



LABORATORY DATA CONSULTANTS, INC.

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Northgate Environmental Management, Inc.
1100 Quail Street Ste. 102
Newport Beach, CA 92660
ATTN: Ms. Cindy Arnold

June 4, 2010

SUBJECT: Tronox LLC Facility, PCS, Henderson, Nevada,
Data Validation

Dear Ms. Arnold,

Enclosed are the final validation reports for the fraction listed below. These SDGs were received on May 14, 2010. Attachment 1 is a summary of the samples that were reviewed for each analysis.

LDC Project # 23188:

SDG #

Fraction

G0D100461, G0D130435, G0D140422	Dioxins/Dibenzofurans
G0D140526, G0D140534, G0D140543	
G0D160437, G0D170488	

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

- Standard Operating Procedures (SOP) 40, Data Review/Validation, BRC 2009
- Quality Assurance Project Plan Tronox LLC Facility, Henderson Nevada, June 2009
- NDEP Guidance, May 2006
- USEPA, Contract Laboratory Program National Functional Guidelines for Polychlorinated Dioxins/Diobenzofurans Data Review, September 2005

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

Sincerely,

Erlinda T. Rauto
Operations Manager/Senior Chemist

EDD CHECKLIST

LDC #: 23188

SDG #: G0D100461, G0D130435, G0D140422, G0D140526

G0D140534, G0D140543, G0D160437, G0D170488

Tronox Northgate Henderson Worksheet

EDD Area	Yes	No	NA	Findings/Comments
I. Completeness				
Is there an EDD for the associated Tronox validation report?	X			
II. EDD Qualifier Population				
Were all qualifiers from the validation report populated into the EDD?	X			
III. EDD Lab Anomalies				
Were EDD anomalies identified?	X			
If yes, were they corrected or documented for the client?	X			See EDD_discrepancy_form_LDC23188_060310.doc
IV. EDD Delivery				
Was the final EDD sent to the client?	X			

**Tronox LLC Facility, PCS, Henderson, Nevada
Data Validation Reports
LDC #23188**

Dioxins/Dibenzofurans

LDC

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

Project/Site Name: Tronox LLC Facility, PCS, Henderson, Nevada
Collection Date: April 8, 2010
LDC Report Date: June 3, 2010
Matrix: Soil
Parameters: Dioxins/Dibenzofurans
Validation Level: Stage 2B & 4
Laboratory: TestAmerica, Inc.

Sample Delivery Group (SDG): G0D100461

Sample Identification

SSAO5-02-1BPC
SSAO5-03-1BPC**
SSAO5-02-1BPC_FD
SSAO5-03-1BPC_FD
SSAO4-02-3BPC

**Indicates sample underwent Stage 4 review

Introduction

This data review covers 5 soil samples listed on the cover sheet including dilutions and reanalysis as applicable. The analyses were per EPA SW 846 Method 8290 for Polychlorinated Dioxins/Dibenzofurans.

This review follows the Standard Operating Procedures (SOP) 40, Data Review/Validation (BRC 2009), the Quality Assurance Project Plan Tronox LLC Facility, Henderson, Nevada (June 2009), NDEP guidance (May 2006), and USEPA Contract Laboratory Program National Functional Guidelines for Polychlorinated Dioxins/Dibenzofurans Data Review (September 2005).

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.

Blank results are summarized in Section V.

Field duplicates are summarized in Section XIV.

Samples indicated by a double asterisk on the front cover underwent a Stage 4 review. A Stage 2B review was performed on all of the other samples. Raw data were not evaluated for the samples reviewed by Stage 2B criteria since this review is based on QC data.

The following are definitions of the data qualifiers:

- J+ Data are qualified as estimated, with a high bias likely to occur. False positives or false negatives are unlikely to have been reported.
- J- Data are qualified as estimated, with a low bias likely to occur. False positives or false negatives are unlikely to have been reported.
- J Data are qualified as estimated; it is not possible to assess the direction of the potential bias. False positives or false negatives are unlikely to have been reported.
- U Indicates the compound or analyte was analyzed for but not detected at or above the stated limit.
- R Data are qualified as rejected. There is a significant potential for the reporting of false negatives or false positives.
- UU Indicates the compound or analyte was analyzed for but not detected. The sample detection limit is an estimated value.
- B The analytical result may be a false positive totally attributable to blank contamination. This qualifier is applicable to radiochemistry analysis only.
- JB The analytical result may be biased high and partially attributable to blank contamination. This qualifier is applicable to radiochemistry analysis only.
- JK The analytical result is an estimated maximum possible concentration (EMPC).
- X The analytical result is not used for reporting because a more accurate and precise result is reported in its place.
- J-TDS The analytical result is estimated based on failure of the Total Dissolved Solids (TDS) correctness check performed in accordance with the Standard Method 1030E.
- J-CAB The analytical result is estimated based on failure of the cation-anion balance correctness check performed in accordance with Standard Method 1030E.
- J-TDS & CAB The analytical result is unreliable based on the failure of the cation-anion balance and TDS correctness check performed in accordance with standard Method 1030E.
- A Indicates the finding is based upon technical validation criteria.
- P Indicates the finding is related to a protocol/contractual deviation.
- None Indicates the data was not significantly impacted by the finding, therefore qualification was not required.

I. Technical Holding Times

All technical holding time requirements were met.

The chain-of-custodies were reviewed for documentation of cooler temperatures. All cooler temperatures met validation criteria.

II. HRGC/HRMS Instrument Performance Check

Instrument performance was checked at the required daily 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 exact mass of 380.9760 of PFK was verified. The static resolving power was at least 10,000 (10% valley definition) for samples on which Stage 4 review was performed. Raw data were not evaluated for the samples reviewed by Stage 2B criteria.

III. Initial Calibration

A five point initial calibration was performed as required by the method.

Percent relative standard deviations (%RSD) were less than or equal to 20.0% for unlabeled compounds and less than or equal to 30.0% for labeled compounds.

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

The minimum S/N ratio for each target compound was greater than or equal to 2.5 and greater than or equal to 10 for each recovery and internal standard compound for samples on which Stage 4 review was performed. Raw data were not evaluated for the samples reviewed by Stage 2B criteria.

IV. Routine Calibration (Continuing)

Routine calibration was performed at the required frequencies.

All of the routine calibration percent differences (%D) between the initial calibration RRF and the routine calibration RRF were less than or equal to 20.0% for unlabeled compounds and less than or equal to 30.0% for labeled compounds with the following exceptions:

Date	Compound	%D	Associated Samples	Affected Compound	Flag	A or P
4/20/10	¹³ C-1,2,3,4,7,8-HxCDF	30.5	SSAO5-03-1BPC** SSAO5-03-1BPC_FD	1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 2,3,4,6,7,8-HxCDF 1,2,3,7,8,9-HxCDF	J+ (all detects) J+ (all detects) J+ (all detects) J+ (all detects)	P

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

V. Blanks

Method blanks were reviewed for each matrix as applicable. No polychlorinated dioxin/dibenzofuran contaminants were found in the method blanks with the following exceptions:

Method Blank ID	Extraction Date	Compound	Concentration	Associated Samples
0103408MB	4/13/10	1,2,3,7,8,9-HxCDD 1,2,3,4,6,7,8-HpCDD OCDD 2,3,7,8-TCDF 1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	0.040 pg/g 0.13 pg/g 0.71 pg/g 0.033 pg/g 0.048 pg/g 0.033 pg/g 0.13 pg/g 0.068 pg/g 0.32 pg/g	SSAO5-02-1BPC SSAO5-03-1BPC** SSAO5-02-1BPC_FD SSAO5-03-1BPC_FD

Sample concentrations were compared to concentrations detected in the method blanks as required by the QAPP. No sample data was qualified.

Sample EB-04072010-RZC (from SDG G0D130519) was identified as an equipment blank. No polychlorinated dioxin/dibenzofuran contaminants were found in this blank with the following exceptions:

Equipment Blank ID	Sampling Date	Compound	Concentration	Associated Samples
EB-04072010-RZC	4/8/10	2,3,7,8-TCDD 1,2,3,6,7,8-HxCDD 1,2,3,7,8,9-HxCDD 1,2,3,4,6,7,8-HpCDD OCDD 2,3,7,8-TCDF 1,2,3,7,8-PeCDF 2,3,4,7,8-PeCDF 1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 2,3,4,6,7,8-HxCDF 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	1.0 pg/L 0.69 pg/L 0.65 pg/L 5.5 pg/L 53 pg/L 2.6 pg/L 1.5 pg/L 1.0 pg/L 1.8 pg/L 1.1 pg/L 0.97 pg/L 4.5 pg/L 1.1 pg/L 12 pg/L	All samples in SDG G0D100461

Sample concentrations were compared to concentrations detected in the equipment blanks as required by the QAPP. No sample data was qualified.

Sample FB-04072010-RZC (from SDG G0D130519) was identified as a field blank. No polychlorinated dioxin/dibenzofuran contaminants were found in this blank with the following exceptions:

Field Blank ID	Sampling Date	Compound	Concentration	Associated Samples
FB-04072010-RZC	4/8/10	1,2,3,4,7,8-HxCDD	0.77 pg/L	All samples in SDG G0D100461
		1,2,3,6,7,8-HxCDD	0.74 pg/L	
		1,2,3,7,8,9-HxCDD	0.82 pg/L	
		1,2,3,4,6,7,8-HpCDD	4.2 pg/L	
		OCDD	37 pg/L	
		2,3,7,8-TCDF	0.57 pg/L	
		1,2,3,7,8-PeCDF	0.96 pg/L	
		2,3,4,7,8-PeCDF	0.67 pg/L	
		1,2,3,4,7,8-HxCDF	1.1 pg/L	
		1,2,3,6,7,8-HxCDF	0.96 pg/L	
		2,3,4,6,7,8-HxCDF	1.0 pg/L	
		1,2,3,7,8,9-HxCDF	1.0 pg/L	
		1,2,3,4,6,7,8-HpCDF	2.1 pg/L	
		1,2,3,4,7,8,9-HpCDF	1.5 pg/L	
		OCDF	6.7 pg/L	

Sample concentrations were compared to concentrations detected in the field blanks as required by the QAPP. No sample data was qualified.

VI. Matrix Spike/Matrix Spike Duplicates

Matrix spike (MS) and matrix spike duplicate (MSD) samples were reviewed for each matrix as applicable. Percent recoveries (%R) and relative percent differences (RPD) were not within QC limits. Since there were no associated samples, no data were qualified.

VII. Laboratory Control Samples (LCS)

Laboratory control samples were reviewed for each matrix as applicable. Percent recoveries (%R) were within QC limits.

VIII. Regional Quality Assurance and Quality Control

Not applicable.

IX. Internal Standards

All internal standard recoveries were within QC limits with the following exceptions:

Sample	Internal Standards	%R (Limits)	Compound	Flag	A or P
SSAO5-02-1BPC	¹³ C-OCDD	31 (40-135)	OCDD OCDF	J (all detects) UJ (all non-detects) J (all detects) UJ (all non-detects)	P
SSAO5-03-1BPC**	¹³ C-OCDD	38 (40-135)	OCDD OCDF	J (all detects) UJ (all non-detects) J (all detects) UJ (all non-detects)	P
SSAO4-02-3BPC	¹³ C-OCDD	36 (40-135)	OCDD OCDF	J (all detects) UJ (all non-detects) J (all detects) UJ (all non-detects)	P

X. Target Compound Identifications

All target compound identifications were within validation criteria for samples on which a Stage 4 review was performed. Raw data were not evaluated for the samples reviewed by Stage 2B criteria.

XI. Project Quantitation Limit

All compound quantitation and CRQLs were within validation criteria.

All compounds reported below the PQL were qualified as follows:

Sample	Finding	Flag	A or P
All samples in SDG G0D100461	All compounds reported below the PQL.	J (all detects)	A

All compounds reported as EMPC were qualified as follows:

Sample	Finding	Flag	A or P
All samples in SDG G0D100461	All compounds reported as estimated maximum possible concentration (EMPC).	JK (all detects)	A

Raw data were not evaluated for the samples reviewed by Stage 2B criteria.

XII. System Performance

The system performance was acceptable for samples on which a Stage 4 review was performed. Raw data were not evaluated for the samples reviewed by Stage 2B criteria.

XIII. Overall Assessment of Data

Data flags are summarized at the end of this report if data has been qualified.

XIV. Field Duplicates

Samples SSAO5-02-1BPC and SSAO5-02-1BPC_FD and samples SSAO5-03-1BPC** and SSAO5-03-1BPC_FD were identified as field duplicates. No polychlorinated dioxins/dibenzofurans were detected in any of the samples with the following exceptions:

Compound	Concentration (pg/g)		RPD (Limits)	Difference (Limits)	Flags	A or P
	SSAO5-02-1BPC	SSAO5-02-1BPC_FD				
2,3,7,8-TCDD	2.5	2.4	-	0.1 (≤ 0.53)	-	-
1,2,3,7,8-PeCDD	7.7	6.8	-	0.9 (≤ 2.6)	-	-
1,2,3,4,7,8-HxCDD	4.7	4.4	-	0.3 (≤ 2.6)	-	-
1,2,3,6,7,8-HxCDD	10	8.7	-	1.3 (≤ 2.6)	-	-
1,2,3,7,8,9-HxCDD	9.2	8.4	-	0.8 (≤ 2.6)	-	-
1,2,3,4,6,7,8-HpCDD	33	30	10 (≤ 50)	-	-	-
OCDD	38	40	5 (≤ 50)	-	-	-
2,3,7,8-TCDF	75	71	5 (≤ 50)	-	-	-
1,2,3,7,8-PeCDF	130	120	8 (≤ 50)	-	-	-
2,3,4,7,8-PeCDF	69	66	4 (≤ 50)	-	-	-
1,2,3,4,7,8-HxCDF	250	240	4 (≤ 50)	-	-	-
1,2,3,6,7,8-HxCDF	120	120	0 (≤ 50)	-	-	-
2,3,4,6,7,8-HxCDF	32	28	13 (≤ 50)	-	-	-
1,2,3,7,8,9-HxCDF	31	29	7 (≤ 50)	-	-	-
1,2,3,4,6,7,8-HpCDF	530	490	8 (≤ 50)	-	-	-
1,2,3,4,7,8,9-HpCDF	290	260	11 (≤ 50)	-	-	-

Compound	Concentration (pg/g)		RPD (Limits)	Difference (Limits)	Flags	A or P
	SSAO5-02-1BPC	SSAO5-02-1BPC_FD				
OCDF	1900	1800	5 (≤ 50)	-	-	-

Compound	Concentration (pg/g)		RPD (Limits)	Difference (Limits)	Flags	A or P
	SSAO5-03-1BPC**	SSAO5-03-1BPC_FD				
2,3,7,8-TCDD	31	33	-	2 (≤ 0.11)	-	-
1,2,3,7,8-PeCDD	110	110	-	0 (≤ 53)	-	-
1,2,3,4,7,8-HxCDD	66	84	-	18 (≤ 53)	-	-
1,2,3,6,7,8-HxCDD	150	150	-	0 (≤ 53)	-	-
1,2,3,7,8,9-HxCDD	130	150	-	20 (≤ 53)	-	-
1,2,3,4,6,7,8-HpCDD	530	630	17 (≤ 50)	-	-	-
OCDD	560	680	19 (≤ 50)	-	-	-
2,3,7,8-TCDF	730	780	7 (≤ 50)	-	-	-
1,2,3,7,8-PeCDF	1500	1600	6 (≤ 50)	-	-	-
2,3,4,7,8-PeCDF	800	820	2 (≤ 50)	-	-	-
1,2,3,4,7,8-HxCDF	3400	3600	6 (≤ 50)	-	-	-
1,2,3,6,7,8-HxCDF	1700	1800	6 (≤ 50)	-	-	-
2,3,4,6,7,8-HxCDF	310	350	12 (≤ 50)	-	-	-
1,2,3,7,8,9-HxCDF	350	370	6 (≤ 50)	-	-	-
1,2,3,4,6,7,8-HpCDF	7600	8200	8 (≤ 50)	-	-	-
1,2,3,4,7,8,9-HpCDF	3600	4000	11 (≤ 50)	-	-	-
OCDF	24000	28000	15 (≤ 50)	-	-	-

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Data Qualification Summary - SDG G0D100461**

SDG	Sample	Compound	Flag	A or P	Reason (Code)
G0D100461	SSAO5-03-1BPC** SSAO5-03-1BPC_FD	1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 2,3,4,6,7,8-HxCDF 1,2,3,7,8,9-HxCDF	J+ (all detects) J+ (all detects) J+ (all detects) J+ (all detects)	P	Routine calibration (%D) (c)
G0D100461	SSAO5-02-1BPC SSAO5-03-1BPC** SSAO4-02-3BPC	OCDD OCDF	J (all detects) UJ (all non-detects) J (all detects) UJ (all non-detects)	P	Internal standards (%R) (i)
G0D100461	SSAO5-02-1BPC SSAO5-03-1BPC** SSAO5-02-1BPC_FD SSAO5-03-1BPC_FD SSAO4-02-3BPC	All compounds reported below the PQL.	J (all detects)	A	Project Quantitation Limit (sp)
G0D100461	SSAO5-02-1BPC SSAO5-03-1BPC** SSAO5-02-1BPC_FD SSAO5-03-1BPC_FD SSAO4-02-3BPC	All compounds reported as EMPC	JK (all detects)	A	Project Quantitation Limit (k)

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Laboratory Blank Data Qualification Summary - SDG
G0D100461**

No Sample Data Qualified in this SDG

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Equipment Blank Data Qualification Summary - SDG
G0D100461**

No Sample Data Qualified in this SDG

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Field Blank Data Qualification Summary - SDG G0D100461**

No Sample Data Qualified in this SDG

Tronox Northgate Henderson

VALIDATION COMPLETENESS WORKSHEET

LDC #: 23188A21
 SDG #: GOD100461
 Laboratory: Test America

Stage 2B

Date: 4/8/10
 Page: 1 of 1
 Reviewer: [Signature]
 2nd Reviewer: [Signature]

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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.	Technical holding times	A	Sampling dates: 4/8/10
II.	HRGC/HRMS Instrument performance check	A	
III.	Initial calibration	A	
IV.	Routine calibration/ICV	SW	
V.	Blanks	SW	
VI.	Matrix spike/Matrix spike duplicates	SW	No sp added - No anal
VII.	Laboratory control samples	A	ICV
VIII.	Regional quality assurance and quality control	N	
IX.	Internal standards	SW	
X.	Target compound identifications	N	
XI.	Compound quantitation and CRQLs	SW	
XII.	System performance	N	
XIII.	Overall assessment of data	A	
XIV.	Field duplicates	SW	D = 1 + 2, 2 + 4
XV.	Field blanks	SW	FB-040710-R2C. EB-040710-R2C (EOD B0519)

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

Validated Samples: * Level 1V

1	SSAO5-02-1BPC	5	11	0103408MB	21	31
2	SSAO5-03-1BPC **		12	0105361MB	22	32
3	SSAO5-02-1BPC_FD		13		23	33
4	SSAO5-03-1BPC_FDD		14		24	34
5	SSAO4-02-3BPC		15		25	35
6			16		26	36
7			17		27	37
8			18		28	38
9			19		29	39
10			20		30	40

Notes: _____

VALIDATION FINDINGS CHECKLIST

Method: Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

Validation Area	Yes	No	NA	Findings/Comments
I. Technical holding times				
All technical holding times were met.	/			
Cooler temperature criteria was met.	/			
II. GC/MS Instrument performance check				
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?	/			
III. Initial calibration				
Was the initial calibration performed at 5 concentration levels?	/			
Were all percent relative standard deviations (%RSD) $\leq 20\%$ for unlabeled standards and $< 30\%$ for labeled standards?	/			
Did all calibration standards meet the Ion Abundance Ratio criteria?	/			
Was the signal to noise ratio for each target compound ≥ 2.5 and for each recovery and internal standard ≥ 10 ?	/			
IV. Continuing calibration				
Was a routine calibration performed at the beginning and end of each 12 hour period?	/			
Were all percent differences (%D) $\leq 20\%$ for unlabeled standards and $\leq 30\%$ for labeled standards?	/			
Did all routine calibration standards meet the Ion Abundance Ratio criteria?	/			
V. Blanks				
Was a method blank associated with every sample in this SDG?	/			
Was a method blank performed for each matrix and concentration?	/			
Was there contamination in the method blanks? If yes, please see the Blanks validation completeness worksheet?	/			
VI. Matrix spike/Matrix spike duplicates				
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.	/	0		
Were the MS/MSD percent recoveries (%R) and the relative percent differences (RPD) within the QC limits?	/			
VII. Laboratory control samples				
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?	/			

LDC #: 231881/1
 SDG #: see cover

VALIDATION FINDINGS CHECKLIST

Page: 2 of 2
 Reviewer: [Signature]
 2nd Reviewer: [Signature]

VIII. Regional Quality Assurance and Quality Control				
Were performance evaluation (PE) samples performed?			<input checked="" type="checkbox"/>	
Were the performance evaluation (PE) samples within the acceptance limits?			<input checked="" type="checkbox"/>	
IX. Internal standards				
Were internal standard recoveries within the 40-135% criteria?			<input checked="" type="checkbox"/>	
Was the minimum S/N ratio of all internal standard peaks > 10?	<input checked="" type="checkbox"/>			
X. Target compound identification				
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?	<input checked="" type="checkbox"/>			
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?	<input checked="" type="checkbox"/>			
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?	<input checked="" type="checkbox"/>			
Did compound spectra contain all characteristic ions listed in the table attached?	<input checked="" type="checkbox"/>			
Was the Ion Abundance Ratio for the two quantitation ions within criteria?	<input checked="" type="checkbox"/>			
Was the signal to noise ratio for each target compound and labeled standard > 2.5?	<input checked="" type="checkbox"/>			
Does the maximum intensity of each specified characteristic ion coincide within ± 2 seconds (includes labeled standards)?	<input checked="" type="checkbox"/>			
For PCDF identification, was any signal (S/N ≥ 2.5, at ± seconds RT) detected in the corresponding PCDPE channel?			<input checked="" type="checkbox"/>	
Was an acceptable lock mass recorded and monitored?	<input checked="" type="checkbox"/>			
XI. Compound quantitation/CRQLs				
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 CRQLs adjusted to reflect all sample dilutions and dry weight factors applicable to level IV validation?	<input checked="" type="checkbox"/>			
XII. System performance				
System performance was found to be acceptable.	<input checked="" type="checkbox"/>			
XIII. Overall assessment of data				
Overall assessment of data was found to be acceptable.	<input checked="" type="checkbox"/>			
XIV. Field duplicates				
Field duplicate pairs were identified in this SDG.	<input checked="" type="checkbox"/>			
Target compounds were detected in the field duplicates.	<input checked="" type="checkbox"/>			
XV. Field blanks				
Field blanks were identified in this SDG.	<input checked="" type="checkbox"/>			
Target compounds were detected in the field blanks.	<input checked="" type="checkbox"/>			

VALIDATION FINDINGS WORKSHEET

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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:

LDC #: 23188A
 SDG #: See Cover

VALIDATION FINDINGS WORKSHEET
Field Blanks

Page: 1 of 1
 Reviewer: [Signature]
 2nd Reviewer: [Signature]

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

NA N/A Were field blanks identified in this SDG?

Blank units: pg/L Associated sample units: pg/g

Sampling date: 4/8/10

Field blank type: (circle one) Field Blank / Rinsate / Other: EB Associated Samples: M (> 5x)

Compound	Blank ID	5X	Sample Identification
	ER-04072010-RZC		
A	1.0	0.005	
D	0.69	0.00345	
E	0.65	0.00325	
F	5.5	0.0275	
G	53	0.265	
H	2.6	0.013	
I	1.5	0.0075	
J	1.0	0.005	
K	1.8	0.009	
L	1.1	0.0055	
M	0.97	0.00485	
O	4.5	0.0225	
P	1.1	0.0055	
Q	12	0.06	
CRQL			

CIRCLED RESULTS WERE NOT QUALIFIED. ALL RESULTS NOT CIRCLED WERE QUALIFIED BY THE FOLLOWING STATEMENT:
 Samples with compound concentrations within five times the associated field blank concentration are listed above, these sample results were qualified as not detected, "U".

LDC #: 23188A
 SDG #: See Cover

VALIDATION FINDINGS WORKSHEET

Field Blanks

Page: 1 of 1
 Reviewer: [Signature]
 2nd Reviewer: [Signature]

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

N/A Were field blanks identified in this SDG?

Blank units: pg/L Associated sample units: pg/g

Sampling date: 4/8/10

Field blank type: (circle one) Field Blank Rinsate / Other: Associated Samples: ML (>5x)

Compound	Blank ID	Blank ID	Sample Identification
	EB-04072010-RZC	5X	
C	0.77	0.00385	
D	0.74	0.0037	
E	0.82	0.0041	
F	4.2	0.021	
G	37	0.185	
H	0.57	0.00285	
I	0.96	0.0048	
J	0.67	0.00335	
K	1.1	0.0055	
L	0.96	0.0048	
M	1.0	0.005	
N	1.0	0.005	
O	2.1	0.0105	
P	1.5	0.0075	
Q	6.7	0.0335	
CRQL			

CIRCLED RESULTS WERE NOT QUALIFIED. ALL RESULTS NOT CIRCLED WERE QUALIFIED BY THE FOLLOWING STATEMENT:
 Samples with compound concentrations within five times the associated field blank concentration are listed above, these sample results were qualified as not detected, "J".

LDC#: 23188A21
 SDG#: See Cover

VALIDATION FINDINGS WORKSHEET
Field Duplicates

Page: 1 of 2
 Reviewer: Q
 2nd Reviewer: f

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

Y N NA
Y N NA

Were field duplicate pairs identified in this SDG?
 Were target analytes detected in the field duplicate pairs?

Compound	Concentration (pg/g)		(<=50) RPD	(pg/g) Difference	(pg/g) Limits	Qualifications (Parent Only)
	1	3				
A	2.5	2.4		0.1	(<=0.53)	
B	7.7	6.8		0.9	(<=2.6)	
C	4.7	4.4		0.3	(<=2.6)	
D	10	8.7		1.3	(<=2.6)	
E	9.2	8.4		0.8	(<=2.6)	
F	33	30	10			
G	38	40	5			
H	75	71	5			
I	130	120	8			
J	69	66	4			
K	250	240	4			
L	120	120	0			
M	32	28	13			
N	31	29	7			
O	530	490	8			
P	290	260	11			
Q	1900	1800	5			

Compound	Concentration (pg/g)		(<=50) RPD	(pg/g) Difference	(pg/g) Limits	Qualifications (Parent Only)
	2	4				
A	31	33		2	(<=11)	
B	110	110		0	(<=53)	
C	66	84		18	(<=53)	
D	150	150		0	(<=53)	
E	130	150		20	(<=53)	
F	530	630	17			
G	560	680	19			

LDC#: 23188A21

SDG#: See Cover

VALIDATION FINDINGS WORKSHEET

Field Duplicates

Page: 2 of 2

Reviewer: *[Signature]*

2nd Reviewer: *[Signature]*

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

Y N NA

Were field duplicate pairs identified in this SDG?

Y N NA

Were target analytes detected in the field duplicate pairs?

Compound	Concentration (pg/g)		(≤50)	(pg/g)	(pg/g)	Qualifications (Parent Only)
	2	4	RPD	Difference	Limits	
H	730	780	7			
I	1500	1600	6			
J	800	820	2			
K	3400	3600	6			
L	1700	1800	6			
M	310	350	12			
N	350	370	6			
O	7600	8200	8			
P	3600	4000	11			
Q	24000	28000	15			

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VALIDATION FINDINGS WORKSHEET
Routine Calibration

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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

N N/A Was a routine calibration performed at the beginning and end of each 12 hour period?

Y N/A Were all percent differences (%D) of RRFs ≤ 20% for unlabeled compounds and ≤ 30% for labeled?

Y N/A Did all routine calibration standards meet the Ion Abundance Ratio criteria?

#	Date	Standard ID	Compound	Finding %D (Limit: ≤30.0%)	Finding Ion Abundance Ratio	Associated Samples	Qualifications (C)	PCDDs		PCDFs	
								Selected Ions (m/z)	Ion Abundance Ratio	Selected Ions (m/z)	Ion Abundance Ratio
	4/24/10	19AP10405.40	13C-K	30.5		2.4	1+detz/p (K-N)	M/M+2	0.65-0.89	M/M+2	0.65-0.89
								M+2/M+4	1.32-1.78	M+2/M+4	1.32-1.78
								M+2/M+4	1.05-1.43	M+2/M+4	1.05-1.43
								M/M+2	0.43-0.59	M/M+2	0.43-0.59
								M/M+2	0.37-0.51	M/M+2	0.37-0.51
								M+2/M+4	0.88-1.20	M+2/M+4	0.88-1.20
								M+2/M+4	0.76-1.02	M+2/M+4	0.76-1.02

VALIDATION FINDINGS WORKSHEET

Blanks

METHOD: HRGC/HRMS Dioxins (EPA Method 8290)

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? If yes, please see qualification below.

Blank extraction date: 4/13/10 **Blank analysis date:** 4/20/10

Conc. units: pg/g **Associated samples:** 1-4 (> 5x)

Compound	Blank ID	Sample Identification
	0103408MB	5X
E	0.040	0.2
F	0.13	0.65
G	0.71	3.55
H	0.033	0.165
K	0.048	0.24
L	0.033	0.165
O	0.13	0.65
P	0.068	0.34
Q	0.32	1.6

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
Internal Standards

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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

Y/N/N/A Are all internal standard recoveries within the 40-135% criteria?

Y/N/N/A Was the S/N ratio all internal standard peaks \geq 10?

#	Date	Lab ID/Reference	Internal Standard	% Recovery (Limit: 40-135%)	Qualifications (if)
		1	I	31	Y/N/A (S.A.)
		2	J	38	
		5	I	36	

	Internal Standards	Check Standard Used	Internal Standards	Check Standard Used
A.	^{13}C -2,3,7,8-TCDF		G.	^{13}C -1,2,3,4,6,7,8-HpCDF
B.	^{13}C -2,3,7,8-TCDD		H.	^{13}C -1,2,3,4,6,7,8-HpCDD
C.	^{13}C -1,2,3,7,8-PeCDF		I.	^{13}C -OCDD
D.	^{13}C -1,2,3,7,8-PeCDD		K.	^{13}C -1,2,3,4-TCDD
E.	^{13}C -1,2,3,4,7,8-HxCDF		L.	^{13}C -1,2,3,7,8,9-HxCDD
F.	^{13}C -1,2,3,6,7,8-HxCDD			

LDC #: 23/88A-1
SDG #: See above

VALIDATION FINDINGS WORKSHEET
Compound Quantitation and Reported CRQLs

Page: 1 of 2
Reviewer: [Signature]
2nd Reviewer: [Signature]

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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

N N/A
Y N N/A

Were the correct internal standard (IS), quantitation ions and relative response factors (RRF) used to quantitate the compound?
Compound quantitation and CRQLs were adjusted to reflect all sample dilutions and dry weight factors (if necessary).

#	Date	Sample ID	Finding	Associated Samples	Qualifications
		M	ZAD results (2 bags)		Y (C F)

Comments: See sample calculation verification worksheet for recalculations

VALIDATION FINDINGS WORKSHEET
Initial Calibration Calculation Verification

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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

$RRF = (A_s)(C_s)/(A_c)(C_c)$
 average RRF = sum of the RRFs/number of standards
 $\%RSD = 100 * (S/X)$

A_s = Area of compound,
 C_s = Concentration of compound,
 S = Standard deviation of the RRFs,
 A_c = Area of associated internal standard
 C_c = Concentration of internal standard
 X = Mean of the RRFs

#	Standard ID	Calibration Date	Compound (Reference Internal Standard)	Reported		Recalculated		Reported		Recalculated	
				Average RRF (initial)	RRF (CS3 std)	Average RRF (initial)	RRF (CS3 std)	%RSD	%RSD	RRF (CS3 std)	%RSD
1	1CAZ (ADS)	4/12/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	0.945	0.98	0.945	0.98	4.44	4.33		
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)	1.021	1.04	1.021	1.04	3.03	2.97		
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)	1.114	1.19	1.114	1.19	5.33	5.25		
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)	1.072	1.11	1.072	1.11	3.60	3.75		
			OCDF (¹³ C-OCDD)	1.445	1.51	1.445	1.51	5.85	5.89		
2	1CAZ	3/10/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	1.000	1.00	1.000	1.00	1.36	1.58		
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)								
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)								
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)								
			OCDF (¹³ C-OCDD)								
3	1CAZ (SD2)	4/21/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	1.088	1.10	1.088	1.10	1.29	1.20		
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)								
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)								
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)								
			OCDF (¹³ 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
Routine Calibration Results Verification

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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 = $100 * (\text{ave. RRF} - \text{RRF}) / \text{ave. RRF}$ Where: ave. RRF = initial calibration average RRF
 RRF = $(A_x)(C_s) / (A_s)(C_x)$ RRF = continuing calibration RRF
 A_x = Area of compound, A_s = Area of associated internal standard
 C_x = Concentration of compound, C_s = Concentration of internal standard

#	Standard ID	Calibration Date	Compound (Reference Internal Standard)	Average RRF (Initial)	Reported		Recalculated	
					RRF (CC)	%D	RRF (CC)	%D
1	19AP10405	4/19/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	0.945	1.00	6.1	1.00	6.1
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)	1.021	1.02	0.2	1.02	0.2
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)	1.114	1.19	6.8	1.19	6.8
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)	1.072	1.07	0.1	1.07	0.1
			OCDF (¹³ C-OCDD)	1.445	1.51	4.8	1.51	4.8
2	19AP10405	4/20/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	0.945	1.06	12.5	1.06	12.5
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)	1.021	1.06	3.9	1.06	3.9
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)	1.114	1.19	7.2	1.19	7.2
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)	1.072	1.12	4.4	1.12	4.4
			OCDF (¹³ C-OCDD)	1.445	1.57	8.3	1.57	8.3
3	20AP10502	4/20/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	1.000	1.05	4.8	1.05	4.8
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)					
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)					
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)					
			OCDF (¹³ C-OCDD)					

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 #: 23188A21
 SDG #: Bel caner

VALIDATION FINDINGS WORKSHEET
Laboratory Control Sample Results Verification

Page: 10 of 10
 Reviewer: _____
 2nd Reviewer: _____

METHOD: GC/MS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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 = $100 \cdot \text{SSC}/\text{SA}$

Where: SSC = Spiked sample concentration
 SA = Spike added

RPD = $100 \cdot \frac{\text{LCS} - \text{LCSD}}{\text{LCS} + \text{LCSD}}$

LCS = Laboratory control sample percent recovery

LCSD = Laboratory control sample duplicate percent recovery

LCS ID: 0103408

Compound	Spike Added (PP5)		Spiked Sample Concentration (PP5)		LCS		LCSD		Percent Recovery		Percent Recovery		RPD	
	LCS	LCSD	LCS	LCSD	Reported	Recalc.	Reported	Recalc.	Reported	Recalc.	Reported	Recalc.	Reported	Recalc.
2,3,7,8-TCDD	20.0	NA	22.0	NA	110	110								
1,2,3,7,8-PeCDD	100		108		108	108								
1,2,3,4,7,8-HxCDD			116		116	116								
1,2,3,4,7,8,9-HpCDF			118		118	118								
OCDF	200		225		113	113								

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.

