

# Data Validation Report

Project: Portland Harbor Pre-Remedial Design Investigation and Baseline Sampling

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Laboratory: Test America, West Sacramento, California

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Laboratory Group: 580-80167-2

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Analyses/Method: Clean Water Act - Dioxins and Furans / CWA1613B

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Validation Level: Stage 2A

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AECOM Project Number: 60566335.2.12

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Prepared by: Peter Fairbanks/AECOM Completed on: 11/07/2018

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Reviewed by: George Kisluk/AECOM File Name: 580-80167-2 DVR

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## SUMMARY

The samples listed below were collected by AECOM in Portland Harbor in Portland, OR on September 5,6, and 7, 2018.

Sample ID	Matrix/Sample Type
PDI-RB-AL-180905	Equipment Blank
PDI-RB-LL-180907	Equipment Blank
PDI-RB-SS-180905	Equipment Blank
PDI-RB-SS-180906	Equipment Blank
PDI-SC-S019-0TO2	Sediment
PDI-SC-S019-10TO12	Sediment
PDI-SC-S019-10TO12D	Field Duplicate of PDI-SC-S019-10TO12
PDI-SC-S019-12TO13.7	Sediment
PDI-SC-S019-13.7TO14.7	Sediment
PDI-SC-S019-2TO4	Sediment
PDI-SC-S019-4TO6	Sediment
PDI-SC-S019-6TO8	Sediment
PDI-SC-S019-8TO10	Sediment
PDI-SC-S112-0TO2	Sediment
PDI-SC-S112-2TO4	Sediment
PDI-SC-S112-4TO6	Sediment
PDI-SC-S113C-0TO1.1	Sediment
PDI-SC-S113C-1.1TO3.1	Sediment
PDI-SC-S113C-3.1TO5.6	Sediment
PDI-SC-S113C-5.6TO6.6	Sediment
PDI-SC-S121-0TO1.8	Sediment

Sample ID	Matrix/Sample Type
PDI-SC-S121-1.8TO3.4	Sediment
PDI-SC-S129-0TO2	Sediment
PDI-SC-S129-2TO4	Sediment
PDI-SC-S129-4TO5.3	Sediment
PDI-SC-S155-0TO2.1	Sediment
PDI-SC-S155-2.1TO4.2	Sediment
PDI-SC-S155-4.2TO5.3	Sediment
PDI-SC-S255-0TO2.1	Sediment
PDI-SC-S255-0TO2.1D	Field Duplicate of PDI-SC-S255-0TO2.1
PDI-SC-S255-2.1TO4.3	Sediment
PDI-SC-S260-0TO1.3	Sediment
PDI-SC-S260-1.3TO2.6	Sediment
PDI-SC-S260-2.6TO4.2	Sediment
PDI-SC-S260-4.2TO6	Sediment
PDI-SC-S260-6TO7	Sediment

Data validation activities were conducted with reference to:

- EPA Method 1613B: *Tetra- through Octa-Chlorinated Dioxins and Furans by Isotope Dilution HRGC/HRMS (October 1994)*,
- USEPA Contract Laboratory Program National Functional Guidelines for High Resolution Superfund Methods Data Review (April 2016),
- Quality Assurance Project Plan, Portland Harbor Pre-Remedial Design Investigation and Baseline Sampling, Portland Harbor Superfund Site (March 2018), and the
- laboratory quality control (QC) limits.

The National Functional Guidelines were modified to accommodate the non-CLP methodologies. In the absence of method-specific information, laboratory QC limits, project-specific requirements and/or AECOM professional judgment were used as appropriate.

## REVIEW ELEMENTS

The data were evaluated based on the following parameters (where applicable to the method):

- ✓ Data completeness [chain-of-custody (COC)/sample integrity]
- ✓ Holding times and sample preservation
- ✗ Laboratory blanks/equipment blanks
- NA Matrix spike (MS) and/or matrix spike duplicate (MSD) results
- ✓ Ongoing precision and recovery (OPR) results
- ✓ Field duplicate results
- ✓ Labeled compound and clean-up standard recoveries
- ✗ Sample results/reporting issues

The symbol (✓) indicates that no validation qualifiers were applied based on this parameter. An NA indicates that the parameter was not included as part of this data set or was not applicable to this validation and therefore not reviewed. The symbol (X) indicates that a QC nonconformance resulted in the qualification of data. Any QC nonconformance that resulted in the qualification of data is discussed below. In addition, nonconformances or other issues that were noted during validation, but did not result in qualification of data, may be discussed for informational purposes only.

The data appear valid as qualified and may be used for decision making purposes. Select data points were qualified as estimated due to nonconformances of certain QC criteria (see discussion below). Qualified sample results are presented in Table 1.

## RESULTS

### Data Completeness (COC)/Sample Integrity

The data package was reviewed and found to meet acceptance criteria for completeness:

- The COCs were reviewed for completeness of information relevant to the samples and requested analyses, and for signatures indicating transfer of sample custody;
- The laboratory sample login sheet(s) were reviewed for issues potentially affecting sample integrity, including the condition of sample containers upon receipt at the laboratory;
- Completeness of analyses was verified by comparing the reported results to the COC requests.

### Holding Times and Sample Preservation

Sample preservation and preparation/analysis holding times were reviewed for conformance with method criteria. All method QC acceptance criteria were met.

