

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 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 # 23104:

SDG # Fraction

280-2143-1 Polynuclear Aromatic Hydrocarbons, Arsenic 280-2306-1

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 Inorganic Data Review, October 2004

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

Sincerely,

Paut

Erlinda T. Rauto Operations Manager/Senior Chemist

lage 2BI4 I.DC: #23104 (Trono sige 2BI4 DATE DATE DATE DATE PAH As SDG# DATE DATE DATE DATE PAH As M S M S reter/Soid RECD DUE (8270C) (6020) S S M S S S S S S S S	Attachment 1 x LLC-Northgate, Henderson NV / Tronox Parcels)		s w s w s w s w s w s w s w s w s w s w																	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
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Bage 2B/4 Date Date	LDC #23104 (Tron	E PAH As (8270C)	W S W S W S	/10 2 2 2 2 2	/10 2 0	//10 0 1														4 3 2 2 0 0
	age 2B/4	SDG# REC'D DUI	ater/Soil	80-2143-1 05/05/10 05/26	:80-2306-1 05/05/10 05/26	<u>280-2306-1 05/05/10 05/26</u>														T/LR

Shaded cells indicate Stage 4 validation (all other cells are Stage 2B validation). These sample counts do not include MS/MSD, and DUPs

23104ST.wpd

Tronox Northgate Henderson Worksheet

EDD Area	Yes	No	NA	Findings/Comments
1. Completeness			1	
Is there an EDD for the associated Tronox validation report?	x			
II. EDD Qualifier Population	r			
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_LDC23104_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 #23104

Polynuclear Aromatic Hydrocarbons

Laboratory Data Consultants, Inc. Data Validation Report

Project/Site Name: Tronox LLC Facility, 2010 Parcels, Henderson, Nevada

Collection Date: April 6, 2010

LDC Report Date: May 17, 2010

Matrix: Soil/Water

Parameters: Polynuclear Aromatic Hydrocarbons

Validation Level: Stage 2B & 4

Laboratory: TestAmerica, Inc.

Sample Delivery Group (SDG): 280-2143-1

Sample Identification

Q3-PF-3-1-0.0** Q3-PF-3-1-0.0FD FB-PARCELS-032910 EB-PARCELS-032910 Q3-PF-3-1-0.0MS Q3-PF-3-1-0.0MSD

**Indicates sample underwent Stage 4 review

Introduction

This data review covers 4 soil samples and 2 water samples listed on the cover sheet including dilutions and reanalysis as applicable. The analyses were per EPA SW 846 Method 8270C for Polynuclear Aromatic Hydrocarbons.

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 XVI.

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. GC/MS Instrument Performance Check

Instrument performance was checked at 12 hour intervals.

All ion abundance requirements were met.

III. Initial Calibration

Initial calibration was performed using required standard concentrations.

Percent relative standard deviations (%RSD) were less than or equal to 30.0% for all compounds.

Average relative response factors (RRF) for all compounds were within method and validation criteria

IV. Continuing Calibration

Continuing calibration was performed at the required frequencies.

Percent differences (%D) between the initial calibration RRF and the continuing calibration RRF were within the method criteria of less than or equal to 20.0% for calibration check compounds (CCCs) and 25.0% for all other compounds.

The percent differences (%D) of the second source calibration standard were less than or equal to 25.0% for all compounds.

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

V. Blanks

Method blanks were reviewed for each matrix as applicable. No polynuclear aromatic hydrocarbon contaminants were found in the method blanks.

Sample EB-PARCELS-032910 was identified as an equipment blank. No polynuclear aromatic hydrocarbon contaminants were found in this blank.

Sample FB-PARCELS-032910 was identified as a field blank. No polynuclear aromatic hydrocarbon 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 some compounds, the MSD or LCS percent recoveries (%R) were 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. Internal Standards

All internal standard areas and retention times were within QC limits with the following exceptions:

Sample	Internal Standards	Area (Limits)	Compound	Flag	A or P
Q3-PF-3-1-0.0**	Perylene-d12	374011 (820545-3282178)	Benzo(b)fluoranthene Benzo(k)fluoranthene Benzo(a)pyrene Indeno(1,2,3-cd)pyrene Dibenz(a,h)anthracene Benzo(g,h,i)perylene	J (all detects) R (all non-detects)	A
Q3-PF-3-1-0.0FD	Perylene-d12	357940 (820545-3282178)	Benzo(b)fluoranthene Benzo(k)fluoranthene Benzo(a)pyrene Indeno(1,2,3-cd)pyrene Dibenz(a,h)anthracene Benzo(g,h,i)perylene	J (all detects) R (all non-detects)	A

XI. 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.

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 280-2143-1	All compounds reported below the PQL.	J (all detects)	A

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

XIII. Tentatively Identified Compounds (TICs)

Tentatively identified compounds were not reported by the laboratory.

XIV. 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.

XV. Overall Assessment

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

XVI. Field Duplicates

Samples Q3-PF-3-1-0.0** and Q3-PF-3-1-0.0FD were identified as field duplicates. No polynuclear aromatic hydrocarbons were detected in any of the samples with the following exceptions:

	Concentrat	ion (ug/Kg)	DDD	Difference		
Compound	Q3-PF-3-1-0.0**	Q3-PF-3-1-0.0FD	(Limits)	(Limits)	Flags	A or P
Phenanthrene	370U	18	-	352 (≤370)	-	-
Pyrene	15	34	-	19 (≤370)	-	-
Benzo(b)fluoranthene	370U	110	-	260 (≤370)	-	-
Chrysene	370U	29	-	341 (≤370)	-	-
Fluoranthene	370U	49	-	321 (≤370)	-	-

Tronox LLC Facility, 2010 Parcels, Henderson, Nevada Polynuclear Aromatic Hydrocarbons - Data Qualification Summary - SDG 280-2143-1

SDG	Sample	Compound	Flag	A or P	Reason (Code)	
280-2143-1	Q3-PF-3-1-0.0** Benzo(b)fluoranthene Q3-PF-3-1-0.0FD Benzo(k)fluoranthene Benzo(a)pyrene Indeno(1,2,3-cd)pyrene Dibenz(a,h)anthracene Benzo(g,h,i)perylene		J (all detects) R (all non-detects)	A	Internal standards (area) (i)	
280-2143-1	Q3-PF-3-1-0.0** Q3-PF-3-1-0.0FD FB-PARCELS-032910 EB-PARCELS-032910	All compounds reported below the PQL.	J (all detects)	A	Project Quantitation Limit (sp)	

Tronox LLC Facility, 2010 Parcels, Henderson, Nevada Polynuclear Aromatic Hydrocarbons - Laboratory Blank Data Qualification Summary - SDG 280-2143-1

No Sample Data Qualified in this SDG

Tronox LLC Facility, 2010 Parcels, Henderson, Nevada Polynuclear Aromatic Hydrocarbons - Equipment Blank Data Qualification Summary - SDG 280-2143-1

No Sample Data Qualified in this SDG

Tronox LLC Facility, 2010 Parcels, Henderson, Nevada Polynuclear Aromatic Hydrocarbons - Field Blank Data Qualification Summary - SDG 280-2143-1

No Sample Data Qualified in this SDG

Tronox Northgate Henderson

VALIDATION COMPLETENESS WORKSHEET

Stage 2B/4

SDG #: 280-2143-1 Laboratory: Test America

LDC #: 23104A2

PAH

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

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
1.	Technical holding times	A	Sampling dates: 4 /6 6 /10
11.	GC/MS Instrument performance check	A	
111.	Initial calibration	A	?, RSD
IV.	Continuing calibration/ICV	A	COV /101 = 257
V.	Blanks	A	
VI.	Surrogate spikes	SW	
VII.	Matrix spike/Matrix spike duplicates	SW	
VIII.	Laboratory control samples	A	us
IX.	Regional Quality Assurance and Quality Control	N	
Х.	Internal standards	W2	
XI.	Target compound identification	A	Not reviewed for Stage 2B validation.
XII.	Compound quantitation/CRQLs	Α	Not reviewed for Stage 2B validation.
XIII.	Tentatively identified compounds (TICs)	N	Not reviewed for Stage 2B validation.
XIV.	System performance	A	Not reviewed for Stage 2B validation.
XV.	Overall assessment of data	A	
XVI.	Field duplicates	SW	$p = 1, \gamma$
XVII.	Field blanks	1,7D	FB = 3 EB = 4

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	Q3-PF-3-1-0.0**	S	11 1	MB280-102524/1-A	21	31
2	Q3-PF-3-1-0.0FD	Ţ	12)	MB280-10253/1-A	22	32
3	FB-PARCELS-032910	W	13		23	33
4,	EB-PARCELS-032910		14		24	34
5	Q3-PE-3-1-0 0MS	Ś	15		25	35
6	03-PE-3-1-0 0MSD	Ť	16		26	36
7		<u>v</u>	17		27	37
			18		28	38
6	······		10		29	39
10		,	20		30	40

Date: 5/13 /6 Page: \of] Reviewer: 5V42nd Reviewer: 5V4

Method: Semivolatiles (EPA SW 846 Method 8270C)

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				
Were the DFTPP performance results reviewed and found to be within the specified criteria?	<	-		
Were all samples analyzed within the 12 hour clock criteria?		-		
III. Initial calibration				
Did the laboratory perform a 5 point calibration prior to sample analysis?				
Were all percent relative standard deviations (%RSD) and relative response factors (RRF) within method criteria for all CCCs and SPCCs?	-			
Was a curve fit used for evaluation?				
Did the initial calibration meet the curve fit acceptance criteria of \geq 0.990?	[
Were all percent relative standard deviations (%RSD) \leq 30% and relative response factors (RRF) \geq 0.05?		-		
Was a continuing calibration standard analyzed at least once over 12 hours for		L		
each instrument?	<			
Were all percent differences (%D) and relative response factors (RRF) within method criteria for all CCCs and SPCCs?	/			
Were all percent differences (%D) \leq 25% and relative response factors (RRF) \geq 0.05?		· ·		
V. Blanks				and the second se
Was a method blank associated with every sample in this SDG?	<			
Was a method blank analyzed for each matrix and concentration?	/			
Was there contamination in the method blanks? If yes, please see the Blanks validation completeness worksheet.			-	
VI. Surrogate spikes				
Were all surrogate %R within QC limits?	+417			
If 2 or more base neutral or acid surrogates were outside QC limits, was a reanalysis performed to confirm %R?			M	r
If any %R was less than 10 percent, was a reanalysis performed to confirm %R?				
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.	/			
Was a MS/MSD analyzed every 20 samples of each matrix?	\leq			
Were the MS/MSD percent recoveries (%R) and the relative percent differences (RPD) within the QC limits?			-	
Vill Laboratory control samples				an a
Was an LCS analyzed for this SDG?				

