2% }			
D7_NMeFOSE_EIS	( 623.0 / 58.9 ) 7334263	( 10.35 , N/A ) ( N/A , 0.00 , N/A )	1741.3	N/A	20.7751 [ 20.0000 ]	103.9% { 98.7% }			
D9_NEtFOSE_EIS	( 639.0 / 58.9 ) 8186054	( 10.51 , N/A ) ( N/A , 0.00 , N/A )	2370.6	N/A	18.1067 [ 20.0000 ]	90.5% { 86.9% }			
13C3_HFPODA_EIS	( 287.0 / 169.0 ) 3594443	( 5.79 , N/A ) ( N/A , 0.01 , N/A )	3327.6	N/A	8.7604 [ 8.0000 ]	109.5% { 100.2% }			



# INITIAL AND CONTINUING CALIBRATION CHECK

## EPA 1633

Laboratory: APPL, LLC

Work Order: 23B0091

Client: AECOM

Project: Red Hill AFFF Assessment Sampling

Instrument ID: Saphira

Calibration: 2307007

Standard ID: 23B0084

Sequence: SC00636

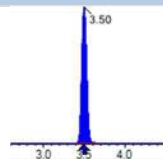
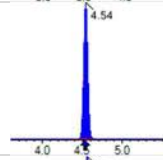
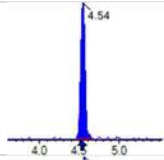
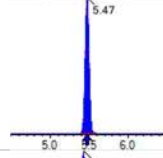
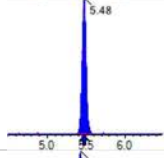
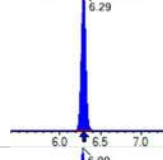
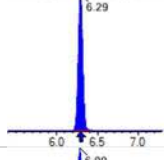
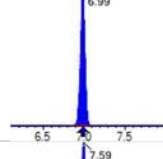
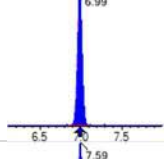
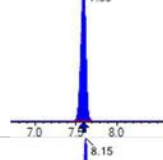
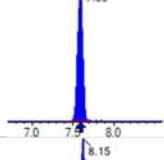
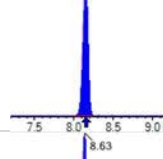
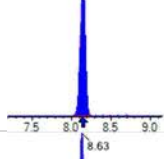
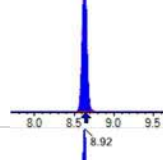
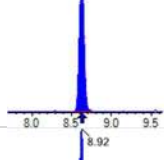
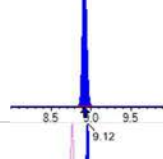
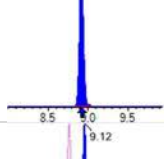
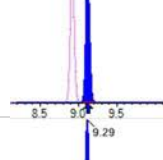
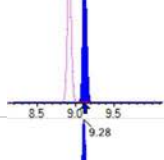
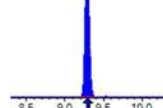
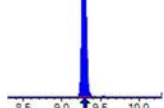
Lab Sample ID	Analyte	True	Found	%R	Units	Control Limit
SC00636-CCV1	PFBA	20.0	19.1	95.5	ng/mL	+/- 30.00%
	PFPEA	10.0	9.85	98.5	ng/mL	+/- 30.00%
	PFHXA	5.00	4.63	92.6	ng/mL	+/- 30.00%
	PFHPA	5.00	4.65	92.9	ng/mL	+/- 30.00%
	PFOA	5.00	4.69	93.9	ng/mL	+/- 30.00%
	PFNA	5.00	5.03	101	ng/mL	+/- 30.00%
	PFDA	5.00	4.89	97.7	ng/mL	+/- 30.00%
	PFUnA	5.00	5.19	104	ng/mL	+/- 30.00%
	PFDOA	5.00	4.57	91.4	ng/mL	+/- 30.00%
	PFTRDA	5.00	4.54	90.7	ng/mL	+/- 30.00%
	PFTEDA	5.00	4.88	97.7	ng/mL	+/- 30.00%
	PFBS	4.42	4.14	93.6	ng/mL	+/- 30.00%
	PFPEs	4.70	4.67	99.3	ng/mL	+/- 30.00%
	PFHXS	4.58	4.27	93.2	ng/mL	+/- 30.00%
	PFHPS	4.78	4.76	99.5	ng/mL	+/- 30.00%
	PFOS	4.65	4.36	93.8	ng/mL	+/- 30.00%
	PFNS	4.80	4.65	96.9	ng/mL	+/- 30.00%
	PFDS	4.82	4.83	100	ng/mL	+/- 30.00%
	PFDOS	4.85	5.11	105	ng/mL	+/- 30.00%
	4:2FTS	18.8	20.2	107	ng/mL	+/- 30.00%
	6:2FTS	19.0	18.6	98.1	ng/mL	+/- 30.00%
	8:2FTS	19.2	18.7	97.5	ng/mL	+/- 30.00%
	PFOSA	5.00	5.13	103	ng/mL	+/- 30.00%
	NMeFOSA	20.0	18.9	94.6	ng/mL	+/- 30.00%
	NEtFOSA	20.0	20.0	100	ng/mL	+/- 30.00%
	NMeFOSAA	5.00	4.81	96.2	ng/mL	+/- 30.00%
	NEtFOSAA	5.00	4.85	97.1	ng/mL	+/- 30.00%
	NMeFOSE	20.0	18.2	91.1	ng/mL	+/- 30.00%
	NEtFOSE	20.0	19.5	97.4	ng/mL	+/- 30.00%
	HFPO-DA	10.0	9.25	92.5	ng/mL	+/- 30.00%

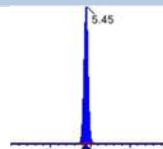
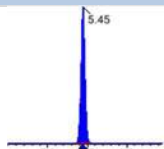
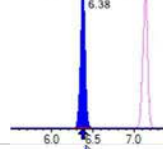
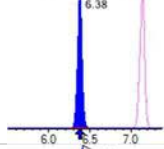
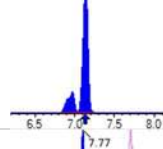
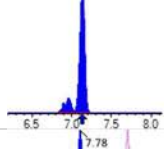
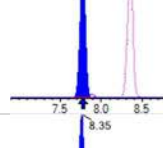
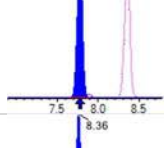
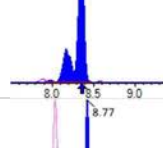
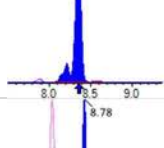
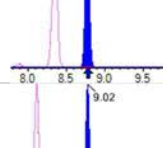
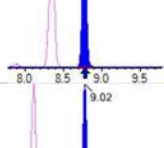
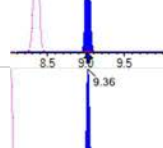
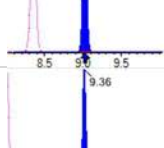
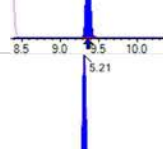
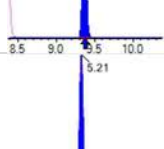
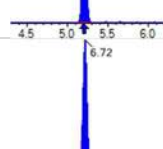
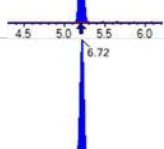
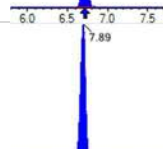
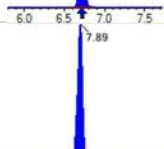

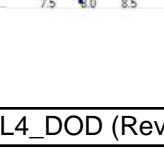
# INITIAL AND CONTINUING CALIBRATION CHECK

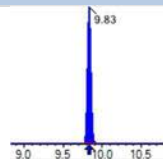
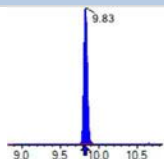
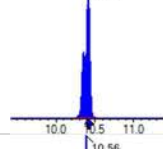
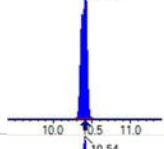
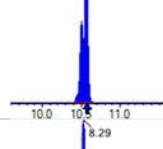
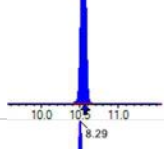
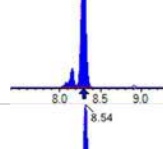
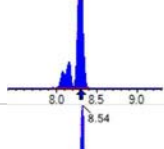
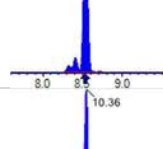
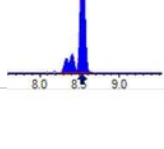
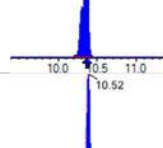
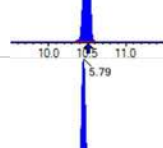
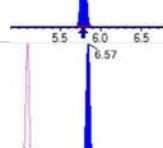
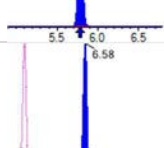
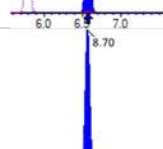
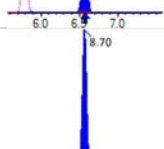
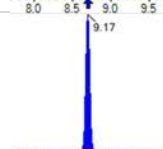
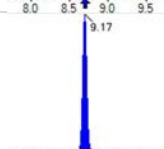
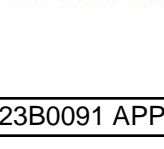
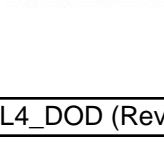
## EPA 1633

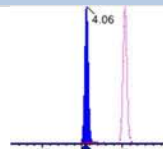
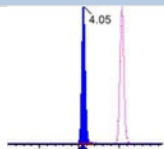
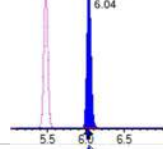
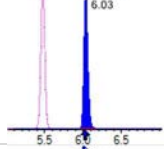
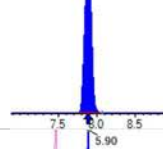
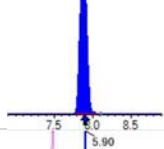
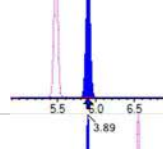
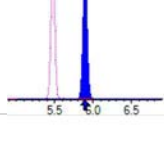
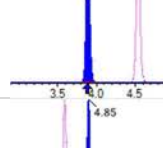
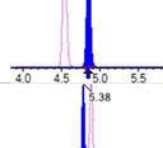
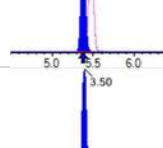
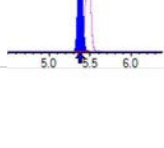
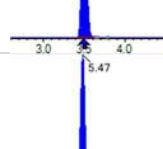
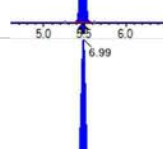
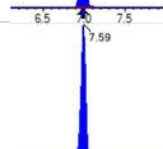

Laboratory:	APPL, LLC	Work Order:	23B0091
Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Instrument ID:	Saphira	Calibration:	2307007
Standard ID:	23B0084	Sequence:	SC00636

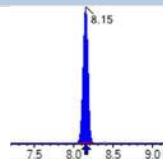
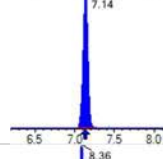
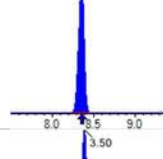
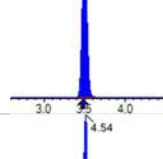
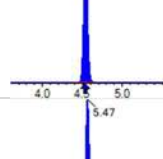
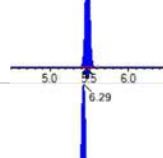
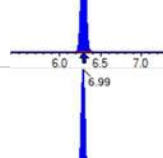
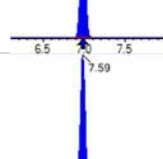
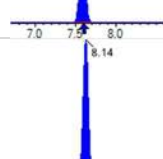
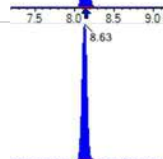
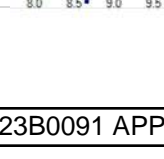
Lab Sample ID	Analyte	True	Found	%R	Units	Control Limit
SC00636-CCV1	ADONA	9.45	9.45	100	ng/mL	+/- 30.00%
	PFEESA	8.90	8.64	97.1	ng/mL	+/- 30.00%
	PFMPA	10.0	9.45	94.5	ng/mL	+/- 30.00%
	PFMBA	10.0	9.43	94.3	ng/mL	+/- 30.00%
	NFDHA	10.0	9.49	94.9	ng/mL	+/- 30.00%
	9CL-PF3ONS	9.35	8.64	92.4	ng/mL	+/- 30.00%
	11CL-PF3OUDS	9.45	9.30	98.4	ng/mL	+/- 30.00%
	3:3FTCA	20.0	18.4	91.8	ng/mL	+/- 30.00%
	5:3FTCA	20.0	19.3	96.6	ng/mL	+/- 30.00%
	7:3FTCA	20.0	19.1	95.3	ng/mL	+/- 30.00%

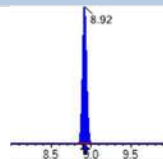
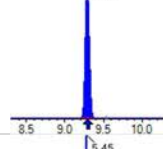
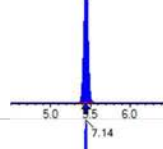
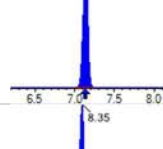
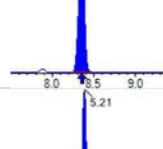
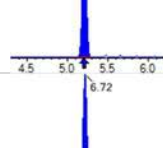
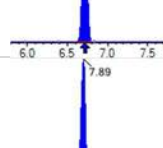
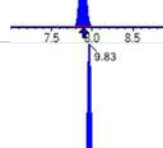
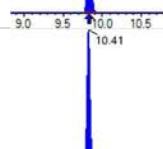
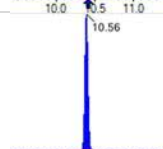
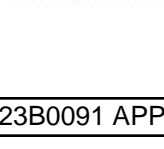
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) ( $\Delta$ RT-I[ $\mu$ min], $\Delta$ RT- CV[ $\mu$ min], $\Delta$ RT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	( 213.0 / 169.0 ) 4545566	( 3.50 , 1.00 ) ( 0.00 , N/A , 0.0 )	211.6	N/A 0.0 0.0	19.1091 [ 20.0000 ]	95.5%			
PFPeA	( 263.0 / 219.0 ) 4668823 ( 263.0 / 69.0 ) 51459	( 4.54 , 1.00 ) ( 0.00 , N/A , -0.1 )	5447.1 388.3	0.0110 102.9 100.0	9.8485 [ 10.0000 ]	98.5%			
PFHxA	( 313.0 / 269.0 ) 2904881 ( 313.0 / 119.0 ) 276451	( 5.47 , 1.00 ) ( 0.00 , N/A , -0.1 )	4635.6 3719.3	0.0952 95.6 100.0	4.6289 [ 5.0000 ]	92.6%			
PFHpA	( 363.0 / 319.0 ) 2311360 ( 363.0 / 169.0 ) 739023	( 6.29 , 1.00 ) ( 0.00 , N/A , 0.1 )	31072.2 10966672.0	0.3197 99.7 100.0	4.6461 [ 5.0000 ]	92.9%			
PFOA	( 413.0 / 369.0 ) 2967294 ( 413.0 / 169.0 ) 1072142	( 6.99 , 1.00 ) ( 0.00 , N/A , 0.0 )	9473.7 6824521.4	0.3613 108.0 100.0	4.6932 [ 5.0000 ]	93.9%			
PFNA	( 463.0 / 419.0 ) 2707898 ( 463.0 / 169.0 ) 598588	( 7.59 , 1.00 ) ( 0.00 , N/A , 0.3 )	11168.3 3567.3	0.2211 98.8 100.0	5.0283 [ 5.0000 ]	100.6%			
PFDA	( 513.0 / 469.0 ) 3205880 ( 513.0 / 169.0 ) 347843	( 8.15 , 1.00 ) ( 0.00 , N/A , 0.0 )	1950.8 1452.8	0.1085 89.2 100.0	4.8871 [ 5.0000 ]	97.7%			
PFUnA	( 563.0 / 519.0 ) 2870912 ( 563.0 / 169.0 ) 335661	( 8.63 , 1.00 ) ( 0.00 , N/A , 0.2 )	1725.9 1273.3	0.1169 111.5 100.0	5.1868 [ 5.0000 ]	103.7%			
PFDaA	( 613.0 / 569.0 ) 2644619 ( 613.0 / 169.0 ) 443798	( 8.92 , 1.00 ) ( 0.00 , N/A , 0.0 )	2500.5 1702.2	0.1678 102.6 100.0	4.5690 [ 5.0000 ]	91.4%			
PFTTrDA	( 663.0 / 619.0 ) 2377082 ( 663.0 / 169.0 ) 639905	( 9.12 , 1.02 ) ( N/A , 0.00 , 0.0 )	2343.7 1278.3	0.2692 112.6 100.0	4.5358 [ 5.0000 ]	90.7%			
PFTeDA	( 713.0 / 669.0 ) 2475442 ( 713.0 / 169.0 ) 511779	( 9.29 , 1.00 ) ( 0.00 , N/A , 0.2 )	1836.5 789.1	0.2067 100.0 100.0	4.8834 [ 5.0000 ]	97.7%			

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 ) 4876321 ( 299.0 / 99.0 ) 3016826	( 5.45 , 1.00 ) ( 0.00 , N/A , 0.0 )	3735.9 3110.7	0.6187 95.7 100.0	4.1381 [ 4.4237 ]	93.5%			
PFPeS	( 349.0 / 80.0 ) 8844504 ( 349.0 / 99.0 ) 3046766	( 6.38 , 0.89 ) ( N/A , 0.00 , 0.0 )	17407.3 180613.2	0.3445 96.5 100.0	4.6667 [ 4.6919 ]	99.5%			
PFHxS	( 399.0 / 80.0 ) 6453411 ( 399.0 / 99.0 ) 2102126	( 7.14 , 1.00 ) ( 0.00 , N/A , 0.1 )	6548.6 9406050.5	0.3257 100.3 100.0	4.2705 [ 4.5549 ]	93.8%			
PFHpS	( 449.0 / 80.0 ) 7680384 ( 449.0 / 99.0 ) 2137538	( 7.77 , 0.93 ) ( N/A , 0.00 , -0.1 )	40655.6 5168.2	0.2783 102.2 100.0	4.7569 [ 4.7570 ]	100.0%			
PFOS	( 499.0 / 80.0 ) 9743530 ( 499.0 / 99.0 ) 2065852	( 8.35 , 1.00 ) ( 0.00 , N/A , 0.0 )	1608.2 1199.8	0.2120 95.8 100.0	4.3606 [ 4.6375 ]	94.0%			
PFNS	( 549.0 / 80.0 ) 11681260 ( 549.0 / 99.0 ) 2686350	( 8.77 , 1.05 ) ( N/A , 0.00 , -0.1 )	10102.5 2802.4	0.2300 104.9 100.0	4.6517 [ 4.7994 ]	96.9%			
PFDS	( 599.0 / 80.0 ) 13182611 ( 599.0 / 99.0 ) 2830382	( 9.02 , 1.08 ) ( N/A , 0.00 , -0.1 )	3451.3 2778.8	0.2147 98.5 100.0	4.8259 [ 4.8155 ]	100.2%			
PFDoS	( 699.0 / 80.0 ) 8900344 ( 699.0 / 99.0 ) 1909885	( 9.36 , 1.12 ) ( N/A , 0.00 , 0.0 )	2811.7 2282.5	0.2146 103.1 100.0	5.1086 [ 4.8478 ]	105.4%			
4:2FTS	( 327.0 / 307.0 ) 4788393 ( 327.0 / 81.0 ) 3282721	( 5.21 , 1.00 ) ( 0.00 , N/A , -0.1 )	3962.1 3508.9	0.6856 101.6 100.0	20.1938 [ 18.6906 ]	108.0%			
6:2FTS	( 427.0 / 407.0 ) 2751008 ( 427.0 / 81.0 ) 2125102	( 6.72 , 1.00 ) ( 0.00 , N/A , 0.0 )	3269.7 4030.0	0.7725 93.3 100.0	18.6396 [ 18.9808 ]	98.2%			
8:2FTS	( 527.0 / 507.0 ) 3074408 ( 527.0 / 81.0 ) 2378995	( 7.89 , 1.00 ) ( 0.00 , N/A , -0.1 )	2078.7 2768.2	0.7738 93.1 100.0	18.7159 [ 19.1658 ]	97.7%			

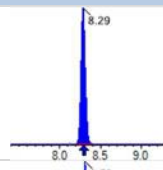
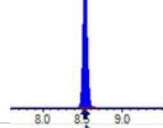
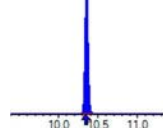
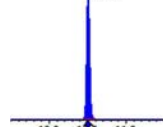
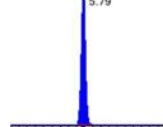
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) ( $\Delta$ RT-[min], $\Delta$ RT- CV[min], $\Delta$ RT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	( 498.0 / 78.0 ) 17051404 ( 498.0 / 478.0 ) 360353	( 9.83 , 1.00 ) ( 0.00 , N/A , 0.1 )	3556.4 1201.3	0.0211 102.9 100.0	5.1326 [ 5.0000 ]	102.7%			
NMeFOSA	( 512.0 / 219.0 ) 15220245 ( 512.0 / 169.0 ) 12464593	( 10.42 , 1.00 ) ( 0.00 , N/A , 1.0 )	3250.2 5589.3	0.8189 100.0 100.0	18.9133 [ 20.0000 ]	94.6%			
NEIFOSA	( 526.0 / 219.0 ) 16668720 ( 526.0 / 169.0 ) 20688606	( 10.56 , 1.00 ) ( 0.00 , N/A , 0.8 )	16050.3 14684.8	1.2412 97.8 100.0	20.0088 [ 20.0000 ]	100.0%			
NMeFOSAA	( 570.0 / 419.0 ) 1249662 ( 570.0 / 483.0 ) 578925	( 8.29 , 1.00 ) ( 0.01 , N/A , -0.1 )	2351.7 250.5	0.4633 97.6 100.0	4.8102 [ 5.0000 ]	96.2%			
NEIFOSAA	( 584.0 / 419.0 ) 1075322 ( 584.0 / 526.0 ) 630191	( 8.54 , 1.00 ) ( 0.01 , N/A , 0.1 )	2559.9 2876.4	0.5860 114.9 100.0	4.8535 [ 5.0000 ]	97.1%			
NMeFOSE	( 616.0 / 59.0 ) 6767825	( 10.36 , 1.00 ) ( 0.01 , N/A , 0.0 )	3231.2	N/A 0.0 0.0	18.2283 [ 20.0000 ]	91.1%			
NEIFOSE	( 630.0 / 59.0 ) 8733335	( 10.52 , 1.00 ) ( 0.01 , N/A , 0.0 )	1739.5	N/A 0.0 0.0	19.4798 [ 20.0000 ]	97.4%			
HFPO-DA	( 285.0 / 169.0 ) 3280605 ( 285.0 / 185.0 ) 9102315	( 5.79 , 1.00 ) ( 0.00 , N/A , 0.0 )	2934.0 4685.7	2.7746 100.7 100.0	9.2539 [ 10.0000 ]	92.5%			
ADONA	( 377.0 / 85.0 ) 11519185 ( 377.0 / 251.0 ) 1075387	( 6.57 , 1.14 ) ( N/A , 0.00 , -0.1 )	4144.5 1985.4	0.0934 105.9 100.0	9.4484 [ 9.4270 ]	100.2%			
9CI-PF3ONS	( 531.0 / 351.0 ) 31391246 ( 533.0 / 353.0 ) 10372156	( 8.70 , 1.50 ) ( N/A , 0.00 , 0.0 )	2709.6 3534.4	0.3304 94.9 100.0	8.6438 [ 9.3325 ]	92.6%			
11CI-PF3OUDS	( 631.0 / 451.0 ) 18470106 ( 633.0 / 453.0 ) 6581198	( 9.17 , 1.59 ) ( N/A , 0.00 , 0.0 )	3005.8 2913.0	0.3563 98.5 100.0	9.2996 [ 9.4321 ]	98.6%			

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 ) 214959 ( 241.0 / 117.0 ) 351740	( 4.06 , 0.89 ) ( N/A , 0.00 , 0.1 )	1494.0 1284.7	1.6363 103.7 100.0	18.3693 [ 20.0000 ]	91.8%			
5:3FTCA	( 341.0 / 236.7 ) 1658127 ( 341.0 / 217.0 ) 2669494	( 6.04 , 1.10 ) ( N/A , 0.00 , 0.1 )	1720.1 2500.5	1.6099 94.4 100.0	19.3216 [ 20.0000 ]	96.6%			
7:3FTCA	( 441.0 / 317.0 ) 3022587 ( 441.0 / 337.0 ) 2457384	( 7.89 , 1.44 ) ( N/A , 0.00 , 0.1 )	1417.2 1129.2	0.8130 94.5 100.0	19.0660 [ 20.0000 ]	95.3%			
PFEESA	( 315.0 / 135.0 ) 6514081 ( 315.0 / 83.0 ) 1794331	( 5.90 , 1.08 ) ( N/A , 0.00 , 0.0 )	4009.4 2055.6	0.2755 107.9 100.0	8.6409 [ 8.9246 ]	96.8%			
PFMPA	( 229.0 / 85.0 ) 978691	( 3.89 , 0.86 ) ( N/A , 0.00 , 0.0 )	4541.3	N/A 0.0 0.0	9.4515 [ 10.0000 ]	94.5%			
PFMBA	( 279.0 / 85.0 ) 4096269	( 4.85 , 1.07 ) ( N/A , 0.00 , 0.0 )	3622.8	N/A 0.0 0.0	9.4331 [ 10.0000 ]	94.3%			
NFDHA	( 295.0 / 201.0 ) 3306050 ( 295.0 / 85.0 ) 3495480	( 5.38 , 0.98 ) ( N/A , 0.00 , 0.0 )	2507.0 3189.8	1.0573 105.6 100.0	9.4913 [ 10.0000 ]	94.9%			
13C3_PFBA_IIS	( 216.0 / 172.0 ) 219434	( 3.50 , N/A ) ( N/A , 0.00 , N/A )	1008.3	N/A	0.9797 [ 1.0000 ]	98.0% { 100.0% }			
13C2_PFHxA_IIS	( 315.0 / 270.0 ) 480590	( 5.47 , N/A ) ( N/A , 0.00 , N/A )	2007.3	N/A	0.9520 [ 1.0000 ]	95.2% { 100.0% }			
13C4_PFOA_IIS	( 417.0 / 372.0 ) 610417	( 6.99 , N/A ) ( N/A , 0.00 , N/A )	8085.2	N/A	0.9631 [ 1.0000 ]	96.3% { 100.0% }			
13C5_PFNA_IIS	( 468.0 / 423.0 ) 563943	( 7.59 , N/A ) ( N/A , 0.00 , N/A )	1793.2	N/A	0.9333 [ 1.0000 ]	93.3% { 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_IIS	( 515.0 / 470.1 ) 535206	( 8.15 , N/A ) ( N/A , 0.00 , N/A )	2299.0	N/A	0.9249 [ 1.0000 ]	92.5% { 100.0% }			
18O2_PFHxS_IIS	( 403.0 / 83.9 ) 842031	( 7.14 , N/A ) ( N/A , 0.00 , N/A )	2141.7	N/A	0.9833 [ 1.0000 ]	98.3% { 100.0% }			
13C4_PFOS_IIS	( 503.0 / 79.9 ) 1340994	( 8.36 , N/A ) ( N/A , 0.00 , N/A )	1367.3	N/A	0.8908 [ 1.0000 ]	89.1% { 100.0% }			
13C4_PFBA_EIS	( 217.0 / 172.0 ) 2297340	( 3.50 , N/A ) ( N/A , 0.00 , N/A )	4721.2	N/A	8.8015 [ 8.0000 ]	110.0% { 100.0% }			
13C5_PFPeA_EIS	( 268.0 / 223.0 ) 2304813	( 4.54 , N/A ) ( N/A , 0.00 , N/A )	2970.6	N/A	4.3832 [ 4.0000 ]	109.6% { 100.0% }			
13C5_PFHxA_EIS	( 318.0 / 273.0 ) 1427673	( 5.47 , N/A ) ( N/A , 0.00 , N/A )	3024.1	N/A	2.1688 [ 2.0000 ]	108.4% { 100.0% }			
13C4_PFHpA_EIS	( 367.0 / 322.0 ) 1178269	( 6.29 , N/A ) ( N/A , 0.00 , N/A )	2645.2	N/A	2.0658 [ 2.0000 ]	103.3% { 100.0% }			
13C8_PFOA_EIS	( 421.0 / 376.0 ) 1420182	( 6.99 , N/A ) ( N/A , 0.00 , N/A )	1708.2	N/A	2.0543 [ 2.0000 ]	102.7% { 100.0% }			
13C9_PFNA_EIS	( 472.0 / 427.0 ) 608665	( 7.59 , N/A ) ( N/A , 0.00 , N/A )	2409.6	N/A	1.0187 [ 1.0000 ]	101.9% { 100.0% }			
13C6_PFDA_EIS	( 519.0 / 474.0 ) 745748	( 8.14 , N/A ) ( N/A , 0.00 , N/A )	1416.3	N/A	1.1051 [ 1.0000 ]	110.5% { 100.0% }			
13C7_PFUnA_EIS	( 570.0 / 525.0 ) 666201	( 8.63 , N/A ) ( N/A , 0.00 , N/A )	1853.1	N/A	0.9928 [ 1.0000 ]	99.3% { 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 ) 658219	( 8.92 , N/A ) ( N/A , 0.00 , N/A )	2474.1	N/A	1.1437 [ 1.0000 ]	114.4% { 100.0% }			
13C2_PFTeDA_EIS	( 715.0 / 670.0 ) 585783	( 9.29 , N/A ) ( N/A , 0.00 , N/A )	1182.1	N/A	1.1108 [ 1.0000 ]	111.1% { 100.0% }			
13C3_PFBs_EIS	( 302.0 / 80.0 ) 3792392	( 5.45 , N/A ) ( N/A , 0.00 , N/A )	4491.4	N/A	2.1520 [ 2.0000 ]	107.6% { 100.0% }			
13C3_PFHxS_EIS	( 402.0 / 80.0 ) 1965463	( 7.14 , N/A ) ( N/A , 0.00 , N/A )	2731.7	N/A	2.1472 [ 2.0000 ]	107.4% { 100.0% }			
13C8_PFOS_EIS	( 507.0 / 80.0 ) 4238028	( 8.35 , N/A ) ( N/A , 0.00 , N/A )	1778.2	N/A	2.3020 [ 2.0000 ]	115.1% { 100.0% }			
13C2_4:2FTS_EIS	( 329.0 / 81.0 ) 347280	( 5.21 , N/A ) ( N/A , 0.00 , N/A )	993.8	N/A	3.5701 [ 4.0000 ]	89.3% { 100.0% }			
13C2_6:2FTS_EIS	( 429.0 / 81.0 ) 416066	( 6.72 , N/A ) ( N/A , 0.00 , N/A )	1373.8	N/A	3.8965 [ 4.0000 ]	97.4% { 100.0% }			
13C2_8:2FTS_EIS	( 529.0 / 81.0 ) 603229	( 7.89 , N/A ) ( N/A , 0.00 , N/A )	2494.6	N/A	3.9987 [ 4.0000 ]	100.0% { 100.0% }			
13C8_PFOsa_EIS	( 506.0 / 78.0 ) 8307470	( 9.83 , N/A ) ( N/A , 0.00 , N/A )	4340.1	N/A	2.4310 [ 2.0000 ]	121.5% { 100.0% }			
D3_NMeFOsa_EIS	( 515.0 / 169.0 ) 1935325	( 10.41 , N/A ) ( N/A , 0.00 , N/A )	3160.3	N/A	2.4037 [ 2.0000 ]	120.2% { 100.0% }			
D5_NEtFOsa_EIS	( 531.0 / 169.0 ) 1783429	( 10.56 , N/A ) ( N/A , 0.00 , N/A )	3099.3	N/A	2.4754 [ 2.0000 ]	123.8% { 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 ) 1324195	( 8.29 , N/A ) ( N/A , 0.00 , N/A )	1316.0	N/A	4.4448 [ 4.0000 ]	111.1% { 100.0% }			
D5_EtFOSAA_EIS	( 589.0 / 419.0 ) 1055150	( 8.53 , N/A ) ( N/A , 0.00 , N/A )	68797.1	N/A	4.3253 [ 4.0000 ]	108.1% { 100.0% }			
D7_NMeFOSE_EIS	( 623.0 / 58.9 ) 7427567	( 10.35 , N/A ) ( N/A , 0.00 , N/A )	2212.4	N/A	25.3045 [ 20.0000 ]	126.5% { 100.0% }			
D9_NEtFOSE_EIS	( 639.0 / 58.9 ) 9422296	( 10.51 , N/A ) ( N/A , 0.00 , N/A )	2106.7	N/A	25.0659 [ 20.0000 ]	125.3% { 100.0% }			
13C3_HFPODA_EIS	( 287.0 / 169.0 ) 3586350	( 5.79 , N/A ) ( N/A , 0.00 , N/A )	3434.0	N/A	8.9478 [ 8.0000 ]	111.8% { 100.0% }			

# INITIAL AND CONTINUING CALIBRATION CHECK

## EPA 1633

Laboratory: APPL, LLC

Work Order: 23B0091

Client: AECOM

Project: Red Hill AFFF Assessment Sampling

Instrument ID: Saphira

Calibration: 2307007

Standard ID: 23B0084

Sequence: SC00636

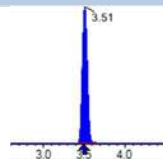
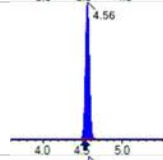
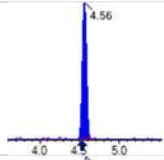
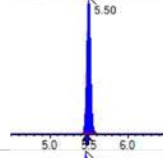
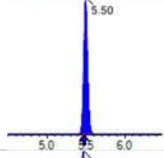
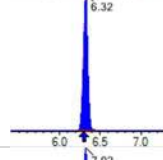
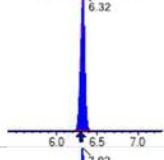
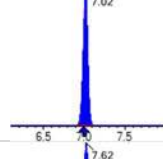
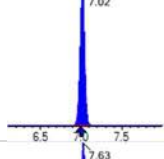
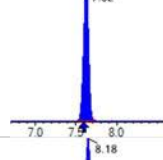
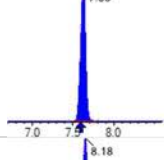
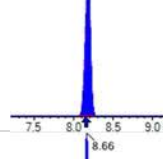
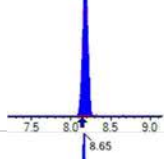
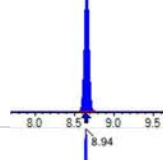
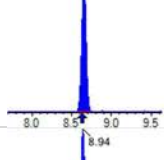
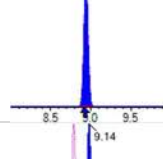
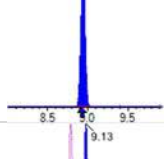
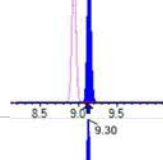
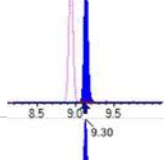
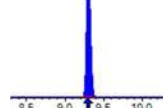
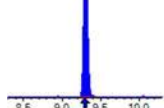
Lab Sample ID	Analyte	True	Found	%R	Units	Control Limit
SC00636-CCV2	PFBA	20.0	19.3	96.4	ng/mL	+/- 30.00%
	PFPEA	10.0	9.48	94.8	ng/mL	+/- 30.00%
	PFHXA	5.00	4.58	91.5	ng/mL	+/- 30.00%
	PFHPA	5.00	4.55	90.9	ng/mL	+/- 30.00%
	PFOA	5.00	4.48	89.6	ng/mL	+/- 30.00%
	PFNA	5.00	4.77	95.3	ng/mL	+/- 30.00%
	PFDA	5.00	4.56	91.1	ng/mL	+/- 30.00%
	PFUnA	5.00	4.34	86.9	ng/mL	+/- 30.00%
	PFDOA	5.00	4.43	88.7	ng/mL	+/- 30.00%
	PFTRDA	5.00	5.13	103	ng/mL	+/- 30.00%
	PFTEDA	5.00	4.98	99.6	ng/mL	+/- 30.00%
	PFBS	4.42	4.20	95.0	ng/mL	+/- 30.00%
	PFPEs	4.70	4.63	98.5	ng/mL	+/- 30.00%
	PFHXS	4.58	4.18	91.3	ng/mL	+/- 30.00%
	PFHPS	4.78	4.69	98.1	ng/mL	+/- 30.00%
	PFOS	4.65	4.29	92.2	ng/mL	+/- 30.00%
	PFNS	4.80	4.57	95.2	ng/mL	+/- 30.00%
	PFDS	4.82	4.94	103	ng/mL	+/- 30.00%
	PFDOS	4.85	5.61	116	ng/mL	+/- 30.00%
	4:2FTS	18.8	18.5	98.5	ng/mL	+/- 30.00%
	6:2FTS	19.0	18.1	95.0	ng/mL	+/- 30.00%
	8:2FTS	19.2	18.8	98.1	ng/mL	+/- 30.00%
	PFOSA	5.00	4.96	99.1	ng/mL	+/- 30.00%
	NMeFOSA	20.0	19.5	97.6	ng/mL	+/- 30.00%
	NEtFOSA	20.0	20.1	101	ng/mL	+/- 30.00%
	NMeFOSAA	5.00	4.91	98.1	ng/mL	+/- 30.00%
	NEtFOSAA	5.00	4.64	92.9	ng/mL	+/- 30.00%
	NMeFOSE	20.0	18.5	92.5	ng/mL	+/- 30.00%
	NEtFOSE	20.0	19.7	98.4	ng/mL	+/- 30.00%
	HFPO-DA	10.0	9.72	97.2	ng/mL	+/- 30.00%

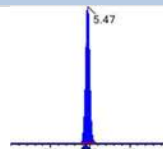
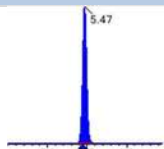
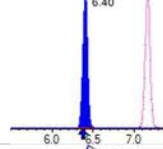
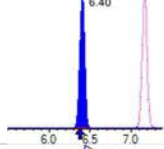
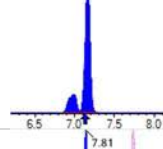
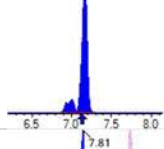
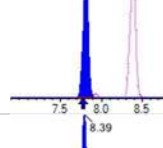
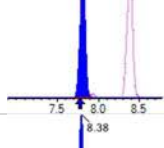
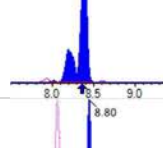
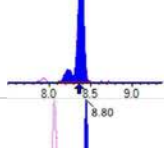
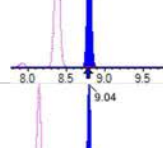
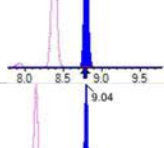
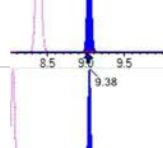
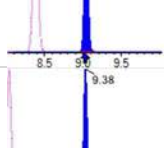
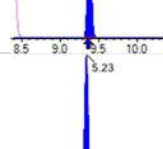
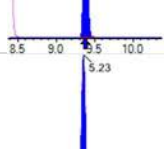
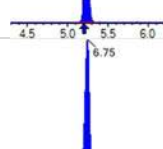
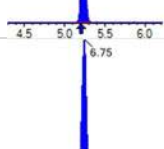
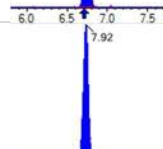
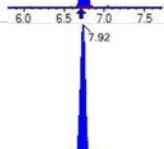

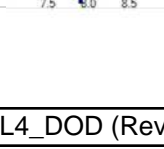
# INITIAL AND CONTINUING CALIBRATION CHECK

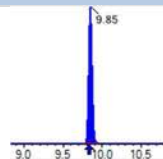
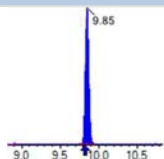
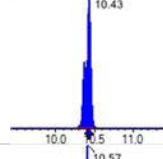
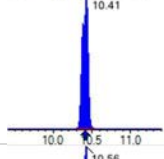
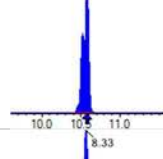
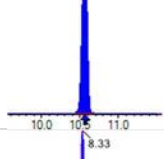
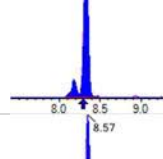
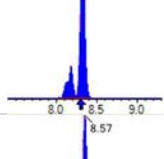
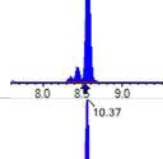
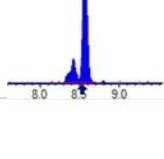
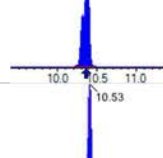
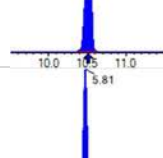
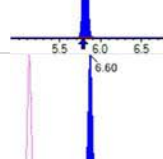
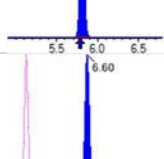
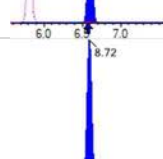
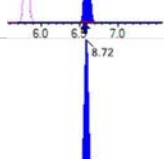
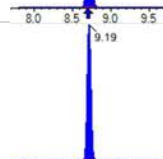
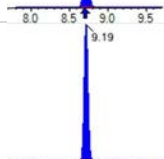
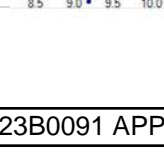
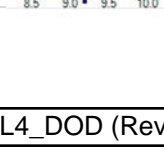
## EPA 1633

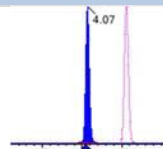
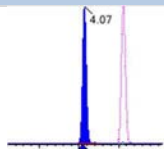
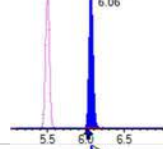
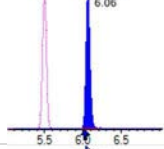
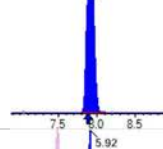
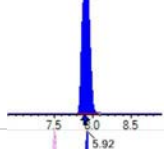
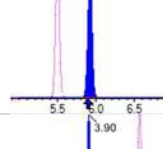
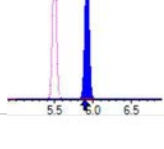
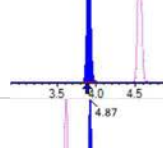
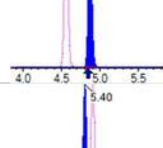
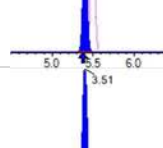
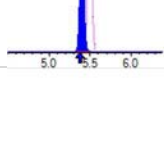
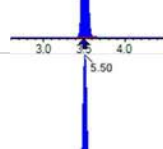
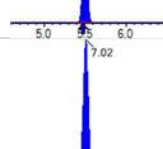
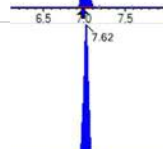

Laboratory:	APPL, LLC	Work Order:	23B0091
Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Instrument ID:	Saphira	Calibration:	2307007
Standard ID:	23B0084	Sequence:	SC00636

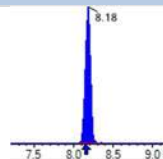
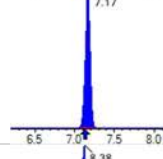
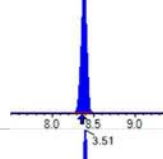
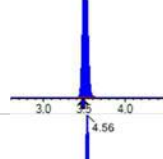
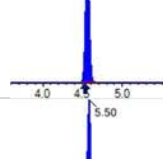
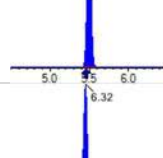
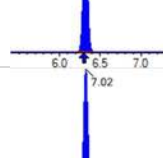
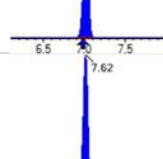
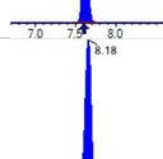
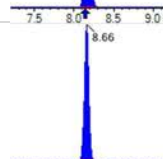
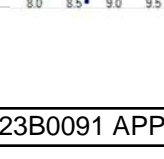
Lab Sample ID	Analyte	True	Found	%R	Units	Control Limit
SC00636-CCV2	ADONA	9.45	9.99	106	ng/mL	+/- 30.00%
	PFEESA	8.90	8.50	95.5	ng/mL	+/- 30.00%
	PFMPA	10.0	9.17	91.7	ng/mL	+/- 30.00%
	PFMBA	10.0	9.44	94.4	ng/mL	+/- 30.00%
	NFDHA	10.0	9.17	91.7	ng/mL	+/- 30.00%
	9CL-PF3ONS	9.35	8.82	94.3	ng/mL	+/- 30.00%
	11CL-PF3OUDS	9.45	9.94	105	ng/mL	+/- 30.00%
	3:3FTCA	20.0	18.2	91.0	ng/mL	+/- 30.00%
	5:3FTCA	20.0	16.5	82.7	ng/mL	+/- 30.00%
	7:3FTCA	20.0	16.7	83.5	ng/mL	+/- 30.00%

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 ) 4582364	( 3.51 , 1.00 ) ( 0.00 , N/A , 0.0 )	227.0	N/A 0.0 0.0	19.2761 [ 20.0000 ]	96.4%			
PFPeA	( 263.0 / 219.0 ) 4672093 ( 263.0 / 69.0 ) 60381	( 4.56 , 1.00 ) ( 0.00 , N/A , 0.0 )	5739.4 436.2	0.0129 120.7 117.3	9.4768 [ 10.0000 ]	94.8%			
PFHxA	( 313.0 / 269.0 ) 3060630 ( 313.0 / 119.0 ) 293387	( 5.50 , 1.00 ) ( 0.00 , N/A , 0.1 )	7180.7 10387.3	0.0959 96.2 100.7	4.5769 [ 5.0000 ]	91.5%			
PFHpA	( 363.0 / 319.0 ) 2553726 ( 363.0 / 169.0 ) 781101	( 6.32 , 1.00 ) ( 0.00 , N/A , -0.2 )	31214.1 5071.0	0.3059 95.4 95.7	4.5473 [ 5.0000 ]	90.9%			
PFOA	( 413.0 / 369.0 ) 3035697 ( 413.0 / 169.0 ) 1024057	( 7.02 , 1.00 ) ( 0.00 , N/A , 0.1 )	5228.3 80814.1	0.3373 100.8 93.4	4.4775 [ 5.0000 ]	89.6%			
PFNA	( 463.0 / 419.0 ) 2648335 ( 463.0 / 169.0 ) 588014	( 7.62 , 1.00 ) ( 0.00 , N/A , -0.1 )	12447.7 5128.0	0.2220 99.2 100.4	4.7675 [ 5.0000 ]	95.3%			
PFDA	( 513.0 / 469.0 ) 3136505 ( 513.0 / 169.0 ) 385057	( 8.18 , 1.00 ) ( 0.00 , N/A , 0.1 )	2494.4 1897.2	0.1228 101.0 113.1	4.5553 [ 5.0000 ]	91.1%			
PFUnA	( 563.0 / 519.0 ) 2869610 ( 563.0 / 169.0 ) 359723	( 8.66 , 1.00 ) ( 0.00 , N/A , 0.1 )	3973.8 1543.7	0.1254 119.6 107.2	4.3430 [ 5.0000 ]	86.9%			
PFDoA	( 613.0 / 569.0 ) 2475060 ( 613.0 / 169.0 ) 423069	( 8.94 , 1.00 ) ( 0.00 , N/A , 0.0 )	1944.0 1789.7	0.1709 104.5 101.9	4.4330 [ 5.0000 ]	88.7%			
PFTTrDA	( 663.0 / 619.0 ) 2591895 ( 663.0 / 169.0 ) 605706	( 9.14 , 1.02 ) ( N/A , 0.02 , 0.1 )	2357.2 1374.5	0.2337 97.8 86.8	5.1272 [ 5.0000 ]	102.5%			
PFTeDA	( 713.0 / 669.0 ) 2614656 ( 713.0 / 169.0 ) 516600	( 9.30 , 1.00 ) ( 0.00 , N/A , -0.1 )	2200.9 1332.8	0.1976 95.6 95.6	4.9813 [ 5.0000 ]	99.6%			

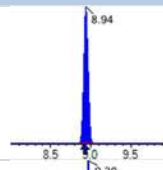
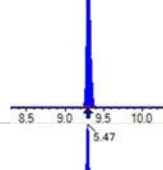
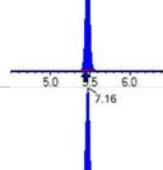
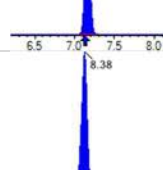
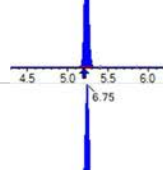
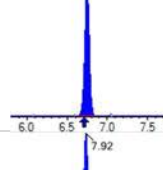
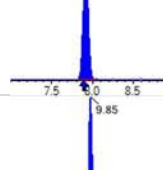
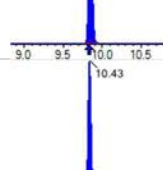
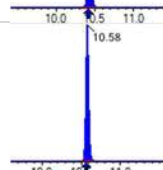
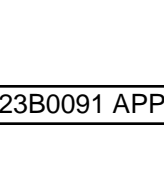
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	( 299.0 / 80.0 ) 5136990 ( 299.0 / 99.0 ) 3190293	( 5.47 , 1.00 ) ( 0.00 , N/A , 0.0 )	3875.9 3252.7	0.6210 96.0 100.4	4.2002 [ 4.4237 ]	94.9%			
PFPeS	( 349.0 / 80.0 ) 8923117 ( 349.0 / 99.0 ) 3069426	( 6.40 , 0.89 ) ( N/A , 0.03 , 0.0 )	17012.8 21400735.1	0.3440 96.4 99.9	4.6303 [ 4.6919 ]	98.7%			
PFHxS	( 399.0 / 80.0 ) 6424149 ( 399.0 / 99.0 ) 2058861	( 7.17 , 1.00 ) ( 0.00 , N/A , -0.1 )	7116.3 4212.7	0.3205 98.7 98.4	4.1809 [ 4.5549 ]	91.8%			
PFHpS	( 449.0 / 80.0 ) 7659655 ( 449.0 / 99.0 ) 2117939	( 7.81 , 0.93 ) ( N/A , 0.03 , 0.0 )	32727.4 8851.7	0.2765 101.6 99.4	4.6869 [ 4.7570 ]	98.5%			
PFOS	( 499.0 / 80.0 ) 9693221 ( 499.0 / 99.0 ) 2107427	( 8.39 , 1.00 ) ( 0.00 , N/A , 0.1 )	1587.9 1721.5	0.2174 98.2 102.5	4.2857 [ 4.6375 ]	92.4%			
PFNS	( 549.0 / 80.0 ) 11621189 ( 549.0 / 99.0 ) 2637729	( 8.80 , 1.05 ) ( N/A , 0.02 , 0.0 )	13633.8 1776613.0	0.2270 103.5 98.7	4.5720 [ 4.7994 ]	95.3%			
PFDS	( 599.0 / 80.0 ) 13668878 ( 599.0 / 99.0 ) 2847746	( 9.04 , 1.08 ) ( N/A , 0.02 , -0.1 )	3151.2 2529.9	0.2083 95.6 97.0	4.9436 [ 4.8155 ]	102.7%			
PFDoS	( 699.0 / 80.0 ) 9887363 ( 699.0 / 99.0 ) 1917055	( 9.38 , 1.12 ) ( N/A , 0.02 , 0.1 )	4094.4 2899.6	0.1939 93.2 90.4	5.6067 [ 4.8478 ]	115.7%			
4:2FTS	( 327.0 / 307.0 ) 4833875 ( 327.0 / 81.0 ) 3207632	( 5.23 , 1.00 ) ( 0.00 , N/A , -0.1 )	4945.9 2648.9	0.6636 98.3 96.8	18.5216 [ 18.6906 ]	99.1%			
6:2FTS	( 427.0 / 407.0 ) 2762871 ( 427.0 / 81.0 ) 2118933	( 6.75 , 1.00 ) ( 0.00 , N/A , 0.1 )	3384.6 2884.1	0.7669 92.7 99.3	18.0571 [ 18.9808 ]	95.1%			
8:2FTS	( 527.0 / 507.0 ) 3014674 ( 527.0 / 81.0 ) 2393308	( 7.92 , 1.00 ) ( 0.00 , N/A , 0.0 )	3357.9 2252.5	0.7939 95.5 102.6	18.8291 [ 19.1658 ]	98.2%			

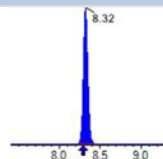
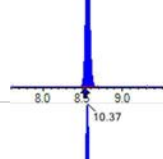
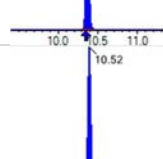
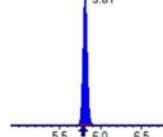

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	( 498.0 / 78.0 ) 16906990 ( 498.0 / 478.0 ) 335614	( 9.85 , 1.00 ) ( 0.00 , N/A , -0.1)	3748.5 1141.3	0.0199 96.6 93.9	4.9561 [ 5.0000 ]	99.1%			
NMeFOSA	( 512.0 / 219.0 ) 15133120 ( 512.0 / 169.0 ) 12600788	( 10.43 , 1.00 ) ( 0.00 , N/A , 0.9)	3076.9 5110.1	0.8327 101.6 101.7	19.5247 [ 20.0000 ]	97.6%			
NEIFOSA	( 526.0 / 219.0 ) 16759887 ( 526.0 / 169.0 ) 21285245	( 10.57 , 1.00 ) ( 0.00 , N/A , 0.8)	14242.1 11718.2	1.2700 100.1 102.3	20.1495 [ 20.0000 ]	100.7%			
NMeFOSAA	( 570.0 / 419.0 ) 1177465 ( 570.0 / 483.0 ) 564289	( 8.33 , 1.00 ) ( 0.01 , N/A , 0.0)	3796.1 54920.4	0.4792 101.0 103.4	4.9057 [ 5.0000 ]	98.1%			
NEIFOSAA	( 584.0 / 419.0 ) 1051835 ( 584.0 / 526.0 ) 570426	( 8.57 , 1.00 ) ( 0.00 , N/A , -0.1)	2348.6 1067.2	0.5423 106.3 92.5	4.6446 [ 5.0000 ]	92.9%			
NMeFOSE	( 616.0 / 59.0 ) 6775594	( 10.37 , 1.00 ) ( 0.01 , N/A , 0.0)	3144.3	N/A 0.0 0.0	18.5030 [ 20.0000 ]	92.5%			
NEtFOSE	( 630.0 / 59.0 ) 8941534	( 10.53 , 1.00 ) ( 0.01 , N/A , 0.0)	1378.2	N/A 0.0 0.0	19.6713 [ 20.0000 ]	98.4%			
HFPO-DA	( 285.0 / 169.0 ) 3396782 ( 285.0 / 185.0 ) 9261306	( 5.81 , 1.00 ) ( 0.00 , N/A , 0.1)	3641.9 4545.3	2.7265 99.0 98.3	9.7203 [ 10.0000 ]	97.2%			
ADONA	( 377.0 / 85.0 ) 12002718 ( 377.0 / 251.0 ) 1058144	( 6.60 , 1.14 ) ( N/A , 0.03 , 0.0)	4129.1 1995.5	0.0882 100.0 94.4	9.9875 [ 9.4270 ]	105.9%			
9CI-PF3ONS	( 531.0 / 351.0 ) 31505365 ( 533.0 / 353.0 ) 10695379	( 8.72 , 1.50 ) ( N/A , 0.02 , 0.0)	3744.2 3924.2	0.3395 97.5 102.7	8.8193 [ 9.3325 ]	94.5%			
11CI-PF3OUDS	( 631.0 / 451.0 ) 19455026 ( 633.0 / 453.0 ) 6758067	( 9.19 , 1.58 ) ( N/A , 0.02 , 0.0)	3418.5 2938.5	0.3474 96.0 97.5	9.9373 [ 9.4321 ]	105.4%			

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 ) 221401 ( 241.0 / 117.0 ) 349819	( 4.07 , 0.89 ) ( N/A , 0.01 , 0.0 )	1341.5 1338.9	1.5800 100.1 96.6	18.2002 [ 20.0000 ]	91.0%			
5:3FTCA	( 341.0 / 236.7 ) 1511872 ( 341.0 / 217.0 ) 2851796	( 6.06 , 1.10 ) ( N/A , 0.03 , 0.1 )	1486.7 2938.1	1.8863 110.6 117.2	16.5333 [ 20.0000 ]	82.7%			
7:3FTCA	( 441.0 / 317.0 ) 2820416 ( 441.0 / 337.0 ) 2545134	( 7.92 , 1.44 ) ( N/A , 0.03 , 0.1 )	1045.8 935.9	0.9024 104.9 111.0	16.6960 [ 20.0000 ]	83.5%			
PFEESA	( 315.0 / 135.0 ) 6824080 ( 315.0 / 83.0 ) 1859084	( 5.92 , 1.08 ) ( N/A , 0.02 , 0.0 )	2637.3 2134.8	0.2724 106.7 98.9	8.4951 [ 8.9246 ]	95.2%			
PFMPA	( 229.0 / 85.0 ) 987277	( 3.90 , 0.86 ) ( N/A , 0.01 , 0.0 )	4349.6	N/A 0.0 0.0	9.1682 [ 10.0000 ]	91.7%			
PFMBA	( 279.0 / 85.0 ) 4265193	( 4.87 , 1.07 ) ( N/A , 0.02 , 0.0 )	3787.0	N/A 0.0 0.0	9.4449 [ 10.0000 ]	94.4%			
NFDHA	( 295.0 / 201.0 ) 3403407 ( 295.0 / 85.0 ) 3632567	( 5.40 , 0.98 ) ( N/A , 0.02 , 0.0 )	4019.2 3880.3	1.0673 106.6 100.9	9.1696 [ 10.0000 ]	91.7%			
13C3_PFBA_IIS	( 216.0 / 172.0 ) 220252	( 3.51 , N/A ) ( N/A , 0.01 , N/A )	1201.0	N/A	0.9833 [ 1.0000 ]	98.3% { 100.4% }			
13C2_PFHxA_IIS	( 315.0 / 270.0 ) 503769	( 5.50 , N/A ) ( N/A , 0.02 , N/A )	2777.7	N/A	0.9979 [ 1.0000 ]	99.8% { 104.8% }			
13C4_PFOA_IIS	( 417.0 / 372.0 ) 603401	( 7.02 , N/A ) ( N/A , 0.03 , N/A )	1353.8	N/A	0.9521 [ 1.0000 ]	95.2% { 98.9% }			
13C5_PFNA_IIS	( 468.0 / 423.0 ) 570570	( 7.62 , N/A ) ( N/A , 0.03 , N/A )	291.9	N/A	0.9442 [ 1.0000 ]	94.4% { 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 ) 530312	( 8.18 , N/A ) ( N/A , 0.03 , N/A )	1914.4	N/A	0.9165 [ 1.0000 ]	91.6% { 99.1% }			
18O2_PFHxS_IIS	( 403.0 / 83.9 ) 835563	( 7.17 , N/A ) ( N/A , 0.03 , N/A )	1644.6	N/A	0.9757 [ 1.0000 ]	97.6% { 99.2% }			
13C4_PFOS_IIS	( 503.0 / 79.9 ) 1384649	( 8.38 , N/A ) ( N/A , 0.03 , N/A )	1552.4	N/A	0.9198 [ 1.0000 ]	92.0% { 103.3% }			
13C4_PFBA_EIS	( 217.0 / 172.0 ) 2295870	( 3.51 , N/A ) ( N/A , 0.01 , N/A )	5201.3	N/A	8.7633 [ 8.0000 ]	109.5% { 99.9% }			
13C5_PFPeA_EIS	( 268.0 / 223.0 ) 2396881	( 4.56 , N/A ) ( N/A , 0.02 , N/A )	3930.3	N/A	4.3486 [ 4.0000 ]	108.7% { 104.0% }			
13C5_PFHxA_EIS	( 318.0 / 273.0 ) 1521284	( 5.50 , N/A ) ( N/A , 0.02 , N/A )	3768.3	N/A	2.2047 [ 2.0000 ]	110.2% { 106.6% }			
13C4_PFHpA_EIS	( 367.0 / 322.0 ) 1330115	( 6.32 , N/A ) ( N/A , 0.03 , N/A )	2179.0	N/A	2.2248 [ 2.0000 ]	111.2% { 112.9% }			
13C8_PFOA_EIS	( 421.0 / 376.0 ) 1522906	( 7.02 , N/A ) ( N/A , 0.03 , N/A )	2039.9	N/A	2.2285 [ 2.0000 ]	111.4% { 107.2% }			
13C9_PFNA_EIS	( 472.0 / 427.0 ) 627849	( 7.62 , N/A ) ( N/A , 0.03 , N/A )	3209.0	N/A	1.0386 [ 1.0000 ]	103.9% { 103.2% }			
13C6_PFDA_EIS	( 519.0 / 474.0 ) 782750	( 8.18 , N/A ) ( N/A , 0.03 , N/A )	3972.2	N/A	1.1707 [ 1.0000 ]	117.1% { 105.0% }			
13C7_PFUnA_EIS	( 570.0 / 525.0 ) 795270	( 8.66 , N/A ) ( N/A , 0.03 , N/A )	3450.2	N/A	1.1961 [ 1.0000 ]	119.6% { 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_EIS	( 615.0 / 570.0 ) 634917	( 8.94 , N/A ) ( N/A , 0.02 , N/A )	2073.4	N/A	1.1134 [ 1.0000 ]	111.3% { 96.5% }			
13C2_PFTeDA_EIS	( 715.0 / 670.0 ) 606570	( 9.30 , N/A ) ( N/A , 0.01 , N/A )	1900.4	N/A	1.1608 [ 1.0000 ]	116.1% { 103.5% }			
13C3_PFBs_EIS	( 302.0 / 80.0 ) 3936105	( 5.47 , N/A ) ( N/A , 0.02 , N/A )	3991.7	N/A	2.2508 [ 2.0000 ]	112.5% { 103.8% }			
13C3_PFHxS_EIS	( 402.0 / 80.0 ) 1998497	( 7.16 , N/A ) ( N/A , 0.03 , N/A )	2199.6	N/A	2.2002 [ 2.0000 ]	110.0% { 101.7% }			
13C8_PFOS_EIS	( 507.0 / 80.0 ) 4289770	( 8.38 , N/A ) ( N/A , 0.03 , N/A )	1442.5	N/A	2.2567 [ 2.0000 ]	112.8% { 101.2% }			
13C2_4:2FTS_EIS	( 329.0 / 81.0 ) 382230	( 5.23 , N/A ) ( N/A , 0.02 , N/A )	1717.6	N/A	3.9598 [ 4.0000 ]	99.0% { 110.1% }			
13C2_6:2FTS_EIS	( 429.0 / 81.0 ) 431339	( 6.75 , N/A ) ( N/A , 0.03 , N/A )	1237.0	N/A	4.0709 [ 4.0000 ]	101.8% { 103.7% }			
13C2_8:2FTS_EIS	( 529.0 / 81.0 ) 587953	( 7.92 , N/A ) ( N/A , 0.03 , N/A )	818.7	N/A	3.9276 [ 4.0000 ]	98.2% { 97.5% }			
13C8_PFOsa_EIS	( 506.0 / 78.0 ) 8530456	( 9.85 , N/A ) ( N/A , 0.02 , N/A )	3619.9	N/A	2.4175 [ 2.0000 ]	120.9% { 102.7% }			
D3_NMeFOSA_EIS	( 515.0 / 169.0 ) 1866344	( 10.43 , N/A ) ( N/A , 0.01 , N/A )	2655.4	N/A	2.2450 [ 2.0000 ]	112.2% { 96.4% }			
D5_NEtFOSA_EIS	( 531.0 / 169.0 ) 1780659	( 10.58 , N/A ) ( N/A , 0.01 , N/A )	3596.3	N/A	2.3936 [ 2.0000 ]	119.7% { 99.8% }			

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 ) 1223396	( 8.32 , N/A ) ( N/A , 0.03 , N/A )	1932.0	N/A	3.9769 [ 4.0000 ]	99.4% { 92.4% }			
D5_EtFOSAA_EIS	( 589.0 / 419.0 ) 1078533	( 8.56 , N/A ) ( N/A , 0.03 , N/A )	118999.3	N/A	4.2818 [ 4.0000 ]	107.0% { 102.2% }			
D7_NMeFOSE_EIS	( 623.0 / 58.9 ) 7325708	( 10.37 , N/A ) ( N/A , 0.01 , N/A )	2685.5	N/A	24.1706 [ 20.0000 ]	120.9% { 98.6% }			
D9_NEtFOSE_EIS	( 639.0 / 58.9 ) 9553012	( 10.52 , N/A ) ( N/A , 0.01 , N/A )	1978.6	N/A	24.6124 [ 20.0000 ]	123.1% { 101.4% }			
13C3_HFPODA_EIS	( 287.0 / 169.0 ) 3535187	( 5.81 , N/A ) ( N/A , 0.02 , N/A )	3501.7	N/A	8.4144 [ 8.0000 ]	105.2% { 98.6% }			

