



EBERLINE
SERVICES

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August 8, 2011

121350

Mr. Michael Neely
CH2M Hill Plateau Remediation Company
P.O. Box 1600
Mail Stop – B6-06
Richland, WA 99352

Reference: P.O. #33677
Eberline Analytical S1-08-007-7752, SDG H4639

Dear Mr. Neely:

Enclosed is a data report for six solid (soil) samples designated under SAF No. F11-127 received at Eberline Analytical on August 1, 2011. The samples were analyzed according to the accompanying chain-of-custody documents.

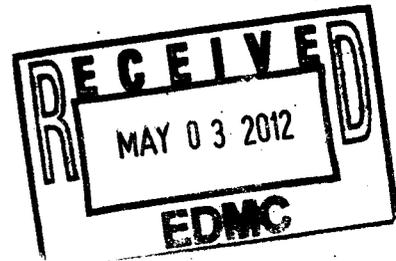
Please call if you have any questions concerning this report.

Sincerely,

N. Joseph Verville
Client Services Manager

NJV/kwp

Enclosure: Data Package



1.0 GENERAL

CH2M Hill Plateau Remediation Company (CHPRC) Sample Delivery Group H4639 was composed of six solid (soil) samples designated under SAF No. F11-127 with a Project Designation of: ARRA Verification Sampling of the 216-S-26 Waste Site and 200-W-148-PL.

The samples were received as stated on the chain-of-custody documents. Any discrepancies are noted on the Eberline Analytical Sample Receipt Checklist.

2.0 ANALYSIS NOTES

2.1 Tritium Analysis

The results for both the original and duplicate analyses were less than their respective MDA's, therefore no RPD or statistical control limit was calculated. No other problems were encountered during the course of the analyses.

3.0 Case Narrative Certification Statement

"I certify that this data package is in compliance with the SOW, both technically and for completeness, for other than the conditions detailed above. Release of the data obtained in this hard copy data package has been authorized by the Laboratory Manager or a designee, as verified by the following signature."



N. Joseph Verville
Client Services Manager

8/8/11

Date

EBERLINE ANALYTICAL / RICHMOND
SAMPLE DELIVERY GROUP H4639

SDG 7752
Contact N. Joseph Verville

Client CHPRC
Contract No. 33677
Case no SDG_H4639

S U M M A R Y D A T A S E C T I O N

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Prepared by


Reviewed by

Lab id EBRLNE
Protocol F11-127
Version Ver 1.0
Form DVD-TOC
Version 3.06
Report date 08/08/11

EBERLINE ANALYTICAL / RICHMOND

SAMPLE DELIVERY GROUP H4639

SDG 7752
Contact N. Joseph Verville

REPORT GUIDE

Client CHPRC
Contract No. 33677
Case no SDG H4639

ABOUT THE DATA SUMMARY SECTION

The Data Summary Section of a Data Package has all data, in several useful orders, necessary for first level, routine review of the data package for a Sample Delivery Group (SDG). This section follows the Data Package Narrative, which has an overview of the data package and a discussion of special problems. It is followed by the Raw Data Section, which has full details.

The Data Summary Section has several groups of reports:

SAMPLE SUMMARIES

The Sample and QC Summary Reports show all samples, including QC samples, reported in one SDG. These reports cross-reference client and lab sample identifiers.

PREPARATION BATCH SUMMARY

The Preparation Batch Summary Report shows all preparation batches (lab groupings reflecting how work was organized) relevant to the reported SDG with information necessary to check the completeness and consistency of the SDG.

WORK SUMMARY

The Work Summary Report shows all samples and work done on them relevant to the reported SDG.

METHOD BLANKS

The Method Blank Reports, one for each Method Blank relevant to the SDG, show all results and primary supporting information for the blanks.

LAB CONTROL SAMPLES

The Lab Control Sample Reports, one for each Lab Control Sample relevant to the SDG, show all results, recoveries and primary supporting information for these QC samples.

REPORT GUIDES

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SUMMARY DATA SECTION

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Lab id EBRLNE
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Form DVD-RG
Version 3.06
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EBERLINE ANALYTICAL / RICHMOND

SAMPLE DELIVERY GROUP H4639

SDG 7752
Contact N. Joseph Verville

GUIDE, cont.

Client CHPRC
Contract No. 33677
Case no SDG H4639

ABOUT THE DATA SUMMARY SECTION

DUPLICATES

The Duplicate Reports, one for each Duplicate and Original sample pair relevant to the SDG, show all results, differences and primary supporting information for these QC samples.

MATRIX SPIKES

The Matrix Spike Reports, one for each Spiked and Original sample pair relevant to the SDG, show all results, recoveries and primary supporting information for these QC samples.

DATA SHEETS

The Data Sheet Reports, one for each client sample in the SDG, show all results and primary supporting information for these samples.

METHOD SUMMARIES

The Method Summary Reports, one for each test used in the SDG, show all results, QC and method performance data for one analyte on one or two pages. (A test is a short code for the method used to do certain work to the client's specification.)

REPORT GUIDES

The Report Guides, one for each of the above groups of reports, have documentation on how to read the associated reports.

REPORT GUIDES

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SUMMARY DATA SECTION

Page 2

Lab id EBRLNE
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EBERLINE ANALYTICAL/RICHMOND

SAMPLE DELIVERY GROUP H4639

SDG 7752
 Contact N. Joseph Verville

Client CHPRC
 Contract No. 33577
 Case no SDG H4639

LAB SAMPLE SUMMARY

LAB SAMPLE ID	CLIENT SAMPLE ID	LOCATION	MATRIX	LEVEL	SAF NO	CHAIN OF CUSTODY	COLLECTED
S108007-01	B2FBT1	216-S-26 IP-1	SOIL		F11-127	F11-127-063	07/27/11 14:40
S108007-02	B2FBT2	216-S-26 IP-2	SOIL		F11-127	F11-127-064	07/27/11 14:44
S108007-03	B2FBT3	216-S-26 IP-3	SOIL		F11-127	F11-127-065	07/27/11 14:47
S108007-04	B2FPF6	216-S-26 IP-4	SOIL		F11-127	F11-127-095	07/27/11 15:03
S108007-05	B2FPF7	216-S-26 IP-5	SOIL		F11-127	F11-127-096	07/27/11 15:08
S108007-06	B2FPF8	216-S-26 IP-6	SOIL		F11-127	F11-127-097	07/27/11 15:12
S108007-07	Lab Control Sample		SOIL		F11-127		
S108007-08	Method Blank		SOIL		F11-127		
S108007-09	Duplicate (S108007-01)	216-S-26 IP-1	SOIL		F11-127		07/27/11 14:40

LAB SUMMARY

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SUMMARY DATA SECTION

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Lab id EBRLNE
 Protocol F11-127
 Version Ver 1.0
 Form DVD-LS
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EBERLINE ANALYTICAL/RICHMOND

SAMPLE DELIVERY GROUP H4639

SDG 7752
 Contact N. Joseph Verville

QC SUMMARY

Client CHPRC
 Contract No. 33677
 Case no SDG H4639

QC BATCH	CHAIN OF CUSTODY	CLIENT SAMPLE ID	MATRIX	* SOLIDS	SAMPLE AMOUNT	BASIS AMOUNT	DAYS SINCE RECEIVED	LAB COLL	LAB SAMPLE ID	DEPARTMENT SAMPLE ID
7752	F11-127-063	B2FBT1	SOIL	93.2	133.0 g		08/01/11 5	S108007-01		7752-001
	F11-127-064	B2FBT2	SOIL	92.9	131.8 g		08/01/11 5	S108007-02		7752-002
	F11-127-065	B2FBT3	SOIL	93.8	162.6 g		08/01/11 5	S108007-03		7752-003
	F11-127-095	B2FPF6	SOIL	92.4	133.1 g		08/01/11 5	S108007-04		7752-004
	F11-127-096	B2FPF7	SOIL	93.2	157.8 g		08/01/11 5	S108007-05		7752-005
	F11-127-097	B2FPF8	SOIL	94.3	136.2 g		08/01/11 5	S108007-06		7752-006
		Method Blank	SOIL					S108007-08		7752-008
		Lab Control Sample	SOIL					S108007-07		7752-007
		Duplicate (S108007-01)	SOIL	93.2	133.0 g		08/01/11 5	S108007-09		7752-009