Validation Area	Yes	No	NA	Findings/Comments
Was an LCS analyzed per extraction batch?	<			
Were the LCS percent recoveries (%R) and relative percent difference (RPD) within the OC limits?	<			
IX. Regional Quality Assurance and Quality Control			Ž.	
Were performance evaluation (PE) samples performed?		/		/
Were the performance evaluation (PE) samples within the acceptance limits?				
X. Internal standards				
Were internal standard area counts within -50% or +100% of the associated calibration standard?	JAR .	/		
Were retention times within ± 30 seconds from the associated calibration standard?		-		
XI. Target compound identification				
Were relative retention times (RRT's) within ± 0.06 RRT units of the standard?	\swarrow			
Did compound spectra meet specified EPA "Functional Guidelines" criteria?	1			
Were chromatogram peaks verified and accounted for?		ſ		
XII. Compound quantitation/CRQLs				
Were the correct internal standard (IS), quantitation ion and relative response factor (RRF) used to quantitate the compound?	1			
Were compound quantitation and CRQLs adjusted to reflect all sample dilutions and dry weight factors applicable to level IV validation?	/	1		
XIII. Tentatively identified compounds (TICs)				
Were the major ions (> 10 percent relative intensity) in the reference spectrum evaluated in sample spectrum?			/	
Were relative intensities of the major ions within \pm 20% between the sample and the reference spectra?			/	-
Did the raw data indicate that the laboratory performed a library search for all required peaks in the chromatograms (samples and blanks)?			-	
XIV. System performance				
System performance was found to be acceptable		-		
XV Overall assessment or data				
Overall assessment of data was found to be acceptable.		ſ		
XVI /Field ouplicates				
Field duplicate pairs were identified in this SDG.	\langle			
Target compounds were detected in the field duplicates.	/			
XVII. Field blanks				
Field blanks were identified in this SDG.	F			
Target compounds were detected in the field blanks.				

VALIDATION FINDINGS WORKSHEET

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

A. Phenol**	P. Bis(2-chloroethoxy)methane	EE. 2,6-Dinitrotoluene	TT. Pentachlorophenol**	III. Benzo(a)pyrene**
B. Bis (2-chloroethyl) ether	Q. 2,4-Dichlorophenol**	FF. 3-Nitroaniline	UU. Phenanthrene	JJJ. Indeno(1,2,3-cd)pyrene
C. 2-Chlorophenol	R. 1,2,4-Trichlorobenzene	GG. Acenaphthene**	VV. Anthracene	KKK. Dibenz(a,h)anthracene
D. 1,3-Dichlorobenzene	S. Naphthalene	HH. 2,4-Dinitrophenol*	WW. Carbazole	LLL. Benzo(g,h,i)perylene
E. 1,4-Dichlorobenzene**	T. 4-Chloroaniline	II. 4-Nitrophenol*	XX. Di-n-butyiphthalate	MMM. Bis(2-Chloroisopropyl)ether
F. 1,2-Dichlorobenzene	U. Hexachlorobutadiene ⁺ *	JJ. Dibenzofuran	YY. Fluoranthene⁺⁺	NNN. Aniline
G. 2-Methyiphenol	V. 4-Chloro-3-methylphenol**	KK. 2,4-Dinitrotoluene	ZZ. Pyrene	000. N-Nitrosodimethylamine
H. 2,2'-Oxybis(1-chloropropane)	W. 2-Methylnaphthalene	LL. Diethylphthalate	AAA. Butylbenzylphthalate	PPP. Benzoic Acid
I. 4-Methylphenol	X. Hexachlorocyclopentadiene*	MM. 4-Chlorophenyl-phenyl ether	BBB. 3,3'-Dichlorobenzidine	QQQ. Benzyl alcohol
J. N-Nitroso-di-n-propylamine*	Y. 2,4,6-Trichlorophenol**	NN. Fluorene	CCC. Benzo(a)anthracene	RRR. Pyridine
K. Hexachloroethane	Z. 2,4,5-Trichlorophenol	00. 4-Nitroaniline	DDD. Chrysene	SSS. Benzidine
L. Nitrobenzene	AA. 2-Chioronaphthalene	PP. 4,6-Dinitro-2-methylphenol	EEE. Bis(2-ethylhexyl)phthalate	TTT.
M. Isophorone	BB. 2-Nitroaniline	QQ. N-Nitrosodiphenylamine (1)**	FFF. Di-n-octylphthalate**	nnn
N. 2-Nitrophenol**	CC. Dimethylphthalate	RR. 4-Bromophenyl-phenylether	GGG. Benzo(b)fluoranthene	ww.
O. 2,4-Dimethylphenol	DD. Acenaphthylene	SS. Hexachlorobenzene	HHH. Benzo(k)fluoranthene	www.

Notes:* = System performance check compound (SPCC) for RRF; ** = Calibration check compound (CCC) for %RSD.

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VALIDATION FINDINGS WORKSHEET

METHOD: GC/MS BNA (EPA SW 846 Method 8270C) Please see qualification below for all questions answered "N". Not applicable questions are identified as "N/A"



F

Y(N NA If 2 or more base neutral or acid s Y N NA If any %R was less than 10 percei	surrogates were outside QC limits, were arready sis performed to co	as a reanalysis performed to confirm %	R?
# Date Sample ID	Surrogate	%R (Limits)	
2	NBZ	42 (52-120)	Qualifications
			June (my Int)
)	
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* QC limits are advisory QC Limits (Soil) QC Limits S1 (NBZ) = Nitrobenzene-d5 23-120 35-114 S2 (FBP) = 2-Fluorobiphenyl 30-115 43-116 S3 (TPH) = Terphenyl-d14 18-137 33-141 S4 (PHL) = Phenol-d5 24-113 10-94	<u>s (Water)</u> S5 (2FP)= 2-Fluoroj S6 (TBP) = 2,4,6-Tr S7 (2CP) = 2-Chlorc S8 (DCB) = 1,2-Dicf	phenol <u>QC Limits (Soil)</u> bromophenol 25-121 phenol-44 19-122 hlorobenzene-d4 20-130*	<u>QC Limits (Water)</u> 21-100 10-123 33-110*

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20-130*

16-110*

VALIDATION FINDINGS WORKSHEET Matrix Spike/Matrix Spike Duplicates

NC ζ lof Page: 2nd Reviewer: Reviewer:_

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

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A". <u>YN N/A</u> Were a matrix spike (MS) and matrix spike duplicate (MSD) analyzed for each matrix in this SDG? If no, indicate which matrix does not have an associated MS/MSD. Soil / Water. Was a MS/MSD analyzed every 20 samples of each matrix?

G NINA

n			\			_	_											_
	Qualifications	No mal (lisin				(medin)												
DC limits?	Associated Samples																	
ces (RPD) within the C	RPD (Limits)	()	()	()	()	()	1	()	()	(()	()	(()	()	()	()	(
ative percent different	MSD %R (Limits)	447 (52-120)	363 54-120)	191 (50-120)	289 (55-120)	()	()	()	()	()	()	()	()	()	()	()	()	()
ries (%R) and the rel	MS %R (Limits)	347 (52-120)	320 (54-120)	157 (52-120)	256 (55-120)	(oc1-h2) 94	()	()	()	()	()	()	()	()	()	()	()	1 1
jercent recove	Compound	606	444	111	KKK	ちっち							-					
Were the MS/MSD	DI DSW/SW	$\frac{2}{b}$																
N/N/A	t Date																	
X	#																	

	Compound	QC Limits (Soil)	RPD (Soil)	QC Limits (Water)	RPD (Water)		Compound	QC Limits (Soil)	RPD (Soil)	QC Limits (Water)	RPD (Water)
	Phenol	26-90%	≤ 35%	12-110%	< 42%	GG	Acenaphthene	31-137%	< 19%	46-118%	< 31%
v	2-Chlorophenol	25-102%	< 50%	27-123%	< 40%	Ξ.	4-Nitrophenol	11-114%	< 50%	10-80%	< 50%
ш	1,4-Dichlorobenzene	28-104%	< 27%	36-97%	< 28%	KK.	2,4-Dinitrotoluene	28-89%	< 47%	24-96%	< 38%
	N-Nitroso-di-n-propylamine	41-126%	< 38%	41-116%	< 38%	Ŀ.	Pentachlorophenol	17-109%	< 47%	9-103%	< 50%
ď	1,2,4-Trichlorobenzene	38-107%	< 23%	39-98%	< 28%	Ż	Pyrene	35-142%	< 36%	26-127%	< 31%
>	4-Chloro-3-methylphenol	26-103%	< 33%	23-97%	< 42%						

424	245
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LDC #:	SDG #:

VALIDATION FINDINGS WORKSHEET Internal Standards

ろ Page: 1 of 1 Reviewer: 2nd Reviewer:

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

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A". <u>Y W/N</u> Were all internal standard area counts within -50 to +100 of the associated calibration standard?

