Test Group Summary Report C39 Inorganics by TCLP January 2025 PT Round

Issued: February 28, 2025



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1.0 The Proficiency Testing Report

The Proficiency Testing Report consists of two parts.

- PTC Proficiency Testing Report: This report contains participant-specific data and other confidential information. This report is emailed to participants at the end of the PT round.
- Test Group Summary Report: A Test Group Summary Report is created for each quantified test group at the end of the PT round. These reports contain more detailed information on the round than is found in the participant-specific PTC Proficiency Testing Report. These reports do not contain any confidential information and are made available on the PTC web site.

2.0 Definitions

The participant-specific PTC Proficiency Testing Report contains some terms that new participants may not be familiar with.

Code: The registration code that is unique to each analyte that a participant is registered for.

App: If a participant is accredited by CALA, this three-digit number is the appendix number that

the accredited method is assigned to.

N: The number of participants results that were used to calculate the summary statistics. This

excludes qualified data (e.g., <) and any results that were flagged as outliers.

Assigned: The Assigned Value is the robust mean of the reported results, outliers excluded. This is

often referred to as the "target" value.

 $\underline{+}u$: The uncertainty of the assigned value.

Reported: The result reported by the participant.

s: The Standard Deviation of Proficiency Assessment (SDPA). This value is used to determine

the acceptance limits for the PT evaluation.

z-Score: A value assigned to each reported result that is a measure of the degree to which it deviates

from the Assigned Value.

Score: The composite score of the four results reported for each analyte. It is normalized to a score

out of 100.

Bias: A flag assigned if bias is detected using the re-scaled z-score procedure.

3.0 Scoring System

Participant performance is evaluated for each proficiency testing sample by a quantitative method that is consistent with ISO/IEC 17043 - Conformity assessment- General requirements for the competence of proficiency testing providers, the International Harmonized Protocol for Proficiency Testing of (Chemical) Analytical Laboratories (2006), and ISO 13528:2015 Statistical methods for use in proficiency testing by interlaboratory comparisons.

The following is a brief description of the evaluation procedure used by PTC. The detailed evaluation procedure is described in PROC09 – PT Evaluation *Procedure*, which is available on the PTC website www.PTCanada.org).

3.1 HOMOGENEITY AND STABILITY ASSESSMENT

Homogeneity and stability are assessed using participant data. Regression analysis is performed on reported result against order of sample production (Homogeneity) and reported result against date of analysis (Stability). If the slope is significantly different than zero for either then the Standard Deviation of Proficiency Assessment (s) is increased to minimize the impact.

3.2 THE Z SCORE

A "z-score" is calculated for each reported result as follows:

$$z-Score = \frac{(x-\bar{X})}{SDPA}$$
 where: $x = \text{participant result};$ $X = \text{the Assigned Value};$ SDPA = the Standard Deviation for Proficiency Assessment.

The assigned value X is generally estimated from the inter-laboratory Robust mean after outliers due to obvious gross errors (e.g., reported in wrong units) have been removed.

The Standard Deviation for Proficiency Assessment, s, is determined as follows:

- The inter-laboratory Robust standard deviation (Stdev_{rob}) is calculated using reported results, obvious outliers removed;
- The regression equation standard deviation (Stdev_{reg}) is estimated from regression equations derived from previous studies (see PROC11- PT Regression Equations for details);
- The SDPA is the higher of Stdev_{rob} and Stdev_{reg};
- When a laboratory reports its detection limit, s will be estimated using a pooled variance procedure that uses both the inter-laboratory data and the reported detection limit.

3.2 COMPOSITE (PT) SCORE

Since each PT round involves four or two separate samples of distinct concentration for each test, it is necessary to calculate a composite PT score for each test to determine overall performance. The composite score is calculated by first averaging the absolute z-scores for the four results and then calculating a final score as $100 + (-15 \times avg |z|)$.

Acceptable PT Scores equal or exceed 70.

3.3 IDENTIFYING BIAS

The proficiency testing report provides flags for bias. These are determined using the re-scaled z-score procedure.

$$RSZ = \frac{\sum z}{\sqrt{N}}$$
 where z = the z- score
N = the number of samples

Flags are assigned for each test group/parameter combination as follows:

 $RSZ \ge -2$ and ≤ 2 no flag assigned RSZ > 2 H (High) RSZ > 3 VH (Very High) RSZ < -2 L (LOW) VL (Very Low)

3.4 DEVIATIONS FROM EVALUATION PROCEDURE

Other than changes to the Standard Deviation of Proficiency Assessment due to homogeneity or stability flags, any deviation from the published evaluation procedure is described on the cover page(s) of the final PTC Proficiency Testing Report.

4.0 PT Round Specific Data Summary

The following pages provide more detailed information about the PT round indicated in the cover page of this report than is found in the participant-specific PTC Proficiency Testing Report. The graphical representations and the statistical summaries are based upon the data after outliers have been removed.

4.1 SUMMARY STATISTICS

In addition to some of the statistics found in the customer reports, this table includes additional summary statistics such as Median, different measures of dispersion, the number of outliers removed, the number of results in the Questionable range (|z| between 2 and 3) and the Unacceptable range (z > 3), and whether a data set was flagged for Homogeneity or Stability. This section also includes sorted scatter plots of the data for each sample.

4.2 z-SCORE PLOTS

The z -scores for each sample are ranked in increasing order and plotted. When the data is normally distributed, the plot should show a slight sigmoidal curve, with an equal number of points above zero as below. Each bar in these plots is colour-coded to indicate the analytical method used by the participant.

4.3 KERNEL DENSITY PLOTS

Kernel density plots are generated for each data set. These plots are a graphical way to represent the overall data distribution and are used to visualize possible deviations from normality and unimodality.

4.4 STABILITY AND HOMOGENEITY PLOTS

Plots of reported result against analysis date, and reported result against order of bottling are displayed, along with the regression line. These regression analyses are used to determine if the SDPA should be adjusted due to homogeneity or stability.

4.5 BOX-AND-WHISKER PLOTS

Box-and-Whisker plots are another way to display the distribution of the data. The box denotes the first and third quartile and the whiskers are the 5^{th} and 95th percentile.

4.6 HISTORIC COMPARISON PLOT

The Historic Comparison Plot is a plot of robust mean against robust standard deviation for the previous ten PT rounds as well as the current PT round. This plot can be used to identify possible changes in the sample formulation.

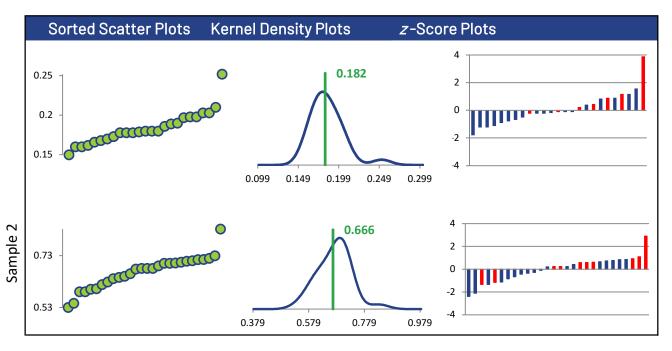
ARSENIC

Summary Statistics

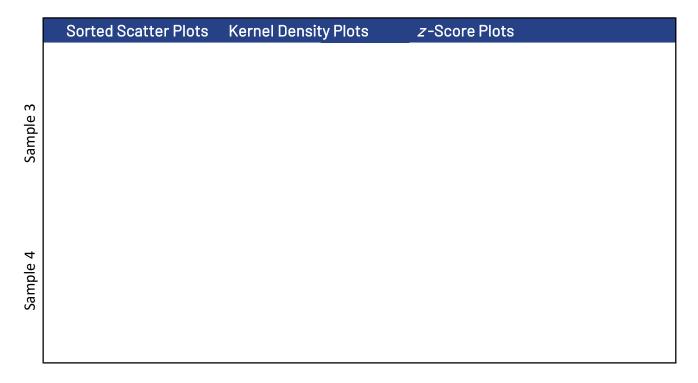
Statistic	C39-1	C39-2	C39-3	C39-4
N	25	28	0	0
Median mg/L	0.180	0.680		
Robust Mean mg/L	0.182	0.666		
U mg/L	0.00450	0.0135		
Robust Standard Deviation mg/L	0.0180	0.0572		
Regression Standard Deviation mg/L				
Stability Flag				
Homogeneity Flag				
Standard Deviation Used (SDPA) mg/L	0.0180	0.0572		
Outliers	0	0	0	0
z >3.0	1	0	0	0
2< z <3	0	3	0	0

