



LABORATORY DATA CONSULTANTS, INC.

7750 El Camino Real, Suite 2L Carlsbad, CA 92009 Phone: 760/634-0437 Fax: 760/634-0439

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

May 20, 2010

SUBJECT: Tronox LLC Facility, 2010 Parcels, Henderson, Nevada,
Data Validation

Dear Ms. Arnold,

Enclosed are the final validation reports for the fractions listed below. These SDGs were received on May 5, 2010. Attachment 1 is a summary of the samples that were reviewed for each analysis.

LDC Project # 23105:

<u>SDG #</u>	<u>Fraction</u>
G0D140559	Polychlorinated Biphenyls, Dioxins/Dibenzofurans
G0D140560	
G0D280454	

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 Superfund Organic Methods Data Review, June 2008
- USEPA, Contract Laboratory Program National Functional Guidelines for Polychlorinated Dioxins/Dibenzofurans 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 #: 23105

SDG #: G0D140559, G0D140560, G0D280454

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_LDC23105_051910.doc
IV. EDD Delivery				
Was the final EDD sent to the client?	X			

Tronox LLC Facility, 2010 Parcels, Henderson, Nevada
Data Validation Reports
LDC #23105

Polychlorinated Biphenyls

LDC

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

Project/Site Name: Tronox LLC Facility, 2010 Parcels, Henderson, Nevada
Collection Date: April 27, 2010
LDC Report Date: May 17, 2010
Matrix: Soil/Water
Parameters: Polychlorinated Biphenyls
Validation Level: Stage 2B & 4
Laboratory: TestAmerica, Inc.

Sample Delivery Group (SDG): G0D280454

Sample Identification

EB-0427200-PF
Q3-PF-3-1-0.0**
Q3-PF-3-1-0.0MS
Q3-PF-3-1-0.0MSD

**Indicates sample underwent Stage 4 review

Introduction

This data review covers 3 soil samples and one water sample listed on the cover sheet including dilutions and reanalysis as applicable. The analyses were per EPA SW 846 Method 8082 for Polychlorinated Biphenyls.

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 a modified outline of the USEPA Contract Laboratory Program National Functional Guidelines for Superfund Organic Methods Data Review (June 2008).

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. GC/ECD Instrument Performance Check

Instrument performance was acceptable unless noted otherwise under initial calibration and continuing calibration sections.

III. Initial Calibration

Initial calibration of multicomponent compounds were performed for the primary (quantitation) column as required by the method.

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

Retention times (RT) of all compounds in the calibration standards were within QC limits.

IV. Continuing Calibration

Continuing calibration was performed at required frequencies.

The percent differences (%D) of calibration factors in continuing standard mixtures were within the 20.0% QC limits.

The percent difference (%D) of the second source calibration standard were less than or equal to 20.0% for all compounds.

Retention times (RT) of all compounds in the calibration standards were within QC limits.

V. Blanks

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

Sample EB-0427200-PF was identified as an equipment blank. No polychlorinated biphenyl contaminants were found in this blank.

VI. Surrogate Spikes

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

VII. 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 one compound, the LCS percent recovery (%R) was within QC limits and no data were qualified.

VIII. Laboratory Control Samples (LCS)

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

IX. Regional Quality Assurance and Quality Control

Not applicable.

X. Pesticide Cleanup Checks

a. Florisil Cartridge Check

Florisil cleanup was not required and therefore not performed in this SDG.

b. GPC Calibration

GPC cleanup was not required and therefore not performed in this SDG.

XI. Target Compound Identification

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.

XII. Project Quantitation Limit

All project quantitation limits were within validation criteria for samples on which a Stage 4 review was performed.

All compounds reported below the PQL were qualified as follows:

Sample	Finding	Flag	A or P
All samples in SDG G0D280454	All compounds reported below the PQL.	J (all detects)	A

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

No field duplicates were identified in this SDG.

**Tronox LLC Facility, 2010 Parcels, Henderson, Nevada
 Polychlorinated Biphenyls - Data Qualification Summary - SDG G0D280454**

SDG	Sample	Compound	Flag	A or P	Reason (Code)
G0D280454	EB-0427200-PF Q3-PF-3-1-0.0**	All compounds reported below the PQL.	J (all detects)	A	Project Quantitation Limit (sp)

**Tronox LLC Facility, 2010 Parcels, Henderson, Nevada
 Polychlorinated Biphenyls - Laboratory Blank Data Qualification Summary - SDG
 G0D280454**

No Sample Data Qualified in this SDG

**Tronox LLC Facility, 2010 Parcels, Henderson, Nevada
 Polychlorinated Biphenyls - Equipment Blank Data Qualification Summary - SDG
 G0D280454**

No Sample Data Qualified in this SDG

LDC #: 23105C3b
 SDG #: G0D280454
 Laboratory: Test America

Tronox Northgate Henderson
VALIDATION COMPLETENESS WORKSHEET
 Stage 2B/4

Date: 5/14/10
 Page: 1 of 1
 Reviewer: [Signature]
 2nd Reviewer: [Signature]

METHOD: GC Polychlorinated Biphenyls (EPA SW 846 Method 8082)

