



FIRMS - Forensic Isotope Ratio Mass Spectrometry PT Scheme

Round: 228

Issue Number 1

Issued 12 June 2015



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Scheme Information

Aims of Scheme

The primary aim of the Forensic Isotope Ratio Mass Spectrometry Proficiency Testing Scheme (FIRMS) is to enable laboratories performing isotope ratio analysis of a range of test materials to monitor their performance and compare it with that of their peers. The FIRMS scheme also aims to provide information to participants on technical issues and methodologies relating to isotope ratio analysis.

Further information on the scheme organisation, the test materials, and the statistical analysis of data are available in the FIRMS Scheme Description and the LGC PT General Protocol.

Performance Assessment

Once a PT round has closed, the results will be analysed and the assigned value determined for each analyte, according to the criteria provided in the Scheme Description. Information regarding the traceability of each calculated assigned value is also provided in the Scheme Description.

For quantitative data, the participant's result, x , (or $\log_{10} x$ for microbiological data) is converted into a z score using the following formula;

$$z = \frac{(x - X)}{\text{SDPA}}$$

X = Assigned value

SDPA = Standard deviation for proficiency assessment

For quantitative data, the uncertainty of the assigned value is calculated to ensure that it would have a negligible effect on participants' performance scores. If the uncertainty of the assigned value is greater than $0.3 \times \text{SDPA}$, then this is not considered negligible. In this situation, a z' (z prime) performance score is automatically calculated rather than a z score, in order to take account of the measurement uncertainty of the assigned value. The z' score is calculated using the following formula;

$$z' = \frac{(x - X)}{\sqrt{(\text{SDPA}^2 + U_x^2)}}$$

X = Assigned value

SDPA = Standard deviation for proficiency assessment

U_x = Uncertainty of the assigned value

Expanded SDPA = $\sqrt{(\text{SDPA}^2 + U_x^2)}$

Trend graphs will use a mixture of z and z' scores, i.e. the 'performance score' for the round.

For quantitative data, gross errors or blunders are removed from the data by removal of any results that are greater than the assigned value $\pm 5 \times \text{SDPA}$. These results are not used in the final calculation of the assigned value and other summary statistics and will be included in the number of 'Excluded Results'. All results, including excluded results, will be given a performance score.

For the purposes of performance assessment for a single round, z and z' scores are interpreted as follows:

z/z' score	Interpretation	Colour coding
$ z \leq 2.00$	Satisfactory result	Green
$2.00 < z \text{ and } < 3.00$	Questionable result	Amber
$ z \geq 3.00$	Unsatisfactory result	Red
No score given	See below	No colour coding

Performance scores will not be given for the following:

- For qualitative results, where satisfactory performance is based on the participants reporting the same result as the assigned result. e.g. detected, not detected. For these results, colour coding of green (satisfactory) or red (unsatisfactory) will apply.
- For results of zero; such a result is not normally appropriate and should not be reported, the result should be reported as less than the detection limit rather than zero.
Note: for a very small number of analytes it may be appropriate to report a result of zero, depending on the type of measurement scale being used.
- For quantitative results where the analyte under test is present in the test material but participants report non-numerical results e.g. 0, <1 , >300 . In these cases, it is not possible to allocate a performance score and participants should assess their performance based on the assigned value and satisfactory range given.
- For quantitative results, for microbiological test materials, where the analyte under test is not present in the test material, the assigned value will be classified as 'Absent'. Results reported as 'less than' at or below the detection level for our method of confirmation will be assessed as satisfactory (green colour code). Results reported at a higher detection level will not be assessed and participants will need to use their own judgement to determine whether their result is fit for its intended use. Results reporting a positive count will be assessed as unsatisfactory (red colour code).

In some cases, performance scores may not be provided or may be provided but with colour coding suspended (indicating that scores need to be interpreted with caution). For example:

- For small data sets where less than 8 results have been submitted and the assigned value is derived using a consensus value from the participants' results. In these circumstances, there may be increased uncertainty of the assigned value, given the low number of participants, and performance scores will be given for information only.
- In cases where the distribution of the results gives cause for concern e.g. bi-modal data sets. These circumstances will be dependent on the statistical design that is in place.

- If the assigned value falls below a concentration threshold (only applies to some schemes).

In these or similar circumstances, further explanation as to the reasons for suspension of performance scoring or colour coding, and on the interpretation of results, will be given in the report.

