

0051663

Thermo Nutech
W.O. No. N9-06-139-7723

Bechtel Hanford Inc.
SDG H0450

Case Narrative

1.0 GENERAL

Bechtel Hanford Inc. Sample Delivery Group H0450 is composed of three solid (filter) samples designated under SAF No. B99-079 with a Project Designation of: 233-S Facility RadCon Air Filter Analysis.

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

2.0 ANALYSIS NOTES

2.1 Isotopic Plutonium Analyses

The samples were reanalyzed due to very low recoveries on the initial analysis. No problems were encountered during the reanalysis.



TMA/RICHMOND
 SAMPLE DELIVERY GROUP H0450

Client Hanford
 Contract TRB-SBB-207925
 Case no SDG-H0450

SDG 7723
 Contact L.A. Johnson

SAMPLE SUMMARY

CLIENT SAMPLE ID	LOCATION	MATRIX	LEVEL	LAB SAMPLE ID	SAF NO	CHAIN OF CUSTODY	COLLECTED
B0VTB5		FILTERS		N906139-01	B99-079	B99-079-01	06/14/99 09:20
B0VTB6		FILTERS		N906139-02	B99-079	B99-079-01	06/14/99 10:00
B0VTB7		FILTERS		N906139-03	B99-079	B99-079-01	06/14/99 15:25
Method Blank		FILTERS		N906139-08	B99-079		
Lab Control Sample		FILTERS		N906139-07	B99-079		
Duplicate (N906139-02)		FILTERS		N906139-09	B99-079		06/14/99 10:00

Lab id TMANC
 Protocol Hanford
 Version Ver 1.0
 Form DVD-CS
 Version 3.06
 Report date 07/08/99

TMA/RICHMOND
SAMPLE DELIVERY GROUP H0450

SDG 7723
Contact L.A. Johnson

QC SUMMARY

Client Hanford
Contract TRB-SBB-207925
Case no SDG-H0450

QC BATCH	CHAIN OF CUSTODY	CLIENT SAMPLE ID	MATRIX	* SOLIDS	SAMPLE AMOUNT	BASIS AMOUNT	DAYS SINCE RECEIVED	LAB COLL SAMPLE ID	DEPARTMENT SAMPLE ID
7723	B99-079-01	B0VTB5	FILTERS				06/24/99 10	N906139-01	7723-001
		B0VTB6	FILTERS				06/24/99 10	N906139-02	7723-002
		B0VTB7	FILTERS				06/24/99 10	N906139-03	7723-003
		Method Blank	FILTERS					N906139-08	7723-008
		Lab Control Sample	FILTERS					N906139-07	7723-007
		Duplicate (N906139-02)	FILTERS				06/24/99 10	N906139-09	7723-009

Lab id TMANC
Protocol Hanford
Version Ver 1.0
Form DVD-QS
Version 3.06
Report date 07/08/99

TMA/RICHMOND

SAMPLE DELIVERY GROUP H0450

SDG 7723
Contact L.A. Johnson

PREP BATCH SUMMARY

Client Hanford
Contract TRB-SBB-207925
Case no SDG-H0450

TEST MATRIX	METHOD	PREPARATION ERROR			PLANCHETS ANALYZED			QUALI-
		BATCH	2σ %	CLIENT MORE	RE BLANK	LCS	DUP/ORIG MS/ORIG	
Alpha Spectroscopy								
PU	FILTERS	Plutonium, Isotopic in Filters	6880-145	5.0	3		1 1	1/1

Duplicates and Matrix Spikes are those with original (Client) sample in this Sample Delivery Group.
Blank and LCS planchets are those in the same preparation batch as some Client, Duplicate or Spike sample.

