



## DATA VALIDATION REPORT

**Red Hill Bulk Fuel Storage Facility  
Joint Base Pearl Harbor-Hickam  
CV 23F0104**

**SDG: 22L0110  
APPL, INC.**

Prepared by  
**ENVIRONMENTAL DATA SERVICES, LTD.**

Prepared for  
**AECOM Environmental**

**Released: 12/23/22**

### Data Validators and Peer Reviewers:

A handwritten signature in black ink, appearing to read "Diane Waldschmidt".

Diane Waldschmidt

A handwritten signature in black ink, appearing to read "Gretchen Phipps".

Gretchen Phipps

A handwritten signature in black ink, appearing to read "Dina Manov".

Dina Manov

A handwritten signature in black ink, appearing to read "Larry Lewis".

Larry Lewis

A handwritten signature in black ink, appearing to read "Paloma Hoelzle".

Paloma Hoelzle

## EXECUTIVE NARRATIVE

**Sample Delivery Group:** 22L0110

**Laboratory:** APPL, Inc.

**Site:** Red Hill Bulk Storage Facility, CV 23F0104

**Sampling dates:** 12/14/2022

**Number of Samples:** 2

**Test Method:** USEPA Method 1633

**Analysis:** per- and polyfluoroalkyl substances (PFAS)

**Quality Assurance Project Plan:** Sampling and Analysis Plan, Investigation and Remediation of Releases and Groundwater Protection and Evaluation, Red Hill Bulk Fuel Storage Facility, Joint Base Pearl Harbor-Hickam, O’ahu, Hawai’i (Revision 01, April 2017); PFAS-Specific Sampling and Analysis plan, Red Hill Bulk Fuel Storage Facility, Adit 6, Joint Base Pearl Harbor-Hickam, O’Ahu, Hawai’i (November 30, 2022) (SAP).

**Validation Guidelines:** United States Department of Defense Data Validation Guidelines Module 6: Data Validation Procedure for Per- and Polyfluoroalkyl Substances analysis by QSM Table B-24, Environmental Data Quality Workgroup, October 18, 2022; United States Department of Defense (DOD) Environmental Data Quality Workgroup (EDQW), General Validation Guidelines, November 2019.

Client Sample Identification	Laboratory Sample Identification	Matrix	Validation Stage
AF-RHMW12A-WGN01B-2212W2	22L0110-01	water	S2BVEM
AF-RHMW06-WGN01B-2212W2	22L0110-02	water	S2BVEM

Table 1 provides a summary of the major and minor data quality issues identified in this data set. All data are acceptable except those results which have been qualified with “X”, rejected. Data validation qualifiers along with associated descriptions are provided in Table 2. All data qualification related to this group of samples is detailed on the attached sheets.

All data users should note two facts. First, an “X” flag means that the associated value is unusable due to significant quality control (QC) problems, the data is invalid and provides no information as to whether the compound is present or not. “X” values should not appear on any data tables even as a last resort. Second, no analyte concentration, even if it passed all QC tests, is guaranteed to be accurate. Strict QC serves to increase confidence in data, but any value potentially contains error.

## DATA ASSESSMENT

### 1. NARRATIVE AND COMPLETENESS REVIEW

*The case narrative was reviewed, and the data package was checked for completeness. No discrepancies were noted.*

### 2. SAMPLE DELIVERY AND CONDITION

*The samples arrived at the laboratory in acceptable condition. Proper custody was documented.*

### 3. HOLDING TIME

**The amount of an analyte in a sample can change with time due to chemical instability, degradation, volatilization, etc. If the specified holding time is exceeded, the data may not be valid. Proper sample handling and preservation also play a role in the chemical stability of analytes in the sample matrix. If samples are not collected and stored using proper containers and/or preservatives, data may not be valid.**

*No problems were found for this criterion.*

### 4. CALIBRATION

**Satisfactory instrument calibration is established to ensure that the instrument can produce acceptable quantitative data. An initial calibration demonstrates that the instrument can give acceptable performance at the beginning of an experimental sequence. The continuing calibration checks document that the instrument is giving satisfactory daily performance. Additionally, a continuing calibration is analyzed at the end of each 12-hour analytical sequence, denoted as a "closing" calibration verification and ascertains acceptable performance at the conclusion of the analytical sequence.**

#### A) Initial Calibration

**Percent relative standard deviation (%RSD) is calculated from the initial calibration and is used to indicate stability of a specific compound over the calibration range.**

**An RSD value outside the initial calibration limit indicates the potential for quantitation errors. For this reason, all positive and non-detected results are qualified as estimated. Severe performance failures (RSD >30%) requires rejection of all results. The following QC criteria have been applied for this project: The %RSD of initial calibration must be <20%.**

*No problems were found for this criterion.*

## B) Continuing Calibration

The Percent Recovery (%R) for all target analytes in the continuing calibration must be within 70-130%. All initial calibration verification (ICV) and continuing calibration verification (CCV) %Rs were with acceptance limits with the following exceptions.