Date	e internal standard	ds within +/- 30 seconds of the reten Area (Limits) 374 oll (820545-3 357940 292305 292305 292305	tion times of the associated calibratio	n standard? J/R/A (1) JJJ, KK, LLL No quel (BC)
		Internal standar Internal Standard	internal standards within +/- 30 seconds of the retering internal standard Area (Limits) PRY 374 oII (820 545 - 3 PRY 3974 fo 292,305 PRY 298,157	internal standards within +/- 30 seconds of the retention times of the associated calibratic Internal Area (Limits) REY 374 oll (8 20 545 - 3 282 178) PRY 392 395 - 3 282 178) 3579 40 - 372 - 328 - 32

* QC limits are advisory IS1 (DCB) = 1,4-Dichlorobenzene-d4 IS2 (NPT) = Naphthalene-d8 IS3 (ANT) = Acenaphthene-d10

IS4 (PHN) = Phenanthrene-d10 IS5 (CRY) = Chrysene-d12 IS6 (PRY) = Perylene-d12

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LDC#: 23104A2b SDG#:See cover

VALIDATION FINDINGS WORKSHEET Field Duplicates

1 of _1 Page: 516 **Reviewer:** 2nd Reviewer:

METHOD: GC/MS PAH (EPA SW 846 Method 8270C) YN NA

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

O and have	Conc (ug/Kg)	BBD	Diff	Diff Limits	Quals
Compound Name	1	2	KFD (≤50%)	Din	Diff Linits	(Parent Only)
Phenanthrene	370U	18		352	≤370	
Pyrene	15	34		19	≤370	
Benzo(b)fluoranthene	370U	110		260	≤370	
Chrysene	370U	29		341	≲370	
Fluoranthene	370U	49		321	≤370	

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LDC #: 73104 A3b SDG #: 54 CV

VALIDATION FINDINGS WORKSHEET Initial Calibration Calculation Verification

Page: 1 of 1 Reviewer: 2nd Reviewer:

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

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,

 $\label{eq:A_s} A_{s} = \mbox{Area of associated internal standard} \\ C_{s} = \mbox{Concentration of internal standard} \\ X = \mbox{Mean of the RRFs} \\ \end{array}$

				Reported	Recalculated	Reported	Recalculated	Reported	Recalculated
		Calibration		RRF	RRF	Average RRF	Average RRF	%RSD	%RSD
#	Standard ID	Date	Compound (Internal Standard)	(50 std)	(50 std)	(Initial)	(Initial)		
-	ICAL	3/18/10	Naphthalene (IS2)	1.1221	1.1221	1.1134	1.1134	3.0	3.0
	MSSD		Fluorene (IS3)	1.3445	1.3445	1.3225	1.3225	5.0	5.0
			Phenanthrene (IS4)	1.1670	1.1670	1.1292	1.1292	6.7	6.7
			Chrysene (IS5)	1.0697	1.0697	1.0065	1.0065	8.2	8.2
			Benzo(a)pyrene (IS6)	1.1044	1.1044	1.0885	1.0885	6.7	6.7

c IS/Cpd	Area cpd	Area IS
10/50	1966829	1402196
10/50	1609531	957696
40/50	2338313	1602894
40/50	2492854	1864283
40/50	2472315	1790944

Benzo(a)py	0.9762	1.0038	1.0453	1.1044	1.1019	1.1505	1.1705	1.1554	1.0885	0.0728	
Chrysene	1.0249	1.0017	1.0484	1.0697	1.0741	1.0559	0.9457	0.8319	1.0065	0.0822	
Phenanth	1.2145	1.1130	1.1499	1.1670	1.1699	1.1804	1.0504	0.9882	1.1292	0.0754	
Fluorene	1.2049	1.2482	1.2972	1.3445	1.3440	1.3805	1.3878	1.3725	1.3225	0.0666	
Naphthalene	1.1382	1.1444	1.1266	1.1221	1.0774	1.1406	1.1068	1.0507	1.1134	0.0335	
Conc	4.00	10.00	20.00	50.00	80.00	120.00	160.00	200.00	×	s =	-

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.

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LDC # 73104 436 SDG # <u>See Cover</u>

VALIDATION FINDINGS WORSHEET

Continuing Calibration Results Verification

× 2 lof Page___ Reviewer:__ 2nd Reviewer:

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

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:

	Where:
% Difference = 100 * (ave. RRF - RRF)/ave. RRF	ave. RRF = initial calibration average RRF
RRF = (Ax)(Cis)/(Ais)(Cx)	RRF = continuing calibration RRF
	Ax = Area of compound

Ais = Area of associated internal standard Cis = Concentration of internal standard Cx = Concentration of compound

		Calibration		Average RRF	Reported	Recalculated	Reported	Recalculated
#	Standard ID	Date	Compound (Ref IS)	(Initial RRF)	(CC RRF)	(CC RRF)	۵%	%D
1	D3873	4/12/10	Naphthalene (IS2)	1.113	1.176	1.176	5.6	5.6
			Fluorene (IS3)	1.323	1.380	1.380	4.3	4.3
			Phenanthrene (iS4)	1.129	1.184	1.184	4.8	4.8
			Chrysene (IS5)	1.007	1.089	1.089	8.2	8.2
			Benzo(a)pyrene (IS6)	1.089	1.178	1.178	8.2	8.2
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			-					

	ccV1		CCV2		CCV3	
Compound	Area Cpd	Area IS	Area Cpd	Area IS	Area Cpd	Area IS
Naphthalene	3148673	1338693				
Fluorene	2644619	958332				
Phenanthrene	3836885	1620877				
Chrysene	4002705	1837231				
Benzo(a)pyrene	3866113	1641089				

LDC #: ~ 3 10 4 A 7 b SDG #: Sre Cover

VALIDATION FINDINGS WORKSHEET Surrogate Results Verification

Page:<u>lof</u> Reviewer:<u>M</u> 2nd reviewer:<u></u>

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

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

	Surrogate Spiked	Surrogate Found	Percent Recovery Reported	Percent Recovery Recalculated	Percent Difference
Nitrobenzene-d5	100	52.6	53	53	0
2-Fluorobiphenyl		64.6	65	65	1
Terphenyl-d14	ł	89.0	89	89	
Phenol-d5					
2-Fluorophenol					
2,4,6-Tribromophenol					
2-Chlorophenol-d4					
1,2-Dichlorobenzene-d4					

Sample ID:

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

Sample ID:

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

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Page: <u>lof 1</u> Reviewer: <u>We</u> 2nd Reviewer: <u>I</u>

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

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

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

Where: SSC = Spiked sample concentration SA = Spike added

MSC = Matrix spike concentration

SC = Sample concentation

MSDC = Matrix spike duplicate concentration

MS/MSD samples:

RPD = I MSC - MSC I * 2/(MSC + MSDC)

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Compound (US) MS Phenol	dded	Concentration								
Ahenol		(MS/K)	עסחניוו (אל /		Percent R	есочегу	Percent R	scovery	RPD	
Phenol	U MSD	0	MS	MSD	Reported	Recalc	Reported	Recalc	Reported	Recalculated
N-Nitroso-di-n-propylamine										
4-Chloro-3-methylphenol										
Acenaphthene 2000	3010	0	1950	2260	ود	ر ک	22	75	15	2
Pentachlorophenol										
Pyrene 3000	30 10	1 2	2680 3	010	89	51	66	99	Σ	2
							~	_		

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.

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	Lab
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boratory Control Sample/Laboratory Control Sample Duplicates Results Verification VALIDATION FINDINGS WORKSHEET

Page: lof 1 Reviewer: <u>WC</u> 2nd Reviewer: 1

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

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

% Recovery = 100 * (SC/SA

Where: SSC = Spike concentration SA = Spike added

RPD = I LCSC - LCSDC I * 2/(LCSC + LCSDC)

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

LCS/LCSD samples: 10 2 8 6- 10 274 /2-A

	ŝ	ika	ß	ike	01	S	Ü T	g		csn
Compound	Ad (کرم	ded Ac	Concet (Idx	ntration /(⊂)	Percent F	Recovery	Percent R	ecovery	R	0
	I CS	ار المحمد 1 CSD	1 CS	ار ۱ CSD	Reported	Recalc	Reported	Recalc	Reported	Recalculated
Phenol										
N-Nitroso-di-n-propylamine										
4-Chloro-3-methylphenol										
Acenaphthene	2650	A N	1970	44	74	74				
Pentachiorophenol										
Pyrene	26 50		2600	~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~	98	86				

Comments: Refer to Laboratory Control Sample/Laboratory Control Sample Duplicates findings worksheet for list of qualifications and associated samples when reported results do not agree within 10.0% of the recalculated results. LDC #: 2304 A 26 SDG #: Sre Cover

VALIDATION FINDINGS WORKSHEET **Sample Calculation Verification**



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

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?

