

LABORATORY DATA CONSULTANTS, INC.

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ERM 2525 Natomas Park Drive, Suite 350

Sacramento, CA 95833

ATTN: Ms. Maria Barajas-Albalawi

SUBJECT: BRC Tronox Parcel H, Data Validation

Dear Ms. Barajas-Albalawi

Enclosed are the final validation reports for the fractions listed below. These SDGs were received on March 21, 2008. Attachment 1 is a summary of the samples that were reviewed for each analysis.

LDC Project # 18484:

SDG # Fraction

204220 Gross Alpha and Beta, Gamma Spectroscopy

The data validation was performed under EPA Level III and Level IV guidelines. The analyses were validated using the following documents, as applicable to each method:

- USEPA, Contract Laboratory Program National Functional Guidelines for Inorganic Data Review, October 2004
- EPA SW 846, Third Edition, Test Methods for Evaluating Solid Waste, update 1, July 1992; update IIA, August 1993; update II, September 1994; update IIB, January 1995; update III, December 1996; update IIIA, April 1998; IIIB, November 2004; Update IV, February 2007

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

Sincerely,

Erlinda T. Rauto

Operations Manager/Senior Chemist

April 28, 2008

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214 Pages-EX	80/20	*SDG*	Water/Soil	204220	204220																T/LR
		LDC	Matrix:	4	4										_					$\frac{1}{1}$	Total

Laboratory Data Consultants, Inc. Data Validation Report

Project/Site Name:

BRC Tronox Parcel H

Collection Date:

March 6, 2008

LDC Report Date:

April 3, 2008

Matrix:

Soil

Parameters:

Gross Alpha and Beta

Validation Level:

EPA Level III & IV

Laboratory:

General Engineering Laboratories, LLC.

Sample Delivery Group (SDG): 204220

Sample Identification

TSB-HJ-10-Surf

TSB-HJ-01-Surf**

TSB-HR-05-Surf

TSB-HJ-10-SurfMS

TSB-HJ-10-SurfMSD

TSB-HJ-10-SurfDUP

^{**}Indicates sample underwent Level IV review

Introduction

This data review covers 6 soil samples listed on the cover sheet. The analyses were per EPA Method 900.0 for Gross Alpha and Beta Radioactivity.

The review follows a modified outline of the USEPA Contract Laboratory Program National Functional Guidelines for Inorganic Data Review (October 2004) as there are no current guidelines for the method stated above.

A qualification summary table is provided at the end of this report if data has been qualified. Flags are classified a 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 III.

Field duplicates are summarized in Section VIII.

Samples indicated by a double asterisk on the front cover underwent a EPA Level IV review. A EPA Level III review was performed on all other samples. Raw data were not evaluated for the samples reviewed by Level III 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.
- R Data are qualified as rejected. There is a significant potential for the reporting of false negatives or false positives.
- U Data are qualified as non-detected, because the analyte was observed in as associated laboratory or field blank.
- 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. Calibration

a. Initial Calibration

All criteria for the initial calibration were met.

Detector efficiency was determined and a self-absorption curve was generated for each radionuclide of interest.

b. Continuing Calibration

Calibration verification and background determination were performed at the required frequencies. Results were within laboratory control limits.

III. Blanks

Method blanks were reviewed for each matrix as applicable. Blank results contained less than the minimum detectable activity (MDA) with the following exceptions:

Method Blank ID	Analyte	Activity	Associated Samples
PB (prep blank)	Gross beta	4.34 pCi/g	All samples in SDG 204220

No sample data were qualified based on the gross alpha or beta contaminants found in the method blanks.

No field blanks were identified in this SDG.

IV. Accuracy and Precision Data

a. Matrix Spike/(Matrix Spike) Duplicate

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.

Duplicate (DUP) sample analyses were reviewed for each matrix as applicable. Results were within QC limits.

b. Laboratory Control Samples

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

V. Minimum Detectable Activity (MDA)

All minimum detectable activities met required detection limits.