Descriptor	Accurate mass ^(a)	Ion ID	Elemental Composition	Analyte	Descriptor	Accurate Mass ^(a)	Ion ID	Elemental Composition	Analyte
1	303.9016 305.8987 315.9419 317.9389 319.8965 321.8936 331.9368 333.9338 375.8964 [354.9792]	M M+2 M M+2 M M+2 M M+2 M+2 LOCK	C ₁₂ H ₄ ³⁵ Cl ₄ O C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ ClO C ₁₂ H ₄ ³⁵ Cl ₃ O C ₁₂ H ₄ ³⁵ Cl ₂ ³⁷ ClO C ₁₂ H ₄ ³⁵ Cl ₂ O C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₄ ³⁵ Cl ₂ ³⁷ ClO ₂ C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₄ ³⁵ Cl ₂ ³⁷ ClO ₂ C ₉ F ₁₃	TCDF TCDF TCDF (S) TCDF (S) TCDD TCDD TCDD (S) TCDD (S) HxCDPE PFK	4	407.7818 409.7788 417.8250 419.8220 423.7767 425.7737 435.8169 437.8140 479.7165 [430.9728]	M+2 M+4 M M+2 M+2 M+4 M+2 M+4 M+4 LOCK	C ₁₂ H ₃ ³⁵ Cl ₄ ³⁷ ClO C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ Cl ₂ O C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO C ₁₂ H ₃ ³⁵ Cl ₂ ³⁷ ClO C ₁₂ H ₃ ³⁵ Cl ₂ O C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₃ ³⁵ Cl ₂ ³⁷ ClO ₂ C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₃ ³⁵ Cl ₂ ³⁷ ClO ₂ C ₉ F ₁₃	OCDF OCDF OCDD OCDD OCDD (S) OCDD (S) DCDPE PFK
2	339.8597 341.8567 351.9000 353.8970 355.8546 357.8516 367.8949 369.8919 409.7974 [354.9792]	M+2 M+4 M+2 M+4 M+2 M+4 M+2 M+4 M+2 LOCK	C ₁₂ H ₂ ³⁵ Cl ₅ ³⁷ ClO C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ Cl ₂ O C ₁₂ H ₂ ³⁵ Cl ₄ O C ₁₂ H ₂ ³⁵ Cl ₃ ³⁷ ClO C ₁₂ H ₂ ³⁵ Cl ₃ O C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ ClO ₂ C ₁₂ H ₂ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ ClO ₂ C ₁₂ H ₂ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₉ F ₁₇	HxCDF HxCDF HxCDF (S) HxCDF (S) HxCDD HxCDD HxCDD (S) HxCDD (S) OCDDPE PFK	5	441.7428 443.7399 457.7377 459.7348 469.7780 471.7750 513.6775 [422.9278]	M+2 M+4 M+2 M+4 M+2 M+4 M+4 M+4 LOCK	C ₁₂ Cl ₇ ³⁷ ClO C ₁₂ Cl ₆ ³⁷ Cl ₂ O C ₁₂ Cl ₆ ³⁷ ClO ₂ C ₁₂ Cl ₆ ³⁷ ClO ₂ C ₁₂ Cl ₆ ³⁷ ClO ₂ C ₁₂ Cl ₆ ³⁷ ClO ₂ C ₁₂ Cl ₆ ³⁷ ClO ₂ C ₁₂ Cl ₆ ³⁷ ClO ₂ C ₁₂ Cl ₆ ³⁷ ClO ₂ C ₁₀ F ₁₇	OCDF OCDF OCDD OCDD OCDD (S) OCDD (S) DCDPE PFK
3	373.8208 375.8178 383.8639 385.8610 389.8156 391.8127 401.8559 403.8529 445.7555 [430.9728]	M+2 M+4 M M+2 M+2 M+4 M+2 M+4 M+4 LOCK	C ₁₂ H ₂ ³⁵ Cl ₅ ³⁷ ClO C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ Cl ₂ O C ₁₂ H ₂ ³⁵ Cl ₄ O C ₁₂ H ₂ ³⁵ Cl ₃ ³⁷ ClO C ₁₂ H ₂ ³⁵ Cl ₃ O C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ ClO ₂ C ₁₂ H ₂ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ ClO ₂ C ₁₂ H ₂ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₉ F ₁₇	HxCDF HxCDF HxCDF (S) HxCDF (S) HxCDD HxCDD HxCDD (S) HxCDD (S) OCDDPE PFK			M+2 M+4 M M+2 M+2 M+4 M+2 M+4 M+4 LOCK	C ₁₂ H ₂ ³⁵ Cl ₅ ³⁷ ClO C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ Cl ₂ O C ₁₂ H ₂ ³⁵ Cl ₄ O C ₁₂ H ₂ ³⁵ Cl ₃ ³⁷ ClO C ₁₂ H ₂ ³⁵ Cl ₃ O C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ ClO ₂ C ₁₂ H ₂ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ ClO ₂ C ₁₂ H ₂ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₉ F ₁₇	HxCDF HxCDF HxCDF (S) HxCDF (S) HxCDD HxCDD HxCDD (S) HxCDD (S) OCDDPE PFK

(a) The following nucleidic masses were used:

H = 1.007825
C = 12.000000
¹³C = 13.003355
F = 18.9984
O = 15.994915
³⁵Cl = 34.968853
³⁷Cl = 36.965903

S = internal/recovery standard

VALIDATION FINDINGS WORKSHEET

Sample Calculation Verification

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

- 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 agree within 10.0% of the reported results?

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

- A_x = Area of the characteristic ion (EICP) for the compound to be measured
- A_s = 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).
- RRF = Relative Response Factor (average) from the initial calibration
- Df = Dilution Factor.
- %S = Percent solids, applicable to soil and solid matrices only.

Example:

Sample I.D. 2, A:

$$\text{Conc.} = \frac{(1302240)(2000)}{(8555920)(1.02)(10.53)(0.92)}$$

= 30.8 ps/g

#	Sample ID	Compound	Reported Concentration ()	Calculated Concentration ()	Qualification

Laboratory Data Consultants, Inc. Data Validation Report

Project/Site Name: Tronox LLC Facility, PCS, Henderson, Nevada
Collection Date: April 9, 2010
LDC Report Date: May 21, 2010
Matrix: Soil
Parameters: Dioxins/Dibenzofurans
Validation Level: Stage 2B & 4
Laboratory: TestAmerica, Inc.

Sample Delivery Group (SDG): G0D130435

Sample Identification

SSAM6-01-1BPC
SA60-3BPC
SSAN6-04-1BPC
SSAN6-04-1BPC_FD
SSAN5-01-1BPC
SSAI2-02-2BPC**
SSAJ3-01-1BPC
SSAO6-04-1BPC_FD
SSAO6-04-1BPC
SSAO7-01-1BPC
SSAO7-01-1BPC_FD
SSAM6-01-1BPCMS
SSAM6-01-1BPCMSD

**Indicates sample underwent Stage 4 review

Introduction

This data review covers 13 soil samples listed on the cover sheet including dilutions and reanalysis as applicable. The analyses were per EPA SW 846 Method 8290 for Polychlorinated Dioxins/Dibenzofurans.

This review follows the Standard Operating Procedures (SOP) 40, Data Review/Validation (BRC 2009), the Quality Assurance Project Plan Tronox LLC Facility, Henderson, Nevada (June 2009), NDEP guidance (May 2006), and USEPA Contract Laboratory Program National Functional Guidelines for Polychlorinated Dioxins/Dibenzofurans Data Review (September 2005).

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.

Blank results are summarized in Section V.

Field duplicates are summarized in Section XIV.

Samples indicated by a double asterisk on the front cover underwent a Stage 4 review. A Stage 2B review was performed on all of the other samples. Raw data were not evaluated for the samples reviewed by Stage 2B criteria since this review is based on QC data.

The following are definitions of the data qualifiers:

- J+ Data are qualified as estimated, with a high bias likely to occur. False positives or false negatives are unlikely to have been reported.
- J- Data are qualified as estimated, with a low bias likely to occur. False positives or false negatives are unlikely to have been reported.
- J Data are qualified as estimated; it is not possible to assess the direction of the potential bias. False positives or false negatives are unlikely to have been reported.
- U Indicates the compound or analyte was analyzed for but not detected at or above the stated limit.
- R Data are qualified as rejected. There is a significant potential for the reporting of false negatives or false positives.
- UU Indicates the compound or analyte was analyzed for but not detected. The sample detection limit is an estimated value.
- B The analytical result may be a false positive totally attributable to blank contamination. This qualifier is applicable to radiochemistry analysis only.
- JB The analytical result may be biased high and partially attributable to blank contamination. This qualifier is applicable to radiochemistry analysis only.
- JK The analytical result is an estimated maximum possible concentration (EMPC).
- X The analytical result is not used for reporting because a more accurate and precise result is reported in its place.
- J-TDS The analytical result is estimated based on failure of the Total Dissolved Solids (TDS) correctness check performed in accordance with the Standard Method 1030E.
- J-CAB The analytical result is estimated based on failure of the cation-anion balance correctness check performed in accordance with Standard Method 1030E.
- J-TDS & CAB The analytical result is unreliable based on the failure of the cation-anion balance and TDS correctness check performed in accordance with standard Method 1030E.
- A Indicates the finding is based upon technical validation criteria.
- P Indicates the finding is related to a protocol/contractual deviation.
- None Indicates the data was not significantly impacted by the finding, therefore qualification was not required.

I. Technical Holding Times

All technical holding time requirements were met.

The chain-of-custodies were reviewed for documentation of cooler temperatures. All cooler temperatures met validation criteria.

II. HRGC/HRMS Instrument Performance Check

Instrument performance was checked at the required daily 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 exact mass of 380.9760 of PFK was verified. The static resolving power was at least 10,000 (10% valley definition) for samples on which Stage 4 review was performed. Raw data were not evaluated for the samples reviewed by Stage 2B criteria.

III. Initial Calibration

A five point initial calibration was performed as required by the method.

Percent relative standard deviations (%RSD) were less than or equal to 20.0% for unlabeled compounds and less than or equal to 30.0% for labeled compounds.

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

The minimum S/N ratio for each target compound was greater than or equal to 2.5 and greater than or equal to 10 for each recovery and internal standard compound for samples on which Stage 4 review was performed. Raw data were not evaluated for the samples reviewed by Stage 2B criteria.

IV. Routine Calibration (Continuing)

Routine calibration was performed at the required frequencies.

All of the routine calibration percent differences (%D) between the initial calibration RRF and the routine calibration RRF were less than or equal to 20.0% for unlabeled compounds and less than or equal to 30.0% for labeled compounds with the following exceptions:

Date	Compound	%D	Associated Samples	Affected Compound	Flag	A or P
4/28/10	¹³ C-1,2,3,6,7,8-HxCDD	30.3	SSAO6-04-1BPC_FD SSAO6-04-1BPC SSAO7-01-1BPC	1,2,3,4,7,8-HxCDD 1,2,3,6,7,8-HxCDD 1,2,3,7,8,9-HxCDD	J+ (all detects) J+ (all detects) J+ (all detects)	P

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

V. Blanks

Method blanks were reviewed for each matrix as applicable. No polychlorinated dioxin/dibenzofuran contaminants were found in the method blanks with the following exceptions:

Method Blank ID	Extraction Date	Compound	Concentration	Associated Samples
0104236MB	4/14/10	1,2,3,6,7,8-HxCDD OCDD 1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 2,3,4,6,7,8-HxCDF 1,2,3,7,8,9-HxCDF 1,2,3,4,6,7,8-HpCDF OCDF	0.11 pg/g 0.68 pg/g 0.19 pg/g 0.13 pg/g 0.098 pg/g 0.13 pg/g 0.15 pg/g 0.41 pg/g	SSAM6-01-1BPC SA60-3BPC SSAN6-04-1BPC SSAN6-04-1BPC_FD SSAN5-01-1BPC SSAI2-02-2BPC** SSAJ3-01-1BPC

Sample concentrations were compared to concentrations detected in the method blanks as required by the QAPP. No sample data was qualified with the following exceptions:

Sample	Compound	Reported Concentration	Modified Final Concentration
SSAN6-04-1BPC	1,2,3,6,7,8-HxCDD 2,3,4,6,7,8-HxCDF 1,2,3,7,8,9-HxCDF	0.22 pg/g 0.27 pg/g 0.34 pg/g	0.22U pg/g 0.27U pg/g 0.34U pg/g
SSAN6-04-1BPC_FD	1,2,3,6,7,8-HxCDD 2,3,4,6,7,8-HxCDF 1,2,3,7,8,9-HxCDF	0.19 pg/g 0.22 pg/g 0.19 pg/g	0.19U pg/g 0.22U pg/g 0.19U pg/g

Samples FB-04072010-RZD (from SDG G0D090441) and FB-04072010-RZC (from SDG G0D130519) were identified as field blanks. No polychlorinated dioxin/dibenzofuran contaminants were found in these blanks with the following exceptions:

Field Blank ID	Sampling Date	Compound	Concentration	Associated Samples
FB-04072010-RZD	4/7/10	1,2,3,4,7,8-HxCDD 1,2,3,7,8,9-HxCDD 1,2,3,4,6,7,8-HpCDD OCDD 1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 2,3,4,6,7,8-HxCDF 1,2,3,7,8,9-HxCDF 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	0.89 pg/L 1.5 pg/L 2.2 pg/L 8.3 pg/L 1.4 pg/L 1.6 pg/L 1.5 pg/L 1.6 pg/L 1.3 pg/L 1.4 pg/L 4.1 pg/L	SSAI2-02-2BPC** SSAJ3-01-1BPC

Field Blank ID	Sampling Date	Compound	Concentration	Associated Samples
FB-04072010-RZC	4/8/10	1,2,3,4,7,8-HxCDD	0.77 pg/L	SSAM6-01-1BPC
		1,2,3,6,7,8-HxCDD	0.74 pg/L	SA60-3BPC
		1,2,3,7,8,9-HxCDD	0.82 pg/L	SSAN6-04-1BPC
		1,2,3,4,6,7,8-HpCDD	4.2 pg/L	SSAN6-04-1BPC_FD
		OCDD	37 pg/L	SSAN5-01-1BPC
		2,3,7,8-TCDF	0.57 pg/L	SSAO6-04-1BPC_FD
		1,2,3,7,8-PeCDF	0.96 pg/L	SSAO6-04-1BPC
		2,3,4,7,8-PeCDF	0.67 pg/L	SSAO7-01-1BPC
		1,2,3,4,7,8-HxCDF	1.1 pg/L	SSAO7-01-1BPC_FD
		1,2,3,6,7,8-HxCDF	0.96 pg/L	
		2,3,4,6,7,8-HxCDF	1.0 pg/L	
		1,2,3,7,8,9-HxCDF	1.0 pg/L	
		1,2,3,4,6,7,8-HpCDF	2.1 pg/L	
		1,2,3,4,7,8,9-HpCDF	1.5 pg/L	
		OCDF	6.7 pg/L	

Sample concentrations were compared to concentrations detected in the field blanks as required by the QAPP. No sample data was qualified.

VI. Matrix Spike/Matrix Spike Duplicates

Matrix spike (MS) and matrix spike duplicate (MSD) samples were reviewed for each matrix as applicable. Although the MS/MSD percent recoveries (%R) were not within QC limits for several compounds, the MS or LCS percent recoveries (%R) were within QC limits and no data were qualified.

VII. Laboratory Control Samples (LCS)

Laboratory control samples were reviewed for each matrix as applicable. Percent recoveries (%R) were within QC limits.

VIII. Regional Quality Assurance and Quality Control

Not applicable.

IX. Internal Standards

All internal standard recoveries were within QC limits with the following exceptions:

Sample	Internal Standards	%R (Limits)	Compound	Flag	A or P
SSAM6-01-1BPC	¹³ C-OCDD	38 (40-135)	OCDD	J (all detects) UJ (all non-detects)	P
			OCDF	J (all detects) UJ (all non-detects)	

Sample	Internal Standards	%R (Limits)	Compound	Flag	A or P
SSAN6-04-1BPC	¹³ C-1,2,3,4,6,7,8-HpCDF ¹³ C-1,2,3,4,6,7,8-HpCDD ¹³ C-OCDD	23 (40-135) 27 (40-135) 15 (40-135)	1,2,3,4,6,7,8-HpCDD OCDD 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	J (all detects) UJ (all non-detects)	P
SSAN6-04-1BPC_FD	¹³ C-1,2,3,4,6,7,8-HpCDF ¹³ C-OCDD	36 (40-135) 27 (40-135)	OCDD 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	J (all detects) UJ (all non-detects)	P
SSAI2-02-2BPC**	¹³ C-1,2,3,4,6,7,8-HpCDF ¹³ C-1,2,3,4,6,7,8-HpCDD ¹³ C-OCDD	33 (40-135) 37 (40-135) 20 (40-135)	1,2,3,4,6,7,8-HpCDD OCDD 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	J (all detects) UJ (all non-detects)	P
SSAJ3-01-1BPC	¹³ C-1,2,3,4,6,7,8-HpCDF ¹³ C-OCDD	38 (40-135) 25 (40-135)	OCDD 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	J (all detects) UJ (all non-detects)	P
SSAO7-01-1BPC	¹³ C-OCDD	27 (40-135)	OCDD OCDF	J (all detects) UJ (all non-detects) J (all detects) UJ (all non-detects)	P
SSAO7-01-1BPC_FD	¹³ C-OCDD	22 (40-135)	OCDD OCDF	J (all detects) UJ (all non-detects) J (all detects) UJ (all non-detects)	P

X. Target Compound Identifications

All target compound identifications were within validation criteria for samples on which a Stage 4 review was performed. Raw data were not evaluated for the samples reviewed by Stage 2B criteria.

XI. Project Quantitation Limit

All compound quantitation and CRQLs were within validation criteria with the following exceptions:

Sample	Compound	Finding	Criteria	Flag	A or P
SSAM6-01-1BPC SA60-3BPC	2,3,7,8-TCDF 1,2,3,7,8-PeCDF 1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	Sample result exceeded calibration range.	Reported result should be within calibration range.	J (all detects) J (all detects) J (all detects) J (all detects) J (all detects) J (all detects) J (all detects)	P
SSAI2-02-2BPC**	2,3,7,8-TCDF 1,2,3,7,8-PeCDF 2,3,4,7,8-PeCDF 1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	Sample result exceeded calibration range.	Reported result should be within calibration range.	J (all detects) J (all detects) J (all detects) J (all detects) J (all detects) J (all detects) J (all detects) J (all detects)	P
SSAJ3-01-1BPC	1,2,3,4,6,7,8-HpCDF OCDF	Sample result exceeded calibration range.	Reported result should be within calibration range.	J (all detects) J (all detects)	P

All compounds reported below the PQL were qualified as follows:

Sample	Finding	Flag	A or P
All samples in SDG G0D130435	All compounds reported below the PQL.	J (all detects)	A

All compounds reported as EMPC were qualified as follows:

Sample	Finding	Flag	A or P
All samples in SDG G0D130435	All compounds reported as estimated maximum possible concentration (EMPC).	JK (all detects)	A

Raw data were not evaluated for the samples reviewed by Stage 2B criteria.

XII. System Performance

The system performance was acceptable for samples on which a Stage 4 review was performed. Raw data were not evaluated for the samples reviewed by Stage 2B criteria.

XIII. Overall Assessment of Data

Data flags are summarized at the end of this report if data has been qualified.

XIV. Field Duplicates

Samples SSAN6-04-1BPC and SSAN6-04-1BPC_FD, samples SSAO6-04-1BPC_FD and SSAO6-04-1BPC, and samples SSAO7-01-1BPC and SSAO7-01-1BPC_FD were identified as field duplicates. No polychlorinated dioxins/dibenzofurans were detected in any of the samples with the following exceptions:

Compound	Concentration (pg/g)		RPD (Limits)	Difference (Limits)	Flags	A or P
	SSAN6-04-1BPC	SSAN6-04-1BPC_FD				
2,3,7,8-TCDD	0.50U	0.041	-	0.459 (≤ 0.50)	-	-
1,2,3,4,7,8-HxCDD	2.5U	0.081	-	2.419 (≤ 2.5)	-	-
1,2,3,6,7,8-HxCDD	0.22	0.19	-	0.03 (≤ 2.5)	-	-
1,2,3,7,8,9-HxCDD	0.30	0.32	-	0.02 (≤ 2.5)	-	-
1,2,3,4,6,7,8-HpCDD	0.80	1.6	-	0.8 (≤ 2.5)	-	-
OCDD	3.7	8.6	-	4.9 (≤ 5.0)	-	-
2,3,7,8-TCDF	0.45	0.46	-	0.01 (≤ 0.50)	-	-
1,2,3,7,8-PeCDF	0.70	0.73	-	0.03 (≤ 2.5)	-	-
2,3,4,7,8-PeCDF	0.45	0.45	-	0 (≤ 2.5)	-	-
1,2,3,4,7,8-HxCDF	1.3	1.5	-	0.2 (≤ 2.5)	-	-
1,2,3,6,7,8-HxCDF	0.68	0.91	-	0.23 (≤ 2.5)	-	-
2,3,4,6,7,8-HxCDF	0.27	0.22	-	0.05 (≤ 2.5)	-	-
1,2,3,7,8,9-HxCDF	0.34	0.19	-	0.15 (≤ 2.5)	-	-
1,2,3,4,6,7,8-HpCDF	2.9	2.8	-	0.1 (≤ 2.5)	-	-
1,2,3,4,7,8,9-HpCDF	1.5	1.6	-	0.1 (≤ 2.5)	-	-
OCDF	5.5	7.3	-	1.8 (≤ 5.0)	-	-

Compound	Concentration (pg/g)		RPD (Limits)	Difference (Limits)	Flags	A or P
	SSAO6-04-1BPC_FD	SSAO6-04-1BPC				
2,3,7,8-TCDD	0.86	0.80	-	0.06 (≤ 0.53)	-	-
1,2,3,7,8-PeCDD	2.1	2.0	-	0.1 (≤ 2.6)	-	-
1,2,3,4,7,8-HxCDD	1.8	1.6	-	0.2 (≤ 2.6)	-	-
1,2,3,6,7,8-HxCDD	3.2	3.1	-	0.1 (≤ 2.6)	-	-
1,2,3,7,8,9-HxCDD	2.8	2.6	-	0.2 (≤ 2.6)	-	-
1,2,3,4,6,7,8-HpCDD	18	12	40 (≤ 50)	-	-	-
OCDD	140	30	129 (≤ 50)	-	J (all detects)	A
2,3,7,8-TCDF	13	13	0 (≤ 50)	-	-	-
1,2,3,7,8-PeCDF	27	25	8 (≤ 50)	-	-	-
2,3,4,7,8-PeCDF	13	12	-	1 (≤ 2.6)	-	-
1,2,3,4,7,8-HxCDF	52	47	10 (≤ 50)	-	-	-
1,2,3,6,7,8-HxCDF	35	32	9 (≤ 50)	-	-	-
2,3,4,6,7,8-HxCDF	8.0	7.6	-	0.4 (≤ 2.6)	-	-
1,2,3,7,8,9-HxCDF	4.5	4.2	-	0.3 (≤ 2.6)	-	-
1,2,3,4,6,7,8-HpCDF	130	120	8 (≤ 50)	-	-	-
1,2,3,4,7,8,9-HpCDF	45	44	2 (≤ 50)	-	-	-
OCDF	300	250	18 (≤ 50)	-	-	-

Compound	Concentration (pg/g)		RPD (Limits)	Difference (Limits)	Flags	A or P
	SSAO7-01-1BPC	SSAO7-01-1BPC_FD				
2,3,7,8-TCDD	1.5	1.6	-	0.1 (≤ 0.52)	-	-

Compound	Concentration (pg/g)		RPD (Limits)	Difference (Limits)	Flags	A or P
	SSAO7-01-1BPC	SSAO7-01-1BPC_FD				
1,2,3,7,8-PeCDD	3.4	4.0	-	0.6 (≤ 2.6)	-	-
1,2,3,4,7,8-HxCDD	2.4	2.6	-	0.2 (≤ 2.6)	-	-
1,2,3,6,7,8-HxCDD	4.2	4.8	-	0.6 (≤ 2.6)	-	-
1,2,3,7,8,9-HxCDD	3.6	4.1	-	0.5 (≤ 2.6)	-	-
1,2,3,4,6,7,8-HpCDD	14	16	13 (≤ 50)	-	-	-
OCDD	22	22	0 (≤ 50)	-	-	-
2,3,7,8-TCDF	160	100	46 (≤ 50)	-	-	-
1,2,3,7,8-PeCDF	58	62	7 (≤ 50)	-	-	-
2,3,4,7,8-PeCDF	24	26	8 (≤ 50)	-	-	-
1,2,3,4,7,8-HxCDF	76	96	23 (≤ 50)	-	-	-
1,2,3,6,7,8-HxCDF	50	63	23 (≤ 50)	-	-	-
2,3,4,6,7,8-HxCDF	12	15	22 (≤ 50)	-	-	-
1,2,3,7,8,9-HxCDF	8.1	9.2	-	1.1 (≤ 2.6)	-	-
1,2,3,4,6,7,8-HpCDF	170	200	16 (≤ 50)	-	-	-
1,2,3,4,7,8,9-HpCDF	110	94	16 (≤ 50)	-	-	-
OCDF	480	940	65 (≤ 50)	-	J (all detects)	A

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Data Qualification Summary - SDG G0D130435**

SDG	Sample	Compound	Flag	A or P	Reason (Code)
G0D130435	SSAO6-04-1BPC_FD SSAO6-04-1BPC SSAO7-01-1BPC	1,2,3,4,7,8-HxCDD 1,2,3,6,7,8-HxCDD 1,2,3,7,8,9-HxCDD	J+ (all detects) J+ (all detects) J+ (all detects)	P	Routine calibration (%D) (c)
G0D130435	SSAM6-01-1BPC SSAO7-01-1BPC SSAO7-01-1BPC_FD	OCDD OCDF	J (all detects) UJ (all non-detects) J (all detects) UJ (all non-detects)	P	Internal standards (%R) (i)
G0D130435	SSAN6-04-1BPC SSAI2-02-2BPC**	1,2,3,4,6,7,8-HpCDD OCDD 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	J (all detects) UJ (all non-detects)	P	Internal standards (%R) (i)
G0D130435	SSAN6-04-1BPC_FD SSAJ3-01-1BPC	OCDD 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	J (all detects) UJ (all non-detects)	P	Internal standards (%R) (i)
G0D130435	SSAM6-01-1BPC SA60-3BPC	2,3,7,8-TCDF 1,2,3,7,8-PeCDF 1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	J (all detects) J (all detects) J (all detects) J (all detects) J (all detects) J (all detects) J (all detects)	P	Compound quantitation and CRQLs (e)
G0D130435	SSAI2-02-2BPC**	2,3,7,8-TCDF 1,2,3,7,8-PeCDF 2,3,4,7,8-PeCDF 1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	J (all detects) J (all detects) J (all detects) J (all detects) J (all detects) J (all detects) J (all detects) J (all detects)	P	Compound quantitation and CRQLs (e)
G0D130435	SSAJ3-01-1BPC	1,2,3,4,6,7,8-HpCDF OCDF	J (all detects) J (all detects)	P	Compound quantitation and CRQLs (e)
G0D130435	SSAM6-01-1BPC SA60-3BPC SSAN6-04-1BPC SSAN6-04-1BPC_FD SSAN5-01-1BPC SSAI2-02-2BPC** SSAJ3-01-1BPC SSAO6-04-1BPC_FD SSAO6-04-1BPC SSAO7-01-1BPC SSAO7-01-1BPC_FD	All compounds reported below the PQL.	J (all detects)	A	Project Quantitation Limit (sp)

SDG	Sample	Compound	Flag	A or P	Reason (Code)
G0D130435	SSAM6-01-1BPC SA60-3BPC SSAN6-04-1BPC SSAN6-04-1BPC_FD SSAN5-01-1BPC SSAI2-02-2BPC** SSAJ3-01-1BPC SSAO6-04-1BPC_FD SSAO6-04-1BPC SSAO7-01-1BPC SSAO7-01-1BPC_FD	All compounds reported as EMPC	JK (all detects)	A	Project Quantitation Limit (k)
G0D130435	SSAO6-04-1BPC_FD SSAO6-04-1BPC	OCDD	J (all detects)	A	Field duplicates (RPD) (fd)
G0D130435	SSAO7-01-1BPC SSAO7-01-1BPC_FD	OCDF	J (all detects)	A	Field duplicates (RPD) (fd)

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Laboratory Blank Data Qualification Summary - SDG
G0D130435**

SDG	Sample	Compound	Modified Final Concentration	A or P	Code
G0D130435	SSAN6-04-1BPC	1,2,3,6,7,8-HxCDD 2,3,4,6,7,8-HxCDF 1,2,3,7,8,9-HxCDF	0.22U pg/g 0.27U pg/g 0.34U pg/g	A	bl
G0D130435	SSAN6-04-1BPC_FD	1,2,3,6,7,8-HxCDD 2,3,4,6,7,8-HxCDF 1,2,3,7,8,9-HxCDF	0.19U pg/g 0.22U pg/g 0.19U pg/g	A	bl

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Field Blank Data Qualification Summary - SDG G0D130435**

No Sample Data Qualified in this SDG

Tronox Northgate Henderson

VALIDATION COMPLETENESS WORKSHEET

LDC #: 23188B21
 SDG #: G0D130435
 Laboratory: Test America

Stage 2B/4

Date: 5/20/10
 Page: 1 of 1
 Reviewer: [Signature]
 2nd Reviewer: _____

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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.	Technical holding times	A	Sampling dates: 4/9/10
II.	HRGC/HRMS Instrument performance check	A	
III.	Initial calibration	A	
IV.	Routine calibration/IX	W	
V.	Blanks	W	
VI.	Matrix spike/Matrix spike duplicates	W	
VII.	Laboratory control samples	A	CCS
VIII.	Regional quality assurance and quality control	N	
IX.	Internal standards	W	
X.	Target compound identifications	A	Not reviewed for Stage 2B validation.
XI.	Compound quantitation and CRQLs	W	Not reviewed for Stage 2B validation.
XII.	System performance	A	Not reviewed for Stage 2B validation.
XIII.	Overall assessment of data	A	
XIV.	Field duplicates	W	D = 3+4, 8+9, 10+11
XV.	Field blanks	W	FB 04072010-R2D (F0D090441), FB 040710-R2C (F0D130519)

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

Validated Samples: ** Indicates sample underwent State 4 validation

1	SSAM6-01-1BPC	11	SSA07-01-1BPC_FD	21	0104236MB	31
2	SA60-3BPC	12	SSAM6-01-1BPCMS	22	0105361MB	32
3	SSAN6-04-1BPC	13	SSAM6-01-1BPCMSD	23		33
4	SSAN6-04-1BPC_FD	14		24		34
5	SSAN5-01-1BPC	15		25		35
6	SSAI2-02-2BPC**	16		26		36
7	SSAJ3-01-1BPC	17		27		37
8	SSAO6-04-1BPC_FD	18		28		38
9	SSAO6-04-1BPC	19		29		39
10	SSA07-01-1BPC	20		30		40

Notes: _____

VALIDATION FINDINGS CHECKLIST

Method: Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

Validation Area	Yes	No	NA	Findings/Comments
I. Technical holding times				
All technical holding times were met.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Cooler temperature criteria was met.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
II. GC/MS Instrument performance check				
Was PFK exact mass 380.9760 verified?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were the retention time windows established for all homologues?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was the chromatographic resolution between 2,3,7,8-TCDD and peaks representing any other unlabeled TCDD isomers $\leq 25\%$?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Is the static resolving power at least 10,000 (10% valley definition)?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was the mass resolution adequately check with PFK?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was the presence of 1,2,8,9-TCDD and 1,3,4,6,8-PeCDF verified?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
III. Initial calibration				
Was the initial calibration performed at 5 concentration levels?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were all percent relative standard deviations (%RSD) $\leq 20\%$ for unlabeled standards and $\leq 30\%$ for labeled standards?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Did all calibration standards meet the Ion Abundance Ratio criteria?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was the signal to noise ratio for each target compound ≥ 2.5 and for each recovery and internal standard ≥ 10 ?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
IV. Continuing calibration				
Was a routine calibration performed at the beginning and end of each 12 hour period?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were all percent differences (%D) $\leq 20\%$ for unlabeled standards and $\leq 30\%$ for labeled standards?	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	
Did all routine calibration standards meet the Ion Abundance Ratio criteria?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
V. Blanks				
Was a method blank associated with every sample in this SDG?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was a method blank performed for each matrix and concentration?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was there contamination in the method blanks? If yes, please see the Blanks validation completeness worksheet?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
VI. Matrix spike/Matrix spike duplicates				
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.	<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"/>	
VII. Laboratory control samples				
Was an LCS analyzed for this SDG?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
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"/>	

VALIDATION FINDINGS CHECKLIST

VIII. Regional Quality Assurance and Quality Control			
Were performance evaluation (PE) samples performed?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Were the performance evaluation (PE) samples within the acceptance limits?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
IX. Internal standards			
Were internal standard recoveries within the 40-135% criteria?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Was the minimum S/N ratio of all internal standard peaks ≥ 10 ?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
X. Target compound identification			
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?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
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?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
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?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Did compound spectra contain all characteristic ions listed in the table attached?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Was the Ion Abundance Ratio for the two quantitation ions within criteria?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Was the signal to noise ratio for each target compound and labeled standard > 2.5 ?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Does the maximum intensity of each specified characteristic ion coincide within ± 2 seconds (includes labeled standards)?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
For PCDF identification, was any signal ($S/N \geq 2.5$, at \pm seconds RT) detected in the corresponding PCDPE channel?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>
Was an acceptable lock mass recorded and monitored?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
XI. Compound quantitation/CRQLs			
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 CRQLs 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"/>
XII. System performance			
System performance was found to be acceptable.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
XIII. Overall assessment of data			
Overall assessment of data was found to be acceptable.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
XIV. Field duplicates			
Field duplicate pairs were identified in this SDG.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Target compounds were detected in the field duplicates.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
XV. Field blanks			
Field blanks were identified in this SDG.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Target compounds were detected in the field blanks.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>

VALIDATION FINDINGS WORKSHEET

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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

Routine Calibration

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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

N N/A Was a routine calibration was performed at the beginning and end of each 12 hour period?