# INITIAL AND CONTINUING CALIBRATION CHECK

## EPA 1633

Laboratory: APPL, LLC

Work Order: 23B0091

Client: AECOM

Project: Red Hill AFFF Assessment Sampling

Instrument ID: Saphira

Calibration: 2307007

Standard ID: 23B0084

Sequence: SC00636

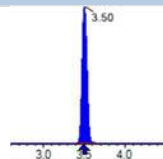
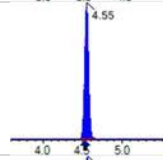
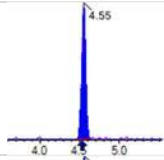
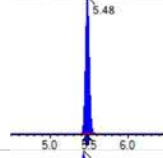
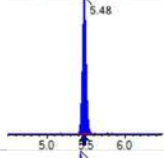
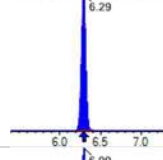
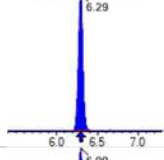
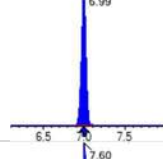
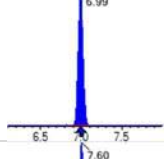
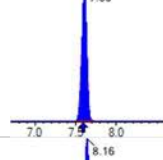
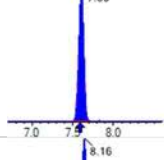
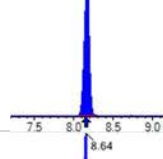
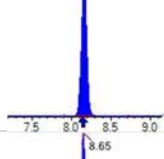
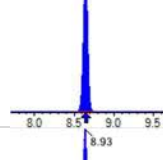
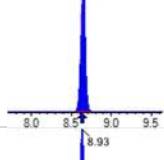
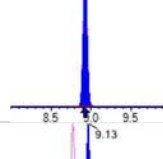
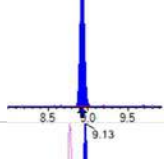
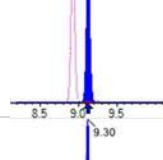
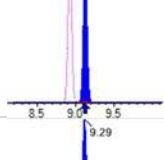
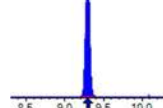
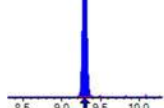
Lab Sample ID	Analyte	True	Found	%R	Units	Control Limit
SC00636-CCV3	PFBA	20.0	18.6	93.1	ng/mL	+/- 30.00%
	PFPEA	10.0	9.58	95.8	ng/mL	+/- 30.00%
	PFHXA	5.00	4.63	92.6	ng/mL	+/- 30.00%
	PFHPA	5.00	4.50	90.0	ng/mL	+/- 30.00%
	PFOA	5.00	4.86	97.2	ng/mL	+/- 30.00%
	PFNA	5.00	4.92	98.3	ng/mL	+/- 30.00%
	PFDA	5.00	4.69	93.7	ng/mL	+/- 30.00%
	PFUnA	5.00	4.59	91.8	ng/mL	+/- 30.00%
	PFDOA	5.00	4.67	93.5	ng/mL	+/- 30.00%
	PFTRDA	5.00	4.39	87.8	ng/mL	+/- 30.00%
	PFTEDA	5.00	4.90	98.0	ng/mL	+/- 30.00%
	PFBS	4.42	4.25	96.1	ng/mL	+/- 30.00%
	PFPEs	4.70	5.05	107	ng/mL	+/- 30.00%
	PFHXS	4.58	4.36	95.1	ng/mL	+/- 30.00%
	PFHPS	4.78	4.93	103	ng/mL	+/- 30.00%
	PFOS	4.65	4.34	93.3	ng/mL	+/- 30.00%
	PFNS	4.80	4.46	92.9	ng/mL	+/- 30.00%
	PFDS	4.82	4.69	97.3	ng/mL	+/- 30.00%
	PFDOS	4.85	4.77	98.3	ng/mL	+/- 30.00%
	4:2FTS	18.8	19.2	102	ng/mL	+/- 30.00%
	6:2FTS	19.0	16.6	87.5	ng/mL	+/- 30.00%
	8:2FTS	19.2	17.7	92.1	ng/mL	+/- 30.00%
	PFOSA	5.00	5.41	108	ng/mL	+/- 30.00%
	NMeFOSA	20.0	19.3	96.5	ng/mL	+/- 30.00%
	NEtFOSA	20.0	20.4	102	ng/mL	+/- 30.00%
	NMeFOSAA	5.00	4.86	97.3	ng/mL	+/- 30.00%
	NEtFOSAA	5.00	4.41	88.3	ng/mL	+/- 30.00%
	NMeFOSE	20.0	18.4	91.8	ng/mL	+/- 30.00%
	NEtFOSE	20.0	18.7	93.5	ng/mL	+/- 30.00%
	HFPO-DA	10.0	9.05	90.5	ng/mL	+/- 30.00%

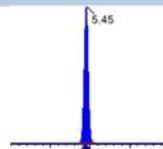
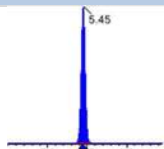
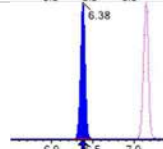
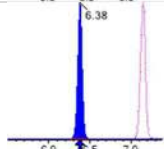
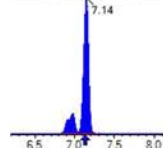
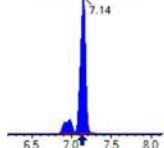
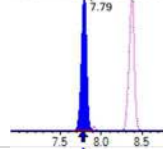
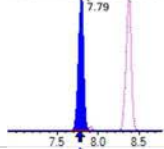
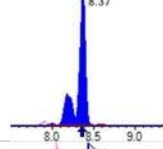
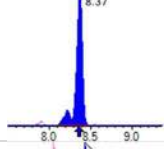
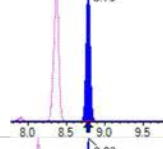
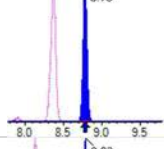
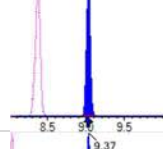
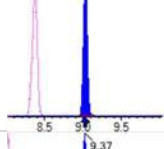
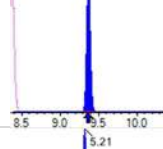
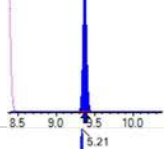
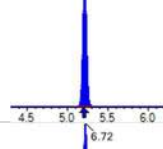
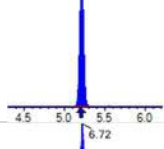
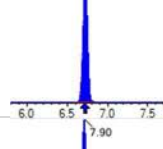
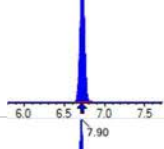
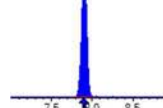
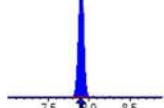
# INITIAL AND CONTINUING CALIBRATION CHECK

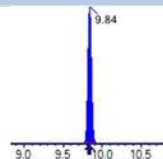
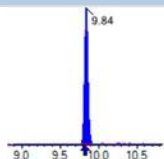
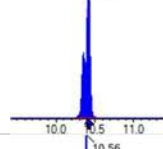
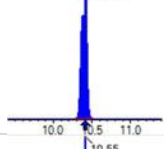
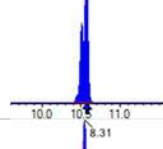
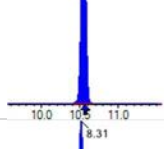
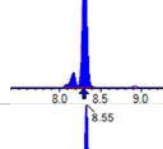
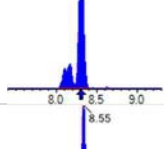
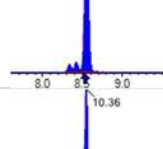
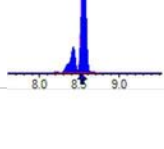
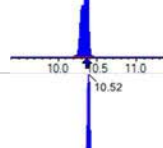
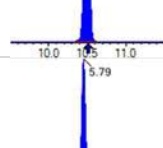
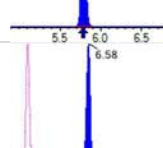
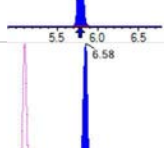
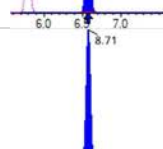
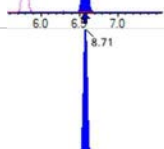
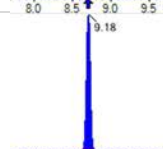
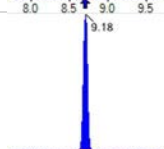
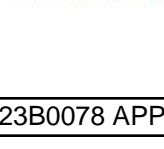
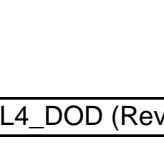
## EPA 1633

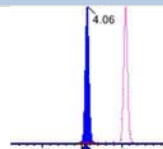
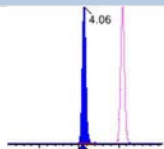
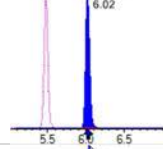
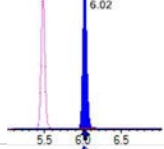
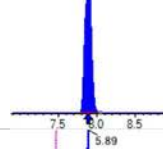
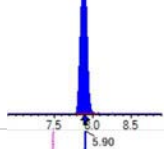
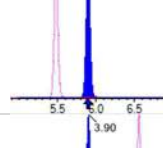
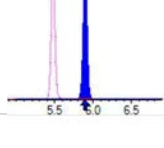
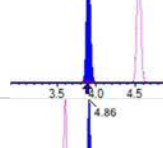
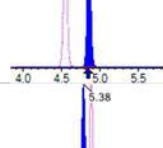
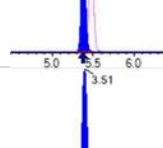
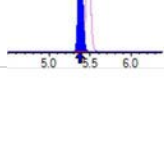
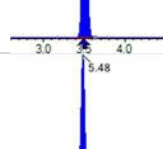
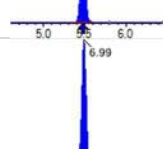
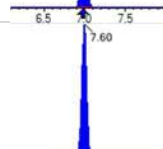

Laboratory:	APPL, LLC	Work Order:	23B0091
Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Instrument ID:	Saphira	Calibration:	2307007
Standard ID:	23B0084	Sequence:	SC00636

Lab Sample ID	Analyte	True	Found	%R	Units	Control Limit
SC00636-CCV3	ADONA	9.45	9.84	104	ng/mL	+/- 30.00%
	PFEESA	8.90	8.64	97.0	ng/mL	+/- 30.00%
	PFMPA	10.0	9.25	92.5	ng/mL	+/- 30.00%
	PFMBA	10.0	9.28	92.8	ng/mL	+/- 30.00%
	NFDHA	10.0	9.50	95.0	ng/mL	+/- 30.00%
	9CL-PF3ONS	9.35	8.70	93.0	ng/mL	+/- 30.00%
	11CL-PF3OUDS	9.45	8.25	87.3	ng/mL	+/- 30.00%
	3:3FTCA	20.0	20.3	102	ng/mL	+/- 30.00%
	5:3FTCA	20.0	19.0	94.8	ng/mL	+/- 30.00%
	7:3FTCA	20.0	20.1	100	ng/mL	+/- 30.00%

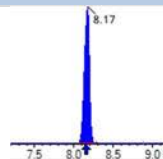
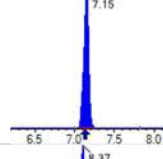
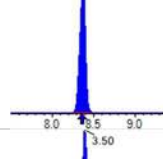
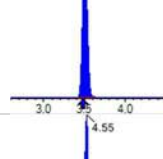
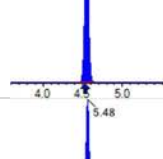
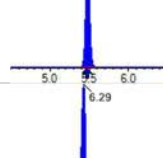
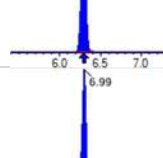
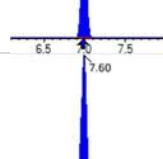
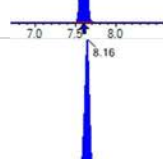
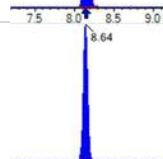
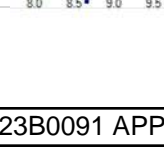
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	( 213.0 / 169.0 ) 4963325	( 3.50 , 1.00 ) ( 0.00 , N/A , 0.0 )	242.5	N/A 0.0 0.0	18.6179 [ 20.0000 ]	93.1%			
PFPeA	( 263.0 / 219.0 ) 5240322 ( 263.0 / 69.0 ) 64108	( 4.55 , 1.00 ) ( 0.00 , N/A , -0.2 )	2825.8 388.0	0.0122 114.2 111.0	9.5774 [ 10.0000 ]	95.8%			
PFHxA	( 313.0 / 269.0 ) 3321861 ( 313.0 / 119.0 ) 354228	( 5.48 , 1.00 ) ( 0.00 , N/A , 0.1 )	6417.9 4645865.7	0.1066 107.1 112.0	4.6298 [ 5.0000 ]	92.6%			
PFHpA	( 363.0 / 319.0 ) 2706089 ( 363.0 / 169.0 ) 825452	( 6.29 , 1.00 ) ( 0.00 , N/A , 0.0 )	11848.0 28415.7	0.3050 95.1 95.4	4.4984 [ 5.0000 ]	90.0%			
PFOA	( 413.0 / 369.0 ) 3564035 ( 413.0 / 169.0 ) 1094448	( 6.99 , 1.00 ) ( 0.00 , N/A , 0.1 )	6607.1 3662.5	0.3071 91.8 85.0	4.8609 [ 5.0000 ]	97.2%			
PFNA	( 463.0 / 419.0 ) 2962201 ( 463.0 / 169.0 ) 710674	( 7.60 , 1.00 ) ( 0.00 , N/A , 0.0 )	41109.4 3704.0	0.2399 107.2 108.5	4.9173 [ 5.0000 ]	98.3%			
PFDA	( 513.0 / 469.0 ) 3622051 ( 513.0 / 169.0 ) 419696	( 8.16 , 1.00 ) ( 0.00 , N/A , 0.1 )	2022.8 1098.4	0.1159 95.3 106.8	4.6864 [ 5.0000 ]	93.7%			
PFUnA	( 563.0 / 519.0 ) 3705669 ( 563.0 / 169.0 ) 434866	( 8.64 , 1.00 ) ( 0.00 , N/A , -0.2 )	2616.3 2743.7	0.1174 111.9 100.4	4.5890 [ 5.0000 ]	91.8%			
PFDaA	( 613.0 / 569.0 ) 3313246 ( 613.0 / 169.0 ) 559562	( 8.93 , 1.00 ) ( 0.00 , N/A , 0.0 )	1829.3 764.0	0.1689 103.3 100.6	4.6748 [ 5.0000 ]	93.5%			
PFTTrDA	( 663.0 / 619.0 ) 2815901 ( 663.0 / 169.0 ) 785313	( 9.13 , 1.02 ) ( N/A , 0.01 , -0.1 )	2729.8 1892.3	0.2789 116.7 103.6	4.3881 [ 5.0000 ]	87.8%			
PFTeDA	( 713.0 / 669.0 ) 2417868 ( 713.0 / 169.0 ) 533016	( 9.30 , 1.00 ) ( 0.00 , N/A , 0.1 )	1912.9 845.6	0.2204 106.7 106.6	4.9022 [ 5.0000 ]	98.0%			

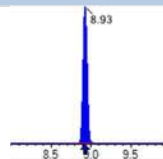
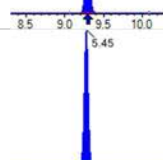
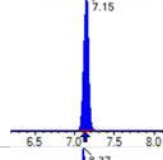
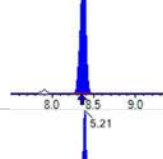
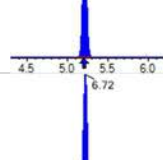
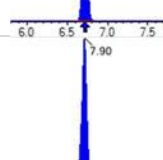
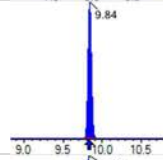
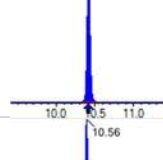
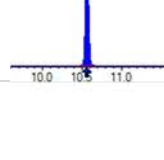
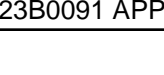
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) ( $\Delta$ RT-[min], $\Delta$ RT- CV[min], $\Delta$ RT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	( 299.0 / 80.0 ) 6016217 ( 299.0 / 99.0 ) 3776191	( 5.45 , 1.00 ) ( 0.00 , N/A , 0.0 )	3540.1 3159.9	0.6277 97.1 101.5	4.2495 [ 4.4237 ]	96.1%			
PFPeS	( 349.0 / 80.0 ) 10596412 ( 349.0 / 99.0 ) 3648414	( 6.38 , 0.89 ) ( N/A , 0.00 , -0.1 )	37081.4 4060.0	0.3443 96.4 99.9	5.0466 [ 4.6919 ]	107.6%			
PFHxS	( 399.0 / 80.0 ) 7295454 ( 399.0 / 99.0 ) 2347674	( 7.14 , 1.00 ) ( 0.00 , N/A , 0.1 )	7298.7 3355.7	0.3218 99.1 98.8	4.3576 [ 4.5549 ]	95.7%			
PFHpS	( 449.0 / 80.0 ) 9564933 ( 449.0 / 99.0 ) 2479968	( 7.79 , 0.93 ) ( N/A , 0.02 , 0.0 )	62996.9 3196.5	0.2593 95.2 93.2	4.9284 [ 4.7570 ]	103.6%			
PFOS	( 499.0 / 80.0 ) 11657549 ( 499.0 / 99.0 ) 2522472	( 8.37 , 1.00 ) ( 0.00 , N/A , 0.1 )	1403.2 2033.4	0.2164 97.8 102.1	4.3403 [ 4.6375 ]	93.6%			
PFNS	( 549.0 / 80.0 ) 13457862 ( 549.0 / 99.0 ) 3201417	( 8.79 , 1.05 ) ( N/A , 0.01 , 0.1 )	21675.3 104183.1	0.2379 108.5 103.4	4.4584 [ 4.7994 ]	92.9%			
PFDS	( 599.0 / 80.0 ) 15403088 ( 599.0 / 99.0 ) 3251219	( 9.03 , 1.08 ) ( N/A , 0.01 , 0.0 )	3770.2 2777.5	0.2111 96.8 98.3	4.6910 [ 4.8155 ]	97.4%			
PFDoS	( 699.0 / 80.0 ) 9986599 ( 699.0 / 99.0 ) 2090771	( 9.37 , 1.12 ) ( N/A , 0.01 , 0.0 )	4168.6 2270.1	0.2094 100.6 97.6	4.7687 [ 4.8478 ]	98.4%			
4:2FTS	( 327.0 / 307.0 ) 6464638 ( 327.0 / 81.0 ) 4437238	( 5.21 , 1.00 ) ( 0.00 , N/A , 0.0 )	5036.6 3839.0	0.6864 101.7 100.1	19.2395 [ 18.6906 ]	102.9%			
6:2FTS	( 427.0 / 407.0 ) 3605437 ( 427.0 / 81.0 ) 3093247	( 6.72 , 1.00 ) ( 0.00 , N/A , 0.0 )	2062.8 2693.1	0.8579 103.7 111.1	16.6236 [ 18.9808 ]	87.6%			
8:2FTS	( 527.0 / 507.0 ) 4083701 ( 527.0 / 81.0 ) 3479454	( 7.90 , 1.00 ) ( 0.00 , N/A , 0.0 )	1541.0 2116.9	0.8520 102.5 110.1	17.6774 [ 19.1658 ]	92.2%			

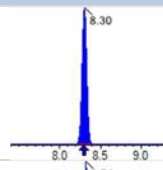




Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	( 498.0 / 78.0 ) 12182616 ( 498.0 / 478.0 ) 246161	( 9.84 , 1.00 ) ( 0.00 , N/A , -0.1)	2738.4 749.0	0.0202 98.4 95.6	5.4111 [ 5.0000 ]	108.2%			
NMeFOSA	( 512.0 / 219.0 ) 7134819 ( 512.0 / 169.0 ) 6177981	( 10.42 , 1.00 ) ( 0.00 , N/A , 1.0)	3741.7 3643.8	0.8659 105.7 105.7	19.2964 [ 20.0000 ]	96.5%			
NEIFOSA	( 526.0 / 219.0 ) 7534863 ( 526.0 / 169.0 ) 9618383	( 10.56 , 1.00 ) ( 0.00 , N/A , 1.0)	6827.8 9516.9	1.2765 100.6 102.8	20.4402 [ 20.0000 ]	102.2%			
NMeFOSAA	( 570.0 / 419.0 ) 1627281 ( 570.0 / 483.0 ) 732054	( 8.31 , 1.00 ) ( 0.00 , N/A , 0.0)	3302.1 6654.3	0.4499 94.8 97.1	4.8635 [ 5.0000 ]	97.3%			
NEIFOSAA	( 584.0 / 419.0 ) 1421409 ( 584.0 / 526.0 ) 801249	( 8.55 , 1.00 ) ( 0.01 , N/A , -0.1)	1720.6 1782.0	0.5637 110.5 96.2	4.4128 [ 5.0000 ]	88.3%			
NMeFOSE	( 616.0 / 59.0 ) 3014212	( 10.36 , 1.00 ) ( 0.01 , N/A , 0.0)	2130.3	N/A 0.0 0.0	18.3567 [ 20.0000 ]	91.8%			
NEtFOSE	( 630.0 / 59.0 ) 3661555	( 10.52 , 1.00 ) ( 0.01 , N/A , 0.0)	1310.6	N/A 0.0 0.0	18.7084 [ 20.0000 ]	93.5%			
HFPO-DA	( 285.0 / 169.0 ) 3543071 ( 285.0 / 185.0 ) 10550482	( 5.79 , 1.00 ) ( 0.00 , N/A , 0.0)	4167.2 4588.0	2.9778 108.1 107.3	9.0496 [ 10.0000 ]	90.5%			
ADONA	( 377.0 / 85.0 ) 13247320 ( 377.0 / 251.0 ) 1214198	( 6.58 , 1.14 ) ( N/A , 0.00 , 0.0)	4299.8 1571.4	0.0917 104.0 98.2	9.8389 [ 9.4270 ]	104.4%			
9CI-Pf3ONS	( 531.0 / 351.0 ) 34867807 ( 533.0 / 353.0 ) 12064687	( 8.71 , 1.50 ) ( N/A , 0.01 , -0.1)	4332.0 2821.3	0.3460 99.4 104.7	8.6994 [ 9.3325 ]	93.2%			
11CI-PF3OUDS	( 631.0 / 451.0 ) 18096158 ( 633.0 / 453.0 ) 6601998	( 9.18 , 1.59 ) ( N/A , 0.01 , -0.1)	4135.0 2491.7	0.3648 100.8 102.4	8.2502 [ 9.4321 ]	87.5%			

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
3:3FTCA	( 241.0 / 177.0 ) 275510 ( 241.0 / 117.0 ) 465756	( 4.06 , 0.89 ) ( N/A , 0.01 , 0.0 )	1457.2 1229.6	1.6905 107.1 103.3	20.3072 [ 20.0000 ]	101.5%			
5:3FTCA	( 341.0 / 236.7 ) 1859884 ( 341.0 / 217.0 ) 3426694	( 6.02 , 1.10 ) ( N/A , -0.02 , -0.2 )	2124.5 2356.4	1.8424 108.0 114.4	18.9559 [ 20.0000 ]	94.8%			
7:3FTCA	( 441.0 / 317.0 ) 3641071 ( 441.0 / 337.0 ) 3247791	( 7.89 , 1.44 ) ( N/A , 0.00 , -0.1 )	1124.4 1562.3	0.8920 103.7 109.7	20.0884 [ 20.0000 ]	100.4%			
PFEESA	( 315.0 / 135.0 ) 7442584 ( 315.0 / 83.0 ) 2211935	( 5.89 , 1.08 ) ( N/A , 0.00 , -0.1 )	3039.2 1918.7	0.2972 116.4 107.9	8.6351 [ 8.9246 ]	96.8%			
PFMPA	( 229.0 / 85.0 ) 1105479	( 3.90 , 0.86 ) ( N/A , 0.01 , 0.0 )	2969.3	N/A 0.0 0.0	9.2498 [ 10.0000 ]	92.5%			
PFMBA	( 279.0 / 85.0 ) 4651363	( 4.86 , 1.07 ) ( N/A , 0.01 , 0.0 )	4393.3	N/A 0.0 0.0	9.2805 [ 10.0000 ]	92.8%			
NFDHA	( 295.0 / 201.0 ) 3784925 ( 295.0 / 85.0 ) 3904070	( 5.38 , 0.98 ) ( N/A , 0.00 , 0.0 )	4828.1 3080.0	1.0315 103.0 97.6	9.5040 [ 10.0000 ]	95.0%			
13C3_PFBa_IIS	( 216.0 / 172.0 ) 235318	( 3.51 , N/A ) ( N/A , 0.00 , N/A )	1137.5	N/A	1.0506 [ 1.0000 ]	105.1% { 107.2% }			
13C2_PFHxA_IIS	( 315.0 / 270.0 ) 527449	( 5.48 , N/A ) ( N/A , 0.00 , N/A )	1147.0	N/A	1.0449 [ 1.0000 ]	104.5% { 109.8% }			
13C4_PFOA_IIS	( 417.0 / 372.0 ) 642931	( 6.99 , N/A ) ( N/A , 0.00 , N/A )	3295.7	N/A	1.0144 [ 1.0000 ]	101.4% { 105.3% }			
13C5_PFNA_IIS	( 468.0 / 423.0 ) 637258	( 7.60 , N/A ) ( N/A , 0.01 , N/A )	1753.8	N/A	1.0546 [ 1.0000 ]	105.5% { 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
13C2_PFDA_IIS	( 515.0 / 470.1 ) 663878	( 8.17 , N/A ) ( N/A , 0.02 , N/A )	1278.6	N/A	1.1473 [ 1.0000 ]	114.7% { 124.0% }			
18O2_PFHxS_IIS	( 403.0 / 83.9 ) 904577	( 7.15 , N/A ) ( N/A , 0.01 , N/A )	1499.6	N/A	1.0563 [ 1.0000 ]	105.6% { 107.4% }			
13C4_PFOS_IIS	( 503.0 / 79.9 ) 1686346	( 8.37 , N/A ) ( N/A , 0.02 , N/A )	1231.1	N/A	1.1202 [ 1.0000 ]	112.0% { 125.8% }			
13C4_PFBA_EIS	( 217.0 / 172.0 ) 2574656	( 3.50 , N/A ) ( N/A , 0.01 , N/A )	4981.9	N/A	9.1982 [ 8.0000 ]	115.0% { 112.1% }			
13C5_PFPeA_EIS	( 268.0 / 223.0 ) 2660176	( 4.55 , N/A ) ( N/A , 0.01 , N/A )	3751.1	N/A	4.6096 [ 4.0000 ]	115.2% { 115.4% }			
13C5_PFHxA_EIS	( 318.0 / 273.0 ) 1632280	( 5.48 , N/A ) ( N/A , 0.00 , N/A )	2799.2	N/A	2.2593 [ 2.0000 ]	113.0% { 114.3% }			
13C4_PFHpA_EIS	( 367.0 / 322.0 ) 1424801	( 6.29 , N/A ) ( N/A , 0.00 , N/A )	2512.3	N/A	2.2761 [ 2.0000 ]	113.8% { 120.9% }			
13C8_PFOA_EIS	( 421.0 / 376.0 ) 1646937	( 6.99 , N/A ) ( N/A , 0.01 , N/A )	2685.4	N/A	2.2618 [ 2.0000 ]	113.1% { 116.0% }			
13C9_PFNA_EIS	( 472.0 / 427.0 ) 680855	( 7.60 , N/A ) ( N/A , 0.01 , N/A )	4200.1	N/A	1.0084 [ 1.0000 ]	100.8% { 111.9% }			
13C6_PFDA_EIS	( 519.0 / 474.0 ) 878634	( 8.16 , N/A ) ( N/A , 0.02 , N/A )	2172.0	N/A	1.0497 [ 1.0000 ]	105.0% { 117.8% }			
13C7_PFUnA_EIS	( 570.0 / 525.0 ) 971932	( 8.64 , N/A ) ( N/A , 0.01 , N/A )	2258.1	N/A	1.1677 [ 1.0000 ]	116.8% { 145.9% }			

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 ) 805973	( 8.93 , N/A ) ( N/A , 0.01 , N/A )	4157.0	N/A	1.1290 [ 1.0000 ]	112.9% { 122.4% }			
13C2_PFTeDA_EIS	( 715.0 / 670.0 ) 569967	( 9.29 , N/A ) ( N/A , 0.01 , N/A )	897.6	N/A	0.8713 [ 1.0000 ]	87.1% { 97.3% }			
13C3_PFBs_EIS	( 302.0 / 80.0 ) 4556259	( 5.45 , N/A ) ( N/A , 0.00 , N/A )	2834.7	N/A	2.4066 [ 2.0000 ]	120.3% { 120.1% }			
13C3_PFHxS_EIS	( 402.0 / 80.0 ) 2177499	( 7.15 , N/A ) ( N/A , 0.01 , N/A )	2722.9	N/A	2.2144 [ 2.0000 ]	110.7% { 110.8% }			
13C8_PFOS_EIS	( 507.0 / 80.0 ) 5094258	( 8.37 , N/A ) ( N/A , 0.02 , N/A )	1454.1	N/A	2.2004 [ 2.0000 ]	110.0% { 120.2% }			
13C2_4:2FTS_EIS	( 329.0 / 81.0 ) 492105	( 5.21 , N/A ) ( N/A , 0.01 , N/A )	2065.9	N/A	4.7092 [ 4.0000 ]	117.7% { 141.7% }			
13C2_6:2FTS_EIS	( 429.0 / 81.0 ) 611422	( 6.72 , N/A ) ( N/A , 0.00 , N/A )	6627.8	N/A	5.3302 [ 4.0000 ]	133.3% { 147.0% }			
13C2_8:2FTS_EIS	( 529.0 / 81.0 ) 848334	( 7.90 , N/A ) ( N/A , 0.02 , N/A )	1307.9	N/A	5.2346 [ 4.0000 ]	130.9% { 140.6% }			
13C8_PFOA_EIS	( 506.0 / 78.0 ) 5629813	( 9.84 , N/A ) ( N/A , 0.01 , N/A )	3394.6	N/A	1.3100 [ 2.0000 ]	65.5% { 67.8% }			
D3_NMeFOA_EIS	( 515.0 / 169.0 ) 889918	( 10.42 , N/A ) ( N/A , 0.00 , N/A )	2153.2	N/A	0.8789 [ 2.0000 ]	43.9% { 46.0% }			
D5_NEtFOA_EIS	( 531.0 / 169.0 ) 789160	( 10.56 , N/A ) ( N/A , 0.00 , N/A )	1856.5	N/A	0.8710 [ 2.0000 ]	43.6% { 44.2% }			

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 ) 1705433	( 8.30 , N/A ) ( N/A , 0.02 , N/A )	2522.8	N/A	4.5521 [ 4.0000 ]	113.8% { 128.8% }			
D5_EtFOSAA_EIS	( 589.0 / 419.0 ) 1534048	( 8.54 , N/A ) ( N/A , 0.01 , N/A )	6684.0	N/A	5.0006 [ 4.0000 ]	125.0% { 145.4% }			
D7_NMeFOSE_EIS	( 623.0 / 58.9 ) 3284906	( 10.35 , N/A ) ( N/A , 0.00 , N/A )	2076.7	N/A	8.8992 [ 20.0000 ]	44.5% { 44.2% }			
D9_NEtFOSE_EIS	( 639.0 / 58.9 ) 4113289	( 10.51 , N/A ) ( N/A , 0.00 , N/A )	1694.3	N/A	8.7015 [ 20.0000 ]	43.5% { 43.7% }			
13C3_HFPODA_EIS	( 287.0 / 169.0 ) 3960709	( 5.79 , N/A ) ( N/A , 0.00 , N/A )	3693.1	N/A	9.0039 [ 8.0000 ]	112.5% { 110.4% }			

# INITIAL AND CONTINUING CALIBRATION CHECK

## EPA 1633

Laboratory: APPL, LLC

Work Order: 23B0091

Client: AECOM

Project: Red Hill AFFF Assessment Sampling

Instrument ID: Saphira

Calibration: 2307007

Standard ID: 23B0084

Sequence: SC00636

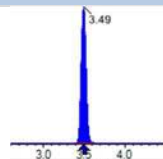
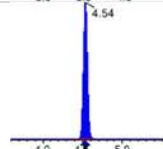
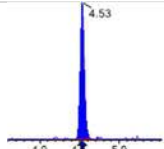
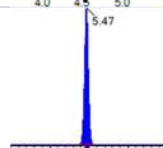
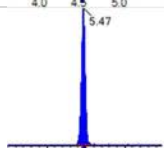
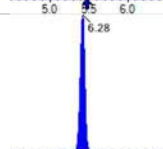
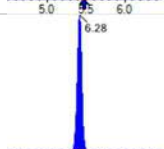
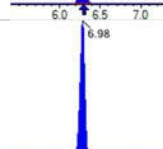
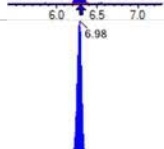
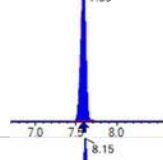
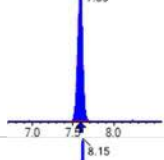
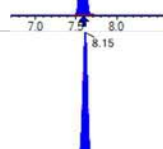
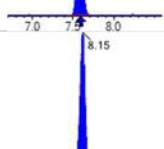
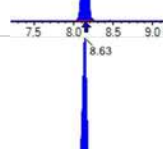
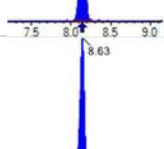
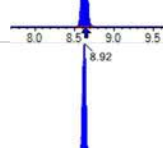
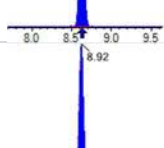
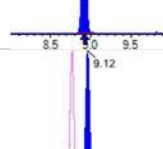
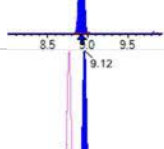
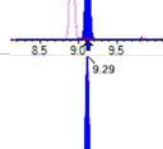
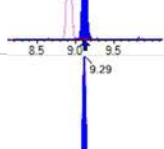
Lab Sample ID	Analyte	True	Found	%R	Units	Control Limit
SC00636-CCV4	PFBA	20.0	19.3	96.7	ng/mL	+/- 30.00%
	PFPEA	10.0	9.34	93.4	ng/mL	+/- 30.00%
	PFHXA	5.00	4.71	94.2	ng/mL	+/- 30.00%
	PFHPA	5.00	4.68	93.5	ng/mL	+/- 30.00%
	PFOA	5.00	5.05	101	ng/mL	+/- 30.00%
	PFNA	5.00	4.94	98.8	ng/mL	+/- 30.00%
	PFDA	5.00	5.38	108	ng/mL	+/- 30.00%
	PFUnA	5.00	4.59	91.9	ng/mL	+/- 30.00%
	PFDOA	5.00	4.29	85.8	ng/mL	+/- 30.00%
	PFTRDA	5.00	4.47	89.3	ng/mL	+/- 30.00%
	PFTEDA	5.00	4.79	95.8	ng/mL	+/- 30.00%
	PFBS	4.42	4.18	94.7	ng/mL	+/- 30.00%
	PFPEs	4.70	4.69	99.9	ng/mL	+/- 30.00%
	PFHXS	4.58	4.40	96.2	ng/mL	+/- 30.00%
	PFHPS	4.78	4.85	101	ng/mL	+/- 30.00%
	PFOS	4.65	4.36	93.9	ng/mL	+/- 30.00%
	PFNS	4.80	4.71	98.1	ng/mL	+/- 30.00%
	PFDS	4.82	4.91	102	ng/mL	+/- 30.00%
	PFDOS	4.85	5.05	104	ng/mL	+/- 30.00%
	4:2FTS	18.8	16.5	87.8	ng/mL	+/- 30.00%
	6:2FTS	19.0	17.3	91.3	ng/mL	+/- 30.00%
	8:2FTS	19.2	18.2	94.6	ng/mL	+/- 30.00%
	PFOSA	5.00	4.98	99.5	ng/mL	+/- 30.00%
	NMeFOSA	20.0	19.8	98.8	ng/mL	+/- 30.00%
	NEtFOSA	20.0	20.7	104	ng/mL	+/- 30.00%
	NMeFOSAA	5.00	4.65	93.0	ng/mL	+/- 30.00%
	NEtFOSAA	5.00	4.31	86.3	ng/mL	+/- 30.00%
	NMeFOSE	20.0	18.1	90.4	ng/mL	+/- 30.00%
	NEtFOSE	20.0	19.7	98.6	ng/mL	+/- 30.00%
	HFPO-DA	10.0	8.39	83.9	ng/mL	+/- 30.00%

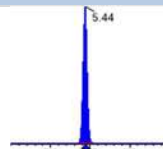
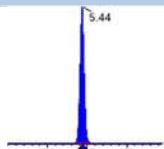
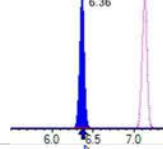
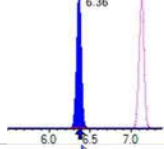
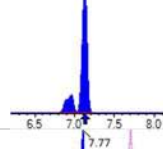
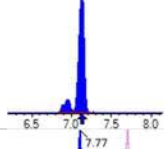
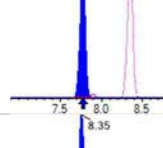
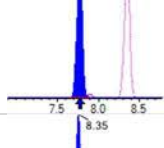
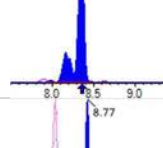
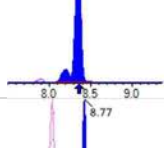
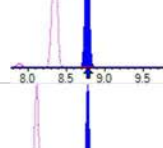
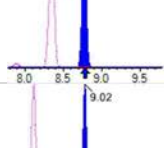
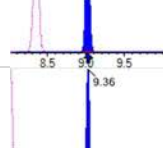
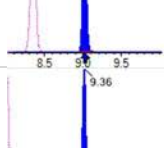
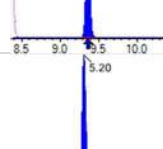
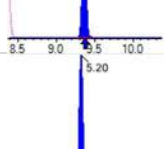
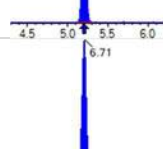
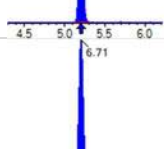
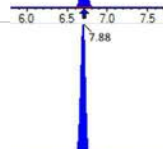
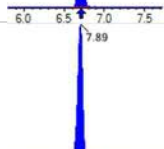

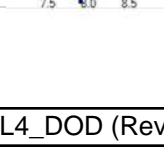
# INITIAL AND CONTINUING CALIBRATION CHECK

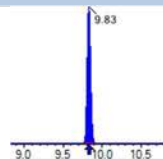
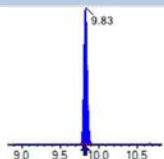
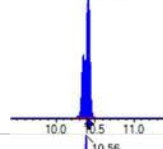
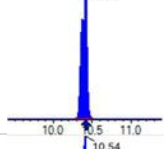
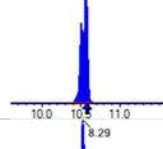
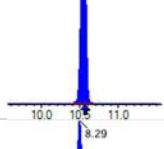
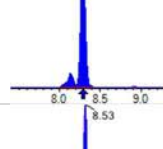
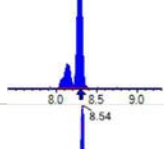
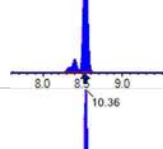
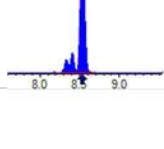
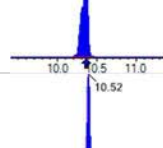
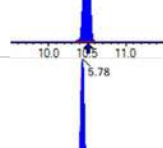
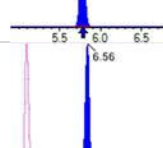
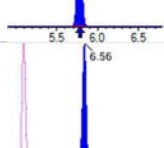
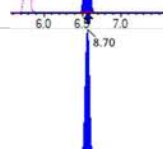
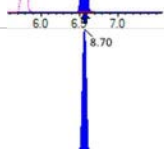
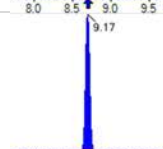
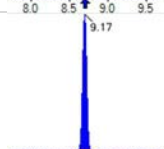
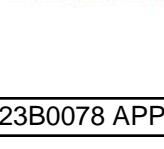
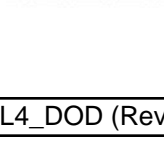
## EPA 1633

Laboratory:	APPL, LLC	Work Order:	23B0091
Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Instrument ID:	Saphira	Calibration:	2307007
Standard ID:	23B0084	Sequence:	SC00636

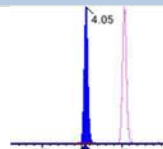
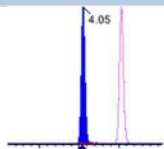
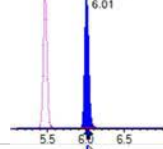
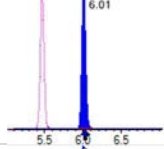
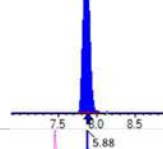
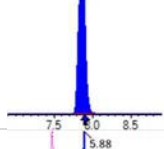
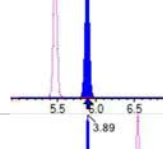
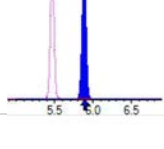
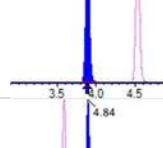
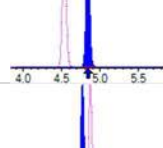
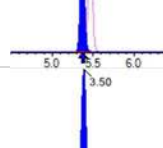
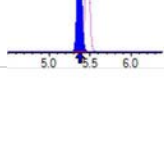
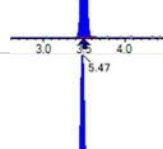
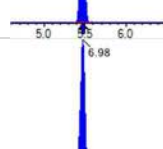
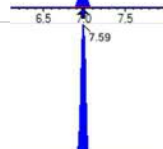

Lab Sample ID	Analyte	True	Found	%R	Units	Control Limit
SC00636-CCV4	ADONA	9.45	8.85	93.6	ng/mL	+/- 30.00%
	PFEESA	8.90	8.86	99.6	ng/mL	+/- 30.00%
	PFMPA	10.0	9.13	91.3	ng/mL	+/- 30.00%
	PFMBA	10.0	9.10	91.0	ng/mL	+/- 30.00%
	NFDHA	10.0	9.61	96.1	ng/mL	+/- 30.00%
	9CL-PF3ONS	9.35	7.60	81.3	ng/mL	+/- 30.00%
	11CL-PF3OUDS	9.45	8.53	90.3	ng/mL	+/- 30.00%
	3:3FTCA	20.0	19.7	98.6	ng/mL	+/- 30.00%
	5:3FTCA	20.0	18.3	91.3	ng/mL	+/- 30.00%
	7:3FTCA	20.0	18.5	92.6	ng/mL	+/- 30.00%

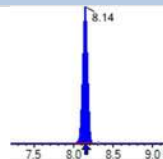
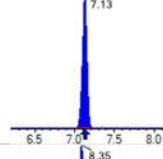
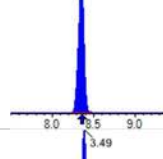
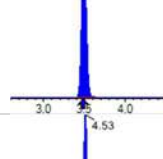
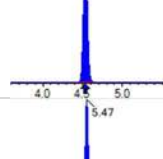
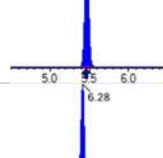
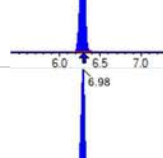
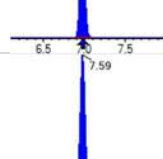
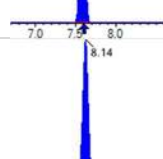
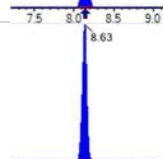
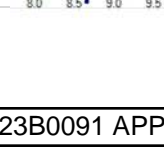
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	( 213.0 / 169.0 ) 5208514	( 3.49, 1.00 ) ( 0.00, N/A, 0.0 )	199.9	N/A 0.0 0.0	19.3379 [ 20.0000 ]	96.7%			
PFPeA	( 263.0 / 219.0 ) 5214775 ( 263.0 / 69.0 ) 62072	( 4.54, 1.00 ) ( 0.00, N/A, 0.2 )	7126.3 480.6	0.0119 111.2 108.0	9.3375 [ 10.0000 ]	93.4%			
PFHxA	( 313.0 / 269.0 ) 3246380 ( 313.0 / 119.0 ) 302957	( 5.47, 1.00 ) ( 0.00, N/A, 0.1 )	7318.4 538137.9	0.0933 93.7 98.1	4.7076 [ 5.0000 ]	94.2%			
PFHpA	( 363.0 / 319.0 ) 2607370 ( 363.0 / 169.0 ) 827512	( 6.28, 1.00 ) ( 0.00, N/A, 0.0 )	14769.4 39080.5	0.3174 99.0 99.3	4.6774 [ 5.0000 ]	93.5%			
PFOA	( 413.0 / 369.0 ) 3486331 ( 413.0 / 169.0 ) 1129429	( 6.98, 1.00 ) ( 0.00, N/A, 0.0 )	36079.6 23411.8	0.3240 96.8 89.7	5.0494 [ 5.0000 ]	101.0%			
PFNA	( 463.0 / 419.0 ) 3098893 ( 463.0 / 169.0 ) 686559	( 7.59, 1.00 ) ( 0.01, N/A, 0.0 )	12438.7 2433.6	0.2215 99.0 100.2	4.9389 [ 5.0000 ]	98.8%			
PFDA	( 513.0 / 469.0 ) 3589477 ( 513.0 / 169.0 ) 419013	( 8.15, 1.00 ) ( 0.00, N/A, 0.0 )	1867.3 805.7	0.1167 96.0 107.6	5.3763 [ 5.0000 ]	107.5%			
PFUnA	( 563.0 / 519.0 ) 3098400 ( 563.0 / 169.0 ) 392269	( 8.63, 1.00 ) ( 0.00, N/A, -0.2 )	2057.4 3539.2	0.1266 120.8 108.3	4.5940 [ 5.0000 ]	91.9%			
PFDoA	( 613.0 / 569.0 ) 2670839 ( 613.0 / 169.0 ) 472563	( 8.92, 1.00 ) ( 0.00, N/A, -0.1 )	2902.0 2179.6	0.1769 108.2 105.4	4.2878 [ 5.0000 ]	85.8%			
PFTTrDA	( 663.0 / 619.0 ) 2518158 ( 663.0 / 169.0 ) 623139	( 9.12, 1.02 ) ( N/A, 0.00, 0.1 )	1940.2 1565.0	0.2475 103.5 91.9	4.4651 [ 5.0000 ]	89.3%			
PFTeDA	( 713.0 / 669.0 ) 2564062 ( 713.0 / 169.0 ) 556382	( 9.29, 1.00 ) ( 0.00, N/A, 0.0 )	2415.7 1070.3	0.2170 105.0 105.0	4.7913 [ 5.0000 ]	95.8%			

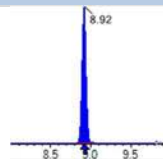
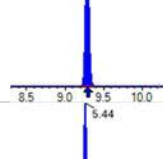
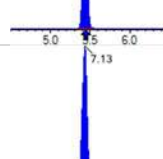
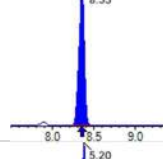
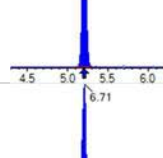
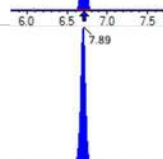
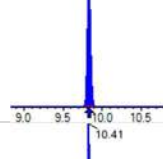
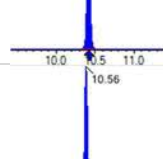
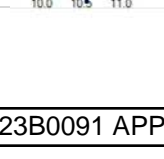
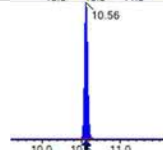
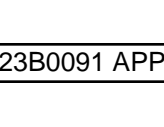
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	( 299.0 / 80.0 ) 5741144 ( 299.0 / 99.0 ) 3586677	( 5.44 , 1.00 ) ( 0.00 , N/A , 0.0 )	2974.5 3992.4	0.6247 96.6 101.0	4.1845 [ 4.4237 ]	94.6%			
PFPeS	( 349.0 / 80.0 ) 10027313 ( 349.0 / 99.0 ) 3304216	( 6.36 , 0.89 ) ( N/A , -0.01 , 0.0 )	24915.1 29207271.2	0.3295 92.3 95.7	4.6930 [ 4.6919 ]	100.0%			
PFHxS	( 399.0 / 80.0 ) 7504463 ( 399.0 / 99.0 ) 2473609	( 7.13 , 1.00 ) ( 0.00 , N/A , 0.0 )	19406.2 1680.2	0.3296 101.5 101.2	4.4050 [ 4.5549 ]	96.7%			
PFHpS	( 449.0 / 80.0 ) 8871524 ( 449.0 / 99.0 ) 2421268	( 7.77 , 0.93 ) ( N/A , 0.00 , 0.0 )	54846.5 1229505.8	0.2729 100.3 98.1	4.8498 [ 4.7570 ]	101.9%			
PFOS	( 499.0 / 80.0 ) 11050055 ( 499.0 / 99.0 ) 2469094	( 8.35 , 1.00 ) ( 0.00 , N/A , 0.1 )	1327.2 2936.2	0.2234 101.0 105.4	4.3649 [ 4.6375 ]	94.1%			
PFNS	( 549.0 / 80.0 ) 13397659 ( 549.0 / 99.0 ) 2958489	( 8.77 , 1.05 ) ( N/A , 0.00 , 0.0 )	14713.0 2015670.7	0.2208 100.7 96.0	4.7090 [ 4.7994 ]	98.1%			
PFDS	( 599.0 / 80.0 ) 15199925 ( 599.0 / 99.0 ) 3168202	( 9.02 , 1.08 ) ( N/A , 0.00 , -0.1 )	4164.2 2666.0	0.2084 95.6 97.1	4.9113 [ 4.8155 ]	102.0%			
PFDoS	( 699.0 / 80.0 ) 9958501 ( 699.0 / 99.0 ) 2093196	( 9.36 , 1.12 ) ( N/A , 0.00 , 0.0 )	3252.9 2373.8	0.2102 101.0 98.0	5.0451 [ 4.8478 ]	104.1%			
4:2FTS	( 327.0 / 307.0 ) 5420520 ( 327.0 / 81.0 ) 3590150	( 5.20 , 1.00 ) ( 0.00 , N/A , -0.1 )	3496.1 2941.3	0.6623 98.2 96.6	16.5026 [ 18.6906 ]	88.3%			
6:2FTS	( 427.0 / 407.0 ) 3272243 ( 427.0 / 81.0 ) 2584014	( 6.71 , 1.00 ) ( 0.00 , N/A , 0.1 )	2447.0 2665.9	0.7897 95.4 102.2	17.3483 [ 18.9808 ]	91.4%			
8:2FTS	( 527.0 / 507.0 ) 3447074 ( 527.0 / 81.0 ) 2906887	( 7.88 , 1.00 ) ( 0.00 , N/A , -0.2 )	2152.6 3163.4	0.8433 101.5 109.0	18.1687 [ 19.1658 ]	94.8%			

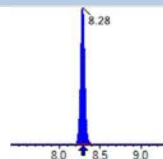
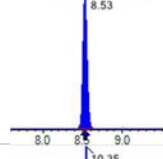
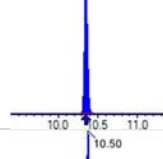
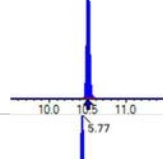
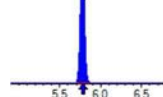
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	( 498.0 / 78.0 ) 17126512 ( 498.0 / 478.0 ) 337736	( 9.83 , 1.00 ) ( 0.00 , N/A , 0.0 )	3408.9 1887.1	0.0197 96.0 93.3	4.9761 [ 5.0000 ]	99.5%			
NMeFOSA	( 512.0 / 219.0 ) 14250142 ( 512.0 / 169.0 ) 11106428	( 10.42 , 1.00 ) ( 0.00 , N/A , 0.9 )	3981.1 2964.4	0.7794 95.1 95.2	19.7563 [ 20.0000 ]	98.8%			
NEIFOSA	( 526.0 / 219.0 ) 15126651 ( 526.0 / 169.0 ) 19297226	( 10.56 , 1.00 ) ( 0.00 , N/A , 0.9 )	15985.8 9835.3	1.2757 100.5 102.8	20.7487 [ 20.0000 ]	103.7%			
NMeFOSAA	( 570.0 / 419.0 ) 1249319 ( 570.0 / 483.0 ) 584209	( 8.29 , 1.00 ) ( 0.00 , N/A , 0.0 )	4153.7 48384.2	0.4676 98.5 100.9	4.6485 [ 5.0000 ]	93.0%			
NEIFOSAA	( 584.0 / 419.0 ) 1044838 ( 584.0 / 526.0 ) 630175	( 8.53 , 1.00 ) ( 0.01 , N/A , -0.1 )	2395.8 1607.2	0.6031 118.2 102.9	4.3149 [ 5.0000 ]	86.3%			
NMeFOSE	( 616.0 / 59.0 ) 5758078	( 10.36 , 1.00 ) ( 0.01 , N/A , 0.0 )	2589.9	N/A 0.0 0.0	18.0739 [ 20.0000 ]	90.4%			
NEtFOSE	( 630.0 / 59.0 ) 7789697	( 10.52 , 1.00 ) ( 0.01 , N/A , 0.0 )	1856.9	N/A 0.0 0.0	19.7106 [ 20.0000 ]	98.6%			
HFPO-DA	( 285.0 / 169.0 ) 3519335 ( 285.0 / 185.0 ) 10525891	( 5.78 , 1.00 ) ( 0.00 , N/A , 0.1 )	2190.3 3783.4	2.9909 108.6 107.8	8.3873 [ 10.0000 ]	83.9%			
ADONA	( 377.0 / 85.0 ) 12764659 ( 377.0 / 251.0 ) 1177712	( 6.56 , 1.14 ) ( N/A , -0.01 , 0.0 )	4167.5 2004.0	0.0923 104.6 98.8	8.8458 [ 9.4270 ]	93.8%			
9CI-Pf3ONS	( 531.0 / 351.0 ) 33090876 ( 533.0 / 353.0 ) 11565356	( 8.70 , 1.51 ) ( N/A , 0.00 , 0.0 )	4114.6 3843.4	0.3495 100.4 105.8	7.6050 [ 9.3325 ]	81.5%			
11CI-PF3OUDS	( 631.0 / 451.0 ) 20060490 ( 633.0 / 453.0 ) 7241534	( 9.17 , 1.59 ) ( N/A , 0.00 , 0.0 )	2368.5 1951.0	0.3610 99.8 101.3	8.5335 [ 9.4321 ]	90.5%			



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
3:3FTCA	( 241.0 / 177.0 ) 272637 ( 241.0 / 117.0 ) 430955	( 4.05 , 0.89 ) ( N/A , 0.00 , 0.0 )	1245.4 1478.6	1.5807 100.2 96.6	19.7151 [ 20.0000 ]	98.6%			
5:3FTCA	( 341.0 / 236.7 ) 1722476 ( 341.0 / 217.0 ) 3232101	( 6.01 , 1.10 ) ( N/A , -0.03 , 0.0 )	2173.8 2527.8	1.8764 110.0 116.6	18.2657 [ 20.0000 ]	91.3%			
7:3FTCA	( 441.0 / 317.0 ) 3227402 ( 441.0 / 337.0 ) 2821278	( 7.86 , 1.44 ) ( N/A , -0.02 , 0.0 )	1190.5 1677.8	0.8742 101.6 107.5	18.5265 [ 20.0000 ]	92.6%			
PFEESA	( 315.0 / 135.0 ) 7342675 ( 315.0 / 83.0 ) 2140583	( 5.88 , 1.08 ) ( N/A , -0.01 , 0.1 )	4188.6 2908.1	0.2915 114.1 105.8	8.8638 [ 8.9246 ]	99.3%			
PFMPA	( 229.0 / 85.0 ) 1113374	( 3.89 , 0.86 ) ( N/A , 0.00 , 0.0 )	4889.6	N/A 0.0 0.0	9.1271 [ 10.0000 ]	91.3%			
PFMBA	( 279.0 / 85.0 ) 4655793	( 4.84 , 1.07 ) ( N/A , -0.01 , 0.0 )	5744.0	N/A 0.0 0.0	9.1011 [ 10.0000 ]	91.0%			
NFDHA	( 295.0 / 201.0 ) 3679232 ( 295.0 / 85.0 ) 3890217	( 5.37 , 0.98 ) ( N/A , -0.01 , 0.0 )	4318.8 3377.1	1.0573 105.6 100.0	9.6124 [ 10.0000 ]	96.1%			
13C3_PFBA_IIS	( 216.0 / 172.0 ) 236097	( 3.50 , N/A ) ( N/A , 0.00 , N/A )	993.1	N/A	1.0540 [ 1.0000 ]	105.4% { 107.6% }			
13C2_PFHxA_IIS	( 315.0 / 270.0 ) 499353	( 5.47 , N/A ) ( N/A , -0.01 , N/A )	3097.2	N/A	0.9892 [ 1.0000 ]	98.9% { 103.9% }			
13C4_PFOA_IIS	( 417.0 / 372.0 ) 648063	( 6.98 , N/A ) ( N/A , -0.01 , N/A )	2626.6	N/A	1.0225 [ 1.0000 ]	102.3% { 106.2% }			
13C5_PFNA_IIS	( 468.0 / 423.0 ) 589008	( 7.59 , N/A ) ( N/A , 0.00 , N/A )	1191.8	N/A	0.9747 [ 1.0000 ]	97.5% { 104.4% }			

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 ) 614149	( 8.14 , N/A ) ( N/A , 0.00 , N/A )	2826.3	N/A	1.0614 [ 1.0000 ]	106.1% { 114.8% }			
18O2_PFHxS_IIS	( 403.0 / 83.9 ) 1001645	( 7.13 , N/A ) ( N/A , -0.01 , N/A )	2308.5	N/A	1.1696 [ 1.0000 ]	117.0% { 119.0% }			
13C4_PFOS_IIS	( 503.0 / 79.9 ) 1580181	( 8.35 , N/A ) ( N/A , 0.00 , N/A )	1107.0	N/A	1.0497 [ 1.0000 ]	105.0% { 117.8% }			
13C4_PFBA_EIS	( 217.0 / 172.0 ) 2601244	( 3.49 , N/A ) ( N/A , 0.00 , N/A )	5650.4	N/A	9.2625 [ 8.0000 ]	115.8% { 113.2% }			
13C5_PFPeA_EIS	( 268.0 / 223.0 ) 2715203	( 4.53 , N/A ) ( N/A , -0.01 , N/A )	4745.1	N/A	4.9696 [ 4.0000 ]	124.2% { 117.8% }			
13C5_PFHxA_EIS	( 318.0 / 273.0 ) 1568808	( 5.47 , N/A ) ( N/A , -0.01 , N/A )	3567.3	N/A	2.2936 [ 2.0000 ]	114.7% { 109.9% }			
13C4_PFHpA_EIS	( 367.0 / 322.0 ) 1320288	( 6.28 , N/A ) ( N/A , -0.01 , N/A )	3350.6	N/A	2.2278 [ 2.0000 ]	111.4% { 112.1% }			
13C8_PFOA_EIS	( 421.0 / 376.0 ) 1550892	( 6.98 , N/A ) ( N/A , 0.00 , N/A )	2590.7	N/A	2.1131 [ 2.0000 ]	105.7% { 109.2% }			
13C9_PFNA_EIS	( 472.0 / 427.0 ) 709160	( 7.59 , N/A ) ( N/A , -0.01 , N/A )	8807329.6	N/A	1.1363 [ 1.0000 ]	113.6% { 116.5% }			
13C6_PFDA_EIS	( 519.0 / 474.0 ) 759008	( 8.14 , N/A ) ( N/A , 0.00 , N/A )	1538.7	N/A	0.9802 [ 1.0000 ]	98.0% { 101.8% }			
13C7_PFUnA_EIS	( 570.0 / 525.0 ) 811762	( 8.63 , N/A ) ( N/A , 0.00 , N/A )	3387.1	N/A	1.0542 [ 1.0000 ]	105.4% { 121.8% }			

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 ) 708335	( 8.92, N/A ) ( N/A, 0.00, N/A )	1277.8	N/A	1.0726 [ 1.0000 ]	107.3% { 107.6% }			
13C2_PFTeDA_EIS	( 715.0 / 670.0 ) 618422	( 9.29, N/A ) ( N/A, 0.00, N/A )	1364.5	N/A	1.0219 [ 1.0000 ]	102.2% { 105.6% }			
13C3_PFBs_EIS	( 302.0 / 80.0 ) 4415445	( 5.44, N/A ) ( N/A, -0.01, N/A )	3168.8	N/A	2.1062 [ 2.0000 ]	105.3% { 116.4% }			
13C3_PFHxS_EIS	( 402.0 / 80.0 ) 2215814	( 7.13, N/A ) ( N/A, -0.01, N/A )	1419.8	N/A	2.0350 [ 2.0000 ]	101.7% { 112.7% }			
13C8_PFOS_EIS	( 507.0 / 80.0 ) 4801577	( 8.35, N/A ) ( N/A, 0.00, N/A )	1635.3	N/A	2.2134 [ 2.0000 ]	110.7% { 113.3% }			
13C2_4:2FTS_EIS	( 329.0 / 81.0 ) 481058	( 5.20, N/A ) ( N/A, -0.01, N/A )	967.3	N/A	4.1573 [ 4.0000 ]	103.9% { 138.5% }			
13C2_6:2FTS_EIS	( 429.0 / 81.0 ) 531735	( 6.71, N/A ) ( N/A, -0.01, N/A )	2590.3	N/A	4.1863 [ 4.0000 ]	104.7% { 127.8% }			
13C2_8:2FTS_EIS	( 529.0 / 81.0 ) 696719	( 7.89, N/A ) ( N/A, 0.00, N/A )	1552.8	N/A	3.8825 [ 4.0000 ]	97.1% { 115.5% }			
13C8_PFOsa_EIS	( 506.0 / 78.0 ) 8606416	( 9.83, N/A ) ( N/A, 0.00, N/A )	2851.3	N/A	2.1372 [ 2.0000 ]	106.9% { 103.6% }			
D3_NMeFOSA_EIS	( 515.0 / 169.0 ) 1737678	( 10.41, N/A ) ( N/A, 0.00, N/A )	2341.7	N/A	1.8316 [ 2.0000 ]	91.6% { 89.8% }			
D5_NEtFOSA_EIS	( 531.0 / 169.0 ) 1560727	( 10.56, N/A ) ( N/A, 0.00, N/A )	3624.2	N/A	1.8384 [ 2.0000 ]	91.9% { 87.5% }			

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 ) 1369876	( 8.28 , N/A ) ( N/A , 0.00 , N/A )	2007.2	N/A	3.9021 [ 4.0000 ]	97.6% { 103.4% }			
D5_EtFOSAA_EIS	( 589.0 / 419.0 ) 1153212	( 8.53 , N/A ) ( N/A , 0.00 , N/A )	4714.1	N/A	4.0118 [ 4.0000 ]	100.3% { 109.3% }			
D7_NMeFOSE_EIS	( 623.0 / 58.9 ) 6373376	( 10.35 , N/A ) ( N/A , 0.00 , N/A )	2432.2	N/A	18.4264 [ 20.0000 ]	92.1% { 85.8% }			
D9_NEtFOSE_EIS	( 639.0 / 58.9 ) 8305805	( 10.50 , N/A ) ( N/A , 0.00 , N/A )	2243.7	N/A	18.7512 [ 20.0000 ]	93.8% { 88.2% }			
13C3_HFPODA_EIS	( 287.0 / 169.0 ) 4244837	( 5.77 , N/A ) ( N/A , -0.01 , N/A )	4379.3	N/A	10.1928 [ 8.0000 ]	127.4% { 118.4% }			

## ANALYSIS SEQUENCE BLANKS

Laboratory: APPL, LLC  
 Client: AECOM  
 Sequence: SC00636  
 Calibration: 2307007

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

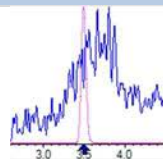
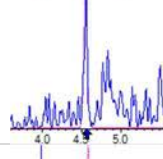
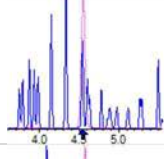
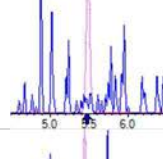
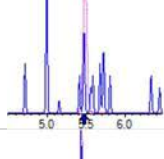
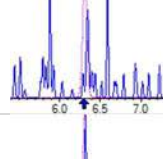
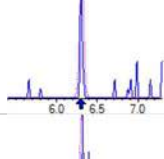
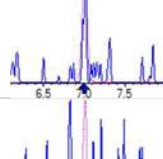
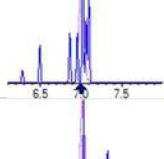
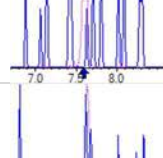
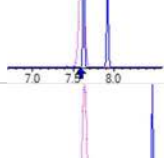
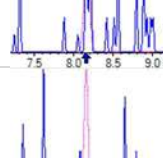
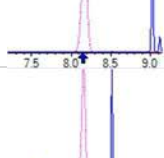
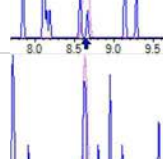
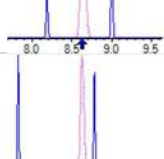
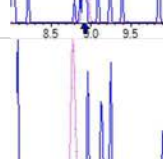
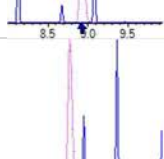
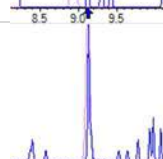
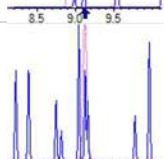
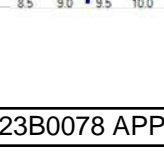
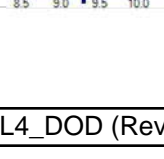
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	PFPEA	0.00	ng/mL	0.20	U
	PFHXA	0.00	ng/mL	0.10	U
	PFHPA	0.00	ng/mL	0.10	U
	PFOA	0.00	ng/mL	0.10	U
	PFNA	0.00	ng/mL	0.10	U
	PFDA	0.00	ng/mL	0.10	U
	PFUnA	0.00	ng/mL	0.10	U
	PFDOA	0.00	ng/mL	0.10	U
	PFTRDA	0.00	ng/mL	0.10	U
	PFTEDA	0.00	ng/mL	0.10	U
	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	0.0748	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
	ADONA	0.00	ng/mL	0.20	U
	PFEESA	0.00	ng/mL	0.20	U
	PFMPA	0.00	ng/mL	0.20	U

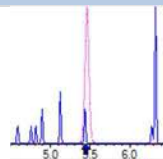
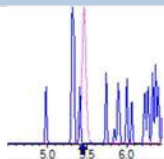
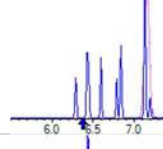
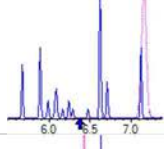
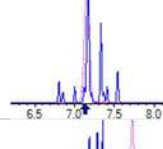
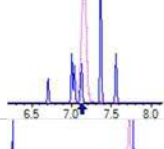
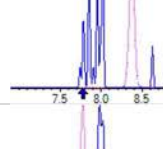
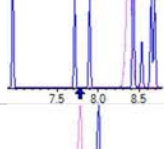
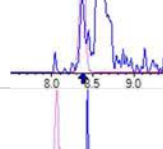
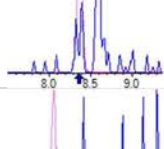
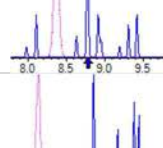
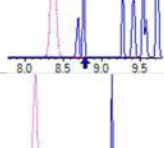
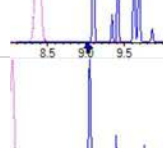
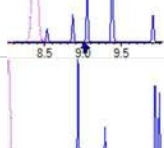
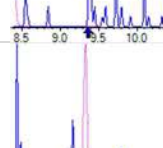
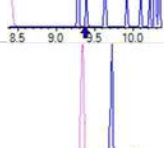
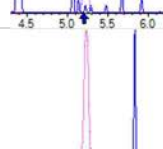
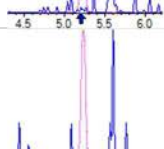
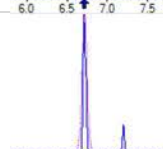
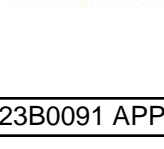
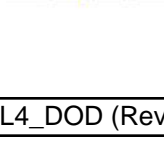
## ANALYSIS SEQUENCE BLANKS

Laboratory: APPL, LLC  
 Client: AECOM  
 Sequence: SC00636  
 Calibration: 2307007

