

Test Group Summary Report

C22 OP Pesticides in Water

January 2024 PT Round

Issued: March 5, 2024

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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.

<i>Code:</i>	The registration code that is unique to each analyte that a participant is registered for.
<i>App:</i>	If a participant is accredited by CALA, this three-digit number is the appendix number that the accredited method is assigned to.
<i>N:</i>	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.
<i>Assigned:</i>	The Assigned Value is the robust mean of the reported results, outliers excluded. This is often referred to as the "target" value.
<i>$\pm u$:</i>	The uncertainty of the assigned value.
<i>Reported:</i>	The result reported by the participant.
<i>s:</i>	The Standard Deviation of Proficiency Assessment (SDPA). This value is used to determine the acceptance limits for the PT evaluation.
<i>z-Score:</i>	A value assigned to each reported result that is a measure of the degree to which it deviates from the Assigned Value.
<i>Score:</i>	The composite score of the four results reported for each analyte. It is normalized to a score out of 100.
<i>Bias:</i>	A flag assigned if bias is detected using the re-scaled <i>z</i> -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:

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

The assigned value \bar{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 $100 + (-15 \times \text{avg } |z|)$.

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.

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

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

$RSZ \geq -2$ and ≤ 2	no flag assigned
$RSZ > 2$	H (High)
$RSZ > 3$	VH (Very High)
$RSZ < -2$	L (Low)
$RSZ < -3$	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 5th 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.

ATRAZINE

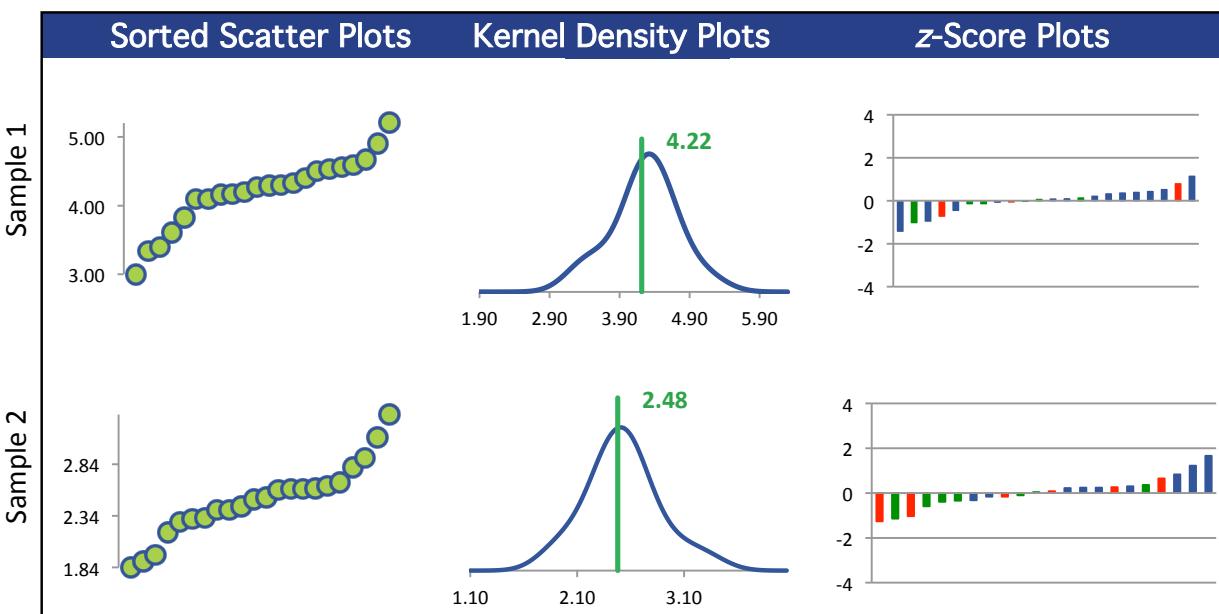
Summary Statistics

Statistic	C22-1	C22-2	C22-3	C22-4
N	22	22	22	22
Median $\mu\text{g/L}$	4.28	2.51	3.58	2.75
Robust Mean $\mu\text{g/L}$	4.22	2.48	3.58	2.70
$\text{U } \mu\text{g/L}$	0.129	0.0911	0.134	0.0640
Robust Standard Deviation $\mu\text{g/L}$	0.483	0.342	0.502	0.240
Regression Standard Deviation $\mu\text{g/L}$	0.862	0.507	0.730	0.552
Stability Flag				
Homogeneity Flag				
Standard Deviation Used (SDPA) $\mu\text{g/L}$	0.862	0.507	0.730	0.552
Outliers	0	0	0	0
$ \text{z} > 3.0$	0	0	0	0
$2 < \text{z} < 3$	0	0	0	0