VI. Sample Result Verification

All sample result verifications were acceptable for samples on which a Level IV review was performed. Raw data were not evaluated for the samples reviewed by Level III criteria.

VII. Overall Assessment of Data

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

VIII. Field Duplicates

No field duplicates were identified in this SDG.

BRC Tronox Parcel H Gross Alpha and Beta - Data Qualification Summary - SDG 204220

No Sample Data Qualified in this SDG

BRC Tronox Parcel H Gross Alpha and Beta - Laboratory Blank Data Qualification Summary - SDG 204220

No Sample Data Qualified in this SDG

BRC Tronox Parcel H Gross Alpha and Beta - Field Blank Data Qualification Summary - SDG 204220

No Sample Data Qualified in this SDG

VALIDATION COMPLETENESS WORKSHEET LDC #: 18484A22

SDG #: 204220

Laboratory: GEL Laboratories LLC

Level III/IV

Date:	4-1-08
Page:_	<u> of </u>
Reviewer:	
2nd Reviewer:	\sim

METHOD: Gross Alpha & Beta (EPA SW 846 Method 900.0)

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	Α	Sampling dates: 3 - 6 - 08
IIa.	Initial calibration	A	
IIb.	Calibration verification	À	
111.	Blanks	SW	
IVa.	Matrix Spike/(Matrix Spike) Duplicates	A	MS/MSD/DUP
IVb.	Laboratory control samples	Α	LCS
V.	Minimum dectectable activity (MDA)	A	
VI.	Sample result verification	A	Not reviewed for Level III validation.
VII.	Overall assessment of data	A	
VIII.	Field duplicates	N	
IX	Field blanks	7	

Note:

A = Acceptable

ND = No compounds detected

D = Duplicate

N = Not provided/applicable SW = See worksheet

R = Rinsate FB = Field blank

TB = Trip blank EB = Equipment blank

Validated Samples: ** Indicates sample underwent Level IV validation

911 5011 11 31 TSB-HJ-10-Surf 21 12 22 32 2 TSB-HJ-01-Surf** TSB-HR-05-Surf 13 23 33 3 34 4 TSB-HJ-10-SurfMS 14 24 TSB-HJ-10-SurfMSD 15 25 35 6 TSB-HJ-10-SurfDUP 16 26 36 PBS 17 37 27 8 18 28 38 19 29 39 20 30 40 10

Notes:		

LDC #: 18484A22 SDG #: 204220

VALIDATION FINDINGS CHECKLIST

Page: 1 of 2 Reviewer: $\mu \sigma$ 2nd Reviewer: $\mu \sigma$

Method:Radiochemistry(EPA Method 900.0)

Method: Radiochemistry (EPA Method 100:0)	Yes	No	NA	Findings/Comments
I. Technical holding times	1 .33	1	1	1
All technical holding times were met.	TZ			
II. Calibration	1		1	
Were all instruments and detectors calibration as required?	1			
Were NIST traceable standards used for all calibrations?	1/			
Was the check source identified by activity and radionuclide?	/			
Were check sources including background counts analyzed at the required frequency and within laboratory control limits?	<i>'</i>			
III, Blanks				
Were blank analyses performed as required?	/			
Were any activities detected in the blanks greater than the minimum detectable activity (MDA)? If yes, please see the Blanks validation completeness worksheet.	/			
IV. Matrix spikes and Duplicates				
Were a matrix spike (MS) 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 percent recoveries (%R) within the QC limits? If the sample concentration exceeded the spike concentration by a factor of 4 or more, no action was taken.	/		ŕ	
Was a duplicate sample anaylzed at the required frequency of 5% in this SDG?	<u>/</u>			
Were all duplicate sample duplicate error rations (DER) ≤1.42?.	. 🗸			
V. Laboratory control samples				
Was an LCS analyzed per analytical batch?	<u> </u>			
Were the LCS percent recoveries (%R) and relative percent difference (RPD) within the 75-125%	/			
VI. Sample Chemical/Carrier Recovery				
Was a tracer/carrier added to each sample?		\checkmark		
Were tracer/carrier recoveries within the QC limits?			<u> </u>	
VII. Regional Quality Assurance and Quality Control		- 7 1		
Were performance evaluation (PE) samples performed?		/		
Were the performance evaluation (PE) samples within the acceptance limits?			<u> </u>	
vIII. Sample Result Verification			 1	
Were activities adjusted to reflect all sample dilutions and dry weight factors applicable to level IV validation?	/			
Were the Minimum Detectable Activities (MDA) < RL?	<u> </u>			