*No problems were found for this criterion.*

## C) Instrument Sensitivity Check

Prior to analysis an instrument sensitivity check (ISC) must be performed. The ISC must be at the limit of quantitation (LOQ). All analyte concentrations must be within  $\pm 30\%$ . Note: the laboratory reports refer to the ISC as Low-Concentration Calibration Verification (LCCV). The validator has determined that the LCCV in the laboratory's report is equivalent to the method required ISC.

*No problems were found for this criterion with the following exceptions.*

*The observed recoveries for PFOA, PFUnA, and NMeFOSAA were higher than 130% for the ISC associated with all samples in this SDG. The results reported for the impacted analytes in the associated samples have been qualified estimated "J+" or "UJ" as appropriate on this basis.*

*The observed recovery for NFDHA was less than 50% for the ISC associated with all samples in this SDG. The non-detected results reported for the impacted analyte in the associated samples have been qualified "UJ" on this basis. It is the data validators recommendation that these results be considered estimated "UJ" when using data as the recovery was less than the lower acceptance limit but greater than 10% rather than applying an "X" qualifier as the validation module instructs.*

## 5. BLANK CONTAMINATION

Quality assurance (QA) blanks, i.e., method, field, or rinse blanks are prepared to identify any contamination which may have been introduced into the samples during sample preparation or field activity. Method blanks measure laboratory contamination. Field and rinse blanks measure cross-contamination of samples during field operations. When an equipment blank, or lab blank has an analyte detection, then all associated field samples are qualified per validation guidance as appropriate.

### A) Method blank contamination:

*No problems were found for this criterion with the following exception. PFOS was positively detected in the method blank associated with all samples in this SDG. Positive results reported for the impacted analyte in the associated samples have been evaluated and qualified per validation guidance as appropriate.*

### B) Instrument blank contamination:

*No problems were found for this criterion.*

**B) Field/Equipment blank contamination:**

*No samples were submitted as a field/equipment blank in association with this SDG.*

**6. EXTRACTED INTERNAL STANDARDS**

All samples are spiked with labeled standard compounds prior to sample preparation and analyses to evaluate overall laboratory performance and efficiency of the analytical technique. The reported project samples had observed surrogate recoveries within the established limits in all cases with the following exceptions.

*No problems requiring result qualification were found for this criterion.*

**7. NON-EXTRACTED INTERNAL STANDARDS**

Non-extracted internal standard peak areas are used to quantify extracted internal standard recoveries. The reported project samples had non-extracted internal standard area counts within the established limits in all cases with the following exceptions.

*No problems were found for this criterion.*

**8. COMPOUND IDENTIFICATION**

The project target analyte compounds are identified on the LC/MS/MS by using the analytes retention time (RT). The retention time of each target analyte should be within  $\pm 0.4$  minutes of the predicted retention. Target analyte detections should display a signal-to-noise of  $\geq 3:1$ , have proper peak integration, and display all ions at the correct retention times.

Target analyte detections should have passing ion ratios (50 - 150% of theoretical). Ion ratio failures could be caused by matrix interference and/or be the result of the presence of isomers in the sample at different ratios than the ratio of isomers present in the calibration standards.

*Target compound identification was verified. No anomalies were identified.*

**9. COMPOUND QUANTIFICATION**

*Target compound quantitation was not verified as part of the Level 2B data validation. No anomalies were identified.*

*Manual integrations were not reviewed at the Stage 2B level.*

**10. MATRIX SPIKE/MATRIX SPIKE DUPLICATE RECOVERY**

Matrix spike/matrix spike duplicate (MS/MSD) data are generated to determine the long-term precision and accuracy of the analytical method in various matrices. The MS/MSD data may be used in conjunction with other quality control criteria for additional qualification of data.

*No samples were submitted for MS/MSD and/or matrix duplicate evaluation in association with this SDG.*

**11. FIELD DUPLICATES/ TRIPLICATES**

Field duplicates may be taken and analyzed as an indication of overall precision. These analyses measure both field and laboratory precision. A control limit of  $\leq 30\%$  for the Relative Percent Difference (RPD) for water samples and  $\leq 50\%$  RPD for solid samples, shall be used when original and duplicate sample values are greater than or equal to the sample specific LOQ. For field duplicate analyses that do not meet the technical criteria, the action was applied to only the parent sample and its duplicate. A control limit of  $\leq 35\%$  RSD was applied for field triplicate samples when original and triplicate sample values are greater than the sample specific LOQ. For field triplicate analyses that do not meet the technical criteria, the action was applied to only the parent sample, duplicate and triplicate.

*No samples were submitted as a field duplicate/triplicate set in association with this SDG.*

**12. LABORATORY CONTROL SAMPLES**

The Laboratory Control Sample (LCS) serves as a monitor of the overall performance of each step during the analysis, including the sample preparation. The LCS results are used to verify that the laboratory can perform the analysis in a clean matrix. Note: in addition to the standard LCS the laboratory has also provided a second LCS referred to as the MRL check in the laboratory report. The validator has determined that the MRL check in the laboratory's report is equivalent to the required low level LCS.