QC SUMMARY

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SUMMARY DATA SECTION

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EBERLINE ANALYTICAL/RICHMOND

SAMPLE DELIVERY GROUP H4639

SDG 7752
 Contact N. Joseph Verville

PREP BATCH SUMMARY

Client CHPRC
 Contract No. 33677
 Case no SDG H4639

TEST MATRIX	METHOD	PREPARATION ERROR		PLANCHETS ANALYZED				QUALI-		
		BATCH	2σ %	CLIENT	MORE	RE	BLANK		LCS	DUP/ORIG
Liquid Scintillation Counting										
H	SOIL	Tritium in Solids	7308-044	10.0	6		1	1	1/1	

Duplicates and Spikes are those with original sample in the QC Batch of some Client sample in this SDG.
 Blank and LCS planchets are those in the same preparation batch as some Client, Duplicate or Spike sample.

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EBERLINE ANALYTICAL/RICHMOND

SAMPLE DELIVERY GROUP H4639

SDG 7752
 Contact N. Joseph Verville

LAB WORK SUMMARY

Client CHPRC
 Contract No. 33677
 Case no SDG H4639

LAB SAMPLE	CLIENT SAMPLE ID	MATRIX	SUF-	REVIEWED BY	METHOD
COLLECTED	LOCATION		FIX	ANALYZED	
RECEIVED	CUSTODY	SAF No	PLANCHET	TEST	
S108007-01	B2FBT1		7752-001	H	08/04/11 08/08/11 BW Tritium in Solids
07/27/11	216-S-26 IP-1	SOIL			
08/01/11	F11-127-063	F11-127			
S108007-02	B2FBT2		7752-002	H	08/04/11 08/08/11 BW Tritium in Solids
07/27/11	216-S-26 IP-2	SOIL			
08/01/11	F11-127-064	F11-127			
S108007-03	B2FBT3		7752-003	H	08/04/11 08/08/11 BW Tritium in Solids
07/27/11	216-S-26 IP-3	SOIL			
08/01/11	F11-127-065	F11-127			
S108007-04	B2FPF6		7752-004	H	08/04/11 08/08/11 BW Tritium in Solids
07/27/11	216-S-26 IP-4	SOIL			
08/01/11	F11-127-095	F11-127			
S108007-05	B2FPF7		7752-005	H	08/04/11 08/08/11 BW Tritium in Solids
07/27/11	216-S-26 IP-5	SOIL			
08/01/11	F11-127-096	F11-127			
S108007-06	B2FPF8		7752-006	H	08/04/11 08/08/11 BW Tritium in Solids
07/27/11	216-S-26 IP-6	SOIL			
08/01/11	F11-127-097	F11-127			
S108007-07	Lab Control Sample	SOIL	7752-007	H	08/04/11 08/08/11 BW Tritium in Solids
		F11-127			
S108007-08	Method Blank	SOIL	7752-008	H	08/04/11 08/08/11 BW Tritium in Solids
		F11-127			
S108007-09	Duplicate (S108007-01)		7752-009	H	08/04/11 08/08/11 BW Tritium in Solids
07/27/11	216-S-26 IP-1	SOIL			
08/01/11		F11-127			

WORK SUMMARY

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SUMMARY DATA SECTION

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EBERLINE ANALYTICAL/RICHMOND

SAMPLE DELIVERY GROUP H4639

SDG 7752
 Contact N. Joseph Verville

WORK SUMMARY, cont.

Client CHPRC
 Contract No. 33677
 Case no SDG H4639

COUNTS OF TESTS BY SAMPLE TYPE											
TEST	SAF No	METHOD	REFERENCE	CLIENT	MORE	RE	BLANK	LCS	DUP	SPIKE	TOTAL
H	F11-127	Tritium in Solids	TRITIUM_COX_LSC	6			1	1	1		9
TOTALS				6			1	1	1		9

Lab id EBRLNE
 Protocol F11-127
 Version Ver 1.0
 Form DVD-LWS
 Version 3.06
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EBERLINE ANALYTICAL / RICHMOND
SAMPLE DELIVERY GROUP H4639

7752-008

Method Blank

METHOD BLANK

SDG <u>7752</u>	Client/Case no <u>CHPRC</u>	<u>SDG H4639</u>
Contact <u>N. Joseph Verville</u>	Contract <u>No. 33677</u>	
Lab sample id <u>S108007-08</u>	Client sample id <u>Method Blank</u>	
Dept sample id <u>7752-008</u>	Material/Matrix <u>SOIL</u>	
	SAF No <u>F11-127</u>	

ANALYTE	CAS NO	RESULT pCi/g	2 σ ERR (COUNT)	MDA pCi/g	RDL pCi/g	QUALI- FIERS	TEST
Tritium	10028-17-8	<u>-7.23</u>	6.8	12.2	30.0	U	H

QC-BLANK #79331

METHOD BLANKS

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Protocol <u>F11-127</u>
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Form <u>DVD-DS</u>
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Report date <u>08/08/11</u>

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SAMPLE DELIVERY GROUP H4639

7752-007

Lab Control Sample

LAB CONTROL SAMPLE

SDG <u>7752</u>	Client/Case no <u>CHPRC</u>	SDG <u>H4639</u>
Contact <u>N. Joseph Verville</u>	Contract No. <u>33677</u>	
Lab sample id <u>S108007-07</u>	Client sample id <u>Lab Control Sample</u>	
Dept sample id <u>7752-007</u>	Material/Matrix <u>SOIL</u>	
	SAF No <u>F11-127</u>	

ANALYTE	RESULT pCi/g	2σ ERR (COUNT)	MDA pCi/g	RDL pCi/g	QUALI- FIERS TEST	ADDED pCi/g	2σ ERR pCi/g	REC %	3σ LMTS (TOTAL)	PROTOCOL LIMITS
Tritium	373	17	14.0	30.0	H	409	16	91	84-116	80-120

QC-LCS #79330

Lab id <u>EBRLNE</u>
Protocol <u>F11-127</u>
Version <u>Ver 1.0</u>
Form <u>DVD-LCS</u>
Version <u>3.06</u>
Report date <u>08/08/11</u>

EBERLINE ANALYTICAL/RICHMOND

SAMPLE DELIVERY GROUP H4639

7752-009

B2FBT1

DUPLICATE

SDG <u>7752</u>	Client/Case no <u>CHPRC</u>	<u>SDG H4639</u>
Contact <u>N. Joseph Verville</u>	Contract No. <u>33677</u>	
<u>DUPLICATE</u>	<u>ORIGINAL</u>	
Lab sample id <u>S108007-09</u>	Lab sample id <u>S108007-01</u>	Client sample id <u>B2FBT1</u>
Dept sample id <u>7752-009</u>	Dept sample id <u>7752-001</u>	Location/Matrix <u>216-S-26 IP-1</u> <u>SOIL</u>
	Received <u>08/01/11</u>	Collected/Weight <u>07/27/11 14:40</u> <u>133.0 g</u>
% solids <u>93.2</u>	% solids <u>93.2</u>	Custody/SAF No <u>F11-127-063</u> <u>F11-127</u>

ANALYTE	DUPLICATE pCi/g	2σ ERR (COUNT)	MDA pCi/g	RDL pCi/g	QUALI- FIERS	TEST	ORIGINAL pCi/g	2σ ERR (COUNT)	MDA pCi/g	QUALI- FIERS	RPD %	3σ TOT	DER σ
Tritium	-1.11	7.0	12.1	30.0	U	H	-3.07	6.7	11.7	U	-		0.4