Y N/A Were all percent differences (%D) of RRFs ≤ 20% for unlabeled compounds and ≤ 30% for labeled?

Y N N/A Did all routine calibration standards meet the Ion Abundance Ratio criteria?

#	Date	Standard ID	Compound	Finding %D (Limit: ≤30.0%)	Finding Ion Abundance Ratio	Associated Samples	Qualifications (C)
	4/28/10	BAF/12375.2	13C-D	30.3		8-10	Y/Dels/P(C-E)

	PCDDs	Selected Ions (m/z)	Ion Abundance Ratio	PCDFs	Selected Ions (m/z)	Ion Abundance Ratio
	Tetra-	M/M+2	0.65-0.89	Tetra-	M/M+2	0.65-0.89
	Penta-	M+2/M+4	1.32-1.78	Penta-	M+2/M+4	1.32-1.78
	Hexa-	M+2/M+4	1.05-1.43	Hexa-	M+2/M+4	1.05-1.43
	Hexa- ¹³ C-HxCDF (IS) only	M/M+2	0.43-0.59	Hexa- ¹³ C-HxCDF (IS) only	M/M+2	0.43-0.59
	Hepta- ¹³ C-HpCDF (IS) only	M/M+2	0.37-0.51	Hepta- ¹³ C-HpCDF (IS) only	M/M+2	0.37-0.51
	Hepta-	M+2/M+4	0.88-1.20	Hepta-	M+2/M+4	0.88-1.20
	Octa-	M+2/M+4	0.76-1.02	Octa-	M+2/M+4	0.76-1.02

LDC #: 2108B-2
 SDG #: See Cover

VALIDATION FINDINGS WORKSHEET

Field Blanks

Page: 1 of 1
 Reviewer: [Signature]
 2nd Reviewer: [Signature]

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

Y N N/A Were field blanks identified in this SDG?

Blank units: pg/L Associated sample units: pg/g

Sampling date: 4/8/10

Field blank type: (circle one) Field Blank Rinsate / Other: 1-5, 8-11 Associated Samples:

Compound	Blank ID	Blank ID	Sample Identification
	FB-04072010-RZC	5X	
C	0.77	0.00385	
D	0.74	0.0037	
E	0.82	0.0041	
F	4.2	0.021	
G	37	0.185	
H	0.57	0.00285	
I	0.96	0.0048	
J	0.67	0.00335	
K	1.1	0.0055	
L	0.96	0.0048	
M	1.0	0.005	
N	1.0	0.005	
O	2.1	0.0105	
P	1.5	0.0075	
Q	6.7	0.0335	
CRQL			

CIRCLED RESULTS WERE NOT QUALIFIED. ALL RESULTS NOT CIRCLED WERE QUALIFIED BY THE FOLLOWING STATEMENT:
 Samples with compound concentrations within five times the associated field blank concentration are listed above, these sample results were qualified as not detected, "U".

LDC #: 23188B21
SDG #: See Cover

VALIDATION FINDINGS WORKSHEET Field Blanks

Page: 1 of 4
Reviewer: _____
2nd Reviewer: _____

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

Y N N/A Were field blanks identified in this SDG?

Blank units: pg/L Associated sample units: pg/g

Sampling date: 4/7/10

Field blank type: (circle one) Field Blank / Rinsate / Other: _____

Associated Samples: _____

6-7 (>5 x)

Compound	Blank ID	5X	Sample Identification
	EB-04072010-RZD		
C	0.89	0.00445	
E	1.5	0.0075	
F	2.2	0.011	
G	8.3	0.0415	
K	1.4	0.007	
L	1.6	0.008	
M	1.5	0.0075	
N	1.6	0.008	
O	1.3	0.0065	
P	1.4	0.007	
Q	4.1	0.0205	
CRQL			

CIRCLED RESULTS WERE NOT QUALIFIED. ALL RESULTS NOT CIRCLED WERE QUALIFIED BY THE FOLLOWING STATEMENT:
Samples with compound concentrations within five times the associated field blank concentration are listed above, these sample results were qualified as not detected, "U".

VALIDATION FINDINGS WORKSHEET
 Matrix Spike/Matrix Spike Duplicates

Page: 1 of 1
 Reviewer: [Signature]
 2nd Reviewer:

LDC #: [Signature]
 SDG #: SEA CONEY

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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? If no, indicate which matrix does not have an associated MS/MSD. Soil / Water.

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

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

#	Date	MS/MSD ID	Compound	MS %R (Limits)	MSD %R (Limits)	RPD (Limits)	Associated Samples	Qualifications
		12/13	70 R	() ()	() ()	() ()	1	No Qual (MS or CS in)

VALIDATION FINDINGS WORKSHEET
Internal Standards

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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

N/A Are all internal standard recoveries within the 40-135% criteria?

N/A Was the S/N ratio all internal standard peaks > 10?

#	Date	Lab ID/Reference	Internal Standard	% Recovery (Limit: 40-135%)	Qualifications (I)
		1	F	38 (40-135)	✓ N/A (F. R)
				()	
				()	
		3	F	23	
			H	27	
			I	15	
				()	
				()	
		4	F	36	
			I	27	(F. O-R)
				()	
				()	
		6	F	33	
			H	37	
			I	20	
				()	
				()	
		7	F	38	
			I	25	
				()	
				()	
		10	I	27	
				()	
				()	
		11	I	22	
				()	
				()	

	Internal Standards	Check Standard Used	Internal Standards	Check Standard Used
A.	¹³ C-2,3,7,8-TCDF		G.	¹³ C-1,2,3,4,6,7,8-HpCDF
B.	¹³ C-2,3,7,8-TCDD		H.	¹³ C-1,2,3,4,6,7,8-HpCDD
C.	¹³ C-1,2,3,7,8-PeCDF		I.	¹³ C-OCDD
D.	¹³ C-1,2,3,7,8-PeCDD		K.	¹³ C-1,2,3,4-TCDD
E.	¹³ C-1,2,3,4,7,8-HxCDF		L.	¹³ C-1,2,3,7,8,9-HxCDD
F.	¹³ C-1,2,3,6,7,8-HxCDD			

VALIDATION FINDINGS WORKSHEET
Field Duplicates

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

Y N NA Were field duplicate pairs identified in this SDG?
Y N NA Were target analytes detected in the field duplicate pairs?

Compound	Concentration (pg/g)		(<50) RPD	(pg/g) Difference	(pg/g) Limits	Qualifications (Parent Only)
	3	4				
A	0.50U	0.041		0.459	(≤0.50)	
C	2.5U	0.081		2.419	(≤2.5)	
D	0.22	0.19		0.03	(≤2.5)	
E	0.30	0.32		0.02	(≤2.5)	
F	0.80	1.6		0.8	(≤2.5)	
G	3.7	8.6		4.9	(≤5.0)	
H	0.45	0.46		0.01	(≤0.50)	
I	0.70	0.73		0.03	(≤2.5)	
J	0.45	0.45		0	(≤2.5)	
K	1.3	1.5		0.2	(≤2.5)	
L	0.68	0.91		0.23	(≤2.5)	
M	0.27	0.22		0.05	(≤2.5)	
N	0.34	0.19		0.15	(≤2.5)	
O	2.9	2.8		0.1	(≤2.5)	
P	1.5	1.6		0.1	(≤2.5)	
Q	5.5	7.3		1.8	(≤5.0)	

Compound	Concentration (pg/g)		(<50) RPD	(pg/g) Difference	(pg/g) Limits	Qualifications (Parent Only)
	8	9				
A	0.86	0.80		0.06	(≤0.53)	
B	2.1	2.0		0.1	(≤2.6)	
C	1.8	1.6		0.2	(≤2.6)	
D	3.2	3.1		0.1	(≤2.6)	
E	2.8	2.6		0.2	(≤2.6)	
F	18	12	40			
G	140	30	129			<u>not/A (total)</u>
H	13	13	0			

LDC#: 23188B21

SDG#: See Cover

VALIDATION FINDINGS WORKSHEET

Field Duplicates

Page: 2 of 3

Reviewer: 9

2nd Reviewer: _____

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

Y N NA
Y N NA

Were field duplicate pairs identified in this SDG?

Were target analytes detected in the field duplicate pairs?

Compound	Concentration (pg/g)		(<50) RPD	(pg/g) Difference	(pg/g) Limits	Qualifications (Parent Only)
	8	9				
I	27	25	8			
J	13	12		1	(≤2.6)	
K	52	47	10			
L	35	32	9			
M	8.0	7.6		0.4	(≤2.6)	
N	4.5	4.2		0.3	(≤2.6)	
O	130	120	8			
P	45	44	2			
Q	300	250	18			

Compound	Concentration (pg/g)		(<50) RPD	(pg/g) Difference	(pg/g) Limits	Qualifications (Parent Only)
	10	11				
A	1.5	1.6		0.1	(≤0.52)	
B	3.4	4.0		0.6	(≤2.6)	
C	2.4	2.6		0.2	(≤2.6)	
D	4.2	4.8		0.6	(≤2.6)	
E	3.6	4.1		0.5	(≤2.6)	
F	14	16	13			
G	22	22	0			
H	160	100	46			
I	58	62	7			
J	24	26	8			
K	76	96	23			
L	50	63	23			
M	12	15	22			
N	8.1	9.2		1.1	(≤2.6)	
O	170	200	16			

LDC#: 23188B21

SDG#: See Cover

VALIDATION FINDINGS WORKSHEET

Field Duplicates

Page: 3 of 3

Reviewer: g

2nd Reviewer: _____

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

Y N NA Were field duplicate pairs identified in this SDG?

Y N NA Were target analytes detected in the field duplicate pairs?

Compound	Concentration (pg/g)		(≤50)	(pg/g)	(pg/g)	Qualifications (Parent Only)
	10	11	RPD	Difference	Limits	
P	110	94	16			
Q	480	940	65			<u>Wt/A</u> (fd)

V:\FIELD DUPLICATES\23188B21.wpd

VALIDATION FINDINGS WORKSHEET I
Initial Calibration Calculation Verification

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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

$RRF = (A_x)(C_s)/(A_s)(C_x)$ A_x = Area of compound, A_s = Area of associated internal standard
 average RRF = sum of the RRFs/number of standards C_x = Concentration of compound, C_s = Concentration of internal standard
 %RSD = $100 * (S/X)$ S = Standard deviation of the RRFs, X = Mean of the RRFs

#	Standard ID	Calibration Date	Compound (Reference Internal Standard)	Reported		Recalculated		Reported		Recalculated	
				Average RRF (Initial)	RRF (CS3 std)	Average RRF (Initial)	RRF (CS3 std)	%RSD	%RSD	RRF (CS3 std)	%RSD
1	1CAR (ADS)	4/12/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	0.945	0.98	0.945	0.98	4.44	4.33	0.98	4.33
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)	1.021	1.04	1.021	1.04	3.03	2.97	1.04	2.97
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)	1.114	1.19	1.114	1.19	5.33	5.25	1.19	5.25
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)	1.072	1.11	1.072	1.11	3.60	3.75	1.11	3.75
			OCDF (¹³ C-OCDD)	1.445	1.51	1.445	1.51	5.85	5.89	1.51	5.89
2	1CAR	3/10/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	1.000	1.00	1.000	1.00	1.36	1.58	1.00	1.58
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)								
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)								
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)								
			OCDF (¹³ C-OCDD)								
3	1CAR (SD-2)	4/21/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	1.088	1.10	1.088	1.10	1.29	1.20	1.10	1.20
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)								
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)								
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)								
			OCDF (¹³ 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
Routine Calibration Results Verification

Page: 1 of 1
 Reviewer: Q
 2nd Reviewer: _____

LDC #: 23/8824
 SDG #: SLCMB

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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 = $100 * (\text{ave. RRF} - \text{RRF}) / \text{ave. RRF}$ Where: ave. RRF = initial calibration average RRF
 RRF = $(A_x)(C_s) / (A_s)(C_x)$ RRF = continuing calibration RRF
 A_x = Area of compound, A_s = Area of associated internal standard
 C_x = Concentration of compound, C_s = Concentration of internal standard

#	Standard ID	Calibration Date	Compound (Reference Internal Standard)	Average RRF (Initial)	Reported		Recalculated	
					RRF (CC)	%D	RRF (CC)	%D
1	247045D	4/24/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	0.945	0.94	0.7	0.94	0.7
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)	1.021	0.96	5.9	0.96	5.9
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)	1.114	1.07	3.8	1.07	3.8
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)	1.072	1.04	3.2	1.04	3.2
			OCDF (¹³ C-OCDD)	1.445	1.41	2.3	1.41	2.3
2	247104DE	4/24/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	0.945	0.90	4.3	0.90	4.3
	19		2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)	1.021	0.96	6.3	0.96	6.3
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)	1.114	1.08	3.1	1.08	3.1
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)	1.072	1.03	3.9	1.03	3.9
			OCDF (¹³ C-OCDD)	1.445	1.36	5.9	1.36	5.9
3	247108BD	4/26/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	1.088	1.02	5.9	1.02	5.9
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)					
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)					
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)					
			OCDF (¹³ C-OCDD)					

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
Matrix Spike/Matrix Spike Duplicates Results Verification

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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 = $100 * (SSR - SR) / SA$ Where: SSR = Spiked sample result, SR = Sample result
 SA = Spike added

RPD = $|MSR - MSDR| * 2 / (MSR + MSDR)$ MSR = Matrix spike percent recovery MSDR = Matrix spike duplicate percent recovery

MS/MSD samples: 12/13

Compound	Spike Added (pg/g)		Sample Concentration (pg/g)	Spiked Sample Concentration (pg/g)		Matrix Spike Percent Recovery		Matrix Spike Duplicate Percent Recovery		Reported RPD	Recalculated RPD
	MS	MSD		MS	MSD	Reported	Recalc	Reported	Recalc		
2,3,7,8-TCDD	19.6	19.9	22	44.7	39.4	121	116	93	87	12	13
1,2,3,7,8-PeCDD	98.2	99.3	72	181	178	114	111	110	107	1.2	1.7
1,2,3,4,7,8-HxCDD	✓	✓	46	168	163	137	124	120	118	2.9	3.0
1,2,3,4,7,8,9-HpCDF	✓	✓	200	274.0	257.0	243	550	44.0	34.2	7.4	7.6
OCDF	196	199	1600	19600	17800	2000	1837	1110	905	9.3	9.6

Comments: Refer to Matrix Spike/Matrix Spike 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
Laboratory Control Sample Results Verification

METHOD: GC/MS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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 = $100 * \frac{LCS - LCSD}{LCS + LCSD}$ LCS = Laboratory control sample percent recovery LCSD = Laboratory control sample duplicate percent recovery

LCS ID: 0104236

Compound	Spike Added (20.0)		Spiked Sample Concentration (99.5)		LCS		LCSD		Percent Recovery		Percent Recovery		RPD	
	LCS	LCSD	LCS	LCSD	Reported	Recalc	Reported	Recalc	Reported	Recalc	Reported	Recalc	Reported	Recalc
2,3,7,8-TCDD	20.0	NA	19.5	NA	97	97								
1,2,3,7,8-PeCDD	100		99.5		100	100								
1,2,3,4,7,8-HxCDD			103		103	103								
1,2,3,4,7,8,9-HpCDF			99.9		100	100								
OCDF	200		197		99	99								

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.

Descriptor	Accurate mass ^(a)	Ion ID	Elemental Composition	Analyte	Descriptor	Accurate Mass ^(a)	Ion ID	Elemental Composition	Analyte			
1	303.9016	M	C ₁₂ H ₄ ³⁵ Cl ₆ O	TCDF	4	407.7818	M+2	C ₁₂ H ³⁵ Cl ₈ ³⁷ ClO	HpCDF			
	305.8987	M+2	C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ Cl ₁₀	TCDF		M+4	409.7788	M+4	C ₁₂ H ³⁵ Cl ₃ ³⁷ Cl ₂ O	HpCDF		
	315.9419	M	¹⁹ C ₁₂ H ₄ ³⁵ Cl ₄ O	TCDF (S)		M	417.8250	M	¹⁹ C ₁₂ H ³⁵ Cl ₇ O	HpCDF (S)		
	317.9399	M+2	¹⁹ C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ ClO	TCDF (S)		M+2	419.8220	M+2	¹⁹ C ₁₂ H ³⁵ Cl ₆ ³⁷ ClO	HpCDF		
	319.8965	M	C ₁₂ H ₄ ³⁵ Cl ₄ O ₂	TCDD		M+2	423.7767	M+2	C ₁₂ H ³⁵ Cl ₈ ³⁷ ClO ₂	HpCDD		
	321.8936	M+2	C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ ClO ₂	TCDD		M+4	425.7737	M+4	C ₁₂ H ³⁵ Cl ₅ ³⁷ Cl ₂ O ₂	HpCDD		
	331.9368	M	¹⁹ C ₁₂ H ₄ ³⁵ Cl ₄ O ₂	TCDD (S)		M+2	435.8169	M+2	¹⁹ C ₁₂ H ³⁵ Cl ₆ ³⁷ ClO ₂	HpCDD (S)		
	333.9338	M+2	¹⁹ C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ ClO ₂	TCDD (S)		M+4	437.8140	M+4	¹⁹ C ₁₂ H ³⁵ Cl ₅ ³⁷ Cl ₂ O ₂	HpCDD (S)		
	375.8364	M+2	C ₁₂ H ₄ ³⁵ Cl ₅ ³⁷ ClO	HxCDFE		M+4	479.7165	M+4	C ₁₂ H ³⁵ Cl ₇ ³⁷ Cl ₂ O	NCDFE		
	[354.9792]	LOCK	C ₉ F ₁₃	PFK		LOCK	[430.9728]	LOCK	C ₉ F ₁₇	PFK		
	2	339.8597	M+2	C ₁₂ H ₃ ³⁵ Cl ₁ ³⁷ ClO		PeCDF	5	441.7428	M+2	C ₁₂ ³⁵ Cl ₁ ³⁷ ClO	OCDF	
		341.8567	M+4	C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ Cl ₂ O		PeCDF		M+4	443.7399	M+4	C ₁₂ ³⁵ Cl ₃ ³⁷ Cl ₂ O	OCDF
		351.9000	M+2	¹⁹ C ₁₂ H ₃ ³⁵ Cl ₄ ³⁷ ClO		PeCDF (S)		M+2	457.7377	M+2	C ₁₂ ³⁵ Cl ₇ ³⁷ ClO ₂	OCDD
		353.8970	M+4	¹⁹ C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ Cl ₂ O		PeCDF (S)		M+4	459.7348	M+4	C ₁₂ ³⁵ Cl ₆ ³⁷ ClO ₂	OCDD
355.8546		M+2	C ₁₂ H ₃ ³⁵ Cl ₄ ³⁷ ClO ₂	PeCDD	M+2	469.7760		M+2	¹⁹ C ₁₂ ³⁵ Cl ₇ ³⁷ ClO ₂	OCDD (S)		
357.8516		M+4	C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ Cl ₂ O ₂	PeCDD	M+4	471.7750		M+4	¹⁹ C ₁₂ ³⁵ Cl ₆ ³⁷ Cl ₂ O ₂	OCDD (S)		
367.8949		M+2	¹⁹ C ₁₂ H ₃ ³⁵ Cl ₄ ³⁷ ClO ₂	PeCDD (S)	M+2	513.6775		M+2	C ₁₂ ³⁵ Cl ₉ ³⁷ ClO	DCDFE		
369.8919		M+4	¹⁹ C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ Cl ₂ O ₂	PeCDD (S)	M+4	[422.9278]		M+4	C ₁₀ F ₁₇	PFK		
409.7974		M+2	C ₁₂ H ₃ ³⁵ Cl ₆ ³⁷ ClO	HxCDFE	LOCK	[430.9728]		LOCK				
[354.9792]		LOCK	C ₉ F ₁₃	PFK								
3		373.8208	M+2	C ₁₂ H ₂ ³⁵ Cl ₁ ³⁷ ClO	HxCDF							
		375.8178	M+4	C ₁₂ H ₂ ³⁵ Cl ₃ ³⁷ Cl ₂ O	HxCDF							
		383.8639	M	¹⁹ C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ ClO	HxCDF (S)							
		385.8610	M+2	¹⁹ C ₁₂ H ₂ ³⁵ Cl ₃ ³⁷ ClO	HxCDF (S)							
	389.8156	M+2	C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ ClO ₂	HxCDD								
	391.8127	M+4	C ₁₂ H ₂ ³⁵ Cl ₃ ³⁷ Cl ₂ O ₂	HxCDD								
	401.8559	M+2	¹⁹ C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ ClO ₂	HxCDD (S)								
	403.8529	M+4	¹⁹ C ₁₂ H ₂ ³⁵ Cl ₃ ³⁷ Cl ₂ O ₂	HxCDD (S)								
	445.7555	M+4	C ₁₂ H ₂ ³⁵ Cl ₆ ³⁷ Cl ₂ O	OCDFE								
	[430.9728]	LOCK	C ₉ F ₁₇	PFK								

(a) The following nucleidic masses were used:

- H = 1.007825
- C = 12.000000
- ¹³C = 13.003355
- F = 18.9984
- O = 15.994915
- ³⁵Cl = 34.968853
- ³⁷Cl = 36.965903

S = internal/recovery standard

VALIDATION FINDINGS WORKSHEET
Sample Calculation Verification

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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 agree within 10.0% of the reported results?

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

- A_x = Area of the characteristic ion (EICP) for the compound to be measured
- A_s = 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).
- RRF = Relative Response Factor (average) from the initial calibration
- Df = Dilution Factor.
- %S = Percent solids, applicable to soil and solid matrices only.

Example:

Sample I.D. 6 A:

$$\text{Conc.} = \frac{(16053/20)(2000)}{(6104/100)(1.021)(10.7)(0.95)}$$

$$= 50.6 \mu\text{g/g}$$

#	Sample ID	Compound	Reported Concentration ()	Calculated Concentration ()	Qualification

Laboratory Data Consultants, Inc. Data Validation Report

Project/Site Name: Tronox LLC Facility, PCS, Henderson, Nevada
Collection Date: April 12, 2010
LDC Report Date: May 24, 2010
Matrix: Soil
Parameters: Dioxins/Dibenzofurans
Validation Level: Stage 2B & 4
Laboratory: TestAmerica, Inc.

Sample Delivery Group (SDG): G0D140422

Sample Identification

SSAJ3-02-1BPC**
SSAH3-01-1BPC**
SSAM7-03-1BPC
SSAM7-04-1BPC
SSAN7-02-1BPC
SSAN6-06-1BPC

**Indicates sample underwent Stage 4 review

Introduction

This data review covers 6 soil samples listed on the cover sheet including dilutions and reanalysis as applicable. The analyses were per EPA SW 846 Method 8290 for Polychlorinated Dioxins/Dibenzofurans.

This review follows the Standard Operating Procedures (SOP) 40, Data Review/Validation (BRC 2009), the Quality Assurance Project Plan Tronox LLC Facility, Henderson, Nevada (June 2009), NDEP guidance (May 2006), and USEPA Contract Laboratory Program National Functional Guidelines for Polychlorinated Dioxins/Dibenzofurans Data Review (September 2005).

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.

Blank results are summarized in Section V.

Field duplicates are summarized in Section XIV.

Samples indicated by a double asterisk on the front cover underwent a Stage 4 review. A Stage 2B review was performed on all of the other samples. Raw data were not evaluated for the samples reviewed by Stage 2B criteria since this review is based on QC data.

The following are definitions of the data qualifiers:

- J+ Data are qualified as estimated, with a high bias likely to occur. False positives or false negatives are unlikely to have been reported.
- J- Data are qualified as estimated, with a low bias likely to occur. False positives or false negatives are unlikely to have been reported.
- J Data are qualified as estimated; it is not possible to assess the direction of the potential bias. False positives or false negatives are unlikely to have been reported.
- U Indicates the compound or analyte was analyzed for but not detected at or above the stated limit.
- R Data are qualified as rejected. There is a significant potential for the reporting of false negatives or false positives.
- UU Indicates the compound or analyte was analyzed for but not detected. The sample detection limit is an estimated value.
- B The analytical result may be a false positive totally attributable to blank contamination. This qualifier is applicable to radiochemistry analysis only.
- JB The analytical result may be biased high and partially attributable to blank contamination. This qualifier is applicable to radiochemistry analysis only.
- JK The analytical result is an estimated maximum possible concentration (EMPC).
- X The analytical result is not used for reporting because a more accurate and precise result is reported in its place.
- J-TDS The analytical result is estimated based on failure of the Total Dissolved Solids (TDS) correctness check performed in accordance with the Standard Method 1030E.
- J-CAB The analytical result is estimated based on failure of the cation-anion balance correctness check performed in accordance with Standard Method 1030E.
- J-TDS & CAB The analytical result is unreliable based on the failure of the cation-anion balance and TDS correctness check performed in accordance with standard Method 1030E.
- A Indicates the finding is based upon technical validation criteria.
- P Indicates the finding is related to a protocol/contractual deviation.
- None Indicates the data was not significantly impacted by the finding, therefore qualification was not required.

I. Technical Holding Times

All technical holding time requirements were met.

The chain-of-custodies were reviewed for documentation of cooler temperatures. All cooler temperatures met validation criteria.

II. HRGC/HRMS Instrument Performance Check

Instrument performance was checked at the required daily 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 exact mass of 380.9760 of PFK was verified. The static resolving power was at least 10,000 (10% valley definition) for samples on which Stage 4 review was performed. Raw data were not evaluated for the samples reviewed by Stage 2B criteria.

III. Initial Calibration

A five point initial calibration was performed as required by the method.

Percent relative standard deviations (%RSD) were less than or equal to 20.0% for unlabeled compounds and less than or equal to 30.0% for labeled compounds.

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

The minimum S/N ratio for each target compound was greater than or equal to 2.5 and greater than or equal to 10 for each recovery and internal standard compound for samples on which Stage 4 review was performed. Raw data were not evaluated for the samples reviewed by Stage 2B criteria.

IV. Routine Calibration (Continuing)

Routine calibration was performed at the required frequencies.

All of the routine calibration percent differences (%D) between the initial calibration RRF and the routine calibration RRF were less than or equal to 20.0% for unlabeled compounds and less than or equal to 30.0% for labeled compounds.

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

V. Blanks

Method blanks were reviewed for each matrix as applicable. No polychlorinated dioxin/dibenzofuran contaminants were found in the method blanks.

Sample EB-04122010-RIG3-RZB (from SDG G0D140534) was identified as an equipment blank. No polychlorinated dioxin/dibenzofuran contaminants were found in this blank with the following exceptions:

Equipment Blank ID	Sampling Date	Compound	Concentration	Associated Samples
EB-04122010-RIG3-RZB	4/12/10	1,2,3,6,7,8-HxCDD	0.64 pg/L	SSAJ3-02-1BPC**
		1,2,3,7,8,9-HxCDD	0.47 pg/L	SSAH3-01-1BPC**
		1,2,3,4,6,7,8-HpCDD	1.9 pg/L	
		OCDD	5.0 pg/L	
		2,3,7,8-TCDF	4.6 pg/L	
		1,2,3,7,8-PeCDF	3.8 pg/L	
		2,3,4,7,8-PeCDF	2.2 pg/L	
		1,2,3,4,7,8-HxCDF	7.7 pg/L	
		1,2,3,6,7,8-HxCDF	4.8 pg/L	
		2,3,4,6,7,8-HxCDF	1.2 pg/L	
		1,2,3,7,8,9-HxCDF	0.59 pg/L	
		1,2,3,4,6,7,8-HpCDF	14 pg/L	
		1,2,3,4,7,8,9-HpCDF	5.0 pg/L	
		OCDF	27 pg/L	

Sample concentrations were compared to concentrations detected in the equipment blanks as required by the QAPP. No sample data was qualified.

Samples FB-04072010-RZD (from SDG G0D090441) and FB-04072010-RZC (from SDG G0D130519) were identified as field blanks. No polychlorinated dioxin/dibenzofuran contaminants were found in these blanks with the following exceptions:

Field Blank ID	Sampling Date	Compound	Concentration	Associated Samples
FB-04072010-RZD	4/7/10	1,2,3,4,7,8-HxCDD	0.89 pg/L	SSAJ3-02-1BPC**
		1,2,3,7,8,9-HxCDD	1.5 pg/L	SSAH3-01-1BPC**
		1,2,3,4,6,7,8-HpCDD	2.2 pg/L	
		OCDD	8.3 pg/L	
		1,2,3,4,7,8-HxCDF	1.4 pg/L	
		1,2,3,6,7,8-HxCDF	1.6 pg/L	
		2,3,4,6,7,8-HxCDF	1.5 pg/L	
		1,2,3,7,8,9-HxCDF	1.6 pg/L	
		1,2,3,4,6,7,8-HpCDF	1.3 pg/L	
		1,2,3,4,7,8,9-HpCDF	1.4 pg/L	
		OCDF	4.1 pg/L	

Field Blank ID	Sampling Date	Compound	Concentration	Associated Samples
FB-04072010-RZC	4/8/10	1,2,3,4,7,8-HxCDD	0.77 pg/L	SSAM7-03-1BPC
		1,2,3,6,7,8-HxCDD	0.74 pg/L	SSAM7-04-1BPC
		1,2,3,7,8,9-HxCDD	0.82 pg/L	SSAN7-02-1BPC
		1,2,3,4,6,7,8-HpCDD	4.2 pg/L	SSAN6-06-1BPC
		OCDD	37 pg/L	
		2,3,7,8-TCDF	0.57 pg/L	
		1,2,3,7,8-PeCDF	0.96 pg/L	
		2,3,4,7,8-PeCDF	0.67 pg/L	
		1,2,3,4,7,8-HxCDF	1.1 pg/L	
		1,2,3,6,7,8-HxCDF	0.96 pg/L	
		2,3,4,6,7,8-HxCDF	1.0 pg/L	
		1,2,3,7,8,9-HxCDF	1.0 pg/L	
		1,2,3,4,6,7,8-HpCDF	2.1 pg/L	
		1,2,3,4,7,8,9-HpCDF	1.5 pg/L	
		OCDF	6.7 pg/L	

Sample concentrations were compared to concentrations detected in the field blanks as required by the QAPP. No sample data was qualified.

VI. Matrix Spike/Matrix Spike Duplicates

Matrix spike (MS) and matrix spike duplicate (MSD) samples were reviewed for each matrix as applicable. Percent recoveries (%R) and relative percent differences (RPD) were not within QC limits. Since there were no associated samples, no data were qualified.

VII. Laboratory Control Samples (LCS)

Laboratory control samples were reviewed for each matrix as applicable. Percent recoveries (%R) were within QC limits.

VIII. Regional Quality Assurance and Quality Control

Not applicable.

IX. Internal Standards

All internal standard recoveries were within QC limits with the following exceptions:

Sample	Internal Standards	%R (Limits)	Compound	Flag	A or P
SSAJ3-02-1BPC**	¹³ C-OCDD	29 (40-135)	OCDD	J (all detects) UJ (all non-detects)	P
			OCDF	J (all detects) UJ (all non-detects)	

Sample	Internal Standards	%R (Limits)	Compound	Flag	A or P
SSAH3-01-1BPC**	¹³ C-1,2,3,4,6,7,8-HpCDF ¹³ C-1,2,3,4,6,7,8-HpCDD ¹³ C-OCDD	27 (40-135) 29 (40-135) 15 (40-135)	1,2,3,4,6,7,8-HpCDD OCDD 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	J (all detects) UJ (all non-detects)	P
SSAM7-03-1BPC	¹³ C-OCDD	25 (40-135)	OCDD OCDF	J (all detects) UJ (all non-detects) J (all detects) UJ (all non-detects)	P
SSAN7-02-1BPC	¹³ C-1,2,3,4,6,7,8-HpCDF ¹³ C-1,2,3,4,6,7,8-HpCDD ¹³ C-OCDD	35 (40-135) 36 (40-135) 26 (40-135)	1,2,3,4,6,7,8-HpCDD OCDD 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	J (all detects) UJ (all non-detects)	P
SSAN6-06-1BPC	¹³ C-1,2,3,4,6,7,8-HpCDF ¹³ C-1,2,3,4,6,7,8-HpCDD ¹³ C-OCDD	31 (40-135) 36 (40-135) 24 (40-135)	1,2,3,4,6,7,8-HpCDD OCDD 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	J (all detects) UJ (all non-detects)	P

X. Target Compound Identifications

All target compound identifications were within validation criteria for samples on which a Stage 4 review was performed. Raw data were not evaluated for the samples reviewed by Stage 2B criteria.

XI. Project Quantitation Limit

All compound quantitation and CRQLs were within validation criteria with the following exceptions:

Sample	Compound	Finding	Criteria	Flag	A or P
SSAM7-03-1BPC	1,2,3,4,6,7,8-HpCDF OCDF	Sample result exceeded calibration range.	Reported result should be within calibration range.	J (all detects) J (all detects)	P
SSAM7-04-1BPC	OCDF	Sample result exceeded calibration range.	Reported result should be within calibration range.	J (all detects)	P
SSAN6-06-1BPC	2,3,7,8-TCDF 1,2,3,7,8-PeCDF 1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	Sample result exceeded calibration range.	Reported result should be within calibration range.	J (all detects) J (all detects) J (all detects) J (all detects) J (all detects) J (all detects) J (all detects)	P

All compounds reported below the PQL were qualified as follows:

Sample	Finding	Flag	A or P
All samples in SDG G0D140422	All compounds reported below the PQL.	J (all detects)	A

All compounds reported as EMPC were qualified as follows:

Sample	Finding	Flag	A or P
All samples in SDG G0D140422	All compounds reported as estimated maximum possible concentration (EMPC).	JK (all detects)	A

Raw data were not evaluated for the samples reviewed by Stage 2B criteria.

XII. System Performance

The system performance was acceptable for samples on which a Stage 4 review was performed. Raw data were not evaluated for the samples reviewed by Stage 2B criteria.

