



# LABORATORY DATA CONSULTANTS, INC.

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March 28, 2022

SUBJECT: Red Hill Bulk Storage Facility, CTO 18F0126 (NOI) - Data Validation

Dear Ramos,

Enclosed are the final validation reports for the fractions listed below. This SDG was received on December 6, 2021. Attachment 1 is a summary of the samples that were reviewed for each analysis.

**LDC Project # 52825B:**

<b><u>SDG #</u></b>	<b><u>Fraction</u></b>
B21110712	Total Petroleum Hydrocarbons as Extractables

The data validation was performed under Stage 2B & 4 guidelines. The analysis were validated using the following documents, as applicable to each method:

- Work Plan/Scope of Work, 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 02, January 2017)
- 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)
- Sampling and Analysis Plan, Addendum 01, 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 00, September 2017)
- Sampling and Analysis Plan, Addendum 03, 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 00, June 2018)
- U.S. Department of Defense (DoD) Quality Systems Manual (QSM) for Environmental Laboratories, Version 5.3 (2019)
- DoD General Validation Guidelines (November 2019)
- U.S. Department of Defense (DoD) Data Validation Guidelines Module 4: Data Validation Procedure for Organic Analysis by GC (March 2021)
- EPA SW 846, Third Edition, Test Methods for Evaluating Solid Waste, update 1, July 1992; update IIA, August 1993; update II, September 1994; update IIB, January 1995; update III, December 1996; update IIIA, April 1998; IIIB, November 2004; update IV, February 2007; update V, July 2014; update VI, July 2018

Please feel free to contact us if you have any questions.

Sincerely,

Stella Cuenco  
[scuenco@lab-data.com](mailto:scuenco@lab-data.com)  
Project Manager/Senior Chemist



## Laboratory Data Consultants, Inc. Data Validation Report

**Project/Site Name:** Red Hill Bulk Storage Facility, CTO 18F0126

**LDC Report Date:** March 21, 2022

**Parameters:** Total Petroleum Hydrocarbons as Extractables

**Validation Level:** Stage 2B & 4

**Laboratory:** Energy Laboratories, Billings, MT

**Sample Delivery Group (SDG):** B21110712

Sample Identification	Laboratory Sample Identification	Matrix	Collection Date
ERH1873(RHMW01R)**	B21110712-001**	Water	11/03/21
ERH1876(RHMW02)	B21110712-002	Water	11/03/21
ERH1879(RHMW03)	B21110712-003	Water	11/03/21
ERH1882(RHMW05)	B21110712-004	Water	11/03/21
ERH1885(RHMW2254-01)	B21110712-005	Water	11/03/21
ERH1888(RHSF)	B21110712-006	Water	11/03/21
ERH1890(RHSF)	B21110712-007	Water	11/03/21
ERH1873(RHMW01R)(SGCU)**	B21110712-001(SGCU)**	Water	11/03/21
ERH1876(RHMW02)(SGCU)	B21110712-002(SGCU)	Water	11/03/21
ERH1879(RHMW03)(SGCU)	B21110712-003(SGCU)	Water	11/03/21
ERH1882(RHMW05)(SGCU)	B21110712-004(SGCU)	Water	11/03/21
ERH1873(RHMW01R)MS	B21110712-001MS	Water	11/03/21
ERH1876(RHMW02)MS	B21110712-002MS	Water	11/03/21
ERH1873(RHMW01R)(SGCU)MS	B21110712-001(SGCU)MS	Water	11/03/21
ERH1876(RHMW02)(SGCU)MS	B21110712-002(SGCU)MS	Water	11/03/21

\*\*Indicates sample underwent Stage 4 validation  
Samples appended with SGCU underwent "Silica Gel Clean Up"

## Introduction

This Data Validation Report (DVR) presents data validation findings and results for the associated samples listed on the cover page. Data validation was performed in accordance with the Work Plan/Scope of Work, 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 02, January 2017), the 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), the Sampling and Analysis Plan, Addendum 01, 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 00, September 2017), the Sampling and Analysis Plan, Addendum 03, 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 00, June 2018), the U.S. Department of Defense (DoD) Quality Systems Manual (QSM) for Environmental Laboratories, Version 5.3 (2019), the DoD General Validation Guidelines (November 2019), and the U.S. Department of Defense (DoD) Data Validation Guidelines Module 4: Data Validation Procedure for Organic Analysis by GC (March 2021). Where specific guidance was not available, the data has been evaluated in a conservative manner consistent with industry standards using professional experience.

