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Results of the Excreta Bioassay Quality Control Program For April 1, 2006 Through March 31, 2007

CL Antonio

January 2008



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RESULTS OF THE EXCRETA BIOASSAY QUALITY CONTROL PROGRAM FOR APRIL 1, 2006 THROUGH MARCH 31, 2007

Cheryl L. Antonio

January 2008

<u>_____</u> Date Peer Reviewed by Jay Maclellan

SUMMARY

A total of 66 urine samples, 6 blank fecal and 6 spiked artificial fecal samples were submitted during the report period (April 1, 2006 through March 31, 2007) to General Engineering Laboratories, South Carolina by the Hanford Internal Dosimetry Program (IDP) to check the accuracy, precision, and detection levels of their analyses. Urine analyses for tritium, Sr, ²³⁸Pu, ²³⁹Pu, ²⁴¹Am, ²⁴³Am ²³⁵U, ²³⁸U, elemental uranium and fecal analyses for ²⁴¹Am, ²³⁸Pu and ²³⁹Pu were tested this year. The number of QC urine samples submitted during the report period represented 1.7% of the total samples submitted.

In addition to the samples provided by IDP, GEL was also required to conduct their own QC program, and submit the results of analyses to IDP. About 36% of the analyses processed by GEL during the second year of this contract were quality control samples. GEL tested the performance of 16 radioisotopes, all of which met or exceeded the specifications in the Statement of Work (Table 4).

IDP concluded that GEL was performing well for all analyses tested, and concerns identified earlier were satisfactorily resolved (see section on Follow-up on Concerns During the Second Contract Year).

The isotopic uranium analysis reports on three uranium isotopes: ²³⁴U, ²³⁵U, and ²³⁸U. The isotopes are differentiated only during counting by alpha spectrometry. GEL reported that the calculated minimum detectable activity (MDA) for ^{233,234}U for the year slightly exceeded the contract required detection limit. This was consistently seen in the first, second and third quarters. The cause was attributed to low tracer recoveries. The standard operating procedure was revised November 30, 2006 to improve the separation chemistry. GEL then reviewed the MDAs for the fourth quarter and showed that all three isotopes met the acceptance criteria.

IDP did not submit isotopic uranium samples to review the ^{233,234}U performance indicators. However, the performance statistics for ²³⁵U and ²³⁸U were reviewed and the MDA for ²³⁵U and the bias and precision for ²³⁸U were acceptable. The percentage of analyses with low yields were reviewed and found not to meet the criteria in the statement of work for the first, second and third quarters. In line with GEL's changes to the standard operating procedure, the low yield rate in the fourth quarter was significantly improved upon and the low yield rate met the criteria in the statement of work. A review of the data from April 1, 2007 through December 31, 2007, also showed improvement in tracer recoveries for isotopic uranium urinalyses. Of the 398 urinalyses processed during this time, all had tracer recoveries at 40% or greater. With the implementation of the revised standard operating procedure, the isotopic uranium analysis program is considered acceptable

No concerns were identified with the elemental uranium urinalysis program and it was considered acceptable. The bias and precision for uranium spiked at 0.06 μ g, as tested by IDP, did not meet the acceptance criteria. However, only 2 samples were tested and environmental contamination was thought to be an interference. Because IDP uses a 0.2 μ g screening level for elemental uranium, samples spiked at 0.06 μ g were discontinued. The MDA at the contractual level of 0.06 μ g was evaluated through GEL's program and was found to be acceptable. The bias and precision was tested by IDP at 0.2 μ g and by GEL at 0.1 μ g and was also found to be acceptable. For the third contract year, GEL will begin evaluating the bias and precision for elemental uranium at levels of 0.06 μ g, in accordance with the contractual level.

The total strontium procedure is used to screen samples to determine which will require analysis for ⁹⁰Sr. Samples with total strontium results less than 15 dpm do not undergo further analysis. Samples with results greater than or equal to 15 dpm may undergo ⁹⁰Y in growth to specifically determine ⁹⁰Sr levels. The calculated MDA, as reported by GEL, for the total strontium part of the analysis was about 30% of the CL. The relative bias and precision, tested by IDP and GEL for the ⁹⁰Sr and total Sr procedures were all within limits. The 14 samples spiked at the contractual level were all detected. The strontium urinalysis procedure was concluded to be acceptable.

Samples spiked with ²³⁸Pu and ²³⁹Pu were analyzed using the same procedures and same reagents. The two isotopes are differentiated only at the end of the procedure by alpha spectrometry. Therefore, laboratory performance is expected to be similar for both isotopes using any of the seven procedures that incorporate plutonium analysis (IPU, IPA, IPS, IPSA, IPSR, IUPU, and ITPAC).

The MDAs and performance statistics for ²³⁹Pu and ²³⁸Pu in urine were acceptable. The 16 samples spiked at the CL for ²³⁹Pu were reported with a result greater than the decision level and the 6 blank samples were reported with results less than the decision level. With the exception of one sample, the 21 blank ²³⁸Pu samples were reported with results less than the decision level, giving a false positive rate less than 5%. Overall the plutonium urinalyses were considered acceptable.

The MDA and performance statistics for ²³⁹Pu and ²³⁸Pu in feces were acceptable. Approximately 15% of the fecal samples analyzed were duplicated to test the consistency of the aliquoting procedure. A review of the duplicate samples determined that the aliquoting procedure produced results within 3 sigma of the initial results. The fecal aliquoting procedure was acceptable. None of the 12 blank ²³⁸Pu or the 6 blank ²³⁹Pu fecal analyses were greater than the decision level. There were no fecal samples spiked at the CL with ²³⁸Pu. The six fecal samples spiked with ²³⁹Pu were reported with a result greater than the decision level. Overall the plutonium fecal analyses were considered acceptable.

The ²⁴¹Am fecal and urine analysis met the acceptance criteria for MDA, relative bias and precision. With regards to the ²⁴¹Am urinalysis program there were only 2 blank samples and the resulting MDA exceeded the acceptance criteria, however, this was most likely a result of low-level counting statistics. The MDA as reported by GEL was 50% of the contractual level. The 17 samples spiked by IDP at the contractual detection level were all greater than decision level. The current AM241 urinalysis procedure was considered acceptable.

All the six blank fecal samples were less than the decision level and the four spiked fecal samples were all greater than the decision. The ²⁴¹Am fecal duplicate samples were evaluated and it was concluded that the aliquoting procedure produced results within the control limits. A review of the routine ²⁴¹Am results reported, however, consistently found more than 10% of the samples with tracer recoveries at or below 40%. GEL reported that the overall low yield for AM241 fecal analyses was 7.4%. However, GEL was including duplicate samples in their tally, a mistake which will be corrected in subsequent QC reports.

In mid-January 2007, GEL identified a cause for the low tracer recoveries in the La(F₃) precipitation step. GEL modified their standard operating procedure to include an additional 1 mL of HF when performing the La(F₃) step. This correction appeared to have improved the tracer recovery rate. However, in early March there was a batch of 10 fecal samples that were declared failed analyses due to a technician error. To better evaluate the efficiency of the procedure modification fecal analyses from the first few quarters of the third contract year (3/1/07 - 12/31/07) were reviewed and only 3% of the samples had tracer recoveries less than 40%. With the implementation of the revised standard operating procedure, the AM241 fecal analysis program is considered acceptable.

The AM243 procedure was identical to the AM241 procedure, except that a different tracer is used (²⁴⁴Cm instead of ²⁴³Am). The 6 blank ²⁴³Am QC samples submitted were all reported with results less than the decision level and the calculated MDA was 50% of the contractual detection level.

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The performance statistics for the AM243, as tested by GEL, met the acceptance criteria. The AM243 procedure was concluded to be acceptable.

IDP did not submit QC samples to test the isotopic curium program, therefore performance statistics were based on the GEL QC results. GEL tested the MDA for ²⁴²Cm and ²⁴⁴Cm and the relative bias and precision for ²⁴⁴Cm. The results met the acceptance criteria and the isotopic curium urinalysis program was considered acceptable.

During the second contract year, no isotopic thorium analyses were requested. Therefore, there were no QC samples to evaluate.

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INTRODUCTION

This report summarizes the results of the excreta bioassay quality control program's monitoring of the performance of General Engineering Laboratories (GEL) for samples submitted from April 1, 2006 through March 31, 2007. During the reporting period GEL analyzed, under the contract with Battelle, 4503 urine and 111 fecal samples for various radionuclides. This is about the same workload as reported in the 2006 report.

The results of the analyses are part of a system of legal records concerning internal deposition of radionuclides for workers at the Hanford Site. GEL is required to have a rigorous quality control (QC) program to ensure the accuracy of its results. In addition, the Pacific Northwest National Laboratory's (PNNL) Hanford Internal Dosimetry Program (IDP) has a QC program in place to independently check the accuracy of the results from GEL. The objective of the PNNL excreta bioassay QC program is to provide quantitative data to support the assessment of performance criteria for excreta bioassay analyses, as specified in the Statement of Work (Battelle 2007).

The reliability of the excreta bioassay program depends, to a significant extent, on the adoption and implementation of performance criteria for laboratory accuracy, precision, and detection levels. Such performance criteria are established in the Statement of Work (Battelle 2007) and include the following:

- Actual minimum detectable activities (MDAs) determined from QC samples for the year shall be equal to or less than the contractual detection level (CL) in the Statement of Work, as calculated from blank QC samples.
- The mean relative bias, B_r, shall fall within ± 20% when calculated from 15 to 50 samples spiked at greater than three times the CL, and within ± 10% when calculated from greater than 50 samples.
- The relative precision statistic, S_B, shall be less than or equal to 0.4 for samples spiked at greater than three times the CL, and less than or equal to 0.5 for samples spiked between one and three times the CL.

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Formulas for MDA, B_r, and S_B, presented in the next section of this report, are based on recommendations in the Health Physics Society (HPS) Standard N13.30 (1996) and are listed in the Statement of Work. In addition to the Statement of Work (SOW) performance criteria, it is expected that the MDA shall also be such that fewer than 10% of the QC samples spiked at the CL shall be reported with values less than the decision level (i.e., twice the total propagated uncertainty of the result)..

METHODS

GENERAL METHODS

Urine collected from PNNL employees who are not occupationally exposed to radioactive material was prepared in the 325 Building as blank and spiked samples by PNNL Radiochemical Processing Group (RPG), according to the directions given by the PNNL Internal Dosimetry Program (IDP), following Procedure PNL-MA-565-800-20, Rev. 2. Most samples were submitted as double-blind samples, with the exception of isotopic uranium urinalyses and the spiked fecal samples. Double blind samples are scheduled with and collected by GEL as if they were personnel samples. The isotopic uranium urinalyses were scheduled as single-blind intercomparisons, which meant that GEL was aware they were intercomparison samples but unaware of the activity. The samples were scheduled as single-blinds because they were spiked with a depleted uranium source. Since depleted uranium exposures at Hanford are rare, the intercomparison samples would stand out and the QC alias names used could become known and compromise the double-blind intercomparison program. The spiked fecal samples were artificial fecal samples consisting of a soil matrix. Blank fecal samples were scheduled as double-blind samples and were actual fecal samples.

GEL analyzed urine samples for tritium, ⁹⁰Sr, ²⁴²Cm, ²⁴⁴Cm, ²³⁸Pu, ^{239,240}Pu, ²⁴¹Pu ²⁴¹Am, ²⁴³Am, ²³⁴U, ²³⁵U, ²³⁸U and elemental uranium and fecal samples for ²³⁸Pu, ^{239,240}Pu, ²⁴¹Am, ²⁴²Cm and ²⁴⁴Cm. To reduce costs in the intercomparison program, plutonium, americium, and strontium analyses were tested using routine sequential procedures when possible (i.e., where one urine sample is analyzed for several radionuclides). The analysis categories specified in the contract with GEL are shown in Table 1. All urinalysis samples contained approximately 1000 ml of urine, except for the samples analyzed for tritium, which contained approximately 100 ml.

GEL's QC sample total is dependent on the number of analytical batches run during the year, and they were well over the 15% criteria specified in the contract.

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		Contractual Detection	Detection	Determination				<u>Oral Reporting Level:</u>	<u>ing Level;</u>
		Level (a) <u>(dpm/sample)</u> Urine	<u>m/sample)</u>	Time (business dave following	I	Remrting Time	¢D-i	(dpm/	(dpm/sample)
<u>Analysis (Code)</u>	Constituents Reported		Fecal	sample receipt)	Oral ⁽⁶⁾	Electronic ^(a)	Written ^(a)	Urine	Fecal
Pu(∝) Isotopic (IPU)	Pu-238, Pu-239, 240	0.02	0.2	ଷ୍ପ	By close of	Within five	Within 10	Eo. 1	Eo. 1
Pu(∝) Isotopic (IPUL)	Pu-238, Pu-239, 240	0.005		30	business on	business	business	с Г Ц	T
Am-241 (AM241)	Am-241	0.02	0.8	20	datarmination	days of determination	days of determination	, t	L L
Am-243 (AM243)	Am-243	0.02	0.8	20					. .
Cm(∞) Isotopic (ICM)	Cm-242, Cm-244(b)	0.02		20					T
U(∞) Isotopic (IU)	U-233, 234, U-235, U-	0.02		20				- Ż	
Th/	238 Th 228 Th 220 Th 220	ł	•	ç				€	
	Th-232	5	-	3				Eo. 1	E. 1
Tritium (H3)	H-3	20 dpm/ml		5				10dnm/ml	Ì
Sr-total (SR)	Sr (sum Sr-89 + Sr-90)	0		20				5	
Sr-90 (SR90) ^(a)	Sr-90	10		30				5 N.C.	
Gamma Spectroscopy (ISPEC)	K-40, Cs-137 + Others(d)	See Table B-5		20				, ,	
Gamma Spectroscopy	Am-241	'n		20				Е ф	
								Еq. 1	
U-nat (U)	Elemental U	0.06 µg⁄sample	0.3 µg/sample	20				0.2	0.2
<u>Sequential Analyses:</u> Pu(∞) Iso and Sr-total	As for individual analyses	As for individual analyses	ial analyses	25	Asf	As for individual analyses	Ses		
(IPS) Pu(x) Iso. Am-241 (IPA)				25					•
Pu(∝) Iso, Am-241, Sr-total (IPSA)	al (IPSA)			3 23					
Pu(x) Iso, U-nat (IUPU)				25					
Actinide(∞) Isotopic (ITPAC) ^(e)	C)*			25					
Pu(∞) Iso and U ISO (IPIU)				25					

Analytical and Reporting Requirements for Routine Processing of Samples

Table B-3

Battelle Contract 11530 - Feb-06

(a) Time allowed following determination of results to receipt of results by Battelle.

(b) Report measured activity for Cm-246, and Cm-248 upon request of the Battelle Technical Administrator.

(c) If total Strontium is less than 15 dpm, Yingrowth is not required.

(d) Report all isotopes present at levels exceeding Equation 5. If ordered by the Battelle Technical Administrator, report results for radionuclides in Table B-5 specified in the processing instruction, regardless of the activity measured.

(e) Pu (\propto) Isotopic, Am-241, and Cm (σ) Isotopic.

(f) 0.16 dpm for U-234, 0.15 dpm for U-238, and the greater of 0.007 dpm and Equation 5 for U-235.

(g) Oral report required only when analytical results exceed level specified. Eq. 1 Lc=2(combined standard uncertainty)

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Procedure	r IK	<u>4/1/05 throu</u>			SECC	<u>4/1/06 thre</u>		
Code ^(a)	Total	IDP QC	%	GEL QC ^(b)	Total	IDP QC	%	GEL QC ^(b)
Urine	10141	IDF QC	/0	<u>GEL QC</u>	Totai	IDF QC	70	GEL QU
H3	795	16	2.0	274	892	2	0.2	274
						3	0.3	276
SR90, SR	202	1	0.5	476	231	3	1.3	482
C14	0	0 3	0.0	0	0	0	0.0	0
AM241	223		1.3	576	103	0	0.0	437
AM243 U235	114	0 0	0.0	130	85	6	7.1	122
	2	0	0.0	0	0	0	0.0	0
ICM IPU	1687	2	0.0	0	13	0	0.0	241
IPUL	0	2	0.1	1555	1243	0	0.0	1152
			0.0	0	1	0	0.0	0
IPA IPS	357	15	4.2	0	293	4	1.4	N/A
	664	12	1.8	0	553	2	0.4	N/A
IPSA	101	1	1.0	0	152	15	9.9	N/A
IPSR	0	0	0.0	0	0	0	0.0	0
ISPEC	0	0	0.0	0	0	0	0.0	0
ITPAC	96	0	0.0	0	90	0	0.0	N/A
ITH	1	0	0.0	9	0	0	0.0	N/A
IUPU	130	0	0.0	0	108	0	0.0	N/A
IPIU	1	0	0.0	0	4	1	25.0	N/A
IU	517	12	2.3	334	500	14	2.8	279
NP237	0	0	0.0	0	0	0	0.0	0
RA226	0	0	0.0	0	0	0	0.0	0
UNAT	354	11	3.1	557	235	18	7.7	339
LEPD	0	0	0.0	0	0	0	0.0	N/A
PU241	0	0	0.0	0	0	0	0.0	N/A
Total	5252	73	1.4	3911	4503	66	1.5	3328
Fecal ^(c)								
U232	0	0	0.0	0	0	0	0.0	0
ICM	ŏ	Õ	0.0	0 0	ů 1	Ő	0.0	0
ITH	ŏ	0	0.0	0 0	0	0	0.0	Ő
AM241	9	1	0.0	103	15	Õ	0.0	133
IPU	16	0	0.0	99	12	0	0.0	133
IPA	72	9	12.5	0	83	12	14.5	N/A
Total	97	10	12.5	202	111	12		
10141	¥/	10	10.5	202	111	12	10.8	271

TABLE 2. Number and Category of Bioassay Samples Analyzed FIRST CONTRACT YEAR - GEL SECOND CONTRACT YEAR - GEL

^(a)Procedures not specifically tested are evaluated with isotopic results

from other procedures. (b) N/A = not available. QC samples are tracked as isotopic analyses not as multiple analyses.

(c) Analyses not analyzed (IPUBA, IRA, ITPAC, IUPU, UNAT, IU, AM243)

Table 2 presents a breakdown of the numbers and categories for all bioassay samples analyzed, including personnel and QC samples. From 66 urine and 12 fecal QC samples submitted by IDP to GEL during the reporting period, GEL reported 136 analytical urine results for 9 different analytes and 36 fecal results for 3 different analytes. The 78 QC samples represent 1.7% of the total analyses performed by GEL. In addition to these samples, GEL analyzed 3,358 internal QC samples. The QC samples analyzed equaled 36% of the samples analyzed by GEL under their contract with Battelle.

GEL's performance was checked by determining detection level, bias, and precision based on the results of blank and spiked samples. Spiked samples fell into two categories: those spiked near the CL and those spiked at equal to or greater than three times the CL. These two categories were necessary to check compliance with the criteria for relative precision (S_B) specified by the Statement of Work. Satisfying these two categories also verified that GEL could detect sample activities near the CL.

DETECTION LEVELS

Various mathematical expressions and terminology can be used to describe a detection level. The statistical approach specified in the Statement of Work basically follows that of Currie (1968) and HPS N13.30 (HPS 1996). However, the HPS N13.30 formulas were modified to account for the difference between a priori estimates of detection levels based on counts (Currie 1968) and a posteriori estimates based on total activity, where chemical yield is determined specifically for each sample.

Two test criteria were used: the decision level (L_c) and the MDA (also called the detection level). The decision level was defined in the Statement of Work as the quantity of radioactivity or mass above which there is at least 95% confidence that the sample is not a blank (Type I error). If the measured value was greater than the L_c, the sample was considered likely to contain the radionuclide of interest. If the measured value was less than L_c, then the result was considered indistinguishable from a blank. The L_c was determined solely by measuring blank samples. Before the L_c was calculated, results that were significant outliers were eliminated from the data set. Outliers were identified by the use of the criteria of ASTM E178-94 (ASTM 1994). Mathematically, L_c is defined by the following equation:

$$L_c = 2s_A$$

where, s_A equals the combined standard uncertainty of the net analyte reported.

The MDA was based on a 95% confidence in detecting activity when the actual activity was equal to the MDA. Conversely, the 95% confidence level is the point at which only 5% of the results for samples containing activity equal to the MDA fall below the L_c and, thus, were judged to contain no activity (Type II error). The MDA, expressed in units of disintegrations per minute, is calculated from the same set of blanks as the L_c (outliers excluded), using the following equation:

$$MDA = \overline{X_o} + 2(t_{n-1}) s_o + \frac{(t_{n-1})^2}{ERT}$$

where E is the typical counter detection efficiency in counts per disintegration, R is the average fractional chemical recovery or yield, and T is the typical counting time. In keeping with the philosophy of HPS N13.30, if t² is less than 3, then 3 is used instead. For elemental uranium analyses, the analytical method does not produce count data; the unit for the analysis result and MDA is micrograms. Thus, the "3" term is not an appropriate part of the equation for the elemental uranium analysis.

The present contract with GEL, implemented on April 1, 2005 with GEL, specifies an operational year that ends March 31st, each year. This QC report covers the second operational year of that contract, and includes samples analyzed by GEL during period of April 1, 2006 through March 31, 2007.

The MDA values GEL calculates for their QC reports are based on mean values for parameters of equation 2 of the contract statement of work, and not replicate measurements. GEL also uses synthetic samples, whereas IDP uses real fecal and urine samples.

The IDP QC samples were evaluated by first calculating the L_c from blank samples, excluding outliers. This L_c was compared with the L_c calculated from GEL's own QC samples. Then, the MDA was calculated and compared with the CL and the MDA calculated from GEL's own QC samples. Values used for E, R, and T in the MDA equation were obtained from the laboratory; they are listed in Table 3. Finally, the percentage of QC samples spiked at the CL that were measured by the laboratory as having less than the decision level (i.e., no activity was detected) was determined; this percentage was then compared with the 5% allowed in the Statement of Work. Outliers were included in this test.

