

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

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

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January 2008

Peer Reviewed by

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### **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, <sup>238</sup>Pu, <sup>239</sup>Pu, <sup>241</sup>Am, <sup>243</sup>Am <sup>235</sup>U, <sup>238</sup>U, elemental uranium and fecal analyses for <sup>241</sup>Am, <sup>238</sup>Pu and <sup>239</sup>Pu 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: <sup>234</sup>U, <sup>235</sup>U, and <sup>238</sup>U. The isotopes are differentiated only during counting by alpha spectrometry. GEL reported that the calculated minimum detectable activity (MDA) for <sup>233,234</sup>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 <sup>233,234</sup>U performance indicators. However, the performance statistics for <sup>235</sup>U and <sup>238</sup>U were reviewed and the MDA for <sup>235</sup>U and the bias and precision for <sup>238</sup>U 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 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.

The total strontium procedure is used to screen samples to determine which will require analysis for  $^{90}$ Sr. 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  $^{90}$ Y in growth to specifically determine  $^{90}$ Sr 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  $^{90}$ Sr 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 <sup>238</sup>Pu and <sup>239</sup>Pu 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 <sup>239</sup>Pu and <sup>238</sup>Pu in urine were acceptable. The 16 samples spiked at the CL for <sup>239</sup>Pu 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 <sup>238</sup>Pu 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 <sup>239</sup>Pu and <sup>238</sup>Pu 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 <sup>238</sup>Pu or the 6 blank <sup>239</sup>Pu fecal analyses were greater than the decision level. There were no fecal samples spiked at the CL with <sup>238</sup>Pu. The six fecal samples spiked with <sup>239</sup>Pu were reported with a result greater than the decision level. Overall the plutonium fecal analyses were considered acceptable.

The <sup>241</sup>Am fecal and urine analysis met the acceptance criteria for MDA, relative bias and precision. With regards to the <sup>241</sup>Am 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 <sup>241</sup>Am fecal duplicate samples were evaluated and it was concluded that the aliquoting procedure produced results within the control limits. A review of the routine <sup>241</sup>Am 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(F<sub>3</sub>) precipitation step. GEL modified their standard operating procedure to include an additional 1 mL of HF when performing the La(F<sub>3</sub>) 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.

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 <sup>242</sup>Cm and <sup>244</sup>Cm and the relative bias and precision for <sup>244</sup>Cm. 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<sub>r</sub>, 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<sub>B</sub>, 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.

Formulas for MDA, B<sub>r</sub>, and S<sub>B</sub>, 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, <sup>90</sup>Sr, <sup>242</sup>Cm, <sup>244</sup>Cm, <sup>238</sup>Pu, <sup>239,240</sup>Pu, <sup>241</sup>Pu <sup>241</sup>Am, <sup>243</sup>Am, <sup>234</sup>U, <sup>235</sup>U, <sup>238</sup>U and elemental uranium and fecal samples for <sup>238</sup>Pu, <sup>239,240</sup>Pu, <sup>241</sup>Am, <sup>242</sup>Cm and <sup>244</sup>Cm. 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.

Battelle Contract 11530 - Feb-06 Table B-3

# Analytical and Reporting Requirements for Routine Processing of Samples

		Contractual Detection	Detection	Determination		; ;		Oral Reporting Level;	ing Level:
		Level (a) <u>(dpm/sample)</u> Urine	m/sample)	Time (business days following	•	Repring Time	<b>d</b> ol		(apm/samble)
Analysis (Code)	Constituents Reported		Fecal	sample receipt)	Oral®	Electronic <sup>(a)</sup>	Written <sup>(x)</sup>	Urine	Feca
Pu(∝) Isotopic (IPU)	Pu-238, Pu-239, 240	0.02	0.2	ଛ	By close of	Within five	Within 10	Ea. 1	Eg. 1
Pu(∝) Isotopic (IPUL)	Pu-238, Pu-239, 240	0.005		30	business on	business	business	E d	-
Am-241 (AM241)	Am-241	0.02	0.8	8	determination	determination	days or		E.
Am-243 (AM243)	Am-243	0.02	9.0	8				· -	
Cm(<) Isotopic (ICM)	Cm-242, Cm-244(b)	0.02		20					· Ť
U(∞) Isotopic (IU)	U-233, 234, U-235, U-	0.02		20				- <del>}</del>	
Th(∞) Isotopic (ITH)	Th-228, Th-229, Th-230,	0.1	-	20				Ê	
Tritium (H3)	In-232 H-3	20 dom/ml		ĸ				Д. -	<u>В</u>
(3)   (4)	(00 20 7 00 20 4117) -0	Ş		, 8				mwdpor	
Sr-total (SR)	Sr (Sum Sr-89 + Sr-90)	2		92				ď	
Sr-90 (SR90) <sup>(a)</sup>	Sr-90	9		8				ĸ	
Gamma Spectroscopy (ISPEC)	K-40, Cs-137 + Others(d)	See Table		20				ı	
Gamma Spectroscopy	Am-241	, 'w		20				-	
(LEPD)								E G	
U-nat (U)	Elemental U	90'0	0.3	20				· Ì	
		ng/sample	ng/sample					0.2	0.2
Sequential Analyses:									
Pu(∞) Iso and Sr-total (IPS)	As for individual analyses	As for individual analyses	ıal analyses	52	Asf	As for individual analyses	ses		
Pu(∝) Iso, Am-241 (IPA)				25					
Pu(≺) Iso, Am-241, Sr-total (IPSA)	al (IPSA)			25					
Pu(∝) Iso, U-nat (IUPU)				25					
Actinide(x) Isotopic (ITPAC) <sup>(e)</sup>	(C)(e)			25					
(IPIU)	1			3					

<sup>(</sup>a) Time allowed following determination of results to receipt of results by Battelle.

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

<sup>(</sup>c) If total Strontlum 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 radionucides in Table B-5 specified in the processing instruction, regardless ofthe activity measured.

<sup>(</sup>e) Pu (∞) Isotopic, Am-241, and Cm (o) Isotopic.

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

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

TABLE 2. Number and Category of Bioassay Samples Analyzed

	FIR	ST CONTRA	CT YEAR	R - GEL	SECOND CONTRACT YEAR - GEL										
Procedure		4/1/05 thro	ugh 3/31/	<u>06</u>		4/1/06 thre	ough 3/3	<u>1/07</u>							
Code(a)	Total	IDP QC	%	GEL QC(b)	Total	IDP QC	%	GEL QC(b)							
Urine		-				•									
H3	795	16	2.0	274	892	3	0.3	276							
SR90, SR	202	1	0.5	476	231	3	1.3	482							
C14	0	0	0.0	0	0	0	0.0	0							
AM241	223	3	1.3	576	103	0	0.0	437							
AM243	114	0	0.0	130	85	6	7.1	122							
U235	2	0	0.0	0	0	0	0.0	0							
ICM	8	0	0.0	0	13	0	0.0	241							
IPU	1687	2	0.1	1555	1243	0	0.0	1152							
IPUL	0	0	0.0	0	1	0	0.0	0							
IPA	357	15	4.2	0	293	4	1.4	N/A							
IPS	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	o	0	0.0	0	0	0	0.0	0							
ICM	ŏ	Ö	0.0	Ö	1	0	0.0	0							
ITH	ő	Ö	0.0	ŏ	0	ō	0.0	0							
AM241	وَ ا	ì	0.0	103	15	ő	0.0	133							
IPU	16	Ô	0.0	99	12	0	0.0	138							
IPA	72	ŷ	12.5	ő	83	12	14.5	N/A							
Total	97	10	10.3	202	111	12	10.8	271							

<sup>(</sup>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.

<sup>(</sup>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<sub>B</sub>) 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<sub>c</sub>) 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<sub>c</sub>, the sample was considered likely to contain the radionuclide of interest. If the measured value was less than L<sub>c</sub>, then the result was considered indistinguishable from a blank. The L<sub>c</sub> was determined solely by measuring blank samples. Before the L<sub>c</sub> 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, Lc is defined by the following equation:

$$L_c = 2s_A$$

where, sA 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<sub>c</sub> 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<sub>c</sub> (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<sub>c</sub> from blank samples, excluding outliers. This L<sub>c</sub> was compared with the L<sub>c</sub> 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.

### **BLAS**

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

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

	Nuclide/	Count	Contract	Counter	Efficiency	Chemi	cal Yield
				<u> 2005-</u>	2006-	2005-	2006-
<u>Matrix</u>	<u>Method</u>	<u>Minutes</u>	Limit(a)	2006	2007	2006	2007
Urine	<sup>3</sup> H	30	20	0.24	0.18		
	Total Sr	60	10	0.379	0.396	0.746	0.774
	SR90	60	10				
	<sup>241</sup> Am	2520	0.02	0.391	0.385	0.883	0.725
	<sup>243</sup> 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
	<sup>237</sup> Np	2520	0.02				
	<sup>239</sup> Pu/ <sup>238</sup> Pu	2520	0.02	0.391	0.385	0.956	0.915
	IPUL	10000	0.005				
	<sup>228</sup> Th/ <sup>230</sup> Th/ <sup>232</sup> 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	<sup>241</sup> Am	960	0.8	0.391	0.385	0.657	0.744
	<sup>238</sup> Pu/ <sup>239</sup> Pu	960	0.2	0.391	0.385	0.88	0.90

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

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.

### **PRECISION**

The precision statistic used for this contract was S<sub>B</sub> from HPS N13.30 (1996), but the limits differ from that standard. S<sub>B</sub> 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 (B<sub>r</sub>).

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

<sup>(</sup>b) Only one sample analyzed

<sup>(</sup>c) NA = Not available. No samples completed.

<sup>(</sup>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.

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.

### **FINDINGS**

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.

TABLE 4. Summary of Statistical Values by Nuclide

	Sample		Blank (dp	m)		Spike	level at CL	. (dpm)	Spike l	Level > 3CI	(dpm)
Isotope(a)	Source	<u>n</u>	<u>L</u> g	<u>MDA</u>	<u>CL</u>	<u>n</u>	<u>B</u> ,	$S_{B}$	<u>n</u>	<u>B</u> ,	<u>S</u> B
<sup>3</sup> H(dpm/mL)					<b>(</b> ()		0.42				
	GEL	138	0.0003	0.6		137	-0.05	0.08			
Total Sr	Taide 2	<del>~~</del> ~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~			10			<b>97</b> 0.23			***
<sup>90</sup> Sr	GEL	16	0.31	7.13		29	0.05	0.14		•••	•••
	GEL	147	0.69	7.95	10	55	-0.03	0.17	231	0.083	0.115
<sup>228</sup> Th	GEL	• • • •	•••	•••	0.02	•••	•••	•••	•••	•••	***
<sup>229</sup> Th	GEL	•••	•••	•••	0.02	•••		•••	•••	•••	***
<sup>230</sup> Th	GEL	***	***	•••	0.02	•••	•••		•	•••	
<sup>242</sup> Cm	GEL	31	0.003	0.010	0.02						•••
<sup>244</sup> Cm	GEL	150	0.003	0.010	0.02		•••		60	0.041	0.101
<sup>238</sup> Pu-urine	-iDE##		in the second second	2018	(141)						3455
	GEL	395	0.003	0.008		1++					
feces	Tide		<b>3</b> 0002	-0.00 Z							
	GEL	49	0.02	0.122					iolidentales		<b>Campus Mil</b> anda puss
<sup>239</sup> Pu-urine	IDP.=2			- 0.000	1919	<b>16</b> -	10.05			8.5	
<b>c</b>	GEL	395	0.003	0.009		219	0.02	0.25	90	-0.056	0.049
feces	IDP.	49	0.02	<b>-0.067</b> 0.131					40	0.054	0.069
<sup>241</sup> Am-urine	Aniber 2	47 31 <b>4 5</b>	0.02	0.131 Mo(89)		 i			49	-0.054	0.058
Am-urine	GEL	150	0.004	0.010	- A. J. (12)	83	0.09	0.37	190	-0.011	0.114
feces	ADD .		0.03	0.010	A D.R	* 35	0.09	0.02	190	· · · · · · · · · · · · · · · · · · ·	0.114
	GEL	41	0.05	0.254	· · · · · · · · · · · · · · · · · · ·		···	***	41	-0.068	0.074
<sup>243</sup> Am-urine	ibr iii		<b>6</b> .0.003	$\mathcal{L}^{0}(0)$	0.02						240
	GEL	42	0.006	0.014		18	0.11	0.31	61	0.026	0.069
$^{234}U$	IDP ***				0.02				eta su	COMP.	
	GEL	94	0.009	0.022 <sup>(e)</sup>					transcer and the second	***	KOLONIA KANTON NILA ()
<sup>235</sup> U	TOPER	112.13		17000	(1)(1)(2)						
•	GEL	94	0.005	0.014	2.500			**************************************	1 a - 100 (10 a 490)		
<sup>238</sup> U	÷ine 🗒				Name N					H0.02 =	0.25
-	GEL	94	0.008	0.019		Like Stabilier andra de		(A) 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	57 <sup>(g)</sup>	-0.04	0.13
									124 <sup>(b)</sup>	0.004	0.084
U-urine (b)	100				OKON.			<b>H</b> ALLY A			60.220
	GEL	198	0.006	0.01		65	-0.04	0.14	67	-0.012	0.076

<sup>(</sup>a) Analyzed in urine matrix unless otherwise noted.

<sup>(</sup>b) Units for L<sub>c</sub>, MDA, and CL are mg per sample.

<sup>(</sup>c) Failed performance criterion.

<sup>(</sup>d) Possible environmental contaminant.