Example:

Conce	ntratio	$n = (A_{,})(I_{s})(V_{i})(DF)(2.0) (A_{s})(RRF)(V_{s})(V_{i})(%S)$
A _x	=	Area of the characteristic ion (EICP) for the compound to be measured
A_{is}	=	Area of the characteristic ion (EICP) for the specific internal standard
l _s	=	Amount of internal standard added in nanograms (ng)
V _°	=	Volume or weight of sample extract in milliliters (ml) or grams (g).
V	=	Volume of extract injected in microliters (ul)
V _t	=	Volume of the concentrated extract in microliters (ul)
Df	=	Dilution Factor.
%S	=	Percent solids, applicable to soil and solid matrices only.

Sample I.D. _____, ____ И И $Conc. = \frac{(19937)(40)(1)(100)}{(357121)(1.1292)(36.2g)(0.93)(0)}$ = 18,5 ug/kg

Factor of 2 to account for GPC cleanup 2.0 =

			Reported Concentration	Calculated Concentration	
#	Sample ID	Compound		()	Qualification
			· · · · · · · · · · · · · · · · · · ·		
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LDC Report# 23104B2b

Laboratory Data Consultants, Inc. Data Validation Report

Project/Site Name: Tronox LLC Facility, 2010 Parcels, Henderson, Nevada

Collection Date: April 8, 2010

LDC Report Date: May 17, 2010

Matrix: Soil/Water

Parameters: Polynuclear Aromatic Hydrocarbons

Validation Level: Stage 2B & 4

Laboratory: TestAmerica, Inc.

Sample Delivery Group (SDG): 280-2306-1

Sample Identification

S3-PG-2-0.0** FB-PARCELS_032910 EB-04082010-PARCELG S3-PG-2-0.0MS S3-PG-2-0.0MSD

**Indicates sample underwent Stage 4 review

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Introduction

This data review covers 3 soil samples and 2 water samples listed on the cover sheet including dilutions and reanalysis as applicable. The analyses were per EPA SW 846 Method 8270C for Polynuclear Aromatic Hydrocarbons.

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 XVI.

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. GC/MS Instrument Performance Check

Instrument performance was checked at 12 hour intervals.

All ion abundance requirements were met.

III. Initial Calibration

Initial calibration was performed using required standard concentrations.

Percent relative standard deviations (%RSD) were less than or equal to 30.0% for all compounds.

Average relative response factors (RRF) for all compounds were within method and validation criteria

IV. Continuing Calibration

Continuing calibration was performed at the required frequencies.

Percent differences (%D) between the initial calibration RRF and the continuing calibration RRF were within the method criteria of less than or equal to 20.0% for calibration check compounds (CCCs) and 25.0% for all other compounds.

The percent differences (%D) of the second source calibration standard were less than or equal to 25.0% for all compounds.

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

V. Blanks

Method blanks were reviewed for each matrix as applicable. No polynuclear aromatic hydrocarbon contaminants were found in the method blanks.

Sample EB-04082010-PARCELG was identified as an equipment blank. No polynuclear aromatic hydrocarbon contaminants were found in this blank.

Sample FB-PARCELS_032910 was identified as a field blank. No polynuclear aromatic hydrocarbon 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. Percent recoveries (%R) and relative percent differences (RPD) were within QC limits.

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. Internal Standards

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

XI. 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.

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 280-2306-1	All compounds reported below the PQL.	J (all detects)	A

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

XIII. Tentatively Identified Compounds (TICs)

Tentatively identified compounds were not reported by the laboratory.

XIV. 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.

XV. Overall Assessment

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

XVI. Field Duplicates

No field duplicates were identified in this SDG.

Tronox LLC Facility, 2010 Parcels, Henderson, Nevada Polynuclear Aromatic Hydrocarbons - Data Qualification Summary - SDG 280-2306-1

SDG	Sample	Compound	Flag	A or P	Reason (Code)
280-2306-1	S3-PG-2-0.0** FB-PARCELS_032910 EB-04082010-PARCELG	All compounds reported below the PQL.	J (all detects)	A	Project Quantitation Limit (sp)

Tronox LLC Facility, 2010 Parcels, Henderson, Nevada Polynuclear Aromatic Hydrocarbons - Laboratory Blank Data Qualification Summary - SDG 280-2306-1

No Sample Data Qualified in this SDG

Tronox LLC Facility, 2010 Parcels, Henderson, Nevada Polynuclear Aromatic Hydrocarbons - Equipment Blank Data Qualification Summary - SDG 280-2306-1

No Sample Data Qualified in this SDG

Tronox LLC Facility, 2010 Parcels, Henderson, Nevada Polynuclear Aromatic Hydrocarbons - Field Blank Data Qualification Summary - SDG 280-2306-1

No Sample Data Qualified in this SDG

Tronox Northgate Henderson

VALIDATION COMPLETENESS WORKSHEET

Stage 2B/4

SDG #: 280-2306-1 Laboratory: Test America

LDC #:

23104B2a

Date: 5/13/10 Page: of Reviewer: NL 2nd Reviewer: $\wedge \land$

P★} METHOD: GC/MS Semivolatiles (EPA SW 846 Method 8270C)

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
	Technical holding times	A	Sampling dates: 4 /o g /10
	GC/MS Instrument performance check	A	
III.	Initial calibration	A	ZKSP
IV.	Continuing calibration/ICV	R	CW/W = 253
V.	Blanks	A	
VI.	Surrogate spikes	Α	
VII.	Matrix spike/Matrix spike duplicates	A	
VIII.	Laboratory control samples	A	us
IX.	Regional Quality Assurance and Quality Control	N	
Х.	Internal standards	A	
XI.	Target compound identification	A	Not reviewed for Stage 2B validation.
XII.	Compound quantitation/CRQLs	A	Not reviewed for Stage 2B validation.
XIII.	Tentatively identified compounds (TICs)	N	Not reviewed for Stage 2B validation.
XIV.	System performance	A	Not reviewed for Stage 2B validation.
XV.	Overall assessment of data	A	
XVI.	Field duplicates	N	
XVII.	Field blanks	ND	FB = 2 $EB = 3$

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: ** Indicates sample underwent State 4 validation

+ 1	G S3-P B -2-0.0**	S	11)	MB 280-10 851 /1-A	21	31
2 7	FB-PARCELS_032910	W	12	MB 280-10934/2-A	22	32
37	EB-04082010-PARCELG	Ţ	13		23	33
4	6 S3-P₽-2-0.0MS	5	14		24	34
5 1	S3-PB-2-0.0MSD	T	15		25	35
6			16		26	36
7			17		27	37
8			18		28	38
9			19		29	39
10			20		30	40

Method: Semivolatiles (EPA SW 846 Method 8270C)

