

DATA VALIDATION REPORT

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

> SDG: 580-133679-1 Eurofins Savannah

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

Prepared for AECOM Environmental

Released: 12/04/23

Data Validators and Peer Reviewers:

Diane Waldschmidt

Gretchen Phipps

Dina Manov

Larry Lewis

Paloma Hoelzle



EXECUTIVE NARRATIVE

Sample Delivery Groups: 580-133679-1

Laboratory: Eurofins, Savannah

Site: Red Hill Bulk Storage Facility, CV 23F0104

Sampling dates: 11/06/2023 Number of Samples: 7 Test Method: SW-846 8015C

Analysis: 2-(2-Butoxyethoxy) ethanol

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 4: Data Validation Procedure for Organic Analysis by GC, Environmental Data Quality Workgroup, March 9, 2021; United States Department of Defense (DOD) General Data Validation Guidelines Environmental Data Quality Workgroup (EDQW), November 2019. United States Department of Defense Data Validation Guidelines Modules 1, 2, 3, and 4 Revised Table for Sample Qualification in the Presence of Blank Contamination, February 09, 2022.

Client Sample Identification	Laboratory Sample Identification	Matrix	Validation Stage
AF-RHMW16-WGN01LF-2311	580-133679-1	water	S2BVEM
AF-RHMW06-WGN01LF-2311	580-133679-2	water	S2BVEM
AF-RHMW04-WGN01LF-2311	580-133679-3	water	S2BVEM
AF-RHMW225401-WGN01B-2311	580-133679-4	water	S2BVEM
AF-HDMW225303-WGN01LF-2311	580-133679-5	water	S2BVEM
AF-RHMW02-WGN01LF-2311	580-133679-6	water	S2BVEM
AF-RHMW02-WGFD01LF-2311	580-133679-7	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. Those analytes detected in the samples whose holding time has been exceeded will be qualified as estimated, "J". The non-detect results will be flagged as not detected at an estimated quantitation limit, "UJ", unless the holding time is grossly exceeded (by more than two times the holding time specified), in which case non-detect results are flagged "X", rejected. Qualifications were applied to the samples and analytes as shown below.

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.

Percent Relative Standard Deviation and Percent Difference

Percent relative standard deviation (%RSD) is calculated from the initial calibration and is used to indicate the stability of the specific compound response factor over increasing concentration. Percent difference (%D) compares the response factor of the continuing calibration check to the mean response factor (RRF) from the initial calibration. Percent difference is a measure of the instrument's daily performance. If the %D exceeds 20% for any analyte, qualify all associated positive results "J" and non-detects "UJ". If %RSD and %D grossly exceed QC criteria, non-detect data may be qualified "X".

A multi-point initial calibration curve was used for the target analyte. The initial calibration demonstrated instrumental linearity. The %RSD was within validation guidelines.

Continuing calibrations verifications (CCVs) were analyzed at the proper frequencies and the observed %D values met quality control criteria with the following exception. One %D value for 2-(2-butoxyethoxy) ethanol was outside acceptance limits for a CCV associated with all samples in this sample delivery group (SDG). The results reported for the impacted analyte in the associated samples have been qualified "UJ" on this basis.

5. BLANK CONTAMINATION

Quality assurance (QA) blanks, i.e., method, trip, 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, equipment and rinse blanks measure cross-contamination of samples during field operations.

A) Method blank contamination

No problems were found for this criterion.

B) Field/Equipment blank contamination

No samples were submitted as field / equipment blanks in association with this SDG.

6. SURROGATES / SYSTEM MONITORING COMPOUNDS

All samples are spiked with surrogate/system monitoring compounds (SMC) prior to sample preparation to evaluate overall laboratory performance and efficiency of the analytical technique. If the measured surrogate/SMC concentrations were outside contract specifications, qualifications were applied to the samples and analytes as shown below.

No surrogates were used for this analysis.

7. COMPOUND IDENTIFICATION

The retention times (RTs) of reported compounds must fall within the calculated retention time windows for chromatographic column.

Retention Time

Target compound identifications were not reviewed at the Stage 2B level.

Relative Percent Difference

No positive results were reported for the samples in this SDG.

8. COMPOUND QUANTIFICATION

Analyte quantitation was not reviewed at the Stage 2B level.

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

9. MATRIX SPIKE / MATRIX SPIKE RECOVERY

Matrix spike / matrix spike duplicate (MS/MSD) data is 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.

Sample AF-RHMW16-WGN01LF-2311 was submitted for MS/MSD pair evaluation in association with this SDG. Upon evaluation all precision and accuracy indicators were favorable.

10. 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. Aqueous/water, soil/sediment, wipe, and filter LCSs shall be analyzed for each analyte utilizing the same sample preparations, analytical methods, and quality assurance/quality control (QA/QC) procedures as employed for the samples. All LCS percent recoveries must fall within the control limits. Qualifications were applied to the samples and analytes as shown below.

No problems were found for this criterion.

11. INTERNAL STANDARDS PERFORMANCE:

Internal standard performance criteria are meant to ensure that the gas chromatograph (GC) sensitivity and response are stable during every experimental run.