<u>BIAS</u>

Relative bias is defined as the mean fractional deviation of the reported results from the true values of spikes added to the samples. The formulas in the Statement of Work used to measure bias in sample results are the same as those in HPS N13.30 (1996). The mean relative bias, Br, is determined using:

$$B_r = \sum_{i=1}^m \sum_{j=1}^n \frac{B_{rij}}{N}$$

where n = number of spike samples in each level

m = number of spike levels

N = total number of spiked samples

 B_{rij} = bias of a single measurement, defined as:

$$B_{rij} = \frac{(A_{ij} - A_{ai})}{A_{ai}}$$

where A_{ij} = the jth measured value of the ith spike level, A_{ai} = the true value of the ith spike level

	Nuclide/	Count	Contract	Counter	Efficiency	Chemi	cal Yield
				<u>2005-</u>	2006-	<u>2005-</u>	<u>2006-</u>
<u>Matrix</u>	<u>Method</u>	Minutes	Limit ^(a)	<u>2006</u>	2007	2006	2007
Urine	³Н	30	20	0.24	0.18		
	Total Sr	60	10	0.379	0.396	0.746	0.774
	SR90	60	10				
	²⁴¹ Am	2520	0.02	0.391	0.385	0.883	0.725
	²⁴³ Am	2520	0.02	0.391	0.385	0.877	0.885
	²⁴² Cm/ ²⁴⁴ Cm	2520	0.02	0.391	0.385	0.883	0.725
	²³⁷ Np	2520	0.02				
	²³⁹ Pu/ ²³⁸ Pu	2520	0.02	0.391	0.385	0.956	0.915
	IPUL	10000	0.005				
	²²⁸ Th/ ²³⁰ Th/ ²³² Th	2520	0.1	0.386	NA	0.913	NA
	²³⁴ U/ ²³⁵ U/ ²³⁸ U	2520	0.02	0.386	0.382	0.718	0.709
	Uranium		0.06	NA	NA	NA	NA
Fecal	²⁴¹ Am	96 0	0.8	0.391	0.385	0.657	0.744
	²³⁸ Pu/ ²³⁹ Pu	96 0	0.2	0.391	0.385	0.88	0.90

<u>TABLE 3.</u> Typical Chemical Yield (R), Typical Detector Efficiencies (E), and Counting Time (T) Values from GEL Quality Control Report

(a) Units dpm/sample except dpm/mL for H3, and µg/sample for U.

(b) Only one sample analyzed

(c) NA = Not available. No samples completed.

(d) GEL combined analysis categories with equivalent procedures (e.g., IPU and IPA for plutonium) to improve statistics. Breakdown by analysis is therefore not available.

Outliers were excluded from the test, but not ignored for the procedure evaluation. As stipulated in the Statement of Work, the mean relative bias shall fall within \pm 20% when calculated from 15 to 50 spiked samples, and within \pm 10% when calculated from over 50 samples.

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PRECISION

The precision statistic used for this contract was S_B from HPS N13.30 (1996), but the limits differ from that standard. S_B is given by:

$$S_{B} = \sqrt{\sum_{i=1}^{m} \sum_{j=1}^{n} \frac{(B_{rij} - B_{r})^{2}}{(N-1)}}$$

where the symbols are the same as for relative bias (Br).

The above equation is valid for samples spiked at one or more levels, subject to the limits for the relative precision, which depend on the activity of the spikes relative to the CL. Specifically, the relative precision statistics shall be less than or equal to 0.4 for samples spiked greater than three times the CL and less than or equal to 0.5 for samples spiked between one and three times the CL. Outliers were not included in the determination of precision.

<u>FINDINGS</u>

Results from three types of QC samples were available: 1) those prepared by GEL and analyzed as single-blinds (spike amount unknown to the analyst), 2) those submitted by IDP and analyzed as single-blinds (spike amount unknown to the analyst), and 3) those submitted by IDP and analyzed as double-blinds (spike amount and sample origin unknown to the analyst).

Single-blind samples this year included 10 urines and 6 artificial fecal samples prepared by RPG. The results of the statistical tests (see Table 4 and Appendix A) are discussed below. Statistical results from the present and previous years are compared in Table 5.

OUTLIERS

Analytical results that are biased by "blunders" during the analysis should not be included in the data set used for the statistical evaluation of the analytical procedure, but too many outliers would indicate poor laboratory performance (see Table 6). GEL (see Appendix B) did not identify any outliers and there were no outliers in the IDP data set.

	Sample		Blank (d	pm)		Spike	e level at CI	. (dpm)	Spike	Level > 3CI	_ (dpm)
Isotope ^(a)	Source	<u>n</u>	<u> </u>	<u>MDA</u>	<u>_CL</u>	<u>n</u>	<u>B</u>	<u>S</u> B	<u>n</u>	<u>B</u> ,	<u>S</u> B
³ H(dpm/mL)		i, i			(0		-0.42				
	GEL	138	0.0003	0.6		137	-0.05	0.08			
Total Sr	TDE 3				10		() () () ()	0.23			••••
	GEL	16	0.31	7.13		29	0.05	0.14		•••	
⁹⁰ Sr	GEL	147	0.69	7.95	10	55	-0.03	0.17	231	0.083	0.115
²²⁸ Th	GEL			•••	0.02						•••
²²⁹ Th	GEL		•••		0.02	•••	,				
²³⁰ Th	GEL				0.02						
²⁴² Cm	GEL	31	0.003	0.010	0.02						
²⁴⁴ Cm	GEL	150	0.003	0.010	0.02		•••		60	0.041	0.101
²³⁸ Pu-urine	- me	2000 - 100 -	Megna ti								
1 0 01110	GEL	395	0.003	0.008				* 1.107.0000			* 54 7 - 5 78 -
feces	1DE		0.02	0.0	4 ULA						
	GEL	49	0.02	0.122				444			
²³⁹ Pu-urine	DP -			0.009	JUK	16	20.05				
	GEL	395	0.003	0.009		219	0.02	0.25	90	-0.056	0.049
feces	· IDP.				0^{2}						0.094
	GEL	49	0.02	0.131				· • •	49	-0.054	0.058
²⁴¹ Am-urine	MOP			068(E)-	(),() 9.	anta pro 200 La 1 ₁ 1 da		• • • • • • • • • • • • • • • • • • •			
<u>^</u>	GEL	150	0.004	0.010	·····	83	0.09	0.37	190	-0.011	0.114
feces	(D)		0.03	<u>)</u> ()))))()) () () () () () () () () () () () () () (£=0.8			0.0205			
243 .	GEL	41	0.05	0.254					41	-0.068	0.074
²⁴³ Am-urine			0.003		FU.02						teration of the second s
²³⁴ LF	GEL	42	0.006	0.014		18	0.11	0.31	61	0.026	0.069
0	DP		- 10		0.02						
	GEL	94	0.009	0.022 ^(e)							
²³⁵ U	DR) (18 12)		102, ^{(*})		1		
270	GEL	94	0.005	0.014	NC						and such a particular
²³⁸ U	TDR				a DAU-					0.02	
	GEL	94	0.008	0.019					57 ^(g)	-0.04	0.13
(P)				an cataoraidh				and the second	124 ^(b)	0.004	0.084
U-urine ⁽⁶⁾	DR				0805			n <u>k</u> (e)			0.220
	GEL	198	0.006	0.01		65	-0.04	0.14	67	-0.012	0.076

TABLE 4. Summary of Statistical Values by Nuclide

(a) Analyzed in urine matrix unless otherwise noted.

(b) Units for L_c, MDA, and CL are mg per sample.

- (c) Failed performance criterion.
- (d) Possible environmental contaminant.

(e) Within statistical uncertainty

(f) Stats for Cm same as Am-241

(g) Spike level 0.15 dpm/S

(h) Spike level 0.4 dpm/S

	II OLL USIN	Report	-P0		lanks		Spike Level	at CL	SI	oike Level >	• 3CL
Nuclide	CL	Year	<u>n</u>	L _c	MDA	n	Br	SB	n	B _r	SB
ЗH	20 dpm/mL	2005	5	0.17	0.97	11	-0.04	0.18			
	-	2006	1			2	0.42	0.63		•••	
									(1996)		
Sr	10 dpm	2005	9	0.43	0.97	5	-0.02	0.15	•••		
		2006	1	 		14	0.12	0.23		•••	
					3 x	ندن . د					
U (element	0.06 mg	2005 2006	5 1	•••	0.3571 (d)	6 2	0.01	0.22	 15	 -0.14	0.22
(elemen	iai)	2000	1 	 • • • • • • • • • • • • • • • • • •		<u>ک</u> ۱۹۹۹ کو کو	-1.6 (c)	2.05 (c)		-0.14	0.22
²³⁵ U	0.02 dpm	2005	2	0.01	0.0687 (c)	10	-0.10	1.0307 (c)	1		
U	0.02 up m	2003	12	0.01	0.0087 (0)	3	-0.10 -0.24(c)	0.3			•••
The second second				(-) (1	0.02						•••
²³⁸ U	0.02 dpm	2005	2	0.15	0.3508 (d)	10	0.15	0.33	a a a a a a a a a a a a a a a a a a a		•
U	0.02 dpm	2005				0	0.00	0.00	 15	0.02	0.23
						· · · · · · · · · · · · · · · · · · ·					
²³⁸ Pu	0.02 dpm	2005	21	0.003	0.009	9	0.17	0.10			
(urine)	F	2006	21	0.004	0.011	ī	-0.18				
	State State										
²³⁹ Pu	0.02 dpm	2005	21	0.003	0.008	9	0.14	0.33			
(urine)	-	2006	6	0.002	0.009	16	0.05	0.23		•••	
						an a					
230_			_	0.02							
²³⁹ Рц	0.2 dpm	2005	5	5 0.02	0.06	•••			4	-0.02	0.12
(fecal)		2006	6	0.02	0.07				6	-0.05	0.09
							 sk 2				
²⁴¹ Am	0.02 dpm	2005	10	0.004	0.012	8	0.05	0.23			
(urine)	0.02 0011	2005	2	0.005	0.068(c)	17	0.19	0.35	••••	•••	•••
									×		
²⁴¹ Am	0.02 dpm	2005	5	0.03	0.085	4	-0.12	0.13			an a ra an b Bhank () dhifte dhifte d
(fecal)		2006	8	0.025	0.063	4	-0.17	0.09			
741				0.02	_						
²⁴³ Am	0.02 dpm	2006	2	0	0.09	0					

TABLE 5. Comparison of Quality Control Statistics Between the First and Second Contract Year with GEL Using QC Samples Submitted by IDP

Note: L_c and MDA units same as CL. B_r and S_B are unitless (fractional values).

			Spikes at	
			Contract	FALSE Negatives
Nuclide	Analyses	Outliers	Level	(%)
Urine				
³ H	3	0 (0)	2	0 (0)
⁹⁰ Sr	15	0 (0)	14	0 (0)
²³⁵ U	15	0 (0)	3	0 (0)
²³⁸ U	0	0 (0)	0	0 (0)
²³⁸ Pu	22	0 (0)	1	0 (0)
²³⁹ Pu	22	0 (0)	16	0 (0)
²⁴¹ Am	19	0 (0)	17	0 (0)
²⁴³ Am	0	0 (0)	0	0 (0)
UNAT	3	0 (0)	15	0 (0)
Total	99	0 (0)	68	0 (0)
Feces				
²⁴¹ Am	12	0 (0)	4	0 (0)
²³⁸ Pu	12	0 (0)		0 (0)
²³⁹ Pu	12	0 (0)	6	0 (0)
Total	36	0 (0)	10	0 (0)

TABLE 6. Other Indicators of Analytical Uncertainty (IDP Samples)

Smillor at

TRITIUM

Effective June 2006, the tritium intercomparison program by IDP was discontinued, performance indicators will be evaluated through GEL's QC program. Prior to June 2006, 3 tritium intercomparison samples were submitted by IDP, 1 blank and 2 spiked at the CL, the relative bias and precision statistics were all within acceptable values. The control samples run by GEL also met all the acceptance criteria tested as part of the quality control program. The tritium analyses were considered acceptable.

STRONTIUM-90 AND TOTAL STRONTIUM

The total strontium procedure is used to screen samples to determine which will require analysis for ⁹⁰Sr. Samples with total strontium results less than 15 dpm do not undergo further analysis. Samples with results greater than or equal to 15 dpm may undergo ⁹⁰Y in growth to specifically determine ⁹⁰Sr levels. The calculated MDA, as reported by GEL, for the total strontium part of the analysis was about 30% of the CL.

The relative bias and precision, tested by IDP and GEL for the ⁹⁰Sr and total Sr procedures were all within limits. The 14 samples spiked at the contractual level were all detected. The strontium urinalysis procedure was concluded to be acceptable.

PLUTONIUM-238 AND -239

Samples spiked with ²³⁸Pu and ²³⁹Pu were analyzed using the same procedures and same reagents. The two isotopes are differentiated only at the end of the procedure by alpha spectrometry. Therefore, laboratory performance is expected to be similar for both isotopes using any of the seven procedures that incorporate plutonium analysis (IPU, IPA, IPS, IPSA, IPSR, IUPU, and ITPAC).

The MDAs and performance statistics for ²³⁹Pu and ²³⁸Pu in urine were acceptable. The 16 samples spiked at the CL for ²³⁹Pu were reported with a result greater than the decision level and the 6 blank samples were reported with results less than the decision level. With the exception of one sample, the 21 blank ²³⁸Pu samples were reported with results less than the decision level, giving a false positive rate less than 5%. Overall the plutonium urinalyses were considered acceptable.

The MDA and performance statistics for ²³⁹Pu and ²³⁸Pu in feces were acceptable. Approximately 15% of the fecal samples analyzed were duplicated to test the consistency of the aliquoting procedure. A review of the duplicate samples determined that the aliquoting procedure produced results within 3 sigma of the initial results. The fecal aliquoting procedure was acceptable. None of the 12 blank ²³⁸Pu or the 6 blank ²³⁹Pu fecal analyses were greater than the decision level. There were no fecal samples spiked at the CL with ²³⁸Pu. The six fecal samples spiked with ²³⁹Pu were reported with a result greater than the decision level. Overall the plutonium fecal analyses were considered acceptable.

URANIUM (UNAT)

No concerns were identified with the elemental uranium urinalysis program and it was considered acceptable. The bias and precision for uranium spiked at 0.06 µg, as tested by IDP, did not meet the acceptance criteria. However, only 2 samples were tested and environmental contamination was thought to be an interference. Because IDP uses a 0.2 µg screening level for elemental uranium, samples spiked at 0.06 µg were discontinued. The MDA at the contractual level of 0.06 µg was evaluated through GEL's program and was found to be acceptable. The bias and precision was tested by IDP at 0.2 µg and by GEL at 0.1 µg and was also found to be acceptable. For the third contract year, GEL will begin evaluating the bias and precision for elemental uranium at levels of 0.06 µg, in accordance with the contractual level.

ISOTOPIC URANIUM

The isotopic uranium analysis reports on three uranium isotopes: ²³⁴U, ²³⁵U, and ²³⁸U. The isotopes are differentiated only during counting by alpha spectrometry. GEL reported that the calculated minimum detectable activity (MDA) for ^{233,234}U for the year slightly exceeded the contract required detection limit. This was consistently seen in the first, second and third quarters. The cause was attributed to low tracer recoveries. The standard operating procedure was revised November 30, 2006 to improve the separation chemistry. GEL then reviewed the MDAs for the fourth quarter and showed that all three isotopes met the acceptance criteria.

IDP did not submit isotopic uranium samples to review the ^{233,234}U performance indicators. However, the performance statistics for ²³⁵U and ²³⁸U were reviewed and the MDA for ²³⁵U and the bias and precision for ²³⁸U were acceptable. The percentage of analyses with low yields were reviewed and found not to meet the criteria in the statement of work for the first, second and third quarters (Table 7). In line with GEL's changes to the standard operating procedure, the low yield rate in the fourth quarter was significantly improved upon and it met the criteria in the statement of work. A review of the data from April 1, 2007 through December 31, 2007, also showed improvement in tracer recoveries for isotopic uranium urinalyses. Of the 398 urinalyses processed during this time, all had tracer recoveries at 40% or greater. With the implementation of the revised standard operating procedure, the isotopic uranium analysis program is considered acceptable.

<u>**TABLE 7**</u>. Low Yields for Isotopic Uranium for the Second Contract Year (4/1/2006 - 3/31/2007) as Broken Down by Quarter. Also included are the summed results for the first few quarters of the Third Contract Year.

First	Quart	er	Secon	d Qua	arter	Third	Quar	ter	Fourt	h quai	rter	3/1/07 -	12/31/	07
Total	Low	Yield	Total	Low	Yield	Total	Low	Yield	Total	Low	Yield	Total	Lo Yie	
Results			%	Results	No.	%	Results	No.	%	Results	No.	%		
121	23	19%	157	35	22%	152	19	13%	148	1	0.7%	398	0	0%

AMERICIUM-241

The ²⁴¹Am fecal and urine analysis met the acceptance criteria for MDA, relative bias and precision. With regards to the ²⁴¹Am urinalysis program there were only 2 blank samples and the resulting MDA exceeded the acceptance criteria, however, this was most likely a result of low-level counting statistics. The MDA as reported by GEL was 50% of the contractual level. The 17 samples spiked by IDP at the contractual detection level were all greater than decision level. The current AM241 urinalysis procedure was considered acceptable. All six blank fecal samples were less than the decision level and the four spiked fecal samples were all greater than the decision. The ²⁴¹Am fecal duplicate samples were evaluated and it was concluded that the aliquoting procedure produced results within the control limits. A review of the routine ²⁴¹Am results reported, however, consistently found more than 10% of the samples with tracer recoveries at or below 40% (Table 8). GEL reported that the overall low yield for AM241 fecal analyses was 7.4%. However, GEL was including duplicate samples in their tally, a mistake which will be corrected in subsequent QC reports.

Table 8. Low Yields for AM241 Fecal Analyses for the Second Contract Year (4/1/2006 - 3/31/2007) as Broken Down by Quarter. Also included are the summed results for the first few quarters of the Third Contract Year.

First	Quart	er	Second	Quarte	er	Third	Quarte	ər	Fourth	n quar	ter	3/1/07 -	12/31/	07
Total	Low	Yield	Total	Low	Yield	Total	Low	Yield	Total	Low	Yield	Total	Lo Yie	
Results	No.	%	Results	Results	No.	%	Results	No.	%	Results	No.	%		
12	3	25%	2	0	0%	20	5	25%	64	16	25%	36	1	3%

In mid-January 2007, GEL identified a cause for the low tracer recoveries in the La(F₃) precipitation step. GEL modified their standard operating procedure to include an additional 1 mL of HF when performing the La(F₃) step. This correction appeared to have improved the tracer recovery rate. However, in early March there was a batch of 10 fecal samples that were declared failed analyses due to a technician error. To better evaluate the efficiency of the procedure modification fecal analyses from the first few quarters of the third contract year (3/1/07 - 12/31/07) were reviewed and only 3% of the samples had tracer recoveries less than 40%. With the implementation of the revised standard operating procedure, the AM241 fecal analysis program is considered acceptable.

AMERICIUM-243

The AM243 procedure was identical to the AM241 procedure, except that a different tracer is used (²⁴⁴Cm instead of ²⁴³Am). The 6 blank ²⁴³Am QC samples submitted were all reported with results less than the decision level and the calculated MDA was 50% of the contractual detection level. The performance statistics for the AM243, as tested by GEL, met the acceptance criteria. The AM243 procedure was concluded to be acceptable.

ISOTOPIC CURIUM

IDP did not submit QC samples to test the isotopic curium program, therefore performance statistics were based on the GEL QC results. GEL tested the MDA for ²⁴²Cm and ²⁴⁴Cm and the relative bias and precision for ²⁴⁴Cm. The results met the acceptance criteria and the isotopic curium urinalysis program was considered acceptable.

ISOTOPIC THORIUM

During the second contract year, no isotopic thorium analyses requested. Therefore, there were no QC samples to evaluate.

FOLLOW-UP ON CONCERNS DURING THE SECOND CONTRACT YEAR

There were a few concerns during the second year with General Engineering Laboratories (GEL) dealing mainly with low tracer recoveries seen in the isotopic uranium urinalyses and AM241 fecal analyses. Steps taken by GEL to address both concerns have been sufficient and tracer recoveries are now within acceptable limits (see discussions above).

A minor observation made was that the GEL calculated MDAs for ²³⁹Pu, ²³⁸Pu and ²⁴¹Am in feces were between 2 to 4 times greater than the IDP's calculated MDAs. The MDAs for urine were similar between the two QC programs. When comparing the urine and fecal calculated MDAs for the 3 isotopes, a factor of 5 difference was expected based on the shorter counting time and aliquoting associated with fecal analyses. However, the fecal MDAs reported by GEL were 15 times greater in fecal analyses than urinalyses for ²³⁹Pu and ²³⁸Pu and about 25 times greater in ²⁴¹Am fecal analyses than urinalyses. GEL is currently investigating this concern and will be evaluated in the third contract year.

A review of Incident reports since the contract with GEL was initiated did not identify a trend or a concern. The majority of incident reports were due to human error and corrective actions were deemed acceptable.

Incident reports issued during the second contract year included:

- 1. Incident Report for Tagword 06E0067, 06E0184, 06E0233, 06E0319, 06E0325, 06E0327 and 06E0338. closed May 24, 2006
 - a. GEL was unable to report the volume received for the above referenced Tagwords as the analyst had recorded the volumes on a loose sheet of paper, which was inadvertently discarded. A bound logbook, RC-O-024, was created to permanently record sample volumes.
- 2. Incident for Tagword 06G0232 and 06G0042, closed August 31, 2006
 - a. The analyst neglected to add the ²⁴⁴Cm tracer to the sample tagwords referenced. The group leader met with the analysts and their team leader and discussed the error and stressed the importance of following the information provided in the que and pull sheets. In addition, the team leader was to use additional identifying methods for infrequent analyte requests.