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LDC #: 18484A22 SDG #: 204320

VALIDATION FINDINGS CHECKLIST

Page: 2 of 2
Reviewer: MG
2nd Reviewer: \(\sum \)

Validation Area	Yes	No	NA	Findings/Comments
X: Overall assessment of data				
Overall assessment of data was found to be acceptable.	/			
X Field duplicates				
Field duplicate pairs were identified in this SDG.		/		ŀ
Target analytes were detected in the field duplicates.			/	
XI: Field blanks				
Field blanks were identified in this SDG.		/		
Target analytes were detected in the field blanks.			V	

LDC #: 18484 A 72 SDG #: 204 230

VALIDATION FINDINGS WORKSHEET BIANKS

Page: 1 of 1 Reviewer: 7 c

METHOD: Radiochemistry (Method: つつこの NA N/A Were blank analyses performed as required? If no

YN N/A

Were any activities detected in the blanks greater than the minimum detectable activity (MDA)? If yes, please see qualifications below. Were blank analyses performed as required? If no, please see qualifications below.

						200	
		Fied					
	ntification	No samples were qualified	7				
ر ا الادن ا ا	Sample Identification	Were		,			
		amples					
all	-	No s					
Samples:							
Associated Samples:							·
	Blank Action	Level	. .				
PC1/9	Blank ID	PBS	4.34				
Units:	Isotope		Gross Bota 4.34				

Units:		,	Associated Samples:	mples:					
Isotope	Blank ID	Blank Action				Sample Identification	ntification		
		Level			-				
		·							
		-							
		·							
							-		

CIRCLED RESULTS WERE NOT QUALIFIED. ALL RESULTS NOT CIRCLED WERE QUALIFIED BY THE FOLLOWING STATEMENT: If there is activity in the blank above the MDA, sample results within 10x the blank activity will be qualified as not detected "U".

LDC #: 18484 A 22 SDG #:

VALIDATION FINDINGS WORKSHEET Level IV Recalculation Worksheet

りぇ 2nd Reviewer: Reviewer:

METHOD: Radiochemistry (Method:___

900.0

Percent recoveries (%R) for a laboratory control sample, a matrix spike and a matrix spike duplicate sample were recaluculated using the following formula:

%R = Found x 100

Found = activity of each analyte measured in the analysis of the sample. True = activity of each analyte in the source.

Where,

A matrix spike and matrix spike duplicate relative percent difference (RPD) was recalculated using the following formula:

S = Original sample activity
D = Duplicate sample activity Where,

RPD = $\frac{|S-D|}{(S+D)/2} \times 100$

					Recalculated	Reported	
Sample ID	Type of Analysis	Analyte	Found/S (units)	True/D (units)	%R or RPD	%R or RPD	Acceptable (Y/N)
	Laboratory control sample						
577		Gross Betg	314 (pc:/4)	314 (pc/4) 301 (pc/4)	h01	HQ1	>
	Matrix spike sample	Gross					
T		Alpha	111.4 (Pci/g)	1.4 (PCi/g) 116 (PCi/g)	96	96	
	Duplicate RPD	Gross					
9		Beta	33.0 (PCi/g)	33.0 (PCilg) 34.3 (PCilg)	7	7	→
	Chemical recovery			1			
1		1	1		1		1