The internal standard area count must not vary by more than a factor of two from the associated continuing calibration standard. The retention time of the internal standard must not vary by more than ± 10 seconds from the associated continuing calibration standard. The area count must be within a (50-200%) range of the associated standard. If the area count is greater than 200%, non-detected results are not qualified and positive results are flagged as estimated with potential negative bias, "J-". If the area count is less than 50%, positive results are flagged as estimated with potential positive bias, "J+", and non-detected results are flagged "UJ". If the area count is less than 20%, positive results and non-detected results will be classified as unusable "X". Qualifications were applied to the samples and analytes as shown below.

No problems were found for this criterion.

12. FIELD DUPLICATE

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 30% for the Relative Percent Difference (RPD) shall be used for original and duplicate sample values greater than the LOQ. A control limit of a difference between results no more than the LOQ shall be used if either the sample or duplicate value is less than the LOQ. For field duplicate analyses that do not meet the technical criteria, the action was applied to only the parent sample and its duplicate.

Samples AF-RHMW02-WGN01LF-2311 and AF-RHMW02-WGFD01LF-2311 were submitted as a field duplicate pair in association with this SDG. Adequate field precision was demonstrated.

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 dilutions, re-extractions, or other re-analyses were performed. The level of quantitation (LOQ) specified in the SAP for the analytes reported has been achieved.

14. OTHER PROBLEMS

None.

Table 1 Major and Minor Findings

	Were acceptance criteria		
	Yes	N	0
2-(2-Butoxyethoxy)ethanol		Major	Minor
Holding Time	Х		
Calibration			Х
Method Blank	Х		
Field/Equipment Blank	NA		
Surrogates/System Monitoring Compounds	NA		
Compound Identification	NA		
Compound Quantitation	NA		
Matrix Spike/Matrix Spike Duplicate	Х		
Internal Standards	Х		
Field Duplicate	Х		
Laboratory Control Samples	Х		
Other Quality Control Data out of Specification	Х		

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.

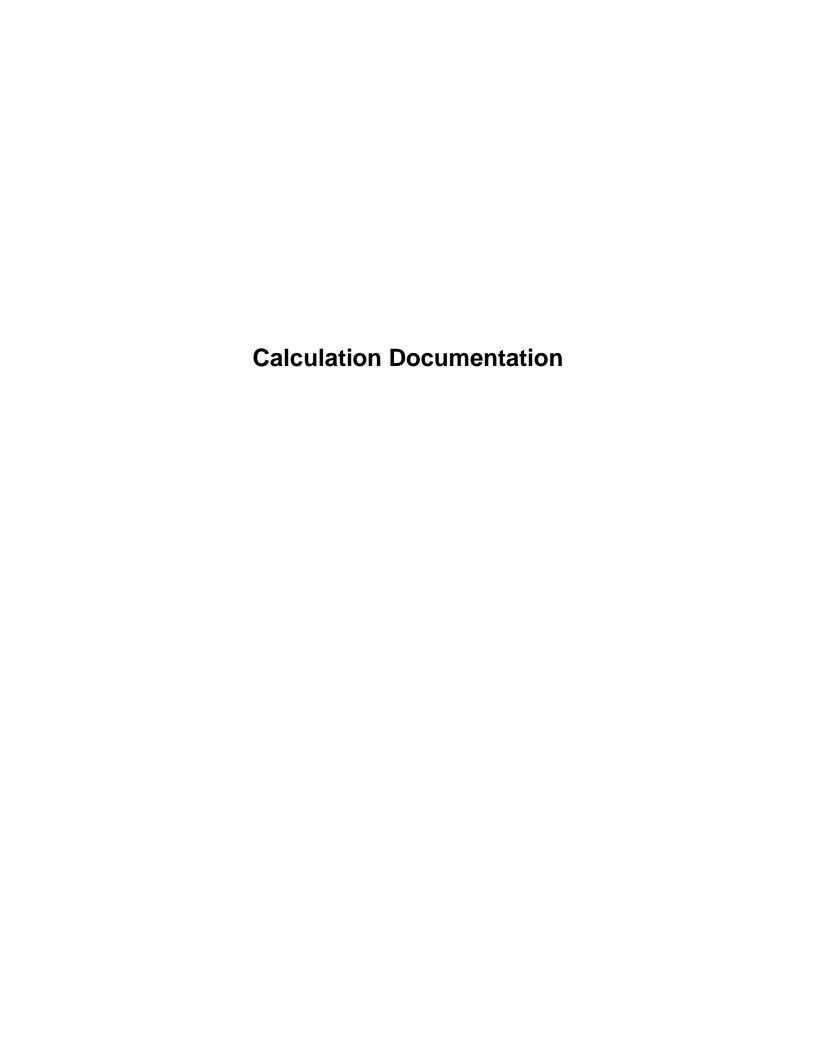
NA = Not applicable

Table 2 Data Validation Qualifiers

Data Qualifier	Definition
U	The analyte was analyzed for but was not detected above the level of the reported sample quantitation limit.
J	The result is an estimated quantity. The associated numerical value is the approximate concentration of the analyte in the sample.
J+	The result is an estimated quantity, but the result may be biased high.
J-	The result is an estimated quantity, but the result may be biased low.
ΩJ	The analyte was analyzed for but was not detected. The reported quantitation limit is approximate and may be inaccurate or imprecise.
X	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.
N	The analysis indicates the presence of an analyte for which there is presumptive evidence to make a tentative identification.
NJ	The analyte was tentatively identified, and the associated numerical value represents its approximate concentration.

