



# LABORATORY DATA CONSULTANTS, INC.

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Windward Environmental, LLC  
200 West Mercer Street, Suite 401  
Seattle, WA 98119  
ATTN: Amara Vandervort  
[amarav@windwardenv.com](mailto:amarav@windwardenv.com)

December 23, 2021

SUBJECT: Duwamish AOC4 - Data Validation

Dear Ms. Vandervort,

Enclosed are the final validation reports for the fractions listed below. These SDGs were received on November 18 and December 1, 2021. Attachment 1 is a summary of the samples that were reviewed for each analysis.

### LDC Project #52703:

<u>SDG #</u>	<u>Fraction</u>
21J0131, 21J0134, 21J0137 21J0142, 21K0332	Semivolatiles, PAHs, Hexachlorobenzene, PCBs, Metals, Dioxins, Wet Chemistry

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

- Final Lower Duwamish Waterway Quality Assurance Project Plan for Remedial Design of Upper Reach: Pre-Design Investigation (May 2020)
- USEPA National Functional Guidelines (NFG) for Organic Superfund Methods Data Review (January 2017)
- USEPA National Functional Guidelines (NFG) for Inorganic Superfund Methods Data Review (January 2017)
- USEPA National Functional Guidelines (NFG) for High Resolution Superfund Methods Data Review (April 2016)
- 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,

Pei Geng  
Project Manager/Senior Chemist  
[pgeng@lab-data.com](mailto:pgeng@lab-data.com)



## Laboratory Data Consultants, Inc. Data Validation Report

**Project/Site Name:** Duwamish AOC4

**LDC Report Date:** December 15, 2021

**Parameters:** Polychlorinated Biphenyls

**Validation Level:** Stage 4

**Laboratory:** Analytical Resources, Inc., Tukwila

**Sample Delivery Group (SDG):** 21J0131

Sample Identification	Laboratory Sample Identification	Matrix	Collection Date
LDW21-SC560F	21J0131-01	Sediment	06/29/21
LDW21-SC509D	21J0131-02	Sediment	07/01/21
LDW21-SC517D	21J0131-03	Sediment	07/01/21
LDW21-SC520A	21J0131-04	Sediment	07/02/21
LDW21-SC527D	21J0131-05	Sediment	07/02/21
LDW21-SC531D	21J0131-06	Sediment	07/02/21
LDW21-SC534D	21J0131-07	Sediment	07/02/21
LDW21-IT601	21J0131-08	Sediment	07/06/21
LDW21-IT592B	21J0131-09	Sediment	07/06/21
LDW21-IT592C	21J0131-10	Sediment	07/06/21
LDW21-IT592D	21J0131-11	Sediment	07/06/21
LDW21-IT592D	21J0131-12	Sediment	07/06/21
LDW21-IT592F	21J0131-13	Sediment	07/06/21
LDW21-IT592G	21J0131-14	Sediment	07/06/21
LDW21-SC510F	21J0131-15	Sediment	07/07/21
LDW21-IT609D	21J0131-16	Sediment	07/07/21
LDW21-SC595	21J0131-18	Sediment	07/08/21
LDW21-SC519B	21J0131-19	Sediment	07/08/21
LDW21-SC519C	21J0131-20	Sediment	07/08/21
LDW21-SC519D	21J0131-21	Sediment	07/08/21
LDW21-SC519E	21J0131-22	Sediment	07/08/21
LDW21-SC519F	21J0131-23	Sediment	07/08/21
LDW21-SC535D	21J0131-24	Sediment	07/08/21
LDW21-SC509DMS	21J0131-02MS	Sediment	07/01/21
LDW21-SC509DMSD	21J0131-02MSD	Sediment	07/01/21
LDW21-IT592CMS	21J0131-10MS	Sediment	07/06/21

<b>Sample Identification</b>	<b>Laboratory Sample Identification</b>	<b>Matrix</b>	<b>Collection Date</b>
LDW21-IT592CMSD	21J0131-10MSD	Sediment	07/06/21

## 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 Final Lower Duwamish Waterway Quality Assurance Project Plan for Remedial Design of Upper Reach: Pre-Design Investigation (May 2020) and a modified outline of the USEPA National Functional Guidelines (NFG) for Organic Superfund Methods Data Review (January 2017). 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:

Polychlorinated Biphenyls (PCBs) by Environmental Protection Agency (EPA) SW 846 Method 8082A

All sample results were subjected to Stage 4 data validation, which is comprised of the quality control (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): 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.
- U (Non-detected): The analyte was analyzed for and positively identified by the laboratory; however the analyte should be considered non-detected at the reported concentration due to the presence of contaminants detected in the associated blank(s).
- UJ (Non-detected estimated): The analyte was reported as not detected by the laboratory; however the reported quantitation/detection limit is estimated due to non-conformances discovered during data validation.
- R (Rejected): The sample results were rejected due to gross non-conformances discovered during data validation. Data qualified as rejected is not usable.
- 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.

### **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.

Retention time windows were established as required by the method.

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 required frequencies.

The percent differences (%D) were less than or equal to 20.0% for all analytes with the following exceptions:

Date	Standard	Column	Analyte	%D	Associated Samples	Flag	A or P
10/27/21	10272103	2C	Aroclor-1260	23.8	LDW21-SC509D	J (all detects)	A

Retention times of all analytes in the calibration standards were within the established retention time windows.

### **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/Internal Standards**

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

All internal standard percent recoveries (%R) were within QC limits with the following exceptions:

Sample	Internal Standards	%R (Limits)	Affected Analyte	Flag	A or P
LDW21-IT592B	Hexabromobiphenyl	49 (50-200)	Aroclor-1260	J (all detects)	A

### VII. Matrix Spike/Matrix Spike Duplicates

Matrix spike (MS) and matrix spike duplicate (MSD) sample analysis was performed on an associated project sample. Percent recoveries (%R) were within QC limits. Relative percent differences (RPD) were within QC limits.

### VIII. Laboratory Control Samples/Standard Reference Materials

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.

Standard reference materials (SRM) were analyzed as required by the method. The results were within QC limits.

### IX. Field Duplicates

No field duplicates were identified in this SDG.

### X. Target Analyte Quantitation

All target analyte quantitations met validation criteria.

The sample results for detected analytes from the two columns were within 40% relative percent difference (RPD) with the following exceptions:

Sample	Analyte	RPD	Flag	A or P
LDW21-SC517D	Aroclor-1248	77.8	J (all detects)	A
LDW21-SC509D	Aroclor-1248 Aroclor-1260	52.9 41.6	J (all detects) J (all detects)	A
LDW21-SC527D	Aroclor-1248	78	J (all detects)	A

### XI. Target Analyte Identification

All target analyte identifications met validation criteria.

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

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

Due to continuing calibration %D, internal standard %R, and RPD between two columns, data were qualified as estimated in four samples.

The quality control criteria reviewed, other than those discussed above, were met and are considered acceptable.



**Duwamish AOC4  
Polychlorinated Biphenyls - Data Qualification Summary - SDG 21J0131**

Sample	Analyte	Flag	A or P	Reason
LDW21-SC509D	Aroclor-1260	J (all detects)	A	Continuing calibration (%D)
LDW21-IT592B	Aroclor-1260	J (all detects)	A	Internal standards (%R)
LDW21-SC517D LDW21-SC527D	Aroclor-1248	J (all detects)	A	Target analyte quantitation (RPD between two columns)
LDW21-SC509D	Aroclor-1248 Aroclor-1260	J (all detects) J (all detects)	A	Target analyte quantitation (RPD between two columns)

**Duwamish AOC4  
Polychlorinated Biphenyls - Laboratory Blank Data Qualification Summary - SDG 21J0131**

No Sample Data Qualified in this SDG

**Duwamish AOC4  
Polychlorinated Biphenyls - Field Blank Data Qualification Summary - SDG 21J0131**

No Sample Data Qualified in this SDG

LDC #: 52703A3b  
 SDG #: 21J0131  
 Laboratory: Analytical Resources, Inc.

**VALIDATION COMPLETENESS WORKSHEET**

Stage 4

Date: 7/10/21  
 Page: 1 of 2  
 Reviewer: [Signature]  
 2nd Reviewer: JG

**METHOD:** GC Polychlorinated Biphenyls (EPA SW846 Method 8082A)

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	A	
II.	Initial calibration/ICV	A/A	REC ≤ 20% ICV ≤ 20%
III.	Continuing calibration	W	CCV ≤ 20%
IV.	Laboratory Blanks	A	
V.	Field blanks	N	
VI.	Surrogate spikes / IS	A/W	
VII.	Matrix spike/Matrix spike duplicates	A	
VIII.	Laboratory control samples / SPM	A	LOS/D
IX.	Field duplicates	N	
X.	Target analyte quantitation	W	
XI.	Target analyte identification	A	
XII.	Overall assessment of data	A	

Note: A = Acceptable      ND = No compounds detected      D = Duplicate      SB=Source blank  
 N = Not provided/applicable      R = Rinsate      TB = Trip blank      OTHER:  
 SW = See worksheet      FB = Field blank      EB = Equipment blank

	Client ID	Lab ID	Matrix	Date
1	LDW21-SC560F	21J0131-01	Sediment	06/29/21
2	LDW21-SC509D	21J0131-02	Sediment	07/01/21
3	LDW21-SC517D	21J0131-03	Sediment	07/01/21
4	LDW21-SC520A	21J0131-04	Sediment	07/02/21
5	LDW21-SC527D	21J0131-05	Sediment	07/02/21
6	LDW21-SC531D	21J0131-06	Sediment	07/02/21
7	LDW21-SC534D	21J0131-07	Sediment	07/02/21
8	LDW21-IT601	21J0131-08	Sediment	07/06/21
9	LDW21-IT592B	21J0131-09	Sediment	07/06/21
10	LDW21-IT592C	21J0131-10	Sediment	07/06/21
11	LDW21-IT592D	21J0131-11	Sediment	07/06/21
12	LDW21-IT592D	21J0131-12	Sediment	07/06/21
13	LDW21-IT592F	21J0131-13	Sediment	07/06/21
14	LDW21-IT592G	21J0131-14	Sediment	07/06/21
15	LDW21-SC510F	21J0131-15	Sediment	07/07/21
16	LDW21-IT609D	21J0131-16	Sediment	07/07/21
17	LDW21-SC595	21J0131-18	Sediment	07/08/21

LDC #: 52703A3b  
 SDG #: 21J0131  
 Laboratory: Analytical Resources, Inc.

**VALIDATION COMPLETENESS WORKSHEET**

Stage 4

Date: 7/10/21  
 Page: 2 of 2  
 Reviewer: [Signature]  
 2nd Reviewer: JG

**METHOD:** GC Polychlorinated Biphenyls (EPA SW846 Method 8082A)

	Client ID	Lab ID	Matrix	Date
18	LDW21-SC519B	21J0131-19	Sediment	07/08/21
19	LDW21-SC519C	21J0131-20	Sediment	07/08/21
20	LDW21-SC519D	21J0131-21	Sediment	07/08/21
21	LDW21-SC519E	21J0131-22	Sediment	07/08/21
22	LDW21-SC519F	21J0131-23	Sediment	07/08/21
23	LDW21-SC535D	21J0131-24	Sediment	07/08/21
24	LDW21-SC509DMS	21J0131-02MS	Sediment	07/01/21
25	LDW21-SC509DMSD	21J0131-02MSD	Sediment	07/01/21
26	LDW21-IT592CMS	21J0131-10MS	Sediment	07/06/21
27	LDW21-IT592CMSD	21J0131-10MSD	Sediment	07/06/21
28				
29				
30				

Notes:

	<u>BH0624-B&amp;K1</u>				
	<u>BH0629-B&amp;K1</u>				
	<u>BH0664-B&amp;K1</u>				

Method: GC HPLC

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

Validation Area	Yes	No	NA	Findings/Comments
<b>IX. Field duplicates</b>				
Were field duplicate pairs identified in this SDG?		/		
Were target compounds detected in the field duplicates?			/	
<b>X. Compound quantitation</b>				
Did the laboratory LOQs/RLs meet the QAPP LOQs/RLs?	/			
Were compound quantitation and RLs adjusted to reflect all sample dilutions and dry weight factors applicable to level IV validation?	/			
<b>XI. Target compound identification</b>				
Were the retention times of reported detects within the RT windows?	/			
<b>XIII. Overall assessment of data</b>				
Overall assessment of data was found to be acceptable.	/			

LDC #: 5-703A26

### VALIDATION FINDINGS WORKSHEET Continuing Calibration

Page: 1 of 1  
Reviewer: [Signature]

METHOD: GC HPLC

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

N N/A Were continuing calibration standards analyzed at the required frequencies?

Y N/A Did the continuing calibration standards meet the %D validation criteria of ≤20.0%?

**Level IV Only**

Y N/A Were the retention times for all calibrated compounds within their respective acceptance windows?

#	Date	Standard ID	Detector/ Column	Compound	%D (Limit)	RT (limit)	Associated Samples	Qualifications
	10/5/21	105T2103	20	BB	23.8	( )	2.24-25 MB (lots)	<u>N/A</u> (10 in)
						( )		
	<del>10/5/21</del>					( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		

**VALIDATION FINDINGS WORKSHEET**  
**Internal Standards**

**METHOD: GC**

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

- Y N N/A Were all internal standard area counts within -50 to +100% of the ICAL midpoint standard?  
Y N N/A Were the retention times of the internal standards within +/- 0.05 min seconds of the retention times of the ICAL midpoint standard?

#	Date	Sample ID	Internal Standard	Area (Limits)	RT (Limits)	Qualifications
		9 (2/15)	HBB (12)	49 (50-200)		✓ (H) / A (BB)

HBB - Hexabromobiphenyl

LDC #: 5703A36

**VALIDATION FINDINGS WORKSHEET**  
**Compound Quantitation and Reported CRQLs**

Page: 1 of 1  
Reviewer: [Signature]

METHOD:  GC  HPLC

**Level IV/D Only**

- Y  N  N/A Were CRQLs adjusted for sample dilutions, dry weight factors, etc.?
- Y  N  N/A Did the reported results for detected target compounds agree within 10.0% of the recalculated results?
- Y  N  N/A Did the relative percent differences of detected compounds between two columns/detectors  $\leq 40\%$ ?  
If no, please see findings below.

#	Compound Name	Sample ID	%RPD Between Two Columns/Detectors Limit ( $\leq 40\%$ )	Qualifications
	Aroclor 1248	3	27.8	↓ det 3/A
	↓			
	Aroclor 1260	2	52.9 41.6	↓
	Aroclor 1248	5	78	↓ det 3/A



LDC #: 5-70-2136

**VALIDATION FINDINGS WORKSHEET**  
**Initial Calibration Calculation Verification**

Page: 1 of 1  
Reviewer: [Signature]

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 (100 std)	CF (100 std)	Ave CF (initial)	Ave CF (initial)	%RSD	%RSD
1	KAC	8/13/21	BB-1 (1c)	0.03587713	0.03587713	0.03599233	0.0359923	2.6	2.6
			BB-1 (2c)	0.0687649	0.068764	0.06650318	0.06650318	7.7	7.8
2									
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 #: 5203A26

**VALIDATION FINDINGS WORKSHEET**  
**Continuing Calibration Results Verification**

Page: 1 of 1  
 Reviewer: 9

METHOD: GC\_HPLC

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

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

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

#	Standard ID	Calibration Date	Compound	Average CF(Ical)/ CCV Conc.	Reported	Recalculated	Reported	Recalculated
					CF/ Conc. CCV	CF/ Conc. CCV	%D	%D
1	10232103	10/23/21	BB-1 (1C)	0.0359923	0.0328101	0.032101	8.8	8.8
	1.3-10.26-27		BB-1 (2C)	0.0665032	0.0550927	0.0550927	17.2	17.2
	MP							
2	10232120	10/24/21	↓	0.0359923	0.0332095	0.0332094	7.6	7.7
	11-17			0.0665032	0.0556153	0.0556153	16.4	16.4
3	10242134	10/25/21	↓	0.0359923	0.0300082	0.0300081	16.8	16.6
	21.23			0.0665032	0.0534658	0.0534657	19.6	19.6
	MP							
4	10252103	10/25/21	↓	0.0359923	0.0299229	0.0299229	16.8	16.9
	8.18-20			0.0665032	0.0542807	0.0542806	18.4	18.4

LDC #: 50703A26

**VALIDATION FINDINGS WORKSHEET**  
**Continuing Calibration Results Verification**

Page: 2 of 2  
 Reviewer: 9

METHOD:  GC\_HPLC

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

% Difference = 100 \* (ave. CF - CF)/ave.CF

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

#	Standard ID	Calibration Date	Compound	Average CF(Ical)/ CCV Conc.	Reported	Recalculated	Reported	Recalculated
					CF/ Conc. CCV	CF/ Conc. CCV	%D	%D
1	<del>1027103</del> MB, 2, 4, 5	10/5/1 T=30	<del>BB-1 (1C)</del>	0.0359923	0.0288124	0.0288124	20.0	20.0
				0.0665032	0.0486648	0.0486647	26.8	26.8
2	<del>10282113</del> 22	10/28/1 50=27	<del>↓</del>	0.0359923	0.0286022		20.4	
				0.0665032	0.0479189		28.0	
3	10282113 22	10/28/1	BB-1 (1C)	0.0359923	0.0320725	0.0320724	10.8	10.9
				BB-1 (2C)	0.0665032	0.0540579	0.0540578	18.8
4								

**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 Difference
				Reported	Recalculated	
DCB	1c	40.0	40.1	100	100	
PMX	↓	↓	30.3	75.7	75.7	

Sample ID: \_\_\_\_\_

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

Sample ID: \_\_\_\_\_

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

## VALIDATION FINDINGS WORKSHEET

### Matrix Spike/Matrix Spike Duplicates Results Verification

METHOD: GC HPLC

The percent recoveries (%R) and relative percent differences (RPD) of the matrix spike and matrix spike duplicate were recalculated for the compounds identified below using the following calculation:

$$\% \text{Recovery} = 100 * (\text{SSC} - \text{SC}) / \text{SA}$$

Where

SSC = Spiked sample concentration

SC = Sample concentration

SA = Spike added

$$\text{RPD} = ((\text{SSCMS} - \text{SSCMSD}) * 2) / (\text{SSCMS} + \text{SSCMSD}) * 100$$

MS = Matrix spike

MSD = Matrix spike duplicate

MS/MSD samples: 26/27

Compound	Spike Added ( <u>µg/g</u> )		Sample Conc. ( <u>µg/g</u> )	Spike Sample Concentration ( <u>µg/g</u> )		Matrix spike		Matrix Spike Duplicate		MS/MSD	
	MS	MSD		MS	MSD	Percent Recovery		Percent Recovery		RPD	
						Reported	Recalc.	Reported	Recalc.	Reported	Recalc.
Gasoline (8015)											
Diesel (8015)											
Benzene (8021B)											
Methane (RSK-175)											
2,4-D (8151)											
Dinoseb (8151)											
Naphthalene (8310)											
Anthracene (8310)											
HMX (8330)											
2,4,6-Trinitrotoluene (8330)											
<del>FB</del>	101	101	20	75.9	74.8	73.4	73.2	72.3	72.1	1.48	1.46

Comments: Refer to Matrix Spike/Matrix Spike Duplicates 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**

**Laboratory Control Sample/Laboratory Control Sample Duplicates Results Verification**

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 compounds identified below using the following calculation:

$\%Recovery = 100 * (SSC - SC) / SA$

Where SSC = Spiked sample concentration  
 SA = Spike added  
 LCS = Laboratory Control Sample

SC = Sample concentration

$RPD = (((SSCLCS - SSCLCSD) * 2) / (SSCLCS + SSCLCSD)) * 100$

LCSD = Laboratory Control Sample duplicate

LCS/LCSD samples: B11067-B51 / RSD1

Compound	Spike Added ( <u>1000</u> )		Spike Sample Concentration ( <u>1000</u> )		LCS		LCSD		LCS/LCSD	
	LCS	LCSD	LCS	LCSD	Percent Recovery		Percent Recovery		RPD	
					Reported	Recalc.	Reported	Recalc.	Reported	Recalc.
Gasoline (8015)										
Diesel (8015)										
Benzene (8021B)										
Methane (RSK-175)										
2,4-D (8151)										
Dinoseb (8151)										
Naphthalene (8310)										
Anthracene (8310)										
HMX (8330)										
2,4,6-Trinitrotoluene (8330)										
<u>BB</u>	<u>101</u>	<u>101</u>	<u>80.0</u>	<u>81.2</u>	<u>81.4</u>	<u>81.1</u>	<u>83.6</u>	<u>83.4</u>	<u>2.67</u>	<u>2.65</u>

Comments: Refer to Laboratory Control Sample/Laboratory Control Sample Duplicate 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**  
**Sample Calculation Verification**

METHOD: 1 GC     HPLC

Y N N/A Were all reported results recalculated and verified for all level IV samples?  
Y N N/A Were all recalculated results for detected target compounds within 10% of the reported results?

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

Example:

Sample ID: 1 Compound Name PCB-1260-1

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

$$\text{Concentration} = \frac{(14889) (80.0)}{(430881) (0.0359923)} = 16.8$$

$$\text{concentration} = \frac{(16.8 + 16.7 + 19.1 + 14.6 + 88.1) (2.5)}{5 \times 14.39 \times 0.8684} = 16.6 \mu\text{g}$$

#	Sample ID	Compound	Reported Concentrations ( <u>μg/g</u> )	Recalculated Results Concentrations ( <u>   </u> )	Qualifications
	<u>1</u>	<u>PCB-1260</u>	<u>16.6</u>		

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

## Laboratory Data Consultants, Inc. Data Validation Report

**Project/Site Name:** Duwamish AOC4  
**LDC Report Date:** December 15, 2021  
**Parameters:** Arsenic  
**Validation Level:** Stage 4  
**Laboratory:** Analytical Resources, Inc., Tukwila, WA

**Sample Delivery Group (SDG):** 21J0131

<b>Sample Identification</b>	<b>Laboratory Sample Identification</b>	<b>Matrix</b>	<b>Collection Date</b>
LDW21-IT601	21J0131-08	Sediment	07/06/21
LDW21-IT592B	21J0131-09	Sediment	07/06/21
LDW21-IT592C	21J0131-10	Sediment	07/06/21
LDW21-IT592D	21J0131-11	Sediment	07/06/21
LDW21-IT592D	21J0131-12	Sediment	07/06/21
LDW21-IT592F	21J0131-13	Sediment	07/06/21
LDW21-IT592G	21J0131-14	Sediment	07/06/21



## 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 Final Lower Duwamish Waterway Quality Assurance Project Plan for Remedial Design of Upper Reach: Pre-Design Investigation (May 2020) and a modified outline of the USEPA National Functional Guidelines (NFG) for Inorganic Superfund Methods Data Review (January 2017). 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:

Arsenic by Environmental Protection Agency (EPA) SW 846 Method 6020B

All sample results were subjected to Stage 4 evaluation, which is comprised of the quality control (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): 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.
- U (Non-detected): The analyte was analyzed for and positively identified by the laboratory; however the analyte should be considered non-detected at the reported concentration due to the presence of contaminants detected in the associated blank(s).
- UJ (Non-detected estimated): The analyte was reported as not detected by the laboratory; however the reported quantitation/detection limit is estimated due to non-conformances discovered during data validation.
- R (Rejected): The sample results were rejected due to gross non-conformances discovered during data validation. Data qualified as rejected is not usable.
- 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.

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

All samples were received in good condition.

All technical holding time requirements were met.

## **II. ICPMS Tune**

The mass calibration was within 0.1 AMU and the percent relative standard deviation (%RSD) was less than or equal to 5%.

## **III. Instrument Calibration**

Initial and continuing calibrations were performed as required by the method.

The initial calibration verification (ICV) and continuing calibration verification (CCV) standards were within QC limits.

## **IV. ICP Interference Check Sample Analysis**

The frequency of interference check sample (ICS) analysis was met. All criteria were within QC limits.

## **V. Laboratory Blanks**

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

## **VI. Field Blanks**

No field blanks were identified in this SDG.

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

The laboratory has indicated that there were no matrix spike (MS) and matrix spike duplicate (MSD) analyses specified for the samples in this SDG, and therefore matrix spike and matrix spike duplicate analyses were not performed for this SDG.

## **VIII. Duplicate Sample Analysis**

The laboratory has indicated that there were no duplicate (DUP) analyses specified for the samples in this SDG, and therefore duplicate analyses were not performed for this SDG.

## **IX. Serial Dilution**

Serial dilution was not performed for this SDG.

## **X. Laboratory Control Samples**

Laboratory control samples (LCS) were analyzed as required by the method. Percent recoveries (%R) were within QC limits.

## **XI. Field Duplicates**

No field duplicates were identified in this SDG.

## **XII. Internal Standards (ICP-MS)**

All internal standard percent recoveries (%R) were within QC limits.

## **XIII. Target Analyte Quantitation**

All target analyte quantitations were within validation criteria.

## **XIV. Overall Assessment of Data**

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

The quality control criteria reviewed were met and are considered acceptable.

**Duwamish AOC4  
Arsenic - Data Qualification Summary - SDG 21J0131**

No Sample Data Qualified in this SDG

**Duwamish AOC4  
Arsenic - Laboratory Blank Data Qualification Summary - SDG 21J0131**

No Sample Data Qualified in this SDG

**Duwamish AOC4  
Arsenic - Field Blank Data Qualification Summary - SDG 21J0131**

No Sample Data Qualified in this SDG

LDC #: 52703A4a

**VALIDATION COMPLETENESS WORKSHEET**

SDG #: 21J0131

Stage 4

Laboratory: Analytical Resources, Inc., Tukwila, WA

Date: 10/9/21

Page: 1 of 1

Reviewer: [Signature]

2nd Reviewer: [Signature]

**METHOD:** Arsenic (EPA SW846 Method 6020B)

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	A, Δ	
II.	ICP/MS Tune	A	
III.	Instrument Calibration	A	
IV.	ICP Interference Check Sample (ICS) Analysis	A	
V.	Laboratory Blanks	A	
VI.	Field Blanks	N	
VII.	Matrix Spike/Matrix Spike Duplicates	N	
VIII.	Duplicate sample analysis	N	
IX.	Serial Dilution	N	
X.	Laboratory control samples	A	LCS
XI.	Field Duplicates	N	
XII.	Internal Standard (ICP-MS)	A	
XIII.	Target Analyte Quantitation	A	
XIV.	Overall Assessment of Data	A	

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:

	Client ID	Lab ID	Matrix	Date
1	LDW21-IT601	21J0131-08	Sediment	07/06/21
2	LDW21-IT592B	21J0131-09	Sediment	07/06/21
3	LDW21-IT592C	21J0131-10	Sediment	07/06/21
4	LDW21-IT592D	21J0131-11	Sediment	07/06/21
5	LDW21-IT592D	21J0131-12	Sediment	07/06/21
6	LDW21-IT592F	21J0131-13	Sediment	07/06/21
7	LDW21-IT592G	21J0131-14	Sediment	07/06/21
8				
9				
10				
11				
12				
13				

Notes: \_\_\_\_\_

METHOD: Trace Metals (EPA SW 846 Methods 6010/6020/7000)				
Validation Area	Yes	No	NA	Comments
<b>I. Technical holding times</b>				
Were all technical holding times met?	X			
Were all water samples preserved to a pH of <2?			X	
<b>II. ICP-MS Tune</b>				
Were mass resolutions within 0.1 amu for all isotopes in the tuning solution?	X			
Were %RSDs of isotopes in the tuning solution ≤5%?	X			
<b>III. Calibration</b>				
Were all instruments calibrated daily?	X			
Were the proper standards used?	X			
Were all initial and continuing calibration verifications within the 90-110% (80-120% for mercury) QC limits?	X			
Were the low level standard checks within 70-130%?			X	
Were all initial calibration correlation coefficients within limits as specified by the method?	X			
<b>IV. Blanks</b>				
Was a method blank associated with every sample in this SDG?	X			
Was there contamination in the method blanks?		X		
Was there contamination in the initial and continuing calibration blanks?		X		
<b>V. Interference Check Sample</b>				
Were the interference check samples performed daily?	X			
Were the AB solution recoveries within 80-120%?	X			
<b>VI. Matrix Spike/Matrix Spike Duplicates/Laboratory Duplicates</b>				
Were MS/MSD recoveries with the QC limits? (If the sample concentration exceeded the spike concentration by a factor of 4, no action was taken.)			X	
Were the MS/MSD or laboratory duplicate relative percent differences (RPDs) within the QC limits?			X	
<b>VII. Laboratory Control Samples</b>				
Was a LCS analyzed for each batch in the SDG?	X			
Were the LCS recoveries and RPDs (if applicable) within QC limits?	X			

METHOD: Trace Metals (EPA SW 846 Methods 6010/6020/7000)				
Validation Area	Yes	No	NA	Comments
<b>VIII. Internal Standards</b>				
Were all percent recoveries within the 30-120% (60-125% for EPA Method 200.8) QC limits?	X			
If the recoveries were outside the limits, was a reanalysis performed?		X		
<b>IX. Serial Dilution</b>				
Were all percent differences <10%?			X	
Was there evidence of negative interference? If yes, professional judgement will be used to qualify the data.			X	
<b>X. Sample Result Verification</b>				
Were all reporting limits adjusted to reflect sample dilutions?	X			
Were all soil samples dry weight corrected?	X			
<b>XI. Overall Assessment of Data</b>				
Was the overall assessment of the data found to be acceptable?	X			
<b>XII. Field Duplicates</b>				
Were field duplicates identified in this SDG?		X		
Were target analytes detected in the field duplicates?			X	
<b>XIII. Field Blanks</b>				
Were field blanks identified in this SDG?		X		
Were target analytes detected in the field blanks?			X	

METHOD: Trace Metals (EPA SW 846 Methods 6010/6020/7000)

An initial calibration verification (ICV), continuing calibration verification (CCV), low level calibration check (LLCC), and interference check sample (ICSAB) percent recovery (%R) was recalculated for each type of analysis using the following formula:

$$\%R = (\text{Found}/\text{True}) \times 100$$

Found = concentration of each analyte measured in the analysis

True = concentration of each analyte in the source

Standard ID	Type of Analysis	Element	Found (ug/L)	True (ug/L)	Recalculated %R	Reported %R	Acceptable (Y/N)
ICV	ICP-MS	As	47.7	50	95.4	95.5	Y
CCV	ICP-MS	As	49.8	50	99.6	99.6	Y
ICSAB	ICP-MS	As	19.283	20	96.4	96.4	Y

ICP-MS Tune	QC Parameter	Mass	Actual	Required
10/28/2021	Mass Axis	115	114.9	± 0.1 amu
10/28/2021	%RSD	115	1	≤ 5%



METHOD: Trace Metals (EPA SW 846 Methods 6010/6020/7000)

Percent recoveries (%R) for the laboratory control sample (LCS), matrix spike (MS), and post digestion spike (PDS) were recalculated using the following formula:

$$\%R = (\text{Found}/\text{True}) \times 100$$

Found = concentration of each analyte measured in the analysis. For the MS calculation, Found = SSR (Spiked Sample Result) - SR (Sample Result)

True = concentration of each analyte in the source

The sample and duplicate relative percent difference (RPD) was recalculated using the following formula:

$$\text{RPD} = (\text{Absolute value}(S-D) \times 200) / (S+D)$$

S = Original sample concentration

D = Duplicate sample concentration

The serial dilution percent difference (%D) was recalculated using the following formula.

$$\%D = (\text{Absolute value}(I - \text{SDR})) \times 100 / (I)$$

I = Initial sample result

SDR = Serial dilution result (with a 5x dilution applied)

Sample ID	Type of Analysis	Element	Found/S/I	True/D/SDR	Recalculated %R/RPD/%D	Reported %R/RPD/%D	Acceptable (Y/N)
LCS	LCS	As	24.1	25	96.4	96.5	Y
	MS						
	Duplicate						
	PDS						
	Serial dilution						



## Laboratory Data Consultants, Inc. Data Validation Report

**Project/Site Name:** Duwamish AOC4

**LDC Report Date:** December 15, 2021

**Parameters:** Wet Chemistry

**Validation Level:** Stage 4

**Laboratory:** Analytical Resources, Inc., Tukwila, WA

**Sample Delivery Group (SDG):** 21J0131

Sample Identification	Laboratory Sample Identification	Matrix	Collection Date
LDW21-SC560F	21J0131-01	Sediment	06/29/21
LDW21-SC509D	21J0131-02	Sediment	07/01/21
LDW21-SC517D	21J0131-03	Sediment	07/01/21
LDW21-SC520A	21J0131-04	Sediment	07/02/21
LDW21-SC527D	21J0131-05	Sediment	07/02/21
LDW21-SC531D	21J0131-06	Sediment	07/02/21
LDW21-SC534D	21J0131-07	Sediment	07/02/21
LDW21-IT601	21J0131-08	Sediment	07/06/21
LDW21-IT592B	21J0131-09	Sediment	07/06/21
LDW21-IT592C	21J0131-10	Sediment	07/06/21
LDW21-IT592D	21J0131-11	Sediment	07/06/21
LDW21-IT592D	21J0131-12	Sediment	07/06/21
LDW21-IT592F	21J0131-13	Sediment	07/06/21
LDW21-IT592G	21J0131-14	Sediment	07/06/21
LDW21-SC510F	21J0131-15	Sediment	07/07/21
LDW21-IT609D	21J0131-16	Sediment	07/07/21
LDW21-SC595	21J0131-18	Sediment	07/08/21
LDW21-SC519B	21J0131-19	Sediment	07/08/21
LDW21-SC519C	21J0131-20	Sediment	07/08/21
LDW21-SC519D	21J0131-21	Sediment	07/08/21
LDW21-SC519E	21J0131-22	Sediment	07/08/21
LDW21-SC519F	21J0131-23	Sediment	07/08/21
LDW21-SC535D	21J0131-24	Sediment	07/08/21
LDW21-SC560FMS	21J0131-01MS	Sediment	06/29/21
LDW21-SC560FDUP1	21J0131-01DUP1	Sediment	06/29/21
LDW21-SC560FDUP2	21J0131-01DUP2	Sediment	06/29/21

<b>Sample Identification</b>	<b>Laboratory Sample Identification</b>	<b>Matrix</b>	<b>Collection Date</b>
LDW21-SC519EMS	21J0131-22MS	Sediment	07/08/21
LDW21-SC519EDUP1	21J0131-22DUP1	Sediment	07/08/21
LDW21-SC519EDUP2	21J0131-22DUP2	Sediment	07/08/21

## 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 Final Lower Duwamish Waterway Quality Assurance Project Plan for Remedial Design of Upper Reach: Pre-Design Investigation (May 2020) and a modified outline of the USEPA National Functional Guidelines (NFG) for Inorganic Superfund Methods Data Review (January 2017). 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 methods:

Total Organic Carbon by Environmental Protection Agency (EPA) SW 846 Method 9060A

Total Solids by Standard Method 2540G

All sample results were subjected to Stage 4 data validation, which is comprised of the quality control (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): 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.
- U (Non-detected): The analyte was analyzed for and positively identified by the laboratory; however the analyte should be considered non-detected at the reported concentration due to the presence of contaminants detected in the associated blank(s).
- UJ (Non-detected estimated): The analyte was reported as not detected by the laboratory; however the reported quantitation/detection limit is estimated due to non-conformances discovered during data validation.
- R (Rejected): The sample results were rejected due to gross non-conformances discovered during data validation. Data qualified as rejected is not usable.
- 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.

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

All samples were received in good condition.

All technical holding time requirements were met.

## **II. Initial Calibration**

All criteria for the initial calibration of each method were met.

## **III. Continuing Calibration**

Continuing calibration frequency and analysis criteria were met for each method when applicable.

## **IV. Laboratory Blanks**

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

## **V. Field Blanks**

No field blanks were identified in this SDG.

## **VI. Matrix Spike/Matrix Spike Duplicates**

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

## **VII. Duplicate Sample Analysis**

Duplicate (DUP) sample analysis was performed on an associated project sample. Results were within QC limits.

## **VIII. Laboratory Control Samples**

Laboratory control samples (LCS) were analyzed as required by the methods. Percent recoveries (%R) were within QC limits.

## **IX. Field Duplicates**

No field duplicates were identified in this SDG.

## **X. Target Analyte Quantitation**

All target analyte quantitations were within validation criteria.

## **XI. Overall Assessment of Data**

The analysis was conducted within all specifications of the methods. No results were rejected in this SDG.

The quality control criteria reviewed were met and are considered acceptable.



**Duwamish AOC4  
Wet Chemistry - Data Qualification Summary - SDG 21J0131**

No Sample Data Qualified in this SDG

**Duwamish AOC4  
Wet Chemistry - Laboratory Blank Data Qualification Summary - SDG 21J0131**

No Sample Data Qualified in this SDG

**Duwamish AOC4  
Wet Chemistry - Field Blank Data Qualification Summary - SDG 21J0131**

No Sample Data Qualified in this SDG

**METHOD: (Analyte) TOC (EPA SW846 Method 9060A), Total Solids (SM2540G)**

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	A, A	
II	Initial calibration	A	
III.	Calibration verification	A	
IV	Laboratory Blanks	A	
V	Field blanks	N	
VI.	Matrix Spike/Matrix Spike Duplicates	A	
VII.	Duplicate sample analysis	A	
VIII.	Laboratory control samples	A	LCS
IX.	Field duplicates	N	
X.	Target Analyte Quantitation	A	
XI	Overall assessment of data	A	

Note: A = Acceptable      ND = No compounds detected      D = Duplicate      SB=Source blank  
 N = Not provided/applicable      R = Rinsate      TB = Trip blank      OTHER:  
 SW = See worksheet      FB = Field blank      EB = Equipment blank

	Client ID	Lab ID	Matrix	Date
1	LDW21-SC560F	21J0131-01	Sediment	06/29/21
2	LDW21-SC509D	21J0131-02	Sediment	07/01/21
3	LDW21-SC517D	21J0131-03	Sediment	07/01/21
4	LDW21-SC520A	21J0131-04	Sediment	07/02/21
5	LDW21-SC527D	21J0131-05	Sediment	07/02/21
6	LDW21-SC531D	21J0131-06	Sediment	07/02/21
7	LDW21-SC534D	21J0131-07	Sediment	07/02/21
8	LDW21-IT601	21J0131-08	Sediment	07/06/21
9	LDW21-IT592B	21J0131-09	Sediment	07/06/21
10	LDW21-IT592C	21J0131-10	Sediment	07/06/21
11	LDW21-IT592D	21J0131-11	Sediment	07/06/21
12	LDW21-IT592D	21J0131-12	Sediment	07/06/21
13	LDW21-IT592F	21J0131-13	Sediment	07/06/21
14	LDW21-IT5492G	21J0131-14	Sediment	07/06/21
15	LDW21-SC510F	21J0131-15	Sediment	07/07/21
16	LDW21-IT609D	21J0131-16	Sediment	07/07/21
17	LDW21-SC595	21J0131-18	Sediment	07/08/21

**METHOD: (Analyte) TOC (EPA SW846 Method 9060A), Total Solids (SM2540G)**

	Client ID	Lab ID	Matrix	Date
18	LDW21-SC519B	21J0131-19	Sediment	07/08/21
19	LDW21-SC519C	21J0131-20	Sediment	07/08/21
20	LDW21-SC519D	21J0131-21	Sediment	07/08/21
21	LDW21-SC519E	21J0131-22	Sediment	07/08/21
22	LDW21-SC519F	21J0131-23	Sediment	07/08/21
23	LDW21-SC535D	21J0131-24	Sediment	07/08/21
24	LDW21-SC560FMS	21J0131-01MS	Sediment	06/29/21
25	LDW21-SC560FDUP 1	21J0131-01DUP	Sediment	06/29/21
26	LDW21-SC560FTRP <u>0.02</u>	21J0131-01TRP	Sediment	06/29/21
27	LDW21-SC519EMS	21J0131-22MS	Sediment	07/08/21
28	LDW21-SC519EDUP 1	21J0131-22DUP	Sediment	07/08/21
29	LDW21-SC519ETRP <u>0.02</u>	21J0131-22TRP	Sediment	07/08/21
30				
31				
32				

Notes: \_\_\_\_\_  
 \_\_\_\_\_  
 \_\_\_\_\_

METHOD: Inorganics				
Validation Area	Yes	No	NA	Comments
<b>I. Technical holding times</b>				
Were all technical holding times were met?	X			Frozen
<b>II. Calibration</b>				
Were all instruments calibrated at the required frequency?	X			
Were the proper number of standards used?	X			
Were all initial and continuing calibration verifications within the QC limits?	X			
Were all initial calibration correlation coefficients within limits as specified by the method?	X			
Were balance checks performed as required?	X			
<b>III. Blanks</b>				
Was a method blank associated with every sample in this SDG?	X			
Was there contamination in the method blanks?		X		
Was there contamination in the initial and continuing calibration blanks?		X		
<b>IV. Matrix Spike/Matrix Spike Duplicates/Laboratory Duplicates</b>				
Were MS/MSD recoveries with the QC limits? (If the sample concentration exceeded the spike concentration by a factor of 4, no action was taken.)	X			
Were the MS/MSD or laboratory duplicate relative percent differences (RPDs) within the QC limits?	X			
<b>V. Laboratory Control Samples</b>				
Was a LCS analyzed for each batch in the SDG?	X			
Were the LCS recoveries and RPDs (if applicable) within QC limits?	X			
<b>X. Sample Result Verification</b>				
Were all reporting limits adjusted to reflect sample dilutions?	X			
Were all soil samples dry weight corrected?	X			
<b>XI. Overall Assessment of Data</b>				
Was the overall assessment of the data found to be acceptable?	X			

METHOD: Inorganics				
Validation Area	Yes	No	NA	Comments
<b>XII. Field Duplicates</b>				
Were field duplicates identified in this SDG?		X		
Were target analytes detected in the field duplicates?			X	
<b>XIII. Field Blanks</b>				
Were field blanks identified in this SDG?		X		
Were target analytes detected in the field blanks?			X	



**Validation Findings Worksheet**  
**Initial and Continuing Calibration Calculation Verification**

**Method:** Inorganics, Method See Cover

An initial or continuing calibration verification percent recovery (%R) was recalculated for each type of analysis using the following formula:

$$\%R = \frac{\text{Found} \times 100}{\text{True}}$$

Where, Found = concentration of each analyte measured in the analysis of the ICV or CCV solution  
 True = concentration of each analyte in the ICV or CCV source

Calibration verification	TOC	ICV	44.446	47.154	106	106	Y
Calibration verification	TOC	CCV	44.446	43.275	97	97	Y
Calibration verification	TOC	CCV	44.446	44.823	101	101	Y

Comments:

METHOD: Inorganics

Percent recoveries (%R) for the laboratory control sample (LCS) and matrix spike (MS) were recalculated using the following formula.

$$\%R = (\text{Found}/\text{True}) \times 100$$

Found = concentration of each analyte measured in the analysis. For the MS calculation, Found = SSR (Spiked Sample Result) - SR (Sample Result)

True = concentration of each analyte in the source

The sample and duplicate relative percent difference (RPD) was recalculated using the following formula.

$$\text{RPD} = (\text{Absolute value}(S-D) \times 200) / (S+D)$$

S = Original sample concentration

D = Duplicate sample concentration

Sample ID	Type of Analysis	Element	Found/S	True/D	Recalculated %R/RPD	Reported %R/RPD	Acceptable (Y/N)
LCS	LCS	TOC	44.1	44.4	99.3	99.3	Y
24	MS	TOC	0.89	0.882	101	101	Y
26	Duplicate	TS	84.21	83.47	0.883	0.883	Y





## Laboratory Data Consultants, Inc. Data Validation Report

**Project/Site Name:** Duwamish AOC4  
**LDC Report Date:** December 15, 2021  
**Parameters:** Polychlorinated Dioxins/Dibenzofurans  
**Validation Level:** Stage 4  
**Laboratory:** Analytical Resources, Inc., Tukwila, WA  
**Sample Delivery Group (SDG):** 21J0131

<b>Sample Identification</b>	<b>Laboratory Sample Identification</b>	<b>Matrix</b>	<b>Collection Date</b>
LDW21-IT627	21J0131-17	Sediment	07/08/21
LDW21-IT627DUP	21J0131-17DUP	Sediment	07/08/21

## 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 Final Lower Duwamish Waterway Quality Assurance Project Plan for Remedial Design of Upper Reach: Pre-Design Investigation (May 2020) and a modified outline of the USEPA National Functional Guidelines (NFG) for High Resolution Superfund Methods Data Review (April 2016). 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:

Polychlorinated Dioxins/Dibenzofurans by Environmental Protection Agency (EPA) Method 1613B

All sample results were subjected to Stage 4 data validation, which is comprised of the quality control (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): 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.
- U (Non-detected): The analyte was analyzed for and positively identified by the laboratory; however the analyte should be considered not detected at the reported concentration due to the presence of contaminants detected in the associated blank(s).
- UJ (Non-detected estimated): The analyte was reported as not detected by the laboratory; however the reported quantitation/detection limit is estimated due to non-conformances discovered during data validation.
- R (Rejected): The sample results were rejected due to gross non-conformances discovered during data validation. Data qualified as rejected is not usable.
- 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.

## **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. HRGC/HRMS Instrument Performance Check**

Instrument performance was checked at the required frequency.

Retention time windows were established for all homologues. The chromatographic resolution between 2,3,7,8-TCDD and peaks representing any other unlabeled TCDD isomer was less than or equal to 25%.

The static resolving power was at least 10,000 (10% valley definition).

## **III. Initial Calibration and Initial Calibration Verification**

A five point 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 and less than or equal to 35.0% for labeled compounds.

The ion abundance ratios for all PCDDs and PCDFs were within validation criteria.

The minimum S/N ratio was greater than or equal to 10 for each analyte and labeled compound.

The percent differences (%D) of the initial calibration verification (ICV) standard were within the QC limits for all analytes and labeled compounds.

## **IV. Continuing Calibration**

Continuing calibration was performed at the required frequencies.

All of the continuing calibration results were within the QC limits for all analytes and labeled compounds.

The ion abundance ratios for all PCDDs and PCDFs were within validation criteria.

The minimum S/N ratio was greater than or equal to 10 for each analyte and labeled compound.

## **V. Laboratory Blanks**

Laboratory blanks were analyzed as required by the method. No contaminants were found in the laboratory blanks with the following exceptions:

Blank ID	Extraction Date	Analyte	Concentration	Associated Samples
BJJ0500-BLK1	10/19/21	OCDD Total HxCDF	0.981 ng/Kg 0.100 ng/Kg	All samples in SDG 21J0131

Sample concentrations were compared to concentrations detected in the laboratory blanks. The sample concentrations were either not detected or were significantly greater than the concentrations found in the associated laboratory blanks.