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

LDC #:_	18484A22
	204220

VALIDATION FINDINGS WORKSHEET Sample Calculation Verification

Page:_	of
Reviewer:	MG
2nd reviewer:	

METHOD: Radiochemistry (Method: 900.0

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

Have results been reported and calculated correctly? Y) N N/A

Are results within the calibrated range of the instruments? N N/A

Gross X reported with a positive detect were recalculated Analyte results for and verified using the following equation:

Activity =

Recalculation:

(cpm - bckgrd cpm) (2.22)(E)(Vol)(CF)

 $\frac{0.5428 - \left(\frac{487}{130} \times 0.01061\right)}{2.92)(0.1355)(0.109g)} = 15.342$

E = Efficiency Vol = Volume

CF = %R, Self-absorbance, abundance, ect.

#	Sample ID	Analyte	Reported Concentration (ア ^{と・} /な)	Calculated Concentration (PCi/g)	Acceptable (Y/N)
,	2	Gross &	15. 3	15.3	· Y
		Gross B	. 33.7	34. 0	1
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Note:	

Laboratory Data Consultants, Inc. Data Validation Report

Project/Site Name:

BRC Tronox Parcel H

Collection Date:

March 6, 2008

LDC Report Date:

April 3, 2008

Matrix:

Soil

Parameters:

Gamma Spectroscopy

Validation Level:

EPA Level III & IV

Laboratory:

General Engineering Laboratories, LLC.

Sample Delivery Group (SDG): 204220

Sample Identification

TSB-HJ-10-Surf TSB-HJ-01-Surf** TSB-HR-05-Surf

TSB-HJ-10-SurfDUP

^{**}Indicates sample underwent Level IV review

Introduction

This data review covers 4 soil samples listed on the cover sheet. The analyses were per EML HASL Method 300.4.5.2.3 for Gamma Spectroscopy.

The review follows a modified outline of the USEPA Contract Laboratory Program National Functional Guidelines for Inorganic Data Review (October 2004) as there are no current guidelines for the method stated above.

A qualification summary table is provided at the end of this report if data has been qualified. Flags are classified a 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 III.

Field duplicates are summarized in Section VIII.

Samples indicated by a double asterisk on the front cover underwent a EPA Level IV review. A EPA Level III review was performed on all other samples. Raw data were not evaluated for the samples reviewed by Level III 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.
- R Data are qualified as rejected. There is a significant potential for the reporting of false negatives or false positives.
- U Data are qualified as non-detected, because the analyte was observed in as associated laboratory or field blank.
- 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. Calibration

a. Initial Calibration

All criteria for the initial calibration were met.

Detector efficiency was determined for each radionuclide of interest.

b. Continuing Calibration

Calibration verification and background determination was performed at the required frequencies.

III. Blanks

Method blanks were reviewed for each matrix as applicable. Blank results contained less than the minimum detectable activity (MDA).

No field blanks were identified in this SDG.

IV. Accuracy and Precision Data

a. Matrix Spike/(Matrix Spike) Duplicates

A matrix spike (MS) analysis was not required by the method.

Duplicate (DUP) sample analyses were reviewed for each matrix as applicable. Results were within QC limits.

b. Laboratory Control Samples

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

V. Minimum Detectable Activity

All minimum detectable activities met required detection limits.

VI. Sample Result Verification

All sample result verifications were acceptable for samples on which a Level IV review was performed. Raw data were not evaluated for the samples reviewed by Level III criteria.

VII. Overall Assessment of Data

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

VIII. Field Duplicates

No field duplicates were identified in this SDG.