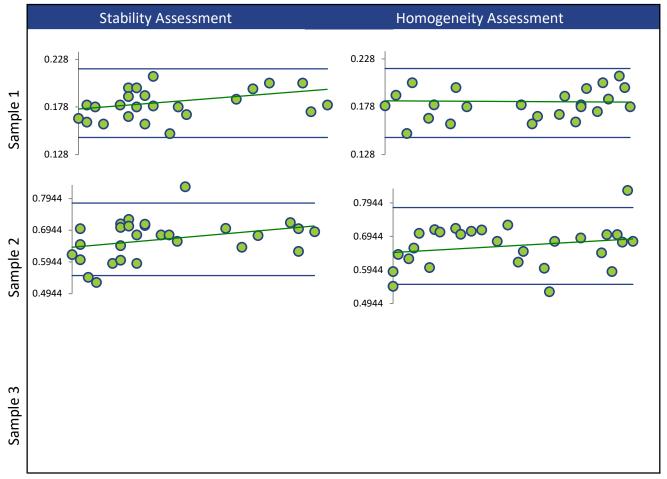
Methods Used

Method	C39-1	C39-2	C39-3	C39-4
ICP/MS (Blue)	18	18	0	0
ICP/0ES(Red)	7	10	0	0

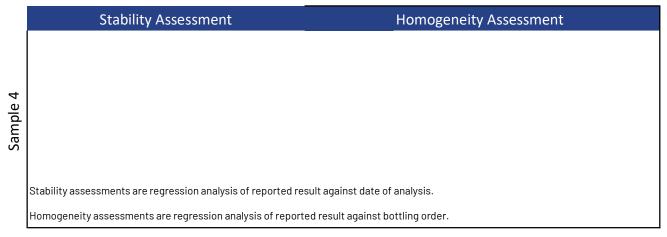


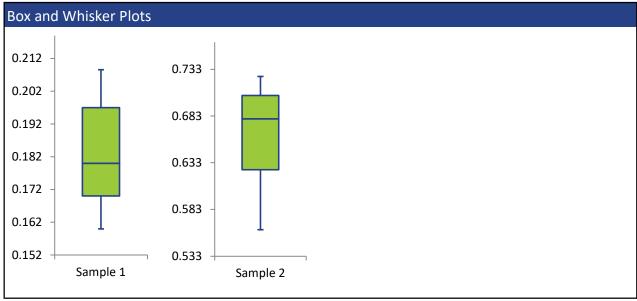
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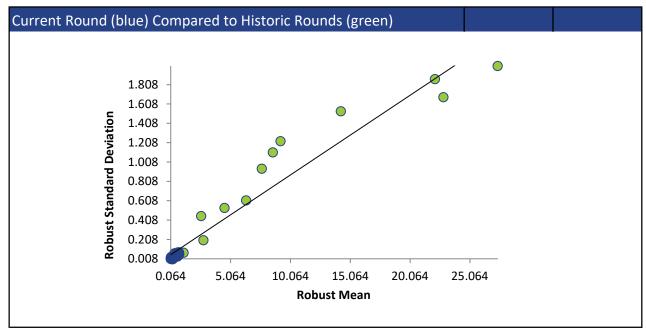




ARSENIC







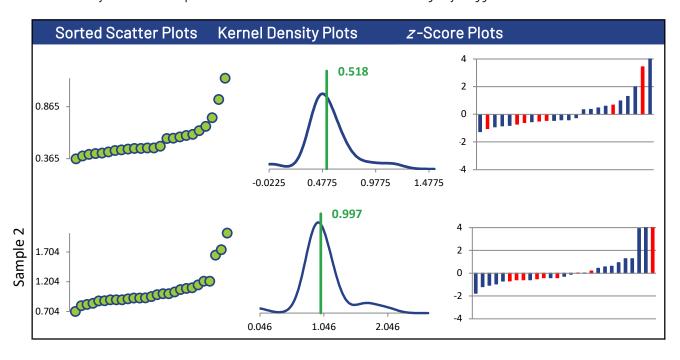
BARIUM

Summary Statistics

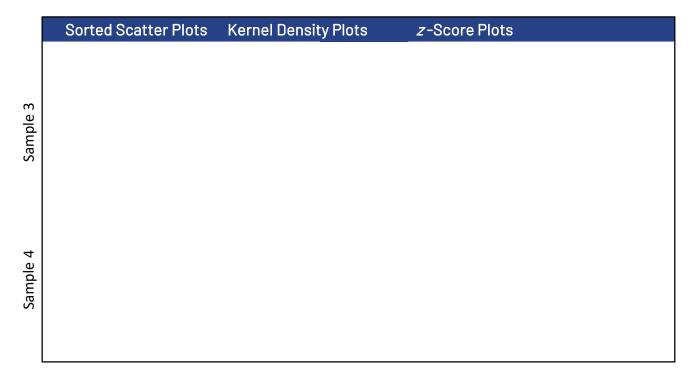
Statistic	C39-1	C39-2	C39-3	C39-4
N	24	27	0	0
Median mg/L	0.470	0.952		
Robust Mean mg/L	0.518	0.997		
U mg/L	0.0309	0.0402		
Robust Standard Deviation mg/L	0.121	0.167		
Regression Standard Deviation mg/L				
Stability Flag				
Homogeneity Flag				
Standard Deviation Used (SDPA) mg/L	0.121	0.167		
Outliers	0	0	0	0
z >3.0	2	3	0	0
2< z <3	0	0	0	0

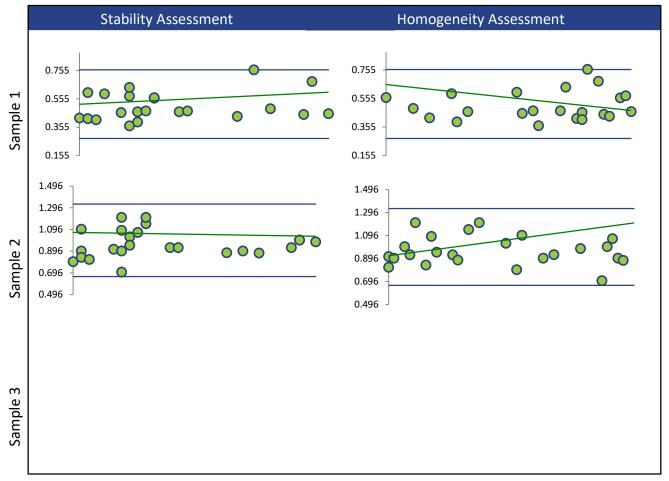
Methods Used

Method	C39-1	C39-2	C39-3	C39-4
ICP/MS(Blue)	17	18	0	0
ICP/0ES(Red)	7	9	0	0

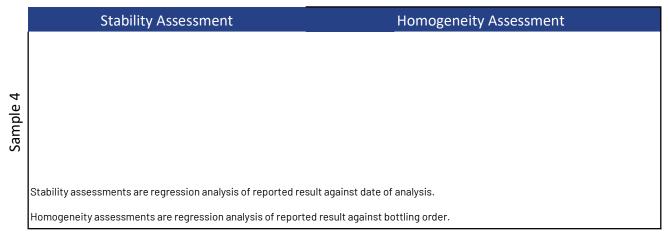


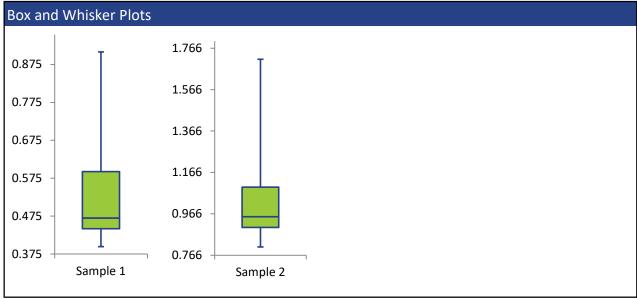
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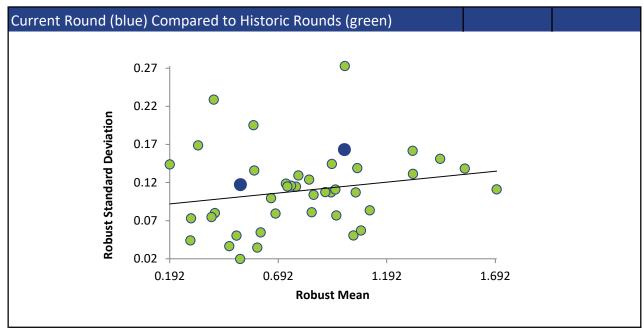




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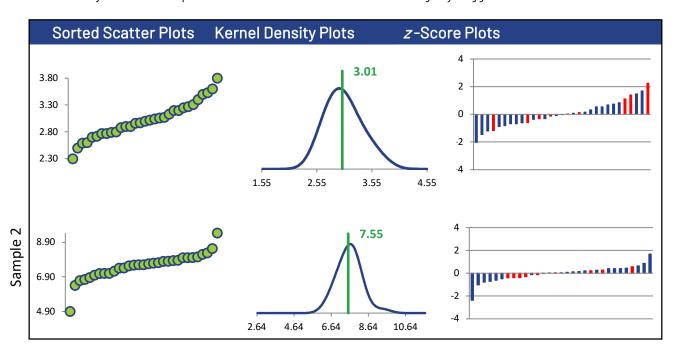
BORON

Summary Statistics

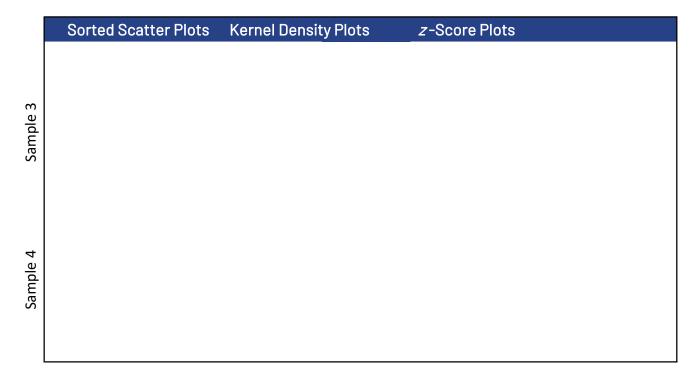
Statistic	C39-1	C39-2	C39-3	C39-4
N	31	31	0	0
Median mg/L	3.00	7.60		
Robust Mean mg/L	3.01	7.55		
U mg/L	0.0781	0.134		
Robust Standard Deviation mg/L	0.348	0.598		
Regression Standard Deviation mg/L				
Stability Flag				
Homogeneity Flag		Homogeneity		
Standard Deviation Used (SDPA) mg/L	0.348	1.12		
Outliers	1	1	0	0
z >3.0	0	0	0	0
2< z <3	2	1	0	0

