



# EBERLINE SERVICES

NOVEMBER 21, 2011

**REV 1**  
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November 3, 2011

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

Reference: **P.O. #33677**  
**Eberline Analytical S1-10-130-7615, SDG H4688**

Dear Mr. Fitzgerald:

Enclosed is a data report for one water sample designated under SAF No. F11-149 received at Eberline Analytical on October 25, 2011. The samples were analyzed according to the accompanying chain-of-custody documents.

Please call if you have any questions concerning this report.

Sincerely,

Joseph Verville  
Client Services Manager

*NJV/kwp*

*Enclosure: Case Narrative*

Eberline Analytical  
Report S1-10-130-7615  
November 3, 2011

CH2M Hill Plateau Remediation Company  
SDG H4688

Case Narrative

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## 1.0 GENERAL

CH2M Hill Plateau Remediation Company (CHPRC) Sample Delivery Group H4688 was composed of one water sample designated under SAF No. F11-149 with a Project Designation of: Liquid Effluent Retention Facility (LERF) Replacement Well C8204 – Water.

The samples were received as stated on the Chain-of-Custody document. Any discrepancies are noted on the Eberline Analytical Sample Receipt Checklist.

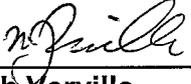
## 2.0 ANALYSIS NOTES

### 2.1 Tritium Analysis

Due to a clerical error, no matrix spike analysis was ordered. 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."**

  
\_\_\_\_\_  
Joseph Verville  
Client Services Manager

11/3/11  
\_\_\_\_\_  
Date

**SAMPLE ISSUE RESOLUTION**

<b>SIR NUM</b>	SDR12-057
<b>REV NUM</b>	0
<b>DATE INITIATED</b>	11/18/2011

**SAMPLE EVENT INFORMATION**

**SAF NUM(S)** F11-149  
**OPERABLE UNIT(S)** 200-PO-1  
**PROJECT(S)** M-24  
**SAMPLE EVENT TITLE(S)** LERF Well  
**LABORATORY** Waste Sampling & Characterization

**SAMPLING INFORMATION**

**NUMBER OF SAMPLES** 1  
**SAMPLE NUMBERS** B2FPH5  
**SAMPLE MATRIX** WATER  
**COLLECTION DATE** 9/22/2011 - 9/22/2011  
**SDG NUM** WSCF113250

**ISSUE BACKGROUND**

**CLASS** Laboratory Issue  
**TYPE** Quality Control Failure  
**DESCRIPTION** Eberline Laboratory did not perform a MS due to a clerical error for the tritium analysis. The laboratory did not issue a SIR to document this anomaly.

**DISPOSITION**

**DESCRIPTION** Document the laboratory anomaly, (MS was not performed for tritium due to a clerical error), and close the SIR.

**JUSTIFICATION** Accepted Resolution: Accept Proposed Resolution.

Submitted by: Michael Baechler / CHPRC Date 11/18/11  
Accepted by: Susan Puckett / CHPRC Date 11/21/11

SDG 7615  
 Contact N. Joseph Verville

Client CHPRC  
 Contract No. 33677  
 Case no SDG\_H4688

SUMMARY DATA SECTION

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 Prepared by \_\_\_\_\_

  
 Reviewed by \_\_\_\_\_

Lab id	<u>EBRLNE</u>
Protocol	<u>Hanford1</u>
Version	<u>Ver 1.0</u>
Form	<u>DVD-TOC</u>
Version	<u>3.06</u>
Report date	<u>11/03/11</u>

SDG 7615Contact N. Joseph Verville

## REPORT GUIDE

Client CHPRCContract No. 33677Case no SDG\_H4688

## 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  
 Protocol Hanford1  
 Version Ver 1.0  
 Form DVD-RG  
 Version 3.06  
 Report date 11/03/11

SDG 7615  
Contact N. Joseph Verville

GUIDE, cont.