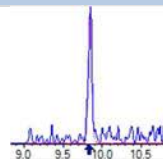
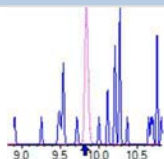
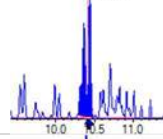
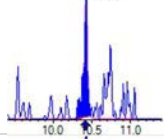
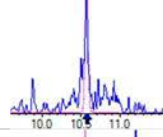
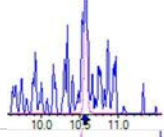
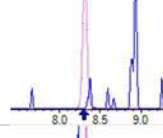
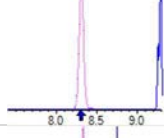
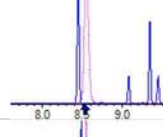
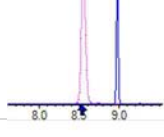
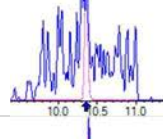
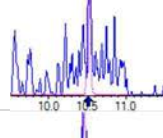
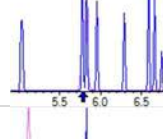
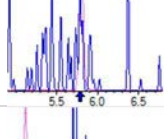
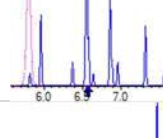
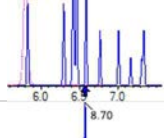
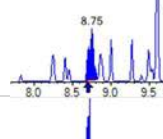
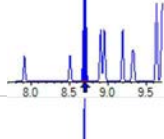
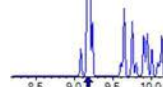
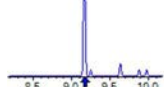
SDG:  
 Project: Red Hill AFFF Assessment Sam  
 Instrument: Saphira

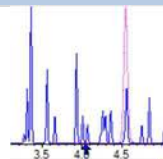
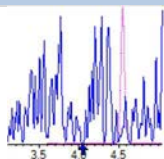
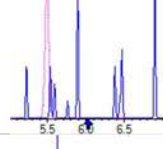
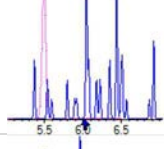
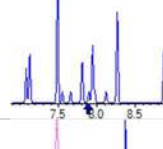
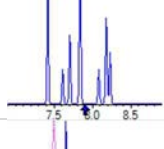
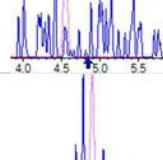
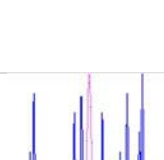
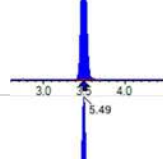
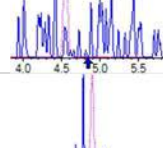
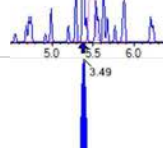
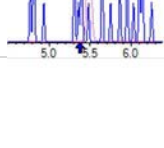
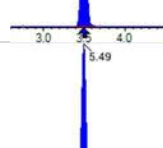
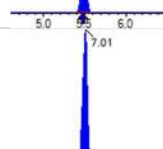
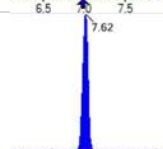
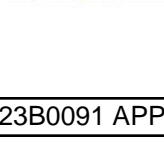
Lab Sample ID	Analyte	Found	Units	RL	C
SC00636-CCB1	PFMBA	0.00	ng/mL	0.20	U
	NFDHA	0.00	ng/mL	0.20	U
	9CL-PF3ONS	0.0520	ng/mL	0.20	U
	11CL-PF3OUDS	0.00	ng/mL	0.20	U
	3:3FTCA	0.00	ng/mL	0.40	U
	5:3FTCA	0.00	ng/mL	0.40	U
	7:3FTCA	0.00	ng/mL	0.40	U
	13C4-PFBA	7.17	ng/mL		
	13C5-PFPEA	3.89	ng/mL		
	13C5-PFHXA	1.76	ng/mL		
	13C4-PFHPA	1.93	ng/mL		
	13C8-PFOA	1.65	ng/mL		
	13C9-PFNA	0.789	ng/mL		
	13C6-PFDA	0.846	ng/mL		
	13C7-PFUnA	0.797	ng/mL		
	13C2-PFDOA	0.905	ng/mL		
	13C2-PFTEDA	0.835	ng/mL		
	13C3-PFBS	1.75	ng/mL		
	13C3-PFHXS	1.66	ng/mL		
	13C8-PFOS	1.76	ng/mL		
	13C2-4:2FTS	3.02	ng/mL		
	13C2-6:2FTS	3.07	ng/mL		
	13C2-8:2FTS	3.22	ng/mL		
	13C8-PFOSA	1.88	ng/mL		
	D3-NMEFOSA	1.84	ng/mL		
	D5-NETFOSA	2.07	ng/mL		
	D3-NMEFOSAA	3.35	ng/mL		
	D5-NETFOSAA	3.18	ng/mL		
	D7-NMEFOSE	19.5	ng/mL		
	D9-NETFOSSE	20.4	ng/mL		
	13C3-HFPO-DA	7.15	ng/mL		

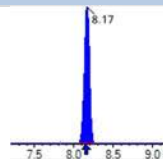
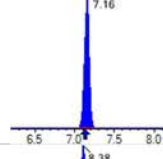
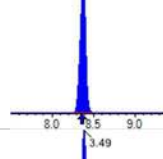
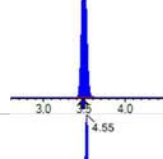
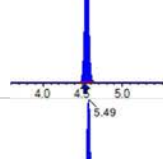
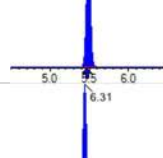
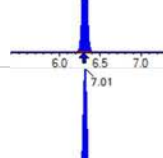
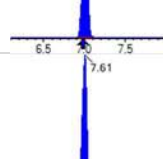
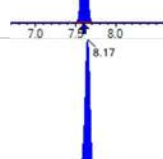
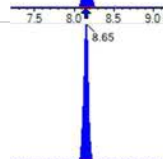
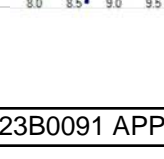
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-Imin, ΔRT-CVmin, ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration True ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	( 213.0 / 169.0 ) N/A	( N/A , N/A ) ( N/A , N/A , N/A )	N/A	N/A 0.0 0.0	0.0000	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			

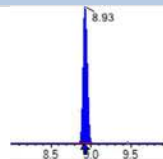
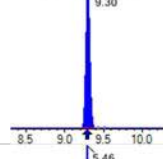
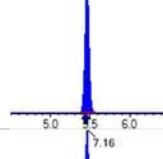
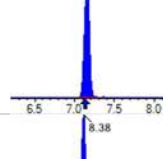
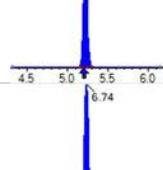
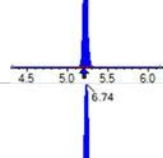
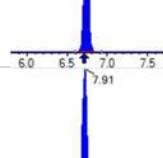
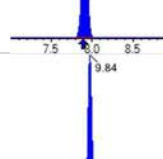
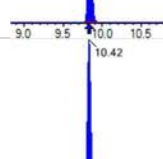
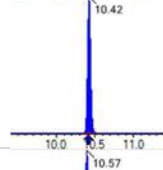
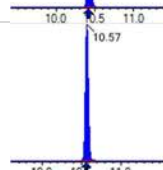
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-Imin, ΔRT-CVmin, ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration True ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	(299.0 / 80.0) N/A (299.0 / 99.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 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			

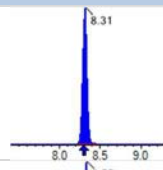
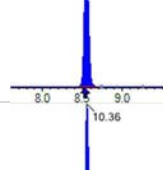
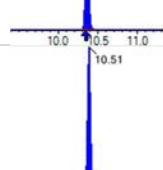




Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-Imin, ΔRT-CVmin, ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration True ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	(498.0 / 78.0) N/A (498.0 / 478.0) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A N/A	N/A 0.0 0.0	0.0000	N/A			
NMeFOSA	(512.0 / 219.0) 5765 (512.0 / 169.0) 3650	(10.42, 1.00) (0.00, N/A, -0.1)	46.2 44.2	0.6331 77.3 77.3	0.0748	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) N/A	(N/A, N/A) (N/A, N/A, N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
NEiFOSE	(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) 4747 (533.0 / 353.0) 1424	(8.75, 1.51) (N/A, 0.05, 2.7)	12.8 38.3	0.2999 86.1 90.8	0.0520	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) 271279	(3.49, N/A) (N/A, -0.01, N/A)	1279.6	N/A	1.2111 [ 1.0000 ]	121.1% { 123.6% }			
13C2_PFHxA_IIS	(315.0 / 270.0) 539694	(5.49, N/A) (N/A, 0.01, N/A)	1915.4	N/A	1.0691 [ 1.0000 ]	106.9% { 112.3% }			
13C4_PFOA_IIS	(417.0 / 372.0) 730884	(7.01, N/A) (N/A, 0.02, N/A)	4731.3	N/A	1.1532 [ 1.0000 ]	115.3% { 119.7% }			
13C5_PFNA_IIS	(468.0 / 423.0) 699898	(7.62, N/A) (N/A, 0.02, N/A)	5068.9	N/A	1.1583 [ 1.0000 ]	115.8% { 124.1% }			

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
13C2_PFDA_IIS	( 515.0 / 470.1 ) 645021	( 8.17 , N/A ) ( N/A , 0.02 , N/A )	3390.1	N/A	1.1147 [ 1.0000 ]	111.5% { 120.5% }			
18O2_PFHxS_IIS	( 403.0 / 83.9 ) 973180	( 7.16 , N/A ) ( N/A , 0.02 , N/A )	1729.1	N/A	1.1364 [ 1.0000 ]	113.6% { 115.6% }			
13C4_PFOS_IIS	( 503.0 / 79.9 ) 1632075	( 8.38 , N/A ) ( N/A , 0.02 , N/A )	1195.5	N/A	1.0842 [ 1.0000 ]	108.4% { 121.7% }			
13C4_PFBA_EIS	( 217.0 / 172.0 ) 2313280	( 3.49 , N/A ) ( N/A , 0.00 , N/A )	6639.6	N/A	7.1689 [ 8.0000 ]	89.6% { 100.7% }			
13C5_PFPeA_EIS	( 268.0 / 223.0 ) 2297283	( 4.55 , N/A ) ( N/A , 0.01 , N/A )	3952.5	N/A	3.8904 [ 4.0000 ]	97.3% { 99.7% }			
13C5_PFHxA_EIS	( 318.0 / 273.0 ) 1301666	( 5.49 , N/A ) ( N/A , 0.01 , N/A )	2869.9	N/A	1.7608 [ 2.0000 ]	88.0% { 91.2% }			
13C4_PFHpA_EIS	( 367.0 / 322.0 ) 1238305	( 6.31 , N/A ) ( N/A , 0.02 , N/A )	2587.3	N/A	1.9333 [ 2.0000 ]	96.7% { 105.1% }			
13C8_PFOA_EIS	( 421.0 / 376.0 ) 1367134	( 7.01 , N/A ) ( N/A , 0.02 , N/A )	3407.5	N/A	1.6516 [ 2.0000 ]	82.6% { 96.3% }			
13C9_PFNA_EIS	( 472.0 / 427.0 ) 585422	( 7.61 , N/A ) ( N/A , 0.02 , N/A )	18410.6	N/A	0.7894 [ 1.0000 ]	78.9% { 96.2% }			
13C6_PFDA_EIS	( 519.0 / 474.0 ) 688105	( 8.17 , N/A ) ( N/A , 0.02 , N/A )	2098.6	N/A	0.8461 [ 1.0000 ]	84.6% { 92.3% }			
13C7_PFUnA_EIS	( 570.0 / 525.0 ) 644329	( 8.65 , N/A ) ( N/A , 0.02 , N/A )	2192.4	N/A	0.7967 [ 1.0000 ]	79.7% { 96.7% }			

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
13C2_PFDa_EIS	( 615.0 / 570.0 ) 627822	( 8.93 , N/A ) ( N/A , 0.01 , N/A )	1802.7	N/A	0.9051 [ 1.0000 ]	90.5% { 95.4% }			
13C2_PFTeDA_EIS	( 715.0 / 670.0 ) 530488	( 9.30 , N/A ) ( N/A , 0.01 , N/A )	1194.5	N/A	0.8347 [ 1.0000 ]	83.5% { 90.6% }			
13C3_PFBs_EIS	( 302.0 / 80.0 ) 3555448	( 5.46 , N/A ) ( N/A , 0.01 , N/A )	3107.4	N/A	1.7456 [ 2.0000 ]	87.3% { 93.8% }			
13C3_PFHxS_EIS	( 402.0 / 80.0 ) 1756343	( 7.16 , N/A ) ( N/A , 0.02 , N/A )	1639.4	N/A	1.6602 [ 2.0000 ]	83.0% { 89.4% }			
13C8_PFOS_EIS	( 507.0 / 80.0 ) 3954484	( 8.38 , N/A ) ( N/A , 0.02 , N/A )	1417.5	N/A	1.7649 [ 2.0000 ]	88.2% { 93.3% }			
13C2_4:2FTS_EIS	( 329.0 / 81.0 ) 339224	( 5.22 , N/A ) ( N/A , 0.01 , N/A )	1013.7	N/A	3.0173 [ 4.0000 ]	75.4% { 97.7% }			
13C2_6:2FTS_EIS	( 429.0 / 81.0 ) 378890	( 6.74 , N/A ) ( N/A , 0.02 , N/A )	1259.7	N/A	3.0702 [ 4.0000 ]	76.8% { 91.1% }			
13C2_8:2FTS_EIS	( 529.0 / 81.0 ) 560745	( 7.91 , N/A ) ( N/A , 0.02 , N/A )	2619.9	N/A	3.2162 [ 4.0000 ]	80.4% { 93.0% }			
13C8_PFOsa_EIS	( 506.0 / 78.0 ) 7831527	( 9.84 , N/A ) ( N/A , 0.01 , N/A )	4111.9	N/A	1.8830 [ 2.0000 ]	94.1% { 94.3% }			
D3_NMeFOsa_EIS	( 515.0 / 169.0 ) 1802752	( 10.42 , N/A ) ( N/A , 0.01 , N/A )	2246.9	N/A	1.8397 [ 2.0000 ]	92.0% { 93.1% }			
D5_NEtFOsa_EIS	( 531.0 / 169.0 ) 1811996	( 10.57 , N/A ) ( N/A , 0.01 , N/A )	3465.9	N/A	2.0665 [ 2.0000 ]	103.3% { 101.6% }			

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) 1215081	(8.31, N/A) (N/A, 0.02, N/A)	1423.5	N/A	3.3511 [4.0000]	83.8% {91.8%}			
D5_EtFOSAA_EIS	(589.0 / 419.0) 944941	(8.55, N/A) (N/A, 0.02, N/A)	12589.6	N/A	3.1827 [4.0000]	79.6% {89.6%}			
D7_NMeFOSE_EIS	(623.0 / 58.9) 6950962	(10.36, N/A) (N/A, 0.01, N/A)	2404.1	N/A	19.4573 [20.0000]	97.3% {93.6%}			
D9_NEtFOSE_EIS	(639.0 / 58.9) 9314191	(10.51, N/A) (N/A, 0.01, N/A)	1779.3	N/A	20.3591 [20.0000]	101.8% {98.9%}			
13C3_HFPODA_EIS	(287.0 / 169.0) 3217560	(5.80, N/A) (N/A, 0.01, N/A)	3609.3	N/A	7.1486 [8.0000]	89.4% {89.7%}			

## ANALYSIS SEQUENCE BLANKS

Laboratory: APPL, LLC  
 Client: AECOM  
 Sequence: SC00636  
 Calibration: 2307007

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

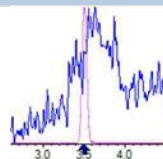
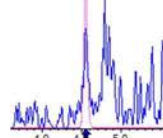
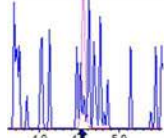
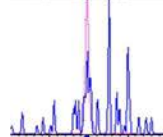
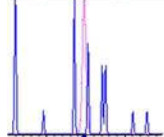
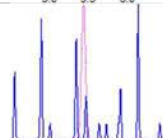
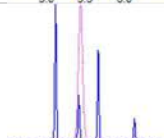
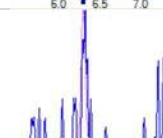
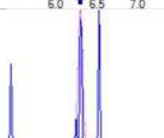
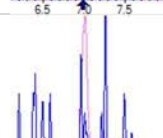
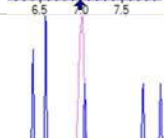
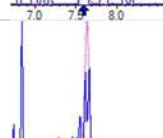
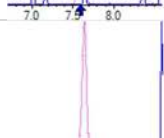
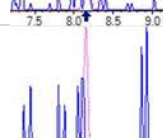
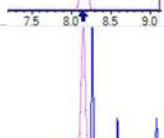
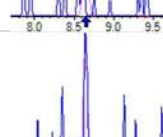
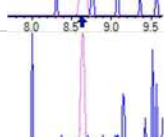
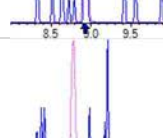
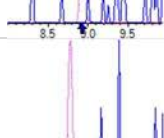
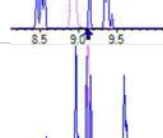
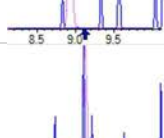
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	PFPEA	0.00	ng/mL	0.20	U
	PFHXA	0.00	ng/mL	0.10	U
	PFHPA	0.00	ng/mL	0.10	U
	PFOA	0.00	ng/mL	0.10	U
	PFNA	0.00	ng/mL	0.10	U
	PFDA	0.00	ng/mL	0.10	U
	PFUnA	0.00	ng/mL	0.10	U
	PFDOA	0.00	ng/mL	0.10	U
	PFTRDA	0.00	ng/mL	0.10	U
	PFTEDA	0.00	ng/mL	0.10	U
	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
	ADONA	0.00	ng/mL	0.20	U
	PFEESA	0.00	ng/mL	0.20	U
	PFMPA	0.00	ng/mL	0.20	U

## ANALYSIS SEQUENCE BLANKS

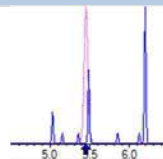
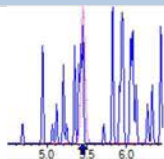
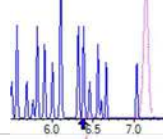
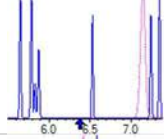
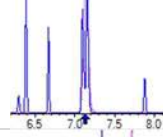
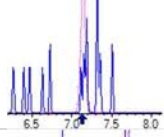
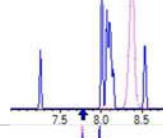
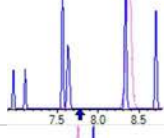
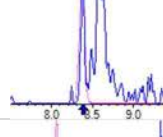
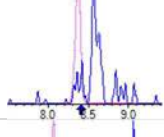
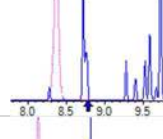
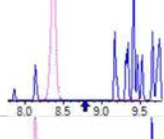
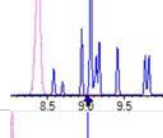
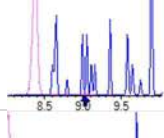
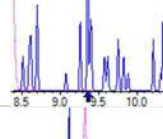
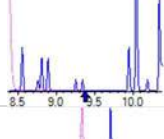
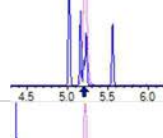
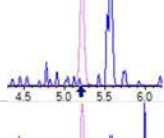
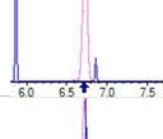
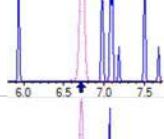
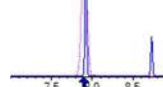
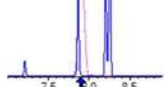
Laboratory: APPL, LLC  
 Client: AECOM  
 Sequence: SC00636  
 Calibration: 2307007

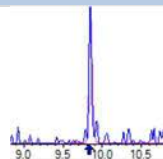
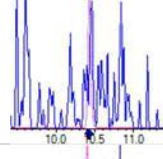
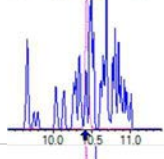
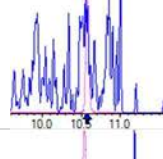
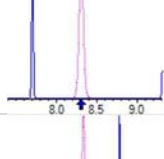
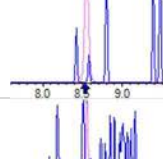
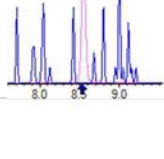
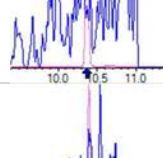
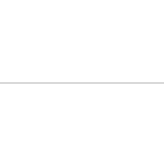
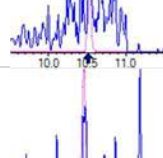
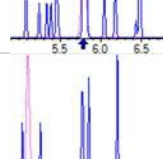
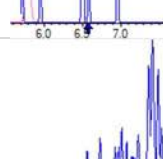
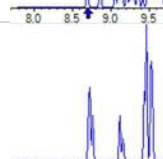
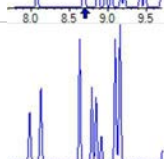
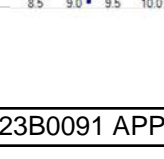
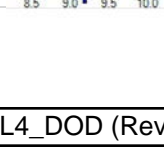
SDG:  
 Project: Red Hill AFFF Assessment Sam  
 Instrument: Saphira

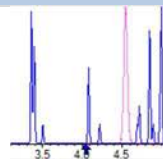
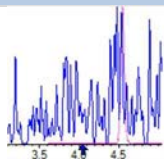
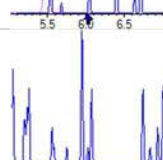
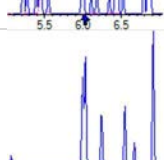
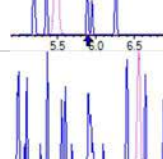
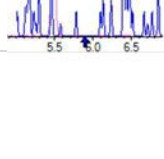
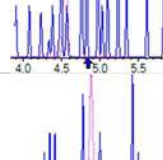
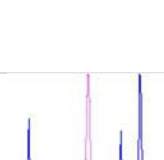
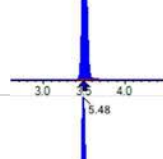
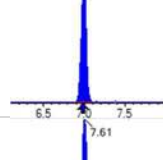
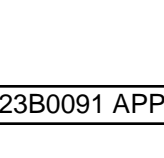
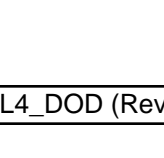
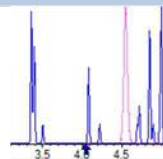
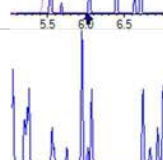
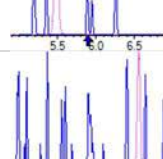
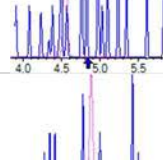
Lab Sample ID	Analyte	Found	Units	RL	C
SC00636-CCB3	PFMBA	0.00	ng/mL	0.20	U
	NFDHA	0.00	ng/mL	0.20	U
	9CL-PF3ONS	0.00	ng/mL	0.20	U
	11CL-PF3OUDS	0.00	ng/mL	0.20	U
	3:3FTCA	0.00	ng/mL	0.40	U
	5:3FTCA	0.00	ng/mL	0.40	U
	7:3FTCA	0.00	ng/mL	0.40	U
	13C4-PFBA	7.10	ng/mL		
	13C5-PFPEA	3.93	ng/mL		
	13C5-PFHXA	1.91	ng/mL		
	13C4-PFHPA	1.89	ng/mL		
	13C8-PFOA	1.68	ng/mL		
	13C9-PFNA	0.787	ng/mL		
	13C6-PFDA	0.833	ng/mL		
	13C7-PFUnA	0.890	ng/mL		
	13C2-PFDOA	0.904	ng/mL		
	13C2-PFTEDA	0.870	ng/mL		
	13C3-PFBS	1.77	ng/mL		
	13C3-PFHXS	1.71	ng/mL		
	13C8-PFOS	1.82	ng/mL		
	13C2-4:2FTS	3.51	ng/mL		
	13C2-6:2FTS	3.51	ng/mL		
	13C2-8:2FTS	3.13	ng/mL		
	13C8-PFOSA	1.30	ng/mL		
	D3-NMEFOSA	0.961	ng/mL		
	D5-NETFOSA	0.985	ng/mL		
	D3-NMEFOSAA	3.24	ng/mL		
	D5-NETFOSAA	3.63	ng/mL		
	D7-NMEFOSE	9.86	ng/mL		
	D9-NETFOSSE	9.66	ng/mL		
	13C3-HFPO-DA	7.67	ng/mL		

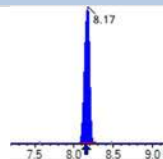
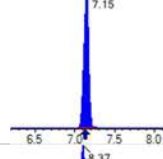
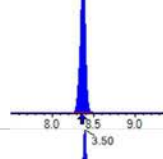
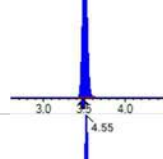
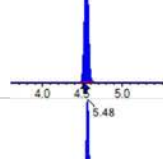
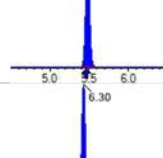
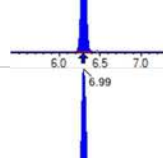
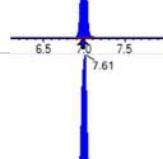
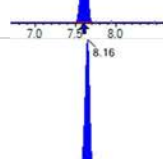
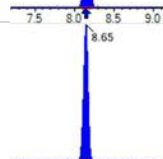
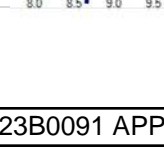
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	( 213.0 / 169.0 ) N/A	( N/A , N/A ) ( N/A , N/A , N/A )	N/A	N/A 0.0 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			
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			

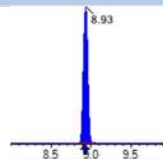
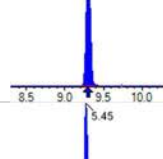
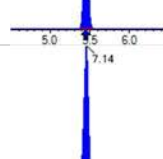
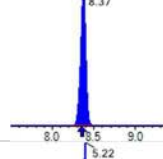
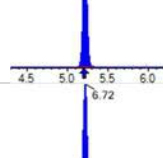
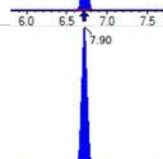
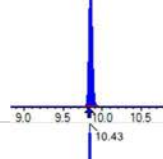
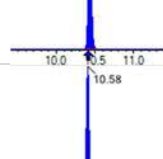
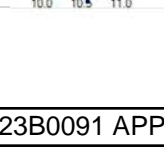
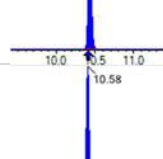
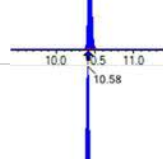


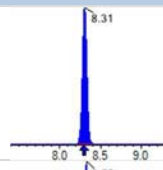




Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	( 299.0 / 80.0 ) N/A ( 299.0 / 99.0 ) N/A	( N/A , N/A ) ( N/A , N/A , 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 ) 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 ) N/A	( N/A , N/A ) ( N/A , N/A , N/A )	N/A	N/A 0.0 0.0	0.0000	N/A			
NEiFOSE	( 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 ) 293514	( 3.51 , N/A ) ( N/A , 0.01 , N/A )	1148.6	N/A	1.3104 [ 1.0000 ]	131.0% { 133.8% }			
13C2_PFHxA_IIS	( 315.0 / 270.0 ) 576625	( 5.48 , N/A ) ( N/A , 0.01 , N/A )	1925.2	N/A	1.1423 [ 1.0000 ]	114.2% { 120.0% }			
13C4_PFOA_IIS	( 417.0 / 372.0 ) 818665	( 6.99 , N/A ) ( N/A , 0.00 , N/A )	1411.5	N/A	1.2917 [ 1.0000 ]	129.2% { 134.1% }			
13C5_PFNA_IIS	( 468.0 / 423.0 ) 768451	( 7.61 , N/A ) ( N/A , 0.01 , N/A )	1526.8	N/A	1.2717 [ 1.0000 ]	127.2% { 136.3% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) ( $\Delta$ RT-I[min], $\Delta$ RT-CV[min], $\Delta$ 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 ) 751010	( 8.17 , N/A ) ( N/A , 0.02 , N/A )	1367.8	N/A	1.2979 [ 1.0000 ]	129.8% { 140.3% }			
18O2_PFHxS_IIS	( 403.0 / 83.9 ) 1073911	( 7.15 , N/A ) ( N/A , 0.01 , N/A )	2908.1	N/A	1.2540 [ 1.0000 ]	125.4% { 127.5% }			
13C4_PFOS_IIS	( 503.0 / 79.9 ) 1849713	( 8.37 , N/A ) ( N/A , 0.02 , N/A )	1518.3	N/A	1.2287 [ 1.0000 ]	122.9% { 137.9% }			
13C4_PFBA_EIS	( 217.0 / 172.0 ) 2479141	( 3.50 , N/A ) ( N/A , 0.01 , N/A )	5112.4	N/A	7.1009 [ 8.0000 ]	88.8% { 107.9% }			
13C5_PFPeA_EIS	( 268.0 / 223.0 ) 2480319	( 4.55 , N/A ) ( N/A , 0.01 , N/A )	4410.2	N/A	3.9314 [ 4.0000 ]	98.3% { 107.6% }			
13C5_PFHxA_EIS	( 318.0 / 273.0 ) 1509029	( 5.48 , N/A ) ( N/A , 0.00 , N/A )	3114.0	N/A	1.9106 [ 2.0000 ]	95.5% { 105.7% }			
13C4_PFHpA_EIS	( 367.0 / 322.0 ) 1294644	( 6.30 , N/A ) ( N/A , 0.01 , N/A )	2834.7	N/A	1.8918 [ 2.0000 ]	94.6% { 109.9% }			
13C8_PFOA_EIS	( 421.0 / 376.0 ) 1554163	( 6.99 , N/A ) ( N/A , 0.01 , N/A )	2078.6	N/A	1.6763 [ 2.0000 ]	83.8% { 109.4% }			
13C9_PFNA_EIS	( 472.0 / 427.0 ) 640443	( 7.61 , N/A ) ( N/A , 0.02 , N/A )	1322.1	N/A	0.7866 [ 1.0000 ]	78.7% { 105.2% }			
13C6_PFDA_EIS	( 519.0 / 474.0 ) 788598	( 8.16 , N/A ) ( N/A , 0.02 , N/A )	912.5	N/A	0.8328 [ 1.0000 ]	83.3% { 105.7% }			
13C7_PFUnA_EIS	( 570.0 / 525.0 ) 837943	( 8.65 , N/A ) ( N/A , 0.02 , N/A )	1679.1	N/A	0.8899 [ 1.0000 ]	89.0% { 125.8% }			

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 ) 730243	( 8.93 , N/A ) ( N/A , 0.01 , N/A )	2545.5	N/A	0.9042 [ 1.0000 ]	90.4% { 110.9% }			
13C2_PFTeDA_EIS	( 715.0 / 670.0 ) 643534	( 9.30 , N/A ) ( N/A , 0.01 , N/A )	1036.8	N/A	0.8696 [ 1.0000 ]	87.0% { 109.9% }			
13C3_PFBs_EIS	( 302.0 / 80.0 ) 3983495	( 5.45 , N/A ) ( N/A , 0.00 , N/A )	3892.6	N/A	1.7723 [ 2.0000 ]	88.6% { 105.0% }			
13C3_PFHxS_EIS	( 402.0 / 80.0 ) 1992365	( 7.14 , N/A ) ( N/A , 0.01 , N/A )	1743.7	N/A	1.7066 [ 2.0000 ]	85.3% { 101.4% }			
13C8_PFOS_EIS	( 507.0 / 80.0 ) 4622985	( 8.37 , N/A ) ( N/A , 0.02 , N/A )	1664.4	N/A	1.8205 [ 2.0000 ]	91.0% { 109.1% }			
13C2_4:2FTS_EIS	( 329.0 / 81.0 ) 435803	( 5.22 , N/A ) ( N/A , 0.01 , N/A )	1326.0	N/A	3.5128 [ 4.0000 ]	87.8% { 125.5% }			
13C2_6:2FTS_EIS	( 429.0 / 81.0 ) 478377	( 6.72 , N/A ) ( N/A , 0.01 , N/A )	1253.0	N/A	3.5128 [ 4.0000 ]	87.8% { 115.0% }			
13C2_8:2FTS_EIS	( 529.0 / 81.0 ) 602294	( 7.90 , N/A ) ( N/A , 0.01 , N/A )	1473.5	N/A	3.1304 [ 4.0000 ]	78.3% { 99.8% }			
13C8_PFOsa_EIS	( 506.0 / 78.0 ) 6104932	( 9.85 , N/A ) ( N/A , 0.02 , N/A )	4173.0	N/A	1.2951 [ 2.0000 ]	64.8% { 73.5% }			
D3_NMeFOSA_EIS	( 515.0 / 169.0 ) 1067792	( 10.43 , N/A ) ( N/A , 0.02 , N/A )	2250.7	N/A	0.9615 [ 2.0000 ]	48.1% { 55.2% }			
D5_NEtFOSA_EIS	( 531.0 / 169.0 ) 978412	( 10.58 , N/A ) ( N/A , 0.02 , N/A )	2150.6	N/A	0.9845 [ 2.0000 ]	49.2% { 54.9% }			

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 ) 1329708	( 8.31 , N/A ) ( N/A , 0.02 , N/A )	1258.5	N/A	3.2357 [ 4.0000 ]	80.9% { 100.4% }			
D5_EtFOSAA_EIS	( 589.0 / 419.0 ) 1222703	( 8.55 , N/A ) ( N/A , 0.02 , N/A )	6946.0	N/A	3.6337 [ 4.0000 ]	90.8% { 115.9% }			
D7_NMeFOSE_EIS	( 623.0 / 58.9 ) 3992437	( 10.37 , N/A ) ( N/A , 0.02 , N/A )	1855.9	N/A	9.8608 [ 20.0000 ]	49.3% { 53.8% }			
D9_NEtFOSE_EIS	( 639.0 / 58.9 ) 5009365	( 10.52 , N/A ) ( N/A , 0.02 , N/A )	1843.9	N/A	9.6612 [ 20.0000 ]	48.3% { 53.2% }			
13C3_HFPODA_EIS	( 287.0 / 169.0 ) 3688614	( 5.79 , N/A ) ( N/A , 0.01 , N/A )	3328.8	N/A	7.6703 [ 8.0000 ]	95.9% { 102.9% }			

## ANALYSIS SEQUENCE BLANKS

Laboratory: APPL, LLC  
 Client: AECOM  
 Sequence: SC00636  
 Calibration: 2307007

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

Lab Sample ID	Analyte	Found	Units	RL	C
SC00636-CCB4	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	0.00	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
	ADONA	0.00	ng/mL	0.20	U
	PFEESA	0.00	ng/mL	0.20	U
	PFMPA	0.00	ng/mL	0.20	U

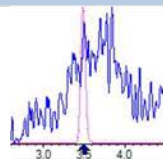
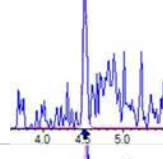
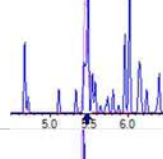
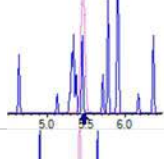
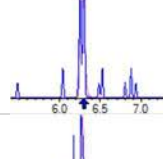
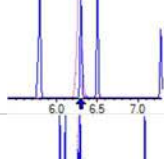
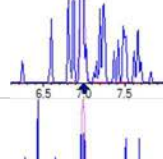
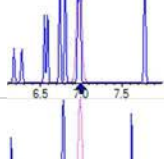
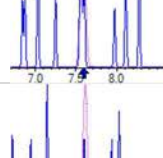
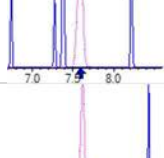
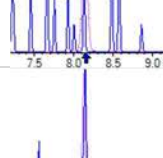
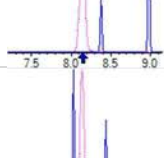
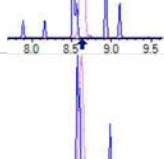
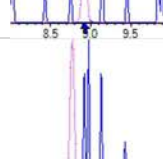
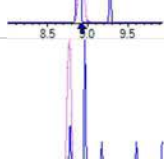
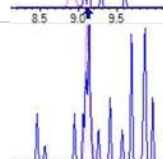
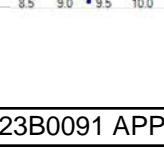
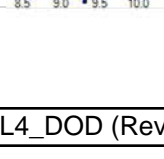
## ANALYSIS SEQUENCE BLANKS

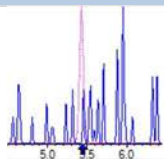
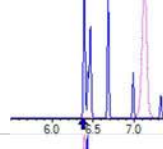
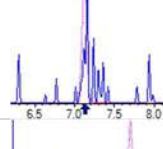
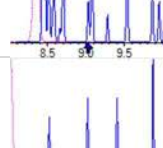
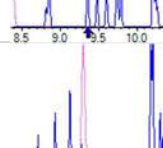
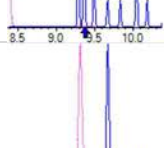
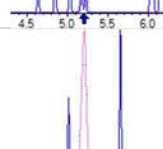
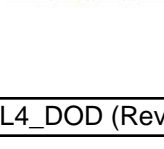
Laboratory: APPL, LLC  
 Client: AECOM  
 Sequence: SC00636  
 Calibration: 2307007

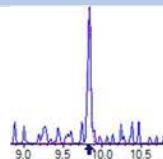
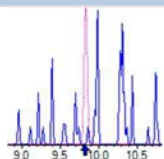
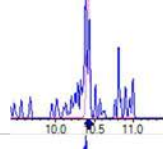
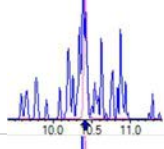
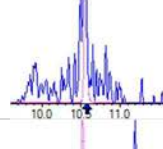
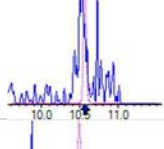
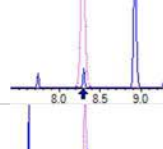
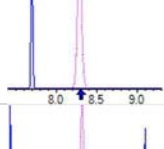
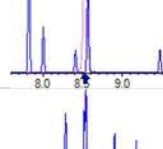
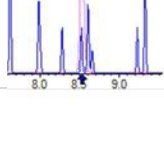
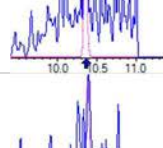
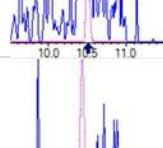
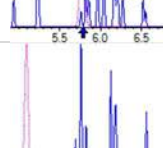
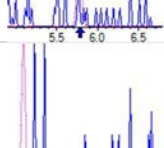
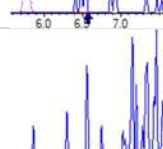
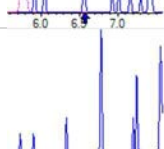
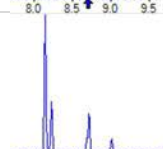
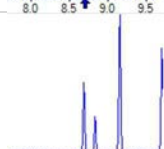
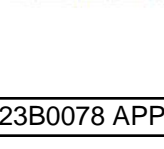
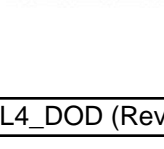
SDG:  
 Project: Red Hill AFFF Assessment Sam  
 Instrument: Saphira

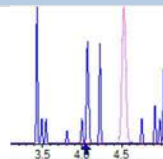
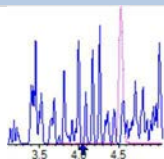
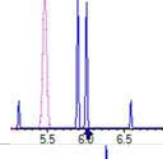
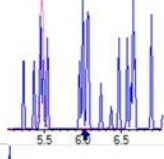
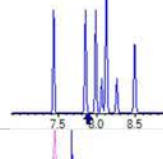
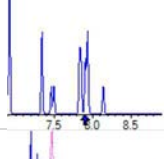
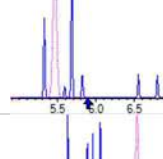
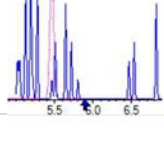
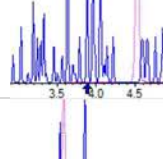
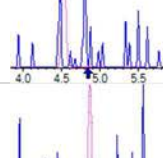
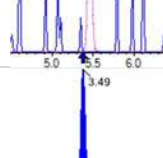
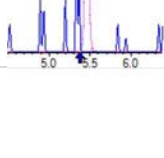
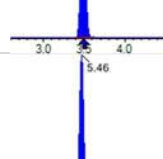
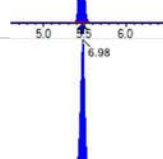
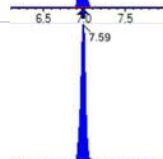

Lab Sample ID	Analyte	Found	Units	RL	C
SC00636-CCB4	PFMBA	0.00	ng/mL	0.20	U
	NFDHA	0.00	ng/mL	0.20	U
	9CL-PF3ONS	0.00	ng/mL	0.20	U
	11CL-PF3OUDS	0.00	ng/mL	0.20	U
	3:3FTCA	0.00	ng/mL	0.40	U
	5:3FTCA	0.00	ng/mL	0.40	U
	7:3FTCA	0.00	ng/mL	0.40	U
	13C4-PFBA	7.00	ng/mL		
	13C5-PFPEA	3.86	ng/mL		
	13C5-PFHXA	1.77	ng/mL		
	13C4-PFHPA	1.78	ng/mL		
	13C8-PFOA	1.80	ng/mL		
	13C9-PFNA	0.789	ng/mL		
	13C6-PFDA	0.941	ng/mL		
	13C7-PFUnA	0.920	ng/mL		
	13C2-PFDOA	0.951	ng/mL		
	13C2-PFTEDA	0.857	ng/mL		
	13C3-PFBS	1.72	ng/mL		
	13C3-PFHXS	1.80	ng/mL		
	13C8-PFOS	1.74	ng/mL		
	13C2-4:2FTS	3.06	ng/mL		
	13C2-6:2FTS	3.07	ng/mL		
	13C2-8:2FTS	2.91	ng/mL		
	13C8-PFOSA	1.72	ng/mL		
	D3-NMEFOSA	1.53	ng/mL		
	D5-NETFOSA	1.63	ng/mL		
	D3-NMEFOSAA	2.71	ng/mL		
	D5-NETFOSAA	2.99	ng/mL		
	D7-NMEFOSE	15.0	ng/mL		
	D9-NETFOSSE	16.1	ng/mL		
	13C3-HFPO-DA	7.41	ng/mL		

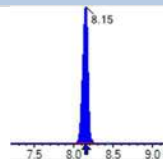
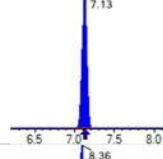
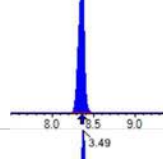
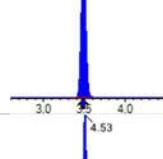
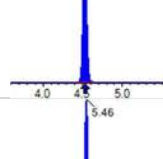
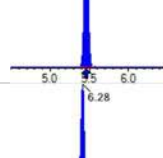
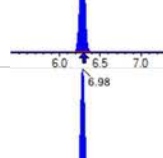
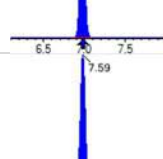
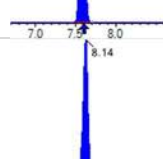
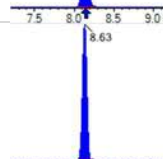
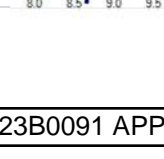


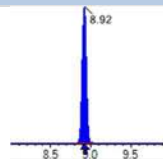
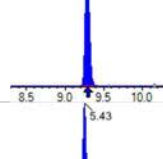
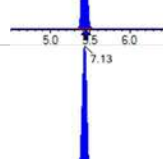
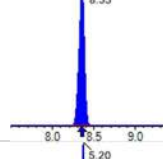
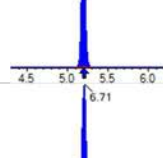
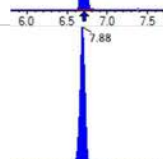
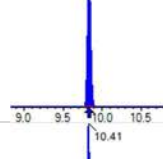
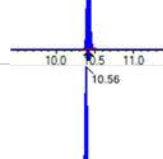
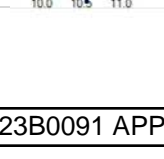
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	( 213.0 / 169.0 ) N/A	( N/A , N/A ) ( N/A , N/A , N/A )	N/A	N/A 0.0 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			
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			

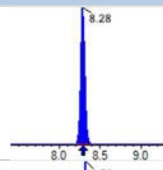
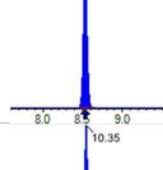
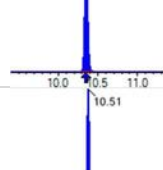
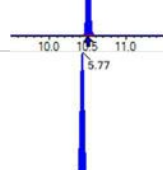

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	( 299.0 / 80.0 ) N/A ( 299.0 / 99.0 ) N/A	( N/A , N/A ) ( N/A , N/A , 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 ) 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 ) N/A	( N/A , N/A ) ( N/A , N/A , N/A )	N/A	N/A 0.0 0.0	0.0000	N/A			
NEiFOSE	( 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 ) 293954	( 3.49 , N/A ) ( N/A , -0.01 , N/A )	1282.6	N/A	1.3123 [ 1.0000 ]	131.2% { 134.0% }			
13C2_PFHxA_IIS	( 315.0 / 270.0 ) 578158	( 5.46 , N/A ) ( N/A , -0.01 , N/A )	4375.3	N/A	1.1453 [ 1.0000 ]	114.5% { 120.3% }			
13C4_PFOA_IIS	( 417.0 / 372.0 ) 742380	( 6.98 , N/A ) ( N/A , -0.01 , N/A )	5946.5	N/A	1.1714 [ 1.0000 ]	117.1% { 121.6% }			
13C5_PFNA_IIS	( 468.0 / 423.0 ) 697422	( 7.59 , N/A ) ( N/A , -0.01 , N/A )	342134.1	N/A	1.1542 [ 1.0000 ]	115.4% { 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
13C2_PFDA_IIS	( 515.0 / 470.1 ) 675811	( 8.15 , N/A ) ( N/A , 0.00 , N/A )	831.2	N/A	1.1679 [ 1.0000 ]	116.8% { 126.3% }			
18O2_PFHxS_IIS	( 403.0 / 83.9 ) 1084294	( 7.13 , N/A ) ( N/A , -0.01 , N/A )	2550.6	N/A	1.2662 [ 1.0000 ]	126.6% { 128.8% }			
13C4_PFOS_IIS	( 503.0 / 79.9 ) 1850045	( 8.36 , N/A ) ( N/A , 0.00 , N/A )	1598.6	N/A	1.2289 [ 1.0000 ]	122.9% { 138.0% }			
13C4_PFBA_EIS	( 217.0 / 172.0 ) 2447711	( 3.49 , N/A ) ( N/A , -0.01 , N/A )	4195.9	N/A	7.0003 [ 8.0000 ]	87.5% { 106.5% }			
13C5_PFPeA_EIS	( 268.0 / 223.0 ) 2442322	( 4.53 , N/A ) ( N/A , -0.01 , N/A )	4139.2	N/A	3.8609 [ 4.0000 ]	96.5% { 106.0% }			
13C5_PFHxA_EIS	( 318.0 / 273.0 ) 1405190	( 5.46 , N/A ) ( N/A , -0.01 , N/A )	2790.8	N/A	1.7744 [ 2.0000 ]	88.7% { 98.4% }			
13C4_PFHpA_EIS	( 367.0 / 322.0 ) 1222503	( 6.28 , N/A ) ( N/A , -0.01 , N/A )	3488.8	N/A	1.7817 [ 2.0000 ]	89.1% { 103.8% }			
13C8_PFOA_EIS	( 421.0 / 376.0 ) 1509238	( 6.98 , N/A ) ( N/A , -0.01 , N/A )	1462.3	N/A	1.7951 [ 2.0000 ]	89.8% { 106.3% }			
13C9_PFNA_EIS	( 472.0 / 427.0 ) 583017	( 7.59 , N/A ) ( N/A , 0.00 , N/A )	1339.8	N/A	0.7890 [ 1.0000 ]	78.9% { 95.8% }			
13C6_PFDA_EIS	( 519.0 / 474.0 ) 801762	( 8.14 , N/A ) ( N/A , 0.00 , N/A )	2626.9	N/A	0.9409 [ 1.0000 ]	94.1% { 107.5% }			
13C7_PFUnA_EIS	( 570.0 / 525.0 ) 779454	( 8.63 , N/A ) ( N/A , 0.00 , N/A )	1740.8	N/A	0.9199 [ 1.0000 ]	92.0% { 117.0% }			

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 ) 691123	( 8.92 , N/A ) ( N/A , 0.00 , N/A )	1589.3	N/A	0.9510 [ 1.0000 ]	95.1% { 105.0% }			
13C2_PFTeDA_EIS	( 715.0 / 670.0 ) 570890	( 9.29 , N/A ) ( N/A , 0.00 , N/A )	701.7	N/A	0.8573 [ 1.0000 ]	85.7% { 97.5% }			
13C3_PFBs_EIS	( 302.0 / 80.0 ) 3894433	( 5.43 , N/A ) ( N/A , -0.02 , N/A )	3657.3	N/A	1.7161 [ 2.0000 ]	85.8% { 102.7% }			
13C3_PFHxS_EIS	( 402.0 / 80.0 ) 2127106	( 7.13 , N/A ) ( N/A , -0.01 , N/A )	2191.1	N/A	1.8046 [ 2.0000 ]	90.2% { 108.2% }			
13C8_PFOS_EIS	( 507.0 / 80.0 ) 4410767	( 8.35 , N/A ) ( N/A , 0.00 , N/A )	2389.1	N/A	1.7366 [ 2.0000 ]	86.8% { 104.1% }			
13C2_4:2FTS_EIS	( 329.0 / 81.0 ) 383437	( 5.20 , N/A ) ( N/A , -0.01 , N/A )	1249.2	N/A	3.0611 [ 4.0000 ]	76.5% { 110.4% }			
13C2_6:2FTS_EIS	( 429.0 / 81.0 ) 421858	( 6.71 , N/A ) ( N/A , -0.01 , N/A )	3592.4	N/A	3.0681 [ 4.0000 ]	76.7% { 101.4% }			
13C2_8:2FTS_EIS	( 529.0 / 81.0 ) 565575	( 7.88 , N/A ) ( N/A , -0.01 , N/A )	1456.9	N/A	2.9114 [ 4.0000 ]	72.8% { 93.8% }			
13C8_PFOsa_EIS	( 506.0 / 78.0 ) 8098767	( 9.83 , N/A ) ( N/A , 0.00 , N/A )	2761.1	N/A	1.7178 [ 2.0000 ]	85.9% { 97.5% }			
D3_NMeFOsa_EIS	( 515.0 / 169.0 ) 1702100	( 10.41 , N/A ) ( N/A , 0.00 , N/A )	3242.0	N/A	1.5324 [ 2.0000 ]	76.6% { 87.9% }			
D5_NEtFOsa_EIS	( 531.0 / 169.0 ) 1624539	( 10.56 , N/A ) ( N/A , 0.00 , N/A )	3106.4	N/A	1.6344 [ 2.0000 ]	81.7% { 91.1% }			

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 ) 1113843	( 8.28 , N/A ) ( N/A , 0.00 , N/A )	2198.5	N/A	2.7100 [ 4.0000 ]	67.7% { 84.1% }			
D5_EtFOSAA_EIS	( 589.0 / 419.0 ) 1006913	( 8.53 , N/A ) ( N/A , 0.00 , N/A )	2871.0	N/A	2.9919 [ 4.0000 ]	74.8% { 95.4% }			
D7_NMeFOSE_EIS	( 623.0 / 58.9 ) 6091745	( 10.35 , N/A ) ( N/A , 0.00 , N/A )	2208.9	N/A	15.0431 [ 20.0000 ]	75.2% { 82.0% }			
D9_NEtFOSE_EIS	( 639.0 / 58.9 ) 8332436	( 10.51 , N/A ) ( N/A , 0.00 , N/A )	2531.6	N/A	16.0673 [ 20.0000 ]	80.3% { 88.4% }			
13C3_HFPODA_EIS	( 287.0 / 169.0 ) 3572258	( 5.77 , N/A ) ( N/A , -0.01 , N/A )	3187.3	N/A	7.4086 [ 8.0000 ]	92.6% { 99.6% }			

## ANALYSIS SEQUENCE BLANKS

Laboratory: APPL, LLC  
 Client: AECOM  
 Sequence: SC00647  
 Calibration: 2307007

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

Lab Sample ID	Analyte	Found	Units	RL	C
SC00647-ICB1	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	0.00	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	0.0123	ng/mL	0.10	U
	NMeFOSA	0.109	ng/mL	0.40	U
	NEtFOSA	0.0498	ng/mL	0.40	U
	NMeFOSAA	0.00	ng/mL	0.10	U
	NEtFOSAA	0.00	ng/mL	0.10	U
	NMeFOSE	0.0837	ng/mL	0.40	U
	NEtFOSE	0.106	ng/mL	0.40	U
	HFPO-DA	0.00	ng/mL	0.20	U
	ADONA	0.00	ng/mL	0.20	U
	PFEESA	0.00	ng/mL	0.20	U
	PFMPA	0.00	ng/mL	0.20	U

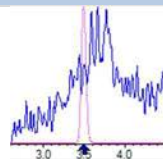
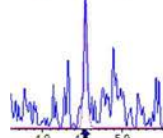
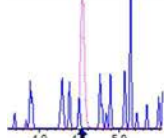
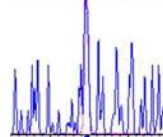
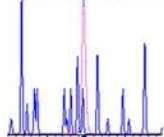
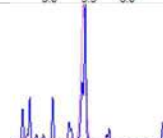
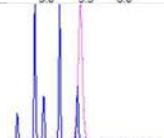
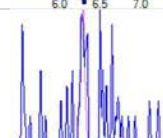
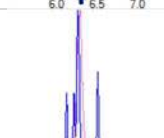
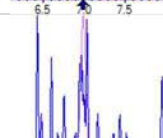
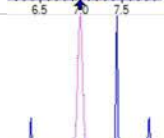
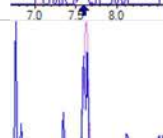
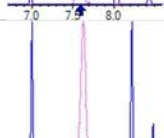
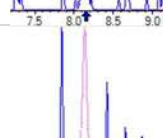
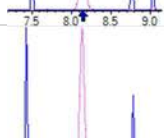
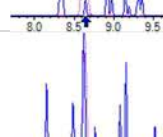
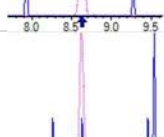
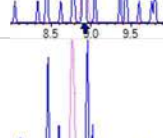
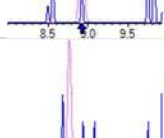
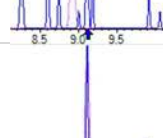
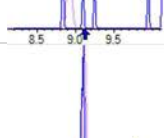


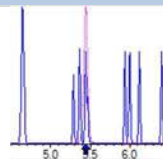
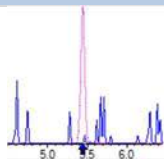
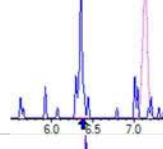
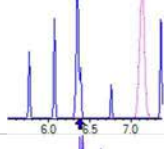
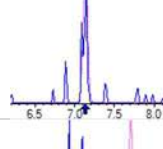
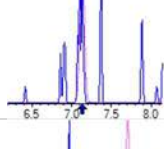
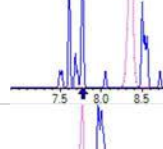
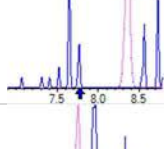
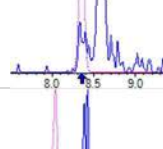
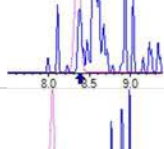
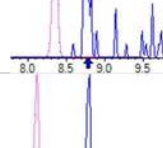
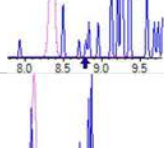
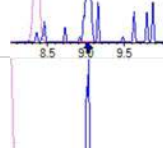
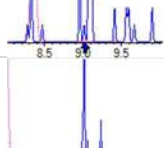
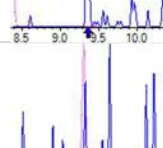
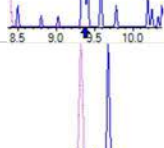
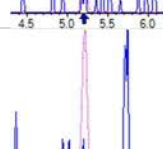
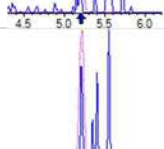
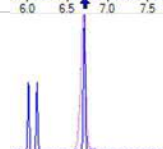
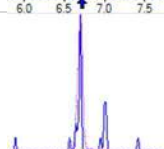
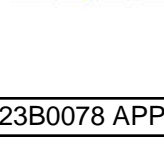
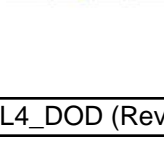
## ANALYSIS SEQUENCE BLANKS

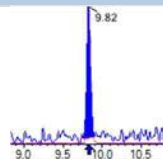
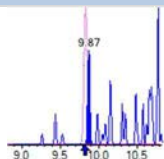
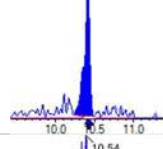
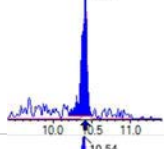
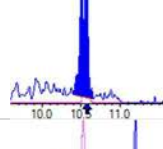
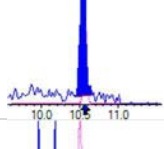
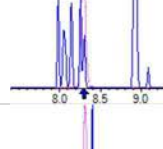
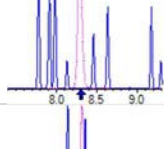
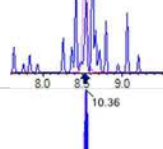
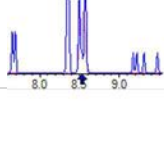
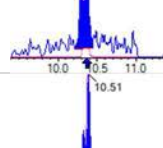
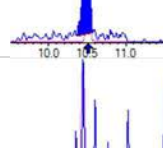
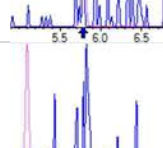
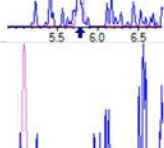
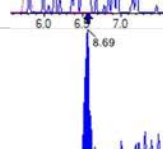
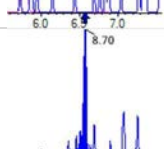
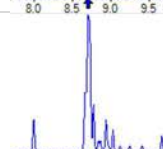
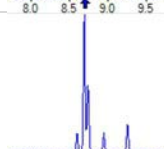
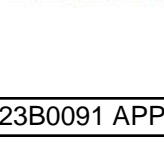
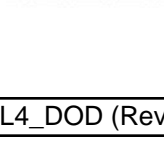
Laboratory: APPL, LLC  
 Client: AECOM  
 Sequence: SC00647  
 Calibration: 2307007

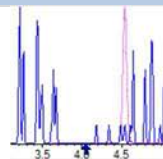
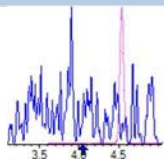
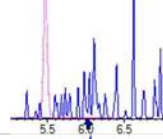
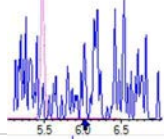
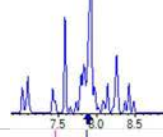
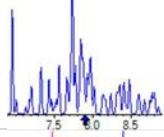
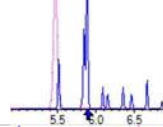
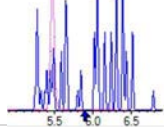
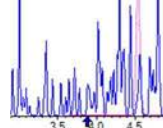
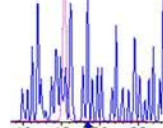
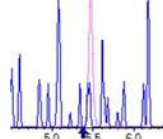
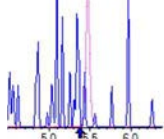
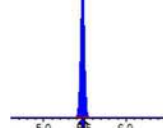
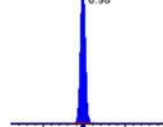
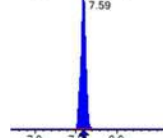
SDG:  
 Project: Red Hill AFFF Assessment Sam  
 Instrument: Saphira

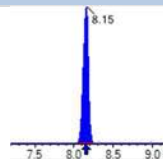
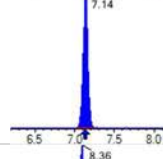
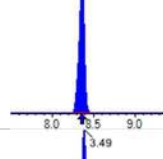
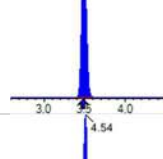
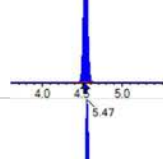
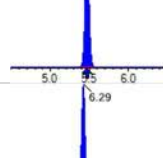
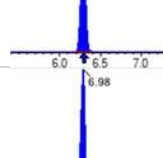
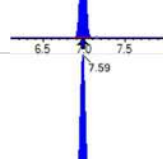
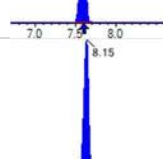
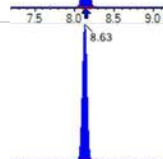
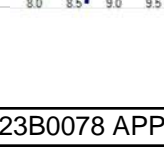
Lab Sample ID	Analyte	Found	Units	RL	C
SC00647-ICB1	PFMBA	0.00	ng/mL	0.20	U
	NFDHA	0.00	ng/mL	0.20	U
	9CL-PF3ONS	0.0571	ng/mL	0.20	U
	11CL-PF3OUDS	0.00	ng/mL	0.20	U
	3:3FTCA	0.00	ng/mL	0.40	U
	5:3FTCA	0.00	ng/mL	0.40	U
	7:3FTCA	0.00	ng/mL	0.40	U
	13C4-PFBA	7.15	ng/mL		
	13C5-PFPEA	3.47	ng/mL		
	13C5-PFHXA	1.78	ng/mL		
	13C4-PFHPA	1.83	ng/mL		
	13C8-PFOA	1.67	ng/mL		
	13C9-PFNA	0.833	ng/mL		
	13C6-PFDA	0.819	ng/mL		
	13C7-PFUnA	0.896	ng/mL		
	13C2-PFDOA	0.867	ng/mL		
	13C2-PFTEDA	0.969	ng/mL		
	13C3-PFBS	1.74	ng/mL		
	13C3-PFHXS	1.69	ng/mL		
	13C8-PFOS	1.74	ng/mL		
	13C2-4:2FTS	3.19	ng/mL		
	13C2-6:2FTS	3.35	ng/mL		
	13C2-8:2FTS	3.12	ng/mL		
	13C8-PFOSA	1.87	ng/mL		
	D3-NMEFOSA	1.87	ng/mL		
	D5-NETFOSA	2.07	ng/mL		
	D3-NMEFOSAA	3.55	ng/mL		
	D5-NETFOSAA	3.73	ng/mL		
	D7-NMEFOSE	19.7	ng/mL		
	D9-NETFOSSE	21.3	ng/mL		
	13C3-HFPO-DA	7.12	ng/mL		

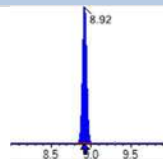
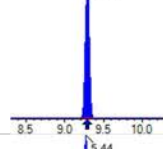
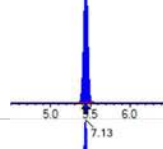
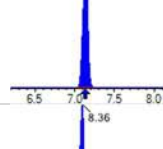
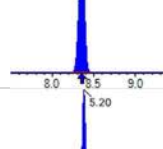
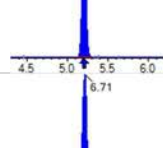
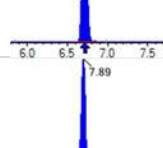
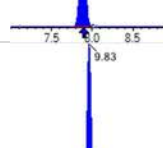
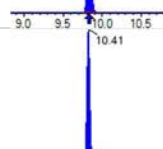
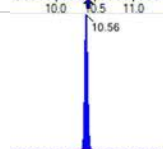
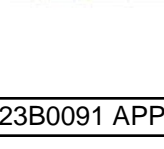
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	( 213.0 / 169.0 ) N/A	( N/A , N/A ) ( N/A , N/A , N/A )	N/A	N/A 0.0 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			

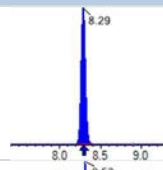
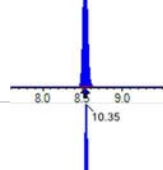
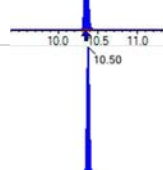


Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	( 299.0 / 80.0 ) N/A ( 299.0 / 99.0 ) N/A	( N/A , N/A ) ( N/A , N/A , 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 ) 36099 ( 498.0 / 478.0 ) 1141	( 9.82 , 1.00 ) ( -0.01 , N/A , -2.9)	51.5 15.8	0.0316 153.8 153.8	0.0123	N/A			
NMeFOSA	( 512.0 / 219.0 ) 31110 ( 512.0 / 169.0 ) 24756	( 10.41 , 1.00 ) ( 0.00 , N/A , 0.2)	86.7 72.0	0.7958 97.1 97.1	0.1090	N/A			
NEIFOSA	( 526.0 / 219.0 ) 39728 ( 526.0 / 169.0 ) 47072	( 10.54 , 1.00 ) ( -0.02 , N/A , 0.0)	78.5 84.5	1.1849 93.3 93.3	0.0498	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 ) 27735	( 10.36 , 1.00 ) ( 0.01 , N/A , 0.0)	48.4	N/A 0.0 0.0	0.0837	N/A			
NEtFOSE	( 630.0 / 59.0 ) 46434	( 10.51 , 1.00 ) ( 0.01 , N/A , 0.0)	78.7	N/A 0.0 0.0	0.1060	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 ) 22131 ( 533.0 / 353.0 ) 7410	( 8.69 , 1.50 ) ( N/A , -0.01 , -0.9)	35.0 34.2	0.3348 96.2 96.2	0.0571	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 ) 241689	( 3.49 , N/A ) ( N/A , 0.01 , N/A )	1351.1	N/A	1.0790 [ 1.0000 ]	107.9% { 107.2% }			
13C2_PFHxA_IIS	( 315.0 / 270.0 ) 514335	( 5.47 , N/A ) ( N/A , 0.01 , N/A )	1753.8	N/A	1.0189 [ 1.0000 ]	101.9% { 107.1% }			
13C4_PFOA_IIS	( 417.0 / 372.0 ) 664713	( 6.98 , N/A ) ( N/A , 0.00 , N/A )	2350.9	N/A	1.0488 [ 1.0000 ]	104.9% { 112.0% }			
13C5_PFNA_IIS	( 468.0 / 423.0 ) 633844	( 7.59 , N/A ) ( N/A , 0.00 , N/A )	2441.3	N/A	1.0489 [ 1.0000 ]	104.9% { 106.9% }			