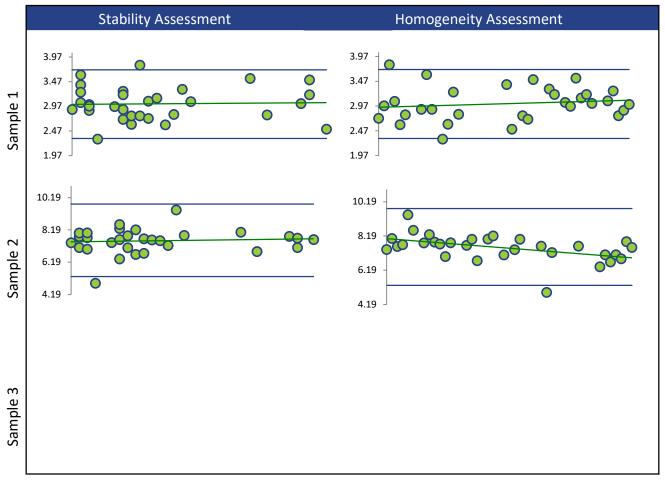
Methods Used

Method	C39-1	C39-2	C39-3	C39-4
ICP/MS (Blue)	22	22	0	0
ICP/0ES(Red)	9	9	0	0

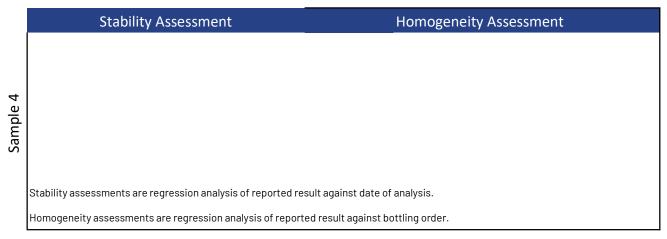


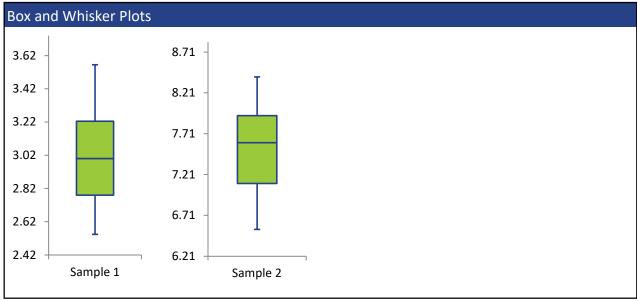
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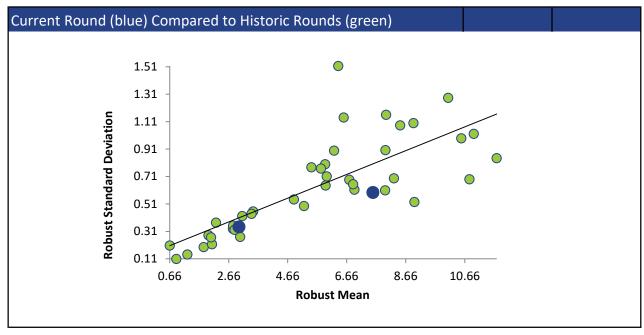




BORON







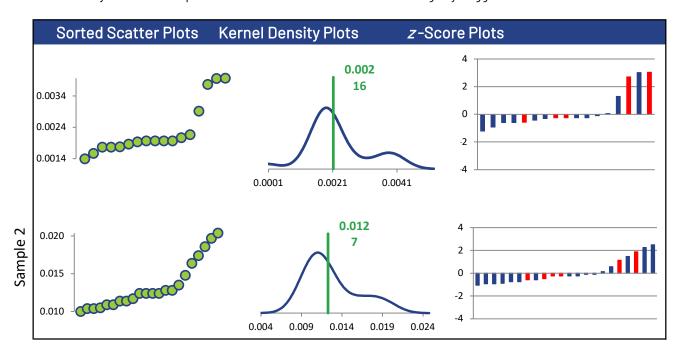
CADMIUM

Summary Statistics

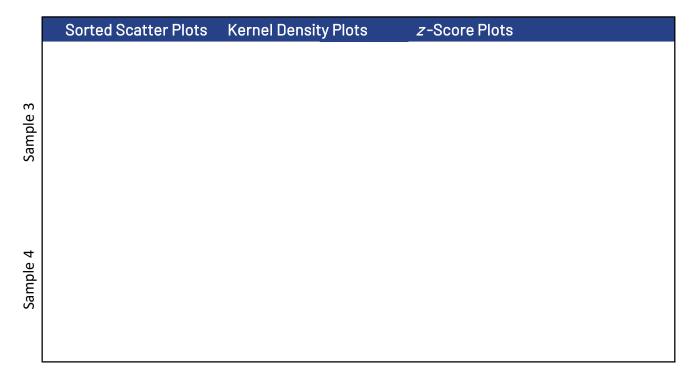
Statistic	C39-1	C39-2	C39-3	C39-4
N	17	22	0	0
Median mg/L	0.00200	0.0120		
Robust Mean mg/L	0.00216	0.0127		
U mg/L	0.000183	0.000778		
Robust Standard Deviation mg/L	0.000604	0.00292		
Regression Standard Deviation mg/L				
Stability Flag				
Homogeneity Flag				
Standard Deviation Used (SDPA) mg/L	0.000604	0.00292		
Outliers	0	0	0	0
z >3.0	2	0	0	0
2< z <3	1	2	0	0

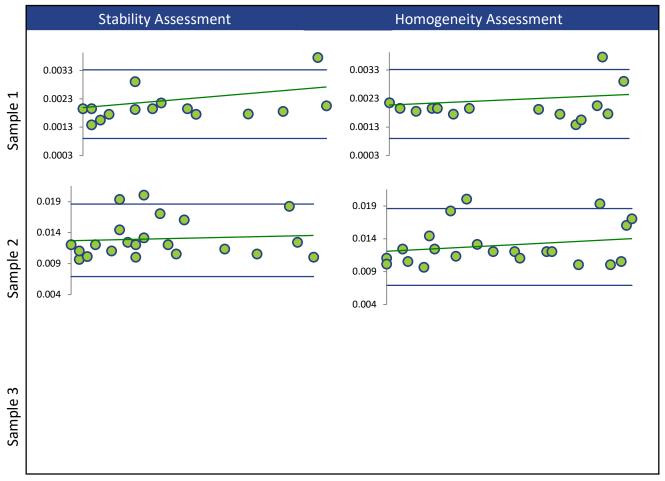
Methods Used

Method	C39-1	C39-2	C39-3	C39-4
ICP/MS (Blue)	12	16	0	0
ICP/0ES(Red)	5	6	0	0

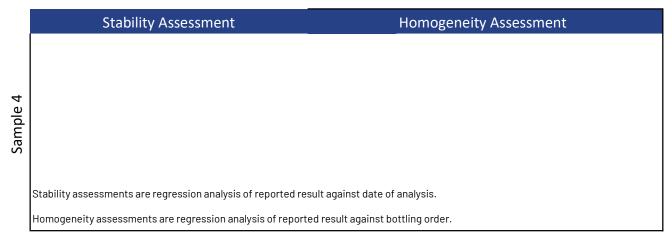


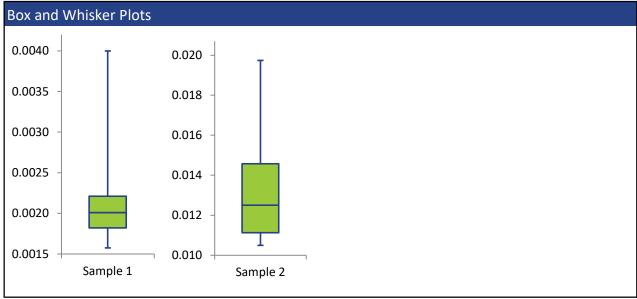
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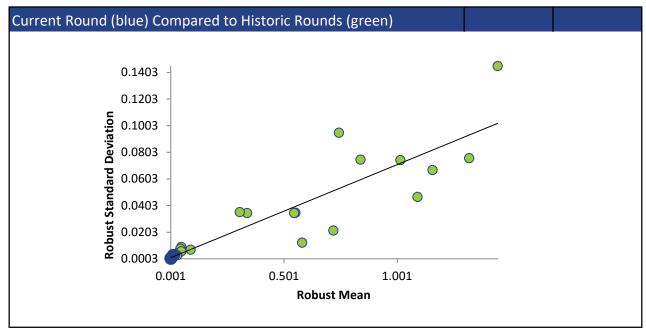




CADMIUM







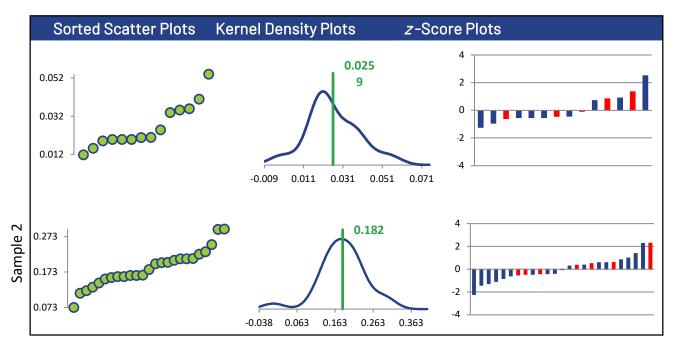
CHROMIUM

Summary Statistics

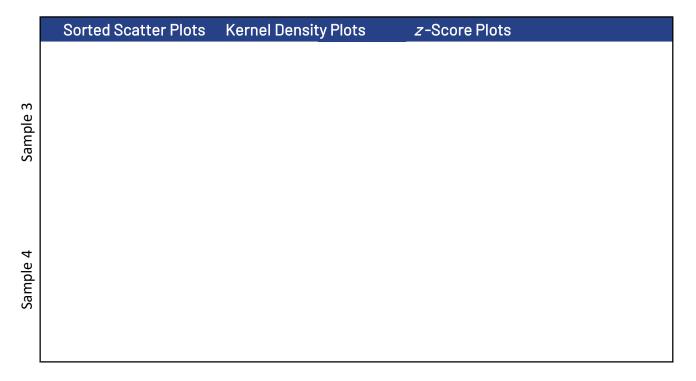
Statistic	C39-1	C39-2	C39-3	C39-4
N	14	25	0	0
Median mg/L	0.0211	0.180		
Robust Mean mg/L	0.0259	0.182		
U mg/L	0.00374	0.0123		
Robust Standard Deviation mg/L	0.0112	0.0492		
Regression Standard Deviation mg/L				
Stability Flag				
Homogeneity Flag				
Standard Deviation Used (SDPA) mg/L	0.0112	0.0492		
Outliers	0	0	0	0
z >3.0	0	0	0	0
2< z <3	1	3	0	0