SUMMARY OF THE BIOASSAY QUALITY CONTROL REPORT FROM GEL INCORPORATED, FOR THE CONTRACT 11530 SECOND YEAR 2006/2007⁽¹⁾

GEL reported all analytical batches were analyzed with a reagent blank, matrix blank or both. GEL considered blanks in control when the calculate MDA was less than the Contract Limit (CL) and the L_e was less than $\frac{1}{2}$ CL (see Appendix B). In addition, the chemical tracer yields were evaluated against the yield requirements stated in the subject contract. Overall, GEL believed that the blank and spike data for each analytical process demonstrated that the analyses were in control.

GEL reported that the duplicate fecal samples were evaluated to determine that the aliquot procedure produces results within control limits. One ²⁴¹Am duplicate was excluded because the results of the parent sample were not reported. Two ²⁴³Am tracers were less than the low yield requirement of 40% for isotopic americium. One of the two tracers was below the minimum yield requirement of 20%; however, the sample was a Laboratory Control Sample (LCS) that met the recovery requirements. No correlation between the samples was observed, therefore, no corrective action was necessary. Two isotopic curium samples were not spiked; however, they were spiked with ²⁴¹Am for the americium portion of the analysis. Since curium and americium are from the same counting source, the curium results were reported based on the acceptable ²⁴¹Am recoveries. A non-conformance reported was created to document this anomaly.

Fourteen tracers were less than the 50% low yield level for isotopic plutonium. No correlation between the samples was observed so no corrective action was necessary. Two ²⁴²Pu tracers were observed to be below the minimum yield requirement of 25%. The failed tracers were for two LCSs with spikes within the recovery requirements, so the results were reported. A non-conformance reports was created to document the anomaly.

Eighteen tracers were less than the 50% low yield for ⁹⁰Sr. No correlation between the samples was observed, so no corrective action was necessary.

Seventy tracers were less than the 40% low yield for isotopic uranium, in addition the MDA for ²³⁴U was slightly greater than the contractual level. The Standard Operating Procedure was revised November 30, 2006 as Revision 25. The revision made involved changing the type of separation chemistry from an AG1X8 anion exchange column to an Eichrom TRU extraction chromatography column. This was discussed in the section on isotopic uranium. Improvements were observed following the corrective actions.

⁽¹⁾ Summaries are taken from Pifer (2007).

RESULTS FROM INTERCOMPARISON PROGRAMS

GEL participated in 3 intercomparison programs (Attachment D – Intercomparison Programs) in the second contract year. On April 1, 2006 they participated in the National Institute of Standards and Technology's program testing the relative bias and precision for ²³⁸Pu, ²³⁹Pu, ²⁴¹Am, ^{230Th}, ²³⁵U, ²³⁸U, ²³⁴U and ⁹⁰Sr in synthetic feces. GEL met the acceptance criteria for relative bias and precision for all isotopes except for ⁹⁰Sr, which failed the portion on relative bias but passed on relative precision. Because Hanford does not use fecal samples for strontium analyses, this was not deemed a concern. GEL also participated in the National Institute of Standards and Technology's program testing the relative bias and precision for ²³⁸Pu, ²³⁹Pu, ²⁴¹Am, ²³⁰Th, ²³⁵U, ²³⁸U, ²³⁴U, ⁹⁰Sr, ⁶⁰Co, ¹³³Ba, ¹³⁷Cs and ¹⁵²Eu in synthetic urine. GEL met the acceptance criteria for relative bias and precision for

On December 1, 2006 GEL participated in the Department of Energy Laboratory Accreditation Program, Session 10. Isotopes tested in a fecal matrix were ²³⁸Pu, ²³⁹Pu, ²⁴¹Am, ²³⁰Th, ²³²Th, ²²²⁸Th ²³⁸U, ²³⁴U, ⁹⁰Sr, ⁶⁰Co, and ¹³⁷Cs. Isotopes tested in a urine matrix were ²³⁸Pu, ²³⁹Pu, ²⁴¹Am, ²³⁰Th, ²³²Th, ²²²⁸Th, ²³⁸U, ²³⁴U, ⁹⁰Sr, ⁶⁰Co, ¹³⁷Cs, tritium and U-total. GEL passed the acceptance criteria for all isotopes in both the fecal and urine matrix for relative bias and precision.

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APPENDIX A

QUALITY CONTROL SAMPLE RESULTS

REL BIAS								960	-0.1033	-0.1600	-0.3000	-0.1033	-0.1667 0.0928															960				
RE								Time	Ŧ	Ŷ	Y	Ŷ	1															Time				
DET -	,		•			,		Chem Yield 0.74 Det Eff 0.39	+	+	+	+	Mean Rel. Bias Mean Rel. Precision			,		·		ı	,	,						Chem Yield 0.90 Det Eff 0.39		ı	ι	ı
<u>UNCERT</u> <u>1</u> 0.0134	0.0198	0.0583	0.0158	1610.0	0.0153	0.0044	0.0034	0.0251 0.0633	0.1800	0.1690	0.1430	0.1800			0000	6400'D	0.0057	0.0069	0.0098	0.0042	0.0052	0.0044	0.0060	0.0182	0.0048	0.0099	0.0069	0.0186 0.0470		0.0086	0.0057	0.0334
RESULT 0.0104	-0.0099	-0.0240	-0.0099	-0.0099	0.0165	0.0053	0.0011	3 MDA	0.8070	0.7560	0.6300	0.8070	120		2000.0	CCUU-0-	-0.0035	-0.0026	0.0144	0.0035	-0.0026	0.0036	0.0066	0.0321	-0.0015	0.0122	0.0076	5 DL 4 MDA		0.0152	-0.0107	-0.0166
<u>MR</u> –	-	ŗ	٦	ŗ	ŗ	ŗ	5	-0.0026 0.0133	5	-	ŗ	-	0.7500		-	-	-	-	ŗ	ŗ	ŗ	ŗ	ŗ	ŗ	ŗ	ŗ	ŗ	0.0055 0.0104		-	ŗ	-
<u>TYPE</u> F	ц	ц	ц	ъ	Ľ.	ц.	ц	sult	ц	ч	ц	ц	ssult		F	L,	ł۲.	ц	ц	ц	ц	ц	ц	ч	ц	ц	ц	sult		٤	<u>11</u>	íL.
<u>UNCERT</u> 0.0000	0.0000	0.0000	0.000	0.0000	0.0000	0.0000	0.0000	Average Result St Dev	0.0450	0.0450	0.0450	0.0450	Average Result St Dev		0000	0.000	0.0000	0.0000	0.000	0.0000	0.0000	0.000	0.0000	0.0000	0.0000	0.0000	0.0000	Average Result St Dev		0.0000	0.0000	0.0000
0.000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.9000	0.9000	0.9000	0.9000	0.9000		00000	0,000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000		0.0000	0.0000	0.0000
67.3 VOL	245.	205.	186.	160	41.3	51.8	51.7	•	50.2	50.3	51.8	51.6			ţ	C.10	245.	205.	160.	41.3	186.	50.2	51.8	51.7	50.3	51.6	51.8			67.3	245.	205.
REQ PA	ΓPA	ΓA	ΓA	ΓA	ГРА	IPA	IPA	AM241 Count	IPA	IPA	ΓPA	ΓA	AM241 Count		·	IFA	ΡA	ΓA	ΓA	IPA	ΡA	IPA	IPA	IPA	ΓA	IPA	IPA	PU238 Count		ΓA	IPA	IPA
	59882	99166	08166	99166	08166	AU001	AU001	 ≞	AU001	AU001	100UA	AU001	 1		00007	79960	59882	99166	99166	99180	08166	AU001	AU001	AU001	AU001	AU001	AU001	н н		59882	59882	99166
TAGWORD PAYID 06K0411 59882	06L0401	06L0414	06L0412	06L0415	06L0413	06L0437	06L0438	90 9	06L0435	06L0433	06L0434	06L0436	4	12		U6KU411	06L0401	06L0414	06L0415	06L0413	06L0412	06L0435	06L0434	06L0438	06L0433	06L0436	06L0437	12	12	06K0411	06L0401	06L0414
ANAL DATE 1 12/13/2006	12/13/2006	01/08/2007	01/08/2007	1/08/2001	01/11/2007	01/16/2007	01/16/2007		01/16/2007	01/16/2007	01/16/2007	01/16/2007		Number of total F AM241		0002/61/21	12/13/2006	01/05/2007	01/05/2007	01/05/2007	01/05/2007	01/11/2007	01/11/2007	01/11/2007	01/11/2007	01/11/2007	01/11/2007		Number of total F PU238	12/13/2006	12/13/2006	01/05/2007
<u>YRMOSEO</u> 0612 17	18	19	21	20	22	15	16		Ξ	12	13	14		r of tot	ţ	1	18	19	20	52	21	П	13	16	12	14	15		er of to	17	81	61
<u>YRM(</u> 0612	0612	0612	0612	0612	0612	0612	0612		0612	0612	0612	0612		Numbe		0612	0612	0612	0612	0612	0612	0612	0612	0612	0612	0612	0612		Numb	0612	0612	0612
<u>ISO CD</u> AM241	AM241	AM241	AM241	AM241	AM241	AM241	AM241		AM241	AM241	AM241	AM241				FU238	PU238			PU239	PU239	PU239										

Run Date 1/29/2008

Analysis dates from 4/1/2006 to 3/31/2007

QC SUMMARY REPORT

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REL BIAS			Time 960	0.0100	-0.1580	0060.0	-0.0560	-0.1430	-0.0200	-0.0462 0.0942				Time 2520	0.1100	0.0600	0.3150	0.3350	0.1600	-0.2050	0.1100	-0.1650	0.0700	-0.3850	0.5250	0.8100	0.6550	0.8500	0.2050	-0.1600	-0.0200	0.1924 0.3544
<u>DET</u>	,	•	Chem Yield 0.90 Ti Det Eff 0.39	+	+	+	÷	+	+	Mean Rel. Bias Mean Rel. Precision		,		Chem Yield 0.73 T Det Eff 0.39	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+	Mean Rel. Bias Mean Rel. Precision
<u>UNCERT</u> <u>D</u> 0.0066	0.0084	0.0052	0.0273	0.1130	01010	0.1200	0.1120	0.1000	0.1260			0.0002	0.0021	0.0054	0.0067	0.0071	0.0074	0.0079	0.0067	0.0053	0.0071	0.0060	0.0065	0.0047	0.0088	0.0096	0.0093	0.0100	0.0071	0.0055	0.0062	
RESULT 0.0096	0.0123	-0.0074	MDA	1.0100	0.8420	0060.1	0.9440	0.8570	0.9800	82		-0000-0	0.0003	03 DL 06 MDA	0.0222	0.0212	0.0263	0.0267	0.0232	0.0159	0.0222	0.0167	0.0214	0.0123	0.0305	0.0362	0.0331	0.0370	0.0241	0.0168	9610.0	38
<u>, MR</u>	7	ŗ	0.0004	٦	~	ŗ	-	-	-	0.9538		-	-	0.0003	7	L	L	L	щ,	Г	Г	Г	Г	Ч	Ч	L	<u>ц</u>	-1	Ч	Г	Г	0.0238
TYPE	μ.	ц	Result	ы	۲.	і т.	ц	щ	Ŀ,	Result		D	Þ	Result	ŋ	D	٦	D	D	D	þ	D	n	D	D	ŋ	n	D	D	D	D	Result
UNCERT 0.0000	0.0000	0.0000	Average Result St Dev	0.0350	0.0350	0.0350	0.0350	0.0350	0.0450	Average Result St Dev		0.0000	0.0000	Average Result St Dev	0.0006	0.0006	0.0008	0.0005	0.0005	0.0005	0.0005	0.0005			0.0006	0.0006	0.0006	0.0006	0.0005	0.0005	0.0005	Average Result St Dev
0.0000	0.0000	0.000	0.0000	0000.1	1.0000	1.0000	1.0000	1.0000	1.0000	1.0000 6		0.0000	0.0000	0.000	0.0200	0.0200	0.0200	0.0200	0.0200	0.0200	0.0200	0.0200	0.0200	0.0200	0.0200	0.0200	0.0200	0.0200	0.0200	0.0200	0.0200	0.0200
107 109	41.3	186.	ł	50.2	51.6	50.3	51.8	51.8	51.7	1		1184	1346	I	0982	2660	1218	1211	1183	1292	1090	1458	1087	1267	1151	1235	1426	1155	1371	1101	1180	1
			PU239 Count							PU239 Count				AM241 Count																		AM241 Count
REQ PANE PANE	IPA	ΓA		IPA	ľΡΑ	IPA	ΓЪ	ΡA	ΡA			ΡA	IPA		IPA	IPA	IPSA															
<u>1 PAVID</u> 99166	08166	08166	۱ ۴	AU001	AU001	AU001	AU001	AU001	AU001	' E		91382	3C135	Þ	59001	3G544	50575	50807	99156	59621	50809	31776	99159	32533	99162	51077	80108	19166	3C134	3C142	99158	Þ
TAGWORD PAVID 061.0415 99166	06L0413	06L0412	¢	061.0435	06L0436	06L0433	06L0437	06L0434	06L0438	Ŷ	12	06E0144	06E0058	4	06C0318	06D0152	06E0102	06F0121	06F0295	06F0294	06F0122	06F0223	06G0241	06G0067	6010H90	06H0140	06H0115	06J0051	07A0418	07A0417	07A0419	17
ANAL <u>DATE</u> <u>T</u> 01/05/2007		01/05/2007		01/11/2007	01/11/2007	01/11/2007	01/11/2007	01/11/2007	01/11/2001		Number of total F PU239	05/26/2006	05/26/2006		04/13/2006	05/01/2006	06/03/2006	01/22/2006	01/22/2006	07/22/2006	07/22/2006	07/30/2006	08/01/2006	08/14/2006	08/28/2006	08/28/2006	08/28/2006	11/06/2006	01/24/2007	01/24/2007	01/24/2007	
<u>YRMO SEO</u> 0612 - 20	ដ ដ	21		Ξ	14	12	15	13	16		er of to	90	07		Π	05	60	05	60	08	90	01	10	60	8	11	01	07	60	10	Π	
<u>YRM(</u>	0612	0612		0612	0612	0612	0612	0612	0612		Numb	0605	0605		0602	0601	0605	000	0606	0000	0006	0090	0607	0607	0608	0608	0608	0610	0701	0201	0701	
<u>ISO CD</u>	PU239	PU239		PU239	PU239	PU239	PU239	PU239	PU239			AM241	AM241		AM241																	

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REL BIAS		Time 2520	Time 20	0.8650 -0.0300	0.6329															
DET		- Chem Yield 0.73 Det Eff 0.39	- Chem Yield 1.00 Det Eff 0.18	+ +	Mean Rel. Bias Mean Rel. Precision	,			,	1 1	,	,	,		+	,		,	ı	
UNCERT D	0.0017 0.0001 0.0001 0.0010 0.0008 0.0024	0.0016 0.0106 0.0106	200	0.7810 0.8110	L	0.0016	0.0022 0.0001	0.0001	0.0021	0.0019	0.0019	0.0023	0:001	0.0021	0.0026	0.0019	0.0023	0.0027	0.0008	0.0001
RESULT	-0.0003 -0.0025 -0.0025 0.0007 0.0001 -0.0023	-0.0011 11 DL 13 MDA		37.3000 19.4000	華 []	0.0013	0.0021 0.0000	0.0000	-0.0012	0.0009	-0.0013	0.0019	-0.0002	0.0028	0.0055	0.0008	0.0023	0.0022	-0.0034	0.0000
MR		0.0013	-0.0351 0.0000		######	5	<u>م</u>	ፈ	-, -		ŗ	Г	L	ц.	Ц	Г	-1	_	Ц	в
IYPE			sult o	n	sult	Ð	n D	n	n II	, D	n	n	D	D :	;	⊃	D	þ	n	D
UNCERT	0.0000 0.0000 0.0000 0.0000 0.0000 0.0000 0.0000	0.0000 U Average Result St Dev 0.0000 U	Average Result St Dev	0.1780 0.1480	Average Result St Dev	0.0000	0.0000 0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
SPIKE	0.0000 0.0000 0.0000 0.0000 0.0000 0.0000	0.0000 0.0000 0.0000 Average 7 St Dev	0.0000		20.0000		0.0000		000000		0.0000									0.0000
TOT	1097 1407 1407 1247 1305 1300	1171		0502 0500		0982	0660	1239	1184	112	1292	183	0601	1458	1087	1267	1151	1235	1426	1281
REQ <u>ANAL</u>	AM243 AM243 AM243 AM243 AM243 AM243	AM243 AM243 Count	H 3 Count		Count Count		PS PA	IPS	PSA PSA	PSA	IPSA					• • •				[DId
<u>OLVA</u>	99120 91386 91386 80098 3C136 32514	99152 U 3 3C136	Þ	32533 3C136	2	59001	32476 3G544	99153	91382 50575	50807	59621	99156	50809	31776 90150	99159 22522	32533	99162	51077	80109	15166
TAGWORD PAYID	06D0145 06D0146 06D0146 06G0247 06H0222 06H0222	0610159 6 06D0142	-	06D0052 06E0059	3	06C0318	06C0067 06D0152	06D0252	06E0144 06E0102	06F0121	06F0294	06F0295	06F0122	06F0223	0600241	06G0067	00H0H00	06H0140	06H0115	06H0304
ANAL <u>DATE</u> al U AM241	04/28/2006 04/28/2006 04/28/2006 07/30/2006 08/28/2006 09/28/2006	0609 02 09/28/2006 Number of totai U AM243 0604 02 04/20/2006		04/20/2006 05/22/2006	Number of total U H 3	04/06/2006	04/19/2006 04/24/2006	04/26/2006	05/19/2006 05/24/2006	07/21/2006	07/21/2006	07/21/2006	07/21/2006	07/28/2006	9007/10/20	08/14/2006	08/28/2006	08/28/2006	08/28/2006	08/31/2006
<u>YRMO SEQ</u> 4umber of tot	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	02 r of totz 02		01	ber of t	= :	88	10	88	05	80	60	8 8	01	2 8	6 8	8:	= :	01	80
<u>YRM</u> Numbe	0604 0604 0604 0607 0608 0609	0609 Numbel 0604		0604 0605	Num	0602	0601 0601	0604	0605 0605	0606	0606	9090	0090	0606	/ 090	1000	0608	0608	8090	0608
ISO CD	AM243 AM243 AM243 AM243 AM243 AM243 AM243	AM243 H 3		H 3 H 3		PU238	PU238 PU238	PU238	PU238 PU238	PU238	PU238	PU238	PU238	PU238	FU258	FU238	PU238	PU238	PU238	PU238

<u>REL BIAS</u>	Time 2520	-0.1750								Time 2520	0.0100	0.0400	0.3550	-0.2050	0.1900	0.1050	-0.3500	-0.2750	0.0100	0.4250	0.1100	0.1150	-0.0600		0.4100	-0.1150	0.2265
<u>DET</u> , ,	Chem Yield 0.92 Det Eff 0.39	⊤ Mean Rel. Bias Mean Rel. Precision			ı		ı	ı		Chem Yield 0.92 ' Det Eff 0.39	+	+	+	+	+	+	+	+	+	+	÷	+	+	+	÷	+	Mean Rel. Bias Mean Rel. Precision
UNCERT 0.0024 0.0019 0.0031 0.0001	0.0036 0.0106			0.0028	0.0019	0.0020	0.0001	0.0001	0.0021	0.0020 0.0087	0.0054	0.0056	0.0058	0.0047	0.0059	0.0051	0.0044	0.0045	0.0056	0.0062	0.0058	0.0055	0.0052	0.0053	0.0064	0.0055	
RESULT 0.0011 0.0059 0.0000	1 MDA			0.0008	0.0003	0.0014	-0.0006	-0.0007	0.0017	S DL	0.0202	0.0208	0.0271	0.0159	0.0238	0.0221	0.0130	0.0145	0.0202	0.0285	0.0222	0.0223	0.0188	0.0200	0.0282	0.0177	100
M H L L L L	0.0010 0.0021	0.0000		'n	<u>م</u> ,	Г	д.	ſ	Г	0.0005	-	Г	Г	Г	Ч	Ч	L	L	L		Ц	Ħ	L	ч	<u>.</u> _	Г	0.0210 0.0045
U U U	sult	sult		n	D	D	Ŋ	n	n	sult	D	n	D	D	D	D	D	D	D	D	D	n	n	D	D	n	sult
UNCERT 0.0000 0.0000 0.0000 0.0000	Average Result St Dev 0.0003 11			0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	Average Result St Dev	0.0004	0.0003	0.0003	0.0003	0.0003	0.0003	0.0059	0.0059	0.0003	0.0003	0.0003	0.0003	0.0003	0.0005	0.0005	0.0005	Average Result St Dev
SPIKE 0.0000 0.0000 0.0000 0.0000	0.0000 21	0.0200		0.000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000 6	0.0200	0.0200	0.0200	0.0200	0.0200	0.0200	0.0200	0.0200	0.0200	0.0200	0.0200	0.0200	0.0200	0.0200	0.0200	0.0200	0.0200
<u>VOL</u> <u>S</u> 1155 1101 1101 1180	1346 0	18		0982	0660			1184	1218	9	346	1211 (1090	1183 (1292 (1458 (1087	1267 (iisi (1235 (1426 (281 (1155 (1371 (101 0	1180 (0
REQ ANAL IPSA IPSA IPSA IPSA IPSA IPSA IPSA IPSA	PU238 Count	PU238 Count		IPA 0	IPS 0			IPA 1	IPSA 1	PU239 Count	IPA 1	IPSA I	IPSA I	IPSA 1	IPSA L	IPSA I	IPSA I	IPSA I	I VSA	IPSA 1.	IPSA I	1 . 1. 1. 1.	IPSA I	IPSA II	IPSA 1	IPSA 1	PU239 Count
PAYID 99161 3C134 3C134 3C142 99158	U 3C135	י ב		59001	32476	3G544	99153	91382	50575		3C135	50807	50809	99156	59621	31776	99159	32533	99162	51077	80109	99151	19166	3C134	3C142	85166	
TAGWORD PA 06J0051 9914 05J0051 9914 07A0418 3C1 07A0417 3C1 07A0419 9914	21 06E0058	1	12	06C0318	06C0067	06D0152	06D0252	06E0144	06E0102	¢	06E0058	06F0121	06F0122	06F0295	06F0294	06F0223	06G0241	06G0067	0010H90	06H0140	06H0115	06H0304	06J0051	07A0418	07A0417	07A0419	16
ANAL DATE DATE 11/03/2006 01/23/2007 01/23/2007	05/19/2006		Number of total U PU238	04/06/2006	04/19/2006	04/24/2006	04/26/2006	05/19/2006	05/24/2006		9002/61/50	07/21/2006	01/21/2006	9007/17/20	9002/12/20	07/28/2006	08/01/2006	08/14/2006	08/28/2006	08/28/2006	08/28/2006	08/31/2006	11/03/2006	01/23/2007	01/23/2007	01/23/2007	
XRMO SEQ 0610 07 0701 09 0701 10 0701 11	01		sr of tot	11	60	3	10	96	60		07	05	90	60	08	07	9	60	60	11	10	80	01	60	10	1	•
YRMG 0610 0701 0701 0701	0605		Numb	0602	0602	0601	0604	0605	0605		0605	0606	0606	9090	9090	9090	0607	0607	0608	0608	0608	0608	0610	0201	0701	0701	
ISO CD PU238 PU238 PU238 PU238	PU238			PU239 ·	PU239	PU239	PU239	PU239	PU239		PU239	PU239	PU239	PU239	PU239												
·																											