Note: Data displayed in the report will have been rounded to the required number of decimal places. However statistical calculations will have been performed on unrounded data. For this reason, there may appear to be differences between displayed data and calculated data, but this does not affect results in any way.

Confidentiality

A unique laboratory reference code is used to report results in order to ensure confidentiality.

Contact details

The Technical Scheme Coordinator is Matthew Whetton

Please contact ptcustomerservices@lqcgroupp.com if you have any questions or comments regarding the scheme.

Authorisation

This report was authorised by Matthew Whetton, Head of Chemistry on the 12 June 2015



Sample Details

Samples were despatched on 27 April 2015.
The reporting deadline was 05 June 2015.

The following samples were distributed in FIRMS Round 228:

- 1: 1 x 0.5g Flour for the determination of delta 2H, 13C and 18O
2: 1 x 0.5g Caffeine for the determination of delta 2H, 13C, 15N and 18O

Further information regarding assigned values, performance assessment and technical comments can be found under the individual sample and analyte results.

Calculated within and between participant standard deviations

Sample 1 (Flour)

Analyte	Within participant SD	Between participant SD
Delta 2H	1.7173	2.1051
Delta 13C	0.1121	0.0461
Delta 18O	0.2247	0.1437

Sample 2 (Caffeine)

Analyte	Within participant SD	Between participant SD
Delta 2H	0.7278	2.2977
Delta 13C	0.0619	0.0269
Delta 15N	0.0938	0.1143
Delta 18O	0.4805	0.2282

Quality Control

All homogeneity assessments have been conducted in accordance with the principles stipulated in ISO 13528^[1] and/or the IUPAC^[2] Harmonized PT Protocol. Further details regarding the assessment of homogeneity can be found in the LGC Standards Proficiency Testing General Protocol.

Sample	Analyte/Test	Result (SD)	Assessment
1 (flour)	delta 13C	-11.35 (0.043)*	Pass
2 (caffeine)	delta 13C	-35.04 (0.029)*	Pass

*Results were scaled to the NBS19-LSVEC scale.

Analysis carried out for the purposes of homogeneity testing was sub-contracted by LGC to an external laboratory.

For quantitative testing in this round, a comparison of the standard deviation of the homogeneity results returned and the SDPA expected for the participant assessment was carried out. The samples were considered to be sufficiently homogeneous for use in the PT scheme, based on the values returned.

For qualitative testing, the target analyte must be detected in 100% of test materials analysed.

For any analyte which has not been proven to be sufficiently homogeneous, and any closely related analytes, the value set for the SDPA may be suspended in order to take account of any potential inhomogeneity. The actual value used for the standard deviation for proficiency assessment is shown at the foot of the results and z-score tables in this report.

Often a particular test material does not require homogeneity assessment prior to distribution. Such sample types include standard solutions and aqueous solutions.

[1] ISO 13528 (2005), 'Statistical methods for use in proficiency testing by inter-laboratory comparisons'.

[2] M Thompson, S L R Ellison, R Wood, 'International Harmonised Protocol for the Proficiency Testing of Analytical Chemistry Laboratories', *Pure Appl. Chem.*, 2006, 78, 145-196.

Sample: 01 - FIRMS sample 1**Analyte: delta 2H (VSMOW)**

Lab ID	Method	Result	z score
FM0002	Isotope Ratio Mass Spectrometry	-5.53	
FM0006	Isotope Ratio Mass Spectrometry	-35.15	
FM0007	Isotope Ratio Mass Spectrometry	4.30	
FM0010	Isotope Ratio Mass Spectrometry	-14.26	
FM0014	Isotope Ratio Mass Spectrometry	5.87	
FM0016	Isotope Ratio Mass Spectrometry	1.18	
FM0018	Isotope Ratio Mass Spectrometry	5.01	
FM0027	Isotope Ratio Mass Spectrometry	3.21	
FM0028	Isotope Ratio Mass Spectrometry	6.30	
FM0029	Isotope Ratio Mass Spectrometry	31.60	
FM0034	Isotope Ratio Mass Spectrometry	-2.45	

Data Statistics

	Value
Number of Results	11
Number of Excluded Results	0
Mean	N/A
Median	N/A
Standard Deviation	N/A
Robust Standard Deviation	N/A
Result Range	-35.15 to 31.60

Comments

The participants in the FIRMS scheme were allowed to report up to 10 results, for the purposes of calculating individual and group summary statistics, plus a mean result, which was used to calculate the summary statistics subsequently applied in the assessment of performance.