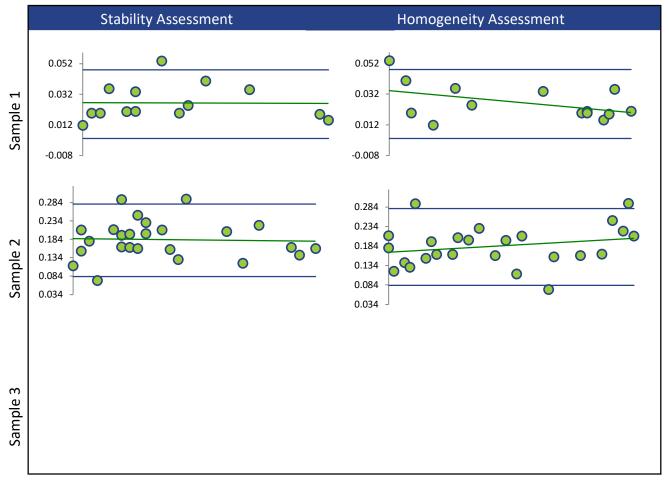
Methods Used

Method	C39-1	C39-2	C39-3	C39-4
ICP/MS(Blue)	9	18	0	0
ICP/0ES(Red)	5	7	0	0

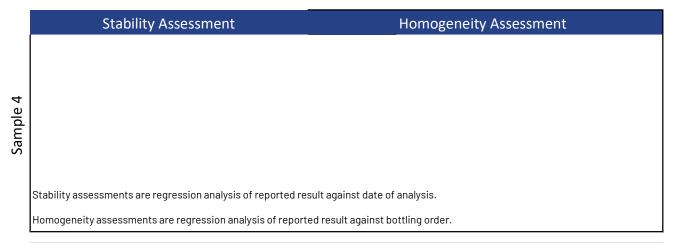


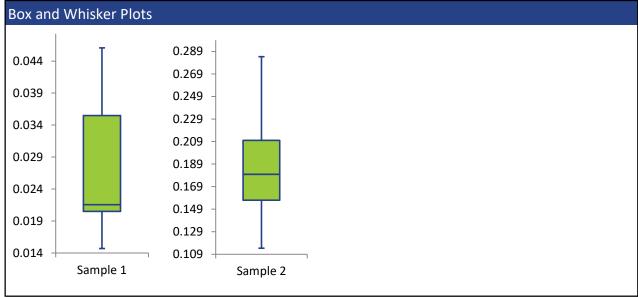
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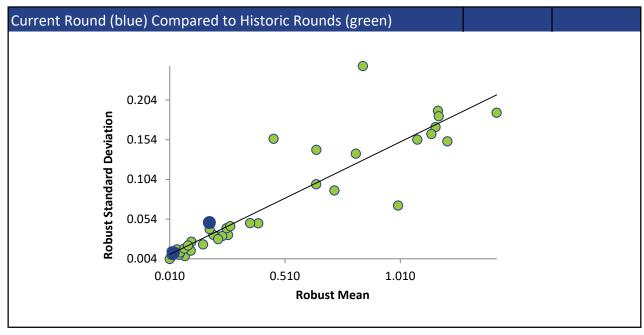




CHROMIUM







CYANIDE (WAD)

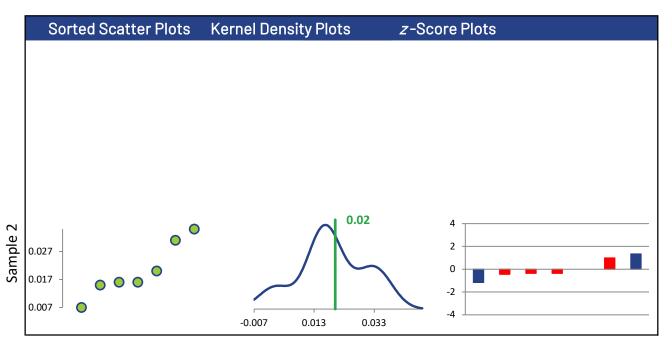
Summary Statistics

Excluded

Statistic	C39-1	C39-2	C39-3	C39-4
N	0	7	0	0
Median mg/L		0.0160		
Robust Mean mg/L		0.0200		
U mg/L		0.00524		
Robust Standard Deviation mg/L		0.0111		
Regression Standard Deviation mg/L				
Stability Flag				
Homogeneity Flag				
Standard Deviation Used (SDPA) mg/L		0.0111		
Outliers	2	0	0	0
z >3.0	0	0	0	0
2< z <3	0	0	0	0

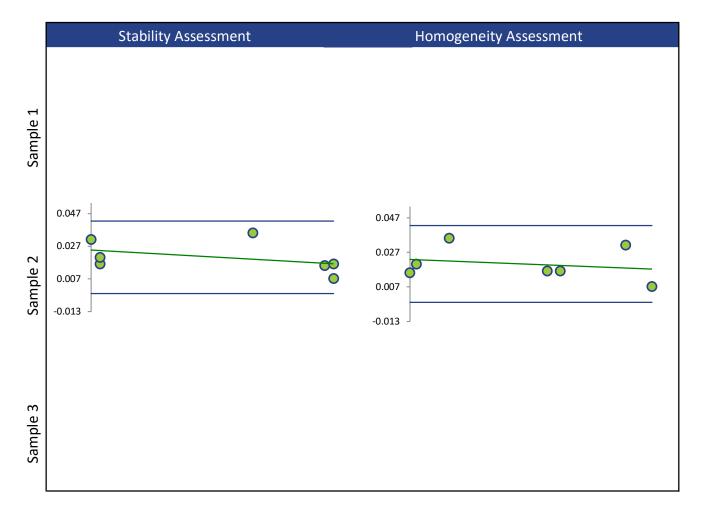
Methods Used

Method	C39-1	C39-2	C39-3	C39-4
AUTO COLOR (Blue)	0	3	0	0
COLORIMETRIC (Red)	0	4	0	0

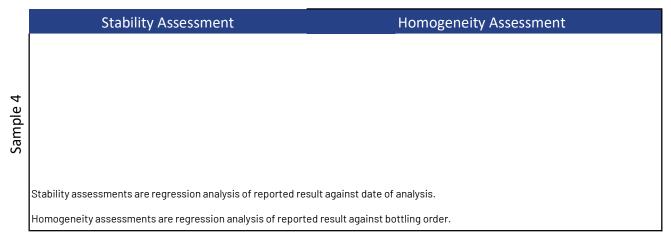


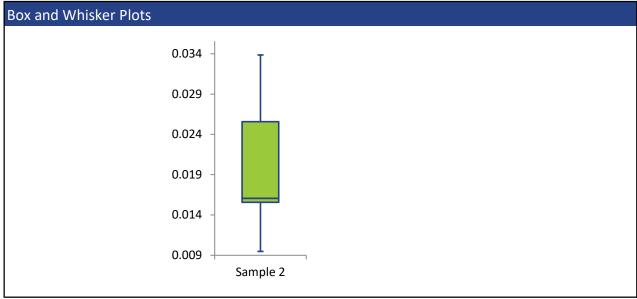
CYANIDE (WAD)

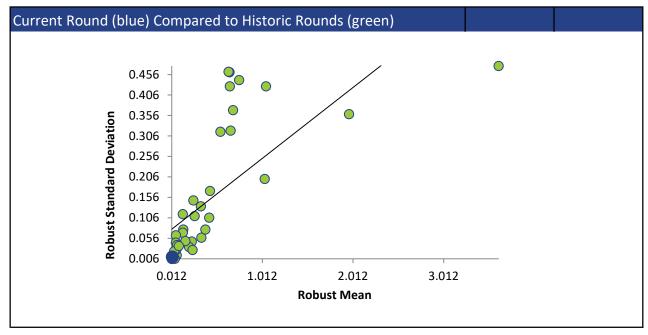
	Sorted Scatter Plots	Kernel Density Plots	z-Score Plots	
Sample 3				
Sample 4				



CYANIDE (WAD)







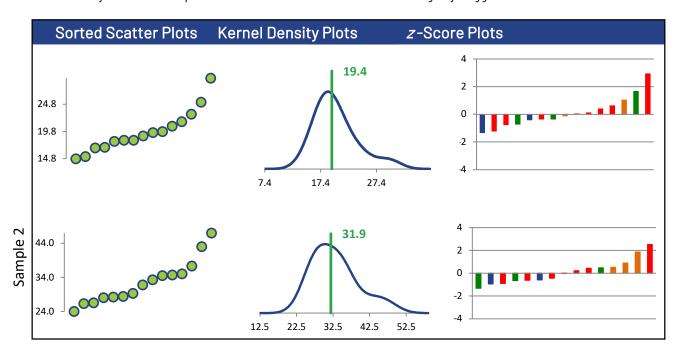
FLUORIDE

Summary Statistics

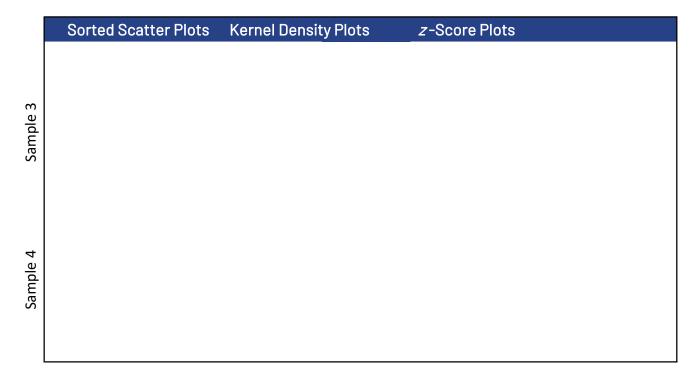
Statistic	C39-1	C39-2	C39-3	C39-4
N	15	15	0	0
Median mg/L	19.0	31.8		
Robust Mean mg/L	19.4	31.9		
U mg/L	1.12	1.93		
Robust Standard Deviation mg/L	3.47	5.97		
Regression Standard Deviation mg/L				
Stability Flag				
Homogeneity Flag				
Standard Deviation Used (SDPA) mg/L	3.47	5.97		
Outliers	0	0	0	0
z >3.0	0	0	0	0
2< z <3	1	1	0	0