XIII. Overall Assessment of Data

Data flags are summarized at the end of this report if data has been qualified.

XIV. Field Duplicates

Samples SSAJ3-02-1BPC** and SSAJ3-02-1BPC_FD (from SDG G0D140453) and samples SSAN7-02-1BPC and SSAN7-02-1BPC_FD (from SDG (G0D140453) and were identified as field duplicates. No polychlorinated dioxins/dibenzofurans were detected in any of the samples with the following exceptions:

Compound	Concentration (pg/g)		RPD (Limits)	Difference (Limits)	Flags	A or P
	SSAJ3-02-1BPC**	SSAJ3-02-1BPC_FD				
2,3,7,8-TCDD	0.22	1.1U	-	0.88 (≤ 1.1)	-	-
1,2,3,7,8-PeCDD	0.90	5.3U	-	4.4 (≤ 5.3)	-	-
1,2,3,4,7,8-HxCDD	0.99	5.3U	-	4.31 (≤ 5.3)	-	-
1,2,3,6,7,8-HxCDD	1.2	0.13	-	1.07 (≤ 5.3)	-	-
1,2,3,7,8,9-HxCDD	1.1	0.12	-	0.98 (≤ 5.3)	-	-

Compound	Concentration (pg/g)		RPD (Limits)	Difference (Limits)	Flags	A or P
	SSAJ3-02-1BPC**	SSAJ3-02-1BPC_FD				
1,2,3,4,6,7,8-HpCDD	1.6	0.41	-	1.19 (≤ 5.3)	-	-
OCDD	5.9	1.4	-	4.5 (≤ 11)	-	-
2,3,7,8-TCDF	0.90	0.60	-	0.3 (≤ 1.1)	-	-
1,2,3,7,8-PeCDF	2.1	0.69	-	1.41 (≤ 5.3)	-	-
2,3,4,7,8-PeCDF	1.5	0.39	-	1.11 (≤ 5.3)	-	-
1,2,3,4,7,8-HxCDF	1.7	1.2	-	0.5 (≤ 5.3)	-	-
1,2,3,6,7,8-HxCDF	2.2	0.76	-	1.44 (≤ 5.3)	-	-
2,3,4,6,7,8-HxCDF	1.5	0.16	-	1.34 (≤ 5.3)	-	-
1,2,3,7,8,9-HxCDF	1.4	0.18	-	1.22 (≤ 5.3)	-	-
1,2,3,4,6,7,8-HpCDF	3.2	2.7	-	0.5 (≤ 5.3)	-	-
1,2,3,4,7,8,9-HpCDF	2.1	0.99	-	1.11 (≤ 5.3)	-	-
OCDF	9.1	7.8	-	1.3 (≤ 11)	-	-

Compound	Concentration (pg/g)		RPD (Limits)	Difference (Limits)	Flags	A or P
	SSAN7-02-1BPC	SSAN7-02-1BPC_FD				
2,3,7,8-TCDD	2.4	2.4	-	0 (≤ 1.1)	-	-
1,2,3,7,8-PeCDD	11	10	-	1 (≤ 5.3)	-	-
1,2,3,4,7,8-HxCDD	6.6	5.3	-	1.3 (≤ 5.3)	-	-
1,2,3,6,7,8-HxCDD	15	12	-	3 (≤ 5.3)	-	-
1,2,3,7,8,9-HxCDD	11	8.6	-	2.4 (≤ 5.3)	-	-
1,2,3,4,6,7,8-HpCDD	39	35	11 (≤ 50)	-	-	-

Compound	Concentration (pg/g)		RPD (Limits)	Difference (Limits)	Flags	A or P
	SSAN7-02-1BPC	SSAN7-02-1BPC_FD				
OCDD	40	35	-	5 (≤ 11)	-	-
2,3,7,8-TCDF	130	170	27 (≤ 50)	-	-	-
1,2,3,7,8-PeCDF	160	210	27 (≤ 50)	-	-	-
2,3,4,7,8-PeCDF	77	82	6 (≤ 50)	-	-	-
1,2,3,4,7,8-HxCDF	230	250	8 (≤ 50)	-	-	-
1,2,3,6,7,8-HxCDF	200	160	22 (≤ 50)	-	-	-
2,3,4,6,7,8-HxCDF	44	35	23 (≤ 50)	-	-	-
1,2,3,7,8,9-HxCDF	42	21	-	21 (≤ 5.3)	J (all detects)	A
1,2,3,4,6,7,8-HpCDF	500	440	13 (≤ 50)	-	-	-
1,2,3,4,7,8,9-HpCDF	270	220	20 (≤ 50)	-	-	-
OCDF	1300	1200	8 (≤ 50)	-	-	-

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Data Qualification Summary - SDG G0D140422**

SDG	Sample	Compound	Flag	A or P	Reason (Code)
G0D140422	SSAJ3-02-1BPC** SSAM7-03-1BPC	OCDD OCDF	J (all detects) UJ (all non-detects) J (all detects) UJ (all non-detects)	P	Internal standards (%R) (i)
G0D140422	SSAH3-01-1BPC** SSAN7-02-1BPC SSAN6-06-1BPC	1,2,3,4,6,7,8-HpCDD OCDD 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	J (all detects) UJ (all non-detects)	P	Internal standards (%R) (i)
G0D140422	SSAM7-03-1BPC	1,2,3,4,6,7,8-HpCDF OCDF	J (all detects) J (all detects)	P	Compound quantitation and CRQLs (e)
G0D140422	SSAM7-04-1BPC	OCDF	J (all detects)	P	Compound quantitation and CRQLs (e)
G0D140422	SSAN6-06-1BPC	2,3,7,8-TCDF 1,2,3,7,8-PeCDF 1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	J (all detects) J (all detects) J (all detects) J (all detects) J (all detects) J (all detects) J (all detects)	P	Compound quantitation and CRQLs (e)
G0D140422	SSAJ3-02-1BPC** SSAH3-01-1BPC** SSAM7-03-1BPC SSAM7-04-1BPC SSAN7-02-1BPC SSAN6-06-1BPC	All compounds reported below the PQL	J (all detects)	A	Project Quantitation Limit (sp)
G0D140422	SSAJ3-02-1BPC** SSAH3-01-1BPC** SSAM7-03-1BPC SSAM7-04-1BPC SSAN7-02-1BPC SSAN6-06-1BPC	All compounds reported as EMPC	JK (all detects)	A	Project Quantitation Limit (k)
G0D140422	SSAN7-02-1BPC	1,2,3,7,8,9-HxCDF	J (all detects)	A	Field duplicates (Difference) (fd)

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Laboratory Blank Data Qualification Summary - SDG
G0D140422**

No Sample Data Qualified in this SDG

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Equipment Blank Data Qualification Summary - SDG
G0D140422**

No Sample Data Qualified in this SDG

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Field Blank Data Qualification Summary - SDG G0D140422**

No Sample Data Qualified in this SDG

Tronox Northgate Henderson

VALIDATION COMPLETENESS WORKSHEET

Stage 2B/4

LDC #: 23188C21
 SDG #: G0D140422
 Laboratory: Test America

Date: 5/21/10
 Page: 1 of 1
 Reviewer: [Signature]
 2nd Reviewer:

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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.	Technical holding times	A	Sampling dates: 4/12/10
II.	HRGC/HRMS Instrument performance check	A	
III.	Initial calibration	A	
IV.	Routine calibration/ICV	A	
V.	Blanks	A	
VI.	Matrix spike/Matrix spike duplicates	SW	No sp added - no anal
VII.	Laboratory control samples	A	ICV
VIII.	Regional quality assurance and quality control	N	
IX.	Internal standards	SW	
X.	Target compound identifications	A	Not reviewed for Stage 2B validation.
XI.	Compound quantitation and CRQLs	SW	Not reviewed for Stage 2B validation.
XII.	System performance	A	Not reviewed for Stage 2B validation.
XIII.	Overall assessment of data	A	
XIV.	Field duplicates	SW	D=1+SSAJ3-02-1BPC_FD, 5+SSAN7-02-1BPC_FD(G0D140543)
XV.	Field blanks	SW	FB-04072010-RZD(G0D090441), FB-040710-RZC(G0D130519) EB-04122010-RIG3-RZD(G0D140534)

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

Validated Samples: ** Indicates sample underwent State 4 validation

1	SSAJ3-02-1BPC**	5	11	0105361MB	21		31
2	SSAH3-01-1BPC**		12		22		32
3	SSAM7-03-1BPC		13		23		33
4	SSAM7-04-1BPC		14		24		34
5	SSAN7-02-1BPC		15		25		35
6	SSAN6-06-1BPC		16		26		36
7			17		27		37
8			18		28		38
9			19		29		39
10			20		30		40

Notes: _____

VALIDATION FINDINGS CHECKLIST

Method: Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

Validation Area	Yes	No	NA	Findings/Comments
I. Technical holding times				
All technical holding times were met.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Cooler temperature criteria was met.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
II. GC/MS Instrument performance check				
Was PFK exact mass 380.9760 verified?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were the retention time windows established for all homologues?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was the chromatographic resolution between 2,3,7,8-TCDD and peaks representing any other unlabeled TCDD isomers $\leq 25\%$?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Is the static resolving power at least 10,000 (10% valley definition)?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was the mass resolution adequately check with PFK?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was the presence of 1,2,8,9-TCDD and 1,3,4,6,8-PeCDF verified?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
III. Initial calibration				
Was the initial calibration performed at 5 concentration levels?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were all percent relative standard deviations (%RSD) $\leq 20\%$ for unlabeled standards and $< 30\%$ for labeled standards?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Did all calibration standards meet the Ion Abundance Ratio criteria?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was the signal to noise ratio for each target compound ≥ 2.5 and for each recovery and internal standard ≥ 10 ?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
IV. Continuing calibration				
Was a routine calibration performed at the beginning and end of each 12 hour period?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were all percent differences (%D) $\leq 20\%$ for unlabeled standards and $\leq 30\%$ for labeled standards?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Did all routine calibration standards meet the Ion Abundance Ratio criteria?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
V. Blanks				
Was a method blank associated with every sample in this SDG?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was a method blank performed for each matrix and concentration?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was there contamination in the method blanks? If yes, please see the Blanks validation completeness worksheet?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
VI. Matrix spike/Matrix spike duplicates				
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.	<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"/>	
VII. Laboratory control samples				
Was an LCS analyzed for this SDG?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
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"/>	

VIII. Regional Quality Assurance and Quality Control			
Were performance evaluation (PE) samples performed?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Were the performance evaluation (PE) samples within the acceptance limits?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
IX. Internal standards			
Were internal standard recoveries within the 40-135% criteria?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Was the minimum S/N ratio of all internal standard peaks ≥ 10 ?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
X. Target compound identification			
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?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
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?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
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?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Did compound spectra contain all characteristic ions listed in the table attached?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Was the Ion Abundance Ratio for the two quantitation ions within criteria?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Was the signal to noise ratio for each target compound and labeled standard ≥ 2.5 ?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Does the maximum intensity of each specified characteristic ion coincide within ± 2 seconds (includes labeled standards)?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
For PCDF identification, was any signal ($S/N \geq 2.5$, at \pm seconds RT) detected in the corresponding PCDPE channel?	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>
Was an acceptable lock mass recorded and monitored?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
XI. Compound quantitation/CRQLs			
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 CRQLs 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"/>
XII. System performance			
System performance was found to be acceptable.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
XIII. Overall assessment of data			
Overall assessment of data was found to be acceptable.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
XIV. Field duplicates			
Field duplicate pairs were identified in this SDG.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Target compounds were detected in the field duplicates.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
XV. Field blanks			
Field blanks were identified in this SDG.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Target compounds were detected in the field blanks.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>

VALIDATION FINDINGS WORKSHEET

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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:

LDC #: 23188C21
SDG #: See Cover

VALIDATION FINDINGS WORKSHEET
Field Blanks

Page: 1 of 1
Reviewer: Y
2nd Reviewer:

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

Y N A Were field blanks identified in this SDG?

Blank units: pg/L Associated sample units: pg/g

Sampling date: 4/12/10

Field blank type: (circle one) Field Blank / Rinsate / Other: Field Blank Associated Samples: 1-2

Compound	Blank ID	Blank Concentration	Sample Identification
	FB-04122010-RIG3-RZD	5X	
D	0.64	0.0032	
E	0.47	0.00235	
F	1.9	0.0095	
G	5.0	0.025	
H	4.6	0.023	
I	3.8	0.019	
J	2.2	0.011	
K	7.7	0.0385	
L	4.8	0.024	
M	1.2	0.006	
N	0.59	0.00295	
O	14	0.07	
P	5.0	0.025	
Q	27	0.135	
CRQL			

CIRCLED RESULTS WERE NOT QUALIFIED. ALL RESULTS NOT CIRCLED WERE QUALIFIED BY THE FOLLOWING STATEMENT:
Samples with compound concentrations within five times the associated field blank concentration are listed above, these sample results were qualified as not detected, "U".

LDC # 2318007
 SDG #: See Cover

VALIDATION FINDINGS WORKSHEET

Field Blanks

Page: 1 of 1
 Reviewer: [Signature]
 2nd Reviewer: [Signature]

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

Y N N/A Were field blanks identified in this SDG?

Blank units: pg/L Associated sample units: pg/g

Sampling date: 4/8/10

Field blank type: (circle one) Field Blank Rinsate / Other: 3-6 (75x) Associated Samples: 3-6 (75x)

Compound	Blank ID	Blank ID	Sample Identification
	EB-04072010-BZC	5X	
C	0.77	0.00385	
D	0.74	0.0037	
E	0.82	0.0041	
F	4.2	0.021	
G	37	0.185	
H	0.57	0.00285	
I	0.96	0.0048	
J	0.67	0.00335	
K	1.1	0.0055	
L	0.96	0.0048	
M	1.0	0.005	
N	1.0	0.005	
O	2.1	0.0105	
P	1.5	0.0075	
Q	6.7	0.0335	
CRQL			

CIRCLED RESULTS WERE NOT QUALIFIED. ALL RESULTS NOT CIRCLED WERE QUALIFIED BY THE FOLLOWING STATEMENT:
 Samples with compound concentrations within five times the associated field blank concentration are listed above, these sample results were qualified as not detected, "U".

VALIDATION FINDINGS WORKSHEET
Internal Standards

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)
 Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".
 Y/N/N/A Are all internal standard recoveries within the 40-135% criteria?
 Y/N/N/A Was the S/N ratio all internal standard peaks ≥ 10 ?

#	Date	Lab ID/Reference	Internal Standard	% Recovery (Limit: 40-135%)	Qualifications
		1	8	29 (40-135)	Y/N/P (S, R)
		0			
		2	9	27	
			11	29	
			1	15	
		3	1	25	(CF-S, O-R)
		4	5	35	
			11	36	
			1	26	
		6	9	31	
			11	36	
			1	24	

Internal Standards		Check Standard Used	Internal Standards	Check Standard Used
A.	¹³ C-2,3,7,8-TCDF		G.	¹³ C-1,2,3,4,6,7,8-HpCDF
B.	¹³ C-2,3,7,8-TCDD		H.	¹³ C-1,2,3,4,6,7,8-HpCDD
C.	¹³ C-1,2,3,7,8-PeCDF		I.	¹³ C-OCDD
D.	¹³ C-1,2,3,7,8-PeCDD		K.	¹³ C-1,2,3,4-TCDD
E.	¹³ C-1,2,3,4,7,8-HxCDF		L.	¹³ C-1,2,3,7,8,9-HxCDD
F.	¹³ C-1,2,3,6,7,8-HxCDD			

LDC #: 31882
 SDG #: 222222

VALIDATION FINDINGS WORKSHEET
Compound Quantitation and Reported CRQLs

Page: 1 of 1
 Reviewer: [Signature]
 2nd Reviewer: _____

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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

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

#	Date	Sample ID	Finding	Associated Samples	Qualifications
		3	spk > calib range		
		4			
		6	H.L.K.L.O.P.R		
		M1	ZMPC results (2 flag)		

Comments: See sample calculation verification worksheet for recalculations

LDC#: 23188C21
 SDG#: See Cover

VALIDATION FINDINGS WORKSHEET
Field Duplicates

Page: 1 of 2
 Reviewer: [Signature]
 2nd Reviewer: _____

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

Y N NA
Y N NA

Were field duplicate pairs identified in this SDG?
 Were target analytes detected in the field duplicate pairs?

Compound	Concentration (pg/g)		(<50) RPD	(pg/g) Difference	(pg/g) Limits	Qualifications (Parent Only)
	1	SSAJ3-02-1BPC_FD				
A	0.22	1.1U		0.88	(≤1.1)	
B	0.90	5.3U		4.4	(≤5.3)	
C	0.99	5.3U		4.31	(≤5.3)	
D	1.2	0.13		1.07	(≤5.3)	
E	1.1	0.12		0.98	(≤5.3)	
F	1.6	0.41		1.19	(≤5.3)	
G	5.9	1.4		4.5	(≤11)	
H	0.90	0.60		0.3	(≤1.1)	
I	2.1	0.69		1.41	(≤5.3)	
J	1.5	0.39		1.11	(≤5.3)	
K	1.7	1.2		0.5	(≤5.3)	
L	2.2	0.76		1.44	(≤5.3)	
M	1.5	0.16		1.34	(≤5.3)	
N	1.4	0.18		1.22	(≤5.3)	
O	3.2	2.7		0.5	(≤5.3)	
P	2.1	0.99		1.11	(≤5.3)	
Q	9.1	7.8		1.3	(≤11)	

Compound	Concentration (pg/g)		(<50) RPD	(pg/g) Difference	(pg/g) Limits	Qualifications (Parent Only)
	5	SSAN7-02-1BPC_FD				
A	2.4	2.4		0	(≤1.1)	
B	11	10		1	(≤5.3)	
C	6.6	5.3		1.3	(≤5.3)	
D	15	12		3	(≤5.3)	
E	11	8.6		2.4	(≤5.3)	
F	39	35	11			
G	40	35		5	(≤11)	

LDC#: 23188C21
 SDG#: See Cover

VALIDATION FINDINGS WORKSHEET
Field Duplicates

Page: 2 of 3
 Reviewer: [Signature]
 2nd Reviewer: _____

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

Y N NA Were field duplicate pairs identified in this SDG?
Y N NA Were target analytes detected in the field duplicate pairs?

Compound	Concentration (pg/g)		(≤50)	(pg/g)	(pg/g)	Qualifications (Parent Only)
	5	SSAN7-02-1BPC_FD	RPD	Difference	Limits	
H	130	170	27			
I	160	210	27			
J	77	82	6			
K	230	250	8			
L	200	160	22			
M	44	35	23			
N	42	21		21	(≤5.3)	<i>data/A (+d)</i>
O	500	440	13			
P	270	220	20			
Q	1300	1200	8			

V:\FIELD DUPLICATES\23188C21.wpd

VALIDATION FINDINGS WORKSHEET
Initial Calibration Calculation Verification

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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

$RRF = (A_x)(C_s)/(A_s)(C_x)$
 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_s = Area of associated internal standard
 C_s = Concentration of internal standard
 X = Mean of the RRFs

#	Standard ID	Calibration Date	Compound (Reference Internal Standard)	Reported		Recalculated		Reported		Recalculated		
				Average RRF (Initial)	(%RSD)	Average RRF (Initial)	(%RSD)	RRF (CS3 std)	(%RSD)	RRF (CS3 std)	(%RSD)	
1	1012 (105)	1/10/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	0.860	10.4	0.87	10.4	0.87	10.4	0.87	10.6	
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)	0.934	12.9	0.95	12.9	0.95	12.9	0.95	12.8	
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)	1.058	11.2	1.09	11.2	1.09	11.2	1.09	11.0	
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)	0.998	12.2	1.05	12.2	1.05	12.2	1.05	12.2	
			OCDF (¹³ C-OCDD)	1.437	14.1	1.52	14.1	1.52	14.1	14.0		
2	1012	4/21/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	1.088	1.20	1.10	1.20	1.10	1.20	1.10	1.20	
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)									
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)									
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)									
			OCDF (¹³ C-OCDD)									
3	1012 (105)	4/12/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	0.945	4.44	0.98	4.44	0.98	4.44	0.98	4.33	
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)	1.021	8.03	1.04	8.03	1.04	8.03	1.04	8.97	
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)	1.114	5.33	1.19	5.33	1.19	5.33	1.19	5.25	
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)	1.072	3.60	1.11	3.60	1.11	3.60	1.11	3.75	
			OCDF (¹³ C-OCDD)	1.445	5.85	1.51	5.85	1.51	5.85	5.89		

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
Routine Calibration Results Verification

LDC #: 2318802
 SDG #: 21.0001

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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 = $100 * (\text{ave. RRF} - \text{RRF}) / \text{ave. RRF}$ Where: ave. RRF = initial calibration average RRF
 RRF = continuing calibration RRF
 $\text{RRF} = (A_x)(C_x) / (A_s)(C_s)$
 A_x = Area of compound, A_s = Area of associated internal standard
 C_x = Concentration of compound, C_s = Concentration of internal standard

#	Standard ID	Calibration Date	Compound (Reference Internal Standard)	Average RRF (Initial)	Reported		Recalculated	
					RRF (CC)	%D	RRF (CC)	%D
1	2AP10A125	4/26/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	0.860	0.92	7.0	0.92	7.0
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)	0.934	0.93	0.1	0.93	0.1
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)	1.058	1.15	8.9	1.15	8.9
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)	0.998	1.06	6.2	1.06	6.2
			OCDF (¹³ C-OCDD)	1.437	1.54	7.2	1.54	7.2
2	2AP10A05	4/27/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	0.945	0.97	3.1	0.97	3.1
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)	1.021	0.94	7.5	0.94	7.5
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)	1.114	1.10	1.2	1.10	1.2
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)	1.072	1.04	2.9	1.04	2.9
			OCDF (¹³ C-OCDD)	1.445	1.42	2.0	1.42	2.0
3	2AP105D2	4/28/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	1.088	1.09	0.3	1.09	0.3
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)					
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)					
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)					
			OCDF (¹³ C-OCDD)					

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: GC/MS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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 = $100 * |LCS - LCSD| / (LCS + LCSD)$ LCS = Laboratory control sample percent recovery LCSD = Laboratory control sample duplicate percent recovery

LCS ID: 010361

Compound	Spike Added		Spiked Sample Concentration		LCS		LCSD		Percent Recovery		Percent Recovery		RPD	
	LCS	LCSD	LCS	LCSD	Reported	Recalc.	Reported	Recalc.	Reported	Recalc.	Reported	Recalc.	Reported	Recalc.
2,3,7,8-TCDD	20.0	NA	20.4	NA	102	102								
1,2,3,7,8-PeCDD	100		109		109	109								
1,2,3,4,7,8-HxCDD			88.7		89	89								
1,2,3,4,7,8,9-HpCDF			88.2		88	88								
OCDF	200		205		102	102								

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.

Descriptor	Accurate mass ^(a)	Ion ID	Elemental Composition	Analyte	Descriptor	Accurate Mass ^(a)	Ion ID	Elemental Composition	Analyte		
1	303.9016	M	C ₁₂ H ₄ ³⁵ Cl ₂ O	TCDF	4	407.7818	M+2	C ₁₂ H ₃₅ Cl ₆ ³⁷ ClO	HpCDF		
	305.8987	M+2	C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ ClO	TCDF		409.7788	M+4	C ₁₂ H ₃₅ Cl ₆ ³⁷ Cl ₂ O	HpCDF		
	315.9419	M	¹³ C ₁₂ H ₄ ³⁵ Cl ₄ O	TCDF (S)		417.8250	M	¹³ C ₁₂ H ₃₅ Cl ₇ O	HpCDF (S)		
	317.9389	M+2	¹³ C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ ClO	TCDF (S)		419.8220	M+2	¹³ C ₁₂ H ₃₅ Cl ₆ ³⁷ ClO	HpCDF		
	319.8965	M	C ₁₂ H ₄ ³⁵ Cl ₄ O ₂	TCDD		423.7767	M+2	C ₁₂ H ₃₅ Cl ₆ ³⁷ ClO ₂	HpCDD		
	321.8936	M+2	C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ ClO ₂	TCDD		425.7737	M+4	C ₁₂ H ₃₅ Cl ₅ ³⁷ Cl ₂ O ₂	HpCDD		
	331.9368	M	¹³ C ₁₂ H ₄ ³⁵ Cl ₄ O ₂	TCDD (S)		435.8169	M+2	¹³ C ₁₂ H ₃₅ Cl ₆ ³⁷ ClO ₂	HpCDD (S)		
	333.9338	M+2	¹³ C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ ClO ₂	TCDD (S)		437.8140	M+4	¹³ C ₁₂ H ₃₅ Cl ₅ ³⁷ Cl ₂ O ₂	HpCDD (S)		
	375.8364	M+2	C ₁₂ H ₄ ³⁵ Cl ₅ ³⁷ ClO	HxCDFE		479.7165	M+4	C ₁₂ H ₃₅ Cl ₇ ³⁷ Cl ₂ O	HxCDFE		
	[354.9792]	LOCK	C ₉ F ₁₃	PFK		[430.9728]	LOCK	C ₉ F ₁₇	PFK		
	2	339.8597	M+2	C ₁₂ H ₃ ³⁵ Cl ₁ ³⁷ ClO		PeCDF	5	441.7428	M+2	C ₁₂ ³⁵ Cl ₇ ³⁷ ClO	OCDF
		341.8567	M+4	C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ Cl ₂ O		PeCDF		443.7399	M+4	C ₁₂ ³⁵ Cl ₆ ³⁷ Cl ₂ O	OCDF
		351.9000	M+2	¹³ C ₁₂ H ₃ ³⁵ Cl ₄ ³⁷ ClO		PeCDF (S)		457.7377	M+2	C ₁₂ ³⁵ Cl ₇ ³⁷ ClO ₂	OCDD
353.8970		M+4	¹³ C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ Cl ₂ O	PeCDF (S)	459.7348	M+4		C ₁₂ ³⁵ Cl ₆ ³⁷ Cl ₂ O ₂	OCDD		
355.8546		M+2	C ₁₂ H ₃ ³⁵ Cl ₄ ³⁷ ClO ₂	PeCDD	469.7780	M+2		¹³ C ₁₂ ³⁵ Cl ₇ ³⁷ ClO ₂	OCDD (S)		
357.8516		M+4	C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ Cl ₂ O ₂	PeCDD	471.7750	M+4		¹³ C ₁₂ ³⁵ Cl ₆ ³⁷ Cl ₂ O ₂	OCDD (S)		
367.8949		M+2	¹³ C ₁₂ H ₃ ³⁵ Cl ₄ ³⁷ ClO ₂	PeCDD (S)	513.6775	M+4		C ₁₂ ³⁵ Cl ₉ ³⁷ ClO	DCDFE		
369.8919		M+4	¹³ C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ Cl ₂ O ₂	PeCDD (S)	[422.9278]	LOCK		C ₁₀ F ₁₇	PFK		
409.7974		M+2	C ₁₂ H ₃ ³⁵ Cl ₅ ³⁷ ClO	HpCDFE							
[354.9792]		LOCK	C ₉ F ₁₃	PFK							
3		373.8208	M+2	C ₁₂ H ₂ ³⁵ Cl ₅ ³⁷ ClO	HxCDF						
		375.8178	M+4	C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ Cl ₂ O	HxCDF						
		383.8639	M	¹³ C ₁₂ H ₂ ³⁵ Cl ₆ O	HxCDF (S)						
	385.8610	M+2	¹³ C ₁₂ H ₂ ³⁵ Cl ₅ ³⁷ ClO	HxCDF (S)							
	389.8156	M+2	C ₁₂ H ₂ ³⁵ Cl ₅ ³⁷ ClO ₂	HxCDD							
	391.8127	M+4	C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ Cl ₂ O ₂	HxCDD							
	401.8559	M+2	¹³ C ₁₂ H ₂ ³⁵ Cl ₅ ³⁷ ClO ₂	HxCDD (S)							
	403.8529	M+4	¹³ C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ Cl ₂ O ₂	HxCDD (S)							
	445.7555	M+4	C ₁₂ H ₂ ³⁵ Cl ₆ ³⁷ Cl ₂ O	OCDFE							
	[430.9728]	LOCK	C ₉ F ₁₇	PFK							

(a) The following nucleidic masses were used:

- H = 1.007825
- C = 12.000000
- ¹³C = 13.003355
- F = 18.9984
- O = 15.994915
- ³⁵Cl = 34.968853
- ³⁷Cl = 36.965903

S = internal/recovery standard

#	Sample ID	Compound	Reported Concentration ()	Calculated Concentration ()	Qualification

- Concentration = $(A_2)(I_1)(DF) / (A_1)(I_2)(DF)$
- A_2 = Area of the characteristic ion (EICP) for the compound to be measured
- A_1 = Area of the characteristic ion (EICP) for the specific internal standard
- I_1 = Amount of internal standard added in nanograms (ng)
- V_0 = Volume or weight of sample extract in milliliters (ml) or grams (g)
- RRF = Relative Response Factor (average) from the initial calibration
- DF = Dilution Factor
- %S = Percent solids, applicable to soil and solid matrices only.

Example: $\frac{765820 (1.000) (10.32)}{1936250 (1.02) (10.32)} = 1.55$ B/d s/g

Conc. = $\frac{765820 (1.000) (10.32)}{1936250 (1.02) (10.32)}$

Sample I.D. _____ F _____

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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

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

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

Project/Site Name: Tronox LLC Facility, PCS, Henderson, Nevada
Collection Date: April 12, 2010
LDC Report Date: May 24, 2010
Matrix: Soil
Parameters: Dioxins/Dibenzofurans
Validation Level: Stage 4
Laboratory: TestAmerica, Inc.
Sample Delivery Group (SDG): GOD140526

Sample Identification

RSAI3-12BPC
RSAI3-12BPCMS
RSAI3-12BPCMSD

Introduction

This data review covers 3 soil samples listed on the cover sheet including dilutions and reanalysis as applicable. The analyses were per EPA SW 846 Method 8290 for Polychlorinated Dioxins/Dibenzofurans.

This review follows the Standard Operating Procedures (SOP) 40, Data Review/Validation (BRC 2009), the Quality Assurance Project Plan Tronox LLC Facility, Henderson, Nevada (June 2009), NDEP guidance (May 2006), and USEPA Contract Laboratory Program National Functional Guidelines for Polychlorinated Dioxins/Dibenzofurans Data Review (September 2005).

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.

Blank results are summarized in Section V.

Field duplicates are summarized in Section XIV.

The following are definitions of the data qualifiers:

- J+ Data are qualified as estimated, with a high bias likely to occur. False positives or false negatives are unlikely to have been reported.
- J- Data are qualified as estimated, with a low bias likely to occur. False positives or false negatives are unlikely to have been reported.
- J Data are qualified as estimated; it is not possible to assess the direction of the potential bias. False positives or false negatives are unlikely to have been reported.
- U Indicates the compound or analyte was analyzed for but not detected at or above the stated limit.
- R Data are qualified as rejected. There is a significant potential for the reporting of false negatives or false positives.
- UU Indicates the compound or analyte was analyzed for but not detected. The sample detection limit is an estimated value.
- B The analytical result may be a false positive totally attributable to blank contamination. This qualifier is applicable to radiochemistry analysis only.
- JB The analytical result may be biased high and partially attributable to blank contamination. This qualifier is applicable to radiochemistry analysis only.
- JK The analytical result is an estimated maximum possible concentration (EMPC).
- X The analytical result is not used for reporting because a more accurate and precise result is reported in its place.
- J-TDS The analytical result is estimated based on failure of the Total Dissolved Solids (TDS) correctness check performed in accordance with the Standard Method 1030E.
- J-CAB The analytical result is estimated based on failure of the cation-anion balance correctness check performed in accordance with Standard Method 1030E.
- J-TDS & CAB The analytical result is unreliable based on the failure of the cation-anion balance and TDS correctness check performed in accordance with standard Method 1030E.
- A Indicates the finding is based upon technical validation criteria.
- P Indicates the finding is related to a protocol/contractual deviation.
- None Indicates the data was not significantly impacted by the finding, therefore qualification was not required.

I. Technical Holding Times

All technical holding time requirements were met.

The chain-of-custodies were reviewed for documentation of cooler temperatures. All cooler temperatures met validation criteria.

II. HRGC/HRMS Instrument Performance Check

Instrument performance was checked at the required daily 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 exact mass of 380.9760 of PFK was verified. The static resolving power was at least 10,000 (10% valley definition).

III. Initial Calibration

A five point initial calibration was performed as required by the method.

Percent relative standard deviations (%RSD) were less than or equal to 20.0% for unlabeled compounds and less than or equal to 30.0% for labeled compounds.

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

The minimum S/N ratio for each target compound was greater than or equal to 2.5 and greater than or equal to 10 for each recovery and internal standard compound for samples.

IV. Routine Calibration (Continuing)

Routine calibration was performed at the required frequencies.

All of the routine calibration percent differences (%D) between the initial calibration RRF and the routine calibration RRF were less than or equal to 20.0% for unlabeled compounds and less than or equal to 30.0% for labeled compounds.

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

V. Blanks

Method blanks were reviewed for each matrix as applicable. No polychlorinated dioxin/dibenzofuran contaminants were found in the method blanks with the following exceptions:

Method Blank ID	Extraction Date	Compound	Concentration	Associated Samples
0109260MB	4/19/10	1,2,3,4,6,7,8-HpCDD OCDD 2,3,7,8-TCDF 1,2,3,7,8-PeCDF 1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 1,2,3,7,8,9-HxCDF 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	0.14 pg/g 0.59 pg/g 0.28 pg/g 0.14 pg/g 0.24 pg/g 0.14 pg/g 0.086 pg/g 0.30 pg/g 0.13 pg/g 0.63 pg/g	All samples in SDG G0D140526

Sample concentrations were compared to concentrations detected in the method blanks as required by the QAPP. No sample data was qualified.

Sample EB-04122010-RIG3-RZD (from SDG G0D140534) was identified as an equipment blank. No polychlorinated dioxin/dibenzofuran contaminants were found in these blanks with the following exceptions:

Equipment Blank ID	Sampling Date	Compound	Concentration	Associated Samples
EB-04122010-RIG3-RZD	4/12/10	1,2,3,6,7,8-HxCDD 1,2,3,7,8,9-HxCDD 1,2,3,4,6,7,8-HpCDD OCDD 2,3,7,8-TCDF 1,2,3,7,8-PeCDF 2,3,4,7,8-PeCDF 1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 2,3,4,6,7,8-HxCDF 1,2,3,7,8,9-HxCDF 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	0.64 pg/L 0.47 pg/L 1.9 pg/L 5.0 pg/L 4.6 pg/L 3.8 pg/L 2.2 pg/L 7.7 pg/L 4.8 pg/L 1.2 pg/L 0.59 pg/L 14 pg/L 5.0 pg/L 27 pg/L	All samples in SDG G0D140526

Sample concentrations were compared to concentrations detected in the equipment blanks as required by the QAPP. No sample data was qualified.