*No problems were found for this criterion.*

**13. DILUTIONS, RE-EXTRACTIONS & REANALYSIS**

Samples may be re-analyzed for dilution, re-extraction and for other QC reasons. In such cases, the best result values are used.

*No samples were re-extracted and/or reanalyzed.*

**14. SYSTEM PERFORMANCE AND OVERALL ASSESSMENT**

*Overall, the laboratory data generated met the project goals and quality control criteria, with the exceptions identified in this report and as summarized in Table 1.*

**Table 1  
Review Elements Summary**

	Were acceptance criteria met?		
	Yes	No	
		Major	Minor
Per-fluorinated Compounds			
Holding Time/Sample Handling	x		
Method Blanks			x
Instrument Blanks	x		
Field Blanks	NA		
Calibration Percent Relative Standard Deviation and Percent Difference	x		
Instrument Sensitivity Check			x
Extracted Internal Standards	x		
Non-Extracted Internal Standards	x		
Compound Identification	x		
Matrix Spike/Matrix Spike Duplicate	NA		
Laboratory Control Sample	x		
Other Quality Control Data out of Specification	x		
Field Duplicate / Triplicate	NA		

Major= Major data quality issue identified resulting in rejection of data.

Minor= Minor data quality issue identified resulting in the qualification of data. Data qualification should be used to inform the data users of data limitations.

NA = Not applicable

**Table 2  
Data Validation Qualifiers**

<b>Data Qualifier</b>	<b>Definition</b>
<b>U</b>	The analyte was analyzed for but was not detected above the level of the reported sample quantitation limit.
<b>J</b>	The result is an estimated quantity. The associated numerical value is the approximate concentration of the analyte in the sample.
<b>J+</b>	The result is an estimated quantity, but the result may be biased high.
<b>J-</b>	The result is an estimated quantity, but the result may be biased low.
<b>UJ</b>	The analyte was analyzed for but was not detected. The reported quantitation limit is approximate and may be inaccurate or imprecise.
<b>X</b>	The sample results (including non-detects) were affected by serious deficiencies in the ability to analyze the sample and to meet published method and project quality control criteria. The presence or absence of the analyte cannot be substantiated by the data provided.
<b>R</b>	The data are unusable. The sample results are rejected due to serious deficiencies in meeting QC criteria. The analyte may or may not be present in the sample.



**Table 3  
PFAS Definitions Table**

<b>NO</b>	<b>CAS #</b>	<b>Target Name</b>	<b>Target Abbreviation</b>
1	763051-92-9	11-Chloroeicosafluoro-3-oxaundecane-1-sulfonic acid	11Cl-PF3OUdS
2	914637-49-3	2H,2H,3H,3H-Perfluorooctanoic acid	5:3FTCA
3	812-70-4	3-Perfluoroheptyl propanoic acid	7:3FTCA
4	356-02-5	3-Perfluoropropyl propanoic acid	3:3FTCA
5	919005-14-4	4,8-Dioxa-3H-perfluorononanoic acid	ADONA
6	757124-72-4	4:2 Fluorotelomer sulfonic acid	4:2 FTS
7	27619-97-2	6:2 Fluorotelomer sulfonic acid	6:2 FTS
8	39108-34-4	8:2 Fluorotelomer sulfonic acid	8:2 FTS
9	756426-58-1	9-Chlorohexadecafluoro-3-oxanone-1-sulfonic acid	9Cl-PF3ONS
10	13252-13-6	Hexafluoropropylene oxide dimer acid	HFPO-DA
11	4151-50-2	N-Ethyl perfluorooctanesulfonamide	NEtFOSA
12	2991-50-6	N-Ethyl perfluorooctanesulfonamidoacetic acid	NEtFOSAA
13	1691-99-2	N-Ethyl perfluorooctanesulfonamidoethanol	NEtFOSE
14	31506-32-8	N-Methyl heptadecafluorooctanesulfonamide	NMeFOSA
15	2355-31-9	N-Methyl perfluorooctanesulfonamidoacetic acid	NMeFOSAA
16	24448-09-7	N-Methyl perfluorooctanesulfonamidoethanol	NMeFOSE
17	151772-58-6	Nonafluoro-3,6-dioxaheptanoic acid	NFDHA
18	113507-82-7	Perfluoro(2-ethoxyethane)sulfonic acid	PFEESA
19	377-73-1	Perfluoro-3-methoxypropanoic acid	PFMPA
20	863090-89-5	Perfluoro-4-methoxybutanoic acid	PFMBA
21	375-73-5	Perfluorobutanesulfonic acid	PFBASA
22	375-22-4	Perfluorobutanoic acid	PFBA
23	335-77-3	Perfluorodecanesulfonic acid	PFDS
24	335-76-2	Perfluorodecanoic acid	PFDA
25	79780-39-5	Perfluorododecanesulfonic acid	PFDoS
26	307-55-1	Perfluorododecanoic acid	PFDoA
27	375-92-8	Perfluoroheptanesulfonic acid	PFHpS
28	375-85-9	Perfluoroheptanoic acid	PFHpA
29	355-46-4	Perfluorohexanesulfonic acid	PFHXSA
30	307-24-4	Perfluorohexanoic acid	PFHxA
31	68259-12-1	Perfluorononanesulfonic acid	PFNS
32	375-95-1	Perfluorononanoic acid	PFNA
33	754-91-6	Perfluorooctanesulfonamide	PFOSA
34	1763-23-1	Perfluorooctanesulfonic acid	PFOS
35	335-67-1	Perfluorooctanoic acid	PFOA
36	2706-91-4	Perfluoropentanesulfonic acid	PFPeS
37	2706-90-3	Perfluoropentanoic acid	PFPeA
38	376-06-7	Perfluorotetradecanoic acid	PFTeDA
39	72629-94-8	Perfluorotridecanoic acid	PFTTrDA
40	2058-94-8	Perfluoroundecanoic acid	PFUnA