BRC Tronox Parcel H
Gamma Spectroscopy - Data Qualification Summary - SDG 204220

No Sample Data Qualified in this SDG

BRC Tronox Parcel H
Gamma Spectroscopy - Laboratory Blank Data Qualification Summary - SDG
204220

No Sample Data Qualified in this SDG

BRC Tronox Parcel H
Gamma Spectroscopy - Field Blank Data Qualification Summary - SDG 204220

No Sample Data Qualified in this SDG

SDG	#:18484A35 #:204220 ratory :_GEL Laborator				PLETEN evel III/I\		EET	Page: 4-1-08 Page: 1 of 1 Reviewer: MG 2nd Reviewer:
The s	HOD: Gamma Spectro camples listed below wation findings workshee	ere revi		•	,	alidation areas. Va	lidation findi	ngs are noted in attached
	Validatio	on Area				C	Comments	
1.	Technical holding times			Α	Sampling of	dates: 3-6	- 03	
IIa.	Initial calibration			A				
IIb.	Calibration verification			A				
10.	Blanks			A				
IVa.	Matrix Spike/(Matrix Spik	e) Duplica	ates	Α	DUP			
IVb.	Laboratory control samp	les		Α	LCS			
V.	Minimum dectectable ac	tivity (MD/	۹)	A				
VI.	Sample result verification	n		Α	Not reviev	ved for Level III validation	on.	
VII.	Overall assessment of d	ata		A				
VIII.	Field duplicates			7				
Lix	Field blanks			7	<u> </u>			
Note: ∕alida	A = Acceptable N = Not provided/applica SW = See worksheet ed Samples: ** Indicates so		R = Rin FB = Fi	eld blank		D = Duplicate TB = Trip blani EB = Equipme		
1	TSB-HJ-10-Surf	11			21		31	
2	TSB-HJ-01-Surf**	12			22		32	
3	TSB-HR-05-Surf	13			23		33	
4	TSB-HJ-10-SurfDUP	14			24		34	
5	PBS	15			25		35	
6		16			26		36	
7		17			27		37	
8		18			28		38	
9		19			29		39	
10		20			30		40	

LDC #: 18404A35 SDG #: 204770

VALIDATION FINDINGS CHECKLIST

Page: 1 of 2
Reviewer: MG
2nd Reviewer:

Method:Radiochemistry(EPA Method EML HASL) 300, 4.5.2.3

Validation Area	Yes	No	NA	Findings/Comments
I. Technical holding times				
All technical holding times were met.	/			
II. Calibration				
Were all instruments and detectors calibration as required?	·/			
Were NIST traceable standards used for all calibrations?	. 🗸			1
Was the check source identified by activity and radionuclide?	/			
Were check sources including background counts analyzed at the requiried frequency and within laboratory control limits?	V			
III. Blanks				
Were blank analyses performed as required?	\checkmark			
Were any activities detected in the blanks greater than the minimum detectable activity (MDA)? If yes, please see the Blanks validation completeness worksheet.		/		
IV. Matrix spikes and Duplicates	- 1			
Were a matrix spike (MS) 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 percent recoveries (%R) within the QC limits? If the sample concentration exceeded the spike concentration by a factor of 4 or more, no action was taken.			/	
Was a duplicate sample anayized at the required frequency of 5% in this SDG?	/			
Were all duplicate sample duplicate error rations (DER) ≤1.42?.	$\sqrt{}$			
y. Laboratory control samples				
Was an LCS analyzed per analytical batch?				
Were the LCS percent recoveries (%R) and relative percent difference (RPD) within the 75-125%				
VI. Sample Chemical/Carrier Recovery				
Was a tracer/carrier added to each sample?		/		
Were tracer/carrier recoveries within the QC limits?			<u> </u>	
VII. Regional Quality Assurance and Quality Control				
Were performance evaluation (PE) samples performed?				
Were the performance evaluation (PE) samples within the acceptance limits?			<u> </u>	
VIII. Sample Result Verification				
Were activities adjusted to reflect all sample dilutions and dry weight factors applicable to level IV validation?	/			
Were the Minimum Detectable Activities (MDA) < RL?	<u> </u>			
A STATE OF THE STA				