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Report date 07/08/99

TMA/RICHMOND
SAMPLE DELIVERY GROUP H0450

SDG 7723
Contact L.A. Johnson

Client Hanford
Contract TRB-SBB-207925
Case no SDG-H0450

WORK SUMMARY

CLIENT SAMPLE ID	LAB SAMPLE ID									
LOCATION	MATRIX	COLLECTED	SUF-							
CUSTODY	SAF No	RECEIVED	PLANCHET	TEST	FIX	ANALYZED	REVIEWED	BY	METHOD	
BOVTB5		N906139-01	7723-001	PU	A1	07/07/99	07/08/99	TAH	Plutonium, Isotopic in Filters	
	FILTERS	06/14/99								
B99-079-01	B99-079	06/24/99								
BOVTB6		N906139-02	7723-002	PU	A1	07/07/99	07/08/99	TAH	Plutonium, Isotopic in Filters	
	FILTERS	06/14/99								
B99-079-01	B99-079	06/24/99								
BOVTB7		N906139-03	7723-003	PU	A1	07/07/99	07/08/99	TAH	Plutonium, Isotopic in Filters	
	FILTERS	06/14/99								
B99-079-01	B99-079	06/24/99								
Method Blank		N906139-08	7723-008	PU		07/07/99	07/08/99	TAH	Plutonium, Isotopic in Filters	
	FILTERS									
	B99-079									
Lab Control Sample		N906139-07	7723-007	PU		07/07/99	07/08/99	TAH	Plutonium, Isotopic in Filters	
	FILTERS									
	B99-079									
Duplicate (N906139-02)		N906139-09	7723-009	PU		07/07/99	07/08/99	TAH	Plutonium, Isotopic in Filters	
	FILTERS	06/14/99								
	B99-079	06/24/99								

COUNTS OF TESTS BY SAMPLE TYPE

TEST	SAF No	METHOD	REFERENCE	CLIENT	MORE	RE	BLANK	LCS	DUP	SPIKE	TOTAL
PU	B99-079	Plutonium, Isotopic in Filters	PUPLATE	3			1	1	1		6
TOTALS				3			1	1	1		6

WORK SUMMARY

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SUMMARY DATA SECTION

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Protocol Hanford
Version Ver 1.0
Form DVD-CWS
Version 3.06
Report date 07/08/99

TMA / RICHMOND
SAMPLE DELIVERY GROUP H0450

N906139-08

Method Blank

METHOD BLANK

SDG <u>7723</u>	Client/Case no <u>Hanford</u>	<u>SDG-H0450</u>
Contact <u>L.A. Johnson</u>	Case no <u>TRB-SBB-207925</u>	
Lab sample id <u>N906139-08</u>	Client sample id <u>Method Blank</u>	
Dept sample id <u>7723-008</u>	Material/Matrix <u>FILTERS</u>	
	SAF No <u>B99-079</u>	

ANALYTE	CAS NO	RESULT pCi/smp	2 σ ERR (COUNT)	MDA pCi/smp	RDL pCi/smp	QUALI- FIERS	TEST
Plutonium 238	13981-16-3	0	0.013	0.049	1.0	U	PU
Plutonium 239/240	PU-239/240	-0.006	0.013	0.049	1.0	U	PU

233-S FACILITY RADCON ARI FILTER ANA

QC-BLANK 31227

Lab id <u>TMANC</u>
Protocol <u>Hanford</u>
Version <u>Ver 1.0</u>
Form <u>DVD-DS</u>
Version <u>3.06</u>
Report date <u>07/08/99</u>

METHOD BLANKS

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TMA/RICHMOND

SAMPLE DELIVERY GROUP H0450

N906139-07

Lab Control Sample

LAB CONTROL SAMPLE

SDG <u>7723</u>	Client/Case no <u>Hanford</u>	<u>SDG-H0450</u>
Contact <u>L.A. Johnson</u>	Case no <u>TRB-SBB-207925</u>	
Lab sample id <u>N906139-07</u>	Client sample id <u>Lab Control Sample</u>	
Dept sample id <u>7723-007</u>	Material/Matrix _____	<u>FILTERS</u>
	SAF No <u>B99-079</u>	

ANALYTE	RESULT pCi/smp	2σ ERR (COUNT)	MDA pCi/smp	RDL pCi/smp	QUALI- FIERS TEST	ADDED pCi/smp	2σ ERR pCi/smp	REC %	3σ LMIS (TOTAL)	PROTOCOL LIMITS
Plutonium 238	6.57	0.51	0.044	1.0	PU	6.28	0.25	105	84-116	80-120
Plutonium 239/240	6.59	0.51	0.044	1.0	PU	6.61	0.26	100	85-115	80-120

233-S FACILITY RADCON ARI FILTER ANA

QC-LCS 31226

LAB CONTROL SAMPLES

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Lab id <u>TMANC</u>
Protocol <u>Hanford</u>
Version <u>Ver 1.0</u>
Form <u>DVD-LCS</u>
Version <u>3.06</u>
Report date <u>07/08/99</u>