Reason Code Reason Code Description A Serial dilution A1 Ambient Blank B The analyte was found in an associated blank as well as in the sample. B2 CCB B3 CCB - Neg B4 Grinding Blank C LCS Recovery C1 Reference Recovery RPD D MS RPD D1 Lab Replicate RPD D2 No precision available D3 Field Duplicate RPD D4 Field Triplicate RSD D5 Laboratory Triplicate RSD F Field Blank F1 Hydrocarbon pattern does not match standard G1 Initial Calibration RRF G2 Initial Calibration RSD/r^2/r G3 ICV RRF H1 Test Hold Time H2 Prep Hold Time I Surrogate recovery outside project limits. J CRA/CRI Recovery K 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. L Lab Blank L1 Lab Blank L1 Lab Blank L1 Lab Blank Neg M MS Recovery M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	Data Qualificati	ion Reason Codes
A Serial dilution At Ambient Blank B The analyte was found in an associated blank as well as in the sample. B2 CCB B3 CCB - Neg B4 Grinding Blank C LCS Recovery C1 Reference Recovery C2 Reference Recovery RPD D MS RPD D1 Lab Replicate RPD D2 No precision available D3 Field Duplicate RSD D5 Laboratory Triplicate RSD D5 Laboratory Triplicate RSD F Field Blank F1 Hydrocarbon pattern does not match standard G1 Initial Calibration RRF G2 Initial Calibration RSD/r^2/r G3 ICV RRF H1 Test Hold Time H2 Prep Hold Time I Surrogate recovery outside project limits. J CRA/CRI Recovery K An analyte (non-common laboratory artifact) was detected in the associated method blank. L Lab Blank L1 Lab Blank L1 Lab Blank Neg M MS Recovery M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.		
A1 Ambient Blank B The analyte was found in an associated blank as well as in the sample. B2 CCB B3 CCB - Neg B4 Grinding Blank C LCS Recovery C1 Reference Recovery C2 Reference Recovery RPD D MS RPD D1 Lab Replicate RPD D2 No precision available D3 Field Duplicate RPD D4 Field Triplicate RSD D5 Laboratory Triplicate RSD F Field Blank F1 Hydrocarbon pattern does not match standard G1 Initial Calibration RRF G2 Initial Calibration RSD/r^2/r G3 ICV RRF H1 Test Hold Time I Surrogate recovery outside project limits. J CRA/CRI Recovery K 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. L Lab Blank L1 Lab Blank L1 Lab Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.		-
B The analyte was found in an associated blank as well as in the sample. B2 CCB B3 CCB - Neg B4 Grinding Blank C LCS Recovery C1 Reference Recovery C2 Reference Recovery RPD D MS RPD D1 Lab Replicate RPD D2 No precision available D3 Field Duplicate RPD D4 Field Triplicate RSD D5 Laboratory Triplicate RSD F Field Blank F1 Hydrocarbon pattern does not match standard G1 Initial Calibration RRF G2 Initial Calibration RRF H1 Test Hold Time H2 Prep Hold Time H2 Prep Hold Time H3 Surrogate recovery outside project limits. J CRA/CRI Recovery K 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. L Lab Blank L1 Lab Blank - Neg M MS Recovery M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.		
B2 CCB - Neg B3 CCB - Neg B4 Grinding Blank C LCS Recovery C1 Reference Recovery C2 Reference Recovery RPD D MS RPD D1 Lab Replicate RPD D2 No precision available D3 Field Duplicate RSD D5 Laboratory Triplicate RSD D5 Laboratory Triplicate RSD F Field Blank F1 Hydrocarbon pattern does not match standard G1 Initial Calibration RRF G2 Initial Calibration RSD/r^2/r G3 ICV RRF H1 Test Hold Time H2 Prep Hold Time H2 Prep Hold Time I Surrogate recovery outside project limits. J CRA/CRI Recovery K 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. L Lab Blank L1 Lab Blank L1 Lab Blank - Neg M MS Recovery M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	A1	
B3 CCB - Neg B4 Grinding Blank C LCS Recovery C1 Reference Recovery C2 Reference Recovery RPD D MS RPD D1 Lab Replicate RPD D2 No precision available D3 Field Duplicate RPD D4 Field Triplicate RSD D5 Laboratory Triplicate RSD F Field Blank F1 Hydrocarbon pattern does not match standard G1 Initial Calibration RRF G2 Initial Calibration RSD/r^2/r G3 ICV RRF H1 Test Hold Time H2 Prep Hold Time I Surrogate recovery outside project limits. J CRA/CRI Recovery K 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. L Lab Blank L1 Lab Blank - Neg M MS Recovery M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	В	The analyte was found in an associated blank as well as in the sample.
