

Data Validation Report

Project:	Portland Harbor Pre-Remedial Design Investigation and Baseline Sampling	
Laboratory:	Test America, West Sacramento, California	
Laboratory Group:	580-77502-2	
Analyses/Method:	Clean Water Act - Dioxins and Furans / CWA1613B	
Validation Level:	Stage 2A	
AECOM Project Number:	60566335.2.12	
Prepared by:	Paula DiMattei/AECOM	Completed on: 07/16/2018
Reviewed by:	Waverly Braunstein/AECOM	File Name: 580-77502-2 DVR

SUMMARY

The samples listed below were collected by AECOM in Portland Harbor in Portland, OR on May 21, 2018 and May 22, 2018.

Sample ID	Matrix/Sample Type
PDI-RB-VV-182105	Equipment Blank
PDI-SG-B234-BL1-D	Field Duplicate of PDI-SG-B234-BL1
PDI-SG-B315-BL1-D	Field Duplicate of PDI-SG-B315-BL1
PDI-SG-B214-BL1	Sediment
PDI-SG-B234-BL1	Sediment
PDI-SG-B246-BL1	Sediment
PDI-SG-B251-BL1	Sediment
PDI-SG-B257-BL1	Sediment
PDI-SG-B298-BL1	Sediment
PDI-SG-B299-BL1	Sediment
PDI-SG-B305-BL1	Sediment
PDI-SG-B315-BL1	Sediment
PDI-SG-B398-BL1	Sediment
PDI-SG-B417-BL1	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 (✗) 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 or negated 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 equipment blank and the results are tabulated below for informational purposes.

Blank ID	Compound	Result	QL	Units
PDI-RB-VV-182105	1,2,3,4,6,7,8-HpCDD	11	49	pg/L
	1,2,3,4,6,7,8-HpCDF	7.2	49	pg/L
	1,2,3,4,7,8,9-HpCDF	8.6	49	pg/L
	1,2,3,4,7,8-HxCDD	4.2	49	pg/L
	1,2,3,6,7,8-HxCDD	2.3	49	pg/L
	1,2,3,7,8,9-HxCDD	5.7	49	pg/L
	1,2,3,7,8,9-HxCDF	6.3	49	pg/L
	1,2,3,7,8-PeCDF	1.1	49	pg/L
	2,3,7,8-TCDF	1.4	9.8	pg/L
	OCDD	30	98	pg/L
	OCDF	22	98	pg/L

Target compounds were detected in the method blanks associated with the samples in this data set. Detected compounds are summarized in Attachment A in Table A-1.

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. All method QC acceptance criteria were met.

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.

All field duplicate precision 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.

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

The sample data was 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. Results were reported from the confirmation column when analysis on a second column was required. Qualification of the data was not required.

Estimated Maximum Possible Concentrations (EMPCs)

The data were reviewed to identify sample results that were indicated by the laboratory to be estimated maximum possible concentrations (EMPCs) because of identification criteria not being met.

The laboratory identified the presence of EMPCs for the samples in this data set. Samples were qualified as follows:

Actions: (Based on AECOM professional judgment)

Criteria	Actions
A native target compound was reported by the laboratory as an EMPC.	Report result as an EMPC and qualify as estimated and presumptively present (JN).
A labeled compound was flagged by the laboratory indicating all identification criteria were not met.	Qualify associated positive and nondetect results as estimated (J/UJ).

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.

Qualified sample results are summarized in Table 1.

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 of the data was not required.