QC-DUP#1 79332

ARRA Verification Sampling of the 216-S-26 Waste Site
and 200-W-148-PL

Lab id EBRLNE
Protocol F11-127
Version Ver 1.0
Form DVD-DUP
Version 3.06
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EBERLINE ANALYTICAL / RICHMOND
 SAMPLE DELIVERY GROUP H4639

7752-001

B2FBT1

DATA SHEET

SDG <u>7752</u>	Client/Case no <u>CHPRC</u>	SDG <u>H4639</u>
Contact <u>N. Joseph Verville</u>	Contract No. <u>33677</u>	
Lab sample id <u>S108007-01</u>	Client sample id <u>B2FBT1</u>	
Dept sample id <u>7752-001</u>	Location/Matrix <u>216-S-26 IP-1</u>	<u>SOIL</u>
Received <u>08/01/11</u>	Collected/Weight <u>07/27/11 14:40</u>	<u>133.0 g</u>
% solids <u>93.2</u>	Custody/SAF No <u>F11-127-063</u>	<u>F11-127</u>

ANALYTE	CAS NO	RESULT pCi/g	2 σ ERR (COUNT)	MDA pCi/g	RDL pCi/g	QUALI- FIERS	TEST
Tritium	10028-17-8	-3.07	6.7	11.7	30.0	U	H

ARRA Verification Sampling of the 216-S-26 Waste Site
 and 200-W-148-PL

Lab id <u>EBRINE</u>
Protocol <u>F11-127</u>
Version <u>Ver 1.0</u>
Form <u>DVD-DS</u>
Version <u>3.06</u>
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EBERLINE ANALYTICAL / RICHMOND
SAMPLE DELIVERY GROUP H4639

7752-002

B2FBT2

DATA SHEET

SDG <u>7752</u>	Client/Case no <u>CHPRC</u>	<u>SDG H4639</u>
Contact <u>N. Joseph Verville</u>	Contract No. <u>33677</u>	
Lab sample id <u>S108007-02</u>	Client sample id <u>B2FBT2</u>	
Dept sample id <u>7752-002</u>	Location/Matrix <u>216-S-26 IP-2</u>	<u>SOIL</u>
Received <u>08/01/11</u>	Collected/Weight <u>07/27/11 14:44</u>	<u>131.8 g</u>
% solids <u>92.9</u>	Custody/SAF No <u>F11-127-064</u>	<u>F11-127</u>

ANALYTE	CAS NO	RESULT pCi/g	2σ ERR (COUNT)	MDA pCi/g	RDL pCi/g	QUALI- FIERS	TEST
Tritium	10028-17-8	16.0	8.1	12.7	30.0		H

ARRA Verification Sampling of the 216-S-26 Waste Site
and 200-W-148-PL

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EBERLINE ANALYTICAL / RICHMOND
 SAMPLE DELIVERY GROUP H4639

7752-003

B2FBT3

DATA SHEET

SDG <u>7752</u>	Client/Case no <u>CHPRC</u>	SDG <u>H4639</u>
Contact <u>N. Joseph Verville</u>	Contract No. <u>33677</u>	
Lab sample id <u>S108007-03</u>	Client sample id <u>B2FBT3</u>	
Dept sample id <u>7752-003</u>	Location/Matrix <u>216-S-26 IP-3</u>	<u>SOIL</u>
Received <u>08/01/11</u>	Collected/Weight <u>07/27/11 14:47</u>	<u>162.6 g</u>
% solids <u>93.8</u>	Custody/SAF No <u>F11-127-065</u>	<u>F11-127</u>

ANALYTE	CAS NO	RESULT pCi/g	2σ ERR (COUNT)	MDA pCi/g	RDL pCi/g	QUALI- FIERS	TEST
Tritium	10028-17-8	13.7	7.8	12.4	30.0		H

ARRA Verification Sampling of the 216-S-26 Waste Site
 and 200-W-148-PL

Lab id <u>EBRINE</u>
Protocol <u>F11-127</u>
Version <u>Ver 1.0</u>
Form <u>DVD-DS</u>
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Report date <u>08/08/11</u>

EBERLINE ANALYTICAL / RICHMOND
 SAMPLE DELIVERY GROUP H4639

7752-004

B2FPF6

DATA SHEET

SDG <u>7752</u>	Client/Case no <u>CHPRC</u>	SDG <u>H4639</u>
Contact <u>N. Joseph Verville</u>	Contract No. <u>33677</u>	
Lab sample id <u>S108007-04</u>	Client sample id <u>B2FPF6</u>	
Dept sample id <u>7752-004</u>	Location/Matrix <u>216-S-26 IP-4</u>	<u>SOIL</u>
Received <u>08/01/11</u>	Collected/Weight <u>07/27/11 15:03</u>	<u>133.1 g</u>
% solids <u>92.4</u>	Custody/SAF No <u>F11-127-095</u>	<u>F11-127</u>

ANALYTE	CAS NO	RESULT pCi/g	2σ ERR (COUNT)	MDA pCi/g	RDL pCi/g	QUALI- FIERS	TEST
Tritium	10028-17-8	-2.22	6.9	12.0	30.0	U	H

ARRA Verification Sampling of the 216-S-26 Waste Site
 and 200-W-148-PL

Lab id <u>EBRLNE</u>
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EBERLINE ANALYTICAL / RICHMOND
 SAMPLE DELIVERY GROUP H4639

7752-005

B2FPF7

DATA SHEET

SDG <u>7752</u>	Client/Case no <u>CHPRC</u>	SDG <u>H4639</u>
Contact <u>N. Joseph Verville</u>	Contract No. <u>33677</u>	
Lab sample id <u>S108007-05</u>	Client sample id <u>B2FPF7</u>	
Dept sample id <u>7752-005</u>	Location/Matrix <u>216-S-26 IP-5</u>	<u>SOIL</u>
Received <u>08/01/11</u>	Collected/Weight <u>07/27/11 15:08</u>	<u>157.8 g</u>
% solids <u>93.2</u>	Custody/SAF No <u>F11-127-096</u>	<u>F11-127</u>

ANALYTE	CAS NO	RESULT pCi/g	2σ ERR (COUNT)	MDA pCi/g	RDL pCi/g	QUALI- FIERS	TEST
Tritium	10028-17-8	2.94	6.7	11.2	30.0	U	H

ARRA Verification Sampling of the 216-S-26 Waste Site
 and 200-W-148-PL

Lab id <u>EBRLNE</u>
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Version <u>Ver 1.0</u>
Form <u>DVD-DS</u>
Version <u>3.06</u>
Report date <u>08/08/11</u>

EBERLINE ANALYTICAL / RICHMOND
SAMPLE DELIVERY GROUP H4639

7752-006

B2FPF8

DATA SHEET

SDG <u>7752</u>	Client/Case no <u>CHPRC</u>	SDG <u>H4639</u>
Contact <u>N. Joseph Verville</u>	Contract No. <u>33677</u>	
Lab sample id <u>S108007-06</u>	Client sample id <u>B2FPF8</u>	
Dept sample id <u>7752-006</u>	Location/Matrix <u>216-S-26 IP-6</u>	<u>SOIL</u>
Received <u>08/01/11</u>	Collected/Weight <u>07/27/11 15:12</u>	<u>136.2 g</u>
% solids <u>94.3</u>	Custody/SAF No <u>F11-127-097</u>	<u>F11-127</u>

ANALYTE	CAS NO	RESULT pCi/g	2σ ERR (COUNT)	MDA pCi/g	RDL pCi/g	QUALI- FIERS	TEST
Tritium	10028-17-8	<u>-14.4</u>	6.0	11.4	30.0	U	H