Schedulation futures All technical hoding times were met. Image: Control temporature criteria was met. Control temporature criteria was met. Image: Control temporature criteria was met. Image: Control temporature criteria was met. Control temporature criteria was met. Image: Control temporature criteria was met. Image: Control temporature criteria was met. Control temporature criteria was met. Image: Control temporature criteria was met. Image: Control temporature criteria was met. Control temporature criteria was met. Image: Control temporature criteria was met. Image: Control temporature criteria was met. View all approximation temporature criteria	Validation Area	Yes	No	NA	Findings/Comments
All technical holding times were met. Coder temperature criteria was met. Coder temperature criteria w	I. Technical holding times				
Cooler temperature criteria was met.	All technical holding times were met.		r		
If (22A) S1 matrixely performance results reviewed and found to be within the specified	Cooler temperature criteria was met.		[
Were the DFTPP performance results reviewed and found to be within the specified	II. GC/MS Instrument performance check	i serie T			
Were all samples analyzed within the 12 hour clock criteria? Image: Control of Contro	Were the DFTPP performance results reviewed and found to be within the specified criteria?	/			
uit. Attait Selection Did the laboratory perform a 5 point calibration plor to sample analysis? Were all percent relative standard deviations (%RSD) and relative response factors (RRP) within method orient for all CCCs and SPCCs? Was a curve fit used for evaluation? Did the initial calibration meet the curve fit acceptance criteria of ≥ 0.990? Were all percent relative standard deviations (%RSD) ≤ 30% and relative response factors (RRF) ≥ 0.05? Was a continuing calibration standard analyzed at least once every 12 hours for each instrument? Was a continuing calibration standard analyzed at least once every 12 hours for each instrument? Were all percent differences (%D) ≤ 25% and relative response factors (RRF) ≥ 0.05? Ware all percent differences (%D) ≤ 25% and relative response factors (RRF) ≥ 0.05? Was a method blank associated with every sample in this SDG? Was a method blank analyzed for each matrix and concentration? Was a method blank analyzed for each matrix and concentration? Was a there contamination in the method blanks? If yes, please see the Blanks validation completeness worksheet. J. surpose pate J. surpose pate Vere all surrogate %R within QC limits? If 2 or more base neutral or acid surcogates were outside QC limits, was a reanalysis performed to confirm %R? If any %R was less than 10 percent, was a reanalysis performed to confirm	Were all samples analyzed within the 12 hour clock criteria?		-		
Did the laboratory perform a 5 point calibration prior to sample analysis? Were all percent relative standard deviations (%RSD) and relative response factors (RRP) within method criteria for all CCCs and SPCCs? Was a curve fit used for evaluation? Did the initial calibration meet the curve fit acceptance criteria of > 0.990? Were all percent relative standard deviations (%RSD) ≤ 30% and relative response factors (RRF) > 0.05? W to Continuing calibration standard analyzed at least once every 12 hours for each instrument? Was a continuing calibration standard analyzed at least once every 12 hours for each instrument? Were all percent differences (%D) and relative response factors (RRF) ≥ 0.05? W. Continuing calibration standard analyzed at least once every 12 hours for each instrument? Was a continuing calibration standard analyzed at least once every 12 hours for each instrument? Were all percent differences (%D) and relative response factors (RRF) ≥ 0.05? Was a method blank associated with every sample in this SDG? Was a method blank analyzed for each matrix and concentration? Was a there contamination in the method blanks? If yes, please see the Blanks validation completeness worksheed. Isomogate systemed to confirm %R? If 2 more base neutral or acid surrogates were outside QC limits, was a reanalysis performed to confirm %R? If any %R was kess than 10 percent, was a reanalysis performed to confirm %R? If any %R was kess than 10 percent, was a reanalysis performed to confirm %R? If any %R was kess than 10 percent, was a reanalysis performed to confirm %R? Was a MSMSD analyzed every 20 samples of each matrix? Was a MMSD analyzed system for the elative percent differences (RPD) within the QC limits? If any %R was kess than 10 percent, was a reanalysis performed to confirm %R? If any %R was kess than 10 percent, was a reanalysis performed to confirm %R? If any %R was kess than 10 percent, was a reanalysis performed to confirm %R? If any %R was kess than 10 percent, was a reanalysis performed to confirm %R? If any	III. Initial calibration				
Were all percent relative standard deviations (%RSD) and relative response factors Image: Constraint of the c	Did the laboratory perform a 5 point calibration prior to sample analysis?	-/			
Was a curve fit used for evaluation?	Were all percent relative standard deviations (%RSD) and relative response factors (RRF) within method criteria for all CCCs and SPCCs?	/			
Did the initial calibration meet the curve fit acceptance criteria of ≥ 0.990? Were all percent relative standard deviations (%RSD) ≤ 30% and relative response factors (RRF) ≥ 0.05? Was a continuing calibration Was a continuing calibration standard analyzed at least once every 12 hours for each instrument? Were all percent differences (%D) and relative response factors (RRF) within method criteria for all CCCs and SPCCs? Were all percent differences (%D) ≤ 25% and relative response factors (RRF) ≥ 0.05? V Barks Was a method blank associated with every sample in this SDG? Was a method blank associated with every sample in this SDG? Was a method blank analyzed for each matrix and concentration? Was there contamination in the method blanks? If yes, please see the Blanks validation completeness worksheet. M. Surrogate spite Were all surcogate %R within QC limits? If any %R was less than 10 percent, was a reanalysis performed to confirm %R? Vil Metho splow/etrip spike (MS) and matrix spike duplicate (MSD) analyzed for each matrix in this SDG? (In this spike duplicate MSD) analyzed for each matrix? Were a matrix spike (MS) and matrix spike duplicate (MSD) analyzed for each matrix? Were a matrix spike (MS) and matrix spike duplicate (MSD) analyzed for each matrix? Were the MS/MSD percent recoveries (%R) and the relative percent differences (RPD) within the QC limits?	Was a curve fit used for evaluation?			[
Were all percent relative standard deviations (%RSD) ≤ 30% and relative response factors (RRF) ≥ 0.05? Image: the imag	Did the initial calibration meet the curve fit acceptance criteria of \geq 0.990?			\leq	
W. Continuing calibration Was a continuing calibration standard analyzed at least once every 12 hours for each instrument? Were all percent differences (%D) and relative response factors (RRF) within method oriteria for all CCCs and SPCCs? Were all percent differences (%D) ≤ 25% and relative response factors (RRF) ≥ 0.05? V. Blanks Was a method blank associated with every sample in this SDG? Was a method blank associated with every sample in this SDG? Was a method blank analyzed for each matrix and concentration? Was a method blank analyzed for each matrix and concentration? Was there contamination in the method blanks? If yes, please see the Blanks validation completeness worksheet. VI. Sortogate spikes Were all surrogate %R within QC limits? If 2 or more base neutral or acid surrogates were outside QC limits, was a reanalysis performed to confirm %R? VI. Metrod standard trains pike duplicate (MSD) analyzed for each matrix n this SDG? Were at matrix spike (MS) and matrix does not have an associated MS/MSD. Soil / Water. Was a MS/MSD percent recoveries (%R) and the relative percent differences (%R) within C it limits? Were the MS/MSD parcent recoveries (%R) and the relative percent differences (RPD) within the QC limits? Were the MS/MSD parcent recoveries (%R) and the relative percent differences (RPD) within the QC limits? Was a a MS/MSD percent recoveries (%R) and the relative	Were all percent relative standard deviations (%RSD) \leq 30% and relative response factors (RRF) > 0.05?	6			
Was a continuing calibration standard analyzed at least once every 12 hours for	IV. Continuing calibration			1./ S.	
Were all percent differences (%D) and relative response factors (RRF) within	Was a continuing calibration standard analyzed at least once every 12 hours for each instrument?	/			
Were all percent differences (%D) ≤ 25% and relative response factors (RRF) ≥ 0.05? Y. Blanks Was a method blank associated with every sample in this SDG? Was a method blank analyzed for each matrix and concentration? Image: Constraint of the method blanks? If yes, please see the Blanks Was there contamination in the method blanks? If yes, please see the Blanks validation completeness worksheet. Image: Constraint of the method blanks? If yes, please see the Blanks YI. Sorrogate sples Image: Constraint of the method blanks? If yes, please see the Blanks validation completeness worksheet. Image: Constraint of the method blanks? If yes, please see the Blanks validation completeness worksheet. YI. Sorrogate sples Image: Constraint of the method blanks? If yes, please see the Blanks validation completeness worksheet. Image: Constraint of the method blanks? If yes, please see the Blanks validation completeness worksheet. YI. Sorrogate sples Image: Constraint of the method blanks? If yes, please see the Constraint of the confirm %R? Image: Constraint of the confirm %R? If any %R was less than 10 percent, was a reanalysis performed to confirm %R? Image: Constraint of the confirm %R? Image: Constraint of the confirm %R? VII. Mathy: splka/Matrix splke duplicate (MSD) analyzed for each matrix in this SDG? If no, indicate which matrix does not have an associated MS/MSD. Soil / Water. Image: Constraint of the confirm %R? Image: Constraint of the confirm %R? Was a MS/MSD analyzed every 2	Were all percent differences (%D) and relative response factors (RRF) within method criteria for all CCCs and SPCCs?	/			
V. Blanks Was a method blank associated with every sample in this SDG? Was a method blank analyzed for each matrix and concentration? Was there contamination in the method blanks? If yes, please see the Blanks validation completeness worksheet. VI. Storogate spikes Were all surrogate %R within QC limits? If 2 or more base neutral or acid surrogates were outside QC limits, was a reanalysis performed to confirm %R? VI. Matrix spike duplicates Vill 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 percent recoveries (%R) and the relative percent differences Was a MS/MSD paralyzed every 20 samples of each matrix? Was a MS/MSD percent recoveries (%R) and the relative percent differences (RPD) within the QC limits? VIII Leboratory control samples	Were all percent differences (%D) \leq 25% and relative response factors (RRF) \geq 0.05?				
Was a method blank associated with every sample in this SDG? / Was a method blank analyzed for each matrix and concentration? ////////////////////////////////////	V. Blanks				
Was a method blank analyzed for each matrix and concentration? Was there contamination in the method blanks? If yes, please see the Blanks validation completeness worksheet. VI. Surogate spikes Were all surrogate %R within QC limits? If 2 or more base neutral or acid surrogates were outside QC limits, was a reanalysis performed to confirm %R? If any %R was less than 10 percent, was a reanalysis performed to confirm %R? VII. Matrix spike (MS) and matrix spike duplicate (MSD) analyzed for each matrix in this SDG? If no, indicate which matrix does not have an associated MS/MSD. Soil / Water. Was a MS/MSD analyzed every 20 samples of each matrix? Were the MS/MSD percent recoveries (%R) and the relative percent differences (RPD) within the QC limits? VIII. Laboratory control samples	Was a method blank associated with every sample in this SDG?				
Was there contamination in the method blanks? If yes, please see the Blanks vil. Surrogate spikes Were all surrogate %R within QC limits? If 2 or more base neutral or acid surrogates were outside QC limits, was a reanalysis performed to confirm %R? If any %R was less than 10 percent, was a reanalysis performed to confirm %R? vil. Matrix spike (MSI) and matrix spike duplicate: Were a matrix spike (MS) and matrix spike duplicate (MSD) analyzed for each matrix in this SDG? If no, indicate which matrix does not have an associated MS/MSD. Soil / Water. Was a MS/MSD analyzed every 20 samples of each matrix? Were the MS/MSD percent recoveries (%R) and the relative percent differences (RPD) within the QC limits? Vill Laboratory control samples	Was a method blank analyzed for each matrix and concentration?				
VI. Surrogate spikes Were all surrogate %R within QC limits? If 2 or more base neutral or acid surrogates were outside QC limits, was a reanalysis performed to confirm %R? If any %R was less than 10 percent, was a reanalysis performed to confirm %R? 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. 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? VIII. Leboratory control samples	Was there contamination in the method blanks? If yes, please see the Blanks validation completeness worksheet.				
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If 2 or more base neutral or acid surrogates were outside QC limits, was a reanalysis performed to confirm %R? If any %R was less than 10 percent, was a reanalysis performed to confirm %R? 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. 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? VII. Leboratory control samples	Were all surrogate %R within QC limits?				
If any %R was less than 10 percent, was a reanalysis performed to confirm %R? 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. 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? VIII. Laboratory control samples	If 2 or more base neutral or acid surrogates were outside QC limits, was a reanalysis performed to confirm %R?			_	
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. 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? VIII. Laboratory control samples	If any %R was less than 10 percent, was a reanalysis performed to confirm %R?				
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. Ms/MSD. Soil / Water. Was a MS/MSD analyzed every 20 samples of each matrix? Were the MS/MSD percent recoveries (%R) and the relative percent differences (RPD) within the QC limits? VIII. Laboratory control samples Matrix in this SDC2	VII. Matrix spike/Matrix spike duplicates				
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? VIII. Leboratory control samples	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? VIII. Leboratory control samples	Was a MS/MSD analyzed every 20 samples of each matrix?	1			
VIII. Leboratory control samples	Were the MS/MSD percent recoveries (%R) and the relative percent differences (RPD) within the QC limits?				
Was as LCS analyzed for this SDC2	VIII. Leboratory control samples				
	Was an LCS analyzed for this SDG?	Ļ			······································