Methods Used

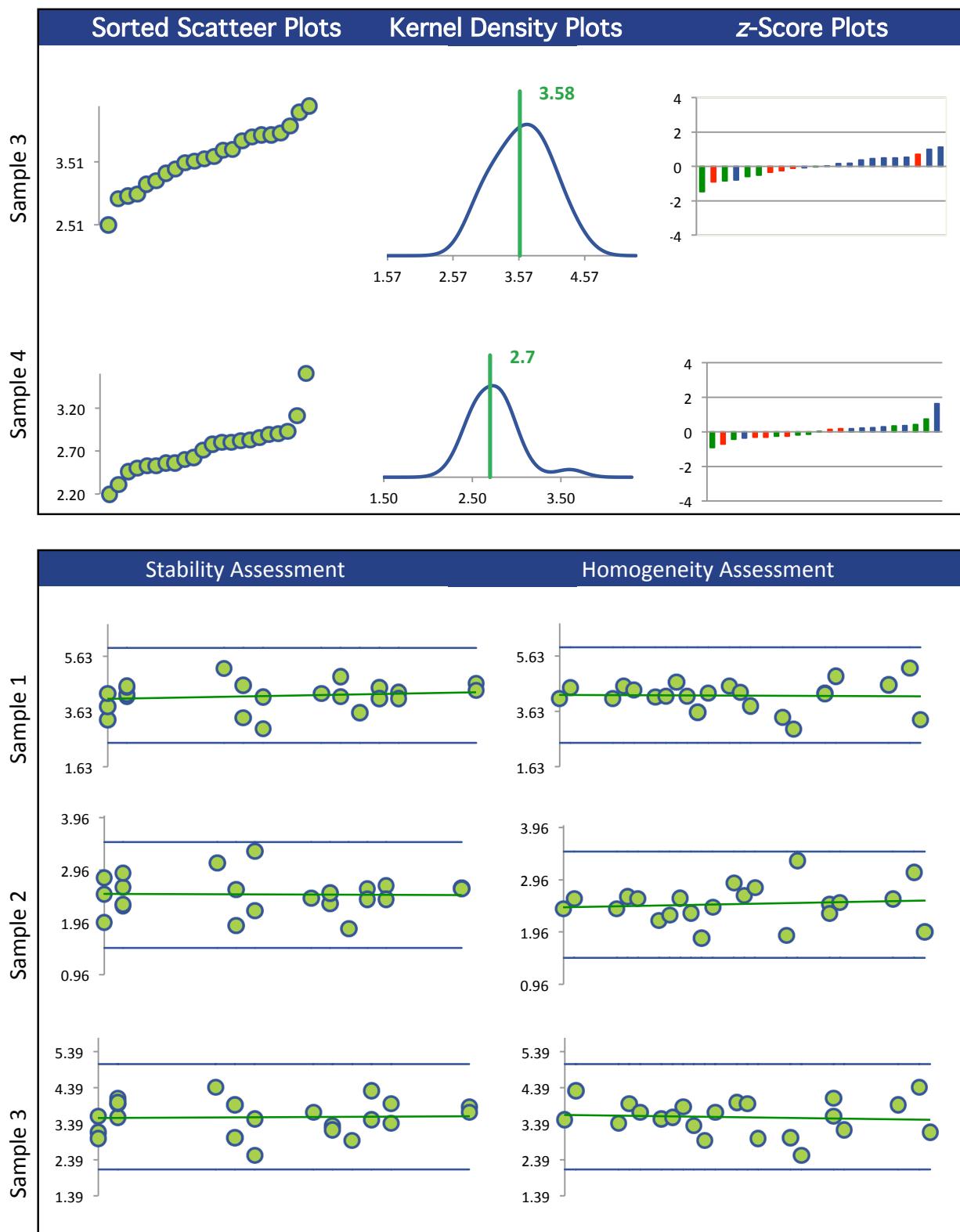
Method	C22-1	C22-2	C22-3	C22-4
GC/MS/MS (Blue)	2	2	2	2
LC-MS/MS (Red)	8	8	8	8
GC/MS (Green)	9	9	9	9
HPLC (Orange)	2	2	2	2
LC/MS (Black)	1	1	1	1

All summary stats and the plots below are based on the data excluding any flagged outliers