B4 Grinding Blank C LCS Recovery C1 Reference Recovery C2 Reference Recovery RPD D MS RPD D1 Lab Replicate RPD D2 No precision available D3 Field Duplicate RPD D4 Field Triplicate RSD D5 Laboratory Triplicate RSD F Field Blank F1 Hydrocarbon pattern does not match standard G1 Initial Calibration RRF G2 Initial Calibration RSD/r^2/r G3 ICV RRF H1 Test Hold Time H2 Prep Hold Time I Surrogate recovery outside project limits. J CRA/CRI Recovery K An analyte (non-common laboratory artifact) was detected in the associated method blank. L Lab Blank L1 Lab Blank L1 Lab Blank Neg M MS Recovery M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	B2	CCB
C LCS Recovery C1 Reference Recovery C2 Reference Recovery RPD D MS RPD D1 Lab Replicate RPD D2 No precision available D3 Field Duplicate RPD D4 Field Triplicate RSD D5 Laboratory Triplicate RSD D6 Laboratory Triplicate RSD D7 Field Blank D8 Field Blank D9 Field Initial Calibration RRF D9 Initial Calibration RRF D9 Initial Calibration RSD/r^2/r D9 Initial Calib	B3	CCB - Neg
C1 Reference Recovery C2 Reference Recovery RPD D MS RPD D MS RPD D1 Lab Replicate RPD D2 No precision available D3 Field Duplicate RPD D4 Field Triplicate RSD D5 Laboratory Triplicate RSD F Field Blank F1 Hydrocarbon pattern does not match standard G1 Initial Calibration RRF G2 Initial Calibration RSD/r^2/r G3 ICV RRF H1 Test Hold Time H2 Prep Hold Time Surrogate recovery outside project limits. J CRA/CRI Recovery K 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. L Lab Blank L1 Lab Blank - Neg M MS Recovery M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	B4	Grinding Blank
C2 Reference Recovery RPD D MS RPD D1 Lab Replicate RPD D2 No precision available D3 Field Duplicate RPD D4 Field Triplicate RSD D5 Laboratory Triplicate RSD F Field Blank F1 Hydrocarbon pattern does not match standard G1 Initial Calibration RRF G2 Initial Calibration RSD/r^2/r G3 ICV RRF H1 Test Hold Time H2 Prep Hold Time I Surrogate recovery outside project limits. J CRA/CRI Recovery K An analyte (non-common laboratory artifact) was detected in the associated method blank. L Lab Blank L1 Lab Blank - Neg M MS Recovery M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	С	LCS Recovery
D MS RPD D1 Lab Replicate RPD D2 No precision available D3 Field Duplicate RPD D4 Field Triplicate RSD D5 Laboratory Triplicate RSD F Field Blank F1 Hydrocarbon pattern does not match standard G1 Initial Calibration RRF G2 Initial Calibration RRF G3 ICV RRF H1 Test Hold Time H2 Prep Hold Time I Surrogate recovery outside project limits. J CRA/CRI Recovery K An analyte (non-common laboratory artifact) was detected in the associated method blank. L Lab Blank L1 Lab Blank Neg M MS Recovery M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	C1	Reference Recovery
D1 Lab Replicate RPD D2 No precision available D3 Field Duplicate RPD D4 Field Triplicate RSD D5 Laboratory Triplicate RSD F Field Blank F1 Hydrocarbon pattern does not match standard G1 Initial Calibration RRF G2 Initial Calibration RSD/r^2/r G3 ICV RRF H1 Test Hold Time H2 Prep Hold Time I Surrogate recovery outside project limits. J CRA/CRI Recovery K 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. L Lab Blank L1 Lab Blank Neg M MS Recovery M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	C2	Reference Recovery RPD
D2 No precision available D3 Field Duplicate RPD D4 Field Triplicate RSD D5 Laboratory Triplicate RSD F Field Blank F1 Hydrocarbon pattern does not match standard G1 Initial Calibration RRF G2 Initial Calibration RSD/r^2/r G3 ICV RRF H1 Test Hold Time H2 Prep Hold Time I Surrogate recovery outside project limits. J CRA/CRI Recovery K 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. L Lab Blank L1 Lab Blank L1 Lab Blank - Neg M MS Recovery M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	D	MS RPD
Field Duplicate RPD D4 Field Triplicate RSD D5 Laboratory Triplicate RSD F Field Blank F1 Hydrocarbon pattern does not match standard G1 Initial Calibration RRF G2 Initial Calibration RSD/r^2/r G3 ICV RRF H1 Test Hold Time H2 Prep Hold Time I Surrogate recovery outside project limits. J CRA/CRI Recovery K 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. L Lab Blank L1 Lab Blank Neg M MS Recovery M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	D1	Lab Replicate RPD
D4 Field Triplicate RSD D5 Laboratory Triplicate RSD F Field Blank F1 Hydrocarbon pattern does not match standard G1 Initial Calibration RRF G2 Initial Calibration RSD/r^2/r G3 ICV RRF H1 Test Hold Time H2 Prep Hold Time I Surrogate recovery outside project limits. J CRA/CRI Recovery K 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. L Lab Blank L1 Lab Blank L1 Lab Blank NM MS Recovery M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	D2	No precision available