Client CHPRC  
Contract No. 33677  
Case no SDG H4688

## 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  
Protocol Hanford1  
Version Ver 1.0  
Form DVD-RG  
Version 3.06  
Report date 11/03/11

SDG 7615  
 Contact N. Joseph Verville

**LAB SAMPLE SUMMARY**

Client CHPRC  
 Contract No. 33677  
 Case no SDG H4688

LAB SAMPLE ID	CLIENT SAMPLE ID	LOCATION	MATRIX	LEVEL	SAF NO	CHAIN OF CUSTODY	COLLECTED
S110130-01	B2FPH5	C8204-002	WATER		F11-149	F11-149-003-C	09/22/11 08:50
S110130-02	Lab Control Sample		WATER		F11-149		
S110130-03	Method Blank		WATER		F11-149		
S110130-04	Duplicate (S110130-01)	C8204-002	WATER		F11-149		09/22/11 08:50

LAB SUMMARY

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SUMMARY DATA SECTION

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Lab id EBRLNE  
 Protocol Hanford1  
 Version Ver 1.0  
 Form DVD-LS  
 Version 3.06  
 Report date 11/03/11

SDG 7615  
 Contact N. Joseph Verville

Client CHPRC  
 Contract No. 33677  
 Case no SDG H4688

QC SUMMARY

QC BATCH	CHAIN OF CUSTODY	CLIENT SAMPLE ID	MATRIX	% SOLIDS	SAMPLE AMOUNT	BASIS AMOUNT	DAYS SINCE RECEIVED	LAB COLL SAMPLE ID	DEPARTMENT SAMPLE ID
7615	F11-149-003-C	B2FPH5	WATER		250 mL		10/25/11 33	S110130-01	7615-001
		Method Blank	WATER					S110130-03	7615-003
		Lab Control Sample	WATER					S110130-02	7615-002
		Duplicate (S110130-01)	WATER		250 mL		10/25/11 33	S110130-04	7615-004

Lab id EBRLNE  
 Protocol Hanford1  
 Version Ver 1.0  
 Form DVD-QS  
 Version 3.06  
 Report date 11/03/11

EBERLINE ANALYTICAL/RICHMOND

SAMPLE DELIVERY GROUP H4688

SDG 7615  
Contact N. Joseph Verville

PREP BATCH SUMMARY

Client CHPRC  
Contract No. 33677  
Case no SDG H4688

TEST	MATRIX	METHOD	PREPARATION ERROR		PLANCHETS ANALYZED				QUALI-	
			BATCH	2σ %	CLIENT	MORE	RE	BLANK		LCS
Liquid Scintillation Counting										
H	WATER	Tritium in Water	7315-151	10.0	1		1	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.

Lab id EBRLNE  
Protocol Hanford1  
Version Ver 1.0  
Form DVD-PBS  
Version 3.06  
Report date 11/03/11

SDG 7615  
 Contact N. Joseph Verville

Client CHPRC  
 Contract No. 33677  
 Case no SDG H4688

**LAB WORK SUMMARY**

LAB SAMPLE	CLIENT SAMPLE ID	COLLECTED	LOCATION	MATRIX	SUF-	RECEIVED	CUSTODY	SAF No	PLANCHET	TEST	FIX	ANALYZED	REVIEWED	BY	METHOD
S110130-01	B2FPH5	09/22/11	C8204-002	WATER		10/25/11	F11-149-003-C	F11-149	7615-001	H		10/31/11	11/03/11	KWP	Tritium in Water
S110130-02	Lab Control Sample			WATER				F11-149	7615-002	H		10/31/11	11/03/11	KWP	Tritium in Water
S110130-03	Method Blank			WATER				F11-149	7615-003	H		10/31/11	11/03/11	KWP	Tritium in Water
S110130-04	Duplicate (S110130-01)	09/22/11	C8204-002	WATER		10/25/11		F11-149	7615-004	H		10/31/11	11/03/11	KWP	Tritium in Water

**COUNTS OF TESTS BY SAMPLE TYPE**

TEST	SAF No	METHOD	REFERENCE	CLIENT	MORE	RE	BLANK	LCS	DUP	SPIKE	TOTAL
H	F11-149	Tritium in Water	906.0_H3_LSC	1			1	1	1		4
TOTALS				1			1	1	1		4

WORK SUMMARY

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SUMMARY DATA SECTION

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Lab id EBRLNE  
 Protocol Hanford1  
 Version Ver 1.0  
 Form DVD-LWS  
 Version 3.06  
 Report date 11/03/11