## VI. Field Blanks

No field blanks were identified in this SDG.

## VII. Matrix Spike/Matrix Spike Duplicates/Duplicate Sample Analysis

The laboratory has indicated that there were no matrix spike (MS) and matrix spike duplicate (MSD) analyses specified for the samples in this SDG, and therefore matrix spike and matrix spike duplicate analyses were not performed for this SDG.

Duplicate (DUP) sample analysis was performed on an associated project sample. Results were within QC limits.

## VIII. Laboratory Control Samples/Standard Reference Materials

Laboratory control samples (LCS) were analyzed as required by the method. Percent recoveries (%R) were within QC limits.

Standard reference materials (SRM) were analyzed as required by the method. The results were within QC limits.

## IX. Field Duplicates

No field duplicates were identified in this SDG.

## X. Internal Standards

All internal standard areas and retention times were within QC limits.

## XI. Target Analyte Quantitation

All target analyte quantitations met validation criteria with the following exceptions:

Sample	Analyte	Flag	A or P
All samples in SDG 21J0131	All analytes reported as estimated maximum possible concentration (EMPC) and greater than the reporting limit (RL).	J (all detects)	A

## **XII. Target Analyte Identification**

All target analyte identifications met validation criteria.

## **XIII. System Performance**

The system performance was acceptable.

## **XIV. Overall Assessment of Data**

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

Due to results reported by the laboratory as EMPCs, data were qualified as estimated in two samples.

The quality control criteria reviewed, other than those discussed above, were met and are considered acceptable.

**Duwamish AOC4  
Polychlorinated Dioxins/Dibenzofurans - Data Qualification Summary - SDG 21J0131**

Sample	Analyte	Flag	A or P	Reason
LDW21-IT627 LDW21-IT627DUP	All analytes reported as estimated maximum possible concentration (EMPC) and greater than the reporting limit (RL).	J (all detects)	A	Target analyte quantitation (EMPC)

**Duwamish AOC4  
Polychlorinated Dioxins/Dibenzofurans - Laboratory Blank Data Qualification Summary - SDG 21J0131**

No Sample Data Qualified in this SDG

**Duwamish AOC4  
Polychlorinated Dioxins/Dibenzofurans - Field Blank Data Qualification Summary - SDG 21J0131**

No Sample Data Qualified in this SDG

LDC #: 52703A21

**VALIDATION COMPLETENESS WORKSHEET**

SDG #: 21J0131

Stage 4

Laboratory: Analytical Resources, Inc., Tukwila, WA

Date: 7/27

Page: 1 of 1

Reviewer: [Signature]

2nd Reviewer: [Signature]

**METHOD:** HRGC/HRMS Polychlorinated Dioxins/Dibenzofurans (EPA Method 1613B)

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	A	
II.	HRGC/HRMS Instrument performance check	A	
III.	Initial calibration/ICV	A/A	RSD ≤ 20/35%. CV ≤ RCLimits
IV.	Continuing calibration	A	CV ≤ RCLimits
V.	Laboratory Blanks	N	
VI.	Field blanks	N	
VII.	Matrix spike/Matrix spike duplicates / DUP	N/A	< 5% PL
VIII.	Laboratory control samples / SRM	A	2 CS
IX.	Field duplicates	N	
X.	Internal standards	A	
XI.	Target analyte quantitation	A/SW	
XII.	Target analyte identification	A	
XIII.	System performance	A	
XIV.	Overall assessment of data	A	

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:

	Client ID	Lab ID	Matrix	Date
1	LDW21-IT627	21J0131-17	Sediment	07/08/21
2	LDW21-IT627DUP	21J0131-17DUP	Sediment	07/08/21
3				
4				
5				
6				
7				
8				
9				
10				

Notes:

	21J0131-17			



**Method:** HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 1613B)

Validation Area	Yes	No	NA	Findings/Comments
<b>I. Technical holding times</b>				
All technical holding times were met.	√			
Cooler temperature criteria were met.	√			
<b>II. GC/MS Instrument performance check</b>				
Was PFK exact mass 380.9760 verified?	√			
Were the retention time windows established for all homologues?	√			
Was the chromatographic resolution between 2,3,7,8-TCDD and peaks representing any other unlabeled TCDD isomers $\leq 25\%$ ?	√			
Is the static resolving power at least 10,000 (10% valley definition)?	√			
Was the mass resolution adequately check with PFK?	√			
Was the presence of 1,2,8,9-TCDD and 1,3,4,6,8-PeCDF verified?	√			
<b>III. Initial calibration and Initial calibration verification</b>				
Was the initial calibration performed at 5 concentration levels?	√			
Were all percent relative standard deviations (%RSD) $\leq 20\%$ for unlabeled compounds and $\leq 35\%$ for unlabeled compounds?	√			
Did all calibration standards meet the Ion Abundance Ratio criteria?	√			
Was the signal to noise ratio for each target compound and labeled compound $\geq 10$ ?	√			
Was an initial calibration verification (ICV) standard analyzed after each initial calibration for each instrument?	√			
Were all ICV concentrations for the unlabeled and labeled compounds within QC limits?	√			
<b>IV. Continuing calibration</b>				
Was a continuing calibration performed at the beginning of each 12-hour period?	√			
Were all continuing calibration concentrations for the unlabeled and labeled compounds within QC limits?	√			
Did all continuing calibration standards meet the Ion Abundance Ratio criteria?	√			
<b>V. Blanks</b>				
Was a method blank associated with every sample in this SDG?	√			
Was a method blank performed for each matrix and whenever a sample extraction was performed?	√			
Was there contamination in the method blanks?	√	∅		
<b>VI. Field blanks</b>				
Were field blanks identified in this SDG?		√		
Were target compounds detected in the field blanks?			√	
<b>VII. Matrix spike/Matrix spike duplicates</b>				
Were matrix spike (MS) and matrix spike duplicate (MSD) analyzed in this SDG?		√		
Were the MS/MSD percent recoveries (%R) and the relative percent differences (RPD) within the QC limits?			√	

Validation Area	Yes	No	NA	Findings/Comments
<b>VIII. Laboratory control samples</b>				
Was an LCS analyzed per extraction batch?	√			
Were the LCS percent recoveries (%R) and relative percent difference (RPD) within the QC limits?	√			
<b>IX. Field duplicates</b>				
Were field duplicate pairs identified in this SDG?		√		
Were target compounds detected in the field duplicates?			√	
<b>X. Labeled Compounds</b>				
Were labeled compounds within QC limits?	√	0		
Was the minimum S/N ratio of all labeled compound peaks $\geq 10$ ?	√			
<b>XI. Compound quantitation</b>				
Did the laboratory LOQs/RLs meet the QAPP LOQs/RLs?	√			
Were the correct labeled compound, quantitation ion and relative response factor (RRF) used to quantitate the compound?	√			
Were compound quantitation and RLs adjusted to reflect all sample dilutions and dry weight factors applicable to level IV validation?	√			
<b>XII. Target compound identification</b>				
For 2,3,7,8 substituted congeners with associated labeled standards, were the retention times of the two quantitation peaks within -1 to 3 sec. of the RT of the labeled standard?	√			
For 2,3,7,8 substituted congeners without associated labeled standards, were the relative retention times of the two quantitation peaks within 0.005 time units of the RRT measured in the routine calibration?	√			
For non-2,3,7,8 substituted congeners, were the retention times of the two quantitation peaks within RT established in the performance check solution?	√			
Did selected ion current profile (SICP) contain all characteristic ions listed in Method 1613B, Table 8?	√			
Was the Ion Abundance Ratio for the two quantitation ions within criteria?		√		
Was the signal to noise ratio for each target compound $\geq 2.5$ and $\geq 10$ for the labeled compound?	√			
Does the maximum intensity of each specified characteristic ion coincide within $\pm 2$ seconds (includes labeled standards)?	√			
For PCDF identification, was any signal ( $S/N \geq 2.5$ , at $\pm$ seconds RT) detected in the corresponding PCDF channel?			√	
Was an acceptable lock mass recorded and monitored?	√			
<b>XIII. System performance</b>				
System performance was found to be acceptable.	√			
<b>XIV. Overall assessment of data</b>				
Overall assessment of data was found to be acceptable.	√			

## VALIDATION FINDINGS WORKSHEET

**METHOD:** HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 1613B)

A. 2,3,7,8-TCDD	F. 1,2,3,4,6,7,8-HpCDD	K. 1,2,3,4,7,8-HxCDF	P. 1,2,3,4,7,8,9-HpCDF	U. Total HpCDD
B. 1,2,3,7,8-PeCDD	G. OCDD	L. 1,2,3,6,7,8-HxCDF	Q. OCDF	V. Total TCDF
C. 1,2,3,4,7,8-HxCDD	H. 2,3,7,8-TCDF	M. 2,3,4,6,7,8-HxCDF	R. Total TCDD	W. Total PeCDF
D. 1,2,3,6,7,8-HxCDD	I. 1,2,3,7,8-PeCDF	N. 1,2,3,7,8,9-HxCDF	S. Total PeCDD	X. Total HxCDF
E. 1,2,3,7,8,9-HxCDD	J. 2,3,4,7,8-PeCDF	O. 1,2,3,4,6,7,8-HpCDF	T. Total HxCDD	Y. Total HpCDF

Notes: \_\_\_\_\_  
 \_\_\_\_\_  
 \_\_\_\_\_

## VALIDATION FINDINGS WORKSHEET

### Blanks

**METHOD:** HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 1613B)

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

- N N/A Were all samples associated with a method blank?
- N N/A Was a method blank performed for each matrix and whenever a sample extraction was performed?
- N N/A Was the method blank contaminated?

Blank extraction date: 10/19/21

Blank analysis date: 10/25/21

Associated samples: All

Conc. units: ng/kg

Compound	Blank ID	Sample Identification							
	<u>B-10500-Blank1</u>								
<u>g</u>	<u>0.981</u>								
<u>X</u>	<u>0.100</u>								

Blank extraction date: \_\_\_\_\_ Blank analysis date: \_\_\_\_\_

Conc. units: \_\_\_\_\_ Associated Samples: \_\_\_\_\_

Compound	Blank ID	Sample Identification							

CIRCLED RESULTS WERE NOT QUALIFIED. ALL RESULTS NOT CIRCLED WERE QUALIFIED BY THE FOLLOWING STATEMENT:  
 All contaminants within five times the method blank concentration were qualified as not detected, "U".

**VALIDATION FINDINGS WORKSHEET**  
**Compound Quantitation and Reported RLs**

**METHOD:** HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 1613B)

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

Y N N/A Were the correct labeled compound, quantitation ions and relative response factors (RRF) used to quantitate the compound?  
Y N N/A Compound quantitation and RLs were adjusted to reflect all sample dilutions and dry weight factors (if necessary).

#	Date	Sample ID	Finding	Associated Samples	Qualifications
		All	All compounds reported as estimated maximum possible concentration (EMPC) > RL		Jdets/A

Comments: See sample calculation verification worksheet for recalculations

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**VALIDATION FINDINGS WORKSHEET**  
**Initial Calibration Calculation Verification**

**METHOD:** HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 1613B)

The Relative Response Factor (RRF), average RRF, and percent relative standard deviation (%RSD) were recalculated for the compounds identified below using the following calculations:

RRF =  $(A_x)(C_{is}) / (A_{is})(C_x)$   
 average RRF = sum of the RRFs/number of standards  
 %RSD =  $100 * (S/X)$

$A_x$  = Area of compound,                       $A_{is}$  = Area of associated internal standard  
 $C_x$  = Concentration of compound,            $C_{is}$  = Concentration of internal standard  
 $S$  = Standard deviation of the RRFs,        $X$  = Mean of the RRFs

#	Standard ID	Calibration Date	Compound (Reference Internal Standard)	Reported	Recalculated	Reported	Recalculated	Reported	Recalculated
				RRF ( 10/50 std)	RRF ( 10/50 std)	Average RRF (initial)	Average RRF (initial)	%RSD	%RSD
1	ICAL 01	8/11/21	2,3,7,8-TCDF ( <sup>13</sup> C-2,3,7,8-TCDF)	1.0832006	1.083746	1.107593	1.107593	3.6	3.6
			2,3,7,8-TCDD ( <sup>13</sup> C-2,3,7,8-TCDD)	0.9085186	0.908390	0.9202875	0.9202874	3.1	3.1
			1,2,3,6,7,8-HxCDD ( <sup>13</sup> C-1,2,3,6,7,8-HxCDD)	1.005616	1.005605	1.00898	1.00898	1.0	1.0
			1,2,3,4,6,7,8-HpCDD ( <sup>13</sup> C-1,2,4,6,7,8,-HpCDD)	1.051009	1.051062	1.068088	1.068088	6.6	6.6
			OCDF ( <sup>13</sup> C-OCDF)	1.440564	1.44059	1.44690	1.44690	5.7	5.7
2			2,3,7,8-TCDF ( <sup>13</sup> C-2,3,7,8-TCDF)						
			2,3,7,8-TCDD ( <sup>13</sup> C-2,3,7,8-TCDD)						
			1,2,3,6,7,8-HxCDD ( <sup>13</sup> C-1,2,3,6,7,8-HxCDD)						
			1,2,3,4,6,7,8-HpCDD ( <sup>13</sup> C-1,2,4,6,7,8,-HpCDD)						
			OCDF ( <sup>13</sup> C-OCDD)						
3			2,3,7,8-TCDF ( <sup>13</sup> C-2,3,7,8-TCDF)						
			2,3,7,8-TCDD ( <sup>13</sup> C-2,3,7,8-TCDD)						
			1,2,3,6,7,8-HxCDD ( <sup>13</sup> C-1,2,3,6,7,8-HxCDD)						
			1,2,3,4,6,7,8-HpCDD ( <sup>13</sup> C-1,2,4,6,7,8,-HpCDD)						
			OCDF ( <sup>13</sup> C-OCDD)						

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.

## VALIDATION FINDINGS WORKSHEET Continuing Calibration Results Verification

**METHOD:** HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 1613B)

The percent difference (%D) of the initial calibration average Relative Response Factors (RRFs) and the continuing calibration RRFs were recalculated for the compounds identified below using the following calculation:

% Difference = 100 \* (ave. RRF - RRF)/ave. RRF  
 $RRF = (A_x)(C_{is}) / (A_{is})(C_x)$

Where: ave. RRF = initial calibration average RRF  
 RRF = continuing calibration RRF  
 $A_x$  = Area of compound,  $A_{is}$  = Area of associated internal standard  
 $C_x$  = Concentration of compound,  $C_{is}$  = Concentration of internal standard

#	Standard ID	Calibration Date	Compound (Reference Internal Standard)	Average RRF (initial)	Reported	Recalculated	Reported	Recalculated
					Conc (CC)	Conc (CC)	%D	%D
1	21102505A	10/5/21	2,3,7,8-TCDF ( <sup>13</sup> C-2,3,7,8-TCDF)	1.107593	1.0745550	1.0746175	3.0	3.0
			2,3,7,8-TCDD ( <sup>13</sup> C-2,3,7,8-TCDD)	0.9202875	1.0081390	1.0081532	9.5	9.5
			1,2,3,6,7,8-HxCDD ( <sup>13</sup> C-1,2,3,6,7,8-HxCDD)	1.00898	1.0688370	1.0683744	5.9	5.9
			1,2,3,4,6,7,8-HpCDD ( <sup>13</sup> C-1,2,4,6,7,8,-HpCDD)	1.068088	1.1679010	1.1678182	9.3	9.3
			OCDF ( <sup>13</sup> C-OCDF)	1.44690	1.3382880	1.338548	7.5	7.5
2			2,3,7,8-TCDF ( <sup>13</sup> C-2,3,7,8-TCDF)					
			2,3,7,8-TCDD ( <sup>13</sup> C-2,3,7,8-TCDD)					
			1,2,3,6,7,8-HxCDD ( <sup>13</sup> C-1,2,3,6,7,8-HxCDD)					
			1,2,3,4,6,7,8-HpCDD ( <sup>13</sup> C-1,2,4,6,7,8,-HpCDD)					
			OCDF ( <sup>13</sup> C-OCDF)					
3			2,3,7,8-TCDF ( <sup>13</sup> C-2,3,7,8-TCDF)					
			2,3,7,8-TCDD ( <sup>13</sup> C-2,3,7,8-TCDD)					
			1,2,3,6,7,8-HxCDD ( <sup>13</sup> C-1,2,3,6,7,8-HxCDD)					
			1,2,3,4,6,7,8-HpCDD ( <sup>13</sup> C-1,2,4,6,7,8,-HpCDD)					
			OCDF ( <sup>13</sup> C-OCDF)					

Comments: Refer to Routine 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 Laboratory Control Sample Results Verification

**METHOD:** HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 1613B)

The percent recoveries (%R) and Relative Percent Difference (RPD) of the laboratory control sample and laboratory control sample duplicate (if applicable) were recalculated for the compounds identified below using the following calculation:

% Recovery =  $100 * \frac{SSC}{SA}$       Where: SSC = Spiked sample concentration  
SA = Spike added

RPD =  $\frac{|LCS - LCSD|}{LCS + LCSD} * 2$       LCS = Laboratory control sample percent recovery      LCSD = Laboratory control sample duplicate percent recovery

LCS ID: BN0500-F35

Compound	Spike Added (NSA)		Spiked Sample Concentration (NSA)		LCS		LCSD		LCS/LCSD	
	LCS	LCSD	LCS	LCSD	Percent Recovery		Percent Recovery		RPD	
					Reported	Recalc	Reported	Recalc	Reported	Recalculated
2,3,7,8-TCDD	20.0	NA	21.0	NA	105	105				
1,2,3,7,8-PeCDD	100	↓	107	↓	107	107				
1,2,3,4,7,8-HxCDD	↓	↓	99.2	↓	99.2	99.2				
1,2,3,4,7,8,9-HpCDF	↓	↓	95.9	↓	95.9	95.9				
OCDF	200	↓	151	↓	75.5	75.5				

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





**Laboratory Data Consultants, Inc.  
Data Validation Report**

**Project/Site Name:** Duwamish AOC4  
**LDC Report Date:** December 15, 2021  
**Parameters:** Semivolatiles  
**Validation Level:** Stage 4  
**Laboratory:** Analytical Resources, Inc., Tukwila, WA  
**Sample Delivery Group (SDG):** 21J0134

<b>Sample Identification</b>	<b>Laboratory Sample Identification</b>	<b>Matrix</b>	<b>Collection Date</b>
LDW21-SS600	21J0134-11	Sediment	07/12/21
LDW21-SS681	21J0134-12	Sediment	07/12/21
LDW21-SC587A	21J0134-14	Sediment	07/12/21
LDW21-SC587F	21J0134-15	Sediment	07/12/21
LDW21-SS600MS	21J0134-11MS	Sediment	07/12/21
LDW21-SS600MSD	21J0134-11MSD	Sediment	07/12/21

## 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 Final Lower Duwamish Waterway Quality Assurance Project Plan for Remedial Design of Upper Reach: Pre-Design Investigation (May 2020) and a modified outline of the USEPA National Functional Guidelines (NFG) for Organic Superfund Methods Data Review (January 2017). 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:

Semivolatile Organic Compounds (SVOCs) by Environmental Protection Agency (EPA) SW 846 Method 8270E

All sample results were subjected to Stage 4 data validation, which is comprised of the quality control (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): 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.
- U (Non-detected): The analyte was analyzed for and positively identified by the laboratory; however the analyte should be considered non-detected at the reported concentration due to the presence of contaminants detected in the associated blank(s).
- UJ (Non-detected estimated): The analyte was reported as not detected by the laboratory; however the reported quantitation/detection limit is estimated due to non-conformances discovered during data validation.
- R (Rejected): The sample results were rejected due to gross non-conformances discovered during data validation. Data qualified as rejected is not usable.
- 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.

## 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. GC/MS Instrument Performance Check

A decafluorotriphenylphosphine (DFTPP) tune was performed at 12 hour intervals.

All ion abundance requirements were met.

## III. Initial Calibration and Initial Calibration Verification

An initial calibration was performed as required by the method.

For analytes where average relative response factors (RRFs) were utilized, percent relative standard deviations (%RSD) were less than or equal to 20.0%.

In the case where the laboratory used a calibration curve to evaluate the analytes, all coefficients of determination ( $r^2$ ) were greater than or equal to 0.990.

Average relative response factors (RRF) for all analytes were within validation criteria.

The percent differences (%D) of the initial calibration verification (ICV) standard were less than or equal to 30.0% for all analytes.

## IV. 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 with the following exceptions:

Date	Analyte	%D	Associated Samples	Flag	A or P
10/30/21	Butylbenzylphthalate	25.7	LDW21-SS600 LDW21-SC587A LDW21-SC587F	J (all detects) UJ (all non-detects)	A

All of the continuing calibration relative response factors (RRF) were within validation criteria.

## V. Laboratory Blanks

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

## VI. Field Blanks

No field blanks were identified in this SDG.

## VII. Surrogates

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

## VIII. Matrix Spike/Matrix Spike Duplicates

Matrix spike (MS) and matrix spike duplicate (MSD) sample analysis was performed on an associated project sample. Percent recoveries (%R) were within QC limits with the following exceptions:

Spike ID (Associated Samples)	Analyte	MS (%R) (Limits)	MSD (%R) (Limits)	Flag	A or P
LDW21-SS600MS/MSD (LDW21-SS600)	Phenanthrene	122 (49-120)	-	J (all detects)	A
	Fluoranthene	174 (53-145)	-	J (all detects)	
	Pyrene	160 (52-134)	-	J (all detects)	
	Butylbenzylphthalate	204 (45-132)	-	J (all detects)	
	Chrysene	131 (47-120)	-	J (all detects)	

Relative percent differences (RPD) were within QC limits with the following exceptions:

Spike ID (Associated Samples)	Analyte	RPD (Limits)	Flag	A or P
LDW21-SS600MS/MSD (LDW21-SS600)	Phenanthrene	44.0 (≤35)	J (all detects)	A
	Anthracene	35.1 (≤35)	J (all detects)	
	Fluoranthene	65.2 (≤35)	J (all detects)	
	Pyrene	62.7 (≤35)	J (all detects)	
	Butylbenzylphthalate	69.0 (≤35)	J (all detects)	
	Benzo(a)anthracene	47.9 (≤35)	J (all detects)	
	Chrysene	38.2 (≤35)	J (all detects)	
	Benzo(a)pyrene	39.5 (≤35)	J (all detects)	
	Benzo(a)fluoranthene, total	38.6 (≤35)	J (all detects)	

## IX. Laboratory Control Samples/Standard Reference Materials

Laboratory control samples (LCS) were analyzed as required by the method. Percent recoveries (%R) were within QC limits.

Standard reference materials (SRM) were analyzed as required by the method. The results were within QC limits with the following exceptions:

SRM ID	Analyte	%R (Limits)	Associated Samples	Flag	A or P
BJJ0826-SRM1	Naphthalene 2-Methylnaphthalene Acenaphthylene Acenaphthene	14.2 (41-159) 26.5 (51-149) 41.6 (57-142) 48.2 (59-141)	LDW21-SS600 LDW21-SS681	J (all detects) UJ (all non-detects)	A

## X. Field Duplicates

No field duplicates were identified in this SDG.

## XI. Internal Standards

All internal standard areas and retention times were within QC limits.

## XII. Target Analyte Quantitation

All target analyte quantitations were within validation criteria.

## XIII. Target Analyte Identification

All target analyte identifications were within validation criteria.

## XIV. System Performance

The system performance was acceptable.

## XV. Overall Assessment of Data

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

Due to continuing calibration %D, MS/MSD %R and RPD, and SRM %R, data were qualified as estimated in four samples.

The quality control criteria reviewed, other than those discussed above, were met and are considered acceptable.

**Duwamish AOC4  
Semivolatiles - Data Qualification Summary - SDG 21J0134**

Sample	Analyte	Flag	A or P	Reason
LDW21-SS600 LDW21-SC587A LDW21-SC587F	Butylbenzylphthalate	J (all detects) UJ (all non-detects)	A	Continuing calibration (%D)
LDW21-SS600	Phenanthrene Fluoranthene Pyrene Butylbenzylphthalate Chrysene	J (all detects) J (all detects) J (all detects) J (all detects) J (all detects)	A	Matrix Spike/Matrix Spike Duplicates (%R)
LDW21-SS600	Phenanthrene Anthracene Fluoranthene Pyrene Butylbenzylphthalate Benzo(a)anthracene Chrysene Benzo(a)pyrene Benzofluoranthenes, total	J (all detects) J (all detects) J (all detects) J (all detects) J (all detects) J (all detects) J (all detects) J (all detects) J (all detects) J (all detects)	A	Matrix Spike/Matrix Spike Duplicates (RPD)
LDW21-SS600 LDW21-SS681	Naphthalene 2-Methylnaphthalene Acenaphthylene Acenaphthene	J (all detects) UJ (all non-detects)	A	Standard reference materials (%R)

**Duwamish AOC4  
Semivolatiles - Laboratory Blank Data Qualification Summary - SDG 21J0134**

No Sample Data Qualified in this SDG

**Duwamish AOC4  
Semivolatiles - Field Blank Data Qualification Summary - SDG 21J0134**

No Sample Data Qualified in this SDG

LDC #: 52703B2a

**VALIDATION COMPLETENESS WORKSHEET**

Date: 12/10/21

SDG #: 21J0134

Stage 4

Page: 1 of 1

Laboratory: Analytical Resources, Inc., Tukwila, WA

Reviewer: [Signature]

2nd Reviewer: [Signature]

**METHOD:** GC/MS Semivolatiles (EPA SW 846 Method 8270E)

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	A	
II.	GC/MS Instrument performance check	A	
III.	Initial calibration/ICV	A/A	RSD ≤ 20%, Y <sup>2</sup> R <sup>2</sup> ≤ 30%
IV.	Continuing calibration	W	CCV ≤ 20%
V.	Laboratory Blanks	A	
VI.	Field blanks	N	
VII.	Surrogate spikes	A	
VIII.	Matrix spike/Matrix spike duplicates	W	
IX.	Laboratory control samples /SRM	W	LCS
X.	Field duplicates	N	
XI.	Internal standards	A	
XII.	Target analyte quantitation	A	
XIII.	Target analyte identification	A	
XIV.	System performance	A	
XV.	Overall assessment of data	A	

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:

	Client ID	Lab ID	Matrix	Date
1	LDW21-SS600	21J0134-11	Sediment	07/12/21
2	LDW21-SS681	21J0134-12	Sediment	07/12/21
3	LDW21-SC587A	21J0134-14	Sediment	07/12/21
4	LDW21-SC587F	21J0134-15	Sediment	07/12/21
5	LDW21-SS600MS	21J0134-11MS	Sediment	07/12/21
6	LDW21-SS600MSD	21J0134-11MSD	Sediment	07/12/21
7				
8				
9				

Notes:

B110826-BK1				



**Method: Semivolatiles (EPA SW 846 Method 8270D)**

Validation Area	Yes	No	NA	Findings/Comments
<b>I. Technical holding times</b>				
Were all technical holding times met?	/			
Was cooler temperature criteria met?	/			
<b>II. GC/MS Instrument performance check</b>				
Were the DFTPP performance results reviewed and found to be within the specified criteria?	/			
Were all samples analyzed within the 12 hour clock criteria?	/			
<b>IIIa. Initial calibration</b>				
Did the laboratory perform a 5 point calibration prior to sample analysis?	/			
Were all percent relative standard deviations (%RSD) ≤ 20% and relative response factors (RRF) within method criteria?	/			
Was a curve fit used for evaluation? If yes, did the initial calibration meet the curve fit acceptance criteria of > 0.990?	/			
<b>IIIb. Initial Calibration Verification</b>				
Was an initial calibration verification standard analyzed after each initial calibration for each instrument?	/			
Were all percent differences (%D) ≤ 30%?	/			
<b>IV. Continuing calibration</b>				
Was a continuing calibration standard analyzed at least once every 12 hours for each instrument?	/			
Were all percent differences (%D) ≤ 20% and relative response factors (RRF) within method criteria?		/		
<b>V. Laboratory Blanks</b>				
Was a laboratory blank associated with every sample in this SDG?	/			
Was a laboratory blank analyzed at least once every 12 hours for each matrix and concentration?	/			
Was there contamination in the laboratory blanks? If yes, please see the blanks validation findings worksheet.		/		
<b>VI. Field blanks</b>				
Were field blanks were identified in this SDG?		/		
Were target compounds detected in the field blanks?			/	
<b>VII. Surrogate spikes</b>				
Were all surrogate percent recovery (%R) within QC limits?	/			
If 2 or more base neutral or acid surrogates were outside QC limits, was a reanalysis performed to confirm %R?			/	
If any percent recoveries (%R) was less than 10%, was a reanalysis performed to confirm %R ?			/	
<b>VIII. Matrix spike/Matrix spike duplicates</b>				
Were matrix spike (MS) and matrix spike duplicate (MSD) analyzed in this SDG?	/			

Validation Area	Yes	No	NA	Findings/Comments
Were the MS/MSD percent recoveries (%R) and the relative percent differences (RPD) within the QC limits?		<input checked="" type="checkbox"/>		
<b>IX. Laboratory control samples</b>				
Was an LCS analyzed per extraction batch?	<input checked="" type="checkbox"/>			
Were the LCS percent recoveries (%R) and relative percent difference (RPD) within the QC limits?		<input checked="" type="checkbox"/>		
<b>X. Field duplicates</b>				
Were field duplicate pairs identified in this SDG?		<input checked="" type="checkbox"/>		
Were target compounds detected in the field duplicates?			<input checked="" type="checkbox"/>	
<b>XI. Internal standards</b>				
Were internal standard area counts within -50% to +100% of the associated calibration standard?	<input checked="" type="checkbox"/>			
Were retention times within + 30 seconds of the associated calibration standard?	<input checked="" type="checkbox"/>			
<b>XII. Compound quantitation</b>				
Did the laboratory LOQs/RLs meet the QAPP LOQs/RLs?	<input checked="" type="checkbox"/>			
Were the correct internal standard (IS), quantitation ion and relative response factor (RRF) used to quantitate the compound?	<input checked="" type="checkbox"/>			
Were compound quantitation and RLs adjusted to reflect all sample dilutions and dry weight factors applicable to level IV validation?	<input checked="" type="checkbox"/>			
<b>XIII. Target compound identification</b>				
Were relative retention times (RRT's) within + 0.06 RRT units of the standard?	<input checked="" type="checkbox"/>			
Did compound spectra meet specified EPA "Functional Guidelines" criteria?	<input checked="" type="checkbox"/>			
Were chromatogram peaks verified and accounted for?	<input checked="" type="checkbox"/>			
<b>XIV. System performance</b>				
System performance was found to be acceptable.	<input checked="" type="checkbox"/>			
<b>XV. Overall assessment of data</b>				
Overall assessment of data was found to be acceptable.	<input checked="" type="checkbox"/>			

## VALIDATION FINDINGS WORKSHEET

**METHOD: GC/MS SVOA**

A. Phenol	CC. Dimethylphthalate	EEE. Bis(2-ethylhexyl)phthalate	GGGG. C30-Hopane	I1. Methyl methanesulfonate
B. Bis (2-chloroethyl) ether	DD. Acenaphthylene	FFF. Di-n-octylphthalate	HHHH. 1-Methylphenanthrene	J1. Ethyl methanesulfonate
C. 2-Chlorophenol	EE. 2,6-Dinitrotoluene	GGG. Benzo(b)fluoranthene	IIII. 1,4-Dioxane	K1. o,o',o''-Triethylphosphorothioate
D. 1,3-Dichlorobenzene	FF. 3-Nitroaniline	HHH. Benzo(k)fluoranthene	JJJJ. Acetophenone	L1. n-Phenylene diamine
E. 1,4-Dichlorobenzene	GG. Acenaphthene	III. Benzo(a)pyrene	KKKK. Atrazine	M1. 1,4-Naphthoquinone
F. 1,2-Dichlorobenzene	HH. 2,4-Dinitrophenol	JJJ. Indeno(1,2,3-cd)pyrene	LLLL. Benzaldehyde	N1. N-Nitro-o-toluidine
G. 2-Methylphenol	II. 4-Nitrophenol	KKK. Dibenz(a,h)anthracene	MMMM. Caprolactam	O1. 1,3,5-Trinitrobenzene
H. 2,2'-Oxybis(1-chloropropane)	JJ. Dibenzofuran	LLL. Benzo(g,h,i)perylene	NNNN. 2,6-Dichlorophenol	P1. Pentachlorobenzene
I. 4-Methylphenol	KK. 2,4-Dinitrotoluene	MMM. Bis(2-Chloroisopropyl)ether	OOOO. 1,2-Diphenylhydrazine	Q1. 4-Aminobiphenyl
J. N-Nitroso-di-n-propylamine	LL. Diethylphthalate	NNN. Aniline	PPPP. 3-Methylphenol	R1. 2-Naphthylamine
K. Hexachloroethane	MM. 4-Chlorophenyl-phenyl ether	OOO. N-Nitrosodimethylamine	QQQQ. 3&4-Methylphenol	S1. Triphenylene
L. Nitrobenzene	NN. Fluorene	PPP. Benzoic Acid	RRRR. 4-Dimethyldibenzothiophene (4MDT)	T1. Octachlorostyrene
M. Isophorone	OO. 4-Nitroaniline	QQQ. Benzyl alcohol	SSSS. 2/3-Dimethyldibenzothiophene (4MDT)	U1. Famphur
N. 2-Nitrophenol	PP. 4,6-Dinitro-2-methylphenol	RRR. Pyridine	TTTT. 1-Methyldibenzothiophene (1MDT)	V1. 1,4-phenylenediamine
O. 2,4-Dimethylphenol	QQ. N-Nitrosodiphenylamine	SSS. Benzidine	UUUU.. 2,3,4,6-Tetrachlorophenol	W1. Methapyriene
P. Bis(2-chloroethoxy)methane	RR. 4-Bromophenyl-phenylether	TTT. 1-Methylnaphthalene	VVVV. 1,2,4,5-Tetrachlorobenzene	X1. Pentachloroethane
Q. 2,4-Dichlorophenol	SS. Hexachlorobenzene	UUU. Benzo(b)thiophene	WWWW.. 2-Picoline	Y1. 3,3'-Dimethylbenzidine
R. 1,2,4-Trichlorobenzene	TT. Pentachlorophenol	VVV. Benzonaphthothiophene	XXXX. 3-Methylcholanthrene	Z1. o-Toluidine
S. Naphthalene	UU. Phenanthrene	WWW. Benzo(e)pyrene	YYYY. a,a-Dimethylphenethylamine	A2. 1-Naphthylamine
T. 4-Chloroaniline	VV. Anthracene	XXX. 2,6-Dimethylnaphthalene	ZZZZ. Hexachloropropene	B2. 4-Aminobiphenyl
U. Hexachlorobutadiene	WW. Carbazole	YYY. 2,3,5-Trimethylnaphthalene	A1. N-Nitrosodiethylamine	C2. 4-Nitroquinoline-1-oxide
V. 4-Chloro-3-methylphenol	XX. Di-n-butylphthalate	ZZZ. Perylene	B1. N-Nitrosodi-n-butylamine	D2. Hexachloropene
W. 2-Methylnaphthalene	YY. Fluoranthene	AAAA. Dibenzothiophene	C1. N-Nitrosomethylethylamine	E2. Bis (2-chloro-1-methylethyl) ether
X. Hexachlorocyclopentadiene	ZZ. Pyrene	BBBB. Benzo(a)fluoranthene	D1. N-Nitrosomorpholine	F2. Bifenthrin
Y. 2,4,6-Trichlorophenol	AAA. Butylbenzylphthalate	CCCC. Benzo(b)fluorene	E1. N-Nitrosopyrrolidine	G2. Cyfluthrin
Z. 2,4,5-Trichlorophenol	BBB. 3,3'-Dichlorobenzidine	DDDD. cis/trans-Decalin	F1. Phenacetin	H2. Cypermethrin
AA. 2-Chloronaphthalene	CCC. Benzo(a)anthracene	EEEE. 1,1'-Biphenyl	G1. 2-Acetylaminofluorene	I2. Permethrin (cis/trans)
BB. 2-Nitroaniline	DDD. Chrysene	FFFF. Retene	H1. Pronamide	J2. 5-Nitro-o-toluidine

**VALIDATION FINDINGS WORKSHEET**  
**Continuing Calibration**

**METHOD:** GC/MS BNA (EPA SW 846 Method 8270D)

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

N N/A Was a continuing calibration standard analyzed at least once every 12 hours for each instrument?

N N/A Were percent differences (%D) ≤ 20 % and relative response factors (RRF) within the method criteria?

#	Date	Standard ID	Compound	Finding %D (Limit: <20.0%)	Finding RRF (Limit)	Associated Samples	Qualifications
	10/30/21	NT1021103002	SA	25.7		1. 3-4. 5-6. ND (213 + ND)	✓/N/A

**VALIDATION FINDINGS WORKSHEET**  
**Matrix Spike/Matrix Spike Duplicates**

**METHOD:** GC/MS BNA (EPA SW 846 Method 8270D)

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

- N N/A Were a matrix spike (MS) and matrix spike duplicate (MSD) analyzed for each matrix in this SDG? If no, indicate which matrix does not have an associated MS/MSD. Soil / Water.
- N N/A Was a MS/MSD analyzed every 20 samples of each matrix?
- N N/A Were the MS/MSD percent recoveries (%R) and the relative percent differences (RPD) within the QC limits?

#	MS/MSD ID	Compound	MS %R (Limits)	MSD %R (Limits)	RPD (Limits)	Associated Samples	Qualifications
	<u>5/6</u>	<u>UU</u>	<u>122 (49-120)</u>	( )	( )	<u>(ND) (ND)</u>	<u>ND/A</u>
		<u>YY</u>	<u>174 (53-145)</u>	( )	( )		
		<u>ZZ</u>	<u>160 (52-134)</u>	( )	( )		
		<u>AAA</u>	<u>204 (45-132)</u>	( )	( )		
		<u>DDD</u>	<u>131 (47-120)</u>	( )	( )		
		<u>UU</u>	( )	( )	<u>44.0 (≤ 35)</u>		
		<u>VV</u>	( )	( )	<u>75.1 ( )</u>		
		<u>YY</u>	( )	( )	<u>65.2 ( )</u>		
		<u>ZZ</u>	( )	( )	<u>63.7 ( )</u>		
		<u>AAA</u>	( )	( )	<u>69.0 ( )</u>		
		<u>CCC</u>	( )	( )	<u>47.9 ( )</u>		
		<u>DDD</u>	( )	( )	<u>38.2 ( )</u>		
		<u>III</u>	( )	( )	<u>39.5 ( )</u>		
		<u>Benzofluoranthenes, Total</u>	( )	( )	<u>38.6 ( )</u>		<u>✓</u>
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
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			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		

## VALIDATION FINDINGS WORKSHEET

### Laboratory Control Samples (LCS)

METHOD: GC/MS BNA (EPA SW 846 Method 8270D)

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

- N N/A Was a LCS required?  
 N N/A Were the LCS/LCSD percent recoveries (%R) and the relative percent differences (RPD) within the QC limits?

#	Date	LCS/LCSD ID	Compound	LCS %R (Limits)	LCSD %R (Limits)	RPD (Limits)	Associated Samples	Qualifications
		R10726-SR11	S	14.2 (41-159)	( )	( )	All (lots + N/D)	↓/U/A
			W	26.5 (51-149)	( )	( )		↓
			DB	41.6 (57-142)	( )	( )		
			GF	48.2 (59-141)	( )	( )		
				( )	( )	( )		
				( )	( )	( )		
				( )	( )	( )		
				( )	( )	( )		
				( )	( )	( )		
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				( )	( )	( )		
				( )	( )	( )		

## VALIDATION FINDINGS WORKSHEET Initial Calibration Calculation Verification

**METHOD:** GC/MS BNA (EPA SW 846 Method 8270D)

The Relative Response Factor (RRF), average RRF, and percent relative standard deviation (%RSD) were recalculated for the compounds identified below using the following calculations:

RRF =  $(A_x)(C_{is}) / (A_{is})(C_x)$   
 average RRF = sum of the RRFs/number of standards  
 %RSD =  $100 * (S/X)$

$A_x$  = Area of compound,  
 $C_x$  = Concentration of compound,  
 $S$  = Standard deviation of the RRFs,  
 $A_{is}$  = Area of associated internal standard  
 $C_{is}$  = Concentration of internal standard  
 $X$  = Mean of the RRFs

#	Standard ID	Calibration Date	Compound (Reference Internal Standard)	Reported	Recalculated	Reported	Recalculated	Reported	Recalculated
				RRF (5 std)	RRF (5 std)	Average RRF (initial)	Average RRF (initial)	%RSD	%RSD
1	KAC	10/25/21	Phenol (1st internal standard)	1.964599	1.964599	2.156898	2.156898	10.1	10.1
			Naphthalene (2nd internal standard)	1.135424	1.135424	1.143444	1.143444	2.1	2.1
			Fluorene (3rd internal standard)	2.097977	2.097977	2.077259	2.077259	7.8	7.8
			Pentachlorophenol (4th internal standard) UU	1.093528	1.093528	1.103879	1.103879	3.1	3.1
			Bis(2-ethylhexyl)phthalate (5th internal standard) AAA	0.913884	0.913884	0.9488406	0.9488406	5.3	5.3
			Benzo(a)pyrene (6th internal standard)	1.452654	1.452654	1.473196	1.473196	2.1	2.1
2			Phenol (1st internal standard)						
			Naphthalene (2nd internal standard)						
			Fluorene (3rd internal standard)						
			Pentachlorophenol (4th internal standard)						
			Bis(2-ethylhexyl)phthalate (5th internal standard)						
			Benzo(a)pyrene (6th internal standard)						
3			Phenol (1st internal standard)						
			Naphthalene (2nd internal standard)						
			Fluorene (3rd internal standard)						
			Pentachlorophenol (4th internal standard)						
			Bis(2-ethylhexyl)phthalate (5th internal standard)						
			Benzo(a)pyrene (6th internal standard)						

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.