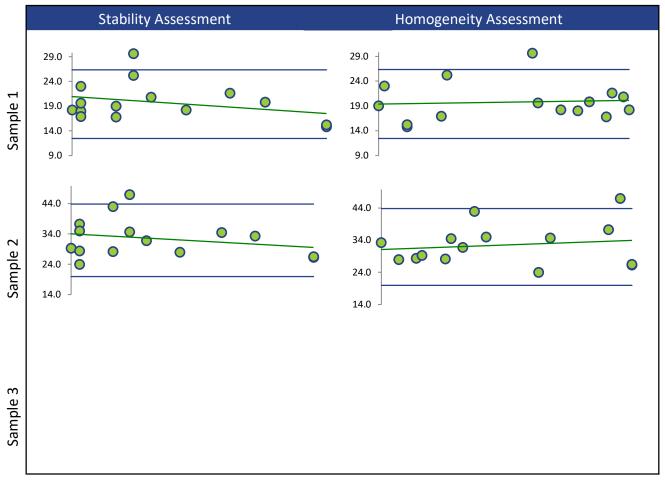
Methods Used

Method	C39-1	C39-2	C39-3	C39-4
TITRIMETRIC (Blue)	2	2	0	0
SELECTIVE ION ELECTRODE (Red)	7	7	0	0
COLORIMETRIC (Green)	3	3	0	0
ION CHROMATOGRAPHY (Orange)	3	3	0	0

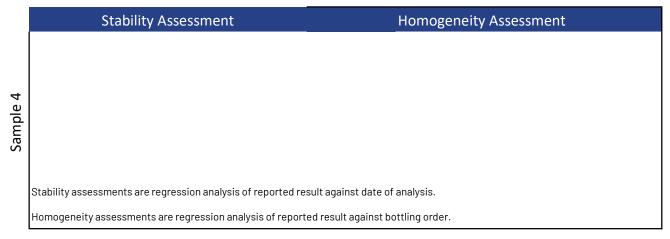


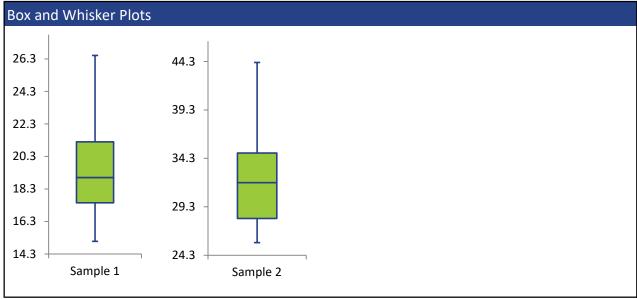
FLUORIDE

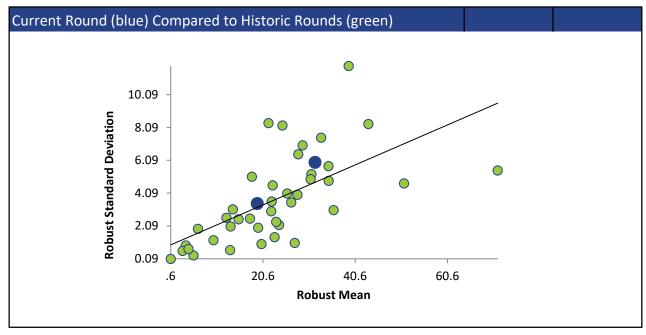




FLUORIDE







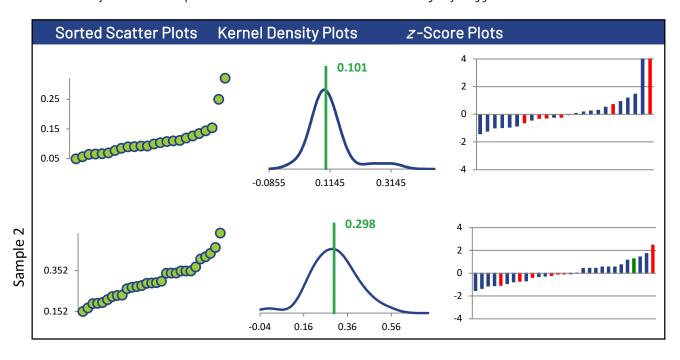
LEAD

Summary Statistics

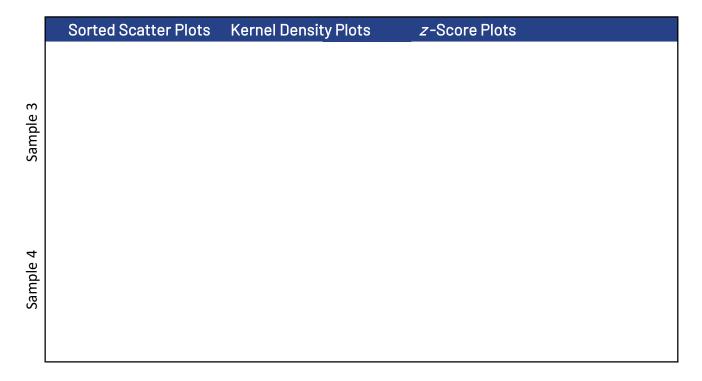
Statistic	C39-1	C39-2	C39-3	C39-4
N	24	29	0	0
Median mg/L	0.0966	0.291		
Robust Mean mg/L	0.101	0.298		
U mg/L	0.00921	0.0224		
Robust Standard Deviation mg/L	0.0361	0.0965		
Regression Standard Deviation mg/L				
Stability Flag				
Homogeneity Flag				
Standard Deviation Used (SDPA) mg/L	0.0361	0.0965		
Outliers	0	1	0	0
z >3.0	2	0	0	0
2< z <3	0	1	0	0

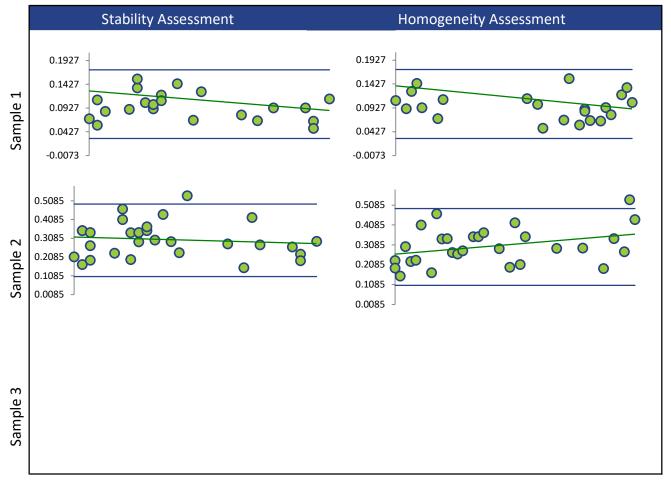
Methods Used

Method	C39-1	C39-2	C39-3	C39-4
ICP/MS (Blue)	18	21	0	0
ICP/0ES(Red)	6	7	0	0
AA FLAME (Green)	0	1	0	0

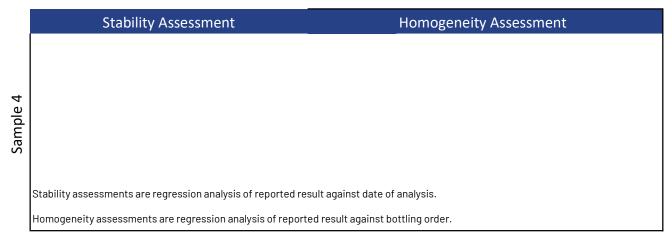


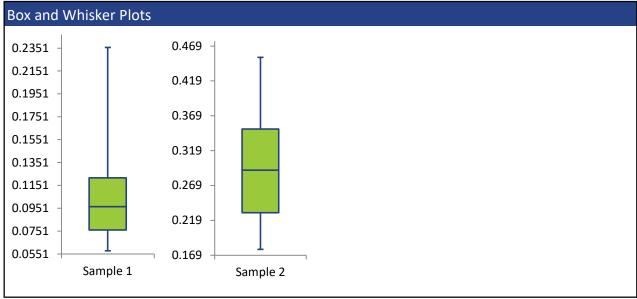
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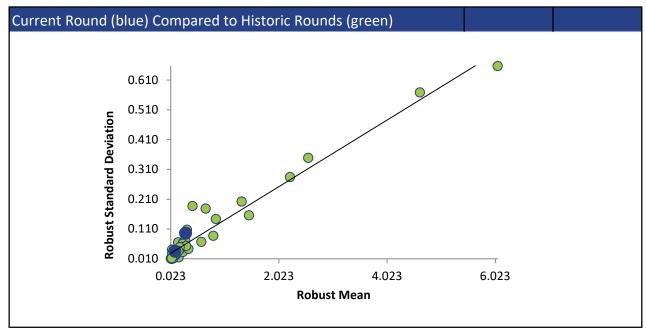