7615-003

Method Blank

METHOD BLANK

SDG <u>7615</u>	Client/Case no <u>CHPRC</u>	SDG <u>H4688</u>
Contact <u>N. Joseph Verville</u>	Contract <u>No. 33677</u>	
Lab sample id <u>S110130-03</u>	Client sample id <u>Method Blank</u>	
Dept sample id <u>7615-003</u>	Material/Matrix <u>WATER</u>	
	SAF No <u>F11-149</u>	

ANALYTE	CAS NO	RESULT pCi/L	2σ ERR (COUNT)	MDA pCi/L	RDL pCi/L	QUALI- FIERS	TEST
Tritium	10028-17-8	4.30	12	13.1	400	U	H

QC-BLANK #80435
-----------------

Lab id <u>EBRLNE</u>
Protocol <u>Hanford1</u>
Version <u>Ver 1.0</u>
Form <u>DVD-DS</u>
Version <u>3.06</u>
Report date <u>11/03/11</u>

EBERLINE ANALYTICAL/RICHMOND

SAMPLE DELIVERY GROUP H4688

7615-002

Lab Control Sample

LAB CONTROL SAMPLE

SDG <u>7615</u>	Client/Case no <u>CHPRC</u>	<u>SDG H4688</u>
Contact <u>N. Joseph Verville</u>	Contract No. <u>33677</u>	
Lab sample id <u>S110130-02</u>	Client sample id <u>Lab Control Sample</u>	
Dept sample id <u>7615-002</u>	Material/Matrix <u>WATER</u>	
	SAF No <u>F11-149</u>	

ANALYTE	RESULT pCi/L	2σ ERR (COUNT)	MDA pCi/L	RDL pCi/L	QUALI- FIERS TEST	ADDED pCi/L	2σ ERR pCi/L	REC %	3σ LMTS (TOTAL)	PROTOCOL LIMITS
Tritium	255	15	13.2	400	H	250	10	102	81-119	80-120

QC-LCS #80434

Lab id <u>EBRLNE</u>
Protocol <u>Hanford1</u>
Version <u>Ver 1.0</u>
Form <u>DVD-LCS</u>
Version <u>3.06</u>
Report date <u>11/03/11</u>

7615-004

B2FPH5

**DUPLICATE**

SDG <u>7615</u>	Client/Case no <u>CHPRC</u>	SDG <u>H4688</u>
Contact <u>N. Joseph Verville</u>	Contract No. <u>33677</u>	
DUPLICATE	ORIGINAL	
Lab sample id <u>S110130-04</u>	Lab sample id <u>S110130-01</u>	Client sample id <u>B2FPH5</u>
Dept sample id <u>7615-004</u>	Dept sample id <u>7615-001</u>	Location/Matrix <u>C8204-002</u> <u>WATER</u>
	Received <u>10/25/11</u>	Collected/Volume <u>09/22/11 08:50</u> <u>250 mL</u>
		Custody/SAF No <u>F11-149-003-C</u> <u>F11-149</u>

ANALYTE	DUPLICATE pCi/L	2σ ERR (COUNT)	MDA pCi/L	RDL pCi/L	QUALI- FIERS	TEST	ORIGINAL pCi/L	2σ ERR (COUNT)	MDA pCi/L	QUALI- FIERS	RPD %	3σ TOT	DER σ
Tritium	144	120	129	400		H	95.7	120	131	U	40	214	0.6

QC-DUP#1 80436

LERF Replacement Well C8204 - Water

DUPLICATES

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SUMMARY DATA SECTION

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Lab id <u>EBRINE</u>
Protocol <u>Hanford1</u>
Version <u>Ver 1.0</u>
Form <u>DVD-DUP</u>
Version <u>3.06</u>
Report date <u>11/03/11</u>

EBERLINE ANALYTICAL / RICHMOND  
 SAMPLE DELIVERY GROUP H4688

7615-001

B2FPH5

DATA SHEET

SDG <u>7615</u>	Client/Case no <u>CHPRC</u>	SDG <u>H4688</u>
Contact <u>N. Joseph Verville</u>	Contract No. <u>33677</u>	
Lab sample id <u>S110130-01</u>	Client sample id <u>B2FPH5</u>	
Dept sample id <u>7615-001</u>	Location/Matrix <u>C8204-002</u>	<u>WATER</u>
Received <u>10/25/11</u>	Collected/Volume <u>09/22/11 08:50</u>	<u>250 mL</u>
	Custody/SAF No <u>F11-149-003-C</u>	<u>F11-149</u>