## VALIDATION FINDINGS WORKSHEET Continuing Calibration Results Verification

**METHOD:** GC/MS BNA (EPA SW 846 Method 8270D)

The percent difference (%D) of the initial calibration average Relative Response Factors (RRFs) and the continuing calibration RRFs were recalculated for the compounds identified below using the following calculation:

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

$$\text{RRF} = (A_x)(C_{is}) / (A_{is})(C_x)$$

Where: ave. RRF = initial calibration average RRF  
RRF = continuing calibration RRF  
A<sub>x</sub> = Area of compound,                      A<sub>is</sub> = Area of associated internal standard  
C<sub>x</sub> = Concentration of compound,        C<sub>is</sub> = Concentration of internal standard

#	Standard ID	Calibration Date	Compound (Reference Internal Standard)	Average RRF (initial)	Reported	Recalculated	Reported	Recalculated
					RRF (CC)	RRF (CC)	%D	%D
1	<u>NT102110302</u>	<u>10/30/21</u>	Phenol (1st internal standard)	<u>2.156898</u>	<u>2.2397960</u>	<u>2.2397962</u>	<u>3.8</u>	<u>3.8</u>
			Naphthalene (2nd internal standard)	<u>1.143444</u>	<u>1.1092970</u>	<u>1.1092967</u>	<u>3.0</u>	<u>3.0</u>
			Fluorene (3rd internal standard)	<u>2.077259</u>	<u>2.0667570</u>	<u>2.0667568</u>	<u>0.5</u>	<u>0.5</u>
			Pentachlorophenol (4th internal standard) <u>UV</u>	<u>1.103879</u>	<u>1.1237720</u>	<u>1.123772</u>	<u>1.8</u>	<u>1.8</u>
			Bis(2-ethylhexyl)phthalate (5th internal standard) <u>AAA</u>	<u>0.9488406</u>	<u>1.1937350</u>	<u>1.1937348</u>	<u>25.7</u>	<u>25.7</u>
			Benzo(a)pyrene (6th internal standard)	<u>1.473196</u>	<u>1.2984050</u>	<u>1.2984052</u>	<u>11.9</u>	<u>11.9</u>
2			Phenol (1st internal standard)					
			Naphthalene (2nd internal standard)					
			Fluorene (3rd internal standard)					
			Pentachlorophenol (4th internal standard)					
			Bis(2-ethylhexyl)phthalate (5th internal standard)					
			Benzo(a)pyrene (6th internal standard)					
3			Phenol (1st internal standard)					
			Naphthalene (2nd internal standard)					
			Fluorene (3rd internal standard)					
			Pentachlorophenol (4th internal standard)					
			Bis(2-ethylhexyl)phthalate (5th internal standard)					
			Benzo(a)pyrene (6th internal standard)					

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/MS Semivolatiles (EPA SW 846 Method 8270D)

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 Spiked	Surrogate Found	Percent Recovery Reported	Percent Recovery Recalculated	Percent Difference
Nitrobenzene-d5	5.0	3.621	72.4	72.4	
2-Fluorobiphenyl	↓	3.848	77.0	77.0	
Terphenyl-d14	↓	3.860	77.2	77.2	
Phenol-d5	7.5	4.372	58.3	58.3	
2-Fluorophenol	↓	4.098	54.6	54.6	
2,4,6-Tribromophenol	↓	7.028	93.7	93.7	
2-Chlorophenol-d4	↓	5.203	69.4	69.4	
1,2-Dichlorobenzene-d4	5.0	3.358	67.2	67.2	

Sample ID: \_\_\_\_\_

	Surrogate Spiked	Surrogate Found	Percent Recovery Reported	Percent Recovery Recalculated	Percent Difference
Nitrobenzene-d5					
2-Fluorobiphenyl					
Terphenyl-d14					
Phenol-d5					
2-Fluorophenol					
2,4,6-Tribromophenol					
2-Chlorophenol-d4					
1,2-Dichlorobenzene-d4					

Sample ID: \_\_\_\_\_

	Surrogate Spiked	Surrogate Found	Percent Recovery Reported	Percent Recovery Recalculated	Percent Difference
Nitrobenzene-d5					
2-Fluorobiphenyl					
Terphenyl-d14					
Phenol-d5					
2-Fluorophenol					
2,4,6-Tribromophenol					
2-Chlorophenol-d4					
1,2-Dichlorobenzene-d4					

**VALIDATION FINDINGS WORKSHEET**  
**Matrix Spike/Matrix Spike Duplicates Results Verification**

**METHOD:** GC/MS BNA (EPA SW 846 Method 8270D)

The percent recoveries (%R) and Relative Percent Difference (RPD) of the matrix spike and matrix spike duplicate were recalculated for the compounds identified below using the following calculation:

% Recovery = 100 \* (SSC - SC)/SA

Where: SSC = Spiked sample concentration  
 SA = Spike added

SC = Sample concentration

RPD = |MSC - MSC| \* 2/(MSC + MSCD)

MSC = Matrix spike concentration

MSCD = Matrix spike duplicate concentration

MS/MSD samples: 5/6

Compound	Spike Added (ng/L)		Sample Concentration (ng/L)	Spiked Sample Concentration (ng/L)		Matrix Spike		Matrix Spike Duplicate		MS/MSD	
	MS	MSD		MS	MSD	Percent Recovery		Percent Recovery		RPD	
						Reported	Recalc	Reported	Recalc	Reported	Recalculated
Phenol	497	495	46.7	457	421	82.7	82.6	75.7	75.6	8.19	8.2
N-Nitroso-di-n-propylamine											
4-Chloro-3-methylphenol											
Acenaphthene	↓	↓	11.8	457	411	89.7	89.6	80.7	80.7	10.7	10.6
Pentachlorophenol											
Pyrene	↓	↓	150	443	493	160	160	69.2	69.3	62.7	62.7

Comments: Refer to Matrix Spike/Matrix Spike Duplicates 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

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

**METHOD:** GC/MS Semivolatiles (EPA SW 846 Method 8270D)

The percent recoveries (%R) and Relative Percent Difference (RPD) of the laboratory control sample and laboratory control sample duplicate were recalculated for the compounds identified below using the following calculation:

% Recovery = 100 \* (SC/SA)

Where: SSC = Spike concentration  
 SA = Spike added

RPD = |LCSC - LCSDC| \* 2 / (LCSC + LCSDC)

LCSC = Laboratory control sample concentration    LCSDC = Laboratory control sample duplicate concentration

LCS/LCSD samples: BH0806-B51

Compound	Spike Added (µg/L)		Spike Concentration (µg/L)		LCS		LCSD		LCS/LCSD	
	LCS	LCSD	LCS	LCSD	Percent Recovery		Percent Recovery		RPD	
					Reported	Recalc	Reported	Recalc	Reported	Recalculated
Phenol	500	NA	380	NA	76.1	76.0				
N-Nitroso-di-n-propylamine										
4-Chloro-3-methylphenol										
Acenaphthene	↓	↓	321	↓	64.3	64.2				
Pentachlorophenol										
Pyrene	↓	↓	366	↓	73.2	73.2				

Comments: Refer to Laboratory Control Sample/Laboratory Control Sample Duplicates 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**  
**Sample Calculation Verification**

**METHOD:** GC/MS SVOA (EPA SW 846 Method 8270D)

(Y) (N) N/A  
(Y) (N) N/A

Were all reported results recalculated and verified for all level IV samples?  
 Were all recalculated results for detected target compounds agree within 10.0% of the reported results?

$$\text{Concentration} = \frac{(A_x)(I_s)(V_i)(DF)(2.0)}{(A_s)(RRF)(V_o)(V_i)(\%S)}$$

- A<sub>x</sub> = Area of the characteristic ion (EICP) for the compound to be measured
- A<sub>s</sub> = Area of the characteristic ion (EICP) for the specific internal standard
- I<sub>s</sub> = Amount of internal standard added in nanograms (ng)
- V<sub>o</sub> = Volume or weight of sample extract in milliliters (ml) or grams (g).
- V<sub>i</sub> = Volume of extract injected in microliters (ul)
- V<sub>t</sub> = Volume of the concentrated extract in microliters (ul)
- Df = Dilution Factor.
- %S = Percent solids, applicable to soil and solid matrices only.
- 2.0 = Factor of 2 to account for GPC cleanup

Example:

Sample I.D. 1, S:

$$\text{Conc.} = \frac{(11949) \times (4.0) \times (1000) \times (1)}{(40518) \times (1.14344) \times (16.04) \times (0.67)} = 10.37 \text{ } \mu\text{g/g}$$

#	Sample ID	Compound	Reported Concentration ( <u>10.3</u> )	Calculated Concentration ( )	Qualification
	<u>1</u>	<u>S</u>	<u>10.3</u>		

## Laboratory Data Consultants, Inc. Data Validation Report

**Project/Site Name:** Duwamish AOC4  
**LDC Report Date:** December 15, 2021  
**Parameters:** Polynuclear Aromatic Hydrocarbons  
**Validation Level:** Stage 4  
**Laboratory:** Analytical Resources, Inc., Tukwila, WA  
**Sample Delivery Group (SDG):** 21J0134

<b>Sample Identification</b>	<b>Laboratory Sample Identification</b>	<b>Matrix</b>	<b>Collection Date</b>
LDW21-SS600	21J0134-11	Sediment	07/12/21
LDW21-SS600MS	21J0134-11MS	Sediment	07/12/21
LDW21-SS600MSD	21J0134-11MSD	Sediment	07/12/21

## 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 Final Lower Duwamish Waterway Quality Assurance Project Plan for Remedial Design of Upper Reach: Pre-Design Investigation (May 2020) and a modified outline of the USEPA National Functional Guidelines (NFG) for Organic Superfund Methods Data Review (January 2017). 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:

Polynuclear Aromatic Hydrocarbons (PAHs) by Environmental Protection Agency (EPA) SW 846 Method 8270E in Selected Ion Monitoring (SIM) mode

All sample results were subjected to Stage 4 data validation, which is comprised of the quality control (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): 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.
- U (Non-detected): The analyte was analyzed for and positively identified by the laboratory; however the analyte should be considered non-detected at the reported concentration due to the presence of contaminants detected in the associated blank(s).
- UJ (Non-detected estimated): The analyte was reported as not detected by the laboratory; however the reported quantitation/detection limit is estimated due to non-conformances discovered during data validation.
- R (Rejected): The sample results were rejected due to gross non-conformances discovered during data validation. Data qualified as rejected is not usable.
- 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.

## 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. GC/MS Instrument Performance Check

A decafluorotriphenylphosphine (DFTPP) tune was performed at 12 hour intervals.

All ion abundance requirements were met.

## III. Initial Calibration and Initial Calibration Verification

An initial calibration was performed as required by the method.

For analytes where average relative response factors (RRFs) were utilized, percent relative standard deviations (%RSD) were less than or equal to 20.0%.

In the case where the laboratory used a calibration curve to evaluate the analytes, all coefficients of determination ( $r^2$ ) were greater than or equal to 0.990.

Average relative response factors (RRF) for all analytes were within validation criteria.

The percent differences (%D) of the initial calibration verification (ICV) standard were less than or equal to 30.0% for all analytes.

## IV. 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 with the following exceptions:

Date	Analyte	%D	Associated Samples	Flag	A or P
10/30/21	Benzoic acid N-Nitrosodiphenylamine Pentachlorophenol	32.7 20.6 36.8	All samples in SDG 21J0134	J (all detects) J (all detects) J (all detects)	A

All of the continuing calibration relative response factors (RRF) were within validation criteria.

## V. Laboratory Blanks

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

## VI. Field Blanks

No field blanks were identified in this SDG.

## VII. Surrogates

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

## VIII. Matrix Spike/Matrix Spike Duplicates

Matrix spike (MS) and matrix spike duplicate (MSD) sample analysis was performed on an associated project sample. Percent recoveries (%R) were within QC limits with the following exceptions:

Spike ID (Associated Samples)	Analyte	MS (%R) (Limits)	MSD (%R) (Limits)	Flag	A or P
LDW21-SS600MS/MSD (LDW21-SS600)	N-Nitrosodiphenylamine	122 (27-120)	-	J (all detects)	A

Relative percent differences (RPD) were within QC limits.

## IX. Laboratory Control Samples/Standard Reference Materials

Laboratory control samples (LCS) were analyzed as required by the method. Percent recoveries (%R) were within QC limits.

Standard reference materials (SRM) were analyzed as required by the method. The results were within QC limits with the following exceptions:

SRM ID	Analyte	%R (Limits)	Associated Samples	Flag	A or P
BJJ0826-SRM2	1,4-Dichlorobenzene 1,2-Dichlorobenzene	10.2 (12-188) 10.2 (17-184)	All samples in SDG 21J0134	J (all detects) J (all detects)	A

## X. Field Duplicates

No field duplicates were identified in this SDG.

## XI. Internal Standards

All internal standard areas and retention times were within QC limits.

## XII. Target Analyte Quantitation

All target analyte quantitations were within validation criteria.



### **XIII. Target Analyte Identification**

All target analyte identifications were within validation criteria.

### **XIV. System Performance**

The system performance was acceptable.

### **XV. Overall Assessment of Data**

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

Due to continuing calibration %D, MS/MSD %R, and SRM %R, data were qualified as estimated in one sample.

The quality control criteria reviewed, other than those discussed above, were met and are considered acceptable.

**Duwamish AOC4  
Polynuclear Aromatic Hydrocarbons - Data Qualification Summary - SDG 21J0134**

Sample	Analyte	Flag	A or P	Reason
LDW21-SS600	Benzoic acid N-Nitrosodiphenylamine Pentachlorophenol	J (all detects) J (all detects) J (all detects)	A	Continuing calibration (%D)
LDW21-SS600	N-Nitrosodiphenylamine	J (all detects)	A	Matrix spike/Matrix spike duplicate (%R)
LDW21-SS600	1,4-Dichlorobenzene 1,2-Dichlorobenzene	J (all detects) J (all detects)	A	Standard reference materials (%R)

**Duwamish AOC4  
Polynuclear Aromatic Hydrocarbons - Laboratory Blank Data Qualification Summary - SDG 21J0134**

No Sample Data Qualified in this SDG

**Duwamish AOC4  
Polynuclear Aromatic Hydrocarbons - Field Blank Data Qualification Summary - SDG 21J0134**

No Sample Data Qualified in this SDG

LDC #: 52703B2b

**VALIDATION COMPLETENESS WORKSHEET**

SDG #: 21J0134

Stage 4

Laboratory: Analytical Resources, Inc., Tukwila, WA

Date: 12/10/21

Page: 1 of 1

Reviewer: [Signature]

2nd Reviewer: [Signature]

**METHOD:** GC/MS Polynuclear Aromatic Hydrocarbons (EPA SW 846 Method 8270E-SIM)

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	A	
II.	GC/MS Instrument performance check	A	
III.	Initial calibration/ICV	A/A	RSD ≤ 20% · γ <sup>2</sup>  CV ≤ 30%
IV.	Continuing calibration	W	CCV ≤ 20%
V.	Laboratory Blanks	A	
VI.	Field blanks	N	
VII.	Surrogate spikes	A	
VIII.	Matrix spike/Matrix spike duplicates	W	
IX.	Laboratory control samples /ERM	W	LCS
X.	Field duplicates	N	
XI.	Internal standards	A	
XII.	Target analyte quantitation	A	
XIII.	Target analyte identification	A	
XIV.	System performance	A	
XV.	Overall assessment of data	A	

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:

	Client ID	Lab ID	Matrix	Date
1	LDW21-SS600	21J0134-11	Sediment	07/12/21
2	LDW21-SS600MS	21J0134-11MS	Sediment	07/12/21
3	LDW21-SS600MSD	21J0134-11MSD	Sediment	07/12/21
4				
5				
6				
7				
8				
9				

Notes:

B110826					

Method: PAH (EPA SW 846 Method 8270D-SIM)

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>II. GC/MS Instrument performance check (Not required)</b>				
Were the DFTPP performance results reviewed and found to be within the specified criteria?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were all samples analyzed within the 12 hour clock criteria?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
<b>IIIa. 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) $\leq$ 20% and relative response factors (RRF) $\geq$ 0.05?	<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 $>$ 0.990?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
<b>IIIb. 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) $\leq$ 30%?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
<b>IV. Continuing calibration</b>				
Was a continuing calibration standard analyzed at least once every 12 hours for each instrument?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were all percent differences (%D) $\leq$ 20% and relative response factors (RRF) $\geq$ 0.05?	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	
<b>V. 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>VI. Field blanks</b>				
Were field blanks identified in this SDG?	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	
Were target compounds detected in the field blanks?	<input type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	
<b>VII. Surrogate spikes</b>				
Were all surrogate percent differences (%R) within QC limits?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
If 2 or more base neutral or acid surrogates were outside QC limits, was a reanalysis performed to confirm %R?	<input type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	
If any percent recoveries (%R) was less than 10 percent, was a reanalysis performed to confirm %R?	<input type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	
<b>VIII. 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 type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	

Validation Area	Yes	No	NA	Findings/Comments
<b>IX. Laboratory control samples</b>				
Was an LCS analyzed per 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"/>	
<b>X. Field duplicates</b>				
Were field duplicate pairs identified in this SDG?	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	
Were target compounds detected in the field duplicates?	<input type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	
<b>XI. Internal standards</b>				
Were internal standard area counts within -50% or +100% of the associated calibration standard?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were retention times within + 30 seconds of the associated calibration standard?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
<b>XII. Compound quantitation</b>				
Did the laboratory LOQs/RLs meet the QAPP LOQs/RLs?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were the correct internal standard (IS), quantitation ion and relative response factor (RRF) used to quantitate the compound?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were compound 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>XIII. Target compound identification</b>				
Were relative retention times (RRT's) within + 0.06 RRT units of the standard?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Did compound spectra meet specified EPA "Functional Guidelines" criteria?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were chromatogram peaks verified and accounted for?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
<b>XIV. System performance</b>				
System performance was found to be acceptable.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
<b>XV. Overall assessment of data</b>				
Overall assessment of data was found to be acceptable.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	

## VALIDATION FINDINGS WORKSHEET

### METHOD: GC/MS SVOA

A. Phenol	AA. 2-Chloronaphthalene	AAA. Butylbenzylphthalate	AAAA. Dibenzothiophene	A1. N-Nitrosodiethylamine
B. Bis (2-chloroethyl) ether	BB. 2-Nitroaniline	BBB. 3,3'-Dichlorobenzidine	BBBB. Benzo(a)fluoranthene	B1. N-Nitrosodi-n-butylamine
C. 2-Chlorophenol	CC. Dimethylphthalate	CCC. Benzo(a)anthracene	CCCC. Benzo(b)fluorene	C1. N-Nitrosomethylethylamine
D. 1,3-Dichlorobenzene	DD. Acenaphthylene	DDD. Chrysene	DDDD. cis/trans-Decalin	D1. N-Nitrosomorpholine
E. 1,4-Dichlorobenzene	EE. 2,6-Dinitrotoluene	EEE. Bis(2-ethylhexyl)phthalate	EEEE. Biphenyl	E1. N-Nitrosopyrrolidine
F. 1,2-Dichlorobenzene	FF. 3-Nitroaniline	FFF. Di-n-octylphthalate	FFFF. Retene	F1. Phenacetin
G. 2-Methylphenol	GG. Acenaphthene	GGG. Benzo(b)fluoranthene	GGGG. C30-Hopane	G1. 2-Acetylaminofluorene
H. 2,2'-Oxybis(1-chloropropane)	HH. 2,4-Dinitrophenol	HHH. Benzo(k)fluoranthene	HHHH. 1-Methylphenanthrene	H1. Pronamide
I. 4-Methylphenol	II. 4-Nitrophenol	III. Benzo(a)pyrene	IIII. 1,4-Dioxane	I1. Methyl methanesulfonate
J. N-Nitroso-di-n-propylamine	JJ. Dibenzofuran	JJJ. Indeno(1,2,3-cd)pyrene	JJJJ. Acetophenone	J1. Ethyl methanesulfonate
K. Hexachloroethane	KK. 2,4-Dinitrotoluene	KKK. Dibenz(a,h)anthracene	KKKK. Atrazine	K1. o,o',o''-Triethylphosphorothioate
L. Nitrobenzene	LL. Diethylphthalate	LLL. Benzo(g,h,i)perylene	LLLL. Benzaldehyde	L1. n-Phenylene diamine
M. Isophorone	MM. 4-Chlorophenyl-phenyl ether	MMM. Bis(2-Chloroisopropyl)ether	MMMM. Caprolactam	M1. 1,4-Naphthoquinone
N. 2-Nitrophenol	NN. Fluorene	NNN. Aniline	NNNN. 2,6-Dichlorophenol	N1. N-Nitro-o-toluidine
O. 2,4-Dimethylphenol	OO. 4-Nitroaniline	OOO. N-Nitrosodimethylamine	OOOO. 1,2-Diphenylhydrazine	O1. 1,3,5-Trinitrobenzene
P. Bis(2-chloroethoxy)methane	PP. 4,6-Dinitro-2-methylphenol	PPP. Benzoic Acid	PPPP. 3-Methylphenol	P1. Pentachlorobenzene
Q. 2,4-Dichlorophenol	QQ. N-Nitrosodiphenylamine	QQQ. Benzyl alcohol	QQQQ. 3&4-Methylphenol	Q1. 4-Aminobiphenyl
R. 1,2,4-Trichlorobenzene	RR. 4-Bromophenyl-phenylether	RRR. Pyridine	RRRR. 4-Dimethyldibenzothiophene (4MDT)	R1. 2-Naphthylamine
S. Naphthalene	SS. Hexachlorobenzene	SSS. Benzidine	SSSS. 2/3-Dimethyldibenzothiophene (4MDT)	S1. Triphenylene
T. 4-Chloroaniline	TT. Pentachlorophenol	TTT. 1-Methylnaphthalene	TTTT. 1-Methyldibenzothiophene (1MDT)	T1. Octachlorostyrene
U. Hexachlorobutadiene	UU. Phenanthrene	UUU. Benzo(b)thiophene	UUUU. 2,3,4,6-Tetrachlorophenol	U1. Famphur
V. 4-Chloro-3-methylphenol	VV. Anthracene	VVV. Benzonaphthothiophene	VVVV. 1,2,4,5-Tetrachlorobenzene	V1. 1,4-phenylenediamine
W. 2-Methylnaphthalene	WW. Carbazole	WWW. Benzo(e)pyrene	WWWWW. 2-Picoline	W1. Methapyrilene
X. Hexachlorocyclopentadiene	XX. Di-n-butylphthalate	XXX. 2,6-Dimethylnaphthalene	XXXX. 3-Methylcholanthrene	X1. Pentachloroethane
Y. 2,4,6-Trichlorophenol	YY. Fluoranthene	YYY. 2,3,5-Trimethylnaphthalene	YYYY. a,a-Dimethylphenethylamine	Y1. 3,3'-Dimethylbenzidine
Z. 2,4,5-Trichlorophenol	ZZ. Pyrene	ZZZ. Perylene	ZZZZ. Hexachloropropene	Z1. o-Toluidine

LDC #: ~~570329~~

### VALIDATION FINDINGS WORKSHEET Continuing Calibration

Page: 1 of 1

Reviewer: Q

2nd Reviewer: \_\_\_\_\_

METHOD: GC/MS BNA (EPA SW 846 Method 8270D)

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

N N/A Was a continuing calibration standard analyzed at least once every 12 hours for each instrument?

Y N/A Were percent differences (%D)  $\leq$  20 % and relative response factors (RRF) within the method criteria?

#	Date	Standard ID	Compound	Finding %D (Limit: $\leq$ 20.0%)	Finding RRF (Limit)	Associated Samples	Qualifications
	10/30/11	NT10211030035	PPP	32.7		#11 (del3)	<del>N/A</del>
			RR	20.6			↓
			TT	36.8			

### VALIDATION FINDINGS WORKSHEET

#### Matrix Spike/Matrix Spike Duplicates

**METHOD:** GC/MS BNA (EPA SW 846 Method 8270D)

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

N N/A Were a matrix spike (MS) and matrix spike duplicate (MSD) analyzed for each matrix in this SDG? If no, indicate which matrix does not have an associated MS/MSD. Soil / Water.

N N/A Was a MS/MSD analyzed every 20 samples of each matrix?

N N/A Were the MS/MSD percent recoveries (%R) and the relative percent differences (RPD) within the QC limits?

#	MS/MSD ID	Compound	MS %R (Limits)	MSD %R (Limits)	RPD (Limits)	Associated Samples	Qualifications
	<u>2/3</u>	<u>RR</u>	<u>122 (27-120)</u>	( )	( )	<u>1 (dots)</u>	<u>dots/A</u>
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
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			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		



LDC #: 5-70306

# VALIDATION FINDINGS WORKSHEET

## Laboratory Control Samples (LCS)

Page: 1 of 1  
Reviewer: [Signature]

METHOD: GC/MS BNA (EPA SW 846 Method 8270D)

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

N N/A Was a LCS required?

N N/A Were the LCS/LCSD percent recoveries (%R) and the relative percent differences (RPD) within the QC limits?

#	Date	LCS/LCSD ID	Compound	LCS %R (Limits)	LCSD %R (Limits)	RPD (Limits)	Associated Samples	Qualifications
		B110826-SM2	E	10.2 (12-188)	( )	( )	All (det)	N/A
			F	10.2 (17-184)	( )	( )		
				( )	( )	( )		
				( )	( )	( )		
				( )	( )	( )		
				( )	( )	( )		
				( )	( )	( )		
				( )	( )	( )		
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				( )	( )	( )		
				( )	( )	( )		
				( )	( )	( )		

## VALIDATION FINDINGS WORKSHEET Initial Calibration Calculation Verification

**METHOD:** GC/MS PCB (EPA SW 846 Method 8270DSIM)

The Relative Response Factor (RRF), average RRF, and percent relative standard deviation (%RSD) were recalculated for the compounds identified below using the following calculations:

RRF =  $(A_x)(C_{is}) / (A_{is})(C_x)$   
 average RRF = sum of the RRFs/number of standards  
 %RSD =  $100 * (S/X)$

$A_x$  = Area of compound,  $A_{is}$  = Area of associated internal standard  
 $C_x$  = Concentration of compound,  $C_{is}$  = Concentration of internal standard  
 S = Standard deviation of the RRFs, X = Mean of the RRFs

#	Standard ID	Calibration Date	Compound (Reference Internal Standard)	Reported	Recalculated	Reported	Recalculated	Reported	Recalculated
				RRF (   std)	RRF (   std)	Average RRF (initial)	Average RRF (initial)	%RSD	%RSD
1	KCA	10/5/21	E (1st internal standard)	1.26600T	1.26600T	1.32508T	1.32508T	11.5	11.5
			O (2nd internal standard)	0.4762963	0.4762963	0.473778	0.473778	14.2	14.2
			QQ (3rd internal standard)	0.781438	0.781438	0.7964012	0.7964012	14.4	14.4
			(4th internal standard)						
			(5th internal standard)						
			(6th internal standard)						
2			(1st internal standard)						
			(2nd internal standard)						
			(3rd internal standard)						
			(4th internal standard)						
			(5th internal standard)						
			(6th internal standard)						
3			(1st internal standard)						
			(2nd internal standard)						
			(3rd internal standard)						
			(4th internal standard)						
			(5th internal standard)						
			(6th internal standard)						

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.

## VALIDATION FINDINGS WORKSHEET Continuing Calibration Results Verification

**METHOD:** GC/MS PCB (EPA SW 846 Method 8270DSIM)

The percent difference (%D) of the initial calibration average Relative Response Factors (RRFs) and the continuing calibration RRFs were recalculated for the compounds identified below using the following calculation:

% Difference =  $100 * (\text{ave. RRF} - \text{RRF}) / \text{ave. RRF}$   
 $\text{RRF} = (A_x)(C_{is}) / (A_{is})(C_x)$

Where: ave. RRF = initial calibration average RRF  
 RRF = continuing calibration RRF  
 $A_x$  = Area of compound,  $A_{is}$  = Area of associated internal standard  
 $C_x$  = Concentration of compound,  $C_{is}$  = Concentration of internal standard

#	Standard ID	Calibration Date	Compound (Reference Internal Standard)	Average RRF (initial)	Reported	Recalculated	Reported	Recalculated
					RRF (CC)	RRF (CC)	%D	%D
1	<u>NT1021070039</u>	<u>10/30/21</u>	<u>E</u> (1st internal standard)	<u>1.32508T</u>	<u>1.2890450</u>	<u>1.2890448</u>	<u>2.7</u>	<u>2.7</u>
			<u>O</u> (2nd internal standard)	<u>0.4131118</u>	<u>0.4562324</u>	<u>0.4562322</u>	<u>3.1</u>	<u>3.1</u>
			<u>QA</u> (3rd internal standard)	<u>0.964012</u>	<u>0.9602963</u>	<u>0.960296</u>	<u>20.6</u>	<u>20.6</u>
			(4th internal standard)					
			(5th internal standard)					
			(6th internal standard)					
2			(1st internal standard)					
			(2nd internal standard)					
			(3rd internal standard)					
			(4th internal standard)					
			(5th internal standard)					
			(6th internal standard)					
3			(1st internal standard)					
			(2nd internal standard)					
			(3rd internal standard)					
			(4th internal standard)					
			(5th internal standard)					
			(6th internal standard)					

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/MS Semivolatiles (EPA SW 846 Method 8270D-SIM)

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 Spiked	Surrogate Found	Percent Recovery Reported	Percent Recovery Recalculated	Percent Difference
Nitrobenzene-d5					
2-Fluorobiphenyl					
Terphenyl-d14	5.0	2.828	56.6	56.6	
Phenol-d5					
2-Fluorophenol	7.5	3.961	52.8	52.8	
2,4,6-Tribromophenol					
2-Chlorophenol-d4					
1,2-Dichlorobenzene-d4					

**Sample ID:** \_\_\_\_\_

	Surrogate Spiked	Surrogate Found	Percent Recovery Reported	Percent Recovery Recalculated	Percent Difference
Nitrobenzene-d5					
2-Fluorobiphenyl					
Terphenyl-d14					
Phenol-d5					
2-Fluorophenol					
2,4,6-Tribromophenol					
2-Chlorophenol-d4					
1,2-Dichlorobenzene-d4					

**Sample ID:** \_\_\_\_\_

	Surrogate Spiked	Surrogate Found	Percent Recovery Reported	Percent Recovery Recalculated	Percent Difference
Nitrobenzene-d5					
2-Fluorobiphenyl					
Terphenyl-d14					
Phenol-d5					
2-Fluorophenol					
2,4,6-Tribromophenol					
2-Chlorophenol-d4					
1,2-Dichlorobenzene-d4					

**VALIDATION FINDINGS WORKSHEET**  
**Matrix Spike/Matrix Spike Duplicates Results Verification**

**METHOD:** GC/MS PAHs (EPA SW 846 Method 8270D-SIM)

The percent recoveries (%R) and Relative Percent Difference (RPD) of the matrix spike and matrix spike duplicate were recalculated for the compounds identified below using the following calculation:

% Recovery =  $100 * (SSC - SC) / SA$

Where: SSC = Spiked sample concentration  
 SA = Spike added

SC = Sample concentration

RPD =  $|MSC - MSC| * 2 / (MSC + MSDC)$

MSC = Matrix spike concentration

MSDC = Matrix spike duplicate concentration

MS/MSD samples: 2/3

Compound	Spike Added ( <u>145</u> )		Sample Concentration ( <u>145</u> )	Spiked Sample Concentration ( <u>145</u> )		Matrix Spike		Matrix Spike Duplicate		MS/MSD	
	MS	MSD		MS	MSD	Percent Recovery		Percent Recovery		RPD	
						Reported	Recalc	Reported	Recalc	Reported	Recalculated
Acenaphthene											
Pyrene											
<u>Σ</u>	<u>497</u>	<u>495</u>	<u>2.4</u>	<u>427</u>	<u>404</u>	<u>85.4</u>	<u>85.4</u>	<u>81.1</u>	<u>81.1</u>	<u>5.56</u>	<u>5.54</u>
<u>TT</u>	<u>1290</u>	<u>1290</u>	<u>2.7</u>	<u>1320</u>	<u>1130</u>	<u>102</u>	<u>102</u>	<u>87.3</u>	<u>87.4</u>	<u>15.6</u>	<u>15.5</u>

Comments: Refer to Matrix Spike/Matrix Spike Duplicates 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

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

**METHOD:** GC/MS PAHs (EPA SW 846 Method 8270D-SIM)

The percent recoveries (%R) and Relative Percent Difference (RPD) of the laboratory control sample and laboratory control sample duplicate were recalculated for the compounds identified below using the following calculation:

% Recovery = 100 \* (SC/SA)

Where: SSC = Spike concentration  
 SA = Spike added

RPD = |LCSC - LCSDC| \* 2 / (LCSC + LCSDC)

LCSC = Laboratory control sample concentration    LCSDC = Laboratory control sample duplicate concentration

LCS/LCSD samples: B-10826-BS2

Compound	Spike Added ( <u>1445</u> )		Spike Concentration ( <u>1415</u> )		LCS		LCSD		LCS/LCSD	
	LCS	LCSD	LCS	LCSD	Percent Recovery		Percent Recovery		RPD	
					Reported	Recalc	Reported	Recalc	Reported	Recalculated
Acenaphthene										
Pyrene										
<u>E</u>	<u>500</u>	<u>500</u>	<u>387</u>	<u>385</u>	<u>77.4</u>	<u>77.4</u>	<u>76.9</u>	<u>77</u>	<u>0.569</u>	<u>0.39</u>
<u>TT</u>	<u>1300</u>	<u>1300</u>	<u>1140</u>	<u>1180</u>	<u>87.3</u>	<u>87.7</u>	<u>90.7</u>	<u>90.8</u>	<u>3.82</u>	<u>3.45</u>

Comments: Refer to Laboratory Control Sample/Laboratory Control Sample Duplicates 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

### Sample Calculation Verification

METHOD: GC/MS PAHs (EPA SW 846 Method 8270D-SIM)

Y N N/A  
Y N N/A

Were all reported results recalculated and verified for all level IV samples?

Were all recalculated results for detected target compounds agree within 10.0% of the reported results?

$$\text{Concentration} = \frac{(A_x)(I_s)(V_i)(DF)(2.0)}{(A_{is})(RRF)(V_o)(V_i)(\%S)}$$

- $A_x$  = Area of the characteristic ion (EICP) for the compound to be measured
- $A_{is}$  = Area of the characteristic ion (EICP) for the specific internal standard
- $I_s$  = Amount of internal standard added in nanograms (ng)
- $V_o$  = Volume or weight of sample extract in milliliters (ml) or grams (g).
- $V_i$  = Volume of extract injected in microliters (ul)
- $V_t$  = Volume of the concentrated extract in microliters (ul)
- Df = Dilution Factor.
- %S = Percent solids, applicable to soil and solid matrices only.
- 2.0 = Factor of 2 to account for GPC cleanup

Example:

Sample I.D. 1, E:

$$\text{Conc.} = \frac{(827) \times (4.0) \times (1000) \times (1)}{(102620) \times (1.32508) \times (16.04) \times (0.672)}$$

$$= 2.42 \text{ ng/g}$$

#	Sample ID	Compound	Reported Concentration ( <u>ng/g</u> )	Calculated Concentration ( )	Qualification
	<u>1</u>	<u>E</u>	<u>2.4</u>		

## Laboratory Data Consultants, Inc. Data Validation Report

**Project/Site Name:** Duwamish AOC4  
**LDC Report Date:** December 15, 2021  
**Parameters:** Hexachlorobenzene  
**Validation Level:** Stage 4  
**Laboratory:** Analytical Resources, Inc., Tukwila, WA  
**Sample Delivery Group (SDG):** 21J0134

<b>Sample Identification</b>	<b>Laboratory Sample Identification</b>	<b>Matrix</b>	<b>Collection Date</b>
LDW21-SS600	21J0134-11	Sediment	07/12/21
LDW21-SS600MS	21J0134-11MS	Sediment	07/12/21
LDW21-SS600MSD	21J0134-11MSD	Sediment	07/12/21



## 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 Final Lower Duwamish Waterway Quality Assurance Project Plan for Remedial Design of Upper Reach: Pre-Design Investigation (May 2020) and a modified outline of the USEPA National Functional Guidelines (NFG) for Organic Superfund Methods Data Review (January 2017). 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:

Hexachlorobenzene by Environmental Protection Agency (EPA) SW 846 Method 8081B

All sample results were subjected to Stage 4 data validation, which is comprised of the quality control (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): 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.
- U (Non-detected): The analyte was analyzed for and positively identified by the laboratory; however the analyte should be considered non-detected at the reported concentration due to the presence of contaminants detected in the associated blank(s).
- UJ (Non-detected estimated): The analyte was reported as not detected by the laboratory; however the reported quantitation/detection limit is estimated due to non-conformances discovered during data validation.
- R (Rejected): The sample results were rejected due to gross non-conformances discovered during data validation. Data qualified as rejected is not usable.
- 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.

## **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. GC Instrument Performance Check**

Instrument performance was checked at 12 hour intervals.

The individual 4,4'-DDT and Endrin breakdowns (%BD) were less than or equal to 15.0%.

## **III. 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%.

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

## **IV. Continuing Calibration**

Continuing calibration was performed at required frequencies.

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

## **V. Laboratory Blanks**

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

## **VI. Field Blanks**

No field blanks were identified in this SDG.

## **VII. Surrogates/Internal Standards**

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

All internal standard areas and retention times were within QC limits.

## **VIII. Matrix Spike/Matrix Spike Duplicates**

Matrix spike (MS) and matrix spike duplicate (MSD) sample analysis was performed on an associated project sample. Percent recoveries (%R) were within QC limits. Relative percent differences (RPD) were within QC limits.

## **IX. 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.

## **X. Field Duplicates**

No field duplicates were identified in this SDG.

## **XI. Target Analyte Quantitation**

All target analyte quantitations met validation criteria.

## **XII. Target Analyte Identification**

All target analyte identifications met validation criteria.

## **XIII. System Performance**

The system performance was acceptable.

## **XIV. Overall Assessment of Data**

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

The quality control criteria reviewed were met and are considered acceptable.

**Duwamish AOC4  
Hexachlorobenzene - Data Qualification Summary - SDG 21J0134**

No Sample Data Qualified in this SDG

**Duwamish AOC4  
Hexachlorobenzene - Laboratory Blank Data Qualification Summary - SDG  
21J0134**

No Sample Data Qualified in this SDG

**Duwamish AOC4  
Hexachlorobenzene - Field Blank Data Qualification Summary - SDG 21J0134**

No Sample Data Qualified in this SDG

LDC #: 52703B3a

**VALIDATION COMPLETENESS WORKSHEET**

Date: 12/10/21

SDG #: 21J0134

Stage 4

Page: 1 of 1

Laboratory: Analytical Resources, Inc., Tukwila, WA

Reviewer: [Signature]

2nd Reviewer: [Signature]

**METHOD:** GC Hexachlorobenzene (EPA SW846 Method 8081B)

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	A	
II.	GC Instrument Performance Check	A	
III.	Initial calibration/ICV	A/A	RSD ≤ 20% 1σV ≤ 20%
IV.	Continuing calibration	A	CCV ≤ 20%
V.	Laboratory Blanks	A	
VI.	Field blanks	N	
VII.	Surrogate spikes / IS	A/A	
VIII.	Matrix spike/Matrix spike duplicates	A	
IX.	Laboratory control samples	A	LCS/D
X.	Field duplicates	N	
XI.	Target analyte quantitation	A	
XII.	Target analyte identification	A	
XIII.	System Performance	A	
XIV.	Overall assessment of data	D	

Note: A = Acceptable ND = No compounds detected D = Duplicate SB=Source blank  
 N = Not provided/applicable R = Rinsate TB = Trip blank OTHER:  
 SW = See worksheet FB = Field blank EB = Equipment blank

	Client ID	Lab ID	Matrix	Date
1	LDW21-SS600	21J0134-11	Sediment	07/12/21
2	LDW21-SS600MS	21J0134-11MS	Sediment	07/12/21
3	LDW21-SS600MSD	21J0134-11MSD	Sediment	07/12/21
4				
5				
6				
7				
8				
9				
10				
11				

Notes:

BH0639				

**Method:** Pesticides (EPA SW 846 Method 8081A)

Validation Area	Yes	No	NA	Findings/Comments
<b><i>I. Technical holding times</i></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><i>II. GC/ECD Instrument performance check</i></b>				
Was the instrument performance found to be acceptable?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were Evaluation mix standards analyzed prior to the initial calibration and at beginning of each 12-hour shift?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were endrin and 4,4'-DDT breakdowns $\leq 15\%$ for individual breakdown in the Evaluation mix standards?	<input type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	
<b><i>IIIa. Initial calibration</i></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) $\leq 20\%$ ?	<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><i>IIIb. Initial calibration verification</i></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) $\leq 20\%$ ?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
<b><i>IV. Continuing calibration</i></b>				
Was a continuing calibration analyzed daily?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were all percent differences (%D) $\leq 20\%$ ?	<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><i>V. Laboratory Blanks</i></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><i>VI. Field blanks</i></b>				
Were field blanks identified in this SDG?	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	
Were target compounds detected in the field blanks?	<input type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	
<b><i>VII. Surrogate spikes/Internal Standards</i></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"/>	

Validation Area	Yes	No	NA	Findings/Comments
If any percent recovery (%R) was less than 10 percent, was a reanalysis performed to confirm %R?			/	
Were internal standard area counts within $\pm 50\%$ of the average area calculated during calibration?	/			
<b>VII. Matrix spike/Matrix spike duplicates</b>				
Were a matrix spike (MS) and matrix spike duplicate (MSD) analyzed for each matrix in this SDG?	/			
Were the MS/MSD percent recoveries (%R) and the relative percent differences (RPD) within the QC limits?	/			
<b>IX. Laboratory control samples</b>				
Was an LCS analyzed per extraction batch?	/			
Were the LCS percent recoveries (%R) and relative percent difference (RPD) within the QC limits?	/			
<b>X. Field duplicates</b>				
Were field duplicate pairs identified in this SDG?		/		
Were target compounds detected in the field duplicates?			/	
<b>XI. Compound quantitation</b>				
Did the laboratory LOQs/RLs meet the QAPP LOQs/RLs?	/			
Were compound quantitation and RLs adjusted to reflect all sample dilutions, dry weight factors, and clean-up activities applicable to level IV validation?	/			
Were relative percent difference (RPD) of the results between two columns $\leq 40\%$ ?			/	
<b>XII. Target compound identification</b>				
Were the retention times of reported detects within the RT windows?	/			
<b>XIII. Overall assessment of data</b>				
Overall assessment of data was found to be acceptable.	/			

## VALIDATION FINDINGS WORKSHEET

**METHOD:** Pesticides

A. alpha-BHC	K. Endrin	U. Toxaphene	EE. 2,4'-DDT	OO. oxy-Chlordane
B. beta-BHC	L. Endosulfan II	V. Aroclor-1016	FF. Hexachlorobenzene	PP. cis-Nonachlor
C. delta-BHC	M. 4,4'-DDD	W. Aroclor-1221	GG. Chlordane	QQ. trans-Nonachlor
D. gamma-BHC	N. Endosulfan sulfate	X. Aroclor-1232	HH. Chlordane (Technical)	RR. cis-Chlordane
E. Heptachlor	O. 4,4'-DDT	Y. Aroclor-1242	II. p,p'-DDE	SS. trans-Chlordane
F. Aldrin	P. Methoxychlor	Z. Aroclor-1248	JJ. p,p'-DDD	TT. alpha-Endosulphan
G. Heptachlor epoxide	Q. Endrin ketone	AA. Aroclor-1254	KK. p,p'-DDT	UU. beta-Endosulphan
H. Endosulfan I	R. Endrin aldehyde	BB. Aroclor-1260	LL. o,p'-DDT	VV. Endosulphan Sulphate
I. Dieldrin	S. alpha-Chlordane	CC. 2,4'-DDD	MM. o,p'-DDE	WW. Mirex
J. 4,4'-DDE	T. gamma-Chlordane	DD. 2,4'-DDE	NN. o,p'-DDD	

Notes: \_\_\_\_\_



**VALIDATION FINDINGS WORKSHEET**  
**Initial Calibration Calculation Verification**

**METHOD:** GC Pesticides (EPA SW 846 Method 8081A)

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 (10 std)	CF (10 std)	Ave CF (initial)	Ave CF (initial)	%RSD	%RSD
1	KAL	6/21/21	FF (10)	1.292649	1.292649	1.29694	1.29694	12.7	12.7
			FF (20)	1.247978	1.247978	1.240281	1.240281	12.5	12.5
2									
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.

**VALIDATION FINDINGS WORKSHEET**  
**Continuing Calibration Results Verification**

**METHOD:** GC Pesticides (EPA SW 846 Method 8081B)

Percent difference (%D) =  $100 * (N - C)/N$

Where: N = Initial Calibration Factor or Nominal Amount (ng)  
 C = Calibration Factor from Continuing Calibration Standard or Calculated Amount (ng)

Standard ID	Calibration Date/Time	Compound	Average CF/ CCV Conc	Reported	Recalculated	Reported	Recalculated
				CF/Conc CCV	CF/Conc CCV	%D	%D
21102805	10/68/21	FF (1C)	1.29694	1.4131260	1.4131260	9.0	9.0
		FF (2C)	1.240281	1.1819270	1.181927	5.0	4.7

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 Pesticides (EPA SW 846 Method 8081A)

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	Surrogate Spiked	Surrogate Found	Percent Recovery		Percent Difference
				Reported	Recalculated	
Tetrachloro-m-xylene	<u>STX-CLP</u>	<u>40.0</u>	<u>30.40</u>	<u>76.0</u>	<u>76.0</u>	
Decachlorobiphenyl	<u>↓</u>	<u>↓</u>	<u>37.17</u>	<u>92.9</u>	<u>92.9</u>	
Tetrachloro-m-xylene						
Decachlorobiphenyl						

**Sample ID:** \_\_\_\_\_

Surrogate	Column	Surrogate Spiked	Surrogate Found	Percent Recovery		Percent Difference
				Reported	Recalculated	
Tetrachloro-m-xylene						
Decachlorobiphenyl						
Tetrachloro-m-xylene						
Decachlorobiphenyl						

**Sample ID:** \_\_\_\_\_

Surrogate	Column	Surrogate Spiked	Surrogate Found	Percent Recovery		Percent Difference
				Reported	Recalculated	
Tetrachloro-m-xylene						
Decachlorobiphenyl						
Tetrachloro-m-xylene						
Decachlorobiphenyl						

**Sample ID:** \_\_\_\_\_

Surrogate	Column	Surrogate Spiked	Surrogate Found	Percent Recovery		Percent Difference
				Reported	Recalculated	
Tetrachloro-m-xylene						
Decachlorobiphenyl						
Tetrachloro-m-xylene						
Decachlorobiphenyl						

Notes: \_\_\_\_\_  
\_\_\_\_\_

## VALIDATION FINDINGS WORKSHEET

### Matrix Spike/Matrix Spike Duplicates Results Verification

**METHOD:** GC Pesticides (EPA SW 846 Method 8081~~8~~)

The percent recoveries (%R) and Relative Percent difference (RPD) of the matrix spike and matrix spike duplicate were recalculated for the compounds identified below using the following calculation:

% Recovery = 100 \* (SSC-SC)/SA

Where: SSC = Spiked sample concentration  
SA = Spike added

SC = Concentration

RPD = | SSCMS - SSCMSD | \* 2 / (SSCMS + SSCMSD)

MS = Matrix spike percent recovery

MSD = Matrix spike duplicate percent recovery

MS/MSD samples: 2/3

Compound	Spike Added ( <u>145</u> )		Sample Concentration ( <u>145</u> )	Spiked Sample Concentration ( <u>145</u> )		Matrix Spike		Matrix Spike Duplicate		MS/MSD	
	MS	MSD		MS	MSD	Percent Recovery		Percent Recovery		RPD	
						Reported	Recalc.	Reported	Recalc.	Reported	Recalculated
gamma-BHC			--								
4,4'-DDT											
Aroclor 1260											
<u>FF</u>	<u>3.99</u>	<u>3.99</u>	<u>ND</u>	<u>3.74</u>	<u>3.38</u>	<u>93.7</u>	<u>93.7</u>	<u>84.7</u>	<u>84.7</u>	<u>10.3</u>	<u>10.1</u>

Comments: Refer to Matrix Spike/Matrix Spike Duplicates 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

### Laboratory Control Sample/Laboratory Control Sample Duplicate Results Verification

**METHOD:** GC Pesticides (EPA SW 846 Method 8081A)

The percent recoveries (%R) and Relative Percent difference (RPD) of the laboratory control sample and laboratory control sample duplicate were recalculated for the compounds identified below using the following calculation:

% Recovery =  $100 * (SSC - SC) / SA$

Where: SSC = Spiked sample concentration  
 SA = Spike added

SC = Concentration

RPD =  $|LCS - LCSD| * 2 / (LCS + LCSD)$

LCS = Laboratory control sample percent recovery

LCSD = Laboratory control sample duplicate percent recovery

LCS/LCSD samples: BJ0639-BS1/-BSD1

Compound	Spike Added (150)		Spiked Sample Concentration (14.8)		LCS		LCSD		LCS/LCSD	
					Percent Recovery		Percent Recovery		RPD	
	LCS	LCSD	LCS	LCSD	Reported	Recalc.	Reported	Recalc.	Reported	Recalc.
gamma-BHC										
4,4'-DDT										
FF	4.00	4.00	3.88	35.1	97.0	97.0	87.8	87.8	9.90	10.0

Comments: Refer to Laboratory Control Sample/Laboratory Control Sample Duplicate 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

### Sample Calculation Verification

**METHOD:** GC Pesticides (EPA SW 846 Method 8081A)

Y N N/A  
Y N N/A

Were all reported results recalculated and verified for all level IV samples?