The analyses were performed by the following method:

Total Petroleum Hydrocarbons (TPH) as Extractables by Environmental Protection Agency (EPA) SW 846 Method 8015C

All sample results were subjected to Stage 2B data validation, which comprises an evaluation of quality control (QC) summary results. Samples appended with a double asterisk on the cover page were subjected to Stage 4 data validation, which is comprised of the QC summary forms as well as the raw data, to confirm sample quantitation and identification.

The following are definitions of the data qualifiers utilized during data validation:

- J+ (Estimated, High Bias): The analyte was analyzed for and positively identified by the laboratory; however the reported concentration is estimated, displaying high bias, due to non-conformances discovered during data validation.
- J- (Estimated, Low Bias): The analyte was analyzed for and positively identified by the laboratory; however the reported concentration is estimated, displaying low bias, due to non-conformances discovered during data validation.
- J (Estimated, Bias Indeterminate): The analyte was analyzed for and positively identified by the laboratory; however the reported concentration is estimated due to non-conformances discovered during data validation. Bias is indeterminate.
- U (Non-detected): The analyte was analyzed for and positively identified by the laboratory; however the analyte should be considered non-detected due to the presence of contaminants detected in the associated blank(s).
- UJ (Non-detected estimated): The analyte was not detected and the associated numerical value is approximate.
- X (Exclusion of data recommended): 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. Exclusion of the data is recommended.
- NA (Not Applicable): The non-conformance discovered during data validation demonstrates a high bias, while the affected analyte in the associated sample(s) was reported as not detected by the laboratory and did not warrant the qualification of the data.

A qualification summary table is provided at the end of this report if data has been qualified. Flags are classified as P (protocol) or A (advisory) to indicate whether the flag is due to a laboratory deviation from a specified protocol or is of technical advisory nature.

## Qualification Code Reference

- a ICP Serial Dilution %D was not within control limits.
- b Presumed contamination from preparation (method blank).
- c Calibration %RSD,  $r$ ,  $r^2$ , %D or %R was noncompliant.
- d The analysis with this flag should not be used because another more technically sound analysis is available.
- e MS/MSD or Duplicate RPD was high.
- f Presumed contamination from FB or ER.
- g ICP ICS results were unsatisfactory.
- h Holding times were exceeded.
- i Internal standard performance was unsatisfactory.
- k Estimated Maximum Possible Concentration (HRGC/HRMS only)
- l LCS/LCSD %R was not within control limits.
- m Result exceeded the calibration range.
- o Cooler temperature or temperature blank was noncompliant and/or sample custody problems.
- p RPD between two columns was high (GC only).
- q MS/MSD recovery was not within control limits.
- s Surrogate recovery was not within control limits.
- t Presumed contamination from trip blank.
- v Unusual problems found with the data not defined elsewhere. Description of the problem can be found in the validation report.
- w LCS/LCSD RPD was high.
- y Chemical recovery was not within control limits (Radiochemistry only).

## **I. Sample Receipt and Technical Holding Times**

All samples were received in good condition and cooler temperatures upon receipt met validation criteria.

All technical holding time requirements were met.

## **II. Initial Calibration and Initial Calibration Verification**

An initial calibration was performed as required by the method.

The percent relative standard deviations (%RSD) were less than or equal to 20.0% for all analytes.

The percent differences (%D) of the initial calibration verification (ICV) standard were less than or equal to 20.0% for all analytes.