ARRA Verification Sampling of the 216-S-26 Waste Site
and 200-W-148-PL

Lab id <u>EBRLNE</u>
Protocol <u>F11-127</u>
Version <u>Ver 1.0</u>
Form <u>DVD-DS</u>
Version <u>3.06</u>
Report date <u>08/08/11</u>

EBERLINE ANALYTICAL/RICHMOND

SAMPLE DELIVERY GROUP H4639

Test H Matrix SOIL
SDG 7752
Contact N. Joseph Verville

LAB METHOD SUMMARY

TRITIUM IN SOLIDS

LIQUID SCINTILLATION COUNTING

Client CHPRC
Contract No. 33677
Contract SDG H4639

RESULTS

LAB	RAW	SUF-		
SAMPLE ID	TEST FIX	PLANCHET	CLIENT SAMPLE ID	Tritium
Preparation batch 7308-044				
S108007-01		7752-001	B2FBT1	U
S108007-02		7752-002	B2FBT2	16.0
S108007-03		7752-003	B2FBT3	13.7
S108007-04		7752-004	B2FPF6	U
S108007-05		7752-005	B2FPF7	U
S108007-06		7752-006	B2FPF8	U
S108007-07		7752-007	Lab Control Sample	ok
S108007-08		7752-008	Method Blank	U
S108007-09		7752-009	Duplicate (S108007-01)	- U

Nominal values and limits from method RDLs (pCi/g) 30.0

METHOD PERFORMANCE

LAB	RAW	SUF-	MDA	ALIQ	PREP	DILU-	YIELD	EFF	COUNT	FWHM	DRIFT	DAYS	ANAL-		
SAMPLE ID	TEST FIX	CLIENT SAMPLE ID	pCi/g	g	FAC	TION	%	%	min	keV	KeV	HELD	PREPARED	YZED	DETECTOR
Preparation batch 7308-044 2σ prep error 10.0 % Reference Lab Notebook 7308 pg. 044															
S108007-01		B2FBT1	11.7	0.213			100		50			8	08/04/11	08/04	LSC-006
S108007-02		B2FBT2	12.7	0.201			100		50			8	08/04/11	08/04	LSC-006
S108007-03		B2FBT3	12.4	0.215			100		50			8	08/04/11	08/04	LSC-006
S108007-04		B2FPF6	12.0	0.207			100		50			8	08/04/11	08/04	LSC-006
S108007-05		B2FPF7	11.2	0.219			100		50			8	08/04/11	08/04	LSC-006
S108007-06		B2FPF8	11.4	0.209			100		50			8	08/04/11	08/04	LSC-006
S108007-07		Lab Control Sample	14.0	0.201			100		50				08/04/11	08/04	LSC-006
S108007-08		Method Blank	12.2	0.201			100		50				08/04/11	08/04	LSC-006
S108007-09		Duplicate (S108007-01)	12.1	0.212			100		50			8	08/04/11	08/04	LSC-006

Nominal values and limits from method 30.0 0.201 50 180

PROCEDURES REFERENCE TRITIUM_COX_LSC
CP-251 Tritium/Carbon-14 Oxidation, rev 11

AVERAGES ± 2 SD MDA 12.2 ± 1.65
FOR 9 SAMPLES YIELD 100 ± 0

METHOD SUMMARIES

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SUMMARY DATA SECTION

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Lab id EBRLNE
Protocol F11-127
Version Ver 1.0
Form DVD-IMS
Version 3.06
Report date 08/08/11

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SAMPLE SUMMARY

The Sample and QC Summary Reports show all samples, including QC samples, reported in one Sample Delivery Group (SDG).

The Sample Summary Report fully identifies client samples and gives the corresponding lab sample identification. The QC Summary Report shows at the sample level how the lab organized the samples into batches and generated QC samples. The Preparation Batch and Method Summary Reports show this at the analysis level.

The following notes apply to these reports:

- * LAB SAMPLE ID is the lab's primary identification for a sample.
- * DEPARTMENT SAMPLE ID is an alternate lab id, for example one assigned by a radiochemistry department in a lab.
- * CLIENT SAMPLE ID is the client's primary identification for a sample. It includes any sample preparation done by the client that is necessary to identify the sample.
- * QC BATCH is a lab assigned code that groups samples to be processed and QCed together. These samples should have similar matrices.

QC BATCH is not necessarily the same as SDG, which reflects samples received and reported together.

- * All Lab Control Samples, Method Blanks, Duplicates and Matrix Spikes are shown that QC any of the samples. Due to possible reanalyses, not all results for all these QC samples may be relevant to the SDG. The Lab Control Sample, Method Blank, Duplicate, Matrix Spike and Method Summary Reports detail these relationships.

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Form DVD-RG
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PREPARATION BATCH SUMMARY

The Preparation Batch Summary Report shows all preparation batches in one Sample Delivery Group (SDG) with information necessary to check the completeness and consistency of the SDG.

The following notes apply to this report:

- * The preparation batches are shown in the same order as the Method Summary Reports are printed.
- * Only analyses of plachets relevant to the SDG are included.
- * Each preparation batch should have at least one Method Blank and LCS in it to validate client sample results.
- * The QUALIFIERS shown are all qualifiers other than U, J, B, L and H that occur on any analysis in the preparation batch. The Method Summary Report has these qualifiers on a per sample basis.

These qualifiers should be reviewed as follows:

- X Some data has been manually entered or modified. Transcription errors are possible.
- P One or more results are 'preliminary'. The data is not ready for final reporting.
- 2 There were two or more results for one analyte on one plachet imported at one time. The results in DVD may not be the same as on the raw data sheets.

Other lab defined qualifiers may occur. In general, these should be addressed in the SDG narrative.

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WORK SUMMARY

The Work Summary Report shows all samples, including QC samples, and all relevant analyses in one Sample Delivery Group (SDG). This report is often useful as supporting documentation for an invoice.

The following notes apply to this report:

- * TEST is a code for the method used to measure associated analytes. Results and related information for each analyte are on the Data Sheet Report. In special cases, a test code used in the summary data section is not the same as in associated raw data. In this case, both codes are shown on the Work Summary.
- * SUFFIX is the lab's code to distinguish multiple analyses (recounts, reworks, reanalyses) of a fraction of the sample. The suffix indicates which result is being reported. An empty suffix normally identifies the first attempt to analyze the sample.
- * The LAB SAMPLE ID, TEST and SUFFIX uniquely identify all supporting data for a result. The Method Summary Report for each TEST has method performance data, such as yield, for each lab sample id and suffix and procedures used in the method.
- * PLANCHET is an alternate lab identifier for work done for one test. It, combined with the TEST and SUFFIX, may be the best link to raw data.
- * For QC samples, only analyses that directly QC some regular sample are shown. The Lab Control Sample, Method Blank, Duplicate, Matrix Spike and Method Summary Reports detail these relationships.
- * The SAS (Special Analytical Services) Number is a client or lab assigned code that reflects special processing for samples, such as rapid turn around. Counts of tests done are lists by SAS number since it is likely to affect prices.

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DATA SHEET

The Data Sheet Report shows all results and primary supporting information for one client sample or Method Blank. This report corresponds to both the CLP Inorganics and Organics Data Sheet.

The following notes apply to this report:

- * TEST is a code for the method used to measure an analyte. If the TEST is empty, no data is available; the analyte was not analyzed for.
- * The LAB SAMPLE ID and TEST uniquely identify work within the Summary Data Section of a Data Package. The Work Summary and Method Summary Reports further identify raw data that underlies this work.

The Method Summary Report for each TEST has method performance data, such as yield, for each Lab Sample ID and a list of procedures used in the method.