Were all recalculated results for detected target compounds agree within 10.0% of the reported results?

$$\text{Concentration} = \frac{(A_x)(I_s)(V_i)(DF)(2.0)}{(A_{is})(RRF)(V_o)(V_1)(\%S)}$$

- A<sub>x</sub> = Area of the characteristic ion (EICP) for the compound to be measured
- A<sub>is</sub> = Area of the characteristic ion (EICP) for the specific internal standard
- I<sub>s</sub> = Amount of internal standard added in nanograms (ng)
- V<sub>o</sub> = Volume or weight of sample extract in milliliters (ml) or grams (g).
- V<sub>i</sub> = Volume of extract injected in microliters (ul)
- V<sub>1</sub> = Volume of the concentrated extract in microliters (ul)
- Df = Dilution Factor.
- %S = Percent solids, applicable to soil and solid matrices only.
- 2.0 = Factor of 2 to account for GPC cleanup

Example:

Sample I.D. NO FF  
BN10639-BS1

$$\text{Conc.} = \frac{(618819)(80.0)(2.5)(1)}{(196778)(1.2969)(12.5)( )}$$

*= 3.88 ng/kg*

#	Sample ID	Compound	Reported Concentration ( <u>ng/kg</u> )	Calculated Concentration ( )	Qualification
	<u>BN10639-BS1</u>	<u>FF</u>	<u>3.88</u>		

## Laboratory Data Consultants, Inc. Data Validation Report

**Project/Site Name:** Duwamish AOC4

**LDC Report Date:** December 15, 2021

**Parameters:** Polychlorinated Biphenyls

**Validation Level:** Stage 4

**Laboratory:** Analytical Resources, Inc., Tukwila

**Sample Delivery Group (SDG):** 21J0134

Sample Identification	Laboratory Sample Identification	Matrix	Collection Date
LDW21-IT669D	21J0134-01	Sediment	07/08/21
LDW21-IT598B	21J0134-02	Sediment	07/08/21
LDW21-IT598C	21J0134-03	Sediment	07/08/21
LDW21-IT598D	21J0134-04	Sediment	07/08/21
LDW21-IT598E	21J0134-05	Sediment	07/08/21
LDW21-IT598F	21J0134-06	Sediment	07/08/21
LDW21-IT598G	21J0134-07	Sediment	07/08/21
LDW21-IT598H	21J0134-08	Sediment	07/08/21
LDW21-SC553D	21J0134-09	Sediment	07/09/21
LDW21-SC554D	21J0134-10	Sediment	07/09/21
LDW21-SS600	21J0134-11	Sediment	07/12/21
LDW21-SS641	21J0134-13	Sediment	07/09/21
LDW21-SC587A	21J0134-14	Sediment	07/12/21
LDW21-SC587F	21J0134-15	Sediment	07/12/21
LDW21-IT660C	21J0134-16	Sediment	07/14/21
LDW21-IT588F	21J0134-17	Sediment	07/14/21
LDW21-SC568F	21J0134-19	Sediment	07/14/21
LDW21-IT598CMS	21J0134-03MS	Sediment	07/08/21
LDW21-IT598CMSD	21J0134-03MSD	Sediment	07/08/21

## 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 Final Lower Duwamish Waterway Quality Assurance Project Plan for Remedial Design of Upper Reach: Pre-Design Investigation (May 2020) and a modified outline of the USEPA National Functional Guidelines (NFG) for Organic Superfund Methods Data Review (January 2017). 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:

Polychlorinated Biphenyls (PCBs) by Environmental Protection Agency (EPA) SW 846 Method 8082A

All sample results were subjected to Stage 4 data validation, which is comprised of the quality control (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): 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.
- U (Non-detected): The analyte was analyzed for and positively identified by the laboratory; however the analyte should be considered non-detected at the reported concentration due to the presence of contaminants detected in the associated blank(s).
- UJ (Non-detected estimated): The analyte was reported as not detected by the laboratory; however the reported quantitation/detection limit is estimated due to non-conformances discovered during data validation.
- R (Rejected): The sample results were rejected due to gross non-conformances discovered during data validation. Data qualified as rejected is not usable.
- 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.



## 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.

Retention time windows were established as required by the method.

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 required frequencies.

The percent differences (%D) were less than or equal to 20.0% for all analytes with the following exceptions:

Date	Standard	Column	Analyte	%D	Associated Samples	Flag	A or P
10/27/21	10272115	2C	Aroclor-1260	24.6	LDW21-IT598G	J (all detects)	A

Retention times of all analytes in the calibration standards were within the established retention time windows.

## 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/Internal Standards

Surrogates were added to all samples as required by the method. Surrogate recoveries (%R) were not within QC limits for sample LDW21-IT598B. No data were qualified for samples analyzed at greater than or equal to 5X dilution.

All internal standard percent recoveries (%R) were within QC limits with the following exceptions:

Sample	Internal Standards	%R (Limits)	Affected Analyte	Flag	A or P
LDW21-IT598F	Hexabromobiphenyl	41 (50-200)	Aroclor-1260	J (all detects)	A
LDW21-SC553D	Hexabromobiphenyl	47 (50-200)	Aroclor-1260	UJ (all non-detects)	A
LDW21-SC587A	Hexabromobiphenyl	45 (50-200)	Aroclor-1260	J (all detects)	A
LDW21-SC587F	Hexabromobiphenyl	40 (50-200)	Aroclor-1260	J (all detects)	A
LDW21-SC568F	Hexabromobiphenyl	49 (50-200)	Aroclor-1260	J (all detects)	A

### VII. Matrix Spike/Matrix Spike Duplicates

Matrix spike (MS) and matrix spike duplicate (MSD) sample analysis was performed on an associated project sample. Percent recoveries (%R) were within QC limits. Relative percent differences (RPD) were within QC limits.

### VIII. Laboratory Control Samples/Standard Reference Materials

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.

Standard reference materials (SRM) were analyzed as required by the method. The results were within QC limits.

### IX. Field Duplicates

No field duplicates were identified in this SDG.

### X. Target Analyte Quantitation

All target analyte quantitations met validation criteria.

The sample results for detected analytes from the two columns were within 40% relative percent difference (RPD) with the following exceptions:

Sample	Analyte	RPD	Flag	A or P
LDW21-IT598D	Aroclor-1260	42.6	J (all detects)	A
LDW21-SC587F	Aroclor-1260	45.1	J (all detects)	A

Sample	Analyte	RPD	Flag	A or P
LDW21-SC568F	Aroclor-1260	40.8	J (all detects)	A

### **XI. Target Analyte Identification**

All target analyte identifications met validation criteria.

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

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

Due to continuing calibration %D, internal standard %R, and RPD between two columns, data were qualified as estimated in seven samples.

The quality control criteria reviewed, other than those discussed above, were met and are considered acceptable.

**Duwamish AOC4  
Polychlorinated Biphenyls - Data Qualification Summary - SDG 21J0134**

Sample	Analyte	Flag	A or P	Reason
LDW21-IT598G	Aroclor-1260	J (all detects)	A	Continuing calibration (%D)
LDW21-IT598F LDW21-SC553D LDW21-SC587A LDW21-SC587F LDW21-SC568F	Aroclor-1260	J (all detects) UJ (all non-detects)	A	Internal standards (%R)
LDW21-IT598D LDW21-SC587F LDW21-SC568F	Aroclor-1260	J (all detects)	A	Target analyte quantitation (RPD between two columns)

**Duwamish AOC4  
Polychlorinated Biphenyls - Laboratory Blank Data Qualification Summary - SDG 21J0134**

No Sample Data Qualified in this SDG

**Duwamish AOC4  
Polychlorinated Biphenyls - Field Blank Data Qualification Summary - SDG 21J0134**

No Sample Data Qualified in this SDG

LDC #: 52703B3b  
 SDG #: 21J0134  
 Laboratory: Analytical Resources, Inc.

**VALIDATION COMPLETENESS WORKSHEET**

Stage 4

Date: 12/10/21  
 Page: 1 of 3  
 Reviewer: Q  
 2nd Reviewer: JLG

**METHOD:** GC Polychlorinated Biphenyls (EPA SW846 Method 8082A)

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	A	
II.	Initial calibration/ICV	A/A	RSD ≤ 20% ICV ≤ 20%
III.	Continuing calibration	W	CCV ≤ 20%
IV.	Laboratory Blanks	A	
V.	Field blanks	N	
VI.	Surrogate spikes / IS	W/W	
VII.	Matrix spike/Matrix spike duplicates	A	
VIII.	Laboratory control samples / SRM	A	LCS/3
IX.	Field duplicates	N	
X.	Target analyte quantitation	W	
XI.	Target analyte identification	A	
XII.	Overall assessment of data	A	

Note: A = Acceptable ND = No compounds detected D = Duplicate SB=Source blank  
 N = Not provided/applicable R = Rinsate TB = Trip blank OTHER:  
 SW = See worksheet FB = Field blank EB = Equipment blank

	Client ID	Lab ID	Matrix	Date
1	LDW21-IT669D	21J0134-01	Sediment	07/08/21
2	LDW21-IT598B	21J0134-02	Sediment	07/08/21
3	LDW21-IT598C	21J0134-03	Sediment	07/08/21
4	LDW21-IT598D	21J0134-04	Sediment	07/08/21
5	LDW21-IT598E	21J0134-05	Sediment	07/08/21
6	LDW21-IT598F	21J0134-06	Sediment	07/08/21
7	LDW21-IT598G	21J0134-07	Sediment	07/08/21
8	LDW21-IT598H	21J0134-08	Sediment	07/08/21
9	LDW21-SC553D	21J0134-09	Sediment	07/09/21
10	LDW21-SC554D	21J0134-10	Sediment	07/09/21
11	LDW21-SS600	21J0134-11	Sediment	07/12/21
12	LDW21-SS641	21J0134-13	Sediment	07/09/21
13	LDW21-SC587A	21J0134-14	Sediment	07/12/21
14	LDW21-SC587F	21J0134-15	Sediment	07/12/21
15	LDW21-IT660C	21J0134-16	Sediment	07/14/21
16	LDW21-IT588F	21J0134-17	Sediment	07/14/21
17	LDW21-SC568F	JLG 21J0134-19	Sediment	07/14/21

LDC #: 52703B3b

# VALIDATION COMPLETENESS WORKSHEET

Date: 12/19/21

SDG #: 21J0134

Stage 4

Page: 2 of 2

Laboratory: Analytical Resources, Inc.

Reviewer: [Signature]

2nd Reviewer: [Signature]

**METHOD:** GC Polychlorinated Biphenyls (EPA SW846 Method 8082A)

	Client ID	Lab ID	Matrix	Date
18	LDW21-IT598CMS	21J0134-03MS	Sediment	07/08/21
19	LDW21-IT598CMSD	21J0134-03MSD	Sediment	07/08/21
20				
21				
22				

Notes:

	BL100-9-BK1				

Method:  GC  HPLC

Validation Area	Yes	No	NA	Findings/Comments
<b><i>I. Technical holding times</i></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><i>Ia. Initial calibration</i></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 ≥ 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><i>Iib. Initial calibration verification</i></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><i>III. Continuing calibration</i></b>				
Was a continuing calibration analyzed daily?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were all percent differences (%D) < 20%?	<input type="checkbox"/>	<input checked="" 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><i>IV. Laboratory Blanks</i></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><i>V. Field Blanks</i></b>				
Were field blanks identified in this SDG?	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	
Were target compounds detected in the field blanks?	<input type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	
<b><i>VI. Surrogate spikes</i></b>				
Were all surrogate percent recovery (%R) within the QC limits?	<input type="checkbox"/>	<input checked="" 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><i>VII. Matrix spike/Matrix spike duplicates</i></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><i>VIII. Laboratory control samples</i></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>IX. Field duplicates</b>				
Were field duplicate pairs identified in this SDG?		/		
Were target compounds detected in the field duplicates?			/	
<b>X. Compound quantitation</b>				
Did the laboratory LOQs/RLs meet the QAPP LOQs/RLs?	/			
Were compound quantitation and RLs adjusted to reflect all sample dilutions and dry weight factors applicable to level IV validation?	/			
<b>XI. Target compound identification</b>				
Were the retention times of reported detects within the RT windows?	/			
<b>XIII. Overall assessment of data</b>				
Overall assessment of data was found to be acceptable.				



LDC #: ~~5-20-2016~~

**VALIDATION FINDINGS WORKSHEET**  
**Continuing Calibration**

Page: 1 of 1  
Reviewer: [Signature]

METHOD:  GC  HPLC

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

Y ~~N~~ N/A Were continuing calibration standards analyzed at the required frequencies?

Y ~~N~~ N/A Did the continuing calibration standards meet the %D validation criteria of ≤20.0%?

Level IV Only

Y ~~N~~ N/A Were the retention times for all calibrated compounds within their respective acceptance windows?

#	Date	Standard ID	Detector/ Column	Compound	%D (Limit)	RT (limit)	Associated Samples	Qualifications
	<del>10/27/16</del>	<del>102T2115</del>	<del>2C</del>	<del>BB</del>	<del>53.7</del>	( )	<del>T</del>	<del>Y/N/A</del>
						( )		
						( )		
	10/27/16	102T2115	2C	BB	24.6	( )	T (dets)	Y/N/A (1c in)
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
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						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		

**VALIDATION FINDINGS WORKSHEET**  
**Surrogate Recovery**

**METHOD:**  GC  HPLC

Are surrogates required by the method? Yes \_\_\_ or No \_\_\_.

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

N N/A Were surrogates spiked into all samples and blanks?  
 N N/A Did all surrogate recoveries (%R) meet the QC limits?

#	Sample ID	Detector/ Column	Surrogate Compound	%R (Limits)	Qualifications
	<u>2</u>	<u>IC</u>	<u>0</u>	<u>138</u> ( <u>40-126</u> )	<u>No qual (10x)</u>
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	

Surrogate Compound	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 1,2-Dinitrobenzene	
C a,a,a-Trifluorotoluene	I Fluorobenzene (FBZ)	O Decachlorobiphenyl (DCB)	U Triphenyltin		
D Bromochlorobenzene	J n-Triacontane	P 1-methylnaphthalene	V Tri-n-propyltin		
E 1,4-Dichlorobutane	K Hexacosane	Q Dichlorophenyl Acetic Acid (DCAA)	W Tributyl Phosphate		
F 1,4-Difluorobenzene (DFB)	L Bromobenzene	R 4-Nitrophenol	X Triphenyl Phosphate		

**VALIDATION FINDINGS WORKSHEET**  
**Internal Standards**

**METHOD:** GC

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

- Y N N/A Were all internal standard area counts within -50 to +100% of the ICAL midpoint standard?  
Y N N/A Were the retention times of the internal standards within +/- 0.05 min seconds of the retention times of the ICAL midpoint standard?

#	Date	Sample ID	Internal Standard	Area (l limits)	RT (l limits)	Qualifications
		6 (dets)	HBB (10)	41 (50 - 200)		↓/U/↓ (BB)
		9 (NO)		47		
		13 (dets)		45		
		14 (dets)		40		
		17 (dets)	↓	49 ↓		↓

HBB - Hexabromobiphenyl

LDC #: 5703B2b

### VALIDATION FINDINGS WORKSHEET Compound Quantitation and Reported CRQLs

METHOD:  GC  HPLC

**Level IV/D Only**

N N/A  
 N N/A  
 N N/A

Were CRQLs adjusted for sample dilutions, dry weight factors, etc.?

Did the reported results for detected target compounds agree within 10.0% of the recalculated results?

Did the relative percent differences of detected compounds between two columns/detectors  $\leq 40\%$ ?

If no, please see findings below.

#	Compound Name	Sample ID	%RPD Between Two Columns/Detectors Limit ( $\leq 40\%$ )	Qualifications
	<u>Aroclor - 1260</u>	<u>A</u>	<u>42.6</u>	<u>Notes/A</u>
	↓	<u>H</u>	<u>45.1</u>	↓
	↓	<u>IT</u>	<u>40.8</u>	↓

**VALIDATION FINDINGS WORKSHEET**  
**Initial Calibration Calculation Verification**

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 (100 std)	CF (100 std)	Ave CF (initial)	Ave CF (initial)	%RSD	%RSD
1	ICAL 7	8/13/21	BB-1 (1c)	0.03587713	0.03587713	0.0359933	0.0359933	2.6	2.6
			BB-1 (2c)	0.06872649	0.0687264	0.06650318	0.06650318	7.7	7.8
2									
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 #: ~~52703330~~

**VALIDATION FINDINGS WORKSHEET**  
**Continuing Calibration Results Verification**

Page: 1 of 1  
 Reviewer: [Signature]

METHOD:  GC\_HPLC

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

% Difference = 100 \* (ave. CF - CF)/ave.CF

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

#	Standard ID	Calibration Date	Compound	Average CF(Ical)/ CCV Conc.	Reported	Recalculated	Reported	Recalculated
					CF/ Conc. CCV	CF/ Conc. CCV	%D	%D
1	<del>10242201</del>	10/25/21	<del>BB1 (1C)</del>	0.03599233	0.0304158	0.0304157	15.2	15.2
	1-2. MB		BB1 (2C)	0.06650318	0.0539459	0.0539459	18.8	18.8
2	<del>1024214T</del>	10/25 12:06	↓	0.03599233	0.0298223	0.0298223	17.2	17.2
	3-6.8-10		↓	0.06650318	0.052425	0.052425	21.2	21.2
3	<del>1024215T</del>	10/25/21 15:31	↓	0.03599233	0.0330805	0.0330804	8.0	8.0
	11-1T		↓	0.06650318	0.0552499	0.0552498	16.8	16.8
4	<del>1024212</del>	10/27/21 10:55	↓	0.03599233	0.0286261	0.028626	20.4	20.5
	T		↓	0.06650318	0.0488190	0.0488189	26.4	26.5

## 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	
DCB	IC	40.0	36.9	92.2	92.2	
TCX	↓	↓	27.7	69.4	69.4	

Sample ID: \_\_\_\_\_

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

Sample ID: \_\_\_\_\_

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

## VALIDATION FINDINGS WORKSHEET

### Matrix Spike/Matrix Spike Duplicates Results Verification

METHOD:  GC  HPLC

The percent recoveries (%R) and relative percent differences (RPD) of the matrix spike and matrix spike duplicate were recalculated for the compounds identified below using the following calculation:

$\% \text{Recovery} = 100 * (\text{SSC} - \text{SC}) / \text{SA}$

Where

SSC = Spiked sample concentration

SC = Sample concentration

SA = Spike added

MS = Matrix spike

MSD = Matrix spike duplicate

$\text{RPD} = ((\text{SSCMS} - \text{SSCMSD}) * 2) / (\text{SSCMS} + \text{SSCMSD}) * 100$

MS/MSD samples: 18/19

Compound	Spike Added ( <u>MS</u> )		Sample Conc. ( <u>MS</u> )	Spike Sample Concentration ( <u>MS</u> )		Matrix spike		Matrix Spike Duplicate		MS/MSD	
	MS	MSD	---	MS	MSD	Percent Recovery		Percent Recovery		RPD	
						Reported	Recalc.	Reported	Recalc.	Reported	Recalc.
Gasoline (8015)											
Diesel (8015)											
Benzene (8021B)											
Methane (RSK-175)											
2,4-D (8151)											
Dinoseb (8151)											
Naphthalene (8310)											
Anthracene (8310)											
HMX (8330)											
2,4,6-Trinitrotoluene (8330)											
<u>FEB-1260</u>	<u>101</u>	<u>101</u>	<u>4.2</u>	<u>86.3</u>	<u>93.8</u>	<u>81.3</u>	<u>81.3</u>	<u>88.7</u>	<u>88.7</u>	<u>8.37</u>	<u>8.3</u>

Comments: Refer to Matrix Spike/Matrix Spike Duplicates 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

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

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 compounds identified below using the following calculation:

$\% \text{Recovery} = 100 * (\text{SSC} - \text{SC}) / \text{SA}$

Where SSC = Spiked sample concentration

SC = Sample concentration

SA = Spike added

$\text{RPD} = ((\text{SSCLCS} - \text{SSCLCSD}) * 2) / (\text{SSCLCS} + \text{SSCLCSD}) * 100$

LCS = Laboratory Control Sample

LCSD = Laboratory Control Sample duplicate

LCS/LCSD samples: ~~BND629-PS1-B501~~

Compound	Spike Added ( <u>101</u> )		Spike Sample Concentration ( <u>101</u> )		LCS		LCSD		LCS/LCSD	
	LCS	LCSD	LCS	LCSD	Percent Recovery		Percent Recovery		RPD	
					Reported	Recalc.	Reported	Recalc.	Reported	Recalc.
Gasoline (8015)										
Diesel (8015)										
Benzene (8021B)										
Methane (RSK-175)										
2,4-D (8151)										
Dinoseb (8151)										
Naphthalene (8310)										
Anthracene (8310)										
HMX (8330)										
2,4,6-Trinitrotoluene (8330)										
<u>PCB-1260</u>	<u>101</u>	<u>101</u>	<u>84.4</u>	<u>87.6</u>	<u>83.7</u>	<u>83.6</u>	<u>82.9</u>	<u>82.8</u>	<u>1.02</u>	<u>1.0</u>

Comments: Refer to Laboratory Control Sample/Laboratory Control Sample Duplicate 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**  
**Sample Calculation Verification**

METHOD:  GC  HPLC

Y N N/A  
Y N N/A

Were all reported results recalculated and verified for all level IV samples?

Were all recalculated results for detected target compounds within 10% of the reported results?

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

Example:

Sample ID. 1 Compound Name PCB-1260-1

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

Concentration =  $\frac{(37659)(80)}{(356030)(0.03599233)} = 235!$

concentration =  $\frac{(235! + 180! + 191! + 202! + 235!)(2.5)}{5 \times 18.28 \times 0.6844} = 41.7 \mu\text{g}$

#	Sample ID	Compound	Reported Concentrations ( <u>14.78</u> )	Recalculated Results Concentrations ( )	Qualifications
	<u>1</u>	<u>PCB-1260</u>	<u>41.7</u>		

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

## Laboratory Data Consultants, Inc. Data Validation Report

**Project/Site Name:** Duwamish AOC4

**LDC Report Date:** December 15, 2021

**Parameters:** Metals

**Validation Level:** Stage 4

**Laboratory:** Analytical Resources, Inc., Tukwila, WA

**Sample Delivery Group (SDG):** 21J0134

<b>Sample Identification</b>	<b>Laboratory Sample Identification</b>	<b>Matrix</b>	<b>Collection Date</b>
LDW21-SS600	21J0134-11	Sediment	07/12/21
LDW21-SC587A	21J0134-14	Sediment	07/12/21
LDW21-SC587F	21J0134-15	Sediment	07/12/21
LDW21-IT588F	21J0134-17	Sediment	07/14/21
LDW21-IT585F	21J0134-18	Sediment	07/14/21
LDW21-SS600MS	21J0134-11MS	Sediment	07/12/21
LDW21-SS600MSD	21J0134-11MSD	Sediment	07/12/21
LDW21-SS600DUP	21J0134-11DUP	Sediment	07/12/21

## 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 Final Lower Duwamish Waterway Quality Assurance Project Plan for Remedial Design of Upper Reach: Pre-Design Investigation (May 2020) and a modified outline of the USEPA National Functional Guidelines (NFG) for Inorganic Superfund Methods Data Review (January 2017). 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 methods:

Arsenic, Cadmium, Chromium, Copper, Lead, Silver, and Zinc by Environmental Protection Agency (EPA) SW 846 Method 6020B  
Mercury by EPA SW 846 Method 7471B

All sample results were subjected to Stage 4 evaluation, which is comprised of the quality control (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): 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.
- U (Non-detected): The analyte was analyzed for and positively identified by the laboratory; however the analyte should be considered non-detected at the reported concentration due to the presence of contaminants detected in the associated blank(s).
- UJ (Non-detected estimated): The analyte was reported as not detected by the laboratory; however the reported quantitation/detection limit is estimated due to non-conformances discovered during data validation.
- R (Rejected): The sample results were rejected due to gross non-conformances discovered during data validation. Data qualified as rejected is not usable.
- 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.

## I. Sample Receipt and Technical Holding Times

All samples were received in good condition.

All technical holding time requirements were met with the following exceptions:

Sample	Analyte	Total Days From Sample Collection Until Analysis	Required Holding Time (in Days) From Sample Collection Until Analysis	Flag	A or P
LDW21-SS600 LDW21-SS600DUP	Mercury	106	28	J (all detects)	P

## II. ICPMS Tune

The mass calibration was within 0.1 AMU and the percent relative standard deviation (%RSD) was less than or equal to 5%.

## III. Instrument Calibration

Initial and continuing calibrations were performed as required by the methods.

The initial calibration verification (ICV) and continuing calibration verification (CCV) standards were within QC limits.

## IV. ICP Interference Check Sample Analysis

The frequency of interference check sample (ICS) analysis was met. All criteria were within QC limits.

## V. Laboratory Blanks

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

## VI. Field Blanks

No field blanks were identified in this SDG.

## VII. Matrix Spike/Matrix Spike Duplicates

Matrix spike (MS) and matrix spike duplicate (MSD) sample analysis was performed on an associated project sample. Percent recoveries (%R) were within QC limits. Relative percent differences (RPD) were within QC limits.

## VIII. Duplicate Sample Analysis

Duplicate (DUP) sample analysis was performed on an associated project sample. Results were within QC limits.

## **IX. Serial Dilution**

Serial dilution was not performed for this SDG.

## **X. Laboratory Control Samples**

Laboratory control samples (LCS) were analyzed as required by the methods. Percent recoveries (%R) were within QC limits.

## **XI. Field Duplicates**

No field duplicates were identified in this SDG.

## **XII. Internal Standards (ICP-MS)**

All internal standard percent recoveries (%R) were within QC limits.

## **XIII. Target Analyte Quantitation**

All target analyte quantitations were within validation criteria.

## **XIV. Overall Assessment of Data**

The analysis was conducted within all specifications of the methods. No results were rejected in this SDG.

Due to technical holding time, data were qualified as estimated in two samples.

The quality control criteria reviewed, other than those discussed above, were met and are considered acceptable.

**Duwamish AOC4  
Metals - Data Qualification Summary - SDG 21J0134**

Sample	Analyte	Flag	A or P	Reason
LDW21-SS600 LDW21-SS600DUP	Mercury	J (all detects)	P	Technical holding times

**Duwamish AOC4  
Metals - Laboratory Blank Data Qualification Summary - SDG 21J0134**

No Sample Data Qualified in this SDG

**Duwamish AOC4  
Metals - Field Blank Data Qualification Summary - SDG 21J0134**

No Sample Data Qualified in this SDG

LDC #: 52703B4a

**VALIDATION COMPLETENESS WORKSHEET**

Date: 2/12/21

SDG #: 21J0134

Stage 4

Page: 1 of 1

Laboratory: Analytical Resources, Inc., Tukwila, WA

Reviewer: [Signature]

2nd Reviewer: [Signature]

**METHOD:** Metals (EPA SW846 Method 6020B) (7471B)

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	A	ASW
II.	ICP/MS Tune	A	
III.	Instrument Calibration	A	
IV.	ICP Interference Check Sample (ICS) Analysis	A	
V.	Laboratory Blanks	A	
VI.	Field Blanks	N	
VII.	Matrix Spike/Matrix Spike Duplicates	A	
VIII.	Duplicate sample analysis	A	
IX.	Serial Dilution	N	
X.	Laboratory control samples	A	LCS
XI.	Field Duplicates	N	
XII.	Internal Standard (ICP-MS)	A	
XIII.	Target Analyte Quantitation	A	
XIV.	Overall Assessment of Data	A	

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:

	Client ID	Lab ID	Matrix	Date
1	LDW21-SS600	21J0134-11	Sediment	07/12/21
2	LDW21-SC587A	21J0134-14	Sediment	07/12/21
3	LDW21-SC587F	21J0134-15	Sediment	07/12/21
4	LDW21-IT588F	21J0134-17	Sediment	07/14/21
5	LDW21-IT585F	21J0134-18	Sediment	07/14/21
6	LDW21-SS600MS	21J0134-11MS	Sediment	07/12/21
7	LDW21-SS600MSD	21J0134-11MSD	Sediment	07/12/21
8	LDW21-SS600DUP	21J0134-11DUP	Sediment	07/12/21
9				
10				
11				
12				
13				

Notes: \_\_\_\_\_



METHOD: Trace Metals (EPA SW 846 Methods 6010/6020/7000)				
Validation Area	Yes	No	NA	Comments
<b>I. Technical holding times</b>				
Were all technical holding times met?		X		
Were all water samples preserved to a pH of <2?			X	
<b>II. ICP-MS Tune</b>				
Were mass resolutions within 0.1 amu for all isotopes in the tuning solution?	X			
Were %RSDs of isotopes in the tuning solution ≤5%?	X			
<b>III. Calibration</b>				
Were all instruments calibrated daily?	X			
Were the proper standards used?	X			
Were all initial and continuing calibration verifications within the 90-110% (80-120% for mercury) QC limits?	X			
Were the low level standard checks within 70-130%?			X	
Were all initial calibration correlation coefficients within limits as specified by the method?	X			
<b>IV. Blanks</b>				
Was a method blank associated with every sample in this SDG?	X			
Was there contamination in the method blanks?		X		
Was there contamination in the initial and continuing calibration blanks?		X		
<b>V. Interference Check Sample</b>				
Were the interference check samples performed daily?	X			
Were the AB solution recoveries within 80-120%?	X			
<b>VI. Matrix Spike/Matrix Spike Duplicates/Laboratory Duplicates</b>				
Were MS/MSD recoveries with the QC limits? (If the sample concentration exceeded the spike concentration by a factor of 4, no action was taken.)	X			
Were the MS/MSD or laboratory duplicate relative percent differences (RPDs) within the QC limits?	X			
<b>VII. Laboratory Control Samples</b>				
Was a LCS analyzed for each batch in the SDG?	X			
Were the LCS recoveries and RPDs (if applicable) within QC limits?	X			

METHOD: Trace Metals (EPA SW 846 Methods 6010/6020/7000)				
Validation Area	Yes	No	NA	Comments
<b>VIII. Internal Standards</b>				
Were all percent recoveries within the 30-120% (60-125% for EPA Method 200.8) QC limits?	X			
If the recoveries were outside the limits, was a reanalysis performed?		X		
<b>IX. Serial Dilution</b>				
Were all percent differences <10%?			X	
Was there evidence of negative interference? If yes, professional judgement will be used to qualify the data.			X	
<b>X. Sample Result Verification</b>				
Were all reporting limits adjusted to reflect sample dilutions?	X			
Were all soil samples dry weight corrected?	X			
<b>XI. Overall Assessment of Data</b>				
Was the overall assessment of the data found to be acceptable?	X			
<b>XII. Field Duplicates</b>				
Were field duplicates identified in this SDG?		X		
Were target analytes detected in the field duplicates?			X	
<b>XIII. Field Blanks</b>				
Were field blanks identified in this SDG?		X		
Were target analytes detected in the field blanks?			X	

All elements are applicable to each sample as noted below.

Sample ID	Target Analyte List
1	As, Cd, Cr, Cu, Pb, Ag, Zn, Hg
2 to 5	As
QC: 6-8	As, Cd, Cr, Cu, Pb, Ag, Zn, Hg

**Analysis Method**

ICP	
ICP-MS	As, Cd, Cr, Cu, Pb, Ag, Zn
CVAA	Hg



METHOD: Trace Metals (EPA SW 846 Methods 6010/6020/7000)

An initial calibration verification (ICV), continuing calibration verification (CCV), low level calibration check (LLCC), and interference check sample (ICSAB) percent recovery (%R) was recalculated for each type of analysis using the following formula:

$$\%R = (\text{Found}/\text{True}) \times 100$$

Found = concentration of each analyte measured in the analysis

True = concentration of each analyte in the source

Standard ID	Type of Analysis	Element	Found (ug/L)	True (ug/L)	Recalculated %R	Reported %R	Acceptable (Y/N)
ICV	ICP-MS	As	47.7	50	95.4	95.5	Y
CCV	ICP-MS	As	50.2	50	100	100	Y
ICSAB	ICP-MS	As	19.283	20	96.4	96.4	Y
ICV	CVAA	Hg	4.1178	4	103	103	Y
CCV	CVAA	Hg	4.0688	4	102	102	Y

ICP-MS Tune	QC Parameter	Mass	Actual	Required
10/28/2021	Mass Axis	115	114.9	± 0.1 amu
10/28/2021	%RSD	115	1	≤ 5%

VALIDATION FINDINGS CHECKLIST  
Quality Control Sample Recalculations

METHOD: Trace Metals (EPA SW 846 Methods 6010/6020/7000)

Percent recoveries (%R) for the laboratory control sample (LCS), matrix spike (MS), and post digestion spike (PDS) were recalculated using the following formula:

$$\%R = (\text{Found}/\text{True}) \times 100$$

Found = concentration of each analyte measured in the analysis. For the MS calculation, Found = SSR (Spiked Sample Result) - SR (Sample Result)

True = concentration of each analyte in the source

The sample and duplicate relative percent difference (RPD) was recalculated using the following formula:

$$\text{RPD} = (\text{Absolute value}(S-D) \times 200) / (S+D)$$

S = Original sample concentration

D = Duplicate sample concentration

The serial dilution percent difference (%D) was recalculated using the following formula.

$$\%D = (\text{Absolute value}(I - \text{SDR})) \times 100 / (I)$$

I = Initial sample result

SDR = Serial dilution result (with a 5x dilution applied)

Sample ID	Type of Analysis	Element	Found/S/I	True/D/SDR	Recalculated %R/RPD/%D	Reported %R/RPD/%D	Acceptable (Y/N)
LCS	LCS	As	24.1	25	96.4	96.5	Y
6	MS	Cd	39.66	39.5	100	100	
8	Duplicate	Cu	28.3	29	2.44	2.54	
	PDS						
	Serial dilution						



## Laboratory Data Consultants, Inc. Data Validation Report

**Project/Site Name:** Duwamish AOC4

**LDC Report Date:** December 15, 2021

**Parameters:** Wet Chemistry

**Validation Level:** Stage 4

**Laboratory:** Analytical Resources, Inc., Tukwila, WA

**Sample Delivery Group (SDG):** 21J0134

Sample Identification	Laboratory Sample Identification	Matrix	Collection Date
LDW21-IT669D	21J0134-01	Sediment	07/08/21
LDW21-IT598B	21J0134-02	Sediment	07/08/21
LDW21-IT598C	21J0134-03	Sediment	07/08/21
LDW21-IT598D	21J0134-04	Sediment	07/08/21
LDW21-IT598E	21J0134-05	Sediment	07/08/21
LDW21-IT598F	21J0134-06	Sediment	07/08/21
LDW21-IT598G	21J0134-07	Sediment	07/08/21
LDW21-IT598H	21J0134-08	Sediment	07/08/21
LDW21-SC553D	21J0134-09	Sediment	07/09/21
LDW21-SC554D	21J0134-10	Sediment	07/09/21
LDW21-SS600	21J0134-11	Sediment	07/12/21
LDW21-SS681	21J0134-12	Sediment	07/12/21
LDW21-SS641	21J0134-13	Sediment	07/09/21
LDW21-SC587A	21J0134-14	Sediment	07/12/21
LDW21-SC587F	21J0134-15	Sediment	07/12/21
LDW21-IT660C	21J0134-16	Sediment	07/14/21
LDW21-IT588F	21J0134-17	Sediment	07/14/21
LDW21-IT585F	21J0134-18	Sediment	07/14/21
LDW21-SC568F	21J0134-19	Sediment	07/14/21
LDW21-IT585FMS	21J0134-18MS	Sediment	07/14/21
LDW21-IT585FDUP1	21J0134-18DUP1	Sediment	07/14/21
LDW21-IT585FDUP2	21J0134-18DUP2	Sediment	07/14/21



## 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 Final Lower Duwamish Waterway Quality Assurance Project Plan for Remedial Design of Upper Reach: Pre-Design Investigation (May 2020) and a modified outline of the USEPA National Functional Guidelines (NFG) for Inorganic Superfund Methods Data Review (January 2017). 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 methods:

Total Organic Carbon by Environmental Protection Agency (EPA) SW 846 Method 9060A

Total Solids by Standard Method 2540G

All sample results were subjected to Stage 4 data validation, which is comprised of the quality control (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): 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.
- U (Non-detected): The analyte was analyzed for and positively identified by the laboratory; however the analyte should be considered non-detected at the reported concentration due to the presence of contaminants detected in the associated blank(s).
- UJ (Non-detected estimated): The analyte was reported as not detected by the laboratory; however the reported quantitation/detection limit is estimated due to non-conformances discovered during data validation.
- R (Rejected): The sample results were rejected due to gross non-conformances discovered during data validation. Data qualified as rejected is not usable.
- 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.

## I. Sample Receipt and Technical Holding Times

All samples were received in good condition.

All technical holding time requirements were met.

## II. Initial Calibration

All criteria for the initial calibration of each method were met.

## III. Continuing Calibration

Continuing calibration frequency and analysis criteria were met for each method when applicable.

## IV. Laboratory Blanks

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

## V. Field Blanks

No field blanks were identified in this SDG.

## VI. Matrix Spike/Matrix Spike Duplicates

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

## VII. Duplicate Sample Analysis

Duplicate (DUP) sample analysis was performed on an associated project sample. Results were within QC limits with the following exceptions:

DUP ID (Associated Samples)	Analyte	RPD (Limits)	Difference (Limits)	Flag	A or P
LDW21-IT585FDUP1 (All samples in SDG 21J0134)	Total organic carbon	22.1 (≤20)	-	J (all detects)	A

## VIII. Laboratory Control Samples

Laboratory control samples (LCS) were analyzed as required by the methods. Percent recoveries (%R) were within QC limits.

## IX. Field Duplicates

No field duplicates were identified in this SDG.

## **X. Target Analyte Quantitation**

All target analyte quantitations were within validation criteria.

## **XI. Overall Assessment of Data**

The analysis was conducted within all specifications of the methods. No results were rejected in this SDG.

Due to DUP RPD, data were qualified as estimated in twenty-one samples.

The quality control criteria reviewed, other than those discussed above, were met and are considered acceptable.

**Duwamish AOC4  
Wet Chemistry - Data Qualification Summary - SDG 21J0134**

Sample	Analyte	Flag	A or P	Reason
LDW21-IT669D LDW21-IT598B LDW21-IT598C LDW21-IT598D LDW21-IT598E LDW21-IT598F LDW21-IT598G LDW21-IT598H LDW21-SC553D LDW21-SC554D LDW21-SS600 LDW21-SS681 LDW21-SS641 LDW21-SC587A LDW21-SC587F LDW21-IT660C LDW21-IT588F LDW21-IT585F LDW21-SC568F LDW21-IT585FDUP1 LDW21-IT585FDUP2	Total organic carbon	J (all detects)	A	Duplicate sample analysis (RPD)

**Duwamish AOC4  
Wet Chemistry - Laboratory Blank Data Qualification Summary - SDG 21J0134**

No Sample Data Qualified in this SDG

**Duwamish AOC4  
Wet Chemistry - Field Blank Data Qualification Summary - SDG 21J0134**

No Sample Data Qualified in this SDG

**METHOD: (Analyte) TOC (EPA SW846 Method 9060A), Total Solids (SM2540G)**

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	AA	
II	Initial calibration	A	
III.	Calibration verification	A	
IV	Laboratory Blanks	A	
V	Field blanks	N	
VI.	Matrix Spike/Matrix Spike Duplicates	A	
VII.	Duplicate sample analysis	SW	
VIII.	Laboratory control samples	A	LCS
IX.	Field duplicates	N	
X.	Target Analyte Quantitation	A	
XI	Overall assessment of data	A	

Note: A = Acceptable ND = No compounds detected D = Duplicate SB=Source blank  
 N = Not provided/applicable R = Rinsate TB = Trip blank OTHER:  
 SW = See worksheet FB = Field blank EB = Equipment blank

	Client ID	Lab ID	Matrix	Date
1	LDW21-IT669D	21J0134-01	Sediment	07/08/21
2	LDW21-IT598B	21J0134-02	Sediment	07/08/21
3	LDW21-IT598C	21J0134-03	Sediment	07/08/21
4	LDW21-IT598D	21J0134-04	Sediment	07/08/21
5	LDW21-IT598E	21J0134-05	Sediment	07/08/21
6	LDW21-IT598F	21J0134-06	Sediment	07/08/21
7	LDW21-IT598G	21J0134-07	Sediment	07/08/21
8	LDW21-IT598H	21J0134-08	Sediment	07/08/21
9	LDW21-SC553D	21J0134-09	Sediment	07/09/21
10	LDW21-SC554D	21J0134-10	Sediment	07/09/21
11	LDW21-SS600	21J0134-11	Sediment	07/12/21
12	LDW21-SS681	21J0134-12	Sediment	07/12/21
13	LDW21-SS641	21J0134-13	Sediment	07/09/21
14	LDW21-SC587A	21J0134-14	Sediment	07/12/21
15	LDW21-SC587F	21J0134-15	Sediment	07/12/21
16	LDW21-IT660C	21J0134-16	Sediment	07/14/21
17	LDW21-IT588F	21J0134-17	Sediment	07/14/21

LDC #: 52703B6      **VALIDATION COMPLETENESS WORKSHEET**  
 SDG #: 21J0134      Stage 4  
 Laboratory: Analytical Resources, Inc., Tukwila, WA

Date: 12/9/21  
 Page: 2 of 2  
 Reviewer: [Signature]  
 2nd Reviewer: [Signature]

**METHOD: (Analyte) TOC (EPA SW846 Method 9060A), Total Solids (SM2540G)**

	Client ID	Lab ID	Matrix	Date
18	LDW21-IT585F	21J0134-18	Sediment	07/14/21
19	LDW21-SC568F	21J0134-19	Sediment	07/14/21
20	LDW21-IT585FMS	21J0134-18MS	Sediment	07/14/21
21	LDW21-IT585FDUP 1	21J0134-18DUP	Sediment	07/14/21
22	LDW21-IT585FTRP <u>DURZ</u>	21J0134-18TRP	Sediment	07/14/21
23				
24				
25				

Notes: \_\_\_\_\_  
 \_\_\_\_\_  
 \_\_\_\_\_

METHOD: Inorganics				
Validation Area	Yes	No	NA	Comments
<b>I. Technical holding times</b>				
Were all technical holding times were met?	X			Frozen
<b>II. Calibration</b>				
Were all instruments calibrated at the required frequency?	X			
Were the proper number of standards used?	X			
Were all initial and continuing calibration verifications within the QC limits?	X			
Were all initial calibration correlation coefficients within limits as specified by the method?	X			
Were balance checks performed as required?	X			
<b>III. Blanks</b>				
Was a method blank associated with every sample in this SDG?	X			
Was there contamination in the method blanks?		X		
Was there contamination in the initial and continuing calibration blanks?		X		
<b>IV. Matrix Spike/Matrix Spike Duplicates/Laboratory Duplicates</b>				
Were MS/MSD recoveries with the QC limits? (If the sample concentration exceeded the spike concentration by a factor of 4, no action was taken.)	X			
Were the MS/MSD or laboratory duplicate relative percent differences (RPDs) within the QC limits?		X		
<b>V. Laboratory Control Samples</b>				
Was a LCS analyzed for each batch in the SDG?	X			
Were the LCS recoveries and RPDs (if applicable) within QC limits?	X			
<b>X. Sample Result Verification</b>				
Were all reporting limits adjusted to reflect sample dilutions?	X			
Were all soil samples dry weight corrected?	X			
<b>XI. Overall Assessment of Data</b>				
Was the overall assessment of the data found to be acceptable?	X			



METHOD: Inorganics				
Validation Area	Yes	No	NA	Comments
<b>XII. Field Duplicates</b>				
Were field duplicates identified in this SDG?		X		
Were target analytes detected in the field duplicates?			X	
<b>XIII. Field Blanks</b>				
Were field blanks identified in this SDG?		X		
Were target analytes detected in the field blanks?			X	

All elements are applicable to each sample as noted below.

Sample ID	Target Analyte List
All	TS, TOC
QC:	
20	TOC
21	TS, TOC
22	TS

Laboratory Duplicates

Reviewer:CR

**METHOD: Inorganics**

Laboratory duplicate analysis was performed by the laboratory. All laboratory duplicates were with the relative percent difference (RPD) for samples >5X the reporting limits with the exceptions listed below. If samples were <5X the reporting limits, the difference was with 1X the reporting limit for water samples and within 2X the reporting limit for soil samples for all samples with the exceptions listed below.

Duplicate ID	Matrix	Analyte	RPD	RPD Limit	Difference (units)	Difference Limit	Associated Samples	Qualification	Det/ND
21	s	TOC	22.1	20			All	J/UJ/A	Det

Comments:

**Validation Findings Worksheet**  
**Initial and Continuing Calibration Calculation Verification**

**Method:** Inorganics, Method See Cover

An initial or continuing calibration verification percent recovery (%R) was recalculated for each type of analysis using the following formula:

$$\%R = \frac{\text{Found} \times 100}{\text{True}}$$

Where, Found = concentration of each analyte measured in the analysis of the ICV or CCV solution  
 True = concentration of each analyte in the ICV or CCV source

Calibration verification	TOC	ICV	44.446	44.742	101	101	Y
Calibration verification	TOC	CCV	44.446	44.325	100	100	Y
Calibration verification	TOC	CCV	44.446	44.814	101	101	Y

Comments:

VALIDATION FINDINGS CHECKLIST  
Quality Control Sample Recalculations

METHOD: Inorganics

Percent recoveries (%R) for the laboratory control sample (LCS) and matrix spike (MS) were recalculated using the following formula.

$$\%R = (\text{Found}/\text{True}) \times 100$$

Found = concentration of each analyte measured in the analysis. For the MS calculation, Found = SSR (Spiked Sample Result) - SR (Sample Result)

True = concentration of each analyte in the source

The sample and duplicate relative percent difference (RPD) was recalculated using the following formula.

$$\text{RPD} = (\text{Absolute value}(S-D) \times 200) / (S+D)$$

S = Original sample concentration

D = Duplicate sample concentration

Sample ID	Type of Analysis	Element	Found/S	True/D	Recalculated %R/RPD	Reported %R/RPD	Acceptable (Y/N)
LCS	LCS	TOC	44.6	44.4	100	100	Y
24	MS	TOC	0.88	0.876	100	101	Y
21	Duplicate	TS	77.89	77.29	0.773	0.775	Y



**Laboratory Data Consultants, Inc.  
Data Validation Report**

**Project/Site Name:** Duwamish AOC4  
**LDC Report Date:** December 15, 2021  
**Parameters:** Polychlorinated Dioxins/Dibenzofurans  
**Validation Level:** Stage 4  
**Laboratory:** Analytical Resources, Inc., Tukwila, WA

**Sample Delivery Group (SDG):** 21J0134

<b>Sample Identification</b>	<b>Laboratory Sample Identification</b>	<b>Matrix</b>	<b>Collection Date</b>
LDW21-IT669D	21J0134-01	Sediment	07/08/21
LDW21-SS641	21J0134-13	Sediment	07/09/21
LDW21-IT660C	21J0134-16	Sediment	07/14/21

## 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 Final Lower Duwamish Waterway Quality Assurance Project Plan for Remedial Design of Upper Reach: Pre-Design Investigation (May 2020) and a modified outline of the USEPA National Functional Guidelines (NFG) for High Resolution Superfund Methods Data Review (April 2016). 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:

Polychlorinated Dioxins/Dibenzofurans by Environmental Protection Agency (EPA) Method 1613B

All sample results were subjected to Stage 4 data validation, which is comprised of the quality control (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): 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.
- U (Non-detected): The analyte was analyzed for and positively identified by the laboratory; however the analyte should be considered not detected at the reported concentration due to the presence of contaminants detected in the associated blank(s).
- UJ (Non-detected estimated): The analyte was reported as not detected by the laboratory; however the reported quantitation/detection limit is estimated due to non-conformances discovered during data validation.
- R (Rejected): The sample results were rejected due to gross non-conformances discovered during data validation. Data qualified as rejected is not usable.
- 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.