<sup>(</sup>e) Within statistical uncertainty

<sup>(</sup>f) Stats for Cm same as Am-241

<sup>(</sup>g) Spike level 0.15 dpm/S

<sup>(</sup>h) Spike level 0.4 dpm/S

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

		Report	1	В	lanks		Spike Level	at CL	Sį	oike Level >	3CL
Nuclide	CL	Year	n	Lc	MDA	n	$\mathbf{B}_{r}$	$S_B$	n	$\mathbf{B}_{r}$	SB
³H	20 dpm/mL	2005	5	0.17	0.97	11	-0.04	0.18		•••	
		2006	1		***	2	0.42	0.63			111
									(er yekka	or int	
Sr	10 dpm	2005	9	0.43	0.97	5	-0.02	0.15	• • •	•••	•••
	* Thirw's	2006	1			14	0.12	0.23			***
U	0.06 mg	2005	5		0.3571 (d)	6 6	0.01	0.22			
(elemen	-	2005	1	•••		2	-1.6 (c)	2.05 (c)	 15	-0.14	0.22
	ar.							7.74			
<sup>235</sup> U	0.02 dpm	2005	2	0.01	0.0687 (c)	10	-0.10	1.0307 (c)			• • •
_	·	2006	12	0.01	0.02	3	-0.24(c)	0.3		•••	•••
A WATER L			, , , , , ,							<b>HAR</b>	
$^{238}U$	0.02 dpm	2005	2	0.15	0.3508 (d)	10	0.15	0.33			***
44.300.1111.4481.3200000.438130		2006			***	0	0.00	0.00	15	0.02	0.23
										k###	
<sup>238</sup> Pu	0.02 dpm	2005	21	0.003	0.009	9	0.17	0.10			
(urine)		2006	21	0.004	0.011	1	-0.18		en e	acameter control (1957)	111
7					- 77				M. C. C. C.		- 17
<sup>239</sup> 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	54, 578, 57 <u>5</u>		
V (2000)				0.02		emines d'Escar				***	
<sup>239</sup> Рц	0.2 dpm	2005	5	5	0.06				4	-0.02	0.12
				0.02					·		•
(fecal)		2006	6	. 7	0.07	Market Committee	***		6	-0.05	0.09
1 44		a specific		ar, <b>53</b> 8.				<b>30.7</b>			
<sup>241</sup> Am	0.02 dpm	2005	10	0.004	0.012	8	0.05	0.23	•••	•••	•••
(urine)		2006	2	0.005	0.068(c)	17	0.19	0.35	•••		
241										A Section 19	A CONTRACTOR
<sup>241</sup> Am	0.02 dpm	2005	5	0.03	0.085	4	-0.12	0.13		•••	•••
(fecal)	4-4-7	2006	8	0.025	0.063	4	-0.17	0.09	•••		
		x (2001) - 14.40 //		0.02	# <del>*</del> ;			* ( TA 22 )		A STATE OF THE STATE OF	en plante de la comunicación de la
<sup>243</sup> Am	0.02 dpm	2006	2	0	0.09	0		•••			***

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

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

			opikes at	
			Contract	FALSE Negatives
Nuclide	Analyses	Outliers	Level	(%)
Urine				
<sup>3</sup> H	3	0 (0)	2	0 (0)
<sup>90</sup> Sr	15	0 (0)	14	0 (0)
$^{235}U$	15	0 (0)	3	0 (0)
<sup>238</sup> U	0	0 (0)	0	0 (0)
<sup>238</sup> Pu	22	0 (0)	1	0 (0)
<sup>239</sup> Pu	22	0 (0)	16	0 (0)
<sup>241</sup> Am	19	0 (0)	17	0 (0)
<sup>243</sup> Am	0	0 (0)	0	0 (0)
UNAT	3	0 (0)	15	0 (0)
Total	99	0 (0)	68	0 (0)
Feces				
<sup>241</sup> Am	12	0 (0)	4	0 (0)
<sup>238</sup> Pu	12	0 (0)	•••	0 (0)
<sup>239</sup> Pu	12	0 (0)	6	0 (0)
Total	36	0 (0)	10	0 (0)

### **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 <sup>90</sup>Sr. 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 <sup>90</sup>Y in growth to specifically determine <sup>90</sup>Sr 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 <sup>90</sup>Sr 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 <sup>238</sup>Pu and <sup>239</sup>Pu 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 <sup>239</sup>Pu and <sup>238</sup>Pu in urine were acceptable. The 16 samples spiked at the CL for <sup>239</sup>Pu 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 <sup>238</sup>Pu 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 <sup>239</sup>Pu and <sup>238</sup>Pu 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 <sup>238</sup>Pu or the 6 blank <sup>239</sup>Pu fecal analyses were greater than the decision level. There were no fecal samples spiked at the CL with <sup>238</sup>Pu. The six fecal samples spiked with <sup>239</sup>Pu were reported with a result greater than the decision level. Overall the plutonium fecal analyses were considered acceptable.

### <u>URANIUM (UNAT)</u>

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: <sup>234</sup>U, <sup>235</sup>U, and <sup>238</sup>U. The isotopes are differentiated only during counting by alpha spectrometry. GEL reported that the calculated minimum detectable activity (MDA) for <sup>233,234</sup>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 <sup>233,234</sup>U performance indicators. However, the performance statistics for <sup>235</sup>U and <sup>238</sup>U were reviewed and the MDA for <sup>235</sup>U and the bias and precision for <sup>238</sup>U 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	ег	Secon	d Qua	ırter	Third	Quar	ter	Fourt	h qua	rter	3/1/07 - 12/31/07					
Total	Low	Yield	Total	Low	Yield	Total	Low	Yield	Total	Low	Yield	Total		ow eld			
Results	No.	%	Results	No.	%	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 <sup>241</sup>Am fecal and urine analysis met the acceptance criteria for MDA, relative bias and precision. With regards to the <sup>241</sup>Am 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 <sup>241</sup>Am fecal duplicate samples were evaluated and it was concluded that the aliquoting procedure produced results within the control limits. A review of the routine <sup>241</sup>Am 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	er	Fourth	quar	ter	3/1/07 -	12/31/	07
Total	Low	ı	Total	ŀ	Yield	Total		Yield	Total	1	Yield	Total	Lo Yie	eld
Results	No.	%	Results	No.	%	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(F<sub>3</sub>) precipitation step. GEL modified their standard operating procedure to include an additional 1 mL of HF when performing the La(F<sub>3</sub>) 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 (<sup>244</sup>Cm instead of <sup>243</sup>Am). The 6 blank <sup>243</sup>Am 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 <sup>242</sup>Cm and <sup>244</sup>Cm and the relative bias and precision for <sup>244</sup>Cm. 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 <sup>239</sup>Pu, <sup>238</sup>Pu and <sup>241</sup>Am 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 <sup>239</sup>Pu and <sup>238</sup>Pu and about 25 times greater in <sup>241</sup>Am 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:

- 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 <sup>244</sup>Cm 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<sup>(1)</sup>

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<sub>c</sub> was less than ½ 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 <sup>241</sup>Am duplicate was excluded because the results of the parent sample were not reported. Two <sup>243</sup>Am 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 <sup>241</sup>Am 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 <sup>241</sup>Am 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 <sup>242</sup>Pu 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 <sup>90</sup>Sr. 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 <sup>234</sup>U 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.

<sup>(1)</sup> 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 <sup>238</sup>Pu, <sup>239</sup>Pu, <sup>241</sup>Am, <sup>230Th</sup>, <sup>235</sup>U, <sup>238</sup>U, <sup>234</sup>U and <sup>90</sup>Sr in synthetic feces. GEL met the acceptance criteria for relative bias and precision for all isotopes except for <sup>90</sup>Sr, 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 <sup>238</sup>Pu, <sup>239</sup>Pu, <sup>241</sup>Am, <sup>230</sup>Th, <sup>235</sup>U, <sup>238</sup>U, <sup>234</sup>U, <sup>90</sup>Sr, <sup>60</sup>Co, <sup>133</sup>Ba, <sup>137</sup>Cs and <sup>152</sup>Eu in synthetic urine. GEL met the acceptance criteria for relative bias and precision on all isotopes.

On December 1, 2006 GEL participated in the Department of Energy Laboratory Accreditation Program, Session 10. Isotopes tested in a fecal matrix were <sup>238</sup>Pu, <sup>239</sup>Pu, <sup>241</sup>Am, <sup>230</sup>Th, <sup>232</sup>Th, <sup>2328</sup>Th, <sup>238</sup>U, <sup>234</sup>U, <sup>90</sup>Sr, <sup>60</sup>Co, and <sup>137</sup>Cs. Isotopes tested in a urine matrix were <sup>238</sup>Pu, <sup>239</sup>Pu, <sup>241</sup>Am, <sup>230</sup>Th, <sup>232</sup>Th, <sup>2328</sup>Th, <sup>238</sup>U, <sup>234</sup>U, <sup>90</sup>Sr, <sup>60</sup>Co, <sup>137</sup>Cs, tritium and U-total. GEL passed the acceptance criteria for all isotopes in both the fecal and urine matrix for relative bias and precision.

### REFERENCES

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# APPENDIX A QUALITY CONTROL SAMPLE RESULTS

REL BIAS	Time 960 -0.1033 -0.1600 -0.3000	-0.1667 0.0928		Time 960
DET	Chem Yield 0.74  Det Eff 0.39  + +	+ Mean Rel. Bias Mean Rel. Precision		Chem Yield 0.99 Det Eff 0.39
UNCERT 0.0134 0.0138 0.0583 0.0158 0.0191 0.00153	ı	0. <b>18</b> 00	0.0049 0.0057 0.0069 0.0042 0.0062 0.0060 0.0182 0.0048 0.0069	0.0186 0.0470 0.0086 0.0057 0.0334
RESULT 0.0104 -0.0099 -0.0099 0.0165 0.0053		0.80/0 1 <b>0</b> 15	0000	5 DL MDA 0.0152 -0.0166
M N N	-0.0026 0.0133 J	0.7500		0.0055 0.0104 1
1 1XPE	Result F F	Result		Result T T T
UNCERT 0.0000 0.0000 0.0000 0.0000 0.0000 0.0000		0.0450 F Average Result St Dev	000000000000000000000000000000000000000	Average Result St Dev 0.0000 F 0.0000 F
SPIKE 0.0000 0.0000 0.0000 0.0000 0.0000	0.9000 0.9000 0.9000	0.9000 <b>0.9000</b> 4	0.0000.0 0.0000.0 0.0000.0 0.0000.0 0.0000.0 0.0000.0 0.0000.0 0.0000.0	0.0000 0.0000 0.0000
VOL 67.3 245. 205. 186. 160. 41.3 51.8		51.6	67.3 245. 205. 160. 41.3 186. 50.2 50.3 50.3 51.8	67.3 245. 205.
REQ	AM241 Count IPA IPA IPA	PA AM241 Count	# # # # # # # # # # # # # # # # # # #	PU238 Count IPA IPA IPA
1 PAVID 59882 59882 99166 99180 99180 AU001	AU001 AU001 AU001	AU001	59882 59882 99166 99166 99180 AU001 AU001 AU001 AU001	F 59882 59882 99166
TAGWORD J 06K0411 06L0401 06L0414 06L0415 06L0413 06L0437	8 061.0435 061.0433 061.0434	06L0436 4	06K0411 06L0401 06L0415 06L0413 06L0412 06L0435 06L0438 06L0433 06L0433	12 12 06K0411 06L0401
ANAL DATE 12/13/2006 12/13/2006 01/08/2007 01/08/2007 01/11/2007 01/16/2007	01/16/2007 01/16/2007 01/16/2007	0612 14 01/16/2007 Number of total F AM241	12/13/2006 112/13/2006 01/05/2007 01/05/2007 01/11/2007 01/11/2007 01/11/2007 01/11/2007 01/11/2007	Number of total F PU238 0612 17 12/13/2006 0612 18 12/13/2006 0612 19 01/05/2007
YRMO SEO 0612 17 0612 18 0612 19 0612 21 0612 20 0612 15	1 2 2 2	14 r of tot	17 18 19 20 22 21 11 11 16 16 17	er of to 17 18 19
YRMI 0612 0612 0612 0612 0612 0612	0612 0612 0612	0612 Numbe	0612 0612 0612 0612 0612 0612 0612 0612	Numb 0612 0612 0612
1SO CD AM241 AM241 AM241 AM241 AM241 AM241 AM241	AM241 AM241 AM241	AM24]	PU238	PU239 PU239 PU239

REL BIAS								Time 2520			Time 20	0.8650	-0.0300	0.4175																		
<u>Tag</u>	1	•	1	,	,	,	•	Chem Yield 0.73 Det Eff 0.39			Chem Yield 1.00 Det Eff 0.18	+	+	Mean Rel. Bias Mean Rel. Precision		ı	,			,	•		1	,	,		+	ı	1	,		
UNCERT	0.0017	0.0001	0.0001	0.0010	0.0008	0.0024	0.0016	0.0026		0.2580		0.7810	0.8110			0.0016	0.0022	0.0001	0.0001	0.0021	0.0001	0.0019	0.0019	0.0023	0.0001	0.0021	0.0026	6100.0	0.0023	0.0027	0.0008	0.0001
RESULT	-0.0003	-0.0025	-0.0025	0.0007	0.0001	-0.0023	-0.0011	111 DL 113 MDA		-0.0351	SI DL MDA	37.3000	19.4000	-#ZZ		0.0013	0.0021	0.0000	0.0000	-0.0012	-0.0002	0.0000	-0.0013	0.0019	-0.0002	0.0028	0.0055	0.0008	0.0023	0.0022	-0.0034	0.0000
AR .								0.0013			-0.0351 0.0000			12.6572		<b>-</b>	a.	1	Д	_	L	-1	ı	Г	L	Т	ı	Г		_	ᆸ	ф
TYPE	n	n	⊋	Þ	n	ם	n	esult		n	esult	n	n	Sault		Þ	Ω	Ω	Ω	n	Ω	n	Ω	)	D	n	Þ	⊃	n	n	n	ם
UNCERT	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	Average Result St Dev		0.0000 0.0000	Average Result St Dev	0.1780	0.1480	Average Result St Dev		0.000.0	0.0000	0.0000	0.000.0	0.0000	0.0000	0.000.0	0.0000	0.0000	0.0000	0.000.0	0.0000	0.000.0	0.000.0	0.0000	0.0000	0.000.0
SPIKE	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000		0.0000	0.0000	20.0000 0.1780	20.0000 0.1480	20.0000		0.0000	0.0000	0.0000	0.0000	0.0000	0.000.0	0.0000	0.000.0	0.000.0	0.0000	0.0000	0.0000	0.000.0	0.0000		0.0000	0.000.0
XOL S	1097	1407	1407	247	1305	1300	1171	İ		0200	°	0502	0200 2	¤		0982	0660	0997	1239 (	1184	1218	1211 (	1292 (	1183 (	0601	1458 (	1087 (	) /97	1151 (		1426 (	1281 0
REQ ANAL	AM243	AM243 Count		н 3 0	Count	Н 3 0	Н 3 0	Count		IPA 0	IPS 0	IPA 0	IPS 1	IPA 1		_		IPSA 1	IPSA 10	IPSA 14	IPSA 1(	IPSA 12	IPSA 1			IPIU 12						
J <u>eayid</u>	99120	91386	91386	86008	3C136	32514	99152	· ->		3C136	ָ ב	32533	3C136	, =		59001	32476	3G544	99153	91382	50575	50807	59621	99156	50809	31776	99159	32533	99162	51077	80109	15166
TAGWORD J <u>PAYID</u> 19	06D0145	06D0146	06D0146	06G0247	06H0222	8600190	0610159	•	•	06D0142	1	06D0052	06E0059	7	3	06C0318	06C0067	06D0152	06D0252	06E0144	06E0102	06F0121	06F0294	06F0295	06F0122	06F0223	06G0241	06G0067	06H0109	06H0140	06H0115	06H0304
XRMOSEQ DATE Number of total U AM241	04/28/2006	04/28/2006	04/28/2006	07/30/2006	08/28/2006	09/28/2006	09/28/2006		Number of total U AM243	04/20/2006 06D0142		04/20/2006	05/22/2006		Number of total UH 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	08/01/2006	08/14/2006	08/28/2006	08/28/2006		08/31/2006
SEO of tota	03	B	65	01	0	01	05		of tota	05		01	02		eroft	Ξ	60	05	10	90				_		0.4	01	60	ි පි			8
YRMO SEQ Number of tota	0604	0604	0604	090	8090	6090	6090		Number	0604		9604	9990		Num	0602	0602	1090	0604	9005	9090	9090	9090	9090	9090	9090	090	2090	9090	8090		8090
ISO CD	AM243			Н 3		Н 3	н 3			PU238		PU238																				