### Laboratory Blanks/Equipment Blanks

Laboratory method blanks and equipment blank results are evaluated as to whether there are contaminants detected above the estimated detection limit (EDL).

Target compounds were detected in the method blanks and the equipment blank associated with the samples in this data set. The equipment blank contamination, after laboratory method blank actions were applied, is summarized below for informational purposes only. Equipment blank contamination was not used to qualify field samples.

Blank ID	Compound	Result	EDL	Units
PDI-RB-LL-180907	1,2,3,6,7,8-HxCDF	0.78	0.12	pg/L
PDI-RB-LL-180907	OCDD	19	0.21	pg/L
PDI-RB-LL-180907	OCDF	4.8	0.16	pg/L
PDI-RB-SS-180905	2,3,7,8-TCDF	2.0	0.095	pg/L
PDI-RB-SS-180905	OCDD	40	0.53	pg/L
PDI-RB-SS-180905	OCDF	5.1	0.40	pg/L
PDI-RB-SS-180906	1,2,3,4,7,8-HxCDF	0.88	0.19	pg/L
PDI-RB-SS-180906	1,2,3,6,7,8-HxCDD	1.0	0.13	pg/L
PDI-RB-SS-180906	1,2,3,6,7,8-HxCDF	1.0	0.19	pg/L
PDI-RB-SS-180906	OCDF	5.3	0.21	pg/L

The NFG guidance stipulates that a conservative approach should be taken with regards to qualification of PCDD/PCDFs due to the toxicity of these compounds and the reporting of false negative results should be avoided.

Therefore, in order to avoid the reporting of false negative results, professional judgment was used to qualify the data in the following manner. As allowed in the NFG, a blank action limit (BAL) was determined as 5 times the blank result:

- When the sample results were  $<$  the blank result, the sample result was qualified as nondetect (U) at the sample result.
- When the sample result was  $\geq$  the blank result and  $\leq$  the BAL, the sample result was qualified as estimated and potentially biased high (J+).
- When the sample result was  $>$  the BAL, sample result was not qualified.

Qualified sample results are summarized in Table 1.

### **MS/MSD Results**

MS/MSD analyses were not performed on a sample in this data set. No data validation actions were taken on this basis.

### **OPR Results**

The OPR %Rs and/or RPDs were reviewed for conformance with the method QC acceptance criteria

### **Field Duplicate Results**

Field duplicate RPDs were reviewed for conformance with the AECOM QC acceptance criteria of  $\leq 50\%$  [if one or both results were greater than five times the quantitation limit (QL)] for solid matrices and  $\leq 30\%$  [if one or both results were greater than five times the QL] for aqueous matrices. All method QC acceptance criteria were met.

### **Labeled Compound and Clean-up Standard Recoveries**

The labeled compounds and labeled clean-up standard %Rs were reviewed for conformance with the QC acceptance criteria. All method QC acceptance criteria were met.

### **Sample Results/Reporting Issues**

All sample results detected at concentrations less than the lowest calibration standard but greater than the EDL are qualified by the laboratory as estimated (J). This "J" qualifier is retained during data validation.

### **Quantitation**

Sample results detected at concentrations greater than the highest calibration standard, qualified by the laboratory with an "E", require secondary dilutions in order to bring the concentrations down within the linear range of calibration, per Method 1613B. This was not done by the laboratory.

It should be noted that according to Section 12.1.7 of the laboratory's SOP, unless the affected peak saturates the instrument detector, secondary dilutions are not performed. Furthermore, *"Historic data indicates that for the isotope dilution method, dilution and re-injection will not produce significantly different results from those reported with the "E" qualifier."*

Despite the laboratory's SOP, NFG guidance stipulates that if a sample is not properly diluted to bring the results within the linear range of calibration, then the results are qualified "J".

Qualified sample results are summarized in Table 1.

#### Compound Identification

The data were reviewed to ensure that:

- the retention time, relative retention time, ion abundance ratios, SIM ion co-maximization, and S/N method acceptance criteria were met for compound identification; and
- the quantitative determination of PCDFs were not affected by the presence of polychlorinated diphenyl ether (PCDPE) interferences detected above the 2.5:1 S/N ratio limit.

All QC acceptance criteria were met with the following exceptions. Sample results which don't meet all of the method stipulated qualitative identification criteria are considered to be Estimated Maximum Possible Concentrations (EMPCs). Details concerning sample results in this data set which did not meet these identification criteria are noted below along with any data qualifications, as applicable.

The laboratory qualified all EMPC sample results with a "q" laboratory qualifier to indicate that the ion ratio criterion was not met. All ion ratios were verified and affected sample results which did not meet the ion ratio criteria were qualified as estimated and tentatively identified (JN). Qualified sample results are shown in Table 1.

It should be noted that the "JN" qualifier was retained rather than replacement with the conventional overall "J" qualifier in instances where sample results were qualified for multiple quality control nonconformances.

#### Compound Quantitation/ Chromatographic Interference

##### Quantitation

Sample results detected at concentrations greater than the highest calibration standard, qualified by the laboratory with an "E", require secondary dilutions in order to bring the concentrations down within the linear range of calibration, per Method 1613B. This was not done by the laboratory.