TMA/RICHMOND
SAMPLE DELIVERY GROUP H0450

N906139-09

B0VTB6

DUPLICATE

SDG <u>7723</u> Contact <u>L.A. Johnson</u> Duplicates Lab sample id <u>N906139-09</u> Dept sample id <u>7723-009</u>	Client/Case no <u>Hanford</u> <u>SDG-H0450</u> Case no <u>TRB-SBB-207925</u> ORIGINAL Lab sample id <u>N906139-02</u> Dept sample id <u>7723-002</u> Received <u>06/24/99</u> Client sample id <u>B0VTB6</u> Location/Matrix <u>FILTERS</u> Collected <u>06/14/99 10:00</u> Custody/SAF No <u>B99-079-01</u> <u>B99-079</u>
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ANALYTE	DUPLICATE		2σ ERR		MDA	RDL	QUALI-	ORIGINAL	2σ ERR		MDA	QUALI-	RPD	3σ	PROT
	pCi/smp	(COUNT)	pCi/smp	pCi/smp	pCi/smp	FIERS	TEST		pCi/smp	(COUNT)	pCi/smp	FIERS	%	TOT	LIMIT
Plutonium 238	0.036	0.14	0.27	1.0	U	PU	0	0.071	0.27	U	-	-	-	-	
Plutonium 239/240	0	0.072	0.27	1.0	U	PU	0.178	0.14	0.27	U	-	-	-	-	

233-S FACILITY RADCON ARI FILTER ANA

QC-DUP#2 31228

DUPLICATES

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Lab id <u>TMANC</u>
Protocol <u>Hanford</u>
Version <u>Ver 1.0</u>
Form <u>DVD-DUP</u>
Version <u>3.06</u>
Report date <u>07/08/99</u>

T M A / R I C H M O N D
S A M P L E D E L I V E R Y G R O U P H 0 4 5 0

N906139-01

B0VTB5

D A T A S H E E T

SDG <u>7723</u>	Client/Case no <u>Hanford</u>	SDG- <u>H0450</u>
Contact <u>L.A. Johnson</u>	Case no <u>TRB-SBB-207925</u>	
Lab sample id <u>N906139-01</u>	Client sample id <u>B0VTB5</u>	
Dept sample id <u>7723-001</u>	Location/Matrix _____	<u>FILTERS</u>
Received <u>06/24/99</u>	Collected <u>06/14/99 09:20</u>	
	Custody/SAF No <u>B99-079-01</u>	<u>B99-079</u>

ANALYTE	CAS NO	RESULT pCi/smp	2σ ERR (COUNT)	MDA pCi/smp	RDL pCi/smp	QUALI- FIERS	TEST
Plutonium 238	13981-16-3	0.065	0.065	0.25	1.0	U	PU
Plutonium 239/240	PU-239/240	0.454	0.26	0.25	1.0	J	PU

233-S FACILITY RADCON ARI FILTER ANA

Lab id <u>TMANC</u>
Protocol <u>Hanford</u>
Version <u>Ver 1.0</u>
Form <u>DVD-DS</u>
Version <u>3.06</u>
Report date <u>07/08/99</u>

TMA / RICHMOND
SAMPLE DELIVERY GROUP H0450

N906139-02

B0VTB6

DATA SHEET

SDG <u>7723</u>	Client/Case no <u>Hanford</u>	SDG-H0450
Contact <u>L.A. Johnson</u>	Case no <u>TRB-SBB-207925</u>	
Lab sample id <u>N906139-02</u>	Client sample id <u>B0VTB6</u>	
Dept sample id <u>7723-002</u>	Location/Matrix _____	<u>FILTERS</u>
Received <u>06/24/99</u>	Collected <u>06/14/99 10:00</u>	
	Custody/SAF No <u>B99-079-01</u>	<u>B99-079</u>

ANALYTE	CAS NO	RESULT pCi/smp	2 σ ERR (COUNT)	MDA pCi/smp	RDL pCi/smp	QUALI- FIERS	TEST
Plutonium 238	13981-16-3	0	0.071	0.27	1.0	U	PU
Plutonium 239/240	PU-239/240	0.178	0.14	0.27	1.0	U	PU

233-S FACILITY RADCON ARI FILTER ANA

Lab id <u>TMANC</u>
Protocol <u>Hanford</u>
Version <u>Ver 1.0</u>
Form <u>DVD-DS</u>
Version <u>3.06</u>
Report date <u>07/08/99</u>