REL BIAS			Time 2520	-0.1750	-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	0.0478	
<u>DET</u>	1		Chem Yield 0.92 Det Eff 0.39	+	Mean Rel. Bias Mean Rel. Precision		ı	,		,	•	1	Chem Yield 0.92 Det Eff 0.39	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+	Mean Rel. Bias Mean Rel. Precision	
<b>UNCERT</b> 0.0024	0.0019	0.0001	0.0036	0.0046			0.0028	0.0019	0.0020	0.0001	0.0001	0.0021	0.0020	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		
<b>RESULT</b> 0.0011	0.0011	0.0000	0 DE	0.0165	%e		0.0008	0.0003	0.0014	-0.0006	-0.0007	0.0017	S DIL	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	le.	
L KR	J J	1	0.0010	<b>-</b> -,	0.0165		-	<u>ن</u> ہ	J	Д	-	1	0.0005	_	Г	1	Г	-1	J	Г	ı	7		J	В	Г	1	_	J	0.0210	
TYPE	מם	n	esult	ם	esult		Þ	Ω	n	n	Ω	Ω	solt	Ω	₽	Ω	Ω	n	n	D	Ω	Ŋ	n	Ω	n	Ω	n	n	Ω	au It	
UNCERT 0.0000	0.0000		Average Result St Dev	0.0003	Average Result St Dev		0.0000	0.0000	0.0000	0.0000	0.000.0	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	0.0200	0.0200		0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	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> 3	1371	1180	1	1346	1-		0982	0660	7660	1239	1184	1218	1	1346	1211	1090	1183	1292	1458	1087	1267	ıisı	1235	1426	1281	1155	1371	1101	1180	١٣	
31			PU238 Count		PU238 Count								PU239 Count																	PU239 Count	
REQ ANAL IPSA	IPSA IPSA	IPSA		IPA	!		IPA	IPS	ΓA	<b>FPS</b>	IPA	IPSA		ΓPA	IPSA	IPSA	IPSA	IPSA	IPSA	IPSA	<b>IPSA</b>	IPSA	IPSA	IPSA	IPIU	IPSA	IPSA	IPSA	IPSA		
<b>PA</b>	3C134 3C142	85166	' n	3C135	n		59001	32476	3G544	99153	91382	50575	ם	3C135	50807	50809	95166	59621	31776	99159	32533	99162	51077	80109	99151	99161	3C134	3C142	85166	נ	
TAGWORD 1 06.10051	07A0418 07A0417	07A0419	21	06E0058	-	22	06C0318	06C0067	06D0152	06D0252	06E0144	06E0102	9	06E0058	06F0121	06F0122	06F0295	06F0294	06F0223	06G0241	06G0067	06H0109	06H0140	06H0115	06H0304	06J0051	07A0418	07A0417	07A0419	16	ţ
ANAL <u>DATE</u> 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	07/21/2006	900Z/1Z//20	9007/17/20	01/28/2006	08/01/2006	08/14/2006	08/28/2006	08/28/2006	900Z/8Z/80	908/31/2000	11/03/2006	01/23/2007	01/23/2007	01/23/2007		Number of total 11 DI 2320
<u>YRMO SEQ</u> 0610 07	<b>8</b> 2	Ξ		01		er of tot	Ξ	60	8	10	90	60		07	0.5	90	60	80	07	2	60	60	Ξ	10	80	03	60	10	<b></b>		e of tot
XRM 0610	0701	0701		0605		Numb	0602	0602	0.601	9604	0605	9090		9090	9090	9090	9090	9090	9090	2090	2090	8090	8090	8090	8090	0610	0701	0701	0701		Numbe
<u>ISO CD</u> PU238	PU238 PU238	PU238		PU238			PU239 ·	PU239	PU239	PU239	PU239	PU239		PU239	PU239	PU239	PU239														

Number of total U PU239 22

PET RIAS	0.1600	-0.3500	-0.0600	-0.2150	-0.3250	-0.3500	-0.1433 0.2198														Time 2520	-0.5686	0.0986	-0.2600	0.2433		-0.1233	0.0616	-0.1370	-0.0137	0.8219	-0.0274	-0.0822	-0.0548
DEF.	<b>‡</b> +	+	+	+	+	+	Mean Rel. Bias Mean Rel. Precision		,	1	ı	•	•	•		1	1	1	ı		Chem Yield 0.71 Det Eff 0.38	,	•	ı	Mean Rel. Bias Mean Rel. Precision		+	+	+	+	+	+	+	+
d Toassil		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		0.0149	0.0170
T 111270	0.2320	0.1300	0.1880	0.1570	0.1350	0.1300	l≘≘		0.0052	-0.0050	0.0056	0.0028	-0.0017	0.0000	0.0049	0.0019	0.0017	0.0000	0.0000	0.0035	16 DL 11 MDA	0.0030	0.0077	0.0052	SS MDA		0.1280	0.1550	0.1260	0.1440	0.2660	0.1420	0.1340	0.1380
<u> </u>							0.1713		n	Ω	Ω	Ω	Ω	Ω	n	n	Ω	Ω	Ω	Ω	0.0016	m	ם	D	0.0053		В	Ω	n	⊃	n	ב	ס	ם
AT TVPF		n	Þ	n	n	n	Average Result St Dev		<u>.</u>	<u>ר</u>	<b>D</b>	<u> </u>	P	Ω	כ	D	Ω	Þ	ם	n	Average Result St Dev	'n	D	⊃	Average Result St Dev		D	ט	Ω	Ω	n	ח	ח	Ω
INCERT		0.0015	0.0023	0.0016	0.0016	0.0016	-Averag 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	Averag St Dev	0.0000	0.000	0.0000	Averag St Dev		0.0010	0.0010	0.0011	0.0011	0.0010	0.0008	0.0008	0.0008
SPIKE	0.2000	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	0.0000	0.0000	0.0000	0.0000	0.0070	0.0070	0.0070	0.0070		0.1460	0.1460	0.1460	0.1460	0.1460	0.1460	0.1460	0.1460
Z	138	1255	1108	1146	1237	1343			1389	1246	1436	1306	1103	1156	1168	1298	1255	1175	1315	1266		1281	1388	1207			1281	1388	1389	1246	1436	1306	1207	1103
REQ		ñ	ū	n	Ω	ם	Count		5	51	5	51	5	5	19	5	10	5	DI	21	U 235 Count	IPIU	2	2	U 235 Count		IPIU	51	12	13	177	2	5	2
1 DAVID	7 1011	50784	99165	99214	81266	99215	5		3C13S	3C137	91384	25166	AU001	D	15166	3C139	AU001	· •		15166	3C139	3C135	3C137	91384	99157	AU001	AU001							
TACWOOD 1 BAVID	06J0285	0610085	07B0220	07C0408	07C0410	07C0412	15	18	0610279	0610280	07A0233	07B0406	07B0606	07B0600	07B0601	07B0608	07B0607	07B0604	07B0605	07B0602	12	06H0304	06H0399	07B0603	m	15	06H0304	06H0399	0610279	0610280	07A0233	07B0406	07B0603	07B0606
ANAL	10/23/2006	10/23/2006	02/26/2007	03/21/2007	03/21/2007	03/21/2007		Number of total U U	10/01/2009	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	09/08/2006	03/19/2007		Number of total U U235	08/31/2006	9002/80/60	10/01/2006	10/01/2006	02/15/2007	02/22/2007	03/19/2007	03/19/2007
OBSOWAA	8	0.5	12	13	14	15		Jumber	8	92	03	0	\$	05	9	11	10	80	40	\$		80	40	90		er of to	80	9	9	02	03	10	90	60
MOV	0610	0190	0702	0702	0702	0702		~	6090	6090	0701	0703	0702	0702	0702	0702	0702	0702	0702	0702		8090	8090	0702		Num.	8090	8090	6090	6090	0701	0702	0702	0702
(C) (S)	n	Ω	Ω	n	Ŋ	n			U 235		U 235	U 235	U 235			U 238																		

	REL BIAS	-0.1370	0680:0-	-0.0753	0.0685	0.1301	0.0548	-0.0274	Mean Rel. Bias 0.0247
	DET		+	+	+	+	+	+	¥,
	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	. خو ا
	MR	þ	D	n	⊃	n	ם	n	0.1496
	TYPE	n	n	Ω	D	n	n	Þ	sult
	UNCERT	0.0008	0.0008	0.0008	0.0008	0.0008	0.0008	0.1460 0.0008	Average Result
	PIKE	0.1460	0.1460	0.1460	0.1460	0.1460	0.1460	0.1460	0.1460
		1156	1298	1255	1175	1266	1315		
REO	ANAL		₽	1 II	II	5	<b>□</b>	121	U 238
	PAYID	AU001	AU001	AU001	AU001	AU001	AU001	AU001	A
	TAGWORD !	07B0600	07B0608	07B0607	07B0604	07B0602 AU001	07B0605	07B0601	15
ANAL	DATE	03/27/2007	03/27/2007	03/27/2007	03/27/2007	03/27/2007	03/27/2007	03/27/2007	
	YRMO SEQ	05	0702 11	0702 10	80	40	40	05	
	YRM	0702	0702	0702	0702	0702	0702	0702	
	SOCD	U 238	U 238	.U 238	U 238	U 238	U 238	U 238	

Number of total U U 238 15

78 172

Total Samples Total Results

# 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
- 7. 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 – <a href="https://www.statsoft.com/textbook/stquacon.html">www.statsoft.com/textbook/stquacon.html</a>

#### PNNL ANNUAL QC PACKAGE

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

# Data was reviewed and found acceptable.

SUPA	5-79-07
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

= a checked box indicates the data was not used in the calculation of the Exclude

mean and control limits

Stdev = Standard Deviation

= Upper Warning Level (plus 2 sigma) UWL UCL = Upper Contol Level (plus 3 sigma)

= the difference of the individual relative bias from the mean Dispersion

Parent Sample = the sample that was duplicated

TPU = Total Proportion Uncertainty (1 sigma combined standard uncertainty) = Relative Error Ratio (the difference of the individual duplicate pairs RER

based on the combined standard uncertainties of the individual analyses)

= the calculated concentration of the spike in the sample geometry Nominal

Result = the actual measured analyte concentration in the sample = the deviation of a measured value from the expected value Bias

## **Duplicate Acceptance Criteria**

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

**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.

Test	Matrix	QC Samples	Total Samples	% QC
		-	•	76 QC
Am	URINE	437	1084	40%
Am-243	URINE	122	213	<b>57%</b>
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	<sup>.</sup> 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							0.385	

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 <sup>(1) (2)</sup>	Average Relative Bias	Relative Precision
Am-241	Urine	High <sup>1</sup>	-0.0108	0.1136
Am-243	Urine	High <sup>1</sup>	0.0263	0.0693
Cm-243/244	Urine	High <sup>1</sup>	0.0405	0.1006
Pu-239/240	Urine	High <sup>1</sup>	-0.0562	0.0487
Sr-90	Urine	High <sup>t</sup>	0.0828	0.1149
Total U	Urine	High <sup>1</sup>	-0.0119	0.0755
<b>⊍-238</b>	Urine	High <sup>2</sup>	0.0037	0.0843
Am-241	Feces	High¹	-0.0349	0.2686
Pu-239/240	Feces	High <sup>1</sup>	-0.0605	0.0729

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

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

			Average	
		Range	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.

#### 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.

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

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Values less than and greater than

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	Cl omas		Run Dete	27	196	Mess	Mismodille	TOT	Evolute	Tapped Co	200	2	Desert Cample	41.400	
-	1201062888	-	11-Apr-06	-0.86	610	1.14	2	ገሔ			247	314	06C0641	. 0069 and 0.0057	0.00412 and 0.00477
2	1201065026	1659	-	8	-19	1.14	1.17	8 1		0	247	3.14	0600330	0.025 and 0.007	0.0148 and 0.00489
3	1201072748 1710	1710	24-APR-06	98:-	<u>.</u>	1.14	.37	.0025 T		-1.2	2.47	3.14	06D0340	0117 and0099	0.00421 and 0.00249
4	1201067780 1637 16-MAY-06	1637	16-MAY-06	98'-	-19	1.14	1.4	T 610.		4	247	3.14	06E0340	0045 and 0.0244	0.00912 and 0.0186
S	1201104166 1635 09-JUN-06	1635	90-NOY-60				0.00E+00	989					06E0534	.0176 and0176	0.0639 and 0.0848
9	1201105097	1649	1649 09-JUN-06	98	- 19	1,14	26	21		6.1	2.47	3.14	06E0532	0137 and0712	0.0629 and 0.209
_	1201120001 1635 26-JUN-06	1635	26-JUN-06	99	-19	1.14	2.48	710.		8	2.47	3.14	06F0561	.0166 and 0.0283	0.00668 and 0.0168
æ		1636		86	- 19	1.14	2.07	.011 T		1.4	2.47	3.14	06G0265	0208 and 0.0151	0.0136 and 0.0107
6		1696	19-JUL-06	98.	-19	1.14	1.78	.0076 F		-	2.47	3.14	06G0Z76	0019 and0179	0.0049 and 0.00755
₽		1711	01-AUG-06	-86	- 19	1.14	92.	.0032		9	2.47	3.14	06G0431	0206 and0053	0.0202 and 0.00318
÷		- - - - - - - - - - - - - - - - - - -	07-AUG-06	-86	19	1.14	2.03	110	  -	1.3	2.47	3.14	0630440	0176 and 0.0167	0.0127 and 0.0112
2			08-AUG-06	8	9	1.14	2.13	_1 9500°		1.5	2.47	3.14	06G0438	031 and0018	0.0125 and 0.00562
2			1682 16-AUG-06	8	÷.	1.14	.54	.0027 □		6.	2.47	3.14	061+0358	0207 and0032	0.0321 and 0.00286
<u>*</u>	_		1690 24-AUG-06	8	<u>۽</u>	1.14	1.19	.0048 T		١.	2.47	3.14	06H0383	0045 and 0.0036	0.00489 and 0.00475
2			21-SEP-06	8	ē.	1.1	œį	.0029 T		5	2.47	3.14	0610319	0145 and0009	0.0166 and 0.00292
اع			1701 12-OCT-06	8	£.	1.14	36.	.37 ┌	П	-1.2	2.47	3.14	0610535	1.6 and 1.8	0.338 and 0.367
=			1691 14-NOV-06	98	-193	1.14	.061	.0037	П	9:1-	2.473	3.14	06K0326	0.0048 and 0.0044	0.00397 and 0.0037
<b>∞</b>	1201224284	1635	1635 16-NOV-06	-86	- 19	1.14	.33	.0094 Г		7	2.47	3.14	06K0330	0187 and0063	0.0362 and 0.00937
6	1201236923 1680 12-DEC-06	<u>88</u>	12-DEC-06	-86	.19	1,14	25.	.043 T		6.	2.47	3.14	06K0409	0.033 and 0.061	0.0233 and 0.043
8	1201239188 1714 13-DEC-06	1714	13-DEC-06	<u>8</u>	6	1.14	.67	9. 44		-7	2.47	3.14	06K0411	0.01 and021	0.0134 and 0.044
2		989	28-DEC-08	8	ē	-1.4	1.73	.05 T		6:	2.47	3.14	06L0442	0.068 and 0.168	0.0282 and 0.0508
23	1201249840	900	1709 29-DEC-06	8	6	1.14	1.71	<u>اه</u>		6.	2.47	3.14	061.0468	0.021 and01	0.0151 and 0.0103
83	1201251761 1640 31-DEC-06	9	31-DEC-06	8	6	7	1.44	.0041 T		3.	2.47	3.14	06L0637	0.031 and 0.004	0.0183 and 0.00414
Ť	1201243298 1700 08-JAN-07	<u> </u>	08-JAN-07	8	<u>6</u>	1.1	1.15	.012 		0	2.47	3.14	06L0412	0099 and 0.0129	0.0158 and 0.012
1			16-JAN-07	88	6	₹.	69	7	7	?	2.47	3.14	061.0433	0.8 and 0.6	0.169 and 0.138
1			24-JAN-07	8	9	7	58	889 F		-1.3	2.47	3.14	07A0462	0076 and0099	0.00587 and 0.00534
7			07-FEB-07	8	6	÷.	1.25	<u>:</u>		.2	2.47	3,14	07A0686	038 and0101	0.0194 and 0.0111
8 8	-		09-FEB-07	<b>8</b>	9	=	1.13	86 7		0	2.47	3.14	07A0697	0.0003 and 0.0135	0.00676 and 0.0095
7	_	01/1	15-FEB-07	8	£:	<u>-</u>	6;	.013		4	2.47	3.14	07B0404	0.0059 and 0.019	0.00595 and 0.0134
<b>₹</b>		1715	27-FEB-07	8	9	1.1	1.51	.0067		9:	2.47	3.14	07B0520	0.0015 and0099	0.00947 and 0.00667
50	1201282350	1703	1703 01-MAR-07	8	2	=	1.46	.0067 T		5.	2.47	3.14	07B0563	0011 and 0.008	0.00269 and 0.00566
1	1201297669 1689 16-MAH-07	1689	(6-MAH-07	8	9	=	1.58	L 5003		.7	2.47	3.14	07C0442	0099 and 0.0008	0.00597 and 0.00328
1	1201282403 1693 16-MAR-07	1693	6-MAR-07	8	6	- 4	1.93	J 110.		1.2	2.47	3.14	07C0448	0042 and 0.0189	0.00381 and 0.0113
8	1201298266 1715 27-MAR-07	1715	27-MAR-07	8	83	1.14	950	.013		-1.6	2,473	3.14	07B0619	0.024 and 0.023	0.0136 and 0.0134
٦	1201298274 1697 28-MAR-07	169713	38-WAR-07	8	-19	1.14	.94	<u>t</u> .		3	2.47	3.14	07B0519	0.6 and 0.5	0.142 and 0.108