LDC #: 18484A35 SDG #: 204820

VALIDATION FINDINGS CHECKLIST

Page: 2 of 2
Reviewer: MG
2nd Reviewer: \(\sum \)

Validation Area	Yes	No	NA	Findings/Comments
IX Overell assessment of data				
Overall assessment of data was found to be acceptable.	/			
X. Field: duplicates				
Field duplicate pairs were identified in this SDG.		/		I
Target analytes were detected in the field duplicates.			/	
XI: Field blanks				
Field blanks were identified in this SDG.		/		
Target analytes were detected in the field blanks.			/	

18434 A 35 LDC #:

VALIDATION FINDINGS WORKSHEET **Level IV Recalculation Worksheet**

Page: Reviewer: 2nd Reviewer:

> 4.5.3.3 METHOD: Radiochemistry (Method: EML HASL 300)

Percent recoveries (%R) for a laboratory control sample, a matrix spike and a matrix spike duplicate sample were recaluculated using the following formula:

%R = Found x 100

Found = activity of each analyte measured in the analysis of the sample. True = activity of each analyte in the source.

Where,

A matrix spike and matrix spike duplicate relative percent difference (RPD) was recalculated using the following formula:

RPD = $\frac{1S-D_1}{(S+D)/2}$ × 100

S = Original sample activity D = Duplicate sample activity Where,

	a J	o I	C/6) 16.0 (PC:/)	(pC,q)	A m , 241		5)7
						Laboratory control sample	
Acce	%R or RPD	%R or RPD	True/D (units)	Found/S (units)	Analyte	Type of Analysis	Sample ID
	Reported	Recalculated					

					Recalculated	Reported	
	Type of Analysis	Analyte	Found/S (units)	True/D (units)	%R or RPD	%R or RPD	Acceptable (Y/N)
	Laboratory control sample						
		Am-241	ফ	,2 (PC/4) 16.0 (PC/4)	95	95	>
	Matrix spike sample						
- 1)	١		١	١	J
	Duplicate RPD						
		Ac-238	1.85 (PCi/q) 1.98 (PCi/q	1.98 (PCi/z)	7	9	7 *
	Chemical recovery	1		i			
					1	1	ĺ

Comments: Refer to appropriate worksheet for list of qualifications and associated samples when reported results do not agree within 10.0% of the recalculated results. 1 mi + 5 5 values 400 h

TOTCLC.35

LDC #:_	18484 A35
SDG #:	204220

VALIDATION FINDINGS WORKSHEET Sample Calculation Verification

Page:	l_of
Reviewer:	MG.
2nd reviewer:	

EML HASL 300, 4.5.2.3 METHOD: Radiochemistry (Method:_

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

Have results been reported and calculated correctly? N/A

Are results within the calibrated range of the instruments? Ý) N N/A

2 Ac-228 reported with a positive detect were recalculated Analyte results for and verified using the following equation:

Activity =

Recalculation:

(cpm - bckgrd cpm) (2.22)(E)(Vol)(CF)

E = Efficiency Vol = Volume

 $\frac{(211/120)}{(2.22)(0.01181)(0.2770)(153.050g)} = 1.582 \text{ pci/g}$

CF = %R, Self-absorbance, abundance, ect.

#	Sample ID	Analyte	Reported Concentration (P ^C /a)	Calculated Concentration	Acceptable (Y/N)
	2	Ac - 228	1.58	1.58	Y
		Bi - 214	. 1-16	1.16	
		Pb-212	1.69	1.68,	
		Pb-214	0.996	0.996	
		Po-212	1.69	1.68	
		Po-214	0.996	0.996	
		Po-216	1.69	1.68	·
		Po-218	0.996	0.996	
		K- 40	21.3	21.3	
		Ra-226	1.16	1.16	
	· ·	Ra - 228	1.58	1.58	
		T1-208	0.613	0.610	
		Th-234	1.23	1.23	
		U- 238	1. 23	1.23	1
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Note:	