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Number of total U PU239 22

Page 4 of 7
REL BIAS	0.1400	0.2600	0.3600	0.3100	0.0900	0.0400	-0.2060	0.2500	0.5500	0.3000	-0.1060	-0.1690	-0.0240	0.1188 0.2288				Time 20			Time 2520	-0.1850	-3.0833	-1.6342	-0.3300	-0.1450	-0.0200	0.1800	0.1950	-0.3450	-0.4150	-0.2350	0.1050
	⊦ +	+	÷	+	+	+	+	· +	+	+	+	+	+	Mean Rel. Bias Mean Rel. Precision		4	-	Chem Yield 0.76 Det Eff 0.40		,	Chem Yield 1.00 Det Eff 0.39	+		Mean Rel. Bias Mean Rel. Precision	+	+	+	+	÷	+	+	+	+
UNCERT 0 5820	0.6460	0.6800	0.7790	0.6840	0.5210	0.7960	0.5170	1.2400	1.4100	1.2900	1.2400	0.7090	1.3000			0.2640				0.0018		0.0134	0.2780	1.5528	0.0412	0.0506	0.0415	0.0164	0.0361	0.0573	0.0321	0.0422	0.0248
RESULT 8.6800	11.4000	12.6000	13.6000	13.1000	10.9000	10.4000	7.9400	12.5000	15.5000	13.0000	8.9400	8.3100	9.7600	 乗₀		0.6780		0 DL		-0.0040	MDA MDA	0.0489	-0.1250	MDA	0.1340	0.1710	0.1960	0.2360	0.2390	0.1310	0.1170	0.1530	0.2210
	ц]	-1	L	Г	Г	۲,	Ц	Г	Ч				###### 2.2876		a .		0.0000			-0.0040			-0.0381 0.1230									
U U		D	D	þ	þ	n	D	D	n	D	D	D	n	esult		D		esult		D	esult	þ	n	esult	D	D	Þ	D	ŋ	Б	D	ņ	D
SPIKE UNCERT 10.0000 0.1980	10.0000 0.0668	10.0000 0.0668	10.0000 0.0668	0.0668	10.0000 0.0668	0.2190	0.2190	0.1120	0.1120	0.1120	10.0000 0.0850	10.0000 0.0850	10.0000 0.0850	Average Result St Dev		0000.0		Average Result St Dev		0.0000	Average Result St Dev	0.0007	0.0007	Average Result St Dev	0.0024	0.0024	0.0026	0.0026	0.0140	0.0140	0.0015	0.0015	0.0140
SPIKE 10.0000	10.0000	10.0000	10.0000	10.0000	10.0000	10.0000	10.0000	10.0000	10.0000	10.0000	10.0000	10.000	10.0000	10.0000 14		0.0000		0.0000		0:0000 0:0000	0.000	0.0600 0.0007	0.0600	0.0600	0.2000	0.2000	0.2000	0.2000	0.2000	0.2000	0.2000		0.2000
<u>VOL</u> 1218	1211	1183	1292	1090	1458	1087	1267	1151	1235	1426	1146	1237	1343	1-		0660	I	-		1410	1-	1193	1335	1-	1143	1124	1398	1087	1205	1323	1358		1116
REQ <u>ANAL</u> IPSA	IPSA	IPSA	IPSA	IPSA	IPSA	IPSA	IPSA	IPSA	IPSA	IPSA	SR	SR	SR	SR Count		IPS		SR 90 Count		ñ	Count	U	U	Count	D	D	U	n	U	U	U	D	D
J <u>PAYID</u> 50575	50807	99156	59621	50809	31776	65166	32533	99162	51077	80109	99214	99218	99215	ا ⊃		32476	ו ו	D		99152	Þ	80076	3G522	ן ב	80108	AU001	99157	99158	59600	19166	91386	99156	80076
TAGWORD J PAYID 06E0102 50575	06F0121	06F0295	06F0294	06F0122	06F0223	06G0241	06G0067	6010H90	06H0140	06H0115	07C0406	07C0409	07C0411	14	14	06C0067	•	-	1	06E0300	1	06D0149	06D0150	7	06F0349	06F0449	06G0234	06G0127	06H0076	06H0108	06I0171	0610265	06J0058
ANAL DATE 06/01/2006	07/23/2006	07/23/2006	07/23/2006	07/23/2006	07/26/2006	08/03/2006	08/13/2006	08/30/2006	08/30/2006	08/30/2006	03/21/2007	03/21/2007	03/21/2007		Number of total U SR	04/17/2006			Number of total U SR 90	05/19/2006		05/01/2006	05/01/2006										10/23/2006
<u>YRMO SEO</u> 0605 09	05	60	08	90	01	0	60	60	=	10	13	14	15		o nadme	60			er of to	8		07	68		10	8	6	03	05	90	90	01	6
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REL BIAS	0.1000	-0.3500	-0.0600	-0.2150	-0.3250	-0.3500	0.2198														Time 2520	-0.5686	0.0986	-0.2600	-0.2433 0.3339		-0.1233	0.0616	-0.1370	-0.0137	0.8219	-0.0274	-0.0822	-0.0548
DET	ł	+	+	+	+	+	Mean Rel. Bias Mean Rel. Precision		,	,	,	ı	ſ	1		ı	ı		ı		Chem Yield 0.71 Det Eff 0.38	ı	ı	ı	Mean Rel. Bias Mean Rel. Precision		+	+	+	+	+	+	+	+
UNCERT !	+CZU.U	0.0173	0.0179	0.0188	0.0149	0.0147			0.0046	0.0028	0.0044	0.0035	0.0045	0.0039	0.0037	0.0033	0.0051	0.0047	0.0031	0.0043	0.0056	0:0080	0.0068	0.0037	0.0261		0.0217	0.0222	0.0166	0.0201	0.0253	0.0161	0.0149	0.0170
RESULT	0.2520	0.1300	0.1880	0.1570	0.1350	0.1300	1-7-0		0.0052	-0.0050	0.0056	0.0028	-0.0017	0.0000	0.0049	0.0019	0.0017	0.000	0.000	0.0035	6 DL	0.0030	0.0077	0.0052	BBA MDA		0.1280	0.1550	0.1260	0.1440	0.2660	0.1420	0.1340	0.1380
EMR							0.1713 0.0440		D	n	D	D	n	D	D	כ	D	D	n	D	0.0016 0.0031	â	D	D	0.0053		В	п	n	D	n	D	D	D
	2	D	D	D	D	ם	Result		n	D	D	D	D	D	C	D	D	D	D	n	Result	D	D	n	Result		D	Э	D	D	Ð	D		ŋ
<u>UNCERT</u>		0.0015	0.0023	0.0016	0.0016	0.0016	Average Result St Dev		0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	Average Result St Dev	0.0000	0.0000	0.0000	Average Result St Dev		0.0010	0.0010	0.0011	0.0011	0.0010	0.0008	0.0008	0.0008
<u>SPIKE</u>	0.200	0.2000	0.2000	0.2000	0.2000	0.2000	0.2000		0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.000	0.0000	0.0000	0.0000	0.0070	0.0070	0.0070	0.0070 3		0.1460	0.1460	0.1460	0.1460	0.1460	0.1460	0.1460	0.1460
TOX		1255	1108	1146	1237	1343	•		1389	1246	1436	1306	1103	1156	1168	1298	1255	1175	1315	1266	I	1281	1388	1207	1		1281	1388	1389	1246	1436	1306	1207	1103
REQ <u>ANAL</u> 	>	n	n	n	U	U	Count		IJ	IJ	JI	IJ	5	IJ	I I	IJ	IU	5	IU	D	U 235 Count	UIAI	5	5	U 235 Count		IP IU	IJ	IJ	IJ	IU	IJ	IJ	5
<u> PAYID</u>		50784	99162	99214	99218	99215	ב		3C135	3C137	91384	99157	AU001	'n	99151	3C139	AU001	•		15166	3C139	3C135	3C137	91384	99157	AU001	AU001							
TAGWORD J PAY	C\$20100	06J0085	07B0220	07C0408	07C0410	07C0412	15	18	0610279	0610280	07A0233	07B0406	07B0606	07B0600	07B0601	07B0608	07B0607	07B0604	07B0605	07B0602	12	06H0304	06H0399	07B0603	ю	15	06H0304	06H0399	0610279	0610280	07A0233	07B0406	07B0603	07B0606
ANAL DATE	0002/22/01	10/23/2006	02/26/2007	03/21/2007	03/21/2007	03/21/2007		Number of total U	9002/10/01	10/01/2006	02/15/2007	02/22/2007	03/19/2007	03/27/2007	03/27/2007	03/27/2007	03/27/2007	03/27/2007	03/27/2007	03/27/2007		08/31/2006	00/08/2006	03/19/2007		Number of total U U 235	08/31/2006	09/08/2006	10/01/2006	10/01/2006	02/15/2007	02/22/2007	03/19/2007	03/19/2007
<u>VRMOSEO</u>	ġ :	05	12	13	14	15		lumber	8	05	03	10	8	02	05	11	10	08	67	4		08	40	90		ber of to	08	04	04	05	03	10	90	60
<u>YRM</u>	0100	0610	0702	0702	0702	0702		2	6090	6090	0701	0702	0702	0702	0702	0702	0702	0702	0702	0702		0608	0608	0702		Numl	0608	0608	0609	090	0701	0702	0702	0702
ISO CD		D	n	D	Ð	D			U 235		U 235	U 235	U 235			U 238	U 238	U 238	U 238	U 238	U 238	U 238	U 238											

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REL BIAS	-0.1370	-0.0890	-0.0753	0.0685	0.1301	0.0548	-0.0274	0.2349	
DET	+	+	+	+	+	+	+	Mean Rel. Bias Mean Rel. Precision	
UNCERT	0.0155	0.0172	0.0166	0.0183	0.0195	0.0177	0.0169		
RESULT	0.1260	0.1330	0.1350	0.1560	0.1650	0.1540	0.1420	20	
MR		D	D	D	n	D	D	0.1496 0.0343	
TYPE	D	D	n	D	D	n	D	sult	
UNCERT	0.0008	0.0008	0.0008	0.0008	0.0008	0.0008	0.0008	Average Result St Dev	
SPIKE	0.1460	0.1460	0.1460	0.1460	0.1460 (0.1460	0.1460	1.1460	
	1156			1175			1168	10	
REQ <u>ANAL</u>	5	IJ	IJ	IU	Ð	I U	UI	U 238 Count	
PAYID	AU001	AU001	AU001	AU001	AU001	AU001	100UA	ا ⊋	
ANAL DATE TAGWORD I PAYID	07B0600	07B0608	07B0607	07B0604	07B0602	07B0605	07B0601	15	15
ANAL DATE	03/27/2007	03/27/2007 07B0608	03/27/2007	03/27/2007	03/27/2007	03/27/2007	03/27/2007		Number of total U U 238 15
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<u>ISO CD</u>					U 238		U 238		

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Total Samples Total Results Page 7 of 7

APPENDIX B

GEL QUALITY CONTROL SAMPLE REPORT SUMMARY

Statistical Parameters Utilized by The GEL Group, Inc

Zone Definitions

Zone A – Area defined as being between 2 and 3 times sigma above the center line Zone B – Area defined as being between 1 and 2 times sigma above the center line Zone C – Area defined as being between the center line and 1 times sigma

Data Flag Definitions

- 1. Nine (9) points on Zone C and beyond on one side of the central line Indicates that the process average may have changed
- 2. Six (6) points in a row steadily increasing or decreasing on one side of the central line Indicates that a drift may be occurring in the process average
- 3. Fourteen (14) points in a row alternating up or down on either side of the center line If this test is positive it indicates that two systematically alternating causes may be producing different results
- 4. Two (2) out of three (3) points in a row are in Zone A or beyond Indicates an early warning of a process shift
- 5. Four (4) out of five (5) points are in Zone B or beyond If positive, this, like flag 4, indicates and early warning of a potential process shift
- 6. Fifteen (15) points are in Zone C above or below the center line Indicates a smaller variability than expected
- Eight (8) points in a row are in Zone B, A or beyond on either side of the center line with no
 points occurring in Zone C Indicates that different samples are affected by different factors
 resulting in bimodal distribution of averages

References

Statistica Software – Data Mining, Statistical Analysis and Quality Control Quality Control Charts – <u>www.statsoft.com/textbook/stquacon.html</u>

PNNL ANNUAL QC PACKAGE

2006/2007 April 1, 2006 – March 31, 2007

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Data was reviewed and found acceptable.

5-29-07 Reviewed By: Date:

Table of Contents

Section 1: Case Narrative

Section 2: Database Results

Urine Data Am-241 - Blank Activity Am-241 - LCS Bias High Am-241 - LCS Bias Low Am-243 - Blank Activity Am-243 - Tracer Yield Am-243 - LCS Bias High Am-243 - LCS Bias Low Cm-242 - Blank Activity Cm-243/244 – Blank Activity Cm-243/244 - Tracer Yield Cm-243/244 - LCS Bias High Pu-238 - Blank Activity Pu-239/240 – Blank Activity Pu-239/240 – LCS Bias High Pu-239/240 - LCS Bias Low Pu-242 – Tracer Yield Sr-90 - Blank Activity Sr-90 - Carrier Yield Sr-90 -- LCS Bias High Sr-90 - LCS Bias Low Total Sr - Blank Activity Total Sr - Carrier Yield Total Sr-LCS Bias Low Total U - Blank Activity Total U – LCS Bias High Total U - LCS Bias Low Tritium – Blank Activity Tritium - LCS Bias Low U-232 - Tracer Yield U-233/234 - Blank Activity U-235/236 - Blank Activity U-238 – Blank Activity U-238 – LCS Bias High U-238 – LCS Bias Low

Fecal Data

Am-241 – Blank Activity Am-241 – Duplicate RER Am-241 – LCS Bias High Am-243 – Tracer Yield Pu-238 – Blank Activity Pu-238 – Duplicate RER Pu-239/240 – Blank Activity Pu-239/240 – Duplicate RER Pu-239/240 – LCS Bias High Pu-242 – Tracer Yield

	Legend
#	= the N-value (number of the samples in the data set)
Samp ID	= GEL laboratory sample identification number
Inst	= the analytical instrument identification number/name
Run Date	= the sample analysis date
LCL	= Lower Control Level (minus 3 sigma)
LWL	= Lower Warning Level (minus 2 sigma)
Mean	= the average value of the data set
Numvalue	= Number Value for parameter being monitored
Exclude	= a checked box indicates the data was not used in the calculation of the
	mean and control limits
Stdev	= Standard Deviation
UWL	= Upper Warning Level (plus 2 sigma)
UCL	= Upper Contol Level (plus 3 sigma)
Dispersion	= the difference of the individual relative bias from the mean
Parent Samp	le = the sample that was duplicated
TPU	= Total Proportion Uncertainty (1 sigma combined standard uncertainty)
RER	= Relative Error Ratio (the difference of the individual duplicate pairs
	based on the combined standard uncertainties of the individual analyses)
Nominal	= the calculated concentration of the spike in the sample geometry
Result	= the actual measured analyte concentration in the sample
Bias	= the deviation of a measured value from the expected value

Duplicate Acceptance Criteria

The RER shall not exceed 3 (standard deviations) in more than one duplicate pairs in a nuclide category.

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SECTION 1

CASE NARRATIVE

Annual - QC Report - Operational Year 2006/2007

This report summarizes Quality Control Samples (QC) analyzed with bioassay samples under Contract 11530 during the Contract Year 2006/2007, beginning April 1, 2006 and ending March 31, 2007. Included in the report are listings for the blank, duplicate and spike results. A description of the attached data is provided below.

During this period, the following numbers of samples were analyzed under this contract. The QC samples include blanks, spikes, and duplicates.

		QC	Total	
Test	Matrix	Samples	Samples	% QC
Am	URINE	437	1084	40%
Am-243	URINE	122	213	57%
H-3	URINE	276	1168	24%
Ρu	URINE	1152	3604	32%
Sr-90	URINE	436	1338	33%
Total Sr	URINE	46	72	64%
Total U	URINE	339	685	49%
U	URINE	279	801	35%
Am	FECAL	133	233	57%
Pu	FECAL	138	241	57%

Blanks

The following table contains the analyses, isotope, matrix, and the calculated MDAs. The alpha spectrometry MDAs are based on the average blank counts and average tracer yields for the year. The Strontium MDAs are adjusted according to the average tracer yield for the year. The total uranium MDAs are based on the standard deviation of the 0.05 ug/L standard analyzed each day throughout the year.

		Number In Set				Sample	Avg.	Detector	Count Time
Isotope	Matrix	(N#)	MDA	Units	Lc	Volume	Yield	Efficiency	(min)
Am-241	Urine	150	0.0103	dpm/s	0.0035	1	0.725	0.385	2520
Am-243	Urine	42	0.0144	dpm/s	0.0059	1	0.885	0.385	2520
Cm-242	Urine	31	0.0095	dpm/s	0.0030	1	0.725	0.385	2520
Cm-243/244	Urine	150	0.0097	dpm/s	0.0030	1	0.725	0.385	2520
Pu-238	Urine	395	0.0083	dpm/s	0.0029	1	0.915	0.385	2520
Pu-239/240	Urine	395	0.0092	dpm/s	0.0034	1	0.915	0.385	2520
U-233/234	Urine	94	0.0216	dpm/s	0.0086	1	0.709	0.382	2520
U-235/236	Urine	94	0.0139	dpm/s	0.0051	່ 1	0.709	0.382	2520
U-238	Urine	94	0.0194	dpm/s	0.0077	1	0.709	0.382	2520
Sr-90	Urine	147	7.95	dpm/s	0.6878	1	0.760	0.396	20
Total Sr	Urine	16	7.13	dpm/s	0.3148	1	0.774	0.396	20
Tritium	Urine	138	610	dpm/L	0.3294	0.01	n/a	0.179	20
Total U	Urine	198	0.0103	ug/s	0.0059	0.05	n/a	n/a	n/a
Am-241	Fecal	41	0.2538	dpm/s	0.0519	0.333	0.744	0.385	960

Pu-238	Fecal	49	0.1220	dpm/s	0.0202	0.333	0.898	0.385	960
Pu-239/240	Fecal	49	0.1306	dpm/s	0.0219	0.333	0.898	0.385	960

The following table contains the analyses, isotope, matrix, and the calculated MDAs for Isotopic Uranium in Urine. The alpha spectrometry MDAs are based on the average blank counts and average tracer yields for the fourth quarter.

		Number In Set				Sample	Avg.	Detector	Count Time
Isotope	Matrix	(N#)	MDA	Units	Lc	Volume	Yield	Efficiency	(min)
U-233/234	Urine	30	0.0198	dpm/s	0.00823	1	0.808	0.382	2520
U-235/236	Urine	30	0.0133	dpm/s	0.00522	1	0.808	0.382	2520
U-238	Urine	30	0.0184	dpm/s	0.00783	1	0.808	0.382	2520

All analytical batches were analyzed with either a reagent blank, matrix blank or both. Blanks are in control when the calculated MDA and blank activity are both less than CRDL (contract required detection limit). In addition, the chemical tracer yields are evaluated against the yield requirements stated in the subject contract.

Overall, the blank data for each analytical process demonstrate the analyses were in control. Processing categories and samples which did not meet contractual requirements are discussed in the **Observations** section of this report.

Laboratory Control Samples (LCS)

The enclosed listing contains the analysis isotope, matrix, average relative bias and the relative precision statistic. One or more LCS sample was analyzed with each batch of samples.