Sample FB-04072010-RZD (from SDG G0D090441) was identified as a field blank. No polychlorinated dioxin/dibenzofuran contaminants were found in this blank with the following exceptions:

Field Blank ID	Sampling Date	Compound	Concentration	Associated Samples
FB-04072010-RZD	4/7/10	1,2,3,4,7,8-HxCDD 1,2,3,7,8,9-HxCDD 1,2,3,4,6,7,8-HpCDD OCDD 1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 2,3,4,6,7,8-HxCDF 1,2,3,7,8,9-HxCDF 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	0.89 pg/L 1.5 pg/L 2.2 pg/L 8.3 pg/L 1.4 pg/L 1.6 pg/L 1.5 pg/L 1.6 pg/L 1.3 pg/L 1.4 pg/L 4.1 pg/L	All samples in SDG G0D140526

Sample concentrations were compared to concentrations detected in the field blanks as required by the QAPP. No sample data was qualified.

VI. Matrix Spike/Matrix Spike Duplicates

Matrix spike (MS) and matrix spike duplicate (MSD) samples were reviewed for each matrix as applicable. Percent recoveries (%R) and relative percent differences (RPD) were not within the QC limits. Since the samples were diluted out, no data were qualified.

VII. Laboratory Control Samples (LCS)

Laboratory control samples were reviewed for each matrix as applicable. Percent recoveries (%R) were within QC limits.

VIII. Regional Quality Assurance and Quality Control

Not applicable.

IX. Internal Standards

All internal standard recoveries were within QC limits with the following exceptions:

Sample	Internal Standards	%R (Limits)	Compound	Flag	A or P
RSAI3-12BPC	¹³ C-1,2,3,4,6,7,8-HpCDF ¹³ C-1,2,3,4,6,7,8-HpCDD ¹³ C-OCDD	31 (40-135) 22 (40-135) 20 (40-135)	1,2,3,4,6,7,8-HpCDD OCDD 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	J (all detects) UJ (all non-detects)	P

X. Target Compound Identifications

All target compound identifications were within validation criteria.

XI. Project Quantitation Limit

All compound quantitation and CRQLs were within validation criteria with the following exceptions:

Sample	Compound	Finding	Criteria	Flag	A or P
RSAI3-12BPC	1,2,3,4,7,8-HxCDF 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	Sample result exceeded calibration range.	Reported result should be within calibration range.	J (all detects) J (all detects) J (all detects) J (all detects)	P

All compounds reported below the PQL were qualified as follows:

Sample	Finding	Flag	A or P
All samples in SDG G0D140526	All compounds reported below the PQL.	J (all detects)	A

All compounds reported as EMPC were qualified as follows:

Sample	Finding	Flag	A or P
All samples in SDG G0D140526	All compounds reported as estimated maximum possible concentration (EMPC).	JK (all detects)	A

XII. System Performance

The system performance was acceptable.

XIII. Overall Assessment of Data

Data flags are summarized at the end of this report if data has been qualified.

XIV. Field Duplicates

No field duplicates were identified in this SDG.

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Data Qualification Summary - SDG G0D140526**

SDG	Sample	Compound	Flag	A or P	Reason (Code)
G0D140526	RSAI3-12BPC	1,2,3,4,6,7,8-HpCDD OCDD 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	J (all detects) UJ (all non-detects)	P	Internal standards (%R) (i)
G0D140526	RSAI3-12BPC	1,2,3,4,7,8-HxCDF 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	J (all detects) J (all detects) J (all detects) J (all detects)	P	Compound quantitation and CRQLs (e)
G0D140526	RSAI3-12BPC	All compounds reported below the PQL.	J (all detects)	A	Project Quantitation Limit (sp)
G0D140526	RSAI3-12BPC	All compounds reported as EMPC	JK (all detects)	A	Project Quantitation Limit (k)

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Laboratory Blank Data Qualification Summary - SDG
G0D140526**

No Sample Data Qualified in this SDG

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Equipment Blank Data Qualification Summary - SDG
G0D140526**

No Sample Data Qualified in this SDG

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Field Blank Data Qualification Summary - SDG G0D140526**

No Sample Data Qualified in this SDG

Tronox Northgate Henderson

VALIDATION COMPLETENESS WORKSHEET

LDC #: 23188D21
 SDG #: G0D140526
 Laboratory: Test America

Stage 4

Date: 5/21/10
 Page: 1 of 1
 Reviewer: [Signature]
 2nd Reviewer:

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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.	Technical holding times	A	Sampling dates: 4/12/10
II.	HRGC/HRMS Instrument performance check	A	
III.	Initial calibration	A	
IV.	Routine calibration/SV	A	
V.	Blanks	TW	
VI.	Matrix spike/Matrix spike duplicates	TW	
VII.	Laboratory control samples	A	LC9
VIII.	Regional quality assurance and quality control	N	
IX.	Internal standards	TW	
X.	Target compound identifications	A	
XI.	Compound quantitation and CRQLs	TW	
XII.	System performance	A	
XIII.	Overall assessment of data	A	
XIV.	Field duplicates	N	
XV.	Field blanks	TW	ZB-04122010-R193-R2D(G0D140534) FB-04072010-R2D(G0D090441)

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

Validated Samples:

1	RSAI3-12BPC	S	11	0109260MB	21		31	
2	RSAI3-12BPCMS		12		22		32	
3	RSAI3-12BPCMSD		13		23		33	
4			14		24		34	
5			15		25		35	
6			16		26		36	
7			17		27		37	
8			18		28		38	
9			19		29		39	
10			20		30		40	

Notes: _____

Method: Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

Validation Area	Yes	No	NA	Findings/Comments
I. Technical holding times				
All technical holding times were met.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Cooler temperature criteria was met.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
II. GC/MS Instrument performance check				
Was PFK exact mass 380.9760 verified?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were the retention time windows established for all homologues?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was the chromatographic resolution between 2,3,7,8-TCDD and peaks representing any other unlabeled TCDD isomers $\leq 25\%$?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Is the static resolving power at least 10,000 (10% valley definition)?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was the mass resolution adequately check with PFK?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was the presence of 1,2,8,9-TCDD and 1,3,4,6,8-PeCDF verified?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
III. Initial calibration				
Was the initial calibration performed at 5 concentration levels?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were all percent relative standard deviations (%RSD) $\leq 20\%$ for unlabeled standards and $< 30\%$ for labeled standards?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Did all calibration standards meet the Ion Abundance Ratio criteria?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was the signal to noise ratio for each target compound ≥ 2.5 and for each recovery and internal standard ≥ 10 ?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
IV. Continuing calibration				
Was a routine calibration performed at the beginning and end of each 12 hour period?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were all percent differences (%D) $\leq 20\%$ for unlabeled standards and $\leq 30\%$ for labeled standards?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Did all routine calibration standards meet the Ion Abundance Ratio criteria?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
V. Blanks				
Was a method blank associated with every sample in this SDG?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was a method blank performed for each matrix and concentration?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was there contamination in the method blanks? If yes, please see the Blanks validation completeness worksheet?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
VI. Matrix spike/Matrix spike duplicates				
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.	<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"/>	
VII. Laboratory control samples				
Was an LCS analyzed for this SDG?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
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"/>	

LDC #: 23188621
 SDG #: SERCONEX

VALIDATION FINDINGS CHECKLIST

Page: 2 of 2
 Reviewer: [Signature]
 2nd Reviewer: _____

VIII. Regional Quality Assurance and Quality Control				
Were performance evaluation (PE) samples performed?			<input checked="" type="checkbox"/>	
Were the performance evaluation (PE) samples within the acceptance limits?			<input checked="" type="checkbox"/>	
IX. Internal standards				
Were internal standard recoveries within the 40-135% criteria?			<input checked="" type="checkbox"/>	
Was the minimum S/N ratio of all internal standard peaks > 10?			<input checked="" type="checkbox"/>	
X. Target compound identification				
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?			<input checked="" type="checkbox"/>	
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?			<input checked="" type="checkbox"/>	
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?			<input checked="" type="checkbox"/>	
Did compound spectra contain all characteristic ions listed in the table attached?			<input checked="" type="checkbox"/>	
Was the Ion Abundance Ratio for the two quantitation ions within criteria?			<input checked="" type="checkbox"/>	
Was the signal to noise ratio for each target compound and labeled standard > 2.5?			<input checked="" type="checkbox"/>	
Does the maximum intensity of each specified characteristic ion coincide within ± 2 seconds (includes labeled standards)?			<input checked="" type="checkbox"/>	
For PCDF identification, was any signal (S/N ≥ 2.5 , at \pm seconds RT) detected in the corresponding PCDPE channel?			<input checked="" type="checkbox"/>	
Was an acceptable lock mass recorded and monitored?			<input checked="" type="checkbox"/>	
XI. Compound quantitation/CRQLs				
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 CRQLs adjusted to reflect all sample dilutions and dry weight factors applicable to level IV validation?			<input checked="" type="checkbox"/>	
XII. System performance				
System performance was found to be acceptable.			<input checked="" type="checkbox"/>	
XIII. Overall assessment of data				
Overall assessment of data was found to be acceptable.			<input checked="" type="checkbox"/>	
XIV. Field duplicates				
Field duplicate pairs were identified in this SDG.			<input checked="" type="checkbox"/>	
Target compounds were detected in the field duplicates.			<input checked="" type="checkbox"/>	
XV. Field blanks				
Field blanks were identified in this SDG.			<input checked="" type="checkbox"/>	
Target compounds were detected in the field blanks.			<input checked="" type="checkbox"/>	

VALIDATION FINDINGS WORKSHEET

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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:

METHOD: HRGC/HRMS Dioxins (EPA Method 8290)

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? If yes, please see qualification below.

Blank extraction date: 4/19/10 Blank analysis date: 4/25/10

Conc. units: pg/L Associated samples: All (>5X)

Compound	Blank ID	5X	Sample Identification																	
	0109260MB	5X																		
F	0.14	0.7																		
G	0.59	2.95																		
H	0.28	1.4																		
I	0.14	0.7																		
K	0.24	1.2																		
L	0.14	0.7																		
N	0.086	0.43																		
O	0.30	1.5																		
P	0.13	0.65																		
Q	0.63	3.15																		

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".

LDC #: 23188D21
 SDG #: See Cover

VALIDATION FINDINGS WORKSHEET

Field Blanks

Page: 1 of 1
 Reviewer: [Signature]
 2nd Reviewer: _____

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

Y N N/A Were field blanks identified in this SDG?

Blank units: pg/L Associated sample units: pg/g

Sampling date: 4/12/10

Field blank type: (circle one) Field Blank / Rinsate / Other: EB

Associated Samples: All (51)

Compound	Blank ID	5X	Sample Identification				
	EB-04122010-RIG3-RZD	5X					
D	0.64	0.0032					
E	0.47	0.00235					
F	1.9	0.0095					
G	5.0	0.025					
H	4.6	0.023					
I	3.8	0.019					
J	2.2	0.011					
K	7.7	0.0385					
L	4.8	0.024					
M	1.2	0.006					
N	0.59	0.00295					
O	14	0.07					
P	5.0	0.025					
Q	27	0.135					
CRQL							

CIRCLED RESULTS WERE NOT QUALIFIED. ALL RESULTS NOT CIRCLED WERE QUALIFIED BY THE FOLLOWING STATEMENT:
 Samples with compound concentrations within five times the associated field blank concentration are listed above, these sample results were qualified as not detected, "U".

LDC #: 23188D21
SDG #: See Cover

VALIDATION FINDINGS WORKSHEET
Field Blanks

Page: 1 of 1
Reviewer: _____
2nd Reviewer: _____

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

Y N N/A Were field blanks identified in this SDG?

Blank units: _____ pg/L Associated sample units: _____ pg/g

Sampling date: 4/7/10

Field blank type: (circle one) Field Blank / Rinsate / Other: _____ Associated Samples: All (✓ SX)

Compound	Blank ID	Blank ID	Sample Identification
C	0.89	FR-04072010-RZD	5X
E	1.5		
F	2.2		
G	8.3		
K	1.4		
L	1.6		
M	1.5		
N	1.6		
O	1.3		
P	1.4		
Q	4.1		
CRQL			

CIRCLED RESULTS WERE NOT QUALIFIED. ALL RESULTS NOT CIRCLED WERE QUALIFIED BY THE FOLLOWING STATEMENT:
Samples with compound concentrations within five times the associated field blank concentration are listed above, these sample results were qualified as not detected, "U".

LDC #: 238827
 SDG #: 210001

VALIDATION FINDINGS WORKSHEET
Compound Quantitation and Reported CRQLs

Page: 1 of 4
 Reviewer: _____
 2nd Reviewer: _____

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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

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

#	Date	Sample ID	Finding	Associated Samples	Qualifications
		1	Spds > Calib curve	1	Blank (e)
			K.O.P. &		
			ZMPC results (2 Yag)		YK (K)

Comments: See sample calculation verification worksheet for recalculations

VALIDATION FINDINGS WORKSHEET
Initial Calibration Calculation Verification

Page: 1 of 1
 Reviewer: Q
 2nd Reviewer: _____

SDG #: 282004
Seawater

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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	
				Average RRF (initial)	(%RSD)	Average RRF (Initial)	(%RSD)	RRF (CS3 std)	(%RSD)	RRF (CS3 std)	(%RSD)
1	ICAR (SDS)	4/12/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	0.945	4.44	0.945	4.33	0.98	4.44	0.98	4.33
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)	1.021	3.03	1.021	2.97	1.04	3.03	1.04	2.97
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)	1.114	5.33	1.114	5.25	1.19	5.33	1.19	5.25
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)	1.072	3.60	1.072	3.75	1.11	3.60	1.11	3.75
			OCDF (¹³ C-OCDF)	1.445	5.85	1.445	5.89	1.51	5.85	1.51	5.89
2	ICAR	3/10/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	1.000	1.36	1.000	1.58	1.00	1.36	1.00	1.58
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)								
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)								
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)								
			OCDF (¹³ C-OCDF)								
3	ICAR (SDS)	4/21/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	1.088	1.29	1.088	1.20	1.10	1.29	1.10	1.20
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)								
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)								
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)								
			OCDF (¹³ C-OCDF)								

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
Routine Calibration Results Verification

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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 = $100 * (\text{ave. RRF} - \text{RRF}) / \text{ave. RRF}$ Where: ave. RRF = initial calibration average RRF
 RRF = $(A_x)(C_{is}) / (A_s)(C_x)$ RRF = continuing calibration RRF
 A_x = Area of compound, A_s = Area of associated internal standard
 C_x = Concentration of compound, C_s = Concentration of internal standard

#	Standard ID	Calibration Date	Compound (Reference Internal Standard)	Average RRF (Initial)	Reported		Recalculated		Reported		Recalculated	
					RRF (CC)	%D	RRF (CC)	%D	RRF (CC)	%D		
1	2378-TCDF	4/24/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	0.945	0.90	4.3	0.90	0.90	0.90	4.3	4.3	4.3
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)	1.027	0.96	6.3	0.96	0.96	0.96	6.3	6.3	6.3
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)	1.114	1.08	3.1	1.08	1.08	1.08	3.1	3.1	3.1
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)	1.072	1.03	3.9	1.03	1.03	1.03	3.9	3.9	3.9
			OCDF (¹³ C-OCDD)	1.445	1.36	5.9	1.36	1.36	1.36	5.9	5.9	5.9
2	2378-TCDF	5/3/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	0.945	0.93	1.3	0.93	0.93	0.93	1.3	1.3	1.3
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)	1.027	0.95	6.8	0.95	0.95	0.95	6.8	6.8	6.8
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)	1.114	1.15	3.4	1.15	1.15	1.15	3.4	3.4	3.4
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)	1.072	1.01	5.4	1.01	1.01	1.01	5.4	5.4	5.4
			OCDF (¹³ C-OCDD)	1.445	1.40	3.4	1.40	1.40	1.40	3.4	3.4	3.4
3	2378-TCDF	5/2/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	1.088	1.12	2.8	1.12	1.12	1.12	2.8	2.8	2.8
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)									
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)									
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)									
	OCDF (¹³ C-OCDD)											

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
Matrix Spike/Matrix Spike Duplicates Results Verification

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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 = $100 * (SSR - SR) / SA$ Where: SSR = Spiked sample result, SR = Sample result
 SA = Spike added

RPD = $|MSR - MSDR| * 2 / (MSR + MSDR)$

MSR = Matrix spike percent recovery MSDR = Matrix spike duplicate percent recovery

MS/MSD samples: 0/3

Compound	Spike Added (ppg)		Sample Concentration (ppg)		Spiked Sample Concentration (ppg)		Matrix Spike Percent Recovery		Matrix Spike Duplicate Percent Recovery		Reported	Recalculated
	MS	MSD	MS	MSD	MS	MSD	Reported	Recalc	Reported	Recalc	RPD	RPD
	2,3,7,8-TCDD	2.1	20.3	51.0		488	436	0	0	0	0	0
1,2,3,7,8-PeCDD	105	101	1700		1740	1530	833	38	0	0	0	13
1,2,3,4,7,8-HxCDD			1400		1570	1180	191	162	0	0	0	28
1,2,3,4,7,8,9-HpCDF			62000		73900	64100	1160	1133	2430	208	14	14
OCDF	211	203	37000		41000	33900	2000	1896	0	0	0	19

Comments: Refer to Matrix Spike/Matrix Spike 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
Laboratory Control Sample Results Verification

METHOD: GC/MS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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 = $100 \cdot \text{SSC}/\text{SA}$ Where: SSC = Spiked sample concentration
 SA = Spike added

RPD = $100 \cdot \text{LCS} - \text{LCSD} / 2(\text{LCS} + \text{LCSD})$ LCS = Laboratory control sample percent recovery LCSD = Laboratory control sample duplicate percent recovery

LCS ID: 0109260

Compound	Spike Added		Spiked Sample Concentration		LCS		LCSD		Percent Recovery		Percent Recovery		RPD	
	LCS	LCSD	LCS	LCSD	Reported	Recalc	Reported	Recalc	Reported	Recalc	Reported	Recalc	Reported	Recalc
	2,3,7,8-TCDD	20.0	NA	19.8	NA	99	99	99	99					
1,2,3,7,8-PeCDD	100	↓	102	↓	102	102	102	102						
1,2,3,4,7,8-HxCDD	↓	↓	108	↓	108	108	108	108						
1,2,3,4,7,8,9-HpCDF	200	↓	104	↓	104	104	104	104						
OCDF			200	↓	101	101	101	101						

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.

Descriptor	Accurate mass ^(a)	Ion ID	Elemental Composition	Analyte	Descriptor	Accurate Mass ^(a)	Ion ID	Elemental Composition	Analyte
1	303.9016 305.8987 315.9419 317.9389 319.8965 321.8936 331.9368 333.9338 375.8364 [354.9792]	M M+2 M M+2 M M+2 M M+2 M+2 LOCK	C ₁₂ H ₂ ³⁵ Cl ₄ O C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ Cl ₁₀ C ₁₂ H ₄ ³⁵ Cl ₄ O ¹³ C ₁₂ H ₂ ³⁵ Cl ₃ ³⁷ ClO C ₁₂ H ₂ ³⁵ Cl ₃ ³⁷ ClO C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ Cl ₂ O ₂ C ₁₂ H ₂ ³⁵ Cl ₃ ³⁷ ClO ₂ ¹³ C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ ClO C ₉ F ₁₃	TCDF TCDF TCDF (S) TCDF (S) TCDD TCDD TCDD (S) TCDD (S) HxCDPE PFK	4	407.7818 409.7788 417.8250 419.8220 423.7767 425.7737 435.8169 437.8140 479.7165 [430.9728]	M+2 M+4 M M+2 M+2 M+4 M+2 M+4 M+4 LOCK	C ₁₂ H ₂ ³⁵ Cl ₉ ³⁷ ClO C ₁₂ H ₂ ³⁵ Cl ₈ ³⁷ Cl ₂ O ¹³ C ₁₂ H ₂ ³⁵ Cl ₇ O ¹³ C ₁₂ H ₄ ³⁵ Cl ₆ ³⁷ ClO C ₁₂ H ₄ ³⁵ Cl ₆ ³⁷ ClO ₂ C ₁₂ H ₂ ³⁵ Cl ₅ ³⁷ ClO ₂ C ₁₂ H ₄ ³⁵ Cl ₅ ³⁷ ClO ₂ ¹³ C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ ClO ₂ ¹³ C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ Cl ₂ O ₂ C ₁₂ H ₂ ³⁵ Cl ₃ ³⁷ ClO C ₉ F ₁₇	HpCDF HpCDF HpCDF (S) HpCDF HpCDD HpCDD HpCDD (S) HpCDD (S) HxCDDPE PFK
2	339.8597 341.8567 351.9000 353.8970 355.8546 357.8516 367.8949 369.8919 409.7974 [354.9792]	M+2 M+4 M+2 M+4 M+2 M+4 M+2 M+4 M+2 LOCK	C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ Cl ₂ O ¹³ C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO ¹³ C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO C ₁₂ H ₃ ³⁵ Cl ₄ ³⁷ ClO C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO ₂ ¹³ C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO ₂ ¹³ C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO C ₉ F ₁₃	PeCDF PeCDF PeCDF (S) PeCDF (S) PeCDD PeCDD PeCDD (S) PeCDD (S) HxCDDPE PFK	5	441.7428 443.7399 457.7377 459.7348 469.7780 471.7750 513.6775 [422.9278]	M+2 M+4 M+2 M+4 M+2 M+4 M+4 M+4 LOCK	C ₁₂ H ₃ ³⁵ Cl ₇ ³⁷ ClO C ₁₂ H ₃ ³⁵ Cl ₆ ³⁷ Cl ₂ O C ₁₂ H ₃ ³⁵ Cl ₇ ³⁷ ClO ₂ C ₁₂ H ₃ ³⁵ Cl ₆ ³⁷ ClO ₂ ¹³ C ₁₂ H ₃ ³⁵ Cl ₇ ³⁷ ClO ¹³ C ₁₂ H ₃ ³⁵ Cl ₆ ³⁷ ClO ₂ C ₁₂ H ₃ ³⁵ Cl ₆ ³⁷ ClO C ₁₀ F ₁₇	OCDF OCDF OCDD OCDD OCDD (S) OCDD (S) DCDDPE PFK
3	373.8208 375.8178 383.8639 385.8610 389.8156 391.8127 401.8559 403.8529 445.7555 [430.9728]	M+2 M+4 M M+2 M+2 M+4 M+2 M+4 M+4 LOCK	C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ ClO C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ Cl ₂ O ¹³ C ₁₂ H ₂ ³⁵ Cl ₃ O ¹³ C ₁₂ H ₂ ³⁵ Cl ₃ ³⁷ ClO C ₁₂ H ₂ ³⁵ Cl ₅ ³⁷ ClO C ₁₂ H ₂ ³⁵ Cl ₅ ³⁷ ClO ₂ ¹³ C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ ClO ₂ ¹³ C ₁₂ H ₂ ³⁵ Cl ₅ ³⁷ ClO ₂ ¹³ C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ Cl ₂ O ₂ C ₁₂ H ₂ ³⁵ Cl ₃ ³⁷ Cl ₂ O C ₉ F ₁₇	HxCDF HxCDF HxCDF (S) HxCDF (S) HxCDD HxCDD HxCDD (S) HxCDD (S) HxCDDPE PFK			M+2 M+4 M M+2 M+2 M+4 M+2 M+4 M+4 LOCK		

(a) The following nuclidic masses were used:

- H = 1.007825
- C = 12.000000
- ¹³C = 13.003355
- F = 18.9984
- O = 15.994915
- ³⁵Cl = 34.968853
- ³⁷Cl = 36.965903

S = internal/recovery standard

LDC #: 2318802
SDG #: SEC-COUB

VALIDATION FINDINGS WORKSHEET
Sample Calculation Verification

Page: 4 of 1
Reviewer: [Signature]
2nd reviewer: _____

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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?

$$\text{Concentration} = \frac{(A_x)(L)(DF)}{(A_s)(RRF)(V_s)(\%S)}$$

- A_x = Area of the characteristic ion (EICP) for the compound to be measured
- A_s = Area of the characteristic ion (EICP) for the specific internal standard
- I_s = Amount of internal standard added in nanograms (ng)
- V_s = Volume or weight of sample extract in milliliters (ml) or grams (g).
- RRF = Relative Response Factor (average) from the initial calibration
- Df = Dilution Factor.
- %S = Percent solids, applicable to soil and solid matrices only.

Example:

Sample I.D. 1, A:

$$\text{Conc.} = \frac{(4113000)(2000)}{(161743)(1.02)(10.56)(0.96)}$$

= 509.4 ppb/g

#	Sample ID	Compound	Reported Concentration ()	Calculated Concentration ()	Qualification

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

Project/Site Name: Tronox LLC Facility, PCS, Henderson, Nevada

Collection Date: April 12, 2010

LDC Report Date: May 26, 2010

Matrix: Water

Parameters: Dioxins/Dibenzofurans

Validation Level: Stage 2B

Laboratory: TestAmerica, Inc.

Sample Delivery Group (SDG): G0D140534

Sample Identification

EB-04122010-RIG3-RZD

EB-04122010-RIG2-RZB

EB-04122010-RIG1-RZB

Introduction

This data review covers 3 water samples listed on the cover sheet including dilutions and reanalysis as applicable. The analyses were per EPA SW 846 Method 8290 for Polychlorinated Dioxins/Dibenzofurans.

This review follows the Standard Operating Procedures (SOP) 40, Data Review/Validation (BRC 2009), the Quality Assurance Project Plan Tronox LLC Facility, Henderson, Nevada (June 2009), NDEP guidance (May 2006), and USEPA Contract Laboratory Program National Functional Guidelines for Polychlorinated Dioxins/Dibenzofurans Data Review (September 2005).

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.

Blank results are summarized in Section V.

Field duplicates are summarized in Section XIV.

Raw data were not reviewed for this SDG. The review was based on QC data.

The following are definitions of the data qualifiers:

- J+ Data are qualified as estimated, with a high bias likely to occur. False positives or false negatives are unlikely to have been reported.
- J- Data are qualified as estimated, with a low bias likely to occur. False positives or false negatives are unlikely to have been reported.
- J Data are qualified as estimated; it is not possible to assess the direction of the potential bias. False positives or false negatives are unlikely to have been reported.
- U Indicates the compound or analyte was analyzed for but not detected at or above the stated limit.
- R Data are qualified as rejected. There is a significant potential for the reporting of false negatives or false positives.
- UJ Indicates the compound or analyte was analyzed for but not detected. The sample detection limit is an estimated value.
- B The analytical result may be a false positive totally attributable to blank contamination. This qualifier is applicable to radiochemistry analysis only.
- JB The analytical result may be biased high and partially attributable to blank contamination. This qualifier is applicable to radiochemistry analysis only.
- JK The analytical result is an estimated maximum possible concentration (EMPC).
- X The analytical result is not used for reporting because a more accurate and precise result is reported in its place.
- J-TDS The analytical result is estimated based on failure of the Total Dissolved Solids (TDS) correctness check performed in accordance with the Standard Method 1030E.
- J-CAB The analytical result is estimated based on failure of the cation-anion balance correctness check performed in accordance with Standard Method 1030E.
- J-TDS & CAB The analytical result is unreliable based on the failure of the cation-anion balance and TDS correctness check performed in accordance with standard Method 1030E.
- A Indicates the finding is based upon technical validation criteria.
- P Indicates the finding is related to a protocol/contractual deviation.
- None Indicates the data was not significantly impacted by the finding, therefore qualification was not required.

I. Technical Holding Times

All technical holding time requirements were met.

The chain-of-custodies were reviewed for documentation of cooler temperatures. All cooler temperatures met validation criteria.

II. HRGC/HRMS Instrument Performance Check

Instrument performance was checked at the required daily 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% .

III. Initial Calibration

A five point initial calibration was performed as required by the method.

Percent relative standard deviations (%RSD) were less than or equal to 20.0% for unlabeled compounds and less than or equal to 30.0% for labeled compounds.

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

IV. Routine Calibration (Continuing)

Routine calibration was performed at the required frequencies.

All of the routine calibration percent differences (%D) between the initial calibration RRF and the routine calibration RRF were less than or equal to 20.0% for unlabeled compounds and less than or equal to 30.0% for labeled compounds with the following exceptions:

Date	Compound	%D	Associated Samples	Affected Compound	Flag	A or P
4/26/10	1,2,3,4,7,8-HxCDD	20.4	0110418MB	1,2,3,4,7,8-HxCDD	J- (all detects) UJ (all non-detects)	P

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

V. Blanks

Method blanks were reviewed for each matrix as applicable. No polychlorinated dioxin/dibenzofuran contaminants were found in the method blanks with the following exceptions:

Method Blank ID	Extraction Date	Compound	Concentration	Associated Samples
0110418MB	4/20/10	1,2,3,4,7,8-HxCDD	1.3 pg/L	All samples in SDG GOD140534
		1,2,3,6,7,8-HxCDD	1.3 pg/L	
		1,2,3,7,8,9-HxCDD	1.8 pg/L	
		1,2,3,4,6,7,8-HpCDD	2.7 pg/L	
		OCDD	11 pg/L	
		2,3,7,8-TCDF	1.8 pg/L	
		1,2,3,7,8-PeCDF	1.8 pg/L	
		2,3,4,7,8-PeCDF	2.2 pg/L	
		1,2,3,4,7,8-HxCDF	1.6 pg/L	
		1,2,3,6,7,8-HxCDF	1.4 pg/L	
		2,3,4,6,7,8-HxCDF	1.8 pg/L	
		1,2,3,7,8,9-HxCDF	1.7 pg/L	
		1,2,3,4,6,7,8-HpCDF	1.9 pg/L	
		1,2,3,4,7,8,9-HpCDF	2.0 pg/L	
		OCDF	4.2 pg/L	

Sample concentrations were compared to concentrations detected in the method blanks as required by the QAPP. No sample data was qualified with the following exceptions:

Sample	Compound	Reported Concentration	Modified Final Concentration
EB-04122010-RIG3-RZD	1,2,3,6,7,8-HxCDD	0.64 pg/L	0.64U pg/L
	1,2,3,7,8,9-HxCDD	0.47 pg/L	0.47U pg/L
	1,2,3,4,6,7,8-HpCDD	1.9 pg/L	1.9U pg/L
	OCDD	5.0 pg/L	5.0U pg/L
	2,3,7,8-TCDF	4.6 pg/L	4.6U pg/L
	1,2,3,7,8-PeCDF	3.8 pg/L	3.8U pg/L
	2,3,4,7,8-PeCDF	2.2 pg/L	2.2U pg/L
	1,2,3,4,7,8-HxCDF	7.7 pg/L	7.7U pg/L
	1,2,3,6,7,8-HxCDF	4.8 pg/L	4.8U pg/L
	2,3,4,6,7,8-HxCDF	1.2 pg/L	1.2U pg/L
	1,2,3,7,8,9-HxCDF	0.59 pg/L	0.59U pg/L
	1,2,3,4,7,8,9-HpCDF	5.0 pg/L	5.0U pg/L
	EB-04122010-RIG2-RZB	1,2,3,4,7,8-HxCDD	0.31 pg/L
1,2,3,6,7,8-HxCDD		0.35 pg/L	0.35U pg/L
1,2,3,7,8,9-HxCDD		0.60 pg/L	0.60U pg/L
1,2,3,4,6,7,8-HpCDD		1.8 pg/L	1.8U pg/L
OCDD		4.0 pg/L	4.0U pg/L
2,3,7,8-TCDF		4.8 pg/L	4.8U pg/L
1,2,3,7,8-PeCDF		4.3 pg/L	4.3U pg/L
2,3,4,7,8-PeCDF		2.2 pg/L	2.2U pg/L
1,2,3,4,7,8-HxCDF		7.3 pg/L	7.3U pg/L
1,2,3,6,7,8-HxCDF		4.6 pg/L	4.6U pg/L
2,3,4,6,7,8-HxCDF		1.3 pg/L	1.3U pg/L
1,2,3,7,8,9-HxCDF		0.85 pg/L	0.85U pg/L
1,2,3,4,7,8,9-HpCDF		5.6 pg/L	5.6U pg/L

Sample	Compound	Reported Concentration	Modified Final Concentration
EB-04122010-RIG1-RZB	1,2,3,4,7,8-HxCDD	0.57 pg/L	0.57U pg/L
	1,2,3,6,7,8-HxCDD	0.59 pg/L	0.59U pg/L
	1,2,3,7,8,9-HxCDD	0.82 pg/L	0.82U pg/L
	1,2,3,4,6,7,8-HpCDD	2.8 pg/L	2.8U pg/L
	OCDD	10 pg/L	10U pg/L
	2,3,7,8-TCDF	2.0 pg/L	2.0U pg/L
	1,2,3,7,8-PeCDF	1.2 pg/L	1.2U pg/L
	2,3,4,7,8-PeCDF	1.3 pg/L	1.3U pg/L
	1,2,3,4,7,8-HxCDF	1.7 pg/L	1.7U pg/L
	1,2,3,6,7,8-HxCDF	1.2 pg/L	1.2U pg/L
	2,3,4,6,7,8-HxCDF	1.1 pg/L	1.1U pg/L
	1,2,3,4,6,7,8-HpCDF	2.0 pg/L	2.0U pg/L
	1,2,3,4,7,8,9-HpCDF	0.95 pg/L	0.95U pg/L
	OCDF	3.6 pg/L	3.6U pg/L

Samples EB-04122010-RIG3-RZD, EB-04122010-RIG2-RZB, and EB-04122010-RIG1-RZB were identified as equipment blanks. No polychlorinated dioxin/dibenzofuran contaminants were found in this blank with the following exceptions:

Equipment Blank ID	Sampling Date	Compound	Concentration	Associated Samples
EB-04122010-RIG3-RZD	4/12/10	1,2,3,6,7,8-HxCDD	0.64 pg/L	No associated samples in this SDG
		1,2,3,7,8,9-HxCDD	0.47 pg/L	
		1,2,3,4,6,7,8-HpCDD	1.9 pg/L	
		OCDD	5.0 pg/L	
		2,3,7,8-TCDF	4.6 pg/L	
		1,2,3,7,8-PeCDF	3.8 pg/L	
		2,3,4,7,8-PeCDF	2.2 pg/L	
		1,2,3,4,7,8-HxCDF	7.7 pg/L	
		1,2,3,6,7,8-HxCDF	4.8 pg/L	
		2,3,4,6,7,8-HxCDF	1.2 pg/L	
		1,2,3,7,8,9-HxCDF	0.59 pg/L	
		1,2,3,4,6,7,8-HpCDF	14 pg/L	
		1,2,3,4,7,8,9-HpCDF	5.0 pg/L	
		OCDF	27 pg/L	
EB-04122010-RIG2-RZB	4/12/10	1,2,3,4,7,8-HxCDD	0.31 pg/L	No associated samples in this SDG
		1,2,3,6,7,8-HxCDD	0.35 pg/L	
		1,2,3,7,8,9-HxCDD	0.960 pg/L	
		1,2,3,4,6,7,8-HpCDD	1.8 pg/L	
		OCDD	4.0 pg/L	
		2,3,7,8-TCDF	4.8 pg/L	
		1,2,3,7,8-PeCDF	4.3 pg/L	
		2,3,4,7,8-PeCDF	2.2 pg/L	
		1,2,3,4,7,8-HxCDF	7.3 pg/L	
		1,2,3,6,7,8-HxCDF	4.6 pg/L	
		2,3,4,6,7,8-HxCDF	1.3 pg/L	
		1,2,3,7,8,9-HxCDF	0.85 pg/L	
		1,2,3,4,6,7,8-HpCDF	13 pg/L	
		1,2,3,4,7,8,9-HpCDF	5.6 pg/L	
OCDF	26 pg/L			

Equipment Blank ID	Sampling Date	Compound	Concentration	Associated Samples
EB-04122010-RIG1-RZB	4/12/10	1,2,3,4,7,8-HxCDD 1,2,3,6,7,8-HxCDD 1,2,3,7,8,9-HxCDD 1,2,3,4,6,7,8-HpCDD OCDD 2,3,7,8-TCDF 1,2,3,7,8-PeCDF 2,3,4,7,8-PeCDF 1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 2,3,4,6,7,8-HxCDF 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	0.57 pg/L 0.59 pg/L 0.82 pg/L 2.8 pg/L 10 pg/L 2.0 pg/L 1.2 pg/L 1.3 pg/L 1.7 pg/L 1.2 pg/L 1.1 pg/L 2.0 pg/L 0.95 pg/L 3.6 pg/L	No associated samples in this SDG

VI. 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.

VII. Laboratory Control Samples (LCS)

Laboratory control samples were reviewed for each matrix as applicable. Percent recoveries (%R) were within QC limits.