## **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. HRGC/HRMS Instrument Performance Check**

Instrument performance was checked at the required frequency.

Retention time windows were established for all homologues. The chromatographic resolution between 2,3,7,8-TCDD and peaks representing any other unlabeled TCDD isomer was less than or equal to 25%.

The static resolving power was at least 10,000 (10% valley definition).

## **III. Initial Calibration and Initial Calibration Verification**

A five point 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 and less than or equal to 35.0% for labeled compounds.

The ion abundance ratios for all PCDDs and PCDFs were within validation criteria.

The minimum S/N ratio was greater than or equal to 10 for each analyte and labeled compound.

The percent differences (%D) of the initial calibration verification (ICV) standard were within the QC limits for all analytes and labeled compounds.

## **IV. Continuing Calibration**

Continuing calibration was performed at the required frequencies.

All of the continuing calibration results were within the QC limits for all analytes and labeled compounds.

The ion abundance ratios for all PCDDs and PCDFs were within validation criteria.

The minimum S/N ratio was greater than or equal to 10 for each analyte and labeled compound.

## **V. Laboratory Blanks**

Laboratory blanks were analyzed as required by the method. No contaminants were found in the laboratory blanks with the following exceptions:

Blank ID	Extraction Date	Analyte	Concentration	Associated Samples
BJJ0500-BLK1	10/19/21	OCDD Total HxCDF	0.981 ng/Kg 0.100 ng/Kg	All samples in SDG 21J0134

Sample concentrations were compared to concentrations detected in the laboratory blanks. The sample concentrations were either not detected or were significantly greater than the concentrations found in the associated laboratory blanks.

## VI. Field Blanks

No field blanks were identified in this SDG.

## VII. Matrix Spike/Matrix Spike Duplicates

The laboratory has indicated that there were no matrix spike (MS) and matrix spike duplicate (MSD) analyses specified for the samples in this SDG, and therefore matrix spike and matrix spike duplicate analyses were not performed for this SDG.

## VIII. Laboratory Control Samples/Standard Reference Materials

Laboratory control samples (LCS) were analyzed as required by the method. Percent recoveries (%R) were within QC limits.

Standard reference materials (SRM) were analyzed as required by the method. The results were within QC limits.

## IX. Field Duplicates

No field duplicates were identified in this SDG.

## X. Internal Standards

All internal standard areas and retention times were within QC limits.

## XI. Target Analyte Quantitation

All target analyte quantitations met validation criteria with the following exceptions:

Sample	Analyte	Flag	A or P
All samples in SDG 21J0134	All analytes reported as estimated maximum possible concentration (EMPC) and greater than the reporting limit (RL).	J (all detects)	A

Sample	Analyte	Flag	A or P
All samples in SDG 21J0134	All analytes reported as estimated maximum possible concentration (EMPC) and less than the reporting limit (RL).	U	A

Sample	Analyte	Finding	Criteria	Flag	A or P
LDW21-IT669D	OCDD	Sample result exceeded calibration range.	Reported result should be within calibration range.	J (all detects)	P

**XII. Target Analyte Identification**

All target analyte identifications met validation criteria.

**XIII. System Performance**

The system performance was acceptable.

**XIV. Overall Assessment of Data**

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

Due to results reported by the laboratory as EMPCs and results exceeding calibration range, data were qualified as estimated in three samples.

The quality control criteria reviewed, other than those discussed above, were met and are considered acceptable.

**Duwamish AOC4  
Polychlorinated Dioxins/Dibenzofurans - Data Qualification Summary - SDG 21J0134**

Sample	Analyte	Flag	A or P	Reason
LDW21-IT669D LDW21-SS641 LDW21-IT660C	All analytes reported as estimated maximum possible concentration (EMPC) and greater than the reporting limit (RL).	J (all detects)	A	Target analyte quantitation (EMPC)
LDW21-IT669D LDW21-SS641 LDW21-IT660C	All analytes reported as estimated maximum possible concentration (EMPC) and less than the reporting limit (RL).	U	A	Target analyte quantitation (EMPC)
LDW21-IT669D	OCDD	J (all detects)	P	Target analyte quantitation (exceeding range)

**Duwamish AOC4  
Polychlorinated Dioxins/Dibenzofurans - Laboratory Blank Data Qualification Summary - SDG 21J0134**

No Sample Data Qualified in this SDG

**Duwamish AOC4  
Polychlorinated Dioxins/Dibenzofurans - Field Blank Data Qualification Summary - SDG 21J0134**

No Sample Data Qualified in this SDG

LDC #: 52703B21  
 SDG #: 21J0134  
 Laboratory: Analytical Resources, Inc., Tukwila, WA

**VALIDATION COMPLETENESS WORKSHEET**

Stage 4

Date: 1/6/21  
 Page: 1 of 1  
 Reviewer: [Signature]  
 2nd Reviewer: JVB

**METHOD:** HRGC/HRMS Polychlorinated Dioxins/Dibenzofurans (EPA Method 1613B)

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	A	
II.	HRGC/HRMS Instrument performance check	A	
III.	Initial calibration/ICV	A/A	RSD < 20/35/70 10% < RCLimits
IV.	Continuing calibration	A	CCV < RCLimits
V.	Laboratory Blanks	W	
VI.	Field blanks	N	
VII.	Matrix spike/Matrix spike duplicates	N	CS
VIII.	Laboratory control samples /SRM	A	LC9
IX.	Field duplicates	N	
X.	Internal standards	A	
XI.	Target analyte quantitation	A	
XII.	Target analyte identification	A	
XIII.	System performance	A	
XIV.	Overall assessment of data	A	

Note: A = Acceptable      ND = No compounds detected      D = Duplicate      SB=Source blank  
 N = Not provided/applicable      R = Rinsate      TB = Trip blank      OTHER:  
 SW = See worksheet      FB = Field blank      EB = Equipment blank

	Client ID	Lab ID	Matrix	Date
1	LDW21-IT669D	21J0134-01	Sediment	07/08/21
2	LDW21-SS641	21J0134-13	Sediment	07/09/21
3	LDW21-IT660C	21J0134-16	Sediment	07/14/21
4				
5				
6				
7				
8				
9				
10				

Notes:

BN 0500-2K1				

**Method:** HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 1613B)

Validation Area	Yes	No	NA	Findings/Comments
<b>I. Technical holding times</b>				
All technical holding times were met.	√			
Cooler temperature criteria were met.	√			
<b>II. GC/MS Instrument performance check</b>				
Was PFK exact mass 380.9760 verified?	√			
Were the retention time windows established for all homologues?	√			
Was the chromatographic resolution between 2,3,7,8-TCDD and peaks representing any other unlabeled TCDD isomers $\leq 25\%$ ?	√			
Is the static resolving power at least 10,000 (10% valley definition)?	√			
Was the mass resolution adequately check with PFK?	√			
Was the presence of 1,2,8,9-TCDD and 1,3,4,6,8-PeCDF verified?	√			
<b>III. Initial calibration and Initial calibration verification</b>				
Was the initial calibration performed at 5 concentration levels?	√			
Were all percent relative standard deviations (%RSD) $\leq 20\%$ for unlabeled compounds and $\leq 35\%$ for unlabeled compounds?	√			
Did all calibration standards meet the Ion Abundance Ratio criteria?	√			
Was the signal to noise ratio for each target compound and labeled compound $\geq 10$ ?	√			
Was an initial calibration verification (ICV) standard analyzed after each initial calibration for each instrument?	√			
Were all ICV concentrations for the unlabeled and labeled compounds within QC limits?	√			
<b>IV. Continuing calibration</b>				
Was a continuing calibration performed at the beginning of each 12-hour period?	√			
Were all continuing calibration concentrations for the unlabeled and labeled compounds within QC limits?	√			
Did all continuing calibration standards meet the Ion Abundance Ratio criteria?	√			
<b>V. Blanks</b>				
Was a method blank associated with every sample in this SDG?	√			
Was a method blank performed for each matrix and whenever a sample extraction was performed?	√			
Was there contamination in the method blanks?	√	∅		
<b>VI. Field blanks</b>				
Were field blanks identified in this SDG?		√		
Were target compounds detected in the field blanks?			√	
<b>VII. Matrix spike/Matrix spike duplicates</b>				
Were matrix spike (MS) and matrix spike duplicate (MSD) analyzed in this SDG?		√		
Were the MS/MSD percent recoveries (%R) and the relative percent differences (RPD) within the QC limits?			√	

Validation Area	Yes	No	NA	Findings/Comments
<b>VIII. Laboratory control samples</b>				
Was an LCS analyzed per extraction batch?	√			
Were the LCS percent recoveries (%R) and relative percent difference (RPD) within the QC limits?	√			
<b>IX. Field duplicates</b>				
Were field duplicate pairs identified in this SDG?		√		
Were target compounds detected in the field duplicates?			√	
<b>X. Labeled Compounds</b>				
Were labeled compounds within QC limits?	√	0		
Was the minimum S/N ratio of all labeled compound peaks $\geq 10$ ?	√			
<b>XI. Compound quantitation</b>				
Did the laboratory LOQs/RLs meet the QAPP LOQs/RLs?	√			
Were the correct labeled compound, quantitation ion and relative response factor (RRF) used to quantitate the compound?	√			
Were compound quantitation and RLs adjusted to reflect all sample dilutions and dry weight factors applicable to level IV validation?	√			
<b>XII. Target compound identification</b>				
For 2,3,7,8 substituted congeners with associated labeled standards, were the retention times of the two quantitation peaks within -1 to 3 sec. of the RT of the labeled standard?	√			
For 2,3,7,8 substituted congeners without associated labeled standards, were the relative retention times of the two quantitation peaks within 0.005 time units of the RRT measured in the routine calibration?	√			
For non-2,3,7,8 substituted congeners, were the retention times of the two quantitation peaks within RT established in the performance check solution?	√			
Did selected ion current profile (SICP) contain all characteristic ions listed in Method 1613B, Table 8?	√			
Was the Ion Abundance Ratio for the two quantitation ions within criteria?		√		
Was the signal to noise ratio for each target compound $\geq 2.5$ and $\geq 10$ for the labeled compound?	√			
Does the maximum intensity of each specified characteristic ion coincide within $\pm 2$ seconds (includes labeled standards)?	√			
For PCDF identification, was any signal ( $S/N \geq 2.5$ , at $\pm$ seconds RT) detected in the corresponding PCDF channel?			√	
Was an acceptable lock mass recorded and monitored?	√			
<b>XIII. System performance</b>				
System performance was found to be acceptable.	√			
<b>XIV. Overall assessment of data</b>				
Overall assessment of data was found to be acceptable.	√			

# VALIDATION FINDINGS WORKSHEET

**METHOD:** HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 1613B)

A. 2,3,7,8-TCDD	F. 1,2,3,4,6,7,8-HpCDD	K. 1,2,3,4,7,8-HxCDF	P. 1,2,3,4,7,8,9-HpCDF	U. Total HpCDD
B. 1,2,3,7,8-PeCDD	G. OCDD	L. 1,2,3,6,7,8-HxCDF	Q. OCDF	V. Total TCDF
C. 1,2,3,4,7,8-HxCDD	H. 2,3,7,8-TCDF	M. 2,3,4,6,7,8-HxCDF	R. Total TCDD	W. Total PeCDF
D. 1,2,3,6,7,8-HxCDD	I. 1,2,3,7,8-PeCDF	N. 1,2,3,7,8,9-HxCDF	S. Total PeCDD	X. Total HxCDF
E. 1,2,3,7,8,9-HxCDD	J. 2,3,4,7,8-PeCDF	O. 1,2,3,4,6,7,8-HpCDF	T. Total HxCDD	Y. Total HpCDF

Notes: \_\_\_\_\_  
\_\_\_\_\_  
\_\_\_\_\_



**VALIDATION FINDINGS WORKSHEET**  
**Blanks**

**METHOD:** HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 1613B)

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

N N/A Were all samples associated with a method blank?

N N/A Was a method blank performed for each matrix and whenever a sample extraction was performed?

Y N/A Was the method blank contaminated?

Blank extraction date: 10/19/21

Blank analysis date: 10/25/21

Associated samples: All

Conc. units: ng/kg

Compound	Blank ID	Sample Identification							
<del>Blank</del>	<del>500-BA1</del>								
G	0.981								
X	0.100								

Blank extraction date: \_\_\_\_\_ Blank analysis date: \_\_\_\_\_

Conc. units: \_\_\_\_\_ Associated Samples: \_\_\_\_\_

Compound	Blank ID	Sample Identification							

CIRCLED RESULTS WERE NOT QUALIFIED. ALL RESULTS NOT CIRCLED WERE QUALIFIED BY THE FOLLOWING STATEMENT:  
All contaminants within five times the method blank concentration were qualified as not detected, "U".

**VALIDATION FINDINGS WORKSHEET**  
**Compound Quantitation and Reported RLs**

**METHOD:** HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 1613B)

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

Y N N/A      Were the correct labeled compound, quantitation ions and relative response factors (RRF) used to quantitate the compound?  
Y N N/A      Compound quantitation and RLs were adjusted to reflect all sample dilutions and dry weight factors (if necessary).

#	Date	Sample ID	Finding	Associated Samples	Qualifications
		All	All compounds reported as estimated maximum possible concentration (EMPC) > RL		Jdets/A
		1	G > calibration range		Jdets/P
		All	All compounds reported as estimated maximum possible concentration (EMPC) < RL		U/A

Comments: See sample calculation verification worksheet for recalculations

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## VALIDATION FINDINGS WORKSHEET

### Initial Calibration Calculation Verification

**METHOD:** HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 1613B)

The Relative Response Factor (RRF), average RRF, and percent relative standard deviation (%RSD) were recalculated for the compounds identified below using the following calculations:

$$RRF = (A_x)(C_{is}) / (A_{is})(C_x)$$

average RRF = sum of the RRFs/number of standards

$$\%RSD = 100 * (S/X)$$

A<sub>x</sub> = Area of compound,

C<sub>x</sub> = Concentration of compound,

S = Standard deviation of the RRFs,

A<sub>is</sub> = Area of associated internal standard

C<sub>is</sub> = Concentration of internal standard

X = Mean of the RRFs

#	Standard ID	Calibration Date	Compound (Reference Internal Standard)	Reported	Recalculated	Reported	Recalculated	Reported	Recalculated
				RRF (10/50 std)	RRF (10/50 std)	Average RRF (initial)	Average RRF (initial)	%RSD	%RSD
1	ICAL 01	8/11/21	2,3,7,8-TCDF ( <sup>13</sup> C-2,3,7,8-TCDF)	1.0832006	1.083746	1.107593	1.107593	3.6	3.6
			2,3,7,8-TCDD ( <sup>13</sup> C-2,3,7,8-TCDD)	0.9085186	0.908390	0.9202875	0.9202874	3.1	3.1
			1,2,3,6,7,8-HxCDD ( <sup>13</sup> C-1,2,3,6,7,8-HxCDD)	1.005616	1.005605	1.00898	1.00898	1.0	1.0
			1,2,3,4,6,7,8-HpCDD ( <sup>13</sup> C-1,2,4,6,7,8,-HpCDD)	1.051009	1.051062	1.068088	1.068088	6.6	6.6
			OCDF ( <sup>13</sup> C-OCDD)	1.440564	1.44059	1.44690	1.44690	5.7	5.7
2			2,3,7,8-TCDF ( <sup>13</sup> C-2,3,7,8-TCDF)						
			2,3,7,8-TCDD ( <sup>13</sup> C-2,3,7,8-TCDD)						
			1,2,3,6,7,8-HxCDD ( <sup>13</sup> C-1,2,3,6,7,8-HxCDD)						
			1,2,3,4,6,7,8-HpCDD ( <sup>13</sup> C-1,2,4,6,7,8,-HpCDD)						
			OCDF ( <sup>13</sup> C-OCDD)						
3			2,3,7,8-TCDF ( <sup>13</sup> C-2,3,7,8-TCDF)						
			2,3,7,8-TCDD ( <sup>13</sup> C-2,3,7,8-TCDD)						
			1,2,3,6,7,8-HxCDD ( <sup>13</sup> C-1,2,3,6,7,8-HxCDD)						
			1,2,3,4,6,7,8-HpCDD ( <sup>13</sup> C-1,2,4,6,7,8,-HpCDD)						
			OCDF ( <sup>13</sup> C-OCDD)						

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.

## VALIDATION FINDINGS WORKSHEET Continuing Calibration Results Verification

**METHOD:** HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 1613B)

The percent difference (%D) of the initial calibration average Relative Response Factors (RRFs) and the continuing calibration RRFs were recalculated for the compounds identified below using the following calculation:

% Difference =  $100 * (\text{ave. RRF} - \text{RRF}) / \text{ave. RRF}$   
 $\text{RRF} = (A_x)(C_{is}) / (A_{is})(C_x)$

Where: ave. RRF = initial calibration average RRF  
 RRF = continuing calibration RRF  
 $A_x$  = Area of compound,  $A_{is}$  = Area of associated internal standard  
 $C_x$  = Concentration of compound,  $C_{is}$  = Concentration of internal standard

#	Standard ID	Calibration Date	Compound (Reference Internal Standard)	Average RRF (initial)	Reported	Recalculated	Reported	Recalculated
					Conc (CC)	Conc (CC)	%D	%D
1	<u>2110555A</u>	<u>10/65/21</u>	2,3,7,8-TCDF ( <sup>13</sup> C-2,3,7,8-TCDF)	<u>1.10593</u>	<u>1.074550</u>	<u>1.074615</u>	<u>3.0</u>	<u>3.0</u>
			2,3,7,8-TCDD ( <sup>13</sup> C-2,3,7,8-TCDD)	<u>0.920285</u>	<u>1.0081390</u>	<u>1.0081532</u>	<u>9.5</u>	<u>9.5</u>
			1,2,3,6,7,8-HxCDD ( <sup>13</sup> C-1,2,3,6,7,8-HxCDD)	<u>1.00898</u>	<u>1.0688370</u>	<u>1.0683744</u>	<u>5.9</u>	<u>5.9</u>
			1,2,3,4,6,7,8-HpCDD ( <sup>13</sup> C-1,2,4,6,7,8,-HpCDD)	<u>1.068088</u>	<u>1.1679010</u>	<u>1.1678182</u>	<u>9.3</u>	<u>9.3</u>
			OCDF ( <sup>13</sup> C-OCDF)	<u>1.44690</u>	<u>1.3382880</u>	<u>1.338548</u>	<u>7.5</u>	<u>7.5</u>
2			2,3,7,8-TCDF ( <sup>13</sup> C-2,3,7,8-TCDF)					
			2,3,7,8-TCDD ( <sup>13</sup> C-2,3,7,8-TCDD)					
			1,2,3,6,7,8-HxCDD ( <sup>13</sup> C-1,2,3,6,7,8-HxCDD)					
			1,2,3,4,6,7,8-HpCDD ( <sup>13</sup> C-1,2,4,6,7,8,-HpCDD)					
			OCDF ( <sup>13</sup> C-OCDF)					
3			2,3,7,8-TCDF ( <sup>13</sup> C-2,3,7,8-TCDF)					
			2,3,7,8-TCDD ( <sup>13</sup> C-2,3,7,8-TCDD)					
			1,2,3,6,7,8-HxCDD ( <sup>13</sup> C-1,2,3,6,7,8-HxCDD)					
			1,2,3,4,6,7,8-HpCDD ( <sup>13</sup> C-1,2,4,6,7,8,-HpCDD)					
			OCDF ( <sup>13</sup> C-OCDF)					

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

LDC #: 52703P21

## VALIDATION FINDINGS WORKSHEET

### Laboratory Control Sample Results Verification

Page: 1 of 1  
Reviewer: 9**METHOD:** HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 1613B)

The percent recoveries (%R) and Relative Percent Difference (RPD) of the laboratory control sample and laboratory control sample duplicate (if applicable) were recalculated for the compounds identified below using the following calculation:

% Recovery =  $100 * SSC/SA$       Where: SSC = Spiked sample concentration  
SA = Spike addedRPD =  $|LCS - LCSD| * 2 / (LCS + LCSD)$       LCS = Laboratory control sample percent recovery      LCSD = Laboratory control sample duplicate percent recoveryLCS ID: B110500-PS

Compound	Spike Added ( <u>NS/IS</u> )		Spiked Sample Concentration ( <u>NS/IS</u> )		LCS		LCSD		LCS/LCSD	
	LCS	LCSD	LCS	LCSD	Percent Recovery		Percent Recovery		RPD	
					Reported	Recalc	Reported	Recalc	Reported	Recalculated
2,3,7,8-TCDD	20.0	NA	21.0	NA	105	105				
1,2,3,7,8-PeCDD	100		107		107	107				
1,2,3,4,7,8-HxCDD	↓	↓	99.2	↓	99.2	99.2				
1,2,3,4,7,8,9-HpCDF	↓	↓	95.9	↓	95.9	95.9				
OCDF	200	↓	151	↓	75.5	75.5				

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



## Laboratory Data Consultants, Inc. Data Validation Report

**Project/Site Name:** Duwamish AOC4

**LDC Report Date:** December 15, 2021

**Parameters:** Polychlorinated Biphenyls

**Validation Level:** Stage 4

**Laboratory:** Analytical Resources, Inc., Tukwila

**Sample Delivery Group (SDG):** 21J0137

Sample Identification	Laboratory Sample Identification	Matrix	Collection Date
LDW21-IT653D	21J0137-01	Sediment	07/12/21
LDW21-IT652A	21J0137-02	Sediment	07/12/21
LDW21-IT632D	21J0137-03	Sediment	07/12/21
LDW21-IT644B	21J0137-04	Sediment	07/12/21
LDW21-IT644C	21J0137-05	Sediment	07/12/21
LDW21-IT644D	21J0137-06	Sediment	07/12/21
LDW21-IT644E	21J0137-07	Sediment	07/12/21
LDW21-SC529B	21J0137-08	Sediment	07/14/21
LDW21-SC529C	21J0137-09	Sediment	07/14/21
LDW21-SC529D	21J0137-10	Sediment	07/14/21
LDW21-SC529E	21J0137-11	Sediment	07/14/21
LDW21-SC529F	21J0137-12	Sediment	07/14/21
LDW21-IT608B	21J0137-13	Sediment	07/13/21
LDW21-IT662A	21J0137-14	Sediment	07/13/21
LDW21-IT658A	21J0137-15	Sediment	07/13/21
LDW21-IT648D	21J0137-16	Sediment	07/13/21
LDW21-IT648E	21J0137-17	Sediment	07/13/21
LDW21-SC596A	21J0137-18	Sediment	07/13/21
LDW21-SC596B	21J0137-19	Sediment	07/13/21
LDW21-SC596C	21J0137-20	Sediment	07/13/21
LDW21-SC596D	21J0137-21	Sediment	07/13/21
LDW21-SC596E	21J0137-22	Sediment	07/13/21
LDW21-SC596F	21J0137-23	Sediment	07/13/21
LDW21-SC562C	21J0137-24	Sediment	07/13/21
LDW21-IT653DMS	21J0137-01MS	Sediment	07/12/21
LDW21-IT653DMSD	21J0137-01MSD	Sediment	07/12/21

<b>Sample Identification</b>	<b>Laboratory Sample Identification</b>	<b>Matrix</b>	<b>Collection Date</b>
LDW21-SC562CMS	21J0137-24MS	Sediment	07/13/21
LDW21-SC562CMSD	21J0137-24MSD	Sediment	07/13/21



## 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 Final Lower Duwamish Waterway Quality Assurance Project Plan for Remedial Design of Upper Reach: Pre-Design Investigation (May 2020) and a modified outline of the USEPA National Functional Guidelines (NFG) for Organic Superfund Methods Data Review (January 2017). 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:

Polychlorinated Biphenyls (PCBs) by Environmental Protection Agency (EPA) SW 846 Method 8082A

All sample results were subjected to Stage 4 data validation, which is comprised of the quality control (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): 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.
- U (Non-detected): The analyte was analyzed for and positively identified by the laboratory; however the analyte should be considered non-detected at the reported concentration due to the presence of contaminants detected in the associated blank(s).
- UJ (Non-detected estimated): The analyte was reported as not detected by the laboratory; however the reported quantitation/detection limit is estimated due to non-conformances discovered during data validation.
- R (Rejected): The sample results were rejected due to gross non-conformances discovered during data validation. Data qualified as rejected is not usable.
- 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.

## 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.

Retention time windows were established as required by the method.

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 required frequencies.

The percent differences (%D) were less than or equal to 20.0% for all analytes with the following exceptions:

Date	Standard	Column	Analyte	%D	Associated Samples	Flag	A or P
10/27/21	10272115	2C	Aroclor-1260	24.6	LDW21-IT653D LDW21-SC596E	J (all detects)	A

Retention times of all analytes in the calibration standards were within the established retention time windows.

## 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/Internal Standards

Surrogates were added to all samples as required by the method. Surrogate recoveries (%R) were not within QC limits for sample LDW21-IT652A. No data were qualified for samples analyzed at greater than or equal to 5X dilution.

All internal standard percent recoveries (%R) were within QC limits with the following exceptions:

Sample	Internal Standards	%R (Limits)	Affected Analyte	Flag	A or P
LDW21-IT644B	Hexabromobiphenyl	45 (50-200)	Aroclor-1260	J (all detects)	A
LDW21-IT644C	Hexabromobiphenyl	48 (50-200)	Aroclor-1260	J (all detects)	A
LDW21-SC596A	Hexabromobiphenyl	48 (50-200)	Aroclor-1260	J (all detects)	A

### VII. Matrix Spike/Matrix Spike Duplicates

Matrix spike (MS) and matrix spike duplicate (MSD) samples were reviewed for each matrix as applicable. Percent recoveries (%R) were not within the QC limits for LDW21-IT653DMS/MSD. No data were qualified for MS/MSD samples analyzed greater than or equal to a 5X dilution. Relative percent differences (RPD) were within the QC limits.

### VIII. Laboratory Control Samples/Standard Reference Materials

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.

Standard reference materials (SRM) were analyzed as required by the method. The results were within QC limits.

### IX. Field Duplicates

No field duplicates were identified in this SDG.

### X. Target Analyte Quantitation

All target analyte quantitations met validation criteria.

The sample results for detected analytes from the two columns were within 40% relative percent difference (RPD) with the following exceptions:

Sample	Analyte	RPD	Flag	A or P
LDW21-IT644C	Aroclor-1260	41.9	J (all detects)	A
LDW21-IT644D	Aroclor-1260 Aroclor-1248	44.8 41.4	J (all detects) J (all detects)	A
LDW21-SC596A	Aroclor-1260	40.5	J (all detects)	A

Sample	Analyte	RPD	Flag	A or P
LDW21-SC596B	Aroclor-1254	49.4	J (all detects)	A
LDW21-SC596D	Aroclor-1260	41.3	J (all detects)	A
LDW21-SC562C	Aroclor-1248	44.6	J (all detects)	A
LDW21-SC529F	Aroclor-1248	58.3	J (all detects)	A

### **XI. Target Analyte Identification**

All target analyte identifications met validation criteria.

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

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

Due to continuing calibration %D, internal standard %R, and RPD between two columns, data were qualified as estimated in ten samples.

The quality control criteria reviewed, other than those discussed above, were met and are considered acceptable.

**Duwamish AOC4**

**Polychlorinated Biphenyls - Data Qualification Summary - SDG 21J0137**

Sample	Analyte	Flag	A or P	Reason
LDW21-IT653D LDW21-SC596E	Aroclor-1260	J (all detects)	A	Continuing calibration (%D)
LDW21-IT644B LDW21-IT644C LDW21-SC596A	Aroclor-1260	J (all detects)	A	Internal standards (%R)
LDW21-IT644C LDW21-SC596A LDW21-SC596D	Aroclor-1260	J (all detects)	A	Target analyte quantitation (RPD between two columns)
LDW21-IT644D	Aroclor-1260 Aroclor-1248	J (all detects) J (all detects)	A	Target analyte quantitation (RPD between two columns)
LDW21-SC596B	Aroclor-1254	J (all detects)	A	Target analyte quantitation (RPD between two columns)
LDW21-SC562C LDW21-SC529F	Aroclor-1248	J (all detects)	A	Target analyte quantitation (RPD between two columns)

**Duwamish AOC4**

**Polychlorinated Biphenyls - Laboratory Blank Data Qualification Summary - SDG 21J0137**

No Sample Data Qualified in this SDG

**Duwamish AOC4**

**Polychlorinated Biphenyls - Field Blank Data Qualification Summary - SDG 21J0137**

No Sample Data Qualified in this SDG

LDC #: 52703C3b

**VALIDATION COMPLETENESS WORKSHEET**

Date: 12/10/21

SDG #: 21J0137

Stage 4

Page: 1 of 2

Laboratory: Analytical Resources, Inc.

Reviewer: [Signature]

2nd Reviewer: IVG

**METHOD:** GC Polychlorinated Biphenyls (EPA SW846 Method 8082A)

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	A	
II.	Initial calibration/ICV	A/A	PSD ≤ 20/0      10V ≤ 20/0
III.	Continuing calibration	W	CCV ≤ 20/0
IV.	Laboratory Blanks	A	
V.	Field blanks	N	
VI.	Surrogate spikes / IS	W/W	
VII.	Matrix spike/Matrix spike duplicates	W	
VIII.	Laboratory control samples / SM	A	LES/D
IX.	Field duplicates	N	
X.	Target analyte quantitation	W	
XI.	Target analyte identification	A	
XII.	Overall assessment of data	D	

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:

	Client ID	Lab ID	Matrix	Date
1	LDW21-IT653D	21J0137-01	Sediment	07/12/21
2	LDW21-IT652A	21J0137-02	Sediment	07/12/21
3	LDW21-IT632D	21J0137-03	Sediment	07/12/21
4	LDW21-IT644B	21J0137-04	Sediment	07/12/21
5	LDW21-IT644C	21J0137-05	Sediment	07/12/21
6	LDW21-IT644D	21J0137-06	Sediment	07/12/21
7	LDW21-IT644E	21J0137-07	Sediment	07/12/21
8	LDW21-SC529B	21J0137-08	Sediment	07/14/21
9	LDW21-SC529C	21J0137-09	Sediment	07/14/21
10	LDW21-SC529D	21J0137-10	Sediment	07/14/21
11	LDW21-SC529E	21J0137-11	Sediment	07/14/21
12	LDW21-SC529F	21J0137-12	Sediment	07/14/21
13	LDW21-IT608B	21J0137-13	Sediment	07/13/21
14	LDW21-IT662A	21J0137-14	Sediment	07/13/21
15	LDW21-IT658A	21J0137-15	Sediment	07/13/21
16	LDW21-IT648D	21J0137-16	Sediment	07/13/21
17	LDW21-IT648E	21J0137-17	Sediment	07/13/21

LDC #: 52703C3b  
 SDG #: 21J0137  
 Laboratory: Analytical Resources, Inc.

**VALIDATION COMPLETENESS WORKSHEET**

Stage 4

Date: 12/10/21  
 Page: 2 of 2  
 Reviewer: [Signature]  
 2nd Reviewer: [Signature]

**METHOD:** GC Polychlorinated Biphenyls (EPA SW846 Method 8082A)

	Client ID	Lab ID	Matrix	Date
18	LDW21-SC596A	21J0137-18	Sediment	07/13/21
19	LDW21-SC596B	21J0137-19	Sediment	07/13/21
20	LDW21-SC596C	21J0137-20	Sediment	07/13/21
21	LDW21-SC596D	21J0137-21	Sediment	07/13/21
22	LDW21-SC596E	21J0137-22	Sediment	07/13/21
23	LDW21-SC596F	21J0137-23	Sediment	07/13/21
24	LDW21-SC562C	21J0137-24	Sediment	07/13/21
25	LDW21-IT653DMS	21J0137-01MS	Sediment	07/12/21
26	LDW21-IT653DMSD	21J0137-01MSD	Sediment	07/12/21
27	LDW21-SC562CMS	21J0137-24MS	Sediment	07/13/21
28	LDW21-SC562CMSD	21J0137-24MSD	Sediment	07/13/21
29				
30				
31				

Notes:


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) $\leq$ 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) $\leq$ 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) $\leq$ 20%?	<input type="checkbox"/>	<input checked="" 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 compounds 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 type="checkbox"/>	<input checked="" 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 checked="" type="checkbox"/>	<input 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 type="checkbox"/>	<input checked="" 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>IX. Field duplicates</b>				
Were field duplicate pairs identified in this SDG?		/		
Were target compounds detected in the field duplicates?			/	
<b>X. Compound quantitation</b>				
Did the laboratory LOQs/RLs meet the QAPP LOQs/RLs?	/			
Were compound quantitation and RLs adjusted to reflect all sample dilutions and dry weight factors applicable to level IV validation?	/			
<b>XI. Target compound identification</b>				
Were the retention times of reported detects within the RT windows?	/			
<b>XIII. Overall assessment of data</b>				
Overall assessment of data was found to be acceptable.	/			

LDC #: 52703036

## VALIDATION FINDINGS WORKSHEET Continuing Calibration

Page: 1 of 1  
Reviewer: 9

METHOD:  GC  HPLC

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

N/A Were continuing calibration standards analyzed at the required frequencies?

N/A Did the continuing calibration standards meet the %D validation criteria of  $\leq 20.0\%$ ?

**Level IV Only**

N/A Were the retention times for all calibrated compounds within their respective acceptance windows?

#	Date	Standard ID	Detector/Column	Compound	%D (Limit)	RT (limit)	Associated Samples	Qualifications
	<u>10/5/11</u>	<u>102T-215</u>	<u>2C</u>	<u>B3</u>	<u>2.6</u>	( )	<u>1.22 (dots)</u>	<u>N/A</u> (1cm)
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
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						( )		
						( )		
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						( )		
						( )		
						( )		

## VALIDATION FINDINGS WORKSHEET Surrogate Recovery

METHOD:  GC  HPLC

Are surrogates required by the method? Yes  or No .

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

- N N/A Were surrogates spiked into all samples and blanks?  
 N N/A Did all surrogate recoveries (%R) meet the QC limits?

#	Sample ID	Detector/ Column	Surrogate Compound	%R (Limits)	Qualifications
	2 <del>126</del>	1e	0	133 (40-126)	<del>Not Spiked</del> No level (25%)
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	
				( )	

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	1,2-Dinitrobenzene
C a,a,a-Trifluorotoluene	I	Fluorobenzene (FBZ)	O	Decachlorobiphenyl (DCB)	U	Triphenyltin		
D Bromochlorobenzene	J	n-Triacontane	P	1-methylnaphthalene	V	Tri-n-propyltin		
E 1,4-Dichlorobutane	K	Hexacosane	Q	Dichlorophenyl Acetic Acid (DCAA)	W	Tributyl Phosphate		
F 1,4-Difluorobenzene (DFB)	L	Bromobenzene	R	4-Nitrophenol	X	Triphenyl Phosphate		

### VALIDATION FINDINGS WORKSHEET Internal Standards

**METHOD:** GC

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

- Y  N  N/A    Were all internal standard area counts within -50 to +100% of the ICAL midpoint standard?  
 Y  N  N/A    Were the retention times of the internal standards within +/- 0.05 min seconds of the retention times of the ICAL midpoint standard?

#	Date	Sample ID	Internal Standard	%R Area (Limits)	RT (Limits)	Qualifications
		4 (dots)	HBB(1c)	45 (50-200)		↓/N/A (BB) ↓ ↓
		5 (dots)	↓	48		
		18 (dots)	↓	48		

### VALIDATION FINDINGS WORKSHEET Matrix Spike/Matrix Spike Duplicates

METHOD:  GC  HPLC

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

Y  N  N/A Were a matrix spike (MS) and matrix spike duplicate (MSD) analyzed for each matrix in this SDG?

Y  N  N/A Was an MS/MSD analyzed every 20 samples for each matrix or whenever a sample extraction was performed?

Y  N  N/A Were the MS/MSD percent recoveries (%R) and relative percent differences (RPD) within QC limits?

#	MS/MSD ID	Compound	MS %R (Limits)	MSD %R (Limits)	RPD (Limits)	Associated Samples	Qualifications
	25/26	70R out	( )	( )	( )	1 (5x)	<del>NR</del> Not Anal
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		

LDC #: 5203036

**VALIDATION FINDINGS WORKSHEET**  
**Compound Quantitation and Reported CRQLs**

Page: 1 of 1  
 Reviewer: [Signature]

METHOD:  GC  HPLC

**Level IV/D Only**

- Y  N  N/A Were CRQLs adjusted for sample dilutions, dry weight factors, etc.?
- Y  N  N/A Did the reported results for detected target compounds agree within 10.0% of the recalculated results?
- Y  N  N/A Did the relative percent differences of detected compounds between two columns/detectors  $\leq 40\%$ ?  
 If no, please see findings below.

#	Compound Name	Sample ID	%RPD Between Two Columns/Detectors Limit ( $\leq 40\%$ )	Qualifications
	<del>Acroloy 1260</del>	5	41.9	<del>↓ det 3/A</del>
	↓	6	44.8	
	<del>Acroloy 1248</del>		41.4	
	<del>Acroloy 1260</del>	18	40.5	
	↓			
	<del>1251</del>	19	49.4	
	↓			
	<del>1260</del>	21	41.3	
	↓			
	<del>1248</del>	24	41.6	↓
	↓			
	<del>1248</del>	12	58.3	<del>↓ det 3/A</del>

**VALIDATION FINDINGS WORKSHEET**  
**Initial Calibration Calculation Verification**

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 (100 std)	CF (100 std)	Ave CF (initial)	Ave CF (initial)	%RSD	%RSD
1	KAC	8/2/1	<del>BB1</del> (1c)	0.0358713	0.0358713	0.0359923	0.059923	2.6	2.6
			<del>BB1</del> (2c)	0.0687269	0.0687264	0.0665078	0.0665072	7.7	7.8
2									
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 #: 5-70303b

## VALIDATION FINDINGS WORKSHEET

### Continuing Calibration Results Verification

Page: 1 of 1  
Reviewer: QMETHOD:  GC\_HPLC

The percent difference (%D) of the initial calibration average Calibration Factors (CF) and the continuing calibration CF were recalculated for the compounds 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 compound  
C = Concentration of compound

#	Standard ID	Calibration Date	Compound	Average CF(Ical)/ CCV Conc.	Reported	Recalculated	Reported	Recalculated
					CF/ Conc. CCV	CF/ Conc. CCV	%D	%D
1	10242104	10/24/21	BB-1 (1c)	0.0359923	0.0301845	0.0301845	16.0	16.1
	2-9.43		BB-1 (2c)	0.0665032	0.053140T	0.053140T	20.0	20.1
2	10242121	10/25/21	↓	0.0359923	0.0304T58	0.0304T5T	15.2	15.3
	10-20			3-14	0.0665032	0.0539459	0.0539459	18.8
3	10252103	10/25/21	↓	0.0359923	0.0299229	0.0299229	16.8	16.9
	21, 23-24			20=10	0.0665032	0.054280T	0.0542806	18.4
4	10272115	10/27/21	↓	0.0359923	0.028410T	0.0284106	21.2	21.1
	1.22			18=25	0.0665032	0.0482295	0.0482294	27.6



**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	
DCB	1C	<del>10.0</del> 8.0	9.1	113	113	
TCMX	↓	↓	6.2	78.0	78	
DCB	2C	↓	6.5	80.7	81	
TCMX	↓	↓	6.1	76.3	76.3	

Sample ID: \_\_\_\_\_

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

Sample ID: \_\_\_\_\_

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

## VALIDATION FINDINGS WORKSHEET Matrix Spike/Matrix Spike Duplicates Results Verification

**METHOD:** GC HPLC

The percent recoveries (%R) and relative percent differences (RPD) of the matrix spike and matrix spike duplicate were recalculated for the compounds identified below using the following calculation:

$\% \text{Recovery} = 100 * (\text{SSC} - \text{SC}) / \text{SA}$

Where

SSC = Spiked sample concentration

SC = Sample concentration

SA = Spike added

$\text{RPD} = ((\text{SSCMS} - \text{SSCMSD}) * 2) / (\text{SSCMS} + \text{SSCMSD}) * 100$

MS = Matrix spike

MSD = Matrix spike duplicate

MS/MSD samples: 25/26

Compound	Spike Added ( <u>MS</u> )		Sample Conc. ( <u>MS</u> )	Spike Sample Concentration ( <u>MS</u> )		Matrix spike		Matrix Spike Duplicate		MS/MSD		
	MS	MSD		---	MS	MSD	Percent Recovery		Percent Recovery		RPD	
							Reported	Recalc.	Reported	Recalc.	Reported	Recalc.
Gasoline (8015)												
Diesel (8015)												
Benzene (8021B)												
Methane (RSK-175)												
2,4-D (8151)												
Dinoseb (8151)												
Naphthalene (8310)												
Anthracene (8310)												
HMX (8330)												
2,4,6-Trinitrotoluene (8330)												
<u>BB</u>	<u>101</u>	<u>101</u>	<u>255</u>	<u>273</u>	<u>297</u>	<u>17.8</u>	<u>17.8</u>	<u>41.6</u>	<u>41.6</u>	<u>8.16</u>	<u>8.4</u>	

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

LDC #: 520303ab

## VALIDATION FINDINGS WORKSHEET

**Laboratory Control Sample/Laboratory Control Sample Duplicates Results Verification**

Page: 1 of 1  
Reviewer: Q

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 compounds identified below using the following calculation:

$$\% \text{Recovery} = 100 * (\text{SSC} - \text{SC}) / \text{SA}$$

Where SSC = Spiked sample concentration  
SA = Spike added  
LCS = Laboratory Control Sample

SC = Sample concentration

$$\text{RPD} = ((\text{SSCLCS} - \text{SSCLCSD}) * 2) / (\text{SSCLCS} + \text{SSCLCSD}) * 100$$

LCSD = Laboratory Control Sample duplicate

LCS/LCSD samples: BN0548

Compound	Spike Added		Spike Sample Concentration		LCS		LCSD		LCS/LCSD	
	(165)		(165)		Percent Recovery		Percent Recovery		RPD	
	LCS	LCSD	LCS	LCSD	Reported	Recalc.	Reported	Recalc.	Reported	Recalc.
Gasoline (8015)										
Diesel (8015)										
Benzene (8021B)										
Methane (RSK-175)										
2,4-D (8151)										
Dinoseb (8151)										
Naphthalene (8310)										
Anthracene (8310)										
HMX (8330)										
2,4,6-Trinitrotoluene (8330)										
<del>BB</del>	101	101	81.7	85.5	81.0	81.0	84.9	84.7	4.61	4.5

Comments: Refer to Laboratory Control Sample/Laboratory Control Sample Duplicate 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**  
**Sample Calculation Verification**

METHOD:  GC  HPLC

Y N N/A  
Y N N/A

Were all reported results recalculated and verified for all level IV samples?  
 Were all recalculated results for detected target compounds within 10% of the reported results?

Concentration =  $\frac{(A)(F_v)(D_f)}{(RF)(V_s \text{ or } W_s)(\%S/100)}$

Example:

Sample ID. 1 Compound Name PCB-1260-1

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

Concentration =  $\frac{(50353)(80.0)}{(384769)(0.0359923)} = 290.9$

con total =  $\frac{(290.9 + 182.1 + 253.1 + 208.7 + 342.3)(2.5)(5)}{5 \times 20.03 \times 0.6243} = 255.3 \text{ } \mu\text{g/g}$

#	Sample ID	Compound	Reported Concentrations ( <u>16755</u> )	Recalculated Results Concentrations ( )	Qualifications
	<u>1</u>	<u>PCB-1260</u>	<u>255</u>		

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

## Laboratory Data Consultants, Inc. Data Validation Report

**Project/Site Name:** Duwamish AOC4

**LDC Report Date:** December 15, 2021

**Parameters:** Wet Chemistry

**Validation Level:** Stage 4

**Laboratory:** Analytical Resources, Inc., Tukwila, WA

**Sample Delivery Group (SDG):** 21J0137

Sample Identification	Laboratory Sample Identification	Matrix	Collection Date
LDW21-IT653D	21J0137-01	Sediment	07/12/21
LDW21-IT652A	21J0137-02	Sediment	07/12/21
LDW21-IT632D	21J0137-03	Sediment	07/12/21
LDW21-IT644B	21J0137-04	Sediment	07/12/21
LDW21-IT644C	21J0137-05	Sediment	07/12/21
LDW21-IT644D	21J0137-06	Sediment	07/12/21
LDW21-IT644E	21J0137-07	Sediment	07/12/21
LDW21-SC529B	21J0137-08	Sediment	07/14/21
LDW21-SC529C	21J0137-09	Sediment	07/14/21
LDW21-SC529D	21J0137-10	Sediment	07/14/21
LDW21-SC529E	21J0137-11	Sediment	07/14/21
LDW21-SC529F	21J0137-12	Sediment	07/14/21
LDW21-IT608B	21J0137-13	Sediment	07/13/21
LDW21-IT662A	21J0137-14	Sediment	07/13/21
LDW21-IT658A	21J0137-15	Sediment	07/13/21
LDW21-IT648D	21J0137-16	Sediment	07/13/21
LDW21-IT648E	21J0137-17	Sediment	07/13/21
LDW21-SC596A	21J0137-18	Sediment	07/13/21
LDW21-SC596B	21J0137-19	Sediment	07/13/21
LDW21-SC596C	21J0137-20	Sediment	07/13/21
LDW21-SC596D	21J0137-21	Sediment	07/13/21
LDW21-SC596E	21J0137-22	Sediment	07/13/21
LDW21-SC596F	21J0137-23	Sediment	07/13/21
LDW21-SC562C	21J0137-24	Sediment	07/13/21
LDW21-IT653DMS	21J0137-01MS	Sediment	07/12/21
LDW21-IT653DDUP1	21J0137-01DUP1	Sediment	07/12/21

<b>Sample Identification</b>	<b>Laboratory Sample Identification</b>	<b>Matrix</b>	<b>Collection Date</b>
LDW21-IT653DDUP2	21J0137-01DUP2	Sediment	07/12/21

## **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 Final Lower Duwamish Waterway Quality Assurance Project Plan for Remedial Design of Upper Reach: Pre-Design Investigation (May 2020) and a modified outline of the USEPA National Functional Guidelines (NFG) for Inorganic Superfund Methods Data Review (January 2017). 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 methods:

Total Organic Carbon by Environmental Protection Agency (EPA) SW 846 Method 9060A

Total Solids by Standard Method 2540G

All sample results were subjected to Stage 4 data validation, which is comprised of the quality control (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): 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.
- U (Non-detected): The analyte was analyzed for and positively identified by the laboratory; however the analyte should be considered non-detected at the reported concentration due to the presence of contaminants detected in the associated blank(s).
- UJ (Non-detected estimated): The analyte was reported as not detected by the laboratory; however the reported quantitation/detection limit is estimated due to non-conformances discovered during data validation.
- R (Rejected): The sample results were rejected due to gross non-conformances discovered during data validation. Data qualified as rejected is not usable.
- 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.