D5 Laboratory Triplicate RSD F Field Blank F1 Hydrocarbon pattern does not match standard G1 Initial Calibration RRF G2 Initial Calibration RSD/r^2/r G3 ICV RRF H1 Test Hold Time H2 Prep Hold Time I Surrogate recovery outside project limits. J CRA/CRI Recovery K 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. L Lab Blank L1 Lab Blank - Neg M MS Recovery M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	D3	Field Duplicate RPD
Field Blank F1 Hydrocarbon pattern does not match standard G1 Initial Calibration RRF G2 Initial Calibration RSD/r^2/r G3 ICV RRF H1 Test Hold Time H2 Prep Hold Time I Surrogate recovery outside project limits. J CRA/CRI Recovery K 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. L Lab Blank L1 Lab Blank L1 Lab Blank - Neg M MS Recovery M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	D4	Field Triplicate RSD
Field Blank F1 Hydrocarbon pattern does not match standard G1 Initial Calibration RRF G2 Initial Calibration RSD/r^2/r G3 ICV RRF H1 Test Hold Time H2 Prep Hold Time I Surrogate recovery outside project limits. J CRA/CRI Recovery K 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. L Lab Blank L1 Lab Blank L1 Lab Blank - Neg M MS Recovery M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	D5	Laboratory Triplicate RSD
G1 Initial Calibration RRF G2 Initial Calibration RSD/r^2/r G3 ICV RRF H1 Test Hold Time H2 Prep Hold Time I Surrogate recovery outside project limits. J CRA/CRI Recovery K 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. L Lab Blank L1 Lab Blank L1 Lab Blank - Neg M MS Recovery M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	F	
G1 Initial Calibration RRF G2 Initial Calibration RSD/r^2/r G3 ICV RRF H1 Test Hold Time H2 Prep Hold Time I Surrogate recovery outside project limits. J CRA/CRI Recovery K 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. L Lab Blank L1 Lab Blank L1 Lab Blank - Neg M MS Recovery M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	F1	Hydrocarbon pattern does not match standard
G3 ICV RRF H1 Test Hold Time H2 Prep Hold Time I Surrogate recovery outside project limits. J CRA/CRI Recovery K 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. L Lab Blank L1 Lab Blank L1 Lab Blank - Neg M MS Recovery M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	G 1	
H1 Test Hold Time H2 Prep Hold Time I Surrogate recovery outside project limits. J CRA/CRI Recovery K 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. L Lab Blank L1 Lab Blank L1 Lab Blank - Neg M MS Recovery M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	G2	Initial Calibration RSD/r^2/r
H2 Prep Hold Time I Surrogate recovery outside project limits. J CRA/CRI Recovery K 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. L Lab Blank L1 Lab Blank - Neg M MS Recovery M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	G3	ICV RRF
I Surrogate recovery outside project limits. J CRA/CRI Recovery K 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. L Lab Blank L1 Lab Blank - Neg M MS Recovery M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	H1	Test Hold Time
I Surrogate recovery outside project limits. J CRA/CRI Recovery K 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. L Lab Blank L1 Lab Blank - Neg M MS Recovery M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	H2	Prep Hold Time
CRA/CRI Recovery 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. L Lab Blank L1 Lab Blank - Neg M MS Recovery M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	1	•
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. L Lab Blank L1 Lab Blank - Neg M MS Recovery M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	J	
L1 Lab Blank - Neg M MS Recovery M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	К	An analyte (non-common laboratory artifact) was detected in the sample at a concentration less than 5X the concentration detected in the associated
M MS Recovery M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	L	Lab Blank
M2 Post Spike N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	L1	Lab Blank - Neg
N Blank - No Action O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	М	MS Recovery
O ICS P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	M2	Post Spike
P Sample preservation/collection requirement not met. P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	N	Blank - No Action
P1 Column RPD P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	0	ICS
P2 Improper preparation/extraction Q Encore sample holding time exceeded by more than 2X.	P	Sample preservation/collection requirement not met.
Q Encore sample holding time exceeded by more than 2X.	P1	
Q Encore sample holding time exceeded by more than 2X.	P2	Improper preparation/extraction
·	Q	
Q'I Materiai Biank	Q1	Material Blank