<b>Data Qualification Reason Codes</b>	
<b>Reason Code</b>	<b>Reason Code Description</b>
<b>A</b>	<b>Serial dilution</b>
<b>A1</b>	<b>Ambient Blank</b>
<b>B</b>	<b>The analyte was found in an associated blank as well as in the sample.</b>
<b>B2</b>	<b>CCB</b>
<b>B3</b>	<b>CCB - Neg</b>
<b>B4</b>	<b>Grinding Blank</b>
<b>C</b>	<b>LCS Recovery</b>
<b>C1</b>	<b>Reference Recovery</b>
<b>C2</b>	<b>Reference Recovery RPD</b>
<b>D</b>	<b>MS RPD</b>
<b>D1</b>	<b>Lab Replicate RPD</b>
<b>D2</b>	<b>No precision available</b>
<b>D3</b>	<b>Field Duplicate RPD</b>
<b>D4</b>	<b>Field Triplicate RSD</b>
<b>D5</b>	<b>Laboratory Triplicate RSD</b>
<b>F</b>	<b>Field Blank</b>
<b>F1</b>	<b>Hydrocarbon pattern does not match standard</b>
<b>G1</b>	<b>Initial Calibration RRF</b>
<b>G2</b>	<b>Initial Calibration RSD/r<sup>2</sup>/r</b>
<b>G3</b>	<b>ICV RRF</b>
<b>H1</b>	<b>Test Hold Time</b>
<b>H2</b>	<b>Prep Hold Time</b>
<b>I</b>	<b>Surrogate recovery outside project limits.</b>
<b>J</b>	<b>CRA/CRI Recovery</b>
<b>K</b>	<b>An analyte (non-common laboratory artifact) was detected in the sample at a concentration less than 5X the concentration detected in the associated method blank.</b>
<b>L</b>	<b>Lab Blank</b>
<b>L1</b>	<b>Lab Blank - Neg</b>
<b>M</b>	<b>MS Recovery</b>
<b>M2</b>	<b>Post Spike</b>
<b>N</b>	<b>Blank - No Action</b>
<b>O</b>	<b>ICS</b>
<b>P</b>	<b>Sample preservation/collection requirement not met.</b>
<b>P1</b>	<b>Column RPD</b>
<b>P2</b>	<b>Improper preparation/extraction</b>
<b>Q</b>	<b>Encore sample holding time exceeded by more than 2X.</b>
<b>Q1</b>	<b>Material Blank</b>

Q2	Encore sample holding time exceeded by less than 2X.
R	Exceeds Linear Calibration Range
S	Internal standard
T	Trip Blank
TI	Tentatively Identified Compound
TR	Trace Level Detect
U	Receipt Temperature
V	Equipment Blank
V1	ICV
V2	CCV
V3	CCV RRF
V4	Sample Receipt Condition
V5	Ending Continuing Calibration Verification
V6	Low Level Calibration Verification
V7	Interference Check Sample A
V8	Interference Check Sample AB
V9	Interference Check Sample A - Negative
W	Column breakdown (pesticides/8270)
X	Raised reporting limit
Y	Cooler temperature greater than 10 degreeec C.
Y1	False Positive
Y2	Data rejected due to radiological anomolies
Y3	Non-accredited analyte/compound. Accreditation not offered at time of analyses for the analyte/compound by the stated method and matrix.
Y4	Performance Check - Degradation of DDT
Y5	Extracted Internal Standard
Y6	Analyte not confirmed on second column.
Y7	Signal to Noise Ratio not met
Z	LCS RPD
Z1	Non-accredited analyte/compound
Z1	Data rejected, more valid data available.
Z2	Detection Level not met uncertainty greater than DL
Z4	MDA Greater than RDL.
Z5	Ion Ratio
Z6	Samples were analyzed past the 12 hour time period from the Tune or opening CCV.