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_IIS	( 515.0 / 470.1 ) 606167	( 8.15 , N/A ) ( N/A , 0.00 , N/A )	14664.1	N/A	1.0476 [ 1.0000 ]	104.8% { 111.3% }			
18O2_PFHxS_IIS	( 403.0 / 83.9 ) 900713	( 7.14 , N/A ) ( N/A , 0.00 , N/A )	1938.3	N/A	1.0518 [ 1.0000 ]	105.2% { 106.7% }			
13C4_PFOS_IIS	( 503.0 / 79.9 ) 1538499	( 8.36 , N/A ) ( N/A , 0.00 , N/A )	1226.8	N/A	1.0220 [ 1.0000 ]	102.2% { 101.7% }			
13C4_PFBA_EIS	( 217.0 / 172.0 ) 2056496	( 3.49 , N/A ) ( N/A , 0.01 , N/A )	5417.6	N/A	7.1533 [ 8.0000 ]	89.4% { 88.1% }			
13C5_PFPeA_EIS	( 268.0 / 223.0 ) 1955249	( 4.54 , N/A ) ( N/A , 0.00 , N/A )	3715.6	N/A	3.4745 [ 4.0000 ]	86.9% { 80.8% }			
13C5_PFHxA_EIS	( 318.0 / 273.0 ) 1255067	( 5.47 , N/A ) ( N/A , 0.00 , N/A )	2609.0	N/A	1.7815 [ 2.0000 ]	89.1% { 81.8% }			
13C4_PFHpA_EIS	( 367.0 / 322.0 ) 1114027	( 6.29 , N/A ) ( N/A , 0.00 , N/A )	2902.8	N/A	1.8250 [ 2.0000 ]	91.3% { 88.2% }			
13C8_PFOA_EIS	( 421.0 / 376.0 ) 1259198	( 6.98 , N/A ) ( N/A , 0.00 , N/A )	2030.9	N/A	1.6727 [ 2.0000 ]	83.6% { 81.7% }			
13C9_PFNA_EIS	( 472.0 / 427.0 ) 559595	( 7.59 , N/A ) ( N/A , 0.00 , N/A )	2630.8	N/A	0.8333 [ 1.0000 ]	83.3% { 79.6% }			
13C6_PFDA_EIS	( 519.0 / 474.0 ) 625878	( 8.15 , N/A ) ( N/A , 0.00 , N/A )	1513.7	N/A	0.8189 [ 1.0000 ]	81.9% { 80.6% }			
13C7_PFUnA_EIS	( 570.0 / 525.0 ) 681091	( 8.63 , N/A ) ( N/A , 0.00 , N/A )	1401.7	N/A	0.8962 [ 1.0000 ]	89.6% { 88.3% }			

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
13C2_PFDa_EIS	( 615.0 / 570.0 ) 565332	( 8.92, N/A ) ( N/A, 0.00, N/A )	1928.8	N/A	0.8673 [ 1.0000 ]	86.7% { 86.1% }			
13C2_PFTeDA_EIS	( 715.0 / 670.0 ) 578714	( 9.29, N/A ) ( N/A, 0.00, N/A )	3045.8	N/A	0.9689 [ 1.0000 ]	96.9% { 94.9% }			
13C3_PFBs_EIS	( 302.0 / 80.0 ) 3288154	( 5.44, N/A ) ( N/A, 0.00, N/A )	3901.4	N/A	1.7443 [ 2.0000 ]	87.2% { 83.2% }			
13C3_PFHxS_EIS	( 402.0 / 80.0 ) 1656509	( 7.13, N/A ) ( N/A, 0.00, N/A )	1729.5	N/A	1.6918 [ 2.0000 ]	84.6% { 83.2% }			
13C8_PFOS_EIS	( 507.0 / 80.0 ) 3665952	( 8.36, N/A ) ( N/A, 0.00, N/A )	2331.1	N/A	1.7357 [ 2.0000 ]	86.8% { 82.9% }			
13C2_4:2FTS_EIS	( 329.0 / 81.0 ) 331778	( 5.20, N/A ) ( N/A, 0.00, N/A )	1227.6	N/A	3.1885 [ 4.0000 ]	79.7% { 81.1% }			
13C2_6:2FTS_EIS	( 429.0 / 81.0 ) 382237	( 6.71, N/A ) ( N/A, 0.00, N/A )	1745.1	N/A	3.3465 [ 4.0000 ]	83.7% { 83.4% }			
13C2_8:2FTS_EIS	( 529.0 / 81.0 ) 504180	( 7.89, N/A ) ( N/A, 0.00, N/A )	943.1	N/A	3.1244 [ 4.0000 ]	78.1% { 83.3% }			
13C8_PFOsa_EIS	( 506.0 / 78.0 ) 7350043	( 9.83, N/A ) ( N/A, 0.00, N/A )	3807.2	N/A	1.8747 [ 2.0000 ]	93.7% { 85.8% }			
D3_NMeFOsa_EIS	( 515.0 / 169.0 ) 1723173	( 10.41, N/A ) ( N/A, 0.00, N/A )	2942.3	N/A	1.8655 [ 2.0000 ]	93.3% { 94.4% }			
D5_NEtFOsa_EIS	( 531.0 / 169.0 ) 1707041	( 10.56, N/A ) ( N/A, 0.00, N/A )	3216.0	N/A	2.0652 [ 2.0000 ]	103.3% { 99.0% }			

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 ) 1214666	( 8.29 , N/A ) ( N/A , 0.00 , N/A )	1780.1	N/A	3.5537 [ 4.0000 ]	88.8% { 81.6% }			
D5_EtFOSAA_EIS	( 589.0 / 419.0 ) 1043394	( 8.53 , N/A ) ( N/A , 0.00 , N/A )	9843.1	N/A	3.7281 [ 4.0000 ]	93.2% { 93.0% }			
D7_NMeFOSE_EIS	( 623.0 / 58.9 ) 6625930	( 10.35 , N/A ) ( N/A , 0.00 , N/A )	2010.5	N/A	19.6755 [ 20.0000 ]	98.4% { 92.2% }			
D9_NEtFOSE_EIS	( 639.0 / 58.9 ) 9203687	( 10.50 , N/A ) ( N/A , 0.00 , N/A )	1865.6	N/A	21.3412 [ 20.0000 ]	106.7% { 96.7% }			
13C3_HFPODA_EIS	( 287.0 / 169.0 ) 3052956	( 5.78 , N/A ) ( N/A , 0.00 , N/A )	3526.4	N/A	7.1173 [ 8.0000 ]	89.0% { 83.4% }			



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# QUALITY CONTROL RAW DATA

# ANALYSIS DATA SHEET

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Laboratory:	APPL, LLC	Work Order:	23B0091
Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Matrix:	Water	Laboratory ID:	BCB0232-BLK1
		File ID:	S2023-02-14D (7)
Sampled:		Prepared:	02/13/23 07:19
		Analyzed:	02/14/23 20:59
Solids:		Preparation:	EPA 1633
		Dilution:	1
Batch:	BCB0232	Sequence:	SC00636
		Calibration:	2307007
Column:	1	Instrument:	Saphira

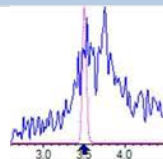
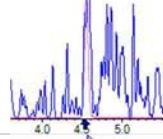
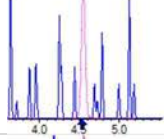
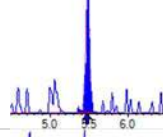
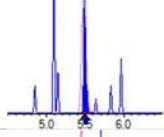
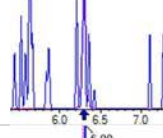
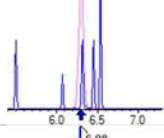
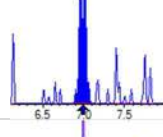
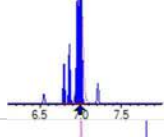
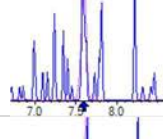
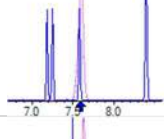
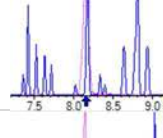
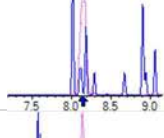
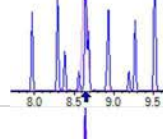
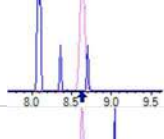
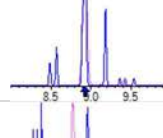
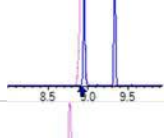
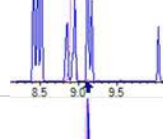
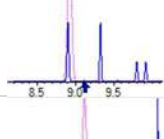
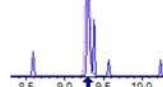
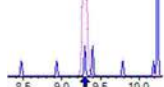
COMPOUND	CONC. (ng/L)	LOQ	LOD	DL	Q
PFBA	0.80 U	1.6	0.80	0.21	U
PFPEA	0.40 U	0.80	0.40	0.065	U
PFHXA	0.0870 J	0.40	0.20	0.055	J
PFHPA	0.20 U	0.40	0.20	0.041	U
PFOA	0.20 U	0.40	0.20	0.15	IR2, U
PFNA	0.20 U	0.40	0.20	0.082	U
PFDA	0.20 U	0.40	0.20	0.10	U
PFUnA	0.20 U	0.40	0.20	0.16	U
PFDOA	0.20 U	0.40	0.20	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	0.112 J	0.40	0.20	0.064	MI5, 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

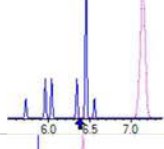
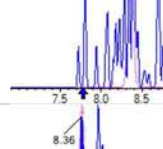
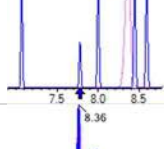
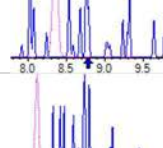
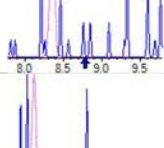
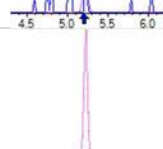
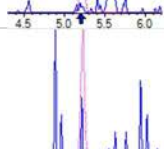
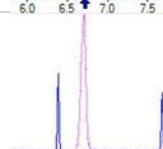
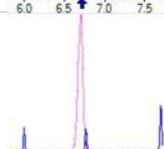
# ANALYSIS DATA SHEET

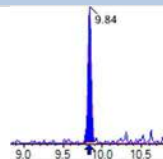
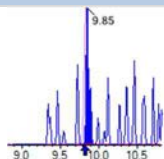
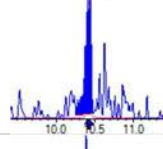
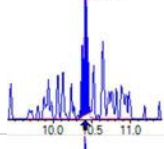
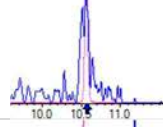
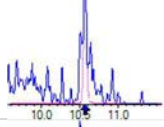
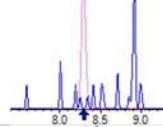
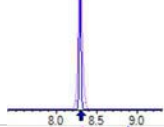
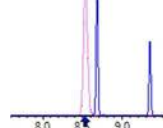
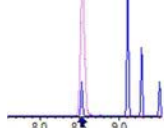
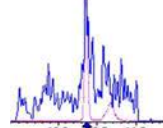
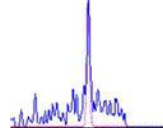
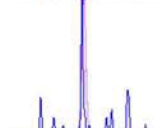
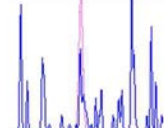
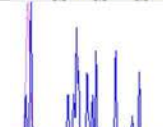
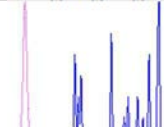
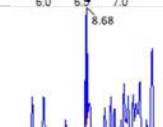
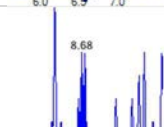
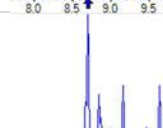
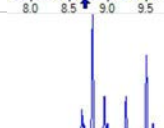
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Laboratory:	APPL, LLC	Work Order:	23B0091
Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Matrix:	Water	Laboratory ID:	BCB0232-BLK1
		File ID:	S2023-02-14D (7)
Sampled:		Prepared:	02/13/23 07:19
		Analyzed:	02/14/23 20:59
Solids:		Preparation:	EPA 1633
		Dilution:	1
Batch:	BCB0232	Sequence:	SC00636
		Calibration:	2307007
Column:	1	Instrument:	Saphira

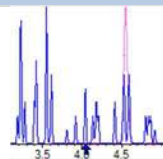
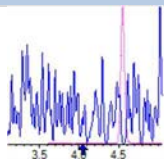
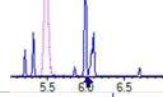
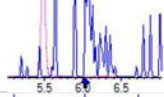
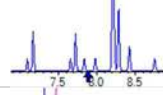
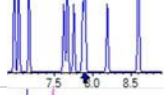
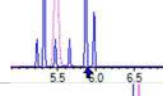
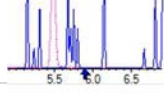
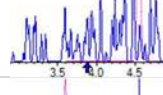
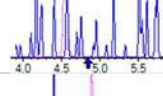
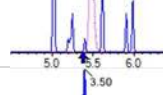
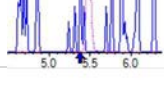
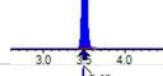
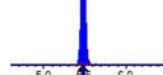

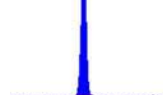
COMPOUND	CONC. (ng/L)	LOQ	LOD	DL	Q
ADONA	0.40 U	0.80	0.40	0.12	U
PFEESA	0.40 U	0.80	0.40	0.11	U
PFMPA	0.40 U	0.80	0.40	0.054	U
PFMBA	0.40 U	0.80	0.40	0.091	U
NFDHA	0.40 U	0.80	0.40	0.30	U
9CL-PF3ONS	0.40 U	0.80	0.40	0.21	IR2, U
11CL-PF3OUDS	0.40 U	0.80	0.40	0.21	U
3:3FTCA	0.80 U	1.6	0.80	0.57	U
5:3FTCA	0.80 U	1.6	0.80	0.44	U
7:3FTCA	0.80 U	1.6	0.80	0.55	U

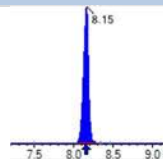
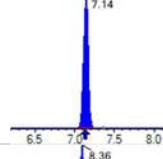
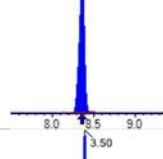
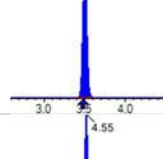
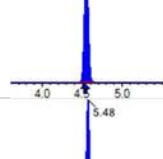
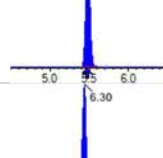
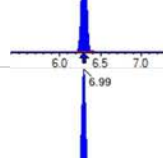
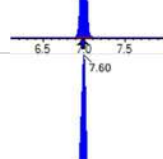
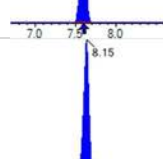
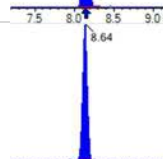
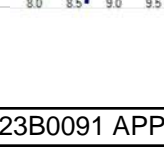
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	( 213.0 / 169.0 ) N/A	( N/A , N/A ) ( N/A , N/A , N/A )	N/A	N/A 0.0 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 ) 10494 ( 313.0 / 119.0 ) 1303	( 5.49 , 1.00 ) ( 0.01 , N/A , -0.3 )	139.1 851.5	0.1242 124.7 130.5	0.0217	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 ) 12380 ( 413.0 / 169.0 ) 7993	( 6.99 , 1.00 ) ( 0.00 , N/A , 0.4 )	250.6 12812.0	0.6456 192.9 178.7	0.0230	N/A			IR2,
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			

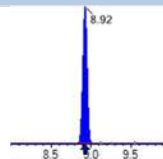
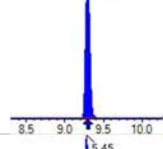
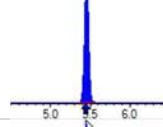
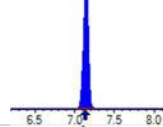
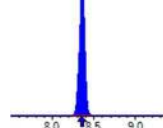
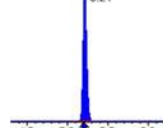
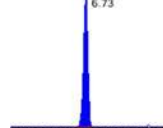
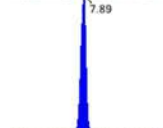
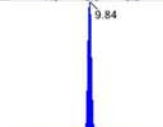
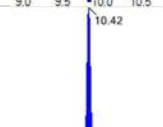
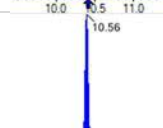
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	( 299.0 / 80.0 ) N/A ( 299.0 / 99.0 ) N/A	( N/A , N/A ) ( N/A , N/A , 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 ) 53215 ( 499.0 / 99.0 ) 12384	( 8.36 , 1.00 ) ( 0.00 , N/A , 0.5 )	46.2 117.8	0.2327 105.2 109.8	0.0280	N/A			MI5 DG 2023-02-15
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			

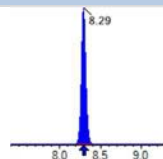
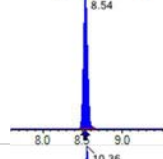
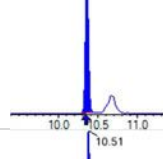
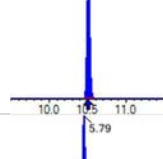
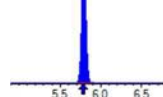
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	( 498.0 / 78.0 ) 46917 ( 498.0 / 478.0 ) 1666	( 9.84 , 1.00 ) ( 0.00 , N/A , -0.9)	108.6 14.4	0.0355 172.8 168.0	0.0172	N/A			
NMeFOSA	( 512.0 / 219.0 ) 5758 ( 512.0 / 169.0 ) 3770	( 10.43 , 1.00 ) ( 0.01 , N/A , 0.6)	35.1 30.9	0.6547 79.9 79.9	0.0799	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 ) N/A	( N/A , N/A ) ( N/A , N/A , N/A)	N/A	N/A 0.0 0.0	0.0000	N/A			
NEiFOSE	( 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 ) 2038 ( 533.0 / 353.0 ) 2050	( 8.68 , 1.50 ) ( N/A , -0.02 , 0.0)	8.2 14.6	1.0060 288.9 304.5	0.0512	N/A			IR2,
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 ) 264564	( 3.50 , N/A ) ( N/A , 0.00 , N/A )	1450.2	N/A	1.1811 [ 1.0000 ]	118.1% { 120.6% }			
13C2_PFHxA_IIS	( 315.0 / 270.0 ) 540893	( 5.48 , N/A ) ( N/A , 0.01 , N/A )	2602.4	N/A	1.0715 [ 1.0000 ]	107.1% { 112.5% }			
13C4_PFOA_IIS	( 417.0 / 372.0 ) 716282	( 6.99 , N/A ) ( N/A , 0.01 , N/A )	1091.6	N/A	1.1302 [ 1.0000 ]	113.0% { 117.3% }			
13C5_PFNA_IIS	( 468.0 / 423.0 ) 701912	( 7.60 , N/A ) ( N/A , 0.00 , N/A )	3219.1	N/A	1.1616 [ 1.0000 ]	116.2% { 124.5% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) ( $\Delta$ RT-I[min], $\Delta$ RT- CV[min], $\Delta$ 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 ) 635802	( 8.15 , N/A ) ( N/A , 0.01 , N/A )	1170.4	N/A	1.0988 [ 1.0000 ]	109.9% { 118.8% }			
18O2_PFHxS_IIS	( 403.0 / 83.9 ) 919443	( 7.14 , N/A ) ( N/A , 0.01 , N/A )	1632.1	N/A	1.0737 [ 1.0000 ]	107.4% { 109.2% }			
13C4_PFOS_IIS	( 503.0 / 79.9 ) 1768239	( 8.36 , N/A ) ( N/A , 0.01 , N/A )	1631.2	N/A	1.1746 [ 1.0000 ]	117.5% { 131.9% }			
13C4_PFBA_EIS	( 217.0 / 172.0 ) 2025069	( 3.50 , N/A ) ( N/A , 0.00 , N/A )	5729.1	N/A	6.4350 [ 8.0000 ]	80.4% { 88.1% }			
13C5_PFPeA_EIS	( 268.0 / 223.0 ) 1976774	( 4.55 , N/A ) ( N/A , 0.01 , N/A )	2543.5	N/A	3.3402 [ 4.0000 ]	83.5% { 85.8% }			
13C5_PFHxA_EIS	( 318.0 / 273.0 ) 1097664	( 5.48 , N/A ) ( N/A , 0.01 , N/A )	2306.2	N/A	1.4816 [ 2.0000 ]	74.1% { 76.9% }			
13C4_PFHpA_EIS	( 367.0 / 322.0 ) 1002220	( 6.30 , N/A ) ( N/A , 0.01 , N/A )	2983.0	N/A	1.5613 [ 2.0000 ]	78.1% { 85.1% }			
13C8_PFOA_EIS	( 421.0 / 376.0 ) 1211634	( 6.99 , N/A ) ( N/A , 0.01 , N/A )	2157.0	N/A	1.4936 [ 2.0000 ]	74.7% { 85.3% }			
13C9_PFNA_EIS	( 472.0 / 427.0 ) 507610	( 7.60 , N/A ) ( N/A , 0.01 , N/A )	3862.3	N/A	0.6825 [ 1.0000 ]	68.3% { 83.4% }			
13C6_PFDA_EIS	( 519.0 / 474.0 ) 603779	( 8.15 , N/A ) ( N/A , 0.01 , N/A )	2011.5	N/A	0.7532 [ 1.0000 ]	75.3% { 81.0% }			
13C7_PFUnA_EIS	( 570.0 / 525.0 ) 605000	( 8.64 , N/A ) ( N/A , 0.01 , N/A )	1289.4	N/A	0.7589 [ 1.0000 ]	75.9% { 90.8% }			

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 ) 619468	( 8.92 , N/A ) ( N/A , 0.01 , N/A )	988.5	N/A	0.9060 [ 1.0000 ]	90.6% { 94.1% }			
13C2_PFTeDA_EIS	( 715.0 / 670.0 ) 515982	( 9.29 , N/A ) ( N/A , 0.01 , N/A )	1520.3	N/A	0.8236 [ 1.0000 ]	82.4% { 88.1% }			
13C3_PFBs_EIS	( 302.0 / 80.0 ) 2883131	( 5.45 , N/A ) ( N/A , 0.01 , N/A )	3581.4	N/A	1.4983 [ 2.0000 ]	74.9% { 76.0% }			
13C3_PFHxS_EIS	( 402.0 / 80.0 ) 1699912	( 7.14 , N/A ) ( N/A , 0.00 , N/A )	2101.3	N/A	1.7007 [ 2.0000 ]	85.0% { 86.5% }			
13C8_PFOS_EIS	( 507.0 / 80.0 ) 3609801	( 8.36 , N/A ) ( N/A , 0.01 , N/A )	1924.2	N/A	1.4870 [ 2.0000 ]	74.4% { 85.2% }			
13C2_4:2FTS_EIS	( 329.0 / 81.0 ) 770186	( 5.21 , N/A ) ( N/A , 0.00 , N/A )	2084.1	N/A	7.2511 [ 4.0000 ]	181.3% { 221.8% }			S2,
13C2_6:2FTS_EIS	( 429.0 / 81.0 ) 318658	( 6.73 , N/A ) ( N/A , 0.01 , N/A )	2664.9	N/A	2.7330 [ 4.0000 ]	68.3% { 76.6% }			
13C2_8:2FTS_EIS	( 529.0 / 81.0 ) 463052	( 7.89 , N/A ) ( N/A , 0.00 , N/A )	6669.7	N/A	2.8111 [ 4.0000 ]	70.3% { 76.8% }			
13C8_PFOsa_EIS	( 506.0 / 78.0 ) 6819838	( 9.84 , N/A ) ( N/A , 0.00 , N/A )	2721.3	N/A	1.5135 [ 2.0000 ]	75.7% { 82.1% }			
D3_NMeFOsa_EIS	( 515.0 / 169.0 ) 1062630	( 10.42 , N/A ) ( N/A , 0.00 , N/A )	1979.7	N/A	1.0009 [ 2.0000 ]	50.0% { 54.9% }			
D5_NEtFOsa_EIS	( 531.0 / 169.0 ) 1008862	( 10.56 , N/A ) ( N/A , 0.00 , N/A )	2813.2	N/A	1.0619 [ 2.0000 ]	53.1% { 56.6% }			

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 ) 1040945	( 8.29 , N/A ) ( N/A , 0.01 , N/A )	2715.8	N/A	2.6498 [ 4.0000 ]	66.2% { 78.6% }			
D5_EtFOSAA_EIS	( 589.0 / 419.0 ) 845671	( 8.54 , N/A ) ( N/A , 0.01 , N/A )	14688.7	N/A	2.6290 [ 4.0000 ]	65.7% { 80.1% }			
D7_NMeFOSE_EIS	( 623.0 / 58.9 ) 3741948	( 10.36 , N/A ) ( N/A , 0.01 , N/A )	628.8	N/A	9.6679 [ 20.0000 ]	48.3% { 50.4% }			
D9_NEtFOSE_EIS	( 639.0 / 58.9 ) 7540801	( 10.51 , N/A ) ( N/A , 0.00 , N/A )	2088.5	N/A	15.2136 [ 20.0000 ]	76.1% { 80.0% }			
13C3_HFPODA_EIS	( 287.0 / 169.0 ) 3009188	( 5.79 , N/A ) ( N/A , 0.01 , N/A )	3498.8	N/A	6.6708 [ 8.0000 ]	83.4% { 83.9% }			

# ANALYSIS DATA SHEET

## LCS

Laboratory:	APPL, LLC	Work Order:	23B0091
Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Matrix:	Water	Laboratory ID:	BCB0232-BS1
		File ID:	S2023-02-14D (8)
Sampled:		Prepared:	02/13/23 07:19
		Analyzed:	02/14/23 21:12
Solids:		Preparation:	EPA 1633
		Dilution:	1
Batch:	BCB0232	Sequence:	SC00636
		Calibration:	2307007
Column:	1	Instrument:	Saphira

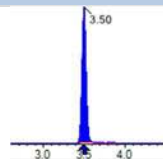
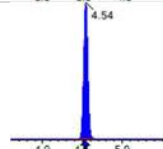
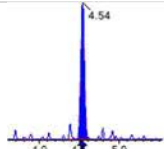
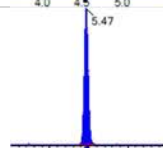
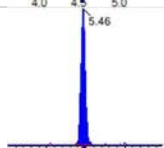
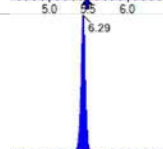
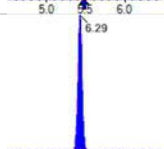
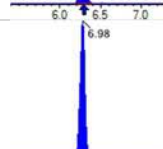
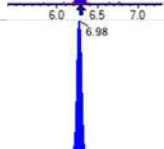
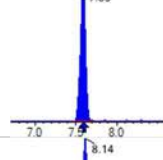
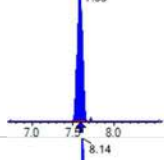
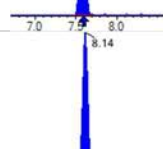
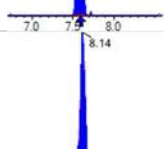
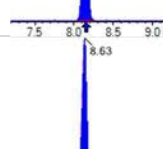
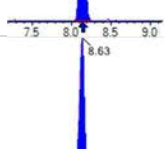
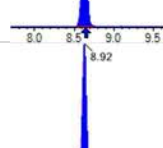
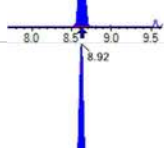
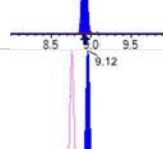
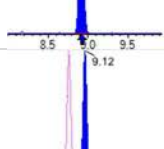
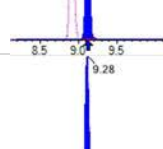
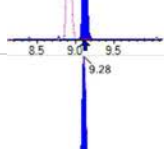
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PFBA	16.7	1.6	0.21	
PFPEA	8.53	0.80	0.065	
PFHXA	4.36	0.40	0.055	
PFHPA	4.01	0.40	0.041	
PFOA	4.30	0.40	0.15	
PFNA	4.02	0.40	0.082	
PFDA	4.62	0.40	0.10	
PFUnA	4.11	0.40	0.16	
PFDOA	3.92	0.40	0.11	
PFTRDA	4.08	0.40	0.20	
PFTEDA	4.34	0.40	0.20	
PFBS	3.92	0.40	0.037	
PFPEs	3.94	0.40	0.063	
PFHXS	3.70	0.40	0.032	
PFHPS	4.10	0.40	0.051	
PFOS	3.80	0.40	0.064	
PFNS	4.06	0.40	0.12	
PFDS	3.98	0.40	0.15	
PFDOS	3.84	0.40	0.12	
4:2FTS	16.0	1.6	0.29	
6:2FTS	18.0	1.6	0.31	
8:2FTS	17.2	1.6	0.082	
PFOSA	4.63	0.40	0.10	
NMeFOSA	16.1	1.6	0.47	
NEtFOSA	18.2	1.6	0.41	
NMeFOSAA	4.43	0.40	0.11	
NEtFOSAA	4.08	0.40	0.11	
NMeFOSE	18.5	1.6	1.0	
NEtFOSE	16.1	1.6	1.0	
HFPO-DA	8.21	0.80	0.17	

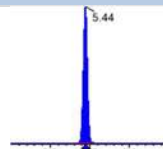
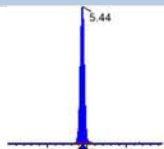
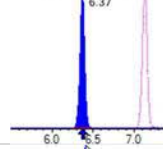
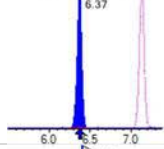
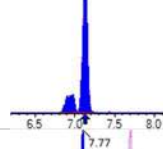
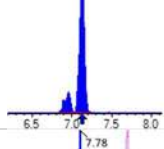
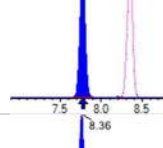
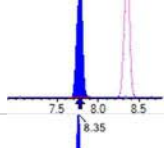
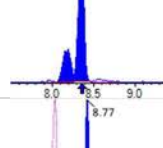
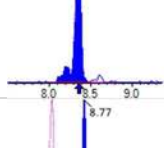
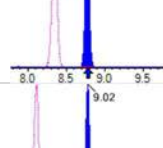
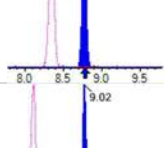
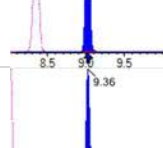
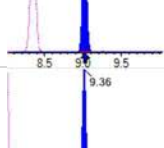
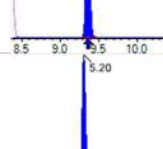
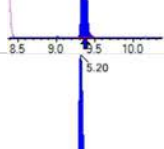
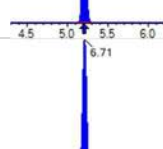
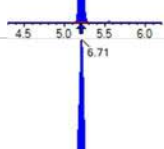
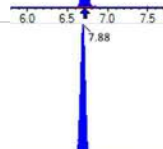
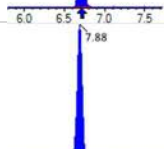
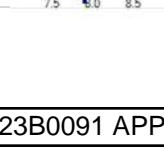
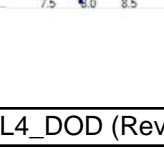
# ANALYSIS DATA SHEET

## LCS

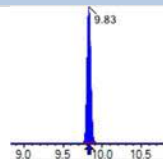
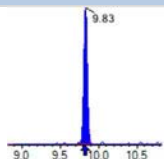
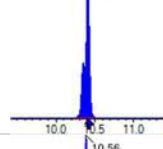
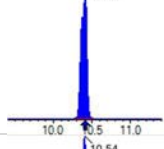
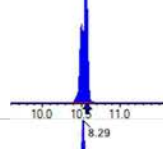
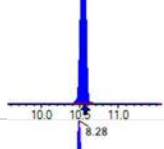
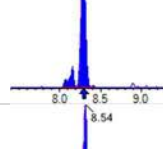
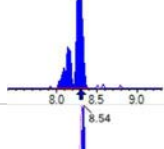
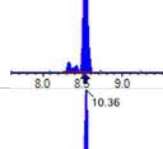
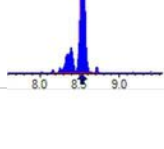
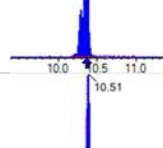
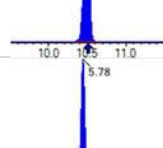
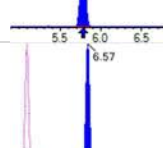
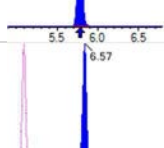
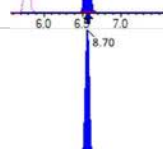
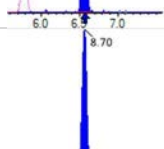
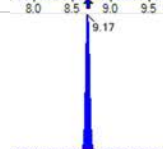
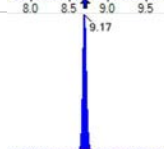
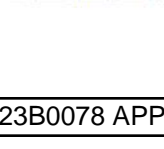
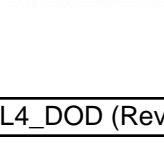
Laboratory:	APPL, LLC	Work Order:	23B0091
Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Matrix:	Water	Laboratory ID:	BCB0232-BS1
		File ID:	S2023-02-14D (8)
Sampled:		Prepared:	02/13/23 07:19
		Analyzed:	02/14/23 21:12
Solids:		Preparation:	EPA 1633
		Dilution:	1
Batch:	BCB0232	Sequence:	SC00636
		Calibration:	2307007
Column:	1	Instrument:	Saphira

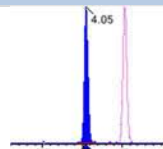
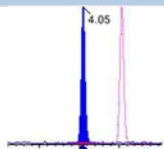
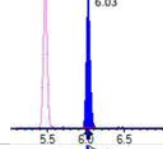
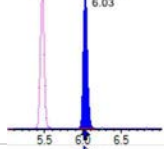
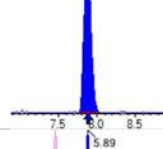
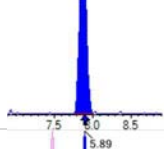
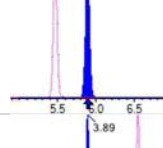
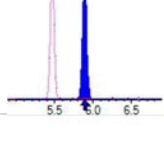
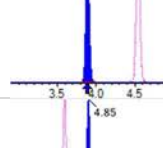
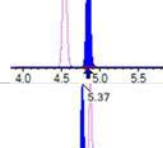
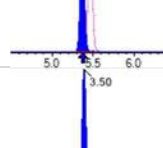
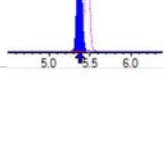
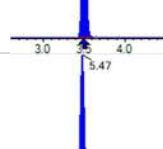
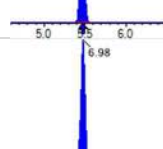
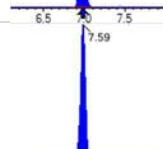

COMPOUND	CONC. (ng/L)	LOQ	DL	Q
ADONA	7.95	0.80	0.12	
PFEESA	8.03	0.80	0.11	
PFMPA	8.03	0.80	0.054	
PFMBA	7.76	0.80	0.091	
NFDHA	8.99	0.80	0.30	
9CL-PF3ONS	7.63	0.80	0.21	
11CL-PF3OUDS	7.70	0.80	0.21	
3:3FTCA	16.9	1.6	0.57	
5:3FTCA	17.6	1.6	0.44	
7:3FTCA	16.0	1.6	0.55	

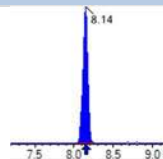
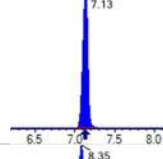
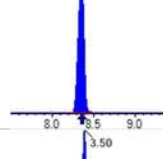
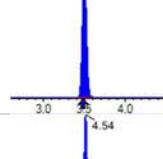
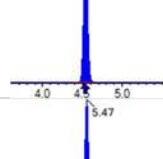
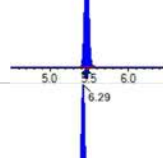
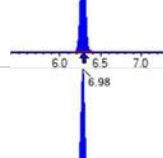
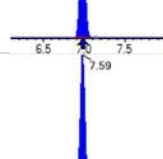
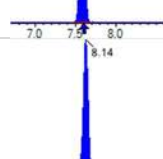
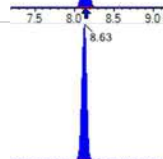
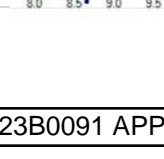
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	( 213.0 / 169.0 ) 1028740	( 3.50 , 1.00 ) ( 0.00 , N/A , 0.0 )	229.9	N/A 0.0 0.0	4.1823 [ 4.0000 ]	104.6%			
PFPeA	( 263.0 / 219.0 ) 1023282 ( 263.0 / 69.0 ) 13702	( 4.54 , 1.00 ) ( 0.00 , N/A , 0.2 )	2573.7 132.6	0.0134 125.0 121.5	2.1335 [ 2.0000 ]	106.7%			
PFHxA	( 313.0 / 269.0 ) 646363 ( 313.0 / 119.0 ) 75028	( 5.47 , 1.00 ) ( 0.00 , N/A , 0.1 )	2464.0 2798.8	0.1161 116.5 122.0	1.0909 [ 1.0000 ]	109.1%			
PFHpA	( 363.0 / 319.0 ) 517688 ( 363.0 / 169.0 ) 153670	( 6.29 , 1.00 ) ( 0.00 , N/A , 0.2 )	8693.5 24718.9	0.2968 92.6 92.8	1.0021 [ 1.0000 ]	100.2%			
PFOA	( 413.0 / 369.0 ) 687919 ( 413.0 / 169.0 ) 201680	( 6.98 , 1.00 ) ( 0.00 , N/A , 0.3 )	9832.7 264925.6	0.2932 87.6 81.1	1.0754 [ 1.0000 ]	107.5%			
PFNA	( 463.0 / 419.0 ) 552486 ( 463.0 / 169.0 ) 133290	( 7.59 , 1.00 ) ( 0.00 , N/A , 0.1 )	2622.2 163028.5	0.2413 107.8 109.1	1.0047 [ 1.0000 ]	100.5%			
PFDA	( 513.0 / 469.0 ) 701206 ( 513.0 / 169.0 ) 91123	( 8.14 , 1.00 ) ( 0.00 , N/A , 0.0 )	1252.1 13815.9	0.1300 106.9 119.8	1.1561 [ 1.0000 ]	115.6%			
PFUnA	( 563.0 / 519.0 ) 616296 ( 563.0 / 169.0 ) 77906	( 8.63 , 1.00 ) ( 0.00 , N/A , -0.1 )	2023.9 2176.3	0.1264 120.6 108.1	1.0268 [ 1.0000 ]	102.7%			
PFDaA	( 613.0 / 569.0 ) 536590 ( 613.0 / 169.0 ) 112621	( 8.92 , 1.00 ) ( 0.00 , N/A , 0.1 )	1477.1 3614.7	0.2099 128.3 125.1	0.9809 [ 1.0000 ]	98.1%			
PFTTrDA	( 663.0 / 619.0 ) 505510 ( 663.0 / 169.0 ) 161483	( 9.12 , 1.02 ) ( N/A , 0.00 , -0.1 )	1670.6 709.1	0.3194 133.7 118.7	1.0206 [ 1.0000 ]	102.1%			
PFTeDA	( 713.0 / 669.0 ) 485519 ( 713.0 / 169.0 ) 117666	( 9.28 , 1.00 ) ( 0.00 , N/A , 0.0 )	943.6 505.3	0.2424 117.3 117.2	1.0841 [ 1.0000 ]	108.4%			

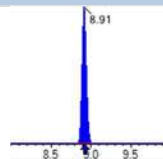
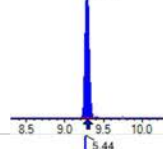
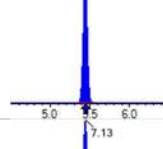
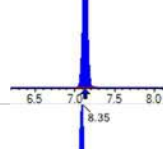
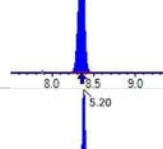
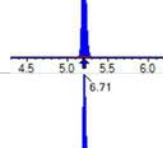
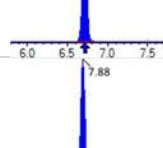
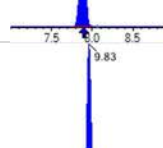
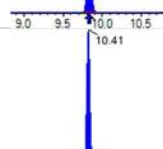
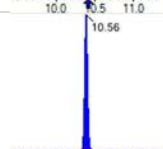
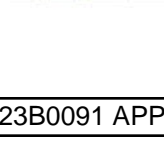
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	( 299.0 / 80.0 ) 1115998 ( 299.0 / 99.0 ) 699946	( 5.44 , 1.00 ) ( 0.00 , N/A , 0.1 )	2887.2 1335.2	0.6272 97.0 101.4	0.9812 [ 0.8847 ]	110.9%			
PFPeS	( 349.0 / 80.0 ) 1807486 ( 349.0 / 99.0 ) 597270	( 6.37 , 0.89 ) ( N/A , -0.01 , -0.1 )	10714532.8 6069763.2	0.3304 92.6 95.9	0.9856 [ 0.9384 ]	105.0%			
PFHxS	( 399.0 / 80.0 ) 1353261 ( 399.0 / 99.0 ) 489677	( 7.13 , 1.00 ) ( 0.00 , N/A , 0.2 )	479293.9 2005.3	0.3618 111.4 111.1	0.9255 [ 0.9110 ]	101.6%			
PFHpS	( 449.0 / 80.0 ) 1597230 ( 449.0 / 99.0 ) 459560	( 7.77 , 0.93 ) ( N/A , 0.00 , -0.3 )	785585.0 73356.9	0.2877 105.7 103.4	1.0257 [ 0.9514 ]	107.8%			
PFOS	( 499.0 / 80.0 ) 2046739 ( 499.0 / 99.0 ) 475253	( 8.36 , 1.00 ) ( 0.00 , N/A , 0.1 )	266.9 202.6	0.2322 104.9 109.5	0.9497 [ 0.9275 ]	102.4%			
PFNS	( 549.0 / 80.0 ) 2459992 ( 549.0 / 99.0 ) 585406	( 8.77 , 1.05 ) ( N/A , 0.00 , 0.0 )	35187.2 75700927.8	0.2380 108.5 103.5	1.0157 [ 0.9599 ]	105.8%			
PFDS	( 599.0 / 80.0 ) 2619461 ( 599.0 / 99.0 ) 523872	( 9.02 , 1.08 ) ( N/A , 0.00 , 0.2 )	2769.2 1476.6	0.2000 91.7 93.1	0.9942 [ 0.9631 ]	103.2%			
PFDoS	( 699.0 / 80.0 ) 1612088 ( 699.0 / 99.0 ) 385963	( 9.36 , 1.12 ) ( N/A , 0.00 , 0.0 )	2308.5 1278.3	0.2394 115.0 111.6	0.9594 [ 0.9696 ]	98.9%			
4:2FTS	( 327.0 / 307.0 ) 2376216 ( 327.0 / 81.0 ) 1565386	( 5.20 , 1.00 ) ( 0.00 , N/A , 0.1 )	5038.8 2723.2	0.6588 97.6 96.1	4.0019 [ 3.7381 ]	107.1%			
6:2FTS	( 427.0 / 407.0 ) 611031 ( 427.0 / 81.0 ) 474564	( 6.71 , 1.00 ) ( 0.00 , N/A , 0.0 )	2036.6 1954.8	0.7767 93.8 100.5	4.5069 [ 3.7962 ]	118.7%			
8:2FTS	( 527.0 / 507.0 ) 613315 ( 527.0 / 81.0 ) 497241	( 7.88 , 1.00 ) ( 0.00 , N/A , 0.1 )	1459.6 1106.3	0.8107 97.5 104.8	4.2875 [ 3.8332 ]	111.9%			

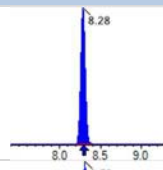
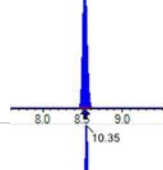
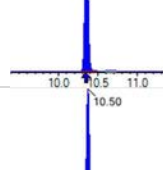
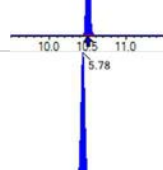



Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min] , R.R.T.) (ΔRT-[min], ΔRT- CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFOSA	( 498.0 / 78.0 ) 3359523 ( 498.0 / 478.0 ) 64647	( 9.83 , 1.00 ) ( 0.00 , N/A , -0.1)	2478.3 503.3	0.0192 93.7 91.1	1.1574 [ 1.0000 ]	115.7%			
NMeFOSA	( 512.0 / 219.0 ) 2010178 ( 512.0 / 169.0 ) 1695361	( 10.41 , 1.00 ) ( 0.00 , N/A , 0.8)	4744.0 2775.0	0.8434 103.0 103.0	4.0292 [ 4.0000 ]	100.7%			
NEIFOSA	( 526.0 / 219.0 ) 2343067 ( 526.0 / 169.0 ) 2986093	( 10.56 , 1.00 ) ( 0.00 , N/A , 0.9)	5219.2 4467.0	1.2744 100.4 102.7	4.5388 [ 4.0000 ]	113.5%			
NMeFOSAA	( 570.0 / 419.0 ) 243929 ( 570.0 / 483.0 ) 107657	( 8.29 , 1.00 ) ( 0.00 , N/A , 0.6)	880.5 78223.0	0.4413 93.0 95.3	1.1064 [ 1.0000 ]	110.6%			
NEIFOSAA	( 584.0 / 419.0 ) 202975 ( 584.0 / 526.0 ) 120806	( 8.54 , 1.00 ) ( 0.01 , N/A , -0.4)	141848.9 8409.5	0.5952 116.6 101.6	1.0204 [ 1.0000 ]	102.0%			
NMeFOSE	( 616.0 / 59.0 ) 884862	( 10.36 , 1.00 ) ( 0.01 , N/A , 0.0)	1155.5	N/A 0.0 0.0	4.6340 [ 4.0000 ]	115.8%			
NEIFOSE	( 630.0 / 59.0 ) 1474308	( 10.51 , 1.00 ) ( 0.01 , N/A , 0.0)	1092.6	N/A 0.0 0.0	4.0359 [ 4.0000 ]	100.9%			
HFPO-DA	( 285.0 / 169.0 ) 708386 ( 285.0 / 185.0 ) 1936370	( 5.78 , 1.00 ) ( 0.00 , N/A , 0.0)	1519.7 2110.7	2.7335 99.2 98.5	2.0514 [ 2.0000 ]	102.6%			
ADONA	( 377.0 / 85.0 ) 2359689 ( 377.0 / 251.0 ) 228515	( 6.57 , 1.14 ) ( N/A , 0.00 , 0.1)	2234.4 632.0	0.0968 109.8 103.7	1.9871 [ 1.8854 ]	105.4%			
9CI-PF3ONS	( 531.0 / 351.0 ) 7171516 ( 533.0 / 353.0 ) 2157893	( 8.70 , 1.51 ) ( N/A , 0.00 , 0.0)	2329.1 1370.9	0.3009 86.4 91.1	1.9084 [ 1.8665 ]	102.2%			
11CI-PF3OUDS	( 631.0 / 451.0 ) 3721907 ( 633.0 / 453.0 ) 1368951	( 9.17 , 1.59 ) ( N/A , 0.00 , 0.1)	2278.4 1642.0	0.3678 101.7 103.2	1.9239 [ 1.8864 ]	102.0%			

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
3:3FTCA	( 241.0 / 177.0 ) 48415 ( 241.0 / 117.0 ) 75897	( 4.05 , 0.89 ) ( N/A , -0.01 , 0.0 )	829.4 388.0	1.5676 99.3 95.8	4.2339 [ 4.0000 ]	105.8%			
5:3FTCA	( 341.0 / 236.7 ) 356077 ( 341.0 / 217.0 ) 653471	( 6.03 , 1.10 ) ( N/A , -0.01 , -0.1 )	935.6 1362.6	1.8352 107.6 114.0	4.3946 [ 4.0000 ]	109.9%			
7:3FTCA	( 441.0 / 317.0 ) 597814 ( 441.0 / 337.0 ) 537524	( 7.88 , 1.44 ) ( N/A , -0.01 , -0.1 )	452.1 541.7	0.8991 104.5 110.6	3.9939 [ 4.0000 ]	99.8%			
PFEESA	( 315.0 / 135.0 ) 1428891 ( 315.0 / 83.0 ) 401065	( 5.89 , 1.08 ) ( N/A , -0.01 , 0.0 )	2473.0 1058.8	0.2807 109.9 101.9	2.0075 [ 1.7849 ]	112.5%			
PFMPA	( 229.0 / 85.0 ) 210277	( 3.89 , 0.86 ) ( N/A , 0.00 , 0.0 )	1582.7	N/A 0.0 0.0	2.0072 [ 2.0000 ]	100.4%			
PFMBA	( 279.0 / 85.0 ) 851818	( 4.85 , 1.07 ) ( N/A , 0.00 , 0.0 )	2063.6	N/A 0.0 0.0	1.9389 [ 2.0000 ]	96.9%			
NFDHA	( 295.0 / 201.0 ) 738788 ( 295.0 / 85.0 ) 706243	( 5.37 , 0.98 ) ( N/A , -0.01 , 0.0 )	1692.1 1822.3	0.9559 95.5 90.4	2.2464 [ 2.0000 ]	112.3%			
13C3_PFBa_IIS	( 216.0 / 172.0 ) 275535	( 3.50 , N/A ) ( N/A , 0.00 , N/A )	1570.1	N/A	1.2301 [ 1.0000 ]	123.0% { 125.6% }			
13C2_PFHxA_IIS	( 315.0 / 270.0 ) 576906	( 5.47 , N/A ) ( N/A , -0.01 , N/A )	3337.6	N/A	1.1428 [ 1.0000 ]	114.3% { 120.0% }			
13C4_PFOA_IIS	( 417.0 / 372.0 ) 712271	( 6.98 , N/A ) ( N/A , 0.00 , N/A )	2180.2	N/A	1.1239 [ 1.0000 ]	112.4% { 116.7% }			
13C5_PFNA_IIS	( 468.0 / 423.0 ) 683819	( 7.59 , N/A ) ( N/A , -0.01 , N/A )	1067.6	N/A	1.1316 [ 1.0000 ]	113.2% { 121.3% }			

Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) ( $\Delta$ RT-I[ $\mu$ min], $\Delta$ RT-CV[ $\mu$ min], $\Delta$ 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 ) 690510	( 8.14 , N/A ) ( N/A , 0.00 , N/A )	1279.2	N/A	1.1933 [ 1.0000 ]	119.3% { 129.0% }			
18O2_PFHxS_IIS	( 403.0 / 83.9 ) 1034664	( 7.13 , N/A ) ( N/A , 0.00 , N/A )	2307.0	N/A	1.2082 [ 1.0000 ]	120.8% { 122.9% }			
13C4_PFOS_IIS	( 503.0 / 79.9 ) 1673461	( 8.35 , N/A ) ( N/A , 0.00 , N/A )	1454.6	N/A	1.1116 [ 1.0000 ]	111.2% { 124.8% }			
13C4_PFBA_EIS	( 217.0 / 172.0 ) 2375575	( 3.50 , N/A ) ( N/A , 0.00 , N/A )	5482.6	N/A	7.2482 [ 8.0000 ]	90.6% { 103.4% }			
13C5_PFPeA_EIS	( 268.0 / 223.0 ) 2331821	( 4.54 , N/A ) ( N/A , 0.00 , N/A )	3952.3	N/A	3.6942 [ 4.0000 ]	92.4% { 101.2% }			
13C5_PFHxA_EIS	( 318.0 / 273.0 ) 1347972	( 5.47 , N/A ) ( N/A , -0.01 , N/A )	2301.1	N/A	1.7058 [ 2.0000 ]	85.3% { 94.4% }			
13C4_PFHpA_EIS	( 367.0 / 322.0 ) 1223554	( 6.29 , N/A ) ( N/A , -0.01 , N/A )	2177.3	N/A	1.7871 [ 2.0000 ]	89.4% { 103.8% }			
13C8_PFOA_EIS	( 421.0 / 376.0 ) 1436829	( 6.98 , N/A ) ( N/A , 0.00 , N/A )	3041.0	N/A	1.7812 [ 2.0000 ]	89.1% { 101.2% }			
13C9_PFNA_EIS	( 472.0 / 427.0 ) 621495	( 7.59 , N/A ) ( N/A , 0.00 , N/A )	4526.4	N/A	0.8578 [ 1.0000 ]	85.8% { 102.1% }			
13C6_PFDA_EIS	( 519.0 / 474.0 ) 689530	( 8.14 , N/A ) ( N/A , 0.00 , N/A )	5175.5	N/A	0.7920 [ 1.0000 ]	79.2% { 92.5% }			
13C7_PFUnA_EIS	( 570.0 / 525.0 ) 722438	( 8.63 , N/A ) ( N/A , 0.00 , N/A )	1272.2	N/A	0.8345 [ 1.0000 ]	83.4% { 108.4% }			

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 ) 622073	( 8.91 , N/A ) ( N/A , -0.01 , N/A )	9584.5	N/A	0.8378 [ 1.0000 ]	83.8% { 94.5% }			
13C2_PFTeDA_EIS	( 715.0 / 670.0 ) 517563	( 9.29 , N/A ) ( N/A , 0.00 , N/A )	1033.3	N/A	0.7607 [ 1.0000 ]	76.1% { 88.4% }			
13C3_PFBs_EIS	( 302.0 / 80.0 ) 3660317	( 5.44 , N/A ) ( N/A , 0.00 , N/A )	3113.9	N/A	1.6903 [ 2.0000 ]	84.5% { 96.5% }			
13C3_PFHxS_EIS	( 402.0 / 80.0 ) 1901767	( 7.13 , N/A ) ( N/A , -0.01 , N/A )	1974.1	N/A	1.6908 [ 2.0000 ]	84.5% { 96.8% }			
13C8_PFOS_EIS	( 507.0 / 80.0 ) 4087528	( 8.35 , N/A ) ( N/A , 0.00 , N/A )	1607.5	N/A	1.7792 [ 2.0000 ]	89.0% { 96.4% }			
13C2_4:2FTS_EIS	( 329.0 / 81.0 ) 869623	( 5.20 , N/A ) ( N/A , -0.01 , N/A )	3060.2	N/A	7.2755 [ 4.0000 ]	181.9% { 250.4% }			S2,
13C2_6:2FTS_EIS	( 429.0 / 81.0 ) 382201	( 6.71 , N/A ) ( N/A , -0.01 , N/A )	1774.5	N/A	2.9130 [ 4.0000 ]	72.8% { 91.9% }			
13C2_8:2FTS_EIS	( 529.0 / 81.0 ) 525301	( 7.88 , N/A ) ( N/A , -0.01 , N/A )	1269.1	N/A	2.8338 [ 4.0000 ]	70.8% { 87.1% }			
13C8_PFOA_EIS	( 506.0 / 78.0 ) 7258433	( 9.83 , N/A ) ( N/A , 0.00 , N/A )	3975.0	N/A	1.7020 [ 2.0000 ]	85.1% { 87.4% }			
D3_NMeFOA_EIS	( 515.0 / 169.0 ) 1176595	( 10.41 , N/A ) ( N/A , 0.00 , N/A )	2442.5	N/A	1.1710 [ 2.0000 ]	58.6% { 60.8% }			
D5_NEtFOA_EIS	( 531.0 / 169.0 ) 1105143	( 10.56 , N/A ) ( N/A , 0.00 , N/A )	3346.9	N/A	1.2292 [ 2.0000 ]	61.5% { 62.0% }			

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 ) 1123740	( 8.28 , N/A ) ( N/A , 0.00 , N/A )	3213.1	N/A	3.0225 [ 4.0000 ]	75.6% { 84.9% }			
D5_EtFOSAA_EIS	( 589.0 / 419.0 ) 947341	( 8.53 , N/A ) ( N/A , 0.00 , N/A )	952923.0	N/A	3.1119 [ 4.0000 ]	77.8% { 89.8% }			
D7_NMeFOSE_EIS	( 623.0 / 58.9 ) 3820002	( 10.35 , N/A ) ( N/A , 0.00 , N/A )	1433.8	N/A	10.4286 [ 20.0000 ]	52.1% { 51.4% }			
D9_NEtFOSE_EIS	( 639.0 / 58.9 ) 7677316	( 10.50 , N/A ) ( N/A , 0.00 , N/A )	2228.1	N/A	16.3662 [ 20.0000 ]	81.8% { 81.5% }			
13C3_HFPODA_EIS	( 287.0 / 169.0 ) 3493284	( 5.78 , N/A ) ( N/A , -0.01 , N/A )	4195.5	N/A	7.2605 [ 8.0000 ]	90.8% { 97.4% }			

# ANALYSIS DATA SHEET

## MRL Check

Laboratory:	APPL, LLC	Work Order:	23B0091
Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Matrix:	Water	Laboratory ID:	BCB0232-MRL1
		File ID:	S2023-02-14D (9)
Sampled:		Prepared:	02/13/23 07:19
		Analyzed:	02/14/23 21:25
Solids:		Preparation:	EPA 1633
		Dilution:	1
Batch:	BCB0232	Sequence:	SC00636
		Calibration:	2307007
Column:	1	Instrument:	Saphira

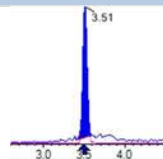
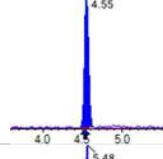
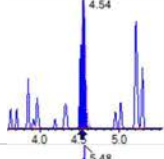
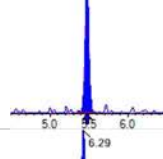
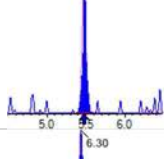
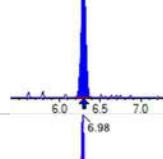
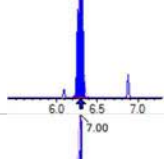
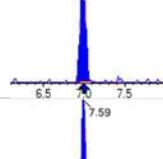
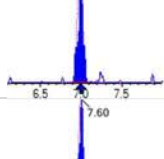
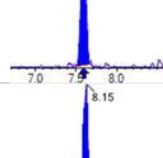
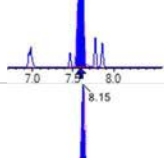
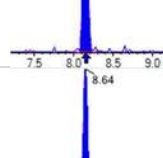
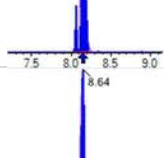
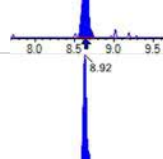
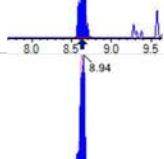
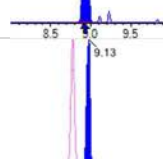
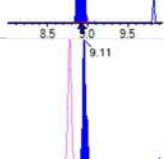
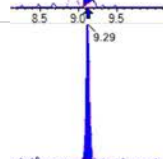
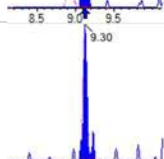
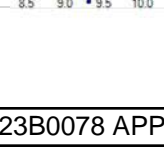
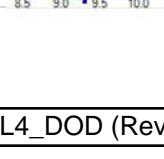
COMPOUND	CONC. (ng/L)	LOQ	DL	Q
PFBA	1.89	1.6	0.21	
PFPEA	1.01	0.80	0.065	
PFHXA	0.544	0.40	0.055	
PFHPA	0.599	0.40	0.041	
PFOA	0.515	0.40	0.15	IR2
PFNA	0.510	0.40	0.082	
PFDA	0.446	0.40	0.10	
PFUnA	0.495	0.40	0.16	
PFDOA	0.475	0.40	0.11	IR2
PFTRDA	0.498	0.40	0.20	
PFTEDA	0.546	0.40	0.20	
PFBS	0.385	0.40	0.037	J
PFPEs	0.477	0.40	0.063	
PFHXS	0.505	0.40	0.032	
PFHPS	0.419	0.40	0.051	
PFOS	0.526	0.40	0.064	
PFNS	0.430	0.40	0.12	
PFDS	0.455	0.40	0.15	
PFDOS	0.418	0.40	0.12	
4:2FTS	1.90	1.6	0.29	
6:2FTS	1.84	1.6	0.31	
8:2FTS	2.25	1.6	0.082	
PFOSA	0.542	0.40	0.10	
NMeFOSA	2.01	1.6	0.47	
NEtFOSA	1.89	1.6	0.41	
NMeFOSAA	0.495	0.40	0.11	
NEtFOSAA	0.488	0.40	0.11	
NMeFOSE	1.97	1.6	1.0	
NEtFOSE	1.77	1.6	1.0	
HFPO-DA	0.940	0.80	0.17	

# ANALYSIS DATA SHEET

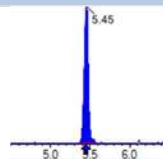
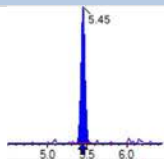
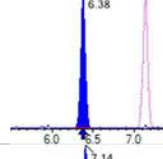
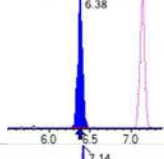
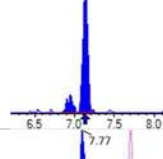
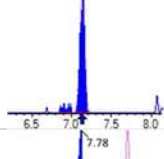
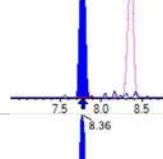
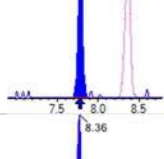
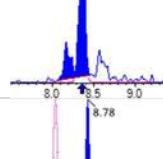
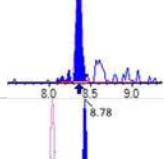
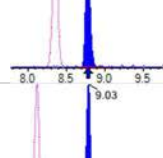
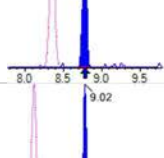
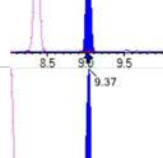
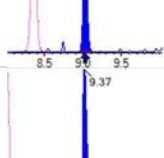
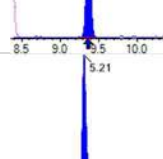
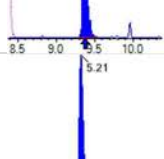
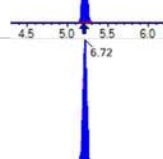
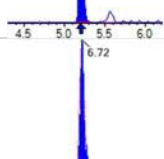
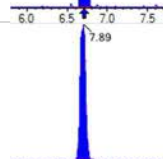
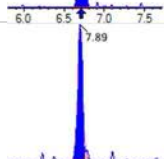
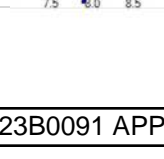
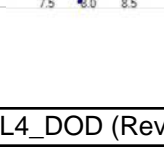
## MRL Check

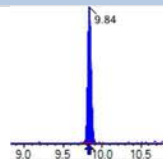
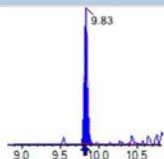
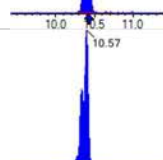
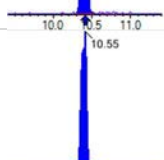
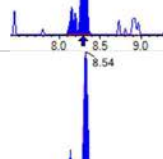
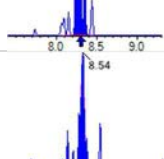
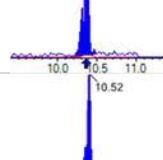
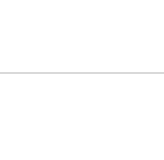
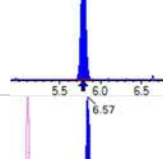
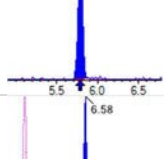
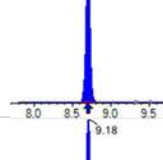
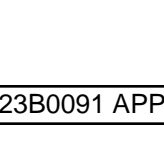
Laboratory:	APPL, LLC	Work Order:	23B0091
Client:	AECOM	Project:	Red Hill AFFF Assessment Sampling
Matrix:	Water	Laboratory ID:	BCB0232-MRL1
		File ID:	S2023-02-14D (9)
Sampled:		Prepared:	02/13/23 07:19
		Analyzed:	02/14/23 21:25
Solids:		Preparation:	EPA 1633
		Dilution:	1
Batch:	BCB0232	Sequence:	SC00636
		Calibration:	2307007
Column:	1	Instrument:	Saphira

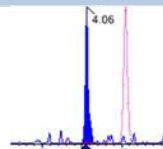
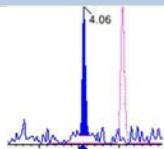
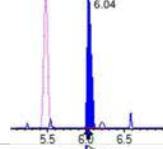
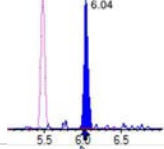
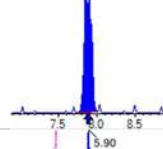
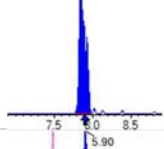
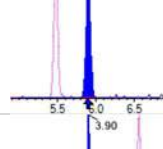
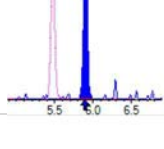
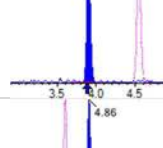
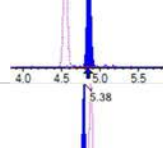
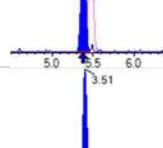
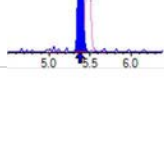
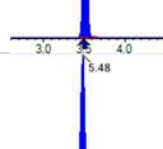
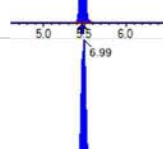
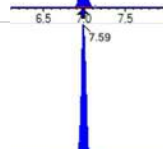

COMPOUND	CONC. (ng/L)	LOQ	DL	Q
ADONA	0.915	0.80	0.12	
PFEESA	0.810	0.80	0.11	
PFMPA	0.996	0.80	0.054	
PFMBA	0.903	0.80	0.091	
NFDHA	0.858	0.80	0.30	
9CL-PF3ONS	0.966	0.80	0.21	
11CL-PF3OUDS	0.935	0.80	0.21	
3:3FTCA	1.90	1.6	0.57	
5:3FTCA	1.66	1.6	0.44	
7:3FTCA	1.48	1.6	0.55	J

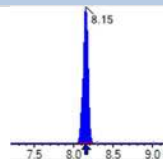
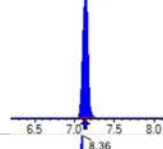
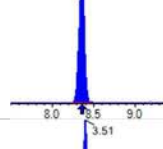
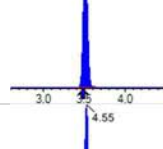
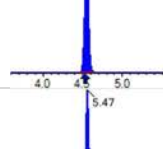
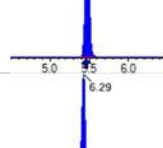
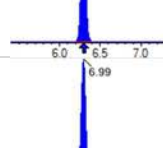
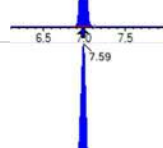
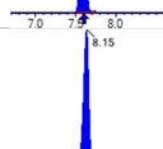
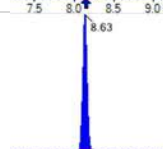
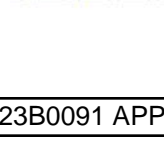
Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) ( $\Delta$ RT-[min], $\Delta$ RT- CV[min], $\Delta$ RT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBA	( 213.0 / 169.0 ) 104294	( 3.51 , 1.00 ) ( 0.00 , N/A , 0.0 )	105.5	N/A 0.0 0.0	0.4718 [ 0.4000 ]	117.9%			
PFPeA	( 263.0 / 219.0 ) 108205 ( 263.0 / 69.0 ) 1436	( 4.55 , 1.00 ) ( 0.00 , N/A , 0.4 )	392.7 34.4	0.0133 123.9 120.4	0.2515 [ 0.2000 ]	125.8%			
PFHxA	( 313.0 / 269.0 ) 77146 ( 313.0 / 119.0 ) 9873	( 5.48 , 1.00 ) ( 0.00 , N/A , -0.4 )	381.1 2323.5	0.1280 128.5 134.5	0.1360 [ 0.1000 ]	136.0%			QC,
PFHpA	( 363.0 / 319.0 ) 66676 ( 363.0 / 169.0 ) 20427	( 6.29 , 1.00 ) ( 0.00 , N/A , -0.2 )	59631.0 2069.4	0.3064 95.6 95.8	0.1499 [ 0.1000 ]	149.9%			QC,
PFOA	( 413.0 / 369.0 ) 70093 ( 413.0 / 169.0 ) 38253	( 6.98 , 1.00 ) ( -0.01 , N/A , -1.1 )	27.9 1132027.5	0.5458 163.1 151.0	0.1289 [ 0.1000 ]	128.9%			IR2,
PFNA	( 463.0 / 419.0 ) 63194 ( 463.0 / 169.0 ) 13481	( 7.59 , 1.00 ) ( 0.00 , N/A , -0.7 )	24.2 1593.1	0.2133 95.3 96.5	0.1276 [ 0.1000 ]	127.6%			
PFDA	( 513.0 / 469.0 ) 67756 ( 513.0 / 169.0 ) 9159	( 8.15 , 1.00 ) ( -0.01 , N/A , 0.0 )	213.6 216.7	0.1352 111.2 124.6	0.1116 [ 0.1000 ]	111.6%			
PFUnA	( 563.0 / 519.0 ) 68897 ( 563.0 / 169.0 ) 9112	( 8.64 , 1.00 ) ( 0.01 , N/A , 0.0 )	313.0 222.3	0.1323 126.2 113.1	0.1237 [ 0.1000 ]	123.7%			
PFDaA	( 613.0 / 569.0 ) 59482 ( 613.0 / 169.0 ) 15143	( 8.92 , 1.00 ) ( 0.00 , N/A , -0.7 )	1663.5 182.0	0.2546 155.6 151.7	0.1189 [ 0.1000 ]	118.9%			IR2,
PFTTrDA	( 663.0 / 619.0 ) 56397 ( 663.0 / 169.0 ) 16442	( 9.13 , 1.02 ) ( N/A , 0.01 , 1.1 )	24.4 192.3	0.2915 122.0 108.3	0.1245 [ 0.1000 ]	124.5%			
PFTeDA	( 713.0 / 669.0 ) 62561 ( 713.0 / 169.0 ) 13973	( 9.29 , 1.00 ) ( -0.01 , N/A , -0.6 )	32.2 116.2	0.2233 108.1 108.0	0.1366 [ 0.1000 ]	136.6%			QC,