Q2	Encore sample holding time exceeded by less than 2X.
R	Exceeds LinearCalibration Range
S	Internal standard
Т	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)
Х	Raised reporting limit
Υ	Cooler temperature greater than 10 degreec 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
Z 1	Non-accredited analyte/compound
Z 1	Data rejected, more valid data available.
Z2	Detection Level not met uncertainty greater than DL
Z4	MDA Greater than RDL.
Z 5	Ion Ratio
Z 6	Samples were analyzed past the 12 hour time period from the Tune or opening CCV.



Field Duplicate Calculations

Parameters	Original Sample	Duplicate Sample	RPD	LOQ
	AF-RHMW02-WGN01LF-2311	AF-RHMW02-WGFD01LF-2311	=	
2-/2-Butowiethow/ethanol	0	1	#DIV/0I	5

Flag



DATA VALIDATION GC (8015C) DOD

Validator: DM

Date Validated: 12/4/23

Reviewer: DLW

Date Reviewed: 12/4/23

Project: Red Hill Bulk Storage Facility, CV 23F0104

SDG: 580-133679-1

LAB: Eurofins, Savannah

Samples Collected: 11/6/2023

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.

No problems found

Holding Times

- ✓ Aqueous samples extracted within 7 days of collection and analyzed within 40 days of extraction
- ✓ Solid samples extracted within 14 days of collection and analyzed within 40 days of extraction.
- ✓ There is no specific holding time for PCB samples
- ✓ If temp of receipt is >6° but ≤15% qualify J-/UJ
- ✓ If temp >15°C qualify X
- ✓ If holding time is exceeded qualify J-/UJ
- ✓ If holding time is grossly exceeded (>2X) qualify J-/X

14-day holding time used for 8015C DAI

Samples collected on 11/06/2023 Samples analyzed on 11/16/2023

ICAL (Form VI)

- \checkmark %RSD ≤ 20% or r > 0.995 or R2>0.990
- ✓ minimum 5 standards for linear; minimum 6 standards for quadratic
- √ 5-pt calibration for multicomponent analytes
- ✓ If %RSD >20% or r<0.995 or r2<0.990 qualify J/UJ</p>
- ✓ If %RSD >40% or r<0.95 or r2<0.90 qualify X

Inst: CVGG2 on 11/15/2023

ICV/CCV (from VII/Analysis Run Log)

- ✓ ICV after ICAL; all %Ds within ± 20%
- ✓ CCVs before sample, every 10 samples
- ✓ all ICV/CCV %D ± 20%
- ✓ RTs within established window
- √ if %D is high then qualify J+
- ✓ if %D is low then qualify J-/UJ
- ✓ if %D is >50% then qualify X

Inst: CVGG2 on 11/15/2023

ICV 680-808404/14 11/15/2023 18:31 1 1IK15014.D J&W DB WAX 0.45(mm) CCVIS 680-808697/7 11/16/2023 18:33 1 1IK16007.D CCV 680-808697/22 11/17/2023 00:25 1 1IK16022.D 22BEE out (30%) sample ND Q UJ (V2) agree with ADR

All samples

CCV 680-808697/37 11/17/2023 06:16 1 1IK16037.D

Surrogate (Form II)

- ✓ if acceptance criteria is not defined by project, use limits in Table C below
- ✓ RTs within range of 5 pt
- ✓ do not evaluate for if diluted out
- ✓ if surrogate recovery <10% then qualify J-/X
- √ if surrogate recovery is low but >10% then qualify J-/UJ
- √ if surrogate recovery is high then qualify J+

No surrogate used

LCS (Form III)

✓ one per prep batch

✓ used QAPP 50-150, RPD 50

- ✓ if recovery low then qualify J-/X
- √ if recovery is high then qualify J+
- ✓ if LCS/LCSD RPD is out then qualify detects J, do not qualify NDs.

	LCS %R	LCSD %R	RPD
LCS/LCSD 680-808697	ok	ok	ok
LCS/LCSD 680-809119	ok	ok	ok

MS/MSD (Form III)

✓ one per prep batch

✓ used QAPP 50-150, RPD 50

- ✓ MS/MSD RPD ≤ 30%
- ✓ if MS recovery <10% then qualify J-/X
- √ if MS recovery is low but >10% then qualify J-/UJ
- √ if MS recovery is high then qualify J+
- ✓ if MS/MSD RPD is out then qualify detects J, do not qualify NDs

	MS	MSD	RPD
AF-RHMW16-WGN01LF-2311	ok	ok	ok

Blanks (Form IV/Form1)

✓ method blank – analyzed one per prep batch

Method Blank

MB 680-808697/11 ND MB 680-809119/8 ND

Field/Equipment Blank

None

Internal Standard Areas and RTs (Form VIII)

- ✓ areas within -50% to +100% of ICAL midpoint standard
- ✓ RTs within 30 seconds of midpoint standard
- ✓ if IS recovery is >200% then detects qualify J (do not qualify NDs)
- √ if IS recovery is <50% but >20% then qualify J/UJ
- ✓ if IS recovery is <20% or RT out then qualify X

Internal standard used – n-Heptyl Alcohol all ok

Identification Summary / Second Column Confirmation (Form X)

- ✓ present for all positive results
- ✓ RTs within range for both columns (not applicable for single column 8015C)
- ✓ RPD ≤ 40% Single col analyses NA
- √ if RPD >40% then qualify J Single col analyses NA