Test	Matrix	Range High ^{(1) (2)}	Average Relative Bias	Relative Precision
Am-241	Urine	High ¹	-0.0108	0.1136
Am-243	Urine	High ¹	0.0263	0.0693
Cm-243/244	Urine	High ¹	0.0405	0.1006
Pu-239/240	Urine	High ¹	-0.0562	0.0487
Sr-90	Urine	High ¹	0.0828	0.1149
Total U	Urine	High ¹	-0.0119	0.0755
U-238	Urine	High ²	0.0037	0.0843
Am-241	Feces	High ¹	-0.0349	0.2686
Pu-239/240	Feces	High ¹	-0.0605	0.0729

(1) High range: nominal > 2 times the Required Detection Limit

(2) High range for U-238: nominal > 0.32 dpm/sample

		Range	Average Relative	Relative
Test	Matrix	Low	Bias	Precision
Am-241	Urine	Low	0.0869	0.3748
Am-243	Urine	Low	0.108	0.3115
Pu-239/240	Urine	Low	0.0173	0.2477
Sr-90	Urine	Low	-0.0294	0.1675
Total Sr	Urine	Low	0.052	0.141
Total U	Urine	Low	-0.0396	0.1404
Tritium	Urine	Low	-0.0506	0.0777
U-238	Urine	Low	-0.0446	0.1251

Overall, the LCS data demonstrates the analytical processes were in control. Any LCS outside the limits is discussed in the **Observations** section of this report.

Duplicate Samples (DUP)

The duplicate samples were evaluated to determine that the aliquot procedure produces results within control limits.

Refer to Narrative Attachments 1-3 for DUP tables.

One Am-241 duplicates was excluded due to not reporting the parent sample.

Sample Summary

Overall, the chemical yields for the analytical processes were greater than the minimum yields required in the SOW. Those not meeting the yield requirements are further discussed in the **Observation** section of this report.

OBSERVATIONS

Section 1:

Am Isotopic in Urine

Two Am-243 tracers (0.185%) were less than the low yield for Isotopic Am this year.

One of the two tracers was below the minimum yield requirement of 20%; however, the sample was a Laboratory Control Sample (LCS) that met the recovery requirements. No correlation between the samples was observed, so no corrective action was necessary.

Cm Isotopic in Urine

Two Cm-243/244 LCS samples were not spiked with Curium; however, they were spiked with Am-241 due to analyzing Curium and Americium together. Since Curium and Americium are from the same counting source, the Curium results were reported based on the acceptable Am-241 recoveries. NCR 346848 was created to document this anomaly.

Pu Isotopic in Urine

Fourteen tracers (0.389%) were less than the low yield for Isotopic Pu this year. No correlation between the samples was observed, so no corrective action was necessary.

Two Pu-242 tracers were observed to be below the minimum yield requirement of 25%. The failed tracers were for two LCSs with spikes within the recovery requirements, so the results were reported. NCRs 312489 and 346865 were created to document this anomaly.

Sr-90/Total Sr in Urine

Eighteen tracers (1.69%) were less than the low yield for Sr-90 this year. No correlation between the samples was observed, so no corrective action was necessary.

Th Isotopic in Urine

There was no Isotopic Th analyzed this year.

Total Uranium in Urine

There were no observations made for Total Uranium this year.

Tritium

There were no observations made for Tritium this year.

Uranium Isotopic in Urine

Seventy tracers (8.97%) were less than the low yield for Isotopic Uranium this year.

The MDA for U-233/234 is greater than CRDL (contract required detection limit) for the year due to the low tracer recoveries observed. The Standard Operating Procedure was revised November 30, 2006 as Revision 25. The revision made involved changing the type of separation chemistry from an AG1X8 anion exchange column to an Eichrom ®TRU extraction chromatography column. Since the MDA was greater than the CRDL for the year, the MDA was also calculated for the fourth quarter alone and was found to be in control. Improvements have been observed following the corrective actions in the 2006/2007 year.

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Isotopic Am in Feces

Fifteen tracers (7.43%) were less than the low yield for Isotopic Am this year.

Due to the low tracers throughout the year, corrective action was necessary and has been taken. The procedure was investigated and revised December 11, 2006 as Revision 10. Improvements have been observed since the end of third quarter.

Isotopic Pu in Feces

Four tracers (1.69%) were less than the low yield for Isotopic Pu this year. No correlation between the samples was observed, so no corrective action was necessary.

Incident Reports

The following incident reports were documented this year:

Incident Report for Tagwords 06E0067, 06E0184, 06E0233, 06E0319, 06E0325, 06E0327, and 06E0338. GEL was unable to report the volume received for the above referenced Tagwords due to the analyst recording the measurements on a loose sheet of paper. The paper was inadvertently discarded prior to the entry of the data into LIMS. A bound logbook, RC-O-024, was created to permanently record the volumes received. The logbook can then be referenced to enter this data into LIMS at a more appropriate time. The incident was closed May 24, 2006.

Incident Report for the Curium spiking error associated with work orders 167241 and 167731. The group leader met with both technicians responsible in the matter and their team leader. The four of them discussed the error and were reminded of the information provided in the que and pull sheets. In addition, the team leader will use additional identifying methods for infrequent analyte requests. The incident was closed August 31,2006.

Corrective Actions

The corrective action regarding low tracer recoveries for Americium in Fecal analyses was closed January 25, 2007 following an investigation and revision of the Standard Operating Procedure. As for the corrective action regarding Uranium in Urine analyses, improvements have been observed following the revision of the procedure and will close during the first quarter of the 2007/2008 year.

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NARRATIVE ATTACHMENT 1 Americium-241 Sample Type: RER Attribute Tracked: RER (dec)

Values less than and greater than

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	0.00477	0.00469	0.00249	0.0186	0.0848	10.209	0.0168	0.0107	0.00755	0.00318	0.0112	0.00562	0.00266	0.00475	0.00292	0.367	0.0037	0.00937	0.043	0.044	0.0508	0.0103	0.00414	0.012	0.138	0.00534	0.0111	0.0095	0.0134	0.00667	0.00566	0.00328	0.0113	0.0134	100
Udi	0.00412 and 0.0047	0.0148 and 0.00469	0.00421 and 0.00249	0.00912 and 0.0186	0.0639 and 0.0848	0.0629 and 0.209	0.00666 and 0.0168	0.0136 and 0.0107	0.0049 and 0.00755	0.0202 and 0.00318	0.0127 and 0.0112	0.0125 and 0.00562	0.0321 and 0.00266	0.00489 and 0.00475	0.0166 and 0.00292	0.338 and 0.367	0.00397 and 0.003	0.0362 and 0.0093	0.0233 and 0.043	0.0134 and 0.044	0.0262 and 0.0508	0.0151 and 0.0103	0.0183 and 0.00414	0.0158 and 0.012	0.169 and 0.138	0:00567 and 0.0053	0.0194 and 0.0111	0.00676 and 0.0095	0.00595 and 0.0134	0.00347 and 0.0066	0.00269 and 0.00566	0.00597 and 0.00328	0.00381 and 0.0113	0.0136 and 0.0134	0 147 and 0 109
Result	0069 and 0.0057	0.025 and 0.007	0117 and0099	0046 and 0.0244	0176 and0176	0137 and0712	0166 and 0.0283	0208 and 0.0151	0019 and0179	0206 and0053	0176 and 0.0167	031 and0018	0207 and0032	0045 and 0.0036	0145 and0009	1.6 and 1.8	0.0048 and 0.0044	0187 and0063	0.033 and 0.061	0.01 and021	0.068 and 0.168	0.021 and01	0.031 and 0.004	0099 and 0.0129	0.8 and 0.6	-0076 and0099	038 and0101	0.0003 and 0.0135	0.0059 and 0.019	0.0015 and0099	- 0011 and 0.008	0099 and 0.0008	0042 and 0.0189	0.024 and 0.023	0.6 and 0.6
Parent Sample	06C0641	0620330	06D0340	06E0340	06E0534	06E0532	06F0561	06G0265	06G0Z76	06G0431	06G0440	06G0438	06140358	06H0383	0610319	0610535	06K0326	06K0330	06K0409	06K0411	06L0442	06L0468			06L0433	07A0462		07A0697	07B0404		0780563		Ì	07B0619	OTENETO 1
UCL	3.14	3.14	3.14	3,14		3.14	3.14	3.14	3.14	3.14	3.14	3.14	3.14	3.14	3.14	3.14	3.14	3.14	3.14	3.14	3.14	3.14	3.14	3.14	3.14	3.14	3.14	3.14	3.14	3.14	3.14	3.14	3.14	3.14	11
UML	2.47	2.47	2.47	2.47		2.47	2.47	2.47	2.47	2.47	2.47	2.47	2.47	2.47	2.47	2.47	2.473	2.47	2.47	2.47	2.47	2.47	2.47	2.47	2.47	2.47	2.47	2.47	2.47	2.47	2.47	2.47	2.47	2.473	2 47
Stdev	1.3	0	-1.2	4		-1,3	2	1.4	-	9	1.3	1.5	6	١.	5	-1.2	-1.6	-12	6	~	o,	σ	'n	•	<u>،</u>	-1.3	Ņ	0	- 4	9	υ	7	+	-1.6	с -
Exclude																																		ĺ	
TPU E	0.0048	-1 -1 -1	0025 7	<u>19</u>	085	21	-017	1 <u>5</u>	0076	0032	011 L	L 9920	0027	0048 1	C 6200	۲ کو	0037	2009 L	۲ 1	튑	۲ کا	티	뒿	<u>.012</u>	L 7	L 8900	티	L 88	013	L 2900	L 2990	L 880	۲ 1	013	L F
Numvatue		1.17	.37	1.4	0.00E+00	- 26	2.48	2.07	1.78	.75		_		1.19	8.		_		.57	-92	1.73	┥		1.15		-	-	-	6		-	-	-	_	9
Mean	1.14	1,14	1.14	1.14		1.14	1.14	1.14	1,14	1.14	1.14	. 4	1.14	1.14	1.14	1.14	1.14	7	1	4	÷	7	¥ 	1.14	7	4	÷.		1.14	1.14	-		-	1,14	1.14
LWL	0.19	<u>.</u>	-19	-19		-19	-19	-19	-19	-19	-19	6	.19	8 <u>-</u>	- 19		<u>8</u>	<u>8</u>		<u>6</u>	6		<u>6</u>	<u>6</u>	-19	<u>6</u>	<u>6</u>	<u>9</u>	- <u>1</u> 9	9	2	8 -	6	8	61-
ಕ್ಷ	-0.86	ŝ	ŝ	8		. 98.	- 66	-86	-86	-86	8 [.]	8	8	8	8	8	ŝ	8	8. 9	8. 8.	8	8	8	8	8	ş	8; 8;	8	8	ģ	ŝ	8	8	ŝ	- 98
Run Dete	11-Apr-06	12-APR-06	24-APR-06	16-MAY-06	1635 09-JUN-06	1649 09-JUN-06	26-JUN-06	17-JUL-06	19-JUL-06	1711 01-AUG-06	1636 07-AUG-06	1714 08-AUG-06	1682 16-AUG-06	1690 24-AUG-06	21-SEP-06	12-OCT-08	14-NOV-06	16-NOV-06	12-DEC-06	13-UEC-06	8-DEC-08	SP-DEC-06	61-DEC-06	1700 08-JAN-07	16-JAN-07	10-NN-02 / 201	07-FEB-07	09-FEB-07	15-FEB-07	27-FEB-07	01-MAH-07	16-MAR-07	6-MAR-07	1715 27-MAR-07	18-MAR-07
	1694	1659	1710	1637			1635	1636		_	_			8	<u>[</u>	1701	1691	1635	8		989	60/1		802				202	1710	1715 2	1703	1689 1	1693 1	1715 2	169712
Samp ID	1201062888	1201065026	1201072748	1201067780	1201104166	1201105097	1201120001	1201128589	1201131872	1201143163	1201146263	1201145207	1201154413	1201159512	1201180963 1687 21-SEP-06	1201198588 1701 12-0CT-06	1201222272 1691 14-NOV-06	1201224284 1635 16-NOV-06	1201236923 1660 12-DEC-06	1201239188 17/14 13-DEC-06	120124/783 1696 26-DEC-06	1201248840 1709 29-DEC-06	1201201/61 1640 31-DEC-06	12/1243298	1201203630 1064		12012/0008	12012/2002	12012/4062	1201280511		1201291669		1201296206	1201298274 1697 28-MAR-07
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, NARRATIVE ATTACHMENT 2 Plutonium-238

Sample Type: RER Attribute Tracked: RER (dec)

Values less than and greater than

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Udit	0.00431 and 0.0141	0.00564 and 0.0149	0.00508 and 0.00507	0.00514 and 0.00445	0.023 and 0.0253	0.00796 and 0.0267	0.00527 and 0.012	0.0101 and 0.0117	0.00578 and 0.0148	0.00755 and 0.00496	0.00584 and 0.0129	0.00623 and 0.0137	0.0133 and 0.0112	0.00665 and 0.0122	0.00468 and 0.0048	0.0207 and 0.00677	0.0231 and 0.0122	0.0133 and 0.0157	0.00684 and 0.0365	0.00493 and 0.00498	0.0156 and 0.0154	0.0161 and 0.0139	0.0103 and 0.0139	0.0186 and 0.0206	0.0052 and 0.0101	0.00479 and 0.0137	0.00473 and 0.00546	0.0105 and 0.00627	0.00512 and 0.00932	0.038 and 0.00811	0.00611 and 0.0161	0.0455 and 0.0441	0.00473 and 0.00639	0.00615 and 0.0127	0.0124 and 0.0219	0.00682 and 0.0043	0.0249 and 0.022
Recut	0.0035 and0056	0003 and0061	0 800 0	0024 and0045	0092 and0099	0138 and0097	0.0043 and 0.0176	0021 and 0.0172	0.0064 and0046	0.0063 and001	0.0048 and0035	0.0051 and0038	0037 and 0.0138	0.0054 and003	0 and 0	10-000 800-	0.036 and 0.009	-,0054 and -,0059	0037 and0142	0035 and0035	0.035 and 0.035	0.033 and 0.023	0.015 and 0.025	0.042 and 0.048	0026 and0039	0015 and0048	0.0039 and0015		0.0017 and 0.0073		0.005 and005	0.2 And 0.2	0012 and 0.0071	0.0066 and0043	0046 and0076	0.0075 and0013	0.07 and 0.055
Perent Semple	0600641	060030	06D0346	06D0340	06E0340	000034	00E0532	06F0560	06G0265	06:00276	06G0431	06G0438	06G0440	06H0358	06H0383	0610319	06/0535	06K0326	06K0409	06K0411	06L0442	06L0465	06L0468	06L0637	06L0412	06L0433	07A0462	07A0697	0780404	07A0702	0780520	07B0518	07B0563	0700425	07C0448	07C0442	0780519
ncr	1.69	1.69	1.6905	1.69	1.69	1.69	1.69	1.69	1.69	1.69	1.69	1.69	1.69	1.69	1.6905	1.69	1.69	1.69	1.69	1.6905	1.69	1.69	1.69	1.69	1.69	1.69	1.69	1.69	1.69	1.69	1.69	1.69	1.69	1.69	1.69	1.69	1.69
UWL	1,29	1.29	1.2911	1.29	1.291	1.29	1.29	1.29	1.29	1.29	1.29	1.29	1.29	1.29	1.2911	1.29	1.29	1.291	1.29	1.2911	1.291	1.29	1.28	1.29	1.29	1.29	1.29	1.29	1.29	1.291	1.29	1.29	1.29	1.29	1.29	1.29	1.29
Stdev -	0.3	6.Q	-1.2	÷.5	42	6'-	1.3	2.8	.5	1.3	Ņ	.3	1.3	Ð,	-1.2	8-	1.3	-1.2	9	-1.2	-1.2	0	ł	- 7	6	-7-	8.	1	•	-1.1	ы	1	1.4	.7	6	1.5	1-
Exclude																																					
TPU E	0.014	0.015	.0051	-0045	.025 T	L 720.	.012	-012	.015 7	- 900.	-013	-014	-011	.012	0048	L 8900	.012	-018 L	-030 -	۲. 80.	.015	L 14	-014 -	18 18	٦. ۲	L 14	0056 7	D063 T	L 6800	0081	018	044 L	D084 T	013	۲ 8	C 100	۲ 8
Numvakue	0.62	0.37	0.00E+00	e,	.02	.15	1.01	1.6	69	1.03	83.	-59	1.01	9	0.00E+00	-19	1.01	.028	27	0.00E+00	.018	.47	.54	.21	-11	23	.74	.87		033	58.	33	1.04	.79		_	.45
Mean	0.49	0.49	4822	.49	.492	.49	.49	.48	.49	.49	49	.49	.49	49	4022	.49	49	492	49	.4822	.492	49	.49	.49	.49	.49	-49	48	49	492	.49	.49	-49	.49	49	49	.49
LWL	-0.31	-0.31	-3066	31	-307	31	31	31	31	31	-31	31	-31	31	3066	-31	31	-307	- 31	- 3066	307	-31	-31	 31	31	-31	-31	-31	.31	307	- 31	31	-31	-31	.31	5.3	-31
LCL	-0.71	-0.71	7061	7	- 706	71	17	-11	17-	71	-1	71	F,	Ч	- 7061	- 71	-71	-708	-,71	- 7061	-,706	-71	71	2	17	-71	Ę	-71	.71	706	71	•.71	17.	71	F.	F.	- 71
Run Date	11-Apr-06	12-Apr-06	13-APR-06	24-APR-06	1201067787 1643 18-MAY-06	00-VIU-00	08-JUN-06	23-JUN-06	17-JUL-06	19-JUL-06	01-AUG-06	04-AUG-06	07-AUG-06	1715 15-AUG-08	24-AUG-06	21-SEP-06	11-OCT-06	13-NOV-06			22-DEC-08	27-DEC-08	29-DEC-06	31-DEC-06	05-JAN-07	11-JAN-07	23-JAN-07	00-FEB-07	15-FEB-07	22-FEB-07	26-FEB-07	26-FEB-07	01-MAR-07	14-MAR-07	15-MAR-07	15-MAR-07	28-MAH-07
Ind	1696	1663	1662	1706	5 <u>5</u>	1640	1645	1836	1648	1702	1715	1715	1637	1715	1687	1662	1663	1683	1670	1708	1703	1692	1669	1636	1675	168	1709	1698	1695				1699	1691	167	1697	1703
Samp ID	1201062891 1696	1201065034 1663	1201065862 1662	1201072761	1201087787	1201104169 1640	1201105100 1645	1201117818 1636	1201128582 1648 17-JUL-06	1201131875 1702 19-JUL-06	1201143166 1715 01-AUG-06	1201145210 1715 04-AUG-06	1201146266 1637	1201154416	1201150519 1087	1201180968 1862	1201196691 1653 11-OCT-06	1201222275 1683 13-NOV-06	1201236936 1670	1201230191] 1708] 13-DEC-08	1201247785 1703 22-DEC-08	1201248653 1692	1201249947 1669 29-DEC-06	1201251768 1638 31-DEC-06	1201243301 1675	1201253833 1696 11-JAN-07	1201261256 1709	1201272605 1698 08-FEB-07	1201274085 1695 15-FEB-07	1201276649 1695	1201280514 1665	1201280274 1678	1201282353	1201290610 1691	1201282406 1671 15-MAR-07	1201291676 1697 15-MAR-07	1201298281 1703 28-MAR-07
•	-	2	0	4	ß	ø	~	æ	ð	ç	÷	12	t 3	4	15	16	17	18	19	ଷ	21	ង	ន	24	8	8	27	8	8	8	31	32	8	8	S	8	37

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NARRATIVE ATTACHMENT 3 Pkutonlum-238/240 Sample Type: RER Attribute Tracked: RER (dec)