## **Data Validation Worksheet**

# DATA VALIDATION PFAS

## Module 6; PFAS by QSM Table 5-24; October 18, 2022

Validator: DM

Reviewer: DLW

Date Validated: 12/22/22

Reviewed: 12/23/22

Project: Red Hill

SDG: 22L0110

LAB: APPL

Samples Collected: 12/14/2022

2 GW Samples

### SAMPLE RECEIPT AND CASE NARRATIVE REVIEW

- ✓ Traffic reports, chain-of-custody forms or SDG narrative do not indicate any problems with sample receipt, condition of the samples, analytical problems or special circumstances affecting the quality of the data.
- ✓ AFFF samples are to be shipped in HDPE containers with an unlined cap
- ✓ Shipment temp 0-6°C: recommended to freeze tissue samples upon receipt
- ✓ If temp upon receipt is greater than 6°C J/UJ all

Received on 12/16 all cooler <6C

### HOLDING TIMES

- ✓ Recommended storage temp is  $\leq -20^{\circ}\text{C}$
- ✓ Per method 1633: aqueous samples may be held in the lab for up to 90 days when stored at recommended temp and protected from light; when stored at 0-6 °C and protected from light samples can be held for up to 28 days (see method for additional details)
- ✓ Per method 1633: solid samples may be held in the lab for up to 90 days when stored at recommended temp or 0-6 °C (see method for additional details)
- ✓ Per method 1633: biosolid samples may be held in the lab for up to 90 days when stored at recommended temp or 0-6 °C; however, freezing is recommended (see method for additional details)
- ✓ Samples extracts should be stored at 0-4°C protected from light and analyzed within 90 days

- ✓ If hold time is exceeded qualify J/UJ
- ✓ If hold time is grossly exceeded (2X hold time) J/X

244 **Table II. Sample Storage and Holding Time Requirements**

Matrix Type	Stored at 0 - 6°C, protected from light		Stored at ≤ -20°C, protected from light	
	Holding Time	Caveat	Holding Time	Caveat
<b>Aqueous</b>	28 days	Precursor degradation occurs after 7 days	90 days	None
<b>Solid and Tissue</b>	90 days	Should be prepared as soon as possible if NFDHA is a target analyte	90 days	Should be prepared as soon as possible if NFDHA is a target analyte
<b>Biosolid</b>	90 days	Not recommended due to the production of gases due to microbiological activity	90 days	None

all inside holding time

#### Extracted Internal STANDARDS

- ✓ Added to all QC and field samples
- ✓ Recoveries are within the limits as defined in QAPP; otherwise QSM criteria (20-150%) should be used
- ✓ Detected for analytes qualified using an EIS percent recovery >200% should be qualified J-. Non-detects should not be qualified.
- ✓ If EIS recovery is <10%; associated detected and non-detects should be qualified X
- ✓ EIS retention times should be within 0.4 minutes of standard; use professional judgment to qualify

**For Red Hill project (see Kristin's email on file in project folder 12/14/22 at 3:25pm)**

**For EIS %Rs >150% J- positive results, no action on non-detects**

**For EIS %Rs between lab limit of 20-150%; no action**

**For EIS %Rs <20% but >10%; J+ positive results, UJ non-detects**

**For EIS %Rs <10% X positive and non-detected (and recommend R of non-detected, J+ of positive results)**

**All ok**

#### **Non-Extracted Internal STANDARDS**

- ✓ Used to quantify EIS
- ✓ If low are counts are reported (<30%) detected and non-detected should be qualified X

#### **Laboratory Control Sample (LCS) and Low-Level Laboratory Control Sample (LLLCS)**

**(MRL in APPL data package)**

- ✓ LCMS Lab Control Recovery (Form III), Form I, prep log, run log
- ✓ LCS prepared, extracted, analyzed, and reported once for every 20 field samples of a similar matrix, per SDG.
- ✓ Laboratory Control Samples were analyzed for all the target analytes that the samples are analyzed for.
- ✓ Use limits as defined in QAPP; otherwise lab limits or QSM criteria of 40-150%.
- ✓ If LCS or LLLCS %R is > upper limit; qualify detects J+; no action on non-detected
- ✓ If LCS or LLLCS %R is < lower limit; qualify detected J- and non-detected X

[Use lab limits \(40-150\) to evaluate  
All 40 compounds included.](#)

**LCS (BBL0322-BS1) all ok**

**MRL Check (BBL0322-MRL1) all ok (QC)**

## MS/MSD and Matrix Duplicate

- ✓ LCMS Matrix Spike Recovery (Form III)
- ✓ The Matrix Spike Samples were spiked and analyzed for all the target analytes that the samples are analyzed for (Same analytes as LCS).
- ✓ Per module 6: MS and MSD are applicable where the spike concentration is at least 3 times greater than the native analyte concentration (**3X rule**)
- ✓ Use limits as defined in QAPP; otherwise lab limits or QSM criteria of 40-150%.
- ✓ If MS or MSD %R is > upper limit; qualify detected J+; no action on non-detected
- ✓ If MS or MSD %R is < lower limit but >10%; qualify detected J- and non-detected UJ
- ✓ If MS or MSD %R is < 10%; qualify detected J- and non-detected X
- ✓ If MS/MSD RPD is out; qualify detected J and non-detected UJ
- ✓ For matrix duplicate; for concentrations of analytes that are equal to or greater than the LOQ, the RPD must be ≤30%; if out qualified detected J; no action on non-detects

Use lab limits to evaluate

Sample: **None**

## BLANKS

- ✓ LCMS Method Blank Summary (Form IV), method blank Form I, prep log, run log
- ✓ Frequency of Analysis: method blank has been analyzed for every 20 (or less) samples of similar matrix or concentration or each extraction batch.
- ✓ Continuing Calibration Blanks (Form I) and run log
- ✓ Frequency of Analysis: immediately following the highest standard analyzed and daily prior to sample analysis.
- ✓ Field/rinse blanks are non-detected for all analytes

If an analyte is detected in the field blank (at any concentration) and in the associated samples, the action taken depends on both the blank and sample concentrations (Table III).