## **III. Continuing Calibration**

Continuing calibration was performed at the required frequencies.

The percent differences (%D) were less than or equal to 20.0% for all analytes.

The percent differences (%D) of the ending continuing calibration verifications (CCVs) were less than or equal to 20.0% for all analytes.

## **IV. Laboratory Blanks**

Laboratory blanks were analyzed as required by the method. No contaminants were found in the laboratory blanks.

## **V. Field Blanks**

No field blanks were identified in this SDG.

## **VI. Surrogates**

Surrogates were added to all samples as required by the method. All surrogate recoveries (%R) were within QC limits.

## **VII. Matrix Spike/Matrix Spike Duplicates**

Matrix spike (MS) sample analysis was performed on an associated project sample. Percent recoveries (%R) were within QC limits.

## **VIII. Laboratory Control Samples**

Laboratory control samples (LCS) and laboratory control samples duplicates (LCSD) were analyzed as required by the method. Percent recoveries (%R) were within QC limits. Relative percent differences (RPD) were within QC limits.

## **IX. Field Duplicates**

Samples ERH1888(RHSF) and ERH1890(RHSF) were identified as field duplicates. No results were detected in any of the samples.

## **X. Target Analyte Quantitation**

All target analyte quantitations met validation criteria for samples which underwent Stage 4 validation. Raw data were not reviewed for Stage 2B validation.

## **XI. Target Analyte Identification**

All target analyte identifications met validation criteria for samples which underwent Stage 4 validation. Raw data were not reviewed for Stage 2B validation.

## **XII. Overall Assessment of Data**

The analysis was conducted within all specifications of the method. No results were rejected or recommended for exclusion in this SDG.



**Red Hill Bulk Storage Facility, CTO 18F0126  
Total Petroleum Hydrocarbons as Extractables - Data Qualification Summary -  
SDG B21110712**

No Sample Data Qualified in this SDG

**Red Hill Bulk Storage Facility, CTO 18F0126  
Total Petroleum Hydrocarbons as Extractables - Laboratory Blank Data  
Qualification Summary - SDG B21110712**

No Sample Data Qualified in this SDG

**Red Hill Bulk Storage Facility, CTO 18F0126  
Total Petroleum Hydrocarbons as Extractables - Field Blank Data Qualification  
Summary - SDG B21110712**

No Sample Data Qualified in this SDG

LDC #: 52825B8

## VALIDATION COMPLETENESS WORKSHEET

Date: 1/6/22

SDG #: B21110712

Stage 2B/4

Page: 1 of 1

Laboratory: Energy Laboratories, Billings, MT

Reviewer:    2nd Reviewer:    **METHOD:** GC TPH as Extractables (EPA SW-846 Method 8015C)

The samples listed below were reviewed for each of the following validation areas. Validation findings are noted in attached validation findings worksheets.

	Validation Area		Comments
I.	Sample receipt/Technical holding times	$\Delta$ , $\Delta$	
II.	Initial calibration/ICV	$\Delta$ , $\Delta$	% PSD = 20    ICV = 20
III.	Continuing calibration	$\Delta$	CCV = 20/20
IV.	Laboratory Blanks	$\Delta$	
V.	Field blanks	N	
VI.	Surrogate spikes	$\Delta$	
VII.	Matrix spike/Matrix spike duplicates	$\Delta$	MS only
VIII.	Laboratory control samples	$\Delta$	ICS 10
IX.	Field duplicates	ND	D = 4.7
X.	Target analyte quantitation	$\Delta$	Not reviewed for Stage 2B validation.
XI.	Target analyte identification	$\Delta$	Not reviewed for Stage 2B validation.
XII.	Overall assessment of data	$\Delta$	

Note: A = Acceptable  
N = Not provided/applicable  
SW = See worksheet

ND = No compounds detected  
R = Rinsate  
FB = Field blank

D = Duplicate  
TB = Trip blank  
EB = Equipment blank

SB=Source blank  
OTHER:

\*\* Indicates sample underwent Stage 4 validation

	Client ID	Lab ID	Matrix	Date
1 <sup>+</sup>	ERH1873(RHMW01R)**	B21110712-001**	Water	11/03/21
2 <sup>+</sup>	ERH1876(RHMW02)	B21110712-002	Water	11/03/21
3 <sup>+</sup>	ERH1879(RHMW03)	B21110712-003	Water	11/03/21
4 <sup>+</sup>	ERH1882(RHMW05)	B21110712-004	Water	11/03/21
5 <sup>-</sup>	ERH1885(RHMW2254-01)	B21110712-005	Water	11/03/21
6 <sup>-</sup>	ERH1888(RHSF) D	B21110712-006	Water	11/03/21
7 <sup>-</sup>	ERH1890(RHSF) D	B21110712-007	Water	11/03/21
8 <sup>+</sup>	ERH1873(RHMW01R)(SGCU)**	B21110712-001(SGCU)**	Water	11/03/21
9 <sup>+</sup>	ERH1876(RHMW02)(SGCU)	B21110712-002(SGCU)	Water	11/03/21
10 <sup>+</sup>	ERH1879(RHMW03)(SGCU)	B21110712-003(SGCU)	Water	11/03/21
11 <sup>+</sup>	ERH1882(RHMW05)(SGCU)	B21110712-004(SGCU)	Water	11/03/21
12	ERH1873(RHMW01R)MS	B21110712-001MS	Water	11/03/21
13	ERH1876(RHMW02)MS	B21110712-002MS	Water	11/03/21
14	ERH1873(RHMW01R)(SGCU)MS	B21110712-001(SGCU)MS	Water	11/03/21
15	ERH1876(RHMW02)(SGCU)MS	B21110712-002(SGCU)MS	Water	11/03/21
16	161122			
17				

Method:  GC  HPLC

Validation Area	Yes	No	NA	Findings/Comments
<b>I. Technical holding times</b>				
Were all technical holding times met?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was cooler temperature criteria met?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
<b>IIa. Initial calibration</b>				
Did the laboratory perform a 5 point calibration prior to sample analysis?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were all percent relative standard deviations (%RSD) < 20%?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was a curve fit used for evaluation? If yes, did the initial calibration meet the curve fit acceptance criteria of $\geq 0.990$ ?	<input type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	
Were the RT windows properly established?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
<b>IIb. Initial calibration verification</b>				
Was an initial calibration verification standard analyzed after each initial calibration for each instrument?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were all percent differences (%D) < 20%?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
<b>III. Continuing calibration</b>				
Was a continuing calibration analyzed daily?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were all percent differences (%D) < 20%?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were all the retention times within the acceptance windows?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
<b>IV. Laboratory Blanks</b>				
Was a laboratory blank associated with every sample in this SDG?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was a laboratory blank analyzed for each matrix and concentration?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was there contamination in the laboratory blanks?	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	
<b>V. Field Blanks</b>				
Were field blanks identified in this SDG?	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	
Were target analytes detected in the field blanks?	<input type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	
<b>VI. Surrogate spikes</b>				
Were all surrogate percent recovery (%R) within the QC limits?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
If the percent recovery (%R) of one or more surrogates was outside QC limits, was a reanalysis performed to confirm %R?	<input type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	
If any %R was less than 10 percent, was a reanalysis performed to confirm %R?	<input type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	
<b>VII. Matrix spike/Matrix spike duplicates</b>				
Were matrix spike (MS) and matrix spike duplicate (MSD) analyzed in this SDG?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were the MS/MSD percent recoveries (%R) and the relative percent differences (RPD) within the QC limits?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
<b>VIII. Laboratory control samples</b>				
Was an LCS analyzed per analytical or extraction batch?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were the LCS percent recoveries (%R) and relative percent difference (RPD) within the QC limits?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	