QUALIFICATION ACTIONS

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

ATTACHMENTS

Attachment A: Nonconformance Summary Tables

Attachment B: Qualifier Codes and Explanations

Attachment C: Reason Codes and Explanations

Table 1 - Data Validation Summary of Qualified Data

Sample ID	Matrix	Compound	Result	EDL	Units	Validation Qualifiers	Validation Reason
PDI-RB-VV-182105	WQ	1,2,3,4,6,7,8-HpCDD	11	0.58	pg/L	J+	bl
PDI-RB-VV-182105	WQ	1,2,3,4,6,7,8-HpCDF	7.2	0.64	pg/L	JN	bl,k
PDI-RB-VV-182105	WQ	1,2,3,4,7,8,9-HpCDF	8.6	0.74	pg/L	J+	bl
PDI-RB-VV-182105	WQ	1,2,3,4,7,8-HxCDD	4.2	0.67	pg/L	J+	bl
PDI-RB-VV-182105	WQ	1,2,3,6,7,8-HxCDD	2.3	0.59	pg/L	J+	bl
PDI-RB-VV-182105	WQ	1,2,3,7,8,9-HxCDD	5.7	0.59	pg/L	J+	bl
PDI-RB-VV-182105	WQ	1,2,3,7,8,9-HxCDF	6.3	0.53	pg/L	J+	bl
PDI-RB-VV-182105	WQ	1,2,3,7,8-PeCDF	1.1	0.45	pg/L	JN	k
PDI-RB-VV-182105	WQ	2,3,7,8-TCDF	1.4	0.19	pg/L	J+	bl
PDI-RB-VV-182105	WQ	OCDD	30	0.46	pg/L	J+	bl
PDI-RB-VV-182105	WQ	OCDF	22	0.46	pg/L	J+	bl
PDI-SG-B214-BL1	SE	1,2,3,4,7,8,9-HpCDF	0.00087	0.00047	ug/kg	J+	bl
PDI-SG-B214-BL1	SE	1,2,3,7,8,9-HxCDF	0.0010	0.00012	ug/kg	J+	bl
PDI-SG-B214-BL1	SE	1,2,3,7,8-PeCDD	0.00041	0.00012	ug/kg	JN	bl,k
PDI-SG-B214-BL1	SE	2,3,4,7,8-PeCDF	0.00026	0.000092	ug/kg	JN	k
PDI-SG-B234-BL1	SE	1,2,3,4,7,8,9-HpCDF	0.0021	0.00076	ug/kg	J+	bl
PDI-SG-B234-BL1	SE	1,2,3,4,7,8-HxCDD	0.00092	0.00018	ug/kg	JN	k
PDI-SG-B234-BL1	SE	1,2,3,6,7,8-HxCDD	0.0035	0.00017	ug/kg	JN	k
PDI-SG-B234-BL1	SE	1,2,3,6,7,8-HxCDF	0.00078	0.00024	ug/kg	JN	k
PDI-SG-B234-BL1	SE	1,2,3,7,8,9-HxCDF	0.0021	0.00020	ug/kg	JN	bl,k
PDI-SG-B234-BL1	SE	1,2,3,7,8-PeCDD	0.00050	0.00020	ug/kg	JN	bl,k
PDI-SG-B234-BL1-D	SE	1,2,3,4,7,8,9-HpCDF	0.0013	0.00078	ug/kg	J+	bl
PDI-SG-B234-BL1-D	SE	1,2,3,4,7,8-HxCDD	0.00098	0.00022	ug/kg	JN	k
PDI-SG-B234-BL1-D	SE	1,2,3,7,8,9-HxCDF	0.0014	0.00018	ug/kg	J+	bl
PDI-SG-B234-BL1-D	SE	1,2,3,7,8-PeCDD	0.00049	0.00020	ug/kg	JN	bl,k
PDI-SG-B234-BL1-D	SE	1,2,3,7,8-PeCDF	0.00061	0.00021	ug/kg	JN	k
PDI-SG-B234-BL1-D	SE	2,3,4,6,7,8-HxCDF	0.00076	0.00019	ug/kg	J+	bl
PDI-SG-B234-BL1-D	SE	2,3,7,8-TCDD	0.0013	0.00018	ug/kg	JN	k
PDI-SG-B246-BL1	SE	1,2,3,4,6,7,8-HpCDF	0.010	0.00042	ug/kg	JN	k
PDI-SG-B246-BL1	SE	1,2,3,4,7,8,9-HpCDF	0.0016	0.00051	ug/kg	JN	bl,k
PDI-SG-B246-BL1	SE	1,2,3,4,7,8-HxCDF	0.0016	0.00025	ug/kg	JN	k
PDI-SG-B246-BL1	SE	1,2,3,6,7,8-HxCDD	0.0027	0.00019	ug/kg	JN	k
PDI-SG-B246-BL1	SE	1,2,3,6,7,8-HxCDF	0.00082	0.00022	ug/kg	JN	k
PDI-SG-B246-BL1	SE	1,2,3,7,8,9-HxCDD	0.0019	0.00018	ug/kg	JN	k
PDI-SG-B246-BL1	SE	1,2,3,7,8,9-HxCDF	0.0018	0.00019	ug/kg	J+	bl
PDI-SG-B246-BL1	SE	1,2,3,7,8-PeCDD	0.00040	0.00019	ug/kg	J+	bl
PDI-SG-B246-BL1	SE	1,2,3,7,8-PeCDF	0.00085	0.00015	ug/kg	JN	k
PDI-SG-B246-BL1	SE	2,3,4,6,7,8-HxCDF	0.0015	0.00021	ug/kg	JN	k