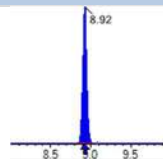
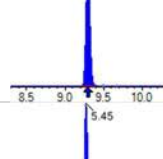
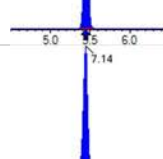
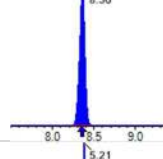
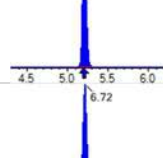
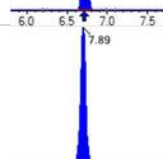
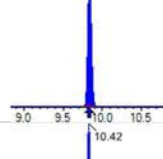
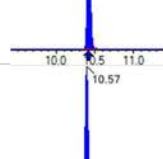
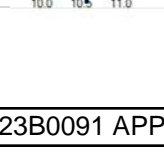
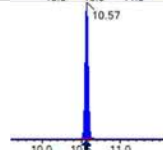
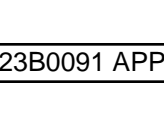


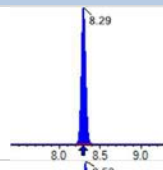
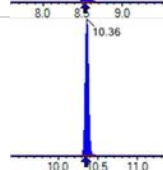
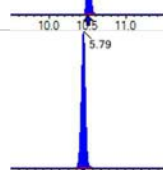


Analyte	(Q1 / Q3) Area Counts*min	R.T. (R.T [min], R.R.T.) (ΔRT-I[min], ΔRT-CV[min], ΔRT ion[s])	S / N	Ion Ratio IR Vs MP% IR Vs CV%	Concentration [ True ] ng/mL	Q.C. Rec. {Area%CV}	Primary Transition	Confirmation Transition	Flag
PFBS	( 299.0 / 80.0 ) 99671 ( 299.0 / 99.0 ) 69989	( 5.45 , 1.00 ) ( 0.00 , N/A , 0.2 )	666.3 325.8	0.7022 108.6 113.5	0.0963 [ 0.0885 ]	108.8%			
PFPeS	( 349.0 / 80.0 ) 191656 ( 349.0 / 99.0 ) 70511	( 6.38 , 0.89 ) ( N/A , 0.00 , -0.1 )	3597675.3 977.9	0.3679 103.1 106.8	0.1193 [ 0.0938 ]	127.2%			
PFHxS	( 399.0 / 80.0 ) 161800 ( 399.0 / 99.0 ) 41261	( 7.14 , 1.00 ) ( 0.00 , N/A , -0.1 )	127433.3 571796.2	0.2550 78.5 78.3	0.1264 [ 0.0911 ]	138.7%			QC,
PFHpS	( 449.0 / 80.0 ) 150068 ( 449.0 / 99.0 ) 43316	( 7.77 , 0.93 ) ( N/A , 0.00 , -0.7 )	449.5 4479.9	0.2886 106.0 103.7	0.1047 [ 0.0951 ]	110.0%			
PFOS	( 499.0 / 80.0 ) 260865 ( 499.0 / 99.0 ) 62580	( 8.36 , 1.00 ) ( 0.00 , N/A , 0.1 )	65.5 139.5	0.2399 108.4 113.1	0.1315 [ 0.0927 ]	141.8%			QC,
PFNS	( 549.0 / 80.0 ) 239893 ( 549.0 / 99.0 ) 63595	( 8.78 , 1.05 ) ( N/A , 0.00 , -0.2 )	2336.8 1612.7	0.2651 120.9 115.3	0.1076 [ 0.0960 ]	112.1%			
PFDS	( 599.0 / 80.0 ) 276029 ( 599.0 / 99.0 ) 65621	( 9.03 , 1.08 ) ( N/A , 0.01 , 0.4 )	619.5 308.9	0.2377 109.1 110.7	0.1138 [ 0.0963 ]	118.2%			
PFDoS	( 699.0 / 80.0 ) 161634 ( 699.0 / 99.0 ) 38804	( 9.37 , 1.12 ) ( N/A , 0.01 , 0.2 )	435.9 334.1	0.2401 115.3 111.9	0.1045 [ 0.0970 ]	107.8%			
4:2FTS	( 327.0 / 307.0 ) 248286 ( 327.0 / 81.0 ) 162566	( 5.21 , 1.00 ) ( 0.00 , N/A , 0.0 )	2396.4 335.6	0.6548 97.0 95.5	0.4756 [ 0.3738 ]	127.2%			
6:2FTS	( 427.0 / 407.0 ) 54026 ( 427.0 / 81.0 ) 48046	( 6.72 , 1.00 ) ( 0.00 , N/A , -0.1 )	2066.9 555.1	0.8893 107.4 115.1	0.4599 [ 0.3796 ]	121.1%			
8:2FTS	( 527.0 / 507.0 ) 71865 ( 527.0 / 81.0 ) 43051	( 7.89 , 1.00 ) ( -0.01 , N/A , -0.5 )	15.9 276.3	0.5991 72.1 77.4	0.5635 [ 0.3833 ]	147.0%			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
PFOSA	( 498.0 / 78.0 ) 367043 ( 498.0 / 478.0 ) 10818	( 9.84 , 1.00 ) ( 0.00 , N/A , 0.2 )	42.7 133.1	0.0295 143.5 139.5	0.1355 [ 0.1000 ]	135.5%			QC,
NMeFOSA	( 512.0 / 219.0 ) 214623 ( 512.0 / 169.0 ) 162033	( 10.42 , 1.00 ) ( 0.00 , N/A , 0.9 )	1199.4 912.9	0.7550 92.2 92.2	0.5023 [ 0.4000 ]	125.6%			
NEIFOSA	( 526.0 / 219.0 ) 242776 ( 526.0 / 169.0 ) 284975	( 10.57 , 1.00 ) ( 0.00 , N/A , 1.0 )	1583.7 1117.1	1.1738 92.5 94.6	0.4728 [ 0.4000 ]	118.2%			
NMeFOSAA	( 570.0 / 419.0 ) 26905 ( 570.0 / 483.0 ) 11209	( 8.31 , 1.00 ) ( 0.01 , N/A , 0.8 )	746.2 180.5	0.4166 87.8 89.9	0.1239 [ 0.1000 ]	123.9%			
NEIFOSAA	( 584.0 / 419.0 ) 24571 ( 584.0 / 526.0 ) 14606	( 8.54 , 1.00 ) ( 0.00 , N/A , 0.2 )	6171.8 771.6	0.5944 116.5 101.4	0.1220 [ 0.1000 ]	122.0%			
NMeFOSE	( 616.0 / 59.0 ) 86764	( 10.37 , 1.00 ) ( 0.01 , N/A , 0.0 )	209.9	N/A 0.0 0.0	0.4913 [ 0.4000 ]	122.8%			
NEtFOSE	( 630.0 / 59.0 ) 150533	( 10.52 , 1.00 ) ( 0.01 , N/A , 0.0 )	313.9	N/A 0.0 0.0	0.4425 [ 0.4000 ]	110.6%			
HFPO-DA	( 285.0 / 169.0 ) 73200 ( 285.0 / 185.0 ) 210658	( 5.79 , 1.00 ) ( 0.00 , N/A , 0.0 )	884.6 339.3	2.8778 104.5 103.7	0.2351 [ 0.2000 ]	117.5%			
ADONA	( 377.0 / 85.0 ) 244992 ( 377.0 / 251.0 ) 24134	( 6.57 , 1.14 ) ( N/A , 0.00 , -0.2 )	862.7 213.7	0.0985 111.7 105.5	0.2288 [ 0.1885 ]	121.3%			
9CI-Pf3ONS	( 531.0 / 351.0 ) 677278 ( 533.0 / 353.0 ) 238455	( 8.70 , 1.50 ) ( N/A , 0.01 , -0.2 )	743.4 526.2	0.3521 101.1 106.6	0.2415 [ 0.1867 ]	129.4%			
11CI-PF3OUDS	( 631.0 / 451.0 ) 407709 ( 633.0 / 453.0 ) 134269	( 9.18 , 1.59 ) ( N/A , 0.01 , -0.1 )	10013.3 2237.5	0.3293 91.0 92.4	0.2337 [ 0.1886 ]	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
3:3FTCA	( 241.0 / 177.0 ) 4742 ( 241.0 / 117.0 ) 5641	( 4.06 , 0.89 ) ( N/A , 0.00 , -0.1 )	141.7 43.3	1.1897 75.4 72.7	0.4749 [ 0.4000 ]	118.7%			
5:3FTCA	( 341.0 / 236.7 ) 32219 ( 341.0 / 217.0 ) 57456	( 6.04 , 1.10 ) ( N/A , 0.00 , 0.0 )	1211.9 178.7	1.7833 104.6 110.8	0.4153 [ 0.4000 ]	103.8%			
7:3FTCA	( 441.0 / 317.0 ) 53166 ( 441.0 / 337.0 ) 52752	( 7.88 , 1.44 ) ( N/A , -0.01 , 1.1 )	232.8 598.6	0.9922 115.3 122.0	0.3710 [ 0.4000 ]	92.8%			
PFEESA	( 315.0 / 135.0 ) 137998 ( 315.0 / 83.0 ) 40206	( 5.90 , 1.08 ) ( N/A , 0.00 , -0.2 )	557.8 136.2	0.2914 114.1 105.8	0.2025 [ 0.1785 ]	113.5%			
PFMPA	( 229.0 / 85.0 ) 23403	( 3.90 , 0.86 ) ( N/A , 0.01 , 0.0 )	278.3	N/A 0.0 0.0	0.2490 [ 0.2000 ]	124.5%			
PFMBA	( 279.0 / 85.0 ) 88998	( 4.86 , 1.07 ) ( N/A , 0.01 , 0.0 )	601.4	N/A 0.0 0.0	0.2258 [ 0.2000 ]	112.9%			
NFDHA	( 295.0 / 201.0 ) 67576 ( 295.0 / 85.0 ) 64958	( 5.38 , 0.98 ) ( N/A , 0.01 , 0.2 )	395.8 179.4	0.9613 96.0 90.9	0.2146 [ 0.2000 ]	107.3%			
13C3_PFBa_IIS	( 216.0 / 172.0 ) 250926	( 3.51 , N/A ) ( N/A , 0.01 , N/A )	1285.1	N/A	1.1203 [ 1.0000 ]	112.0% { 114.4% }			
13C2_PFHxA_IIS	( 315.0 / 270.0 ) 489204	( 5.48 , N/A ) ( N/A , 0.00 , N/A )	3066.1	N/A	0.9691 [ 1.0000 ]	96.9% { 101.8% }			
13C4_PFOA_IIS	( 417.0 / 372.0 ) 642508	( 6.99 , N/A ) ( N/A , 0.00 , N/A )	1340.6	N/A	1.0138 [ 1.0000 ]	101.4% { 105.3% }			
13C5_PFNAl_IIS	( 468.0 / 423.0 ) 604551	( 7.59 , N/A ) ( N/A , 0.00 , N/A )	3991.4	N/A	1.0005 [ 1.0000 ]	100.0% { 107.2% }			

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 ) 609817	( 8.15 , N/A ) ( N/A , 0.01 , N/A )	2081.7	N/A	1.0539 [ 1.0000 ]	105.4% { 113.9% }			
18O2_PFHxS_IIS	( 403.0 / 83.9 ) 932166	( 7.14 , N/A ) ( N/A , 0.00 , N/A )	2167.8	N/A	1.0885 [ 1.0000 ]	108.9% { 110.7% }			
13C4_PFOS_IIS	( 503.0 / 79.9 ) 1492538	( 8.36 , N/A ) ( N/A , 0.00 , N/A )	1336.1	N/A	0.9915 [ 1.0000 ]	99.1% { 111.3% }			
13C4_PFBA_EIS	( 217.0 / 172.0 ) 2134906	( 3.51 , N/A ) ( N/A , 0.01 , N/A )	4661.6	N/A	7.1527 [ 8.0000 ]	89.4% { 92.9% }			
13C5_PFPeA_EIS	( 268.0 / 223.0 ) 2091640	( 4.55 , N/A ) ( N/A , 0.01 , N/A )	3649.5	N/A	3.9078 [ 4.0000 ]	97.7% { 90.8% }			
13C5_PFHxA_EIS	( 318.0 / 273.0 ) 1290515	( 5.47 , N/A ) ( N/A , 0.00 , N/A )	1967.2	N/A	1.9259 [ 2.0000 ]	96.3% { 90.4% }			
13C4_PFHpA_EIS	( 367.0 / 322.0 ) 1053792	( 6.29 , N/A ) ( N/A , 0.00 , N/A )	7669.0	N/A	1.8150 [ 2.0000 ]	90.8% { 89.4% }			
13C8_PFOA_EIS	( 421.0 / 376.0 ) 1221794	( 6.99 , N/A ) ( N/A , 0.00 , N/A )	1352.8	N/A	1.6791 [ 2.0000 ]	84.0% { 86.0% }			
13C9_PFNA_EIS	( 472.0 / 427.0 ) 559706	( 7.59 , N/A ) ( N/A , 0.00 , N/A )	1823.7	N/A	0.8738 [ 1.0000 ]	87.4% { 92.0% }			
13C6_PFDA_EIS	( 519.0 / 474.0 ) 690264	( 8.15 , N/A ) ( N/A , 0.01 , N/A )	2544.6	N/A	0.8977 [ 1.0000 ]	89.8% { 92.6% }			
13C7_PFUnA_EIS	( 570.0 / 525.0 ) 670521	( 8.63 , N/A ) ( N/A , 0.01 , N/A )	2226.9	N/A	0.8770 [ 1.0000 ]	87.7% { 100.6% }			

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 ) 569029	( 8.92, N/A ) ( N/A, 0.01, N/A )	5649.4	N/A	0.8677 [ 1.0000 ]	86.8% { 86.4% }			
13C2_PFTeDA_EIS	( 715.0 / 670.0 ) 529347	( 9.30, N/A ) ( N/A, 0.01, N/A )	814.2	N/A	0.8810 [ 1.0000 ]	88.1% { 90.4% }			
13C3_PFBs_EIS	( 302.0 / 80.0 ) 3332569	( 5.45, N/A ) ( N/A, 0.00, N/A )	4564.5	N/A	1.7082 [ 2.0000 ]	85.4% { 87.9% }			
13C3_PFHxS_EIS	( 402.0 / 80.0 ) 1665499	( 7.14, N/A ) ( N/A, 0.00, N/A )	1639.4	N/A	1.6436 [ 2.0000 ]	82.2% { 84.7% }			
13C8_PFOS_EIS	( 507.0 / 80.0 ) 3763047	( 8.36, N/A ) ( N/A, 0.00, N/A )	1642.7	N/A	1.8365 [ 2.0000 ]	91.8% { 88.8% }			
13C2_4:2FTS_EIS	( 329.0 / 81.0 ) 764534	( 5.21, N/A ) ( N/A, 0.00, N/A )	1682.9	N/A	7.0996 [ 4.0000 ]	177.5% { 220.1% }			S2,
13C2_6:2FTS_EIS	( 429.0 / 81.0 ) 331179	( 6.72, N/A ) ( N/A, 0.00, N/A )	1339.7	N/A	2.8017 [ 4.0000 ]	70.0% { 79.6% }			
13C2_8:2FTS_EIS	( 529.0 / 81.0 ) 468317	( 7.89, N/A ) ( N/A, 0.00, N/A )	1472.4	N/A	2.8042 [ 4.0000 ]	70.1% { 77.6% }			
13C8_PFOsa_EIS	( 506.0 / 78.0 ) 6773226	( 9.84, N/A ) ( N/A, 0.01, N/A )	3187.9	N/A	1.7808 [ 2.0000 ]	89.0% { 81.5% }			
D3_NMeFOsa_EIS	( 515.0 / 169.0 ) 1135756	( 10.42, N/A ) ( N/A, 0.01, N/A )	2190.5	N/A	1.2674 [ 2.0000 ]	63.4% { 58.7% }			
D5_NEtFOsa_EIS	( 531.0 / 169.0 ) 1099261	( 10.57, N/A ) ( N/A, 0.01, N/A )	2782.7	N/A	1.3708 [ 2.0000 ]	68.5% { 61.6% }			

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 ) 1107071	( 8.29 , N/A ) ( N/A , 0.01 , N/A )	1746.6	N/A	3.3387 [ 4.0000 ]	83.5% { 83.6% }			
D5_EtFOSAA_EIS	( 589.0 / 419.0 ) 958945	( 8.53 , N/A ) ( N/A , 0.01 , N/A )	65933.1	N/A	3.5318 [ 4.0000 ]	88.3% { 90.9% }			
D7_NMeFOSE_EIS	( 623.0 / 58.9 ) 3532800	( 10.36 , N/A ) ( N/A , 0.01 , N/A )	1989.1	N/A	10.8136 [ 20.0000 ]	54.1% { 47.6% }			
D9_NEtFOSE_EIS	( 639.0 / 58.9 ) 7150182	( 10.51 , N/A ) ( N/A , 0.01 , N/A )	2028.4	N/A	17.0901 [ 20.0000 ]	85.5% { 75.9% }			
13C3_HFPODA_EIS	( 287.0 / 169.0 ) 3150017	( 5.79 , N/A ) ( N/A , 0.00 , N/A )	3732.5	N/A	7.7208 [ 8.0000 ]	96.5% { 87.8% }			

## PREPARATION BATCH SUMMARY

## EPA 1633

Laboratory: APPL, LLC

Work Order: 23B0091

Client: AECOM

Project: Red Hill AFFF Assessment Sampling

Batch: BCB0232

Batch Matrix: Water

Preparation: EPA 1633

SAMPLE NAME	LAB SAMPLE ID	DATE PREPARED	INITIAL VOL./WEIGHT mL	FINAL VOL. mL
AF-RHMW17-WGN01LF-2302W1	23B0091-01	02/13/23 07:19	566.32	2.00
AF-RHMW17D-WGN01LF-2302W1	23B0091-02	02/13/23 07:19	569.47	2.00
AF-RHMW17D-WGN01LF-2302W1	23B0091-02RE1	02/13/23 07:19	569.47	2.00
AF-RHMW17D-WQFB01-2302W1	23B0091-03	02/13/23 07:19	556.90	2.00
AF-RHMW225401-WGN01B-2302W1	23B0091-04	02/13/23 07:19	527.55	2.00
Blank	BCB0232-BLK1	02/13/23 07:19	500.00	2.00
LCS	BCB0232-BS1	02/13/23 07:19	500.00	2.00
MRL Check	BCB0232-MRL1	02/13/23 07:19	500.00	2.00

# PREPARATION BENCH SHEET

## Organics

BCB0232

Print Date/Time: 02/16/2023 11:10 am

Matrix: Water

Prepared using: PFAS - EPA 1633

Analyses		Spiking Solution(s)				Surrogate Solution(s)			
1633		23B0196 PFAS - MIX 1633 10ng/mL				23B0135 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
23B0073-01	AF-RHWW16-WGN01LF-2302W1	02/16/2023	03/08/2023	2/13/2023 7:19:00AM	562.3	2		200	
23B0073-01RE1	AF-RHWW16-WGN01LF-2302W1	02/16/2023	03/08/2023	2/13/2023 7:19:00AM	562.3	2		200	Added 2/14/2023 by PAF
23B0073-02	AF-RHWW12A-WGN01LF-2302W1	02/16/2023	03/08/2023	2/13/2023 7:19:00AM	552.35	2		200	
23B0073-02RE1	AF-RHWW12A-WGN01LF-2302W1	02/16/2023	03/08/2023	2/13/2023 7:19:00AM	552.35	2		200	Added 2/14/2023 by PAF
23B0073-03	AF-RHWW12A-WGFD01LF-2302W1	02/16/2023	03/08/2023	2/13/2023 7:19:00AM	564.77	2		200	
23B0073-03RE1	AF-RHWW12A-WGFD01LF-2302W1	02/16/2023	03/08/2023	2/13/2023 7:19:00AM	564.77	2		200	Added 2/14/2023 by PAF
23B0078-01	AF-RHWW06-WGN01LF-2302W1	02/17/2023	03/09/2023	2/13/2023 7:19:00AM	519.05	2		200	"Report relevant surrogates"
23B0078-01RE1	AF-RHWW06-WGN01LF-2302W1	02/17/2023	03/09/2023	2/13/2023 7:19:00AM	519.05	2		200	"Report relevant surrogates"
23B0078-02	AF-RHWW04-WGN01LF-2302W1	02/17/2023	03/09/2023	2/13/2023 7:19:00AM	558.31	2		200	"Report relevant surrogates"
23B0078-02RE1	AF-RHWW04-WGN01LF-2302W1	02/17/2023	03/09/2023	2/13/2023 7:19:00AM	558.31	2		200	"Report relevant surrogates"
23B0091-01	AF-RHWW17-WGN01LF-2302W1	02/20/2023	03/10/2023	2/13/2023 7:19:00AM	566.32	2		200	"Report relevant surrogates"
23B0091-01RE1	AF-RHWW17-WGN01LF-2302W1	02/20/2023	03/10/2023	2/13/2023 7:19:00AM	566.32	2		200	"Report relevant surrogates"
23B0091-02	AF-RHWW17D-WGN01LF-2302W1	02/20/2023	03/10/2023	2/13/2023 7:19:00AM	569.47	2		200	"Report relevant surrogates"
23B0091-02RE1	AF-RHWW17D-WGN01LF-2302W1	02/20/2023	03/10/2023	2/13/2023 7:19:00AM	569.47	2		200	"Report relevant surrogates"
23B0091-03	AF-RHWW17D-WQFB01-2302W1	02/20/2023	03/10/2023	2/13/2023 7:19:00AM	556.9	2		200	"Report relevant surrogates"
23B0091-03RE1	AF-RHWW17D-WQFB01-2302W1	02/20/2023	03/10/2023	2/13/2023 7:19:00AM	556.9	2		200	"Report relevant surrogates"
23B0091-04	AF-RHWW225401-WGN01B-2302W1	02/20/2023	03/10/2023	2/13/2023 7:19:00AM	527.55	2		200	"Report relevant surrogates"
23B0091-04RE1	AF-RHWW225401-WGN01B-2302W1	02/20/2023	03/10/2023	2/13/2023 7:19:00AM	527.55	2		200	"Report relevant surrogates"



PREPARATION BENCH SHEET

Organics

Print Date/Time: 02/16/2023 11:10 am

BCB0232

(Continued)

Matrix: Water

Prepared using: PFAS - EPA 1633

Analyses		Spiking Solution(s)				Surrogate Solution(s)	
1633		23B0196	PFAS - MIX 1633 10ng/mL			23B0135	MPFAC-HIF-ES 20.0ng/mL
BCB0232-BLK1	Blank		2/13/2023 7:19:00AM	500	2	0	200
BCB0232-BS1	LCS		2/13/2023 7:19:00AM	500	2	200	200
BCB0232-MRL1	MRL Check		2/13/2023 7:19:00AM	500	2	20	200

Start Date/Time

Stop Date/Time

Reagents		
Standard	Description	LotNum
22C0296	Envi-Carb	122395
22L0094	Reagent - 0.05MFA wash	x
23A0369	Reagent -0.3M Formic Acid	M13H051
23A0442	Am. Ac. preservative	*

Batch Comments:

Spiked by: LYA 2/13/23 8:25am W/tness: DAG

Balance #: WB2

Cartridge: Oasis Waters

Concentration: 2/13/23 425pm - 5:14pm

Preparation Reviewed By

Date

Extracts Received By

Date

Spiking Witnessed By

## INJECTION LOG - ANALYSIS SEQUENCE SUMMARY

EPA 1633

Laboratory: APPL, LLC

SDG:

Client: AECOM

Project: Red Hill AFFF Assessment Sampling

Sequence: SC00636

Instrument: Saphira

Calibration: 2307007

Sample Name	Lab Sample ID	Lab File ID	Analysis Date/Time
Calibration Blank	SC00636-CCB1	S2023-02-14D (1)	02/14/23 19:41
Low Cal Check	SC00636-LCV1	S2023-02-14D (2)	02/14/23 19:54
Calibration Check	SC00636-CCV1	S2023-02-14D (3)	02/14/23 20:07
Calibration Check	SC00636-CCV2	S2023-02-14D (6)	02/14/23 20:46
Blank	BCB0232-BLK1	S2023-02-14D (7)	02/14/23 20:59
LCS	BCB0232-BS1	S2023-02-14D (8)	02/14/23 21:12
MRL Check	BCB0232-MRL1	S2023-02-14D (9)	02/14/23 21:25
AF-RHMW17-WGN01LF-2302W1	23B0091-01	S2023-02-14D (20)	02/14/23 23:47
AF-RHMW17D-WGN01LF-2302W1	23B0091-02	S2023-02-14D (22)	02/15/23 00:13
AF-RHMW17D-WGN01LF-2302W1	23B0091-02RE1	S2023-02-14D (23)	02/15/23 00:26
AF-RHMW17D-WQFB01-2302W1	23B0091-03	S2023-02-14D (24)	02/15/23 00:39
AF-RHMW225401-WGN01B-2302W1	23B0091-04	S2023-02-14D (26)	02/15/23 01:04
Calibration Check	SC00636-CCV3	S2023-02-14D (28)	02/15/23 10:11
Calibration Blank	SC00636-CCB3	S2023-02-14D (29)	02/15/23 10:24
Calibration Check	SC00636-CCV4	S2023-02-14D (35)	02/15/23 11:41
Calibration Blank	SC00636-CCB4	S2023-02-14D (36)	02/15/23 11:54

## INJECTION LOG - ANALYSIS SEQUENCE SUMMARY

EPA 1633

Laboratory: APPL, LLC

SDG:

Client: AECOM

Project: Red Hill AFFF Assessment Sampling

Sequence: SC00647

Instrument: Saphira

Calibration: 2307007

Sample Name	Lab Sample ID	Lab File ID	Analysis Date/Time
Cal Standard	SC00647-CAL1	S2023-02-14C (1)	02/14/23 17:19
Cal Standard	SC00647-CAL2	S2023-02-14C (2)	02/14/23 17:32
Cal Standard	SC00647-CAL3	S2023-02-14C (3)	02/14/23 17:45
Cal Standard	SC00647-CAL4	S2023-02-14C (4)	02/14/23 17:58
Cal Standard	SC00647-CAL5	S2023-02-14C (5)	02/14/23 18:10
Cal Standard	SC00647-CAL6	S2023-02-14C (6)	02/14/23 18:23
Cal Standard	SC00647-CAL7	S2023-02-14C (7)	02/14/23 18:36
Cal Standard	SC00647-CAL8	S2023-02-14C (8)	02/14/23 18:49
Initial Cal Blank	SC00647-ICB1	S2023-02-14C (9)	02/14/23 19:02
Secondary Cal Check	SC00647-SCV1	S2023-02-14C (10)	02/14/23 19:15



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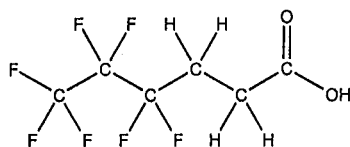
## CERTIFICATE OF ANALYSIS DOCUMENTATION

**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  $^{19}\text{F}$  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:**

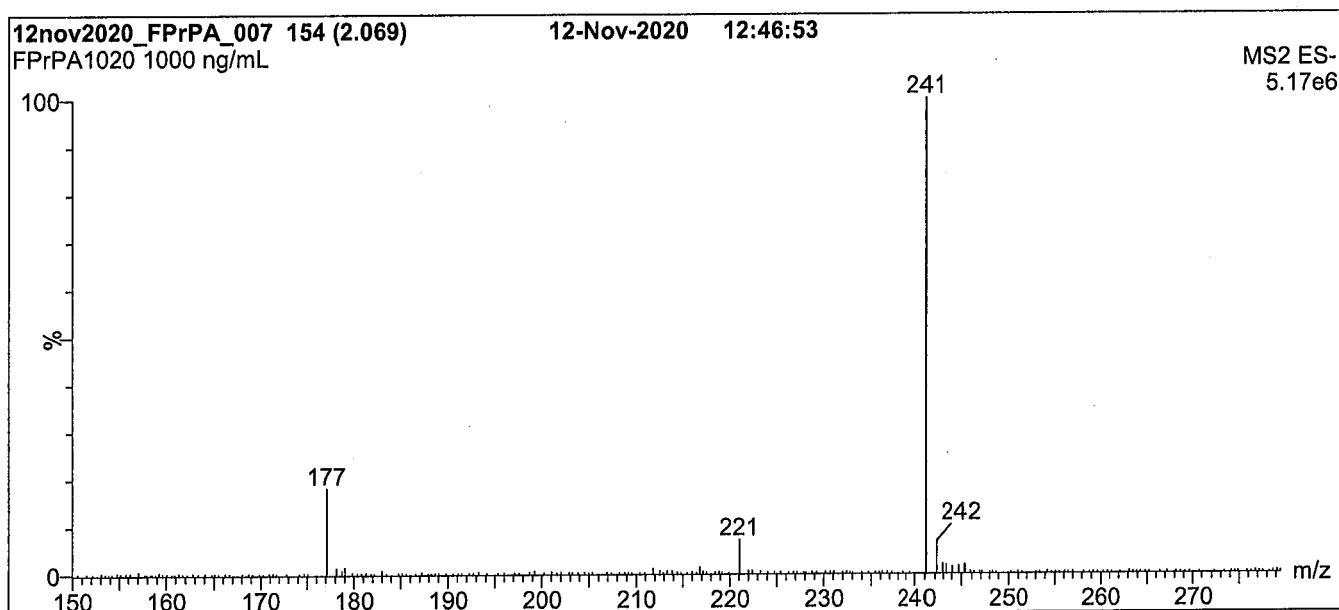
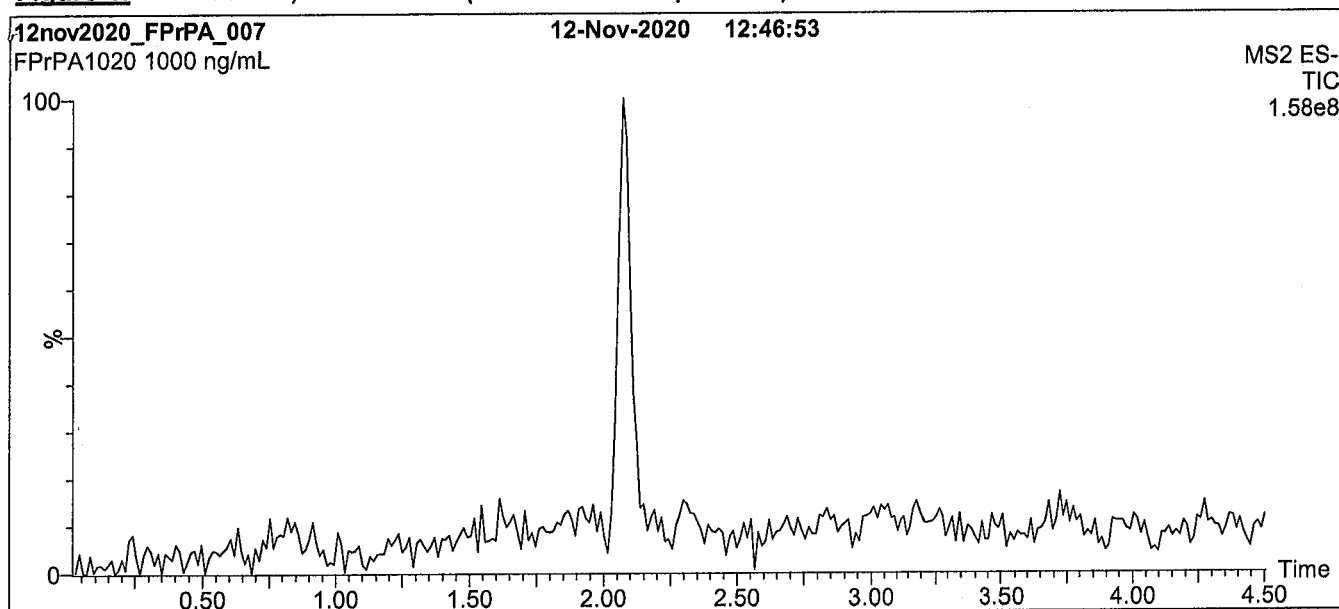
At the time of shipment, all products are warranted to be free of defects in material and workmanship and to conform to the stated technical and purity specifications.

**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](http://www.well-labs.com) or contact us directly at [info@well-labs.com](mailto: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 RP<sub>18</sub>  
1.7  $\mu$ m, 2.1 x 100 mm

Mobile phase: Gradient

Start: 60% H<sub>2</sub>O / 40% (80:20 MeOH:ACN)  
(both with 10 mM NH<sub>4</sub>OAc 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  $\mu$ L/min

**MS Parameters:**

Experiment: Full Scan (150 - 850 amu)

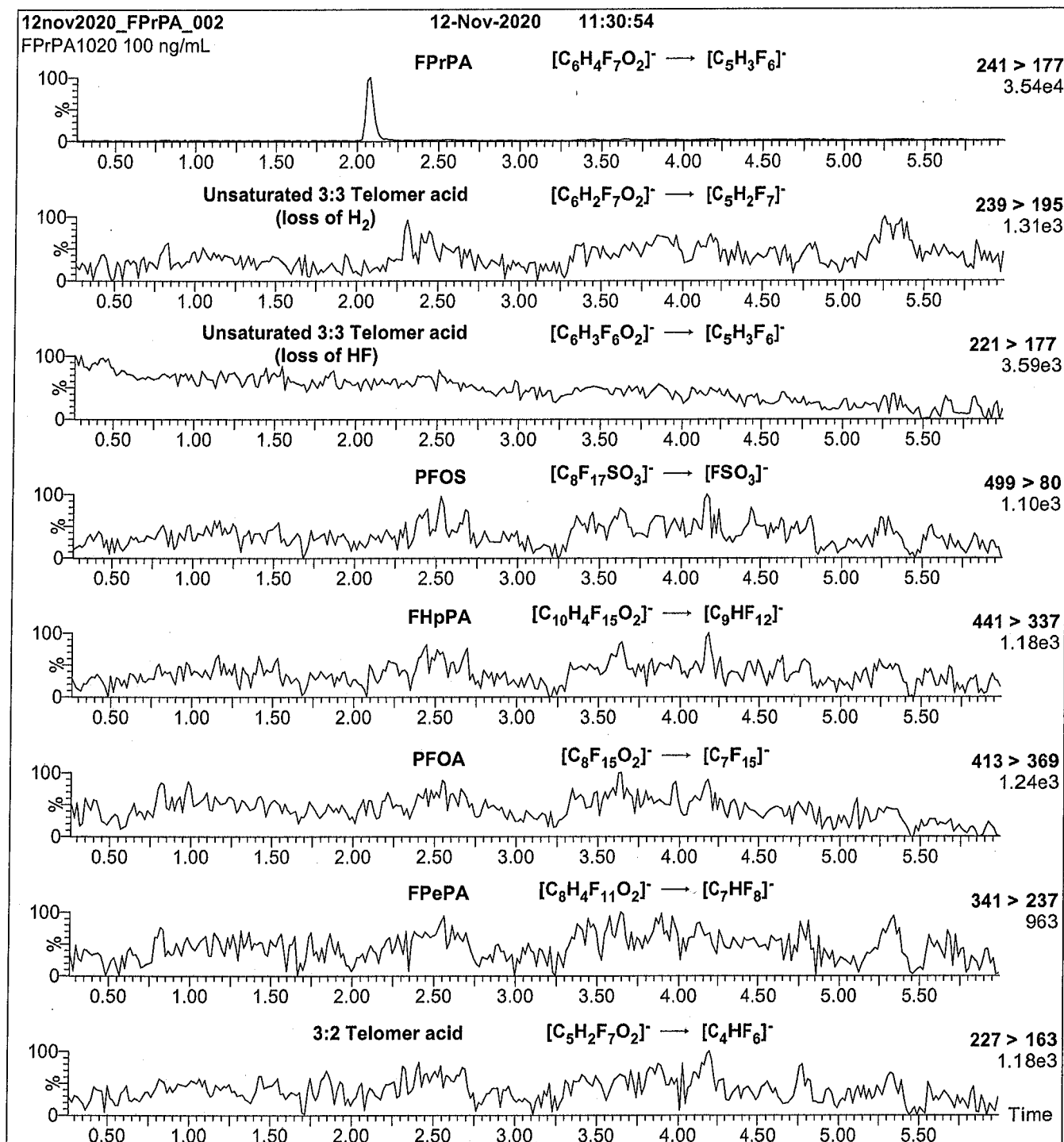
Source: Electrospray (negative)

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  $\mu$ 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

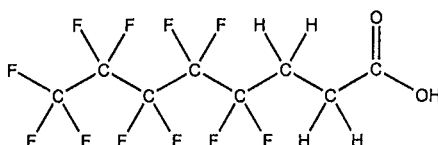
## CERTIFICATE OF ANALYSIS DOCUMENTATION

**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  $^{19}\text{F}$  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:**

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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:**

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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**EXPIRY DATE / PERIOD OF VALIDITY:**

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**LIMITED WARRANTY:**

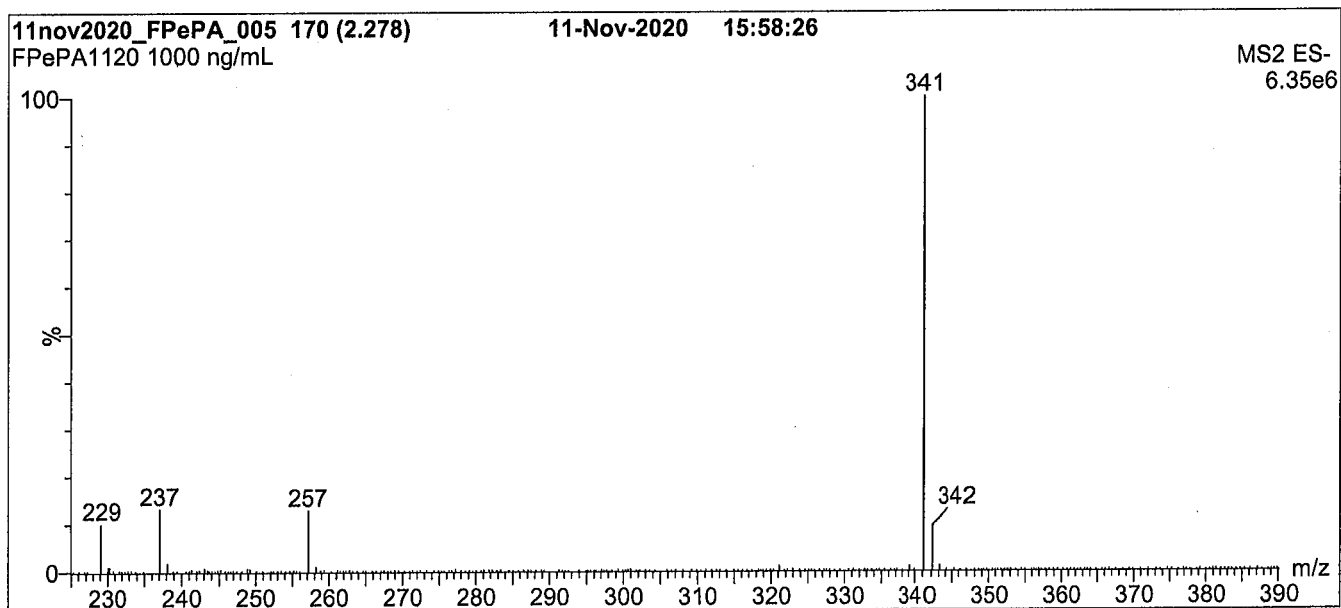
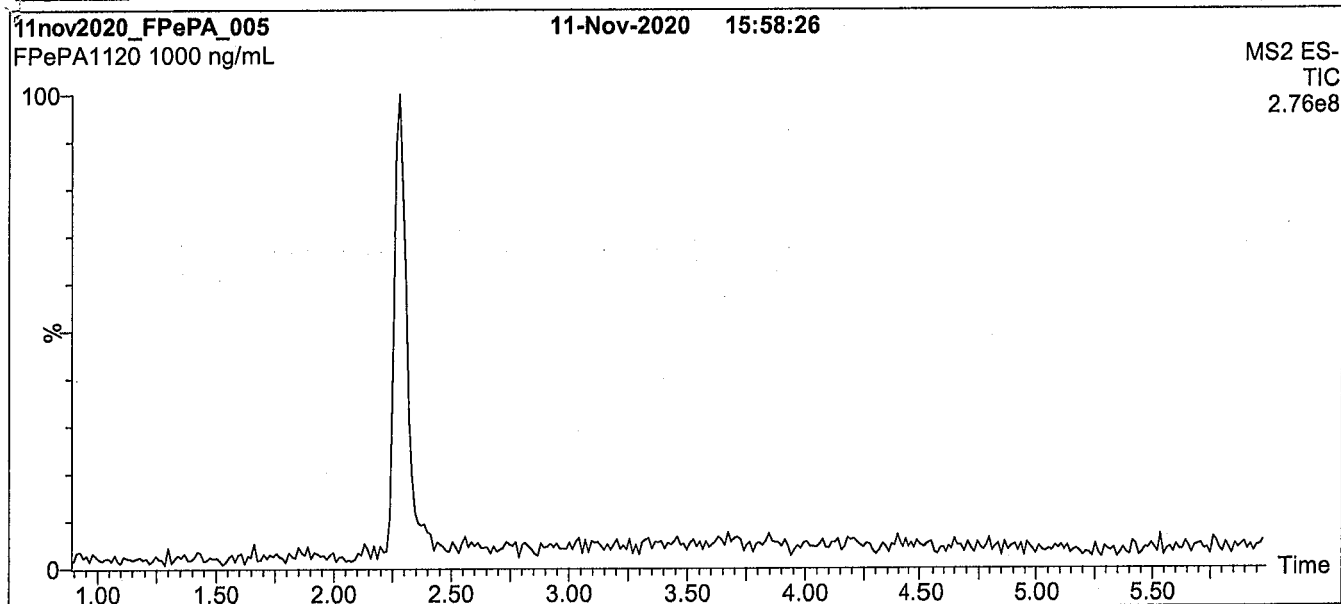
At the time of shipment, all products are warranted to be free of defects in material and workmanship and to conform to the stated technical and purity specifications.

**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).



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**Figure 1:** FPePA; 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 RP<sub>18</sub>  
1.7  $\mu$ m, 2.1 x 100 mm

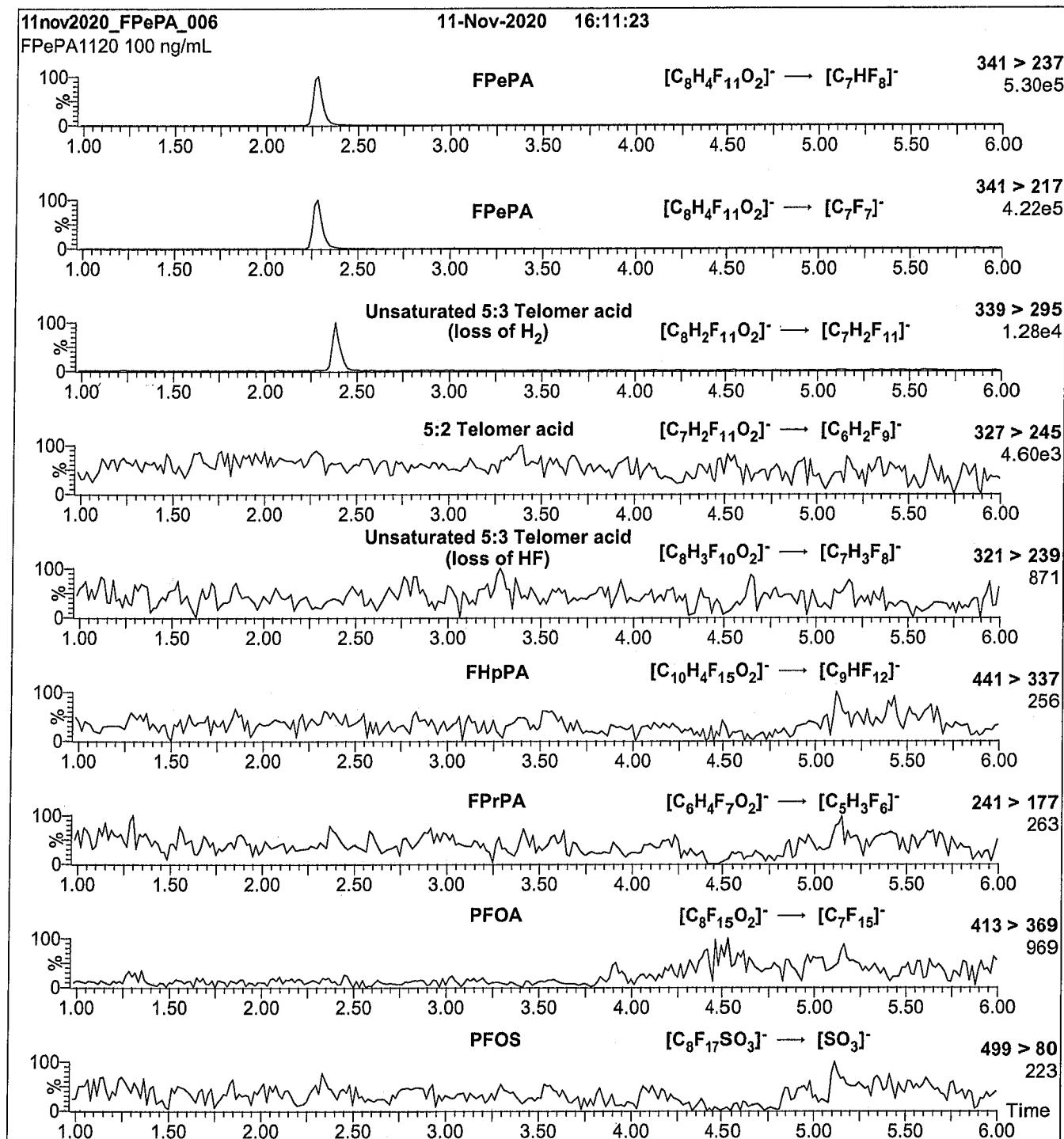
Mobile phase: Gradient  
Start: 45% H<sub>2</sub>O / 55% (80:20 MeOH:ACN)  
(both with 10 mM NH<sub>4</sub>OAc buffer)  
Ramp to 90% organic over 8 min and hold for  
2 min before returning to initial conditions in 0.75 min.  
Time: 12 min

Flow: 300  $\mu$ 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  $\mu$ 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



# WELLINGTON LABORATORIES

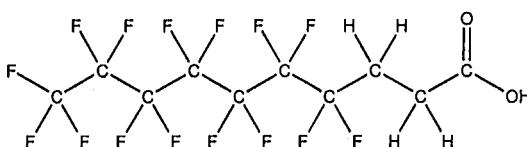
## CERTIFICATE OF ANALYSIS DOCUMENTATION

**PRODUCT CODE:** FHpPA  
**COMPOUND:** 3-Perfluoroheptyl propanoic acid

**LOT NUMBER:** FHpPA1020

**STRUCTURE:**

**CAS #:** 812-70-4



**MOLECULAR FORMULA:**  $C_{10}H_6F_{16}O_2$   
**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.

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Certified By:   
B.G. Chittim, General Manager

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

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**LIMITED WARRANTY:**

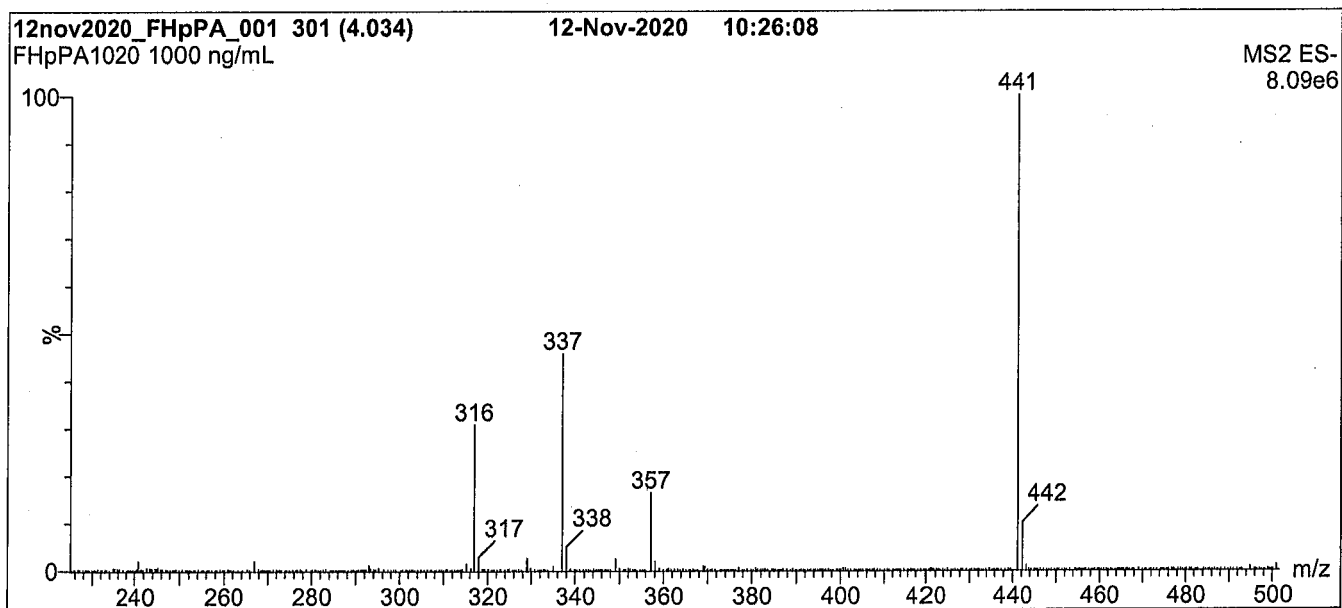
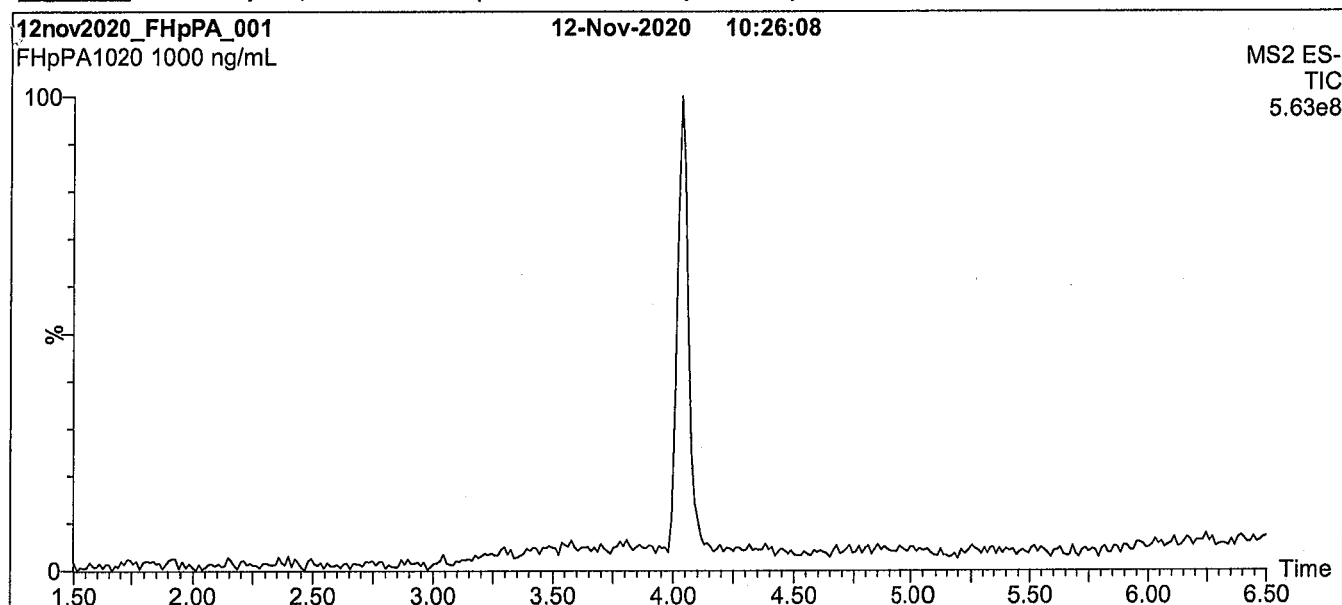
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**Figure 1:** FHpPA; LC/MS Data (TIC and Mass Spectrum)**Conditions for Figure 1:**

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

**Chromatographic Conditions:**

Column: Acquity UPLC BEH Shield RP<sub>18</sub>  
1.7  $\mu$ m, 2.1 x 100 mm

Mobile phase: Gradient

Start: 45% H<sub>2</sub>O / 55% (80:20 MeOH:ACN)  
(both with 10 mM NH<sub>4</sub>OAc 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  $\mu$ L/min

**MS Parameters:**

Experiment: Full Scan (225 - 850 amu)

Source: Electrospray (negative)

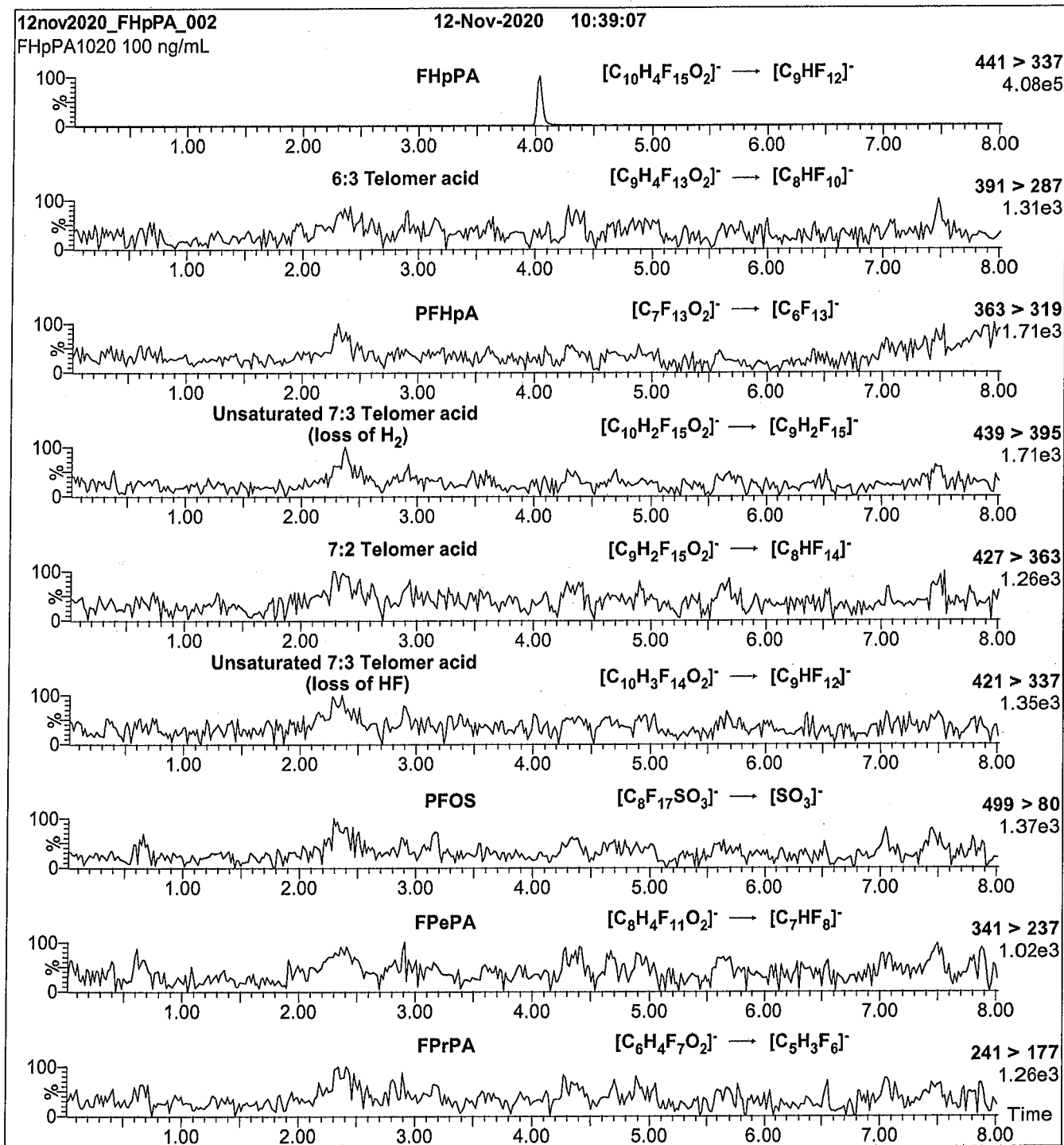
Capillary Voltage (kV) = 0.50

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  $\mu$ 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



# WELLINGTON LABORATORIES

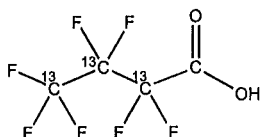
## CERTIFICATE OF ANALYSIS DOCUMENTATION

**PRODUCT CODE:** M3PFBA  
**COMPOUND:** Perfluoro-n-(2,3,4-<sup>13</sup>C<sub>3</sub>)butanoic acid

**LOT NUMBER:** M3PFBA0721

**STRUCTURE:**

**CAS #:** Not available



**MOLECULAR FORMULA:** <sup>13</sup>C<sub>3</sub><sup>12</sup>CHF<sub>7</sub>O<sub>2</sub>  
**CONCENTRATION:** 50.0 ± 2.5 µg/mL

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

**CHEMICAL PURITY:** >98%

**ISOTOPIC PURITY:** ≥99%<sup>13</sup>C  
(2,3,4-<sup>13</sup>C<sub>3</sub>)

**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-(<sup>13</sup>C<sub>3</sub>)propanoic acid and also contains ~1.0% of perfluoro-n-(1,2,3,4-<sup>13</sup>C<sub>4</sub>)butanoic acid due to the naturally occurring isotopic abundance of <sup>13</sup>C in the unlabelled carbon atom.