- * ERRORS can be labeled TOTAL or COUNT. TOTAL implies a preparation (non-counting method) error has been added, as square root of sum of squares, to the counting error denoted by COUNT. The preparation errors, which may vary by preparation batch, are shown on the Method Summary Report.
- * A RESULT can be 'N.R.' (Not Reported). This means the lab did this work but chooses not to report it now, possibly because it was reported at another time.
- * When reporting a Method Blank, a RESULT can be 'N.A.' (Not Applicable). This means there is no reported client sample work in the same preparation batch as the Blank's result. This is likely to occur when the Method Blank is associated with reanalyses of selected work for a few samples in the SDG.

The following qualifiers are defined by the DVD system:

- U The RESULT is less than the MDA (Minimum Detectable Activity).

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DATA SHEET

If the MDA is blank, the ERROR is used as the limit.

J The RESULT is less than the RDL (Required Detection Limit) and no U qualifier is assigned.

B A Method Blank associated with this sample had a result without a U flag and, after correcting for possibly different aliquots, that result is greater than or equal to the MDA for this sample.

Normally, B is not assigned if U is. When method blank subtraction is shown on this report, B flags are assigned based on the unsubtracted values while U's are assigned based on the subtracted ones. Both flags can be assigned in this case.

For each sample result, all Method Blank results in the same preparation batch are compared. The Method Summary Report documents this and other QC relationships.

L Some Lab Control Sample that QC's this sample had a low recovery. The lab can disable assignment of this qualifier.

H Similar to 'L' except the recovery was high.

P The RESULT is 'preliminary'.

X Some data necessary to compute the RESULT, ERROR or MDA was manually entered or modified.

2 There were two or more results available for this analyte. The reported result may not be the same as in the raw data.

Other qualifiers are lab defined. Definitions should be in the SDG narrative.

The following values are underlined to indicate possible problems:

* An MDA is underlined if it is bigger than its RDL.

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DATA SHEET

- * An ERROR is underlined if the 1.645 sigma counting error is bigger than both the MDA and the RESULT, implying that the MDA may not be a good estimate of the 'real' minimum detectable activity.
- * A negative RESULT is underlined if it is less than the negative of its 2 sigma counting ERROR.
- * When reporting a Method Blank, a RESULT is underlined if greater than its MDA. If the MDA is blank, the 2 sigma counting error is used in the comparison.

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LAB CONTROL SAMPLE

The Lab Control Sample Report shows all results, recoveries and primary supporting information for one Lab Control Sample.

The following notes apply to this report:

- * All fields in common with the Data Sheet Report have similar usage. Refer to its Report Guide for details.
- * An amount ADDED is the lab's value for the actual amount spiked into this sample with its ERROR an estimate of the error of this amount.

An amount added is underlined if its ratio to the corresponding RDL is outside protocol specified limits.

- * REC (Recovery) is RESULT divided by ADDED expressed as a percent.
- * The first, computed limits for the recovery reflect:
 1. The error of RESULT, including that introduced by rounding the result prior to printing.

If the limits are labeled (TOTAL), they include preparation error in the result. If labeled (COUNT), they do not.
 2. The error of ADDED.
 3. A lab specified, per analyte bias. The bias changes the center of the computed limits.
- * The second limits are protocol defined upper and lower QC limits for the recovery.
- * The recovery is underlined if it is outside either of these ranges.

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DUPLICATE

The Duplicate Report shows all results, differences and primary supporting information for one Duplicate and associated Original sample.

The following notes apply to this report:

- * All fields in common with the Data Sheet Report have similar usage. This applies both to the Duplicate and Original sample data. Refer to the Data Sheet Report Guide for details.

If the Duplicate has data for a TEST and the lab did not do this test to the Original, the Original's RESULTS are underlined.

- * The RPD (Relative Percent Difference) is the absolute value of the difference of the RESULTS divided by their average expressed as a percent.

If both RESULTS are less than their MDAs, no RPD is computed and a '-' is printed.

For an analyte, if the lab did work for both samples but has data for only one, the MDA from the sample with data is used as the other's result in the RPD.

- * The first, computed limit is the sum, as square root of sum of squares, of the errors of the results divided by the average result as a percent, hence the relative error of the difference rather than the error of the relative difference. The errors include those introduced by rounding the RESULTS prior to printing.

If this limit is labeled TOT, it includes the preparation error in the RESULTS. If labeled CNT, it does not.

This value reported for this limit is at most 999.

- * The second limit for the RPD is the larger of:
 1. A fixed percentage specified in the protocol.

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DUPLICATE

2. A protocol factor (typically 2) times the average MDA as a percent of the average result. This limit applies when the results are close to the MDAs.

- * The RPD is underlined if it is greater than either limit.
- * If specified by the lab, the second limit column is replaced by the Difference Error Ratio (DER), which is the absolute value of the difference of the results divided by the quadratic sum of their one sigma errors, the same errors as used in the first limit.

Except for differences due to rounding, the DER is the same as the RPD divided by the first RPD limit with the limit scaled to 1 sigma.

- * The DER is underlined if it is greater than the sigma factor, typically 2 or 3, shown in the header for the first RPD limit..

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MATRIX SPIKE

The Matrix Spike Report shows all results, recoveries and primary supporting information for one Matrix Spike and associated Original sample.

The following notes apply to this report:

- * All fields in common with the Data Sheet Report have similar usage. This applies both to the Spiked and Original sample data. Refer to the Data Sheet Report Guide for details.

If the Spike has data for a TEST and the lab did not do this test to the Original, the Original's RESULTS are underlined.

- * An amount ADDED is the lab's value for the actual amount spiked into the Spike sample with its ERROR an estimate of the error of this amount.

An amount is underlined if its ratio to the corresponding RDL is outside protocol specified limits.

- * REC (Recovery) is the Spike RESULT minus the Original RESULT divided by ADDED expressed as a percent.

- * The first, computed limits for the recovery reflect:

1. The errors of the two RESULTS, including those introduced by rounding them prior to printing.

If the limits are labeled (TOTAL), they include preparation error in the result. If labeled (COUNT), they do not.

2. The error of ADDED.

3. A lab specified, per analyte bias. The bias changes the center of the computed limits.

- * The second limits are protocol defined upper and lower QC limits

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MATRIX SPIKE

for the recovery.

These limits are left blank if the Original RESULT is more than a protocol defined factor (typically 4) times ADDED. This is a way of accounting for that when the spike is small compared to the amount in the original sample, the recovery is unreliable.

* The recovery is underlined (out of spec) if it is outside either of these ranges.

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METHOD SUMMARY

The Method Summary Report has two tables. One shows up to five results measured using one method. The other has performance data for the method. There is one report for each TEST, as used on the Data Sheet Report.

The following notes apply to this report:

- * Each table is subdivided into sections, one for each preparation batch. A preparation batch is a group of aliquots prepared at roughly the same time in one work area of the lab using the same method.

There should be Lab Control Sample and Method Blank results in each preparation batch since this close correspondence makes the QC meaningful. Depending on lab policy, Duplicates need not occur in each batch since they QC sample dependencies such as matrix effects.

- * The RAW TEST column shows the test code used in the raw data to identify a particular analysis if it is different than the test code in the header of the report. This occurs in special cases due to method specific details about how the lab labels work.

The Lab Sample or Planchet ID combined with the (Raw) Test Code and Suffix uniquely identify the raw data for each analysis.

- * If a result is less than both its MDA and RDL, it is replaced by just 'U' on this report. If it is greater than or equal to the RDL but less than the MDA, the result is shown with a 'U' flag.

The J and X flags are as on the data sheet.

- * Non-U results for Method Blanks are underlined to indicate possible contamination of other samples in the preparation batch. The Method Blank Report has supporting data.
- * Lab Control Sample and Matrix Spike results are shown as: ok, No data, LOW or HIGH, with the last two underlined. 'No data'

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means no amount ADDED was specified. 'LOW' and 'HIGH' correspond to when the recovery is underlined on the Lab Control Sample or Matrix Spike Report. See these reports for supporting data.