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>4/27/10</u>
II.	GC/ECD Instrument Performance Check	-	
III.	Initial calibration	A	
IV.	Continuing calibration/ICV	A	<u>ICV/CCV ≤ 20%</u>
V.	Blanks	A	
VI.	Surrogate spikes	A	
VII.	Matrix spike/Matrix spike duplicates	<u>SW</u>	
VIII.	Laboratory control samples	A	<u>LCS</u>
IX.	Regional quality assurance and quality control	N	
Xa.	Florisil cartridge check	N	
Xb.	GPC Calibration	N	
XI.	Target compound identification	A	Not reviewed for Stage 2B validation.
XII.	Compound quantitation and reported CRQLs	A	Not reviewed for Stage 2B validation.
XIII.	Overall assessment of data	A	
XIV.	Field duplicates	N	
XV.	Field blanks	<u>ND</u>	<u>EB=1</u>

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	EB-0427200-PF	<u>W</u>	11	<u>0118266MB</u>	21		31	
2	Q3-PF-3-1-0.0**	<u>S</u>	12	<u>0118268MB</u>	22		32	
3	Q3-PF-3-1-0.0MS	<u>I</u>	13		23		33	
4	Q3-PF-3-1-0.0MSD	<u>I</u>	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	

VALIDATION FINDINGS WORKSHEET

METHOD: Pesticide/PCBs (EPASW 846 Method 8081/8082)

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

Notes:

LDC #: 23105C3b
 SDG #: see cover

VALIDATION FINDINGS CHECKLIST

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

Method: GC HPLC

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. Initial calibration				
Did the laboratory perform a 5 point calibration prior to sample analysis?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were all percent relative standard deviations (%RSD) < 20%?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Was a curve fit used for evaluation?	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	
Did the initial calibration meet the curve fit acceptance criteria of > 0.990?	<input type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	
Were the RT windows properly established?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
IV. Continuing calibration				
Was a continuing calibration analyzed daily?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were all percent differences (%D) < 15% ^{20%?} or percent recoveries 85-115%?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
Were all the retention times within the acceptance windows?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
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 analyzed 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 type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	
VI. Surrogate spikes				
Were all surrogate %R within the QC limits?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	
If the percent recovery (%R) for one or more surrogates was out of QC limits, was a reanalysis performed to confirm samples with %R outside of criteria?	<input type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	
VII. 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"/>	
Was a MS/MSD analyzed every 20 samples of each matrix?	<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"/>	
VIII. 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"/>	
IX. Regional Quality Assurance and Quality Control				
Were performance evaluation (PE) samples performed?	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	
Were the performance evaluation (PE) samples within the acceptance limits?	<input type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	

LDC #: 20105036
 SDG #: SEICONE

VALIDATION FINDINGS CHECKLIST

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

Validation Area	Yes	No	NA	Findings/Comments
X. Target compound identification				
Were the retention times of reported detects within the RT windows?	/			
XI. Compound quantitation/CRQLs				
Were compound quantitation and CRQLs adjusted to reflect all sample dilutions and dry weight factors applicable to level IV validation?	/			
XII. System performance				
System performance was found to be acceptable.	/			
XIII. Overall assessment of data				
Overall assessment of data was found to be acceptable.	/			
XIV. Field duplicates				
Field duplicate pairs were identified in this SDG.		/		
Target compounds were detected in the field duplicates.			/	
XV. Field blanks				
Field blanks were identified in this SDG.		/		
Target compounds were detected in the field blanks.			/	

LDC #: 2905C36
 SDG #: BA CON

VALIDATION FINDINGS WORKSHEET
Continuing Calibration Results Verification

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

METHOD: GC ✓ HPLC _____

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

% Difference = $100 * (\text{ave. CF} - \text{CF}) / \text{ave. CF}$ Where: ave. CF = initial calibration average CF
 CF = A/C CF = continuing calibration CF
 A = Area of compound
 C = Concentration of compound

#	Standard ID	Calibration Date	Compound	Average CF (cal)/ CCV Conc.	Reported		Recalculated	
					CF/Conc. CCV	%D	CF/Conc. CCV	%D
1	<u>CCV (11-49)</u>	<u>4/28/10</u>	<u>BB</u>	<u>300</u>	<u>297</u>	<u>1</u>	<u>297</u>	<u>1</u>
2								
3								
4								

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

LDC #: 23105C36
 SDG #: See copy

VALIDATION FINDINGS WORKSHEET
Initial Calibration Calculation Verification

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

METHOD: GC ✓ HPLC _____

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

- CF = A/C
- average CF = sum of the CF/number of standards
- %RSD = $100 * (S/X)$
- A = Area of compound,
- C = Concentration of compound,
- S = Standard deviation of the CF
- X = Mean of the CFs

#	Standard ID	Calibration Date	Compound	Reported		Recalculated		Reported		Recalculated	
				CF (100 std)	CF (100 std)	Average CF (initial)	Average CF (initial)	%RSD	%RSD	%RSD	%RSD
1	1CAZ	4/7/10	BB	96.9119	96.9119	947048	947048	7.615	7.615	7.615	7.615
2											
3											
4											

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

LDC # 23105C3b
 SDG # Reednet

VALIDATION FINDINGS WORKSHEET
Surrogate Results Verification

Page: 1 of 1
 Reviewer: g
 2nd reviewer: l

METHOD: GC HPLC

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

% Recovery: SF/SS * 100
 Where: SF = Surrogate Found
 SS = Surrogate Spiked

Sample ID: 2

Surrogate	Column/Detector	Surrogate Spiked	Surrogate Found	Percent Recovery		Percent Difference
				Reported	Recalculated	
<u>DCB</u>	<u>DB-1701</u>	<u>13.3508</u>	<u>10.9800</u>	<u>82</u>	<u>82</u>	<u>0</u>
<u>TCMX</u>	<u>↓</u>	<u>↓</u>	<u>10.8200</u>	<u>81</u>	<u>81</u>	<u>0</u>