All samples ND NA

Sample Data (Form I)

- ✓ Chromatogram acceptable
- ✓ manual integrations acceptable

N/A-Level 2B

Field Duplicates

- ✓ no criteria per QSM; use project specific criteria when available in QAPP
- ✓ per Module 4: if RPD > QAPP limit qualify J, no Q for non-detects
- ✓ See field duplicate worksheet

use 30% for aqueous and solids

AF-RHMW02-WGN01LF-2311 AF-RHMW02-WGFD01LF-2311 OK

Facility: RH PFAS Sampling

Event: AFFF Assessment Sampling GW 2023 November

SDG: 5801336791

Guidance Document: RHS PFAS UFP-QAPP

Prime Contractor: AECOM, Honolulu, HI

Project Manager: Kristin Rutherford

Contract Laboratory(ies): Eurofins Environment Testing TestAmerica, Savannah, GA

Data Review Contractor: Environmental Data Services Ltd.

Data Review Level: S2BVEM

Primary Data Reviewer: Dina Manov, Technical Specialist

Date Submitted: December 04, 2023

Field Sample ID	Lab Sample ID	Matrix	Type/Type Code	SW8015C
AF-HDMW225303-WGN01LF- 2311	580-133679-5	Water	Field Sample/N	Χ
AF-RHMW02-WGFD01LF- 2311	580-133679-7	Water	Field Duplicate/FD	Χ
AF-RHMW02-WGN01LF-2311	580-133679-6	Water	Field Sample/N	Χ
AF-RHMW04-WGN01LF-2311	580-133679-3	Water	Field Sample/N	Χ
AF-RHMW06-WGN01LF-2311	580-133679-2	Water	Field Sample/N	Χ
AF-RHMW16-WGN01LF-2311	580-133679-1	Water	Field Sample/N	Χ
AF-RHMW225401-WGN01B- 2311	580-133679-4	Water	Field Sample/N	Χ

This report assesses the analytical data quality associated with the analyses listed on the preceding cover page at S2BVEM data validation level. This assessment has been made through a combination of automated data review (ADR) and supplemental manual review, the details of which are described below. The approach taken in the review of this data set is consistent with the requirements contained in the RHS PFAS UFP-QAPP and the additional guidance documents incorporated by reference to the extent possible. Where definitive guidance is not provided, results have been evaluated in a conservative manner using professional judgment.

Sample collection was managed and directed by AECOM, Honolulu, HI; analyses were performed by Eurofins Environment Testing TestAmerica, Savannah, GA and were reported under sample delivery group (SDG) 5801336791. Data have been evaluated electronically based on electronic data deliverables (EDDs) provided by the laboratory, and hard copy data summary forms have also been reviewed during this effort and compared to the automated review output by the reviewers whose signatures appear on the following page. Findings based on the automated data submission and manual data verification processes are detailed in the ADR narrative and throughout this report.

All quality control (QC) elements associated with this SDG have been reviewed by a project chemist in accordance with the requirements defined for the project. This review is documented in the attached Data Review Checklists. The QC elements listed below were supported by the electronic deliverable and were evaluated using ADR processes.

Continuing Calibration Verification

Field Duplicate RPD

Lab Blank

LCS Recovery

LCS RPD

MS Recovery

MS RPD

Prep Hold Time

Surrogate

Test Hold Time

Results of the ADR process were subsequently reviewed and updated as applicable by the data review chemists identified on the signature page. Quality control elements that were not included in the electronic deliverable were reviewed manually and findings are documented within this report. Summaries of findings and associated qualified results are documented throughout this report.

A total of 7 results (100.00%) out of the 7 results (sample and field QC samples) reported are qualified based on review and 0 results (0.00%) have been rejected or deemed a serious deficiency (X qualifier). Trace values, defined as results that are qualified as estimated because they fall between the detection limit and the reporting limit/limit of quantitation, are not counted as qualified results in the above count. The qualified results are detailed throughout this report and discussed in the narrative below, where appropriate.

Narrative Comments

Analytical Method	Data Reviewer Comment
SW8015C	No additional comments; see Checklist for detail.

December 04, 2023

Reviewed by Dina Manov, Technical Specialist, Environmental Data Services Ltd.

As the Reviewer, I certify that I have performed a data review process in accordance with the requirements of the project guidance document, and have compared the electronic data to the laboratory's hard copy report and have verified the consistency of the reported sample results and method quality control data between the two deliverables.

Quality Control Outliers for test method SW8015C, Continuing Calibration Verification

Compliance requirements for satisfactory continuing calibration are established to ensure that the instrument is capable of producing acceptable qualitative and quantitative data. Continuing calibration is performed to verify and evaluate instrument performance during sample analysis. Summary forms were evaluated against project acceptance criteria, and any associated qualified results, are listed below.