- * Duplicate sample results are shown as: ok, No data, or OUT, with the last two underlined. 'No data' means there was no original sample data found for this duplicate. 'OUT' corresponds to when the RPD is underlined on the Duplicate Report. See this report for supporting data.
 - * If the MDA column is labeled 'MAX MDA', there was more than one result measured by the reported method and the MDA shown is the largest MDA. If not all these results have the same RDL, the MAX MDA reflects only those results with RDL equal to the smallest one.
- MDAs are underlined if greater than the printed RDL.
- * Aliquots are underlined if less than the nominal value specified for the method.
 - * Preparation factors are underlined if greater than the nominal value specified for the method.
 - * Dilution factors are underlined if greater than the nominal value specified for the method.
 - * Residues are underlined if outside the range specified for the method. Residues are not printed if yields are.
 - * Yields, which may be gravimetric, radiometric or some type of recovery depending on the method, are underlined if outside the range specified for the method.
 - * Efficiencies are underlined if outside the range specified for the method. Efficiencies are detector and geometry dependent so this test is only approximate.

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- * Count times are underlined if less than the nominal value specified for the method.
- * Resolutions (as FWHM; Full Width at Half Max) are underlined if greater than the method specified limit.
- * Tracer drifts are underlined if their absolute values are greater than the method specified limit. Tracer drifts are not printed if percent moistures are.
- * Days Held are underlined if greater than the holding time specified in the protocol.
- * Analysis dates are underlined if before their planchet's preparation date or, if a limit is specified, too far after it.

For some methods, ratios as percentages and error estimates for them are computed for pairs of results. A ratio column header like '1+3' means the ratio of the first result column and the third result column.

Ratios are not computed for Lab Control Sample, Method Blank or Matrix Spike results since their matrices are not necessarily similar to client samples'.

The error estimate for a ratio of results from one planchet reflects only counting errors since other errors should be correlated. For a ratio involving different planchets, if QC limits are computed based on total errors, the error for the ratio allows for the preparation errors for the planchets.

The ratio is underlined (out of spec) if the absolute value of its difference from the nominal value is greater than its error estimate. If no nominal value is specified, this test is not done.

For Gross Alpha or Gross Beta results, there may be a column showing the sum of other Alpha or Beta emitters. This sum includes all relevant

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results in the DVD database, whether reported or not. Results in the sum are weighted by a particles/decay value specified by the lab for each relevant analyte. Results less than their MDA are not included. No sums are computed for Lab Control, Method Blank or Matrix Spike samples since their various planchets may not be physically related.

If a ratio of total isotopic to Gross Alpha or Beta is shown, the error for the ratio reflects both the error in the Gross result and the sum, as square root of sum of squares, of the errors in the isotopic results.

For total elemental uranium or thorium results, there may be a column showing the total weight computed from associated isotopic results. Ignoring results less than their MDAs, this is a weighted sum of the isotopic results. The weights depend on the molecular weight and half-life of each isotope so as to convert activities (decays) to weight (atoms).

If a ratio of total computed to measured elemental uranium or thorium is shown, the error for the ratio reflects the errors in all the measurements.

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COLLECTOR
Chamberlain

COMPANY CONTACT
BAUER, RG **14639**

TELEPHONE NO.
373-3931 **(7752)**

PROJECT COORDINATOR
BAUER, RG

PRICE CODE
8B

DATA TURNAROUND
7 Days / 15 Days

SAMPLING LOCATION
216-S-26 1p-2

PROJECT DESIGNATION
ARRA Verification Sampling of the 216-S-26 Waste Site and 200-W-148-PL

SAFE NO.
F11-127

AIR QUALITY

METHOD OF SHIPMENT
FEDERAL EXPRESS

ICE CHEST NO.

FIELD LOGBOOK NO.
HNF-N-507-19-31

COA
30222ES10

METHOD OF SHIPMENT
FEDERAL EXPRESS

ORIGINAL

SHIPPED TO
Eberline Services

OFFSITE PROPERTY NO.
SEE PTR **2903**

BILL OF LADING/AIR BILL NO.
SEE PTR **7973 5379 3581**

MATRIX*
A=Air
DL=Drum
Liquids
DS=Drum
Solids
L=Liquid
O=Oil
S=Soil
SE=Sediment
T=Tissue
V=Vegetation
W=Water
WI=Wipe
X=Other

POSSIBLE SAMPLE HAZARDS/ REMARKS
Contains Radioactive Material at concentrations that may or may not be regulated for transportation per 49 CFR / IATA Dangerous Goods Regulations but are not releasable per DOE Order 5400.5 (1990/1993)

PRESERVATION
None

HOLDING TIME
6 Months

TYPE OF CONTAINER
G/P

NO. OF CONTAINERS(S)
1

SPECIAL HANDLING AND/OR STORAGE

SAMPLE ANALYSIS
Tritium - H3 (Tribium);

VOLUME
120ml

SAMPLE NO.	MATRIX*	SAMPLE DATE	SAMPLE TIME
B2FBT2	SOIL	JUL 27 2011	1444 ✓

CHAIN OF POSSESSION

SIGN/ PRINT NAMES

SPECIAL INSTRUCTIONS

RELINQUISHED BY/REMOVED FROM	DATE/TIME	RECEIVED BY/STORED IN	DATE/TIME
RELINQUISHED BY/REMOVED FROM <i>Chamberlain</i>	JUL 27 2011	RECEIVED BY/STORED IN <i>SAF #1</i>	JUL 27 2011
RELINQUISHED BY/REMOVED FROM <i>SSW</i>	JUL 28 2011	RECEIVED BY/STORED IN KC Patternson	JUL 28 2011
RELINQUISHED BY/REMOVED FROM KC Patternson	JUL 28 2011	RECEIVED BY/STORED IN FEDER	JUL 28 2011
RELINQUISHED BY/REMOVED FROM	DATE/TIME	RECEIVED BY/STORED IN	DATE/TIME
RELINQUISHED BY/REMOVED FROM	DATE/TIME	RECEIVED BY/STORED IN	DATE/TIME
RELINQUISHED BY/REMOVED FROM	DATE/TIME	RECEIVED BY/STORED IN	DATE/TIME
RELINQUISHED BY/REMOVED FROM	DATE/TIME	RECEIVED BY/STORED IN	DATE/TIME
RELINQUISHED BY/REMOVED FROM	DATE/TIME	RECEIVED BY/STORED IN	DATE/TIME

** The 200 Area S&GRP Characterization and Monitoring Sampling and Analysis GKI applies to this SAF.

** The laboratory shall achieve a RDL of 30 pCi/g for Tritium.

** The laboratory shall report only the requested constituents of concern in all final data reports, and EDDs. QC results for non-requested constituents need not be reported, but samples chosen for matrix spike, and all spike compounds shall be identified in the final data report case narrative. No TICs shall be reported.