Sample ID: _____

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

Sample ID: _____

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

VALIDATION FINDINGS WORKSHEET
Matrix Spike/Matrix Spike Duplicates Results Verification

METHOD: GC HPLC

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

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

SSC = Spiked sample concentration

SA = Spike added

MSD = Matrix spike duplicate

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

MS/MSD samples: 2/4

Compound	Spike Added (<u>MS</u>)		Spike Sample Concentration (<u>MS</u>)		Sample Conc. (<u>MS</u>)		Matrix spike Percent Recovery		Matrix Spike Duplicate Percent Recovery		MS/MSD RPD	
	MS	MSD	MS	MSD	MS	MSD	Reported	Recalc.	Reported	Recalc.	Reported	Recalc.
Gasoline (8015)												
Diesel (8015)												
Benzene (8021B)												
Methane (RSK-175)												
2,4-D (8151)												
Dinoseb (8151)												
Naphthalene (8310)												
Anthracene (8310)												
HMX (8330)												
2,4,6-Trinitrotoluene (8330)												
<u>BB</u>	<u>66.3</u>	<u>65.9</u>	<u>74.6</u>	<u>73.2</u>	<u>ND</u>	<u>ND</u>	<u>112</u>	<u>112</u>	<u>111</u>	<u>111</u>	<u>1.9</u>	<u>1.9</u>

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

METHOD: GC HPLC

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

% Recovery = $100 * (SSC - SC) / SA$ Where: SSC = Spiked sample concentration SC = Concentration
 SA = Spike added
 RPD = $100 * |SSCLCS - SSCLCSD| / (SSCLCS + SSCLCSD)$ LCS = Laboratory control sample percent recovery
 LCSD = Laboratory control sample duplicate percent recovery

LCS/LCSD samples: 0118-266

Compound	Spike Added (100%)		Spiked Sample Concentration (100%)		LCS		LCSD	
	LCS	LCSD	LCS	LCSD	Reported	Recalc.	Reported	Recalc.
Gasoline (8015)								
Diesel (8015)								
Benzene (8021B)								
Methane (RSK-175)								
2,4-D (8151)								
Dinoseb (8151)								
Naphthalene (8310)								
Anthracene (8310)								
HMX (8330)								
2,4,6-Trinitrotoluene (8330)								
BB	66.7	NA	60.0	NA	90	90		

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

VALIDATION FINDINGS WORKSHEET

Sample Calculation Verification

METHOD: GC HPLC

Y/N N/A
Y/N N/A

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

Concentration = $\frac{A_i(F_v)(D_f)}{(RF)(V_s \text{ or } W_s)(\%S/100)}$

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

Example:
Sample ID: 2 Compound Name ND

Concentration = _____

#	Sample ID	Compound	Reported Concentrations ()	Recalculated Results Concentrations ()	Qualifications

Comments: _____

**Tronox LLC Facility, 2010 Parcels, Henderson, Nevada
Data Validation Reports
LDC #23105**

Dioxins/Dibenzofurans

LDC

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

Project/Site Name: Tronox LLC Facility, 2010 Parcels, Henderson, Nevada
Collection Date: April 13, 2010
LDC Report Date: May 17, 2010
Matrix: Water
Parameters: Dioxins/Dibenzofurans
Validation Level: Stage 2B
Laboratory: TestAmerica, Inc.
Sample Delivery Group (SDG): G0D140559

Sample Identification

EB-PARCELC_033110
FB-PARCELC_033110

Introduction

This data review covers 2 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.
- 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% .

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
4/22/10	¹³ C-1,2,3,4,6,7,8-HpCDD	30.1	All samples in SDG G0D140559	1,2,3,4,6,7,8-HpCDD	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
0113332MB	4/20/10	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	0.49 pg/L 0.64 pg/L 0.75 pg/L 1.3 pg/L 6.0 pg/L 1.5 pg/L 0.91 pg/L 0.63 pg/L 1.1 pg/L 0.48 pg/L 0.46 pg/L 0.56 pg/L 1.2 pg/L 0.96 pg/L 2.8 pg/L	All samples in SDG G0D140559

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-PARCELC_033110	1,2,3,4,7,8-HxCDD 2,3,7,8-TCDF 1,2,3,7,8-PeCDF 2,3,4,7,8-PeCDF 1,2,3,4,7,8-HxCDF 2,3,4,6,7,8-HxCDF 1,2,3,7,8,9-HxCDF 1,2,3,4,7,8,9-HpCDF	1.9 pg/L 2.6 pg/L 1.2 pg/L 1.0 pg/L 2.4 pg/L 1.7 pg/L 1.4 pg/L 2.2 pg/L	1.9U pg/L 2.6U pg/L 1.2U pg/L 1.0U pg/L 2.4U pg/L 1.7U pg/L 1.4U pg/L 2.2U pg/L
FB-PARCELC_033110	1,2,3,4,7,8-HxCDD 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 1,2,3,4,7,8,9-HpCDF	1.1 pg/L 2.2 pg/L 0.67 pg/L 0.73 pg/L 1.6 pg/L 1.7 pg/L 0.99 pg/L 0.71 pg/L	1.1U pg/L 2.2U pg/L 0.67U pg/L 0.73U pg/L 1.6U pg/L 1.7U pg/L 0.99U pg/L 0.71U pg/L