Sample ID/ Lab Sample ID	Analyte	Result	Warning Limits	Control Limits	Units	Qualifier	Reason Code	Comment
CCV68080869722 (CV)	2-(2- Butoxyethoxy)etha nol	130	80 - 120	50 - 150	percent	J/UJ	V2	
CCV68080911910 (CV)	2-(2- Butoxyethoxy)etha nol	76.0	80 - 120	50 - 150	percent	J/UJ	V2	

Where two qualifiers are listed, such as 'J/UJ', the first applies to positive results, and the second to non-detect results. Upper and Lower Warning and Control Limits are abbreviated UWL, LWL, UCL, and LCL in the Comment field.

Qualified Results associated with the Continuing Calibration Verification for SW8015C

FieldSample ID	Туре	Analyte	LOQ	Lab Result	Qualified Result	Bias	Units	Reason
AF-HDMW225303- WGN01LF-2311 580-133679-5	N	2-(2-Butoxyethoxy)ethanol	5.00	3.00 U Q	3.00 UJ		mg/l	V2
AF-RHMW02- WGFD01LF-2311 580-133679-7	FD	2-(2-Butoxyethoxy)ethanol	5.00	3.00 U Q	3.00 UJ		mg/l	V2
AF-RHMW02- WGN01LF-2311 580-133679-6	N	2-(2-Butoxyethoxy)ethanol	5.00	3.00 U Q	3.00 UJ		mg/l	V2
AF-RHMW04- WGN01LF-2311 580-133679-3	N	2-(2-Butoxyethoxy)ethanol	5.00	3.00 U M Q	3.00 UJ		mg/l	V2
AF-RHMW06- WGN01LF-2311 580-133679-2	N	2-(2-Butoxyethoxy)ethanol	5.00	3.00 U Q	3.00 UJ		mg/l	V2
AF-RHMW16- WGN01LF-2311 580-133679-1	N	2-(2-Butoxyethoxy)ethanol	5.00	3.00 U Q	3.00 UJ		mg/l	V2
AF-RHMW225401- WGN01B-2311 580-133679-4	N	2-(2-Butoxyethoxy)ethanol	5.00	3.00 U M Q	3.00 UJ		mg/l	V2

Analytes not found in project samples are reported as not detected at the limit of detection (LOD) unless blank contamination occurs and then the sample may be reported as not detected at the (LOD) or (LOQ) based on the sample concentration and the validation guidance. In instances where no LOD is provided, results are reported down to the LOQ.

Table of All Qualified Results

Test Method: SW8015C		Extraction Method: METHOD						
FieldSample ID / LabSample ID	Туре	Analyte	LOQ	Lab Result	Qualified Result	Bias	Units	Reason
AF-HDMW225303- WGN01LF-2311 580-133679-5	N	2-(2-Butoxyethoxy)ethanol	5.00	3.00 U Q	3.00 UJ		mg/l	V2
AF-RHMW02- WGFD01LF-2311 580-133679-7	FD	2-(2-Butoxyethoxy)ethanol	5.00	3.00 U Q	3.00 UJ		mg/l	V2
AF-RHMW02- WGN01LF-2311 580-133679-6	N	2-(2-Butoxyethoxy)ethanol	5.00	3.00 U Q	3.00 UJ		mg/l	V2
AF-RHMW04- WGN01LF-2311 580-133679-3	N	2-(2-Butoxyethoxy)ethanol	5.00	3.00 U M Q	3.00 UJ		mg/l	V2
AF-RHMW06- WGN01LF-2311 580-133679-2	N	2-(2-Butoxyethoxy)ethanol	5.00	3.00 U Q	3.00 UJ		mg/l	V2
AF-RHMW16- WGN01LF-2311 580-133679-1	N	2-(2-Butoxyethoxy)ethanol	5.00	3.00 U Q	3.00 UJ		mg/l	V2
AF-RHMW225401- WGN01B-2311 580-133679-4	N	2-(2-Butoxyethoxy)ethanol	5.00	3.00 U M Q	3.00 UJ		mg/l	V2

Analytes not found in project samples are reported as not detected at the limit of detection (LOD) unless blank contamination occurs and then the sample may be reported as not detected at the (LOQ) based on the sample concentration. In instances where no LOD is provided, results are reported down to the LOQ.

Trace values are not included in the qualified results table unless additional reason codes are associated.

Results with Modified Qualifiers

No qualifiers associated with this sample delivery group were modified manually.

Reason Code Definitions					
Code	Definition				
V2	CCV				
Flag Cod	e and Definitions				
Flag	Definition				
J	Estimated Value				
N	The analysis indicates the presence of an analyte for which there was presumptive evidence to make a tentative identification.				
NJ	The analyte has been tentatively identified or presumptively as present and the associated numerical value was the estimated concentration in the sample.				
R	The data are rejected due to deficiencies in meeting QC criteria and may not be used for decision making.				
U	Undetected: The analyte was analyzed for, but not detected.				
UJ	The analyte was not detected; however, the result is estimated due to discrepancies in meeting certain analyte-specific quality control criteria.				
X	Result may require rejection; PDT attention required				
Bias					
-	The result may be biased low				
+	The result may be biased high				
Note - The	e bias field is a separate field; however, it is an integral part of the final flag (qualifier) on the sample result				

Page 6 of 8