LEAD







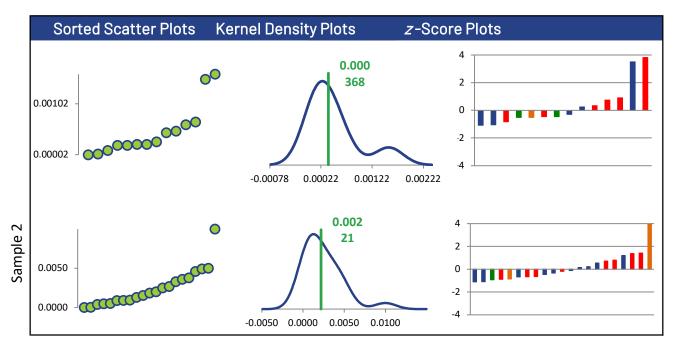
MERCURY

Summary Statistics

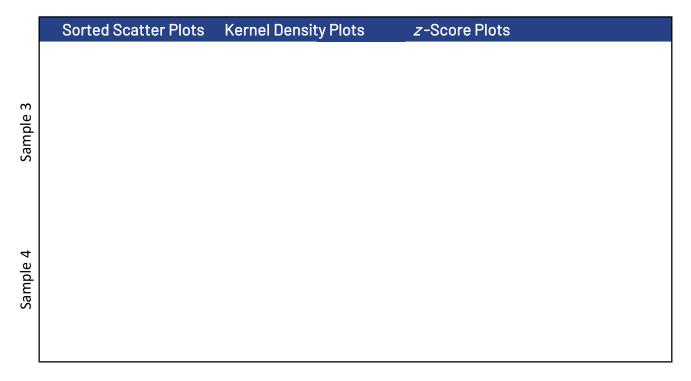
Statistic	C39-1	C39-2	C39-3	C39-4
N	14	21	0	0
Median mg/L	0.000245	0.00185		
Robust Mean mg/L	0.000368	0.00221		
U mg/L	0.000108	0.000537		
Robust Standard Deviation mg/L	0.000322	0.00197		
Regression Standard Deviation mg/L				
Stability Flag				
Homogeneity Flag				
Standard Deviation Used (SDPA) mg/L	0.000322	0.00197		
Outliers	1	2	0	0
z >3.0	2	1	0	0
2< z <3	0	0	0	0

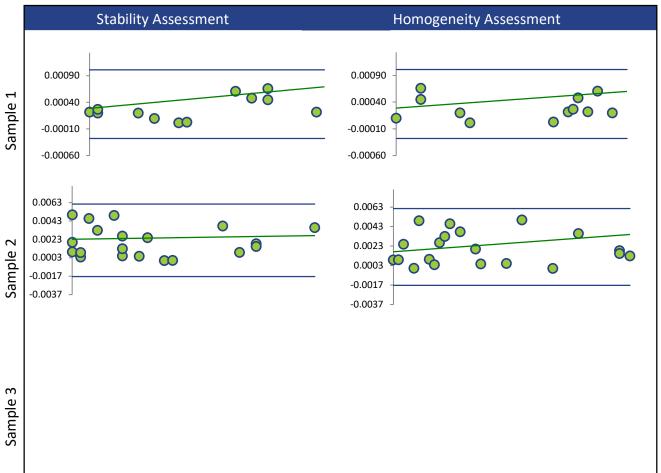
Methods Used

	Method	C39-1	C39-2	C39-3	C39-4
	COLD VAPOUR AA (Blue)	5	10	0	0
	ICP/MS(Red)	6	8	0	0
≀ ATO	MIC FLUORESCENCE SPECTROPHOTOMETRY (Green)	2	1	0	0
	ICP/0ES (Orange)	1	2	0	0

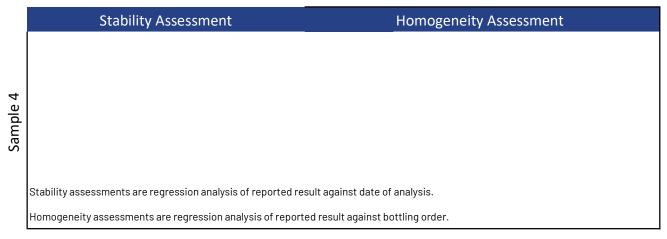


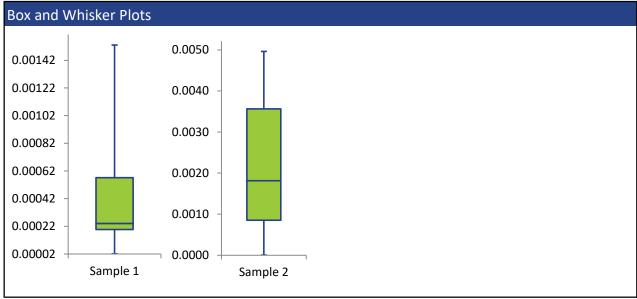
MERCURY

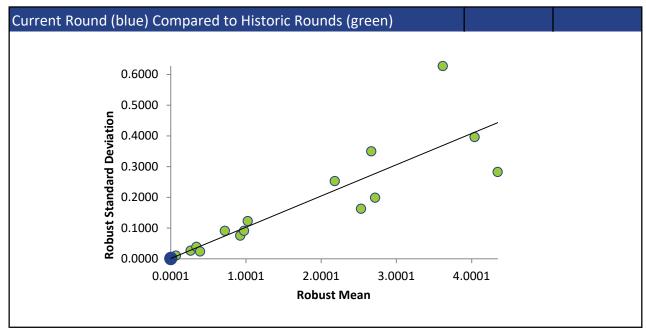




MERCURY







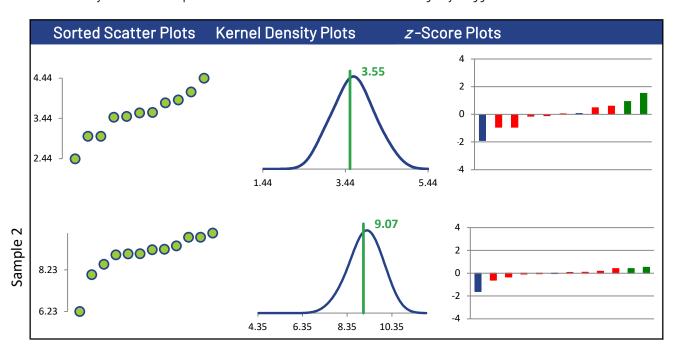
NITRATE

Summary Statistics

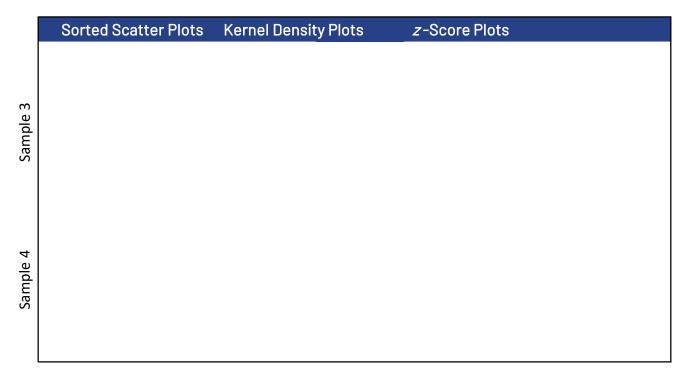
Statistic	C39-1	C39-2	C39-3	C39-4
N	11	12	0	0
Median mg/L	3.58	9.11		
Robust Mean mg/L	3.55	9.07		
U mg/L	0.220	0.271		
Robust Standard Deviation mg/L	0.583	0.752		
Regression Standard Deviation mg/L				
Stability Flag				
Homogeneity Flag		Homogeneity		
Standard Deviation Used (SDPA) mg/L	0.583	1.78		
Outliers	0	0	0	0
z >3.0	0	0	0	0
2< z <3	0	0	0	0

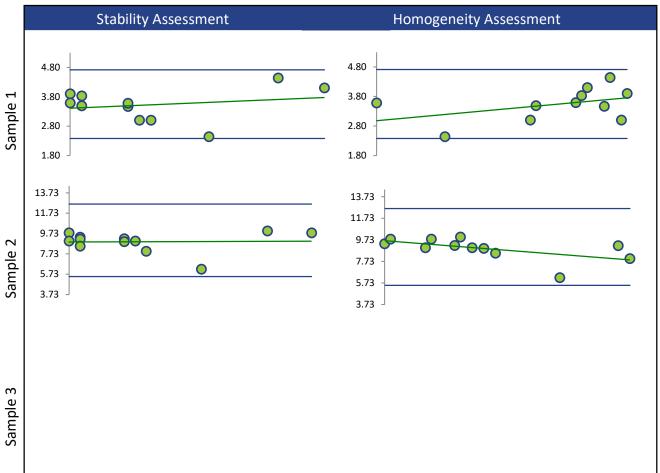
Methods Used

Method	C39-1	C39-2	C39-3	C39-4
COLORIMETRIC (Blue)	2	2	0	0
ION CHROMATOGRAPHY (Red)	7	8	0	0
AUTO COLOR (Green)	2	2	0	0