Sample EB-PARCELC_033110 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-PARCELC_033110	4/13/10	2,3,7,8-TCDD 1,2,3,7,8-PeCDD 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	0.68 pg/L 2.2 pg/L 1.9 pg/L 7.7 pg/L 4.5 pg/L 100 pg/L 360 pg/L 2.6 pg/L 1.2 pg/L 1.0 pg/L 2.4 pg/L 2.6 pg/L 1.7 pg/L 1.4 pg/L 24 pg/L 2.2 pg/L 23 pg/L	No associated samples in this SDG

Sample FB-PARCELC_033110 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-PARCELC_033110	4/13/10	2,3,7,8-TCDD 1,2,3,7,8-PeCDD 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	0.34 pg/L 1.3 pg/L 1.1 pg/L 5.9 pg/L 4.0 pg/L 91 pg/L 350 pg/L 2.2 pg/L 0.67 pg/L 0.73 pg/L 1.6 pg/L 1.7 pg/L 0.99 pg/L 20 pg/L 0.71 pg/L 18 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. Although the LCS percent recovery (%R) was not within QC limits for one compound, the LCSD percent recovery (%R) was within QC limits and no data were qualified.

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

Sample	Compound	Finding	Criteria	Flag	A or P
All samples in SDG G0D140559	2,3,7,8-TCDF	2nd column confirmation was not performed for this compound.	This compound must be confirmed on the 2nd column per the method.	None	P

All compounds reported below the PQL were qualified as follows:

Sample	Finding	Flag	A or P
All samples in SDG G0D140559	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 G0D140559	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, 2010 Parcels, Henderson, Nevada
Dioxins/Dibenzofurans - Data Qualification Summary - SDG G0D140559**

SDG	Sample	Compound	Flag	A or P	Reason (Code)
G0D140559	EB-PARCELC_033110 FB-PARCELC_033110	1,2,3,4,6,7,8-HpCDD	J+ (all detects)	P	Routine calibration (%D) (c)
G0D140559	EB-PARCELC_033110 FB-PARCELC_033110	2,3,7,8-TCDF	None	P	Compound quantitation and CRQLs (o)
G0D140559	EB-PARCELC_033110 FB-PARCELC_033110	All compounds reported below the PQL.	J (all detects)	A	Project Quantitation Limit (sp)
G0D140559	EB-PARCELC_033110 FB-PARCELC_033110	All compounds reported as EMPC	JK (all detects)	A	Project Quantitation Limit (k)

**Tronox LLC Facility, 2010 Parcels, Henderson, Nevada
Dioxins/Dibenzofurans - Laboratory Blank Data Qualification Summary - SDG
G0D140559**

SDG	Sample	Compound	Modified Final Concentration	A or P	Code
G0D140559	EB-PARCELC_033110	1,2,3,4,7,8-HxCDD 2,3,7,8-TCDF 1,2,3,7,8-PeCDF 2,3,4,7,8-PeCDF 1,2,3,4,7,8-HxCDF 2,3,4,6,7,8-HxCDF 1,2,3,7,8,9-HxCDF 1,2,3,4,7,8,9-HpCDF	1.9U pg/L 2.6U pg/L 1.2U pg/L 1.0U pg/L 2.4U pg/L 1.7U pg/L 1.4U pg/L 2.2U pg/L	A	bl
G0D140559	FB-PARCELC_033110	1,2,3,4,7,8-HxCDD 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 1,2,3,4,7,8,9-HpCDF	1.1U pg/L 2.2U pg/L 0.67U pg/L 0.73U pg/L 1.6U pg/L 1.7U pg/L 0.99U pg/L 0.71U pg/L	A	bl

**Tronox LLC Facility, 2010 Parcels, Henderson, Nevada
Dioxins/Dibenzofurans - Equipment Blank Data Qualification Summary - SDG
G0D140559**

No Sample Data Qualified in this SDG

**Tronox LLC Facility, 2010 Parcels, Henderson, Nevada
Dioxins/Dibenzofurans - Field Blank Data Qualification Summary - SDG G0D140559**

No Sample Data Qualified in this SDG

LDC #: 23105A21
 SDG #: GOD140559
 Laboratory: Test America

Tronox Northgate Henderson
VALIDATION COMPLETENESS WORKSHEET
 Stage 2B

Date: 5/14/10
 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: 4/13/10
II.	HRGC/HRMS Instrument performance check	A	
III.	Initial calibration	A	
IV.	Routine calibration/ICV	W	
V.	Blanks	W	
VI.	Matrix spike/Matrix spike duplicates	N	client specified
VII.	Laboratory control samples	W	LOS/D
VIII.	Regional quality assurance and quality control	N	
IX.	Internal standards	A	
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	W	FB = 1 . FB = 2 .