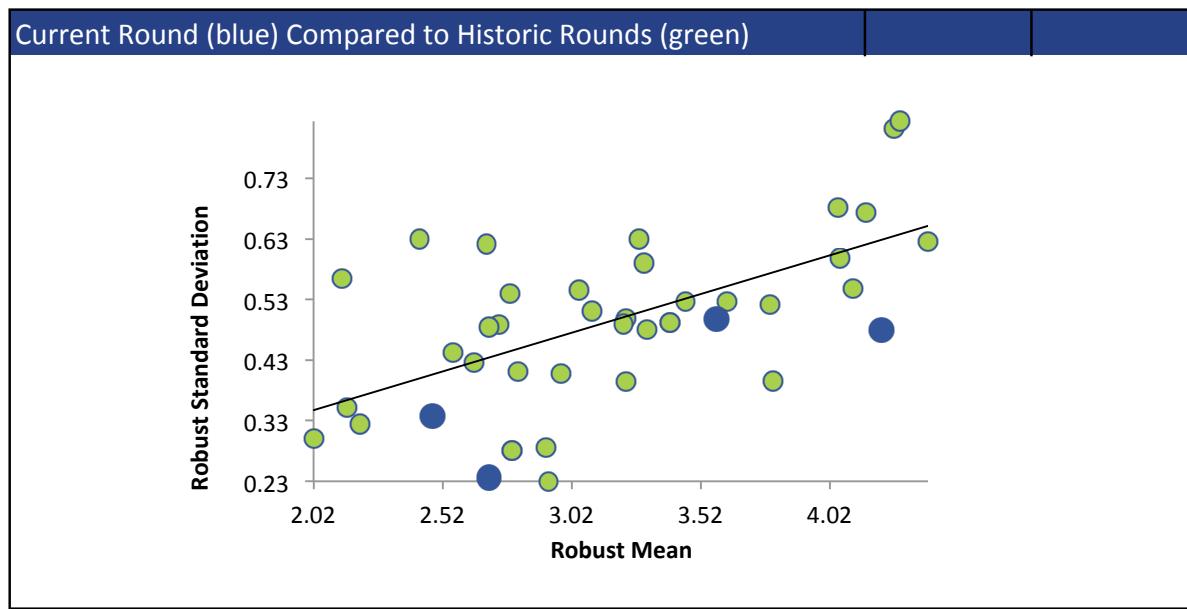
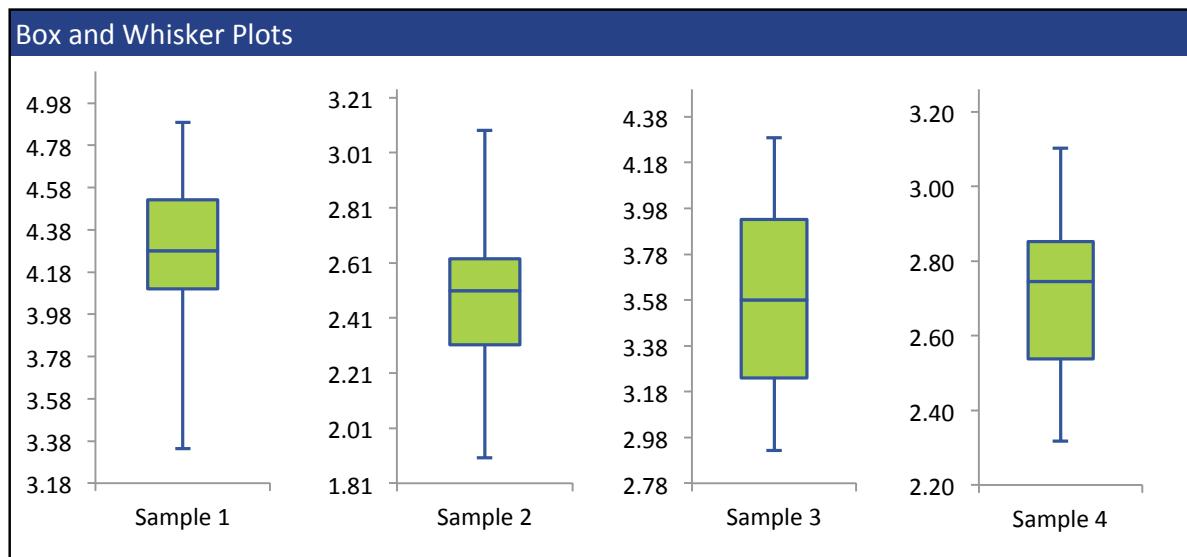
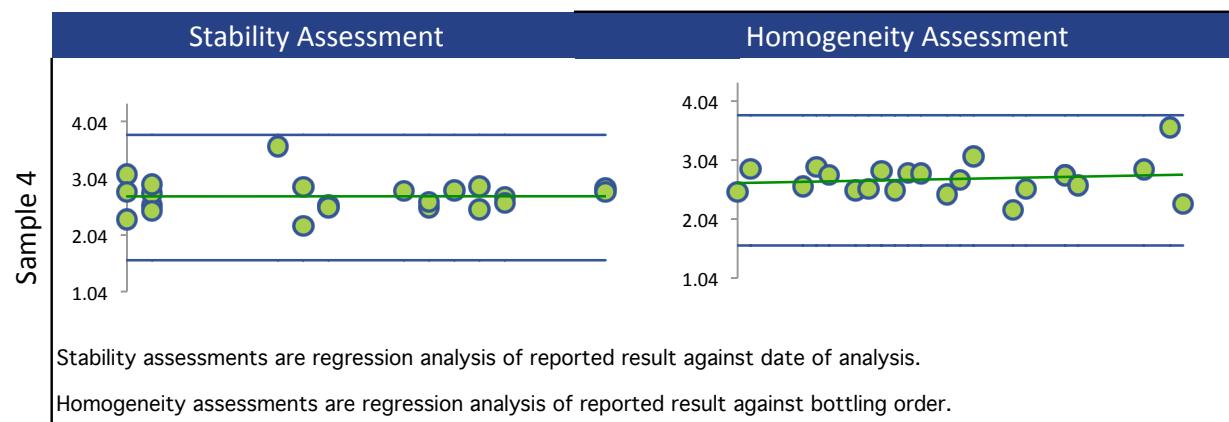
Annex A Summary by Analyte