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

All samples were received in good condition.

All technical holding time requirements were met.

## **II. Initial Calibration**

All criteria for the initial calibration of each method were met.

## **III. Continuing Calibration**

Continuing calibration frequency and analysis criteria were met for each method when applicable.

## **IV. Laboratory Blanks**

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

## **V. Field Blanks**

No field blanks were identified in this SDG.

## **VI. Matrix Spike/Matrix Spike Duplicates**

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

## **VII. Duplicate Sample Analysis**

Duplicate (DUP) sample analysis was performed on an associated project sample. Results were within QC limits.

## **VIII. Laboratory Control Samples**

Laboratory control samples (LCS) were analyzed as required by the methods. Percent recoveries (%R) were within QC limits.

## **IX. Field Duplicates**

No field duplicates were identified in this SDG.

## **X. Target Analyte Quantitation**

All target analyte quantitations were within validation criteria.

## **XI. Overall Assessment of Data**

The analysis was conducted within all specifications of the methods. No results were rejected in this SDG.

The quality control criteria reviewed were met and are considered acceptable.

**Duwamish AOC4  
Wet Chemistry - Data Qualification Summary - SDG 21J0137**

No Sample Data Qualified in this SDG

**Duwamish AOC4  
Wet Chemistry - Laboratory Blank Data Qualification Summary - SDG 21J0137**

No Sample Data Qualified in this SDG

**Duwamish AOC4  
Wet Chemistry - Field Blank Data Qualification Summary - SDG 21J0137**

No Sample Data Qualified in this SDG

LDC #: 52703C6

**VALIDATION COMPLETENESS WORKSHEET**

SDG #: 21J0137

Stage 4

Laboratory: Analytical Resources, Inc., Tukwila, WA

Date: 12/19/21

Page: 1 of 2

Reviewer: [Signature]

2nd Reviewer: [Signature]

**METHOD: (Analyte) TOC (EPA SW846 Method 9060A), Total Solids (SM2540G)**

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	AA	
II.	Initial calibration	A	
III.	Calibration verification	A	
IV.	Laboratory Blanks	A	
V.	Field blanks	N	
VI.	Matrix Spike/Matrix Spike Duplicates	A	
VII.	Duplicate sample analysis	A	
VIII.	Laboratory control samples	A	LCS
IX.	Field duplicates	N	
X.	Target Analyte Quantitation	A	
XI.	Overall assessment of data	A	

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:

	Client ID	Lab ID	Matrix	Date
1	LDW21-IT653D	21J0137-01	Sediment	07/12/21
2	LDW21-IT652A	21J0137-02	Sediment	07/12/21
3	LDW21-IT632D	21J0137-03	Sediment	07/12/21
4	LDW21-IT644B	21J0137-04	Sediment	07/12/21
5	LDW21-IT644C	21J0137-05	Sediment	07/12/21
6	LDW21-IT644D	21J0137-06	Sediment	07/12/21
7	LDW21-IT644E	21J0137-07	Sediment	07/12/21
8	LDW21-SC529B	21J0137-08	Sediment	07/14/21
9	LDW21-SC529C	21J0137-09	Sediment	07/14/21
10	LDW21-SC529D	21J0137-10	Sediment	07/14/21
11	LDW21-SC529E	21J0137-11	Sediment	07/14/21
12	LDW21-SC529F	21J0137-12	Sediment	07/14/21
13	LDW21-IT608B	21J0137-13	Sediment	07/13/21
14	LDW21-IT662A	21J0137-14	Sediment	07/13/21
15	LDW21-IT658A	21J0137-15	Sediment	07/13/21
16	LDW21-IT648D	21J0137-16	Sediment	07/13/21
17	LDW21-IT648E	21J0137-17	Sediment	07/13/21

LDC #: 52703C6  
SDG #: 21J0137  
Laboratory: Analytical Resources, Inc., Tukwila, WA

### VALIDATION COMPLETENESS WORKSHEET

Stage 4

Date: 10/9/21  
Page: 2 of 2  
Reviewer: [Signature]  
2nd Reviewer: [Signature]

**METHOD: (Analyte) TOC (EPA SW846 Method 9060A), Total Solids (SM2540G)**

	Client ID	Lab ID	Matrix	Date
18	LDW21-SC596A	21J0137-18	Sediment	07/13/21
19	LDW21-SC596B	21J0137-19	Sediment	07/13/21
20	LDW21-SC596C	21J0137-20	Sediment	07/13/21
21	LDW21-SC596D	21J0137-21	Sediment	07/13/21
22	LDW21-SC596E	21J0137-22	Sediment	07/13/21
23	LDW21-SC596F	21J0137-23	Sediment	07/13/21
24	LDW21-SC562C	21J0137-24	Sediment	07/13/21
25	LDW21-IT653DMS	21J0137-01MS	Sediment	07/12/21
26	LDW21-IT653DDUP \	21J0137-01DUP \	Sediment	07/12/21
27	LDW21-IT653DTRP DupZ	21J0137-01TRP DupZ	Sediment	07/12/21
28				
29				
30				

Notes:  
\_\_\_\_\_  
\_\_\_\_\_

METHOD: Inorganics				
Validation Area	Yes	No	NA	Comments
<b>I. Technical holding times</b>				
Were all technical holding times were met?	X			Frozen
<b>II. Calibration</b>				
Were all instruments calibrated at the required frequency?	X			
Were the proper number of standards used?	X			
Were all initial and continuing calibration verifications within the QC limits?	X			
Were all initial calibration correlation coefficients within limits as specified by the method?	X			
Were balance checks performed as required?	X			
<b>III. Blanks</b>				
Was a method blank associated with every sample in this SDG?	X			
Was there contamination in the method blanks?		X		
Was there contamination in the initial and continuing calibration blanks?		X		
<b>IV. Matrix Spike/Matrix Spike Duplicates/Laboratory Duplicates</b>				
Were MS/MSD recoveries with the QC limits? (If the sample concentration exceeded the spike concentration by a factor of 4, no action was taken.)	X			
Were the MS/MSD or laboratory duplicate relative percent differences (RPDs) within the QC limits?	X			
<b>V. Laboratory Control Samples</b>				
Was a LCS analyzed for each batch in the SDG?	X			
Were the LCS recoveries and RPDs (if applicable) within QC limits?	X			
<b>X. Sample Result Verification</b>				
Were all reporting limits adjusted to reflect sample dilutions?	X			
Were all soil samples dry weight corrected?	X			
<b>XI. Overall Assessment of Data</b>				
Was the overall assessment of the data found to be acceptable?	X			

METHOD: Inorganics				
Validation Area	Yes	No	NA	Comments
<b>XII. Field Duplicates</b>				
Were field duplicates identified in this SDG?		X		
Were target analytes detected in the field duplicates?			X	
<b>XIII. Field Blanks</b>				
Were field blanks identified in this SDG?		X		
Were target analytes detected in the field blanks?			X	

All elements are applicable to each sample as noted below.

Sample ID	Target Analyte List
All	TS, TOC
QC:	
25	TOC
26	TS, TOC
27	TS



**Validation Findings Worksheet**  
**Initial and Continuing Calibration Calculation Verification**

**Method:** Inorganics, Method See Cover

An initial or continuing calibration verification percent recovery (%R) was recalculated for each type of analysis using the following formula:

$$\%R = \frac{\text{Found} \times 100}{\text{True}}$$

Where,

Found = concentration of each analyte measured in the analysis of the ICV or CCV solution

True = concentration of each analyte in the ICV or CCV source

Calibration verification	TOC	ICV	44.446	47.154	106	106	Y
Calibration verification	TOC	CCV	44.446	44.629	100	100	Y
Calibration verification	TOC	CCV	44.446	44.357	100	100	Y

Comments:

VALIDATION FINDINGS CHECKLIST  
Quality Control Sample Recalculations

METHOD: Inorganics

Percent recoveries (%R) for the laboratory control sample (LCS) and matrix spike (MS) were recalculated using the following formula.

$$\%R = (\text{Found}/\text{True}) \times 100$$

Found = concentration of each analyte measured in the analysis. For the MS calculation, Found = SSR (Spiked Sample Result) - SR (Sample Result)

True = concentration of each analyte in the source

The sample and duplicate relative percent difference (RPD) was recalculated using the following formula.

$$\text{RPD} = (\text{Absolute value}(S-D) \times 200) / (S+D)$$

S = Original sample concentration

D = Duplicate sample concentration

Sample ID	Type of Analysis	Element	Found/S	True/D	Recalculated %R/RPD	Reported %R/RPD	Acceptable (Y/N)
LCS	LCS	TOC	45.3	44.4	102	102	Y
24	MS	TOC	1.31	1.16	113	112	Y
26	Duplicate	TS	64.1	63.39	1.11	1.11	Y

VALIDATION FINDINGS CHECKLIST  
Sample Calculation Verification

METHOD: Inorganics

Analytes were recalculated and verified using the following equation.

Concentration = (Result from raw data x Final volume x Dilution factor) / (Percent solids (if applicable) x Initial weight or volume)

Sample ID	Analyte	Raw Data (%)	Dry (g)	Sample Dry (g)	Tare (g)	Percent solids (%)	Reported Result (%)	Recalculated Result (mg/Kg)	Acceptable (Y/N)
1	TOC	1.75				64.1	2.73	2.73	Y
2	TOC	0.919				60.23	1.53	1.53	Y
3	TOC	0.027				81.62	0.03	0.03	Y
4	TOC	1.48				65.21	2.27	2.27	Y
5	TOC	1.556				62.73	2.48	2.48	Y
6	TOC	1.055				66.96	1.58	1.58	Y
7	TOC	0.038				91.18	0.04	0.04	Y
8	TOC	1.117				58.44	1.91	1.91	Y
9	TOC	1.01				60.68	1.66	1.66	Y
10	TOC	0.871				64.17	1.36	1.36	Y
11	TOC	0.617				67.65	0.91	0.91	Y
12	TOC	0.961				63.19	1.52	1.52	Y
13	TOC	0.523				69.94	0.75	0.75	Y
14	TOC	1.067				55.34	1.93	1.93	Y
15	TS		3.936	5.7969	0.8112		62.68	62.68	Y
16	TS		3.4649	4.8347	0.8033		66.02	66.02	Y
17	TS		3.8553	4.1942	0.8078		89.4	89.99	Y
18	TS		3.2583	4.8916	0.7968		60.11	60.11	Y
19	TS		3.4221	4.4786	0.7955		71.31	71.31	Y
20	TS		5.4429	7.0638	0.7986		74.13	74.13	Y
21	TS		4.0123	5.7007	0.7972		65.57	65.57	Y
22	TS		2.8332	4.2395	0.8143		58.94	58.94	Y
23	TS		3.7536	5.2064	0.8098		66.96	66.96	Y
24	TS		4.4018	6.7354	0.7919		60.74	60.74	Y

## Laboratory Data Consultants, Inc. Data Validation Report

**Project/Site Name:** Duwamish AOC4  
**LDC Report Date:** December 15, 2021  
**Parameters:** Polychlorinated Dioxins/Dibenzofurans  
**Validation Level:** Stage 4  
**Laboratory:** Analytical Resources, Inc., Tukwila, WA  
**Sample Delivery Group (SDG):** 21J0137

<b>Sample Identification</b>	<b>Laboratory Sample Identification</b>	<b>Matrix</b>	<b>Collection Date</b>
LDW21-IT653D	21J0137-01	Sediment	07/12/21
LDW21-IT652A	21J0137-02	Sediment	07/12/21
LDW21-IT662A	21J0137-14	Sediment	07/13/21
LDW21-IT658A	21J0137-15	Sediment	07/13/21
LDW21-IT648D	21J0137-16	Sediment	07/13/21
LDW21-IT648E	21J0137-17	Sediment	07/13/21

## 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 Final Lower Duwamish Waterway Quality Assurance Project Plan for Remedial Design of Upper Reach: Pre-Design Investigation (May 2020) and a modified outline of the USEPA National Functional Guidelines (NFG) for High Resolution Superfund Methods Data Review (April 2016). 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:

Polychlorinated Dioxins/Dibenzofurans by Environmental Protection Agency (EPA) Method 1613B

All sample results were subjected to Stage 4 data validation, which is comprised of the quality control (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): 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.
- U (Non-detected): The analyte was analyzed for and positively identified by the laboratory; however the analyte should be considered not detected at the reported concentration due to the presence of contaminants detected in the associated blank(s).
- UJ (Non-detected estimated): The analyte was reported as not detected by the laboratory; however the reported quantitation/detection limit is estimated due to non-conformances discovered during data validation.
- R (Rejected): The sample results were rejected due to gross non-conformances discovered during data validation. Data qualified as rejected is not usable.
- 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.

## **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. HRGC/HRMS Instrument Performance Check**

Instrument performance was checked at the required frequency.

Retention time windows were established for all homologues. The chromatographic resolution between 2,3,7,8-TCDD and peaks representing any other unlabeled TCDD isomer was less than or equal to 25%.

The static resolving power was at least 10,000 (10% valley definition).

## **III. Initial Calibration and Initial Calibration Verification**

A five point 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 and less than or equal to 35.0% for labeled compounds.

The ion abundance ratios for all PCDDs and PCDFs were within validation criteria.

The minimum S/N ratio was greater than or equal to 10 for each analyte and labeled compound.

The percent differences (%D) of the initial calibration verification (ICV) standard were within the QC limits for all analytes and labeled compounds.

## **IV. Continuing Calibration**

Continuing calibration was performed at the required frequencies.

All of the continuing calibration results were within the QC limits for all analytes and labeled compounds.

The ion abundance ratios for all PCDDs and PCDFs were within validation criteria.

The minimum S/N ratio was greater than or equal to 10 for each analyte and labeled compound.

## **V. Laboratory Blanks**

Laboratory blanks were analyzed as required by the method. No contaminants were found in the laboratory blanks with the following exceptions:

Blank ID	Extraction Date	Analyte	Concentration	Associated Samples
BJJ0500-BLK1	10/19/21	OCDD Total HxCDF	0.981 ng/Kg 0.100 ng/Kg	All samples in SDG 21J0137

Sample concentrations were compared to concentrations detected in the laboratory blanks. The sample concentrations were either not detected or were significantly greater than the concentrations found in the associated laboratory blanks.

## VI. Field Blanks

No field blanks were identified in this SDG.

## VII. Matrix Spike/Matrix Spike Duplicates

The laboratory has indicated that there were no matrix spike (MS) and matrix spike duplicate (MSD) analyses specified for the samples in this SDG, and therefore matrix spike and matrix spike duplicate analyses were not performed for this SDG.

## VIII. Laboratory Control Samples/Standard Reference Materials

Laboratory control samples (LCS) were analyzed as required by the method. Percent recoveries (%R) were within QC limits.

Standard reference materials (SRM) were analyzed as required by the method. The results were within QC limits.

## IX. Field Duplicates

No field duplicates were identified in this SDG.

## X. Internal Standards

All internal standard areas and retention times were within QC limits.

## XI. Target Analyte Quantitation

All target analyte quantitations met validation criteria with the following exceptions:

Sample	Analyte	Flag	A or P
All samples in SDG 21J0137	All analytes reported as estimated maximum possible concentration (EMPC) and less than the reporting limit (RL).	U	A
All samples in SDG 21J0137	All analytes flagged "X" due to chlorinated diphenyl ether (CDPE) interference.	J (all detects)	A

## **XII. Target Analyte Identification**

All target analyte identifications met validation criteria.

## **XIII. System Performance**

The system performance was acceptable.

## **XIV. Overall Assessment of Data**

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

Due to results reported by the laboratory as EMPCs and CDPE interference, data were qualified as estimated in six samples.

The quality control criteria reviewed, other than those discussed above, were met and are considered acceptable.



**Duwamish AOC4  
Polychlorinated Dioxins/Dibenzofurans - Data Qualification Summary - SDG 21J0137**

Sample	Analyte	Flag	A or P	Reason
LDW21-IT653D LDW21-IT652A LDW21-IT662A LDW21-IT658A LDW21-IT648D LDW21-IT648E	All analytes reported as estimated maximum possible concentration (EMPC) and less than the reporting limit (RL).	U	A	Target analyte quantitation (EMPC)
LDW21-IT653D LDW21-IT652A LDW21-IT662A LDW21-IT658A LDW21-IT648D LDW21-IT648E	All analytes flagged "X" due to chlorinated diphenyl ether (CDPE) interference.	J (all detects)	A	Target analyte quantitation (CDPE interference)

**Duwamish AOC4  
Polychlorinated Dioxins/Dibenzofurans - Laboratory Blank Data Qualification Summary - SDG 21J0137**

No Sample Data Qualified in this SDG

**Duwamish AOC4  
Polychlorinated Dioxins/Dibenzofurans - Field Blank Data Qualification Summary - SDG 21J0137**

No Sample Data Qualified in this SDG

LDC #: 52703C21

**VALIDATION COMPLETENESS WORKSHEET**

Date: 07/21

SDG #: 21J0137

Stage 4

Page: 1 of 1

Laboratory: Analytical Resources, Inc., Tukwila, WA

Reviewer: [Signature]

2nd Reviewer: [Signature]

**METHOD:** HRGC/HRMS Polychlorinated Dioxins/Dibenzofurans (EPA Method 1613B)

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	A	
II.	HRGC/HRMS Instrument performance check	A	
III.	Initial calibration/ICV	A/A	FSB ≤ 20/3570    KCV ≤ QC limits
IV.	Continuing calibration	A	CCV ≤ QC limits
V.	Laboratory Blanks	SW	
VI.	Field blanks	N	
VII.	Matrix spike/Matrix spike duplicates	N	CS
VIII.	Laboratory control samples /SRM	A	LCs
IX.	Field duplicates	N	
X.	Internal standards	A	
XI.	Target analyte quantitation	SW	
XII.	Target analyte identification	A	
XIII.	System performance	A	
XIV.	Overall assessment of data	A	

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:

	Client ID	Lab ID	Matrix	Date
1	LDW21-IT653D	21J0137-01	Sediment	07/12/21
2	LDW21-IT652A	21J0137-02	Sediment	07/12/21
3	LDW21-IT662A	21J0137-14	Sediment	07/13/21
4	LDW21-IT658A	21J0137-15	Sediment	07/13/21
5	LDW21-IT648D	21J0137-16	Sediment	07/13/21
6	LDW21-IT648E	21J0137-17	Sediment	07/13/21
7				
8				
9				
10				

Notes:

B N0500- <del>21</del>				

**Method:** HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 1613B)

Validation Area	Yes	No	NA	Findings/Comments
<b>I. Technical holding times</b>				
All technical holding times were met.	√			
Cooler temperature criteria were met.	√			
<b>II. GC/MS Instrument performance check</b>				
Was PFK exact mass 380.9760 verified?	√			
Were the retention time windows established for all homologues?	√			
Was the chromatographic resolution between 2,3,7,8-TCDD and peaks representing any other unlabeled TCDD isomers $\leq 25\%$ ?	√			
Is the static resolving power at least 10,000 (10% valley definition)?	√			
Was the mass resolution adequately check with PFK?	√			
Was the presence of 1,2,8,9-TCDD and 1,3,4,6,8-PeCDF verified?	√			
<b>III. Initial calibration and Initial calibration verification</b>				
Was the initial calibration performed at 5 concentration levels?	√			
Were all percent relative standard deviations (%RSD) $\leq 20\%$ for unlabeled compounds and $\leq 35\%$ for unlabeled compounds?	√			
Did all calibration standards meet the Ion Abundance Ratio criteria?	√			
Was the signal to noise ratio for each target compound and labeled compound $\geq 10$ ?	√			
Was an initial calibration verification (ICV) standard analyzed after each initial calibration for each instrument?	√			
Were all ICV concentrations for the unlabeled and labeled compounds within QC limits?	√			
<b>IV. Continuing calibration</b>				
Was a continuing calibration performed at the beginning of each 12-hour period?	√			
Were all continuing calibration concentrations for the unlabeled and labeled compounds within QC limits?	√			
Did all continuing calibration standards meet the Ion Abundance Ratio criteria?	√			
<b>V. Blanks</b>				
Was a method blank associated with every sample in this SDG?	√			
Was a method blank performed for each matrix and whenever a sample extraction was performed?	√			
Was there contamination in the method blanks?	√	0		
<b>VI. Field blanks</b>				
Were field blanks identified in this SDG?		√		
Were target compounds detected in the field blanks?			√	
<b>VII. Matrix spike/Matrix spike duplicates</b>				
Were matrix spike (MS) and matrix spike duplicate (MSD) analyzed in this SDG?		√		
Were the MS/MSD percent recoveries (%R) and the relative percent differences (RPD) within the QC limits?			√	

Validation Area	Yes	No	NA	Findings/Comments
<b>VIII. Laboratory control samples</b>				
Was an LCS analyzed per extraction batch?	√			
Were the LCS percent recoveries (%R) and relative percent difference (RPD) within the QC limits?	√			
<b>IX. Field duplicates</b>				
Were field duplicate pairs identified in this SDG?		√		
Were target compounds detected in the field duplicates?			√	
<b>X. Labeled Compounds</b>				
Were labeled compounds within QC limits?	√	Ⓟ		
Was the minimum S/N ratio of all labeled compound peaks $\geq 10$ ?	√			
<b>XI. Compound quantitation</b>				
Did the laboratory LOQs/RLs meet the QAPP LOQs/RLs?	√			
Were the correct labeled compound, quantitation ion and relative response factor (RRF) used to quantitate the compound?	√			
Were compound quantitation and RLs adjusted to reflect all sample dilutions and dry weight factors applicable to level IV validation?	√			
<b>XII. Target compound identification</b>				
For 2,3,7,8 substituted congeners with associated labeled standards, were the retention times of the two quantitation peaks within -1 to 3 sec. of the RT of the labeled standard?	√			
For 2,3,7,8 substituted congeners without associated labeled standards, were the relative retention times of the two quantitation peaks within 0.005 time units of the RRT measured in the routine calibration?	√			
For non-2,3,7,8 substituted congeners, were the retention times of the two quantitation peaks within RT established in the performance check solution?	√			
Did selected ion current profile (SICP) contain all characteristic ions listed in Method 1613B, Table 8?	√			
Was the Ion Abundance Ratio for the two quantitation ions within criteria?		√		
Was the signal to noise ratio for each target compound $\geq 2.5$ and $\geq 10$ for the labeled compound?	√			
Does the maximum intensity of each specified characteristic ion coincide within $\pm 2$ seconds (includes labeled standards)?	√			
For PCDF identification, was any signal ( $S/N \geq 2.5$ , at $\pm$ seconds RT) detected in the corresponding PCDPE channel?			√	
Was an acceptable lock mass recorded and monitored?	√			
<b>XIII. System performance</b>				
System performance was found to be acceptable.	√			
<b>XIV. Overall assessment of data</b>				
Overall assessment of data was found to be acceptable.	√			

## VALIDATION FINDINGS WORKSHEET

**METHOD:** HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 1613B)

A. 2,3,7,8-TCDD	F. 1,2,3,4,6,7,8-HpCDD	K. 1,2,3,4,7,8-HxCDF	P. 1,2,3,4,7,8,9-HpCDF	U. Total HpCDD
B. 1,2,3,7,8-PeCDD	G. OCDD	L. 1,2,3,6,7,8-HxCDF	Q. OCDF	V. Total TCDF
C. 1,2,3,4,7,8-HxCDD	H. 2,3,7,8-TCDF	M. 2,3,4,6,7,8-HxCDF	R. Total TCDD	W. Total PeCDF
D. 1,2,3,6,7,8-HxCDD	I. 1,2,3,7,8-PeCDF	N. 1,2,3,7,8,9-HxCDF	S. Total PeCDD	X. Total HxCDF
E. 1,2,3,7,8,9-HxCDD	J. 2,3,4,7,8-PeCDF	O. 1,2,3,4,6,7,8-HpCDF	T. Total HxCDD	Y. Total HpCDF

Notes: \_\_\_\_\_  
\_\_\_\_\_  
\_\_\_\_\_

**VALIDATION FINDINGS WORKSHEET**  
**Blanks**

**METHOD:** HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 1613B)

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

- Y N N/A Were all samples associated with a method blank?
- Y N N/A Was a method blank performed for each matrix and whenever a sample extraction was performed?
- Y N N/A Was the method blank contaminated?

Blank extraction date: 10/19/21      Blank analysis date: 10/25/21

Associated samples: All

Conc. units: ng/kg

Compound	Blank ID	Sample Identification							
	<u>BL 0520 B K1</u>								
<u>☒</u>	<u>0.981</u>								
<u>X</u>	<u>0.100</u>								

Blank extraction date: \_\_\_\_\_ Blank analysis date: \_\_\_\_\_

Conc. units: \_\_\_\_\_ Associated Samples: \_\_\_\_\_

Compound	Blank ID	Sample Identification							

CIRCLED RESULTS WERE NOT QUALIFIED. ALL RESULTS NOT CIRCLED WERE QUALIFIED BY THE FOLLOWING STATEMENT:  
All contaminants within five times the method blank concentration were qualified as not detected, "U".

**VALIDATION FINDINGS WORKSHEET**  
**Compound Quantitation and Reported RLs**

**METHOD:** HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 1613B)

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

Y N N/A Were the correct labeled compound, quantitation ions and relative response factors (RRF) used to quantitate the compound?  
Y N N/A Compound quantitation and RLs were adjusted to reflect all sample dilutions and dry weight factors (if necessary).

#	Date	Sample ID	Finding	Associated Samples	Qualifications
		All	All compounds reported as estimated maximum possible concentration (EMPC) < RL		U/A
		All	All compounds flagged "X" due to chlorinated diphenyl ether interference		Jdets/A

Comments: See sample calculation verification worksheet for recalculations

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### VALIDATION FINDINGS WORKSHEET Initial Calibration Calculation Verification

**METHOD:** HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 1613B)

The Relative Response Factor (RRF), average RRF, and percent relative standard deviation (%RSD) were recalculated for the compounds identified below using the following calculations:

RRF =  $(A_x)(C_{is}) / (A_{is})(C_x)$   
 average RRF = sum of the RRFs/number of standards  
 %RSD =  $100 * (S/X)$

$A_x$  = Area of compound,                       $A_{is}$  = Area of associated internal standard  
 $C_x$  = Concentration of compound,            $C_{is}$  = Concentration of internal standard  
 S = Standard deviation of the RRFs,        X = Mean of the RRFs

#	Standard ID	Calibration Date	Compound (Reference Internal Standard)	Reported	Recalculated	Reported	Recalculated	Reported	Recalculated
				RRF (10/50 std)	RRF (10/50 std)	Average RRF (initial)	Average RRF (initial)	%RSD	%RSD
1	ICAL 01	8/11/21	2,3,7,8-TCDF ( <sup>13</sup> C-2,3,7,8-TCDF)	1.0832006	1.083746	1.107593	1.107593	3.6	3.6
			2,3,7,8-TCDD ( <sup>13</sup> C-2,3,7,8-TCDD)	0.9085186	0.908390	0.9202875	0.9202874	3.1	3.1
			1,2,3,6,7,8-HxCDD ( <sup>13</sup> C-1,2,3,6,7,8-HxCDD)	1.005616	1.005605	1.00898	1.00898	1.0	1.0
			1,2,3,4,6,7,8-HpCDD ( <sup>13</sup> C-1,2,4,6,7,8,-HpCDD)	1.051009	1.051062	1.068088	1.068088	6.6	6.6
			OCDF ( <sup>13</sup> C-OCDD)	1.440564	1.44059	1.44690	1.44690	5.7	5.7
2			2,3,7,8-TCDF ( <sup>13</sup> C-2,3,7,8-TCDF)						
			2,3,7,8-TCDD ( <sup>13</sup> C-2,3,7,8-TCDD)						
			1,2,3,6,7,8-HxCDD ( <sup>13</sup> C-1,2,3,6,7,8-HxCDD)						
			1,2,3,4,6,7,8-HpCDD ( <sup>13</sup> C-1,2,4,6,7,8,-HpCDD)						
			OCDF ( <sup>13</sup> C-OCDD)						
3			2,3,7,8-TCDF ( <sup>13</sup> C-2,3,7,8-TCDF)						
			2,3,7,8-TCDD ( <sup>13</sup> C-2,3,7,8-TCDD)						
			1,2,3,6,7,8-HxCDD ( <sup>13</sup> C-1,2,3,6,7,8-HxCDD)						
			1,2,3,4,6,7,8-HpCDD ( <sup>13</sup> C-1,2,4,6,7,8,-HpCDD)						
			OCDF ( <sup>13</sup> C-OCDD)						

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.



**VALIDATION FINDINGS WORKSHEET**  
**Continuing Calibration Results Verification**

**METHOD:** HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 1613B)

The percent difference (%D) of the initial calibration average Relative Response Factors (RRFs) and the continuing calibration RRFs were recalculated for the compounds identified below using the following calculation:

% Difference = 100 \* (ave. RRF - RRF)/ave. RRF  
 RRF = (A<sub>x</sub>)(C<sub>is</sub>)/(A<sub>is</sub>)(C<sub>x</sub>)

Where: ave. RRF = initial calibration average RRF  
 RRF = continuing calibration RRF  
 A<sub>x</sub> = Area of compound, A<sub>is</sub> = Area of associated internal standard  
 C<sub>x</sub> = Concentration of compound, C<sub>is</sub> = Concentration of internal standard

#	Standard ID	Calibration Date	Compound (Reference Internal Standard)	Average RRF (initial)	Reported	Recalculated	Reported	Recalculated
					Conc (CC)	Conc (CC)	%D	%D
1	21102505A	10/5/21	2,3,7,8-TCDF ( <sup>13</sup> C-2,3,7,8-TCDF)	1.107593	1.0745550	1.0746175	3.0	3.0
			2,3,7,8-TCDD ( <sup>13</sup> C-2,3,7,8-TCDD)	0.920875	1.0081390	1.0081532	9.5	9.5
			1,2,3,6,7,8-HxCDD ( <sup>13</sup> C-1,2,3,6,7,8-HxCDD)	1.00898	1.0688370	1.0683744	5.9	5.9
			1,2,3,4,6,7,8-HpCDD ( <sup>13</sup> C-1,2,4,6,7,8,-HpCDD)	1.068088	1.1679010	1.1678182	9.3	9.3
			OCDF ( <sup>13</sup> C-OCDF)	1.44690	1.3382880	1.338548	7.5	7.5
2	21102518	10/26/21	2,3,7,8-TCDF ( <sup>13</sup> C-2,3,7,8-TCDF)	1.107593	1.0713550	1.0713484	3.3	3.3
			2,3,7,8-TCDD ( <sup>13</sup> C-2,3,7,8-TCDD)	0.920875	1.0205990	1.0206664	10.9	10.9
			1,2,3,6,7,8-HxCDD ( <sup>13</sup> C-1,2,3,6,7,8-HxCDD)	1.00898	1.0288700	1.028884	2.0	2.0
			1,2,3,4,6,7,8-HpCDD ( <sup>13</sup> C-1,2,4,6,7,8,-HpCDD)	1.068088	1.1328670	1.1328398	6.1	6.1
			OCDF ( <sup>13</sup> C-OCDF)	1.44690	1.3304760	1.3304528	8.0	8.0
3			2,3,7,8-TCDF ( <sup>13</sup> C-2,3,7,8-TCDF)					
			2,3,7,8-TCDD ( <sup>13</sup> C-2,3,7,8-TCDD)					
			1,2,3,6,7,8-HxCDD ( <sup>13</sup> C-1,2,3,6,7,8-HxCDD)					
			1,2,3,4,6,7,8-HpCDD ( <sup>13</sup> C-1,2,4,6,7,8,-HpCDD)					
			OCDF ( <sup>13</sup> C-OCDF)					

Comments: Refer to Routine 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 Laboratory Control Sample Results Verification

**METHOD:** HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 1613B)

The percent recoveries (%R) and Relative Percent Difference (RPD) of the laboratory control sample and laboratory control sample duplicate (if applicable) were recalculated for the compounds identified below using the following calculation:

% Recovery =  $100 * SSC/SA$       Where: SSC = Spiked sample concentration  
SA = Spike added

RPD =  $|LCS - LCSD| * 2 / (LCS + LCSD)$       LCS = Laboratory control sample percent recovery      LCSD = Laboratory control sample duplicate percent recovery

LCS ID: BH0500-BS

Compound	Spike Added (NSF)		Spiked Sample Concentration (NSF)		LCS		LCSD		LCS/LCSD	
	LCS	LCSD	LCS	LCSD	Percent Recovery		Percent Recovery		RPD	
					Reported	Recalc	Reported	Recalc	Reported	Recalculated
2,3,7,8-TCDD	20.0	NA	21.0	NA	105	105				
1,2,3,7,8-PeCDD	100	↓	107	↓	107	107				
1,2,3,4,7,8-HxCDD	↓	↓	99.2	↓	99.2	99.2				
1,2,3,4,7,8,9-HpCDF	↓	↓	95.9	↓	95.9	95.9				
OCDF	200	↓	151	↓	75.5	75.5				

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



**Laboratory Data Consultants, Inc.  
Data Validation Report**

**Project/Site Name:** Duwamish AOC4  
**LDC Report Date:** December 15, 2021  
**Parameters:** Semivolatiles  
**Validation Level:** Stage 4  
**Laboratory:** Analytical Resources, Inc., Tukwila, WA  
**Sample Delivery Group (SDG):** 21J0142

<b>Sample Identification</b>	<b>Laboratory Sample Identification</b>	<b>Matrix</b>	<b>Collection Date</b>
LDW21-IT621B	21J0142-23	Sediment	08/02/21
LDW21-IT621BMS	21J0142-23MS	Sediment	08/02/21
LDW21-IT621BMSD	21J0142-23MSD	Sediment	08/02/21

## 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 Final Lower Duwamish Waterway Quality Assurance Project Plan for Remedial Design of Upper Reach: Pre-Design Investigation (May 2020) and a modified outline of the USEPA National Functional Guidelines (NFG) for Organic Superfund Methods Data Review (January 2017). 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:

Semivolatile Organic Compounds (SVOCs) by Environmental Protection Agency (EPA) SW 846 Method 8270E

All sample results were subjected to Stage 4 data validation, which is comprised of the quality control (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): 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.
- U (Non-detected): The analyte was analyzed for and positively identified by the laboratory; however the analyte should be considered non-detected at the reported concentration due to the presence of contaminants detected in the associated blank(s).
- UJ (Non-detected estimated): The analyte was reported as not detected by the laboratory; however the reported quantitation/detection limit is estimated due to non-conformances discovered during data validation.
- R (Rejected): The sample results were rejected due to gross non-conformances discovered during data validation. Data qualified as rejected is not usable.
- 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.

## 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. GC/MS Instrument Performance Check

A decafluorotriphenylphosphine (DFTPP) tune was performed at 12 hour intervals.

All ion abundance requirements were met.

## III. 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.

Average relative response factors (RRF) for all analytes were within validation criteria.

The percent differences (%D) of the initial calibration verification (ICV) standard were less than or equal to 30.0% for all analytes.

## IV. 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 with the following exceptions:

Date	Analyte	%D	Associated Samples	Flag	A or P
10/30/21	Butylbenzylphthalate	25.7	All samples in SDG 21J0142	UJ (all non-detects)	A

All of the continuing calibration relative response factors (RRF) were within validation criteria.

## V. Laboratory Blanks

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

## VI. Field Blanks

No field blanks were identified in this SDG.

## **VII. Surrogates**

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

## **VIII. Matrix Spike/Matrix Spike Duplicates**

Matrix spike (MS) and matrix spike duplicate (MSD) sample analysis was performed on an associated project sample. Percent recoveries (%R) were within QC limits. Relative percent differences (RPD) were within QC limits.

## **IX. 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.

## **X. Field Duplicates**

No field duplicates were identified in this SDG.

## **XI. Internal Standards**

All internal standard areas and retention times were within QC limits.

## **XII. Target Analyte Quantitation**

All target analyte quantitations were within validation criteria.

## **XIII. Target Analyte Identification**

All target analyte identifications were within validation criteria.

## **XIV. System Performance**

The system performance was acceptable.

## **XV. Overall Assessment of Data**

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

Due to continuing calibration %D, data were qualified as estimated in one sample.

The quality control criteria reviewed, other than those discussed above, were met and are considered acceptable.

**Duwamish AOC4  
Semivolatiles - Data Qualification Summary - SDG 21J0142**

<b>Sample</b>	<b>Analyte</b>	<b>Flag</b>	<b>A or P</b>	<b>Reason</b>
LDW21-IT621B	Butylbenzylphthalate	UJ (all non-detects)	A	Continuing calibration (%D)

**Duwamish AOC4  
Semivolatiles - Laboratory Blank Data Qualification Summary - SDG 21J0142**

No Sample Data Qualified in this SDG

**Duwamish AOC4  
Semivolatiles - Field Blank Data Qualification Summary - SDG 21J0142**

No Sample Data Qualified in this SDG



LDC #: 52703D2a

**VALIDATION COMPLETENESS WORKSHEET**

Date: 1/7/21

SDG #: 21J0142

Stage 4

Page: 1 of 1

Laboratory: Analytical Resources, Inc., Tukwila, WA

Reviewer: [Signature]

2nd Reviewer: [Signature]

**METHOD:** GC/MS Butylbenzylphthalate (EPA SW 846 Method 8270E)

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	A	
II.	GC/MS Instrument performance check	A	
III.	Initial calibration/ICV	A/A	RSD ≤ 20%    1CV ≤ 20%
IV.	Continuing calibration	W	CCV ≤ 20%
V.	Laboratory Blanks	A	
VI.	Field blanks	N	
VII.	Surrogate spikes	A	
VIII.	Matrix spike/Matrix spike duplicates	A	
IX.	Laboratory control samples	A	LCS/D
X.	Field duplicates	N	
XI.	Internal standards	A	
XII.	Target analyte quantitation	A	
XIII.	Target analyte identification	A	
XIV.	System performance	A	
XV.	Overall assessment of data	D	

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:

	Client ID	Lab ID	Matrix	Date
1	LDW21-IT621B	21J0142-23	Sediment	08/02/21
2	LDW21-IT621BMS	21J0142-23MS	Sediment	08/02/21
3	LDW21-IT621BMSD	21J0142-23MSD	Sediment	08/02/21
4				
5				
6				
7				
8				
9				

Notes:

<u>BW07947K1</u>				

**Method: Semivolatiles (EPA SW 846 Method 8270D)**

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>II. GC/MS Instrument performance check</b>				
Were the DFTPP performance results reviewed and found to be within the specified criteria?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were all samples analyzed within the 12 hour clock criteria?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
<b>IIIa. 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) $\leq$ 20% and relative response factors (RRF) within method criteria?	<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 $> 0.990$ ?	<input type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	
<b>IIIb. 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) $\leq$ 30%?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
<b>IV. Continuing calibration</b>				
Was a continuing calibration standard analyzed at least once every 12 hours for each instrument?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were all percent differences (%D) $\leq$ 20% and relative response factors (RRF) within method criteria?	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	
<b>V. 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 at least once every 12 hours for each matrix and concentration?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was there contamination in the laboratory blanks? If yes, please see the blanks validation findings worksheet.	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	
<b>VI. Field blanks</b>				
Were field blanks were identified in this SDG?	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	
Were target compounds detected in the field blanks?	<input type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	
<b>VII. Surrogate spikes</b>				
Were all surrogate percent recovery (%R) within QC limits?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
If 2 or more base neutral or acid surrogates were outside QC limits, was a reanalysis performed to confirm %R?	<input type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	
If any percent recoveries (%R) was less than 10%, was a reanalysis performed to confirm %R ?	<input type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	
<b>VIII. 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"/>	

Validation Area	Yes	No	NA	Findings/Comments
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>IX. Laboratory control samples</b>				
Was an LCS analyzed per 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"/>	
<b>X. Field duplicates</b>				
Were field duplicate pairs identified in this SDG?	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	
Were target compounds detected in the field duplicates?	<input type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	
<b>XI. Internal standards</b>				
Were internal standard area counts within -50% to +100% of the associated calibration standard?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were retention times within + 30 seconds of the associated calibration standard?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
<b>XII. Compound quantitation</b>				
Did the laboratory LOQs/RLs meet the QAPP LOQs/RLs?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were the correct internal standard (IS), quantitation ion and relative response factor (RRF) used to quantitate the compound?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were compound 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>XIII. Target compound identification</b>				
Were relative retention times (RRT's) within + 0.06 RRT units of the standard?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Did compound spectra meet specified EPA "Functional Guidelines" criteria?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were chromatogram peaks verified and accounted for?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
<b>XIV. System performance</b>				
System performance was found to be acceptable.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
<b>XV. Overall assessment of data</b>				
Overall assessment of data was found to be acceptable.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	

## VALIDATION FINDINGS WORKSHEET

**METHOD: GC/MS SVOA**

A. Phenol	CC. Dimethylphthalate	EEE. Bis(2-ethylhexyl)phthalate	GGGG. C30-Hopane	I1. Methyl methanesulfonate
B. Bis (2-chloroethyl) ether	DD. Acenaphthylene	FFF. Di-n-octylphthalate	HHHH. 1-Methylphenanthrene	J1. Ethyl methanesulfonate
C. 2-Chlorophenol	EE. 2,6-Dinitrotoluene	GGG. Benzo(b)fluoranthene	IIII. 1,4-Dioxane	K1. o,o',o''-Triethylphosphorothioate
D. 1,3-Dichlorobenzene	FF. 3-Nitroaniline	HHH. Benzo(k)fluoranthene	JJJJ. Acetophenone	L1. n-Phenylene diamine
E. 1,4-Dichlorobenzene	GG. Acenaphthene	III. Benzo(a)pyrene	KKKK. Atrazine	M1. 1,4-Naphthoquinone
F. 1,2-Dichlorobenzene	HH. 2,4-Dinitrophenol	JJJ. Indeno(1,2,3-cd)pyrene	LLLL. Benzaldehyde	N1. N-Nitro-o-toluidine
G. 2-Methylphenol	II. 4-Nitrophenol	KKK. Dibenz(a,h)anthracene	MMMM. Caprolactam	O1. 1,3,5-Trinitrobenzene
H. 2,2'-Oxybis(1-chloropropane)	JJ. Dibenzofuran	LLL. Benzo(g,h,i)perylene	NNNN. 2,6-Dichlorophenol	P1. Pentachlorobenzene
I. 4-Methylphenol	KK. 2,4-Dinitrotoluene	MMM. Bis(2-Chloroisopropyl)ether	OOOO. 1,2-Diphenylhydrazine	Q1. 4-Aminobiphenyl
J. N-Nitroso-di-n-propylamine	LL. Diethylphthalate	NNN. Aniline	PPPP. 3-Methylphenol	R1. 2-Naphthylamine
K. Hexachloroethane	MM. 4-Chlorophenyl-phenyl ether	OOO. N-Nitrosodimethylamine	QQQQ. 3&4-Methylphenol	S1. Triphenylene
L. Nitrobenzene	NN. Fluorene	PPP. Benzoic Acid	RRRR. 4-Dimethyldibenzothiophene (4MDT)	T1. Octachlorostyrene
M. Isophorone	OO. 4-Nitroaniline	QQQ. Benzyl alcohol	SSSS. 2/3-Dimethyldibenzothiophene (4MDT)	U1. Famphur
N. 2-Nitrophenol	PP. 4,6-Dinitro-2-methylphenol	RRR. Pyridine	TTTT. 1-Methyldibenzothiophene (1MDT)	V1. 1,4-phenylenediamine
O. 2,4-Dimethylphenol	QQ. N-Nitrosodiphenylamine	SSS. Benzidine	UUUU.. 2,3,4,6-Tetrachlorophenol	W1. Methapyrilene
P. Bis(2-chloroethoxy)methane	RR. 4-Bromophenyl-phenylether	TTT. 1-Methylnaphthalene	VVVV. 1,2,4,5-Tetrachlorobenzene	X1. Pentachloroethane
Q. 2,4-Dichlorophenol	SS. Hexachlorobenzene	UUU. Benzo(b)thiophene	WWWW.. 2-Picoline	Y1. 3,3'-Dimethylbenzidine
R. 1,2,4-Trichlorobenzene	TT. Pentachlorophenol	VVV. Benzonaphthothiophene	XXXX. 3-Methylcholanthrene	Z1. o-Toluidine
S. Naphthalene	UU. Phenanthrene	WWW. Benzo(e)pyrene	YYYY. a, a-Dimethylphenethylamine	A2. 1-Naphthylamine
T. 4-Chloroaniline	VV. Anthracene	XXX. 2,6-Dimethylnaphthalene	ZZZZ. Hexachloropropene	B2. 4-Aminobiphenyl
U. Hexachlorobutadiene	WW. Carbazole	YYY. 2,3,5-Trimethylnaphthalene	A1. N-Nitrosodiethylamine	C2. 4-Nitroquinoline-1-oxide
V. 4-Chloro-3-methylphenol	XX. Di-n-butylphthalate	ZZZ. Perylene	B1. N-Nitrosodi-n-butylamine	D2. Hexachloropene
W. 2-Methylnaphthalene	YY. Fluoranthene	AAAA. Dibenzothiophene	C1. N-Nitrosomethylethylamine	E2. Bis (2-chloro-1-methylethyl) ether
X. Hexachlorocyclopentadiene	ZZ. Pyrene	BBBB. Benzo(a)fluoranthene	D1. N-Nitrosomorpholine	F2. Bifenthrin
Y. 2,4,6-Trichlorophenol	AAA. Butylbenzylphthalate	CCCC. Benzo(b)fluorene	E1. N-Nitrosopyrrolidine	G2. Cyfluthrin
Z. 2,4,5-Trichlorophenol	BBB. 3,3'-Dichlorobenzidine	DDDD. cis/trans-Decalin	F1. Phenacetin	H2. Cypermethrin
AA. 2-Chloronaphthalene	CCC. Benzo(a)anthracene	EEEE. 1,1'-Biphenyl	G1. 2-Acetylaminofluorene	I2. Permethrin (cis/trans)
BB. 2-Nitroaniline	DDD. Chrysene	FFFF. Retene	H1. Pronamide	J2. 5-Nitro-o-toluidine

VALIDATION FINDINGS WORKSHEET  
Continuing Calibration

METHOD: GC/MS BNA (EPA SW 846 Method 8270D)

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

 N N/A Was a continuing calibration standard analyzed at least once every 12 hours for each instrument? N N/A Were percent differences (%D)  $\leq 20\%$  and relative response factors (RRF) within the method criteria?