It should be noted that according to Section 12.1.7 of the laboratory's SOP, unless the affected peak saturates the instrument detector, secondary dilutions are not performed. Furthermore, *"Historic data indicates that for the isotope dilution method, dilution and re-injection will not produce significantly different results from those reported with the "E" qualifier."*

Despite the laboratory's SOP, NFG guidance stipulates that if a sample is not properly diluted to bring the results within the linear range of calibration, then the results are qualified "J".

Qualified sample results are summarized in Table 1.

### Chromatographic Interference

Samples exhibiting elevated background noise or matrix interference resulting in elevation of EDLs are qualified with a “G” by the laboratory. Professional judgement was used to qualify affected results “J”.

Qualified sample results are summarized in Table 1.

### Second Column Confirmation (2,3,7,8-TCDF)

The sample data were reviewed to ensure that results for 2,3,7,8-TCDF when analyzed on a DB-5 (or equivalent) column were confirmed on a second column ( i.e., DB-225 or equivalent) when isomer specificity is not achieved. All sample results requiring confirmation were confirmed and results were reported from the confirmation column.

It should be noted that according to Section 11.3.5 of the laboratory's SOP, *“Any sample which 2,3,7,8-TCDF is identified above the lower calibration limit must be confirmed on a DB-225 column, SP-2331, or equivalent GC column.”* This suggests that 2,3,7,8-TCDF results detected below the lower calibration limit (i.e., “J” values) are not confirmed on a secondary column by the laboratory. Professional judgment was used to take no action in instances where 2,3,7,8-TCDF was detected as “J” values on the primary column (i.e., DB-5).

### Percent Solids Content

The percent solids data were reviewed since the amount of moisture in a solid sample may have an impact on data representativeness. Due to the extremely low solubility of dioxins and furans in water, these analytes should be contained in the solid phase. Consequently, the NFG guidance does not stipulate a percent solids criterion. If applicable, EPA Regional guidance is used when assessing percent solids content. In the absence of EPA Regional guidance, AECOM uses 30% solids (from the NFG semivolatile guidance) as a benchmark to evaluate the percent solids content and professional judgment is used to determine the necessity to qualify data. Qualification on this basis was not required.

Verification of calculations was performed on a subset of the data as deemed appropriate. No discrepancies were noted.

### Dilutions

Sample ID	Compound	Dilution Factor
PDI-SC-S113C-0TO1.1	All results, except 2,3,7,8-TCDF	20

## **QUALIFICATION ACTIONS**

Sample results qualified as a result of validation actions are summarized in Table 1. All actions are described above.

**ATTACHMENTS**

Attachment A: Qualifier Codes and Explanations

Attachment B: Reason Codes and Explanations

Table 1 - Data Validation Summary of Qualified Data

Sample ID	Matrix	Compound	Result	RDL	EDL	Units	Validation Qualifiers	Validation Reason
PDI-RB-AL-180905	WQ	1,2,3,4,6,7,8-HpCDD		1.0	0.11	pg/L	U	bl
PDI-RB-AL-180905	WQ	1,2,3,4,6,7,8-HpCDF		0.70	0.15	pg/L	U	bl
PDI-RB-AL-180905	WQ	1,2,3,4,7,8,9-HpCDF		1.6	0.19	pg/L	U	bl
PDI-RB-AL-180905	WQ	1,2,3,4,7,8-HxCDD		1.3	0.11	pg/L	U	bl
PDI-RB-AL-180905	WQ	1,2,3,4,7,8-HxCDF		0.48	0.16	pg/L	U	bl
PDI-RB-AL-180905	WQ	1,2,3,6,7,8-HxCDD		0.27	0.11	pg/L	U	bl
PDI-RB-AL-180905	WQ	1,2,3,6,7,8-HxCDF		0.49	0.16	pg/L	U	bl
PDI-RB-AL-180905	WQ	1,2,3,7,8,9-HxCDF		5.1	0.12	pg/L	U	bl
PDI-RB-AL-180905	WQ	1,2,3,7,8-PeCDF		1.1	0.15	pg/L	U	bl
PDI-RB-AL-180905	WQ	2,3,4,6,7,8-HxCDF		0.20	0.12	pg/L	U	bl
PDI-RB-AL-180905	WQ	2,3,7,8-TCDF		1.1	0.081	pg/L	U	bl
PDI-RB-AL-180905	WQ	OCDD		11	0.20	pg/L	U	bl
PDI-RB-AL-180905	WQ	OCDF		2.1	0.19	pg/L	U	bl
PDI-RB-LL-180907	WQ	1,2,3,4,6,7,8-HpCDD		1.9	0.10	pg/L	U	bl
PDI-RB-LL-180907	WQ	1,2,3,4,6,7,8-HpCDF		1.8	0.13	pg/L	U	bl
PDI-RB-LL-180907	WQ	1,2,3,4,7,8,9-HpCDF		1.9	0.17	pg/L	U	bl
PDI-RB-LL-180907	WQ	1,2,3,4,7,8-HxCDD		1.4	0.10	pg/L	U	bl
PDI-RB-LL-180907	WQ	1,2,3,4,7,8-HxCDF		0.81	0.13	pg/L	U	bl
PDI-RB-LL-180907	WQ	1,2,3,6,7,8-HxCDD		0.62	0.098	pg/L	U	bl
PDI-RB-LL-180907	WQ	1,2,3,6,7,8-HxCDF	0.78	0.12	0.12	pg/L	J+	bl
PDI-RB-LL-180907	WQ	1,2,3,7,8,9-HxCDD		0.70	0.092	pg/L	U	bl
PDI-RB-LL-180907	WQ	1,2,3,7,8,9-HxCDF		4.4	0.094	pg/L	U	bl
PDI-RB-LL-180907	WQ	1,2,3,7,8-PeCDD		0.30	0.13	pg/L	U	bl
PDI-RB-LL-180907	WQ	1,2,3,7,8-PeCDF		1.3	0.14	pg/L	U	bl
PDI-RB-LL-180907	WQ	2,3,4,6,7,8-HxCDF		0.50	0.10	pg/L	U	bl
PDI-RB-LL-180907	WQ	2,3,4,7,8-PeCDF		0.68	0.15	pg/L	U	bl
PDI-RB-LL-180907	WQ	2,3,7,8-TCDF		1.2	0.073	pg/L	U	bl
PDI-RB-LL-180907	WQ	OCDD	19	0.21	0.21	pg/L	J+	bl
PDI-RB-LL-180907	WQ	OCDF	4.8	0.16	0.16	pg/L	J+	bl
PDI-RB-SS-180905	WQ	1,2,3,4,6,7,8-HpCDD		1.9	0.36	pg/L	U	bl
PDI-RB-SS-180905	WQ	1,2,3,4,6,7,8-HpCDF		0.87	0.26	pg/L	U	bl
PDI-RB-SS-180905	WQ	1,2,3,4,7,8,9-HpCDF		2.1	0.34	pg/L	U	bl
PDI-RB-SS-180905	WQ	1,2,3,4,7,8-HxCDD		1.3	0.18	pg/L	U	bl
PDI-RB-SS-180905	WQ	1,2,3,4,7,8-HxCDF		0.34	0.23	pg/L	U	bl
PDI-RB-SS-180905	WQ	1,2,3,6,7,8-HxCDF		0.30	0.22	pg/L	U	bl
PDI-RB-SS-180905	WQ	1,2,3,7,8,9-HxCDD		0.42	0.16	pg/L	U	bl
PDI-RB-SS-180905	WQ	1,2,3,7,8,9-HxCDF		5.7	0.16	pg/L	U	bl
PDI-RB-SS-180905	WQ	1,2,3,7,8-PeCDF		1.1	0.16	pg/L	U	bl