Sample ID	Matrix	Compound	Result	EDL	Units	Validation Qualifiers	Validation Reason
PDI-SG-B246-BL1	SE	2,3,4,7,8-PeCDF	0.00052	0.00016	ug/kg	JN	k
PDI-SG-B251-BL1	SE	1,2,3,4,7,8,9-HpCDF	0.0016	0.00069	ug/kg	J+	bl
PDI-SG-B251-BL1	SE	1,2,3,6,7,8-HxCDF	0.0012	0.00023	ug/kg	JN	k
PDI-SG-B251-BL1	SE	1,2,3,7,8,9-HxCDF	0.0015	0.00021	ug/kg	JN	bl,k
PDI-SG-B251-BL1	SE	1,2,3,7,8-PeCDD	0.00049	0.00026	ug/kg	J+	bl
PDI-SG-B251-BL1	SE	1,2,3,7,8-PeCDF	0.00081	0.00017	ug/kg	JN	k
PDI-SG-B251-BL1	SE	2,3,4,6,7,8-HxCDF	0.00058	0.00022	ug/kg	J+	bl
PDI-SG-B257-BL1	SE	1,2,3,4,6,7,8-HpCDD	0.00078	0.000075	ug/kg	J+	bl
PDI-SG-B257-BL1	SE	1,2,3,4,7,8,9-HpCDF		0.00019	ug/kg	U	bl
PDI-SG-B257-BL1	SE	1,2,3,7,8,9-HxCDF		0.00050	ug/kg	U	bl
PDI-SG-B257-BL1	SE	OCDD	0.0052	0.00011	ug/kg	J+	bl
PDI-SG-B298-BL1	SE	1,2,3,4,7,8,9-HpCDF	0.0014	0.00071	ug/kg	J+	bl
PDI-SG-B298-BL1	SE	1,2,3,4,7,8-HxCDF	0.0014	0.00021	ug/kg	JN	k
PDI-SG-B298-BL1	SE	1,2,3,6,7,8-HxCDF	0.00067	0.00020	ug/kg	JN	k
PDI-SG-B298-BL1	SE	1,2,3,7,8,9-HxCDF		0.00043	ug/kg	U	bl
PDI-SG-B298-BL1	SE	1,2,3,7,8-PeCDD	0.00056	0.00011	ug/kg	JN	bl,k
PDI-SG-B298-BL1	SE	2,3,4,6,7,8-HxCDF	0.00057	0.00017	ug/kg	J+	bl
PDI-SG-B298-BL1	SE	2,3,7,8-TCDD	0.00071	0.000092	ug/kg	JN	k
PDI-SG-B299-BL1	SE	1,2,3,4,7,8,9-HpCDF	0.0039	0.0017	ug/kg	J+	bl
PDI-SG-B299-BL1	SE	1,2,3,7,8,9-HxCDF	0.0019	0.00025	ug/kg	JN	bl,k
PDI-SG-B299-BL1	SE	1,2,3,7,8-PeCDD	0.0011	0.00023	ug/kg	JN	bl,k
PDI-SG-B299-BL1	SE	2,3,4,6,7,8-HxCDF	0.0013	0.00024	ug/kg	J+	bl
PDI-SG-B305-BL1	SE	1,2,3,4,6,7,8-HpCDF	0.0081	0.00044	ug/kg	JN	k
PDI-SG-B305-BL1	SE	1,2,3,4,7,8,9-HpCDF	0.0013	0.00059	ug/kg	J+	bl
PDI-SG-B305-BL1	SE	1,2,3,4,7,8-HxCDD	0.00059	0.00014	ug/kg	JN	k
PDI-SG-B305-BL1	SE	1,2,3,7,8,9-HxCDF	0.0017	0.00017	ug/kg	J+	bl
PDI-SG-B305-BL1	SE	1,2,3,7,8-PeCDD	0.00031	0.00017	ug/kg	JN	bl,k
PDI-SG-B305-BL1	SE	2,3,4,7,8-PeCDF	0.00042	0.00016	ug/kg	JN	k
PDI-SG-B315-BL1	SE	1,2,3,4,7,8,9-HpCDF	0.0014	0.00059	ug/kg	J+	bl
PDI-SG-B315-BL1	SE	1,2,3,6,7,8-HxCDF	0.0011	0.00022	ug/kg	JN	k
PDI-SG-B315-BL1	SE	1,2,3,7,8,9-HxCDF	0.0012	0.00019	ug/kg	J+	bl
PDI-SG-B315-BL1	SE	1,2,3,7,8-PeCDD	0.00049	0.00016	ug/kg	JN	bl,k
PDI-SG-B315-BL1-D	SE	1,2,3,4,7,8,9-HpCDF	0.0011	0.00059	ug/kg	JN	bl,k
PDI-SG-B315-BL1-D	SE	1,2,3,7,8,9-HxCDD	0.0021	0.00013	ug/kg	JN	k
PDI-SG-B315-BL1-D	SE	1,2,3,7,8-PeCDD	0.00060	0.00012	ug/kg	JN	bl,k
PDI-SG-B315-BL1-D	SE	1,2,3,7,8-PeCDF	0.00046	0.00016	ug/kg	JN	k
PDI-SG-B315-BL1-D	SE	2,3,4,6,7,8-HxCDF	0.00054	0.00017	ug/kg	JN	bl,k
PDI-SG-B315-BL1-D	SE	2,3,7,8-TCDD	0.00075	0.00014	ug/kg	JN	k
PDI-SG-B398-BL1	SE	1,2,3,4,6,7,8-HpCDF	0.014	0.00051	ug/kg	JN	k
PDI-SG-B398-BL1	SE	1,2,3,4,7,8,9-HpCDF	0.0011	0.00059	ug/kg	JN	bl,k