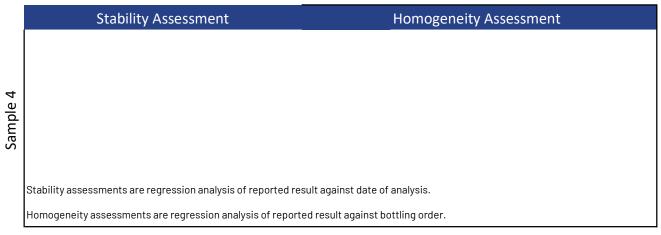


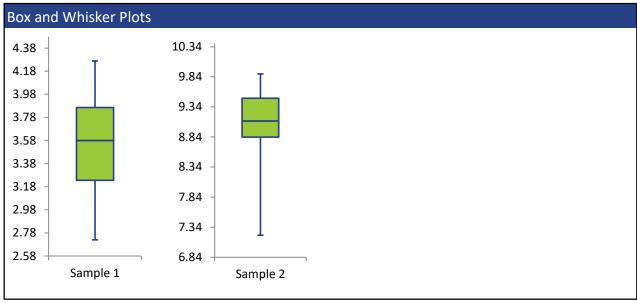
NITRATE

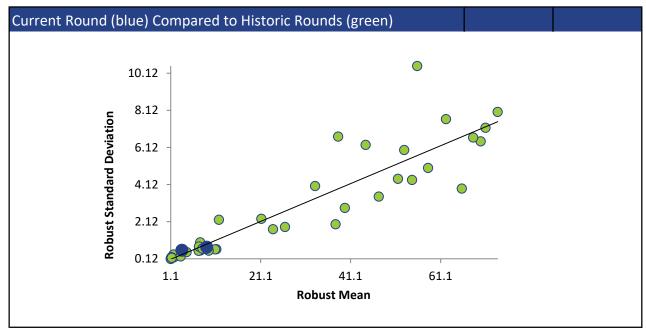




NITRATE







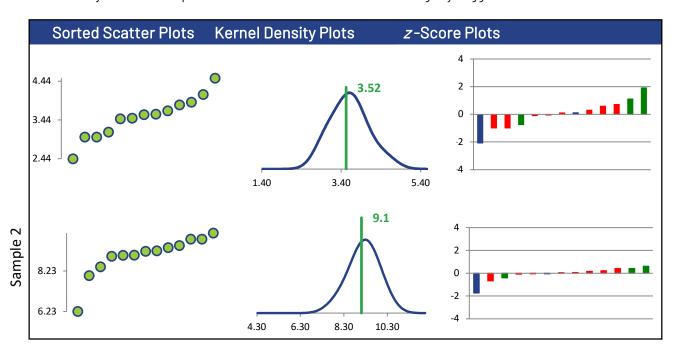
NITRATE PLUS NITRITE

Summary Statistics

Statistic	C39-1	C39-2	C39-3	C39-4
N	13	13	0	0
Median mg/L	3.58	9.20		
Robust Mean mg/L	3.52	9.10		
U mg/L	0.180	0.258		
Robust Standard Deviation mg/L	0.520	0.744		
Regression Standard Deviation mg/L				
Stability Flag				
Homogeneity Flag		Homogeneity		
Standard Deviation Used (SDPA) mg/L	0.520	1.65		
Outliers	0	0	0	0
z >3.0	0	0	0	0
2< z <3	1	0	0	0

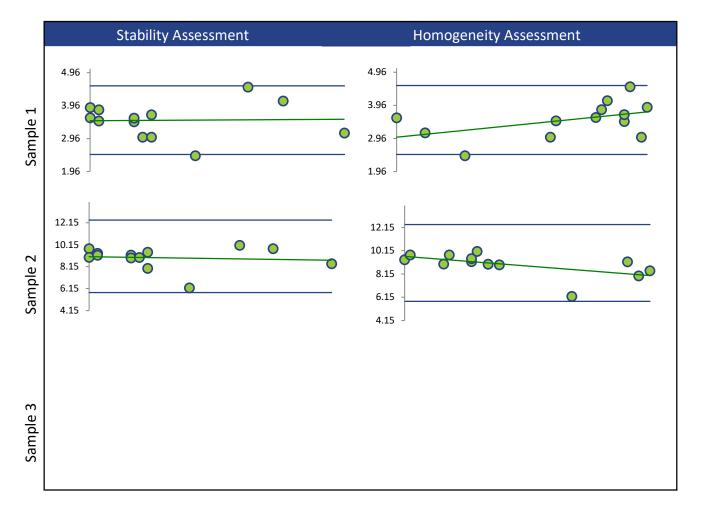
Methods Used

Method	C39-1	C39-2	C39-3	C39-4
COLORIMETRIC (Blue)	2	2	0	0
ION CHROMATOGRAPHY (Red)	8	8	0	0
AUTO COLOR (Green)	3	3	0	0

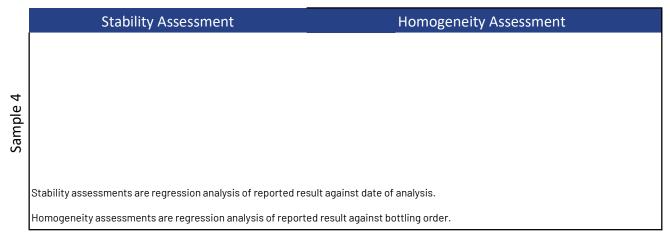


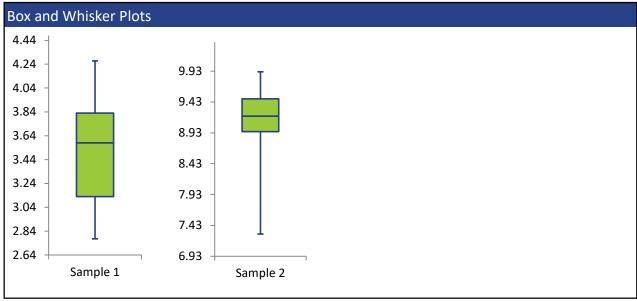
NITRATE PLUS NITRITE

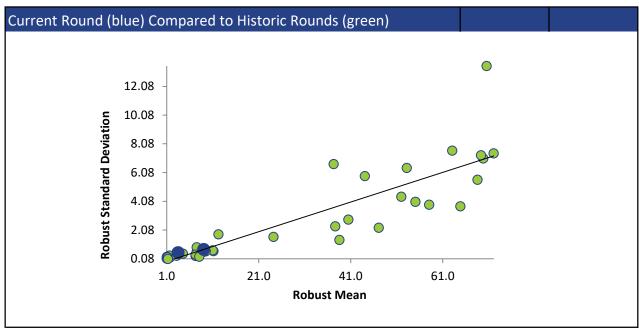
	Sorted Scatter Plots	Kernel Density Plots	z-Score Plots	
Sample 3				
Sample 4				



NITRATE PLUS NITRITE







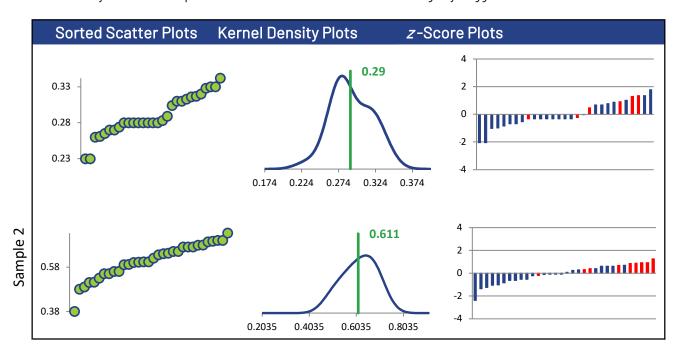
SELENIUM

Summary Statistics

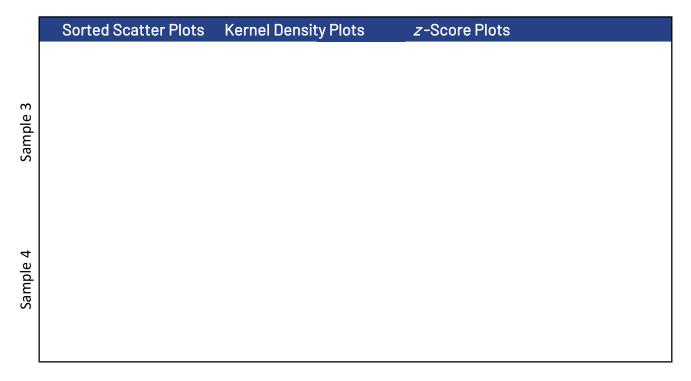
Statistic	C39-1	C39-2	C39-3	C39-4
N	29	32	0	0
Median mg/L	0.280	0.612		
Robust Mean mg/L	0.290	0.611		
U mg/L	0.00675	0.0175		
Robust Standard Deviation mg/L	0.0291	0.0793		
Regression Standard Deviation mg/L				
Stability Flag		Stability		
Homogeneity Flag				
Standard Deviation Used (SDPA) mg/L	0.0291	0.0966		
Outliers	1	0	0	0
z >3.0	0	0	0	0
2< z <3	2	1	0	0