ATRAZINE



Annex A Summary by Analyte

ATRAZINE



AZINPHOS-METHYL

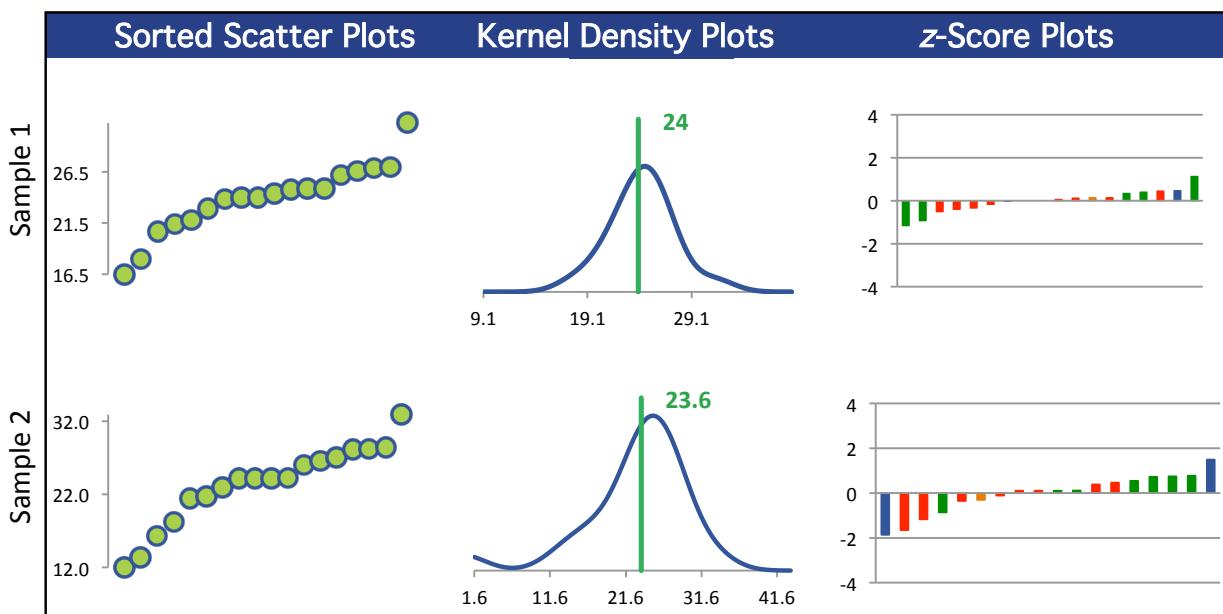
Summary Statistics

Statistic	C22-1	C22-2	C22-3	C22-4
N	18	18	18	18
Median $\mu\text{g/L}$	24.2	24.2	9.65	15.0
Robust Mean $\mu\text{g/L}$	24.0	23.6	9.67	14.9
U $\mu\text{g/L}$	0.860	1.57	0.489	0.595
Robust Standard Deviation $\mu\text{g/L}$	2.92	5.33	1.66	2.02
Regression Standard Deviation $\mu\text{g/L}$	6.36	6.25	2.77	4.07
Stability Flag				
Homogeneity Flag	Homogeneity			
Standard Deviation Used (SDPA) $\mu\text{g/L}$	6.44	6.25	2.77	4.07
Outliers	0	0	0	0
$ z > 3.0$	0	0	0	1
$2 < z < 3$	0	0	0	0