VIII. Regional Quality Assurance and Quality Control

Not applicable.

IX. Internal Standards

All internal standard recoveries were within QC limits.

X. Target Compound Identifications

Raw data were not reviewed for this SDG.

XI. Project Quantitation Limit

All compounds reported below the PQL were qualified as follows:

Sample	Finding	Flag	A or P
All samples in SDG G0D140534	All compounds reported below the PQL.	J (all detects)	A

All compounds reported as EMPC were qualified as follows:

Sample	Finding	Flag	A or P
All samples in SDG G0D140534	All compounds reported as estimated maximum possible concentration (EMPC).	JK (all detects)	A

Raw data were not reviewed for this SDG.

XII. System Performance

Raw data were not reviewed for this SDG.

XIII. Overall Assessment of Data

Data flags are summarized at the end of this report if data has been qualified.

XIV. Field Duplicates

No field duplicates were identified in this SDG.

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Data Qualification Summary - SDG G0D140534**

SDG	Sample	Compound	Flag	A or P	Reason (Code)
G0D140534	EB-04122010-RIG3-RZD EB-04122010-RIG2-RZB EB-04122010-RIG1-RZB	All compounds reported below the PQL.	J (all detects)	A	Project Quantitation Limit (sp)
G0D140534	EB-04122010-RIG3-RZD EB-04122010-RIG2-RZB EB-04122010-RIG1-RZB	All compounds reported as EMPC	JK (all detects)	A	Project Quantitation Limit (k)

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Laboratory Blank Data Qualification Summary - SDG G0D140534**

SDG	Sample	Compound	Modified Final Concentration	A or P	Code
G0D140534	EB-04122010-RIG3-RZD	1,2,3,6,7,8-HxCDD 1,2,3,7,8,9-HxCDD 1,2,3,4,6,7,8-HpCDD OCDD 2,3,7,8-TCDF 1,2,3,7,8-PeCDF 2,3,4,7,8-PeCDF 1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 2,3,4,6,7,8-HxCDF 1,2,3,7,8,9-HxCDF 1,2,3,4,7,8,9-HpCDF	0.64U pg/L 0.47U pg/L 1.9U pg/L 5.0U pg/L 4.6U pg/L 3.8U pg/L 2.2U pg/L 7.7U pg/L 4.8U pg/L 1.2U pg/L 0.59U pg/L 5.0U pg/L	A	bl
G0D140534	EB-04122010-RIG2-RZB	1,2,3,4,7,8-HxCDD 1,2,3,6,7,8-HxCDD 1,2,3,7,8,9-HxCDD 1,2,3,4,6,7,8-HpCDD OCDD 2,3,7,8-TCDF 1,2,3,7,8-PeCDF 2,3,4,7,8-PeCDF 1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 2,3,4,6,7,8-HxCDF 1,2,3,7,8,9-HxCDF 1,2,3,4,7,8,9-HpCDF	0.31U pg/L 0.35U pg/L 0.60U pg/L 1.8U pg/L 4.0U pg/L 4.8U pg/L 4.3U pg/L 2.2U pg/L 7.3U pg/L 4.6U pg/L 1.3U pg/L 0.85U pg/L 5.6U pg/L	A	bl

SDG	Sample	Compound	Modified Final Concentration	A or P	Code
G0D140534	EB-04122010-RIG1-RZB	1,2,3,4,7,8-HxCDD 1,2,3,6,7,8-HxCDD 1,2,3,7,8,9-HxCDD 1,2,3,4,6,7,8-HpCDD OCDD 2,3,7,8-TCDF 1,2,3,7,8-PeCDF 2,3,4,7,8-PeCDF 1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 2,3,4,6,7,8-HxCDF 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	0.57U pg/L 0.59U pg/L 0.82U pg/L 2.8U pg/L 10U pg/L 2.0U pg/L 1.2U pg/L 1.3U pg/L 1.7U pg/L 1.2U pg/L 1.1U pg/L 2.0U pg/L 0.95U pg/L 3.6U pg/L	A	bl

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Equipment Blank Data Qualification Summary - SDG
G0D140534**

No Sample Data Qualified in this SDG

Tronox Northgate Henderson

VALIDATION COMPLETENESS WORKSHEET

LDC #: 23188E21

SDG #: G0D140534

Laboratory: Test America

Stage 2B

Date: 4/20/10

Page: 1 of 1

Reviewer: [Signature]

2nd Reviewer: _____

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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.	Technical holding times	A	Sampling dates: 4/12/10
II.	HRGC/HRMS Instrument performance check	A	
III.	Initial calibration	A	
IV.	Routine calibration/IC	SW	
V.	Blanks	SW	
VI.	Matrix spike/Matrix spike duplicates	N	client specified
VII.	Laboratory control samples	A	LCS
VIII.	Regional quality assurance and quality control	N	
IX.	Internal standards	A	
X.	Target compound identifications	N	
XI.	Compound quantitation and CRQLs	SN	
XII.	System performance	N	
XIII.	Overall assessment of data	A	
XIV.	Field duplicates	N	
XV.	Field blanks	SW	ZB = 1, 2, 3

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

Validated Samples:

1	EB-04122010-RIG3-RZD	W	11	0110418MB	21		31
2	EB-04122010-RIG2-RZB		12		22		32
3	EB-04122010-RIG1-RZB		13		23		33
4			14		24		34
5			15		25		35
6			16		26		36
7			17		27		37
8			18		28		38
9			19		29		39
10			20		30		40

Notes: _____

VALIDATION FINDINGS WORKSHEET

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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:

METHOD: HRGC/HRMS Dioxins (EPA Method 8290)

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? If yes, please see qualification below.

Blank extraction date: 4/20/10 Blank analysis date: 4/26/10 Associated samples: All (68)
 Conc. units: pg/L

Compound	Blank ID	Sample Identification				
		5X	1	2	3	
	0110418MR					
C	1.3	6.5		0.31/U	0.57/U	
D	1.3	6.5	0.64/U	0.35/U	0.59/U	
E	1.8	9	0.47/U	0.60/U	0.82/U	
F	2.7	13.5	1.9/U	1.8/U	2.8/U	
G	11	55	5.0/U	4.0/U	10/U	
H	1.8	9	4.6/U	4.8/U	2.0/U	
I	1.8	9	3.8/U	4.3/U	1.2/U	
J	2.2	11	2.2/U	2.2/U	1.3/U	
K	1.6	8	7.7/U	7.3/U	1.7/U	
L	1.4	7	4.8/U	4.6/U	1.2/U	
M	1.8	9	1.2/U	1.3/U	1.1/U	
N	1.7	8.5	0.59/U	0.85/U		
O	1.9	9.5	-	-	2.0/U	
P	2.0	10	5.0/U	5.6/U	0.95/U	
Q	4.2	21	-	-	3.6/U	

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
Field Blanks

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

N/A Were field blanks identified in this SDG?

Blank units: pg/L Associated sample units: pg/g

Sampling date: 4/12/10

Field blank type: (circle one) EB Field Blank / Rinsate / Other: None

Associated Samples: _____

Compound	Blank ID	Sample Identification				
	EB-04122010-RIG3-RZD		5X			
D	0.64		0.0032			
E	0.47		0.00235			
F	1.9		0.0095			
G	5.0		0.025			
H	4.6		0.023			
I	3.8		0.019			
J	2.2		0.011			
K	7.7		0.0385			
L	4.8		0.024			
M	1.2		0.006			
N	0.59		0.00295			
O	14		0.07			
P	5.0		0.025			
Q	27		0.135			
CRQL						

CIRCLED RESULTS WERE NOT QUALIFIED. ALL RESULTS NOT CIRCLED WERE QUALIFIED BY THE FOLLOWING STATEMENT:

Samples with compound concentrations within five times the associated field blank concentration are listed above, these sample results were qualified as not detected, "U".

VALIDATION FINDINGS WORKSHEET
Field Blanks

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

Y/N N/A Were field blanks identified in this SDG?

Blank units: pg/L Associated sample units: pg/g

Sampling date: 4/12/10

Field blank type: (circle one) Field Blank / Rinsate / Other: Field Blank Associated Samples: None

Compound	Blank ID	5X	Sample Identification			
	ER-04122010-RIG2-RZR	5X				
C	0.31	0.00155				
D	0.35	0.00175				
E	0.960	0.0048				
F	1.8	0.009				
G	4.0	0.02				
H	4.8	0.024				
I	4.3	0.0215				
J	2.2	0.011				
K	7.3	0.0365				
L	4.6	0.023				
M	1.3	0.0065				
N	0.85	0.00425				
O	13	0.065				
P	5.6	0.028				
Q	26	0.13				
CRQL						

CIRCLED RESULTS WERE NOT QUALIFIED. ALL RESULTS NOT CIRCLED WERE QUALIFIED BY THE FOLLOWING STATEMENT:
Samples with compound concentrations within five times the associated field blank concentration are listed above, these sample results were qualified as not detected, "U".

VALIDATION FINDINGS WORKSHEET
Field Blanks

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

N/A Were field blanks identified in this SDG?

Blank units: pg/L Associated sample units: pg/g

Sampling date: 4/12/10

Field blank type: (circle one) Field Blank / Rinsate / Other: EB Associated Samples: None

Compound	Blank ID	Blank ID	Sample Identification
	FER-04122010-RIG4-RZR	5X	
C	0.57	0.00285	
D	0.59	0.00295	
E	0.82	0.0041	
F	2.8	0.014	
G	10	0.05	
H	2.0	0.01	
I	1.2	0.006	
J	1.3	0.0065	
K	1.7	0.0085	
L	1.2	0.006	
M	1.1	0.0055	
O	2.0	0.01	
P	0.95	0.00475	
Q	3.6	0.018	
CRQL			

CIRCLED RESULTS WERE NOT QUALIFIED. ALL RESULTS NOT CIRCLED WERE QUALIFIED BY THE FOLLOWING STATEMENT:
 Samples with compound concentrations within five times the associated field blank concentration are listed above, these sample results were qualified as not detected, "U".

VALIDATION FINDINGS WORKSHEET
Compound Quantitation and Reported CRQLs

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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

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

#	Date	Sample ID	Finding	Associated Samples	Qualifications
		M1	ZMPC results (Q-tag)		(K)(e)
		M1	No confirmation for 2,3,7,8-TCDF		None

Comments: See sample calculation verification worksheet for recalculations

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

Project/Site Name: Tronox LLC Facility, PCS, Henderson, Nevada
Collection Date: April 12, 2010
LDC Report Date: May 24, 2010
Matrix: Soil
Parameters: Dioxins/Dibenzofurans
Validation Level: Stage 2B & 4
Laboratory: TestAmerica, Inc.

Sample Delivery Group (SDG): GOD140543

Sample Identification

SSAN6-02-3BPC
SSAN7-02-1BPC_FD
SSAJ3-02-1BPC_FD
SA156-3BPC_FD**
SSAN6-02-3BPCMS
SSAN6-02-3BPCMSD

**Indicates sample underwent Stage 4 review

Introduction

This data review covers 6 soil samples listed on the cover sheet including dilutions and reanalysis as applicable. The analyses were per EPA SW 846 Method 8290 for Polychlorinated Dioxins/Dibenzofurans.

This review follows the Standard Operating Procedures (SOP) 40, Data Review/Validation (BRC 2009), the Quality Assurance Project Plan Tronox LLC Facility, Henderson, Nevada (June 2009), NDEP guidance (May 2006), and USEPA Contract Laboratory Program National Functional Guidelines for Polychlorinated Dioxins/Dibenzofurans Data Review (September 2005).

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.

Blank results are summarized in Section V.

Field duplicates are summarized in Section XIV.

Samples indicated by a double asterisk on the front cover underwent a Stage 4 review. A Stage 2B review was performed on all of the other samples. Raw data were not evaluated for the samples reviewed by Stage 2B criteria since this review is based on QC data.

The following are definitions of the data qualifiers:

- J+ Data are qualified as estimated, with a high bias likely to occur. False positives or false negatives are unlikely to have been reported.
- J- Data are qualified as estimated, with a low bias likely to occur. False positives or false negatives are unlikely to have been reported.
- J Data are qualified as estimated; it is not possible to assess the direction of the potential bias. False positives or false negatives are unlikely to have been reported.
- U Indicates the compound or analyte was analyzed for but not detected at or above the stated limit.
- R Data are qualified as rejected. There is a significant potential for the reporting of false negatives or false positives.
- UU Indicates the compound or analyte was analyzed for but not detected. The sample detection limit is an estimated value.
- B The analytical result may be a false positive totally attributable to blank contamination. This qualifier is applicable to radiochemistry analysis only.
- JB The analytical result may be biased high and partially attributable to blank contamination. This qualifier is applicable to radiochemistry analysis only.
- JK The analytical result is an estimated maximum possible concentration (EMPC).
- X The analytical result is not used for reporting because a more accurate and precise result is reported in its place.
- J-TDS The analytical result is estimated based on failure of the Total Dissolved Solids (TDS) correctness check performed in accordance with the Standard Method 1030E.
- J-CAB The analytical result is estimated based on failure of the cation-anion balance correctness check performed in accordance with Standard Method 1030E.
- J-TDS & CAB The analytical result is unreliable based on the failure of the cation-anion balance and TDS correctness check performed in accordance with standard Method 1030E.
- A Indicates the finding is based upon technical validation criteria.
- P Indicates the finding is related to a protocol/contractual deviation.
- None Indicates the data was not significantly impacted by the finding, therefore qualification was not required.

I. Technical Holding Times

All technical holding time requirements were met.

The chain-of-custodies were reviewed for documentation of cooler temperatures. All cooler temperatures met validation criteria.

II. HRGC/HRMS Instrument Performance Check

Instrument performance was checked at the required daily 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 exact mass of 380.9760 of PFK was verified. The static resolving power was at least 10,000 (10% valley definition) for samples on which Stage 4 review was performed. Raw data were not evaluated for the samples reviewed by Stage 2B criteria.

III. Initial Calibration

A five point initial calibration was performed as required by the method.

Percent relative standard deviations (%RSD) were less than or equal to 20.0% for unlabeled compounds and less than or equal to 30.0% for labeled compounds.

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

The minimum S/N ratio for each target compound was greater than or equal to 2.5 and greater than or equal to 10 for each recovery and internal standard compound for samples on which Stage 4 review was performed. Raw data were not evaluated for the samples reviewed by Stage 2B criteria.

IV. Routine Calibration (Continuing)

Routine calibration was performed at the required frequencies.

All of the routine calibration percent differences (%D) between the initial calibration RRF and the routine calibration RRF were less than or equal to 20.0% for unlabeled compounds and less than or equal to 30.0% for labeled compounds.

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

V. Blanks

Method blanks were reviewed for each matrix as applicable. No polychlorinated dioxin/dibenzofuran contaminants were found in the method blanks with the following exceptions:

Method Blank ID	Extraction Date	Compound	Concentration	Associated Samples
0110455MB	4/20/10	2,3,7,8-TCDF 1,2,3,7,8-PeCDF 1,2,3,4,6,7,8-HpCDF	0.31 pg/g 0.31 pg/g 0.30 pg/g	SSAN6-02-3BPC
0109260MB	4/19/10	1,2,3,4,6,7,8-HpCDD OCDD 2,3,7,8-TCDF 1,2,3,7,8-PeCDF 1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 1,2,3,7,8,9-HxCDF 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	0.14 pg/g 0.59 pg/g 0.28 pg/g 0.14 pg/g 0.24 pg/g 0.14 pg/g 0.086 pg/g 0.30 pg/g 0.13 pg/g 0.63 pg/g	SSAN7-02-1BPC_FD SSAJ3-02-1BPC_FD SA156-3BPC_FD**

Sample concentrations were compared to concentrations detected in the method blanks as required by the QAPP. No sample data was qualified with the following exceptions:

Sample	Compound	Reported Concentration	Modified Final Concentration
SSAJ3-02-1BPC_FD	1,2,3,4,6,7,8-HpCDD OCDD 2,3,7,8-TCDF 1,2,3,7,8-PeCDF 1,2,3,4,7,8-HxCDF 1,2,3,7,8,9-HxCDF	0.41 pg/g 1.4 pg/g 0.60 pg/g 0.69 pg/g 1.2 pg/g 0.18 pg/g	0.41U pg/g 1.4U pg/g 0.60U pg/g 0.69U pg/g 1.2U pg/g 0.18U pg/g
SA156-3BPC_FD**	2,3,7,8-TCDF 1,2,3,4,6,7,8-HpCDF	0.16 pg/g 1.2 pg/g	0.16U pg/g 1.2U pg/g

Samples EB-04122010-RIG1-RZB, EB-04122010-RIG2-RZB, and EB-04122010-RIG3-RZB (all from SDG GOD140534) were identified as equipment blanks. No polychlorinated dioxin/dibenzofuran contaminants were found in these blanks with the following exceptions:

Equipment Blank ID	Sampling Date	Compound	Concentration	Associated Samples
EB-04122010-RIG1-RZB	4/12/10	1,2,3,4,7,8-HxCDD 1,2,3,6,7,8-HxCDD 1,2,3,7,8,9-HxCDD 1,2,3,4,6,7,8-HpCDD OCDD 2,3,7,8-TCDF 1,2,3,7,8-PeCDF 2,3,4,7,8-PeCDF 1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 2,3,4,6,7,8-HxCDF 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	0.57 pg/L 0.59 pg/L 0.82 pg/L 2.8 pg/L 10 pg/L 2.0 pg/L 1.2 pg/L 1.3 pg/L 1.7 pg/L 1.2 pg/L 1.1 pg/L 2.0 pg/L 0.95 pg/L 3.6 pg/L	SA156-3BPC_FD**
EB-04122010-RIG2-RZB	4/12/10	1,2,3,4,7,8-HxCDD 1,2,3,6,7,8-HxCDD 1,2,3,7,8,9-HxCDD 1,2,3,4,6,7,8-HpCDD OCDD 2,3,7,8-TCDF 1,2,3,7,8-PeCDF 2,3,4,7,8-PeCDF 1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 2,3,4,6,7,8-HxCDF 1,2,3,7,8,9-HxCDF 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	0.31 pg/L 0.35 pg/L 0.60 pg/L 1.8 pg/L 4.0 pg/L 4.8 pg/L 4.3 pg/L 2.2 pg/L 7.3 pg/L 4.6 pg/L 1.3 pg/L 0.85 pg/L 13 pg/L 5.6 pg/L 26 pg/L	SA156-3BPC_FD**
EB-04122010-RIG3-RZB	4/12/10	1,2,3,6,7,8-HxCDD 1,2,3,7,8,9-HxCDD 1,2,3,4,6,7,8-HpCDD OCDD 2,3,7,8-TCDF 1,2,3,7,8-PeCDF 2,3,4,7,8-PeCDF 1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 2,3,4,6,7,8-HxCDF 1,2,3,7,8,9-HxCDF 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	0.64 pg/L 0.47 pg/L 1.9 pg/L 5.0 pg/L 4.6 pg/L 3.8 pg/L 2.2 pg/L 7.7 pg/L 4.8 pg/L 1.2 pg/L 0.59 pg/L 14 pg/L 5.0 pg/L 27 pg/L	SSAJ3-02-1BPC_FD

Sample concentrations were compared to concentrations detected in the equipment blanks as required by the QAPP. No sample data was qualified.

Samples FB04062010-RZB (from SDG G0D120488), FB-04072010-RZD (from SDG G0D090441), and FB-04072010-RZC (from SDG G0D130519) were identified as field blanks. No polychlorinated dioxin/dibenzofuran contaminants were found in these blanks with the following exceptions:

Field Blank ID	Sampling Date	Compound	Concentration	Associated Samples
FB-04072010-RZD	4/7/10	1,2,3,4,7,8-HxCDD 1,2,3,7,8,9-HxCDD 1,2,3,4,6,7,8-HpCDD OCDD 1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 2,3,4,6,7,8-HxCDF 1,2,3,7,8,9-HxCDF 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	0.89 pg/L 1.5 pg/L 2.2 pg/L 8.3 pg/L 1.4 pg/L 1.6 pg/L 1.5 pg/L 1.6 pg/L 1.3 pg/L 1.4 pg/L 4.1 pg/L	SSAJ3-02-1BPC_FD
FB04062010-RZB	4/6/10	1,2,3,7,8,9-HxCDD 1,2,3,4,6,7,8-HpCDD OCDD 2,3,7,8-TCDF 1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 1,2,3,7,8,9-HxCDF 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	0.68 pg/L 2.5 pg/L 6.2 pg/L 2.7 pg/L 1.4 pg/L 0.82 pg/L 0.94 pg/L 1.8 pg/L 1.2 pg/L 4.4 pg/L	SA156-3BPC_FD**
FB-04072010-RZC	4/8/10	1,2,3,4,7,8-HxCDD 1,2,3,6,7,8-HxCDD 1,2,3,7,8,9-HxCDD 1,2,3,4,6,7,8-HpCDD OCDD 2,3,7,8-TCDF 1,2,3,7,8-PeCDF 2,3,4,7,8-PeCDF 1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 2,3,4,6,7,8-HxCDF 1,2,3,7,8,9-HxCDF 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	0.77 pg/L 0.74 pg/L 0.82 pg/L 4.2 pg/L 37 pg/L 0.57 pg/L 0.96 pg/L 0.67 pg/L 1.1 pg/L 0.96 pg/L 1.0 pg/L 1.0 pg/L 2.1 pg/L 1.5 pg/L 6.7 pg/L	SSAN6-02-3BPC SSAN7-02-1BPC_FD

Sample concentrations were compared to concentrations detected in the field blanks as required by the QAPP. No sample data was qualified.

VI. Matrix Spike/Matrix Spike Duplicates

Matrix spike (MS) and matrix spike duplicate (MSD) samples were reviewed for each matrix as applicable. Although the MS/MSD percent recoveries (%R) were not within QC limits for several compounds, the LCS percent recoveries (%R) were within QC limits and no data were qualified.

VII. Laboratory Control Samples (LCS)

Laboratory control samples were reviewed for each matrix as applicable. Percent recoveries (%R) were within QC limits.

VIII. Regional Quality Assurance and Quality Control

Not applicable.

IX. Internal Standards

All internal standard recoveries were within QC limits with the following exceptions:

Sample	Internal Standards	%R (Limits)	Compound	Flag	A or P
SSAN6-02-3BPC	¹³ C-OCDD	34 (40-135)	OCDD OCDF	J (all detects) UJ (all non-detects) J (all detects) UJ (all non-detects)	P
SA156-3BPC_FD**	¹³ C-OCDD	32 (40-135)	OCDD OCDF	J (all detects) UJ (all non-detects) J (all detects) UJ (all non-detects)	P

X. Target Compound Identifications

All target compound identifications were within validation criteria for samples on which a Stage 4 review was performed. Raw data were not evaluated for the samples reviewed by Stage 2B criteria.

XI. Project Quantitation Limit

All compound quantitation and CRQLs were within validation criteria with the following exceptions:

Sample	Compound	Finding	Criteria	Flag	A or P
SSAN6-02-3BPC	1,2,3,4,6,7,8-HpCDF OCDF	Sample result exceeded calibration range.	Reported result should be within calibration range.	J (all detects) J (all detects)	P
SSAN7-02-1BPC_FD	2,3,7,8-TCDF	Sample result exceeded calibration range.	Reported result should be within calibration range.	J (all detects)	P

All compounds reported below the PQL were qualified as follows:

Sample	Finding	Flag	A or P
All samples in SDG G0D140543	All compounds reported below the PQL.	J (all detects)	A

All compounds reported as EMPC were qualified as follows:

Sample	Finding	Flag	A or P
All samples in SDG G0D140543	All compounds reported as estimated maximum possible concentration (EMPC).	JK (all detects)	A

Raw data were not evaluated for the samples reviewed by Stage 2B criteria.

XII. System Performance

The system performance was acceptable for samples on which a Stage 4 review was performed. Raw data were not evaluated for the samples reviewed by Stage 2B criteria.

XIII. Overall Assessment of Data

Data flags are summarized at the end of this report if data has been qualified.

XIV. Field Duplicates

Samples SSAN7-02-1BPC_FD and SSAN7-02-1BPC (from SDG (G0D140422)), samples SSAJ3-02-1BPC_FD and SSAJ3-02-1BPC (From SDG G0D140422), and samples SA156-3BPC_FD** and SA156-3BPC (from SDG G0D140435) were identified as field duplicates. No polychlorinated dioxins/dibenzofurans were detected in any of the samples with the following exceptions:

Compound	Concentration (pg/g)		RPD (Limits)	Difference (Limits)	Flags	A or P
	SSAJ3-02-1BPC	SSAJ3-02-1BPC_FD				
2,3,7,8-TCDD	0.22	1.1U	-	0.88 (≤1.1)	-	-
1,2,3,7,8-PeCDD	0.90	5.3U	-	4.4 (≤5.3)	-	-
1,2,3,4,7,8-HxCDD	0.99	5.3U	-	4.31 (≤5.3)	-	-
1,2,3,6,7,8-HxCDD	1.2	0.13	-	1.07 (≤5.3)	-	-
1,2,3,7,8,9-HxCDD	1.1	0.12	-	0.98 (≤5.3)	-	-
1,2,3,4,6,7,8-HpCDD	1.6	0.41	-	1.19 (≤5.3)	-	-
OCDD	5.9	1.4	-	4.5 (≤11)	-	-
2,3,7,8-TCDF	0.90	0.60	-	0.3 (≤1.1)	-	-

Compound	Concentration (pg/g)		RPD (Limits)	Difference (Limits)	Flags	A or P
	SSAJ3-02-1BPC	SSAJ3-02-1BPC_FD				
1,2,3,7,8-PeCDF	2.1	0.69	-	1.41 (≤ 5.3)	-	-
2,3,4,7,8-PeCDF	1.5	0.39	-	1.11 (≤ 5.3)	-	-
1,2,3,4,7,8-HxCDF	1.7	1.2	-	0.5 (≤ 5.3)	-	-
1,2,3,6,7,8-HxCDF	2.2	0.76	-	1.44 (≤ 5.3)	-	-
2,3,4,6,7,8-HxCDF	1.5	0.16	-	1.34 (≤ 5.3)	-	-
1,2,3,7,8,9-HxCDF	1.4	0.18	-	1.22 (≤ 5.3)	-	-
1,2,3,4,6,7,8-HpCDF	3.2	2.7	-	0.5 (≤ 5.3)	-	-
1,2,3,4,7,8,9-HpCDF	2.1	0.99	-	1.11 (≤ 5.3)	-	-
OCDF	9.1	7.8	-	1.3 (≤ 11)	-	-

Compound	Concentration (pg/g)		RPD (Limits)	Difference (Limits)	Flags	A or P
	SSAN7-02-1BPC	SSAN7-02-1BPC_FD				
2,3,7,8-TCDD	2.4	2.4	-	0 (≤ 1.1)	-	-
1,2,3,7,8-PeCDD	11	10	-	1 (≤ 5.3)	-	-
1,2,3,4,7,8-HxCDD	6.6	5.3	-	1.3 (≤ 5.3)	-	-
1,2,3,6,7,8-HxCDD	15	12	-	3 (≤ 5.3)	-	-
1,2,3,7,8,9-HxCDD	11	8.6	-	2.4 (≤ 5.3)	-	-
1,2,3,4,6,7,8-HpCDD	39	35	11 (≤ 50)	-	-	-
OCDD	40	35	-	5 (≤ 11)	-	-
2,3,7,8-TCDF	130	170	27 (≤ 50)	-	-	-
1,2,3,7,8-PeCDF	160	210	27 (≤ 50)	-	-	-

Compound	Concentration (pg/g)		RPD (Limits)	Difference (Limits)	Flags	A or P
	SSAN7-02-1BPC	SSAN7-02-1BPC_FD				
2,3,4,7,8-PeCDF	77	82	6 (≤ 50)	-	-	-
1,2,3,4,7,8-HxCDF	230	250	8 (≤ 50)	-	-	-
1,2,3,6,7,8-HxCDF	200	160	22 (≤ 50)	-	-	-
2,3,4,6,7,8-HxCDF	44	35	23 (≤ 50)	-	-	-
1,2,3,7,8,9-HxCDF	42	21	-	21 (≤ 5.3)	J (all detects)	A
1,2,3,4,6,7,8-HpCDF	500	440	13 (≤ 50)	-	-	-
1,2,3,4,7,8,9-HpCDF	270	220	20 (≤ 50)	-	-	-
OCDF	1300	1200	8 (≤ 50)	-	-	-

Compound	Concentration (pg/g)		RPD (Limits)	Difference (Limits)	Flags	A or P
	SA156-3BPC	SA156-3BPC_FD**				
2,3,7,8-TCDD	0.15	0.54U	-	0.39 (≤ 0.54)	-	-
1,2,3,7,8-PeCDD	0.56	2.7U	-	2.14 (≤ 2.7)	-	-
1,2,3,4,7,8-HxCDD	0.84	2.7U	-	1.86 (≤ 2.7)	-	-
1,2,3,6,7,8-HxCDD	0.68	2.7U	-	2.02 (≤ 2.7)	-	-
1,2,3,7,8,9-HxCDD	0.81	2.7U	-	1.89 (≤ 2.7)	-	-
1,2,3,4,6,7,8-HpCDD	1.4	2.7U	-	1.3 (≤ 2.7)	-	-
OCDD	5.6	5.4U	-	0.2 (≤ 5.4)	-	-
2,3,7,8-TCDF	0.51	0.16	-	0.35 (≤ 0.54)	-	-
1,2,3,7,8-PeCDF	1.7	2.7U	-	1 (≤ 2.7)	-	-
1,2,3,4,7,8-HxCDF	4.1	2.7U	-	1.4 (≤ 2.7)	-	-

Compound	Concentration (pg/g)		RPD (Limits)	Difference (Limits)	Flags	A or P
	SA156-3BPC	SA156-3BPC_FD**				
1,2,3,6,7,8-HxCDF	3.3	2.7U	-	0.6 (≤ 2.7)	-	-
2,3,4,6,7,8-HxCDF	1.4	2.7U	-	1.3 (≤ 2.7)	-	-
1,2,3,7,8,9-HxCDF	1.7	2.7U	-	1 (≤ 2.7)	-	-
1,2,3,4,6,7,8-HpCDF	10	1.2	-	8.8 (≤ 2.7)	J (all detects)	A
1,2,3,4,7,8,9-HpCDF	5.1	2.7U	-	2.4 (≤ 2.7)	-	-
OCDF	25	5.4	-	19.6 (≤ 5.4)	J (all detects)	A

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Data Qualification Summary - SDG G0D140543**

SDG	Sample	Compound	Flag	A or P	Reason (Code)
G0D140543	SSAN6-02-3BPC SA156-3BPC_FD**	OCDD OCDF	J (all detects) UJ (all non-detects) J (all detects) UJ (all non-detects)	P	Internal standards (%R) (i)
G0D140543	SSAN6-02-3BPC	1,2,3,4,6,7,8-HpCDF OCDF	J (all detects) J (all detects)	P	Compound quantitation and CRQLs (e)
G0D140543	SSAN7-02-1BPC_FD	2,3,7,8-TCDF	J (all detects)	P	Compound quantitation and CRQLs (e)
G0D140543	SSAN6-02-3BPC SSAN7-02-1BPC_FD SSAJ3-02-1BPC_FD SA156-3BPC_FD**	All compounds reported below the PQL.	J (all detects)	A	Project Quantitation Limit (sp)
G0D140543	SSAN6-02-3BPC SSAN7-02-1BPC_FD SSAJ3-02-1BPC_FD SA156-3BPC_FD**	All compounds reported as EMPC	JK (all detects)	A	Project Quantitation Limit (k)
G0D140543	SSAN7-02-1BPC_FD	1,2,3,7,8,9-HxCDF	J (all detects)	A	Field duplicates (Difference) (fd)
G0D140543	SA156-3BPC_FD**	1,2,3,4,6,7,8-HpCDF OCDF	J (all detects) J (all detects)	A	Field duplicates (Difference) (fd)