NARRATIVE ATTACHMENT 2 Plutonium-238 Sample Type: RER Attribute Tracked: RER (dec)

Values less than and greater than

or equal to

	Semon III	3	400	2	33	Magn	Photograph and	TON! Evaluate	_ [7	Obdess	in the second	3	Derrord Compile	#ineng	T TOP	
	12040000	+					AND A DESCRIPTION OF THE PERSON OF THE PERSO	╸.		_	+	3 3	TOTAL SOUTH	TOTAL PORT	OLI CONTRACTOR	
-[	ZVII02031	-4		7	200	<b>?</b>	7.62	0.014 T	1	5.0	87.	1.69	06535641	0.0035 and0056	0.00431 and 0.0141	
2	1201065034			-0.71	<del>0</del> .31	0.49	0.37	0.015 T	_	0.3	8	1.69	0600330	0003 and0061	0.00554 and 0.0149	
3	1201065862	1662		- 7061	-,3068	.4822	0.00E+00	.0051 r		-1.2 1	2911	1.6905	06D0346	0 pure 0	0.00508 and 0.00507	
4	1201072761	1706	24-APR-06	14.	31	64	8.	.0045 r		. 9-	.29	1.69	06D0340	0024 and0045	0.00514 and 0.00449	
5	1201087787	1643	18-MAY-06	- 706	-307	.492	.02	.025 F	Ė	12 1	1.291	1.69	06E0340	-,0092 and -,0099	0.023 and 0.0253	
9	1201104169	1640		14.	.31	.49	.15	.027	$\vdash$	6	83	1.69	08E0534	0138 and0097	0.00798 and 0.0267	
7	1201105100 1645	1645	90-XIN7-60	17.	15.	49	101	.012 T		1.3	53	1.69	08E0532	0.0043 and 0.0176	0.00527 and 0.012	
8	1201117818 1836	1836	23-JUN-06	1.2	31	84.	1.6	7 510.	-	2.8	82	1.69	06F0580	0021 and 0.0172	0.0101 and 0.0117	
6	1201128592 1648 17-JUL-06	1648	17-JUL-08	12.	31	64.	89:	.015 T	-	ις	Šį.	1.68	08G0265	0.0064 and0046	0.00578 and 0.0148	
10	1201131875 1702 19-JUL-06	1702	19-JUL-06	12'-	-31	48	1.03	500:	-	6.	1.29	1.69	06G0276	0.0083 and001	0.00755 and 0.00496	
11	1201143166 1715 01-AUG-06	1715	01-AUG-06	12.	31	€	85	.013 T	┝	Q	1.29	1.69	06G0431	0.0048 and0035	0.00564 and 0.0129	
12	1201145210 1715 04-AUG-06	1715	04-AUG-06	-71	15:	9	85	1 7	$\vdash$	H	1.29	1.69	06G0438	0.0051 and0038	0.00823 and 0.0137	
13	1201146286 1637 07-AUG-06	1637	07-AUG-06	12.	15.	49	1.01	  -  -	┝	6.	1.29	1.69	06G0440	0037 and 0.0138	0.0133 and 0.0112	
14	1201154416 1715 15-AUG-06	1715	15-AUG-08	71	31	64.	9	.012 F		ω.	1.29	1.69	06H0358	0.0054 and003	0.00665 and 0.0122	
15	1201159519] 1887  24-AUG-06	1687	24-AUG-08	1907	-3066	4922	0.00E+00	.0048 T	ŀ	12	1.2911	1,6905	06H0383	0 and 0	0.00468 and 0.0048	
16	1201180968	1882	21-SEP-06	12:-	31	64	19	T 8900.	-	8	53	1.89	0610319	10 bra 9200	0.0207 and 0.00677	
17	1201198591	1653	11-OCT-06	12.	31	.49	1.01	.012	-	8.1	84	69.	0610535	0.036 and 0.009	0.0231 and 0.0122	
18	80-VON-61 683 13-NOV-06	1683	13-NOV-06	- 708	-307	28	829:	10. 10.	-	-1.2	1.291	1,69	06K0326	0054 and0059	0.0133 and 0.0157	
19	1201238838	1670	1670 12-DEC-06	14:-	31	67	27	989.	$\vdash$	9	8	1.69	08K0409	0037 and0142	0.00684 and 0.0385	
	1201239191	1708	1708 13-DEC-08	-,7061	-3088	.4822	0.00E+00	L.	Ė	12 1	1.2911	3008	06K0411	0035 and0035	0.00493 and 0.00498	
	1201247785 1703 22-DEC-08	1703	22-DEC-08	708	307	.482	.018	- 1910.	ŀ	1.2.1	1.291	1.69	061.0442	0.035 and 0.035	0.0156 and 0.0154	
	1201248653 1692 27-DEC-06	1692	27-DEC-08	71	31	64.	.47	1 1 10.	┞		23	1.68	061.0465	0.033 and 0.023	0.0161 and 0.0139	
	120(249947   1669   29-DEC-06	1669	29-DEC-06	71	31	49	.54	.014 ⊓	$\vdash$	-	8	69.	061.0468	0,015 and 0,025	0.0103 and 0.0139	
24	1201251768 1636 31-DEC-06	1636	31-DEC-06	-71	31	49	.21	120.		.7	53	69.	061.0637	0.042 and 0.048	0.0186 and 0.0208	
	1201243301 1675 05-JAN-07	1675	05-JAN-07	-71	31	.49	.11	10. F	_	Ĥ	1.29	69.1	06L0412	0026 and0039	0.0052 and 0.0101	
	1201253833 1696 11-JAN-07	1696	11-JAN-07	-71	31	64	.23	.014 □		-7.	1.29	69.1	06L0433	0015 and0048	0.00479 and 0.0137	
	1201261256 1709 23-JAN-07	1709	23-JAN-07	71	-31	-49	.74	.0055 ⊤	_	.8	.29	1.69	07A0462	0.0039 and0015	0.00473 and 0.00546	
	1201272605 1698 08-FEB-07	1696	08-FEB-07	71	-31	-48	.87	L 8900		1 1	- 58	1.69	07A0897	0038 gmd 0.0069	0.0105 and 0.00627	
	1201274085 1695 15 FEB-07	1695	15-FEB-07	.71	31	-49	.53	∟ 2800.	-	-  -	53	69	0780404	0.0017 and 0.0073	0.00512 and 0.00932	
	1201278649 1695 22-FEB-07	1695	22-FEB-07	706	307	492	530.	1900:	-		291	1.69	07A0702	0113 and -,0126	0.038 and 0.00811	
	1201280514 1665 26-FEB-07	1665	26-FEB-07	.71	-31	.49	85.	910.	-	1	1.29	69	07B0520	0.005 and005	0.00611 and 0.0161	
٦	1201280274 1678 26-FEB-07	1678	28-FEB-07	.71	31	.49	.33	.044 T	F	•	1.29	1.69	07B0518	0.2 And 0.2	0.0455 and 0.0441	
	1201282353 1699 01-MAR-07	1699	01-MAR-07	-71	-:31	-49	1.04	.0064 ⊤	_	1.4	1.29	1.69	07B0563	0012 and 0.0071	0.00473 and 0.00639	
	1201280610 1691 14-MAR-07		14-MAR-07	71	31	.49	.79	.013 ⊤	H	7 1 1	1.29	1.69	07C0425	0.0068 and0043	0.00615 and 0.0127	
32	1201282406 1671		15-MAR-07	71	-:31	.49	.12	. 220.	_	9	1.29	1.69	07C0448	0046 and0078	0.0124 and 0.0219	
	1201291676 1697	1697	15-MAR-07	•.71	31	.49	1.09	0043 □		1.5	1.29	1.69	07C0442	0.0075 and0013	0.00682 and 0.0043	
	1201298281 1703 28-MAR-07	1703	28-MAH-07	-71	16.	64'	<del>.</del> 5	220.	ŀ	-	ŀ	1.69	0780519	0.07 and 0.055	0.0249 and 0.022	

# 

# NARRATIVE ATTACHMENT 3 Pktonlum-238/240 Sample Type: RER Attribute Trecked: RER (dec)

Values less than and greater than

or equal to

	Ci omes	1	Ram Oete	101	T.M.	Mean	Numvative	TPG	Exclude	Stoley	<b>₹</b>	2	Parent Sample	Result	UAL
-	1201062891	+	11-Apr-06	1.19	-0.46	66.0	0.63	יייו	[	6.5	2.44	3.16	06C0641	0.06 and 0.083	0.024 and 0.0267
2	1201065034 1663	-	12-Apr-06	-1.19	9	0.99	1.6	0.015		0.8	2.44	3.16	0600390	0.022 and009	0.0126 and 0.0149
8	1201065862	1662		-1.19	-0.46	96.0	19.0	0.014		-0.5	2.44	3.16	06D0346	0007 and0096	0.00359 and 0.014
4	1201072751	_		-1.19	.46	66:	1.02	.0081		0	2.44	3.16	0600340	006 and 0.0099	0.0135 and 0.00807
9	1201087787	-	1643 16-MAY-06	1.1876	.4628	8986	0:00E+00	.0052	֚֡֞֜֝֟֜֜֟֝֜֜֟֜֜֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֜֜֜֓֓֓֓֓֡֓֜֜֜֓֓֓֡֓֜֓֡֓֜֡֡	-1.4	2.4365	3.1613	06E0340	0026 and0026	0.00474 and 0.00523
8	1201104169	-	1640 09-JUN-06	-1.19	-48	86.	2	7,700.		1.4	2.44	3.16	06E0534	0.0013 and016	0.00407 and 0.00765
7	1201105100	-	1645 09-JUN-06	-1.19	94.	86.	86.	.015		6	2.44	3.16	06E0532	.0024 and0082	0.00549 and 0.0149
8	1201117818	1636	23-JUN-06	-1.19	946	<b>6</b> 6:	1.5	.014		1	2.44	3.16	06F0560	0105 and 0.0255	0,0143 and 0,0139
6	1201128592	٠		-1.19	.46	8	2.46	7500.	$\prod_{\mathbf{i}}$	2	2.44	3.16	06G0265	0.029 and006	0.0128 and 0.00565
2	1201131875	┺		-1,19	94.	86.	78.	.0035		-2	2.44	3.16	06G0276	0.0008 and0031	0.00279 and 0.00351
=	1201143166		1715 01-AUG-08	-1.19	94	66:	1.91	.0096 1	<u> </u>	1.3	2.44	3.16	06G0431	0025 and 0.017	0.00333 and 0.00963
72	1201145210	1715	1201145210 1715 04-AUG-06	-1.19	46	66	99.	.0026 r	ָ 	.4	2.44	3.16	06G0438	0006 and 0.0025	0.00367 and 0.0026
5	1201146266	1637	1201146266 1637 07-AUG-06	-1.19	94	86	28:	.013 r	_	-2	2.44	3.16	06G0440	0.0093 and 0.021	0.00631 and 0.0128
4	1201154416	1715	1201154416 1715 15-AUG-06	-1.19	-46	89	29	.0023	•	-1	2.44	3.16	06H0358	0.0008 and0002	0,00263 and 0.00232
12	1201159519	1687	1687 24-AUG-06	-1.19	94.	68.	1.9'	.013		ç	2.44	3.16	06H0383	0041 and0124	0.00331 and 0.0133
9	1201180968	1682	21-SEP-06	-1.19	94.	85	.25	906. -	•	ŀ	2.44	3.16	0610319	0.012 and 0.009	0.00765 and 0.0064
12	1201198591	1653	11-OCT-06	-1.19	97	86	12.	.034		٠١ ا	2.44	3.16	0610535	0.08 and 0.093	0.0346 and 0.0335
18	1201222275	1683	13-NOV-06	-1.19	-46	66:	89	.0088	Π	Ψ-	2.44	3.16	06K0326	0.024 and 0.013	0.0128 and 0.00875
19	1201238838	1670	12-DEC-08	-1.19	94.	66	1.62	.0056 r		6.	2.44	3.16	06K0409	0.02 and003	0.0128 and 0.00563
ន	1201239191	1708	13-DEC-06	61.1-	-46	66:	2.6	5	,	2.2	2.44	3.16	06K0411	0.015 and011	0.00861 and 0.00498
21	1201247785	1703	22-DEC-08	-1.19	46	66	62'	7 660.		6	2.44	3.16	061.0442	0.1 and 0.2	0.0344 and 0.039
8	1201248653	1692	27-DEC-06	-1.1876	-4628	9996	00+300*0	11.	,	-1.4	2.4365	3.1613	061.0465	1 and 1	0.113 and 0.114
ន	1201249947	1669	1669 29-DEC-06	-1.19	94.	66.	1.85	.0055 -		1.2	2.44	3.16	06L0468	0.014 and008	0.0103 and 0.00549
2	1201251768		1636 31-DEC-06	-1.19	.48	66:	1.66	410.	_	6.	2.44	3.16	06L0637	0.022 and011	0.0134 and 0.0143
83	1201243301		1675 05-JAN-07	-1.19	46	66	2.04	.013		1.5	2.44	3.16	06L0412	0074 and 0.0218	0.0052 and 0.0133
92	1201253833	_	1696 11-JAN-07	-1.19	-,46	66	1.49	<u>ا</u> ۔		.7	2.44	3.16	06L0433	1.1 and 0.9	0.12 and 0.103
27	1201261256		1709 23-JAN-07	-1.19	-46	66.	29	.015 r	, 	9'-	2.44	3.16	07A0462	0.0039 and0051	0.00474 and 0.0151
83	1201272605	J	1696 08 FEB-07	-1.19	48	66	.12	.015 T		-12	2.44	3.16	07A0697	.0089 and0053	0.026 and 0.0147
83	1201274085		1695 15-FEB-07	-1.19	48	66	99	.0081		ιĊ	2.44	3.16	0780404	0,0023 and 0,0081	0,00344 and 0,00807
8	1201278649	1695	22-FEB-07	-1.19	94.	66:	£9°	.0062		ď.	2.44	3.16	07A0702	0.006 and 0	0.0074 and 0.00615
31	1201280514	1665	26-FEB-07	-1.19	-46	66.	.83	.016		-:2	2.44	3.16	0780520	0.0034 and0103	0.00306 and 0.0161
g	1201280274	1678	28-FEB-07	-1.19	46	66'	87.	4.		6	2.44	3.16	0780518	1.6 and 1.4	0.149 and 0.141
ន	1201282353	1688	01-MAR-07	-1.19	-,46	66	98.	.013 r	,	- 2	2.44	3.16	0780563	0.0028 and - 0086	0.00255 and 0.013
¥	1201290610	1691	14-MAR-07	-1.19	46	66	64	.011		8.	2.44	3.16	07C0425	0.013 and 0.019	0.00957 and 0.0109
જ	1201292406	1671	15-MAR-07	-1.19	46	66	2.23	.0063		1.7	2.44	3.16	07C0448	0.0009 and0149	0.00334 and 0.00627
ဗ္တ	1201291676	1697	15-MAR-07	188	- 463	286	.026	.012		-1.3	2.436	3.161	07C0442	-,0092 and -,0087	0.0138 and 0.0119
37	1201298281 1703 28-MAR-07	1703	28-MAR-07	-1.19	.46	66.	.74	7,70.	$\prod$	.3	2.44	3.16	07B0519	0.6 and 0.6	0.0862 and 0.0768