Sample ID	Matrix	Compound	Result	RDL	EDL	Units	Validation Qualifiers	Validation Reason
PDI-RB-SS-180905	WQ	2,3,4,7,8-PeCDF		0.58	0.18	pg/L	U	bl
PDI-RB-SS-180905	WQ	2,3,7,8-TCDF	2.0	0.095	0.095	pg/L	J+	bl
PDI-RB-SS-180905	WQ	OCDD	40	0.53	0.53	pg/L	J+	bl
PDI-RB-SS-180905	WQ	OCDF	5.1	0.40	0.40	pg/L	JN	bl,k
PDI-RB-SS-180906	WQ	1,2,3,4,6,7,8-HpCDD		1.7	0.12	pg/L	U	bl
PDI-RB-SS-180906	WQ	1,2,3,4,6,7,8-HpCDF		1.5	0.15	pg/L	U	bl
PDI-RB-SS-180906	WQ	1,2,3,4,7,8,9-HpCDF		2.6	0.20	pg/L	U	bl
PDI-RB-SS-180906	WQ	1,2,3,4,7,8-HxCDD		1.5	0.14	pg/L	U	bl
PDI-RB-SS-180906	WQ	1,2,3,4,7,8-HxCDF	0.88	0.19	0.19	pg/L	J+	bl
PDI-RB-SS-180906	WQ	1,2,3,6,7,8-HxCDD	1.0	0.13	0.13	pg/L	J+	bl
PDI-RB-SS-180906	WQ	1,2,3,6,7,8-HxCDF	1.0	0.19	0.19	pg/L	J+	bl
PDI-RB-SS-180906	WQ	1,2,3,7,8,9-HxCDD		0.93	0.12	pg/L	U	bl
PDI-RB-SS-180906	WQ	1,2,3,7,8,9-HxCDF		6.0	0.14	pg/L	U	bl
PDI-RB-SS-180906	WQ	1,2,3,7,8-PeCDF		1.5	0.15	pg/L	U	bl
PDI-RB-SS-180906	WQ	2,3,4,6,7,8-HxCDF		0.60	0.15	pg/L	U	bl
PDI-RB-SS-180906	WQ	2,3,4,7,8-PeCDF		0.79	0.17	pg/L	U	bl
PDI-RB-SS-180906	WQ	2,3,7,8-TCDF		1.6	0.080	pg/L	U	bl
PDI-RB-SS-180906	WQ	OCDD		14	0.29	pg/L	U	bl
PDI-RB-SS-180906	WQ	OCDF	5.3	0.21	0.21	pg/L	J+	bl
PDI-SC-S019-0TO2	SE	1,2,3,7,8,9-HxCDF		0.0017	0.00017	ug/kg	U	bl
PDI-SC-S019-10TO12	SE	1,2,3,7,8,9-HxCDF		0.0016	0.00031	ug/kg	U	bl
PDI-SC-S019-10TO12	SE	2,3,7,8-TCDD	0.00048	0.000063	0.000063	ug/kg	JN	k
PDI-SC-S019-10TO12D	SE	1,2,3,7,8,9-HxCDF		0.0012	0.00019	ug/kg	U	bl
PDI-SC-S019-12TO13.7	SE	1,2,3,4,7,8-HxCDD	0.00058	0.00011	0.00011	ug/kg	J+	bl
PDI-SC-S019-12TO13.7	SE	1,2,3,7,8,9-HxCDF		0.0013	0.00033	ug/kg	U	bl
PDI-SC-S019-12TO13.7	SE	1,2,3,7,8-PeCDF	0.0014	0.00036	0.00036	ug/kg	J+	bl
PDI-SC-S019-12TO13.7	SE	2,3,7,8-TCDD	0.00031	0.000053	0.000053	ug/kg	JN	k
PDI-SC-S019-13.7TO14.7	SE	1,2,3,7,8,9-HxCDF		0.00094	0.00032	ug/kg	U	bl
PDI-SC-S019-2TO4	SE	1,2,3,7,8,9-HxCDF	0.0025	0.00039	0.00039	ug/kg	J+	bl
PDI-SC-S019-2TO4	SE	OCDD	6.4	0.0014	0.0014	ug/kg	J	q
PDI-SC-S019-4TO6	SE	1,2,3,7,8,9-HxCDF		0.0017	0.00051	ug/kg	U	bl
PDI-SC-S019-4TO6	SE	OCDD	9.0	0.0018	0.0018	ug/kg	J	q
PDI-SC-S019-6TO8	SE	1,2,3,7,8,9-HxCDF	0.0018	0.00039	0.00039	ug/kg	J+	bl
PDI-SC-S019-6TO8	SE	OCDD	5.5	0.0010	0.0010	ug/kg	J	q
PDI-SC-S019-8TO10	SE	1,2,3,7,8,9-HxCDF		0.0017	0.00056	ug/kg	U	bl
PDI-SC-S112-0TO2	SE	1,2,3,7,8,9-HxCDF	0.0033	0.00045	0.00045	ug/kg	J+	bl
PDI-SC-S112-0TO2	SE	OCDD	7.5	0.0032	0.0032	ug/kg	J	q
PDI-SC-S112-2TO4	SE	1,2,3,7,8,9-HxCDF	0.0029	0.00088	0.00088	ug/kg	J+	bl
PDI-SC-S112-2TO4	SE	OCDD	9.1	0.0038	0.0038	ug/kg	J	q
PDI-SC-S112-4TO6	SE	1,2,3,7,8,9-HxCDF	0.0030	0.0011	0.0011	ug/kg	J+	bl