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Laboratory Blank Data Qualification Summary - SDG
G0D140543**

SDG	Sample	Compound	Modified Final Concentration	A or P	Code
G0D140543	SSAJ3-02-1BPC_FD	1,2,3,4,6,7,8-HpCDD OCDD 2,3,7,8-TCDF 1,2,3,7,8-PeCDF 1,2,3,4,7,8-HxCDF 1,2,3,7,8,9-HxCDF	0.41U pg/g 1.4U pg/g 0.60U pg/g 0.69U pg/g 1.2U pg/g 0.18U pg/g	A	bl
G0D140543	SA156-3BPC_FD**	2,3,7,8-TCDF 1,2,3,4,6,7,8-HpCDF	0.16U pg/g 1.2U pg/g	A	bl

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Equipment Blank Data Qualification Summary - SDG
G0D140543**

No Sample Data Qualified in this SDG

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Field Blank Data Qualification Summary - SDG G0D140543**

No Sample Data Qualified in this SDG

Tronox Northgate Henderson

VALIDATION COMPLETENESS WORKSHEET

Stage 2B/4

LDC #: 23188F21

SDG #: G0D140543

Laboratory: Test America

Date: 5/11/10

Page: 1 of 1

Reviewer: [Signature]

2nd Reviewer: _____

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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.	Technical holding times	A	Sampling dates: <u>4/12/10</u>
II.	HRGC/HRMS Instrument performance check	A	
III.	Initial calibration	A	
IV.	Routine calibration/ ICX	A	
V.	Blanks	SW	
VI.	Matrix spike/Matrix spike duplicates	SW	
VII.	Laboratory control samples	A	<u>LOS</u>
VIII.	Regional quality assurance and quality control	N	
IX.	Internal standards	SW	
X.	Target compound identifications	A	Not reviewed for Stage 2B validation.
XI.	Compound quantitation and CRQLs	SW	Not reviewed for Stage 2B validation.
XII.	System performance	A	Not reviewed for Stage 2B validation.
XIII.	Overall assessment of data	A	
XIV.	Field duplicates	SW	D=2+SSAN7-02-1BPC, 3+SSAJ3-02-1BPC(G0D140422), D=4+SA156-3BPC(G0D140435)
XV.	Field blanks	SW	FB04062010-RZB(G0D120488)FB-04072010-RZD(G0D090441) FB-040710-RZC(G0D130519), EB-04122010-RIG3-RZD (G0D140534) EB-04122010-RIG2-RZB, EB-04122010-RIG1-RZB (G0D140534)

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

Validated Samples: ** Indicates sample underwent State 4 validation

1	SSAN6-02-3BPC	5	11	<u>0110445MB</u>	21		31
2	SSAN7-02-1BPC_FD		12	<u>0109260MB</u>	22		32
3	SSAJ3-02-1BPC_FD		13		23		33
4	SA156-3BPC_FD**		14		24		34
5	SSAN6-02-3BPCMS		15		25		35
6	SSAN6-02-3BPCMSD		16		26		36
7			17		27		37
8			18		28		38
9			19		29		39
10			20		30		40

Notes: _____

LDC #: 23188F21
 SDG #: 22 cones

VALIDATION FINDINGS CHECKLIST

Page: 1 of 4
 Reviewer: 9
 2nd Reviewer: _____

Method: Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

Validation Area	Yes	No	NA	Findings/Comments
I. Technical holding times				
All technical holding times were met.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Cooler temperature criteria was met.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
II. GC/MS Instrument performance check				
Was PFK exact mass 380.9760 verified?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were the retention time windows established for all homologues?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was the chromatographic resolution between 2,3,7,8-TCDD and peaks representing any other unlabeled TCDD isomers $\leq 25\%$?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Is the static resolving power at least 10,000 (10% valley definition)?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was the mass resolution adequately check with PFK?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was the presence of 1,2,8,9-TCDD and 1,3,4,6,8-PeCDF verified?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
III. Initial calibration				
Was the initial calibration performed at 5 concentration levels?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were all percent relative standard deviations (%RSD) $\leq 20\%$ for unlabeled standards and $< 30\%$ for labeled standards?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Did all calibration standards meet the Ion Abundance Ratio criteria?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was the signal to noise ratio for each target compound ≥ 2.5 and for each recovery and internal standard > 10 ?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
IV. Continuing calibration				
Was a routine calibration performed at the beginning and end of each 12 hour period?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were all percent differences (%D) $\leq 20\%$ for unlabeled standards and $\leq 30\%$ for labeled standards?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Did all routine calibration standards meet the Ion Abundance Ratio criteria?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
V. Blanks				
Was a method blank associated with every sample in this SDG?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was a method blank performed for each matrix and concentration?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was there contamination in the method blanks? If yes, please see the Blanks validation completeness worksheet?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
VI. Matrix spike/Matrix spike duplicates				
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.	<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"/>	
VII. Laboratory control samples				
Was an LCS analyzed for this SDG?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
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"/>	

VIII. Regional Quality Assurance and Quality Control			
Were performance evaluation (PE) samples performed?	<input checked="" type="checkbox"/>		
Were the performance evaluation (PE) samples within the acceptance limits?	<input checked="" type="checkbox"/>		
IX. Internal standards			
Were internal standard recoveries within the 40-135% criteria?	<input checked="" type="checkbox"/>		
Was the minimum S/N ratio of all internal standard peaks ≥ 10 ?	<input checked="" type="checkbox"/>		
X. Target compound identification			
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?	<input checked="" type="checkbox"/>		
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?	<input checked="" type="checkbox"/>		
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?	<input checked="" type="checkbox"/>		
Did compound spectra contain all characteristic ions listed in the table attached?	<input checked="" type="checkbox"/>		
Was the Ion Abundance Ratio for the two quantitation ions within criteria?	<input checked="" type="checkbox"/>		
Was the signal to noise ratio for each target compound and labeled standard ≥ 2.5 ?	<input checked="" type="checkbox"/>		
Does the maximum intensity of each specified characteristic ion coincide within ± 2 seconds (includes labeled standards)?	<input checked="" type="checkbox"/>		
For PCDF identification, was any signal ($S/N \geq 2.5$, at \pm seconds RT) detected in the corresponding PCDPE channel?	<input checked="" type="checkbox"/>		
Was an acceptable lock mass recorded and monitored?	<input checked="" type="checkbox"/>		
XI. Compound quantitation/CRQLs			
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 CRQLs adjusted to reflect all sample dilutions and dry weight factors applicable to level IV validation?	<input checked="" type="checkbox"/>		
XII. System performance			
System performance was found to be acceptable.	<input checked="" type="checkbox"/>		
XIII. Overall assessment of data			
Overall assessment of data was found to be acceptable.	<input checked="" type="checkbox"/>		
XIV. Field duplicates			
Field duplicate pairs were identified in this SDG.	<input checked="" type="checkbox"/>		
Target compounds were detected in the field duplicates.	<input checked="" type="checkbox"/>		
XV. Field blanks			
Field blanks were identified in this SDG.	<input checked="" type="checkbox"/>		
Target compounds were detected in the field blanks.	<input checked="" type="checkbox"/>		

VALIDATION FINDINGS WORKSHEET

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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

LDC #: 231887-1
 SDG #: see cone

Page: 1 of 1
 Reviewer: 9
 2nd Reviewer: _____

Blanks

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 8290)

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

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

Were all samples associated with a method blank?

Was a method blank analyzed for each matrix?

Was the blank contaminated? If yes, please see qualification below.

Blank extraction date: 4/20/12 Blank analysis date: 4/26/12

Conc. units: pg/g Associated Samples: 1 (75X)

Compound	Blank ID	Sample Identification
	01107455MB	P
A	0.31	
I	0.31	
O	0.30	

Blank extraction date: _____ Blank analysis date: _____ 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

Blanks

METHOD: HRGC/HRMS Dioxins (EPA Method 8290)

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? If yes, please see qualification below.

Blank extraction date: 4/19/10 Blank analysis date: 4/25/10

Conc. units: pg/kg Associated samples: 2-4 (bl)

Compound	Blank ID	Sample Identification				
		5X	3	4		
	0109260MB					
F	0.14	0.7	0.41/U			
G	0.59	2.95	1.4/U			
H	0.28	1.4	0.60/U	0.16/U		
I	0.14	0.7	0.69/U			
K	0.24	1.2	1.2/U			
L ✓	0.14	0.7	-			
N	0.086	0.43	0.18/U			
O	0.30	1.5	-	1.2/U		
P	0.13	0.65	-			
Q	0.63	3.15	-	-		

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".

LDC #: 23188EF21
SDG #: See Cover

VALIDATION FINDINGS WORKSHEET

Field Blanks

Page: 1 of 1
Reviewer: [Signature]
2nd Reviewer: _____

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

Y/N N/A Were field blanks identified in this SDG? Y

Blank units: pg/L **Associated sample units:** pg/g

Sampling date: 4/12/10

Field blank type: (circle one) Field Blank / Rinsate / Other: EB Associated Samples: 4

Compound	Blank ID	Blank ID	Sample Identification
	EB-04122010-RIG1-RZB	5X	
C	0.57	0.00285	
D	0.59	0.00295	
E	0.82	0.0041	
F	2.8	0.014	
G	10	0.05	
H	2.0	0.01	
I	1.2	0.006	
J	1.3	0.0065	
K	1.7	0.0085	
L	1.2	0.006	
M	1.1	0.0055	
O	2.0	0.01	
P	0.95	0.00475	
Q	3.6	0.018	
CRQL			

CIRCLED RESULTS WERE NOT QUALIFIED. ALL RESULTS NOT CIRCLED WERE QUALIFIED BY THE FOLLOWING STATEMENT:
Samples with compound concentrations within five times the associated field blank concentration are listed above, these sample results were qualified as not detected, "U".

VALIDATION FINDINGS WORKSHEET
Field Blanks

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

Y N N/A Were field blanks identified in this SDG?

Blank units: pg/L Associated sample units: pg/g

Sampling date: 4/12/10

Field blank type: (circle one) Field Blank / Rinsate / Other: EB Associated Samples: 4

Compound	Blank ID	5X	Sample Identification			
	EB-04122010-RIG2-RZR	5X				
C	0.31	0.00155				
D	0.35	0.00175				
E	0.60	0.003				
F	1.8	0.009				
G	4.0	0.02				
H	4.8	0.024				
I	4.3	0.0215				
J	2.2	0.011				
K	7.3	0.0365				
L	4.6	0.023				
M	1.3	0.0065				
N	0.85	0.00425				
O	13	0.065				
P	5.6	0.028				
Q	26	0.13				
CRQL						

CIRCLED RESULTS WERE NOT QUALIFIED. ALL RESULTS NOT CIRCLED WERE QUALIFIED BY THE FOLLOWING STATEMENT:
Samples with compound concentrations within five times the associated field blank concentration are listed above, these sample results were qualified as not detected,

VALIDATION FINDINGS WORKSHEET
Field Blanks

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

N/A Were field blanks identified in this SDG?

Blank units: pg/L **Associated sample units:** pg/g

Sampling date: 4/12/10

Field blank type: (circle one) Field Blank / Rinsate / Other: EB Associated Samples: 3

Compound	Blank ID	5X	Sample Identification
	ER-04122010-RIG3-RZD	5X	
D	0.64	0.0032	
E	0.47	0.00235	
F	1.9	0.0095	
G	5.0	0.025	
H	4.6	0.023	
I	3.8	0.019	
J	2.2	0.011	
K	7.7	0.0385	
L	4.8	0.024	
M	1.2	0.006	
N	0.59	0.00295	
O	14	0.07	
P	5.0	0.025	
Q	27	0.135	
CRQL			

CIRCLED RESULTS WERE NOT QUALIFIED. ALL RESULTS NOT CIRCLED WERE QUALIFIED BY THE FOLLOWING STATEMENT:
Samples with compound concentrations within five times the associated field blank concentration are listed above, these sample results were qualified as not detected, "U".

LDC #: 23/888
 SDG #: See Cover

VALIDATION FINDINGS WORKSHEET
Field Blanks

Page: 1 of 1
 Reviewer: [Signature]
 2nd Reviewer: [Signature]

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

N/A Were field blanks identified in this SDG?

Blank units: pg/L **Associated sample units:** pg/g

Sampling date: 4/8/10

Field blank type: (circle one) Field Blank Rinsate / Other: Associated Samples: 1-2 (> 5x)

Compound	Blank ID	Blank ID	Sample Identification
	ER-04072010-RZC	5X	
C	0.77	0.00385	
D	0.74	0.0037	
E	0.82	0.0041	
F	4.2	0.021	
G	37	0.185	
H	0.57	0.00285	
I	0.96	0.0048	
J	0.67	0.00335	
K	1.1	0.0055	
L	0.96	0.0048	
M	1.0	0.005	
N	1.0	0.005	
O	2.1	0.0105	
P	1.5	0.0075	
Q	6.7	0.0335	
CRQL			

CIRCLED RESULTS WERE NOT QUALIFIED. ALL RESULTS NOT CIRCLED WERE QUALIFIED BY THE FOLLOWING STATEMENT:
 Samples with compound concentrations within five times the associated field blank concentration are listed above, these sample results were qualified as not detected, "U".

LDC #: 23188F21
 SDG #: See Cover

VALIDATION FINDINGS WORKSHEET

Field Blanks

Page: 1 of 1
 Reviewer: Q
 2nd Reviewer: _____

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

N/A Were field blanks identified in this SDG?

Blank units: pg/L Associated sample units: pg/g

Sampling date: 4/6/10

Field blank type: (circle one) Field Blank / Rinsate / Other: _____ Associated Samples: 4

Compound	Blank ID	5X	Sample Identification				
	ES04062010-RZR	5X					
E	0.68	0.0034					
F	2.5	0.0125					
G	6.2	0.031					
H	2.7	0.0135					
K	1.4	0.007					
L	0.82	0.0041					
N	0.94	0.0047					
O	1.8	0.009					
P	1.2	0.006					
Q	4.4	0.022					
CRQL							

CIRCLED RESULTS WERE NOT QUALIFIED. ALL RESULTS NOT CIRCLED WERE QUALIFIED BY THE FOLLOWING STATEMENT:
 Samples with compound concentrations within five times the associated field blank concentration are listed above, these sample results were qualified as not detected, "U".

VALIDATION FINDINGS WORKSHEET
Matrix Spike/Matrix Spike Duplicates

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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? If no, indicate which matrix does not have an associated MS/MSD, Soil / Water.

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

#	Date	MS/MSD ID	Compound	MS %R (Limits)	MSD %R (Limits)	RPD (Limits)	Associated Samples	Qualifications
		5/6	Soil	() ()	several () ()	() ()	1	No Qual (6.0.5 in)

LDC #: 231088
 SDG #: 5810200

VALIDATION FINDINGS WORKSHEET
Compound Quantitation and Reported CRQLs

Page: 10 of 19
 Reviewer: _____
 2nd Reviewer: _____

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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

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

#	Date	Sample ID	Finding	Associated Samples	Qualifications
		1	<i>gals > calib range</i>		<i>NR (R)</i>
		2	<i>H</i>		
		All	<i># empc results (R flag)</i>		<i>NR (R)</i>

Comments: See sample calculation verification worksheet for recalculations

VALIDATION FINDINGS WORKSHEET
Field Duplicates

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

Y N NA Were field duplicate pairs identified in this SDG?
Y N NA Were target analytes detected in the field duplicate pairs?

Compound	Concentration (pg/g)		(≤50)	(pg/g)	(pg/g)	Qualifications (Parent Only)
	SSAJ3-02-1BPC	3	RPD	Difference	Limits	
A	0.22	1.1U		0.88	(≤1.1)	
B	0.90	5.3U		4.4	(≤5.3)	
C	0.99	5.3U		4.31	(≤5.3)	
D	1.2	0.13		1.07	(≤5.3)	
E	1.1	0.12		0.98	(≤5.3)	
F	1.6	0.41		1.19	(≤5.3)	
G	5.9	1.4		4.5	(≤11)	
H	0.90	0.60		0.3	(≤1.1)	
I	2.1	0.69		1.41	(≤5.3)	
J	1.5	0.39		1.11	(≤5.3)	
K	1.7	1.2		0.5	(≤5.3)	
L	2.2	0.76		1.44	(≤5.3)	
M	1.5	0.16		1.34	(≤5.3)	
N	1.4	0.18		1.22	(≤5.3)	
O	3.2	2.7		0.5	(≤5.3)	
P	2.1	0.99		1.11	(≤5.3)	
Q	9.1	7.8		1.3	(≤11)	

Compound	Concentration (pg/g)		(≤50)	(pg/g)	(pg/g)	Qualifications (Parent Only)
	SSAN7-02-1BPC	2	RPD	Difference	Limits	
A	2.4	2.4		0	(≤1.1)	
B	11	10		1	(≤5.3)	
C	6.6	5.3		1.3	(≤5.3)	
D	15	12		3	(≤5.3)	
E	11	8.6		2.4	(≤5.3)	
F	39	35	11			
G	40	35		5	(≤11)	

LDC#: 23188F21

SDG#: See Cover

VALIDATION FINDINGS WORKSHEET

Field Duplicates

Page: 2 of 3

Reviewer: *[Signature]*

2nd Reviewer: _____

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

Y N NA
Y N NA

Were field duplicate pairs identified in this SDG?

Were target analytes detected in the field duplicate pairs?

Compound	Concentration (pg/g)		(<50) RPD	(pg/g) Difference	(pg/g) Limits	Qualifications (Parent Only)
	SSAN7-02-1BPC	2				
H	130	170	27			
I	160	210	27			
J	77	82	6			
K	230	250	8			
L	200	160	22			
M	44	35	23			
N	42	21		21	(≤5.3)	<i>ndets/A (fd)</i>
O	500	440	13			
P	270	220	20			
Q	1300	1200	8			

Compound	Concentration (pg/g)		(<50) RPD	(pg/g) Difference	(pg/g) Limits	Qualifications (Parent Only)
	SA156-3BPC	4				
A	0.15	0.54U		0.39	(≤0.54)	
B	0.56	2.7U		2.14	(≤2.7)	
C	0.84	2.7U		1.86	(≤2.7)	
D	0.68	2.7U		2.02	(≤2.7)	
E	0.81	2.7U		1.89	(≤2.7)	
F	1.4	2.7U		1.3	(≤2.7)	
G	5.6	5.4U		0.2	(≤5.4)	
H	0.51	0.16		0.35	(≤0.54)	
I	1.7	2.7U		1	(≤2.7)	
K	4.1	2.7U		1.4	(≤2.7)	
L	3.3	2.7U		0.6	(≤2.7)	
M	1.4	2.7U		1.3	(≤2.7)	
N	1.7	2.7U		1	(≤2.7)	
O	10	1.2		8.8	(≤2.7)	<i>ndets/A (fd)</i>

LDC#: 23188F21
SDG#: See Cover

VALIDATION FINDINGS WORKSHEET
Field Duplicates

Page: 3 of 3
Reviewer: [Signature]
2nd Reviewer: _____

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

Y N NA Were field duplicate pairs identified in this SDG?
Y N NA Were target analytes detected in the field duplicate pairs?

Compound	Concentration (pg/g)		(≤ 50)	(pg/g)	(pg/g)	Qualifications (Parent Only)
	SA156-3BPC	4	RPD	Difference	Limits	
P	5.1	2.7U		2.4	(≤ 2.7)	
Q	25	5.4		19.6	(≤ 5.4)	Not A (fd)

V:\FIELD DUPLICATES\23188F21.wpd

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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	
				Average RRF (initial)	RRF (S/Std)	Average RRF (initial)	RRF (S/Std)	%RSD	%RSD		
1	1CAT (303)	3/4/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	0.98315	1.01872	0.98315	1.01872	4.65926	4.65926	4.659	
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)	1.05105	1.11325	1.05105	1.11325	7.43940	7.43940	7.440	
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)	1.08249	1.09101	1.08249	1.09101	5.63096	5.63096	5.631	
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)	1.07268	1.08562	1.07268	1.08562	4.49467	4.49467	4.495	
			OCDF (¹³ C-OCDD)	1.42582	1.57900	1.42582	1.57900	8.93881	8.93881	8.939	
2	1CAT (AD5)	4/12/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	0.945	0.98	0.945	0.98	4.44	4.44	4.33	
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)	1.021	1.04	1.021	1.04	3.03	3.03	2.97	
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)	1.114	1.19	1.114	1.19	5.33	5.33	5.25	
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)	1.072	1.11	1.072	1.11	3.60	3.60	3.75	
			OCDF (¹³ C-OCDD)	1.445	1.51	1.445	1.51	5.83	5.83	5.89	
3	1CAT	4/21/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	1.088	1.10	1.088	1.10	1.29	1.29	1.20	
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)								
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)								
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)								
			OCDF (¹³ 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.

LDC #: 23108f2
 SDG #: 2210001

VALIDATION FINDINGS WORKSHEET
Routine Calibration Results Verification

Page: 6 of 1
 Reviewer: _____
 2nd Reviewer: _____

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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 = $100 * (\text{ave. RRF} - \text{RRF}) / \text{ave. RRF}$ Where: ave. RRF = initial calibration average RRF
 RRF = $(A_x)(C_s) / (A_s)(C_x)$ RRF = continuing calibration RRF
 A_x = Area of compound, A_s = Area of associated internal standard
 C_x = Concentration of compound, C_s = Concentration of internal standard

#	Standard ID	Calibration Date	Compound (Reference Internal Standard)	Average RRF (Initial)	Reported		Recalculated	
					RRF (CC)	%D	RRF (CC)	%D
1	23AP10432	4/08/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	0.98315	0.92144	6.3	0.92144	6.3
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)	1.05705	1.10087	4.7	1.10087	4.7
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)	1.082449	1.13255	4.6	1.13255	4.6
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)	1.03068	1.00820	2.2	1.00820	2.2
			OCDF (¹³ C-OCDD)	1.42582	1.38551	2.8	1.38551	2.8
2	23AP10405	4/24/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	0.945	0.90	4.3	0.90	4.3
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)	1.02	0.96	6.3	0.96	6.3
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)	1.112	1.08	3.1	1.08	3.1
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)	1.072	1.03	3.9	1.03	3.9
			OCDF (¹³ C-OCDD)	1.445	1.36	5.9	1.36	5.9
3	23AP10432	5/3/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	1.088	1.00	8.1	1.00	8.1
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)					
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)					
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)					
			OCDF (¹³ C-OCDD)					

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
Matrix Spike/Matrix Spike Duplicates Results Verification

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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 = $100 * (SSR - SR) / SA$

Where: SSR = Spiked sample result, SR = Sample result
SA = Spike added

RPD = $|MSR - MSDR| * 2 / (MSR + MSDR)$

MSR = Matrix spike percent recovery MSDR = Matrix spike duplicate percent recovery

MS/MSD samples: 5/6

Compound	Spike Added (ppg)		Sample Concentration (ppg)	Spiked Sample Concentration (ppg)		Matrix Spike Percent Recovery		Matrix Spike Duplicate Percent Recovery		Reported RPD	Recalculated RPD
	MS	MSD		MS	MSD	Reported	Recalc.	Reported	Recalc.		
2,3,7,8-TCDD	212	210	5.1	29.1	31.6	113	113	126	126	8.7	8.7
1,2,3,7,8-PeCDD	106	105	1.6	139	152	116	116	130	130	9.1	8.9
1,2,3,4,7,8-HxCDD	↓	↓	9.6	116	142	101	100	126	126	20	20
1,2,3,4,7,8,9-HpCDF	↓	↓	490	715	733	209	212	229	231	20	20.5
OCDF	212	210	3900	4340	5000	210	208	551	548	15	15

Comments: Refer to Matrix Spike/Matrix Spike 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
Laboratory Control Sample Results Verification

METHOD: GC/MS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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 = $|(LCS - LCSD) / ((LCS + LCSD) / 2)| * 100$ LCS = Laboratory control sample percent recovery LCSD = Laboratory control sample duplicate percent recovery

LCS ID: 010920

Compound	Spike Added (PPM)		Spiked Sample Concentration		LCS		LCSD		Percent Recovery		Percent Recovery		RPD	
	LCS	LCSD	LCS	LCSD	Reported	Recalc	Reported	Recalc	Reported	Recalc	Reported	Recalc	Reported	Recalc
2,3,7,8-TCDD	200	NA	19.8	NA	99	99								
1,2,3,7,8-PeCDD	100		10.2		102	102								
1,2,3,4,7,8-HxCDD	↓		10.8		108	108								
1,2,3,4,7,8,9-HpCDF	↓		10.4		104	104								
OCDF	200	↓	20.2	↓	101	101								

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.

Descriptor	Accurate mass ^(b)	Ion ID	Elemental Composition	Analyte	Descriptor	Accurate Mass ^(a)	Ion ID	Elemental Composition	Analyte		
1	303.9016	M	C ₁₂ H ₄ ³⁵ Cl ₄ O	TCDF	4	407.7818	M+2	C ₁₂ H ³⁵ Cl ₃ ³⁷ ClO	HpCDF		
	305.8987	M+2	C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ Cl ₁₀	TCDF		409.7788	M+4	C ₁₂ H ³⁵ Cl ₂ ³⁷ Cl ₂ O	HpCDF		
	315.9419	M	¹³ C ₁₂ H ₄ ³⁵ Cl ₄ O	TCDF (S)		417.8250	M	¹³ C ₁₂ H ³⁵ Cl ₃ O	HpCDF (S)		
	317.9389	M+2	¹³ C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO	TCDF (S)		419.8220	M+2	¹³ C ₁₂ H ³⁵ Cl ₂ ³⁷ ClO	HpCDF		
	319.8965	M	C ₁₂ H ₄ ³⁵ Cl ₄ O ₂	TCDD		423.7767	M+2	C ₁₂ H ³⁵ Cl ₃ ³⁷ ClO ₂	HpCDD		
	321.8936	M+2	C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO ₂	TCDD		425.7737	M+4	C ₁₂ H ³⁵ Cl ₂ ³⁷ Cl ₂ O ₂	HpCDD		
	331.9368	M	¹³ C ₁₂ H ₄ ³⁵ Cl ₄ O ₂	TCDD (S)		435.8169	M+2	¹³ C ₁₂ H ³⁵ Cl ₃ ³⁷ ClO ₂	HpCDD (S)		
	333.9338	M+2	¹³ C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO ₂	TCDD (S)		437.8140	M+4	¹³ C ₁₂ H ³⁵ Cl ₂ ³⁷ Cl ₂ O ₂	HpCDD (S)		
	375.8364	M+2	C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ ClO	HxCDFE		479.7165	M+4	C ₁₂ H ³⁵ Cl ₂ ³⁷ Cl ₂ O	NCDFE		
	[354.9792]	LOCK	C ₉ F ₁₃	PFK		[430.9728]	LOCK	C ₉ F ₁₇	PFK		
	2	339.8597	M+2	C ₁₂ H ₃ ³⁵ Cl ₄ ³⁷ ClO		PeCDF	5	441.7428	M+2	C ₁₂ ³⁵ Cl ₇ ³⁷ ClO	OCDF
		341.8567	M+4	C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ Cl ₂ O		PeCDF		443.7399	M+4	C ₁₂ ³⁵ Cl ₆ ³⁷ Cl ₂ O	OCDF
		351.9000	M+2	¹³ C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ Cl ₁₀		PeCDF (S)		457.7377	M+2	C ₁₂ ³⁵ Cl ₃ ³⁷ ClO ₂	OCDD
		353.8970	M+4	¹³ C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ Cl ₂ O		PeCDF (S)		459.7348	M+4	C ₁₂ ³⁵ Cl ₂ ³⁷ Cl ₂ O ₂	OCDD
355.8546		M+2	C ₁₂ H ₃ ³⁵ Cl ₄ ³⁷ ClO ₂	PeCDD	469.7780	M+2		¹³ C ₁₂ ³⁵ Cl ₇ ³⁷ ClO ₂	OCDD (S)		
357.8516		M+4	C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ Cl ₂ O ₂	PeCDD	471.7750	M+4		¹³ C ₁₂ ³⁵ Cl ₆ ³⁷ Cl ₂ O ₂	OCDD (S)		
367.8949		M+2	¹³ C ₁₂ H ₃ ³⁵ Cl ₄ ³⁷ ClO ₂	PeCDD (S)	513.6775	M+4		C ₁₂ ³⁵ Cl ₈ ³⁷ Cl ₂ O	DCDFE		
369.8919		M+4	¹³ C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ Cl ₂ O ₂	PeCDD (S)	[422.9278]	LOCK		C ₁₀ F ₁₇	PFK		
409.7974		M+2	C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO	HxCDFE							
[354.9792]		LOCK	C ₉ F ₁₃	PFK							
3		373.8208	M+2	C ₁₂ H ₂ ³⁵ Cl ₅ ³⁷ ClO	HxCDF						
		375.8178	M+4	C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ Cl ₂ O	HxCDF						
		383.8639	M	¹³ C ₁₂ H ₂ ³⁵ Cl ₆ O	HxCDF (S)						
		385.8610	M+2	¹³ C ₁₂ H ₂ ³⁵ Cl ₅ ³⁷ ClO	HxCDF (S)						
	389.8156	M+2	C ₁₂ H ₂ ³⁵ Cl ₃ ³⁷ ClO ₂	HxCDD							
	391.8127	M+4	C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ ClO ₂	HxCDD							
	401.8559	M+2	¹³ C ₁₂ H ₂ ³⁵ Cl ₅ ³⁷ ClO ₂	HxCDD (S)							
	403.8529	M+4	¹³ C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ Cl ₂ O ₂	HxCDD (S)							
	445.7555	M+4	C ₁₂ H ₂ ³⁵ Cl ₆ ³⁷ Cl ₂ O	OCDFE							
	[430.9728]	LOCK	C ₉ F ₁₇	PFK							

(a) The following nucleic masses were used:

- H = 1.007825
- C = 12.000000
- ¹³C = 13.003355
- F = 18.9984
- O = 15.994915
- ³⁵Cl = 34.968853
- ³⁷Cl = 36.965903

S = internal/recovery standard

VALIDATION FINDINGS WORKSHEET
Sample Calculation Verification

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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 agree within 10.0% of the reported results?

$$\text{Concentration} = \frac{A_x(L)(DF)}{(A_s)(RRF)(V_s)(\%S)}$$

- A_x = Area of the characteristic ion (EICP) for the compound to be measured
- A_s = Area of the characteristic ion (EICP) for the specific internal standard
- I_s = Amount of internal standard added in nanograms (ng)
- V_s = Volume or weight of sample extract in milliliters (ml) or grams (g).
- RRF = Relative Response Factor (average) from the initial calibration
- Df = Dilution Factor.
- %S = Percent solids, applicable to soil and solid matrices only.

Example:

Sample I.D. 4, 2:

$$\text{Conc.} = \frac{(1453245)(4000)}{(8188738)(1.42582)(10)(0.918)}$$

= 5.4 ppm/g

#	Sample ID	Compound	Reported Concentration ()	Calculated Concentration ()	Qualification

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

Project/Site Name: Tronox LLC Facility, PCS, Henderson, Nevada

Collection Date: April 14, 2010

LDC Report Date: May 24, 2010

Matrix: Soil

Parameters: Dioxins/Dibenzofurans

Validation Level: Stage 4

Laboratory: TestAmerica, Inc.

Sample Delivery Group (SDG): GOD160437

Sample Identification

SSA04-01-1BPC

SA106-3BPC

Introduction

This data review covers 2 soil samples listed on the cover sheet including dilutions and reanalysis as applicable. The analyses were per EPA SW 846 Method 8290 for Polychlorinated Dioxins/Dibenzofurans.

This review follows the Standard Operating Procedures (SOP) 40, Data Review/Validation (BRC 2009), the Quality Assurance Project Plan Tronox LLC Facility, Henderson, Nevada (June 2009), NDEP guidance (May 2006), and USEPA Contract Laboratory Program National Functional Guidelines for Polychlorinated Dioxins/Dibenzofurans Data Review (September 2005).

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.

Blank results are summarized in Section V.

Field duplicates are summarized in Section XIV.

The following are definitions of the data qualifiers:

- J+ Data are qualified as estimated, with a high bias likely to occur. False positives or false negatives are unlikely to have been reported.
- J- Data are qualified as estimated, with a low bias likely to occur. False positives or false negatives are unlikely to have been reported.
- J Data are qualified as estimated; it is not possible to assess the direction of the potential bias. False positives or false negatives are unlikely to have been reported.
- U Indicates the compound or analyte was analyzed for but not detected at or above the stated limit.
- R Data are qualified as rejected. There is a significant potential for the reporting of false negatives or false positives.
- UU Indicates the compound or analyte was analyzed for but not detected. The sample detection limit is an estimated value.
- B The analytical result may be a false positive totally attributable to blank contamination. This qualifier is applicable to radiochemistry analysis only.
- JB The analytical result may be biased high and partially attributable to blank contamination. This qualifier is applicable to radiochemistry analysis only.
- JK The analytical result is an estimated maximum possible concentration (EMPC).
- X The analytical result is not used for reporting because a more accurate and precise result is reported in its place.
- J-TDS The analytical result is estimated based on failure of the Total Dissolved Solids (TDS) correctness check performed in accordance with the Standard Method 1030E.
- J-CAB The analytical result is estimated based on failure of the cation-anion balance correctness check performed in accordance with Standard Method 1030E.
- J-TDS & CAB The analytical result is unreliable based on the failure of the cation-anion balance and TDS correctness check performed in accordance with standard Method 1030E.
- A Indicates the finding is based upon technical validation criteria.
- P Indicates the finding is related to a protocol/contractual deviation.
- None Indicates the data was not significantly impacted by the finding, therefore qualification was not required.

I. Technical Holding Times

All technical holding time requirements were met.

The chain-of-custodies were reviewed for documentation of cooler temperatures. All cooler temperatures met validation criteria.

II. HRGC/HRMS Instrument Performance Check

Instrument performance was checked at the required daily 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 exact mass of 380.9760 of PFK was verified. The static resolving power was at least 10,000 (10% valley definition).

III. Initial Calibration

A five point initial calibration was performed as required by the method.

Percent relative standard deviations (%RSD) were less than or equal to 20.0% for unlabeled compounds and less than or equal to 30.0% for labeled compounds.

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

The minimum S/N ratio for each target compound was greater than or equal to 2.5 and greater than or equal to 10 for each recovery and internal standard compound for samples.

IV. Routine Calibration (Continuing)

Routine calibration was performed at the required frequencies.