LABORATORY SECTION	RECEIVED BY	DATE/TIME	TITLE	DATE/TIME
FINAL SAMPLE DISPOSITION	DISPOSAL METHOD			

COLLECTOR <i>C. Chamberlain</i>	COMPANY CONTACT BAUER, RG	TELEPHONE NO. 373-3931	PROJECT COORDINATOR BAUER, RG	F11-127-065	PAGE 1 OF 1
SAMPLING LOCATION 216-S-26 JP-3	PROJECT DESIGNATION ARRA Verification Sampling of the 216-S-26 Waste Site and 200-W-148-PL	FIELD LOGBOOK NO. HNF-N-507-19-31	SAF NO. F11-127	PRICE CODE 88	DATA TURNAROUND 7 Days / 15 Days
ICE CHEST NO.	ACTUAL SAMPLE DEPTH 20 FT.	COA 3022225510	METHOD OF SHIPMENT FEDERAL EXPRESS	ORIGINAL	

SHIPPED TO Eberline Services	OFFSITE PROPERTY NO. SEE PTR	2963	BILL OF LADING/AIR BILL NO. SEE PTR	7978	5374	3581
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MATRIX* A=Air DL=Drum Liquids DS=Drum Solids L=Liquid O=Oil S=Sediment SE=Seepage T=Tissue V=Vegetation W=Water WI=Wipe X=Other	POSSIBLE SAMPLE HAZARDS/ REMARKS Contains Radioactive Material at concentrations that may or may not be regulated for transportation per 49 CFR / IATA Dangerous Goods Regulations but are not releasable per DOE Order 5400.5 (1990/1993)	PRESERVATION	None
		HOLDING TIME	6 Months
		TYPE OF CONTAINER	G/P
		NO. OF CONTAINER(S)	1
		VOLUME	120mL

SPECIAL HANDLING AND/OR STORAGE	SAMPLE ANALYSIS	Tritium - H3 (Tritium);
SAMPLE NO.	MATRIX*	
B2FBT3	SOIL	
SAMPLE DATE	SAMPLE TIME	
JUL 27 2011	1447	✓

CHAIN OF POSSESSION	SIGN/ PRINT NAMES	SPECIAL INSTRUCTIONS
REINQUISHED BY/REMOVED FROM <i>Chamberlain</i>	RECEIVED BY/STORED IN SQL# 1	<p>SPECIAL INSTRUCTIONS</p> <p>** The 200 Area S&GRP Characterization and Monitoring Sampling and Analysis GKI applies to this SAF.</p> <p>** The laboratory shall achieve a RDL of 30 pCi/g for Tritium.</p> <p>** The laboratory shall report only the requested constituents of concern in all final data reports, and EDDS. QC results for non-requested constituents need not be reported, but samples chosen for matrix spike, and all spike compounds shall be identified in the final data report case narrative. No TICs shall be reported.</p>
DATE/TIME JUL 27 2011	DATE/TIME JUL 27 2011	
REINQUISHED BY/REMOVED FROM SSU-1	RECEIVED BY/STORED IN KC Partnership	
DATE/TIME JUL 28 2011	DATE/TIME JUL 28 2011	
REINQUISHED BY/REMOVED FROM KC Partnership	RECEIVED BY/STORED IN FEDERATION	
DATE/TIME JUL 28 2011	DATE/TIME JUL 28 2011	
REINQUISHED BY/REMOVED FROM FEDERATION	RECEIVED BY/STORED IN FE WATERS	
DATE/TIME JUL 28 2011	DATE/TIME JUL 28 2011	
REINQUISHED BY/REMOVED FROM	RECEIVED BY/STORED IN	
DATE/TIME	DATE/TIME	
REINQUISHED BY/REMOVED FROM	RECEIVED BY/STORED IN	
DATE/TIME	DATE/TIME	

LABORATORY SECTION	RECEIVED BY	DATE/TIME
FINAL SAMPLE DISPOSITION	DISPOSAL METHOD	DATE/TIME

COLLECTOR
C. Chamberlain

SAMPLING LOCATION
216-S-26 IP-4

COMPANY CONTACT
BAUER, RG

PROJECT DESIGNATION
ARRA Verification Sampling of the 216-S-26 Waste Site and 200-W-148-PL

TELEPHONE NO.
373-3931

PROJECT COORDINATOR
BAUER, RG

PRICE CODE
8B

DATA TURNAROUND
7 Days / 15 Days

ICE CHEST NO.

FIELD LOGBOOK NO.
HNF-N-507-19-34

ACTUAL SAMPLE DEPTH
16 ft.

SAF NO.
F11-127

COA
302222ES10

METHOD OF SHIPMENT
FEDERAL EXPRESS

ORIGINAL

SHIPPED TO
Eberline Services

OFFSITE PROPERTY NO.
SEE PTR

PRESERVATION
None

BILL OF LADING/AIR BILL NO.
7973

5379

3581

MATRIX*
A=Air
DL=Drum
Liquids
DS=Drum
Solids
L=Liquid
O=Oil
S=Soil
SF=Sediment
T=Tissue
V=Vegetation
W=Water
WI=Wipe
X=Other

POSSIBLE SAMPLE HAZARDS/ REMARKS
Contains Radioactive Material at concentrations that may or may not be regulated for transportation per 49 CFR / IATA Dangerous Goods Regulations but are not releasable per DOE Order 5400.5 (1990/1993)

HOLDING TIME
6 Months

TYPE OF CONTAINER
G/P

NO. OF CONTAINERS(S)
1

VOLUME
120ml

SPECIAL HANDLING AND/OR STORAGE

SAMPLE ANALYSIS
Tritium - H3 (Tridium);

SAMPLE NO. B2FPF6

MATRIX* SOIL

SAMPLE DATE
JUL 27 2011

SAMPLE TIME
1503

✓

CHAIN OF POSSESSION

SIGN/ PRINT NAMES

RELINQUISHED BY/REMOVED FROM
C. Chamberlain

RECEIVED BY/STORED IN
SSU

DATE/TIME
JUL 27 2011

DATE/TIME
JUL 27 2011

RELINQUISHED BY/REMOVED FROM
SSU

RECEIVED BY/STORED IN
KC Patterson

DATE/TIME
JUL 28 2011

DATE/TIME
JUL 28 2011

RELINQUISHED BY/REMOVED FROM
KC Patterson

RECEIVED BY/STORED IN
FEDEX

DATE/TIME
JUL 28 2011

DATE/TIME
JUL 28 2011

RELINQUISHED BY/REMOVED FROM

RECEIVED BY/STORED IN

DATE/TIME

DATE/TIME

RELINQUISHED BY/REMOVED FROM

RECEIVED BY/STORED IN

DATE/TIME

DATE/TIME

RELINQUISHED BY/REMOVED FROM

RECEIVED BY/STORED IN

DATE/TIME

DATE/TIME

LABORATORY SECTION

RECEIVED BY

DATE/TIME

DATE/TIME

FINAL SAMPLE DISPOSITION

DISPOSAL METHOD

DATE/TIME

DATE/TIME

SPECIAL INSTRUCTIONS

** The 200 Area S&GRP Characterization and Monitoring Sampling and Analysis GKI applies to this SAF.
 ** The laboratory shall achieve a RDL of 30 pCi/g for Tritium.
 ** The laboratory shall report only the requested constituents of concern in all final data reports, and EDDs. QC results for non-requested constituents need not be reported, but samples chosen for matrix spike, and all spike compounds shall be identified in the final data report case narrative. No TICs shall be reported.

COLLECTOR <i>C. Chamberlain</i>	COMPANY CONTACT BAUER, RG	TELEPHONE NO. 373-3931	PROJECT COORDINATOR BAUER, RG	F11-127-096	PRICE CODE 8B	DATA TURNAROUND 7 Days / 15 Days
SAMPLING LOCATION 216-S-26 1P-5	PROJECT DESIGNATION ARRA Verification Sampling of the 216-S-26 Waste Site and 200-W-148-PL	FIELD LOGBOOK NO. HNF-N-507-B.351	SAF NO. F11-127	F11-127-096	AIR QUALITY <input type="checkbox"/>	METHOD OF SHIPMENT FEDERAL EXPRESS
ICE CHEST NO.	ACTUAL SAMPLE DEPTH 18 ft	COA 30222ES10	BILL OF LADING/AIR BILL NO. SEE PTR	7973	5379	5581
SHIPPED TO Eberline Services	OFFSITE PROPERTY NO. SEE PTR	NO. OF CONTAINER(S) 1	SEE PTR	ORIGINAL		