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:

1	EB-PARCELC_033110	W	11	0113332MB	21		31	
2	FB-PARCELC_033110	W	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:

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: 4/22/10

Conc. units: pg/L Associated samples: All (bl)

Compound	Blank ID	Sample Identification				
		5X	1	2		
	0113332MR					
C	0.49	2.45	1.9/U	1.1/U		
D	0.64	3.2	-	-		
E	0.75	3.75	-	-		
F	1.3	6.5	-	-		
G	6.0	30	-	-		
H	1.5	7.5	2.6/U	2.2/U		
I	0.91	4.55	1.2/U	0.67/U		
J	0.63	3.15	1.0/U	0.73/U		
K	1.1	5.5	2.4/U	1.6/U		
L	0.48	2.4	-	1.7/U		
M	0.46	2.3	1.7/U	0.99/U		
N	0.56	2.8	1.4/U			
O	1.2	6	-	-		
P	0.96	4.8	2.2/U	0.71/U		
Q	2.8	14	-	-		

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)

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

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

Sampling date: 4/13/10

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

Compound	Blank ID	Sample Identification
	FB-PARCELC 033110	5X
A	0.68	0.0034
B	2.2	0.011
C	1.9	0.0095
D	7.7	0.0385
E	4.5	0.0225
F	100	0.5
G	360	1.8
H	2.6	0.013
I	1.2	0.006
J	1.0	0.005
K	2.4	0.012
L	2.6	0.013
M	1.7	0.0085
N	1.4	0.007
O	24	0.12
P	2.2	0.011
Q	23	0.115
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 #: 23105A21
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:** -

Sampling date: 4/13/10

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

Associated Samples:

None

Compound	Blank ID	Sample Identification
	ER-PARCEL 033110	5X
A	0.34	0.0017
B	1.3	0.0065
C	1.1	0.0055
D	5.9	0.0295
E	4.0	0.02
F	91	0.455
G	350	1.75
H	2.2	0.011
I	0.67	0.00335
J	0.73	0.00365
K	1.6	0.008
L	1.7	0.0085
M	0.99	0.00495
O	20	0.1
P	0.71	0.00355
Q	18	0.09
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 Laboratory Control Samples (LCS)

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 Was a LCS required?

Y N N/A Was a LCS analyzed every 20 samples for each matrix or whenever a sample extraction was performed?

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

#	Date	Lab ID/Reference	Compound	LCS %R (Limits)	LCS %R (Limits)	RPD (Limits)	Associated Samples	Qualifications
		011333203	M	78 (80-130)	() ()	() ()	M1 + B1	No Qual (LCSO in)
					() ()	() ()		
					() ()	() ()		
					() ()	() ()		
					() ()	() ()		
					() ()	() ()		
					() ()	() ()		
					() ()	() ()		
					() ()	() ()		
					() ()	() ()		
					() ()	() ()		
					() ()	() ()		
					() ()	() ()		
					() ()	() ()		
					() ()	() ()		
					() ()	() ()		
					() ()	() ()		
					() ()	() ()		
					() ()	() ()		
					() ()	() ()		
					() ()	() ()		
					() ()	() ()		
					() ()	() ()		
					() ()	() ()		
					() ()	() ()		
					() ()	() ()		
					() ()	() ()		
					() ()	() ()		

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
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
		<u>All</u>	<u>All EMP C (R flag) results</u>	<u>ML</u>	<u>NA ✓K (K)</u>
		<u>ad</u>	<u>No confirmation</u>	<u>ML</u>	<u>None (10)</u>
			<u>YW 2.3.7.8-TCDF</u>		<u>P</u>

Comments: See sample calculation verification worksheet for recalculations

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

Project/Site Name: Tronox LLC Facility, 2010 Parcels, Henderson, Nevada
Collection Date: April 13, 2010
LDC Report Date: May 17, 2010
Matrix: Soil
Parameters: Dioxins/Dibenzofurans
Validation Level: Stage 2B & 4
Laboratory: TestAmerica, Inc.
Sample Delivery Group (SDG): G0D140560

Sample Identification

I6-PC-1-0.0**
I6-PC-1-0.0FD
I6-PC-1-0.0MS
I6-PC-1-0.0MSD

**Indicates sample underwent Stage 4 review

Introduction

This data review covers 4 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.
- 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% .

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
0106187MB	4/16/10	2,3,7,8-TCDD 1,2,3,4,6,7,8-HpCDD OCDD 2,3,7,8-TCDF 1,2,3,4,7,8-HxCDF 1,2,3,4,6,7,8-HpCDF OCDF	0.036 pg/g 0.14 pg/g 0.45 pg/g 0.088 pg/g 0.054 pg/g 0.076 pg/g 0.15 pg/g	All samples in SDG G0D140560

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
I6-PC-1-0.0**	1,2,3,4,6,7,8-HpCDD 2,3,7,8-TCDF	0.40 pg/g 0.12 pg/g	0.40U pg/g 0.12U pg/g
I6-PC-1-0.0FD	2,3,7,8-TCDF	0.14 pg/g	0.14U pg/g

Sample EB-PARCELC_033110 (from SDG G0D140559) 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-PARCELC_033110	4/13/10	2,3,7,8-TCDD 1,2,3,7,8-PeCDD 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	0.68 pg/L 2.2 pg/L 1.9 pg/L 7.7 pg/L 4.5 pg/L 100 pg/L 360 pg/L 2.6 pg/L 1.2 pg/L 1.0 pg/L 2.4 pg/L 2.6 pg/L 1.7 pg/L 1.4 pg/L 24 pg/L 2.2 pg/L 23 pg/L	All samples in SDG G0D140560