In this round most participants obtained satisfactory performance scores for most parameters. Performance scores are currently calculated using the robust standard deviation obtained from the data reported for delta 2H and delta 18O; however a fixed SDPA has been implemented for delta 13C and delta 15N.

Generally the data reported was good for most analytes; however a wide range of values from -35.15 to 31.60 were reported for delta 2H for sample 1. As there is little comparability an assigned value cannot be provided and so performance scores have been removed and the results have been included in the report for information purposes only.

In cases where the results returned are more varied, the robust standard deviation will also be larger and so a wider range of results will therefore be assessed as satisfactory for delta 2H and delta 18O. Participants are advised to consider the magnitude of the robust standard deviations used, and the absolute difference of their result from the assigned value, when interpreting their performance.

Sample: 01 - FIRMS sample 1

Analyte: delta 13C (VPDB)

Lab ID	Method	Result	z score
FM0002	Isotope Ratio Mass Spectrometry	-11.38	0.27
FM0003	Isotope Ratio Mass Spectrometry	-11.42	0.00
FM0006	Isotope Ratio Mass Spectrometry	-11.42	0.00
FM0007	Isotope Ratio Mass Spectrometry	-10.88	3.60
FM0010	Isotope Ratio Mass Spectrometry	-11.47	-0.33
FM0011	Isotope Ratio Mass Spectrometry	-11.52	-0.67
FM0014	Isotope Ratio Mass Spectrometry	-11.45	-0.20
FM0016	Isotope Ratio Mass Spectrometry	-11.22	1.33
FM0018	Isotope Ratio Mass Spectrometry	-11.07	2.33
FM0023	Isotope Ratio Mass Spectrometry	-11.53	-0.73
FM0027	Isotope Ratio Mass Spectrometry	-11.45	-0.20
FM0028	Isotope Ratio Mass Spectrometry	-11.59	-1.13
FM0029	Isotope Ratio Mass Spectrometry	-11.30	0.80
FM0033	Isotope Ratio Mass Spectrometry	-11.48	-0.40
FM0034	Isotope Ratio Mass Spectrometry	-11.36	0.40
FM0035	Isotope Ratio Mass Spectrometry	-11.33	0.60
FM0036	Isotope Ratio Mass Spectrometry	-11.39	0.20

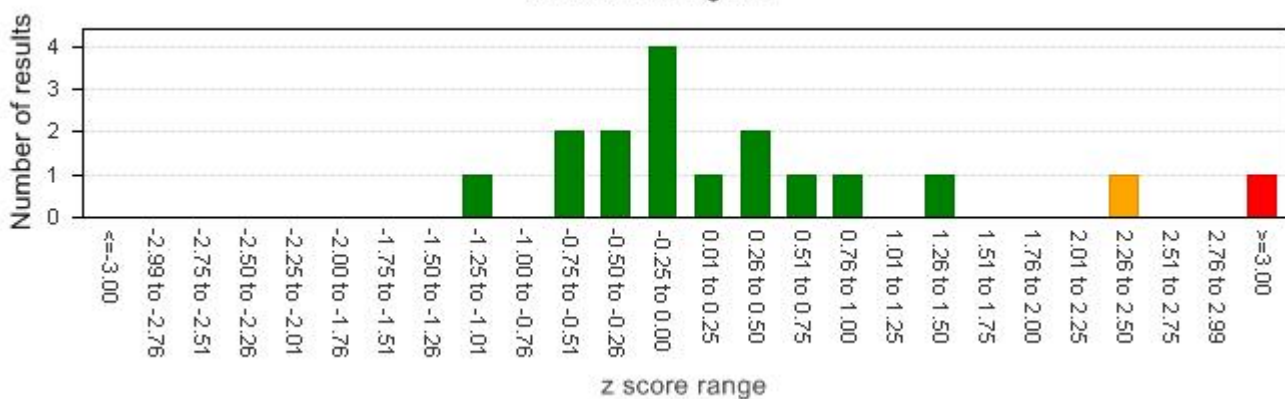
Data Statistics

	Value
Number of Results	17
Number of Excluded Results	0
Mean	-11.37
Median	-11.42
Standard Deviation	0.176
Robust Standard Deviation	0.089
Result Range	-11.59 to -10.88