A.5.,

Validation Area	Yes	No	NA	Findings/Comments
Was an LCS analyzed per extraction batch?	\square			
Were the LCS percent recoveries (%R) and relative percent difference (RPD) within the QC limits?				
IX. Regional Quality Assurance and Quality Control				
Were performance evaluation (PE) samples performed?				۷
Were the performance evaluation (PE) samples within the acceptance limits?				
X. Internal standards			** A(\$2) 1	
Were internal standard area counts within -50% or +100% of the associated calibration standard?				
Were retention times within ± 30 seconds from the associated calibration standard?				
XI. Target compound identification		· · · ·	1	
Were relative retention times (RRTs) within ± 0.06 RRT units of the standard?	[ļ		
Did compound spectra meet specified EPA "Functional Guidelines" criteria?	/			
Were chromatogram peaks verified and accounted for?		<u> </u>		
XII. Compound quantitation/CRQLs	1	1	le se è T	
Were the correct internal standard (IS), quantitation ion and relative response factor (RRF) used to quantitate the compound?				
Were compound quantitation and CRQLs adjusted to reflect all sample dilutions and dry weight factors applicable to level IV validation?				
XIII. Tentatively identified compounds (TICs)	-		en en	
Were the major ions (> 10 percent relative intensity) in the reference spectrum evaluated in sample spectrum?			/	
Were relative intensities of the major ions within \pm 20% between the sample and the reference spectra?				/
Did the raw data indicate that the laboratory performed a library search for all required peaks in the chromatograms (samples and blanks)?		/	ł	
XIV. System performance				
System performance was found to be acceptable.		ľ		
XV Overall assessment of data and the second s				
Overall assessment of data was found to be acceptable.				
XVI: Field duplicates	(₁ .			
Field duplicate pairs were identified in this SDG.	-		Z	
Target compounds were detected in the field duplicates.				
XVII. Field blanks				
Field blanks were identified in this SDG.	/	1		
Target compounds were detected in the field blanks.		1		

VALIDATION FINDINGS WORKSHEET

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

A. Phenol**	P. Bis(2-chloroethoxy)methane	EE. 2,6-Dinitrotoluene	TT. Pentachlorophenol⁺*	III. Benzo(a)pyrene**
B. Bis (2-chloroethyl) ether	Q. 2,4-Dichlorophenol**	FF. 3-Nitroaniline	UU. Phenanthrene	JJJ. Indeno(1,2,3-cd)pyrene
C. 2-Chlorophenoł	R. 1,2,4-Trichlorobenzene	GG. Acenaphthene**	VV. Anthracene	KKK. Dibenz(a,h)anthracene
D. 1,3-Dichlorobenzene	S. Naphthalene	HH. 2,4-Dinitrophenol*	WW. Carbazole	LLL. Benzo(g,h,i)perylene
E. 1,4-Dichlorobenzene**	T. 4-Chloroaniline	ll. 4-Nitrophenol*	XX. Di-n-butyiphthalate	MMM. Bis(2-Chloroisopropyl)ether
F. 1,2-Dichlorobenzene	U. Hexachlorobutadiene**	JJ. Dibenzofuran	YY. Fluoranthene**	NNN. Aniline
G. 2-Methylphenol	V. 4-Chloro-3-methylphenol**	KK. 2,4-Dinitrotoluene	ZZ. Pyrene	000. N-Nitrosodimethylamine
H. 2,2'-Oxybis(1-chloropropane)	W. 2-Methylnaphthalene	LL. Diethylphthalate	AAA. Butylbenzylphthalate	PPP. Benzoic Acid
I. 4-Methylphenof	X. Hexachlorocyclopentadiene*	MM. 4-Chlorophenyl-phenyl ether	BBB. 3,3'-Dichlorobenzidine	QQQ. Benzyl alcohol
J. N-Nitroso-di-n-propylamine*	Y. 2,4,6-Trichlorophenol**	NN. Fluorene	CCC. Benzo(a)anthracene	RRR. Pyridine
K. Hexachloroethane	Z. 2,4,5-Trichlorophenol	00. 4-Nitroaniline	DDD. Chrysene	SSS. Benzidine
L. Nitrobenzene	AA. 2-Chloronaphthalene	PP. 4,6-Dinitro-2-methylphenol	EEE. Bis(2-ethylhexyl)phthalate	ТТТ.
M. Isophorone	BB. 2-Nitroaniline	QQ. N-Nitrosodiphenylamine (1)**	FFF. Di-n-octylphthalate**	ΛΠΩ
N. 2-Nitrophenol**	CC. Dimethylphthalate	RR. 4-Bromophenyl-phenylether	GGG. Benzo(b)fluoranthene	ww.
0. 2,4-Dimethylphenol	DD. Acenaphthylene	SS. Hexachlorobenzene	HHH. Benzo(k)fluoranthene	www.

Notes:* = System performance check compound (SPCC) for RRF; ** = Calibration check compound (CCC) for %RSD.

COMPNDL

LDC #: <u>~~3/04</u> B 2b SDG #: <u>______</u>Cm_/

VALIDATION FINDINGS WORKSHEET Initial Calibration Calculation Verification

Page: _____ of <u>/</u>______ Reviewer: ______<u>0/0</u> 2nd Reviewer: ________

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

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

				Reported	Recalculated	Reported	Recalculated	Reported	Recalculated
		Calibration		RRF	RRF	Average RRF	Average RRF	%RSD	%RSD
#	Standard ID	Date	Compound (Internal Standard)	(50 std)	(50 std)	(Initial)	(Initial)		1
۲	ICAL	4/13/10	Naphthalene (IS2)	1.0312	1.0312	0.9822	0.9822	11.7	11.7
	MSSK		Fluorene (IS3)	1.3050	1.3050	1.2461	1.2462	11.2	11.2
			Phenanthrene (IS4)	1.0729	1.0729	1.0336	1.0336	13.4	13.4
			Chrysene (IS5)	1.0610	1.0610	1.0410	1.0410	10.7	10.7
			Benzo(a)pyrene (IS6)	1.1036	1.1036	1.0281	1.0281	5.7	5.7

			_	_		
Area IS	792159	483840	845901	981110	997687	
Area cpd	1021070	789238	1134473	1301240	1376307	
Conc IS/Cpd	40/50	40/50	40/50	40/50	40/50	

Benzo(a)py	0.9292	1.0099	1.0972	1.1036	1.0631	1.0333	0.9979	0.9906	1.0281	0.0587
Chrysene	1.1807	1.1392	1.1556	1.0610	1.0154	0.9617	0.9268	0.8874	1.0410	0.1110
Phenanth	1.2180	1.1707	1.1386	1.0729	1.0070	0.9388	0.8778	0.8449	1.0336	0.1388
Fluorene	1.3747	1.3919	1.3882	1.3050	1.2199	1.1599	1.0870	1.0426	1.2462	0.1395
Naphthalene	1.1409	1.0702	1.0728	1.0312	0.9598	0.9097	0.8529	0.8202	0.9822	0.1148
Conc	4.00	10.00	20.00	50.00	80.00	120.00	160.00	200.00	×	S II

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.

 $\frac{23/64}{\text{SDG } \# \frac{23/64}{\text{See Cover}}} \text{ } \beta 2 \beta$

VALIDATION FINDINGS WORSHEET Continuing Calibration Results Verification

Page 1 of 1 R J Reviewer:__ 2nd Reviewer:

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

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:

	Where:
% Difference = 100 * (ave. RRF - RRF)/ave. RRF	ave. RRF = initial calibi
RRF = (Ax)(Cis)/(Ais)(Cx)	RRF = continuing calib

ave. RRF = initial calibration average RRF RRF = continuing calibration RRF Ax = Area of compound

Cx = Concentration of compound Ais = Area of associated internal standard Cis = Concentration of internal standard

						_				_	 	
Recalculated %D	2.6	1.1	2.0	1.1	1.9							
Reported %D	2.6	1.1	2.0	1.1	1.9							
Recalculated (CC RRF)	0.956	1.233	1.013	1.029	1.047							
Reported (CC RRF)	0.956	1.233	1.013	1.029	1.047							
Average RRF (Initial RRF)	0.982	1.246	1.034	1.041	1.028							
Compound (Ref IS)	Naphthalene (IS2)	Fluorene (IS3)	Phenanthrene (IS4)	Chrysene (IS5)	Benzo(a)pyrene (IS6)							
Calibration Date	4/15/10											
Standard ID	K2738											
#	-					2			3			

	ccv1		CCV2		CCV3	
Compound	Area Cpd	Area IS	Area Cpd	Area IS	Area Cpd	Area IS
Naphthalene	1474236	770758				
Fluorene	1162445	471515				
Phenanthrene	1685735	831764				
Chrysene	2026913	984781				
Benzo(a)pyrene	2337667	1116096				