Methods Used

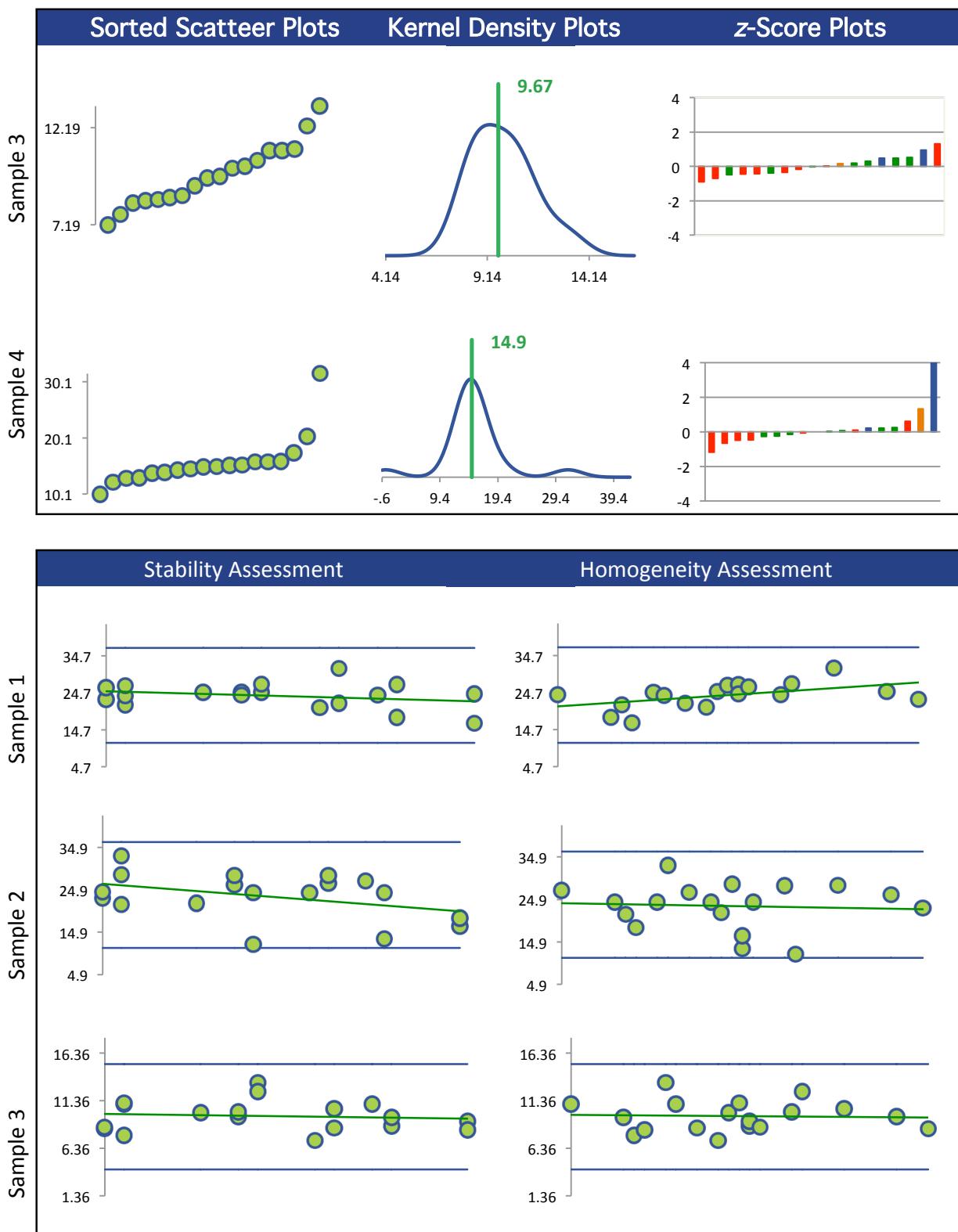
Method	C22-1	C22-2	C22-3	C22-4
GC/MS (Blue)	7	7	7	7
LC-MS/MS (Red)	8	8	8	8
GC/MS/MS (Green)	2	2	2	2
HPLC (Orange)	1	1	1	1

All summary stats and the plots below are based on the data excluding any flagged outliers