NCR Report No.: 312489

Revision No.: 1

COMPANY - WIDE NONCONFORMANCE REPORT					
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 on Application lesues: Failed Recovery for Surrogate	der(s)(SDG):159049,160202,160212	·			
Specification and Requirem Nonconformance Description	ents Nn:	NAG Disposition:			
1. 1201058578 LCS failed to	acer yield.	Because the LCS recove	ary is acceptable, the results are reported.		
	·				
·					

Originator's Name;

Anson Walsh

01-MAY-06

Data Validator/Group Leader:

Robert Timm

05-MAY-06

Quality Review:

Lonnie Morris

03-JUL-06

Director:

Robert Timm

10-NOV-06

Director:

NCR Report No.: 346848
Revision No.: 4

	COMPANY - WIDE NO	CONFORMANCE REPOR	सर
Mo.Day Yr. 17-AUG-06	Division: Bioassay	Quality Criteria: SOP	Type: Process
Instrument Type: BIOASSAY ALPHA	Test / Method: GL-RAD-B-001	Matrix Type: Liquid	Cilent Code: PNNL
Batch ID: 549167	Sample Numbers: 1201137898, 1201137897		
otentially affected wo application issues: failed RPD for LCS/LCS	rk order(s)(SDG):157241,167731		
specification and Requienconformance Desc	rirements ription:	NRG Disposition:	
QCs 1201137896 (L splke recovery requiren splke.	.CS) and 1201137897 (LCS) did not meet the nents for Curium due to missing the Cm-244	The PM was informed, a excepted and is reported duantlyses.	nd the client was contacted. The data was as to no positive results within the Curium
riginator's Name:		Data Validator/Group Lead	er:

rwy. durod

NCR Report No.: 346865 Revision No.: 2

	COMPANY - WIDE N	ONCONFORMANCE REPOR	π
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		
Potentially affected work order(a Application lesues: Failed Recovery for Surrogate or T	itacer		
Specification and Requirements Nonconformance Description:		NRG Disposition:	
1. QC 1201143409 (LCS)'s trace	r yield was less than 25%.	The LCS was recounted.  The results were reported.	The recount confirmed the low tracer yield.

Originator's Name:

Anson Walsh

17-AUG-06

Data Validator/Group Leader:

Robert Timm

28-AUG-06

Quality Review:

Director:

#### SECTION 2

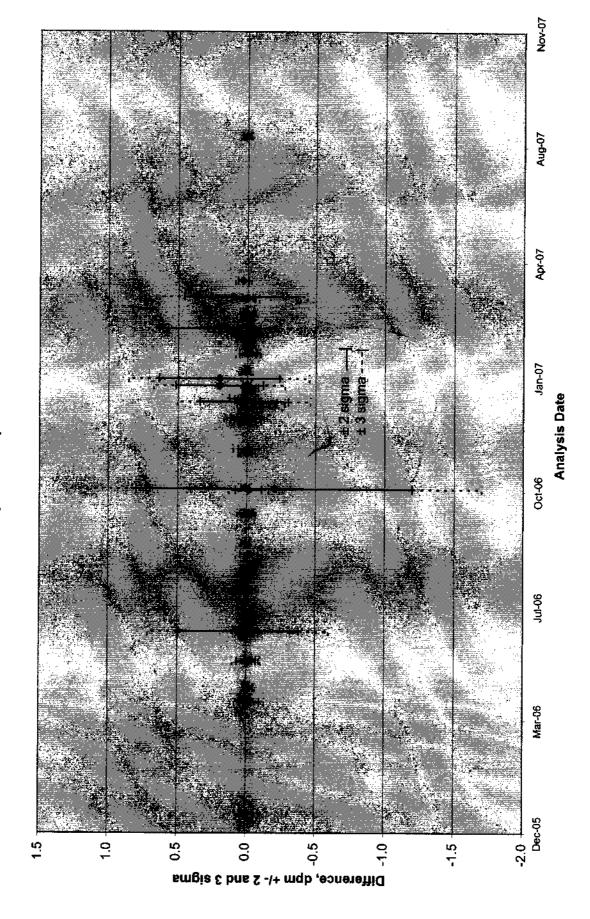
#### DATABASE RESULTS

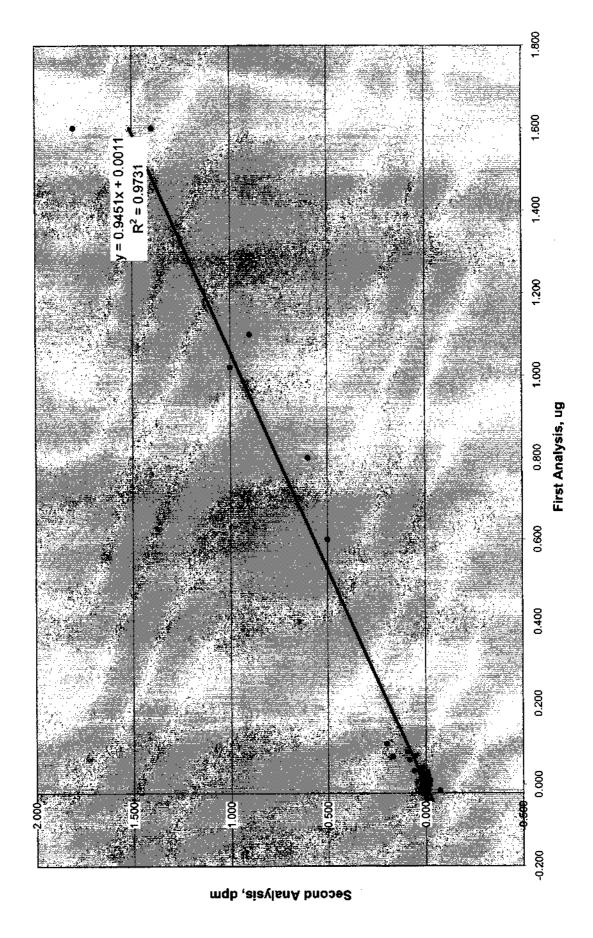
(Maintained electronically in Hanford Internal Dosimetry Program records)

#### APPENDIX C

GEL Duplicates

GEL Fecal Duplicates Apr. 2006 - Mar. 2007





#### 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:

NRIP-06-SF  $^{241} Am, ^{238} Pu, ^{230} Th, ^{238} U, ^{235} U, ^{234} U, ^{90} Sr, ^{60} Co, ^{57} Co, ^{133} Ba, ^{137} Cs, ^{134} Cs, ^{54} Mn$  Synthetic Feces  $^{1}$ 

Matrix Description:

Test Activity Range:

30mBq•sample<sup>-1</sup> to 300mBq•sample<sup>-1</sup>

Reference Time:

12:00 EST, April 1, 2006

#### Measurement Results

Nuclide	NIS	ΓValue <sup>2,3</sup>	Repo	rted Value <sup>4</sup>	Difference <sup>5</sup>
	Massic Activity	Relative Expanded	Massic Activity	Relative Expanded	1
	Bq•g⁻¹	Uncertainty (%, k=2)	Bq•g⁻¹	Uncertainty (%, k=2)	(±% Bias)
<sup>241</sup> Am	1.081	0.95	0.945	14.10	-12.6
<sup>238</sup> Pu	0.938	0.68	0.877	14.72	-6.5
<sup>230</sup> Th	1.630	0.58	1.371	43.91	-15.9
<sup>238</sup> U	4.255	0.62	4.258	10.88	0.1
<sup>234</sup> U	4.098	0.98	4.024	11.04	-1.8
<sup>235</sup> U	0.196	0.64	0.212	31.99	8.1
<sup>90</sup> Sr	3.920	0.74	2.520	54.48	-35,7
NR= Not Rep	orted			NA= Not	Applicable
		Method	\$		
		NIST	<del>.</del>	Reporting Labor	atory <sup>7</sup>
Activity Me	asurements	Alpha- and Beta-Spectrometry Mass Spectrometry		Aipha, Beta, and Gamma Spectrometry	

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

Nuclide	N42.22 <sup>8</sup>		N13.3	09
	ANSI N42.22 Traceable	Traceability Limit	Results Acceptable pe (Pass/F	
		(±Percent)	Bias	Precision
<sup>241</sup> Am	Yes	19	Pass	Pass
<sup>238</sup> Pu	Yes	21	Pass	Pass
<sup>230</sup> Th	Yes	55	Pass	Pass
$^{238}U$	Yes	16	Pass	Pass
<sup>234</sup> U	Yes	16	Pass	Pass
<sup>235</sup> U	Yes	52	Pass	Pass
<sup>90</sup> Sr	Yes	53	Fail	Pass

Samples Distributed Reporting Data Received

October 21, 2006 December 21, 2006 For the Director

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:

NRIP-06-SF  $^{241}Am,\,^{238}Pu,\,^{230}Th,\,^{238}U,\,^{235}U,\,^{234}U,\,^{90}Sr,\,^{60}Co,\,^{57}Co,\,^{133}Ba,\,^{137}Cs,\,^{134}Cs,\,^{54}Mn$ 

Matrix Description:

Synthetic Feces<sup>1</sup>

Test Activity Range:

30mBq+sample<sup>-1</sup> to 300mBq+sample<sup>-1</sup>

Reference Time:

12:00 EST, April 1, 2006

#### Measurement Results

Nuclide	NIS'	Γ Value <sup>2,3</sup>	Repo	rted Value <sup>4</sup>	Difference <sup>5</sup>
	Massic Activity	Relative Expanded	Massic Activity	Relative Expanded	1
	Bq•g⁻¹	Uncertainty (%, k=2)	Bq•g⁻¹	Uncertainty (%, k=2)	(±% Bias)
241Am	1.081	0.95	0.987	14.50	-8.7
<sup>238</sup> Рц	0.938	0.68	0.852	13.48	-9.2
<sup>230</sup> 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
<sup>235</sup> U	0.196	0.64	0.224	23.16	14.6
<sup>90</sup> Sr	3.920	0.74	3.376	20.62	-13.9
NR= Not Rep	orted		·	NA= Not	Applicable
		Method	S		
		NIST <sup>6</sup>		Reporting Labor	atory'
Activity Measurements		Alpha- and Beta-Spectrometry  Mass Spectrometry		Alpha, Beta, and Gamma Spectrometry	

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

Nuclide	N42.22*		N13.3	09
	ANSI N42.22 Traceable	Traceability Limit	Results Acceptable pe (Pass/F	
	(±Percent)		Bias	Precision
<sup>241</sup> Am	Yes	20	Pass	Pass
<sup>236</sup> Pu	Yes	18	Pass	Pass
<sup>230</sup> Th	Yes	15	Pass	Pass
$^{238}U$	Yes	16	Pass	Pass
<sup>234</sup> U	Yes	16	Pass	Pass
<sup>235</sup> U	Yes	40	Pass	Pass
<sup>90</sup> Sr	Yes	27	Pass	Pass

Samples Distributed Reporting Data Received October 21, 2006

December 21, 2006

For the Director

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

Composition of the Syn	thetic 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 (distifled)	65

- (1b) The test samples were prepared by depositing a known amount of a NIST calibrated "spike" solution (aqueous solution containing known quantities of <sup>241</sup>Am, <sup>238</sup>Pu, <sup>236</sup>Th, <sup>238</sup>U, <sup>235</sup>U, <sup>234</sup>U, <sup>90</sup>Sr, <sup>57</sup>Co, <sup>60</sup>Co, <sup>54</sup>Mn, <sup>137</sup>Cs, <sup>134</sup>Cs, and <sup>133</sup>Ba 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.

(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)	<sup>241</sup> Am (4322B)	0.48
b)	<sup>238</sup> Pu (4323A)	0.34
c)	<sup>230</sup> Th (4342)	0.29
d)	<sup>238</sup> U (4321C)	0.31
e)	<sup>235</sup> U (4321C)	0.32
f)	<sup>234</sup> U (4321C)	0.49
g)	<sup>90</sup> Sr (4919H)	0.37
h)	<sup>60</sup> Со (4915Е)	0.27
i)	<sup>57</sup> Co (Analytics, NIST calibration)	0.35
j)	<sup>133</sup> Ba (4251C)	0.26
k)	<sup>137</sup> Cs (4233D)	0.34
1)	<sup>134</sup> Cs (4370C)	0.37
m)	54Mn (Analytics, NIST calibration)	0.34
	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

	<u>Nuclide</u>	Half-life
a)	<sup>241</sup> Am	432.2±0.5 y
b)	<sup>238</sup> Pu	87.74±0.04 y
c)	<sup>230</sup> <b>T</b> h	75380±30 y
d)	$^{238}\mathrm{U}$	$(4.468\pm0.003) \times 10^9 \text{ y}$
e)	<sup>235</sup> U	$(7.038\pm0.005) \times 10^8 \text{ y}$
f)	<sup>234</sup> U	$(2.455\pm0.006) \times 10^5 \text{ y}$
g)	90Sr	28.78±0.04 y
h)	<sup>60</sup> Co	5.2713 <u>+</u> 0.0008 y
i)	<sup>57</sup> Co	271.79±0.09 d
j)	<sup>133</sup> Ba	10.52±0.13 y
k)	<sup>137</sup> Cs	30.07±0.03
1)	<sup>134</sup> Cs	2.0648±0.001 y
m)	<sup>54</sup> 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.