LDC #: 23/04 826 SDG #: Ste Cover

VALIDATION FINDINGS WORKSHEET Surrogate Results Verification



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

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

% Recovery: SF/SS * 100

#1

Where: SF = Surrogate Found SS = Surrogate Spiked

	Surrogate Spiked	Surrogate Found	Percent Recovery Reported	Percent Recovery Recalculated	Percent Difference
Nitrobenzene-d5	100	86.3	86	86	0
2-Fluorobiphenyl	1	84.2	84	8 F	
Terphenyl-d14		86.0	86	56	ł
Phenol-d5					
2-Fluorophenol					
2,4,6-Tribromophenol					
2-Chlorophenol-d4					
1,2-Dichlorobenzene-d4					

Sample ID:_____

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

Sample ID:

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

5	
DC # 23 104 1	DG# See Con
	0

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

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:

RPD = I MSC - MSC I * 2/(MSC + MSDC)

す

MS/MSD samples:

Where: SSC = Spiked sample concentration SA = Spike added

MSC = Matrix spike concentration

SC = Sample concentation

MSDC = Matrix spike duplicate concentration

Recalculated 4 ñ **MS/MSD** RPD Reported 5 N Matrix Spike Duplicate 105 Recalc و 8 Percent Recovery Reported. se) 20 も 84 Recalc Percent Recovery **Matrix Spike** Reported 54 53 2393 2962 MSD Spiked Sample Concentration (F) (F) 2290 2390 -202-Ś Sample Concentration (V5 /5,) 3 0 2110 2790 MSD Spike Addgd ž 28 00 28 00 NS N-Nitroso-di-n-propylamine 4-Chloro-3-methylphenol Compound Pentachlorophenol Acenaphthene Pyrene Phenol

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

	Labora
LDC #: 23164 830	SDG #: See Cover

atory Control Sample/Laboratory Control Sample Duplicates Results Verification VALIDATION FINDINGS WORKSHEET

Page: lof l Reviewer: JrC 2nd Reviewer:

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

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

% Recovery = 100 * (SC/SA

Where: SSC = Spike concentration SA = Spike added

RPD = I LCSC - LCSDC I * 2/(LCSC + LCSDC)

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

LCS/LCSD samples: 10 280- 10 YS1 /2-A

	ŝ	ike	S	ike	J J	s	U T	g	I CS/I	CSD
Compound	PA (۹۹)	ded /k)	Concel (MS	htration /c、)	Percent R	ecovery	Percent F	Recovery	RF	
	1 CS		1 CS	ار ۱ CSD	Reported	Recalc	Renorted	Recalc	Reported	Recalculated
Phenol										
N-Nitroso-di-n-propylamine										
4-Chloro-3-methylphenol										
Acenaphthene	515	ΨĄ	مكحر	tsy.	89	69				
Pentachlorophenol					-					
Pyrene	es 7c	\rightarrow	08 pc	~	94	94				

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

LDC #: ~310 + \$ 26 VALIDATION FINDINGS WORKSHEET SDG #: <u>Sre Cover</u> <u>Sample Calculation Verification</u>



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

Y) 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?

Conce	entratio	$n = (A_{,})(I_{,})(V_{,})(DF)(2.0) - (A_{,})(RRF)(V_{,})(V_{,})(%S)$	Example:
A _x	=	Area of the characteristic ion (EICP) for the compound to be measured	Sample I.D. #,
A_{is}	=	Area of the characteristic ion (EICP) for the specific internal standard	stand the second
l _s	=	Amount of internal standard added in nanograms (ng)	Conc. = (14008)(10)(1)(1000)(0)(0)(0)(0)(0)(0)(0)(0)(0)(0)(0)(0)
V _o	=	Volume or weight of sample extract in milliliters (ml) or grams (g).	
V,	=	Volume of extract injected in microliters (ul)	= 20.7
V,	=	Volume of the concentrated extract in microliters (ul)	
Df	=	Dilution Factor.	~ 21 us/kg
%S	=	Percent solids, applicable to soil and solid matrices only.	

Factor of 2 to account for GPC cleanup =

2.0 Reported Calculated Concentration Concentration Compound Qualification # Sample ID) 1)

Tronox LLC Facility,2010 Parcels, Henderson, Nevada Data Validation Reports LDC #23104

Arsenic



LDC Report# 23104A4

Laboratory Data Consultants, Inc. Data Validation Report

Project/Site Name: Tronox LLC Fac

Tronox LLC Facility, 2010 Parcels, Henderson, Nevada

Collection Date: April 6, 2010

LDC Report Date: May 18, 2010

Matrix: Soil/Water

Parameters: Arsenic

Validation Level: Stage 2B & 4

Laboratory: TestAmerica, Inc.

Sample Delivery Group (SDG): 280-2143-1

Sample Identification

P3-PF-2-1-0.0** P3-PF-2-1-0.0FD FB-PARCELS-032910 EB-PARCELS-032910 P3-PF-2-1-0.0MS P3-PF-2-1-0.0MSD EB-PARCELS-032910MS EB-PARCELS-032910MSD

**Indicates sample underwent Stage 4 review

Introduction

This data review covers 4 soil samples and 4 water samples listed on the cover sheet including dilutions and reanalysis as applicable. The analyses were per EPA SW 846 Methods 6020 for Arsenic.

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 Inorganic Data Review (October 2004).

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.

Blanks are summarized in Section IV.

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. ICPMS Tune

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

III. Calibration

An initial calibration was performed.

The frequency and analysis criteria of the initial calibration verification (ICV) and continuing calibration verification (CCV) were met.

IV. Blanks

Method blanks were reviewed for each matrix as applicable. No arsenic was found in the initial, continuing and preparation blanks.

Sample EB-PARCELS-032910 was identified as an equipment blank. No arsenic was found in this blank.

Sample FB-PARCELS-032910 was identified as a field blank. No arsenic was found in this blank.

V. ICP Interference Check Sample (ICS) Analysis

The frequency of analysis was met.

The criteria for analysis were met.

VI. Matrix Spike Analysis

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 within QC limits.

VII. Duplicate Sample Analysis

Duplicate (DUP) sample analyses were reviewed for each matrix as applicable.

VIII. Laboratory Control Samples (LCS)

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

IX. Internal Standards

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

X. Furnace Atomic Absorption QC

Graphite furnace atomic absorption was not utilized in this SDG.

XI. ICP Serial Dilution

ICP serial dilution analysis was performed by the laboratory. The analysis criteria were met.

XII. Sample Result Verification and Project Quantitation Limit

All sample result verifications were acceptable for samples on which a Stage 4 review was performed.

All analytes reported below the PQL were qualified as follows:

Sample	Finding	Finding Flag				
All samples in SDG 280-2143-1	All analytes 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

Samples P3-PF-2-1-0.0** and P3-PF-2-1-0.0FD were identified as field duplicates. No metals were detected in any of the samples with the following exceptions:

	Concentrat	ion (mg/Kg)	000	Difference			
Compound	P3-PF-2-1-0.0**	P3-PF-2-1-0.0FD	(Limits)	(Limits)	Flags	A or P	
Arsenic	3.1	3.2	3 (≤50)	-	-	-	

Tronox LLC Facility, 2010 Parcels, Henderson, Nevada Arsenic - Data Qualification Summary - SDG 280-2143-1

SDG	Sample	Analyte	Flag	A or P	Reason (Code)
280-2143-1	P3-PF-2-1-0.0** P3-PF-2-1-0.0FD FB-PARCELS-032910 EB-PARCELS-032910	All analytes reported below the PQL.	J (all detects)	A	Sample result verification (PQL) (sp)

Tronox LLC Facility, 2010 Parcels, Henderson, Nevada Arsenic - Laboratory Blank Data Qualification Summary - SDG 280-2143-1

No Sample Data Qualified in this SDG

Tronox LLC Facility, 2010 Parcels, Henderson, Nevada Arsenic - Equipment Blank Data Qualification Summary - SDG 280-2143-1

No Sample Data Qualified in this SDG

Tronox LLC Facility, 2010 Parcels, Henderson, Nevada Arsenic - Field Blank Data Qualification Summary - SDG 280-2143-1

No Sample Data Qualified in this SDG

Tronox Northgate Henderson

VALIDATION COMPLETENESS WORKSHEET

Stage 2B/4

SDG #: 280-2143-1 Laboratory: Test America

23104A4

LDC #:

Date: 5- 10-10 Page: __of___ Reviewer: ______ 2nd Reviewer: ______

METHOD: Arsenic (EPA SW 846 Method 6020)

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
1.	Technical holding times	Α	Sampling dates: 4-6-10
11.	ICP/MS Tune	A	
111.	Calibration	A	
IV.	Blanks	A	
V.	ICP Interference Check Sample (ICS) Analysis	A	
VI.	Matrix Spike Analysis	A	MS/MSD
VII.	Duplicate Sample Analysis	2	
VIII.	Laboratory Control Samples (LCS)	A	LCS
IX.	Internal Standard (ICP-MS)	A	
Х.	Furnace Atomic Absorption QC	N	not utilized
XI.	ICP Serial Dilution	A	
XII.	Sample Result Verification	A	Not reviewed for Stage 2B validation.
XIII	Overall Assessment of Data	A	
XIV.	Field Duplicates	รพ	D = 1 + 2
xv	Field Blanks	ND	FB=3 $EB=4$