Sample ID	Matrix	Compound	Result	RDL	EDL	Units	Validation Qualifiers	Validation Reason
PDI-SC-S112-4TO6	SE	OCDD	8.9	0.0025	0.0025	ug/kg	J	q
PDI-SC-S113C-0TO1.1	SE	1,2,3,7,8,9-HxCDF	0.0044	0.0011	0.0011	ug/kg	J+	bl
PDI-SC-S113C-1.1TO3.1	SE	1,2,3,4,7,8,9-HpCDF	0.0012	0.00015	0.00015	ug/kg	J+	bl
PDI-SC-S113C-1.1TO3.1	SE	1,2,3,4,7,8-HxCDD	0.00031	0.00026	0.00026	ug/kg	JN	k
PDI-SC-S113C-1.1TO3.1	SE	1,2,3,7,8,9-HxCDF	0.0014	0.00019	0.00019	ug/kg	J+	bl
PDI-SC-S113C-3.1TO5.6	SE	1,2,3,4,6,7,8-HpCDF	0.00050	0.000071	0.000071	ug/kg	J+	bl
PDI-SC-S113C-3.1TO5.6	SE	1,2,3,4,7,8,9-HpCDF	0.00042	0.000075	0.000075	ug/kg	J+	bl
PDI-SC-S113C-3.1TO5.6	SE	1,2,3,4,7,8-HxCDD	0.00012	0.000031	0.000031	ug/kg	J+	bl
PDI-SC-S113C-3.1TO5.6	SE	1,2,3,6,7,8-HxCDD	0.00014	0.000031	0.000031	ug/kg	JN	bl,k
PDI-SC-S113C-3.1TO5.6	SE	1,2,3,7,8,9-HxCDD	0.00016	0.000028	0.000028	ug/kg	J+	bl
PDI-SC-S113C-3.1TO5.6	SE	1,2,3,7,8,9-HxCDF	0.0013	0.000038	0.000038	ug/kg	J+	bl
PDI-SC-S113C-3.1TO5.6	SE	1,2,3,7,8-PeCDF	0.00037	0.000039	0.000039	ug/kg	JN	bl,k
PDI-SC-S113C-3.1TO5.6	SE	2,3,4,7,8-PeCDF	0.00014	0.000039	0.000039	ug/kg	JN	bl,k
PDI-SC-S113C-3.1TO5.6	SE	2,3,7,8-TCDF	0.00022	0.000019	0.000019	ug/kg	J+	bl
PDI-SC-S113C-5.6TO6.6	SE	1,2,3,4,6,7,8-HpCDF	0.00037	0.000022	0.000022	ug/kg	J+	bl
PDI-SC-S113C-5.6TO6.6	SE	1,2,3,4,7,8,9-HpCDF	0.00045	0.000023	0.000023	ug/kg	J+	bl
PDI-SC-S113C-5.6TO6.6	SE	1,2,3,4,7,8-HxCDD	0.00019	0.000016	0.000016	ug/kg	JN	bl,k
PDI-SC-S113C-5.6TO6.6	SE	1,2,3,6,7,8-HxCDD	0.00020	0.000016	0.000016	ug/kg	JN	k
PDI-SC-S113C-5.6TO6.6	SE	1,2,3,7,8,9-HxCDD	0.00031	0.000014	0.000014	ug/kg	J+	bl
PDI-SC-S113C-5.6TO6.6	SE	1,2,3,7,8,9-HxCDF		0.0012	0.000016	ug/kg	U	bl
PDI-SC-S113C-5.6TO6.6	SE	1,2,3,7,8-PeCDF	0.00036	0.000016	0.000016	ug/kg	J+	bl
PDI-SC-S113C-5.6TO6.6	SE	2,3,4,7,8-PeCDF	0.000071	0.000016	0.000016	ug/kg	JN	bl,k
PDI-SC-S113C-5.6TO6.6	SE	2,3,7,8-TCDF		0.00011	0.0000058	ug/kg	U	bl
PDI-SC-S113C-5.6TO6.6	SE	OCDF	0.0013	0.000024	0.000024	ug/kg	J+	bl
PDI-SC-S121-0TO1.8	SE	1,2,3,4,7,8,9-HpCDF	0.00036	0.00018	0.00018	ug/kg	J+	bl
PDI-SC-S121-0TO1.8	SE	1,2,3,4,7,8-HxCDD	0.00039	0.000040	0.000040	ug/kg	J+	bl
PDI-SC-S121-0TO1.8	SE	1,2,3,7,8,9-HxCDF		0.0011	0.000038	ug/kg	U	bl
PDI-SC-S121-0TO1.8	SE	1,2,3,7,8-PeCDF	0.00040	0.000029	0.000029	ug/kg	J+	bl
PDI-SC-S121-0TO1.8	SE	2,3,4,7,8-PeCDF	0.00019	0.000031	0.000031	ug/kg	J+	bl
PDI-SC-S121-0TO1.8	SE	2,3,7,8-TCDF	0.00035	0.000014	0.000014	ug/kg	J+	bl
PDI-SC-S121-1.8TO3.4	SE	1,2,3,4,7,8,9-HpCDF	0.0015	0.00044	0.00044	ug/kg	J+	bl
PDI-SC-S121-1.8TO3.4	SE	1,2,3,7,8,9-HxCDF	0.0014	0.00011	0.00011	ug/kg	J+	bl
PDI-SC-S121-1.8TO3.4	SE	OCDD	3.1	0.0011	0.0011	ug/kg	J	q
PDI-SC-S129-0TO2	SE	1,2,3,7,8,9-HxCDF	0.0026	0.00011	0.00011	ug/kg	J+	bl
PDI-SC-S129-0TO2	SE	1,2,3,7,8-PeCDF	0.0013	0.00012	0.00012	ug/kg	J+	bl
PDI-SC-S129-0TO2	SE	2,3,7,8-TCDF	0.00086	0.00064	0.00064	ug/kg	JN	k
PDI-SC-S129-2TO4	SE	1,2,3,4,6,7,8-HpCDD	0.72	0.0063	0.0063	ug/kg	J	d
PDI-SC-S129-2TO4	SE	1,2,3,7,8,9-HxCDF	0.0019	0.00042	0.00042	ug/kg	J+	bl
PDI-SC-S129-2TO4	SE	OCDD	6.2	0.0023	0.0023	ug/kg	J	q
PDI-SC-S129-4TO5.3	SE	1,2,3,7,8,9-HxCDF	0.0022	0.00051	0.00051	ug/kg	J+	bl