- (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:

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

u<sub>c</sub>(N)= standard combine uncertainty of the NIST value, V<sub>N</sub>;

u<sub>c</sub>(R)= standard combine uncertainty of the Laboratory value, V<sub>R</sub>; 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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#### 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 239Pu, 239Pu, 230Th, 238U, 235U, 234U, 90Sr, 60Co, 133Ba, 137Cs, 152Eu Synthetic Urine 30mBq\*sample to 300mBq\*sample 1 to 300mBq\*sample 1 12:00 EST, April 1, 2006

Measurement Results

Nuclide	NIS'	ľ Value <sup>2,3</sup>	Repo	Reported Value			
	Massic Activity	Relative Expanded	Massic Activity	Relative Expanded	]		
	Bq•g¹	Uncertainty (%, k=2)	Bq•g⁻¹	Uncertainty (%, k=2)	(±% Bias)		
241 Am	1.610	0,95	1.540	10.92	-4.4		
<sup>238</sup> Pu	1.767	0.68	1.624	11.18	-8.1		
<sup>239</sup> Pu	1.685	0.68	1.528	11.28	-9.3		
<sup>230</sup> Th	2.009	0.58	1.870	10.70	-6.9		
<sup>238</sup> U	4.156	0.60	3.980	10.20	-4.2		
<sup>234</sup> U	4.003	0.98	3.764	10.26	-6.0		
<sup>235</sup> U	0.191	0.62	0.220	19.02	15.1		
90Sr	3.869	0.74	5.176	18.88	33.8		
<sup>60</sup> Co	692.7	0.54	677.4	11.40	-2.2		
<sup>133</sup> Ba	939.3	0.52	881.6	12.48	-6.1		
137 <b>C</b> s	1022.0	0.68	1027.2	11.08	0.5		
<sup>152</sup> Eu	857.6	0.73	785.6	12.86	-8.4		
NR= Not Rep	NR= Not Reported NA= Not Applicable						
	-	Method	1				
NIST <sup>6</sup> Reporting Laboratory							

Reporting Laboratory **Activity Measurements** Alpha- and Beta-Spectrometry Alpha, Beta, and Gamma Spectrometry Mass Spectrometry

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

Nuclide	N42.22 <sup>8</sup>		N13.30 <sup>9</sup>		
	ANSI N42.22 Traceable	· I	Results Acceptable per (Pass/Fa		
			Bias	Precision	
241Am	Yes	16	Pass	Pass	
<sup>238</sup> Pu	Yes	15	Pass	Pass	
<sup>239</sup> Pu	Yes	15	Pass	Pass	
<sup>230</sup> Th	Yes	15	Pass	Pass	
<sup>238</sup> U	Yes	15	Pass	Pass	
$^{234}U$	Yes	15	Pass	Pass	
<sup>235</sup> U	Yes	33	Pass	Pass	
%Sr	Yes	38	Pass	Pass	
<sup>60</sup> Co	Yes	17	Pass	Pass	
<sup>133</sup> Ba	Yes	18	Pass	Pass	
<sup>137</sup> Cs	Yes	17	Pass	Pass	
152 <b>E</b> u	Yes	18	Pass	Pass	

Samples Distributed

May 16, 2006 Reporting Data Received July 14, 2006

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-SU 239 Pu, 239 Pu, 230 Th, 238 U, 235 U, 234 U, 90 Sr, 60 Co, 133 Ba, 137 Cs, 152 Eu Synthetic Urine 1 30 mBq • sample 1 to 300 mBq • sample 1 12:00 EST, April 1, 2006

Measurement Results

Nuclide	NIST Value 2,3		Reported Value <sup>4</sup>		Difference <sup>5</sup>	
	Massic Activity	Relative Expanded	Massic Activity	Relative Expanded		
	Bq•g <sup>-1</sup>	Uncertainty (%, k=2)	Bq•g⁻¹	Uncertainty (%, k=2)	(±% Bias)	
<sup>241</sup> Am	1.610	0.95	1.474	10.94	-8.5	
<sup>23‡</sup> Рц	1.767	0.68	1.702	10.90	-3.7	
<sup>239</sup> Pu	1.685	0.68	1.622	10.96	-3.7	
<sup>230</sup> Th	2.009	0.58	1.904	10.72	-5.2	
<sup>238</sup> U	4.156	0.60	4.046	10.28	-2.7	
<sup>234</sup> U	4.003	0.98	3.892	10.30	-2.8	
<sup>235</sup> U	0.191	0.62	0.206	20.36	7.5	
<sup>90</sup> Sr	3.869	0.74	5,490	22.04	41.7	
<sup>60</sup> Co	692.7	0.54	710.2	11.44	2.5	
<sup>133</sup> Ba	939.3	0.52	936.4	13.28	-0.3	
<sup>137</sup> Cs	1022.0	0.68	1062.0	11.44	3.9	
<sup>152</sup> Eu	857.6	0.73	820.6	12.60	-4.3	
NR= Not Reported NA= Not Applicable					Applicable	

		Methods	
		NIST	Reporting Laboratory
	Activity Measurements	Alpha- and Beta-Spectrometry	Alpha, Beta, and Gamma Spectrometry
- 1		Mass Spectrometry	· · · · · · · · · · · · · · · · · · ·

Evaluation (ner ANSI N42 22 and N12 20)

Nuclide	N4	2,22	N13.30°		
• •	ANSI N42.22 Traceable		Results Acceptable per N13.30 Criteria (Pass/Fail).		
		(±Percent)	Bias	Precision	
<sup>241</sup> Am	Yes	15	Pass	Pass	
<sup>238</sup> Pu	Yes	16	Pass	Pass	
<sup>239</sup> Pu	Yes	16	Pass	Pass	
<sup>230</sup> Th	Yes	15	Pass	Pass	
<sup>238</sup> U	Yes	15	Pass	Pass	
<sup>234</sup> U	Yes	15	Pass	Pass	
<sup>235</sup> U	Yes	33	Pass	Pass	
<sup>90</sup> Sr	Yes	47	Pass	Pass	
<sup>60</sup> Co	Yes	18	Pass	Pass	
<sup>133</sup> Ba	Yes	20	Pass	Pass	
<sup>237</sup> Cs	Yes	18	Pass	Pass	
<sup>152</sup> Eu	Yes	18	Pass	Pass	

Samples Distributed Reporting Data Received July 14, 2006

May 16, 2006

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.

(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.

#### Composition of the Synthetic Urine

	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
С5Н11О5СНО	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
KCl	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

- (1b) The test samples were prepared by depositing a known amount of a NIST calibrated "spike" solution (aqueous solution containing known quantities of <sup>241</sup>Am, <sup>238</sup>Pu, <sup>239</sup>Pu, <sup>239</sup>Pu, <sup>236</sup>U, <sup>235</sup>U, <sup>234</sup>U, <sup>90</sup>Sr, <sup>60</sup>Co, <sup>133</sup>Ba, <sup>137</sup>Cs, and <sup>152</sup>Eu) 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)	<sup>60</sup> Co (4915E)	0.35
b)	<sup>90</sup> Sт (4919Н)	0.37
c)	<sup>133</sup> Ba (4251Ć)	0.26
ď)	<sup>137</sup> Cs (4233D)	0.34
e)	<sup>152</sup> Eu (4370C)	0.37
f)	<sup>230</sup> Th (4342)	0.29
g)	<sup>234</sup> U (4321C)	0.49
h)	<sup>235</sup> U (4321C)	0.31
i)	<sup>238</sup> U (4321C)	0.30
j)	<sup>238</sup> Pu (4323A)	0.34
k)	<sup>239</sup> Pu (4330B)	0.34
1)	<sup>241</sup> 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 Nuclide	<u> Half-life (years)</u>
	5.2713 <u>+</u> 0.0008
<sup>90</sup> Sr	28.78±0.04
<sup>133</sup> Ba	10.51±0.05
<sup>137</sup> Cs	30.07±0.03
<sup>152</sup> Eu	15.537±0.006
<sup>230</sup> Th	75380±30
	$(2.455\pm0.006) \times 10^{5}$
<sup>235</sup> U	$(7.038\pm0.005) \times 10^{8}$
<sup>238</sup> U	$(4.468\pm0.003) \times 10^9$
<sup>238</sup> Pu	87.74±0.04
<sup>239</sup> Pu	24110±30
<sup>241</sup> Am	432.2±0.5
	60°Co 90°Sr 133°Ba 137°Cs 152°Eu 230° Th 234°U 235°U 238°U 238°Pu

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	<u>Methodology</u>
a)	<sup>60</sup> Co	Pressurized "4π" γ ionization chamber "A" calibrated using a
,		cobalt-60 solution whose activity was determined by " $4\pi$ "- $(\beta+\gamma)$ -
		coincidence and anti-coincidence counting
<b>b</b> )	90Sr	NIST 4πβ liquid-scintillation counting system
c)	<sup>133</sup> Ba	Pressurized "4π" γ ionization chamber "A" calibrated using a
		barium-133 solution whose activity was determined by "4π"-(e
	117	+ X)-gamma-anti-coincidence counting
ď)	<sup>137</sup> Cs	Pressurized "4π"-γ-ionization chamber "A" calibrated using a
		cesium-137 solution whose activity was determined by " $4\pi$ "-(c +
		X)-γ-anti-coincidence counting
		Pressurized "4π" gamma ionization chamber "A" calibrated
		using a barium-133 solution whose number of cesium-137 atoms
		was determined by isotope-dilution mass spectrometry
e)	<sup>152</sup> Eu	Pressurized "4π" gamma ionization chamber "A" previously
		standardized by "47" counting with the NBS 8"x8" NaI(Tl)
		crystals
f)	<sup>230</sup> Th	Two 4πα liquid scintillation counting systems
g)	<sup>234</sup> U, <sup>235</sup> U, <sup>238</sup> U	Mass spectrometry, silicon surface barrier alpha-detection, and
		$4\pi (\alpha + \beta)$ liquid-scintillation counting systems
h)	<sup>238</sup> Pu	NIST "0.1π"α defined-solid-angle scintillation detector
,		Two 4πα liquid scintillation counting systems
i)	<sup>239</sup> Pu	Two 4πα liquid scintillation counting systems
j)	<sup>241</sup> Am	4πα liquid-scintillation counting system
27		d

- (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<sub>R</sub> = Reported Value;

u<sub>c</sub>(N)= standard combine uncertainty of the NIST value, V<sub>N</sub>;

u<sub>c</sub>(R)= standard combine uncertainty of the Laboratory value, V<sub>R</sub>; 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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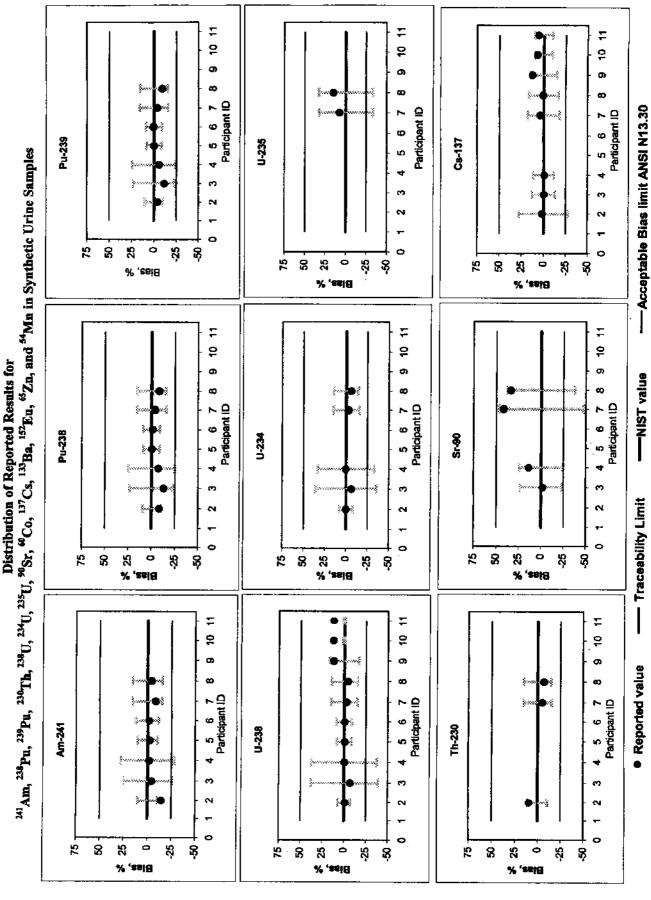
kenneth.inn@nist.gov

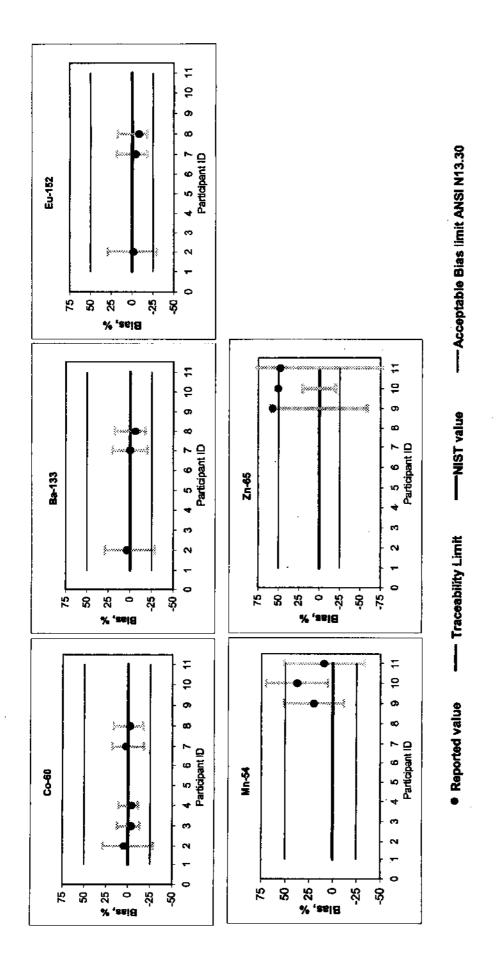
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NIST Radiochemistry Intercomparison Program NRIP-2006