TMA / RICHMOND
 SAMPLE DELIVERY GROUP H0450

N906139-03

B0VTB7

DATA SHEET

SDG <u>7723</u>	Client/Case no <u>Hanford</u>	<u>SDG-H0450</u>
Contact <u>L.A. Johnson</u>	Case no <u>TRB-SBB-207925</u>	
Lab sample id <u>N906139-03</u>	Client sample id <u>B0VTB7</u>	
Dept sample id <u>7723-003</u>	Location/Matrix _____	<u>FILTERS</u>
Received <u>06/24/99</u>	Collected <u>06/14/99 15:25</u>	
	Custody/SAF No <u>B99-079-01</u>	<u>B99-079</u>

ANALYTE	CAS NO	RESULT pCi/smp	2 σ ERR (COUNT)	MDA pCi/smp	RDL pCi/smp	QUALI- FIERS	TEST
Plutonium 238	13981-16-3	0	0.071	0.27	1.0	U	PU
Plutonium 239/240	PU-239/240	0.035	0.14	0.27	1.0	U	PU

233-S FACILITY RADCON ARI FILTER ANA

Lab id <u>TMANC</u>
Protocol <u>Hanford</u>
Version <u>Ver 1.0</u>
Form <u>DVD-DS</u>
Version <u>3.06</u>
Report date <u>07/08/99</u>

TMA/RICHMOND

SAMPLE DELIVERY GROUP H0450

METHOD SUMMARY

PLUTONIUM, ISOTOPIC IN FILTERS

ALPHA SPECTROSCOPY

Test PU Matrix FILTERS

SDG 7723

Contact L.A. Johnson

Client Hanford

Contract TRB-SBB-207925

Case no SDG-H0450

RESULTS

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW TEST	SUF- FIX	PLANCHET	Plutonium 238	Plutonium 239/240
Preparation batch 6880-145						
B0VTB5	N906139-01	A1	7723-001		U	0.454 J
B0VTB6	N906139-02	A1	7723-002		U	U
B0VTB7	N906139-03	A1	7723-003		U	U
BLK (QC ID=31227)	N906139-08		7723-008		U	U
LCS (QC ID=31226)	N906139-07		7723-007		ok	ok
Duplicate (N906139-02)	N906139-09		7723-009		- U	- U

Nominal values and limits from method RDLs (pCi/smp) 1.0 1.0
233-S FACILITY RADCON ARI FILTER ANA

METHOD PERFORMANCE

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW TEST	SUF- FIX	MAX MDA pCi/smp	ALIQ smp	PREP PAC	DILU- TION	YIELD %	EFF %	COUNT min	FWHM keV	DRIFT KeV	DAYS HELD	ANAL- PREPARED	YZED	DETECTOR
Preparation batch 6880-145 2σ prep error 5.0 % Reference Lab Notebook 6880 pg.144																
B0VTB5	N906139-01	A1	0.25	<u>0.200</u>				82		212			23	07/07/99	07/07	SS-055
B0VTB6	N906139-02	A1	0.27	<u>0.200</u>				77		212			23	07/07/99	07/07	SS-056
B0VTB7	N906139-03	A1	0.27	<u>0.200</u>				79		212			23	07/07/99	07/07	SS-057
BLK (QC ID=31227)	N906139-08		0.049	1.00				83		211				07/07/99	07/07	SS-060
LCS (QC ID=31226)	N906139-07		0.044	1.00				96		211				07/07/99	07/07	SS-059
Duplicate (N906139-02) (QC ID=31228)	N906139-09		0.27	<u>0.200</u>				76		211			23	07/07/99	07/07	SS-061

Nominal values and limits from method 1.0 1.00 20-105 10 100 180

PROCEDURES	REFERENCE	PUPLATE
EP-060		Soil Preparation, rev 0
EP-070		Soil Dissolution, rev 0
EP-940		Plutonium Purification, rev 0
EP-008		Heavy Elements Electroplating, rev 0

AVERAGES ± 2 SD	MDA <u>0.19</u> ± <u>0.23</u>
FOR 6 SAMPLES	YIELD <u>82</u> ± <u>15</u>

METHOD SUMMARIES

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SUMMARY DATA SECTION

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Lab id TMANC

Protocol Hanford

Version Ver 1.0

Form DVD-CMS

Version 3.06

Report date 07/08/99

TMA / RICHMOND
SAMPLE DELIVERY GROUP H0450

SDG 7723
Contact L.A. Johnson

REPORT GUIDE

Client Hanford
Contract TRB-SBB-207925
Case no SDG-H0450

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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Lab id TMANC
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TMA / RICHMOND
SAMPLE DELIVERY GROUP H0450

SDG 7723
Contact L.A. Johnson

R E P O R T G U I D E

Client Hanford
Contract TRB-SBB-207925
Case no SDG-H0450

P R E P A R A T I O N B A T C H S U M M A R Y

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.