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

Sample	Compound	Reported Concentration	Modified Final Concentration
I6-PC-1-0.0**	1,2,3,4,6,7,8-HpCDD	0.40 pg/g	0.40U pg/g

Sample FB-PARCELC_033110 (from SDG G0D140559) 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-PARCELC_033110	4/13/10	2,3,7,8-TCDD 1,2,3,7,8-PeCDD 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	0.34 pg/L 1.3 pg/L 1.1 pg/L 5.9 pg/L 4.0 pg/L 91 pg/L 350 pg/L 2.2 pg/L 0.67 pg/L 0.73 pg/L 1.6 pg/L 1.7 pg/L 0.99 pg/L 20 pg/L 0.71 pg/L 18 pg/L	All samples in SDG G0D140560

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

Sample	Compound	Reported Concentration	Modified Final Concentration
I6-PC-1-0.0**	1,2,3,4,6,7,8-HpCDD	0.40 pg/g	0.40U pg/g

VI. Matrix Spike/Matrix Spike Duplicates

Matrix spike (MS) and matrix spike duplicate (MSD) samples were reviewed for each matrix as applicable. Although the MSD percent recovery (%R) was not within QC limits for one compound, the MS percent recovery (%R) was 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.

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
All samples in SDG G0D140560	2,3,7,8-TCDF	2nd column confirmation was not performed for this compound.	This compound must be confirmed on the 2nd column per the method.	None	P

All compounds reported below the PQL were qualified as follows:

Sample	Finding	Flag	A or P
All samples in SDG G0D140560	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 G0D140560	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 I6-PC-1-0.0** and I6-PC-1-0.0FD 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
	I6-PC-1-0.0**	I6-PC-1-0.0FD				
1,2,3,4,6,7,8-HpCDD	0.40	0.91	-	0.51 (≤ 5.2)	-	-
OCDD	2.6	13	-	10.4 (≤ 10)	J (all detects)	A
2,3,7,8-TCDF	0.12	0.14	-	0.02 (≤ 1.0)	-	-

**Tronox LLC Facility, 2010 Parcels, Henderson, Nevada
Dioxins/Dibenzofurans - Data Qualification Summary - SDG G0D140560**

SDG	Sample	Compound	Flag	A or P	Reason (Code)
G0D140560	I6-PC-1-0.0** I6-PC-1-0.0FD	2,3,7,8-TCDF	None	P	Compound quantitation and CRQLs (o)
G0D140560	I6-PC-1-0.0** I6-PC-1-0.0FD	All compounds reported below the PQL.	J (all detects)	A	Project Quantitation Limit (sp)
G0D140560	I6-PC-1-0.0** I6-PC-1-0.0FD	All compounds reported as EMPC	JK (all detects)	A	Project Quantitation Limit (k)
G0D140560	I6-PC-1-0.0** I6-PC-1-0.0FD	OCDD	J (all detects)	A	Field duplicates (Difference) (fd)

**Tronox LLC Facility, 2010 Parcels, Henderson, Nevada
Dioxins/Dibenzofurans - Laboratory Blank Data Qualification Summary - SDG G0D140560**

SDG	Sample	Compound	Modified Final Concentration	A or P	Code
G0D140560	I6-PC-1-0.0**	1,2,3,4,6,7,8-HpCDD 2,3,7,8-TCDF	0.40U pg/g 0.12U pg/g	A	bl
G0D140560	I6-PC-1-0.0FD	2,3,7,8-TCDF	0.14U pg/g	A	bl

**Tronox LLC Facility, 2010 Parcels, Henderson, Nevada
Dioxins/Dibenzofurans - Equipment Blank Data Qualification Summary - SDG G0D140560**

SDG	Sample	Compound	Modified Final Concentration	A or P	Code
G0D140560	I6-PC-1-0.0**	1,2,3,4,6,7,8-HpCDD	0.40U pg/g	A	be

**Tronox LLC Facility, 2010 Parcels, Henderson, Nevada
Dioxins/Dibenzofurans - Field Blank Data Qualification Summary - SDG G0D140560**

SDG	Sample	Compound	Modified Final Concentration	A or P	Code
G0D140560	I6-PC-1-0.0**	1,2,3,4,6,7,8-HpCDD	0.40U pg/g	A	bf

Tronox Northgate Henderson

VALIDATION COMPLETENESS WORKSHEET

Stage 2B/4

LDC #: 23105B21
 SDG #: GOD140560
 Laboratory: Test America

Date: 4/14/10
 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: 4/13/10
II.	HRGC/HRMS Instrument performance check	A	
III.	Initial calibration	A	
IV.	Routine calibration/ISV	A	
V.	Blanks	W	
VI.	Matrix spike/Matrix spike duplicates	SW	direct spiked
VII.	Laboratory control samples	A	LCs
VIII.	Regional quality assurance and quality control	N	
IX.	Internal standards	A	
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 = 1+2
XV.	Field blanks	W	ZB-PARCEL C-033110, FB-PARCEL C-033110

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

(GOD140559)

Validated Samples: ** Indicates sample underwent State 4 validation

1	I6-PC-1-0.0 **	5	11	0/06187MB	21		31
2	I6-PC-1-0.0FD	↓	12		22		32
3	I6-PC-1-0.0ALS		13		23		33
4	↓ ALS D		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: _____