#	Date	Standard ID	Compound	Finding %D (Limit: $\leq 20.0\%$ )	Finding RRF (Limit)	Associated Samples	Qualifications
	10/20/11	NT102110302	AAA	25.7		All (ND)	√/N/A

## VALIDATION FINDINGS WORKSHEET Initial Calibration Calculation Verification

**METHOD:** GC/MS BNA (EPA SW 846 Method 8270D)

The Relative Response Factor (RRF), average RRF, and percent relative standard deviation (%RSD) were recalculated for the compounds identified below using the following calculations:

$$RRF = (A_x)(C_{is}) / (A_{is})(C_x)$$

average RRF = sum of the RRFs/number of standards  
%RSD = 100 \* (S/X)

$A_x$  = Area of compound,  
 $C_x$  = Concentration of compound,  
S = Standard deviation of the RRFs,

$A_{is}$  = Area of associated internal standard  
 $C_{is}$  = Concentration of internal standard  
X = Mean of the RRFs

#	Standard ID	Calibration Date	Compound (Reference Internal Standard)	Reported	Recalculated	Reported	Recalculated	Reported	Recalculated
				RRF ( <u>5</u> std)	RRF ( <u>5</u> std)	Average RRF (initial)	Average RRF (initial)	%RSD	%RSD
1	ICAC	10/25/21	Phenol (1st internal standard) <u>AAA</u>	0.9138844	0.9138844	0.9488406	0.9488406	5.3	5.3
			Naphthalene (2nd internal standard)						
			Fluorene (3rd internal standard)						
			Pentachlorophenol (4th internal standard)						
			Bis(2-ethylhexyl)phthalate (5th internal standard)						
			Benzo(a)pyrene (6th internal standard)						
2			Phenol (1st internal standard)						
			Naphthalene (2nd internal standard)						
			Fluorene (3rd internal standard)						
			Pentachlorophenol (4th internal standard)						
			Bis(2-ethylhexyl)phthalate (5th internal standard)						
			Benzo(a)pyrene (6th internal standard)						
3			Phenol (1st internal standard)						
			Naphthalene (2nd internal standard)						
			Fluorene (3rd internal standard)						
			Pentachlorophenol (4th internal standard)						
			Bis(2-ethylhexyl)phthalate (5th internal standard)						
			Benzo(a)pyrene (6th internal standard)						

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.

## VALIDATION FINDINGS WORKSHEET Continuing Calibration Results Verification

**METHOD:** GC/MS BNA (EPA SW 846 Method 8270D)

The percent difference (%D) of the initial calibration average Relative Response Factors (RRFs) and the continuing calibration RRFs were recalculated for the compounds identified below using the following calculation:

% Difference = 100 \* (ave. RRF - RRF)/ave. RRF  
 $RRF = (A_x)(C_{is}) / (A_{is})(C_x)$

Where: ave. RRF = initial calibration average RRF  
 RRF = continuing calibration RRF  
 $A_x$  = Area of compound,  $A_{is}$  = Area of associated internal standard  
 $C_x$  = Concentration of compound,  $C_{is}$  = Concentration of internal standard

#	Standard ID	Calibration Date	Compound (Reference Internal Standard)	Average RRF (Initial)	Reported	Recalculated	Reported	Recalculated
					RRF (CC)	RRF (CC)	%D	%D
1	<del>NT1021103002</del>	<del>10/30/21</del>	Phenol (1st internal standard) <del>AAA</del>	<del>0.9488406</del>	1.1931350	1.1931348	25.7	25.7
			Naphthalene (2nd internal standard)					
			Fluorene (3rd internal standard)					
			Pentachlorophenol (4th internal standard)					
			Bis(2-ethylhexyl)phthalate (5th internal standard)					
			Benzo(a)pyrene (6th internal standard)					
2			Phenol (1st internal standard)					
			Naphthalene (2nd internal standard)					
			Fluorene (3rd internal standard)					
			Pentachlorophenol (4th internal standard)					
			Bis(2-ethylhexyl)phthalate (5th internal standard)					
			Benzo(a)pyrene (6th internal standard)					
3			Phenol (1st internal standard)					
			Naphthalene (2nd internal standard)					
			Fluorene (3rd internal standard)					
			Pentachlorophenol (4th internal standard)					
			Bis(2-ethylhexyl)phthalate (5th internal standard)					
			Benzo(a)pyrene (6th internal standard)					

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/MS Semivolatiles (EPA SW 846 Method 8270D)

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 Spiked	Surrogate Found	Percent Recovery Reported	Percent Recovery Recalculated	Percent Difference
Nitrobenzene-d5					
2-Fluorobiphenyl					
Terphenyl-d14	5.0	3.94505	78.9	78.9	
Phenol-d5					
2-Fluorophenol					
2,4,6-Tribromophenol					
2-Chlorophenol-d4					
1,2-Dichlorobenzene-d4					

Sample ID: \_\_\_\_\_

	Surrogate Spiked	Surrogate Found	Percent Recovery Reported	Percent Recovery Recalculated	Percent Difference
Nitrobenzene-d5					
2-Fluorobiphenyl					
Terphenyl-d14					
Phenol-d5					
2-Fluorophenol					
2,4,6-Tribromophenol					
2-Chlorophenol-d4					
1,2-Dichlorobenzene-d4					

Sample ID: \_\_\_\_\_

	Surrogate Spiked	Surrogate Found	Percent Recovery Reported	Percent Recovery Recalculated	Percent Difference
Nitrobenzene-d5					
2-Fluorobiphenyl					
Terphenyl-d14					
Phenol-d5					
2-Fluorophenol					
2,4,6-Tribromophenol					
2-Chlorophenol-d4					
1,2-Dichlorobenzene-d4					



**VALIDATION FINDINGS WORKSHEET**  
**Matrix Spike/Matrix Spike Duplicates Results Verification**

**METHOD:** GC/MS BNA (EPA SW 846 Method 8270D)

The percent recoveries (%R) and Relative Percent Difference (RPD) of the matrix spike and matrix spike duplicate were recalculated for the compounds identified below using the following calculation:

% Recovery =  $100 * (SSC - SC) / SA$

Where: SSC = Spiked sample concentration  
 SA = Spike added

SC = Sample concentration

RPD =  $|MSC - MSC| * 2 / (MSC + MSC)$

MSC = Matrix spike concentration

MSDC = Matrix spike duplicate concentration

MS/MSD samples: 2/3

Compound	Spike Added (µg/L)		Sample Concentration (µg/L)	Spiked Sample Concentration (µg/L)		Matrix Spike		Matrix Spike Duplicate		MS/MSD	
	MS	MSD		MS	MSD	Percent Recovery		Percent Recovery		RPD	
						Reported	Recalc	Reported	Recalc	Reported	Recalculated
Phenol											
N-Nitroso-di-n-propylamine											
4-Chloro-3-methylphenol											
Acenaphthene											
Pentachlorophenol											
Pyrene											
<u>AAA</u>	<u>500</u>	<u>500</u>	<u>ND</u>	<u>405</u>	<u>430</u>	<u>81.0</u>	<u>81.0</u>	<u>85.9</u>	<u>86.0</u>	<u>3.93</u>	<u>5.8</u>

Comments: Refer to Matrix Spike/Matrix Spike Duplicates 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

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

**METHOD:** GC/MS Semivolatiles (EPA SW 846 Method 8270D)

The percent recoveries (%R) and Relative Percent Difference (RPD) of the laboratory control sample and laboratory control sample duplicate were recalculated for the compounds identified below using the following calculation:

% Recovery = 100 \* (SC/SA)

Where: SSC = Spike concentration  
 SA = Spike added

RPD = | LCSC - LCSDC | \* 2 / (LCSC + LCSDC)

LCSC = Laboratory control sample concentration    LCSDC = Laboratory control sample duplicate concentration

LCS/LCSD samples: B110794-B51 / -B501

Compound	Spike Added ( <u>µg/L</u> )		Spike Concentration ( <u>µg/L</u> )		LCS		LCSD		LCS/LCSD	
	LCS	LCSD	LCS	LCSD	Percent Recovery		Percent Recovery		RPD	
					Reported	Recalc	Reported	Recalc	Reported	Recalculated
Phenol										
N-Nitroso-di-n-propylamine										
4-Chloro-3-methylphenol										
Acenaphthene										
Pentachlorophenol										
Pyrene										
<u>AAA</u>	<u>500</u>	<u>500</u>	<u>458</u>	<u>472</u>	<u>91.5</u>	<u>91.5</u>	<u>94.3</u>	<u>94.3</u>	<u>3.01</u>	<u>2.9</u>

Comments: Refer to Laboratory Control Sample/Laboratory Control Sample Duplicates 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 Sample Calculation Verification

**METHOD:** GC/MS SVOA (EPA SW 846 Method 8270D)

Y N N/A  
Y N N/A

Were all reported results recalculated and verified for all level IV samples?

Were all recalculated results for detected target compounds agree within 10.0% of the reported results?

$$\text{Concentration} = \frac{(A_x)(I_s)(V_i)(DF)(2.0)}{(A_s)(RRF)(V_o)(V_i)(\%S)}$$

- A<sub>x</sub> = Area of the characteristic ion (EICP) for the compound to be measured
- A<sub>s</sub> = Area of the characteristic ion (EICP) for the specific internal standard
- I<sub>s</sub> = Amount of internal standard added in nanograms (ng)
- V<sub>o</sub> = Volume or weight of sample extract in milliliters (ml) or grams (g).
- V<sub>i</sub> = Volume of extract injected in microliters (ul)
- V<sub>t</sub> = Volume of the concentrated extract in microliters (ul)
- Df = Dilution Factor.
- %S = Percent solids, applicable to soil and solid matrices only.
- 2.0 = Factor of 2 to account for GPC cleanup

Example:

Sample I.D. NO AAA  
BNO794-B501

$$\text{Conc.} = \frac{(29041) \times (4.0) \times (1000) \times (1)}{(195770) \times (0.9488406) \times (10) \times ( )}$$

= 471.6  $\mu\text{g/L}$

#	Sample ID	Compound	Reported Concentration <u><math>\mu\text{g/L}</math></u>	Calculated Concentration ( )	Qualification
	<u>BNO794-B501</u>	<u>AAA</u>	<u>472</u>		

## Laboratory Data Consultants, Inc. Data Validation Report

**Project/Site Name:** Duwamish AOC4

**LDC Report Date:** December 15, 2021

**Parameters:** Polychlorinated Biphenyls

**Validation Level:** Stage 4

**Laboratory:** Analytical Resources, Inc., Tukwila

**Sample Delivery Group (SDG):** 21J0142

Sample Identification	Laboratory Sample Identification	Matrix	Collection Date
LDW21-SC525	21J0142-01	Sediment	07/15/21
LDW21-SS500	21J0142-02	Sediment	07/16/21
LDW21-SS501	21J0142-03	Sediment	07/16/21
LDW21-SS502	21J0142-04	Sediment	07/16/21
LDW21-IT579C	21J0142-06	Sediment	07/16/21
LDW21-IT597A	21J0142-07	Sediment	07/16/21
LDW21-IT597D	21J0142-08	Sediment	07/16/21
LDW21-SC673A	21J0142-09	Sediment	07/19/21
LDW21-IT600	21J0142-10	Sediment	07/19/21
LDW21-IT665D	21J0142-11	Sediment	07/19/21
LDW21-IT666D	21J0142-12	Sediment	07/19/21
LDW21-SS541	21J0142-13	Sediment	07/21/21
LDW21-IT512	21J0142-14	Sediment	07/19/21
LDW21-IT663D	21J0142-15	Sediment	07/19/21
LDW21-SC500	21J0142-16	Sediment	07/20/21
LDW21-SC501	21J0142-17	Sediment	07/20/21
LDW21-SC502	21J0142-18	Sediment	07/20/21
LDW21-SC563A	21J0142-19	Sediment	07/20/21
LDW21-SC628A	21J0142-20	Sediment	07/20/21
LDW21-IT670A	21J0142-22	Sediment	07/20/21
LDW21-IT621B	21J0142-23	Sediment	08/02/21
LDW21-SC628AMS	21J0142-20MS	Sediment	07/20/21
LDW21-SC628AMSD	21J0142-20MSD	Sediment	07/20/21

## 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 Final Lower Duwamish Waterway Quality Assurance Project Plan for Remedial Design of Upper Reach: Pre-Design Investigation (May 2020) and a modified outline of the USEPA National Functional Guidelines (NFG) for Organic Superfund Methods Data Review (January 2017). 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:

Polychlorinated Biphenyls (PCBs) by Environmental Protection Agency (EPA) SW 846 Method 8082A

All sample results were subjected to Stage 4 data validation, which is comprised of the quality control (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): 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.
- U (Non-detected): The analyte was analyzed for and positively identified by the laboratory; however the analyte should be considered non-detected at the reported concentration due to the presence of contaminants detected in the associated blank(s).
- UJ (Non-detected estimated): The analyte was reported as not detected by the laboratory; however the reported quantitation/detection limit is estimated due to non-conformances discovered during data validation.
- R (Rejected): The sample results were rejected due to gross non-conformances discovered during data validation. Data qualified as rejected is not usable.
- 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.

### **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.

Retention time windows were established as required by the method.

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 required frequencies.

The percent differences (%D) were less than or equal to 20.0% for all analytes with the following exceptions:

Date	Standard	Column	Analyte	%D	Associated Samples	Flag	A or P
10/27/21	10272115	2C	Aroclor-1260	24.6	LDW21-SC525	J (all detects)	A

Retention times of all analytes in the calibration standards were within the established retention time windows.

### **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/Internal Standards**

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

All internal standard percent recoveries (%R) were within QC limits.

### VII. Matrix Spike/Matrix Spike Duplicates

Matrix spike (MS) and matrix spike duplicate (MSD) sample analysis was performed on an associated project sample. Percent recoveries (%R) were within QC limits with the following exceptions:

Spike ID (Associated Samples)	Analyte	MS (%R) (Limits)	MSD (%R) (Limits)	Flag	A or P
LDW21-SC628AMS/MSD (LDW21-SC628A)	Aroclor-1260	49.5 (58-120)	-	J (all detects)	A

Relative percent differences (RPD) were within QC limits.

### VIII. Laboratory Control Samples/Standard Reference Materials

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.

Standard reference materials (SRM) were analyzed as required by the method. The results were within QC limits.

### IX. Field Duplicates

No field duplicates were identified in this SDG.

### X. Target Analyte Quantitation

All target analyte quantitations met validation criteria.

The sample results for detected analytes from the two columns were within 40% relative percent difference (RPD) with the following exceptions:

Sample	Analyte	RPD	Flag	A or P
LDW21-IT512	Aroclor-1260	42.3	J (all detects)	A
LDW21-SC563A	Aroclor-1254	41.5	J (all detects)	A

### XI. Target Analyte Identification

All target analyte identifications met validation criteria.

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

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

Due to continuing calibration %D, MS/MSD %R, and RPD between two columns, data were qualified as estimated in four samples.

The quality control criteria reviewed, other than those discussed above, were met and are considered acceptable.



**Duwamish AOC4  
Polychlorinated Biphenyls - Data Qualification Summary - SDG 21J0142**

Sample	Analyte	Flag	A or P	Reason
LDW21-SC525	Aroclor-1260	J (all detects)	A	Continuing calibration (%D)
LDW21-SC628A	Aroclor-1260	J (all detects)	A	Matrix spike/Matrix spike duplicate (%R)
LDW21-IT512	Aroclor-1260	J (all detects)	A	Target analyte quantitation (RPD between two columns)
LDW21-SC563A	Aroclor-1254	J (all detects)	A	Target analyte quantitation (RPD between two columns)

**Duwamish AOC4  
Polychlorinated Biphenyls - Laboratory Blank Data Qualification Summary - SDG 21J0142**

No Sample Data Qualified in this SDG

**Duwamish AOC4  
Polychlorinated Biphenyls - Field Blank Data Qualification Summary - SDG 21J0142**

No Sample Data Qualified in this SDG

LDC #: 52703D3b

**VALIDATION COMPLETENESS WORKSHEET**

Date: 7/16/21

SDG #: 21J0142

Stage 4

Page: 1 of 2

Laboratory: Analytical Resources, Inc.

Reviewer: G

2nd Reviewer: JVB

**METHOD:** GC Polychlorinated Biphenyls (EPA SW846 Method 8082A)

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	A	
II.	Initial calibration/ICV	A/A	FSO ≤ 20%      CV ≤ 20%
III.	Continuing calibration	W	CV ≤ 20%
IV.	Laboratory Blanks	A	
V.	Field blanks	N	
VI.	Surrogate spikes / IS	A/A	
VII.	Matrix spike/Matrix spike duplicates	W	
VIII.	Laboratory control samples / EB	A	LES/O
IX.	Field duplicates	N	
X.	Target analyte quantitation	W	
XI.	Target analyte identification	A	
XII.	Overall assessment of data	A	

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:

	Client ID	Lab ID	Matrix	Date
1	LDW21-SC525	21J0142-01	Sediment	07/15/21
2	LDW21-SS500	21J0142-02	Sediment	07/16/21
3	LDW21-SS501	21J0142-03	Sediment	07/16/21
4	LDW21-SS502	21J0142-04	Sediment	07/16/21
5	LDW21-IT579C	21J0142-06	Sediment	07/16/21
6	LDW21-IT597A	21J0142-07	Sediment	07/16/21
7	LDW21-IT597D	21J0142-08	Sediment	07/16/21
8	LDW21-SC673A	21J0142-09	Sediment	07/19/21
9	LDW21-IT600	21J0142-10	Sediment	07/19/21
10	LDW21-IT665D	21J0142-11	Sediment	07/19/21
11	LDW21-IT666D	21J0142-12	Sediment	07/19/21
12	LDW21-SS541	21J0142-13	Sediment	07/21/21
13	LDW21-IT512	21J0142-14	Sediment	07/19/21
14	LDW21-IT663D	21J0142-15	Sediment	07/19/21
15	LDW21-SC500	21J0142-16	Sediment	07/20/21
16	LDW21-SC501	21J0142-17	Sediment	07/20/21
17	LDW21-SC502	21J0142-18	Sediment	07/20/21

LDC #: 52703D3b

### VALIDATION COMPLETENESS WORKSHEET

SDG #: 21J0142

Stage 4

Laboratory: Analytical Resources, Inc.

Date: 7/20/21

Page: 2 of 2

Reviewer: [Signature]

2nd Reviewer: [Signature]

**METHOD:** GC Polychlorinated Biphenyls (EPA SW846 Method 8082A)

	Client ID	Lab ID	Matrix	Date
18	LDW21-SC563A	21J0142-19	Sediment	07/20/21
19	LDW21-SC628A	21J0142-20	Sediment	07/20/21
20	LDW21-IT670A	21J0142-22	Sediment	07/20/21
21	LDW21-IT621B	21J0142-23	Sediment	08/02/21
22	LDW21-SC628AMS	21J0142-20MS	Sediment	07/20/21
23	LDW21-SC628AMSD	21J0142-20MSD	Sediment	07/20/21
24				
25				
26				

Notes:

B-10633				
B-10636				

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) $\leq 20\%$ ?	<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) $\leq 20\%$ ?	<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) $\leq 20\%$ ?	<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 compounds 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 type="checkbox"/>	<input checked="" 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>IX. Field duplicates</b>				
Were field duplicate pairs identified in this SDG?		/		
Were target compounds detected in the field duplicates?			/	
<b>X. Compound quantitation</b>				
Did the laboratory LOQs/RLs meet the QAPP LOQs/RLs?	/			
Were compound quantitation and RLs adjusted to reflect all sample dilutions and dry weight factors applicable to level IV validation?	/			
<b>XI. Target compound identification</b>				
Were the retention times of reported detects within the RT windows?	/			
<b>XIII. Overall assessment of data</b>				
Overall assessment of data was found to be acceptable.	/			

LDC #: 5-103036

## VALIDATION FINDINGS WORKSHEET Continuing Calibration

Page: 1 of 1  
Reviewer: [Signature]

METHOD:  GC  HPLC

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

N  N/A Were continuing calibration standards analyzed at the required frequencies?

Y  N  N/A Did the continuing calibration standards meet the %D validation criteria of  $\leq 20.0\%$ ?

Level IV Only

Y  N  N/A Were the retention times for all calibrated compounds within their respective acceptance windows?

#	Date	Standard ID	Detector/ Column	Compound	%D (Limit)	RT (limit)	Associated Samples	Qualifications
	<u>10/5/11</u>	<u>10-T-115</u>	<u>2C</u>	<u>BB</u>	<u>24.6</u>	( )	<u>1. MB (dots)</u>	<u>✓ N/A</u> <u>(12 in)</u>
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		
						( )		

# VALIDATION FINDINGS WORKSHEET

## Matrix Spike/Matrix Spike Duplicates

METHOD:  GC  HPLC

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

N/A

Were a matrix spike (MS) and matrix spike duplicate (MSD) analyzed for each matrix in this SDG?

N/A

Was an MS/MSD analyzed every 20 samples for each matrix or whenever a sample extraction was performed?

N/A

Were the MS/MSD percent recoveries (%R) and relative percent differences (RPD) within QC limits?

#	MS/MSD ID	Compound	MS %R (Limits)	MSD %R (Limits)	RPD (Limits)	Associated Samples	Qualifications
	<u>22/23</u>	<u>BB</u>	<u>49.5 (58-120)</u>	( )	( )	<u>19 (det's)</u>	<u>N/A</u>
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		
			( )	( )	( )		

### VALIDATION FINDINGS WORKSHEET Compound Quantitation and Reported CRQLs

METHOD:  GC  HPLC

**Level IV/D Only**

Y  N  N/A  
 Y  N  N/A  
 Y  N  N/A

Were CRQLs adjusted for sample dilutions, dry weight factors, etc.?

Did the reported results for detected target compounds agree within 10.0% of the recalculated results?

Did the relative percent differences of detected compounds between two columns/detectors  $\leq$ 40%?

If no, please see findings below.

#	Compound Name	Sample ID	%RPD Between Two Columns/Detectors Limit ( $\leq$ 40%)	Qualifications
	Aroclor 1260	13	42.3	Jdets/A
	Aroclor 1254	18	41.5	Jdets/A



LDC #: 5203D30

**VALIDATION FINDINGS WORKSHEET**  
**Initial Calibration Calculation Verification**

Page: 1 of 1  
 Reviewer: 9

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 (100 std)	CF (100 std)	Ave CF (initial)	Ave CF (initial)	%RSD	%RSD
1	KAC	8/13/21	BB-1 (1c)	0.03587713	0.03587713	0.0359933	0.0359933	2.6	2.6
			BB-1 (2c)	0.06872649	0.0687264	0.06650318	0.0665032	T.T	T.8
2									
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 #: 5203D26

**VALIDATION FINDINGS WORKSHEET**  
**Continuing Calibration Results Verification**

Page: 1 of 1  
 Reviewer: 9

METHOD: GC\_HPLC

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

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

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

#	Standard ID	Calibration Date	Compound	Average CF(Ical)/ CCV Conc.	Reported	Recalculated	Reported	Recalculated
					CF/ Conc. CCV	CF/ Conc. CCV	%D	%D
1	<del>1025212</del>	10/6/21 2=19	BB-1 (1C)	0.0359923	0.0311367	0.0311366	12.0	11.8
	<del>3-4.6-13</del>			0.0665032	0.0543594	0.0543593	18.4	18.3
	<del>7-12</del>							
2	<del>10252133</del>	10/24/21 6=24	↓	0.0359923	0.0307250	0.0307249	14.8	14.6
	<del>13-23 MB</del>			0.0665032	0.0539980	0.0539979	18.8	18.8
3	<del>1025215</del>	10/27/21 18=25	↓	0.0359923	0.0284107	0.0284106	21.2	21.1
	1			0.0665032	0.0482295	0.0482294	27.6	27.5
4	<del>1025203</del>	10/25/21 20=10	↓	0.0359923	0.0299229	0.0299229	16.8	16.9
	2			0.0665032	0.0542807	0.0542806	18.4	18.4

**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>DCB</u>	<u>IC</u>	<u>8.0</u>	<u>8.3</u>	<u>104</u>	<u>104</u>	
<u>TELX</u>	<u>↓</u>	<u>↓</u>	<u>6.3</u>	<u>78.9</u>	<u>78.8</u>	

Sample ID: \_\_\_\_\_

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

Sample ID: \_\_\_\_\_

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

## VALIDATION FINDINGS WORKSHEET Matrix Spike/Matrix Spike Duplicates Results Verification

METHOD: GC HPLC

The percent recoveries (%R) and relative percent differences (RPD) of the matrix spike and matrix spike duplicate were recalculated for the compounds identified below using the following calculation:

$$\% \text{Recovery} = 100 * (\text{SSC} - \text{SC}) / \text{SA}$$

Where

SSC = Spiked sample concentration

SC = Sample concentration

SA = Spike added

$$\text{RPD} = ((\text{SSCMS} - \text{SSCMSD}) * 2) / (\text{SSCMS} + \text{SSCMSD}) * 100$$

MS = Matrix spike

MSD = Matrix spike duplicate

MS/MSD samples: 22/23

Compound	Spike Added ( <u>MS</u> )		Sample Conc. ( <u>MS</u> )	Spike Sample Concentration ( <u>MS</u> )		Matrix spike		Matrix Spike Duplicate		MS/MSD	
	MS	MSD		MS	MSD	Percent Recovery		Percent Recovery		RPD	
						Reported	Recalc.	Reported	Recalc.	Reported	Recalc.
Gasoline (8015)											
Diesel (8015)											
Benzene (8021B)											
Methane (RSK-175)											
2,4-D (8151)											
Dinoseb (8151)											
Naphthalene (8310)											
Anthracene (8310)											
HMX (8330)											
2,4,6-Trinitrotoluene (8330)											
<u>FB</u>	<u>101</u>	<u>101</u>	<u>15.4</u>	<u>65.4</u>	<u>67.1</u>	<u>49.5</u>	<u>49.5</u>	<u>51.2</u>	<u>51.2</u>	<u>2.50</u>	<u>2.57</u>

Comments: Refer to Matrix Spike/Matrix Spike Duplicates 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**  
**Laboratory Control Sample/Laboratory Control Sample Duplicates Results Verification**

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 compounds identified below using the following calculation:

$\% \text{Recovery} = 100 * (\text{SSC} - \text{SC}) / \text{SA}$

Where SSC = Spiked sample concentration  
 SA = Spike added  
 LCS = Laboratory Control Sample

SC = Sample concentration

$\text{RPD} = (((\text{SSCLCS} - \text{SSCLCSD}) * 2) / (\text{SSCLCS} + \text{SSCLCSD})) * 100$

LCSD = Laboratory Control Sample duplicate

LCS/LCSD samples: B10683

Compound	Spike Added (100)		Spike Sample Concentration (100)		LCS		LCSD		LCS/LCSD	
	LCS	LCSD	LCS	LCSD	Percent Recovery		Percent Recovery		RPD	
					Reported	Recalc.	Reported	Recalc.	Reported	Recalc.
Gasoline (8015)										
Diesel (8015)										
Benzene (8021B)										
Methane (RSK-175)										
2,4-D (8151)										
Dinoseb (8151)										
Naphthalene (8310)										
Anthracene (8310)										
HMX (8330)										
2,4,6-Trinitrotoluene (8330)										
<del>BD</del>	101	101	84.8	83.3	84.1	84.0	82.6	82.5	1.81	1.8

Comments: Refer to Laboratory Control Sample/Laboratory Control Sample Duplicate 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**  
**Sample Calculation Verification**

METHOD: ✓ GC     HPLC

✓ N N/A  
✓ N N/A

Were all reported results recalculated and verified for all level IV samples?  
 Were all recalculated results for detected target compounds within 10% of the reported results?

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

Example:

Sample ID. 1 Compound Name PCB-1260-1

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

Concentration =  $\frac{(20224) (80.0)}{(398163) (0.0759923)} = 112.9$

Concentration =  $\frac{(112.9 + 102.0 + 107.5 + 40.6 + 96.9) (2.5) (5)}{5 \times 20.93 \times 0.6001} = 111.4 \mu\text{g/g}$

#	Sample ID	Compound	Reported Concentrations ( <u>μg/g</u> )	Recalculated Results Concentrations ( <u>   </u> )	Qualifications
	<u>1</u>	<u>PCB-1260</u>	<u>111</u>		

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

**Laboratory Data Consultants, Inc.  
Data Validation Report**

**Project/Site Name:** Duwamish AOC4  
**LDC Report Date:** December 15, 2021  
**Parameters:** Arsenic  
**Validation Level:** Stage 4  
**Laboratory:** Analytical Resources, Inc., Tukwila, WA  
**Sample Delivery Group (SDG):** 21J0142

<b>Sample Identification</b>	<b>Laboratory Sample Identification</b>	<b>Matrix</b>	<b>Collection Date</b>
LDW21-IT582F	21J0142-05	Sediment	07/16/21
LDW21-IT600	21J0142-10	Sediment	07/19/21

## 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 Final Lower Duwamish Waterway Quality Assurance Project Plan for Remedial Design of Upper Reach: Pre-Design Investigation (May 2020) and a modified outline of the USEPA National Functional Guidelines (NFG) for Inorganic Superfund Methods Data Review (January 2017). 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:

Arsenic by Environmental Protection Agency (EPA) SW 846 Method 6020B

All sample results were subjected to Stage 4 evaluation, which is comprised of the quality control (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): 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.
- U (Non-detected): The analyte was analyzed for and positively identified by the laboratory; however the analyte should be considered non-detected at the reported concentration due to the presence of contaminants detected in the associated blank(s).
- UJ (Non-detected estimated): The analyte was reported as not detected by the laboratory; however the reported quantitation/detection limit is estimated due to non-conformances discovered during data validation.
- R (Rejected): The sample results were rejected due to gross non-conformances discovered during data validation. Data qualified as rejected is not usable.
- 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.



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

All samples were received in good condition.

All technical holding time requirements were met.

## **II. ICPMS Tune**

The mass calibration was within 0.1 AMU and the percent relative standard deviation (%RSD) was less than or equal to 5%.

## **III. Instrument Calibration**

Initial and continuing calibrations were performed as required by the method.

The initial calibration verification (ICV) and continuing calibration verification (CCV) standards were within QC limits.

## **IV. ICP Interference Check Sample Analysis**

The frequency of interference check sample (ICS) analysis was met. All criteria were within QC limits.

## **V. Laboratory Blanks**

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

## **VI. Field Blanks**

No field blanks were identified in this SDG.

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

The laboratory has indicated that there were no matrix spike (MS) and matrix spike duplicate (MSD) analyses specified for the samples in this SDG, and therefore matrix spike and matrix spike duplicate analyses were not performed for this SDG.

## **VIII. Duplicate Sample Analysis**

The laboratory has indicated that there were no duplicate (DUP) analyses specified for the samples in this SDG, and therefore duplicate analyses were not performed for this SDG.

## **IX. Serial Dilution**

Serial dilution was not performed for this SDG.

## **X. Laboratory Control Samples**

Laboratory control samples (LCS) were analyzed as required by the method. Percent recoveries (%R) were within QC limits.

## **XI. Field Duplicates**

No field duplicates were identified in this SDG.

## **XII. Internal Standards (ICP-MS)**

All internal standard percent recoveries (%R) were within QC limits.

## **XIII. Target Analyte Quantitation**

All target analyte quantitations were within validation criteria.

## **XIV. Overall Assessment of Data**

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

The quality control criteria reviewed were met and are considered acceptable.

**Duwamish AOC4  
Arsenic - Data Qualification Summary - SDG 21J0142**

No Sample Data Qualified in this SDG

**Duwamish AOC4  
Arsenic - Laboratory Blank Data Qualification Summary - SDG 21J0142**

No Sample Data Qualified in this SDG

**Duwamish AOC4  
Arsenic - Field Blank Data Qualification Summary - SDG 21J0142**

No Sample Data Qualified in this SDG

LDC #: 52703D4a

**VALIDATION COMPLETENESS WORKSHEET**

Date: 12/9/21

SDG #: 21J0142

Stage 4

Page: of

Laboratory: Analytical Resources, Inc., Tukwila, WA

Reviewer: [Signature]

2nd Reviewer: [Signature]

**METHOD:** Arsenic (EPA SW846 Method 6020B)

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	A/A	
II.	ICP/MS Tune	A	
III.	Instrument Calibration	A	
IV.	ICP Interference Check Sample (ICS) Analysis	A	
V.	Laboratory Blanks	A	
VI.	Field Blanks	N	
VII.	Matrix Spike/Matrix Spike Duplicates	N	
VIII.	Duplicate sample analysis	N	
IX.	Serial Dilution	N	
X.	Laboratory control samples	A	LCS
XI.	Field Duplicates	N	
XII.	Internal Standard (ICP-MS)	A	
XIII.	Target Analyte Quantitation	A	
XIV.	Overall Assessment of Data	A	

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:

	Client ID	Lab ID	Matrix	Date
1	LDW21-IT582F	21J0142-05	Sediment	07/16/21
2	LDW21-IT600	21J0142-10	Sediment	07/19/21
3				
4				
5				
6				
7				
8				
9				
10				
11				
12				
13				

Notes: \_\_\_\_\_  
 \_\_\_\_\_  
 \_\_\_\_\_

METHOD: Trace Metals (EPA SW 846 Methods 6010/6020/7000)				
Validation Area	Yes	No	NA	Comments
<b>I. Technical holding times</b>				
Were all technical holding times met?	X			
Were all water samples preserved to a pH of <2?			X	
<b>II. ICP-MS Tune</b>				
Were mass resolutions within 0.1 amu for all isotopes in the tuning solution?	X			
Were %RSDs of isotopes in the tuning solution ≤5%?	X			
<b>III. Calibration</b>				
Were all instruments calibrated daily?	X			
Were the proper standards used?	X			
Were all initial and continuing calibration verifications within the 90-110% (80-120% for mercury) QC limits?	X			
Were the low level standard checks within 70-130%?			X	
Were all initial calibration correlation coefficients within limits as specified by the method?	X			
<b>IV. Blanks</b>				
Was a method blank associated with every sample in this SDG?	X			
Was there contamination in the method blanks?		X		
Was there contamination in the initial and continuing calibration blanks?		X		
<b>V. Interference Check Sample</b>				
Were the interference check samples performed daily?	X			
Were the AB solution recoveries within 80-120%?	X			
<b>VI. Matrix Spike/Matrix Spike Duplicates/Laboratory Duplicates</b>				
Were MS/MSD recoveries with the QC limits? (If the sample concentration exceeded the spike concentration by a factor of 4, no action was taken.)			X	
Were the MS/MSD or laboratory duplicate relative percent differences (RPDs) within the QC limits?			X	
<b>VII. Laboratory Control Samples</b>				
Was a LCS analyzed for each batch in the SDG?	X			
Were the LCS recoveries and RPDs (if applicable) within QC limits?	X			

METHOD: Trace Metals (EPA SW 846 Methods 6010/6020/7000)				
Validation Area	Yes	No	NA	Comments
<b>VIII. Internal Standards</b>				
Were all percent recoveries within the 30-120% (60-125% for EPA Method 200.8) QC limits?	X			
If the recoveries were outside the limits, was a reanalysis performed?		X		
<b>IX. Serial Dilution</b>				
Were all percent differences <10%?			X	
Was there evidence of negative interference? If yes, professional judgement will be used to qualify the data.			X	
<b>X. Sample Result Verification</b>				
Were all reporting limits adjusted to reflect sample dilutions?	X			
Were all soil samples dry weight corrected?	X			
<b>XI. Overall Assessment of Data</b>				
Was the overall assessment of the data found to be acceptable?	X			
<b>XII. Field Duplicates</b>				
Were field duplicates identified in this SDG?		X		
Were target analytes detected in the field duplicates?			X	
<b>XIII. Field Blanks</b>				
Were field blanks identified in this SDG?		X		
Were target analytes detected in the field blanks?			X	

METHOD: Trace Metals (EPA SW 846 Methods 6010/6020/7000)

An initial calibration verification (ICV), continuing calibration verification (CCV), low level calibration check (LLCC), and interference check sample (ICSAB) percent recovery (%R) was recalculated for each type of analysis using the following formula:

$$\%R = (\text{Found}/\text{True}) \times 100$$

Found = concentration of each analyte measured in the analysis

True = concentration of each analyte in the source

Standard ID	Type of Analysis	Element	Found (ug/L)	True (ug/L)	Recalculated %R	Reported %R	Acceptable (Y/N)
ICV	ICP-MS	As	47.7	50	95.4	95.5	Y
CCV	ICP-MS	Cd	49.8	50	99.6	99.6	Y
ICSAB	ICP-MS	As	19.283	20	96.4	96.4	Y

ICP-MS Tune	QC Parameter	Mass	Actual	Required
10/28/2021	Mass Axis	115	114.9	± 0.1 amu
10/28/2021	%RSD	115	1	≤ 5%

METHOD: Trace Metals (EPA SW 846 Methods 6010/6020/7000)

Percent recoveries (%R) for the laboratory control sample (LCS), matrix spike (MS), and post digestion spike (PDS) were recalculated using the following formula:

$$\%R = (\text{Found}/\text{True}) \times 100$$

Found = concentration of each analyte measured in the analysis. For the MS calculation, Found = SSR (Spiked Sample Result) - SR (Sample Result)

True = concentration of each analyte in the source

The sample and duplicate relative percent difference (RPD) was recalculated using the following formula:

$$\text{RPD} = (\text{Absolute value}(S-D) \times 200) / (S+D)$$

S = Original sample concentration

D = Duplicate sample concentration

The serial dilution percent difference (%D) was recalculated using the following formula.

$$\%D = (\text{Absolute value}(I - \text{SDR})) \times 100 / (I)$$

I = Initial sample result

SDR = Serial dilution result (with a 5x dilution applied)

Sample ID	Type of Analysis	Element	Found/S/I	True/D/SDR	Recalculated %R/RPD/%D	Reported %R/RPD/%D	Acceptable (Y/N)
LCS	LCS	As	24.1	25	96.4	96.5	Y
	MS						
	Duplicate						
	PDS						
	Serial dilution						





## Laboratory Data Consultants, Inc. Data Validation Report

**Project/Site Name:** Duwamish AOC4

**LDC Report Date:** December 15, 2021

**Parameters:** Wet Chemistry

**Validation Level:** Stage 4

**Laboratory:** Analytical Resources, Inc., Tukwila, WA

**Sample Delivery Group (SDG):** 21J0142

Sample Identification	Laboratory Sample Identification	Matrix	Collection Date
LDW21-SC525	21J0142-01	Sediment	07/15/21
LDW21-SS500	21J0142-02	Sediment	07/16/21
LDW21-SS501	21J0142-03	Sediment	07/16/21
LDW21-SS502	21J0142-04	Sediment	07/16/21
LDW21-IT582F	21J0142-05	Sediment	07/16/21
LDW21-IT579C	21J0142-06	Sediment	07/16/21
LDW21-IT597A	21J0142-07	Sediment	07/16/21
LDW21-IT597D	21J0142-08	Sediment	07/16/21
LDW21-SC673A	21J0142-09	Sediment	07/19/21
LDW21-IT600	21J0142-10	Sediment	07/19/21
LDW21-IT665D	21J0142-11	Sediment	07/19/21
LDW21-IT666D	21J0142-12	Sediment	07/19/21
LDW21-SS541	21J0142-13	Sediment	07/21/21
LDW21-IT512	21J0142-14	Sediment	07/19/21
LDW21-IT663D	21J0142-15	Sediment	07/19/21
LDW21-SC500	21J0142-16	Sediment	07/20/21
LDW21-SC501	21J0142-17	Sediment	07/20/21
LDW21-SC502	21J0142-18	Sediment	07/20/21
LDW21-SC563A	21J0142-19	Sediment	07/20/21
LDW21-SC628A	21J0142-20	Sediment	07/20/21
LDW21-IT664A	21J0142-21	Sediment	07/20/21
LDW21-IT670A	21J0142-22	Sediment	07/20/21
LDW21-IT621B	21J0142-23	Sediment	08/02/21
LDW21-IT665DMS	21J0142-11MS	Sediment	07/19/21
LDW21-IT665DDUP	21J0142-11DUP	Sediment	07/19/21
LDW21-SC563ADUP1	21J0142-19DUP1	Sediment	07/20/21

<b>Sample Identification</b>	<b>Laboratory Sample Identification</b>	<b>Matrix</b>	<b>Collection Date</b>
LDW21-SC563ADUP2	21J0142-19DUP2	Sediment	07/20/21

## 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 Final Lower Duwamish Waterway Quality Assurance Project Plan for Remedial Design of Upper Reach: Pre-Design Investigation (May 2020) and a modified outline of the USEPA National Functional Guidelines (NFG) for Inorganic Superfund Methods Data Review (January 2017). 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 methods:

Total Organic Carbon by Environmental Protection Agency (EPA) SW 846 Method 9060A

Total Solids by Standard Method 2540G

All sample results were subjected to Stage 4 data validation, which is comprised of the quality control (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): 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.
- U (Non-detected): The analyte was analyzed for and positively identified by the laboratory; however the analyte should be considered non-detected at the reported concentration due to the presence of contaminants detected in the associated blank(s).
- UJ (Non-detected estimated): The analyte was reported as not detected by the laboratory; however the reported quantitation/detection limit is estimated due to non-conformances discovered during data validation.
- R (Rejected): The sample results were rejected due to gross non-conformances discovered during data validation. Data qualified as rejected is not usable.
- 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.

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

All samples were received in good condition.

All technical holding time requirements were met.

## **II. Initial Calibration**

All criteria for the initial calibration of each method were met.

## **III. Continuing Calibration**

Continuing calibration frequency and analysis criteria were met for each method when applicable.

## **IV. Laboratory Blanks**

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

## **V. Field Blanks**

No field blanks were identified in this SDG.

## **VI. Matrix Spike/Matrix Spike Duplicates**

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

## **VII. Duplicate Sample Analysis**

Duplicate (DUP) sample analysis was performed on an associated project sample. Results were within QC limits.

## **VIII. Laboratory Control Samples**

Laboratory control samples (LCS) were analyzed as required by the methods. Percent recoveries (%R) were within QC limits.

## **IX. Field Duplicates**

No field duplicates were identified in this SDG.

## **X. Target Analyte Quantitation**

All target analyte quantitations were within validation criteria.

## **XI. Overall Assessment of Data**

The analysis was conducted within all specifications of the methods. No results were rejected in this SDG.

The quality control criteria reviewed were met and are considered acceptable.

**Duwamish AOC4  
Wet Chemistry - Data Qualification Summary - SDG 21J0142**

No Sample Data Qualified in this SDG

**Duwamish AOC4  
Wet Chemistry - Laboratory Blank Data Qualification Summary - SDG 21J0142**

No Sample Data Qualified in this SDG

**Duwamish AOC4  
Wet Chemistry - Field Blank Data Qualification Summary - SDG 21J0142**

No Sample Data Qualified in this SDG



**METHOD: (Analyte) TOC (EPA SW846 Method 9060A), Total Solids (SM2540G)**

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	A	
II	Initial calibration	A	
III.	Calibration verification	A	
IV	Laboratory Blanks	A	
V	Field blanks	N	
VI.	Matrix Spike/Matrix Spike Duplicates	A	
VII.	Duplicate sample analysis	A	
VIII.	Laboratory control samples	A	LCS
IX.	Field duplicates	N	
X.	Target Analyte Quantitation	A	
XI	Overall assessment of data	A	

Note: A = Acceptable      ND = No compounds detected      D = Duplicate      SB=Source blank  
 N = Not provided/applicable      R = Rinsate      TB = Trip blank      OTHER:  
 SW = See worksheet      FB = Field blank      EB = Equipment blank

	Client ID	Lab ID	Matrix	Date
1	LDW21-SC525	21J0142-01	Sediment	07/15/21
2	LDW21-SS500	21J0142-02	Sediment	07/16/21
3	LDW21-SS501	21J0142-03	Sediment	07/16/21
4	LDW21-SS502	21J0142-04	Sediment	07/16/21
5	LDW21-IT582F	21J0142-05	Sediment	07/16/21
6	LDW21-IT579C	21J0142-06	Sediment	07/16/21
7	LDW21-IT597A	21J0142-07	Sediment	07/16/21
8	LDW21-IT597D	21J0142-08	Sediment	07/16/21
9	LDW21-SC673A	21J0142-09	Sediment	07/19/21
10	LDW21-IT600	21J0142-10	Sediment	07/19/21
11	LDW21-IT665D	21J0142-11	Sediment	07/19/21
12	LDW21-IT666D	21J0142-12	Sediment	07/19/21
13	LDW21-SS541	21J0142-13	Sediment	07/21/21
14	LDW21-IT512	21J0142-14	Sediment	07/19/21
15	LDW21-IT663D	21J0142-15	Sediment	07/19/21
16	LDW21-SC500	21J0142-16	Sediment	07/20/21
17	LDW21-SC501	21J0142-17	Sediment	07/20/21

**METHOD: (Analyte) TOC (EPA SW846 Method 9060A), Total Solids (SM2540G)**

	Client ID	Lab ID	Matrix	Date
18	LDW21-SC502	21J0142-18	Sediment	07/20/21
19	LDW21-SC563A	21J0142-19	Sediment	07/20/21
20	LDW21-SC628A	21J0142-20	Sediment	07/20/21
21	LDW21-IT664A	21J0142-21	Sediment	07/20/21
22	LDW21-IT670A	21J0142-22	Sediment	07/20/21
23	LDW21-IT621B	21J0142-23	Sediment	08/02/21
24	LDW21-IT665DMS	21J0142-11MS	Sediment	07/19/21
25	LDW21-IT665DDUP	21J0142-11DUP	Sediment	07/19/21
26	LDW21-SC563ADUP \	21J0142-19DUP \	Sediment	07/20/21
27	LDW21-SC563ATRP <i>over</i>	21J0142-19TRP <i>over</i>	Sediment	07/20/21
28				
29				
30				

Notes: \_\_\_\_\_

\_\_\_\_\_

\_\_\_\_\_

METHOD: Inorganics				
Validation Area	Yes	No	NA	Comments
<b>I. Technical holding times</b>				
Were all technical holding times were met?	X			Frozen
<b>II. Calibration</b>				
Were all instruments calibrated at the required frequency?	X			
Were the proper number of standards used?	X			
Were all initial and continuing calibration verifications within the QC limits?	X			
Were all initial calibration correlation coefficients within limits as specified by the method?	X			
Were balance checks performed as required?	X			
<b>III. Blanks</b>				
Was a method blank associated with every sample in this SDG?	X			
Was there contamination in the method blanks?		X		
Was there contamination in the initial and continuing calibration blanks?		X		
<b>IV. Matrix Spike/Matrix Spike Duplicates/Laboratory Duplicates</b>				
Were MS/MSD recoveries with the QC limits? (If the sample concentration exceeded the spike concentration by a factor of 4, no action was taken.)	X			
Were the MS/MSD or laboratory duplicate relative percent differences (RPDs) within the QC limits?	X			
<b>V. Laboratory Control Samples</b>				
Was a LCS analyzed for each batch in the SDG?	X			
Were the LCS recoveries and RPDs (if applicable) within QC limits?	X			
<b>X. Sample Result Verification</b>				
Were all reporting limits adjusted to reflect sample dilutions?	X			
Were all soil samples dry weight corrected?	X			
<b>XI. Overall Assessment of Data</b>				
Was the overall assessment of the data found to be acceptable?	X			

METHOD: Inorganics				
Validation Area	Yes	No	NA	Comments
<b>XII. Field Duplicates</b>				
Were field duplicates identified in this SDG?		X		
Were target analytes detected in the field duplicates?			X	
<b>XIII. Field Blanks</b>				
Were field blanks identified in this SDG?		X		
Were target analytes detected in the field blanks?			X	

All elements are applicable to each sample as noted below.