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**Certified By:**   
B.G. Chittim, General Manager

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

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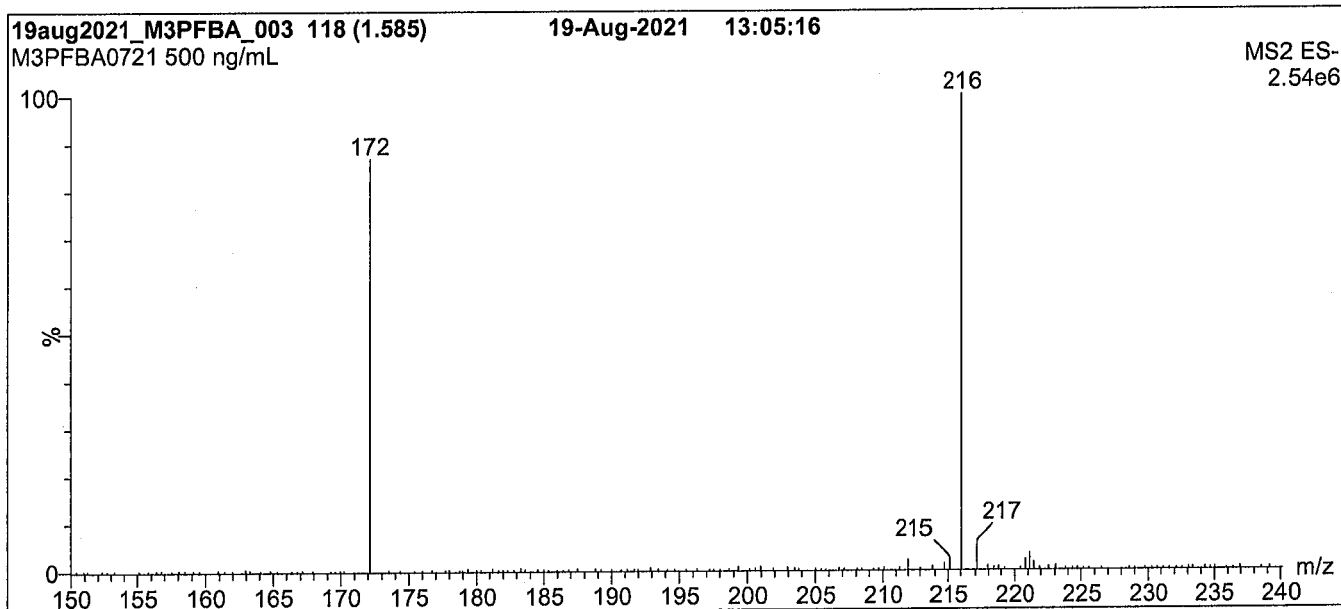
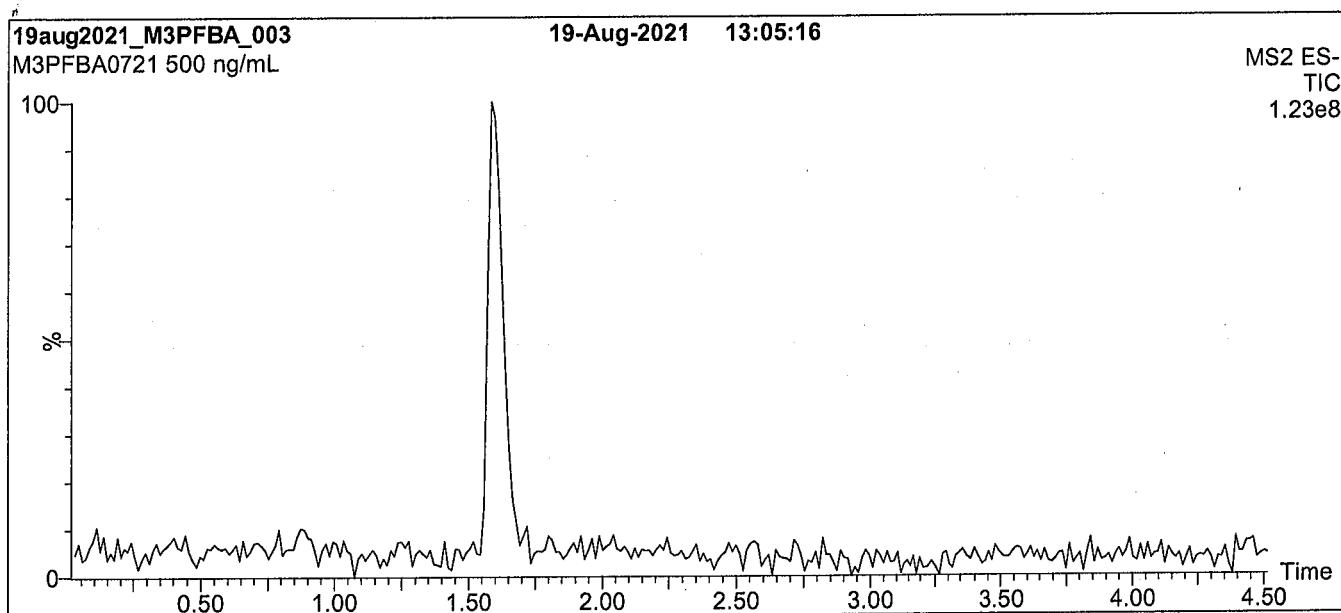
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**Figure 1: M3PFBA; LC/MS Data (Full Scan and Mass Spectrum)****Conditions for Figure 1:**

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

**Chromatographic Conditions:**

Column: Acquity UPLC BEH Shield RP<sub>18</sub>  
1.7  $\mu$ m, 2.1 x 100 mm

Mobile phase: Gradient

Start: 60% H<sub>2</sub>O / 40% (80:20 MeOH:ACN)  
(both with 10 mM NH<sub>4</sub>OAc 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  $\mu$ L/min

**MS Parameters:**

Experiment: Full Scan (150 - 850 amu)

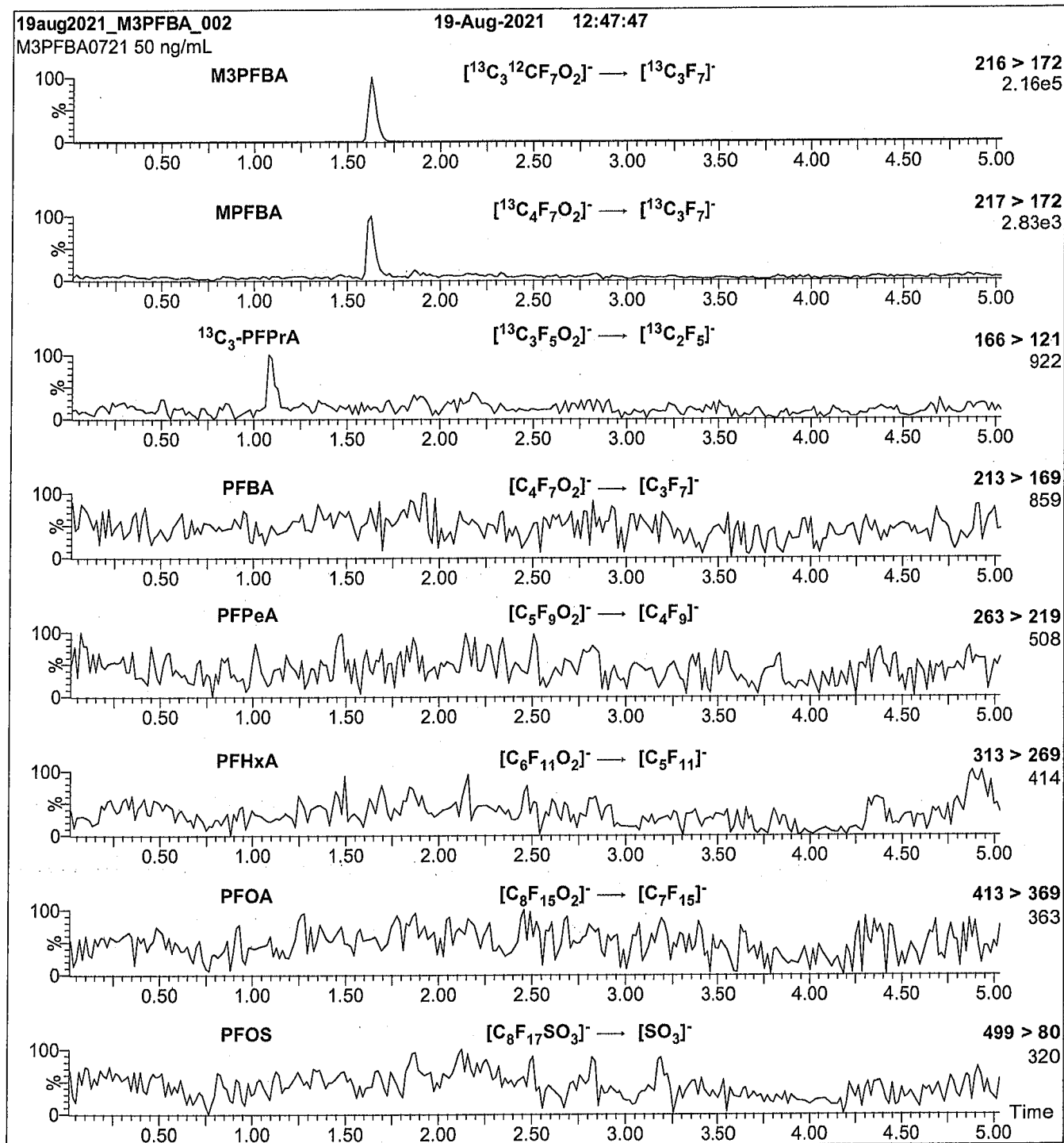
Source: Electrospray (negative)

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

Expires: 08/19/2026

Standard Type: Analyte Spike

Prepared: 08/19/2021

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
13C3-PFBA		13C3-PFBA	50	ug/mL



# WELLINGTON LABORATORIES

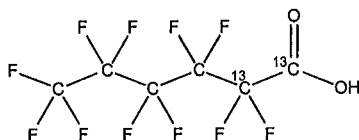
## CERTIFICATE OF ANALYSIS DOCUMENTATION

**PRODUCT CODE:** MPFHxA  
**COMPOUND:** Perfluoro-n-(1,2-<sup>13</sup>C<sub>2</sub>)hexanoic acid

**LOT NUMBER:** MPFHxA0921

**STRUCTURE:**

**CAS #:** 960315-47-3



**MOLECULAR FORMULA:** <sup>13</sup>C<sub>2</sub><sup>12</sup>C<sub>4</sub>HF<sub>11</sub>O<sub>2</sub>  
**CONCENTRATION:** 50.0 ± 2.5 µg/mL

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

**CHEMICAL PURITY:** >98%

**ISOTOPIC PURITY:** ≥99% <sup>13</sup>C  
(1,2-<sup>13</sup>C<sub>2</sub>)

**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:**

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

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

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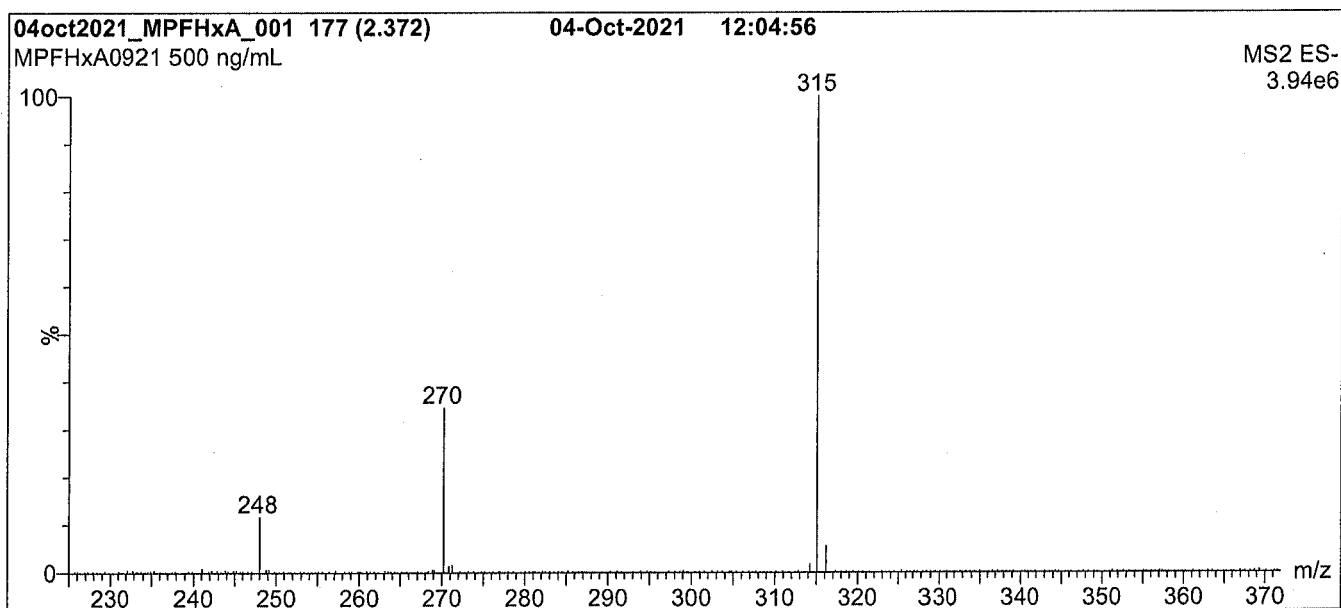
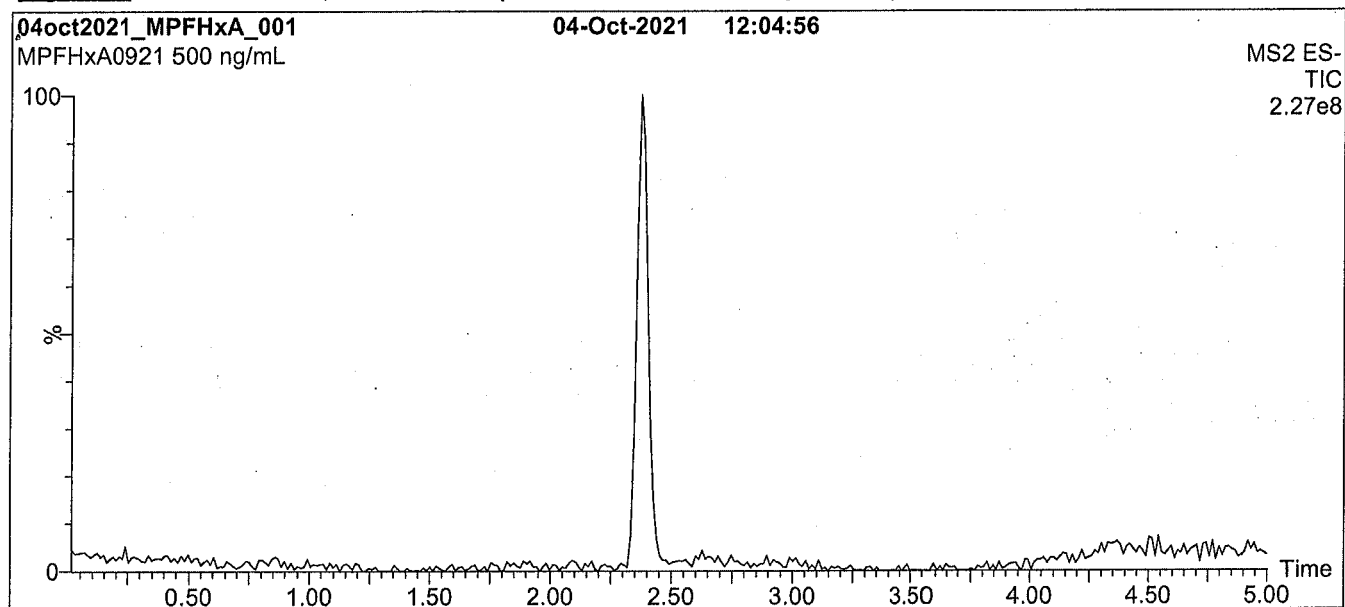
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**Figure 1: MPFHxA; LC/MS Data (Full Scan and Mass Spectrum)****Conditions for Figure 1:**

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

**Chromatographic Conditions:**

Column: Acquity UPLC BEH Shield RP<sub>18</sub>  
1.7  $\mu$ m, 2.1 x 100 mm

Mobile phase: Gradient

Start: 50% H<sub>2</sub>O / 50% (80:20 MeOH:ACN)  
(both with 10 mM NH<sub>4</sub>OAc 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  $\mu$ L/min

**MS Parameters:**

Experiment: Full Scan (225 - 850 amu)

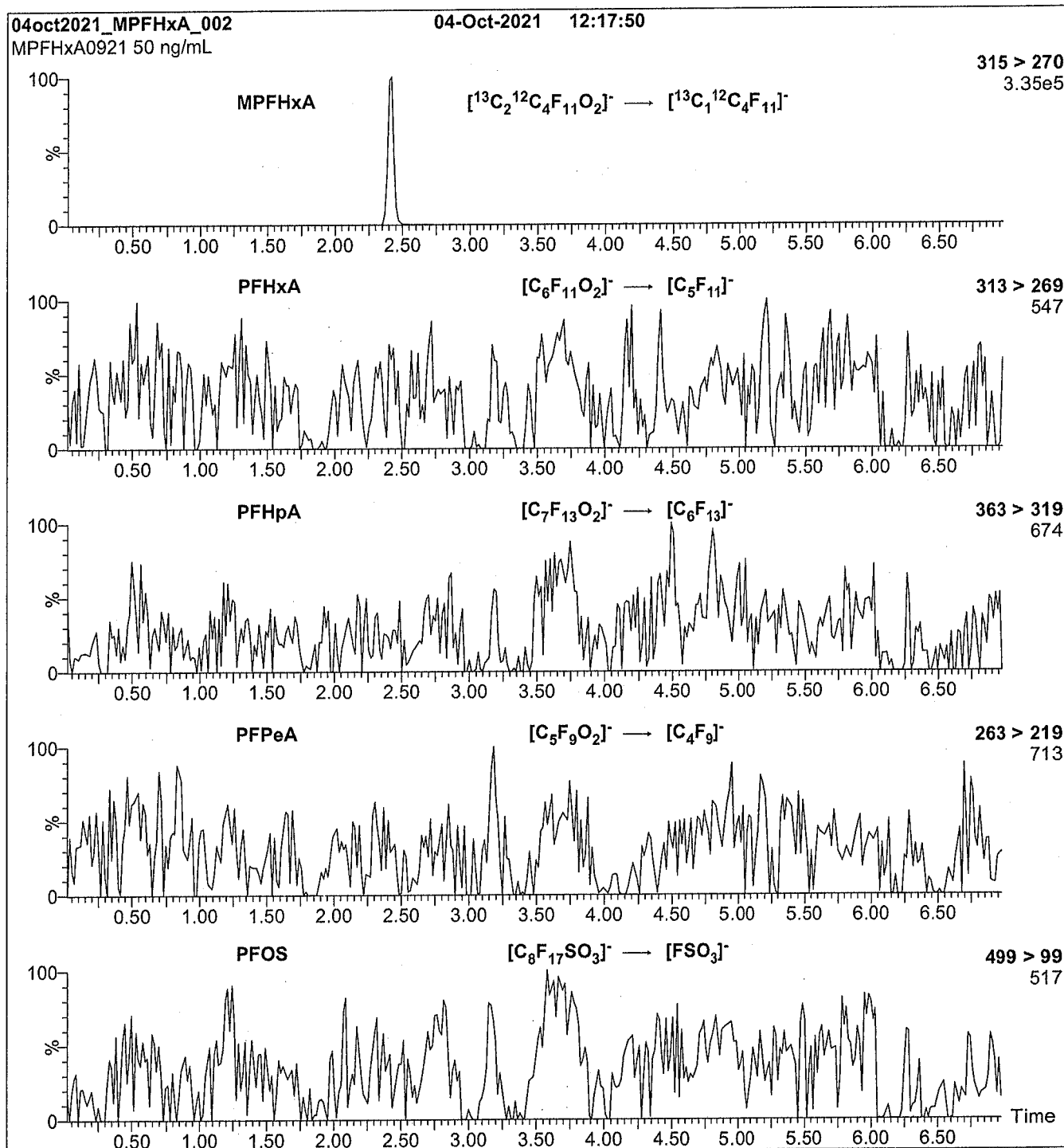
Source: Electrospray (negative)

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

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Description: PFAS - IIS MPFHxA 50ug/mL  
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# WELLINGTON LABORATORIES

## CERTIFICATE OF ANALYSIS DOCUMENTATION

**PRODUCT CODE:**

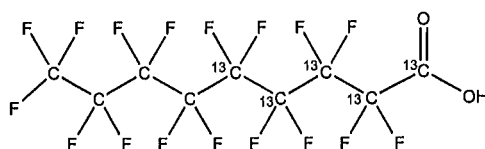
MPFNA

**LOT NUMBER:**

MPFNA1021

**COMPOUND:**Perfluoro-n-(1,2,3,4,5-<sup>13</sup>C<sub>5</sub>)nonanoic acid**STRUCTURE:****CAS #:**

960315-49-5

**MOLECULAR FORMULA:**<sup>13</sup>C<sub>5</sub><sup>12</sup>C<sub>4</sub>HF<sub>17</sub>O<sub>2</sub>**MOLECULAR WEIGHT:**

469.04

**CONCENTRATION:**

50.0 ± 2.5 µg/mL

**SOLVENT(S):**

Methanol

Water (&lt;1%)

**CHEMICAL PURITY:**

&gt;98%

**ISOTOPIC PURITY:**≥99% <sup>13</sup>C(1,2,3,4,5-<sup>13</sup>C<sub>5</sub>)**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.

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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

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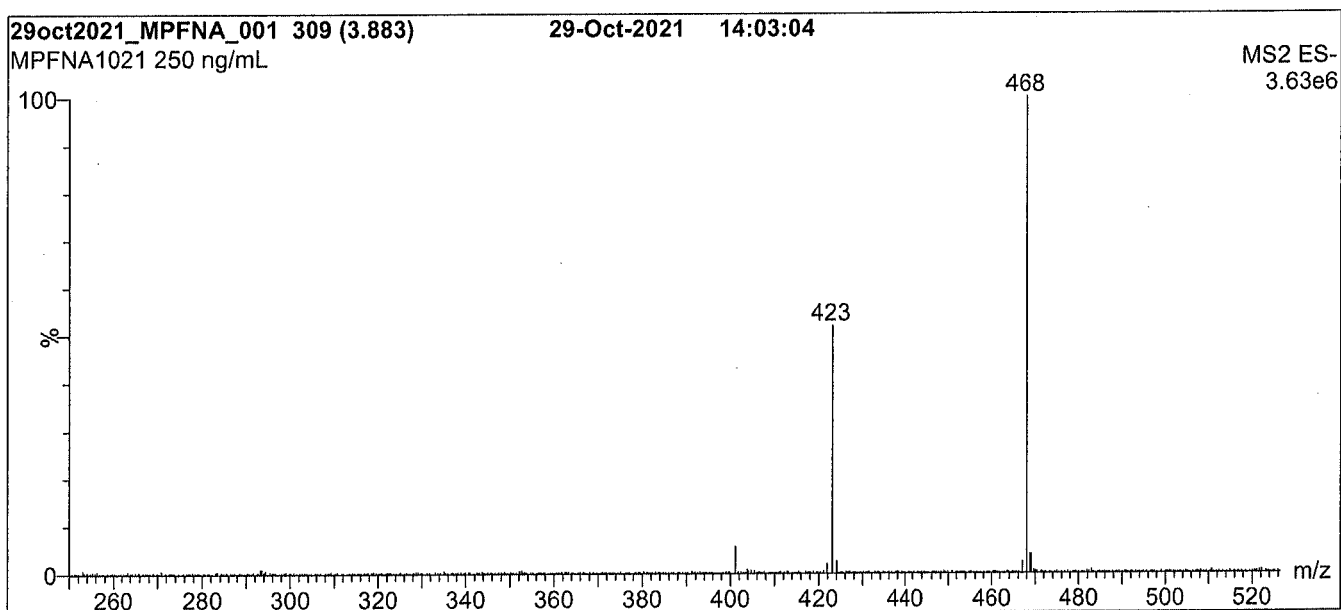
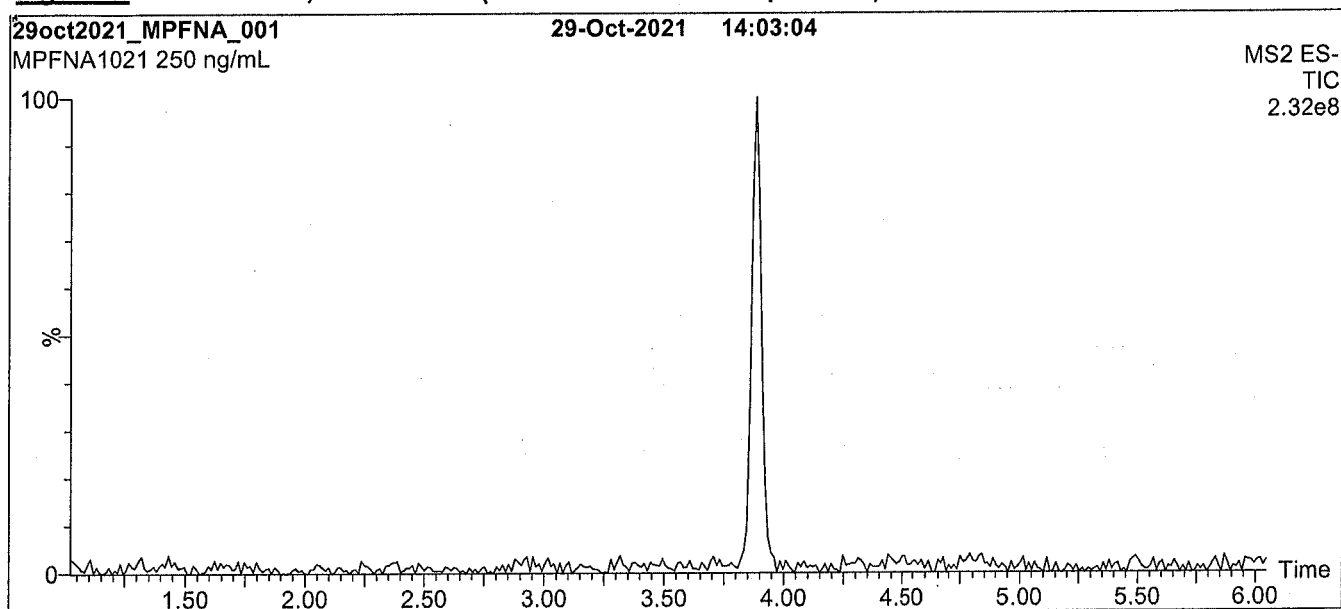
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**Figure 1: MPFNA; LC/MS Data (Full Scan and Mass Spectrum)****Conditions for Figure 1:**

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

**Chromatographic Conditions:**

Column: Acquity UPLC BEH Shield RP<sub>18</sub>  
1.7  $\mu$ m, 2.1 x 100 mm

Mobile phase: Gradient

Start: 40% H<sub>2</sub>O / 60% (80:20 MeOH:ACN)  
(both with 10 mM NH<sub>4</sub>OAc 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  $\mu$ L/min

**MS Parameters:**

Experiment: Full Scan (250 - 850 amu)

Source: Electrospray (negative)

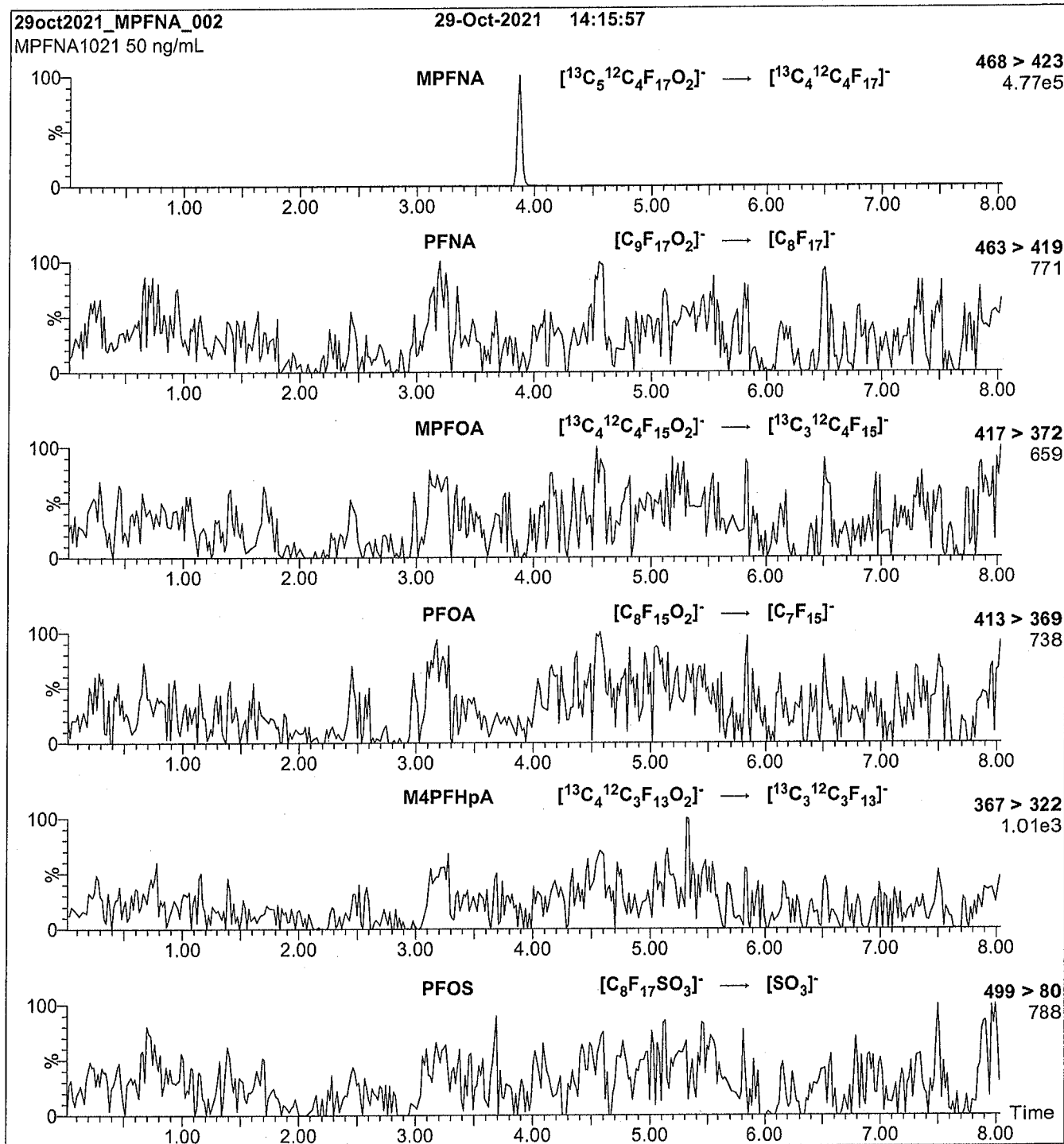
Capillary Voltage (kV) = 2.00

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	Expires:	10/29/2026
Standard Type:	Analyte Spike	Prepared:	10/29/2021
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
13C5-PFNA		13C5-PFNA	50	ug/mL

# 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



# WELLINGTON LABORATORIES

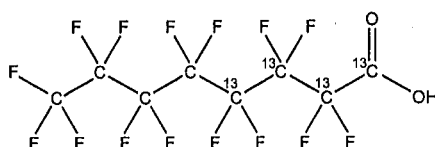
## CERTIFICATE OF ANALYSIS DOCUMENTATION

**PRODUCT CODE:** MPFOA  
**COMPOUND:** Perfluoro-n-(1,2,3,4-<sup>13</sup>C<sub>4</sub>)octanoic acid

**LOT NUMBER:** MPFOA1121

**STRUCTURE:**

**CAS #:** 960315-48-4



**MOLECULAR FORMULA:** <sup>13</sup>C<sub>4</sub><sup>12</sup>C<sub>4</sub>HF<sub>15</sub>O<sub>2</sub>  
**CONCENTRATION:** 50.0 ± 2.5 µg/mL

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

**CHEMICAL PURITY:** >98%

**ISOTOPIC PURITY:** ≥99% <sup>13</sup>C  
(1,2,3,4-<sup>13</sup>C<sub>4</sub>)

**LAST TESTED:** (mm/dd/yyyy) 12/07/2021

**EXPIRY DATE:** (mm/dd/yyyy) 12/07/2026

**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:** 12/20/2021  
(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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where  $x$  is expressed as a relative standard uncertainty of the individual parameter.

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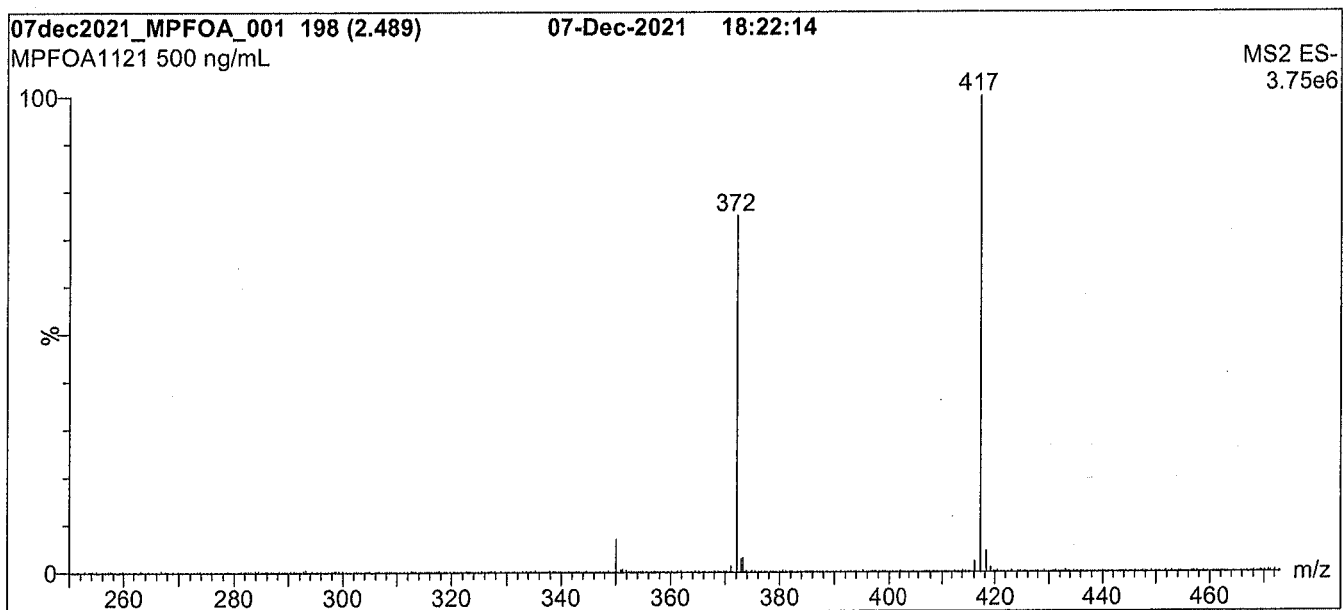
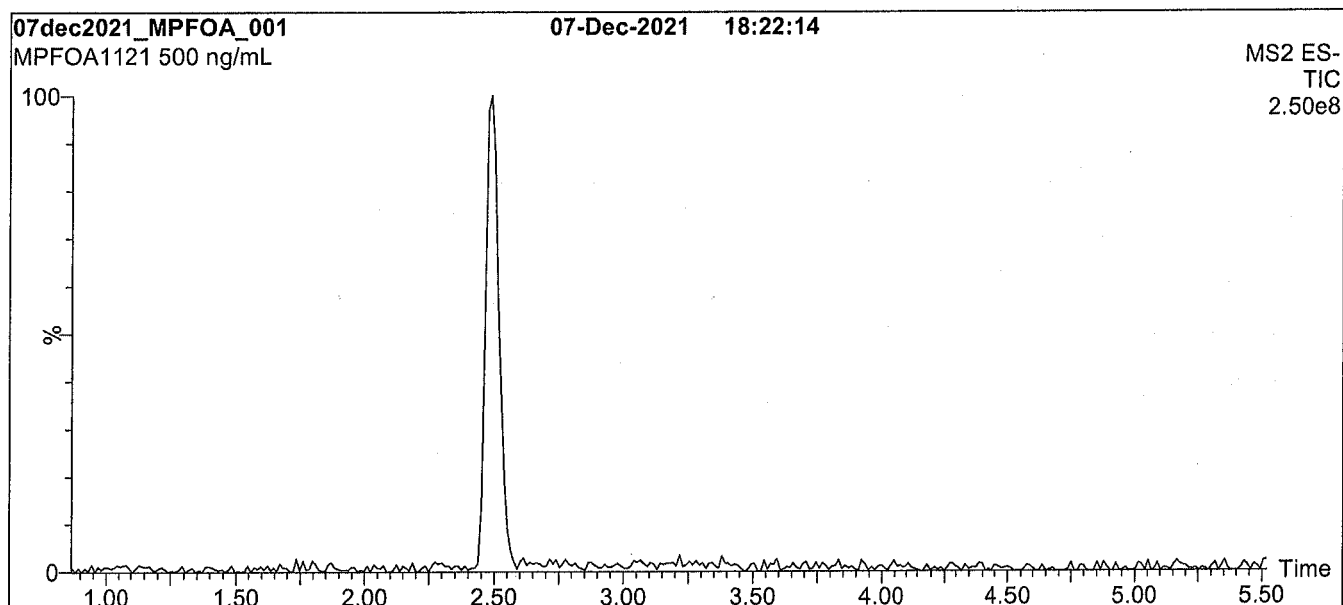
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**Figure 1: MPFOA; LC/MS Data (Full Scan and Mass Spectrum)****Conditions for Figure 1:**

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

**Chromatographic Conditions:**

Column: Acquity UPLC BEH Shield RP<sub>18</sub>  
1.7  $\mu$ m, 2.1 x 100 mm

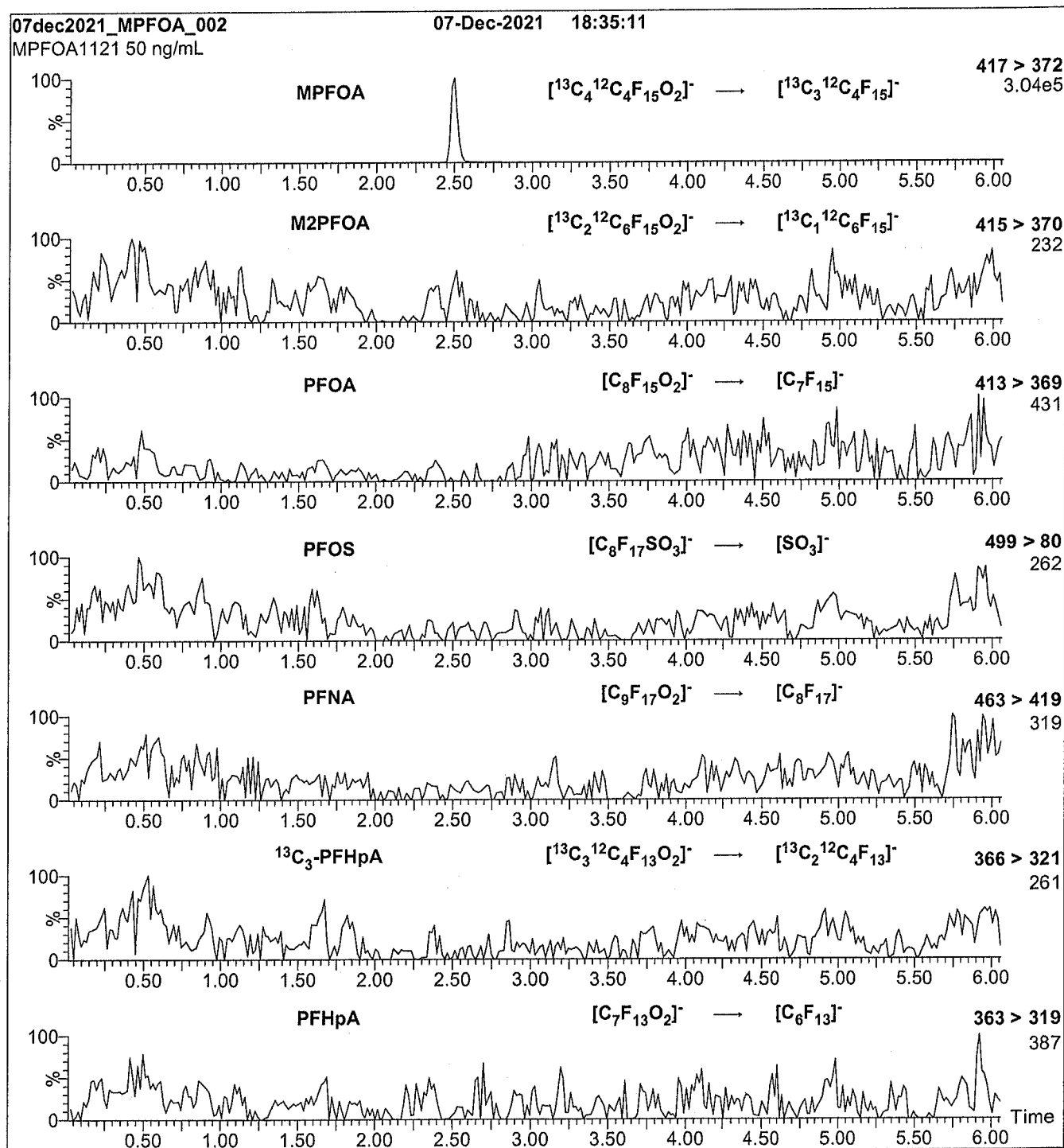
Mobile phase: Gradient  
Start: 40% H<sub>2</sub>O / 60% (80:20 MeOH:ACN)  
(both with 10 mM NH<sub>4</sub>OAc buffer)  
Ramp to 90% organic over 8 min and hold for 2 min  
before returning to initial conditions in 0.75 min.  
Time: 12 min

Flow: 300  $\mu$ 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  
Standard Type: Analyte Spike  
Solvent: MeOH  
Final Volume (mLs): 1.2  
Vials: 1

Expires: 12/07/2026  
Prepared: 12/07/2021  
Prepared By: Dipti Gokal  
Department: PFAS  
Last Edit: 01/20/2022 15:48 by HGH

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





# WELLINGTON LABORATORIES

## CERTIFICATE OF ANALYSIS DOCUMENTATION

**PRODUCT CODE:**

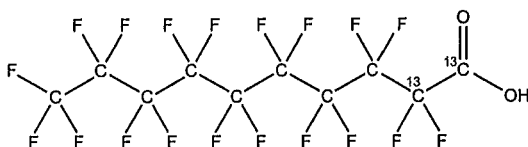
MPFDA

**LOT NUMBER:**

MPFDA1221

**COMPOUND:**Perfluoro-n-(1,2-<sup>13</sup>C<sub>2</sub>)decanoic acid**STRUCTURE:****CAS #:**

960315-50-8

**MOLECULAR FORMULA:**<sup>13</sup>C<sub>2</sub><sup>12</sup>C<sub>8</sub>H<sub>19</sub>O<sub>2</sub>**MOLECULAR WEIGHT:**

516.07

**CONCENTRATION:**

50.0 ± 2.5 µg/mL

**SOLVENT(S):**

Methanol

Water (&lt;1%)

**CHEMICAL PURITY:**

&gt;98%

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

12/08/2021

(1,2-<sup>13</sup>C<sub>2</sub>)**EXPIRY DATE:** (mm/dd/yyyy)

12/08/2026

**RECOMMENDED STORAGE:**

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Date:

12/13/2021

(mm/dd/yyyy)

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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.

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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**EXPIRY DATE / PERIOD OF VALIDITY:**

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**LIMITED WARRANTY:**

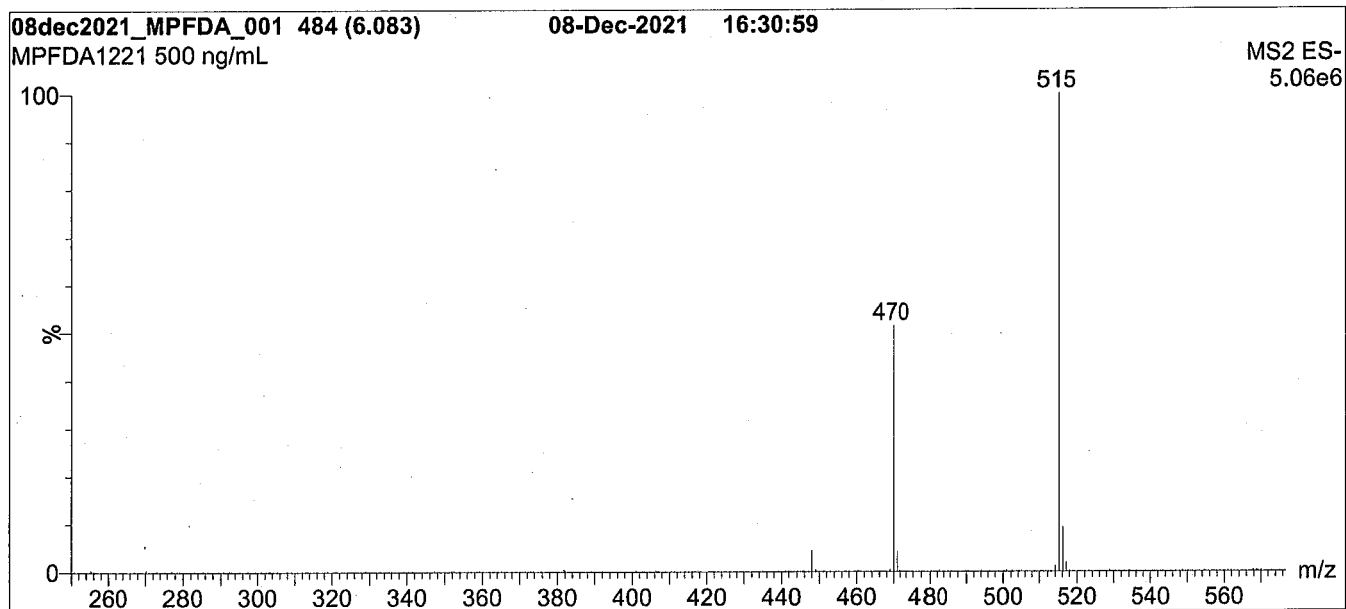
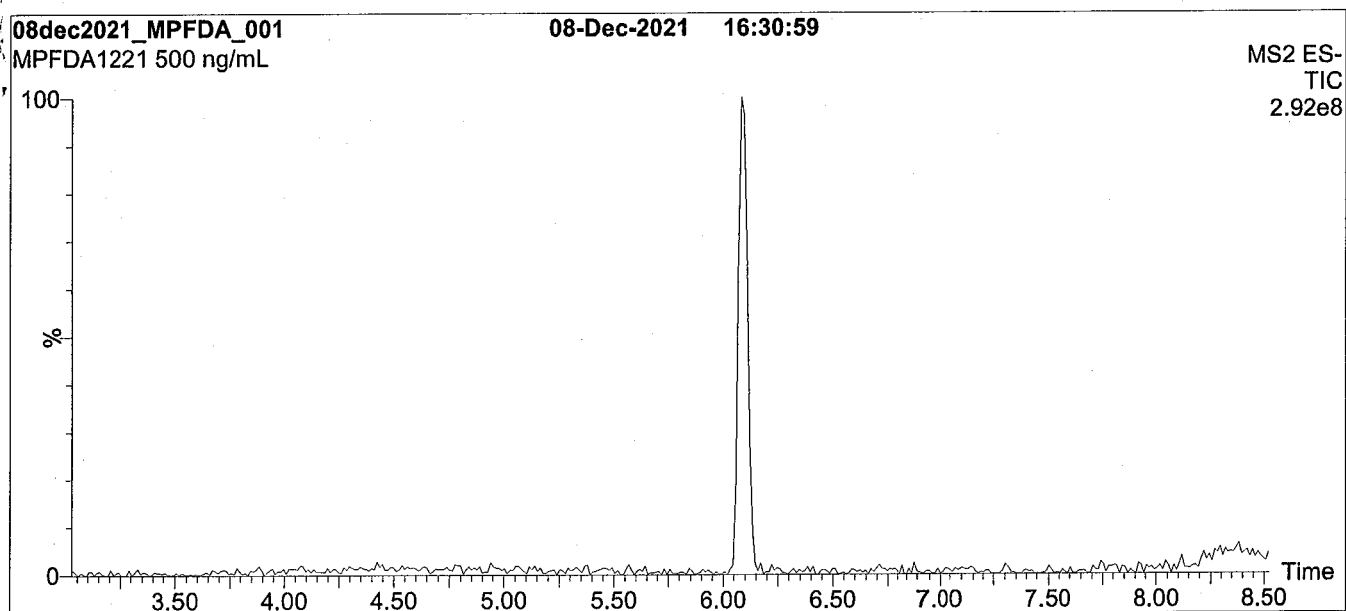
At the time of shipment, all products are warranted to be free of defects in material and workmanship and to conform to the stated technical and purity specifications.

**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).



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**Figure 1: MPFDA; 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 RP<sub>18</sub>  
1.7  $\mu$ m, 2.1 x 100 mm

Mobile phase: Gradient

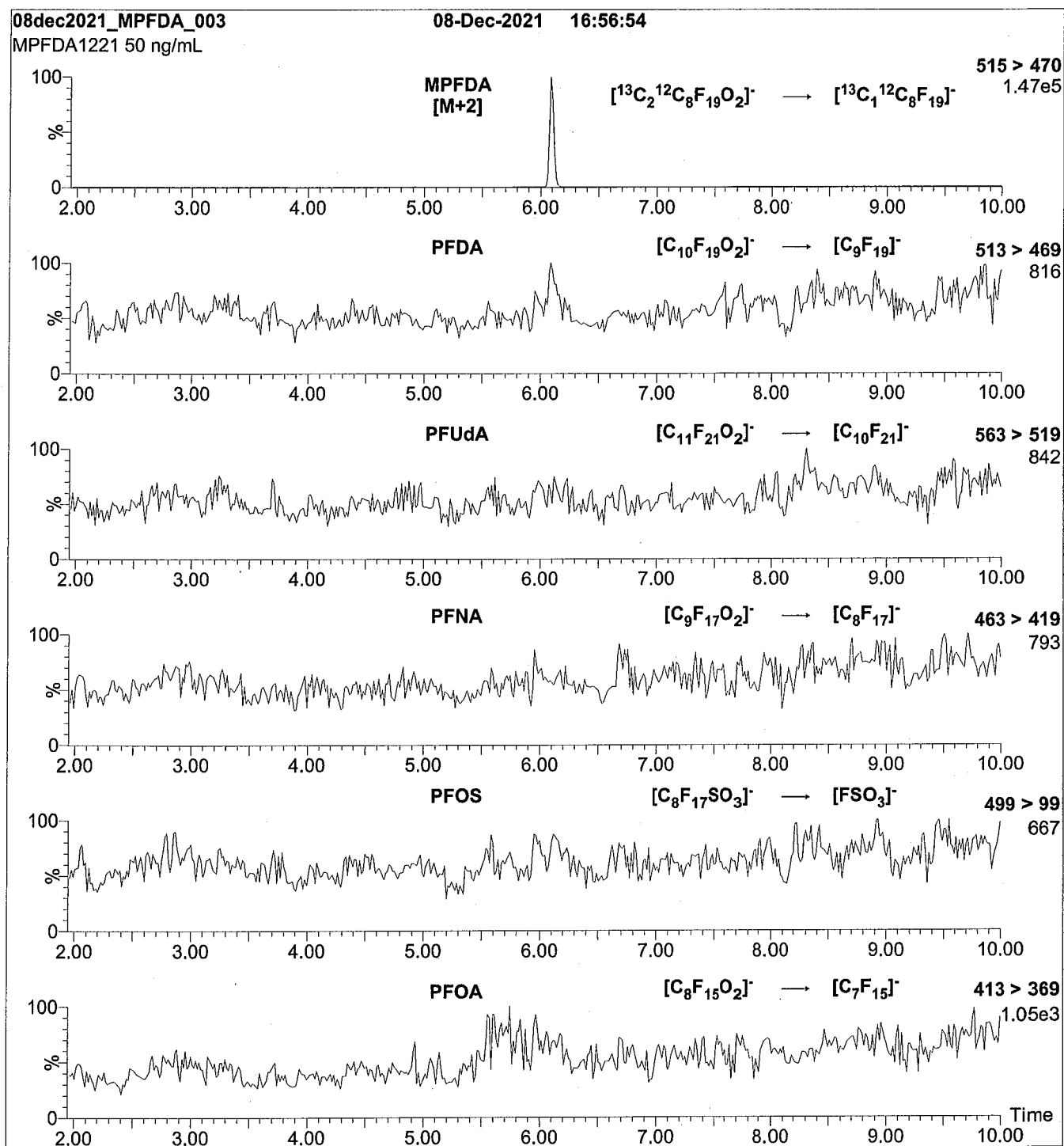
Start: 50% H<sub>2</sub>O / 50% (80:20 MeOH:ACN)  
(both with 10 mM NH<sub>4</sub>OAc buffer)  
Ramp to 90% organic over 9 min and hold for  
1 min before returning to initial conditions in 0.75 min.  
Time: 12 min

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

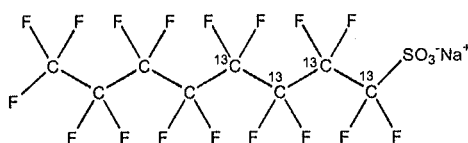
Analyte	Parent	CAS Number	Concentration	Units
13C2-PFDA		13C2-PFDA	50	ug/mL



# WELLINGTON LABORATORIES

## CERTIFICATE OF ANALYSIS DOCUMENTATION

**PRODUCT CODE:** MPFOS **LOT NUMBER:** MPFOS0821  
**COMPOUND:** Sodium perfluoro-1-(1,2,3,4-<sup>13</sup>C<sub>4</sub>)octanesulfonate  
**STRUCTURE:** **CAS #:** 960315-53-1



**MOLECULAR FORMULA:** <sup>13</sup>C<sub>4</sub><sup>12</sup>C<sub>4</sub>F<sub>17</sub>SO<sub>3</sub>Na **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% <sup>13</sup>C  
**LAST TESTED:** (mm/dd/yyyy) 08/18/2021 (1,2,3,4-<sup>13</sup>C<sub>4</sub>)  
**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)  
 Figure 2: LC/MS/MS Data (Selected MRM Transitions)

### ADDITIONAL INFORMATION:

- See page 2 for further details.
- Contains ~0.4% sodium perfluoro-1-(<sup>13</sup>C<sub>3</sub>)heptanesulfonate.

**FOR LABORATORY USE ONLY: NOT FOR HUMAN OR DRUG USE**

**Certified By:**   
 B.G. Chittim, General Manager

**Date:** 08/19/2021  
 (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.

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**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.

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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:**

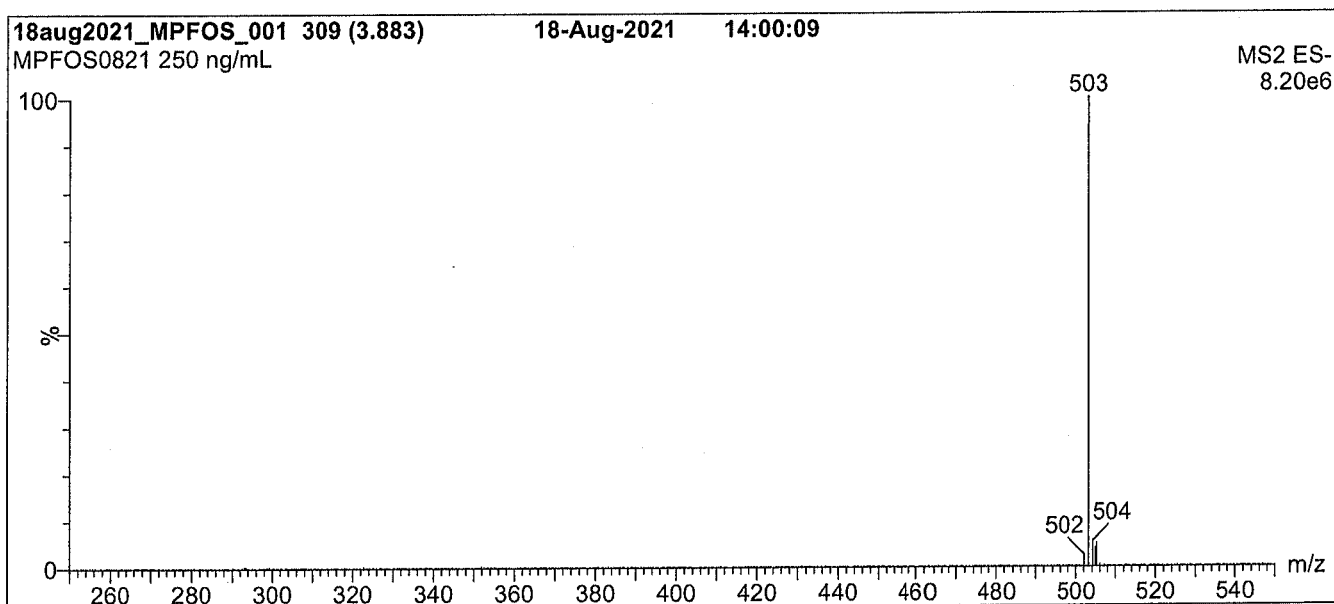
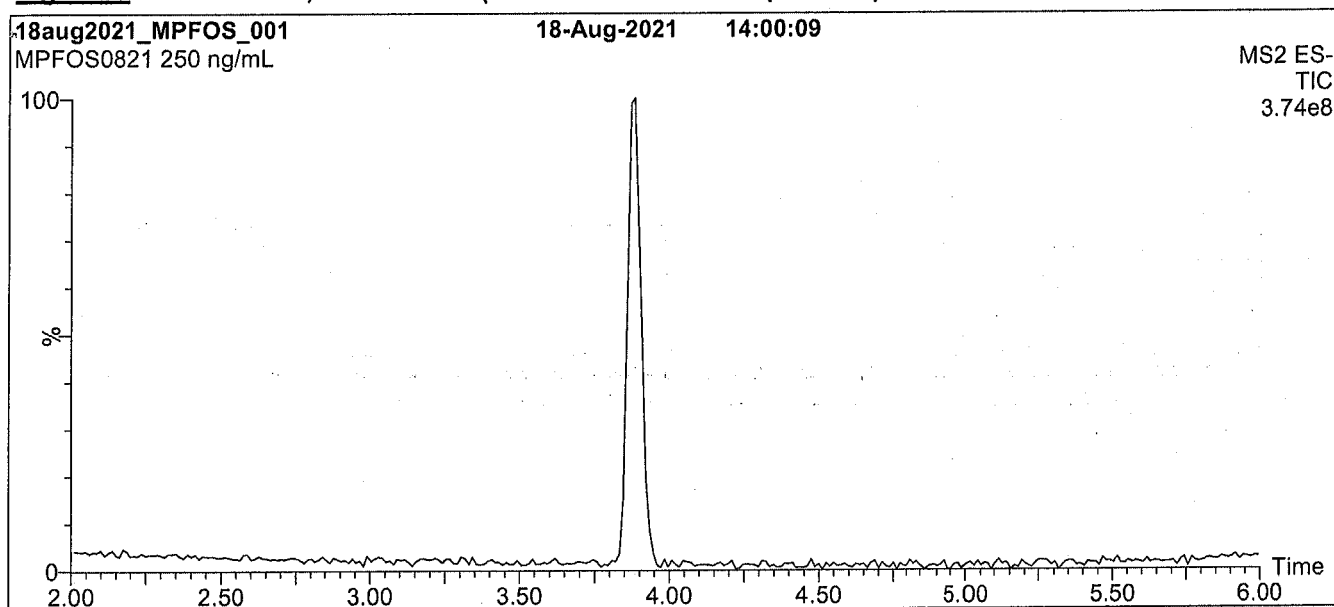
At the time of shipment, all products are warranted to be free of defects in material and workmanship and to conform to the stated technical and purity specifications.

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**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 RP<sub>18</sub>  
1.7  $\mu$ m, 2.1 x 100 mm

Mobile phase: Gradient

Start: 40% H<sub>2</sub>O / 60% (80:20 MeOH:ACN)  
(both with 10 mM NH<sub>4</sub>OAc 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  $\mu$ L/min

**MS Parameters:**

Experiment: Full Scan (250 - 850 amu)

Source: Electrospray (negative)

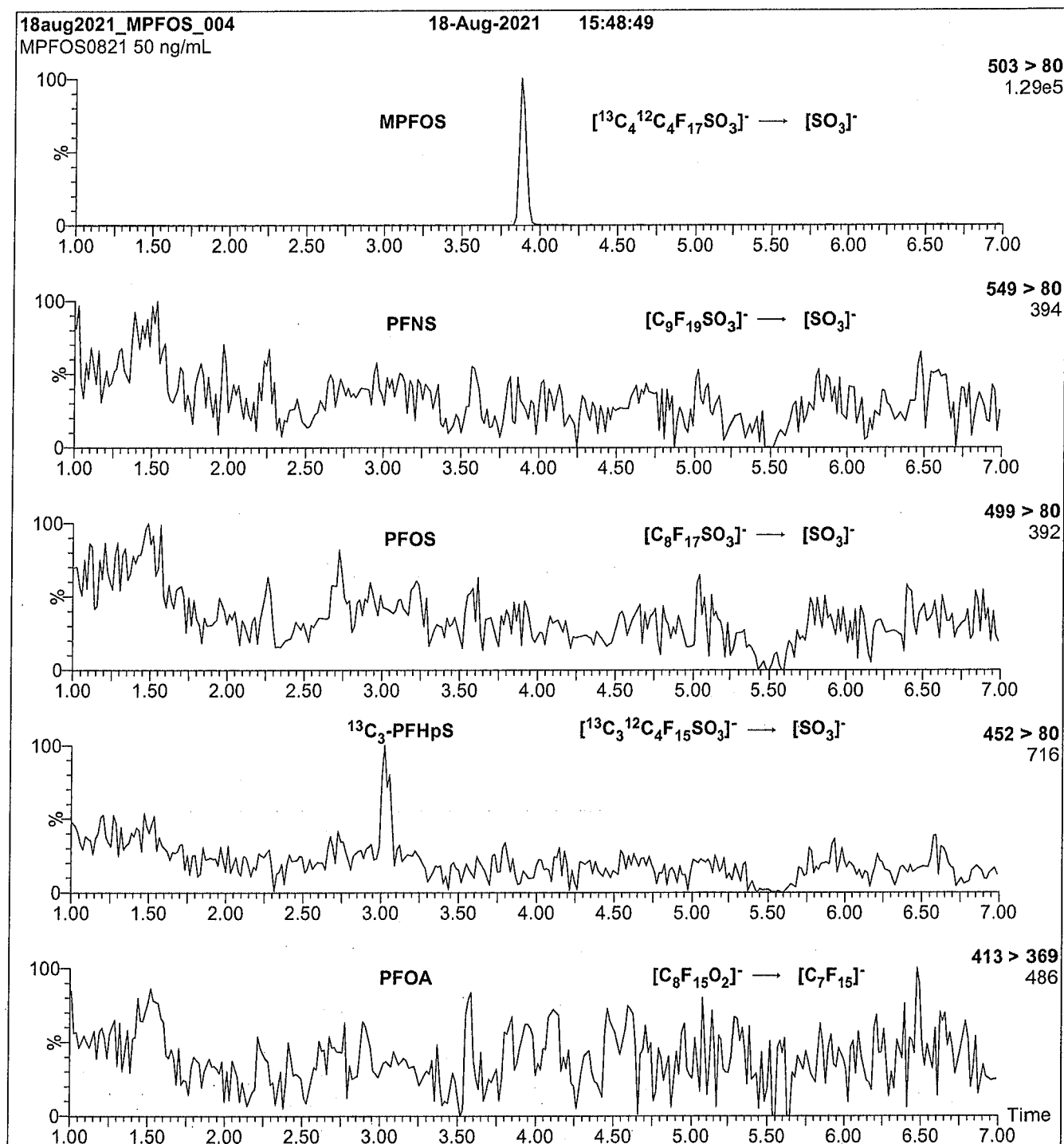
Capillary Voltage (kV) = 2.00

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



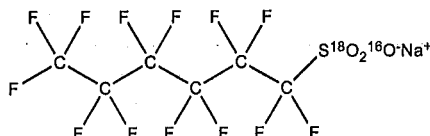
# WELLINGTON LABORATORIES

## CERTIFICATE OF ANALYSIS DOCUMENTATION

**PRODUCT CODE:** MPFHxS  
**COMPOUND:** Sodium perfluoro-1-hexane(<sup>18</sup>O<sub>2</sub>)sulfonate

**LOT NUMBER:** MPFHxS1021

**STRUCTURE:**



**CAS #:** 1585941-14-5

**MOLECULAR FORMULA:** C<sub>6</sub>F<sub>13</sub>S<sup>18</sup>O<sub>2</sub><sup>16</sup>ONa  
**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% (<sup>18</sup>O<sub>2</sub>)

**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 (C<sub>6</sub>F<sub>13</sub>S<sup>18</sup>O<sub>2</sub><sup>16</sup>O) has been observed to be up to 10% lower than for PFHxS (C<sub>6</sub>F<sub>13</sub>S<sup>18</sup>O<sub>3</sub>) when both compounds are injected together. This difference may vary between instruments.
- Contains ~0.6% of sodium perfluoro-1-octane(<sup>18</sup>O<sub>2</sub>)sulfonate (<sup>18</sup>O<sub>2</sub>-PFOS) and ~0.3% of sodium perfluoro-1-heptane(<sup>18</sup>O<sub>2</sub>)sulfonate (<sup>18</sup>O<sub>2</sub>-PFHpS).
- Due to the isotopic purity of the starting material (<sup>18</sup>O<sub>2</sub> >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

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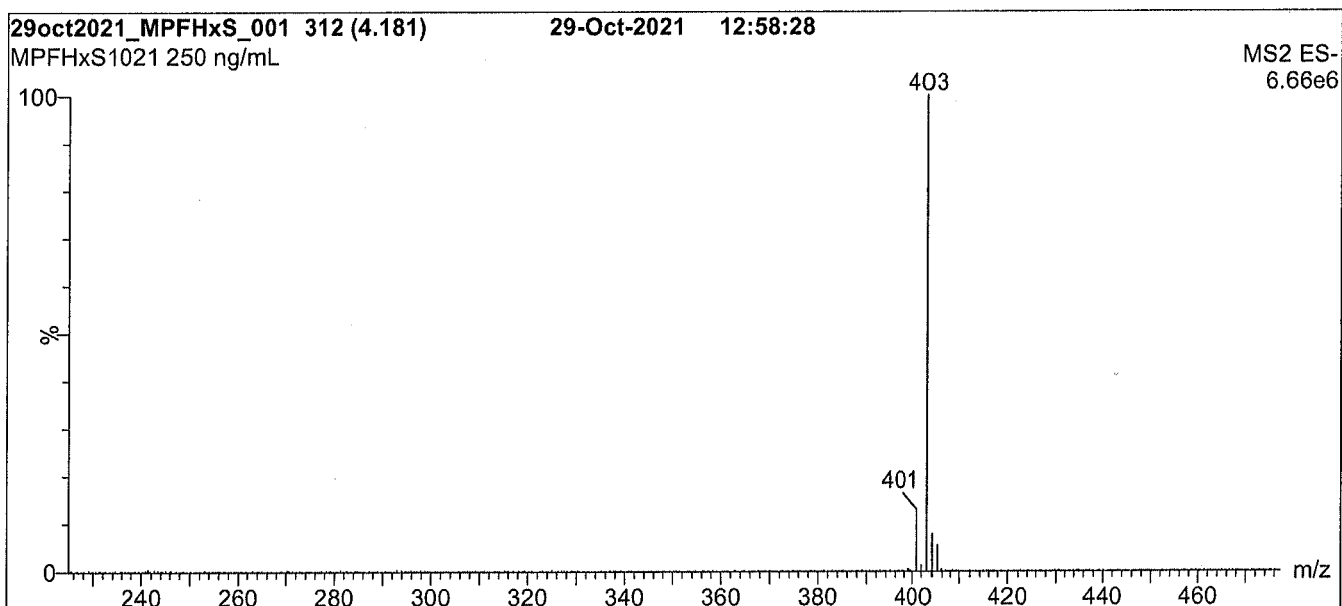
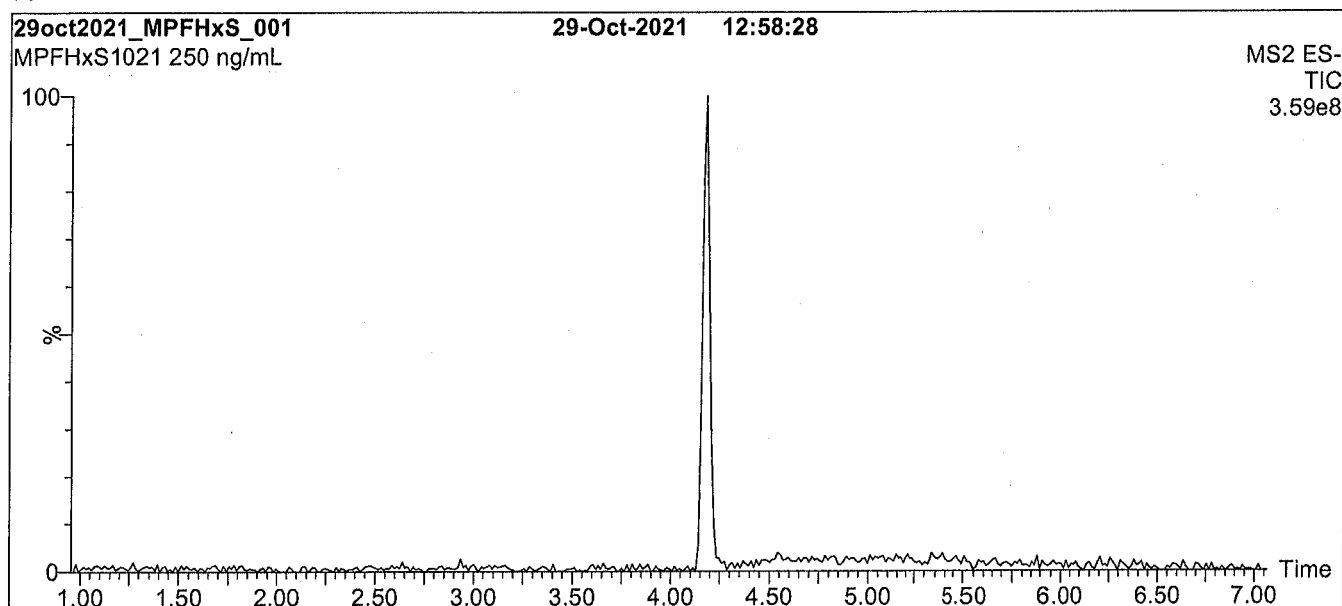
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**Figure 1: MPFHxS; LC/MS Data (Full Scan and Mass Spectrum)****Conditions for Figure 1:**

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

**Chromatographic Conditions:**

Column: Acquity UPLC BEH Shield RP<sub>18</sub>  
1.7  $\mu$ m, 2.1 x 100 mm

Mobile phase: Gradient

Start: 50% H<sub>2</sub>O / 50% (80:20 MeOH:ACN)  
(both with 10 mM NH<sub>4</sub>OAc 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  $\mu$ L/min

**MS Parameters:**

Experiment: Full Scan (225 - 850 amu)

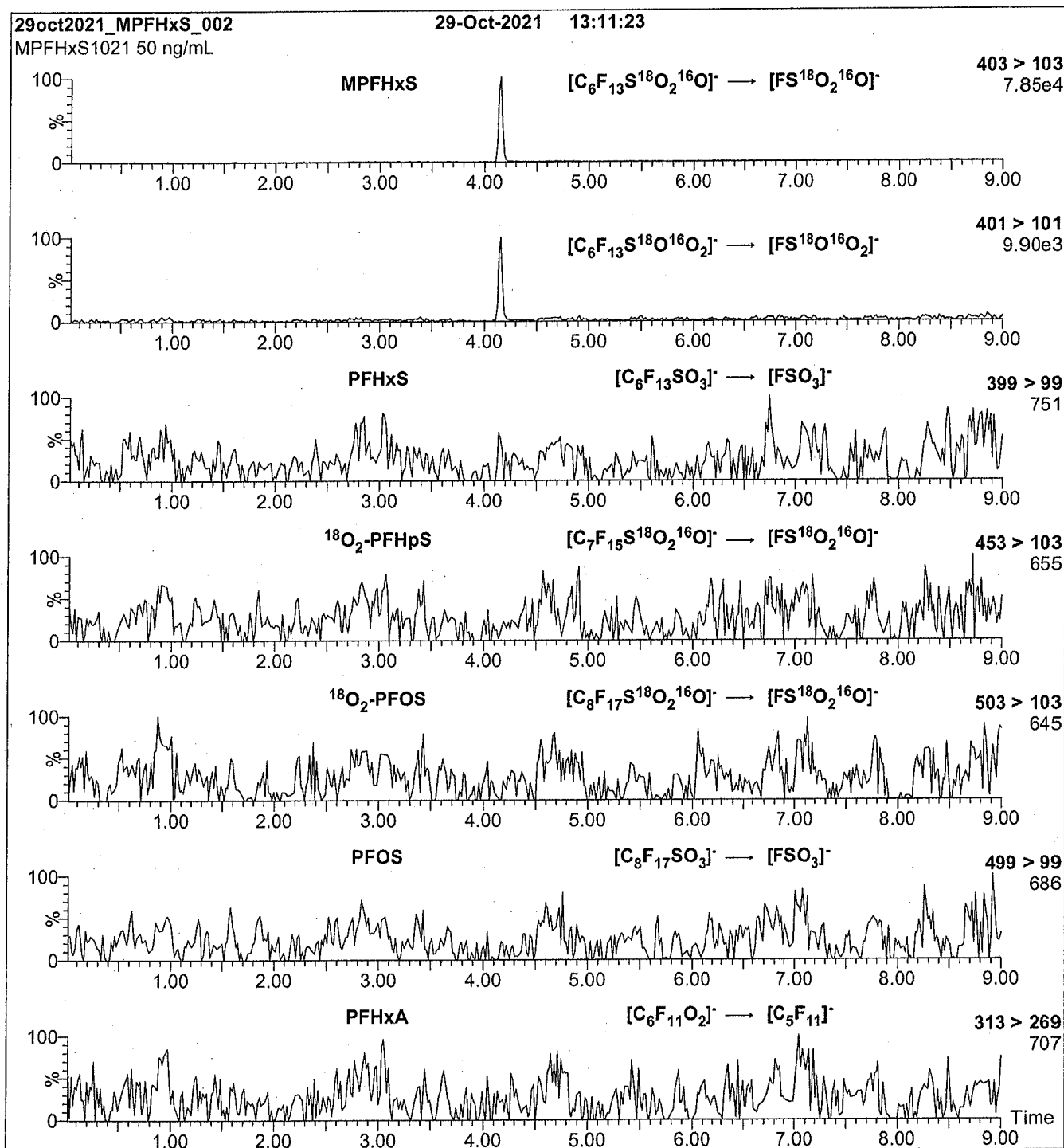
Source: Electrospray (negative)

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  $\mu$ 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		18O2-PFHXS	50	ug/mL

# Analytical Standard Record

**22A0122**

Description: PFAS - IIS MPFHxS 50ug/mL

Expires: 10/29/2026

Standard Type: Analyte Spike

Prepared: 10/29/2021

Solvent: MeOH

Prepared By: Dipti Gokal

Final Volume (mLs): 1.2

Department: PFAS

Vials: 1

Last Edit: 01/20/2022 15:49 by HGH

Analyte	Parent	CAS Number	Concentration	Units
18O2-PFHXS		18O2-PFHXS	50	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

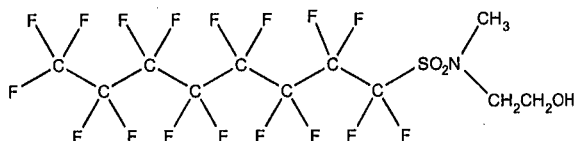
**LOT NUMBER:** NMeFOSE0921M**COMPOUND:**

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:**

&gt;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  
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:**

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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:**

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:**

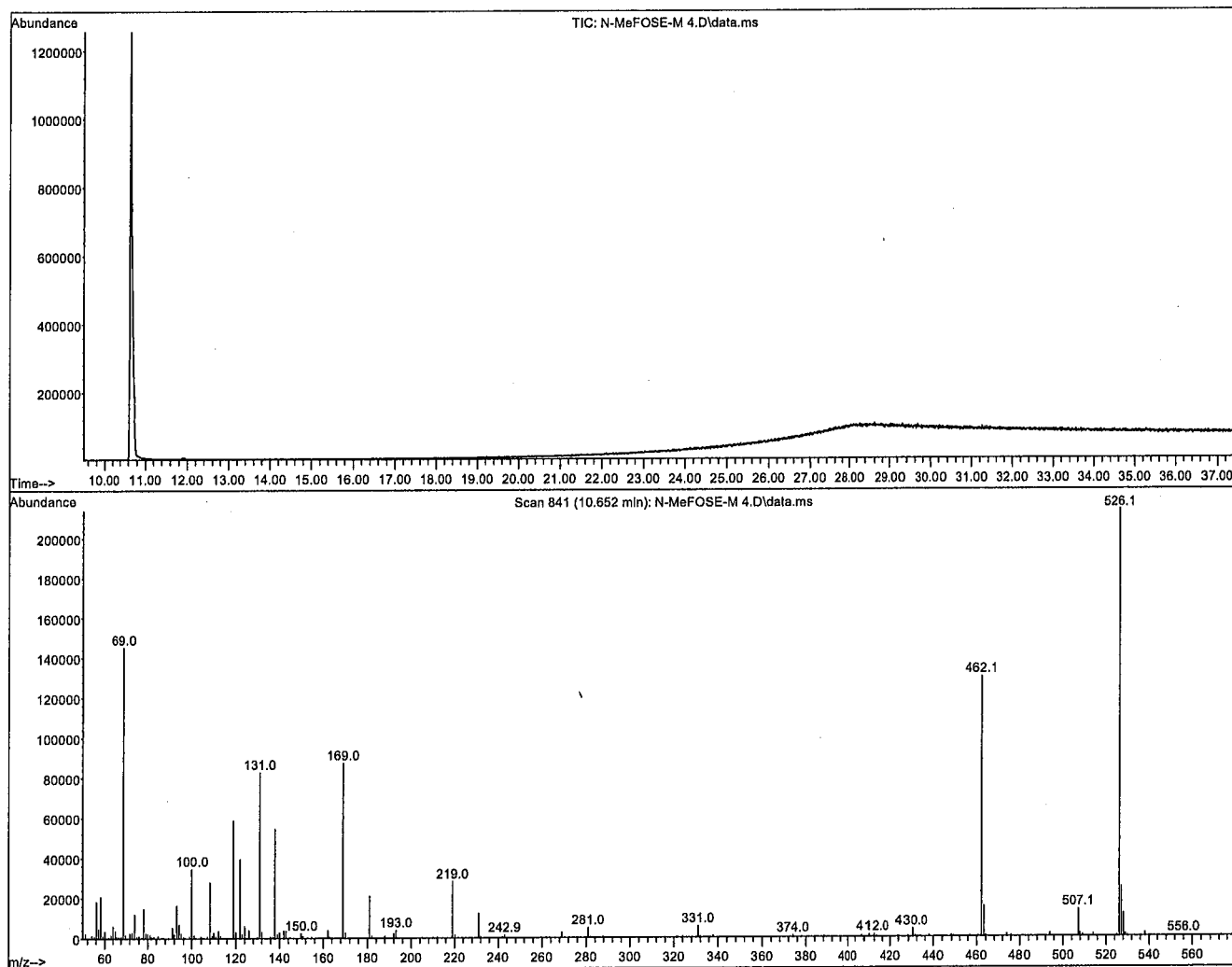
At the time of shipment, all products are warranted to be free of defects in material and workmanship and to conform to the stated technical and purity specifications.