MATRIX* A=Air L=Liquids DS=Drum S=Solids O=Oil SE=Sediment T=Tissue V=Vegetation W=Water WI=Wipe X=Other	POSSIBLE SAMPLE HAZARDS/ REMARKS Contains Radioactive Material at concentrations that may or may not be regulated for transportation per 49 CFR / IATA Dangerous Goods Regulations but are not releasable per DOE Order 5400.5 (1990/1993)	PRESERVATION None	HOLDING TIME 6 Months	TYPE OF CONTAINER G/P	VOLUME 120mL	SPECIAL HANDLING AND/OR STORAGE Tritium - H3 (Tritium)
SAMPLE NO. B2FPF7	MATRIX* SOIL	SAMPLE DATE JUL 27 2011	SAMPLE TIME 1508	✓		

CHAIN OF POSSESSION	SIGN/ PRINT NAMES	DATE/TIME	DATE/TIME	SPECIAL INSTRUCTIONS
RELINQUISHED BY/REMOVED FROM <i>C. Chamberlain</i>	RECEIVED BY/STORED IN <i>SSU-1</i>	JUL 27 2011	JUL 27 2011	** The 200 Area S&GRP Characterization and Monitoring Sampling and Analysis GKI applies to this SAF.
RELINQUISHED BY/REMOVED FROM <i>SSU-1</i>	RECEIVED BY/STORED IN <i>SSU-1</i>	JUL 28 2011	JUL 28 2011	<input type="checkbox"/> ** The laboratory shall achieve a RDL of 30 pCi/g for Tritium.
RELINQUISHED BY/REMOVED FROM <i>KC Patrao</i>	RECEIVED BY/STORED IN <i>FEDEK</i>	JUL 28 2011	JUL 28 2011	<input type="checkbox"/> ** The laboratory shall report only the requested constituents of concern in all final data reports, and EDDs. QC results for non-requested constituents need not be reported, but samples chosen for matrix spike, and all spike compounds shall be identified in the final data report case narrative. No TICs shall be reported.
RELINQUISHED BY/REMOVED FROM <i>CHPRC</i>	RECEIVED BY/STORED IN <i>FE-LEATHRYAN</i>	JUL 28 2011	JUL 28 2011	
RELINQUISHED BY/REMOVED FROM	RECEIVED BY/STORED IN	DATE/TIME	DATE/TIME	
RELINQUISHED BY/REMOVED FROM	RECEIVED BY/STORED IN	DATE/TIME	DATE/TIME	
RELINQUISHED BY/REMOVED FROM	RECEIVED BY/STORED IN	DATE/TIME	DATE/TIME	
LABORATORY SECTION	RECEIVED BY	DATE/TIME	DATE/TIME	TITLE
FINAL SAMPLE DISPOSITION	DISPOSAL METHOD	DATE/TIME	DATE/TIME	DISPOSED BY

COLLECTOR <i>C. Chamberlain</i>	COMPANY CONTACT BAUER, RG	TELEPHONE NO. 4639 373-3931	PROJECT COORDINATOR BAUER, RG	PRICE CODE 8B	DATA TURNAROUND 7 Days / 15 Days
SAMPLING LOCATION 216-S-26 Jp-6	PROJECT DESIGNATION ARRA Verification Sampling of the 216-S-26 Waste Site and 200-W-148-PL	FIELD LOGBOOK NO. HNF-N-507-19-24	SAF NO. F11-127	AIR QUALITY <input type="checkbox"/>	
ICE CHEST NO.	ACTUAL SAMPLE DEPTH 20 ft.	COA 302222ES10	METHOD OF SHIPMENT FEDERAL EXPRESS	ORIGINAL	

SHIPPED TO Eberline Services	OFFSITE PROPERTY NO. SEE PTR 2903	BILL OF LADING/AIR BILL NO. SEE PTR 7973	5374	3581
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MATRIX*	POSSIBLE SAMPLE HAZARDS/ REMARKS	PRESERVATION	HOLDING TIME	TYPE OF CONTAINER	NO. OF CONTAINER(S)	VOLUME	SPECIAL HANDLING AND/OR STORAGE	SAMPLE NO.	MATRIX*	SAMPLE DATE	SAMPLE TIME
A=Air DL=Drum Liquids DS=Drum Solids L=Liquid O=Oil S=Soil SE=Sediment T=Tissue V=Vegetation W=Water WI=Wipe X=Other	Contains Radioactive Material at concentrations that may or may not be regulated for transportation per 49 CFR / IATA Dangerous Goods Regulations but are not releasable per DOE Order 5400.5 (1990/1993)	None	6 Months	G/P	1	120mL		B2PFF8	SOIL	JUL 27 2011	1512

CHAIN OF POSSESSION	SIGN/ PRINT NAMES	SPECIAL INSTRUCTIONS
RELINQUISHED BY/REMOVED FROM <i>C. Chamberlain</i>	RECEIVED BY/STORED IN <i>SSU#1</i>	** The 200 Area S&GRP Characterization and Monitoring Sampling and Analysis GKI applies to this SAF.
DATE/TIME JUL 27 2011	DATE/TIME JUL 27 2011	
RELINQUISHED BY/REMOVED FROM <i>SSU#1</i>	RECEIVED BY/STORED IN KC Patterson	<input type="checkbox"/> ** The laboratory shall achieve a RDL of 30 pCi/g for Tritium.
DATE/TIME JUL 28 2011/103	DATE/TIME JUL 28 2011/103	<input type="checkbox"/> ** The laboratory shall achieve a RDL of 30 pCi/g for Tritium.
RELINQUISHED BY/REMOVED FROM CHPRC	RECEIVED BY/STORED IN EDDER	** The laboratory shall report only the requested constituents of concern in all final data reports, and EDDs. QC results for non-requested constituents need not be reported, but samples chosen for matrix spike, and all spike compounds shall be identified in the final data report case narrative. No TICs shall be reported.
DATE/TIME JUL 28 2011/103	DATE/TIME JUL 28 2011/103	
RELINQUISHED BY/REMOVED FROM	RECEIVED BY/STORED IN	
DATE/TIME	DATE/TIME	
RELINQUISHED BY/REMOVED FROM	RECEIVED BY/STORED IN	
DATE/TIME	DATE/TIME	
LABORATORY SECTION	RECEIVED BY	TITLE
FINAL SAMPLE DISPOSITION	DISPOSAL METHOD	DATE/TIME



RICHMOND, CA LABORATORY
SAMPLE RECEIPT CHECKLIST

7752

H4639

Client: CHPRC City RICHMOND State WA
 Date/Time received 08/01/11 0855 CoC No. FM-127-063,064,065,095,096,097
 Container I.D. No. GWS-003 Requested TAT (Days) 7/15 P.O. Received Yes [] No []

INSPECTION

1. Custody seals on shipping container intact? Yes No [] N/A []
2. Custody seals on shipping container dated & signed? Yes No [] N/A []
3. Custody seals on sample containers intact? Yes No [] N/A []
4. Custody seals on sample containers dated & signed? Yes No [] N/A []
5. Packing material is: Wet [] Dry
6. Number of samples in shipping container: 6 Sample Matrix S
7. Number of containers per sample: 1 (Or see CoC _____)
8. Samples are in correct container Yes No []
9. Paperwork agrees with samples? Yes No []
10. Samples have: Tape [] Hazard labels [] Rad labels [] Appropriate sample labels
11. Samples are: In good condition Leaking [] Broken Container [] Missing []
12. Samples are: Preserved [] Not preserved [] pH N/A Preservative _____
13. Describe any anomalies:

14. Was P.M. notified of any anomalies? Yes [] No [] Date _____
15. Inspected by Tracy Date: 08/01/11 Time: 0930

Customer Sample No.	Beta/Gamma com	Ion Chamber mR/hr	Wide	Customer Sample No.	Beta/Gamma com	Ion Chamber mR/hr	Wide
<u>All Sources</u>	<u>LSD</u>						

Ion Chamber Ser. No. _____
 Alpha Meter Ser. No. _____
 Beta/Gamma Meter Ser. No. 100482

Calibration date _____
 Calibration date _____
 Calibration date 24 SEP 10