Performance Statistics

	Value
Assigned Value	-11.42
Uncertainty of Assigned Value	0.03
SDPA	0.15
Satisfactory Range	-11.72 to -11.12
Satisfactory z scores	88.2%
Questionable z scores	5.9%
Unsatisfactory z scores	5.9%

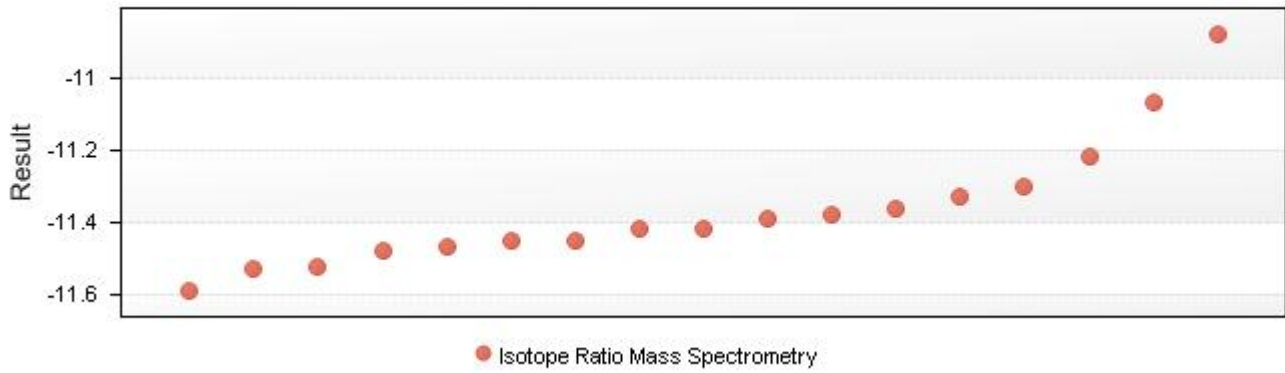
z score Histogram



Sample: 01 - FIRMS sample 1

Analyte: delta 13C (VPDB)

Distribution Graph



Methodology Summary

Method	Number of Results	Excluded Results	% of Total	Median	Robust SD	Range	Sat.
							%
Isotope Ratio Mass Spectrometry	17	0	100	-11.42	0.089	-11.59 to -10.88	88.2
All	17	0	100	-11.42	0.089	-11.59 to -10.88	88.2

Comments

Sample: 01 - FIRMS sample 1

Analyte: delta 18O (VSMOW)

Lab ID	Method	Result	z' score*
FM0002	Isotope Ratio Mass Spectrometry	31.17	0.00
FM0006	Isotope Ratio Mass Spectrometry	29.92	-3.24
FM0007	Isotope Ratio Mass Spectrometry	32.75	4.09
FM0011	Isotope Ratio Mass Spectrometry	30.93	-0.62
FM0016	Isotope Ratio Mass Spectrometry	31.22	0.13
FM0018	Isotope Ratio Mass Spectrometry	31.54	0.96
FM0027	Isotope Ratio Mass Spectrometry	31.03	-0.36
FM0029	Isotope Ratio Mass Spectrometry	31.30	0.34
FM0033	Isotope Ratio Mass Spectrometry	26.10	-13.12
FM0034	Isotope Ratio Mass Spectrometry	29.18	-5.15