**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](http://www.well-labs.com) or contact us directly at [info@well-labs.com](mailto: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  $\mu$ m film thickness) Agilent J&W

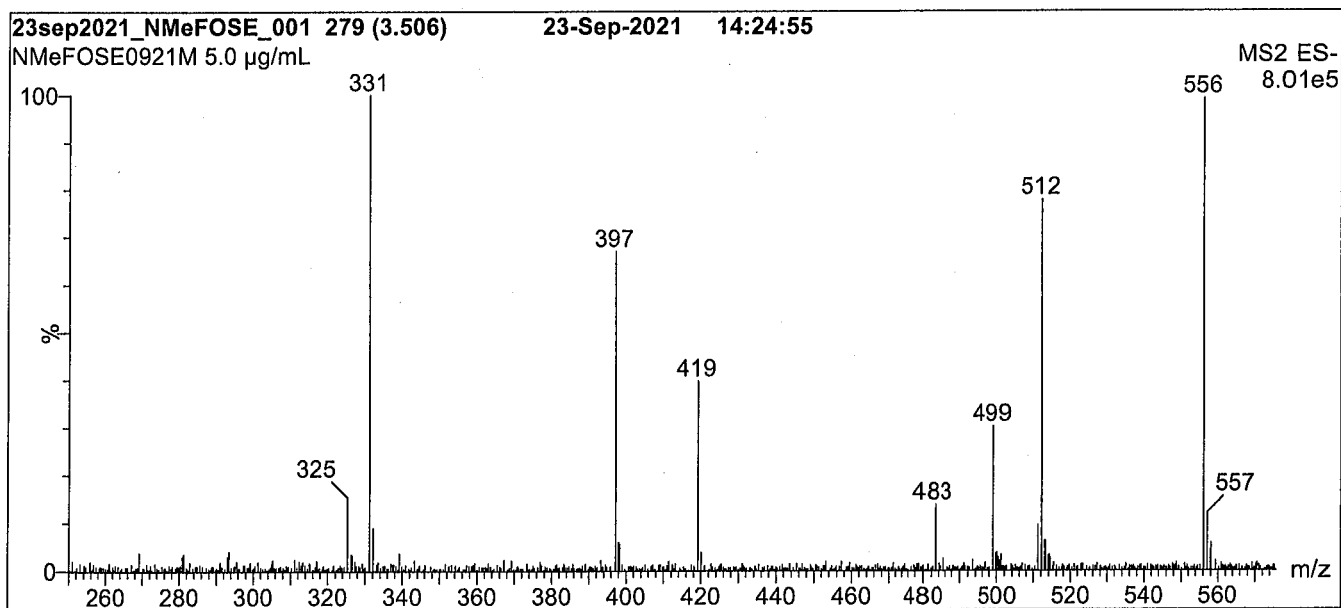
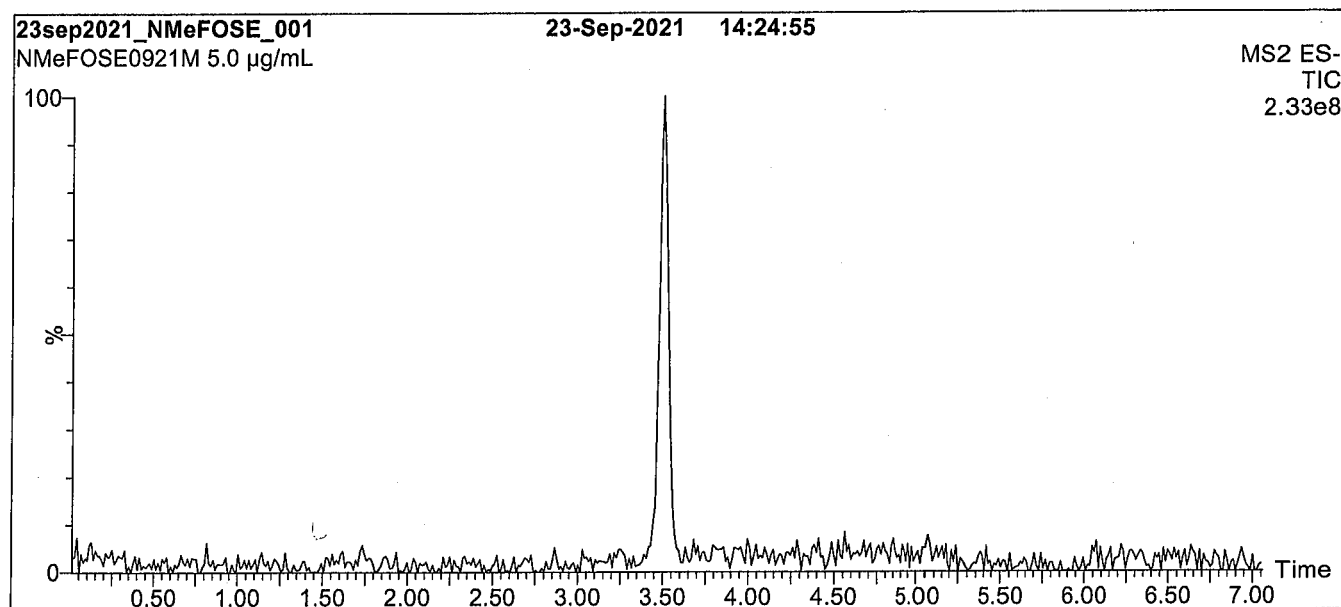
Flow: Constant at 1 mL/min

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 RP<sub>18</sub>  
1.7 µm, 2.1 x 100 mm

Mobile phase: Gradient

Start: 30% H<sub>2</sub>O / 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)

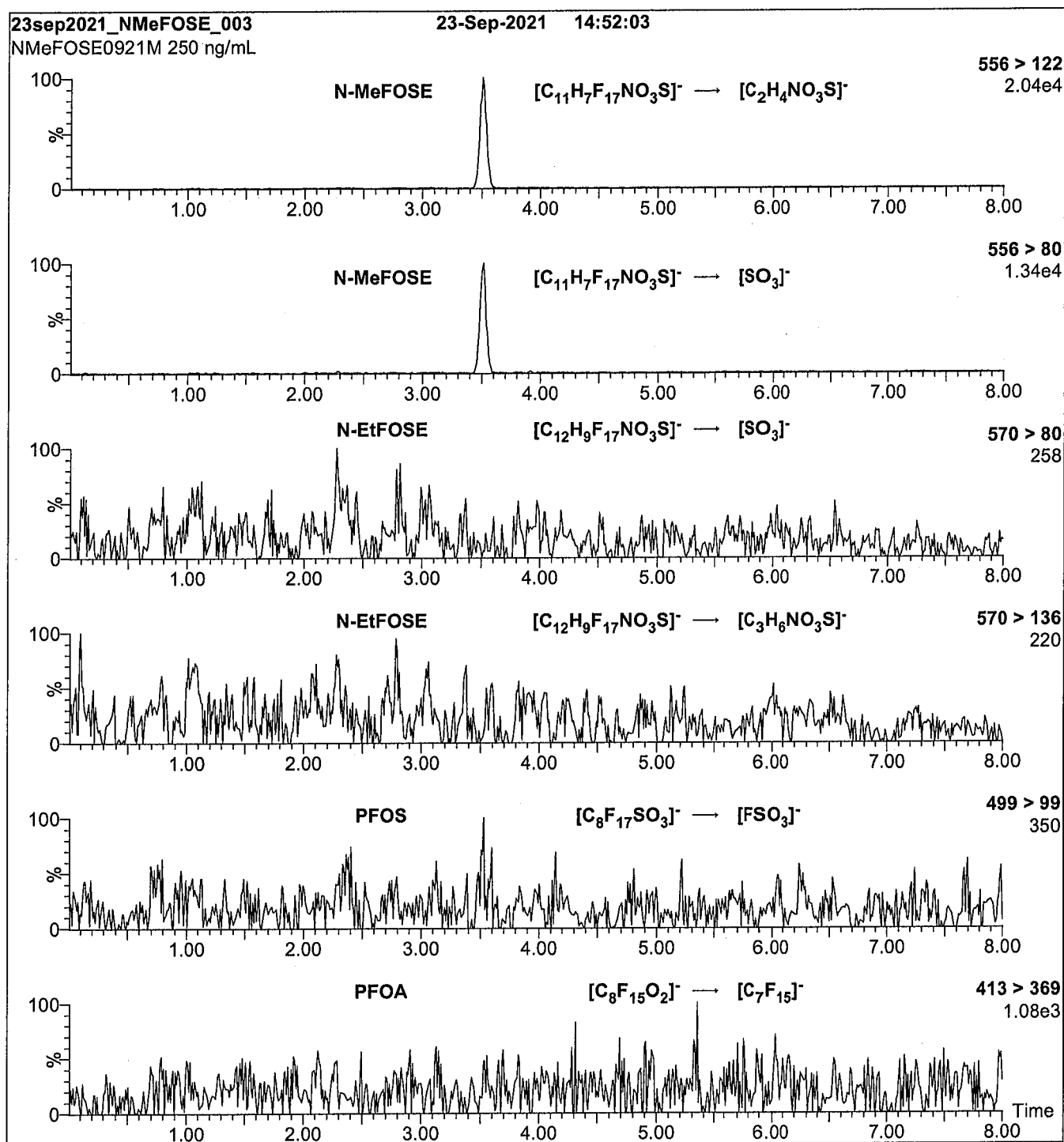
Source: Electrospray (negative)

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  $\mu$ 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



# WELLINGTON LABORATORIES

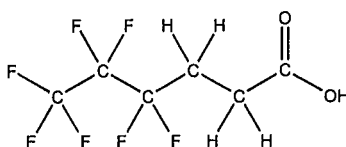
## CERTIFICATE OF ANALYSIS DOCUMENTATION

**PRODUCT CODE:** FPrPA  
**COMPOUND:** 3-Perfluoropropyl propanoic acid

**LOT NUMBER:** FPrPA0122  
22C0308

**STRUCTURE:**

**CAS #:** 356-02-5



**MOLECULAR FORMULA:**  $C_8H_6F_7O_2$   
**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  $^{19}\text{F}$  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:**

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**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.

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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.

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:**

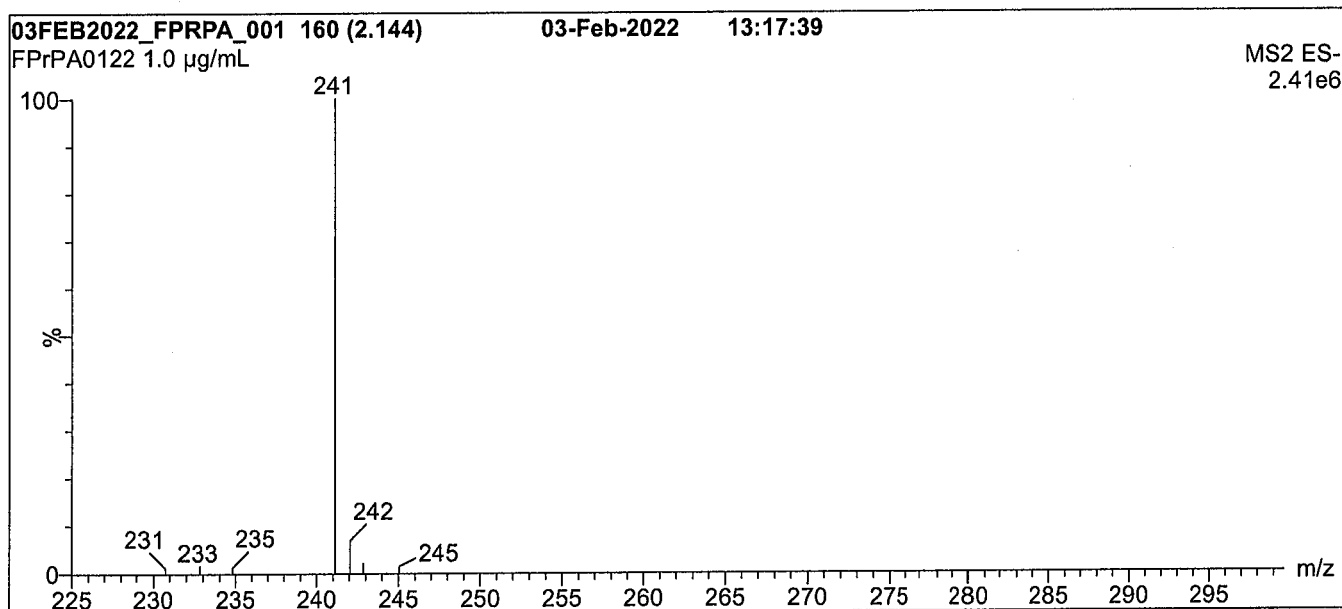
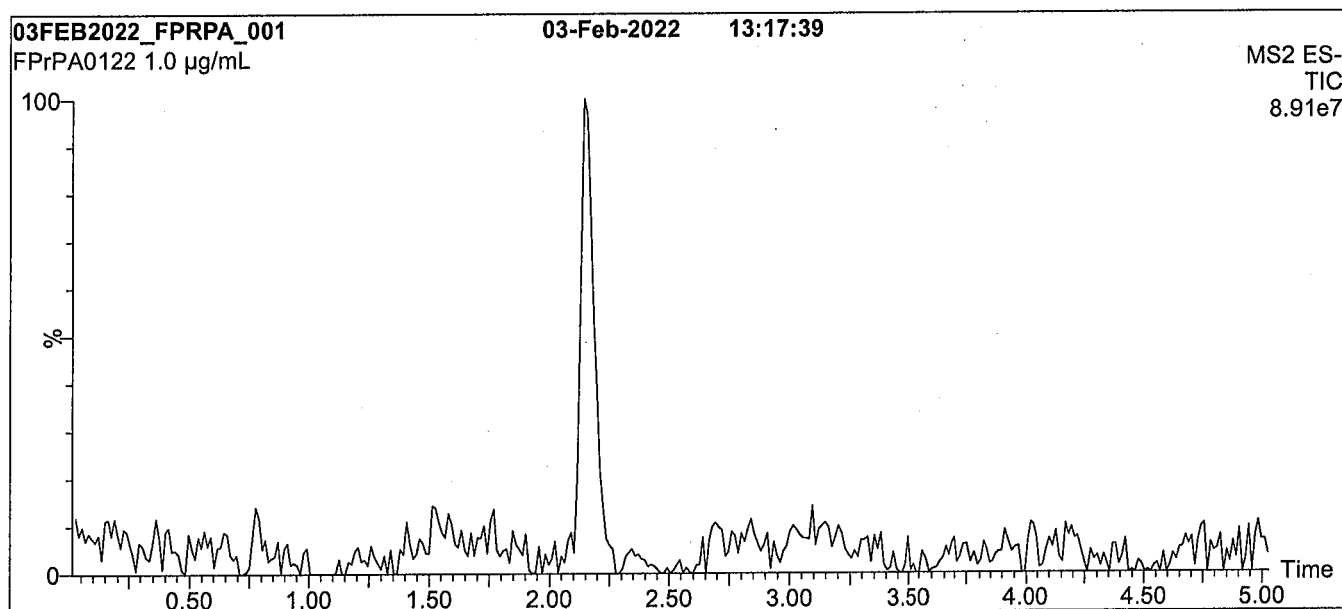
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**QUALITY MANAGEMENT:**

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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 RP<sub>18</sub>  
1.7 µm, 2.1 x 100 mm

Mobile phase: Gradient

Start: 60% H<sub>2</sub>O / 40% (80:20 MeOH:ACN)  
(both with 10 mM NH<sub>4</sub>OAc 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)

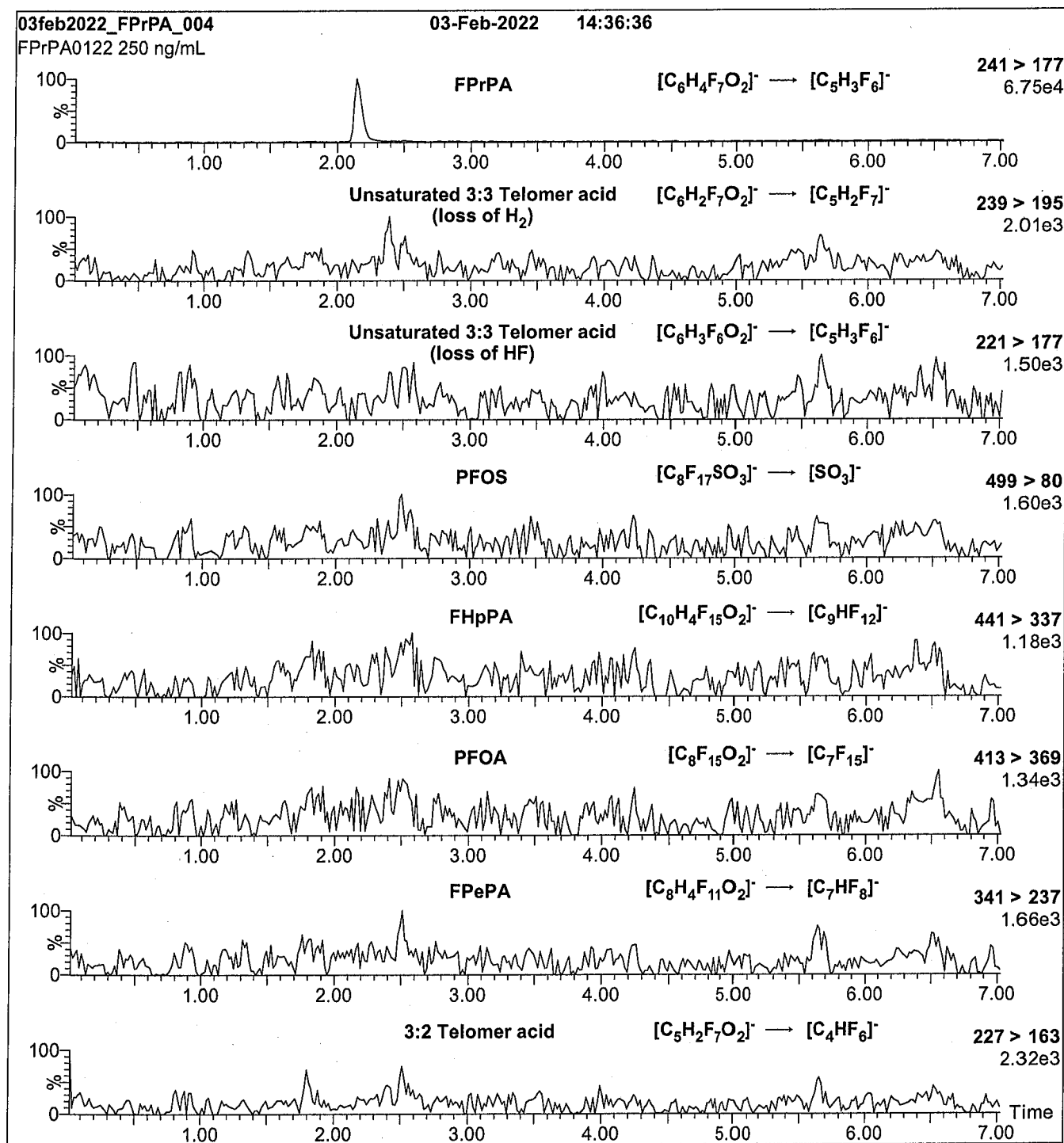
Source: Electrospray (negative)

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  $\mu$ 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



# WELLINGTON LABORATORIES

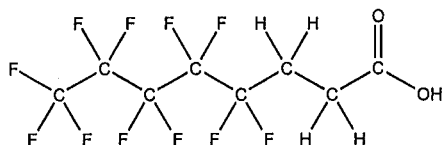
## CERTIFICATE OF ANALYSIS DOCUMENTATION

**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

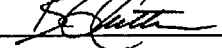
**DOCUMENTATION/ DATA ATTACHED:**

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  $^1\text{H}$  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



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**LIMITED WARRANTY:**

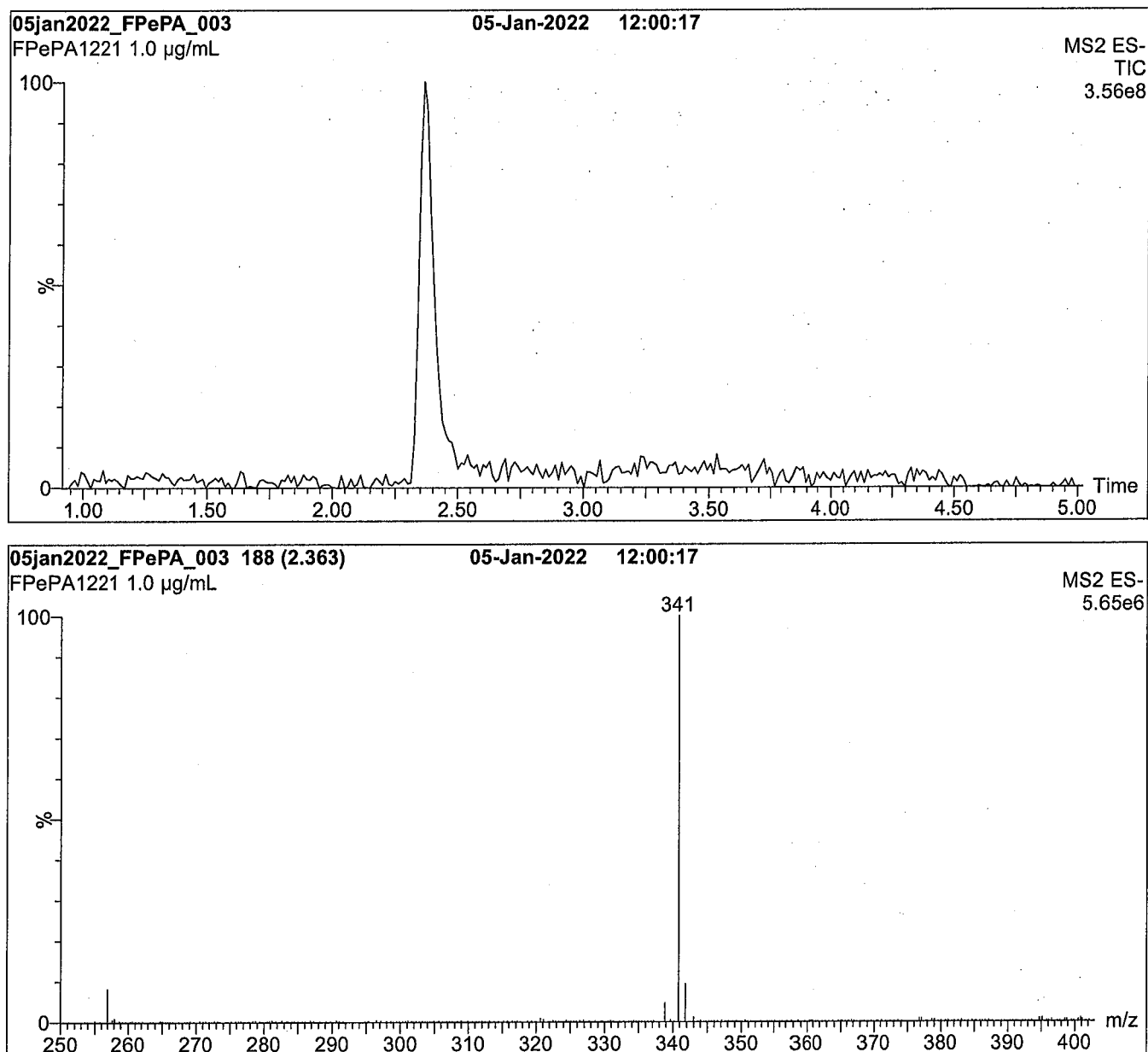
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**Figure 1: FPePA; LC/MS Data (Full Scan and Mass Spectrum)****Conditions for Figure 1:**

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

**Chromatographic Conditions:**

Column: Acquity UPLC BEH Shield RP<sub>18</sub>  
1.7 µm, 2.1 x 100 mm

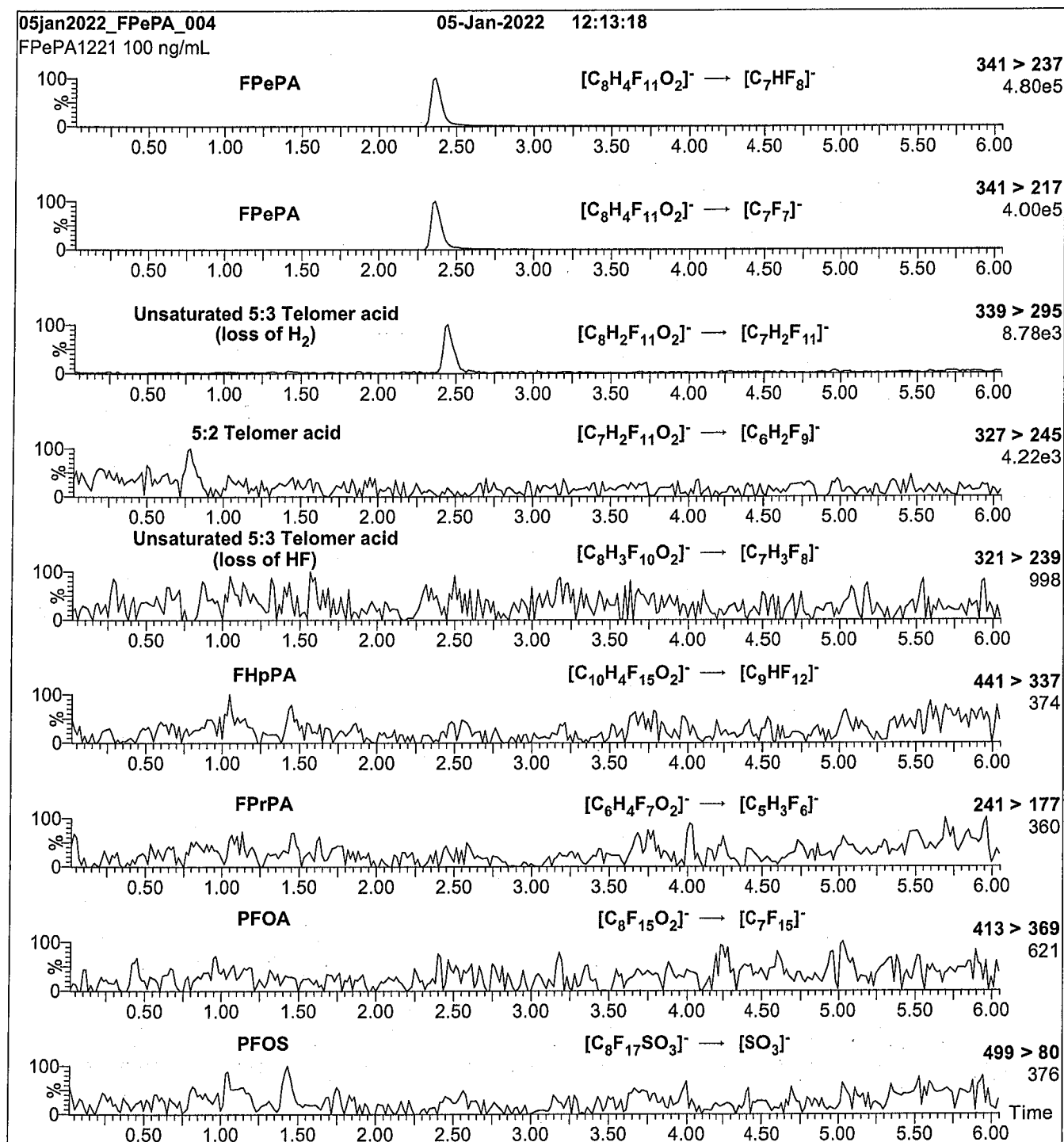
Mobile phase: Gradient  
Start: 45% H<sub>2</sub>O / 55% (80:20 MeOH:ACN)  
(both with 10 mM NH<sub>4</sub>OAc buffer)  
Ramp to 90% organic over 7 min and hold for  
3 min before returning to initial conditions in 0.75 min.  
Time: 12 min

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  $\mu$ 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

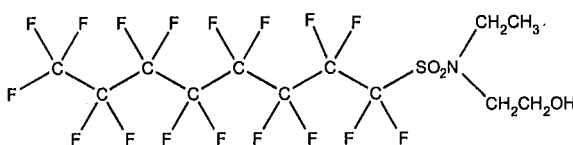
Analyte	Parent	CAS Number	Concentration	Units
5:3FTCA		914637-49-3	50	ug/mL



# WELLINGTON LABORATORIES

## 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

**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:**

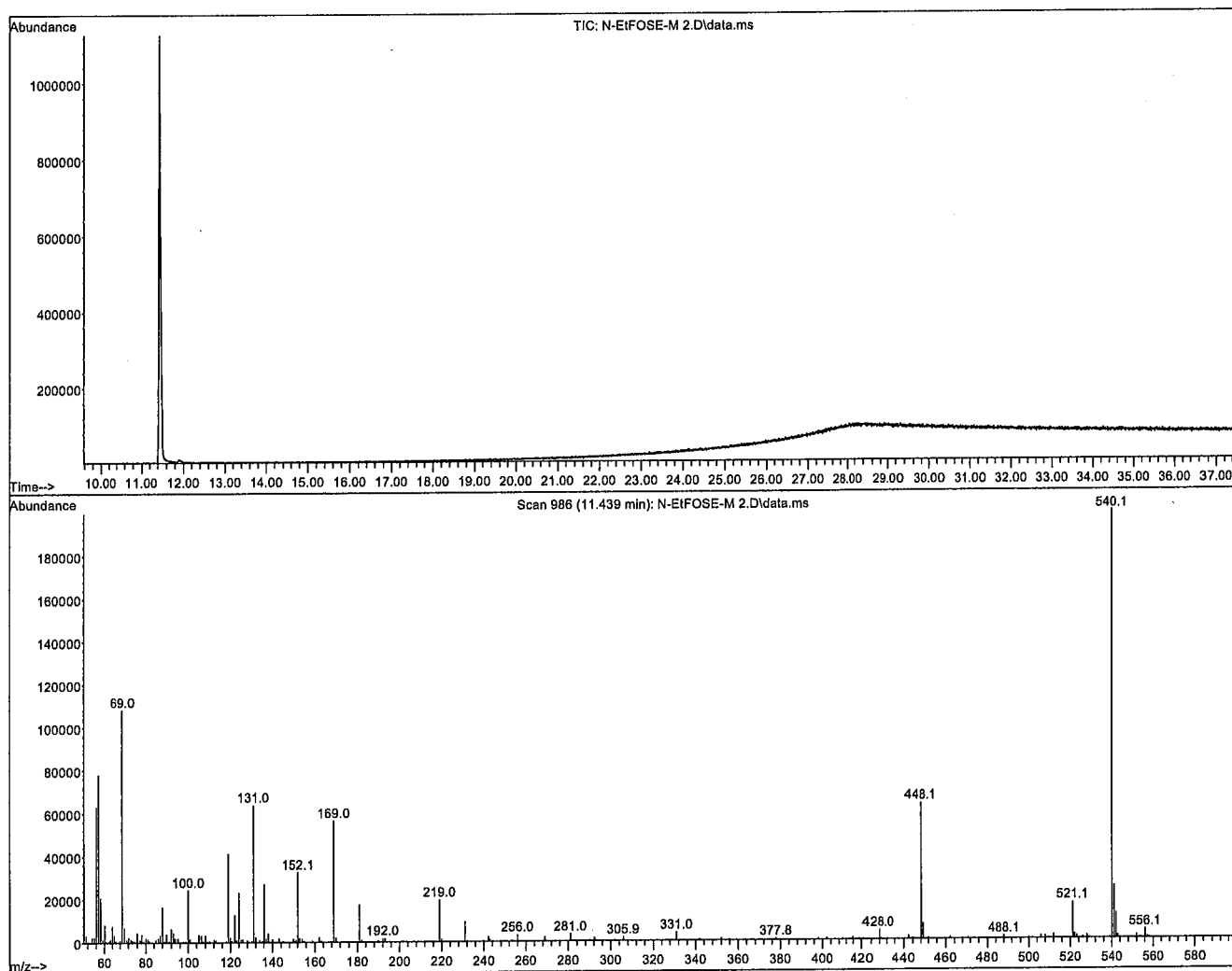
At the time of shipment, all products are warranted to be free of defects in material and workmanship and to conform to the stated technical and purity specifications.

**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 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](http://www.well-labs.com) or contact us directly at [info@well-labs.com](mailto: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  $\mu$ m film thickness) Agilent J&W

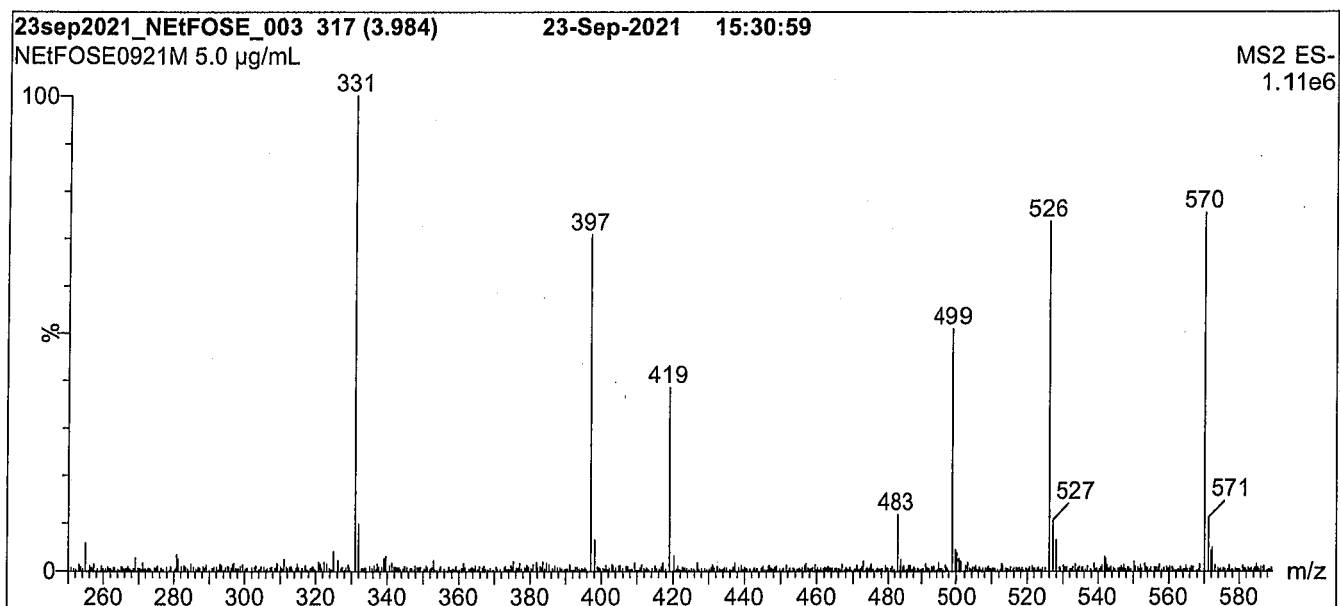
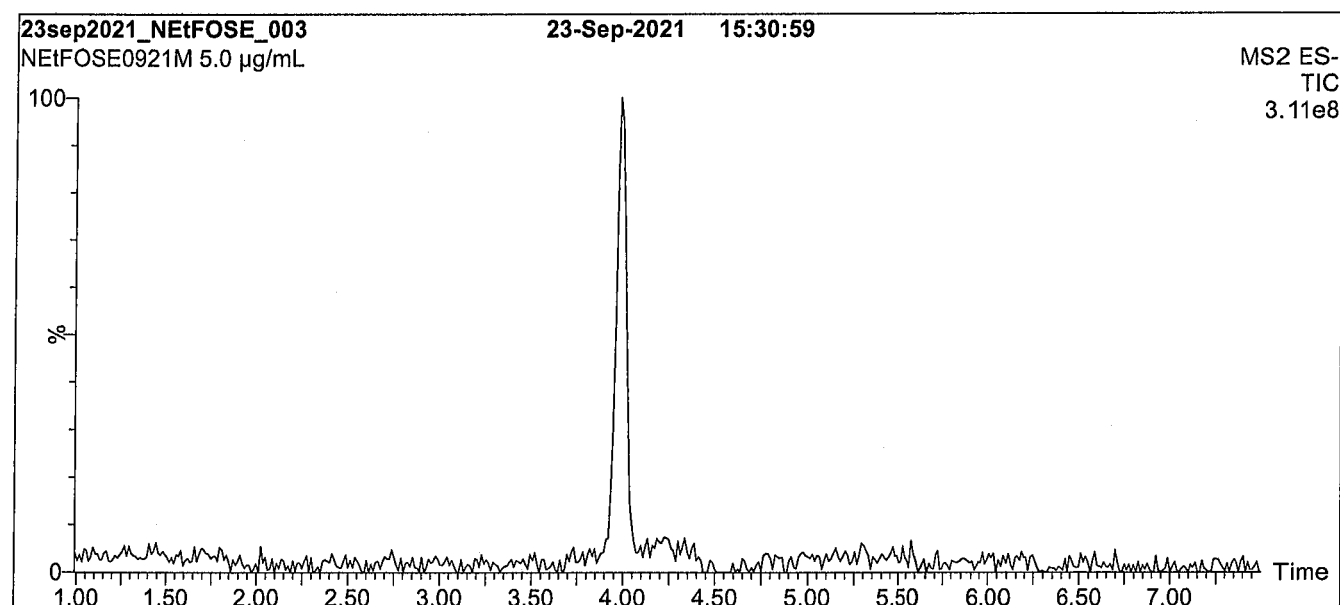
Flow: Constant at 1 mL/min

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 RP<sub>18</sub>  
1.7 µm, 2.1 x 100 mm

Mobile phase: Gradient

Start: 30% H<sub>2</sub>O / 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)

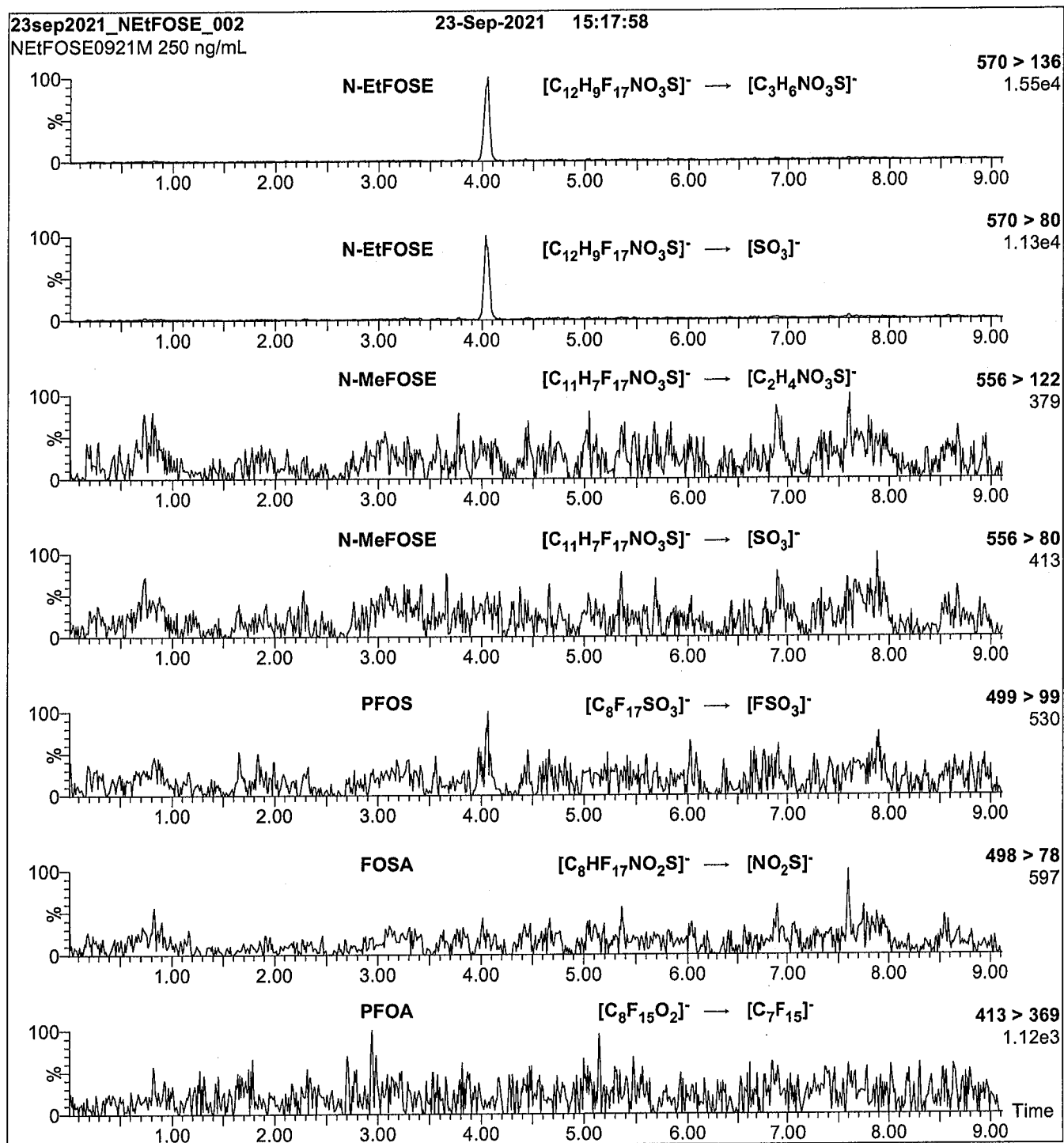
Capillary Voltage (kV) = 2.00

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  $\mu$ 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#: NETFOSE0921M)
Final Volume (mLs):	1	Department:	PFAS
Vials:	1	Last Edit:	03/15/2022 15:59 by DAG

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



# WELLINGTON LABORATORIES

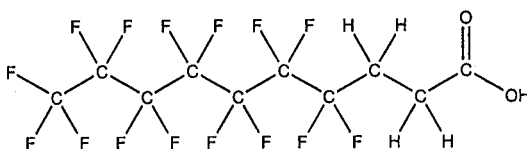
## CERTIFICATE OF ANALYSIS DOCUMENTATION

**PRODUCT CODE:** FHpPA  
**COMPOUND:** 3-Perfluoroheptyl propanoic acid

**LOT NUMBER:** FHpPA1020  
22C0311

**STRUCTURE:**

**CAS #:** 812-70-4



**MOLECULAR FORMULA:**  $C_{19}H_{15}F_{13}O_2$   
**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](mailto:info@well-labs.com)

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**SYNTHESIS / CHARACTERIZATION:**

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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.

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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:**

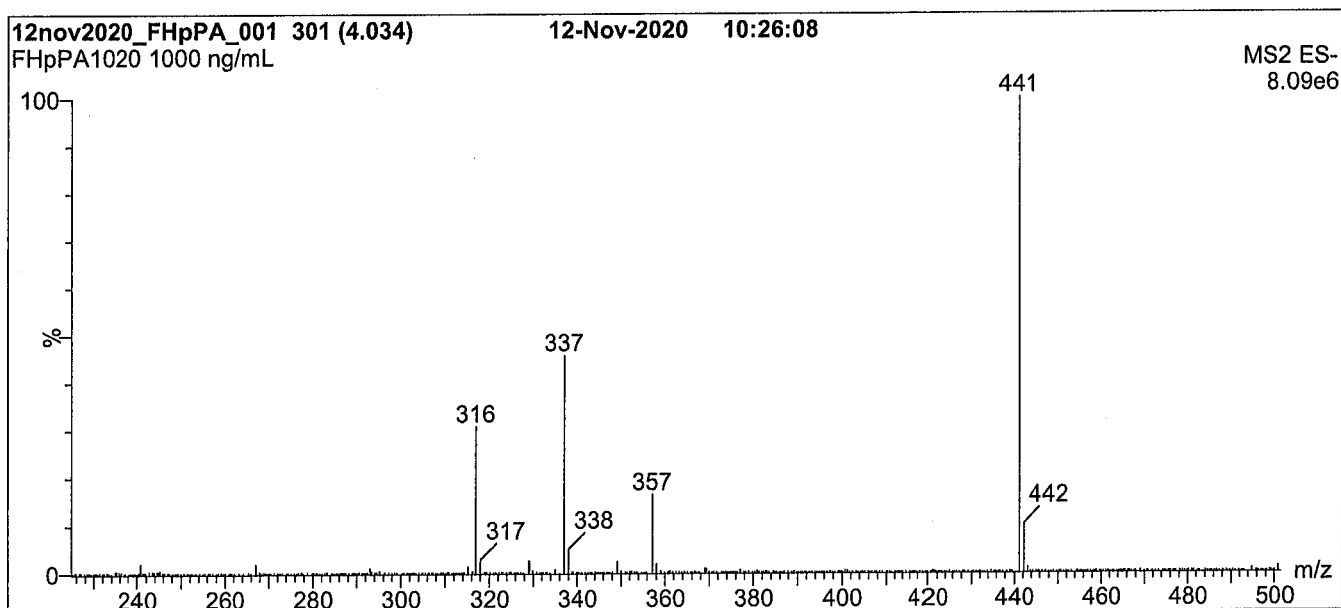
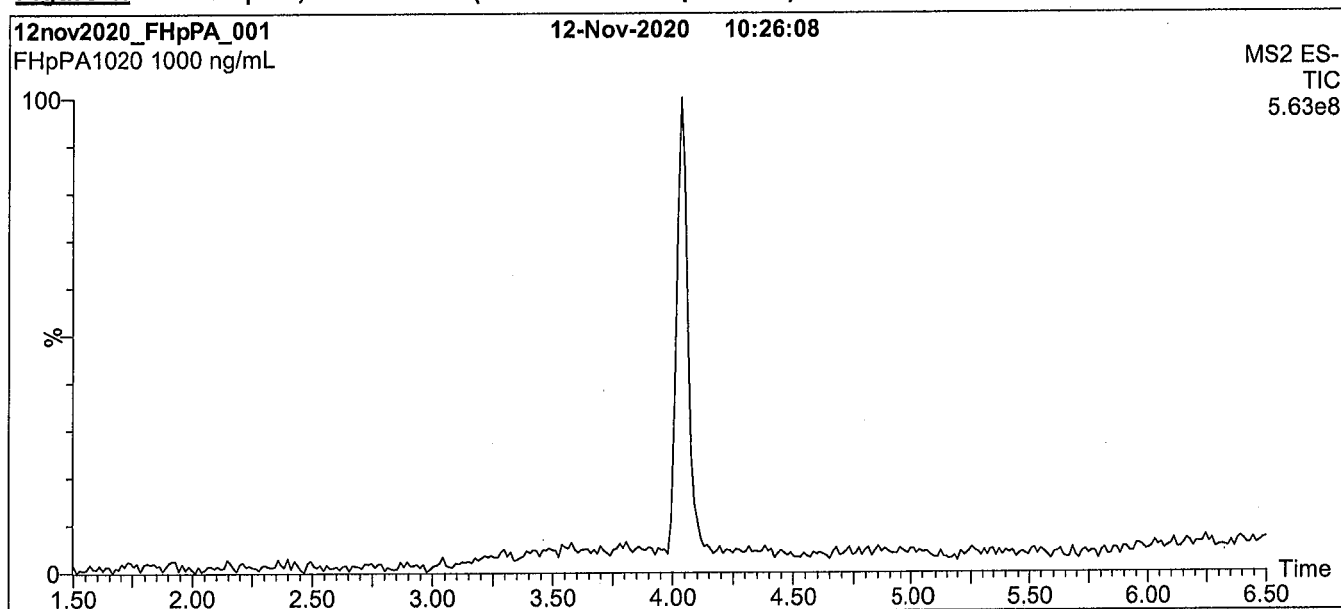
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**Figure 1: FHpPA; LC/MS Data (TIC and Mass Spectrum)****Conditions for Figure 1:**

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

**Chromatographic Conditions:**

Column: Acquity UPLC BEH Shield RP<sub>18</sub>  
1.7  $\mu$ m, 2.1 x 100 mm

Mobile phase: Gradient

Start: 45% H<sub>2</sub>O / 55% (80:20 MeOH:ACN)  
(both with 10 mM NH<sub>4</sub>OAc 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  $\mu$ L/min

**MS Parameters:**

Experiment: Full Scan (225 - 850 amu)

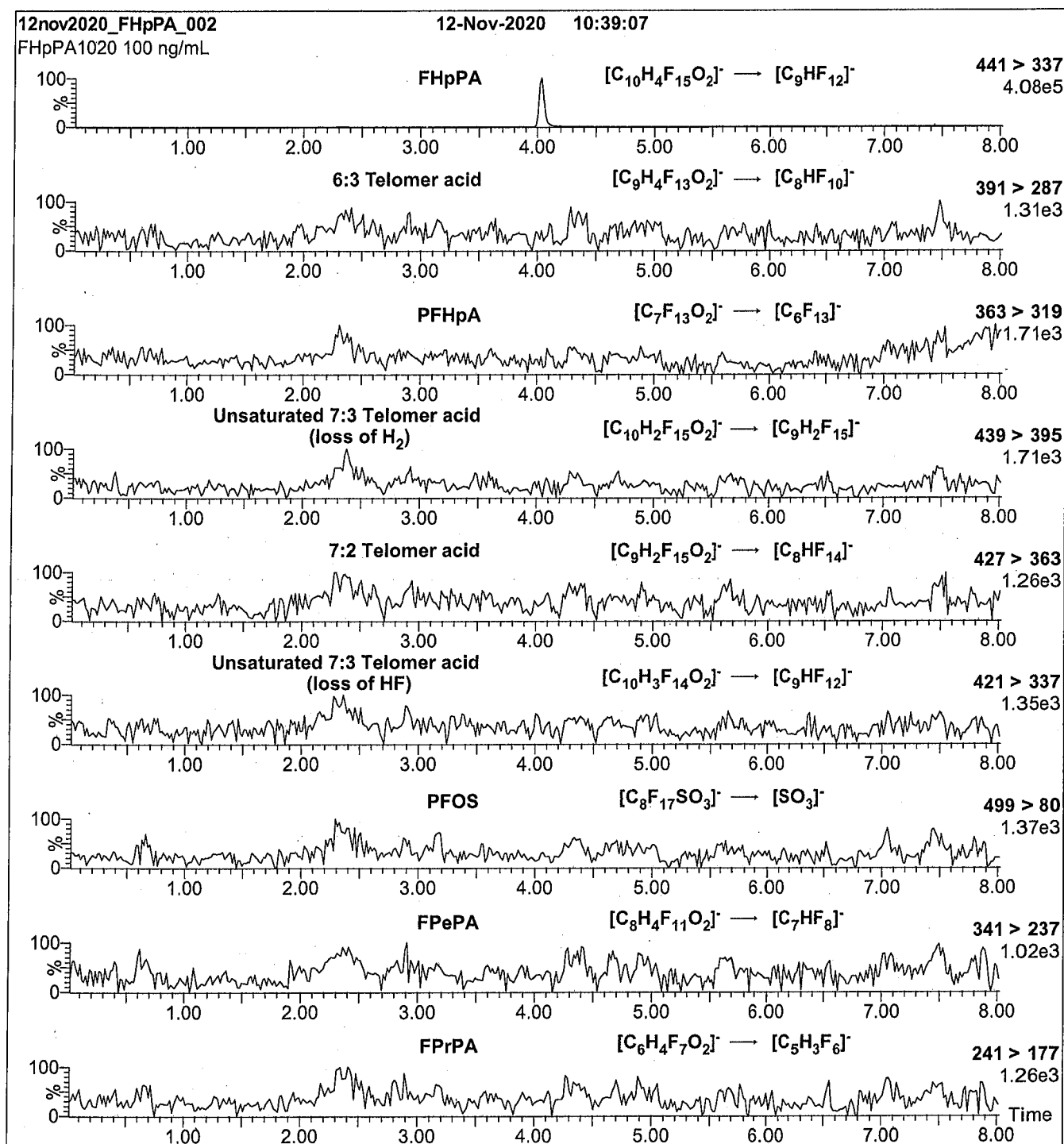
Source: Electrospray (negative)

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  $\mu$ 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#:
Final Volume (mLs):	1	Department:	PFAS (HFA1020)
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



# WELLINGTON LABORATORIES

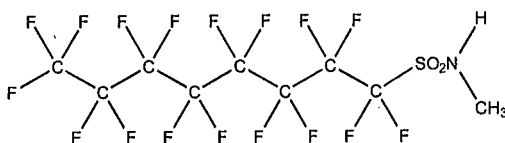
## CERTIFICATE OF ANALYSIS DOCUMENTATION

**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

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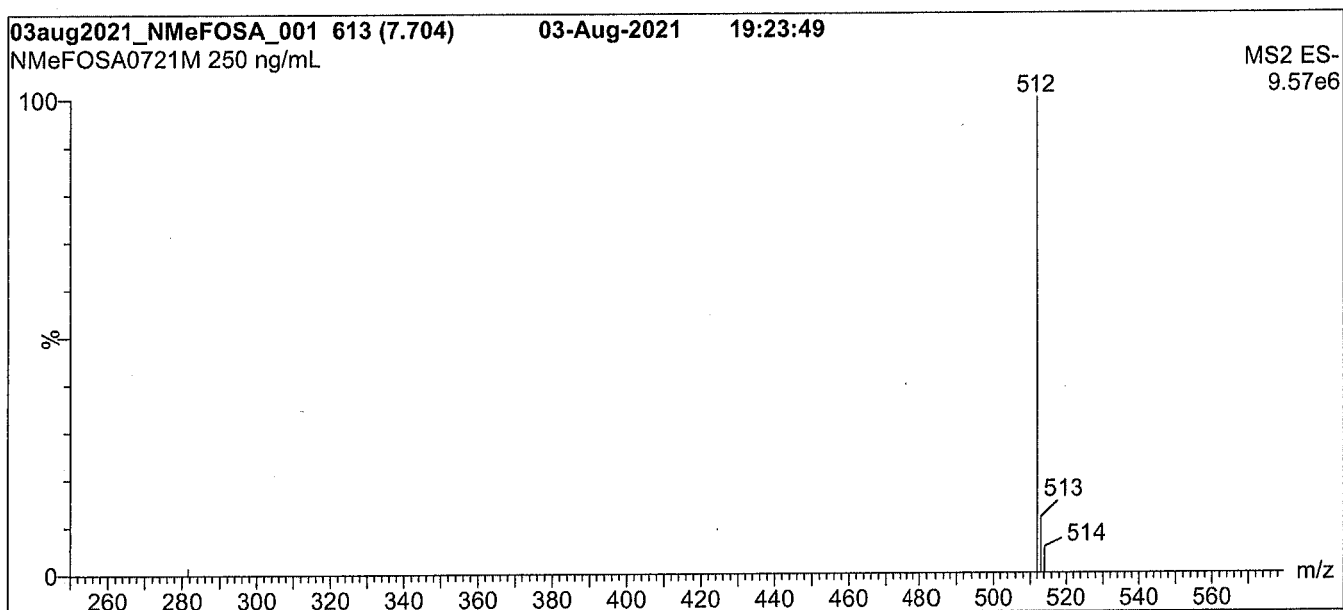
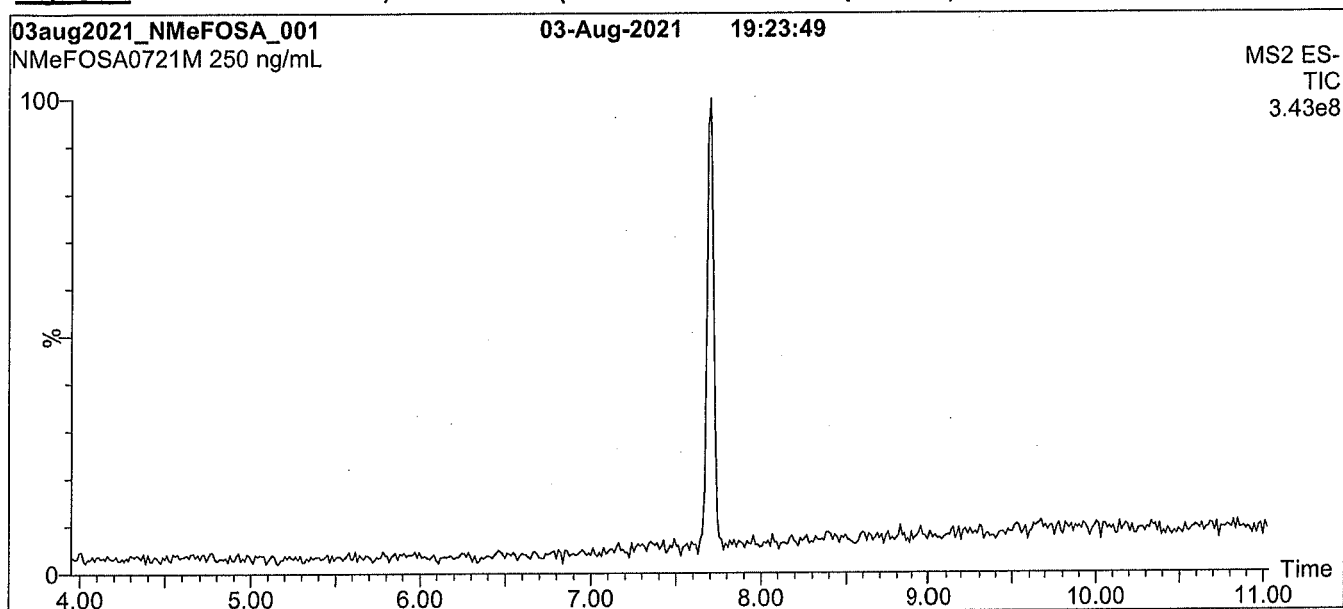
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**Figure 1: N-MeFOSA-M; LC/MS Data (Full Scan and Mass Spectrum)****Conditions for Figure 1:**

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

**Chromatographic Conditions:**

Column: Acquity UPLC BEH Shield RP<sub>18</sub>  
1.7  $\mu$ m, 2.1 x 100 mm

Mobile phase: Gradient

Start: 40% H<sub>2</sub>O / 60% (80:20 MeOH:ACN)  
(both with 10 mM NH<sub>4</sub>OAc 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  $\mu$ L/min

**MS Parameters:**

Experiment: Full Scan (250 - 850 amu)

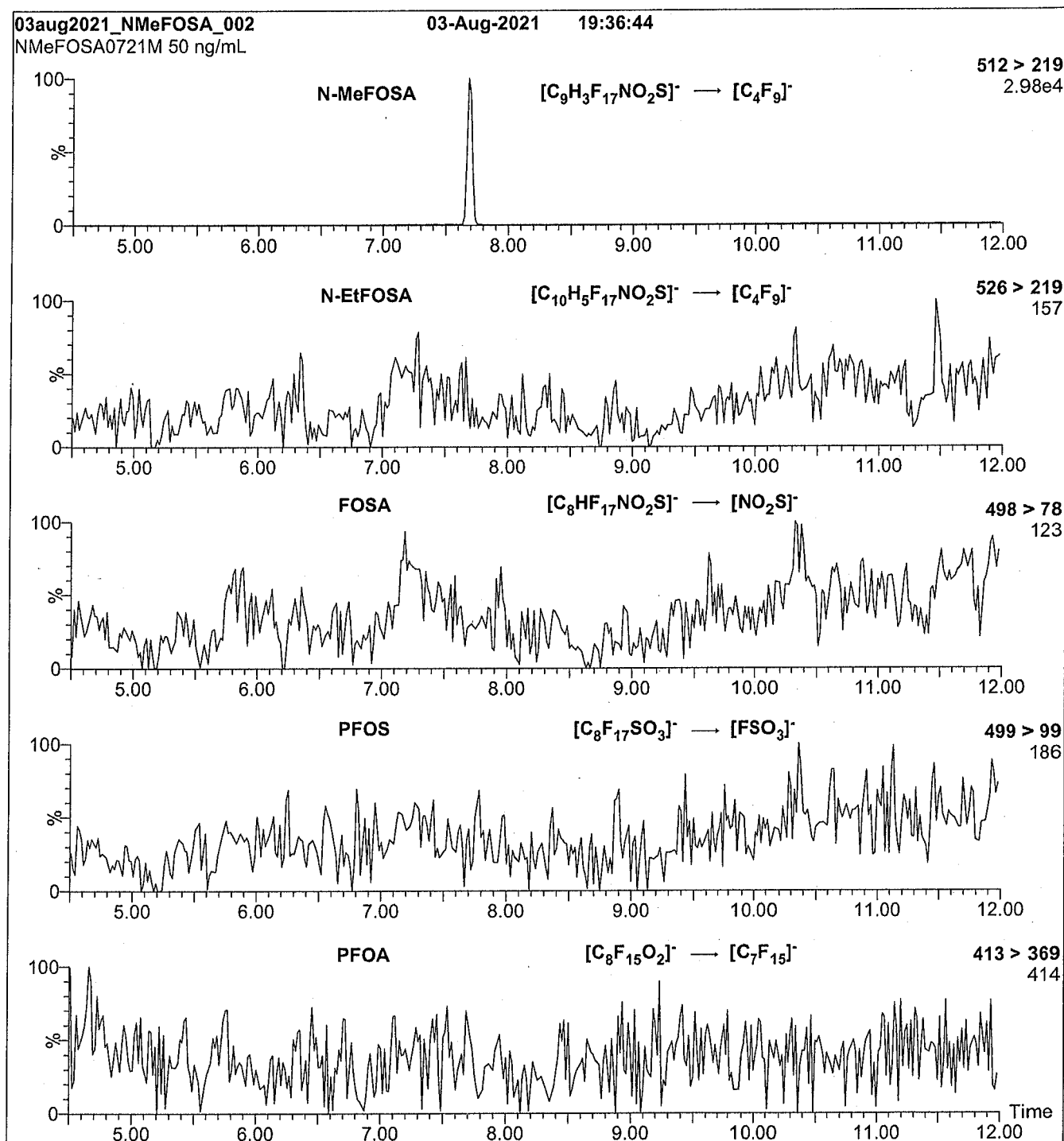
Source: Electrospray (negative)

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  $\mu$ 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