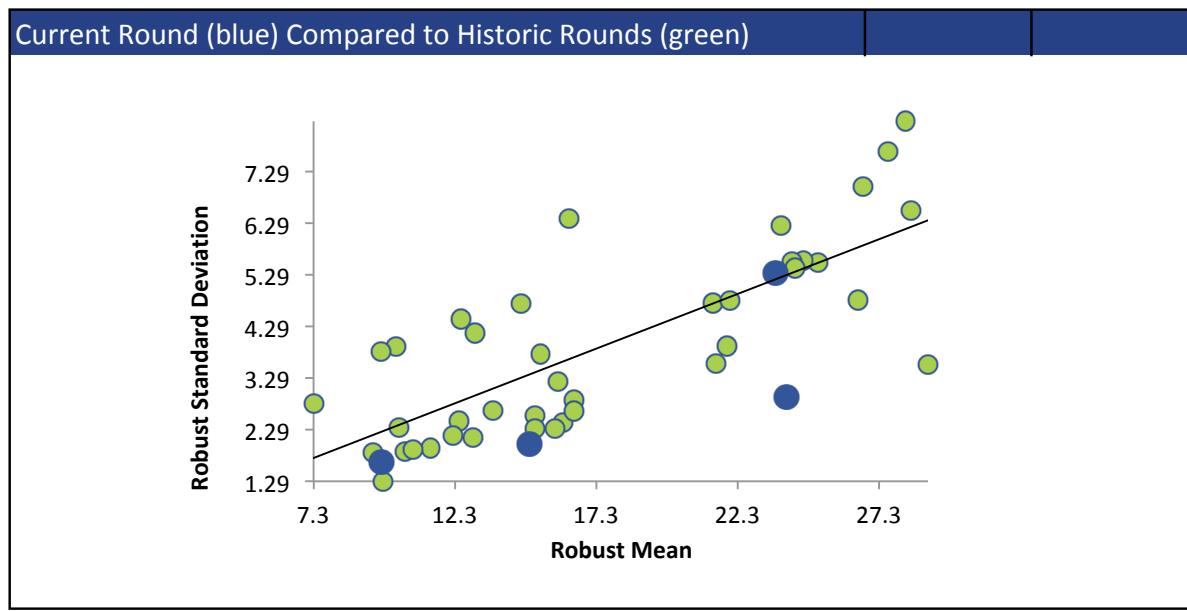
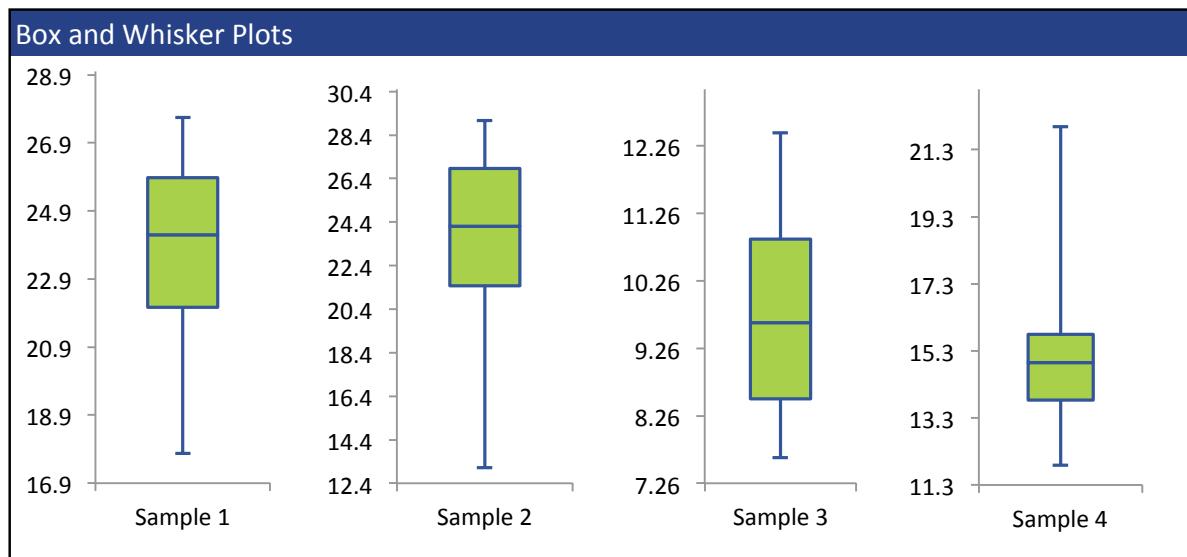
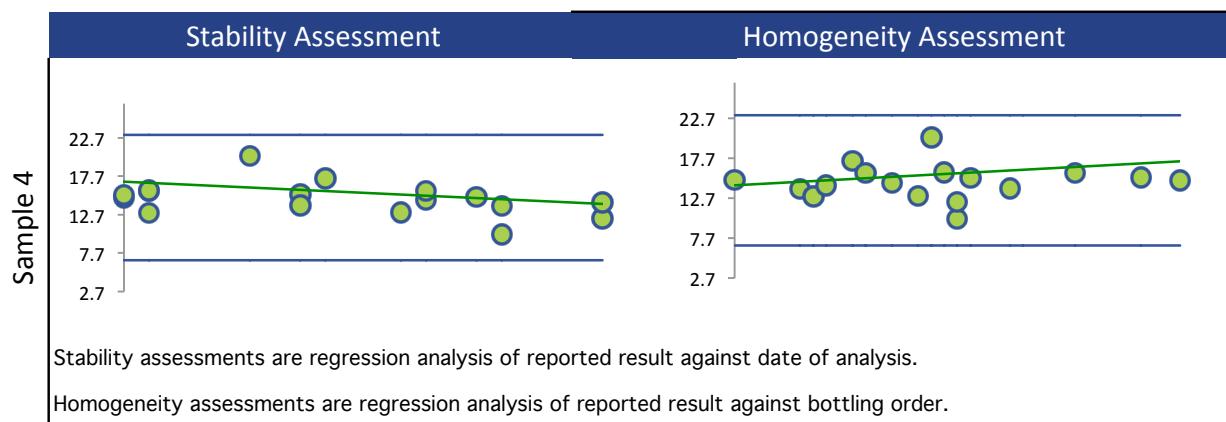
Annex A Summary by Analyte