Sample ID	Matrix	Compound	Result	RDL	EDL	Units	Validation Qualifiers	Validation Reason
PDI-SC-S129-4TO5.3	SE	OCDD	8.6	0.0027	0.0027	ug/kg	J	q
PDI-SC-S155-0TO2.1	SE	1,2,3,4,6,7,8-HpCDF	0.012	0.00020	0.00020	ug/kg	JN	k
PDI-SC-S155-0TO2.1	SE	1,2,3,4,7,8,9-HpCDF	0.0015	0.00024	0.00024	ug/kg	J+	bl
PDI-SC-S155-0TO2.1	SE	1,2,3,7,8,9-HxCDF	0.0013	0.000063	0.000063	ug/kg	J+	bl
PDI-SC-S155-0TO2.1	SE	2,3,7,8-TCDD	0.00018	0.000029	0.000029	ug/kg	JN	k
PDI-SC-S155-2.1TO4.2	SE	1,2,3,7,8,9-HxCDF	0.0015	0.00077	0.00077	ug/kg	J+	bl
PDI-SC-S155-4.2TO5.3	SE	1,2,3,4,6,7,8-HpCDF	0.00042	0.000044	0.000044	ug/kg	J+	bl
PDI-SC-S155-4.2TO5.3	SE	1,2,3,4,7,8,9-HpCDF		0.00027	0.000054	ug/kg	U	bl
PDI-SC-S155-4.2TO5.3	SE	1,2,3,4,7,8-HxCDD	0.00013	0.000023	0.000023	ug/kg	J+	bl
PDI-SC-S155-4.2TO5.3	SE	1,2,3,6,7,8-HxCDD	0.00017	0.000023	0.000023	ug/kg	J+	bl
PDI-SC-S155-4.2TO5.3	SE	1,2,3,7,8,9-HxCDD	0.00021	0.000021	0.000021	ug/kg	JN	bl,k
PDI-SC-S155-4.2TO5.3	SE	1,2,3,7,8,9-HxCDF		0.00085	0.000023	ug/kg	U	bl
PDI-SC-S155-4.2TO5.3	SE	1,2,3,7,8-PeCDF	0.00029	0.000022	0.000022	ug/kg	J+	bl
PDI-SC-S155-4.2TO5.3	SE	2,3,4,7,8-PeCDF	0.000087	0.000024	0.000024	ug/kg	J+	bl
PDI-SC-S155-4.2TO5.3	SE	2,3,7,8-TCDF	0.00016	0.000015	0.000015	ug/kg	J+	bl
PDI-SC-S155-4.2TO5.3	SE	OCDF	0.0019	0.000050	0.000050	ug/kg	J+	bl
PDI-SC-S255-0TO2.1	SE	1,2,3,4,7,8,9-HpCDF	0.00056	0.00010	0.00010	ug/kg	J+	bl
PDI-SC-S255-0TO2.1	SE	1,2,3,4,7,8-HxCDD	0.00041	0.000032	0.000032	ug/kg	J+	bl
PDI-SC-S255-0TO2.1	SE	1,2,3,7,8,9-HxCDF		0.0010	0.000037	ug/kg	U	bl
PDI-SC-S255-0TO2.1	SE	1,2,3,7,8-PeCDF	0.00036	0.000038	0.000038	ug/kg	J+	bl
PDI-SC-S255-0TO2.1	SE	2,3,4,7,8-PeCDF	0.00023	0.000042	0.000042	ug/kg	J+	bl
PDI-SC-S255-0TO2.1	SE	2,3,7,8-TCDD	0.000061	0.000025	0.000025	ug/kg	JN	k
PDI-SC-S255-0TO2.1	SE	2,3,7,8-TCDF	0.00048	0.000052	0.000052	ug/kg	J+	bl
PDI-SC-S255-0TO2.1D	SE	1,2,3,4,7,8,9-HpCDF	0.00049	0.00010	0.00010	ug/kg	J+	bl
PDI-SC-S255-0TO2.1D	SE	1,2,3,4,7,8-HxCDD	0.00039	0.000030	0.000030	ug/kg	J+	bl
PDI-SC-S255-0TO2.1D	SE	1,2,3,7,8,9-HxCDF		0.00097	0.000030	ug/kg	U	bl
PDI-SC-S255-0TO2.1D	SE	1,2,3,7,8-PeCDD	0.00017	0.000033	0.000033	ug/kg	JN	k
PDI-SC-S255-0TO2.1D	SE	1,2,3,7,8-PeCDF	0.00037	0.000033	0.000033	ug/kg	J+	bl
PDI-SC-S255-0TO2.1D	SE	2,3,4,7,8-PeCDF	0.00019	0.000036	0.000036	ug/kg	JN	bl,k
PDI-SC-S255-0TO2.1D	SE	2,3,7,8-TCDD	0.000079	0.000023	0.000023	ug/kg	JN	k
PDI-SC-S255-0TO2.1D	SE	2,3,7,8-TCDF	0.00048	0.000056	0.000056	ug/kg	J+	bl
PDI-SC-S255-2.1TO4.3	SE	1,2,3,7,8,9-HxCDF		0.00085	0.00014	ug/kg	U	bl
PDI-SC-S255-2.1TO4.3	SE	1,2,3,7,8-PeCDF	0.00043	0.00018	0.00018	ug/kg	J+	bl
PDI-SC-S255-2.1TO4.3	SE	2,3,7,8-TCDD	0.00016	0.000028	0.000028	ug/kg	JN	k
PDI-SC-S255-2.1TO4.3	SE	2,3,7,8-TCDF	0.00051	0.00012	0.00012	ug/kg	J+	bl
PDI-SC-S260-0TO1.3	SE	1,2,3,4,7,8-HxCDD	0.0019	0.00026	0.00026	ug/kg	JN	k
PDI-SC-S260-1.3TO2.6	SE	2,3,7,8-TCDF	0.61	0.0016	0.0016	ug/kg	J	d,q
PDI-SC-S260-2.6TO4.2	SE	1,2,3,4,7,8-HxCDD	0.00020	0.000039	0.000039	ug/kg	J+	bl
PDI-SC-S260-2.6TO4.2	SE	1,2,3,7,8,9-HxCDF	0.0020	0.00034	0.00034	ug/kg	J+	bl
PDI-SC-S260-4.2TO6	SE	1,2,3,4,6,7,8-HpCDD	0.0010	0.000041	0.000041	ug/kg	J+	bl