ANALYTE	CAS NO	RESULT pCi/L	2σ ERR (COUNT)	MDA pCi/L	RDL pCi/L	QUALI- FIERS	TEST
Tritium	10028-17-8	95.7	120	131	400	U	H

LERF Replacement Well C8204 - Water

Lab id <u>EBRLNE</u>
Protocol <u>Hanford1</u>
Version <u>Ver 1.0</u>
Form <u>DVD-DS</u>
Version <u>3.06</u>
Report date <u>11/03/11</u>

SAMPLE DELIVERY GROUP H4688

Test H          Matrix WATER  
 SDG 7615  
 Contact N. Joseph Verville

Client CHPRC  
 Contract No. 33677  
 Contract SDG H4688

**LAB METHOD SUMMARY**

TRITIUM IN WATER  
 LIQUID SCINTILLATION COUNTING

**RESULTS**

LAB            RAW    SUF-  
 SAMPLE ID    TEST FIX    PLANCHET    CLIENT SAMPLE ID            Tritium

Preparation batch 7315-151

S110130-01	7615-001	B2FPH5	U
S110130-02	7615-002	Lab Control Sample	ok
S110130-03	7615-003	Method Blank	U
S110130-04	7615-004	Duplicate (S110130-01)	ok

Nominal values and limits from method            RDLs (pCi/L)            400

**METHOD PERFORMANCE**

LAB            RAW    SUF-            MDA    ALIQ    PREP    DILU-    YIELD    EFF    COUNT    FWHM    DRIFT    DAYS            ANAL-  
 SAMPLE ID    TEST FIX    CLIENT SAMPLE ID            pCi/L    L            FAC    TION            %            %            min    keV    KeV    HELD    PREPARED    YZED    .DETECTOR

Preparation batch 7315-151            2σ prep error 10.0 %            Reference Lab Notebook 7315 pg. 151

S110130-01	B2FPH5	131	0.0100	100	300	39	10/28/11	10/31	LSC-005
S110130-02	Lab Control Sample	13.2	1.00	10	300		10/28/11	10/31	LSC-005
S110130-03	Method Blank	13.1	1.00	10	300		10/28/11	10/31	LSC-005
S110130-04	Duplicate (S110130-01)	129	0.0100	100	300	39	10/28/11	10/31	LSC-005

Nominal values and limits from method            400            0.0100            25            180

PROCEDURES    REFERENCE    906.0\_H3\_LSC  
 CP-210            Tritium in Water Samples by Distillation, rev 11

AVERAGES ± 2 SD            MDA 71.6 ± 135  
 FOR 4 SAMPLES            YIELD 55 ± 104

Lab id EBRLNE  
 Protocol Hanford1  
 Version Ver 1.0  
 Form DVD-LMS  
 Version 3.06  
 Report date 11/03/11

SDG 7615Contact N. Joseph Verville

## REPORT GUIDE

Client CHPRCContract No. 33677Case no SDG\_H4688

## 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.

## REPORT GUIDES

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## SUMMARY DATA SECTION

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Lab id EBRLNEProtocol Hanford1Version Ver 1.0Form DVD-RGVersion 3.06Report date 11/03/11

SDG 7615  
 Contact N. Joseph Verville

## REPORT GUIDE

Client CHPRC  
 Contract No. 33677  
 Case no SDG H4688

## 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 planchets 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 planchet 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.

REPORT GUIDES

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SUMMARY DATA SECTION

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Lab id EBRLNE  
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SDG 7615  
 Contact N. Joseph Verville

REPORT GUIDE

Client CHPRC  
 Contract No. 33677  
 Case no SDG H4688

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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 Protocol Hanford1  
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## REPORT GUIDE

Client CHPRC  
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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).

## REPORT GUIDES

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## SUMMARY DATA SECTION

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SDG 7615  
Contact N. Joseph Verville

G U I D E ,   c o n t .

Client CHPRC  
Contract No. 33677  
Case no SDG H4688

## D A T A   S H E E T

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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## SUMMARY DATA SECTION

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SDG 7615Contact N. Joseph Verville

GUIDE, cont.