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TMA / RICHMOND
SAMPLE DELIVERY GROUP H0450

SDG 7723
Contact L.A. Johnson

R E P O R T G U I D E

Client Hanford
Contract TRB-SBB-207925
Case no SDG-H0450

W O R K S U M M A R Y

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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TMA / RICHMOND
SAMPLE DELIVERY GROUP H0450

SDG 7723
Contact L.A. Johnson

REPORT GUIDE

Client Hanford
Contract TRB-SBB-207925
Case no SDG-H0450

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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TMA / RICHMOND
SAMPLE DELIVERY GROUP H0450

SDG 7723
Contact L.A. Johnson

GUIDE , c o n t .

Client Hanford
Contract TRB-SBB-207925
Case no SDG-H0450

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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TMA / RICHMOND
SAMPLE DELIVERY GROUP H0450

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Contract TRB-SBB-207925
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GUIDE , c o n t .

D A T A S H E E T

- * 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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Version Ver 1.0
Form DVD-RG
Version 3.06
Report date 07/08/99

TMA / RICHMOND
SAMPLE DELIVERY GROUP H0450

SDG 7723
Contact L.A. Johnson

REPORT GUIDE

Client Hanford
Contract TRB-SBB-207925
Case no SDG-H0450

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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SAMPLE DELIVERY GROUP H0450

SDG 7723
Contact L.A. Johnson

R E P O R T G U I D E

Client Hanford
Contract TRB-SBB-207925
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D U P L I C A T E

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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T M A / R I C H M O N D
S A M P L E D E L I V E R Y G R O U P H 0 4 5 0

SDG 7723
Contact L.A. Johnson

R E P O R T G U I D E

Client Hanford
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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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G U I D E , c o n t .

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M A T R I X S P I K E

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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Contact L.A. Johnson

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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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Version 3.06
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SDG 7723
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M E T H O D S U M M A R Y

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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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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Lab id TMANC
Protocol Hanford
Version Ver 1.0
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Collector R Fahlberg / <i>Randy</i>	Company Contact K Mathews	Telephone No. 531-0709	Project Coordinator TRENT, SJ	Price Code 6F	Data Turnaround 7 Days
Project Designation 233-S Facility RadCon Air Filter Analysis	Sampling Location 233-S 200 Area	SAF No. B99-079			
Ice Chest No. <i>SML-69</i>	Field Logbook No. <i>EL 1435</i>	Method of Shipment <i>Fed Ex</i>			
Shipped To TMA/REGA <i>6-22-99</i>	Offsite Property No. <i>A990172</i>	Bill of Lading/Air Bill No. <i>423579527124</i>		COA <i>R233SG2300</i>	

POSSIBLE SAMPLE HAZARDS/REMARKS	Preservation	None																		
	Type of Container	Poly Bag																		
	No. of Container(s)	1																		
	Special Handling and/or Storage	Volume	10g																	
SAMPLE ANALYSIS		Isotopic Plutonium																		

Sample No.	Matrix *	Sample Date	Sample Time																		
B0VTB5	Other Solid	6-14-99	0920	X																	tie to RCF 6198
B0VTB6	Other Solid	6-14-99	1000	X																	RCF 6199
B0VTB7	Other Solid	6-14-99	1525	X																	RCF 6197

CHAIN OF POSSESSION	Sign/Print Names		SPECIAL INSTRUCTIONS										Matrix * Soil Water Vapor Other Solid Other Liquid		
	Relinquished By <i>R. Fahlberg / R. Fahlberg</i>	Date/Time <i>6-22-99 1400</i>	Received By <i>Ref 1-B</i>	Date/Time <i>6-22-99 1400</i>											
	Relinquished By <i>R. Fahlberg</i>	Date/Time <i>6-23-99 1500</i>	Received By <i>Fed Ex</i>	Date/Time <i>6-23-99 9:40</i>											
	Relinquished By <i>Fed Ex</i>	Date/Time <i>6-24-99 9:40</i>	Received By <i>TNU H. Goldenberg</i>	Date/Time <i>6-24-99 9:40</i>											

LABORATORY SECTION	Received By	Title	Date/Time
FINAL SAMPLE DISPOSITION	Disposal Method	Disposed By	Date/Time