Sample ID	Matrix	Compound	Result	RDL	EDL	Units	Validation Qualifiers	Validation Reason
PDI-SC-S260-4.2TO6	SE	1,2,3,4,6,7,8-HpCDF	0.00029	0.000042	0.000042	ug/kg	J+	bl
PDI-SC-S260-4.2TO6	SE	1,2,3,4,7,8,9-HpCDF	0.00036	0.000052	0.000052	ug/kg	J+	bl
PDI-SC-S260-4.2TO6	SE	1,2,3,4,7,8-HxCDD		0.000094	0.000021	ug/kg	U	bl
PDI-SC-S260-4.2TO6	SE	1,2,3,6,7,8-HxCDF	0.00021	0.000047	0.000047	ug/kg	J+	bl
PDI-SC-S260-4.2TO6	SE	1,2,3,7,8,9-HxCDD	0.00013	0.000019	0.000019	ug/kg	J+	bl
PDI-SC-S260-4.2TO6	SE	1,2,3,7,8,9-HxCDF		0.0012	0.000034	ug/kg	U	bl
PDI-SC-S260-4.2TO6	SE	1,2,3,7,8-PeCDF		0.00036	0.000028	ug/kg	U	bl
PDI-SC-S260-4.2TO6	SE	2,3,7,8-TCDF	0.00021	0.000025	0.000025	ug/kg	J+	bl
PDI-SC-S260-4.2TO6	SE	OCDF	0.00083	0.000053	0.000053	ug/kg	J+	bl
PDI-SC-S260-6TO7	SE	1,2,3,4,6,7,8-HpCDF	0.00023	0.000022	0.000022	ug/kg	J+	bl
PDI-SC-S260-6TO7	SE	1,2,3,4,7,8,9-HpCDF	0.00045	0.000028	0.000028	ug/kg	J+	bl
PDI-SC-S260-6TO7	SE	1,2,3,4,7,8-HxCDD		0.00013	0.000012	ug/kg	U	bl
PDI-SC-S260-6TO7	SE	1,2,3,6,7,8-HxCDD	0.000085	0.000013	0.000013	ug/kg	JN	k
PDI-SC-S260-6TO7	SE	1,2,3,6,7,8-HxCDF	0.000098	0.000029	0.000029	ug/kg	J+	bl
PDI-SC-S260-6TO7	SE	1,2,3,7,8,9-HxCDD	0.00017	0.000011	0.000011	ug/kg	J+	bl
PDI-SC-S260-6TO7	SE	1,2,3,7,8,9-HxCDF		0.0011	0.000022	ug/kg	U	bl
PDI-SC-S260-6TO7	SE	1,2,3,7,8-PeCDD	0.000040	0.000023	0.000023	ug/kg	JN	k
PDI-SC-S260-6TO7	SE	1,2,3,7,8-PeCDF		0.00021	0.000017	ug/kg	U	bl
PDI-SC-S260-6TO7	SE	2,3,4,6,7,8-HxCDF	0.000034	0.000025	0.000025	ug/kg	JN	k
PDI-SC-S260-6TO7	SE	2,3,4,7,8-PeCDF	0.00011	0.000020	0.000020	ug/kg	J+	bl
PDI-SC-S260-6TO7	SE	2,3,7,8-TCDF	0.00040	0.000018	0.000018	ug/kg	J+	bl
PDI-SC-S260-6TO7	SE	OCDF	0.00088	0.000033	0.000033	ug/kg	J+	bl