312 **Table III: Sample Qualification in the Presence of Blank Contamination**

	Sample		
Row Number	Result	Validated Result	Validation Qualifier
1	Non-detect or detect $\leq$ LOD	Report at LOD	U
2	> LOQ but $\leq$ 5x blank	Report at Sample Result	J+
3	> LOQ and > 5x blank	Report at Sample Result	None

313 LOD = Limit of Detection

Multiple blank contaminations (such as a batch with field blanks and a method blank) does not establish a 'hierarchy' of one blank over another. Each blank must be evaluated 695 individually. Blanks should not be qualified due to the results of other blanks.

Method blank:

BBL0322-BLK1  
PFOS 0.111J,

SAMPLE 22L0110-02 > LOD FLAG J+  
SAMPLE 22L0110-01 ND no Q

ICBs/CCBs see below

### MASS CALIBRATION

- ✓ Verified to be  $\pm 0.2$  amu of true value

### Bile Salt Interference Check and Qualitative Identification Standard

- ✓ Provided and requirements met
- ✓ See Module 6

acceptable

### ICAL

- ✓ Initial Calibration Data Curve Evaluation (Form VI) and run log
- ✓ Lowest standard should be at or below LOQ
- ✓ %RSD <20% or relative standard error (RSE) <20%
- ✓ If %RSD > 20% but <30% J/UJ
- ✓ If %RSD >30% J/R

See below

### **INSTRUMENT PERFORMANCE CHECK PER DRAFT METHOD 1633 (LCV in APPL data package)**

- ✓ Concentration equal to LOQ
- ✓ Analyzed after ICAL and daily before samples
- ✓ If not analyzed all associated data should be qualified X
- ✓ The %R for ICV and CCV 30%; if out >130% qualify positive J+ and nondetected UJ; if out <70% qualify positives J- and nondetects UJ
- ✓ Per module if gross exceedances of recoveries <50% or >150%; qualify all associate data X

### **CCAL**

- ✓ Continuing Calibration Data (Form VII) and run log
- ✓ Continuing calibration standard analyzed on each working day, prior to sample analyses.
- ✓ Calibration verification/continuing calibration standard been analyzed after every 10 samples and at the end of each analytical sequence
- ✓ If not analyzed all associated data should be qualified X
- ✓ The %R for ICV and CCV +/- 30%; if out >130% qualify positive J+ and nondetected UJ; if out <70% qualify positives J- and nondetects UJ
- ✓ Per module if gross exceedances of recoveries <50% or >150%; qualify all associate data X

LCV is the method required ISC

Instrument Saphira

12/13/22 All %RSE <20%

Initial Cal Blank SB03856-ICB1 S2022-12-15A (9) 12/15/22 14:15 all ND

Secondary Cal Check SB03856-SCV1 S2022-12-15A (10) 12/15/22 14:28 all ok

Calibration Blank SB03886-CCB1 S2022-12-19B (1) 12/19/22 15:06 all ND

Low Cal Check SB03886-LCV1 S2022-12-19B (2) 12/19/22 15:19

PFOA 135%R FLAG J

PFOA 131.2 %R FLAG UJ

NMeFOSAA 187.7 %R FLAG UJ

NFDHA 11.8 %R grossly exceeded recommended to qualify UJ since above 10% per module X

Calibration Check SB03886-CCV1 S2022-12-19B (3) 12/19/22 15:32 all ok

Calibration Blank SB03886-CCB2 S2022-12-19B (4) 12/19/22 16:10 all ND

Calibration Check SB03886-CCV2 S2022-12-19B (16) 12/19/22 18:42 all ok

Calibration Blank SB03886-CCB3 S2022-12-19B (17) 12/19/22 18:55 all ND

**COMPOUND IDENTIFICATION**

- ✓ RT within  $\pm 0.4$  RRT units (review for Level 4)
- ✓ S/N ration 3:1 (review for Level 4)
- ✓ Ion response ratio with  $\pm 50\%$  (review for Level 2B)
- ✓ If ion ratio is outside limit; qualify J

Use J flag for module 6

Reason Code: Z5

Ion ratio:

AF-RHMW12A-WGN01B-2212W2  
PFOS out ND no Q

**FIELD DUPLICATES**

- ✓ Use QAPP defined criteria
- ✓ If outside acceptance criteria qualify J/UJ (MODULE FLAGS NONDETECTS TOO)

**For field triplicates use 35% RSD per Kristin's email on file from 12/14/22**

**NONE**

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The following analytes were >35%