AZINPHOS-METHYL



Annex A Summary by Analyte

AZINPHOS-METHYL



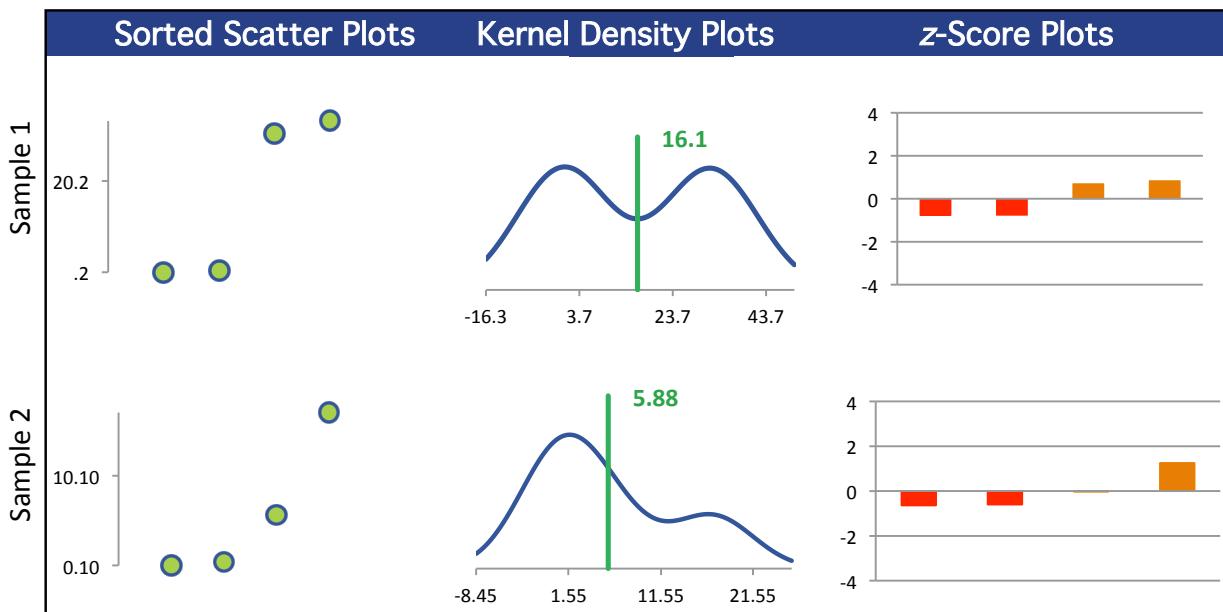
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Summary Statistics	Excluded	Excluded	Excluded	Excluded
Statistic	C22-1	C22-2	C22-3	C22-4
N	4	4	3	4
Median $\mu\text{g/L}$	15.5	3.10	2.96	4.76
Robust Mean $\mu\text{g/L}$	16.1	5.88	10.3	6.10
U $\mu\text{g/L}$	12.9	5.65	12.2	5.04
Robust Standard Deviation $\mu\text{g/L}$	20.6	9.04	16.9	8.07
Regression Standard Deviation $\mu\text{g/L}$	4.18	1.91	2.89	1.96
Stability Flag				
Homogeneity Flag				
Standard Deviation Used (SDPA) $\mu\text{g/L}$	20.6	9.04	16.9	8.07
Outliers	0	0	0	0
$ z > 3.0$	0	0	0	0
$2 < z < 3$	0	0	0	0