Validation Area	Yes	No	NA	Findings/Comments
<b><i>IX. Field duplicates</i></b>				
Were field duplicate pairs identified in this SDG?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were target analytes detected in the field duplicates?	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	
<b><i>X. Target analyte quantitation</i></b>				
Did the laboratory LOQs/RLs meet the QAPP LOQs/RLs?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were analyte quantitation and RLs adjusted to reflect all sample dilutions and dry weight factors applicable to level IV validation?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
<b><i>XI. Target analyte identification</i></b>				
Were the retention times of reported detects within the RT windows?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were manual integrations reviewed and found acceptable?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Did the laboratory provide before and after integration printouts?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
<b><i>XIII. Overall assessment of data</i></b>				
Overall assessment of data was found to be acceptable.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	

LDC #: 52 825 BY

## VALIDATION FINDINGS WORKSHEET Initial Calibration Calculation Verification

Page: 1 of 1  
Reviewer: FT

METHOD: GC  HPLC

The calibration factors (CF) and relative standard deviation (%RSD) were recalculated using the following calculations:

CF = A/C  
Average CF = sum of the CF/number of standards  
%RSD = 100 \* (S/X)

Where: A = Area of compound  
C = Concentration of compound  
S = Standard deviation of calibration factors  
X = Mean of calibration factors

#	Standard ID	Calibration Date	Compound	Reported	Recalculated	Reported	Recalculated	Reported	Recalculated
				CF (15000 std)	CF (15000 std)	CF (initial)	CF (initial)	%RSD	%RSD
1	ICAL For DRU TEH	11/2/21	DRU	<del>47852.77 FT</del> 4108 → 31901.86 5000	31901.86 5000	31353.19	31353.19	2.487	2.487
2	ICAL FOR RRU	2/18/21	DRU	28746.28	28746.28	28542	28542	4.497	4.497
3									
4									

Comments: Refer to Initial Calibration findings worksheet for list of qualifications and associated samples when reported results do not agree within 10.0% of the recalculated results.

LDC #: 52825 BX

## VALIDATION FINDINGS WORKSHEET Continuing Calibration Results Verification

Page: 1 of 1  
Reviewer: FT

METHOD: GC  HPLC

The percent difference (%D) of the initial calibration average Calibration Factors (CF) and the continuing calibration CF were recalculated for the target analytes identified below using the following calculation:

$$\% \text{ Difference} = 100 * (\text{ave. CF} - \text{CF}) / \text{ave. CF}$$

Where: ave. CF = initial calibration average CF  
CF = continuing calibration CF  
A = Area of target analyte  
C = Concentration of target analyte

#	Standard ID	Calibration Date	Target Analyte	Average CF(Ical)/ CCV Conc.	Reported	Recalculated	Reported	Recalculated
					CF/ Conc. CCV	CF/ Conc. CCV	%R	%R
1	CCV 504r	11/11/21 0649	DRO C <sub>10</sub> -E <sub>24</sub>	15.0	15.0	15.0	100	100
						15.583	103	104
2	CCV 520r	11/11/21 1738	↓	15.0	14.0	14.394	96.0	96.0
3	CCV 533r-W	11/12/21 0338	↓	15.0	15.0	14.893	99.0	99.0
4								

Comments: Refer to Continuing Calibration findings worksheet for list of qualifications and associated samples when reported results do not agree within 10.0% of the recalculated results.

**VALIDATION FINDINGS WORKSHEET**  
**Surrogate Results Verification**

METHOD:  GC  HPLC

The percent recoveries (%R) of surrogates were recalculated for the compounds identified below using the following calculation:

% Recovery: SF/SS \* 100

Where: SF = Surrogate Found  
SS = Surrogate Spiked

Sample ID: #1

Surrogate	Column/Detector	Surrogate Spiked	Surrogate Found	Percent Recovery	Percent Recovery	Percent Difference
				Reported	Recalculated	
<u>o-Terphenyl</u>		<u>0.194</u>	<u>0.146</u>	<u>75.0</u>	<u>75</u>	<u>0</u>
<u>n-Triacontane</u>		<u>0.097</u>	<u>0.085</u>	<u>88.0</u>	<u>88</u>	<u>0</u>