LC #: 23105A-1
 SDG #: See cover

VALIDATION FINDINGS CHECKLIST

Page: 1 of 3
 Reviewer: 9
 2nd Reviewer: h

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 $\leq 30\%$ 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.			/	
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?	/			

LDC #: 23105B24
 SDG #: See EDU01

VALIDATION FINDINGS CHECKLIST

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

Validation Area	Yes	No	NA	Findings/Comments
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 type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	
Were the performance evaluation (PE) samples within the acceptance limits?	<input type="checkbox"/>	<input type="checkbox"/>	<input checked="" 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"/>	

LDC #: 23105 B2
SDG #: see cover

VALIDATION FINDINGS CHECKLIST

Page: 7 of 3
Reviewer: Q
2nd Reviewer: W

Validation Area	Yes	No	NA	Findings/Comments
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:

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: 4/16/10 Blank analysis date: 4/21/10 Associated samples: All (bl)
 Conc. units: pg/g

Compound	Blank ID	Sample Identification				
		5X	1	2		
A	010618ZMB	0.18				
F	0.14	0.7	0.40/U	-		
G	0.45	2.25	-	-		
H	0.088	0.44	0.12/U	0.14/U		
K	0.054	0.27				
O	0.076	0.38				
Q	0.15	0.75				

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)

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

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

Sampling date: 4/13/10

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

Compound	Blank ID	1	2	Sample Identification
	EB-PARCELF 033110	5X		
A	0.68	0.0034		
B	2.2	0.011		
C	1.9	0.0095		
D	7.7	0.0385		
E	4.5	0.0225		
F	100	0.5	0.40/U	
G	360	1.8	-	
H	2.6	0.013	-	
I	1.2	0.006		
J	1.0	0.005		
K	2.4	0.012		
L	2.6	0.013		
M	1.7	0.0085		
N	1.4	0.007		
O	24	0.12		
P	2.2	0.011		
Q	23	0.115		
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 #: 23105B21
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: 4/13/10

Field blank type: (circle one) Field Blank (bf) Rinsate / Other: Associated Samples: All (bf)

Compound	Blank ID	5X	1	2	Sample Identification
	FB-PARCEL 033110				
A	0.34	0.0017			
B	1.3	0.0065			
C	1.1	0.0055			
D	5.9	0.0295			
E	4.0	0.02			
F	91	0.455	0.40/U	-	
G	350	1.75	-	-	
H	2.2	0.011	-	-	
I	0.67	0.00335			
J	0.73	0.00365			
K	1.6	0.008			
L	1.7	0.0085			
M	0.99	0.00495			
O	20	0.1			
P	0.71	0.00355			
Q	18	0.09			
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

LDC #: 23105B24
 SDG #: See earlier

METHOD: HRGC/HRMS Polychlorinated Biphenyls (EPA Method 1668)

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

Were a MS/MSD analyzed every 20 samples of each matrix?
 Were the MS/MSD concentrations and the relative percent differences (RPD) within the QC limits stated below?

#	Date	MS/MSD ID	Compound	MS %R (Limits)	MSD %R (Limits)	RPD (Limits)	Associated Samples	Qualifications
		3/4	Σ	()	76 (80-113)	()	1	No Qual (MS in)
				()	()	()		
				()	()	()		
				()	()	()		
				()	()	()		
				()	()	()		
				()	()	()		
				()	()	()		
				()	()	()		
				()	()	()		
				()	()	()		
				()	()	()		
				()	()	()		
				()	()	()		
				()	()	()		
				()	()	()		
				()	()	()		
				()	()	()		
				()	()	()		
				()	()	()		
				()	()	()		
				()	()	()		
				()	()	()		
				()	()	()		
				()	()	()		
				()	()	()		
				()	()	()		
				()	()	()		
				()	()	()		

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
		A11	EMPC results (& flag)	M1	NK (K)
		A11	No confirmation in 2.3.7.8 - TCOF	M1	None (10) A

Comments: See sample calculation verification worksheet for recalculations

LDC#: 23105B21
SDG#: See Cover

VALIDATION FINDINGS WORKSHEET
Field Duplicates

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

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)
	1	2	RPD	Difference	Limits	
F	0.40	0.91		0.51	(≤5.2)	
G	2.6	13		10.4	(≤10)	<u>Not/A</u> (fd)
H	0.12	0.14		0.02	(<1.0)	

V:\FIELD DUPLICATES\23105B21.wpd

VALIDATION FINDINGS WORKSHEET
Initial Calibration Calculation Verification

SDG #: [Signature]

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

#	Standard ID	Calibration Date	Compound (Reference Internal Standard)	Reported		Recalculated		Reported		Recalculated	
				Average RRF (Initial)	Average RRF (Initial)	RRF (CS3 std)	RRF (CS3 std)	RRF (CS3 std)	RRF (CS3 std)	%RSD	%RSD
1	1CAR	3/4/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	0.98315	0.98315	1.01872	1.01872	4.65926	4.65926	4.65926	4.65926
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)	1.05705	1.05705	1.11325	1.11325	7.43940	7.43940	7.43940	7.43940
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)	1.08449	1.08449	1.09101	1.09101	5.63094	5.63094	5.63094	5.63094
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)	1.03068	1.03068	1.08562	1.08562	4.49467	4.49467	4.49467	4.49467
			OCDF (¹³ C-OCDD)	1.42582	1.42582	1.57902	1.57902	8.93881	8.93881	8.93881	8.93881
2	1CAR	4/12/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	0.945	0.945	0.98	0.98	4.44	4.44	4.44	4.44
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)	1.02	1.02	1.04	1.04	3.03	3.03	3.03	3.03
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)	1.114	1.114	1.19	1.19	5.33	5.33	5.33	5.33
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)	1.07	1.07	1.11	1.11	3.60	3.60	3.60	3.60
			OCDF (¹³ C-OCDD)	1.445	1.445	1.51	1.51	5.85	5.85	5.85	5.85
3	1CAR		2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)								
			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

LDC #: 23105B21
 SDG #: see cont.