SEE FIELD DUPLICATE WORKSHEET

**Automated Data Review Detail Report for 22L0110**  
 RH Fire Suppression System  
 RHS PFAS UFP-QAPP

Sample Summary								E1633DR	
Location	Field Sample ID	Date	Time	Sample Type	Matrix	SBD	SED		
RHMW12A	AF-RHMW12A-WGN01B-2212W2	12-14-2022	1250	N	WG	0.00	0.00		X
RHMW06	AF-RHMW06-WGN01B-2212W2	12-14-2022	1725	N	WG	0.00	0.00		X
<b>Total</b>								<b>2</b>	

**Automated Data Review Detail Report for 22L0110**  
 RH Fire Suppression System  
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Batch Report

Test Method: E1633DR		Analysis Batch: SB03886								
Location	Matrix	Field Sample ID	Lab Sample ID	Calibration Ref	Run#/Dil'n	Collection Date/Time	Extraction Date/Time	Analysis Date/Time	Prep/Leach Batch	Sample Type
LABQC	WQ	LABQC	BBL0322-BLK1	2251019	1/1	12/15/2022 08:32	12/15/2022 08:32	12/19/2022 16:23	BBL0322/	LB
LABQC	WQ	LABQC	BBL0322-BS1	2251019	1/1	12/15/2022 08:32	12/15/2022 08:32	12/19/2022 16:35	BBL0322/	BS
RHMW12A	WG	AF-RHMW12A-WGN01B-2212W2	22L0110-01	2251019	1/1	12/14/2022 12:50	12/15/2022 08:32	12/19/2022 17:52	BBL0322/	N
RHMW06	WG	AF-RHMW06-WGN01B-2212W2	22L0110-02	2251019	1/1	12/14/2022 17:25	12/15/2022 08:32	12/19/2022 18:17	BBL0322/	N

**Automated Data Review Detail Report for 22L0110**  
RH Fire Suppression System  
RHS PFAS UFP-QAPP

Field Batch Report

--No Records Found--

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MS Mismatch Report

--No Records Found--

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Section to identify Matrix Spike mismatches where parent sample differs from MS by dilution.

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**Automated Data Review Detail Report for 22L0110**  
 RH Fire Suppression System  
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QC Outlier Report

Test Method: E1633DR		Extraction Method: METHOD		Leach Method: NONE							
QC Element	Sample ID/ Lab Sample ID	Run#/ Dil'n	Analyte	Result (Units)	Qualifier	Warning Limits	Control Limits	Reason	Comment	Rule	Action Level
Lab Blank	BBL0322-BLK1 (LB) / BBL0322-BLK1	1 / 1.00	Perfluorooctanesulfonic acid (PFOS)	0.1110 (ng/l)	U/None*	< 0.064	< 0.4	L		5	0.555

\*Blank flags displayed in the above table identify qualification of the sample result when it is less than or equal to the LOQ/RL. Sample results above the LOQ will be qualified based on the validation type such as J+ at the sample result.

Rule is the multiplier used when blank contamination occurs to determine action level.

\*Blank qualification may apply a J flag when the sample result is greater than the reporting limit for NFG and when the sample result is between 5 and 20 times the blank result for EM200-1-10.

**Automated Data Review Detail Report for 22L0110**  
 RH Fire Suppression System  
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Qualified Results

Test Method: E1633DR		Extraction Method: METHOD		Leach Method: NONE						
FieldSample ID	LabSample ID	Matrix	Type	Analyte	LOQ	Lab Result	Qualified Result	Bias	Units	Reason
AF-RHMW06-WGN01B-2212W2	22L0110-02	W	N	N-Methyl perfluorooctanesulfonamidoacetic acid (NMeFOSAA)	0.380	0.190 U	0.190 UJ		ng/l	V6
AF-RHMW06-WGN01B-2212W2	22L0110-02	W	N	Nonafluoro-3,6-dioxahheptanoic acid (NFDHA)	0.760	0.380 U	0.380 UJ		ng/l	V6
AF-RHMW06-WGN01B-2212W2	22L0110-02	W	N	Perfluorobutanoic acid (PFBA)	1.50	1.00 FJ	1.00 J		ng/l	TR
AF-RHMW06-WGN01B-2212W2	22L0110-02	W	N	Perfluorodecanoic acid (PFDA)	0.380	0.170 FJ	0.170 J		ng/l	TR
AF-RHMW06-WGN01B-2212W2	22L0110-02	W	N	Perfluorononanoic acid (PFNA)	0.380	0.230 FJ	0.230 J		ng/l	TR
AF-RHMW06-WGN01B-2212W2	22L0110-02	W	N	Perfluorooctanesulfonic acid (PFOS)	0.380	0.200 FJ	0.200 J	+	ng/l	L/TR
AF-RHMW06-WGN01B-2212W2	22L0110-02	W	N	Perfluorooctanoic acid (PFOA)	0.380	0.410 MI	0.410 J	+	ng/l	V6
AF-RHMW06-WGN01B-2212W2	22L0110-02	W	N	Perfluoroundecanoic acid (PFUnA)	0.380	0.420	0.420 J	+	ng/l	V6
AF-RHMW12A-WGN01B-2212W2	22L0110-01	W	N	N-Methyl perfluorooctanesulfonamidoacetic acid (NMeFOSAA)	0.370	0.190 U	0.190 UJ		ng/l	V6
AF-RHMW12A-WGN01B-2212W2	22L0110-01	W	N	Nonafluoro-3,6-dioxahheptanoic acid (NFDHA)	0.740	0.370 U	0.370 UJ		ng/l	V6
AF-RHMW12A-WGN01B-2212W2	22L0110-01	W	N	Perfluorobutanesulfonic acid	0.370	0.0510 FJ	0.0510 J		ng/l	TR
AF-RHMW12A-WGN01B-2212W2	22L0110-01	W	N	Perfluorooctanoic acid (PFOA)	0.370	0.350 FMI J	0.350 J	+	ng/l	TR/V6
AF-RHMW12A-WGN01B-2212W2	22L0110-01	W	N	Perfluoroundecanoic acid (PFUnA)	0.370	0.190 U	0.190 UJ		ng/l	V6