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

GEL 7 and 8

#### DEPARTMENT OF ENERGY LABORATORY ACCREDITATION PROGRAM

#### Performance Evaluation Results

Session 10 In Vitro

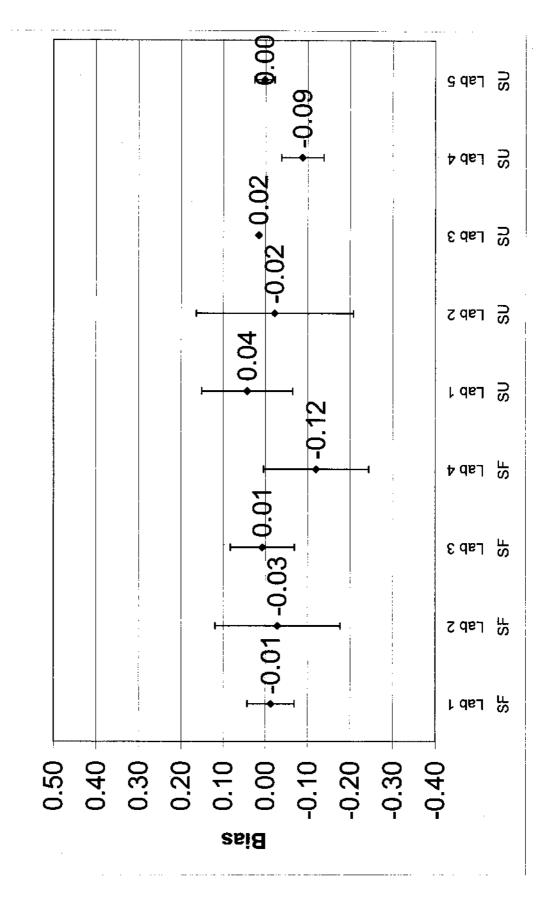
#### **DEPARTMENT OF ENERGY** LABORATORY ACCREDITATION PROGRAM

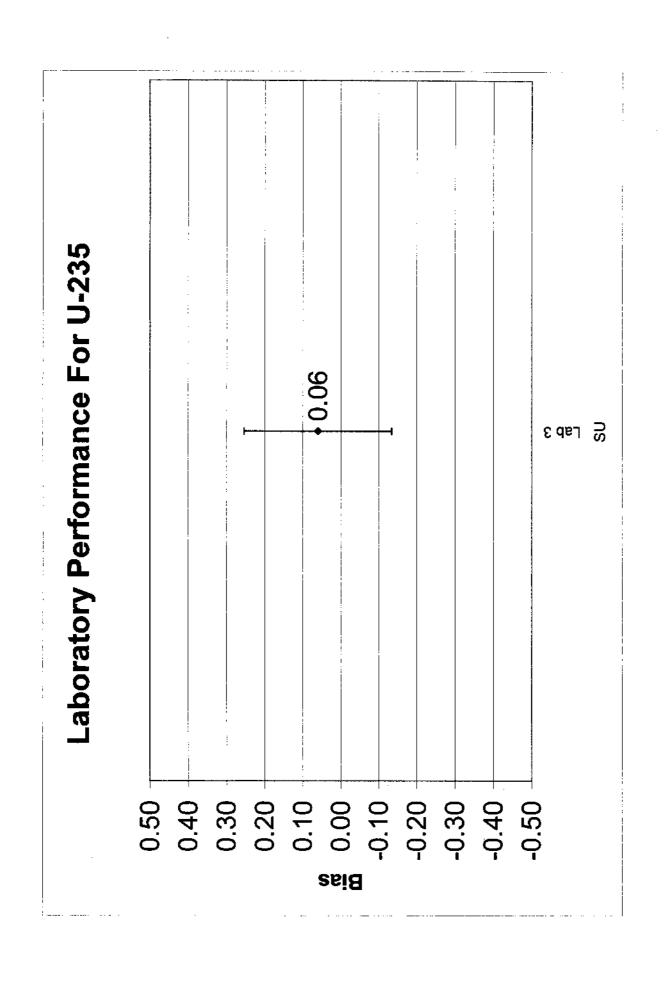
#### **Summary Report**

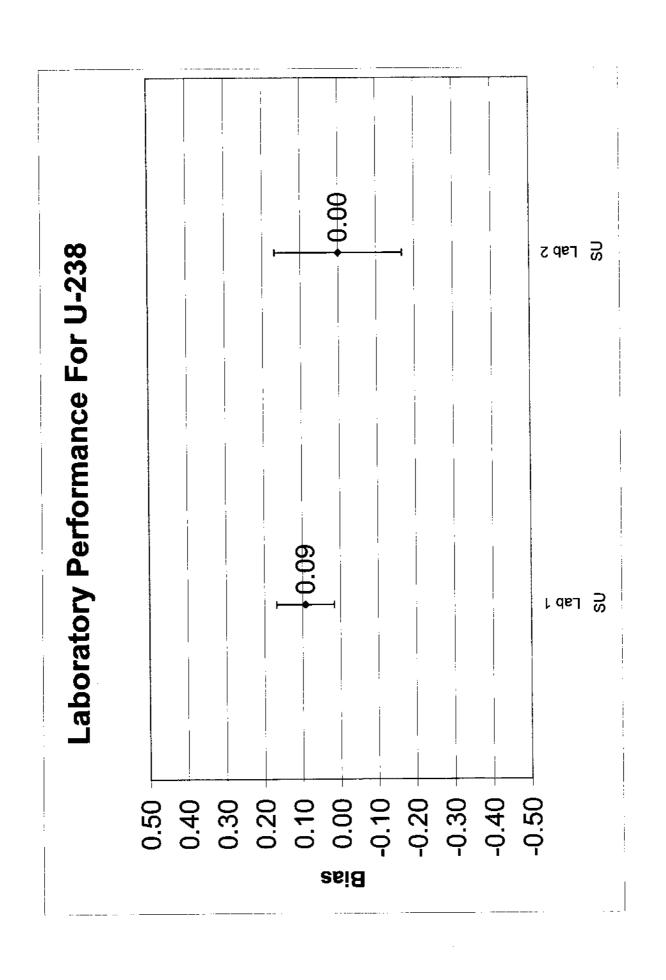
		171181-1		
 Laboratory	r: GEL		Session: 1006	
Matrix:	SF			
RESL LogNo	Radionuclide	Mean Bias (Br):	St. Dev Bias (Sb)	Pass / Fail
DL191	Am-241	-0.162	0.045	Pass
DL191	Co-60	0.089	0.034	Pass
DL191	Cs-137	0.032	0.022	Pass
DL191	Pu-238	-0.085	0.019	Pass
DL191	Pu-239	-0.060	0.022	Pass
DL191	Sr-90	0.028	0.051	Pass
DL191	Th-228	-0.065	0.014	Pass
DL191	Th-230	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

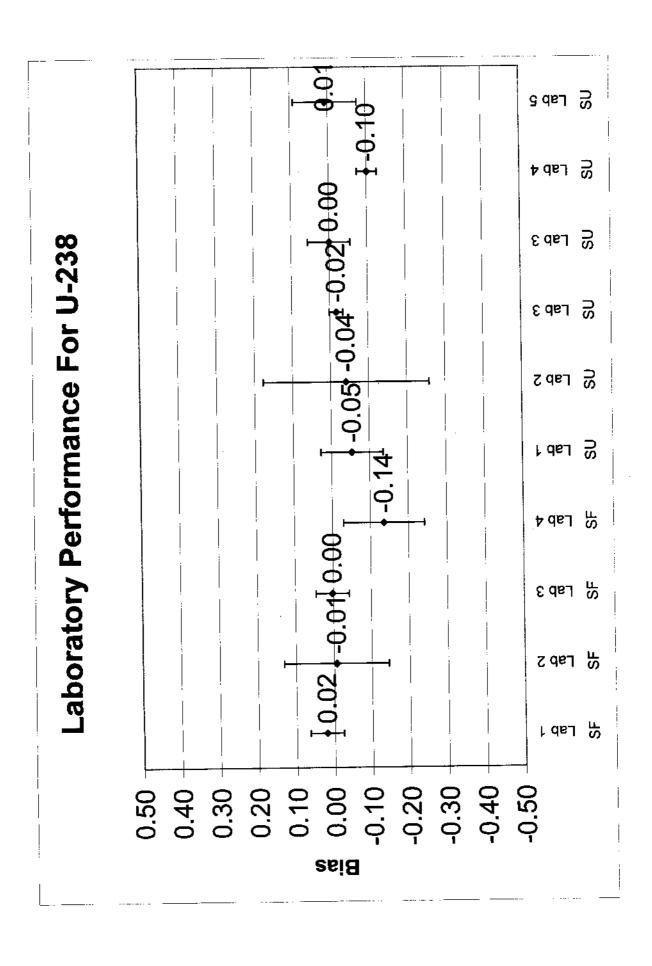
Acceptance Criteria: -0.25 = Sb =< 0.4

# **Laboratory Performance For U-234**









#### DEPARTMENT OF ENERGY LABORATORY ACCREDITATION PROGRAM

#### Session Performance Graphs for Each Radionuclide

Reference Date: 12/1/2006	/2006 Radionuclide: U-Tot	≥: U-Tot	RESL Log# DL188	DL188
Laboratory: GEL		Matrix: SU	Session: 1006	1006
Sample #	Known Values and 1s Uncertainty	Experimental Results and 1s Uncertainty	Bias	
DL1SU1006EL-1	1.24 +/- 0.04 E0 ug/L	1.32 +/- 0.03 E0 ug/L	0.065	
DL1SU1006EL-2	1.24 +/- 0.04 E0 ug/L	1.32 +/- 0.03 E0 ug/L	0.065	
DL1SU1006EL-3	1.24 +/- 0.04 E0 ug/L	1.33 +/- 0.03 E0 ug/L	0.073	
DL1SU1006EL-4	1.24 +/- 0.04 E0 ug/L	1.37 +/- 0.03 E0 ug/L	0.105	
DL1SU1006EL-6	1.24 +/- 0.04 E0 ug/L	1.43 +/- 0.03 E0 ug/L	0.153	
	Mean (Br) = 0.092	St. Dev (Sb) = 0.038		

Sb = < 0.4

-0.25 =< Br <= 0.50

Acceptance Criteria:

Reference Date: 12/1/2006	12/1/2006	Radionuclide: U-238	238	RESL Log # DL180	DL180
Laboratory: GEL	GEL	Matrix: SU	_	Session:	1006
Sample #		Known Values and 1s Uncertainty	Experimental Results and 1s Uncertainty	Bias	
DL1SU1006MR-1		4.08 +/- 0.13 E0 pCi/L	3.9 +/- 0.3 E0 pCi/L	-0.044	
DL1SU1006MR-2		4.08 +/- 0.13 E0 pCi/L	3.6 +/- 0.2 E0 pCi/L	-0.118	
DL1SU1006MR-4		4.08 +/- 0.13 E0 pCi/L	4.0 +/- 0.3 E0 pCi/L	-0.020	
DL1SU1006MR-5		4.08 +/- 0.13 E0 pCi/L	3.8 +/- 0.3 E0 pCi/L	-0.069	
DL1SU1006MR-6		4.08 +/- 0.13 E0 pCi/L	4.0 +/- 0.3 E0 pCi/L	-0.020	
		Mean (Br) = -0.054	St. Dev (Sb) = 0.041		
Acc	Acceptance Criteria:	riteria: -0.25 =< Br <= 0.50	Sb =< 0.4		

RESL Log # DL18	•
Radionuclide: U-234	
Reference Date: 12/1/2006	

Laboratory: GEL	Matrix: SU	: su	Session: 1006
Sample #	Known Values and 1s Uncertainty	Experimental Results and 1s Uncertainty	Bias
DL1SU1006MR-1	3.93 +/- 0.12 E0 pCi/L	4.0 +/- 0.3 E0 pCi/L	0.018
DL1SU1006MR-3	3.93 +/- 0.12 E0 pCi/L	3.8 +/- 0.2 E0 pCi/L	-0.033
DL1SU1006MR-4	3.93 +/- 0.12 E0 pCi/L	4.3 +/- 0.3 E0 pCi/L	0.094
DL1SU1006MR-5	3.93 +/- 0.12 E0 pCi/L	4.3 +/- 0.3 E0 pCi/L	0.094
DL1SU1006MR-6	3.93 +/- 0.12 E0 pCi/L	4.1 +/- 0.3 E0 pCi/L	0.043

Acceptance Criteria: -0

-0.25 =< Br <= 0.50 Sb =< (

St. Dev (Sb) = 0.054

Mean (Br) = 0.043

RESL Log # DL180	Session: 1006
Radionuclide: Th-232	Matrix: SU
Reference Date: 12/1/2006	Laboratory: GEL

Sample #	Known Values and 1s Uncertainty	Experimental Results and 1s Uncertainty	Bias
DL1SU1006MR-2	3.46 +/- 0.11 E0 pCi/L	3.19 +/- 0.13 E0 pCi/L	-0.078
DL1SU1006MR-3	3.46 +/- 0.11 E0 pCi/L	3.23 +/- 0.13 E0 pCi/L	-0.066
DL1SU1006MR-4	3.46 +/- 0.11 E0 pCi/L	3.23 +/- 0.13 E0 pCi/L	-0.066
DL1SU1006MR-5	3.46 +/- 0.11 E0 pCi/L	3.44 +/- 0.13 E0 pCi/L	-0.006
DL1SU1006MR-6	3.46 +/- 0.11 E0 pCi/L	3.25 +/- 0.13 E0 pCi/L	-0.061

Mean (Br) = -0.055

-0.25 =< Br <= 0.50

Acceptance Criteria:

St. Dev (Sb) = 0.028

Sb = < 0.4

Reference Date: 12/1/2006	12/1/2006	Radionuclide: Th-230	Th-230	RESL Log # DL18(	DL 18(
Laboratory: GEL	GEL	Matrix: SU	ns	Session: 1006	1006
Sample #		Known Values and 1s Uncertainty	Experimental Results and 1s Uncertainty	Bias	
DL1SU1006MR-1		1.82 +/- 0.04 E0 pCi/L	2.22 +/- 0.10 E0 pCi/L	0.220	
DL1SU1006MR-2		1.82 +/- 0.04 E0 pCi/L	2.11 +/- 0.10 E0 pCi/L	0.159	
DL1SU1006MR-3		1.82 +/- 0.04 E0 pCi/L	2.22 +/- 0.10 E0 pCi/L	0.220	
DL1SU1006MR-5		1.82 +/- 0.04 E0 pCi/L	2.22 +/- 0.10 E0 pCi/L	0.220	
DL1SU1006MR-6		1.82 +/- 0.04 E0 pCi/L	2.25 +/- 0.10 E0 pCi/L	0.236	

St. Dev (Sb) = 0.030

Mean (Br) = 0.211

Sb = < 0.4

-0.25 =< Br <= 0.50

# DL180	Session: 1006		<b>~</b>		~		*	
RESL Log # DL180	Session	Bias	-0.023	-0.014	-0.003	0.032	-0.014	
Th-228	ns	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	
Radionuclide: Th-228	Matrix: SU	Known Values and 1s Uncertainty	3.46 +/- 0.11 E0 pCi/L					
Date: 12/1/2006	GEL							
Reference Date:	Laboratory: GEL	Sample #	DL1SU1006MR-1	DL1SU1006MR-2	DL1SU1006MR-3	DL1SU1006MR-4	DL1SU1006MR-6	