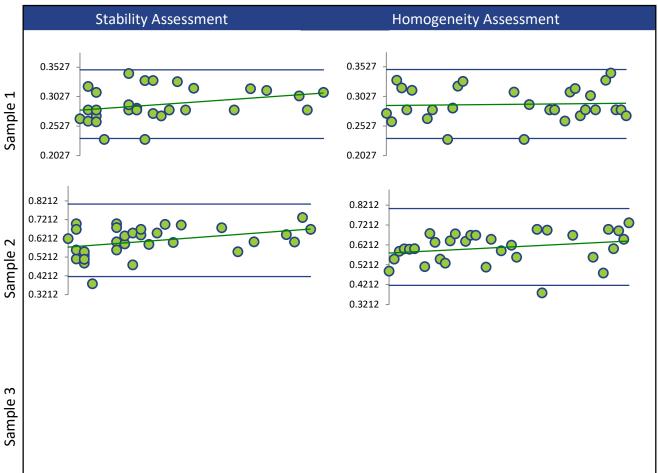
Methods Used

Method	C39-1	C39-2	C39-3	C39-4
ICP/MS (Blue)	23	23	0	0
ICP/0ES(Red)	6	9	0	0

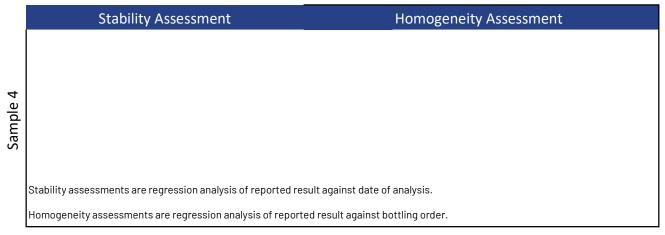


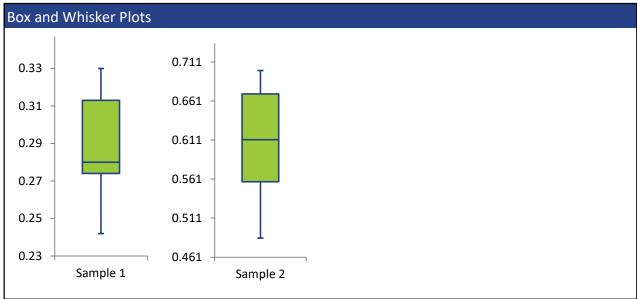
SELENIUM

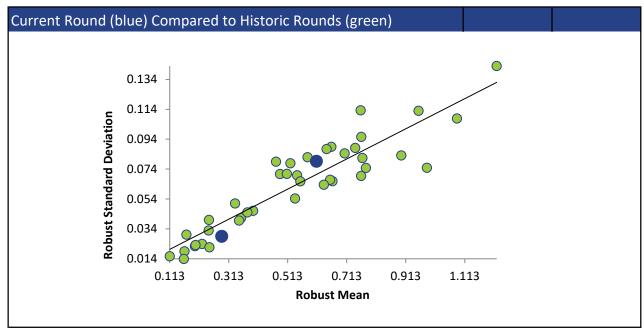




SELENIUM







SILVER

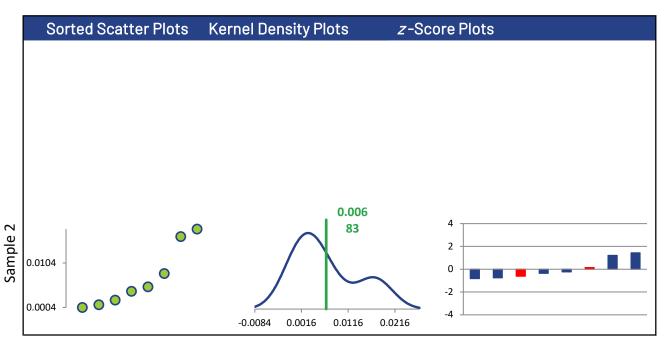
Summary Statistics

Excluded

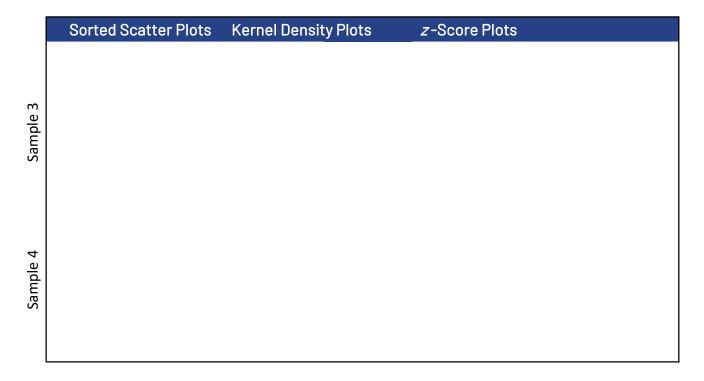
Statistic	C39-1	C39-2	C39-3	C39-4
N	0	8	0	0
Median mg/L		0.00450		
Robust Mean mg/L		0.00683		
U mg/L		0.00343		
Robust Standard Deviation mg/L		0.00775		
Regression Standard Deviation mg/L				
Stability Flag				
Homogeneity Flag				
Standard Deviation Used (SDPA) mg/L		0.00775		
Outliers	4	1	0	0
z >3.0	0	0	0	0
2< z <3	0	0	0	0

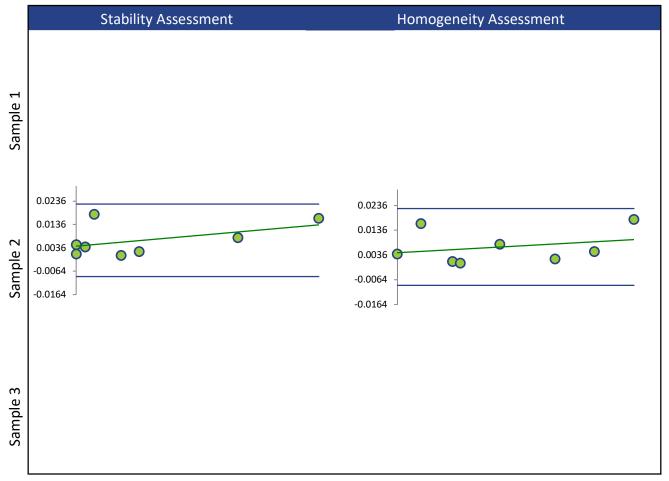
Methods Used

Method	C39-1	C39-2	C39-3	C39-4
ICP/MS (Blue)	0	6	0	0
ICP/0ES(Red)	0	2	0	0

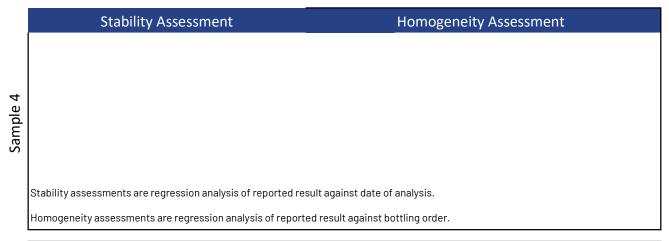


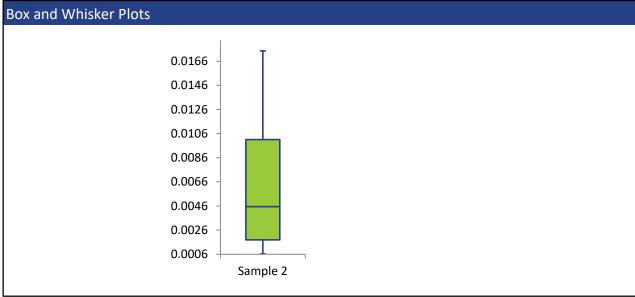
SILVER

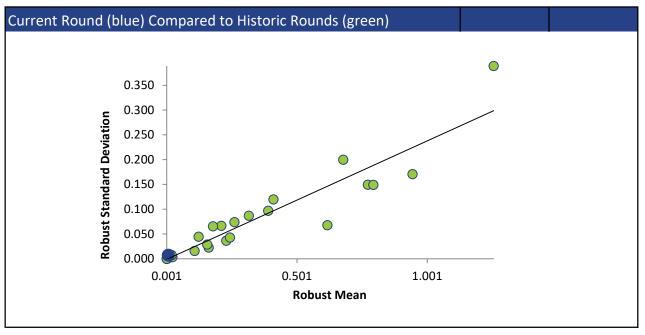




SILVER







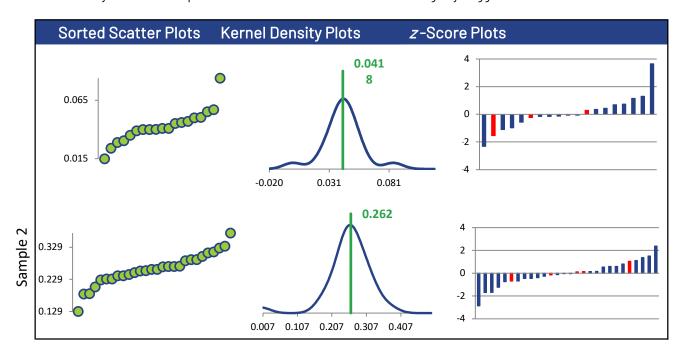
URANIUM

Summary Statistics

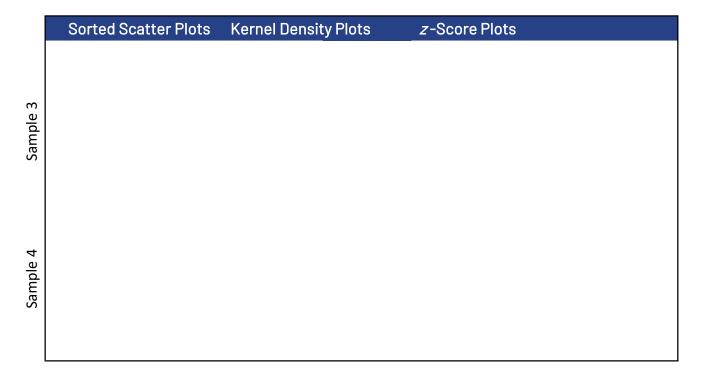
Statistic	C39-1	C39-2	C39-3	C39-4
N	19	28	0	0
Median mg/L	0.0410	0.260		
Robust Mean mg/L	0.0418	0.262		
U mg/L	0.00330	0.0110		
Robust Standard Deviation mg/L	0.0115	0.0465		
Regression Standard Deviation mg/L				
Stability Flag				
Homogeneity Flag				
Standard Deviation Used (SDPA) mg/L	0.0115	0.0465		
Outliers	0	1	0	0
z >3.0	1	0	0	0
2< z <3	1	2	0	0

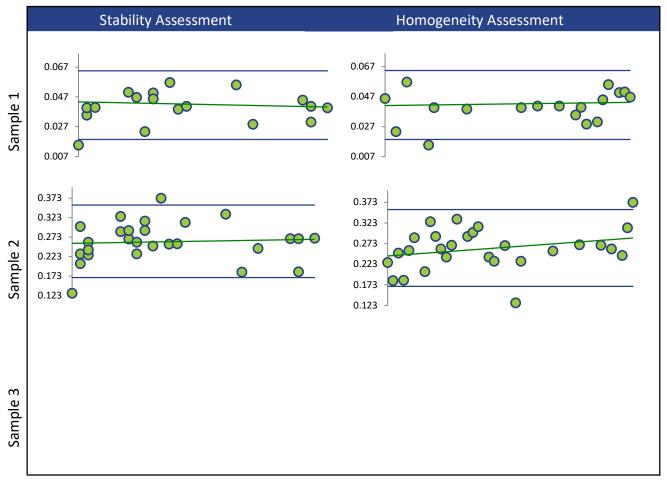
Methods Used

Method	C39-1	C39-2	C39-3	C39-4
ICP/MS (Blue)	16	23	0	0
ICP/0ES(Red)	3	5	0	0



URANIUM





URANIUM