Sample ID	Matrix	Compound	Result	EDL	Units	Validation Qualifiers	Validation Reason
PDI-SG-B398-BL1	SE	1,2,3,7,8,9-HxCDF	0.0017	0.00019	ug/kg	JN	bl,k
PDI-SG-B398-BL1	SE	1,2,3,7,8-PeCDD	0.00058	0.00021	ug/kg	J+	bl
PDI-SG-B398-BL1	SE	2,3,4,7,8-PeCDF	0.00034	0.00017	ug/kg	JN	k
PDI-SG-B417-BL1	SE	1,2,3,4,6,7,8-HpCDD	0.0030	0.00018	ug/kg	JN	k
PDI-SG-B417-BL1	SE	1,2,3,4,7,8,9-HpCDF	0.00059	0.00014	ug/kg	J+	bl
PDI-SG-B417-BL1	SE	1,2,3,6,7,8-HxCDD	0.00021	0.000096	ug/kg	JN	k
PDI-SG-B417-BL1	SE	1,2,3,6,7,8-HxCDF	0.00013	0.000088	ug/kg	JN	k
PDI-SG-B417-BL1	SE	1,2,3,7,8,9-HxCDD	0.00018	0.000087	ug/kg	JN	k
PDI-SG-B417-BL1	SE	1,2,3,7,8,9-HxCDF	0.00070	0.000068	ug/kg	JN	bl,k
PDI-SG-B417-BL1	SE	2,3,7,8-TCDF	0.00018	0.000058	ug/kg	JN	k
PDI-SG-B417-BL1	SE	OCDF	0.0015	0.00018	ug/kg	J+	bl

Attachment A

Nonconformance Summary Tables

Table A-1 - Lab Blanks

Blank ID	Compound	Result	QL	BAL	Units	Associated Samples
MB 320-227727/1-A	1,2,3,4,6,7,8-HpCDD	3.66	50	18.3	pg/L	PDI-RB-VV-182105
	1,2,3,4,6,7,8-HpCDF	2.21	50	11.0	pg/L	
	1,2,3,4,7,8,9-HpCDF	3.35	50	16.8	pg/L	
	1,2,3,4,7,8-HxCDD	1.61	50	8.05	pg/L	
	1,2,3,6,7,8-HxCDD	0.920	50	4.60	pg/L	
	1,2,3,7,8,9-HxCDD	1.13	50	5.65	pg/L	
	1,2,3,7,8,9-HxCDF	3.12	50	15.6	pg/L	
	2,3,4,6,7,8-HxCDF	0.889	50	4.44	pg/L	
	2,3,7,8-TCDF	0.976	10	4.88	pg/L	
	OCDD	14.5	100	72.5	pg/L	
OCDF	8.46	100	42.3	pg/L		
MB 320-228667/1-A	1,2,3,4,6,7,8-HpCDD	0.000269	0.0050	0.001345	ug/kg	PDI-SG-B214-BL1 PDI-SG-B234-BL1 PDI-SG-B234-BL1-D PDI-SG-B246-BL1 PDI-SG-B251-BL1 PDI-SG-B257-BL1 PDI-SG-B298-BL1 PDI-SG-B299-BL1 PDI-SG-B305-BL1 PDI-SG-B315-BL1 PDI-SG-B315-BL1-D PDI-SG-B398-BL1 PDI-SG-B417-BL1
	1,2,3,4,7,8,9-HpCDF	0.000554	0.0050	0.00277	ug/kg	
	1,2,3,7,8,9-HxCDF	0.000646	0.0050	0.00323	ug/kg	
	1,2,3,7,8-PeCDD	0.000252	0.0050	0.00126	ug/kg	
	2,3,4,6,7,8-HxCDF	0.000158	0.0050	0.00079	ug/kg	
	OCDD	0.000682	0.010	0.00341	ug/kg	
	OCDF	0.000268	0.010	0.00134	ug/kg	

Attachment B**Qualifier Codes and Explanations**

Qualifier	Explanation
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 C

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
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