St. Dev (Sb) = 0.022

Mean (Br) = -0.005

Sb = < 0.4

-0.25 =< Br <= 0.50

Reference Date: 12/1/2006	2/1/2006	Radionuclide: Sr-90	90	RESL Log # DL180	)L180
Laboratory: GEL	EL	Matrix: SU	<b>-</b>	Session: 1006	900
Sample #	Knc and 13	Known Values and 1s Uncertainty	Experimental Results and 1s Uncertainty	Bias	
DL1SU1006MR-1	2.02 +/-	2.02 +/- 0.05 E2 pCi/L	1.99 +/- 0.06 E2 pCi/L	-0.015	
DL1SU1006MR-3	2.02 +/-	2.02 +/- 0.05 E2 pCi/L	2.16 +/- 0.07 E2 pCi/L	0.069	
DL1SU1006MR-4	2.02 +/-	2.02 +/- 0.05 E2 pCi/L	2.13 +/- 0.07 E2 pCi/L	0.054	
DL1SU1006MR-5	2.02 +/-	2.02 +/- 0.05 E2 pCi/L	2.04 +/- 0.06 E2 pCi/L	0.010	
DL1SU1006MR-6	2.02 +/-	2.02 +/- 0.05 E2 pCi/L	2.03 +/- 0.06 E2 pCi/L	0.005	
	Mean	Mean (Br) = 0.025	St. Dev (Sb) = 0.036		
Accel	Acceptance Criteria:	-0.25 =< Br <= 0.50	Sb =< 0.4		

Reference Date: 12/1/2006	12/1/2006	Radionuclide: Pu-239		RESL Log # DL180	DL180
Laboratory: GEL	GEL	Matrix: SU		Session: 1006	1006
Sample #		Known Values Exp and 1s Uncertainty and	Experimental Results and 1s Uncertainty	Bias	
DL1SU1006MR-1		1.64 +/- 0.04 E0 pCi/L 1.54	1.54 +/- 0.10 E0 pCi/L	-0.061	
DL1SU1006MR-2		1.64 +/- 0.04 E0 pCi/L 1.55	1.55 +/- 0.10 E0 pCi/L	-0.055	
DL1SU1006MR-3		1.64 +/- 0.04 E0 pCi/L 1.60	1.60 +/- 0.10 E0 pCi/L	-0.024	
DL1SU1006MR-5		1.64 +/- 0.04 E0 pCi/L 1.59	1.59 +/- 0.10 E0 pCi/L	-0.030	
DL1SU1006MR-6		1.64 +/- 0.04 E0 pCi/L 1.57	1.57 +/- 0.10 E0 pCi/L	-0.043	

St. Dev (Sb) = 0.016

Mean (Br) = -0.043

Sb = < 0.4

-0.25 =< Br <= 0.50

Reference Date: 12/1/2006	12/1/2006	Radionuclide: Pu-238	Pu-238	RESL Log # DL180	DL180
Laboratory: GEL	GEL	Matrix: SU	ns	Session:	1006
Sample #		Known Values and 1s Uncertainty	Experimental Results and 1s Uncertainty	Bias	
DL1SU1006MR-1		3.02 +/- 0.07 E0 pCi/L	2.73 +/- 0.15 E0 pCi/L	-0.096	
DL1SU1006MR-2		3.02 +/- 0.07 E0 pCi/L	2.84 +/- 0.16 E0 pCi/L	-0.060	
DL1SU1006MR-3		3.02 +/- 0.07 E0 pCi/L	2.77 +/- 0.15 E0 pCi/L	-0.083	
DL1SU1006MR-5		3.02 +/- 0.07 E0 pCi/L	2.89 +/- 0.16 E0 pCi/L	-0.043	
DL1SU1006MR-6		3.02 +/- 0.07 E0 pCi/L	2.83 +/- 0.16 E0 pCi/L	-0.063	
		Mean (Br) = -0.069	St. Dev (Sb) = 0.021		

-0.25 =< Br <= 0.50

Reference Date: 12/1/2006	12/1/2006	Radionuclide: Cs-137	\s-137	RESL Log # DL180	DL180
Laboratory:	GEL	Matrix: SU		Session:	1006
Sample #		Known Values and 1s Uncertainty	Experimental Results and 1s Uncertainty	Bias	
DL1SU1006MR-1		1.79 +/- 0.05 E2 pCi/L	1.57 +/- 0.06 E2 pCi/L	-0.123	
DL1SU1006MR-2		1.79 +/- 0.05 E2 pCi/L	1.65 +/- 0.06 E2 pCi/L	-0.078	
DL1SU1006MR-3		1.79 +/- 0.05 E2 pCi/L	1.60 +/- 0.05 E2 pCi/L	-0.106	
DL1SU1006MR-4		1.79 +/- 0.05 E2 pCi/L	1.42 +/- 0.05 E2 pCi/L	-0.207	
DL1SU1006MR-6		1.79 +/- 0.05 E2 pCi/L	1,75 +/- 0.05 E2 pCi/L	-0.022	
		Mean (Br) = -0.107	St. Dev (Sb) = 0.067		

Sb = < 0.4

-0.25 =< Br <= 0.50

Reference Date: 12/1/2006	12/1/2006	Radionuclide: Co-60	Co-60	RESL Log # DL180	DL180
Laboratory: GEL	GEL	Matrix: SU	SU	Session: 1006	1006
Sample #		Known Values and 1s Uncertainty	Experimental Results and 1s Uncertainty	Bias	
DL1SU1006MR-1		1.44 +/- 0.05 E2 pCi/L	1.35 +/- 0.07 E2 pCi/L	-0.063	
DL1SU1006MR-2		1.44 +/- 0.05 E2 pCi/L	1.27 +/- 0.06 E2 pCi/L	-0.118	
DL1SU1006MR-3		1.44 +/- 0.05 E2 pCi/L	1,44 +/- 0.06 E2 pCi/L	0.000	
DL1SU1006MR-4		1.44 +/- 0.05 E2 pCi/L	1.39 +/- 0.06 E2 pCi/L	-0.035	
DL1SU1006MR-5		1.44 +/- 0.05 E2 pCi/L	1.21 +/- 0.06 E2 pCi/L	-0.160	

St. Dev (Sb) = 0.064

Mean (Br) = -0.075

Sb = < 0.4

-0.25 =< Br <= 0.50

Reference Date: 12/1/2006	12/1/2006	Radionuclide: Am-241	Am-241	RESL Log # DL180	DL 180
Laboratory: GEL	GEL	Matrix: SU	ns	Session: 1006	1006
Sample #		Known Values and 1s Uncertainty	Experimental Results and 1s Uncertainty	Bias	
DL1SU1006MR-1		2.23 +/- 0.06 E0 pCi/L	2.06 +/- 0.16 E0 pCi/L	-0.076	
DL1SU1006MR-2		2.23 +/- 0.06 E0 pCi/L	2.08 +/- 0.16 E0 pCi/L	-0.067	
DL1SU1006MR-4		2.23 +/- 0.06 E0 pCi/L	2.10 +/- 0.17 E0 pCi/L	-0.058	
DL1SU1006MR-5		2.23 +/- 0.06 E0 pCi/L	1.93 +/- 0.15 E0 pCi/L	-0.135	
DL1SU1006MR-6		2.23 +/- 0.06 E0 pCi/L	2.13 +/- 0.17 E0 pCi/L	-0.045	

St. Dev (Sb) = 0.035

Mean (Br) = -0.076

RESL Log #
Radionuclide: H-3
Reference Date: 12/1/2006

DL177

Laboratory: GEL	Matri	Matrix: SU	Session: 1006	1006
Sample #	Known Values and 1s Uncertainty	Experimental Results and 1s Uncertainty	Bias	
DL1SU1006LB-1	7.1 +/- 0.2 E4 pCi/L	6.76 +/- 0.10 E4 pCi/L	-0.048	
DL1SU1006LB-2	7.1 +/- 0.2 E4 pCi/L	6.80 +/- 0.10 E4 pCi/L	-0.042	
DL1SU1006LB-3	7.1 +/- 0.2 E4 pCi/L	6.79 +/- 0.10 E4 pCi/L	-0.044	
DL1SU1006LB-5	7.1 +/- 0.2 E4 pCi/l	6 72 +/- 0 10 E4 pCi/l	-0.054	

Acceptance Criteria: -0.25 =< Br <= 0.50

Sb =< 0.4

6.68 +/- 0.10 E4 pCi/L

St. Dev (Sb) = 0.007

Mean (Br) = -0.049

7.1 +/- 0.2 E4 pCi/L

DL1SU1006LB-6

#### **DEPARTMENT OF ENERGY** LABORATORY ACCREDITATION PROGRAM

#### **Summary Report**

Laborato	ory: GEL		Session: 1006	4 6 6
Matrix:	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

Acceptance Criteria: -0.25 = Sb =< 0.4

#### RADIOLOGICAL AND ENVIRONMENTAL SCIENCES LABORATORY

<sup>\*</sup> Mass Determination

RESL Log # DL191	
Radionuclide: Sr-90	
Reference Date: 12/1/2006	

Laboratory: GEL	Matri	Matrix: SF	Session: 1006	1006	
Sample #	Known Values and 1s Uncertainty	Experimental Results and 1s Uncertainty	Bías		
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		

Acceptance Criteria: -0.25 =< Br <= 0.50

St. Dev (Sb) = 0.051

Mean (Br) = 0.028

Sb = < 0.4

DL191	1006						
RESL Log # DL191	Session: 1006	Bías	0.005	0.005	0.054	0.029	0.005
J-238	70	Experimental Results and 1s Uncertainty	4.1 +/- 0.3 E0 pCi	4.1 +/- 0.3 E0 pCi	4.3 +/- 0.3 E0 pCi	4.2 +/- 0.3 E0 pCi	4.1 +/- 0.3 E0 pCi
Radionuclide: U-238	Matrix: SF	Known Values and 1s Uncertainty	4.08 +/- 0.12 E0 pCi				
Date: 12/1/2006	GEL						
Reference Date:	Laboratory: GEL	Sample #	DL1SF1006MR-1	DL1SF1006MR-2	DL1SF1006MR-4	DL1SF1006MR-5	DL1SF1006MR-6

Acceptance Criteria: -0.25 =< Br <= 0.50

.50 Sb =< 0.4

St. Dev (Sb) = 0.022

Mean (Br) = 0.020

RESL Log # DL191	Session: 1006	ults ty Bias
0		Experimental Results and 1s Uncertainty
Radionuclide: Co-60	Matrix: SF	Known Values and 1s Uncertainty
Reference Date: 12/1/2006	Laboratory: GEL	Sample #

Reference Date: 12/1/2006	12/1/2006	Radionuclide: U-234	U-23 <b>4</b>	RESt Log # DL191	มนายา
Laboratory: GEL	GEL	Matrix: SF	RS	Session:	1006
Sample #		Known Values and 1s Uncertainty	Experimental Results and 1s Uncertainty	Bias	
DL1SF1006MR-2		3.93 +/- 0.11 E0 pCi	3.7 +/- 0.3 E0 pCi	-0.059	
DL1SF1006MR-3		3.93 +/- 0.11 E0 pCi	3.9 +/- 0.3 E0 pCi	-0.008	
DL1SF1006MR-4		3,93 +/- 0,11 E0 pCi	4.0 +/- 0.3 E0 pCi	0.018	
DL1SF1006MR-5		3.93 +/- 0.11 E0 pCi	3.9 +/- 0.3 E0 pCi	-0.008	
DL1SF1006MR-6		3.93 +/- 0.11 E0 pCi	3.9 +/- 0.3 E0 pCi	-0.008	

-0.25 = Sr <= 0.50 Sb =< 0.4

Acceptance Criteria:

St. Dev (Sb) = 0.028

Mean (Br) = -0.013

Reference Date: 12/1/2006	12/1/2006	Radionuclide: Th-232	Th-232	RESL Log # DL191	DL 191
Laboratory: GEL	GEL	Matrix: SF	SF	Session: 1006	1006
Sample #		Known Values and 1s Uncertainty	Experimental Results 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	

Acceptance Criteria:

-0.25 =< Br <= 0.50 Sb =< 0.4

St. Dev (Sb) = 0.017

Mean (Br) = -0.100

Reference Date: 12/1/2006	12/1/2006	Radionucl	Radionuclide: Th-230	RESL Log # DL191	DL191
Laboratory: GEL	GEL	Ma	Matrix: SF	Session: 1006	1006
Sample #		Known Values and 1s Uncertainty	Experimental Results and 1s Uncertainty	is 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	690.0	
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		

Sb = < 0.4

-0.25 =< Br <= 0.50

RESL Log # DL191	
Radionuclide: Th-228	
12/1/2006	
Reference Date:	

Laboratory: GEL	Matrix: SF	: SF	Session: 1006	1006	
Sample #	Known Values and 1s Uncertainty	Experimental Results and 1s Uncertainty	Bias		
DL1SF1006MR-2	3.03 +/- 0.10 E0 pCi	2.79 +/- 0.11 E0 pCi	-0.079		
DL1SF1006MR-3	3.03 +/- 0.10 E0 pCi	2.89 +/- 0.12 E0 pCi	-0.046		
DL1SF1006MR-4	3.03 +/- 0.10 E0 pCi	2.86 +/- 0.11 E0 pCi	-0.056		
DL1SF1006MR-5	3.03 +/- 0.10 E0 pCi	2.81 +/- 0.11 E0 pCi	-0.073		
DL1SF1006MR-6	3.03 +/- 0.10 E0 pCi	2.81 +/- 0.11 E0 pCi	-0.073		

Acceptance Criteria: -0.

 $-0.25 \approx 8r \le 0.50$  Sb = < 0.4

St. Dev (Sb) = 0.014

Mean (Br) = -0.065

Reference Date: 12/1/2006	Radionuclide: Pu-239	u-239	RESL Log # DL191	_
Laboratory: GEL	Matrix: SF	<u>ı</u>	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	

Acceptance Criteria: -0.25 =< Br <= 0.50

Sb =< 0.4

St. Dev (Sb) = 0.022

Mean (Br) = -0.060

Reference Date: 12/1/2006	12/1/2006	Radionuclide: Pu-238	-238	RESL Log # DL191	DL 191
Laboratory: GEL	GEL	Matrix: SF		Session:	1006
Sample #		Known Values and 1s Uncertainty	Experimental Results and 1s Uncertainty	Bias	
DL1SF1006MR-1		3.02 +/- 0.07 E0 pCi	2.77 +/- 0.16 E0 pCi	-0.083	
DL1SF1006MR-2		3.02 +/- 0.07 E0 pCi	2.77 +/- 0.17 E0 pCi	-0.083	
DL1SF1006MR-3		3.02 +/- 0.07 E0 pCi	2.83 +/- 0.17 E0 pCi	-0.063	
DL1SF1006MR-4		3.02 +/- 0.07 E0 pCi	2.77 +/- 0.17 E0 pCi	-0.083	
DL1SF1006MR-5		3.02 +/- 0.07 E0 pCi	2.67 +/- 0.16 E0 pCi	-0.116	
		Mean (Br) = -0.085	St. Dev (Sb) = 0.019		

-0.25 =< Br <= 0.50

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

RESL Log # DL191	
Radionuclide: Am-241	
Reference Date: 12/1/2006	

Laboratory: GEL	Matri	Matrix: SF	Session: 1006	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		

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

Mean (Br) = -0.162

St. Dev (Sb) = 0.045

# Laboratory Performance For Th-232