All of the routine calibration percent differences (%D) between the initial calibration RRF and the routine calibration RRF were less than or equal to 20.0% for unlabeled compounds and less than or equal to 30.0% for labeled compounds.

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

V. Blanks

Method blanks were reviewed for each matrix as applicable. No polychlorinated dioxin/dibenzofuran contaminants were found in the method blanks with the following exceptions:

Method Blank ID	Extraction Date	Compound	Concentration	Associated Samples
0110455MB	4/20/10	2,3,7,8-TCDF 1,2,3,7,8-PeCDF 1,2,3,4,6,7,8-HpCDF	0.31 pg/g 0.31 pg/g 0.30 pg/g	All samples in SDG G0D160437

Sample concentrations were compared to concentrations detected in the method blanks as required by the QAPP. No sample data was qualified with the following exceptions:

Sample	Compound	Reported Concentration	Modified Final Concentration
SA106-3BPC	2,3,7,8-TCDF	1.4 pg/g	1.4U pg/g

Sample FB-04072010-RZC (from SDG G0D130519) was identified as a field blank. No polychlorinated dioxin/dibenzofuran contaminants were found in this blank with the following exceptions:

Field Blank ID	Sampling Date	Compound	Concentration	Associated Samples
FB-04072010-RZC	4/8/10	1,2,3,4,7,8-HxCDD 1,2,3,6,7,8-HxCDD 1,2,3,7,8,9-HxCDD 1,2,3,4,6,7,8-HpCDD OCDD 2,3,7,8-TCDF 1,2,3,7,8-PeCDF 2,3,4,7,8-PeCDF 1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 2,3,4,6,7,8-HxCDF 1,2,3,7,8,9-HxCDF 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	0.77 pg/L 0.74 pg/L 0.82 pg/L 4.2 pg/L 37 pg/L 0.57 pg/L 0.96 pg/L 0.67 pg/L 1.1 pg/L 0.96 pg/L 1.0 pg/L 1.0 pg/L 2.1 pg/L 1.5 pg/L 6.7 pg/L	All samples in SDG G0D160437

Sample concentrations were compared to concentrations detected in the field blanks as required by the QAPP. No sample data was qualified.

VI. Matrix Spike/Matrix Spike Duplicates

Matrix spike (MS) and matrix spike duplicate (MSD) samples were reviewed for each matrix as applicable. Percent recoveries (%R) and relative percent differences (RPD) were not within QC limits. Since there were no associated samples, no data were qualified.

VII. Laboratory Control Samples (LCS)

Laboratory control samples were reviewed for each matrix as applicable. Percent recoveries (%R) were within QC limits.

VIII. Regional Quality Assurance and Quality Control

Not applicable.

IX. Internal Standards

All internal standard recoveries were within QC limits with the following exceptions:

Sample	Internal Standards	%R (Limits)	Compound	Flag	A or P
SSAO4-01-1BPC	¹³ C-OCDD	34 (40-135)	OCDD OCDF	J (all detects) UJ (all non-detects) J (all detects) UJ (all non-detects)	P

X. Target Compound Identifications

All target compound identifications were within validation criteria.

XI. Project Quantitation Limit

All compound quantitation and CRQLs were within validation criteria with the following exceptions:

Sample	Compound	Finding	Criteria	Flag	A or P
SSAO4-01-1BPC	OCDF	Sample result exceeded calibration range.	Reported result should be within calibration range.	J (all detects)	P

All compounds reported below the PQL were qualified as follows:

Sample	Finding	Flag	A or P
All samples in SDG G0D160437	All compounds reported below the PQL.	J (all detects)	A

All compounds reported as EMPC were qualified as follows:

Sample	Finding	Flag	A or P
All samples in SDG G0D160437	All compounds reported as estimated maximum possible concentration (EMPC).	JK (all detects)	A

XII. System Performance

The system performance was acceptable.

XIII. Overall Assessment of Data

Data flags are summarized at the end of this report if data has been qualified.

XIV. Field Duplicates

No field duplicates were identified in this SDG.

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Data Qualification Summary - SDG G0D160437**

SDG	Sample	Compound	Flag	A or P	Reason (Code)
G0D160437	SSAO4-01-1BPC	OCDD OCDF	J (all detects) UJ (all non-detects) J (all detects) UJ (all non-detects)	P	Internal standards (%R) (l)
G0D160437	SSAO4-01-1BPC	OCDF	J (all detects)	P	Compound quantitation and CRQLs (e)
G0D160437	SSAO4-01-1BPC SA106-3BPC	All compounds reported below the PQL.	J (all detects)	A	Project Quantitation Limit (sp)
G0D160437	SSAO4-01-1BPC SA106-3BPC	All compounds reported as EMPC	JK (all detects)	A	Project Quantitation Limit (k)

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Laboratory Blank Data Qualification Summary - SDG
G0D160437**

SDG	Sample	Compound	Modified Final Concentration	A or P	Code
G0D160437	SA106-3BPC	2,3,7,8-TCDF	1.4U pg/g	A	bl

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Field Blank Data Qualification Summary - SDG G0D160437**

No Sample Data Qualified in this SDG

Tronox Northgate Henderson

VALIDATION COMPLETENESS WORKSHEET

LDC #: 23188G21
 SDG #: G0D160437
 Laboratory: Test America

Stage 4

Date: 5/24/10
 Page: 1 of 1
 Reviewer: [Signature]
 2nd Reviewer: _____

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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.	Technical holding times	A	Sampling dates: 4/14/10
II.	HRGC/HRMS Instrument performance check	A	
III.	Initial calibration	A	
IV.	Routine calibration/ICV	A	
V.	Blanks	SW	
VI.	Matrix spike/Matrix spike duplicates	SW	No sp/ass'd - No Qual
VII.	Laboratory control samples	A	100%
VIII.	Regional quality assurance and quality control	N	
IX.	Internal standards	SW	
X.	Target compound identifications	A	
XI.	Compound quantitation and CRQLs	SW	
XII.	System performance	A	
XIII.	Overall assessment of data	A	
XIV.	Field duplicates	N	
XV.	Field blanks	SW	FB-040710-RZC (FOD130519)

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

Validated Samples:

1	SSAO4-01-1BPC	S	11	0110455MB	21		31	
2	SA106-3BPC	V	12		22		32	
3			13		23		33	
4			14		24		34	
5			15		25		35	
6			16		26		36	
7			17		27		37	
8			18		28		38	
9			19		29		39	
10			20		30		40	

Notes: _____

Method: Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

Validation Area	Yes	No	NA	Findings/Comments
I. Technical holding times				
All technical holding times were met.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Cooler temperature criteria was met.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
II. GC/MS Instrument performance check				
Was PFK exact mass 380.9760 verified?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were the retention time windows established for all homologues?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was the chromatographic resolution between 2,3,7,8-TCDD and peaks representing any other unlabeled TCDD isomers $\leq 25\%$?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Is the static resolving power at least 10,000 (10% valley definition)?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was the mass resolution adequately check with PFK?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was the presence of 1,2,8,9-TCDD and 1,3,4,6,8-PeCDF verified?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
III. Initial calibration				
Was the initial calibration performed at 5 concentration levels?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were all percent relative standard deviations (%RSD) $\leq 20\%$ for unlabeled standards and $\leq 30\%$ for labeled standards?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Did all calibration standards meet the Ion Abundance Ratio criteria?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was the signal to noise ratio for each target compound ≥ 2.5 and for each recovery and internal standard > 10 ?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
IV. Continuing calibration				
Was a routine calibration performed at the beginning and end of each 12 hour period?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were all percent differences (%D) $\leq 20\%$ for unlabeled standards and $\leq 30\%$ for labeled standards?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Did all routine calibration standards meet the Ion Abundance Ratio criteria?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
V. Blanks				
Was a method blank associated with every sample in this SDG?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was a method blank performed for each matrix and concentration?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was there contamination in the method blanks? If yes, please see the Blanks validation completeness worksheet?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
VI. Matrix spike/Matrix spike duplicates				
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.	<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"/>	
VII. Laboratory control samples				
Was an LCS analyzed for this SDG?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
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"/>	

VIII. Regional Quality Assurance and Quality Control			
Were performance evaluation (PE) samples performed?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Were the performance evaluation (PE) samples within the acceptance limits?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
IX. Internal standards			
Were internal standard recoveries within the 40-135% criteria?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Was the minimum S/N ratio of all internal standard peaks > 10?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
X. Target compound identification			
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?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
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?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
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?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Did compound spectra contain all characteristic ions listed in the table attached?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Was the Ion Abundance Ratio for the two quantitation ions within criteria?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Was the signal to noise ratio for each target compound and labeled standard > 2.5?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Does the maximum intensity of each specified characteristic ion coincide within ± 2 seconds (includes labeled standards)?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
For PCDF identification, was any signal (S/N ≥ 2.5, at ± seconds RT) detected in the corresponding PCDPE channel?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>
Was an acceptable lock mass recorded and monitored?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
XI. Compound quantitation/CRQLs			
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 CRQLs 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"/>
XII. System performance			
System performance was found to be acceptable.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
XIII. Overall assessment of data			
Overall assessment of data was found to be acceptable.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
XIV. Field duplicates			
Field duplicate pairs were identified in this SDG.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Target compounds were detected in the field duplicates.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
XV. Field blanks			
Field blanks were identified in this SDG.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Target compounds were detected in the field blanks.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>

VALIDATION FINDINGS WORKSHEET

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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:

LDC # 23188421
SDG # See Cover

VALIDATION FINDINGS WORKSHEET
Field Blanks

Page: 1 of 1
Reviewer: [Signature]
2nd Reviewer: [Signature]

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

Y N N/A Were field blanks identified in this SDG?

Blank units: pg/L **Associated sample units:** pg/g

Sampling date: 4/8/10

Field blank type: (circle one) Field Blank Rinsate / Other: 5X Associated Samples: AM (>5x)

Compound	Blank ID	Sample Identification
	ER-04072010-RZC	5X
C	0.77	0.00385
D	0.74	0.0037
E	0.82	0.0041
F	4.2	0.021
G	37	0.185
H	0.57	0.00285
I	0.96	0.0048
J	0.67	0.00335
K	1.1	0.0055
L	0.96	0.0048
M	1.0	0.005
N	1.0	0.005
O	2.1	0.0105
P	1.5	0.0075
Q	6.7	0.0335
CRQL		

CIRCLED RESULTS WERE NOT QUALIFIED. ALL RESULTS NOT CIRCLED WERE QUALIFIED BY THE FOLLOWING STATEMENT:
Samples with compound concentrations within five times the associated field blank concentration are listed above, these sample results were qualified as not detected, "U".

VALIDATION FINDINGS WORKSHEET
Initial Calibration Calculation Verification

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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

$RRF = (A_x)(C_w)/(A_w)(C_x)$
 average RRF = sum of the RRFs/number of standards
 $\%RSD = 100 * (S/X)$
 A_x = Area of compound,
 A_w = Area of associated internal standard
 C_x = Concentration of compound,
 C_w = 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	
				Average RRF (Initial)	RRF (Initial)	Average RRF (Initial)	RRF (CS3 std)	%RSD	%RSD	RRF (CS3 std)	%RSD
1	1012 (105)	1/10/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	0.860	0.87	0.860	0.87	10.4	10.6	0.87	10.6
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)	0.934	0.95	0.934	0.95	10.9	12.8	0.95	12.8
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)	1.058	1.09	1.058	1.09	11.2	11.0	1.09	11.0
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)	0.998	1.05	0.998	1.05	12.2	12.2	1.05	12.2
			OCDF (¹³ C-OCDD)	1.437	1.52	1.437	1.52	14.1	14.0	1.52	14.0
2	1012	4/21/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	1.088	1.10	1.088	1.10	12.9	1.20	1.10	1.20
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)								
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)								
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)								
			OCDF (¹³ C-OCDD)								
3	1012 (405)	4/21/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	0.945	0.98	0.945	0.98	4.44	4.33	0.98	4.33
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)	1.021	1.04	1.021	1.04	8.03	8.97	1.04	8.97
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)	1.114	1.19	1.114	1.19	5.33	5.25	1.19	5.25
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)	1.072	1.11	1.072	1.11	3.60	3.75	1.11	3.75
			OCDF (¹³ C-OCDD)	1.445	1.51	1.445	1.51	5.85	5.89	1.51	5.89

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
Routine Calibration Results Verification

LDC #: 2318852
 SDG #: See below

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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 = $100 * (\text{ave. RRF} - \text{RRF}) / \text{ave. RRF}$ Where: ave. RRF = initial calibration average RRF
 RRF = $(A_x)(C_x) / (A_s)(C_s)$ RRF = continuing calibration RRF
 A_x = Area of compound, A_s = Area of associated internal standard
 C_x = Concentration of compound, C_s = Concentration of internal standard

#	Standard ID	Calibration Date	Compound (Reference Internal Standard)	Average RRF (Initial)	Reported		Recalculated		Reported		Recalculated		
					RRF (CC)	%D	RRF (CC)	%D	RRF (CC)	%D			
1	<u>2318852</u>	<u>4/27/10</u>	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	<u>0.860</u>	<u>0.92</u>	<u>7.0</u>	<u>0.92</u>	<u>7.0</u>	<u>0.92</u>	<u>7.0</u>	<u>7.0</u>	<u>7.0</u>	
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)	<u>0.932</u>	<u>0.93</u>	<u>0.1</u>	<u>0.93</u>	<u>0.1</u>	<u>0.93</u>	<u>0.1</u>	<u>0.93</u>	<u>0.1</u>	<u>0.1</u>
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)	<u>1.058</u>	<u>1.15</u>	<u>8.9</u>	<u>1.15</u>	<u>8.9</u>	<u>1.15</u>	<u>8.9</u>	<u>1.15</u>	<u>8.9</u>	<u>8.9</u>
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)	<u>0.998</u>	<u>1.06</u>	<u>6.2</u>	<u>1.06</u>	<u>6.2</u>	<u>1.06</u>	<u>6.2</u>	<u>1.06</u>	<u>6.2</u>	<u>6.2</u>
			OCDF (¹³ C-OCDD)	<u>1.437</u>	<u>1.52</u>	<u>7.2</u>	<u>1.52</u>	<u>7.2</u>	<u>1.52</u>	<u>7.2</u>	<u>1.52</u>	<u>7.2</u>	<u>7.2</u>
2	<u>2318852</u>	<u>5/3/10</u>	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	<u>0.945</u>	<u>0.938</u>	<u>1.3</u>	<u>0.93</u>	<u>1.3</u>	<u>0.93</u>	<u>1.3</u>	<u>1.3</u>	<u>1.3</u>	
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)	<u>1.021</u>	<u>0.95</u>	<u>6.8</u>	<u>0.95</u>	<u>6.8</u>	<u>0.95</u>	<u>6.8</u>	<u>0.95</u>	<u>6.8</u>	
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)	<u>1.114</u>	<u>1.15</u>	<u>3.4</u>	<u>1.15</u>	<u>3.4</u>	<u>1.15</u>	<u>3.4</u>	<u>1.15</u>	<u>3.4</u>	
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)	<u>1.072</u>	<u>1.01</u>	<u>5.4</u>	<u>1.01</u>	<u>5.4</u>	<u>1.01</u>	<u>5.4</u>	<u>1.01</u>	<u>5.4</u>	
			OCDF (¹³ C-OCDD)	<u>1.445</u>	<u>0.97</u>	<u>3.4</u>	<u>1.40</u>	<u>3.4</u>	<u>1.40</u>	<u>3.4</u>	<u>1.40</u>	<u>3.4</u>	
3	<u>2318852</u>	<u>5/4/10</u>	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	<u>1.088</u>	<u>1.15</u>	<u>5.8</u>	<u>1.15</u>	<u>5.8</u>	<u>1.15</u>	<u>5.8</u>	<u>5.8</u>		
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)										
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)										
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)										
			OCDF (¹³ C-OCDD)										

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.

Descriptor	Accurate mass ^(a)	Ion ID	Elemental Composition	Analyte	Descriptor	Accurate Mass ^(a)	Ion ID	Elemental Composition	Analyte		
1	303.9016	M	C ₁₂ H ₄ ³⁵ Cl ₄ O	TODF	4	407.7818	M+2	C ₁₂ H ³⁵ Cl ₆ ³⁷ ClO	HpCDF		
	305.8987	M+2	C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ Cl ₁ O	TODF		409.7788	M+4	C ₁₂ H ³⁵ Cl ₅ ³⁷ Cl ₂ O	HpCDF		
	315.9419	M	¹³ C ₁₂ H ₄ ³⁵ Cl ₄ O	TODF (S)		417.8250	M	¹³ C ₁₂ H ³⁵ Cl ₇ O	HpCDF (S)		
	317.9389	M+2	¹³ C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ ClO	TODF (S)		419.8220	M+2	¹³ C ₁₂ H ³⁵ Cl ₆ ³⁷ ClO	HpCDF		
	319.8965	M	C ₁₂ H ₄ ³⁵ Cl ₄ O ₂	TCDD		423.7767	M+2	C ₁₂ H ³⁵ Cl ₆ ³⁷ ClO ₂	HpCDD		
	321.8936	M+2	C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ ClO ₂	TCDD		425.7737	M+4	C ₁₂ H ³⁵ Cl ₅ ³⁷ Cl ₂ O ₂	HpCDD		
	331.9368	M	¹³ C ₁₂ H ₄ ³⁵ Cl ₄ O ₂	TCDD (S)		435.8169	M+2	¹³ C ₁₂ H ³⁵ Cl ₆ ³⁷ ClO ₂	HpCDD (S)		
	333.9338	M+2	¹³ C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ ClO ₂	TCDD (S)		437.8140	M+4	¹³ C ₁₂ H ³⁵ Cl ₅ ³⁷ Cl ₂ O ₂	HpCDD (S)		
	375.8364	M+2	C ₁₂ H ₄ ³⁵ Cl ₅ ³⁷ ClO	HxCDFE		479.7165	M+4	C ₁₂ H ³⁵ Cl ₇ ³⁷ ClO	NCDFE		
	[354.9792]	LOCK	C ₉ F ₁₃	PFK		[430.9728]	LOCK	C ₉ F ₁₇	PFK		
	2	339.8597	M+2	C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO		PeCDF	5	441.7428	M+2	C ₁₂ ³⁵ Cl ₇ ³⁷ ClO	OCDF
		341.8567	M+4	C ₁₂ H ₃ ³⁵ Cl ₂ ³⁷ Cl ₂ O		PeCDF		443.7399	M+4	C ₁₂ ³⁵ Cl ₆ ³⁷ Cl ₂ O	OCDF
		351.9000	M+2	¹³ C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO		PeCDF (S)		457.7377	M+2	¹³ C ₁₂ ³⁵ Cl ₇ ³⁷ ClO ₂	OCDD
353.8970		M+4	¹³ C ₁₂ H ₃ ³⁵ Cl ₂ ³⁷ Cl ₂ O	PeCDF (S)	459.7348	M+4		¹³ C ₁₂ ³⁵ Cl ₆ ³⁷ Cl ₂ O ₂	OCDD		
355.8546		M+2	C ₁₂ H ₃ ³⁵ Cl ₄ ³⁷ ClO ₂	PeCDD	469.7780	M+2		¹³ C ₁₂ ³⁵ Cl ₆ ³⁷ ClO ₂	OCDD (S)		
357.8516		M+4	C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ Cl ₂ O ₂	PeCDD	471.7750	M+4		¹³ C ₁₂ ³⁵ Cl ₅ ³⁷ Cl ₂ O ₂	OCDD (S)		
367.8949		M+2	¹³ C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO ₂	PeCDD (S)	513.6775	M+4		C ₁₂ ³⁵ Cl ₉ ³⁷ ClO	DCDFE		
369.8919		M+4	¹³ C ₁₂ H ₃ ³⁵ Cl ₂ ³⁷ Cl ₂ O ₂	PeCDD (S)	[422.9278]	LOCK		C ₁₀ F ₁₇	PFK		
409.7974		M+2	C ₁₂ H ₃ ³⁵ Cl ₆ ³⁷ ClO	HxCDFE							
[354.9792]		LOCK	C ₉ F ₁₃	PFK							
3		373.8208	M+2	C ₁₂ H ₂ ³⁵ Cl ₆ ³⁷ ClO	HxCDF						
		375.8178	M+4	C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ Cl ₂ O	HxCDF						
		383.8639	M	¹³ C ₁₂ H ₂ ³⁵ Cl ₆ O	HxCDF (S)						
	385.8610	M+2	¹³ C ₁₂ H ₂ ³⁵ Cl ₅ ³⁷ ClO	HxCDF (S)							
	389.8156	M+2	C ₁₂ H ₂ ³⁵ Cl ₆ ³⁷ ClO ₂	HxCDD							
	391.8127	M+4	C ₁₂ H ₂ ³⁵ Cl ₅ ³⁷ Cl ₂ O ₂	HxCDD							
	401.8559	M+2	¹³ C ₁₂ H ₂ ³⁵ Cl ₆ ³⁷ ClO ₂	HxCDD (S)							
	403.8529	M+4	¹³ C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ Cl ₂ O ₂	HxCDD (S)							
	445.7555	M+4	C ₁₂ H ₂ ³⁵ Cl ₈ ³⁷ ClO	OCDFE							
	[430.9728]	LOCK	C ₉ F ₁₇	PFK							

(a) The following nucleidic masses were used:

H = 1.007825
 C = 12.000000
¹³C = 13.003355
 F = 18.9984
 O = 15.994915
³⁵Cl = 34.968853
³⁷Cl = 36.965903

S = internal/recovery standard

LDC #: 2318842
SDG #: See cover

VALIDATION FINDINGS WORKSHEET

Sample Calculation Verification

Page: 1 of 1
Reviewer: [Signature]
2nd reviewer: _____

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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 agree within 10.0% of the reported results?

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

- A_x = Area of the characteristic ion (EICP) for the compound to be measured
- A_s = 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).
- RRF = Relative Response Factor (average) from the initial calibration
- Df = Dilution Factor.
- %S = Percent solids, applicable to soil and solid matrices only.

Example:

Sample I.D. L F:

$$\text{Conc.} = \frac{(19461360)(2000)}{(8279220)(1.072)(10.08)(0.927)}$$

= 46.9 mg/g

#	Sample ID	Compound	Reported Concentration ()	Calculated Concentration ()	Qualification

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

Project/Site Name: Tronox LLC Facility, PCS, Henderson, Nevada
Collection Date: April 15, 2010
LDC Report Date: May 21, 2010
Matrix: Soil
Parameters: Dioxins/Dibenzofurans
Validation Level: Stage 2B
Laboratory: TestAmerica, Inc.
Sample Delivery Group (SDG): G0D170488

Sample Identification

SSAN6-03-1BPC
SA58-3BPC

Introduction

This data review covers 2 soil samples listed on the cover sheet including dilutions and reanalysis as applicable. The analyses were per EPA SW 846 Method 8290 for Polychlorinated Dioxins/Dibenzofurans.

This review follows the Standard Operating Procedures (SOP) 40, Data Review/Validation (BRC 2009), the Quality Assurance Project Plan Tronox LLC Facility, Henderson, Nevada (June 2009), NDEP guidance (May 2006), and USEPA Contract Laboratory Program National Functional Guidelines for Polychlorinated Dioxins/Dibenzofurans Data Review (September 2005).

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.

Blank results are summarized in Section V.

Field duplicates are summarized in Section XIV.

Raw data were not reviewed for this SDG. The review was based on QC data.

The following are definitions of the data qualifiers:

- J+ Data are qualified as estimated, with a high bias likely to occur. False positives or false negatives are unlikely to have been reported.
- J- Data are qualified as estimated, with a low bias likely to occur. False positives or false negatives are unlikely to have been reported.
- J Data are qualified as estimated; it is not possible to assess the direction of the potential bias. False positives or false negatives are unlikely to have been reported.
- U Indicates the compound or analyte was analyzed for but not detected at or above the stated limit.
- R Data are qualified as rejected. There is a significant potential for the reporting of false negatives or false positives.
- UJ Indicates the compound or analyte was analyzed for but not detected. The sample detection limit is an estimated value.
- B The analytical result may be a false positive totally attributable to blank contamination. This qualifier is applicable to radiochemistry analysis only.
- JB The analytical result may be biased high and partially attributable to blank contamination. This qualifier is applicable to radiochemistry analysis only.
- JK The analytical result is an estimated maximum possible concentration (EMPC).
- X The analytical result is not used for reporting because a more accurate and precise result is reported in its place.
- J-TDS The analytical result is estimated based on failure of the Total Dissolved Solids (TDS) correctness check performed in accordance with the Standard Method 1030E.
- J-CAB The analytical result is estimated based on failure of the cation-anion balance correctness check performed in accordance with Standard Method 1030E.
- J-TDS & CAB The analytical result is unreliable based on the failure of the cation-anion balance and TDS correctness check performed in accordance with standard Method 1030E.
- A Indicates the finding is based upon technical validation criteria.
- P Indicates the finding is related to a protocol/contractual deviation.
- None Indicates the data was not significantly impacted by the finding, therefore qualification was not required.

I. Technical Holding Times

All technical holding time requirements were met.

The chain-of-custodies were reviewed for documentation of cooler temperatures. All cooler temperatures met validation criteria.

II. HRGC/HRMS Instrument Performance Check

Instrument performance was checked at the required daily 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% .

III. Initial Calibration

A five point initial calibration was performed as required by the method.

Percent relative standard deviations (%RSD) were less than or equal to 20.0% for unlabeled compounds and less than or equal to 30.0% for labeled compounds.

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

IV. Routine Calibration (Continuing)

Routine calibration was performed at the required frequencies.

All of the routine calibration percent differences (%D) between the initial calibration RRF and the routine calibration RRF were less than or equal to 20.0% for unlabeled compounds and less than or equal to 30.0% for labeled compounds.

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

V. Blanks

Method blanks were reviewed for each matrix as applicable. No polychlorinated dioxin/dibenzofuran contaminants were found in the method blanks with the following exceptions:

Method Blank ID	Extraction Date	Compound	Concentration	Associated Samples
0110455MB	4/20/10	2,3,7,8-TCDF 1,2,3,7,8-PeCDF 1,2,3,4,6,7,8-HpCDF	0.31 pg/g 0.31 pg/g 0.30 pg/g	All samples in SDG G0D170488

Sample concentrations were compared to concentrations detected in the method blanks as required by the QAPP. No sample data was qualified with the following exceptions:

Sample	Compound	Reported Concentration	Modified Final Concentration
SA58-3BPC	1,2,3,7,8-PeCDF 1,2,3,4,6,7,8-HpCDF	0.49 pg/g 1.1 pg/g	0.49U pg/g 1.1U pg/g

Sample FB-04072010-RZC (from SDG G0D130519) was identified as a field blank. No polychlorinated dioxin/dibenzofuran contaminants were found in this blank with the following exceptions:

Field Blank ID	Sampling Date	Compound	Concentration	Associated Samples
FB-04072010-RZC	4/8/10	1,2,3,4,7,8-HxCDD 1,2,3,6,7,8-HxCDD 1,2,3,7,8,9-HxCDD 1,2,3,4,6,7,8-HpCDD OCDD 2,3,7,8-TCDF 1,2,3,7,8-PeCDF 2,3,4,7,8-PeCDF 1,2,3,4,7,8-HxCDF 1,2,3,6,7,8-HxCDF 2,3,4,6,7,8-HxCDF 1,2,3,7,8,9-HxCDF 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	0.77 pg/L 0.74 pg/L 0.82 pg/L 4.2 pg/L 37 pg/L 0.57 pg/L 0.96 pg/L 0.67 pg/L 1.1 pg/L 0.96 pg/L 1.0 pg/L 1.0 pg/L 2.1 pg/L 1.5 pg/L 6.7 pg/L	All samples in SDG G0D170488

Sample concentrations were compared to concentrations detected in the field blanks as required by the QAPP. No sample data was qualified.

VI. Matrix Spike/Matrix Spike Duplicates

Matrix spike (MS) and matrix spike duplicate (MSD) samples were reviewed for each matrix as applicable. Percent recoveries (%R) and relative percent differences (RPD) were not within QC limits. Since there were no associated samples, no data were qualified.

VII. Laboratory Control Samples (LCS)

Laboratory control samples were reviewed for each matrix as applicable. Percent recoveries (%R) were within QC limits.

VIII. Regional Quality Assurance and Quality Control

Not applicable.

IX. Internal Standards

All internal standard recoveries were within QC limits with the following exceptions:

Sample	Internal Standards	%R (Limits)	Compound	Flag	A or P
SSAN6-03-1BPC	¹³ C-1,2,3,4,6,7,8-HpCDF ¹³ C-OCDD	38 (40-135) 28 (40-135)	OCDD 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	J (all detects) UJ (all non-detects)	P
SA58-3BPC	¹³ C-1,2,3,4,6,7,8-HpCDF ¹³ C-OCDD	38 (40-135) 20 (40-135)	OCDD 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	J (all detects) UJ (all non-detects)	P

X. Target Compound Identifications

Raw data were not reviewed for this SDG.

XI. Project Quantitation Limit

All compounds reported below the PQL were qualified as follows:

Sample	Finding	Flag	A or P
All samples in SDG G0D170488	All compounds reported below the PQL.	J (all detects)	A

All compounds reported as EMPC were qualified as follows:

Sample	Finding	Flag	A or P
All samples in SDG G0D170488	All compounds reported as estimated maximum possible concentration (EMPC).	JK (all detects)	A

Raw data were not reviewed for this SDG.

XII. System Performance

Raw data were not reviewed for this SDG.

XIII. Overall Assessment of Data

Data flags are summarized at the end of this report if data has been qualified.

XIV. Field Duplicates

No field duplicates were identified in this SDG.

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Data Qualification Summary - SDG G0D170488**

SDG	Sample	Compound	Flag	A or P	Reason (Code)
G0D170488	SSAN6-03-1BPC SA58-3BPC	OCDD 1,2,3,4,6,7,8-HpCDF 1,2,3,4,7,8,9-HpCDF OCDF	J (all detects) UJ (all non-detects)	P	Internal standards (%R) (i)
G0D170488	SSAN6-03-1BPC SA58-3BPC	All compounds reported below the PQL.	J (all detects)	A	Project Quantitation Limit (sp)
G0D170488	SSAN6-03-1BPC SA58-3BPC	All compounds reported as EMPC	JK (all detects)	A	Project Quantitation Limit (k)

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Laboratory Blank Data Qualification Summary - SDG
G0D170488**

SDG	Sample	Compound	Modified Final Concentration	A or P	Code
G0D170488	SA58-3BPC	1,2,3,7,8-PeCDF 1,2,3,4,6,7,8-HpCDF	0.49U pg/g 1.1U pg/g	A	bl

**Tronox LLC Facility, PCS, Henderson, Nevada
Dioxins/Dibenzofurans - Field Blank Data Qualification Summary - SDG G0D170488**

No Sample Data Qualified in this SDG

Tronox Northgate Henderson

VALIDATION COMPLETENESS WORKSHEET

Stage 2B

LDC #: 23188H21
 SDG #: G0D170488
 Laboratory: Test America

Date: 4/10/10
 Page: 1 of 1
 Reviewer: Q
 2nd Reviewer:

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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.	Technical holding times	A	Sampling dates: 4/15/10
II.	HRGC/HRMS Instrument performance check	A	
III.	Initial calibration	A	
IV.	Routine calibration/ICV	A	
V.	Blanks	SW	
VI.	Matrix spike/Matrix spike duplicates	SW	No spl ass'd - No Anal
VII.	Laboratory control samples	A	ICS
VIII.	Regional quality assurance and quality control	N	
IX.	Internal standards	SW	
X.	Target compound identifications	N	
XI.	Compound quantitation and CRQLs	SW	
XII.	System performance	N	
XIII.	Overall assessment of data	A	
XIV.	Field duplicates	N	
XV.	Field blanks	SW	FB-040710 RZC (G0D130519)

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

Validated Samples:

1	SSAN6-03-1BPC	S	11	0110435NB	21		31
2	SA58-3BPC	V	12		22		32
3			13		23		33
4			14		24		34
5			15		25		35
6			16		26		36
7			17		27		37
8			18		28		38
9			19		29		39
10			20		30		40

Notes: _____

VALIDATION FINDINGS WORKSHEET

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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:

LDC #: 2318541
 SDG #: 225042

VALIDATION FINDINGS WORKSHEET

Blanks

Page: 1 of 1
 Reviewer: _____
 2nd Reviewer: _____

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA Method 8290)
 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?
- Y N N/A Was a method blank analyzed for each matrix?
- Y N N/A Was the blank contaminated? If yes, please see qualification below.

Blank extraction date: 4/26/10 Blank analysis date: 4/26/10
 Conc. units: PPB Associated Samples: 175X

Compound	Blank ID	Sample Identification
	0110455NR	
H	0.31	
I	0.31	
D	0.30	

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".

LDC #: 231887
 SDG #: See Cover

VALIDATION FINDINGS WORKSHEET

Field Blanks

Page: 1 of 1
 Reviewer: [Signature]
 2nd Reviewer: [Signature]

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

Y/N N/A Were field blanks identified in this SDG?

Blank units: pg/L Associated sample units: pg/g

Sampling date: 4/8/10

Field blank type: (circle one) Field Blank Rinsate / Other: ML (> 5X) Associated Samples:

Compound	Blank ID	5X	Sample Identification
	EB-04072010-RZC	5X	
C	0.77	0.00385	
D	0.74	0.0037	
E	0.82	0.0041	
F	4.2	0.021	
G	37	0.185	
H	0.57	0.00285	
I	0.96	0.0048	
J	0.67	0.00335	
K	1.1	0.0055	
L	0.96	0.0048	
M	1.0	0.005	
N	1.0	0.005	
O	2.1	0.0105	
P	1.5	0.0075	
Q	6.7	0.0335	
CRQL			

CIRCLED RESULTS WERE NOT QUALIFIED. ALL RESULTS NOT CIRCLED WERE QUALIFIED BY THE FOLLOWING STATEMENT:
 Samples with compound concentrations within five times the associated field blank concentration are listed above, these sample results were qualified as not detected, "U".

LDC #: 23188421
SDG #: see cover

Page: 1 of 9
Reviewer: [Signature]
2nd Reviewer: _____

VALIDATION FINDINGS WORKSHEET

Compound Quantitation and Reported CRQLs

METHOD: HRGC/HRMS Dioxins/Dibenzofurans (EPA SW 846 Method 8290)

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

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

#	Date	Sample ID	Finding	Associated Samples	Qualifications
		41	ZMPC results (R flag)	AM	N (CK)

Comments: See sample calculation verification worksheet for recalculations