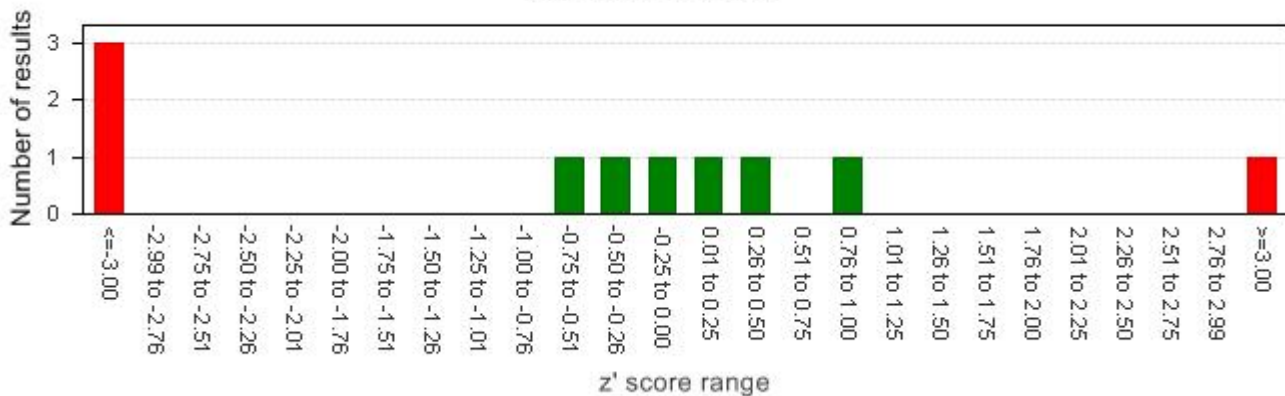
Data Statistics

	Value
Number of Results	10
Number of Excluded Results	1
Mean	31.00
Median	31.17
Standard Deviation	1.000
Robust Standard Deviation	0.356
Result Range	29.18 to 32.75

Performance Statistics

	Value
Assigned Value	31.17
Uncertainty of Assigned Value	0.15
SDPA	0.356
Expanded SDPA	0.386
Satisfactory Range	30.40 to 31.94
Satisfactory z' scores	60.0%
Questionable z' scores	0.0%
Unsatisfactory z' scores	40.0%

z' score Histogram



Methodology Summary

Method	Number of Results	Excluded Results	% of Total	Median	Robust SD	Range	Sat.
							%
Isotope Ratio Mass Spectrometry	10	1	100	31.17	0.356	29.18 to 32.75	60.0
All	10	1	100	31.17	0.356	29.18 to 32.75	60.0

Comments

*Please note, participant performance for this analyte has been assessed using a z' score, rather than a z score, in order to account for the measurement uncertainty of the assigned value which is not negligible when compared to the SDPA.

Sample: 02 - FIRMS sample 2

Analyte: delta 2H (VSMOW)

Lab ID	Method	Result	z' score*
FM0002	Isotope Ratio Mass Spectrometry	-134.81	0.81
FM0006	Isotope Ratio Mass Spectrometry	-158.57	-0.90
FM0007	Isotope Ratio Mass Spectrometry	-149.20	-0.23
FM0010	Isotope Ratio Mass Spectrometry	-162.41	-1.18
FM0011	Isotope Ratio Mass Spectrometry	-134.42	0.84
FM0014	Isotope Ratio Mass Spectrometry	-153.29	-0.52
FM0016	Isotope Ratio Mass Spectrometry	-149.06	-0.22
FM0018	Isotope Ratio Mass Spectrometry	-134.76	0.82
FM0027	Isotope Ratio Mass Spectrometry	-135.71	0.75
FM0028	Isotope Ratio Mass Spectrometry	-151.80	-0.41
FM0029	Isotope Ratio Mass Spectrometry	-143.10	0.22
FM0034	Isotope Ratio Mass Spectrometry	-140.19	0.43

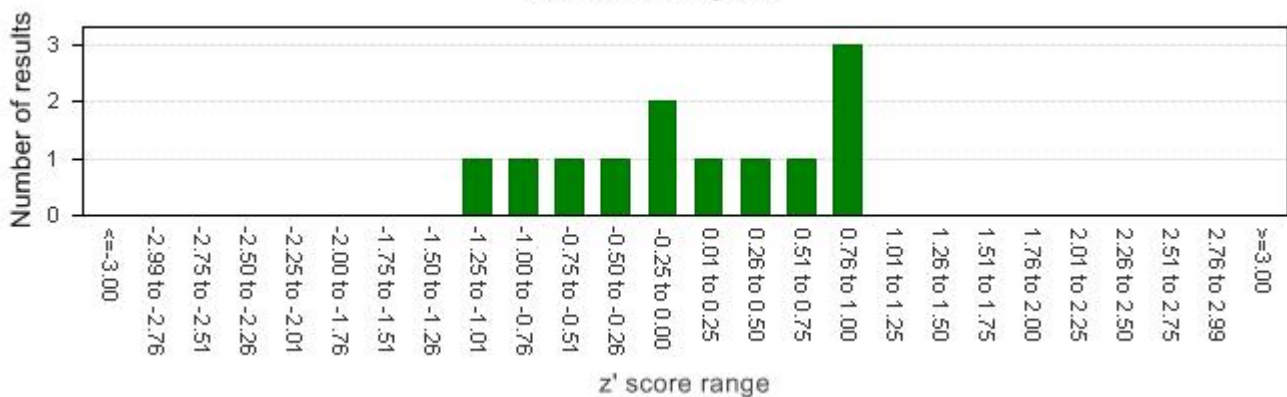
Data Statistics

	Value
Number of Results	12
Number of Excluded Results	0
Mean	-145.61
Median	-146.08
Standard Deviation	9.842
Robust Standard Deviation	13.036
Result Range	-162.41 to -134.42