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	P3-PF-2-1-0.0** S	11	21		31	
2	P3-PF-2-1-0.0FD	12	22		32	
3	FB-PARCELS-032910	13	23		33	
4	EB-PARCELS-032910	14	24		34	
5 I	P3-PF-2-1-0.0MS \$	15	25	······································	35	
6	P3-PF-2-1-0.0MSD	16	26		36	
7	EB-PARCELS-032910MS	17	27		37	
8	EB-PARCELS-032910MSD	18	28		38	
9	PBS	19	29		39	
10 2	PBW	20	30		40	

Notes:

Method:Metals (EPA SW 846 Method 6010/7000/6020)

Validation Area	Yes	No	NA	Findings/Comments
Ji Technical holding times				
All technical holding times were met.		· ·		·
Cooler temperature criteria was met.				
1. Calibration	i de la composición d T	аса —) Т		
Were all isotopes in the tuning solution mass resolution within 0.1 amu?	1	ļ		
Were %RSD of isotopes in the tuning solution < 5%?				
Were all instruments calibrated daily, each set-up time?			ļ	
Were the proper number of standards used?	1		ļ	
Were all initial and continuing calibration verification %Rs within the 90-110% (80- 120% for mercury and 85-115% for cyanide) QC limits?	1		ļ	
Were all initial calibration correlation coefficients > 0.995?			1912-00070	
JII, Blanks				
Was a method blank associated with every sample in this SDG?		<u> </u>		
Was there contamination in the method blanks? If yes, please see the Blanks validation completeness worksheet.		/		
IV JCP Interfetence Check Sample				
Were ICP interference check samples performed daily?				
Were the AB solution percent recoveries (%R) with the 80-120% QC limits?		1. Sector Sector		
IV Matrix spike/Matrix spike duplicates				
Were a matrix spike (MS) and duplicate (DUP) analyzed for each matrix in this SDG? If no, indicate which matrix does not have an associated MS/MSD or MS/DUP. Soil / Water.	/			
Were the MS/MSD percent recoveries (%R) and the relative percent differences (RPD) within the 75-125 QC limits? If the sample concentration exceeded the spike concentration by a factor of 4 or more, no action was taken.	/			
Were the MS/MSD or duplicate relative percent differences (RPD) \leq 20% for waters and \leq 35% for soil samples? A control limit of +/- RL(+/-2X RL for soil) was used for samples that were \leq 5X the RL, including when only one of the duplicate sample values were \leq 5X the RL.	/			
W. Laboratory control samples				
Was an LCS anaylzed for this SDG?	/			
Was an LCS analyzed per extraction batch?	\checkmark			
Were the LCS percent recoveries (%R) and relative percent difference (RPD) within the 80-120% QC limits for water samples and laboratory established QC limits for soils?	/			

Validation Area	Yes	No	NA	Findings/Comments
VI. Fumace Atomic Absorption QC				
If MSA was performed, was the correlation coefficients > 0.995?			\checkmark	
Do all applicable analysies have duplicate injections? (Level IV only)				
For sample concentrations > RL, are applicable duplicate injection RSD values < 20%? (Level IV only)			/	
Were analytical spike recoveries within the 85-115% QC limits?	023365270			
VII. ICP Senal Dilution				and the second
Was an ICP serial dilution analyzed if analyte concentrations were > 50X the IDL?	\checkmark			
Were all percent differences (%Ds) < 10%?	\checkmark			
Was there evidence of negative interference? If yes, professional judgement will be used to qualify the data.	-Southern Street			
VIII Internal Standards (EPA SW/846 Method 6020)				
Were all the percent recoveries (%R) within the 30-120% of the intensity of the internal standard in the associated initial calibration?				
If the %Rs were outside the criteria, was a reanalysis performed?		ana		
IX Regional Quality Assurance and Quality Control	1 1			
Were performance evaluation (PE) samples performed?				
Were the performance evaluation (PE) samples within the acceptance limits?				
X Sample Result Venification			<u>.</u>	
Were RLs adjusted to reflect all sample dilutions and dry weight factors applicable to level IV validation?				
XI: Overalli assessment of cata				
Overall assessment of data was found to be acceptable.	1			
XII. Field ddplicates				
Field duplicate pairs were identified in this SDG.	/			
Target analytes were detected in the field duplicates.	1			
ANE Frequioring				
Field blanks were identified in this SDG.	/			
Target analytes were detected in the field blanks.		\backslash		

LDC#: 23104A4 SDG#: 280-2143-1

VALIDATION FINDINGS WORKSHEET Field Duplicates

Page: | of | Reviewer: MG 2nd Reviewer:

METHOD: Metals (EPA Method 6010B/6020/7000)

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

	Concentratio	on (mg/kg)	(≤50)	(mg/Kg) (mg/Kg)		Qualifications
Analyte	1	2	RPD	Difference	Limits	(Parent Only)
Arsenic	3.1	3.2	3			

V:\FIELD DUPLICATES\FD_inorganic\263104A4.wpd

LDC #. 23104A4 SDG #. 280-2143-1

VALIDATION FINDINGS WORKSHEET Initial and Continuing Calibration Calculation Verification

Reviewer: MG Page: Lof 2nd Reviewer:___

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

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

Where, Found = concentration (In ug/L) of each analyte measured in the analysis of the ICV or CCV solution True = concentration (in ug/L) of each analyte in the ICV or CCV source %R = <u>Found</u> x 100 True

					Recalculated	Reported	
Standard ID	Type of Analysis	Element	Found (ug/L)	True (ug/L)	%R	%R	Acceptable (Y/N)
	ICP (Initial calibration)						
	GFAA (Initial calibration)						
	CVAA (Initial calibration)						
	ICP (Continuing calibration)						
	GFAA (Continuing calibration)						
	CVAA (Continuing calibration)						
1995 ICV	ICP/MS (Initial calibration)	As	40.57	40.0	101	101	Y
0144 CCV	ICP/MS (Continuing callbation)	As	51, 89	50.0	104	104	~

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

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LDC # <u>33104</u> SDG # <u>780-21</u>	Ач 43-1		VALIDA7 Level	rion FIN IV Recal	IDINGS WO culation Wo	RKSHEI <u>orkshee</u> t	h	Ñ	Page: of Reviewer: 서중 nd Reviewer:
METHOD: Trace Me	etals (EPA SW 846 Metho	d 6010/7	(000						
Percent recoveries ((%R) for an ICP interferen	ce check	sample, a labo	ratory cont	rol sample and	a matrix s	spike sample were r	ecalculated using th	e following formula:
%R = Found x 100 True	Where, Found = Conce True :	entration of Found = Concei	each analyte <u>meas</u> = SSR (spiked sam ntration of each an	ured in the a tple result) - a alyte in the so	nalysis of the sam SR (sample result) Nrce.	ple. For the r.	natrix spike calculation,		
A sample and duplic	cate relative percent differ	ence (RP	D) was recalcul	lated using	the following f	ormula:			
RPD = <u>[S-D]</u> x 100 (S+D)/2	Where, S = C D = D	higinal sam Juplicate sa	ple concentration imple concentration	_					
An ICP serial dilutio	in percent difference (%D)) was rec	alculated using	the followi	ng formula:				
%D = <u>1-SDR1</u> x 100 1	Where, I = In SDR = Serial D	itial Sample flution Rest	e Result (mg/L) ult (mg/L) (Instrume	ent Reading x	(2)				
							Recalculated	Reported	
Sample ID	Type of Analysis	Element	Found / 5 (units)	3/1	True / D / SDR	(units)	%R / RPD / %D	%R / RPD / %D	Acceptable (Y/N)
1958 ICSAB	ICP Interference check	As	104.20	(mg/r)	100.	(ng/r)	104	104	· /
2 C S	Laboratory control sample	Å۶	19.63	(mg/ky)	0'08	(mg/kg)	98	86	
7000 5	Matrix spike	As	(ssr.sr) 18.23	(mg/kg	18.5	(mg/kg)	66	86	
0207/0210 5/6	Duplicate	AS	91.34	(mg/kg)	23.14	(m3/ba)	8	8	
0158 / 0201 	ICP serial dilution	As	3.115	(mg/kg)	3.096	mg/kg)	19.0 *	0.53	
Comments: Refer	to appropriate worksheet	for list of	qualifications a	0 Ind associ	ated samples v	vhen repor	ted results do not a	gree within 10.0% of	the recalculated results.
1 2601 X X	>; both in lim.	× •	o gual.						

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LDC #: 2310444 SDG #: 280-2143-1

VALIDATION FINDINGS WORKSHEET Sample Calculation Verification

Page:	of
Reviewer:	MG
2nd reviewer:	\sim

were recalculated and verified using the

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

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".(Y) N N/AHave results been reported and calculated correctly?(Y) N N/AAre results within the calibrated range of the instruments and within the linear range of the ICP?(Y) N N/AAre all detection limits below the CRDL?

As

Detected analyte results for \pm , following equation:

Concentration = <u>(RD)(FV)(Dil)</u> (In. Vol.)(%S) Recalculation:

RD FV	= =	Raw data concentration Final volume (ml)	(6.79 mg/L)(0	100 1) (5)	2115	мд	mg
in. vol. Dil %S	# #	Dilution factor Decimal percent solids	1.09 g	wet weight	5.115	g or	Kg

Sample ID	Analyte	Reported Concentration (M & /kg)	Calculated Concentration (Mg./kg)	Acceptable (Y/N)
1	As	3.1	3.1	Ý
			· · · ·	
	· · · · · · · · · · · · · · · · · · ·			
	· · · · · · · · · · · · · · · · · · ·			· · · · _ ·
	· · · · · · · · · · · · · · · · · · ·			
		······		
			<u>.</u>	

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