**Attachment A**  
**Qualifier Codes and Explanations**

<b>Qualifier</b>	<b>Explanation</b>
J	The analyte was positively identified; the associated numerical value is the approximate concentration of the analyte in the sample.
J-	The analyte was positively identified; the associated numerical value is the approximate concentration of the analyte in the sample with a potential low bias.
J+	The analyte was positively identified; the associated numerical value is the approximate concentration of the analyte in the sample with a potential high bias.
JN	The analyte was tentatively identified; the associated numerical value is the approximate concentration of the analyte in the sample.
UJ	The analyte was not detected above the reported sample quantitation limit. However, the reported quantitation limit is approximate and may or may not represent the actual limit of quantitation necessary to accurately and precisely measure the analyte in the sample.
U	The analyte was analyzed for, but was not detected above the reported sample quantitation limit.
R	The sample results are rejected due to serious deficiencies in the ability to analyze the sample and meet quality control criteria. The presence or absence of the analyte cannot be verified.

## Attachment B

### Reason Codes and Explanations

Reason Code	Explanation
be	Equipment blank contamination
bf	Field blank contamination
bl	Laboratory blank contamination
c	Calibration issue
cl	Clean-up standard recovery
d	Reporting limit raised due to chromatographic interference
fd	Field duplicate RPDs
h	Holding times
i	Internal standard areas
k	Estimated Maximum Possible Concentration (EMPC)
l	LCS or OPR recoveries
lc	Labeled compound recovery
ld	Laboratory duplicate RPDs
lp	Laboratory control sample/laboratory control sample duplicate RPDs
m	Matrix spike recovery
md	Matrix spike/matrix spike duplicate RPDs
ma	Multiple analyses, sample analyzed more than once, a value from another analysis should be used
nb	Negative laboratory blank contamination
p	Chemical preservation issue
r	Dual column RPD
q	Quantitation issue
s	Surrogate recovery
su	Ion suppression
t	Temperature preservation issue
x	Percent solids
y	Serial dilution results
z	ICS results