Performance Statistics

	Value
Assigned Value	-146.08
Uncertainty of Assigned Value	4.70
SDPA	13.036
Expanded SDPA	13.857
Satisfactory Range	-173.79 to -118.37
Satisfactory z' scores	100.0%
Questionable z' scores	0.0%
Unsatisfactory z' scores	0.0%

z' score Histogram



Methodology Summary

Method	Number of Results	Excluded Results	% of Total	Median	Robust SD	Range	Sat.
							%
Isotope Ratio Mass Spectrometry	12	0	100	-146.08	13.036	-162.41 to -134.42	100.0
All	12	0	100	-146.08	13.036	-162.41 to -134.42	100.0

Comments

*Please note, participant performance for this analyte has been assessed using a z' score, rather than a z score, in order to account for the measurement uncertainty of the assigned value which is not negligible when compared to the SDPA.

Sample: 02 - FIRMS sample 2

Analyte: delta 13C (VPDB)

Lab ID	Method	Result	z score
FM0002	Isotope Ratio Mass Spectrometry	-38.41	-0.53
FM0003	Isotope Ratio Mass Spectrometry	-38.16	1.13
FM0006	Isotope Ratio Mass Spectrometry	-38.33	0.00
FM0007	Isotope Ratio Mass Spectrometry	-38.35	-0.13
FM0010	Isotope Ratio Mass Spectrometry	-38.37	-0.27
FM0011	Isotope Ratio Mass Spectrometry	-38.43	-0.67
FM0014	Isotope Ratio Mass Spectrometry	-38.31	0.13
FM0016	Isotope Ratio Mass Spectrometry	-38.30	0.20
FM0018	Isotope Ratio Mass Spectrometry	-38.41	-0.53
FM0023	Isotope Ratio Mass Spectrometry	-38.07	1.73
FM0027	Isotope Ratio Mass Spectrometry	-38.33	0.00
FM0028	Isotope Ratio Mass Spectrometry	-38.33	0.00
FM0029	Isotope Ratio Mass Spectrometry	-38.10	1.53
FM0033	Isotope Ratio Mass Spectrometry	-38.27	0.40
FM0034	Isotope Ratio Mass Spectrometry	-38.41	-0.53
FM0035	Isotope Ratio Mass Spectrometry	-38.45	-0.80
FM0036	Isotope Ratio Mass Spectrometry	-38.34	-0.07

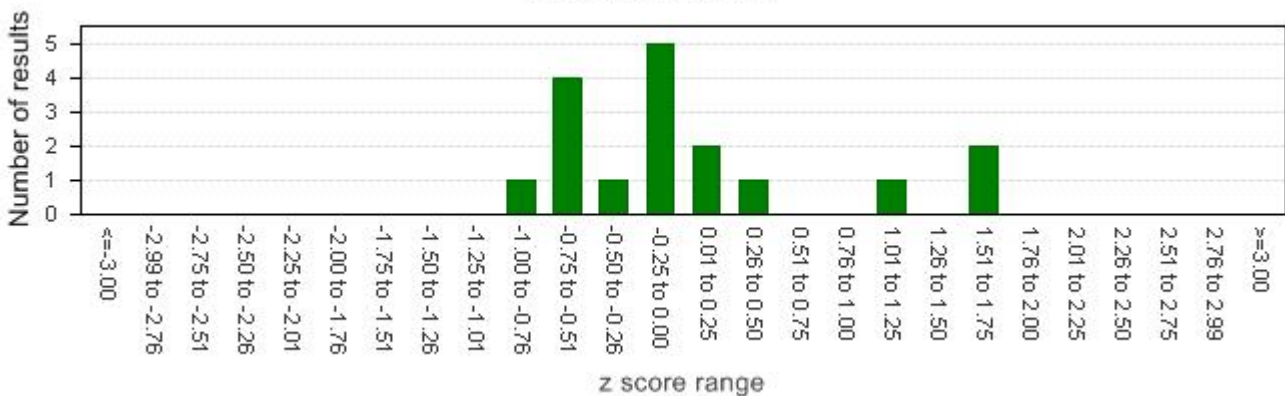
Data Statistics

	Value
Number of Results	17
Number of Excluded Results	0
Mean	-38.32
Median	-38.33
Standard Deviation	0.111
Robust Standard Deviation	0.089
Result Range	-38.45 to -38.07