Values less than and greater than

or annal to

	TPU	0.024 and 0.0267	0.0126 and 0.0149	0.00359 and 0.014	0.0135 and 0.00807	0.00474 and 0.00523	0.00407 and 0.00765	0.00549 and 0.0149	0.0143 and 0.0139	0.0128 and 0.00565	0.00279 and 0.00351	0.00333 and 0.00963	0.00367 and 0.0026	0.00631 and 0.0128	0.00263 and 0.00232	0.00331 and 0.0133	0.00765 and 0.0064	0.0346 and 0.0335	0.0128 and 0.00875	0.0128 and 0.00563	0.00861 and 0.00498	0.0344 and 0.039	0.113 and 0.114	0.0103 and 0.00549	0.0134 and 0.0143	0.0052 and 0.0133	0.12 and 0.103	0.00474 and 0.0151	0.026 and 0.0147	0.00344 and 0.00807	0.0074 and 0.00615	0.00306 and 0.0161	0.149 and 0.141	0.00255 and 0.013	0.00957 and 0.0109	0.00334 and 0.00627	0.0138 and 0.0119	0.0862 and 0.0768
	Result	0.06 and 0.083	0.022 and009	0007 and0096	006 and 0.0099	0026 and0026	0.0013 and016	-,0024 and -,0082	0105 and 0.0255	0.029 and006	0.0008 and - 0031	0025 and 0.017	0006 and 0.0025	0.0093 and 0.021	0.0008 and - 0002	0041 and0124	0.012 and 0.009	0.08 and 0.093	0.024 and 0.013	0.02 and003	0.015 and011	0.1 and 0.2	1 and 1	0.014 and008	0.022 and011	0074 and 0.0218	1.1 and 0.9	0.0039 and0051	0069 and0053	0.0023 and 0.0081	0.006 and 0	0.0034 and0103	1.6 and 1.4	0.0028 and0086	0.013 and 0.019	0.0009 and0149	-,0092 and -,0087	0.6 and 0.6
	Parent Sample	06C0641	06D0330	06D0346	06D0340	06E0340	06E0534	06E0532	06F0560	06G0265	06G0276	06G0431	0600438	0660440	06H0358	06H0383	0610319	0610635	06K0326	06K0409	06K0411	06L0442	06L0465	06L0468	06L0637	06L0412	06L0433	07A0462	07A0697	0780404	07A0702	0780520	0780518	0780563	07C0425	07C0448	07C0442	0780519
	л С	3.16	3.16	3.16	3.16	3.1613	3.16	3.16	3.16	3.16	3.16	3.16	3.16	3.16	3.16	3.16	3.16	3.16	3.16	3.16	3.16	3.16	3.1613	3.16	3.16	3.16	3.16	3.16	3.16	3.16	3.16	3.16	3.16	3.16	3.16	3.16	3.161	3.16
	MS	2.44	2.44	2.44	2.44	2.4365	2.44	2.44	2.44	24	2.44	2.44	2.44	2.44	2.44	2.44	2.44	2.44	2.44	2.44	2.44	2.44	2.4365	2.44	2.44	2.44	2.44	2.44	2.44	2.44	2.44	2.44	2.44	2.44	2.44	2.44	2,436	2.44
	Stdev	9 9	0.8	-0.5	0	-1.4	1.4	6'-	£,	~	-7	1.3	4	¢,		ς; •	÷	•	- A	6.	2.2	6	-1.4	1.2	¢.	1.5	۲.	9.	-1,2	ŝ	5	-2	6	-2	8.	1.7	е Г	ę
or a qual lo	Exclude					: ,							,				 •				 1		 ,		1									 ,				
5		יייי ר	0.015	0.014	0081	.0052	1400	015	014	.0057	.0035	9600.	.0026	.013	0023	013	.0064	.034	0088	.0056 Г	.005	<u>, 039</u>	111	.0055 F	.014 L	.013 r	-	.015 Г	.015 Г	.0081	.0062 Г	.016	14	-013	-011	.0063	.012	770.
	Numvalue	0.63	1.6	0.61	1.02	0.00E+00	2	8	1.5	2.46	- 18.	1.91	8 9.	2 <u>3</u>	8	19 [.]	-25	-27	89	1.62	2.6	.79	0.00E+00	1.85	1.66	2.04	1.49	- 22	.12	99	.63	63.	82.	98.	43	2.23	.026	74
	Mean	96.0	66.0	0.99	66	9968	8	6 6'	8 i	8	8 ;	66 .	8	8 6.	6.	8	6 6.	66`	8	8	66.	66.	9996	66'	66'	66	66	66	66	66	6 6	8	66'	6 ,	66	- 68	.967	66.
	ž	-0.46	0.46	-0.46	- 46	- 4628	-46	-46	-46	46	46	46	-46	46	-46	46	- 4 6	91	-46	-46	-46	-46	-4628	46	- 46	46	-46	46	-,46	- 46	- 46	-46	- 46	-46	46	46	- 463	46
	Ę	-1.19	-1.19	-1.19	-1.19	1.1676	-1.19	-1.19	-1.19	-1.19	-1,19	-1.19	-1.19	1.19	-1.19	-1.19	-1.19	-1.19	-1.19	-1.19	-1.19	-1.19	-1.1876	-1.18	-1.19	-1.19	-1.19	-1.19	-1.19	-1.19	-1.19	-1.19	-1.19	-1.19	-1.19	-1.19	-1.168	-1.19
	Run Oate	11-Apr-06		13-Apr-06			90-NUL-20	90-NUL-60	1636 23-JUN-06	17-701-06	19-JUL-06	1715 01-AUG-06	1715 04-AUG-06	1637 07-AUG-06	1715 15-AUG-06	1687 24-AUG-06	21-SEP-06	1653 11-OCT-06	13-NOV-06	1670 12-DEC-06	1708 13-0EC-06	1703 22-DEC-06	27-DEC-06	29-DEC-06	1636 31-DEC-06	1675 05-JAN-07	11-JAN-07	1709 23-JAN-07	1696 08-FEB-07	15-FEB-07 1	22.FEB-07	26-FEB-07	26-FEB-07	1699 01-MAR-07	1691 14-MAR-07	1671 15-MAR-07	1697 15-MAR-07	1703 28-MAR-07
	Inst	+	1983	1662	1706	1643	9 9	÷		1648	1702		_	_	1715	1687	_	_	1683	_			1692	1669	_	1675	1696			1695	1695	1665	1678	_				_
1	l Samo ID	1201062891	1201065034	1201065862	1201072751	1201087787	1201104169	1201105100	1201117818	1201128582 1648	1201131875	1201143166	1201145210	1201146266	1201154416	1201159519	1201180968	1201198591	1201222275	1201238938	1201239191	1201247785	1201248863	1201249947	1201251768	1201243301	1201253633	1201261256	1201272605	1201274085	1201278649	1201280614	1201280274	1201282353	1201290610	1201292406	1201291676	1201298281
	i,	-	5	6	4	s	8	7	80	6	₽	11	12	13	14	15	16	17	18	19	8	21	ន	ន	24	*8	8	27	83	8	8	31	8	ß	æ	35	36	37

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	COMPANY - WIDE NO	NCONFORMANCE RE	PORT
Mo.Day Yr. 01-MAY-06	Division: Bioassay	Quality Criteria: Client Contract	Type: Material
instrument Type: BIOASSAY ALPHA	Test / Method: GL-RAD-B-001	Matrix Type: Liquid	Client Code: PNNL/160202
Batch ID: 515206	Sample Numbers: 1201058578		
Potentially affected work order(s)(SDG): 159049,160202,160212		
Application issues:			
Failed Recovery for Surrogate or Tra			
Specification and Requirements Nonconformance Description:		NRG Disposition:	
1. 1201058578 LCS failed tracer yi	ełd.	1. Because the LCS re	ecovery is acceptable, the results are reported.
			1
Originator's Name;		Data Validator/Group	I saviar
Anson Walsh 01-MAY-06		Robert Timm	05-MAY-06
Quality Review: Lonnie Morris 03-JUL-06			
Director:			
Robert Timm 10-NOV-06			

I.

Revision No.: 4

IDASSAY ALPHA IDASSAY ALPHA IDASSAY ALPHA IDASSAY ALPHA ICL ID: IDASSAY ALPHA ICL ID: IDE ID: IDE	Matrix Type: Liquid	Cilent Code: PNNL
49167 1201137896, 1201137897 otentially affected work order(s)(SDG): 157241,167731 pplication issues: ailed RPD for LCS/LCSD pecification and Requirements		I
pplication issues: ailed RPD for LCS/LCSD pecification and Requirements		······································
ailed RPD for LCS/LCSD		
pecification and Requirements		
pecification and Requirements		
	NRG Disposition:	
 QCs 1201137896 (LCS) and 1201137897 (LCS) did not meet the splke recovery requirements for Curlum due to missing the Cm-244 splke. 	1. The PM was informed, a excepted and is reported of analyses.	and the client was contacted. The data was us to no positive results within the Curium
Iginator's Name:	Data Validator/Group Lee	der:
aon Walah 17-AUG-06	-	3-AUG-08
uality Review;		

15

;

GEL Laboratories LLC Form GEL-NCR Rev. 06/05

NCR Report No.: 346865 Revision No.: 2

	COMPANY - WID	E NONCONFORMANCE REPOR	r
Mo.Day Yr. 17-AUG-06	Division: Bioassay	Quality Criteria: Client Contract	Type: Process
Instrument Type: BIOASSAY ALPHA	Test / Method: GL-RAD-B-001	Matrix Type: Liquid	Client Code: PNNL
Batch ID: 551506	Sample Numbers: 1201143409		
Application issues: Failed Recovery for Su	-	0	
Specification and Re Nonconformance De	quirements scription:	NRQ Disposition:	
1. QC 1201143409 (LCS)'s tracer yield was less than 25%.	1. The LCS was recounted. The results were reported.	The recount confirmed the low tracer yield.
Originator's Name: Anson Walsh Quality Review:	17-AUG-08	Data Validator/Group Leade Robert Timm 28-/	vr: NUG-06

Director:

Page 1

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I.

SECTION 2

DATABASE RESULTS

(Maintained electronically in Hanford Internal Dosimetry Program records)

APPENDIX C

GEL Duplicates



GEL Fecal Duplicates Apr. 2006 - Mar. 2007

GEL Fecal Duplicates, April 2006 - March 2007



Second Analysis, dpm

APPENDIX D

QUALITY CONTROL INTERCOMPARISON PARTICIPATION RESULTS



U.S. DEPARTMENT OF COMMERCE

National Institute of Standards and Technology Gaithersburg, MD

REPORT OF TRACEABILITY

General Engineering Laboratories, LLC Charleston, South Caroline

Test Identification: Test Radionuclides: Matrix Description: Test Activity Range: Reference Time: NRIP-06-SF ²⁴¹Am, ²³⁸Pu, ²³⁰Th, ²³⁸U, ²³⁵U, ²³⁴U, ⁹⁰Sr, ⁶⁰Co, ⁵⁷Co, ¹³³Ba, ¹³⁷Cs, ¹³⁴Cs, ⁵⁴Mn Synthetic Feces¹ 30mBq•sample⁻¹ to 300mBq•sample⁻¹ 12:00 EST, April 1, 2006

Measurement Results

Nuclide	NIS	T Value ^{2,3}	Repo	rted Value ⁴	Difference ⁵
	Massic Activity	Relative Expanded	Massic Activity	Relative Expanded	
	Bq∙g ⁻¹	Uncertainty (%, k=2)	Bq•g ⁻¹	Uncertainty (%, k=2)	(±% Bias)
²⁴¹ Am	1.081	0.95	0.945	14.10	-12.6
²³⁸ Pu	0.938	0.68	0.877	14.72	-6.5
230Th	1.630	0.58	1.371	43.91	-15.9
²³⁸ U	4.255	0.62	4.258	10.88	0.1
²³⁴ Ŭ	4.098	0.98	4.024	11.04	-1.8
²³⁵ U	0.196	0.64	0.212	31.99	8.1
⁹⁰ Sr	3.920	0.74	2.520	54.48	-35.7
NR= Not Rep	orted			NA= Not	Applicable
		Method	\$		•
		NIST	· ·	Reporting Labor	atory ⁷
Activity Measurements		Alpha- and Beta-S Mass Spectro		Aipha, Beta, and Gamma	Spectrometry

Evaluation (per ANSI N42.22 and N13.30)

Nuclide	N4	2.22*	NI	3.309
	ANSI N42.22 Traceable	Traceability Limit	-	e per N13.30 Criteria is/Fail)
		(±Percent)	Bias	Precision
²⁴¹ Am	Yes	19	Pass	Pass
²³⁸ Pu	Yes	21	Pass	Pass
²³⁰ Th	Yes	55	Pass	Pass
238U	Yes	16	Pass	Pass
²³⁴ U	Yes	16	Pass	Pass
²³⁵ U	Yes	52	Pass	Pass
⁹⁰ Sr	Yes	53	Fail	Pass

Samples Distributed O Reporting Data Received D

October 21, 2006 December 21, 2006

For the Director

Mehael

Michael Unterweger, Acting Leader Radioactivity Group Physics Laboratory (Continued)



U.S. DEPARTMENT OF COMMERCE

National Institute of Standards and Technology Gaithersburg, MD

REPORT OF TRACEABILITY

General Engineering Laboratories, LLC Charleston, South Caroline

Test Identification: Test Radionuclides: Matrix Description: Test Activity Range: Reference Time: NRIP-06-SF ²⁴¹Am, ²³⁸Pu, ²³⁰Th, ²³⁸U, ²³⁵U, ²³⁴U, ⁹⁰Sr, ⁶⁰Co, ⁵⁷Co, ¹³³Ba, ¹³⁷Cs, ¹³⁴Cs, ⁵⁴Mn Synthetic Feces¹ 30mBq•sample⁻¹ to 300mBq•sample⁻¹ 12:00 EST, April 1, 2006

Measurement Results

Nuclide	NIS	ſ Value ^{2,3}	Repo	rted Value ⁴	Difference
	Massic Activity	Relative Expanded	Massic Activity	Relative Expanded	1
	Bq•g ⁻¹	Uncertainty (%, k=2)	Bq•g ⁻¹	Uncertainty (%, k=2)	(±% Bias)
²⁴¹ Am	1.081	0.95	0.987	14,50	-8.7
²³⁸ Pu	0.938	0.68	0.852	13.48	-9.2
²³⁰ Th	1.630	0.58	1.410	11.80	-13.5
²³⁸ U	4.255	0.62	4.186	10.56	-1.6
²³⁴ U	4.098	0.98	3.996	10.60	-2.5
²³⁵ U	0.196	0.64	0.224	23.16	14.6
⁹⁰ Sr	3.920	0.74	3.376	20.62	-13.9
NR= Not Rep	orted		•	NA= Not	Applicable
		Method	5	· · · · · · · · · · · · · · · · · · ·	· · · · · · · · · · · · · · · · · · ·
		NIST ⁶		Reporting Labor	atory ⁷
Activity Me	asurements	Alpha- and Beta-S Mass Spectro		Alpha, Beta, and Gamma	Spectrometry

Evaluation (per ANSI N42.22 and N13.30)

Nuclide	N4	2.22*	N1	3.309					
	ANSI N42.22 Traceable	Traceability Limit	Results Acceptable per N13.30 Criteri (Pass/Fail)						
		(±Percent)	Bias	Precision					
²⁴¹ Am	Yes	20	Pass	Pass					
²³⁸ Pu	Yes	18	Pass	Pass					
²³⁰ Th	Yes	15	Pass	Pass					
²³⁸ U	Yes	16	Pass	Pass					
234U	Yes	16	Pass	Pass					
²³⁵ U	Yes	40	Pass	Pass					
⁹⁰ Sr	Yes	27	Pass	Pass					

Samples Distributed Octobe Reporting Data Received Decemi

October 21, 2006 December 21, 2006 For the Director

Michael Unterway

Michael Unterweger, Acting Leader Radioactivity Group Physics Laboratory (Continued)

As guidance for the proper use of this Report, it should be emphasized that the National Institute of Standards and Technology is concerned only with fostering good measurement capability and consistency with the national measurements system. The assurance of the proper application of that capability to the ultimate consumer products is the responsibility of each manufacturer and of the Federal regulatory agencies.

A continuing traceability program in radioactivity demonstrates, to the degree established by the periodic assays of calibrated radioactivity samples, a continuing competence to maintain the instrument systems and standards necessary for accurate measurement. Such a program cannot, however, endorse each and every measurement nor the final product, any more than a spot check can vouch for every unchecked item. Care should be taken, therefore, not to imply such endorsement. The proper use of this Report is governed by section 200.114 of Title 15 of the Code of Federal Regulations. These regulations may be met if Reports are quoted only in their entirety. Excerpts out of context may be misleading.

Notes

(1a) Five test-samples and three (identical matrix) blanks were provided for this test. Each sample consisted of approximately 100 grams of synthetic fecal material contained in a plastic zip-lock bag that was packed in a plastic container.

Composition of the Synthetic Feces							
Reagent	g/sample						
Calcium Nitrate	0.97						
Ferric Ammonium Sulfate	0.04						
Magnesium Carbonate	0.61						
Potassium Carbonate	0.83						
Ammonium Dihydrogen Phosphate	2.1						
Sodium Sulfate	0.37						
Ammonium Chloride	0.04						
Zinc Sulfide	0.01						
Stannous Chloride	0.03						
Leucine	7.1						
Lysine	5,1						
Methionine	0.8						
Threonine	2						
Palmitic Acid	3						
Stearic Acid	2						
Cellulose	4						
Gelatin	5						
Oleic Acid (Liquid)	1						
Peanut Oil	1.5						
Water (distilled)	65						

Composition of the Synthetic Feces

- (1b) The test samples were prepared by depositing a known amount of a NIST calibrated "spike" solution (aqueous solution containing known quantities of ²⁴¹Am, ²³⁸Pu, ²¹⁰Th, ²³⁸U, ²³⁵U, ²³⁴U, ⁹⁰Sr, ⁵⁷Co, ⁶⁰Co, ⁵⁴Mn, ¹³⁷Cs, ¹³⁴Cs, and ¹³³Ba to the center of individual ashless paper filters (37 mm diameter). After deposition of this solution, filters were dried overnight. Once dry, each filter was sandwiched between two unspiked filters. Each sandwich was then slipped into a low-density polyethylene sleeve (wall density ~0.1 mm) and sealed for counting. After confirmation measurement, each spiked filter pack was placed inside of the matrix contained plastic zip-lock bag (1a) for packaging and shipment.
- (2a) Solutions of tracers were prepared by gravimetric dilutions of NIST Standard Reference Material SRM's or NIST calibrated solutions. The dilution factors at each step were confirmed by radioactivity measurements.

The analysis methodology and nomenclature used for the NIST uncertainties are based on uniform guidelines [cf., B.N. Taylor and C. E. Kuyatt, NIST Technical Note 1297 (1994)] and are compatible with those adopted by the principal international metrology standardization bodies. Individual uncertainties have the significance of one standard deviation of the mean, or an approximation thereof. The relative combined uncertainty is the quadratic combination of the standard deviation (or standard deviation of the mean where appropriate), or approximation thereof, for the following component uncertainties:

	Nuclide (SRM Identification)	Uncertainty (%, 1s)
a)	²⁴¹ Am (4322B)	0.48
b)	²³⁸ Pu (4323A)	0.34
c)	²³⁰ Th (4342)	0.29
d)	²³⁸ U (4321C)	0.31
e)	²³⁵ U (4321C)	0.32
ſ)	²³⁴ U (4321C)	0.49
g)	⁹⁰ St (4919H)	0.37
ĥ)	⁶⁰ Co (4915E)	0.27
i)	⁵⁷ Co (Analytics, NIST calibration)	0.35
Ď	¹³³ Ba (4251C)	0.26
k)	¹³⁷ Cs (4233D)	0.34
l)	¹³⁴ Cs (4370C)	0.37
m)	⁵⁴ Mn (Analytics, NIST calibration)	0.34
	Other Sources	
5		

i) Gravimetrics (dilutions) 0.05

The individual certified uncertainties of standard reference materials are based on the quadratic combination of all sources of uncertainty manifested in the preparation of the material. These uncertainties may result from uncertainties from any or all of the following: alpha-decay emission rate, background, balance calibration, decay corrections, decay-scheme data, extrapolation of alpha-particle-count-rate-versus-energy to zero energy, live time, alpha-particle detection efficiency, alpha-emitting impurities, gamma-emitting impurities.

The Relative Expanded Uncertainty is obtained by multiplying the standard uncertainty by a coverage factor of k-2 and is assumed to provide an uncertainty interval of approximately 95 percent confidence.

(3)Half-lives used

	Nuclide	<u>Half-life</u>
a)	²⁴¹ Am	432.2±0.5 y
b)	²³⁸ Pu	87.74±0.04 y
c)	²³⁰ Th	75380±30 y
d)	²³⁸ U	(4.468±0.003) x 10 ⁹ y
e)	²³⁵ U	$(7.038\pm0.005) \times 10^8 \text{ y}$
f)	²³⁴ U	(2.455±0.006) x 10 ⁵ y
g)	[%] Sr	28.78±0.04 y
h)	^{ee} Co	5.2713 <u>+</u> 0.0008 y
i)	⁵⁷ Co	271.79 <u>+</u> 0.09 d
j)	¹³³ Ba	10.52±0.13 y
k)	¹³⁷ Cs	30.07±0.03
1)	¹³⁴ Cs	2.0648±0.001 y
m)	⁵⁴ Mn	312.3±0.4 d

Note: Half-life data are based on NIST certificates (Note 2b) or Evaluated Nuclear Structure Data File (ENSDF 2006). Uncertainties quoted at one sigma level.

(4) The laboratory value represents the mean of five replicate measurements. The reported uncertainty was multiplied by a coverage factor of k=2.

(2b)

- (5) The Difference quoted is the difference between the NIST Value and Reported Value, expressed as a percent relative to the NIST Value.
- (6) Test samples were prepared by gravimetric dilutions of NIST calibrated solutions and SRM's. These solutions and SRM's were calibrated using the following activity measurement methodologies:

a)	Nuclide ²⁴¹ Am	<u>Methodology</u> 4πα liquid-scintillation counting system
b)	²³⁸ Pu	NIST "0.1 π " a defined-solid-angle scintillation detector
		Two 4πα liquid scintillation counting systems
c)	²³⁰ Th ²³⁴ U, ²³⁵ U, ²³⁸ U	Two $4\pi\alpha$ liquid scintillation counting systems
d)	²³⁴ U, ²³⁵ U, ²³⁸ U	Mass spectrometry, silicon surface barrier alpha-detection, and
		$4\pi (\alpha + \beta)$ liquid-scintillation counting systems
e)	⁹⁰ Sr ⁶⁰ Co, ⁵⁷ Co,	NIST 4πβ liquid-scintillation counting system
Ŋ	⁶⁰ Co, ⁵⁷ Co,	Pressurized " 4π " y ionization chamber "A" calibrated using a
		cobalt-60 solution whose activity was determined by " 4π "-(β + γ)- coincidence and anti-coincidence counting
g)	¹³³ Ba	Pressurized " 4π " γ ionization chamber "A" calibrated using a
		barium-133 solution whose activity was determined by " 4π "-(e + X)-gamma-anti-coincidence counting
h)	¹³⁷ Cs, ¹³⁴ Cs,	Pressurized " 4π "- γ -ionization chamber "A" calibrated using a cesium-137 solution whose activity was determined by " 4π "-(e +
		X)-y-anti-coincidence counting
i)	⁵⁴ Mn	Pressurized " 4π " gamma ionization chamber "A" previously standardized by " 4γ " counting with the NBS 8"x8" NaI(TI) crystals

(7) Summary of the reporting laboratory methodologies.

(8) ANSI N42.22 defines the acceptance criteria for verification testing by NIST as:

$$|V_R - V_N| < 3 * \sqrt{u_c^2(N) + u_c^2(R)}$$

Where: $V_N = NIST Value;$

 V_R = Reported Value;

 $u_c(N)$ = standard combine uncertainty of the NIST value, V_N ;

 $u_c(R)$ = standard combine uncertainty of the Laboratory value, V_R ; and

 $3*\sqrt{u_c^2(N)+u_c^2(R)}$ = Traceability Limit (limit to which measurement traceability may be claimed with 99% confidence)

(9) ANSI N13.30 defines criteria for acceptable bias between -25 and +50 percent, and acceptable precision between -40 and +40 percent, 1 sigma total propagated uncertainty.

Reference:

ANSI National Standards Institute, ANSI N42.22-1995, "Traceability of Radioactive Sources to the National Institute of Standards and Technology (NIST) and Associated Instrument Quality Control."