Sample ID:

Surrogate	Column/Detector	Surrogate Spiked	Surrogate Found	Percent Recovery	Percent Recovery	Percent Difference
				Reported	Recalculated	

	Surrogate Compound		Surrogate Compound		Surrogate Compound		Surrogate Compound		Surrogate Compound
A	Chlorobenzene (CBZ)	G	Octacosane	M	Benzo(e)Pyrene	S	1-Chloro-3-Nitrobenzene	Y	Tetrachloro-m- xylene
B	4-Bromofluorobenzene (BFB)	H	Ortho-Terphenyl	N	Terphenyl-D14	T	3,4-Dinitrotoluene	Z	2-Bromonaphthalene
C	a,a,a-Trifluorotoluene	I	Fluorobenzene (FBZ)	O	Decachlorobiphenyl (DCB)	U	Triphenyltin	AA	Chloro-octadecane
D	Bromochlorobenzene	J	n-Triacontane	P	1-methylnaphthalene	V	Tri-n-propyltin	BB	2,4-Dichlorophenylacetic acid
E	1,4-Dichlorobutane	K	Hexacosane	Q	Dichlorophenyl Acetic Acid (DCAA)	W	Tributyl Phosphate	CC	2,5-Dibromotoluene
F	1,4-Difluorobenzene (DFB)	L	Bromobenzene	R	4-Nitrophenol	X	Triphenyl Phosphate		





LDC #: 52825PY

## VALIDATION FINDINGS WORKSHEET

### Laboratory Control Sample/Laboratory Control Sample Duplicates Results Verification

Page: 1 of 1  
Reviewer: FT

METHOD: GC HPLC

The percent recoveries (%R) and relative percent differences (RPD) of the laboratory control sample and laboratory control sample duplicate were recalculated for the target analytes identified below using the following calculation:

%Recovery =  $100 * (SSC/SA)$   
 RPD =  $((SSCLCS - SSCLCSD) * 2) / (SSCLCS + SSCLCSD) * 100$

Where SSC = Spiked sample concentration  
 LCS = Laboratory Control Sample

SA = Spike added  
 LCSD = Laboratory Control Sample duplicate

LCS/LCSD samples: LCS 10 - 161122

Compound	Spike Added (mg/L)		Spike Sample Concentration (mg/L)		LCS		LCSD		LCS/LCSD	
	LCS	LCSD	LCS	LCSD	Percent Recovery		Percent Recovery		RPD	
					Reported	Recalc.	Reported	Recalc.	Reported	Recalc.
DRU	15	15	13.3259	13 (12.72302)	89.0	89	85	85	4.6	4.6

Comments: \_\_\_\_\_

LDC #: 52825 P8

### VALIDATION FINDINGS WORKSHEET Sample Calculation Verification

Page: 1 of 1  
Reviewer: FT

METHOD:  GC  HPLC

The concentration of the sample was calculated for the target analyte identified below using the following calculation:

$$\text{Concentration} = \frac{(A)(Fv)(Df)}{(RF)(Vs \text{ or } Ws)(\%S/100)}$$

Example:

Sample ID. #1 : DRU C10-C24

- A= Area or height of the target analyte to be measured
- Fv= Final Volume of extract
- Df= Dilution Factor
- RF= Average response factor of the target analyte  
In the initial calibration
- Vs= Initial volume of the sample
- Ws= Initial weight of the sample
- %S= Percent Solid

$$\text{Concentration} = \frac{1.51811 \times 10^7 (1)}{31353.19 (1030)} = 0.4701 \text{ mg/L}$$

*SGT analysis 11/12/21 1123*

*Analysis date 11/11/21*

#	Sample ID	Target analyte	Reported Concentrations (mg/L)	Recalculated Results Concentrations (mg/L)	Qualifications @ 1947
	#1	DRU C10-C24	0.47	0.4701	

Comments: \_\_\_\_\_  
\_\_\_\_\_  
\_\_\_\_\_