Performance Statistics

	Value
Assigned Value	-38.33
Uncertainty of Assigned Value	0.03
SDPA	0.15
Satisfactory Range	-38.63 to -38.03
Satisfactory z scores	100.0%
Questionable z scores	0.0%
Unsatisfactory z scores	0.0%

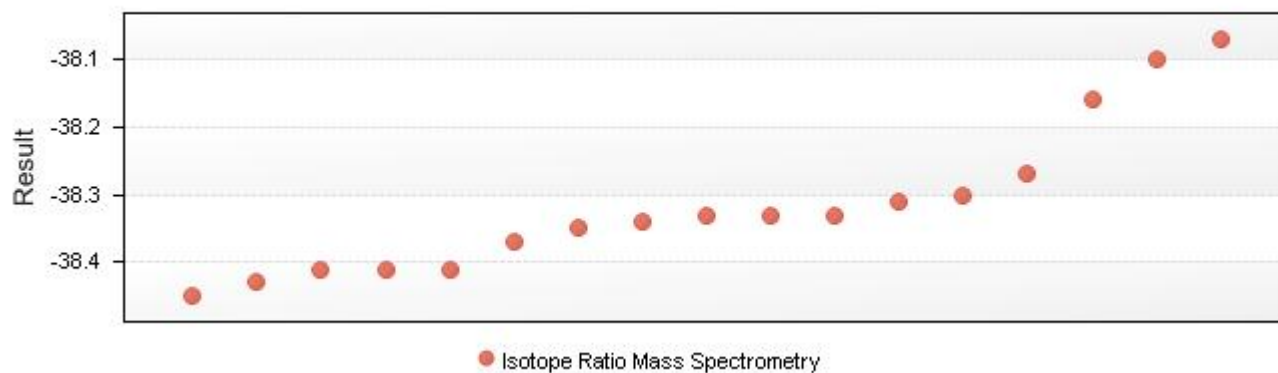
z score Histogram



Sample: 02 - FIRMS sample 2

Analyte: delta 13C (VPDB)

Distribution Graph



Methodology Summary

Method	Number of Results	Excluded Results	% of Total	Median	Robust SD	Range	Sat.
							%
Isotope Ratio Mass Spectrometry	17	0	100	-38.33	0.089	-38.45 to -38.07	100.0
All	17	0	100	-38.33	0.089	-38.45 to -38.07	100.0

Comments

Sample: 02 - FIRMS sample 2

Analyte: delta 15N (AIR)

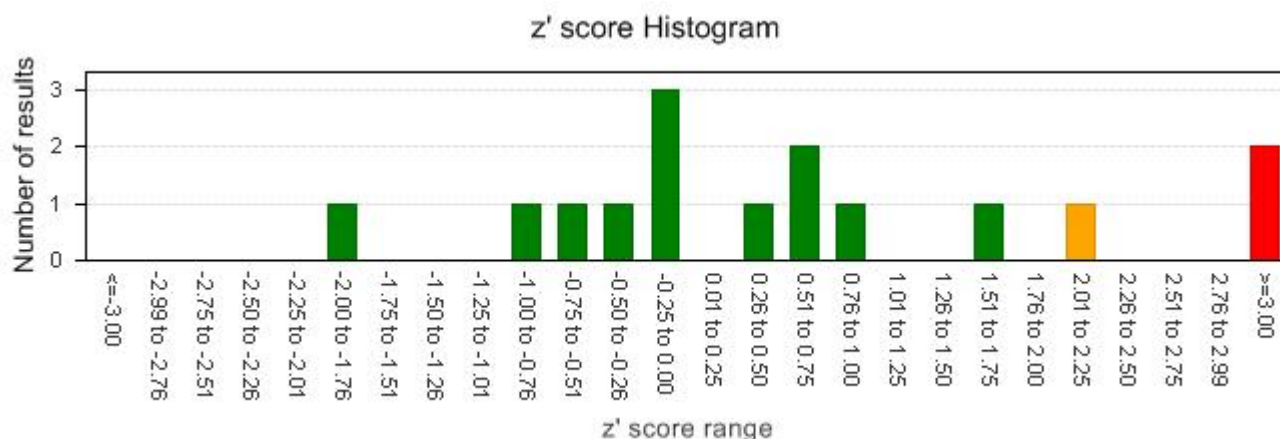
Lab ID	Method	Result	z' score*
FM0002	Isotope Ratio Mass Spectrometry	-8.17	-0.12
FM0006	Isotope Ratio Mass Spectrometry	-8.22	-0.43
FM0007	Isotope Ratio Mass Spectrometry	-8.25	-0.62
FM0011	Isotope Ratio Mass Spectrometry	-8.03	0.74
FM0014	Isotope Ratio Mass Spectrometry	-8.17	-0.12
FM0016	Isotope Ratio Mass Spectrometry	-6.62	9.47
FM0018	Isotope Ratio Mass Spectrometry	-8.29	-0.87
FM0023	Isotope Ratio Mass Spectrometry	-6.66	9.22
FM0027	Isotope Ratio Mass Spectrometry	-8.07	0.50
FM0028	Isotope Ratio Mass Spectrometry	-8.47	-1.98
FM0029	Isotope Ratio Mass Spectrometry	-8.00	0.93
FM0033	Isotope Ratio Mass Spectrometry	-7.88	1.67
FM0034	Isotope Ratio Mass Spectrometry	-7.82	2.04
FM0035	Isotope Ratio Mass Spectrometry	-8.03	0.74
FM0036	Isotope Ratio Mass Spectrometry	-8.15	0.00