# WELLINGTON LABORATORIES

## CERTIFICATE OF ANALYSIS DOCUMENTATION

**PRODUCT CODE:**

N-EtFOSA-M

**LOT NUMBER:**

NEtFOSA0821M

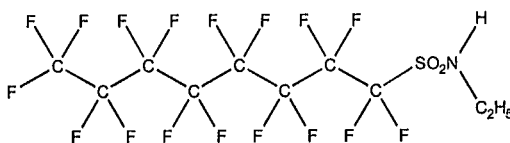
**COMPOUND:**

N-ethylperfluoro-1-octanesulfonamide

22C0313

**STRUCTURE:****CAS #:**

4151-50-2

**MOLECULAR FORMULA:** $C_{10}H_{17}F_{17}NO_2S$ **MOLECULAR WEIGHT:**

527.20

**CONCENTRATION:** $50.0 \pm 2.5 \mu\text{g/mL}$ **SOLVENT(S):**

Methanol

**CHEMICAL PURITY:**

&gt;98%

**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.

**FOR LABORATORY USE ONLY: NOT FOR HUMAN OR DRUG USE**

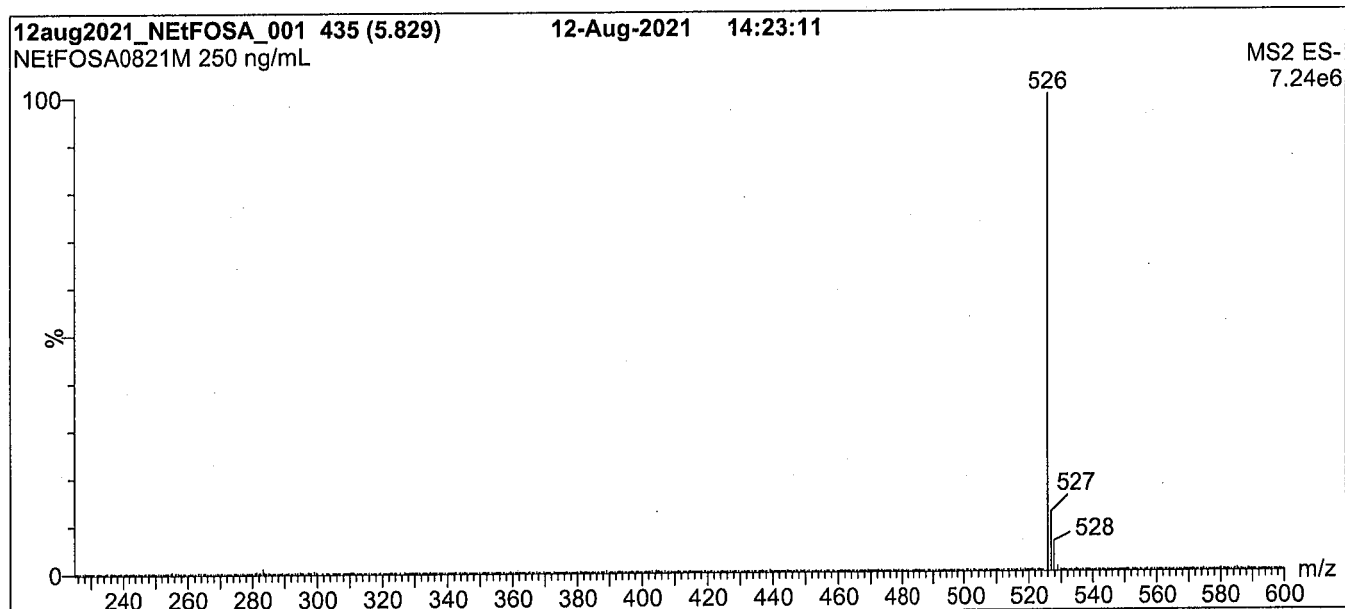
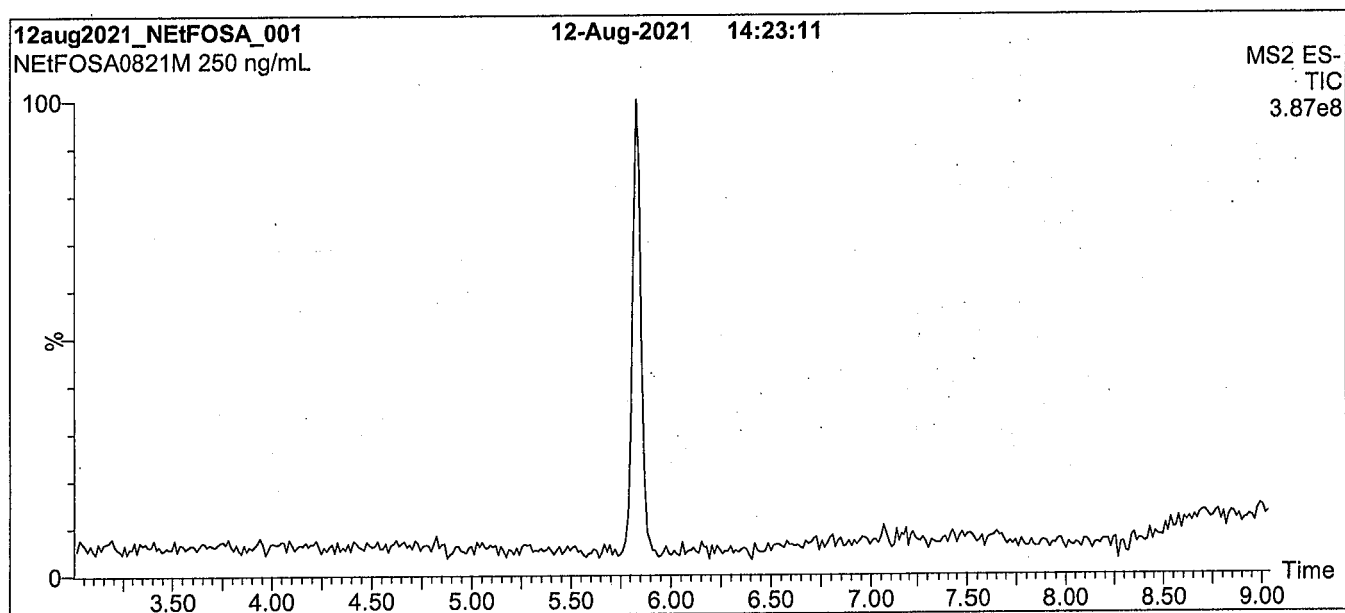
Certified By:

B.G. Chittim, General Manager

Date: 08/16/2021

(mm/dd/yyyy)

Wellington Laboratories Inc., 345 Southgate Dr. Guelph ON N1G 3M5 CANADA  
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**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 RP<sub>18</sub>  
1.7  $\mu$ m, 2.1 x 100 mm

Mobile phase: Gradient

Start: 30% H<sub>2</sub>O / 70% (80:20 MeOH:ACN)  
(both with 10 mM NH<sub>4</sub>OAc 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  $\mu$ 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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**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](http://www.well-labs.com) or contact us directly at [info@well-labs.com](mailto:info@well-labs.com)\*\*

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

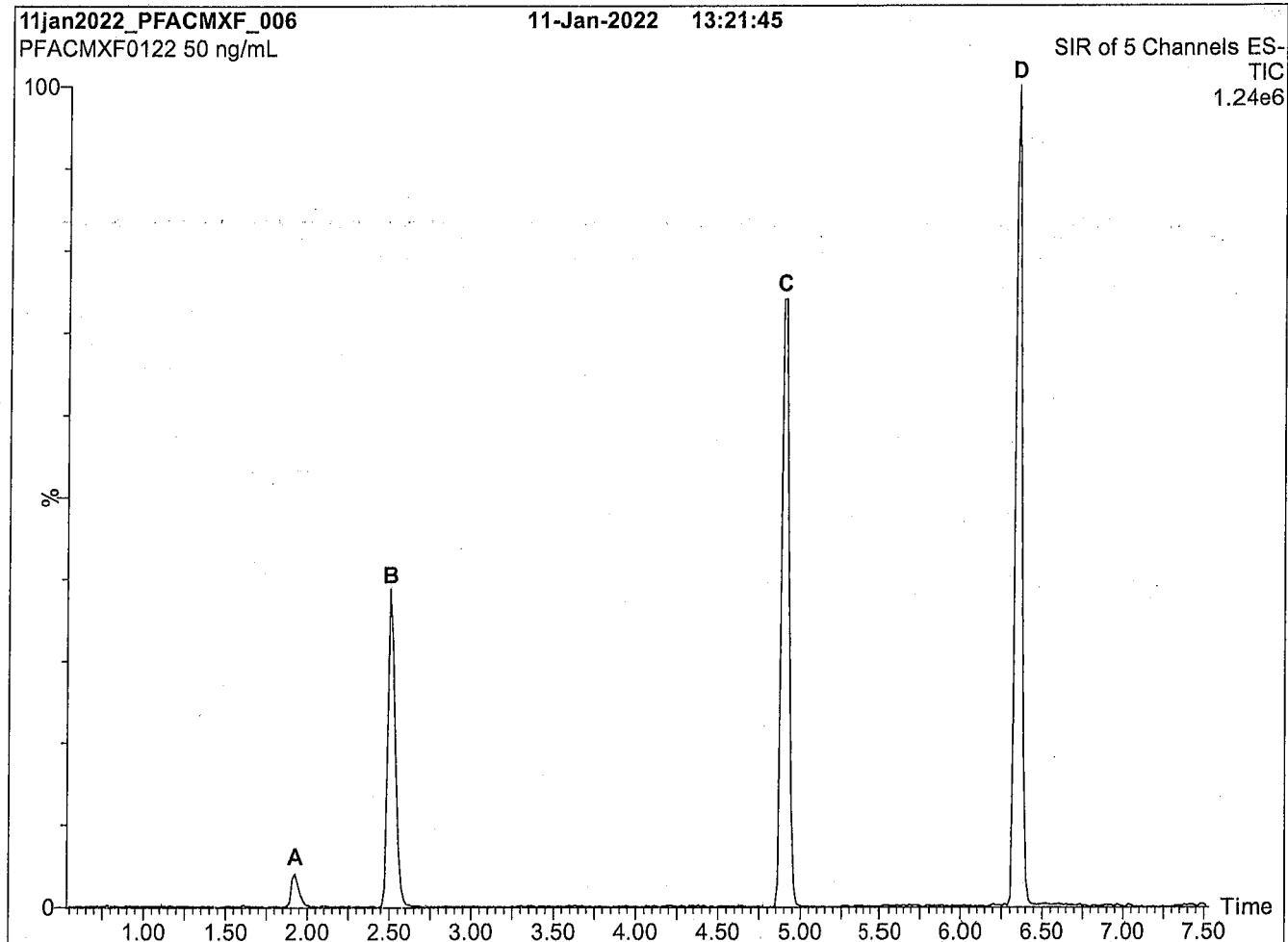
Compound	Acronym	Concentration* (ng/ml)		Peak Assignment in Figure 1
2,3,3,3-Tetrafluoro-2-(1,1,2,2,3,3,3-heptafluoropropoxy)-propanoic acid	HFPO-DA	2000		A
Compound	Acronym	Concentration* (ng/mL)		Peak Assignment in Figure 1
		as the salt	as the acid	
Sodium dodecafluoro-3H-4,8-dioxanonanoate	NaDONA	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 RP<sub>18</sub>  
1.7  $\mu$ m, 2.1 x 100 mm

Mobile phase: Gradient

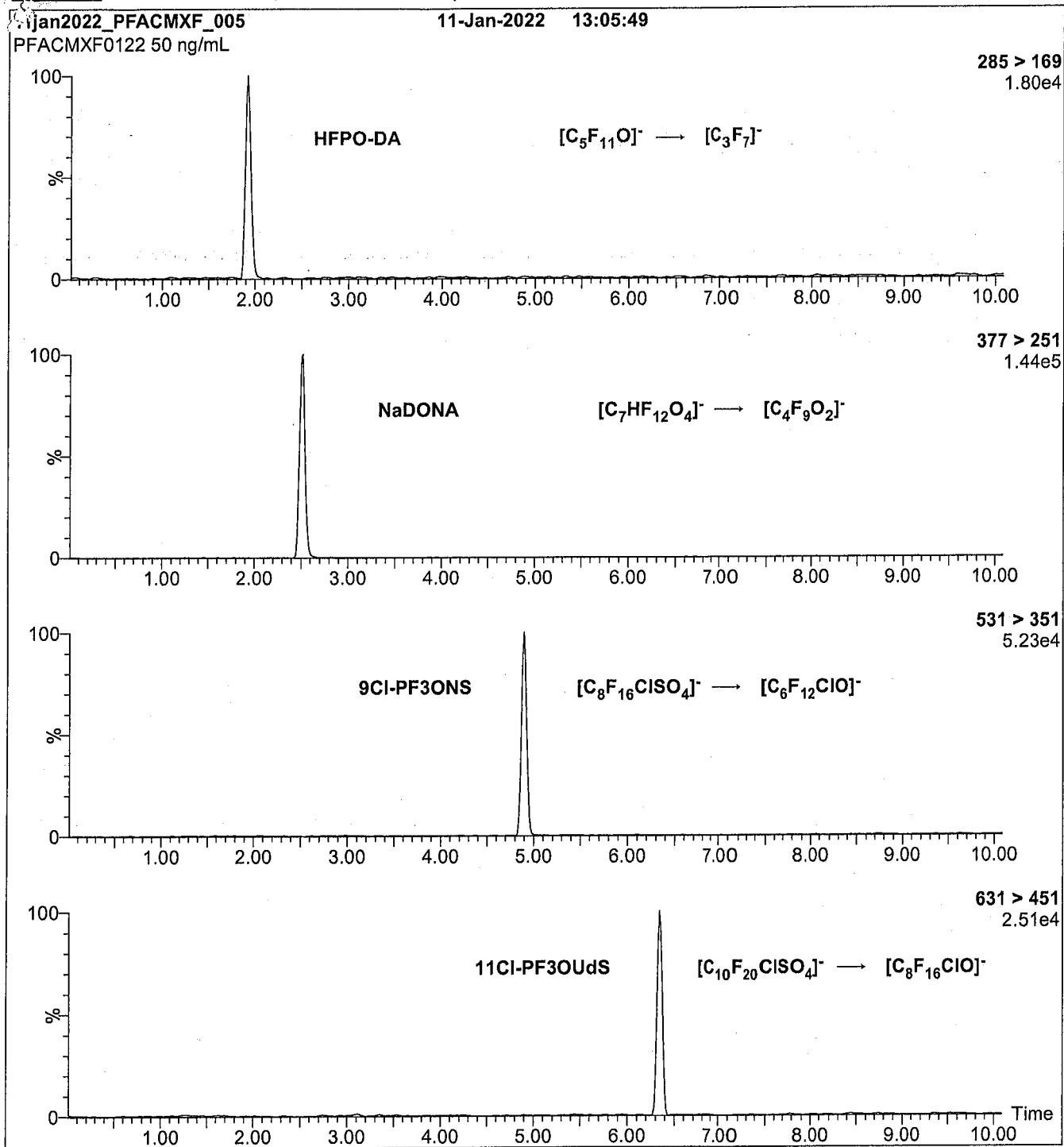
Start: 45% H<sub>2</sub>O / 55% (80:20 MeOH:ACN)  
(both with 10 mM NH<sub>4</sub>OAc buffer)  
Ramp to 90% organic over 8 min and hold for 2 min  
before returning to initial conditions in 0.75 min.  
Time: 12 min

Flow: 300  $\mu$ 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  $\mu$ 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 (mL): 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 ( $C_4$ ,  $C_5$ ,  $C_7$ ,  $C_9$ ,  $C_{10}$  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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**SYNTHESIS / CHARACTERIZATION:**

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**HOMOGENEITY:**

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

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

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**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 <sup>a</sup>	N-MeFOSAA: linear isomer	0.760		20
	N-MeFOSAA: ∑ branched isomers	0.240		17
N-ethylperfluorooctanesulfonamidoacetic acid <sup>b</sup>	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 <sup>c</sup>	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 <sup>d</sup>	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

<sup>a</sup> See Table B for percent composition of linear and branched N-MeFOSAA isomers.

<sup>b</sup> See Table C for percent composition of linear and branched N-EtFOSAA isomers.

<sup>c</sup> See Table D for percent composition of linear and branched PFHxSK isomers.

<sup>d</sup> 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  $^{19}\text{F}$ -NMR)\*

Isomer	Compound	Structure	Percent Composition by $^{19}\text{F}$ -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  $^{19}\text{F}$ -NMR)\***

Isomer	Compound	Structure	Percent Composition by $^{19}\text{F}$ -NMR	
1	N-ethylperfluoro-1-octanesulfonamidoacetic acid	$\text{CF}_3(\text{CF}_2)_7\text{SO}_2\text{NCH}_2\text{CO}_2\text{H}$ $\text{C}_2\text{H}_5$	77.5	77.5
2	N-ethylperfluoro-3-methylheptanesulfonamidoacetic acid	$\text{CF}_3(\text{CF}_2)_3\text{CF}(\text{CF}_2)_2\text{SO}_2\text{NCH}_2\text{CO}_2\text{H}$ $\text{CF}_3$ $\text{C}_2\text{H}_5$	2.3	22.5
3	N-ethylperfluoro-4-methylheptanesulfonamidoacetic acid	$\text{CF}_3(\text{CF}_2)_2\text{CF}(\text{CF}_2)_3\text{SO}_2\text{NCH}_2\text{CO}_2\text{H}$ $\text{CF}_3$ $\text{C}_2\text{H}_5$	2.2	
4	N-ethylperfluoro-5-methylheptanesulfonamidoacetic acid	$\text{CF}_3\text{CF}_2\text{CF}(\text{CF}_2)_4\text{SO}_2\text{NCH}_2\text{CO}_2\text{H}$ $\text{CF}_3$ $\text{C}_2\text{H}_5$	5.4	
5	N-ethylperfluoro-6-methylheptanesulfonamidoacetic acid	$\text{CF}_3\text{CF}(\text{CF}_2)_5\text{SO}_2\text{NCH}_2\text{CO}_2\text{H}$ $\text{CF}_3$ $\text{C}_2\text{H}_5$	10.4	
6	N-ethylperfluoro-5,5-dimethylhexanesulfonamidoacetic acid	$\text{CF}_3$ $\text{CF}_3\text{C}(\text{CF}_2)_4\text{SO}_2\text{NCH}_2\text{CO}_2\text{H}$ $\text{CF}_3$ $\text{C}_2\text{H}_5$	0.3	
7	N-ethylperfluoro-4,5-dimethylhexanesulfonamidoacetic acid	$\text{CF}_3$ $\text{CF}_3\text{CFCF}(\text{CF}_2)_3\text{SO}_2\text{NCH}_2\text{CO}_2\text{H}$ $\text{CF}_3$ $\text{C}_2\text{H}_5$	0.3	
8	N-ethylperfluoro-3,5-dimethylhexanesulfonamidoacetic acid	$\text{CF}_3$ $\text{CF}_3\text{CFCF}_2\text{CF}(\text{CF}_2)_2\text{SO}_2\text{NCH}_2\text{CO}_2\text{H}$ $\text{CF}_3$ $\text{C}_2\text{H}_5$	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  $^{19}\text{F}$ -NMR)\***

Isomer	Compound	Structure	Percent Composition by $^{19}\text{F}$ -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.

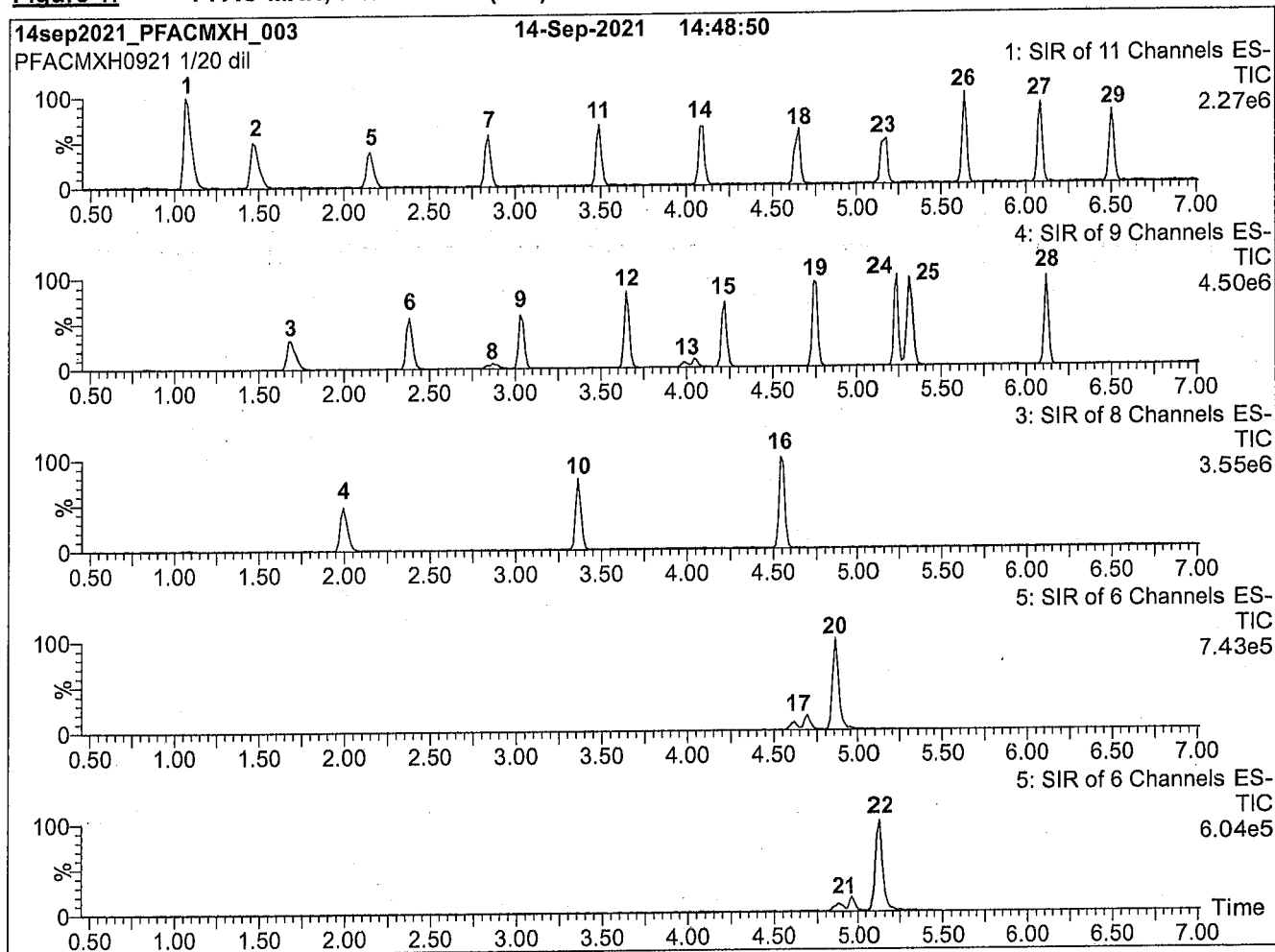
\*\* Systematic Name: Potassium perfluorohexane-2-sulfonate.

**Table E: PFOSK; Isomeric Components and Percent Composition (by  $^{19}\text{F}$ -NMR)\***

Isomer	Compound	Structure	Percent Composition by $^{19}\text{F}$ -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**	$\begin{array}{c} \text{CF}_3\text{CF}_2\text{CF}_2\text{CF}_2\text{CF}_2\text{CF}_2\text{CFSO}_3^-\text{K}^+ \\   \\ \text{CF}_3 \end{array}$	1.2	21.1
3	Potassium 2-trifluoromethylperfluoroheptanesulfonate	$\begin{array}{c} \text{CF}_3\text{CF}_2\text{CF}_2\text{CF}_2\text{CF}_2\text{CFCF}_2\text{SO}_3^-\text{K}^+ \\   \\ \text{CF}_3 \end{array}$	0.6	
4	Potassium 3-trifluoromethylperfluoroheptanesulfonate	$\begin{array}{c} \text{CF}_3\text{CF}_2\text{CF}_2\text{CF}_2\text{CFCF}_2\text{CF}_2\text{SO}_3^-\text{K}^+ \\   \\ \text{CF}_3 \end{array}$	1.9	
5	Potassium 4-trifluoromethylperfluoroheptanesulfonate	$\begin{array}{c} \text{CF}_3\text{CF}_2\text{CF}_2\text{CFCF}_2\text{CF}_2\text{CF}_2\text{SO}_3^-\text{K}^+ \\   \\ \text{CF}_3 \end{array}$	2.2	
6	Potassium 5-trifluoromethylperfluoroheptanesulfonate	$\begin{array}{c} \text{CF}_3\text{CF}_2\text{CFCF}_2\text{CF}_2\text{CF}_2\text{CF}_2\text{SO}_3^-\text{K}^+ \\   \\ \text{CF}_3 \end{array}$	4.5	
7	Potassium 6-trifluoromethylperfluoroheptanesulfonate	$\begin{array}{c} \text{CF}_3\text{CFCF}_2\text{CF}_2\text{CF}_2\text{CF}_2\text{CF}_2\text{SO}_3^-\text{K}^+ \\   \\ \text{CF}_3 \end{array}$	10.0	
8	Potassium 5,5-di(trifluoromethyl)perfluorohexanesulfonate	$\begin{array}{c} \text{CF}_3 \\   \\ \text{CF}_3\text{CCF}_2\text{CF}_2\text{CF}_2\text{CF}_2\text{SO}_3^-\text{K}^+ \\   \\ \text{CF}_3 \end{array}$	0.2	
9	Potassium 4,4-di(trifluoromethyl)perfluorohexanesulfonate	$\begin{array}{c} \text{CF}_3 \\   \\ \text{CF}_3\text{CF}_2\text{CCF}_2\text{CF}_2\text{CF}_2\text{SO}_3^-\text{K}^+ \\   \\ \text{CF}_3 \end{array}$	0.03	
10	Potassium 4,5-di(trifluoromethyl)perfluorohexanesulfonate	$\begin{array}{c} \text{CF}_3 \\   \\ \text{CF}_3\text{CFCF}_2\text{CF}_2\text{CF}_2\text{CF}_2\text{SO}_3^-\text{K}^+ \\   \\ \text{CF}_3 \end{array}$	0.4	
11	Potassium 3,5-di(trifluoromethyl)perfluorohexanesulfonate	$\begin{array}{c} \text{CF}_3 \\   \\ \text{CF}_3\text{CFCF}_2\text{CFCF}_2\text{CF}_2\text{SO}_3^-\text{K}^+ \\   \\ \text{CF}_3 \end{array}$	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 RP<sub>18</sub>  
1.7  $\mu$ m, 2.1 x 100 mm

Mobile phase: Gradient

Start: 50% H<sub>2</sub>O / 50% (80:20 MeOH:ACN)  
(both with 10 mM NH<sub>4</sub>OAc buffer)  
Ramp to 90% organic over 9 min and hold for 2 min  
before returning to initial conditions in 1 min.  
Time: 15 min

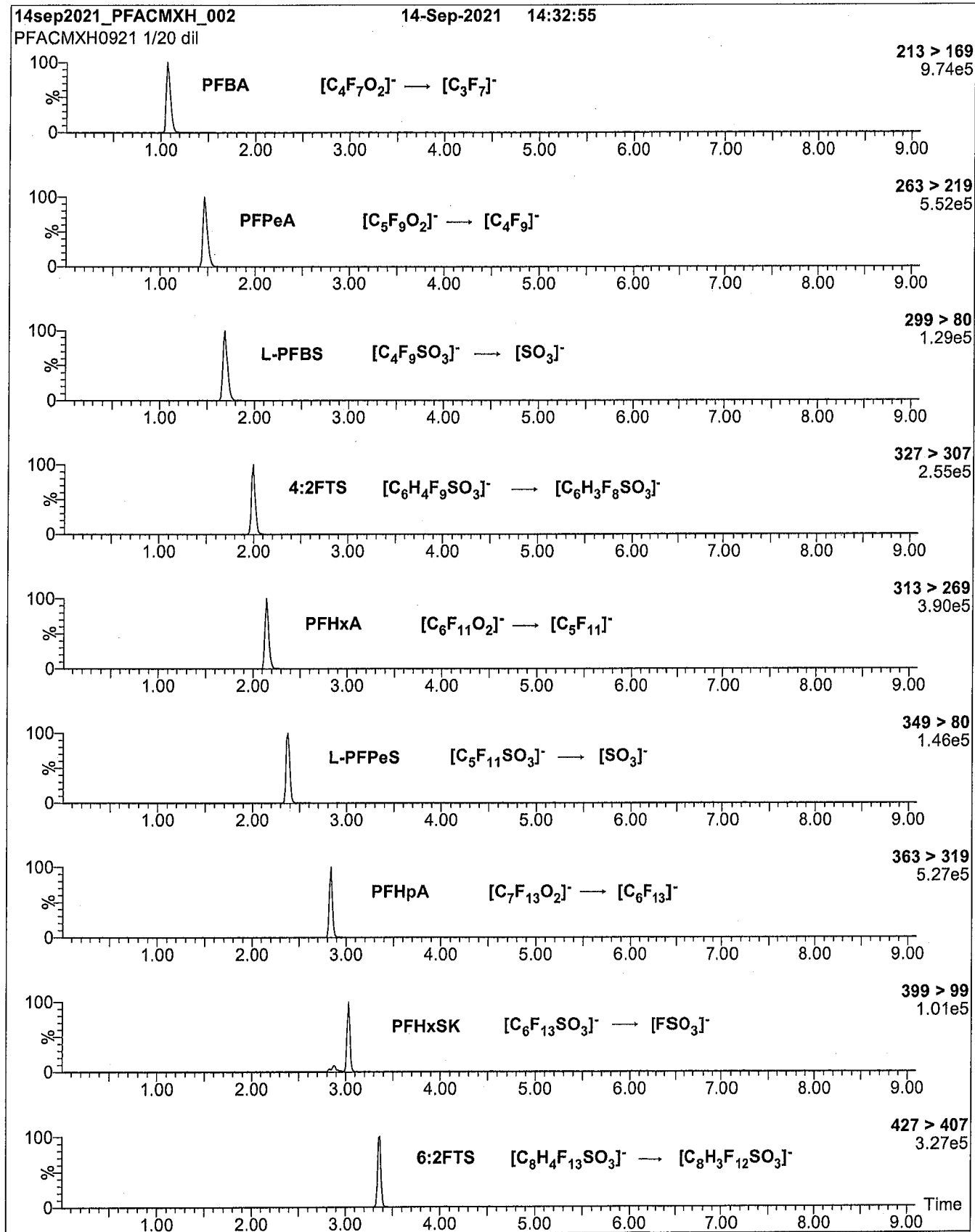
Flow: 300  $\mu$ L/min

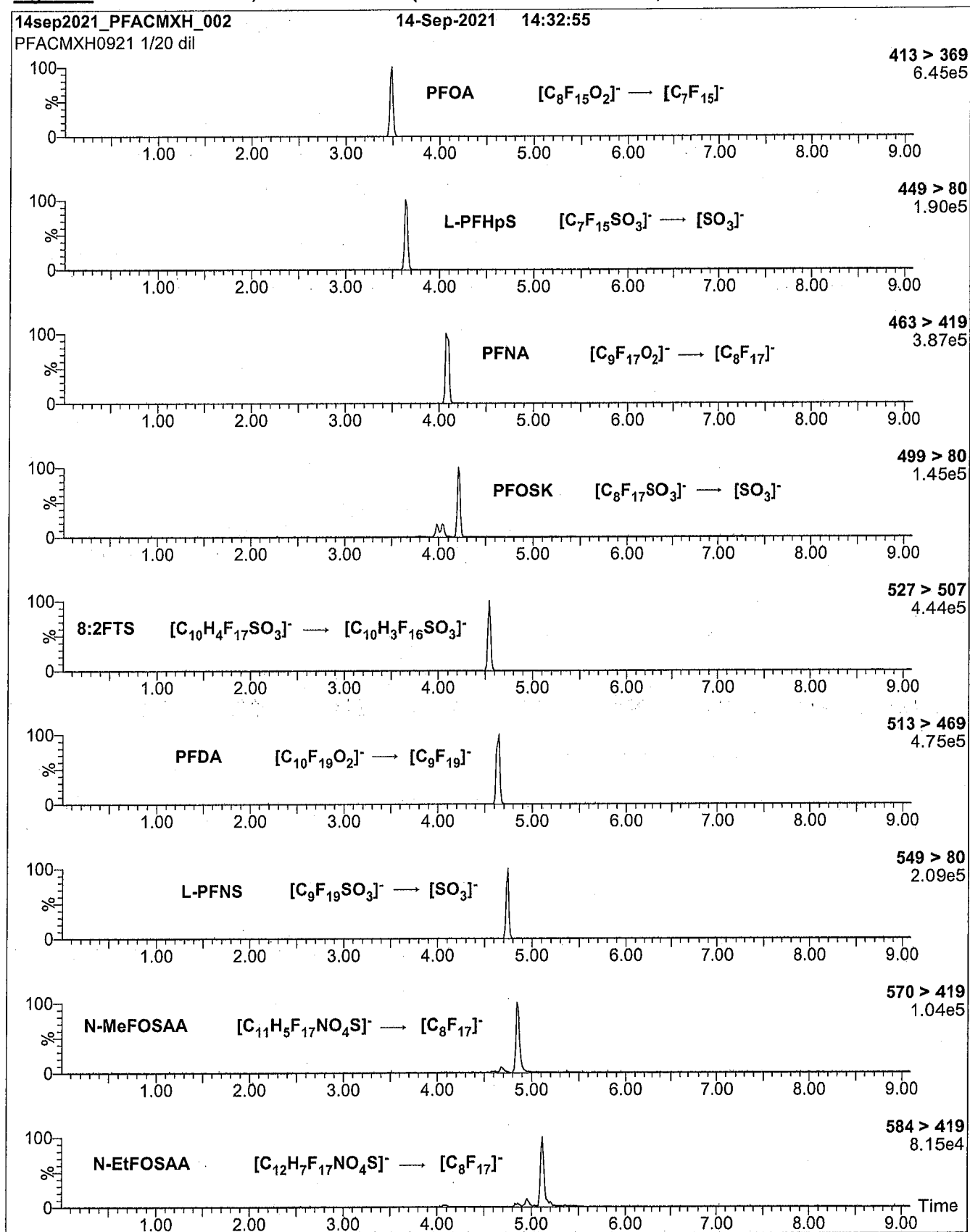
**MS Parameters:**

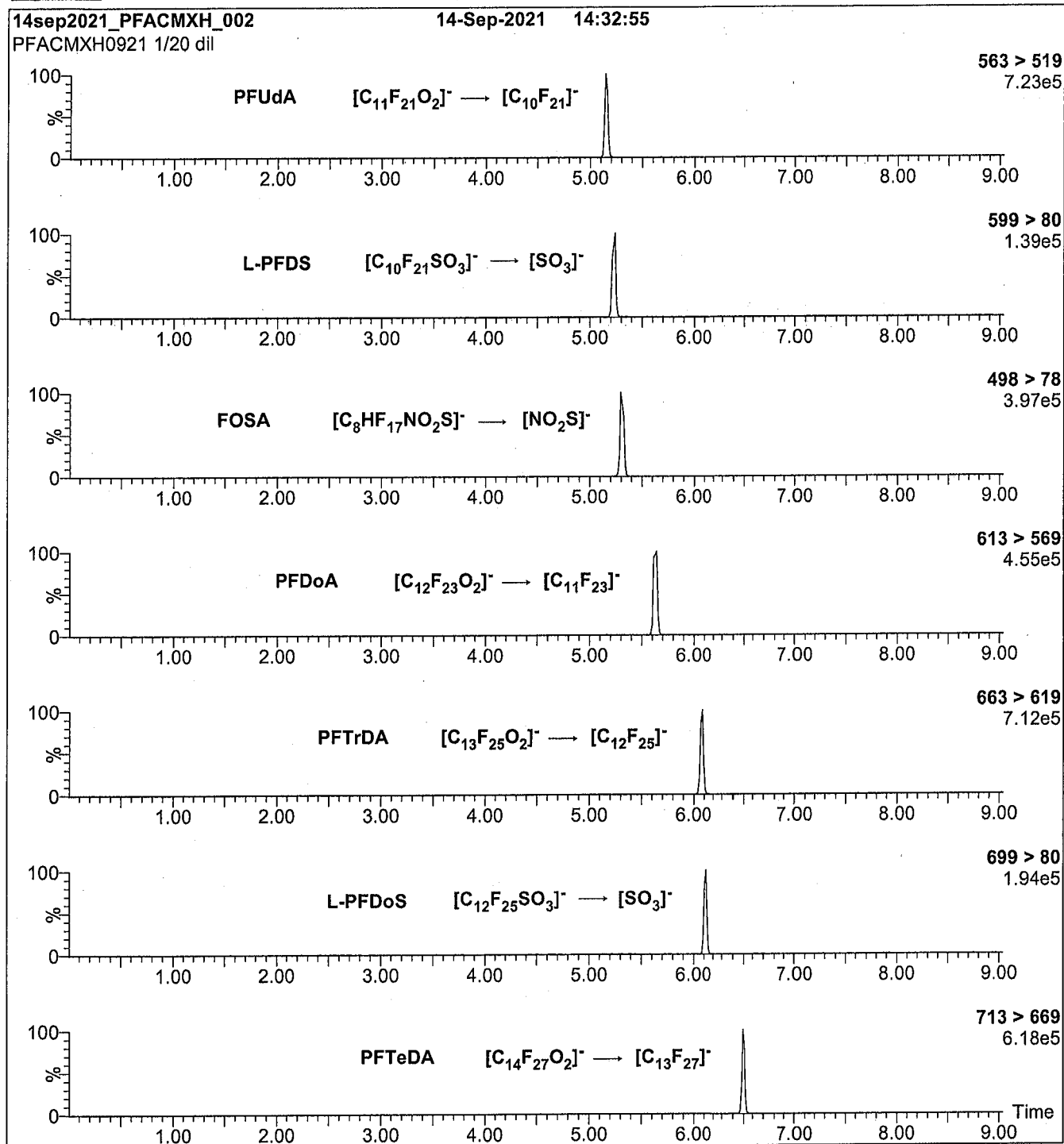
Experiment: SIR

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  $\mu$ 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.

**FOR LABORATORY USE ONLY: NOT FOR HUMAN OR DRUG USE**

**Wellington Laboratories Inc., 345 Southgate Dr. Guelph ON N1G 3M5 CANADA**  
**519-822-2436 • Fax: 519-822-2849 • [Info@well-labs.com](mailto: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 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](http://www.well-labs.com) or contact us directly at [info@well-labs.com](mailto: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

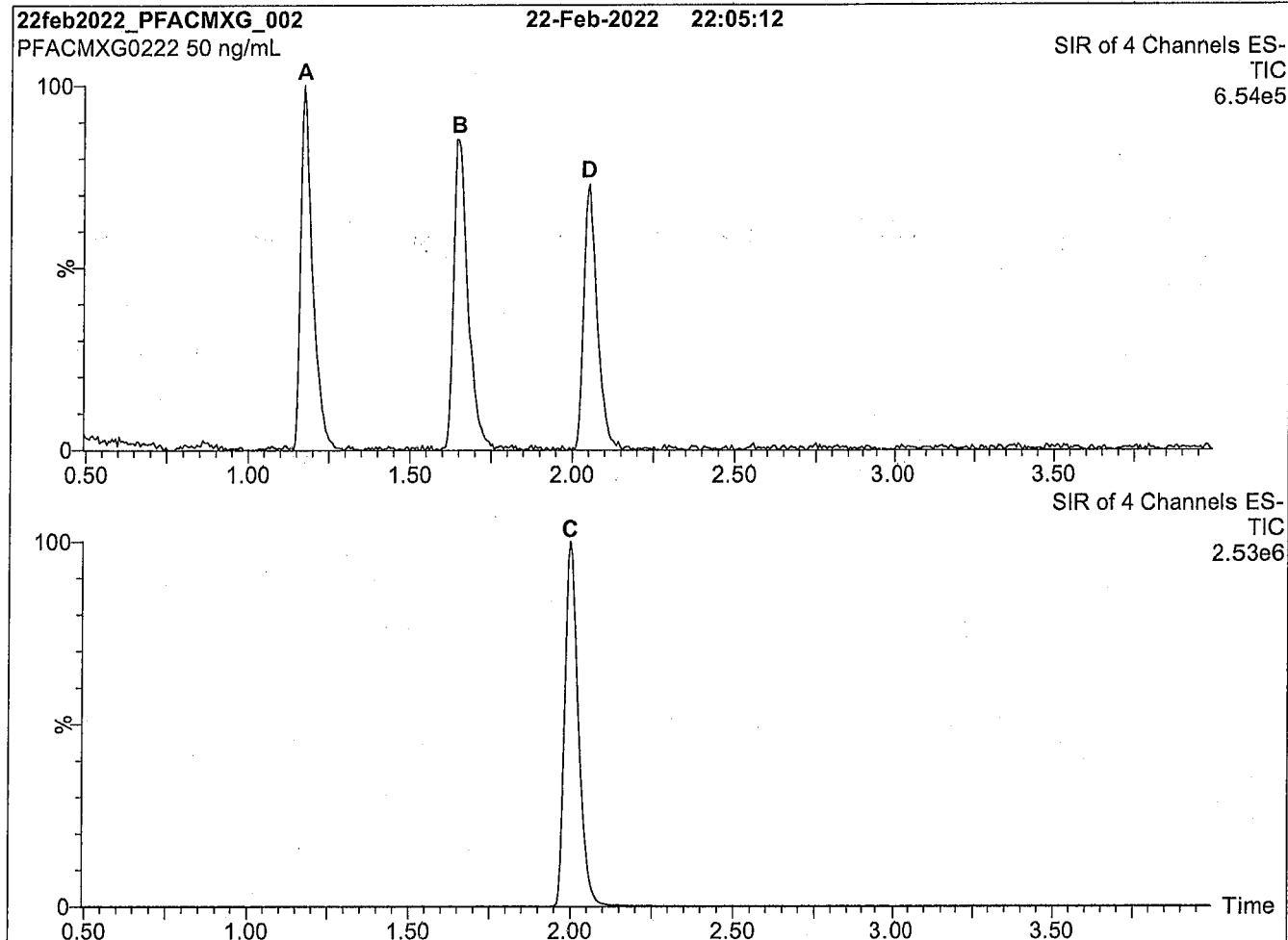
\* Concentrations have been rounded to three significant figures.

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 RP<sub>18</sub>  
1.7  $\mu$ m, 2.1 x 100 mm

**Mobile phase:** Gradient

Start: 50% H<sub>2</sub>O / 50% (80:20 MeOH:ACN)  
(both with 10 mM NH<sub>4</sub>OAc 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  $\mu$ L/min

**MS Parameters:**

Experiment: SIR

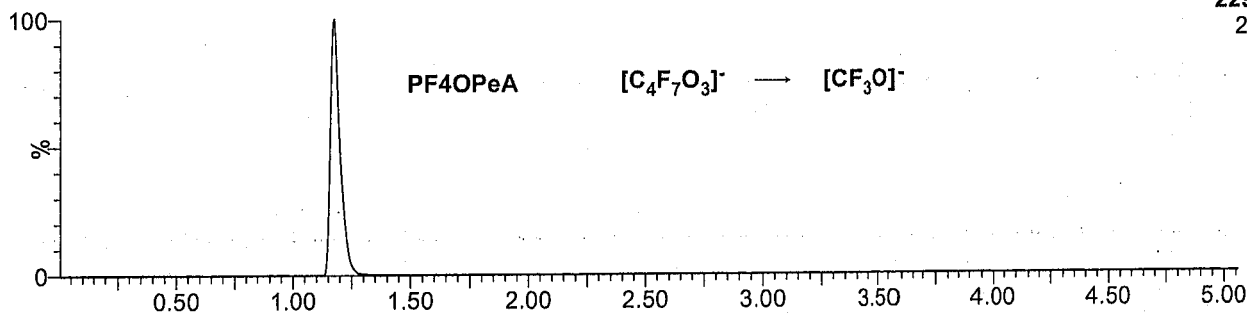
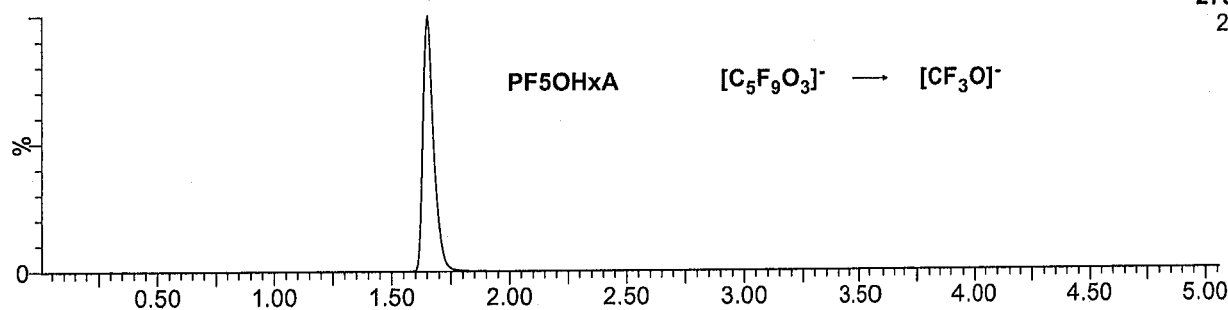
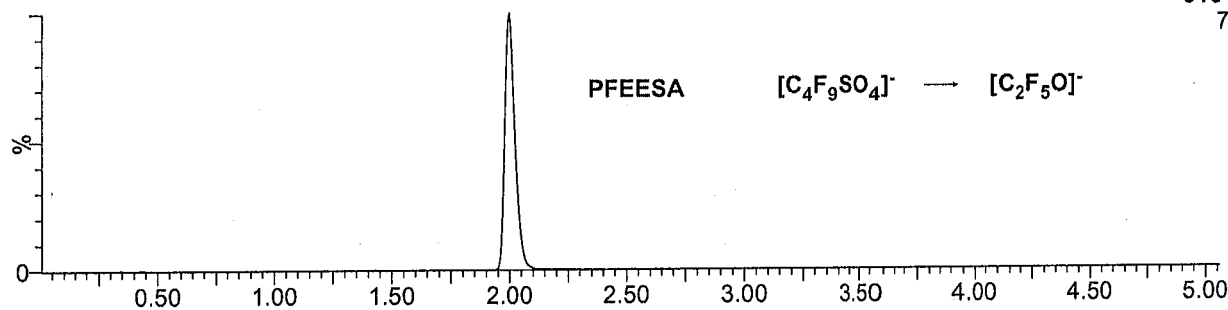
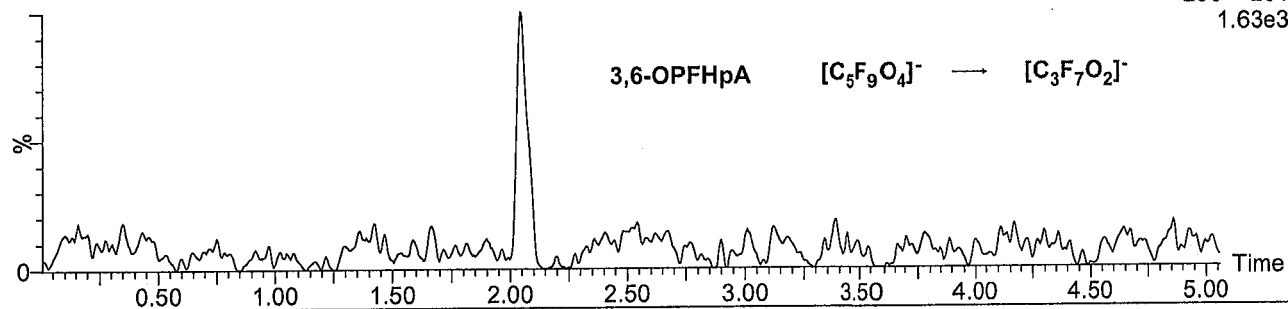
Source: Electrospray (negative)  
Capillary Voltage (kV) = 1.00  
Cone Voltage (V) = variable (15-35)  
Desolvation Temperature ( $^{\circ}$ C) = 500  
Desolvation Gas Flow (L/hr) = 1000

**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  $\mu$ 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

**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



# WELLINGTON LABORATORIES

## 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:**  $C_8F_{16}O_2NH_4$   
**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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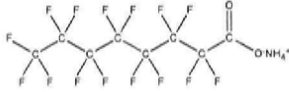
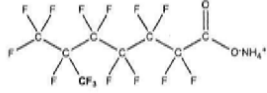
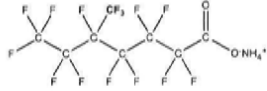
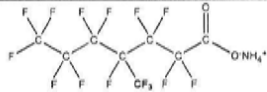
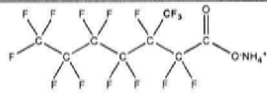
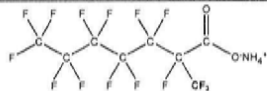
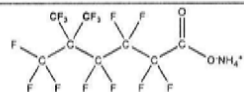
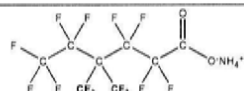
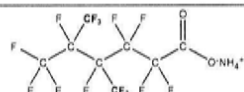
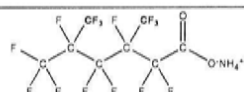
Certified By:   
 B.G. Chittim, General Manager

Date: 01/27/2022  
 (mm/dd/yyyy)

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**Table A:** T-PFOA; Isomeric Components and Percent Composition (by  $^{19}\text{F}$ -NMR)\*

Isomer	Compound	Structure	Percent Composition by $^{19}\text{F}$ -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  $^{19}\text{F}$ -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#: 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

**WELLINGTON  
LABORATORIES****CERTIFICATE OF ANALYSIS  
DOCUMENTATION****br-FOSA****Perfluorooctanesulfonamide  
Isomeric Mix**

**PRODUCT CODE:** br-FOSA  
**LOT NUMBER:** brFOSA0922  
**CONCENTRATION:** 50.0 ± 2.5 µg/mL  
**SOLVENT(S):** Isopropanol  
**DATE PREPARED:** (mm/dd/yyyy) 09/14/2022  
**LAST TESTED:** (mm/dd/yyyy) 10/07/2022  
**EXPIRY DATE:** (mm/dd/yyyy) 10/07/2027  
**RECOMMENDED STORAGE:** 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 <sup>19</sup>F-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  $^{19}\text{F}$ -NMR)\***

Isomer	Compound	Structure	Percent Composition by $^{19}\text{F}$ -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**	$\text{CF}_3(\text{CF}_2)_5\text{CF}(\text{SO}_2\text{NH}_2)\text{CF}_3$	0.8
3	Perfluoro-2-methyl-1-heptanesulfonamide	$\text{CF}_3(\text{CF}_2)_4\text{CF}(\text{CF}_3)\text{CF}_2\text{SO}_2\text{NH}_2$	0.3
4	Perfluoro-3-methyl-1-heptanesulfonamide	$\text{CF}_3(\text{CF}_2)_3\text{CF}(\text{CF}_3)\text{CF}_2\text{SO}_2\text{NH}_2$	4.2
5	Perfluoro-4-methyl-1-heptanesulfonamide	$\text{CF}_3(\text{CF}_2)_2\text{CF}(\text{CF}_3)\text{CF}_2\text{SO}_2\text{NH}_2$	3.5
6	Perfluoro-5-methyl-1-heptanesulfonamide	$\text{CF}_3\text{CF}_2\text{CF}(\text{CF}_3)\text{CF}_2\text{SO}_2\text{NH}_2$	7.8
7	Perfluoro-6-methyl-1-heptanesulfonamide	$\text{CF}_3\text{CF}(\text{CF}_3)\text{CF}_2\text{SO}_2\text{NH}_2$	16.8
8	Perfluoro-5,5-dimethyl-1-hexanesulfonamide	$\text{CF}_3\text{C}(\text{CF}_3)_2(\text{CF}_2)_4\text{SO}_2\text{NH}_2$	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#: 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



# WELLINGTON LABORATORIES

## CERTIFICATE OF ANALYSIS DOCUMENTATION

### br-NMeFOSA

#### N-Methylperfluorooctanesulfonamide Isomeric Mix

<b><u>PRODUCT CODE:</u></b>	br-NMeFOSA
<b><u>LOT NUMBER:</u></b>	brNMeFOSA0822
<b><u>CONCENTRATION:</u></b>	50.0 ± 2.5 µg/mL
<b><u>SOLVENT(S):</u></b>	Methanol
<b><u>DATE PREPARED:</u></b> (mm/dd/yyyy)	08/18/2022
<b><u>LAST TESTED:</u></b> (mm/dd/yyyy)	08/23/2022
<b><u>EXPIRY DATE:</u></b> (mm/dd/yyyy)	08/23/2027
<b><u>RECOMMENDED STORAGE:</u></b>	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 <sup>19</sup>F-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  $^{19}\text{F}$ -NMR)\***

Isomer	Compound	Structure	Percent Composition by $^{19}\text{F}$ -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#: bENETfOSA0922)
Final Volume (mls):	1	Department:	PPAS
Vials:	1	Last Edit:	10/18/2022 13:38 by HGH

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



# WELLINGTON LABORATORIES

## CERTIFICATE OF ANALYSIS DOCUMENTATION

### br-NEtFOSA

#### N-Ethylperfluorooctanesulfonamide Isomeric Mix

<b><u>PRODUCT CODE:</u></b>	br-NEtFOSA
<b><u>LOT NUMBER:</u></b>	brNEtFOSA0922
<b><u>CONCENTRATION:</u></b>	50.0 ± 2.5 µg/mL
<b><u>SOLVENT(S):</u></b>	Methanol
<b><u>DATE PREPARED:</u></b> (mm/dd/yyyy)	08/23/2022
<b><u>LAST TESTED:</u></b> (mm/dd/yyyy)	10/07/2022
<b><u>EXPIRY DATE:</u></b> (mm/dd/yyyy)	10/07/2027
<b><u>RECOMMENDED STORAGE:</u></b>	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 <sup>19</sup>F-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  $^{19}\text{F}$ -NMR)\***

Isomer	Compound	Structure	Percent Composition by $^{19}\text{F}$ -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	$\text{CF}_3\text{C}(\text{CF}_3)_2(\text{CF}_2)_4\text{SO}_2\text{NHCH}_2\text{CH}_3$	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

<b><u>PRODUCT CODE:</u></b>	br-NMeFOSE
<b><u>LOT NUMBER:</u></b>	brNMeFOSE0922
<b><u>CONCENTRATION:</u></b>	50.0 ± 2.5 µg/mL
<b><u>SOLVENT(S):</u></b>	Methanol
<b><u>DATE PREPARED:</u></b> (mm/dd/yyyy)	09/02/2022
<b><u>LAST TESTED:</u></b> (mm/dd/yyyy)	09/07/2022 (HRGC/LRMS) 10/07/2022 (LC/MS)
<b><u>EXPIRY DATE:</u></b> (mm/dd/yyyy)	10/07/2027
<b><u>RECOMMENDED STORAGE:</u></b>	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 <sup>19</sup>F-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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\* Percent of total 2-(N-methylperfluorooctanesulfonamido)ethanol isomers only.

Date: 11/14/2022  
(mm/dd/yyyy)

# 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:	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

<b>PRODUCT CODE:</b>	br-NEtFOSE
<b>LOT NUMBER:</b>	brNEtFOSE1022
<b>CONCENTRATION:</b>	50.0 ± 2.5 µg/mL
<b>SOLVENT(S):</b>	Methanol
<b>DATE PREPARED:</b> (mm/dd/yyyy)	09/12/2022
<b>LAST TESTED:</b> (mm/dd/yyyy)	09/12/2022 (HRGC/LRMS) 10/07/2022 (LC/MS)
<b>EXPIRY DATE:</b> (mm/dd/yyyy)	10/07/2027
<b>RECOMMENDED STORAGE:</b>	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 <sup>19</sup>F-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  $^{19}\text{F}$ -NMR)\***

Isomer	Compound	Structure	Percent Composition by $^{19}\text{F}$ -NMR
1	2-(N-Ethylperfluoro-1-octanesulfonamido)ethanol	$\text{CF}_3(\text{CF}_2)_7\text{SO}_2\text{NCH}_2\text{CH}_2\text{OH}$ $\text{CH}_2\text{CH}_3$	64.6
2	2-(N-Ethylperfluoro-2-methyl-1-heptanesulfonamido)ethanol	$\text{CF}_3(\text{CF}_2)_4\text{CFCF}_2\text{SO}_2\text{NCH}_2\text{CH}_2\text{OH}$ $\text{CF}_3 \quad \text{CH}_2\text{CH}_3$	0.2
3	2-(N-Ethylperfluoro-3-methyl-1-heptanesulfonamido)ethanol	$\text{CF}_3(\text{CF}_2)_3\text{CF}(\text{CF}_2)_2\text{SO}_2\text{NCH}_2\text{CH}_2\text{OH}$ $\text{CF}_3 \quad \text{CH}_2\text{CH}_3$	4.1
4	2-(N-Ethylperfluoro-4-methyl-1-heptanesulfonamido)ethanol	$\text{CF}_3(\text{CF}_2)_2\text{CF}(\text{CF}_2)_3\text{SO}_2\text{NCH}_2\text{CH}_2\text{OH}$ $\text{CF}_3 \quad \text{CH}_2\text{CH}_3$	4.3
5	2-(N-Ethylperfluoro-5-methyl-1-heptanesulfonamido)ethanol	$\text{CF}_3\text{CF}_2\text{CF}(\text{CF}_2)_4\text{SO}_2\text{NCH}_2\text{CH}_2\text{OH}$ $\text{CF}_3 \quad \text{CH}_2\text{CH}_3$	8.8
6	2-(N-Ethylperfluoro-6-methyl-1-heptanesulfonamido)ethanol	$\text{CF}_3\text{CF}(\text{CF}_2)_5\text{SO}_2\text{NCH}_2\text{CH}_2\text{OH}$ $\text{CF}_3 \quad \text{CH}_2\text{CH}_3$	17.8
7	2-(N-Ethylperfluoro-5,5-dimethyl-1-hexanesulfonamido)ethanol	$\text{CF}_3$ $\text{CF}_3\text{C}(\text{CF}_2)_4\text{SO}_2\text{NCH}_2\text{CH}_2\text{OH}$ $\text{CF}_3 \quad \text{CH}_2\text{CH}_3$	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:**

<b>Standard</b>	<b>Description</b>	<b>Prepared</b>	<b>Prepared By</b>	<b>Lot Nbr</b>	<b>Expires</b>	<b>Last Edit</b>	<b>(mls)</b>
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  
 Standard Type: Analyte Spike  
 Solvent: MeOH  
 Final Volume (mLs): 1.2  
 Vials: 1

Expires: 08/08/2027  
 Prepared: 08/05/2022  
 Prepared By: Wellington Laboratories (Lot#:  
 Department: PFAS MXH0822)  
 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

**23A0180**

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:	01/11/2023 14:40 by PAF
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 ( $^{13}\text{C}$ ) perfluoroalkylcarboxylic acids ( $\text{C}_4$ - $\text{C}_{12}$ ,  $\text{C}_{14}$ ), three mass-labelled ( $^{13}\text{C}$ ) perfluoroalkanesulfonates ( $\text{C}_4$ ,  $\text{C}_6$ , and  $\text{C}_8$ ), three mass-labelled (one  $^{13}\text{C}$  and two  $^2\text{H}$ ) perfluoro-1-octanesulfonamides, three mass-labelled ( $^{13}\text{C}$ ) fluorotelomer sulfonates (4:2, 6:2, and 8:2), two mass-labelled ( $^2\text{H}$ ) perfluorooctanesulfonamidoacetic acids, two mass-labelled ( $^2\text{H}$ ) perfluorooctanesulfonamidoethanols, and mass-labelled ( $^{13}\text{C}$ ) hexafluoropropylene oxide dimer acid (GenX, M3HFPO-DA). The components and their concentrations are given in Table A.

The individual  $^{13}\text{C}$ -labelled components all have chemical purities >98% and isotopic purities of  $\geq 99\%$ . The individual  $^2\text{H}$ -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.

**FOR LABORATORY USE ONLY: NOT FOR HUMAN OR DRUG USE**

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

**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-( <sup>13</sup> C <sub>4</sub> )butanoic acid	MPFBA	2000	1	
Perfluoro-n-( <sup>13</sup> C <sub>5</sub> )pentanoic acid	M5PFPeA	1000	2	
Perfluoro-n-(1,2,3,4,6- <sup>13</sup> C <sub>5</sub> )hexanoic acid	M5PFHxA	500	5	
Perfluoro-n-(1,2,3,4- <sup>13</sup> C <sub>6</sub> )heptanoic acid	M4PFHpA	500	7	
Perfluoro-n-( <sup>13</sup> C <sub>8</sub> )octanoic acid	M8PFOA	500	10	
Perfluoro-n-( <sup>13</sup> C <sub>9</sub> )nonanoic acid	M9PFNA	250	11	
Perfluoro-n-(1,2,3,4,5,6- <sup>13</sup> C <sub>6</sub> )decanoic acid	M6PFDA	250	14	
Perfluoro-n-(1,2,3,4,5,6,7- <sup>13</sup> C <sub>7</sub> )undecanoic acid	M7PFUdA	250	18	
Perfluoro-n-(1,2- <sup>13</sup> C <sub>2</sub> )dodecanoic acid	MPFDoA	250	19	
Perfluoro-n-(1,2- <sup>13</sup> C <sub>2</sub> )tetradecanoic acid	M2PFTeDA	250	22	
Perfluoro-1-( <sup>13</sup> C <sub>8</sub> )octanesulfonamide	M8FOSA	500	17	
N-methyl-d <sub>3</sub> -perfluoro-1-octanesulfonamide	d-N-MeFOSA	500	21	
N-ethyl-d <sub>5</sub> -perfluoro-1-octanesulfonamide	d-N-EtFOSA	500	24	
N-methyl-d <sub>3</sub> -perfluoro-1-octanesulfonamidoacetic acid	d3-N-MeFOSAA	1000	15	
N-ethyl-d <sub>5</sub> -perfluoro-1-octanesulfonamidoacetic acid	d5-N-EtFOSAA	1000	16	
2-(N-methyl-d <sub>3</sub> -perfluoro-1-octanesulfonamido)ethan-d <sub>4</sub> -ol	d7-N-MeFOSE	5000	20	
2-(N-ethyl-d <sub>5</sub> -perfluoro-1-octanesulfonamido)ethan-d <sub>4</sub> -ol	d9-N-EtFOSE	5000	23	
2,3,3,3-Tetrafluoro-2-(1,1,2,2,3,3,3-heptafluoropropoxy)( <sup>13</sup> C <sub>3</sub> )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- <sup>13</sup> C <sub>3</sub> )butanesulfonate	M3PFBS	500	466	3
Sodium perfluoro-1-(1,2,3- <sup>13</sup> C <sub>3</sub> )hexanesulfonate	M3PFHxS	500	474	8
Sodium perfluoro-1-( <sup>13</sup> C <sub>8</sub> )octanesulfonate	M8PFOS	500	479	12
Sodium 1H,1H,2H,2H-perfluoro-(1,2- <sup>13</sup> C <sub>2</sub> )hexanesulfonate	M2-4:2FTS	1000	938	4
Sodium 1H,1H,2H,2H-perfluoro-(1,2- <sup>13</sup> C <sub>2</sub> )octanesulfonate	M2-6:2FTS	1000	951	9
Sodium 1H,1H,2H,2H-perfluoro-(1,2- <sup>13</sup> C <sub>2</sub> )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

**23A0180**

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:	01/11/2023 14:40 by PAF
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

# 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 ( $^{13}\text{C}$ ) perfluoroalkylcarboxylic acids ( $\text{C}_4$ - $\text{C}_{12}$ ,  $\text{C}_{14}$ ), three mass-labelled ( $^{13}\text{C}$ ) perfluoroalkanesulfonates ( $\text{C}_4$ ,  $\text{C}_6$ , and  $\text{C}_8$ ), three mass-labelled (one  $^{13}\text{C}$  and two  $^2\text{H}$ ) perfluoro-1-octanesulfonamides, three mass-labelled ( $^{13}\text{C}$ ) fluorotelomer sulfonates (4:2, 6:2, and 8:2), two mass-labelled ( $^2\text{H}$ ) perfluorooctanesulfonamidoacetic acids, two mass-labelled ( $^2\text{H}$ ) perfluorooctanesulfonamidoethanols, and mass-labelled ( $^{13}\text{C}$ ) hexafluoropropylene oxide dimer acid (GenX, M3HFPO-DA). The components and their concentrations are given in Table A.

The individual  $^{13}\text{C}$ -labelled components all have chemical purities >98% and isotopic purities of  $\geq 99\%$ . The individual  $^2\text{H}$ -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.

**FOR LABORATORY USE ONLY: NOT FOR HUMAN OR DRUG USE**

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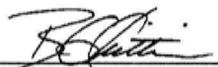


**Table A:** MPFAC-HIF-ES; Components and Concentrations  
(ng/mL,  $\pm$  5% in methanol/isopropanol (1%)/water (<1%))

Compound	Acronym	Concentration (ng/mL)	Peak Assignment in Figure 1	
Perfluoro-n-( <sup>13</sup> C <sub>4</sub> )butanoic acid	MPFBA	2000	1	
Perfluoro-n-( <sup>13</sup> C <sub>5</sub> )pentanoic acid	M5PFPeA	1000	2	
Perfluoro-n-(1,2,3,4,6- <sup>13</sup> C <sub>6</sub> )hexanoic acid	M5PFHxA	500	5	
Perfluoro-n-(1,2,3,4- <sup>13</sup> C <sub>7</sub> )heptanoic acid	M4PFHpA	500	7	
Perfluoro-n-( <sup>13</sup> C <sub>8</sub> )octanoic acid	M8PFOA	500	10	
Perfluoro-n-( <sup>13</sup> C <sub>9</sub> )nonanoic acid	M9PFNA	250	11	
Perfluoro-n-(1,2,3,4,5,6- <sup>13</sup> C <sub>10</sub> )decanoic acid	M6PFDA	250	14	
Perfluoro-n-(1,2,3,4,5,6,7- <sup>13</sup> C <sub>11</sub> )undecanoic acid	M7PFUdA	250	18	
Perfluoro-n-(1,2- <sup>13</sup> C <sub>12</sub> )dodecanoic acid	MPFDoA	250	19	
Perfluoro-n-(1,2- <sup>13</sup> C <sub>14</sub> )tetradecanoic acid	M2PFTeDA	250	22	
Perfluoro-1-( <sup>13</sup> C <sub>8</sub> )octanesulfonamide	M8FOSA	500	17	
N-methyl-d <sub>3</sub> -perfluoro-1-octanesulfonamide	d-N-MeFOSA	500	21	
N-ethyl-d <sub>5</sub> -perfluoro-1-octanesulfonamide	d-N-EtFOSA	500	24	
N-methyl-d <sub>3</sub> -perfluoro-1-octanesulfonamidoacetic acid	d3-N-MeFOSAA	1000	15	
N-ethyl-d <sub>5</sub> -perfluoro-1-octanesulfonamidoacetic acid	d5-N-EtFOSAA	1000	16	
2-(N-methyl-d <sub>3</sub> -perfluoro-1-octanesulfonamido)ethan-d <sub>4</sub> -ol	d7-N-MeFOSE	5000	20	
2-(N-ethyl-d <sub>5</sub> -perfluoro-1-octanesulfonamido)ethan-d <sub>4</sub> -ol	d9-N-EtFOSE	5000	23	
2,3,3,3-Tetrafluoro-2-(1,1,2,2,3,3,3-heptafluoropropoxy)( <sup>13</sup> C <sub>3</sub> )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- <sup>13</sup> C <sub>3</sub> )butanesulfonate	M3PFBS	500	466	3
Sodium perfluoro-1-(1,2,3- <sup>13</sup> C <sub>3</sub> )hexanesulfonate	M3PFHxS	500	474	8
Sodium perfluoro-1-( <sup>13</sup> C <sub>8</sub> )octanesulfonate	M8PFOS	500	479	12
Sodium 1H,1H,2H,2H-perfluoro-(1,2- <sup>13</sup> C <sub>2</sub> )hexanesulfonate	M2-4:2FTS	1000	938	4
Sodium 1H,1H,2H,2H-perfluoro-(1,2- <sup>13</sup> C <sub>2</sub> )octanesulfonate	M2-6:2FTS	1000	951	9
Sodium 1H,1H,2H,2H-perfluoro-(1,2- <sup>13</sup> C <sub>2</sub> )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)