

The IAEA-TEL-2016-03 world wide open proficiency test on the determination of anthropogenic and natural radionuclides in water and biota samples

Laboratory's Individual Evaluation Report Laboratory Code: 99 (CuNo: 13949) Total Pages (with cover): 7



Individual Evaluation Report

for the World-Wide Open Proficiency Test IAEA-TEL-2016-03

Individual Evaluation Report

for Laboratory Nr. 99

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http://nucleus.iaea.org/rpst/

Description of the evaluation criterias:

The data is evaluated according to the following steps:

The relative bias between the reported and the target value (the best estimation of the true value) is expressed by the following equation:

$$Bias_{relative} = \frac{Value_{reported} - Value_{target}}{Value_{target}} \times 100\%$$

The relative bias will be compared to the Maximum Acceptable Relative Bias (**MARB**) which has been determined for each measurand, considering the physical background of radioanalytical methods, including the level of the radioactivity and the complexity of the task.

If the **Bias**_{relative} ≤ **MARB** value the result will be "Accepted" for accuracy.

Based on fit for purpose and the good laboratory practice principles the expanded relative combined uncertainty should cover the relative bias:

$$P = \sqrt[2]{\left(\frac{u_{target}}{A_{target}}\right)^2 + \left(\frac{u_{reported}}{A_{reported}}\right)^2} \times 100$$

$$Bias_{relative} \leq k * P$$

where k is the coverage factor, for the 95% confidential level k is 2.56. If the reported result is between the \pm MARB values, but it is not overlapping with the target value within their uncertainties, this equation helps to decide whether they are significantly different or not.

The **P** value will be compared to the **MARB** also. If both the **P** \leq **MARB** and **Bias**_{relative} \leq **k*****P** are fulfilled the reported result will be "Accepted" for the precision. If one of them is insufficient the result will be assigned the "Not accepted" status for precision.

The final score according to the above detailed evaluation:

"Accepted" when both accuracy and precision achieved "Accepted" status,

"Not Accepted" when the accuracy is "Not accepted" and

"Warning" when accuracy is "Accepted", but the precision is "Not accepted".

Target Value Tables for the proficiency test parameters:

Sample Code	Analyte	Target Value	Uncertainty	MARB
1	Na-22	53.2	1.5	15 %
1	Sr-90	14.7	0.5	15 %
1	Cs-134	19.9	0.6	15 %
1	Cs-137	39.6	1	15 %

Table of Target Values and Evaluation Parameters (MARB) for Sample 1

Table of Target Values and Evaluation Parameters (MARB) for Sample 2

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Sample Code	Analyte	Target Value	Uncertainty	MARB
2	Sr-89	373	15	30 %
2	Sr-90	20.5	0.5	20 %
2	Am-241	26.7	0.7	15 %

#### Table of Target Values and Evaluation Parameters (MARB) for Sample 4

Sample Code	Analyte	Target Value	Uncertainty	MARB
4	K-40	216	13	25 %
4	Sr-90	17	2	30 %
4	Cs-137	209	11	20 %

# Evaluation Tables for Labcode 99.

Sample Code	Analyte	Target Value	Target Unc.	MARB	Rep. Value	Rep. Unc	Rel. Bias	Robust SD	Z-Score	U-Test	Accuracy	Р	Precision	Final Score
1	Cs-134	19.9	0.6	15 %	22.8	1.1	14.57 %	1	2.90	2.31	А	5.69	N	W
1	Cs-137	39.6	1	15 %	38.0	1.6	-4.04 %	1.5	-1.07	-0.85	А	4.91	А	А
1	Na-22	53.2	1.5	15 %	51.4	3.2	-3.38 %	3.5	-0.51	-0.51	А	6.83	А	А

### Evaluation Result Table for Sample 1

### Evaluation Result Table for Sample 2

Sample Code	Analyte	Target Value	Target Unc.	MARB	Rep. Value	Rep. Unc	Rel. Bias	Robust SD	Z-Score	U-Test	Accuracy	Р	Precision	Final Score
2	Am-241	26.7	0.7	15 %	26.7	2.1	0.00 %	1.9	0.00	0.00	А	8.29	А	А

### Evaluation Result Table for Sample 4

Sample Code	Analyte	Target Value	Target Unc.	MARB	Rep. Value	Rep. Unc	Rel. Bias	Robust SD	Z-Score	U-Test	Accuracy	Р	Precision	Final Score
4	Cs-137	209	11	20 %	205	16	-1.91 %	17	-0.24	-0.21	А	9.41	А	А

Intercomparison Parameter Evaluation:

Calculation of Evaluation Parameters

robust average: x*=median of xi (i=1,2..p), robust standart deviation s*=1.483 * median of |xi-x*|, z-score: z=(Reported-Value - TargetValue)/s* z-score evaluation: z<2: accepted, 2<=z<=3: warning, z>3: not accepted

The analytes listed in the table below have been identified but are not present in the samples (false positive):

Sample Code	Analyte	Reported Value