Data Statistics

	Value
Number of Results	15
Number of Excluded Results	2
Mean	-8.12
Median	-8.15
Standard Deviation	0.174
Robust Standard Deviation	0.178
Result Range	-8.47 to -7.82

Performance Statistics

	Value
Assigned Value	-8.15
Uncertainty of Assigned Value	0.06
SDPA	0.15
Expanded SDPA	0.162
Satisfactory Range	-8.47 to -7.83
Satisfactory z' scores	80.0%
Questionable z' scores	6.7%
Unsatisfactory z' scores	13.3%



Methodology Summary

Method	Number of Results	Excluded Results	% of Total	Median	Robust SD	Range	Sat.
							%
Isotope Ratio Mass Spectrometry	15	2	100	-8.15	0.178	-8.47 to -7.82	80.0
All	15	2	100	-8.15	0.178	-8.47 to -7.82	80.0

Comments

*Please note, participant performance for this analyte has been assessed using a z' score, rather than a z score, in order to account for the measurement uncertainty of the assigned value which is not negligible when compared to the SDPA.

Sample: 02 - FIRMS sample 2

Analyte: delta 18O (VSMOW)

Lab ID	Method	Result	z' score*
FM0002	Isotope Ratio Mass Spectrometry	17.36	0.72
FM0006	Isotope Ratio Mass Spectrometry	19.96	5.37
FM0007	Isotope Ratio Mass Spectrometry	18.47	2.70
FM0016	Isotope Ratio Mass Spectrometry	16.93	-0.05
FM0018	Isotope Ratio Mass Spectrometry	15.88	-1.93
FM0027	Isotope Ratio Mass Spectrometry	16.96	0.00
FM0029	Isotope Ratio Mass Spectrometry	17.30	0.61
FM0033	Isotope Ratio Mass Spectrometry	16.64	-0.57
FM0034	Isotope Ratio Mass Spectrometry	12.87	-7.33

Due to the low number of results returned, performance scores are shown for information purposes only

Data Statistics

	Value
Number of Results	9
Number of Excluded Results	2
Mean	17.08
Median	16.96
Standard Deviation	0.788
Robust Standard Deviation	0.504
Result Range	15.88 to 18.47

Performance Statistics

	Value
Assigned Value	16.96
Uncertainty of Assigned Value	0.24
SDPA	0.504
Expanded SDPA	0.558
Satisfactory Range	15.84 to 18.08
Satisfactory z' scores	66.7%
Questionable z' scores	11.1%
Unsatisfactory z' scores	22.2%

Methodology Summary

Method	Number of Results	Excluded Results	% of Total	Median	Robust SD	Range	Sat.
							%
Isotope Ratio Mass Spectrometry	9	2	100	16.96	0.504	15.88 to 18.47	66.7
All	9	2	100	16.96	0.504	15.88 to 18.47	66.7

Comments

*Please note, participant performance for this analyte has been assessed using a z' score, rather than a z score, in order to account for the measurement uncertainty of the assigned value which is not negligible when compared to the SDPA.