Client CHPRCContract No. 33677Case no SDG\_H4688

## 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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 Contact N. Joseph Verville

REPORT GUIDE

Client CHPRC  
 Contract No. 33677  
 Case no SDG H4688

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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 Contact N. Joseph Verville

## R E P O R T   G U I D E

Client CHPRC  
 Contract No. 33677  
 Case no SDG H4688

## M A T R I X   S P I K E

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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Client CHPRC  
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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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## METHOD SUMMARY

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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Client CHPRCContract No. 33677Case no SDG\_H4688

## METHOD SUMMARY

- \* 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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## METHOD SUMMARY

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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CH2M Hill Plateau Remediation Company

CHAIN OF CUSTODY/SAMPLE ANALYSIS REQUEST

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COLLECTOR

COMPANY CONTACT

TELEPHONE NO.

PROJECT COORDINATOR

PRICE CODE

DATA TURNAROUND

Yvette Rust

LUKE, SN H4688

372-1667 (7615)

LUKE, SN

7H

30 Days / 30 Days

ICE CHEST NO.

6005-233

FIELD LOGBOOK NO. 49131

ACTUAL SAMPLE DEPTH

COA

AIR QUALITY

ORIGINAL

SHIPPED TO EBERLINE

OFFSITE PROPERTY NO.

226.7 to 240.25 FT

BILL OF LADING/AIR BILL NO.

7976 5898 5008

Waste Sampling & Characterization

FARE

3239

880, 10/24/11

7976 5898 5008

MATRIX\* POSSIBLE SAMPLE HAZARDS/ REMARKS

PRESERVATION

HOLDING TIME

TYPE OF CONTAINER

NO. OF CONTAINER(S)

VOLUME

SAMPLE ANALYSIS

SPECIAL HANDLING AND/OR STORAGE

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)

HNO3 to pH <2 Cool-4C Cool-4C None

Months 48 Hours 14 Days 6 Months

G/A G/A G/P G

1 1 1 1

500ml 500ml 250ml 250ml

SEE ITEM (1) IN SPECIAL INSTRUCTIONS (FAMSHIP); SEE ITEM (2) IN SPECIAL INSTRUCTIONS (FAMSHIP); 220 ALKALIN Ex (Tribium);

11325D WATER 9-22-11 0850

CHAIN OF POSSESSION

SIGN/ PRINT NAMES

SPECIAL INSTRUCTIONS

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RECEIVED BY/STORED IN

\*\* The CACN for all analytical work at WSCF laboratory is 402089E520.

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

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(1) ICP Metals - 6010 (TAL) {Calcium, Magnesium, Potassium, Sodium};

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(2) IC Anions - 300.0 {Chloride, Nitrogen in Nitrate, Sulfate};

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Please email copies of chain of custody, sample receipt form and final report to Marisol Avila (Marisol\_Avila@rl.gov) and Heather Medley (Heather\_A\_Medley@rl.gov), WSCF Project Managers.

RELINQUISHED BY/REMOVED FROM

RECEIVED BY/STORED IN

DATE/TIME

RELINQUISHED BY/REMOVED FROM

RECEIVED BY/STORED IN

DATE/TIME

RELINQUISHED BY/REMOVED FROM

RECEIVED BY/STORED IN

DATE/TIME

LABORATORY SECTION

RECEIVED BY

DATE/TIME

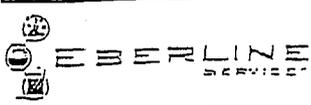
FINAL SAMPLE DISPOSITION

DISPOSAL METHOD

DATE/TIME

PRINTED ON 7/19/2011

A-6003-618 (REV 2)



**RICHMOND, CA LABORATORY**  
SAMPLE RECEIPT CHECKLIST

7615

Client: CHPRC City RICHMOND State WA  
 Date/Time received 10/25/11 1045 CoC No. F11-K9-003-C  
 Container I.D. No. GWS-233 Requested TAT (Days) 30 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: 1 Sample Matrix W
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 [Signature] Date: 10/25/11 Time: 1115

Customer Sample No.	Beta/Gamma com	Ion Chamber mR/hr	Wide	Customer Sample No.	Beta/Gamma com	Ion Chamber mR/hr	Wide
B2FPH5	280						

Ion Chamber Ser. No. \_\_\_\_\_ Calibration date \_\_\_\_\_  
 Alpha Meter Ser. No. \_\_\_\_\_ Calibration date \_\_\_\_\_  
 Beta/Gamma Meter Ser. No. 99574 Calibration date 15 JUL 11