Sample ID	Target Analyte List
All	TS, TOC
QC:	
24, 25	TOC
26, 27	TS

**Validation Findings Worksheet**  
**Initial and Continuing Calibration Calculation Verification**

**Method:** Inorganics, Method See Cover

An initial or continuing calibration verification percent recovery (%R) was recalculated for each type of analysis using the following formula:

$$\%R = \frac{\text{Found} \times 100}{\text{True}}$$

Where,

Found = concentration of each analyte measured in the analysis of the ICV or CCV solution

True = concentration of each analyte in the ICV or CCV source

Calibration verification	TOC	ICV	44.446	47.154	106	106	Y
Calibration verification	TOC	CCV	44.446	45.03	101	101	Y
Calibration verification	TOC	CCV	44.446	46.741	105	105	Y

Comments:

VALIDATION FINDINGS CHECKLIST  
Quality Control Sample Recalculations

METHOD: Inorganics

Percent recoveries (%R) for the laboratory control sample (LCS) and matrix spike (MS) were recalculated using the following formula.

$$\%R = (\text{Found}/\text{True}) \times 100$$

Found = concentration of each analyte measured in the analysis. For the MS calculation, Found = SSR (Spiked Sample Result) - SR (Sample Result)

True = concentration of each analyte in the source

The sample and duplicate relative percent difference (RPD) was recalculated using the following formula.

$$\text{RPD} = (\text{Absolute value}(S-D) \times 200) / (S+D)$$

S = Original sample concentration

D = Duplicate sample concentration

Sample ID	Type of Analysis	Element	Found/S	True/D	Recalculated %R/RPD	Reported %R/RPD	Acceptable (Y/N)
LCS	LCS	TOC	44.8	44.4	101	101	Y
24	MS	TOC	1.03	1.04	99	99	Y
26	Duplicate	TS	64	63.25	1.18	1.18	Y





## Laboratory Data Consultants, Inc. Data Validation Report

**Project/Site Name:** Duwamish AOC4  
**LDC Report Date:** December 15, 2021  
**Parameters:** Polychlorinated Dioxins/Dibenzofurans  
**Validation Level:** Stage 4  
**Laboratory:** Analytical Resources, Inc., Tukwila, WA  
**Sample Delivery Group (SDG):** 21J0142

<b>Sample Identification</b>	<b>Laboratory Sample Identification</b>	<b>Matrix</b>	<b>Collection Date</b>
LDW21-IT665D	21J0142-11	Sediment	07/19/21
LDW21-IT663D	21J0142-15	Sediment	07/19/21
LDW21-IT664A	21J0142-21	Sediment	07/20/21
LDW21-IT621B	21J0142-23	Sediment	08/02/21

## 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 Final Lower Duwamish Waterway Quality Assurance Project Plan for Remedial Design of Upper Reach: Pre-Design Investigation (May 2020) and a modified outline of the USEPA National Functional Guidelines (NFG) for High Resolution Superfund Methods Data Review (April 2016). 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:

Polychlorinated Dioxins/Dibenzofurans by Environmental Protection Agency (EPA) Method 1613B

All sample results were subjected to Stage 4 data validation, which is comprised of the quality control (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): 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.
- U (Non-detected): The analyte was analyzed for and positively identified by the laboratory; however the analyte should be considered not detected at the reported concentration due to the presence of contaminants detected in the associated blank(s).
- UJ (Non-detected estimated): The analyte was reported as not detected by the laboratory; however the reported quantitation/detection limit is estimated due to non-conformances discovered during data validation.
- R (Rejected): The sample results were rejected due to gross non-conformances discovered during data validation. Data qualified as rejected is not usable.
- 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.

## **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. HRGC/HRMS Instrument Performance Check**

Instrument performance was checked at the required frequency.

Retention time windows were established for all homologues. The chromatographic resolution between 2,3,7,8-TCDD and peaks representing any other unlabeled TCDD isomer was less than or equal to 25%.

The static resolving power was at least 10,000 (10% valley definition).

## **III. Initial Calibration and Initial Calibration Verification**

A five point 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 and less than or equal to 35.0% for labeled compounds.

The ion abundance ratios for all PCDDs and PCDFs were within validation criteria.

The minimum S/N ratio was greater than or equal to 10 for each analyte and labeled compound.

The percent differences (%D) of the initial calibration verification (ICV) standard were within the QC limits for all analytes and labeled compounds.

## **IV. Continuing Calibration**

Continuing calibration was performed at the required frequencies.

All of the continuing calibration results were within the QC limits for all analytes and labeled compounds.

The ion abundance ratios for all PCDDs and PCDFs were within validation criteria.

The minimum S/N ratio was greater than or equal to 10 for each analyte and labeled compound.

## **V. Laboratory Blanks**

Laboratory blanks were analyzed as required by the method. No contaminants were found in the laboratory blanks with the following exceptions:

Blank ID	Extraction Date	Analyte	Concentration	Associated Samples
BJJ0500-BLK1	10/19/21	OCDD Total HxCDF	0.981 ng/Kg 0.100 ng/Kg	All samples in SDG 21J0142

Sample concentrations were compared to concentrations detected in the laboratory blanks. The sample concentrations were either not detected or were significantly greater than the concentrations found in the associated laboratory blanks with the following exceptions:

Sample	Analyte	Reported Concentration	Modified Final Concentration
LDW21-IT621B	Total HxCDF	0.184 ng/Kg	0.184J ng/Kg

## VI. Field Blanks

No field blanks were identified in this SDG.

## VII. Matrix Spike/Matrix Spike Duplicates

The laboratory has indicated that there were no matrix spike (MS) and matrix spike duplicate (MSD) analyses specified for the samples in this SDG, and therefore matrix spike and matrix spike duplicate analyses were not performed for this SDG.

## VIII. Laboratory Control Samples/Standard Reference Materials

Laboratory control samples (LCS) were analyzed as required by the method. Percent recoveries (%R) were within QC limits.

Standard reference materials (SRM) were analyzed as required by the method. The results were within QC limits.

## IX. Field Duplicates

No field duplicates were identified in this SDG.

## X. Internal Standards

All internal standard areas and retention times were within QC limits.

## XI. Target Analyte Quantitation

All target analyte quantitations met validation criteria with the following exceptions:

Sample	Analyte	Flag	A or P
All samples in SDG 21J0142	All analytes reported as estimated maximum possible concentration (EMPC) and less than the reporting limit (RL).	U	A
All samples in SDG 21J0142	All analytes flagged "X" due to chlorinated diphenyl ether (CDPE) interference.	J (all detects)	A

## XII. Target Analyte Identification

All target analyte identifications met validation criteria.

## XIII. System Performance

The system performance was acceptable.

## XIV. Overall Assessment of Data

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

Due to results reported by the laboratory as EMPCs and CDPE interference, data were qualified as estimated in four samples.

Due to laboratory blank contamination, data were qualified as not detected in one sample.

The quality control criteria reviewed, other than those discussed above, were met and are considered acceptable.

**Duwamish AOC4  
Polychlorinated Dioxins/Dibenzofurans - Data Qualification Summary - SDG 21J0142**

Sample	Analyte	Flag	A or P	Reason
LDW21-IT665D LDW21-IT663D LDW21-IT664A LDW21-IT621B	All analytes reported as estimated maximum possible concentration (EMPC) and less than the reporting limit (RL).	U	A	Target analyte quantitation (EMPC)
LDW21-IT665D LDW21-IT663D LDW21-IT664A LDW21-IT621B	All analytes flagged "X" due to chlorinated diphenyl ether (CDPE) interference.	J (all detects)	A	Target analyte quantitation (CDPE interference)

**Duwamish AOC4  
Polychlorinated Dioxins/Dibenzofurans - Laboratory Blank Data Qualification Summary - SDG 21J0142**

Sample	Analyte	Modified Final Concentration	A or P
LDW21-IT621B	Total HxCDF	0.184J ng/Kg	A

**Duwamish AOC4  
Polychlorinated Dioxins/Dibenzofurans - Field Blank Data Qualification Summary - SDG 21J0142**

No Sample Data Qualified in this SDG

LDC #: 52703D21

### VALIDATION COMPLETENESS WORKSHEET

Date: 12/9/21

SDG #: 21J0142

Stage 4

Page: 1 of 1

Laboratory: Analytical Resources, Inc., Tukwila, WA

Reviewer: [Signature]

2nd Reviewer: [Signature]

**METHOD:** HRGC/HRMS Polychlorinated Dioxins/Dibenzofurans (EPA Method 1613B)

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	A	
II.	HRGC/HRMS Instrument performance check	A	
III.	Initial calibration/ICV	A/A	RSD ≤ 20/35/70 CV ≤ 20 limits
IV.	Continuing calibration	A	CCV ≤ 20 limits
V.	Laboratory Blanks	N	
VI.	Field blanks	N	
VII.	Matrix spike/Matrix spike duplicates	N	CS
VIII.	Laboratory control samples / SW	A	LES
IX.	Field duplicates	N	
X.	Internal standards	A	
XI.	Target analyte quantitation	A	
XII.	Target analyte identification	A	
XIII.	System performance	A	
XIV.	Overall assessment of data	A	

Note: A = Acceptable      ND = No compounds detected      D = Duplicate      SB=Source blank  
 N = Not provided/applicable      R = Rinsate      TB = Trip blank      OTHER:  
 SW = See worksheet      FB = Field blank      EB = Equipment blank

	Client ID	Lab ID	Matrix	Date
1	LDW21-IT665D	21J0142-11	Sediment	07/19/21
2	LDW21-IT663D	21J0142-15	Sediment	07/19/21
3	LDW21-IT664A	21J0142-21	Sediment	07/20/21
4	LDW21-IT621B	21J0142-23	Sediment	08/02/21
5				
6				
7				
8				
9				
10				

Notes:

BL0500 - [Signature]				

**Method:** HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 1613B)

Validation Area	Yes	No	NA	Findings/Comments
<b>I. Technical holding times</b>				
All technical holding times were met.	√			
Cooler temperature criteria were met.	√			
<b>II. GC/MS Instrument performance check</b>				
Was PFK exact mass 380.9760 verified?	√			
Were the retention time windows established for all homologues?	√			
Was the chromatographic resolution between 2,3,7,8-TCDD and peaks representing any other unlabeled TCDD isomers $\leq 25\%$ ?	√			
Is the static resolving power at least 10,000 (10% valley definition)?	√			
Was the mass resolution adequately check with PFK?	√			
Was the presence of 1,2,8,9-TCDD and 1,3,4,6,8-PeCDF verified?	√			
<b>III. Initial calibration and Initial calibration verification</b>				
Was the initial calibration performed at 5 concentration levels?	√			
Were all percent relative standard deviations (%RSD) $\leq 20\%$ for unlabeled compounds and $\leq 35\%$ for unlabeled compounds?	√			
Did all calibration standards meet the Ion Abundance Ratio criteria?	√			
Was the signal to noise ratio for each target compound and labeled compound $\geq 10$ ?	√			
Was an initial calibration verification (ICV) standard analyzed after each initial calibration for each instrument?	√			
Were all ICV concentrations for the unlabeled and labeled compounds within QC limits?	√			
<b>IV. Continuing calibration</b>				
Was a continuing calibration performed at the beginning of each 12-hour period?	√			
Were all continuing calibration concentrations for the unlabeled and labeled compounds within QC limits?	√			
Did all continuing calibration standards meet the Ion Abundance Ratio criteria?	√			
<b>V. Blanks</b>				
Was a method blank associated with every sample in this SDG?	√			
Was a method blank performed for each matrix and whenever a sample extraction was performed?	√			
Was there contamination in the method blanks?	√	0		
<b>VI. Field blanks</b>				
Were field blanks identified in this SDG?		√		
Were target compounds detected in the field blanks?			√	
<b>VII. Matrix spike/Matrix spike duplicates</b>				
Were matrix spike (MS) and matrix spike duplicate (MSD) analyzed in this SDG?		√		
Were the MS/MSD percent recoveries (%R) and the relative percent differences (RPD) within the QC limits?			√	



Validation Area	Yes	No	NA	Findings/Comments
<b>VIII. Laboratory control samples</b>				
Was an LCS analyzed per extraction batch?	√			
Were the LCS percent recoveries (%R) and relative percent difference (RPD) within the QC limits?	√			
<b>IX. Field duplicates</b>				
Were field duplicate pairs identified in this SDG?		√		
Were target compounds detected in the field duplicates?			√	
<b>X. Labeled Compounds</b>				
Were labeled compounds within QC limits?	√	∅		
Was the minimum S/N ratio of all labeled compound peaks $\geq 10$ ?	√			
<b>XI. Compound quantitation</b>				
Did the laboratory LOQs/RLs meet the QAPP LOQs/RLs?	√			
Were the correct labeled compound, quantitation ion and relative response factor (RRF) used to quantitate the compound?	√			
Were compound quantitation and RLs adjusted to reflect all sample dilutions and dry weight factors applicable to level IV validation?	√			
<b>XII. Target compound identification</b>				
For 2,3,7,8 substituted congeners with associated labeled standards, were the retention times of the two quantitation peaks within -1 to 3 sec. of the RT of the labeled standard?	√			
For 2,3,7,8 substituted congeners without associated labeled standards, were the relative retention times of the two quantitation peaks within 0.005 time units of the RRT measured in the routine calibration?	√			
For non-2,3,7,8 substituted congeners, were the retention times of the two quantitation peaks within RT established in the performance check solution?	√			
Did selected ion current profile (SICP) contain all characteristic ions listed in Method 1613B, Table 8?	√			
Was the Ion Abundance Ratio for the two quantitation ions within criteria?		√		
Was the signal to noise ratio for each target compound $\geq 2.5$ and $\geq 10$ for the labeled compound?	√			
Does the maximum intensity of each specified characteristic ion coincide within $\pm 2$ seconds (includes labeled standards)?	√			
For PCDF identification, was any signal ( $S/N \geq 2.5$ , at $\pm$ seconds RT) detected in the corresponding PCDPE channel?			√	
Was an acceptable lock mass recorded and monitored?	√			
<b>XIII. System performance</b>				
System performance was found to be acceptable.	√			
<b>XIV. Overall assessment of data</b>				
Overall assessment of data was found to be acceptable.	√			

## VALIDATION FINDINGS WORKSHEET

**METHOD:** HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 1613B)

A. 2,3,7,8-TCDD	F. 1,2,3,4,6,7,8-HpCDD	K. 1,2,3,4,7,8-HxCDF	P. 1,2,3,4,7,8,9-HpCDF	U. Total HpCDD
B. 1,2,3,7,8-PeCDD	G. OCDD	L. 1,2,3,6,7,8-HxCDF	Q. OCDF	V. Total TCDF
C. 1,2,3,4,7,8-HxCDD	H. 2,3,7,8-TCDF	M. 2,3,4,6,7,8-HxCDF	R. Total TCDD	W. Total PeCDF
D. 1,2,3,6,7,8-HxCDD	I. 1,2,3,7,8-PeCDF	N. 1,2,3,7,8,9-HxCDF	S. Total PeCDD	X. Total HxCDF
E. 1,2,3,7,8,9-HxCDD	J. 2,3,4,7,8-PeCDF	O. 1,2,3,4,6,7,8-HpCDF	T. Total HxCDD	Y. Total HpCDF

Notes: \_\_\_\_\_  
 \_\_\_\_\_  
 \_\_\_\_\_

**VALIDATION FINDINGS WORKSHEET**  
**Blanks**

**METHOD:** HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 1613B)

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

N N/A Were all samples associated with a method blank?

N N/A Was a method blank performed for each matrix and whenever a sample extraction was performed?

N N/A Was the method blank contaminated?

Blank extraction date: 10/19/21

Blank analysis date: 10/25/21

Associated samples: All

Conc. units: ug/g

Compound	Blank ID	Sample Identification							
	<u>BL10500-B4-1</u>		<u>A</u>						
<u>Q</u>	<u>0.981</u>								
<u>X</u>	<u>0.100</u>		<u>0.184/J</u>						

Blank extraction date: \_\_\_\_\_ Blank analysis date: \_\_\_\_\_

Conc. units: \_\_\_\_\_ Associated Samples: \_\_\_\_\_

Compound	Blank ID	Sample Identification							

CIRCLED RESULTS WERE NOT QUALIFIED. ALL RESULTS NOT CIRCLED WERE QUALIFIED BY THE FOLLOWING STATEMENT:  
All contaminants within five times the method blank concentration were qualified as not detected, "U".

**VALIDATION FINDINGS WORKSHEET**  
**Compound Quantitation and Reported RLs**

**METHOD:** HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 1613B)

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

N N/A      Were the correct labeled compound, quantitation ions and relative response factors (RRF) used to quantitate the compound?  
 N N/A      Compound quantitation and RLs were adjusted to reflect all sample dilutions and dry weight factors (if necessary).

#	Date	Sample ID	Finding	Associated Samples	Qualifications
		All	All compounds reported as estimated maximum possible concentration (EMPC) < RL		U/A
		All	All compounds flagged "X" due to chlorinated diphenyl ether interference		Jdets/A

Comments: See sample calculation verification worksheet for recalculations

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### VALIDATION FINDINGS WORKSHEET Initial Calibration Calculation Verification

**METHOD:** HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 1613B)

The Relative Response Factor (RRF), average RRF, and percent relative standard deviation (%RSD) were recalculated for the compounds identified below using the following calculations:

$$RRF = (A_x)(C_{is}) / (A_{is})(C_x)$$

average RRF = sum of the RRFs/number of standards

$$\%RSD = 100 * (S/X)$$

A<sub>x</sub> = Area of compound,

C<sub>x</sub> = Concentration of compound,

S = Standard deviation of the RRFs,

A<sub>is</sub> = Area of associated internal standard

C<sub>is</sub> = Concentration of internal standard

X = Mean of the RRFs

#	Standard ID	Calibration Date	Compound (Reference Internal Standard)	Reported	Recalculated	Reported	Recalculated	Reported	Recalculated
				RRF (10/50 std)	RRF (10/50 std)	Average RRF (initial)	Average RRF (initial)	%RSD	%RSD
1	ICAL 01	8/11/21	2,3,7,8-TCDF ( <sup>13</sup> C-2,3,7,8-TCDF)	1.0832006	1.083746	1.107593	1.107593	3.6	3.6
			2,3,7,8-TCDD ( <sup>13</sup> C-2,3,7,8-TCDD)	0.9085186	0.908390	0.9202875	0.9202874	3.1	3.1
			1,2,3,6,7,8-HxCDD ( <sup>13</sup> C-1,2,3,6,7,8-HxCDD)	1.005616	1.005605	1.00898	1.00898	1.0	1.0
			1,2,3,4,6,7,8-HpCDD ( <sup>13</sup> C-1,2,4,6,7,8,-HpCDD)	1.051009	1.051062	1.068088	1.068088	6.6	6.6
			OCDF ( <sup>13</sup> C-OCDD)	1.440564	1.44059	1.44690	1.44690	5.7	5.7
2			2,3,7,8-TCDF ( <sup>13</sup> C-2,3,7,8-TCDF)						
			2,3,7,8-TCDD ( <sup>13</sup> C-2,3,7,8-TCDD)						
			1,2,3,6,7,8-HxCDD ( <sup>13</sup> C-1,2,3,6,7,8-HxCDD)						
			1,2,3,4,6,7,8-HpCDD ( <sup>13</sup> C-1,2,4,6,7,8,-HpCDD)						
			OCDF ( <sup>13</sup> C-OCDD)						
3			2,3,7,8-TCDF ( <sup>13</sup> C-2,3,7,8-TCDF)						
			2,3,7,8-TCDD ( <sup>13</sup> C-2,3,7,8-TCDD)						
			1,2,3,6,7,8-HxCDD ( <sup>13</sup> C-1,2,3,6,7,8-HxCDD)						
			1,2,3,4,6,7,8-HpCDD ( <sup>13</sup> C-1,2,4,6,7,8,-HpCDD)						
			OCDF ( <sup>13</sup> C-OCDD)						

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.

**VALIDATION FINDINGS WORKSHEET**  
**Continuing Calibration Results Verification**

**METHOD:** HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 1613B)

The percent difference (%D) of the initial calibration average Relative Response Factors (RRFs) and the continuing calibration RRFs were recalculated for the compounds identified below using the following calculation:

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

$$\text{RRF} = (A_x)(C_{is}) / (A_{is})(C_x)$$

Where: ave. RRF = initial calibration average RRF

RRF = continuing calibration RRF

 $A_x$  = Area of compound, $A_{is}$  = Area of associated internal standard $C_x$  = Concentration of compound, $C_{is}$  = Concentration of internal standard

#	Standard ID	Calibration Date	Compound (Reference Internal Standard)	Average RRF (initial)	Reported	Recalculated	Reported	Recalculated
					Conc (CC)	Conc (CC)	%D	%D
1	2110505A	10/5/01	2,3,7,8-TCDF ( <sup>13</sup> C-2,3,7,8-TCDF)	1.107593	1.0745550	1.074615	3.0	3.0
			2,3,7,8-TCDD ( <sup>13</sup> C-2,3,7,8-TCDD)	0.9202875	1.0081390	1.0081532	9.5	9.5
			1,2,3,6,7,8-HxCDD ( <sup>13</sup> C-1,2,3,6,7,8-HxCDD)	1.00898	1.0688370	1.0683744	5.9	5.9
			1,2,3,4,6,7,8-HpCDD ( <sup>13</sup> C-1,2,4,6,7,8,-HpCDD)	1.068088	1.1679010	1.1678182	9.3	9.3
			OCDF ( <sup>13</sup> C-OCDF)	1.44690	1.3382880	1.338548	7.5	7.5
2	2110518	10/6/01	2,3,7,8-TCDF ( <sup>13</sup> C-2,3,7,8-TCDF)	1.107593	1.0713550	1.0713484	3.3	3.3
			2,3,7,8-TCDD ( <sup>13</sup> C-2,3,7,8-TCDD)	0.9202875	1.0205990	1.0206664	10.9	10.9
			1,2,3,6,7,8-HxCDD ( <sup>13</sup> C-1,2,3,6,7,8-HxCDD)	1.00898	1.0288700	1.028884	2.0	2.0
			1,2,3,4,6,7,8-HpCDD ( <sup>13</sup> C-1,2,4,6,7,8,-HpCDD)	1.068088	1.1328670	1.1328398	6.1	6.1
			OCDF ( <sup>13</sup> C-OCDF)	1.44690	1.3304760	1.3304528	8.0	8.0
3			2,3,7,8-TCDF ( <sup>13</sup> C-2,3,7,8-TCDF)					
			2,3,7,8-TCDD ( <sup>13</sup> C-2,3,7,8-TCDD)					
			1,2,3,6,7,8-HxCDD ( <sup>13</sup> C-1,2,3,6,7,8-HxCDD)					
			1,2,3,4,6,7,8-HpCDD ( <sup>13</sup> C-1,2,4,6,7,8,-HpCDD)					
			OCDF ( <sup>13</sup> C-OCDF)					

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







## Laboratory Data Consultants, Inc. Data Validation Report

**Project/Site Name:** Duwamish AOC4  
**LDC Report Date:** December 15, 2021  
**Parameters:** Polychlorinated Biphenyls  
**Validation Level:** Stage 4  
**Laboratory:** Analytical Resources, Inc., Tukwila  
**Sample Delivery Group (SDG):** 21K0332

<b>Sample Identification</b>	<b>Laboratory Sample Identification</b>	<b>Matrix</b>	<b>Collection Date</b>
LDW21-IT637A	21K0332-01	Sediment	07/06/21
LDW21-IT637AMS	21K0332-01MS	Sediment	07/06/21
LDW21-IT637AMSD	21K0332-01MSD	Sediment	07/06/21

## 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 Final Lower Duwamish Waterway Quality Assurance Project Plan for Remedial Design of Upper Reach: Pre-Design Investigation (May 2020) and a modified outline of the USEPA National Functional Guidelines (NFG) for Organic Superfund Methods Data Review (January 2017). 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:

Polychlorinated Biphenyls (PCBs) by Environmental Protection Agency (EPA) SW 846 Method 8082A

All sample results were subjected to Stage 4 data validation, which is comprised of the quality control (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): 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.
- U (Non-detected): The analyte was analyzed for and positively identified by the laboratory; however the analyte should be considered non-detected at the reported concentration due to the presence of contaminants detected in the associated blank(s).
- UJ (Non-detected estimated): The analyte was reported as not detected by the laboratory; however the reported quantitation/detection limit is estimated due to non-conformances discovered during data validation.
- R (Rejected): The sample results were rejected due to gross non-conformances discovered during data validation. Data qualified as rejected is not usable.
- 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.

## **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.

Retention time windows were established as required by the method.

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 required frequencies.

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

Retention times of all analytes in the calibration standards were within the established retention time windows.

## **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/Internal Standards**

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

All internal standard percent recoveries (%R) were within QC limits.

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

Matrix spike (MS) and matrix spike duplicate (MSD) sample analysis was performed on an associated project sample. Percent recoveries (%R) were within QC limits. Relative percent differences (RPD) were within QC limits.

## **VIII. Laboratory Control Samples/Standard Reference Materials**

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.

Standard reference materials (SRM) were analyzed as required by the method. The results were within QC limits.

## **IX. Field Duplicates**

No field duplicates were identified in this SDG.

## **X. Target Analyte Quantitation**

All target analyte quantitations met validation criteria.

## **XI. Target Analyte Identification**

All target analyte identifications met validation criteria.

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

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

The quality control criteria reviewed were met and are considered acceptable.

**Duwamish AOC4  
Polychlorinated Biphenyls - Data Qualification Summary - SDG 21K0332**

No Sample Data Qualified in this SDG

**Duwamish AOC4  
Polychlorinated Biphenyls - Laboratory Blank Data Qualification Summary - SDG  
21K0332**

No Sample Data Qualified in this SDG

**Duwamish AOC4  
Polychlorinated Biphenyls - Field Blank Data Qualification Summary - SDG  
21K0332**

No Sample Data Qualified in this SDG

LDC #: 52703E3b

**VALIDATION COMPLETENESS WORKSHEET**

Date: 12/10/21

SDG #: 21K0332

Stage 4

Page: 1 of 1

Laboratory: Analytical Resources, Inc., Tukwila, WA

Reviewer: PG

2nd Reviewer: JLB

**METHOD:** GC Polychlorinated Biphenyls (EPA SW846 Method 8082A)

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	A		
II.	Initial calibration/ICV	A / A	RSD < 20 %	ICV < 20 %
III.	Continuing calibration	A	CCV < 20 %	
IV.	Laboratory Blanks	A		
V.	Field blanks	N		
VI.	Surrogate spikes / IS	A / A		
VII.	Matrix spike/Matrix spike duplicates	A		
VIII.	Laboratory control samples / SRM	A	LCS / LCSD	
IX.	Field duplicates	N		
X.	Target analyte quantitation	A		
XI.	Target analyte identification	A		
XII.	Overall assessment of data	A		

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:

	Client ID	Lab ID	Matrix	Date
1	LDW21-IT637A	21K0332-01	Sediment	07/06/21
2	LDW21-IT637AMS	21K0332-01MS	Sediment	07/06/21
3	LDW21-IT637AMSD	21K0332-01MSD	Sediment	07/06/21
4				
5				
6				
7				
8				
9				
10				
11				
12				
13				

Notes:


Method: GC HPLC

Validation Area	Yes	No	NA	Findings/Comments
<b>I. Technical holding times</b>				
Were all technical holding times met?	✓			
Was cooler temperature criteria met?	✓			
<b>IIa. Initial calibration</b>				
Did the laboratory perform a 5 point calibration prior to sample analysis?	✓			
Were all percent relative standard deviations (%RSD) < 20%?	✓			
Was a curve fit used for evaluation? If yes, did the initial calibration meet the curve fit acceptance criteria of $\geq 0.990$ ?		✓		
Were the RT windows properly established?	✓			
<b>IIb. Initial calibration verification</b>				
Was an initial calibration verification standard analyzed after each initial calibration for each instrument?	✓			
Were all percent differences (%D) < 20%?	✓			
<b>III. Continuing calibration</b>				
Was a continuing calibration analyzed daily?	✓			
Were all percent differences (%D) < 20%?	✓			
Were all the retention times within the acceptance windows?	✓			
<b>IV. Laboratory Blanks</b>				
Was a laboratory blank associated with every sample in this SDG?	✓			
Was a laboratory blank analyzed for each matrix and concentration?	✓			
Was there contamination in the laboratory blanks? If yes, please see the Blanks validation completeness worksheet.		✓		
<b>V. Field Blanks</b>				
Were field blanks identified in this SDG?		✓		
Were target compounds detected in the field blanks?			✓	
<b>VI. Surrogate spikes</b>				
Were all surrogate percent recovery (%R) within the QC limits?	✓			
If the percent recovery (%R) of one or more surrogates was outside QC limits, was a reanalysis performed to confirm %R?			✓	
If any %R was less than 10 percent, was a reanalysis performed to confirm %R?			✓	
<b>VII. Matrix spike/Matrix spike duplicates</b>				
Were a matrix spike (MS) and matrix spike duplicate (MSD) analyzed for each matrix in this SDG? If no, indicate which matrix does not have an associated MS/MSD. Soil / Water.	✓			
Was a MS/MSD analyzed every 20 samples of each matrix?	✓			
Were the MS/MSD percent recoveries (%R) and the relative percent differences (RPD) within the QC limits?	✓			

Validation Area	Yes	No	NA	Findings/Comments
<b>VIII. Laboratory control samples</b>				
Was an LCS analyzed for this SDG?	✓			
Was an LCS analyzed per extraction batch?	✓			
Were the LCS percent recoveries (%R) and relative percent difference (RPD) within the QC limits?	✓			
<b>IX. Field duplicates</b>				
Were field duplicate pairs identified in this SDG?		✓		
Were target compounds detected in the field duplicates?			✓	
<b>X. Compound quantitation</b>				
Were compound quantitation and RLs adjusted to reflect all sample dilutions and dry weight factors applicable to level IV validation?	✓			
<b>XI. Target compound identification</b>				
Were the retention times of reported detects within the RT windows?	✓			
<b>XIII. Overall assessment of data</b>				
Overall assessment of data was found to be acceptable.	✓			



LDC #: 570323b

**VALIDATION FINDINGS WORKSHEET**  
**Initial Calibration Calculation Verification**

Page: 1 of 1  
 Reviewer: 9

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 (100 std)	CF (100 std)	Ave CF (initial)	Ave CF (initial)	%RSD	%RSD
1	KAZ	11/26/21	BBT (1C)	0.05637804	0.056378	0.0553271	0.0553271	6.1	6.1
			BBT (2C)	0.05869047	0.0586904	0.05758081	0.05758081	3.1	3.1
2									
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 #: 5203230

### VALIDATION FINDINGS WORKSHEET Continuing Calibration Results Verification

Page: 1 of 1  
Reviewer: [Signature]

METHOD:  GC\_HPLC

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

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

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

#	Standard ID	Calibration Date	Compound	Average CF(Ical)/ CCV Conc.	Reported	Recalculated	Reported	Recalculated
					CF/ Conc. CCV	CF/ Conc. CCV	%D	%D
1	<del>ISA</del>	11/27/21	BB-1 (1c)	0.0553271	0.0528161	0.052816	4.4	4.5
	112703		BB-1 (2c)	0.0575808	0.0546505	0.0546504	5.2	5.1
2								
3								
4								

**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>DCB</u>	<u>IC</u>	<u>40.0</u>	<u>39.4</u>	<u>98.5</u>	<u>98.5</u>	
<u>TCMX</u>	<u>↓</u>	<u>↓</u>	<u>34.5</u>	<u>86.2</u>	<u>86.2</u>	

Sample ID: \_\_\_\_\_

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

Sample ID: \_\_\_\_\_

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

## VALIDATION FINDINGS WORKSHEET

### Matrix Spike/Matrix Spike Duplicates Results Verification

METHOD: GC HPLC

The percent recoveries (%R) and relative percent differences (RPD) of the matrix spike and matrix spike duplicate were recalculated for the compounds identified below using the following calculation:

$$\% \text{Recovery} = 100 * (\text{SSC} - \text{SC}) / \text{SA}$$

Where

SSC = Spiked sample concentration

SC = Sample concentration

SA = Spike added

MS = Matrix spike

MSD = Matrix spike duplicate

$$\text{RPD} = ((\text{SSCMS} - \text{SSCMSD}) * 2) / (\text{SSCMS} + \text{SSCMSD}) * 100$$

MS/MSD samples: 2/3

Compound	Spike Added		Sample Comp.	Spike Sample Concentration		Matrix spike		Matrix Spike Duplicate		MS/MSD	
	(MS)			(MS)		Percent Recovery		Percent Recovery		RPD	
	MS	MSD		---	MS	MSD	Reported	Recalc.	Reported	Recalc.	Reported
Gasoline (8015)											
Diesel (8015)											
Benzene (8021B)											
Methane (RSK-175)											
2,4-D (8151)											
Dinoseb (8151)											
Naphthalene (8310)											
Anthracene (8310)											
HMX (8330)											
2,4,6-Trinitrotoluene (8330)											
<u>PCB-160</u>	<u>101</u>	<u>101</u>	<u>5.0</u>	<u>73.8</u>	<u>75.1</u>	<u>68.1</u>	<u>68.1</u>	<u>69.4</u>	<u>69.4</u>	<u>1.75</u>	<u>1.75</u>

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

LDC #: 520323b

## VALIDATION FINDINGS WORKSHEET

### Laboratory Control Sample/Laboratory Control Sample Duplicate Results Verification

Page: 1 of 1  
 Reviewer: 9

**METHOD:** GC Pesticides (EPA SW 846 Method 8081A)

The percent recoveries (%R) and Relative Percent difference (RPD) of the laboratory control sample and laboratory control sample duplicate were recalculated for the compounds identified below using the following calculation:

% Recovery = 100 \* (SSC-SC)/SA

Where: SSC = Spiked sample concentration  
 SA = Spike added

SC = Concentration

RPD = |LCS - LCSD| \* 2 / (LCS + LCSD)

LCS = Laboratory control sample percent recovery

LCSD = Laboratory control sample duplicate percent recovery

LCS/LCSD samples: BK0603

Compound	Spike Added		Spiked Sample Concentration		LCS		LCSD		LCS/LCSD	
	LCS	LCSD	LCS	LCSD	Percent Recovery		Percent Recovery		RPD	
					Reported	Recalc.	Reported	Recalc.	Reported	Recalc.
gamma-BHC										
4,4'-DDT										
<del>BB</del>	101	NA	88.4	NA	87.7	87.5				

Comments: Refer to Laboratory Control Sample/Laboratory Control Sample Duplicate 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**  
**Sample Calculation Verification**

METHOD: GC HPLC

Y N N/A  
Y N N/A

Were all reported results recalculated and verified for all level IV samples?  
 Were all recalculated results for detected target compounds within 10% of the reported results?

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

Example:

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

Sample ID. 1 Compound Name PCB-1260-1  
 LCS, Methane

Concentration =  $\frac{(5818)(80.0)}{(335270)(0.0553271)} = 25.1$

Concentration(total) =  $\frac{(25.1+19.3+26.9+24.1+29.6)(2.5)(1)}{5 \times 15.93 \times 0.785} = 5.0 \text{ ug kg}$

#	Sample ID	Compound	Reported Concentrations (Ug/kg)	Recalculated Results Concentrations	Qualifications
	1	PCB - 1260	5.0		

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

**Laboratory Data Consultants, Inc.  
Data Validation Report**

**Project/Site Name:** Duwamish AOC4

**LDC Report Date:** December 15, 2021

**Parameters:** Wet Chemistry

**Validation Level:** Stage 4

**Laboratory:** Analytical Resources, Inc., Tukwila, WA

**Sample Delivery Group (SDG):** 21K0332

<b>Sample Identification</b>	<b>Laboratory Sample Identification</b>	<b>Matrix</b>	<b>Collection Date</b>
LDW21-IT637A	21K0332-01	Sediment	07/06/21
LDW21-IT637AMS	21K0332-01MS	Sediment	07/06/21
LDW21-IT637ADUP1	21K0332-01DUP1	Sediment	07/06/21
LDW21-IT637ADUP2	21K0332-01DUP2	Sediment	07/06/21

## 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 Final Lower Duwamish Waterway Quality Assurance Project Plan for Remedial Design of Upper Reach: Pre-Design Investigation (May 2020) and a modified outline of the USEPA National Functional Guidelines (NFG) for Inorganic Superfund Methods Data Review (January 2017). 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 methods:

Total Organic Carbon by Environmental Protection Agency (EPA) SW 846 Method 9060A

Total Solids by Standard Method 2540G

All sample results were subjected to Stage 4 data validation, which is comprised of the quality control (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): 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.
- U (Non-detected): The analyte was analyzed for and positively identified by the laboratory; however the analyte should be considered non-detected at the reported concentration due to the presence of contaminants detected in the associated blank(s).
- UJ (Non-detected estimated): The analyte was reported as not detected by the laboratory; however the reported quantitation/detection limit is estimated due to non-conformances discovered during data validation.
- R (Rejected): The sample results were rejected due to gross non-conformances discovered during data validation. Data qualified as rejected is not usable.
- 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.

## I. Sample Receipt and Technical Holding Times

All samples were received in good condition.

All technical holding time requirements were met.

## II. Initial Calibration

All criteria for the initial calibration of each method were met.

## III. Continuing Calibration

Continuing calibration frequency and analysis criteria were met for each method when applicable.

## IV. Laboratory Blanks

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

## V. Field Blanks

No field blanks were identified in this SDG.

## VI. Matrix Spike/Matrix Spike Duplicates

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

## VII. Duplicate Sample Analysis

Duplicate (DUP) sample analysis was performed on an associated project sample. Results were within QC limits with the following exceptions:

DUP ID (Associated Samples)	Analyte	RPD (Limits)	Difference (Limits)	Flag	A or P
LDW21-IT637ADUP1 (All samples in SDG 21K0332)	Total organic carbon	29.6 (≤20)	-	J (all detects)	A

## VIII. Laboratory Control Samples

Laboratory control samples (LCS) were analyzed as required by the methods. Percent recoveries (%R) were within QC limits.

## IX. Field Duplicates

No field duplicates were identified in this SDG.

## **X. Target Analyte Quantitation**

All target analyte quantitations were within validation criteria.

## **XI. Overall Assessment of Data**

The analysis was conducted within all specifications of the methods. No results were rejected in this SDG.

Due to DUP RPD, data were qualified as estimated in three samples.

The quality control criteria reviewed, other than those discussed above, were met and are considered acceptable.

**Duwamish AOC4**

**Wet Chemistry - Data Qualification Summary - SDG 21K0332**

<b>Sample</b>	<b>Analyte</b>	<b>Flag</b>	<b>A or P</b>	<b>Reason</b>
LDW21-IT637A LDW21-IT637ADUP1 LDW21-IT637ADUP2	Total organic carbon	J (all detects)	A	Duplicate sample analysis (RPD)

**Duwamish AOC4**

**Wet Chemistry - Laboratory Blank Data Qualification Summary - SDG 21K0332**

No Sample Data Qualified in this SDG

**Duwamish AOC4**

**Wet Chemistry - Field Blank Data Qualification Summary - SDG 21K0332**

No Sample Data Qualified in this SDG

**METHOD: (Analyte) TOC (EPA SW846 Method 9060A), Total Solids (SM2540G)**

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	A, A	
II	Initial calibration	A	
III.	Calibration verification	A	
IV	Laboratory Blanks	A	
V	Field blanks	N	
VI.	Matrix Spike/Matrix Spike Duplicates	A	
VII.	Duplicate sample analysis	SW	
VIII.	Laboratory control samples	A	LCS
IX.	Field duplicates	N	
X.	Target Analyte Quantitation	A	
XI	Overall assessment of data	A	

Note: A = Acceptable ND = No compounds detected D = Duplicate SB=Source blank  
 N = Not provided/applicable R = Rinsate TB = Trip blank OTHER:  
 SW = See worksheet FB = Field blank EB = Equipment blank

	Client ID	Lab ID	Matrix	Date
1	LDW21-IT637A	21K0332-01	Sediment	07/06/21
2	LDW21-IT637AMS	21K0332-01MS	Sediment	07/06/21
3	LDW21-IT637ADUP \	21K0332-01DUP	Sediment	07/06/21
4	LDW21-IT637ATRP <i>DUP2</i>	21K0332-01TRP	Sediment	07/06/21
5				
6				
7				
8				
9				
10				
11				
12				
13				
14				
15				
16				

Notes: \_\_\_\_\_

METHOD: Inorganics				
Validation Area	Yes	No	NA	Comments
<b>I. Technical holding times</b>				
Were all technical holding times were met?	X			Frozen
<b>II. Calibration</b>				
Were all instruments calibrated at the required frequency?	X			
Were the proper number of standards used?	X			
Were all initial and continuing calibration verifications within the QC limits?	X			
Were all initial calibration correlation coefficients within limits as specified by the method?	X			
Were balance checks performed as required?	X			
<b>III. Blanks</b>				
Was a method blank associated with every sample in this SDG?	X			
Was there contamination in the method blanks?		X		
Was there contamination in the initial and continuing calibration blanks?		X		
<b>IV. Matrix Spike/Matrix Spike Duplicates/Laboratory Duplicates</b>				
Were MS/MSD recoveries with the QC limits? (If the sample concentration exceeded the spike concentration by a factor of 4, no action was taken.)	X			
Were the MS/MSD or laboratory duplicate relative percent differences (RPDs) within the QC limits?		X		
<b>V. Laboratory Control Samples</b>				
Was a LCS analyzed for each batch in the SDG?	X			
Were the LCS recoveries and RPDs (if applicable) within QC limits?	X			
<b>X. Sample Result Verification</b>				
Were all reporting limits adjusted to reflect sample dilutions?	X			
Were all soil samples dry weight corrected?	X			
<b>XI. Overall Assessment of Data</b>				
Was the overall assessment of the data found to be acceptable?	X			

METHOD: Inorganics				
Validation Area	Yes	No	NA	Comments
<b>XII. Field Duplicates</b>				
Were field duplicates identified in this SDG?		X		
Were target analytes detected in the field duplicates?			X	
<b>XIII. Field Blanks</b>				
Were field blanks identified in this SDG?		X		
Were target analytes detected in the field blanks?			X	

All elements are applicable to each sample as noted below.

Sample ID	Target Analyte List
All	TS, TOC
QC:	
2	TOC
3	TOC, TS
4	TS





**Validation Findings Worksheet**  
**Initial and Continuing Calibration Calculation Verification**

**Method:** Inorganics, Method See Cover

An initial or continuing calibration verification percent recovery (%R) was recalculated for each type of analysis using the following formula:

$$\%R = \frac{\text{Found} \times 100}{\text{True}}$$

Where,

Found = concentration of each analyte measured in the analysis of the ICV or CCV solution

True = concentration of each analyte in the ICV or CCV source

Calibration verification	TOC	ICV	44.446	45.388	102	102	Y
Calibration verification	TOC	CCV	44.446	43.406	98	98	Y
Calibration verification							

Comments:

METHOD: Inorganics

Percent recoveries (%R) for the laboratory control sample (LCS) and matrix spike (MS) were recalculated using the following formula.

$$\%R = (\text{Found}/\text{True}) \times 100$$

Found = concentration of each analyte measured in the analysis. For the MS calculation, Found = SSR (Spiked Sample Result) - SR (Sample Result)

True = concentration of each analyte in the source

The sample and duplicate relative percent difference (RPD) was recalculated using the following formula.

$$\text{RPD} = (\text{Absolute value}(S-D) \times 200) / (S+D)$$

S = Original sample concentration

D = Duplicate sample concentration

Sample ID	Type of Analysis	Element	Found/S	True/D	Recalculated %R/RPD	Reported %R/RPD	Acceptable (Y/N)
LCS	LCS	TOC	45.5	44.4	102	102	Y
24	MS	TOC	1.03	1.04	99	99	Y
26	Duplicate	TS	64	63.25	1.18	1.18	Y

VALIDATION FINDINGS CHECKLIST  
Sample Calculation Verification

METHOD: Inorganics

Analytes were recalculated and verified using the following equation.

$$\text{Concentration} = (\text{Result from raw data} \times \text{Final volume} \times \text{Dilution factor}) / (\text{Percent solids (if applicable)} \times \text{Initial weight or volume})$$

Sample ID	Analyte	Raw Data (%)	Dry (g)	Sample Dry (g)	Tare (g)	Percent solids (%)	Reported Result (%)	Recalculated Result (mg/Kg)	Acceptable (Y/N)
1	TOC	0.818				76.76	1.07	1.07	Y
	TS		5.3841	6.7644	0.8261		76.76	76.76	Y