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

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	23105B21 -30	4/23/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	0.98315	0.88010	10.5	0.88010	10.5
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)	1.05705	1.07014	1.8	1.07014	1.8
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)	1.082449	1.11017	2.4	1.11017	2.4
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)	1.03068	1.02319	0.7	1.02319	0.7
			OCDF (¹³ C-OCDD)	1.42502	1.42460	0.1	1.42460	0.1
2	23105B21 -1	4/21/10	2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)	0.945	1.04	9.6	1.04	9.6
			2,3,7,8-TCDD (¹³ C-2,3,7,8-TCDD)	1.021	1.03	1.2	1.03	1.2
			1,2,3,6,7,8-HxCDD (¹³ C-1,2,3,6,7,8-HxCDD)	1.114	1.26	13.5	1.26	13.5
			1,2,3,4,6,7,8-HpCDD (¹³ C-1,2,4,6,7,8-HpCDD)	1.07	1.09	2.0	1.09	2.0
			OCDF (¹³ C-OCDD)	1.445	1.50	3.9	1.50	3.9
3			2,3,7,8-TCDF (¹³ C-2,3,7,8-TCDF)					
			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 = $1 MSR - MSDR$ $1 * 2 / (MSR + MSDR)$ MSR = Matrix spike percent recovery MSDR = Matrix spike duplicate percent recovery

MS/MSD samples: 3/4

Compound	Spike Added		Sample Concentration	Spiked Sample Concentration		Matrix Spike		Matrix Spike Duplicate		Reported RPD	Recalculated RPD
	MS	MSD		MS	MSD	Percent Recovery		Percent Recovery			
						Reported	Recalc	Reported	Recalc		
2,3,7,8-TCDD	19.361	20.00	NO	20.8	21.5	107	107	108	108	3	3
1,2,3,7,8-PeCDD	96.87	100.00		97.8	98.8	101	101	99	99	1	1
1,2,3,4,7,8-HxCDD	↓	↓		102.6	84.7	106	106	85	85	19	19
1,2,3,4,7,8,9-HpCDF	193.61	200.00		122.3	92.6	126	126	93	93	28	28
OCDF				251.2	178.0	130	130	89	89	34	34

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.

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 = $100 \cdot \frac{SSC}{SA}$ Where: SSC = Spiked sample concentration
 SA = Spike added
 RPD = $100 \cdot \frac{LCS - LCSD}{LCS + LCSD}$ LCS = Laboratory control sample percent recovery
 LCSD = Laboratory control sample duplicate percent recovery

LCS ID: 0106157

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	20.2	NA	101	101								
1,2,3,7,8-PeCDD	100	↓	103	↓	103	103								
1,2,3,4,7,8-HxCDD	↓	↓	96.0	↓	96	96								
1,2,3,4,7,8,9-HpCDF	200	↓	110	↓	110	110								
OCDF			204	↓	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 ^(b)	Ion ID	Elemental Composition	Analyte	Descriptor	Accurate Mass ^(b)	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 ₄ ³⁵ Cl ₄ O C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO C ₁₂ H ₃ ³⁵ Cl ₃ O C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ ClO C ₁₂ H ₄ ³⁵ Cl ₃ O C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₄ ³⁵ Cl ₃ O ₂ C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₃ ³⁵ Cl ₃ O ₂ C ₉ F ₁₃	TCDF TCDF TCDF (S) TCDF (S) TCDD TCDD TCDD (S) TCDD (S) HxCDFE 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 ₃ ³⁷ ClO C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ Cl ₂ O C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ Cl ₂ O C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ Cl ₂ O C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₉ F ₁₇	HpCDF HpCDF HpCDF (S) HpCDF HpCDD HpCDD HpCDD (S) HpCDD (S) HpCDD (S) NCDPE 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 C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ Cl ₂ O C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ Cl ₂ O C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₉ F ₁₃	PeCDF PeCDF PeCDF (S) PeCDF (S) PeCDD PeCDD PeCDD (S) PeCDD (S) HxCDFE 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 LOCK	C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ Cl ₂ O C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ Cl ₂ O C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ Cl ₂ O C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₄ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₃ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₉ F ₁₃	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 ₄ ³⁷ ClO C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ Cl ₂ O C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ ClO C ₁₂ H ₂ ³⁵ Cl ₃ ³⁷ ClO C ₁₂ H ₂ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₂ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₂ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₂ ³⁵ Cl ₃ ³⁷ ClO ₂ C ₁₂ H ₂ ³⁵ Cl ₄ ³⁷ ClO ₂ C ₉ F ₁₇	HxCDF HxCDF HxCDF (S) HxCDF (S) HxCDD HxCDD HxCDD (S) HxCDD (S) OCDPE 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