Qualified analytes in samples are reported as estimated, not detected (UJ) at the Limit of Detection (LOD).



**Automated Data Review Detail Report for 22L0110**  
 RH Fire Suppression System  
 RHS PFAS UFP-QAPP

Detected Results

Test Method: E1633DR		Extraction Method: METHOD			Leach Method: NONE					
FieldSample ID	LabSample ID	Matrix	Type	Dilution	Analyte	LOQ	Lab Result	Qualified Result	Units	Reason
AF-RHMW06-WGN01B-2212W2	22L0110-02	W	N	1	6:2 Fluorotelomer sulfonic acid (6:2 FTS)	1.50	2.10	2.10	ng/l	
AF-RHMW06-WGN01B-2212W2	22L0110-02	W	N	1	Perfluorobutanoic acid (PFBA)	1.50	1.00 FJ	1.00 J	ng/l	TR
AF-RHMW06-WGN01B-2212W2	22L0110-02	W	N	1	Perfluorodecanoic acid (PFDA)	0.380	0.170 FJ	0.170 J	ng/l	TR
AF-RHMW06-WGN01B-2212W2	22L0110-02	W	N	1	Perfluoroheptanoic acid (PFHpA)	0.380	4.90	4.90	ng/l	
AF-RHMW06-WGN01B-2212W2	22L0110-02	W	N	1	Perfluorohexanoic acid (PFHxA)	0.380	0.730	0.730	ng/l	
AF-RHMW06-WGN01B-2212W2	22L0110-02	W	N	1	Perfluorononanoic acid (PFNA)	0.380	0.230 FJ	0.230 J	ng/l	TR
AF-RHMW06-WGN01B-2212W2	22L0110-02	W	N	1	Perfluorooctanesulfonic acid (PFOS)	0.380	0.200 FJ	0.200 J	ng/l	L/TR
AF-RHMW06-WGN01B-2212W2	22L0110-02	W	N	1	Perfluorooctanoic acid (PFOA)	0.380	0.410 MI	0.410 J	ng/l	V6
AF-RHMW06-WGN01B-2212W2	22L0110-02	W	N	1	Perfluoropentanoic acid (PFPeA)	0.760	2.00	2.00	ng/l	
AF-RHMW06-WGN01B-2212W2	22L0110-02	W	N	1	Perfluoroundecanoic acid (PFUnA)	0.380	0.420	0.420 J	ng/l	V6
AF-RHMW12A-WGN01B-2212W2	22L0110-01	W	N	1	6:2 Fluorotelomer sulfonic acid (6:2 FTS)	1.50	18.0	18.0	ng/l	
AF-RHMW12A-WGN01B-2212W2	22L0110-01	W	N	1	Perfluorobutanesulfonic acid	0.370	0.0510 FJ	0.0510 J	ng/l	TR
AF-RHMW12A-WGN01B-2212W2	22L0110-01	W	N	1	Perfluorobutanoic acid (PFBA)	1.50	13.0	13.0	ng/l	
AF-RHMW12A-WGN01B-2212W2	22L0110-01	W	N	1	Perfluoroheptanoic acid (PFHpA)	0.370	1.20	1.20	ng/l	
AF-RHMW12A-WGN01B-2212W2	22L0110-01	W	N	1	Perfluorohexanoic acid (PFHxA)	0.370	5.50	5.50	ng/l	
AF-RHMW12A-WGN01B-2212W2	22L0110-01	W	N	1	Perfluorooctanoic acid (PFOA)	0.370	0.350 FMI J	0.350 J	ng/l	TR/V6
AF-RHMW12A-WGN01B-2212W2	22L0110-01	W	N	1	Perfluoropentanoic acid (PFPeA)	0.740	22.0	22.0	ng/l	

**Automated Data Review Detail Report for 22L0110**  
RH Fire Suppression System  
RHS PFAS UFP-QAPP

Rejected Results

--No Records Found--

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Anomalies Count

--No Records Found--

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Reporting Anomalies

--No Records Found--

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