ANSI National Standards Institute, ANSI N13.30-1996, "Performance Criteria for Radiobioassay."

Information contacts:	Dr. Kenneth G. W. Inn	(301) 975-5541	kenneth.inn@nist.gov
	Ms. Svetlana Nour	(301) 975-4927	Svetlana.nour@nist.gov



U.S. DEPARTMENT OF COMMERCE

National Institute of Standards and Technology Gaithersburg, MD

REPORT OF TRACEABILITY

General Engineering Laboratories, LLC Charleston, South Caroline

Test Identification: Test Radionuclides: Matrix Description: Test Activity Range: Reference Time: NRIP-06-SU ²⁴¹Am, ²²⁸Pu, ²³⁹Pu, ²³⁰Th, ²³⁸U, ²³⁵U, ²³⁴U, ⁹⁰Sr, ⁶⁰Co, ¹³³Ba, ¹³⁷Cs, ¹⁵²Eu Synthetic Urine¹ 30mBq•sample⁻¹ to 300mBq•sample⁻¹ 12:00 EST, April 1, 2006

Measurement Results

Nuclide	NIST Value ^{2,3}		Repo	Difference ⁵	
	Massic Activity	Relative Expanded	Massic Activity	Relative Expanded	
	, Bq∙g ^{-t}	Uncertainty (%, k=2)	Bq•g ⁻¹	Uncertainty (%, k=2)	(±% Bias)
²⁴³ Am	1.610	0,95	1.540	10.92	-4.4
^{23#} Pu	1.767	0.68	1.624	11.18	-8.1
²³⁹ Pu	1.685	0.68	1.528	11.28	-9.3
²³⁰ Th	2.009	0.58	1.870	10.70	-6.9
²³⁸ U	4.156	0.60	3.980	10.20	-4.2
234U	4.003	0.98	3.764	10.26	-6.0
235U	0.191	0.62	0.220	19.02	15.1
⁹⁰ Sr	3.869	0.74	5.176	18.88	33.8
60Co	692.7	0.54	677.4	11,40	-2.2
¹³³ Ba	939.3	0.52	881.6	12.48	-6.1
¹³⁷ Cs	1022.0	0.68	1027.2	11.08	0.5
¹⁵² Eu	857.6	0.73	785.6	12.86	-8.4
NR= Not Rep	orted	· · · · · · · · · · · · · · · · · · ·	•	NA= Not	Applicable
		Method	e		
NIST			IST ⁶ Reporting Labo		atory ⁷
Activity Measurements		Alpha- and Beta-S Mass Spectri			Spectrometry

Evaluation (per ANSI N42.22 and N13.30)

Nuclide	N42.22 ⁸		N13.30 ⁵ Results Acceptable per N13.30 Criteria (Pass/Fail)	
	ANSI N42.22 Traceability Traceable Limit			
		(±Percent)	Bias	Precision
²⁴¹ Am	Yes	16	Pass	Pass
²³⁸ Pu	Yes	15	Pass	Pass
²³⁹ Pu	Yes	15	Pass	Pass
²³⁰ Th	Yes	15	Pass	Pass
²³⁸ U	Yes	15	Pass	Pass
²³⁴ U	Ycs	15	Pass	Pass
235U	Yes	33	Pass	Pass
⁹⁰ Sr	Yes	38	Pass	Pass
⁶⁰ Co	Yes	17	Pass	Pass
¹³³ Ba	Yes	18	Pass	Pass
137Cs	Yes	17	Pass	Pass
¹⁵² Eu	Yes	18	Pass	Pass

Samples Distributed May 16, 2006 Reporting Data Received July 14, 2006 For the Director

Acting Leader **Radioactivity Group Physics Laboratory** (Continued)



U.S. DEPARTMENT OF COMMERCE

National Institute of Standards and Technology Gaithersburg, MD

REPORT OF TRACEABILITY

General Engineering Laboratories, LLC Charleston, South Caroline

Test Identification: Test Radionuclides: Matrix Description: Test Activity Range: Reference Time: NRIP-06-SU ²⁴¹Am, ²³⁸Pu, ²³⁹Pu, ²³⁰Th, ²³⁸U, ²³⁵U, ²³⁴U, ⁹⁰Sr, ⁶⁰Co, ¹³³Ba, ¹³⁷Cs, ¹⁵²Eu Synthetic Urine¹ 30mBq•sample⁻¹ to 300mBq•sample⁻¹ 12:00 EST, April 1, 2006

Measurement Results

Nuclide	NIST Value 23 Reported Value ⁴			Difference ⁵	
	Massic Activity	Relative Expanded	Massic Activity	Relative Expanded	
	Bq∙g ⁻¹	Uncertainty (%, k=2)	Bq•g ⁻¹	Uncertainty (%, k=2)	(±% Bias)
²⁴¹ Am	1.610	0.95	1.474	10,94	-8.5
^{23¥} ₽u	1.767	0.68	1.702	10.90	-3.7
²³⁹ Pu	1.685	0.68	1.622	10.96	-3.7
²³⁰ Th	2.009	0.58	1.904	10.72	-5.2
238U	4.156	0.60	4.046	10.28	-2.7
234U	4.003	0.98	3.892	10.30	-2.8
²³⁵ U	0.191	0.62	0.206	20.36	7.5
⁹⁰ Sr	3.869	0.74	5,490	22.04	41.7
⁶⁰ Co	692.7	0.54	710.2	11.44	2.5
¹³³ Ba	939.3	0.52	936.4	13.28	-0.3
¹³⁷ Cs	1022.0	0.68	1062.0	11.44	3.9
¹⁵² Eu	857.6	0.73	820.6	12.60	-4.3
NR= Not Rep	R= Not Reported			NA- Not	Applicable
		Method			
		NIST		Reporting Labor	atory

		Tepering Encorner)
Activity Measurements	Alpha- and Beta-Spectrometry	Alpha, Beta, and Gamma Spectrometry
	Mass Spectrometry	

Nuclide	N4	2.22	N1.	3.309
	ANSI N42.22 Traceable	Traceability Limit		per N13.30 Criteria s/Fail).
		(±Percent)	Bias	Precision
²⁴¹ Am	Yes	15	Pass	Pass
²³⁸ Pu	Yes	16	Pass	Pass
²³⁹ Pu	Yes	16	Pass	Pass
²³⁰ Th	Yes	15	Pass	Pass
238 U	Yes	15	Pass	Pass
²³⁴ U	Yes	15	Pass	Pass
235U	Yes	33	Pass	Pass
⁹⁰ Sr	Yes	47	Pass	Pass
⁶⁰ Co	Yes	18	Pass	Pass
¹³³ Ba	Yes	20	Pass	Pass
²³⁷ Cs	Yes	18	Pass	Pass
¹⁵² Eu	Yes	18	Pass	Pass

Samples Distributed Reporting Data Received July 14, 2006

May 16, 2006

For the Director Telo Tules Michael Unterweger,

Acting Leader Radioactivity Group **Physics Laboratory** (Continued)

As guidance for the proper use of this Report, it should be emphasized that the National Institute of Standards and Technology is concerned only with fostering good measurement capability and consistency with the national measurements system. The assurance of the proper application of that capability to the ultimate consumer products is the responsibility of each manufacturer and of the Federal regulatory agencies.

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Notes

(1a) Five test-samples and three (identical matrix) blanks were provided for this test. Each sample consisted of approximately 1000 grams of synthetic urine material contained in a 1-L size polyethylene bottle.

	Reagent	Weight/Sample (g)
H2C2O4*2H2O	Oxalic Acid	0.02
Pepsin	Pepsin	0.029
СН3СНОНСО2Н	Lactic Acid (Liguid)	0.094
MgSO4*7H2O	Magnesium Sulfate	0.46
C5H11O5CHO	Glucose(dextrose)	0.48
Citric Acid	Citric Acid	0.54
CaCl2*2H2O	Calcium Chloride	0.63
C9H9NO3, 98%	Hippuric Acid	0.63
Na2SiO3*9H2O	Sodium Silicate	0.071
NH4C1, 99%	Ammonium Chloride	1.06
C4H9N3O2*H2O	Creatine	1.1
NaCl, 99+%	Sodium Chloride	2.32
NaH2PO4*H2O	Sodium Dihydrogen Phosphate	2.73
KC1	Potassium Chloride	3.43
Na2SO4	Sodium Sulfate	4.31
CH4N2O, 98%	Urea	16
HNO3	Concentrated nitric acid (50 mL)	70.67
H2O	Water	950
Total		1054.6

Composition of the Synthetic Urine

(1b) The test samples were prepared by depositing a known amount of a NIST calibrated "spike" solution (aqueous solution containing known quantities of ²⁴¹Am, ²³⁸Pu, ²³⁹Pu, ²³⁹Du, ²³⁶U, ²³⁵U, ²³⁴U, ⁹⁰Sr, ⁶⁰Co, ¹³³Ba, ¹³⁷Cs, and ¹⁵²Eu) to the bottle with the urine matrix.

(2a) Solutions of tracers were prepared by gravimetric dilutions of NIST Standard Reference Material SRM's or NIST calibrated solutions. The dilution factors at each step were confirmed by radioactivity measurements.

(2b) The analysis methodology and nomenclature used for the NIST uncertainties are based on uniform guidelines [cf., B.N. Taylor and C. E. Kuyatt, NIST Technical Note 1297 (1994)] and are compatible with those adopted by the principal international metrology standardization bodies.
Individual uncertainties have the significance of one standard deviation of the mean, or an approximation thereof. The relative combined uncertainty is the quadratic combination of the standard deviation (or standard deviation of the mean where appropriate), or approximation thereof, for the following component uncertainties:

	Nuclide (SRM Identification)	Uncertainty (%, 1s)
a)	⁶⁰ Co (4915E)	0.35
b)	⁹⁰ Sr (4919H)	0.37
c)	¹³³ Ba (4251C)	0.26
ď)	¹³⁷ Cs (4233D)	0.34
e)	¹⁵² Eu (4370C)	0.37
f)	²³⁰ Th (4342)	0.29
g)	²³⁴ U (4321C)	0.49
ĥ)	²³⁵ U (4321C)	0.31
i)	²³⁸ U (4321C)	0.30
j) –	²³⁸ Pu (4323A)	0.34
k)	²³⁹ Pu (4330B)	0.34
1)	²⁴¹ Am (4322B)	0.48
	Other Sources	
i)	Gravimetrics (dilutions)	0.05

The individual certified uncertainties of standard reference materials are based on the quadratic combination of all sources of uncertainty manifested in the preparation of the material. These uncertainties may result from uncertainties from any or all of the following: alpha-decay emission rate, background, balance calibration, decay corrections, decay-scheme data, extrapolation of alpha-particle-count-rate-versus-energy to zero energy, live time, alpha-particle detection efficiency, alpha-emitting impurities, gamma-emitting impurities.

The Relative Expanded Uncertainty is obtained by multiplying the standard uncertainty by a coverage factor of k=2 and is assumed to provide an uncertainty interval of approximately 95 percent confidence.

(3) Half-lives used

	Nuclide	Half-life (years)
a)	⁶⁰ Co	5.2713 <u>+</u> 0.0008
b)	⁹⁰ Sr	28.78±0.04
c)	¹³³ Ba	10.51±0.05
d)	¹³⁷ Cs	30.07±0.03
e)	¹⁵² Eu	15.537±0.006
f)	²³⁰ Th	75380±30
g)	²³⁴ U	$(2.455\pm0.006) \times 10^{5}$
h)	235U	$(7.038\pm0.005) \ge 10^8$
i)	²³⁸ U	$(4.468\pm0.003) \ge 10^9$
j)	²³⁸ Pu	87.74±0.04
k) ·	²³⁹ Pu	24110±30
1)	²⁴¹ Am	432.2±0.5

Note: Half-life data are based on NIST certificates (Note 2b). Uncertainties quoted at one sigma level.

- (4) The laboratory value represents the mean of five replicate measurements. The reported uncertainty was multiplied by a coverage factor of k=2.
- (5) The Difference quoted is the difference between the NIST Value and Reported Value, expressed as a percent relative to the NIST Value.
- (6) Test samples were prepared by gravimetric dilutions of NIST calibrated solutions and SRM's. These solutions and SRM's were calibrated using the following activity measurement methodologies:

	Nuclide	Methodology
a)	⁶⁰ Co	Pressurized " 4π " y ionization chamber "A" calibrated using a
		cobalt-60 solution whose activity was determined by " 4π "-(β + γ)-
		coincidence and anti-coincidence counting
b)	⁹⁰ Sr	NIST 4πβ liquid-scintillation counting system
c)	¹³³ Ba	Pressurized " 4π " γ ionization chamber "A" calibrated using a
		barium-133 solution whose activity was determined by " 4π "-(e
		+ X)-gamma-anti-coincidence counting
ď)	¹³⁷ Cs	Pressurized " 4π "- γ -ionization chamber "A" calibrated using a
		cesium-137 solution whose activity was determined by " 4π "-(c +
		X)-y-anti-coincidence counting
		Pressurized " 4π " gamma ionization chamber "A" calibrated
		using a barium-133 solution whose number of cesium-137 atoms
	163	was determined by isotope-dilution mass spectrometry
e)	¹⁵² Eu	Pressurized " 4π " gamma ionization chamber "A" previously
		standardized by "47" counting with the NBS 8"x8" Nal(Tl)
		crystals
f)	²³⁶ Th ²³⁴ U, ²³⁵ U, ²³⁸ U	Two $4\pi \alpha$ liquid scintillation counting systems
g)	²³⁴ U, ²³⁵ U, ²³⁶ U	Mass spectrometry, silicon surface barrier alpha-detection, and
		$4\pi (\alpha + \beta)$ liquid-scintillation counting systems
h)	²³⁸ Pu	NIST "0.1 π " defined-solid-angle scintillation detector
		Two 4na liquid scintillation counting systems
i)	²³⁹ Pu	Two 4na liquid scintillation counting systems
j)	²⁴¹ Am	4πα liquid-scintillation counting system

(7) Summary of the reporting laboratory methodologies.

(8)

ANSI N42.22 defines the acceptance criteria for verification testing by NIST as:

$$|V_R - V_N| < 3 * \sqrt{u_c^2(N) + u_c^2(R)}$$

Where: $V_N = NIST$ Value;

V_R = Reported Value;

 $u_c(N)$ = standard combine uncertainty of the NIST value, V_N ;

 $u_c(R)$ = standard combine uncertainty of the Laboratory value, V_R ; and

 $3*\sqrt{u_c^2(N)+u_c^2(R)}$ = Traceability Limit (limit to which measurement traceability may be claimed with 99% confidence)

(9) ANSI N13.30 defines criteria for acceptable bias between -25 and +50 percent, and acceptable precision between -40 and +40 percent, 1 sigma total propagated uncertainty.

Reference:

ANSI National Standards Institute, ANSI N42.22-1995, "Traceability of Radioactive Sources to the National Institute of Standards and Technology (NIST) and Associated Instrument Quality Control."

ANSI National Standards Institute, ANSI N13.30-1996, "Performance Criteria for Radiobioassay."

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NIST Radiochemistry Intercomparison Program NRIP-2006 Distribution of Reported Results for





Note: Uncertainty bars represent acceptance criteria defined by ANSI N42.22. Participant ID:

GEL 7 and 8

Performance Evaluation Results

Session 10 In Vitro

La	boratory: GEL	Se	ession:	1006
Ma	atrix: SF			
RESL LogNo		Mean Bias Si (Br):	t. Dev Bias (Sb)	Pass / Fail
DL191	Am-241	-0.162	0.045	Pass
DL191	Co-60	0.089	0.034	Pass
DL191	Cs-137	0.032	0.022	Pass
DL191	Pu-238	-0.085	0.019	Pass
DL191	Pu-239	-0.060	0.022	Pass
DL191	Sr-90	0.028	0.051	Pass
DL191	Th-228	-0.065	0.014	Pass
DL191	T h-23 0	0.076	0.035	Pass
DL191	Th-232	-0.100	0.017	Pass
DL191	U-234	-0.013	0.028	Pass
DL191	U-238	0.020	0.022	Pass
	Acceptance Criteria:	-0.25 =< Br <= 0.50	Sb =	-< 0.4

Summary Report









Session Performance Graphs for Each Radionuclide

OGRAM		RESL Log # DL188	Session: 1006	Bias 0.065 0.073 0.105 0.153
TATION PR	IDE REPORT	ot		Experimental Results and 1s Uncertainty 1.32 +/- 0.03 E0 ug/L 1.33 +/- 0.03 E0 ug/L 1.37 +/- 0.03 E0 ug/L 1.43 +/- 0.03 E0 ug/L 1.43 +/- 0.03 E0 ug/L 3t. Dev (Sb) = 0.038 St. Dev (Sb) = 0.038
ABORATORY ACCREDITATION PROGRAM	INDIVIDUAL NUCLIDE REPORT	Radionuclide: U-Tot	Matrix: SU	Known Values and 1s Uncertainty 1.24 +/- 0.04 E0 ug/L 1.24 +/- 0.04 E0 ug/L 1.24 +/- 0.04 E0 ug/L 1.24 +/- 0.04 E0 ug/L 1.24 +/- 0.04 E0 ug/L Mean (Br) = 0.092 Mean (Br) = 0.092
LABORA		Reference Date: 12/1/2006	Laboratory: GEL	Ki Sample # and DL1SU1006EL-1 1.24 + DL1SU1006EL-2 1.24 + DL1SU1006EL-3 1.24 + DL1SU1006EL-6 1.24 + DL1SU1006EL-6 1.24 + DL1SU1006EL-6 1.24 +

Mean (Br) = -0.054 St. Dev (Sb) = 0.041 Acceptance Criteria: -0.25 =< Br <= 0.50 Sb =< 0.4	Reference Date: 12/1/2006 Reference Date: 12/1/2006 Laboratory: GEL Sample # Sample # DL1SU1006MR-1 DL1SU1006MR-4 DL1SU1006MR-5 DL1SU1006MR-6 DL1SU1006MR-6	LABOKATOKKUDITATION PROGRAM INDIVIDUAL NUCLIDE REPORT E Date: 12/1/2006 RESL Log # DL Fatory: GEL Radionuclide: U-238 RESL Log # DL oratory: GEL Matrix: SU Session: 100 ratory: GEL 4.08 +/- 0.13 E0 pC/L -0.044 ratory: R-5 4.08 +/- 0.13 E0 pC/L -0.020 ratory: R-5 4.08 +/- 0.13 E0 pC/L -0.020 ratory: R-6 4.0 +/- 0.3 E	LIDE REPORT 238 238 Experimental Results and 1s Uncertainty 3.9 +/- 0.3 E0 pCi/L 3.6 +/- 0.2 E0 pCi/L 3.8 +/- 0.3 E0 pCi/L 3.8 +/- 0.3 E0 pCi/L 4.0 +/- 0.3 E0 pCi/L	RESL Log # DL180 Session: 1006 Blas -0.044 -0.118 -0.020 -0.020 -0.020	
-0.25 =< Br <= 0.50		Mean (Br) = <i>−</i> 0.054	St. Dev (Sb) = 0.041		
	Acceptance Cri		Sb =< 0.4		

PROGRAM	L	RESL Log # DL180	Session: 1006	lits y Bias	i/L -0.078	i/L -0.066	i/L -0.066	i/L -0.006	i/L -0.061	ß	
ITATION I	LIDE REPOR	h-232	Ŋ	Experimental Results and 1s Uncertainty	3.19 +/- 0.13 E0 pCi/L	3.23 +/- 0.13 E0 pCi/L	3.23 +/- 0.13 E0 pCi/L	3.44 +/- 0.13 E0 pCi/L	3.25 +/- 0.13 E0 pCi/L	St. Dev (Sb) = 0.028	0 Sb =< 0.4
ABORATORY ACCREDITATION PROGRAM	INDIVIDUAL NUCLIDE REPORT	Radionuclide: Th-232	Matrix: SU	Known Values and 1s Uncertainty	3.46 +/- 0.11 E0 pCi/L	Mean (Br) = -0.055	iteria: -0.25 =< Br <= 0.50				
LABORA		Reference Date: 12/1/2006	Laboratory: GEL	Sample #	DL1SU1006MR-2	DL1SU1006MR-3	DL1SU1006MR-4	DL1SU1006MR-5	DL1SU1006MR-6		Acceptance Criteria:

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ROGRAM		RESL Log # DL180	Session: 1006	Bias	0.220	0.159	0.220	0.220	0.236		
TATION PI	LIDE REPORT	-230	_	Experimental Results and 1s Uncertainty	2.22 +/- 0.10 E0 pCi/L	2.11 +/- 0.10 E0 pCi/L	2.22 +/- 0.10 E0 pCi/L	2.22 +/- 0.10 E0 pCi/L	2.25 +/- 0.10 E0 pCi/L	St. Dev (Sb) = 0.030	Sb =< 0.4
ABORATORY ACCREDITATION PROGRAM	INDIVIDUAL NUCLIDE REPORT	Radionuclide: Th-230	Matrix: SU	Known Values and 1s Uncertainty	1.82 +/- 0.04 E0 pCi/L	Mean (Br) = 0.211	sria: −0.25 =< Br <= 0.50				
LABORAT		Reference Date: 12/1/2006	Laboratory: GEL	Sample #	DL1SU1006MR-1 1.	DL1SU1006MR-2 1.	DL1SU1006MR-3 1.	DL1SU1006MR-5 1.	DL1SU1006MR-6 1.		Acceptance Criteria:

ROGRAM		RESL Log # DL180	Session: 1006	Bias	-0.023	-0.014	-0.003	0.032	-0.014		
TATION PR	IDE REPORT	-228		Experimental Results and 1s Uncertainty	3.38 +/- 0.14 E0 pCi/L	3.41 +/- 0.14 E0 pCi/L	3.45 +/- 0.14 E0 pCi/L	3.57 +/- 0.14 E0 pCi/L	3.41 +/- 0.14 E0 pCi/L	St. Dev (Sb) = 0.022	Sb =< 0.4
ABORATORY ACCREDITATION PROGRAM	INDIVIDUAL NUCLIDE REPORT	Radionuclide: Th-228	Matrix: SU	Known Values and 1s Uncertainty	3.46 +/- 0.11 E0 pCi/L	Mean (Br) = -0.005	iteria: -0.25 =< Br <≂ 0.50				
LABORA		Reference Date: 12/1/2006	Laboratory: GEL	Sample #	DL1SU1006MR-1	DL1SU1006MR-2	DL1SU1006MR-3	DL1SU1006MR-4	DL1SU1006MR-6		Acceptance Criteria:

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OGRAM		RESL Log # DL180	Session: 1006	Bias	-0.015	0.069	0.054	0.010	0.005		
TATION PR	IDE REPORT	06		Experimental Results and 1s Uncertainty	1.99 +/- 0.06 E2 pCi/L	2.16 +/- 0.07 E2 pCi/L	2.13 +/- 0.07 E2 pCi/L	2.04 +/- 0.06 E2 pCi/L	2.03 +/- 0.06 E2 pCi/L	St. Dev (Sb) = 0.036	Sb =< 0,4
BORATORY ACCREDITATION PROGRAM	INDIVIDUAL NUCLIDE REPORT	Radionuclide: Sr-90	Matrix: SU	Known Values and 1s Uncertainty	2.02 +/- 0.05 E2 pCi/L	Mean (Br) = 0.025	eria: -0.25 =< Br <≕ 0.50				
LABORAT		Reference Date: 12/1/2006	Laboratory: GEL	Sample #		DL1SU1006MR-3 2	DL1SU1006MR-4	DL1SU1006MR-5	DL1SU1006MR-6 2		Acceptance Criteria:

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ROGRAM		RESL Log # DL180	Session: 1006	Bias	-0.061	-0.055	-0.024	-0.030	-0.043		
TATION PR	IDE REPORT	-239		Experimental Results and 1s Uncertainty	1.54 +/- 0.10 E0 pCi/L	1.55 +/- 0.10 E0 pCi/L	1.60 +/- 0.10 E0 pCi/L	1.59 +/- 0.10 E0 pCi/L	1.57 +/- 0.10 E0 pCi/L	St. Dev (Sb) = 0.016	Sb =< 0.4
ABORATORY ACCREDITATION PROGRAM	INDIVIDUAL NUCLIDE REPORT	Radionuclide: Pu-239	Matrix: SU	Known Values and 1s Uncertainty	1.64 +/- 0.04 E0 pCi/L	Mean (Br) = <i>-</i> 0.043	a: -0.25 =< Br <= 0.50				
BORAT	IJ	: 12/1/2006	: GEL	το	1.6	1.6	1.6	1.6	1.6	2	Acceptance Criteria:
ΓV		Reference Date: 12/1/2006	Laboratory: GEL	Sample #	DL1SU1006MR-1	DL1SU1006MR-2	DL1SU1006MR-3	DL1SU1006MR-5	DL1SU1006MR-6		A

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ROGRAM		RESL Log # DL180	Session: 1006	Bias	-0.096	-0.060	-0.083	-0.043	-0.063		
TATION PR	LIDE REPORT	-238		Experimental Results and 1s Uncertainty	2.73 +/- 0.15 E0 pCi/L	2.84 +/- 0.16 E0 pCi/L	2.77 +/- 0.15 E0 pCi/L	2.89 +/- 0.16 E0 pCi/L	2.83 +/- 0.16 E0 pCi/L	St. Dev (Sb) = 0.021	Sb =< 0.4
ABORATORY ACCREDITATION PROGRAM	INDIVIDUAL NUCLIDE REPORT	Radionuclide: Pu-238	Matrix: SU	Known Values and 1s Uncertainty	3.02 +/- 0.07 E0 pCi/L	Mean (Br) = -0.069	riteria: -0.25 =< Br <= 0.50				
LABORA		Reference Date: 12/1/2006	Laboratory: GEL	Sample #	DL1SU1006MR-1	DL1SU1006MR-2	DL1SU1006MR-3	DL1SU1006MR-5	DL1SU1006MR-6		Acceptance Criteria:

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OGRA		RESL Log # DL180	Session: 1006	Bias	-0.123	-0.078	-0.106	-0.207	-0.022		
TATION PR	IDE REPORT	-137		Experimental Results and 1s Uncertainty	1.57 +/- 0.06 E2 pCi/L	1.65 +/- 0.06 E2 pCi/L	1.60 +/- 0.05 E2 pCi/L	1.42 +/- 0.05 E2 pCi/L	1.75 +/- 0.05 E2 pCi/L	St. Dev (Sb) = 0.067	Sb =< 0.4
ABORATORY ACCREDITATION PROGRAM	INDIVIDUAL NUCLIDE REPORT	Radionuclide: Cs-137	Matrix: SU	Known Values and 1s Uncertainty	1.79 +/- 0.05 E2 pCi/L	Mean (Br) = -0.107	ria: -0.25 =< Br <= 0.50				
LABORAT	I	Reference Date: 12/1/2006	Laboratory: GEL	Sample #	DL1SU1006MR-1	DL1SU1006MR-2 1.	DL1SU1006MR-3 1.	DL1SU1006MR-4 1.	DL1SU1006MR-6 1.		Acceptance Criteria:

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ROGRAM		RESL Log # DL180	Session: 1006	Bias	-0.063	-0.118	0.000	-0.035	-0.160		
TATION PR	IDE REPORT	60		Experimental Results and 1s Uncertainty	1.35 +/- 0.07 E2 pCi/L	1.27 +/- 0.06 E2 pCi/L	1.44 +/- 0.06 E2 pCi/L	1.39 +/- 0.06 E2 pCi/L	1.21 +/- 0.06 E2 pCi/L	St. Dev (Sb) = 0.064	Sb =< 0.4
ABORATORY ACCREDITATION PROGRAM	INDIVIDUAL NUCLIDE REPORT	Radionuclide: Co-60	Matrix: SU	Known Values and 1s Uncertainty	1.44 +/- 0.05 E2 pCi/L	Mean (Br) = -0.075	iteria: -0.25 =< Br <= 0.50				
LABORAT		Reference Date: 12/1/2006	Laboratory: GEL	Sample #	DL1SU1006MR-1	DL1SU1006MR-2	DL1SU1006MR-3	DL1SU1006MR-4	DL1SU1006MR-5		Acceptance Criteria:

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ROGRAM	Å	RESL Log # DL180	Session: 1006	ts / Bias	/L -0.076	/L -0.067	/L -0.058	/L -0.135	/L -0.045		
ITATION P	LIDE REPOR	lm-241	Ŋ	Experimental Results and 1s Uncertainty	2.06 +/- 0.16 E0 pCi/L	2.08 +/- 0.16 E0 pCi/L	2.10 +/- 0.17 E0 pCi/L	1.93 +/- 0.15 E0 pCi/L	2.13 +/- 0.17 E0 pCi/L	St. Dev (Sb) = 0.035	0 Sb =< 0.4
ABORATORY ACCREDITATION PROGRAM	INDIVIDUAL NUCLIDE REPORT	Radionuclide: Am-241	Matrix: SU	Known Values and 1s Uncertainty	2.23 +/- 0.06 E0 pCi/L	Mean (Br) = -0.076	iteria: -0.25 =< Br <= 0.50				
LABORA'		Reference Date: 12/1/2006	Laboratory: GEL	Sample #	DL1SU1006MR-1	DL1SU1006MR-2	DL1SU1006MR-4	DL1SU1006MR-5	DL1SU1006MR-6		Acceptance Criteria:

ROGRAM		RESL Log # DL177	Session: 1006	Bias	-0.048	-0.042	-0.044	-0.054	-0.059		
TATION PR	IDE REPORT	8		Experimental Results and 1s Uncertainty	6.76 +/- 0.10 E4 pCi/L	6.80 +/- 0.10 E4 pCi/L	6.79 +/- 0.10 E4 pCi/L	6.72 +/- 0.10 E4 pCi/L	6.68 +/- 0.10 E4 pCi/L	St. Dev (Sb) = 0.007	Sb =< 0.4
ABORATORY ACCREDITATION PROGRAM	INDIVIDUAL NUCLIDE REPORT	Radionuclide: H-3	Matrix: SU	Known Values and 1s Uncertainty	7.1 +/- 0.2 E4 pCi/L	Mean (Br) = -0.049	eria: -0.25 =< Br <= 0.50				
LABORAT		Reference Date: 12/1/2006	Laboratory: GEL	Sample #	DL1SU1006LB-1 7	DL1SU1006LB-2 7	DL1SU1006LB-3 7	DL1SU1006LB-5 7	DL1SU1006LB-6 7		Acceptance Criteria:

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Lab	oratory: GEL		Session: 100)6
Mat	rix: SU			
RESL LogNo	Radionuclide	Mean Bias (Br):	St. Dev Bias (Sb)	Pass / Fail
DL177	H-3	-0.049	0.007	Pass
DL180	Am-241	-0.076	0.035	Pass
DL180	Co-60	-0.075	0.064	Pass
DL180	Cs-137	-0.107	0.067	Pass
DL180	Pu-238	-0.069	0.021	Pass
DL180	Pu-239	-0.043	0.016	Pass
DL180	Sr-90	0.025	0.036	Pass
DL180	Th-228	-0.005	0.022	Pass
DL180	Th-230	0.211	0.030	Pass
DL180	Th-232	-0.055	0.028	Pass
DL180	U-234	0.043	0.054	Pass
DL180	U-238	-0.054	0.041	Pass
DL188	* U-Tot	0.092	0.038	Pass

Summary Report

Acceptance Criteria: -0.25 =< Br <= 0.50 Sb =< 0.4

* Mass Determination

LABOR	ABORATORY ACCREDITATION PROGRAM	TATION PF	ROGRAM
ana ana ao aminina dia mampione aminina dia mandritry dia mandritry dia mandritry dia mandritry dia mandritry d	INDIVIDUAL NUCLIDE REPORT	LIDE REPORT	
Reference Date: 12/1/2006	6 Radionuclide: Sr-90	06-	RESL Log # DL191
Laboratory: GEL	Matrix: SF		Session: 1006
Sample #	Known Values and 1s Uncertainty	Experimental Results and 1s Uncertainty	Bias
DL1SF1006MR-1	2.02 +/- 0.05 E2 pCi	2.10 +/- 0.05 E2 pCi	0.040
DL1SF1006MR-2	2.02 +/- 0.05 E2 pCi	1.98 +/- 0.04 E2 pCi	-0.020
DL1SF1006MR-3	2.02 +/- 0.05 E2 pCi	2.23 +/- 0.05 E2 pCi	0.104
DL1SF1006MR-4	2.02 +/- 0.05 E2 pCi	1.98 +/- 0.04 E2 pCi	-0.020
DL1SF1006MR-6	2.02 +/- 0.05 E2 pCi	2.09 +/- 0.04 E2 pCi	0.035
	Mean (Br) = 0.028	St. Dev (Sb) = 0.051	
Acceptance Criteria:	Criteria: -0.25 =< Br <= 0.50	Sb =< 0.4	

INDIVIDUAL NUCLIDE REPORT

Reference Date: 12/1/2006	12/1/2006	Radionuclide: U-238	U-238	RESL Log # DL191	DL191
Laboratory: GEL	GEL	Matrix: SF	SF	Session: 1006	1006
Sample #		Known Values and 1s Uncertainty	Experimental Results and 1s Uncertainty	Bias	
DL1SF1006MR-1		4.08 +/- 0.12 E0 pCi	4.1 +/- 0.3 E0 pCi	0.005	
DL1SF1006MR-2		4.08 +/- 0.12 E0 pCi	4.1 +/- 0.3 E0 pCi	0.005	
DL1SF1006MR-4		4.08 +/- 0.12 E0 pCi	4.3 +/- 0.3 E0 pCi	0.054	
DL1SF1006MR-5		4.08 +/- 0.12 E0 pCi	4.2 +/- 0.3 E0 pCi	0.029	
DL1SF1006MR-6		4.08 +/- 0.12 E0 pCi	4.1 +/- 0.3 E0 pCi	0.005	
		Mean (Br) = 0.020	St. Dev (Sb) = 0.022		
Ac	Acceptance Criteria:	riteria: -0.25 =< Br <= 0.50	50 Sb =< 0.4		

OGRAM		RESL Log # DL191	Session: 1006	Bias	0.132	060.0	0.111	0.049	0.063		
TATION PR	IDE REPORT	60		Experimental Results and 1s Uncertainty	1.63 +/- 0.04 E2 pCi	1.57 +/- 0.04 E2 pCi	1.60 +/- 0.04 E2 pCi	1.51 +/- 0.05 E2 pCi	1.53 +/- 0.04 E2 pCi	St. Dev (Sb) = 0.034	Sb =< 0.4
ABORATORY ACCREDITATION PROGRAM	INDIVIDUAL NUCLIDE REPORT	Radionuclide: Co-60	Matrix: SF	Known Values and 1s Uncertainty	1.44 +/- 0.04 E2 pCi	Mean (Br) = 0.089	iteria: -0.25 =< Br <= 0.50				
LABORAT		Reference Date: 12/1/2006	Laboratory: GEL	Sample #	DL1SF1006MR-1	2	DL1SF1006MR-4	DL1SF1006MR-5	DL1SF1006MR-6		Acceptance Criteria:

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LABORATORY ACCREDITATION PROGRAM	INDIVIDUAL NUCLIDE REPORT	Reference Date: 12/1/2006 Radionuclide: Th-232 RESL Log # DL191	Laboratory: GEL Matrix: SF Session: 1006	Known Values Experimental Results ample # and 1s Uncertainty Bias	DL1SF1006MR-1 3.03 +/- 0.10 E0 pCi 2.72 +/- 0.11 E0 pCi -0.102	DL1SF1006MR-2 3.03 +/- 0.10 E0 pCi 2.69 +/- 0.10 E0 pCi -0.112	DL1SF1006MR-3 3.03 +/- 0.10 E0 pCi 2.82 +/- 0.11 E0 pCi -0.069	DL1SF1006MR-4 3.03 +/- 0.10 E0 pCi 2.71 +/- 0.11 E0 pCi -0.106	DL1SF1006MR-6 3.03 +/- 0.10 E0 pCi 2.70 +/- 0.11 E0 pCi -0.109	Mean (Br) = -0.100 St. Dev (Sb) = 0.017	Acceptance Criteria: -0.25 =< Br <= 0.50 Sb =< 0.4
		Referenc	Labo	Sample #	DL1SF1006N	DL1SF1006N	DL1SF1006N	DL1SF1006N	DL1SF1006N		

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LABORA	ABORATORY ACCREDITATION PROGRAM INDIVIDUAL NUCLIDE REPORT	ITATION PF	ROGRAM
Reference Date: 12/1/2006	Radionuclide: Th-230	-230	RESL Log # DL191
Laboratory: GEL	Matrix: SF		Session: 1006
Sample #	Known Values and 1s Uncertainty	Experimental Results and 1s Uncertainty	Bias
DL1SF1006MR-2	1.60 +/- 0.04 E0 pCi	1.70 +/- 0.10 E0 pCi	0.062
DL1SF1006MR-3	1.60 +/- 0.04 E0 pCi	1.71 +/- 0.10 E0 pCi	0.069
DL1SF1006MR-4	1.60 +/- 0.04 E0 pCi	1.80 +/- 0.10 E0 pCi	0.125
DL1SF1006MR-5	1.60 +/- 0.04 E0 pCi	1.75 +/- 0.10 E0 pCi	0.094
DL1SF1006MR-6	1.60 +/- 0.04 E0 pCi	1.65 +/- 0.10 E0 pCi	0.031
	Mean (Br) = 0.076	St. Dev (Sb) = 0.035	
Acceptance Criteria:	≿riteria: -0.25 =< Br <= 0.50	Sb =< 0.4	

OGRAM		RESL Log # DL191	Session: 1006	Bias	-0.079	-0.046	-0.056	-0.073	-0.073		
TATION PR	IDE REPORT	-228		Experimental Results and 1s Uncertainty	2.79 +/- 0.11 E0 pCi	2.89 +/- 0.12 E0 pCi	2.86 +/- 0.11 E0 pCi	2.81 +/- 0.11 E0 pCi	2.81 +/- 0.11 E0 pCi	St. Dev (Sb) = 0.014	Sb =< 0.4
ABORATORY ACCREDITATION PROGRAM	INDIVIDUAL NUCLIDE REPORT	Radionuclide: Th-228	Matrix: SF	Known Values and 1s Uncertainty	3.03 +/- 0.10 E0 pCi	Mean (Br) = -0.065	:riteria: -0.25 ≖< Br <= 0.50				
LABORA	, manna a *1 − 1484 * an a Malana ann an 24 Yur taraon mar a a an ar taraon ann an 1800.	Reference Date: 12/1/2006	Laboratory: GEL	Sample #	DL1SF1006MR-2	DL1SF1006MR-3	DL1SF1006MR-4	DL1SF1006MR-5	DL1SF1006MR-6		Acceptance Criteria:

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[ABO]	LABORATORY ACCREDITATION PROGRAM	NITATION PR	ROGRAM
	INDIVIDUAL NUCLIDE REPORT	CLIDE REPORT	
Reference Date: 12/1/2006	/2006 Radionuclide: Pu-239	Pu-239	RESL Log # DL191
Laboratory: GEL	Matrix: SF	SF	Session: 1006
Sample #	Known Values and 1s Uncertainty	Experimental Results and 1s Uncertainty	Bias
DL1SF1006MR-1	1.64 +/- 0.04 E0 pCi	1.53 +/- 0.10 E0 pCi	-0.067
DL1SF1006MR-3	1.64 +/- 0.04 E0 pCi	1.55 +/- 0.10 E0 pCi	-0.055
DL1SF1006MR-4	1.64 +/- 0.04 E0 pCi	1.55 +/- 0.10 E0 pCi	-0.055
DL1SF1006MR-5	1.64 +/- 0.04 E0 pCi	1.49 +/- 0.10 E0 pCi	-0.091
DL1SF1006MR-6	1.64 +/- 0.04 E0 pCi	1.59 +/- 0.11 E0 pCi	-0.030
	Mean (Br) = -0.060	St. Dev (Sb) = 0.022	
Acceptance	ance Criteria: -0.25 =< Br <= 0.50	50 Sb =< 0.4	

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# DL191	1: 1006								
RESL Log # DL191	Session:	Bias	-0.083	-0.083	-0.063	-0.083	-0.116		
-238		Experimental Results and 1s Uncertainty	2.77 +/- 0.16 E0 pCi	2.77 +/- 0.17 E0 pCi	2.83 +/- 0.17 E0 pCi	2.77 +/- 0.17 E0 pCi	2.67 +/- 0.16 E0 pCi	St. Dev (Sb) = 0.019	Sb =< 0.4
Radionuclide: Pu-238	Matrix: SF	Known Values and 1s Uncertainty	3.02 +/- 0.07 E0 pCi	Mean (Br) = -0.085	teria: -0.25 =< Br <= 0.50				
: 12/1/2006	: GEL		.,						Acceptance Criteria:
Reference Date: 12/1/2006	Laboratory: GEL	Sample #	DL1SF1006MR-1	DL1SF1006MR-2	DL1SF1006MR-3	DL1SF1006MR-4	DL1SF1006MR-5		A

	RESL Log # DL191	Session: 1006	Bias	0.039 0.067	0.017	0.011	0.028		
IDE REPORT			Experimental Results and 1s Uncertainty	1.86 +/- 0.05 E2 pCi 1.91 +/- 0.05 E2 pCi	1.82 +/- 0.04 E2 pCi	1.81 +/- 0.03 E2 pCi	1.84 +/- 0.05 E2 pCi	St. Dev (Sb) = 0.022	Sb =< 0.4
INDIVIDUAL NUCLIDE REPORT	Radionuclide: Cs-137	Matrix: SF	Known Values and 1s Uncertainty	1.79 +/- 0.04 E2 pCi 1.79 +/- 0.04 E2 pCi	1.79 +/- 0.04 E2 pCi	1.79 +/- 0.04 E2 pCi	1.79 +/- 0.04 E2 pCi	Mean (Br) = 0.032	riteria: -0.25 =< Br <= 0.50
	Reference Date: 12/1/2006	Laboratory: GEL	Sample #	DL1SF1006MR-1 DI 1SF1006MR-2	DL1SF1006MR-3	DL1SF1006MR-4	DL1SF1006MR-6		Acceptance Criteria:

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INDIVIDUAL NUCLIDE REPORT

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Reference Date: 12/1/2006	Radionuclide: Am-241	n-241	RESL Log # DL191	0L191
Laboratory: GEL	Matrix: SF		Session: 1	1006
Sample #	Known Values and 1s Uncertainty	Experimental Results and 1s Uncertainty	Bias	
DL1SF1006MR-1	2.23 +/- 0.05 E0 pCi	1.88 +/- 0.13 E0 pCi	-0.157	
DL1SF1006MR-2	2.23 +/- 0.05 E0 pCi	1.96 +/- 0.13 E0 pCi	-0.121	
DL1SF1006MR-4	2.23 +/- 0.05 E0 pCi	1.88 +/- 0.13 E0 pCi	-0.157	
DL1SF1006MR-5	2.23 +/- 0.05 E0 pCi	1.92 +/- 0.13 E0 pCi	-0.139	
DL1SF1006MR-6	2.23 +/- 0.05 E0 pCi	1.70 +/- 0.12 E0 pCi	-0.238	
	Mean (Br) = -0.162	St. Dev (Sb) = 0.045		
Acceptance Criteria:	riteria: -0.25 =< Br <= 0.50	Sb =< 0.4		



















з авј 🖉 0.18 S Lab 4 0.25 Laboratory Performance For Co-60 2 Lab 3 -0.10 С гар 2 -0.08 rds. Lab↑ -0.32 לא Lab 4 д гар 2 60.0 rdej R ssi8