Methods Used

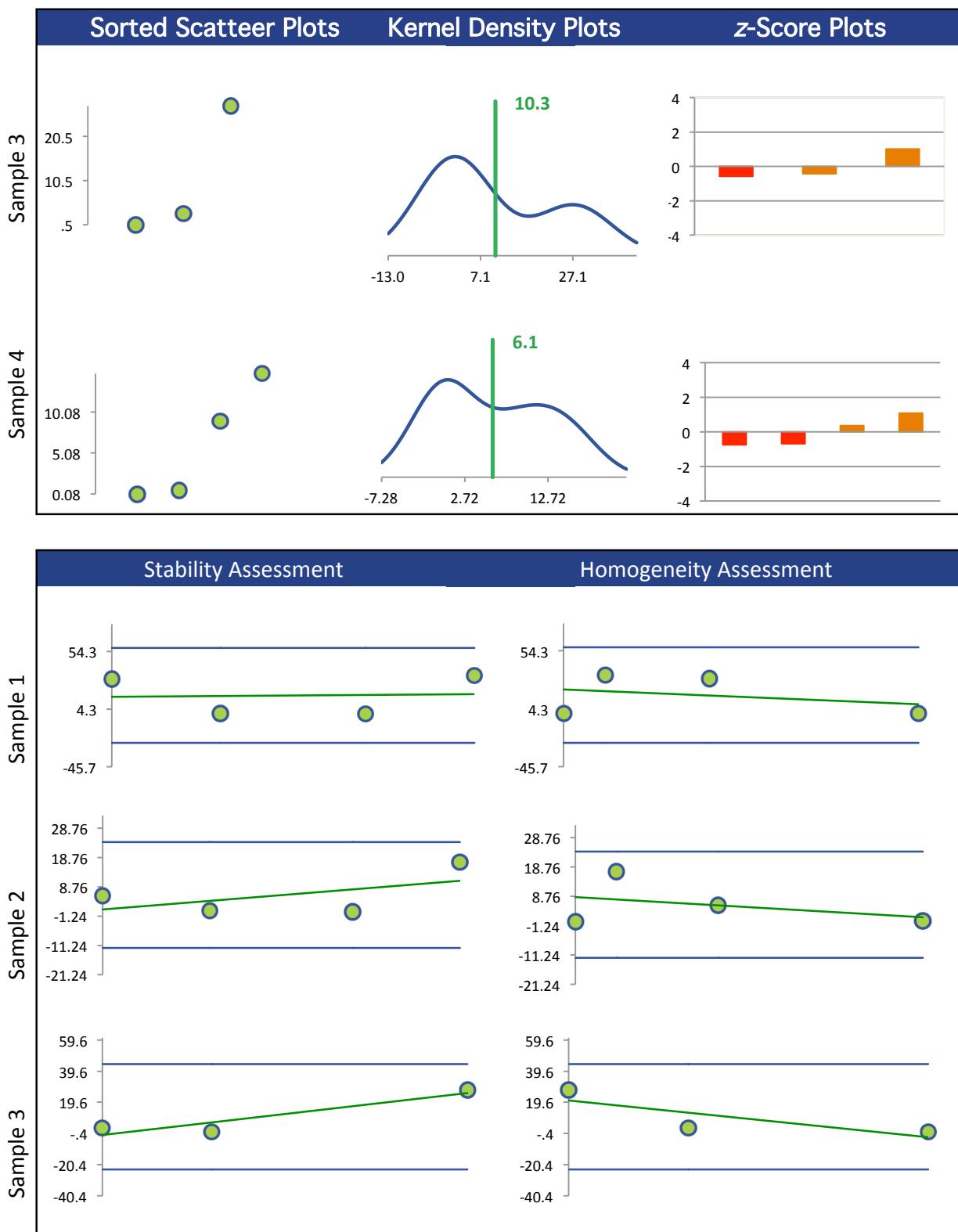
Method	C22-1	C22-2	C22-3	C22-4
LC-MS/MS (Blue)	2	2	1	2
HPLC (Red)	2	2	2	2

All summary stats and the plots below are based on the data excluding any flagged outliers



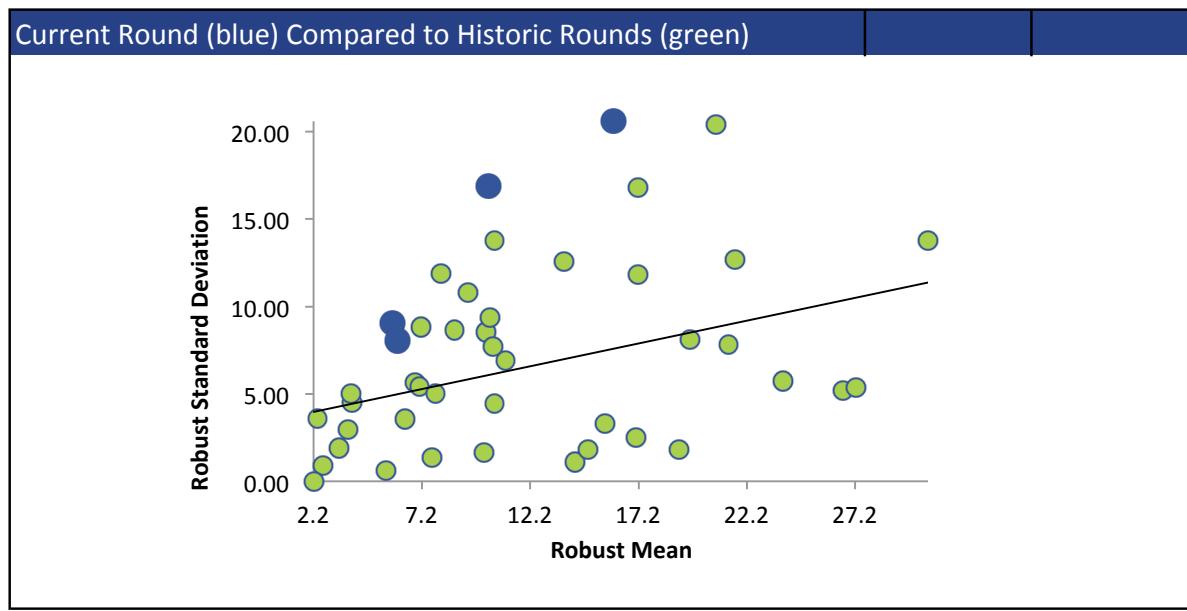
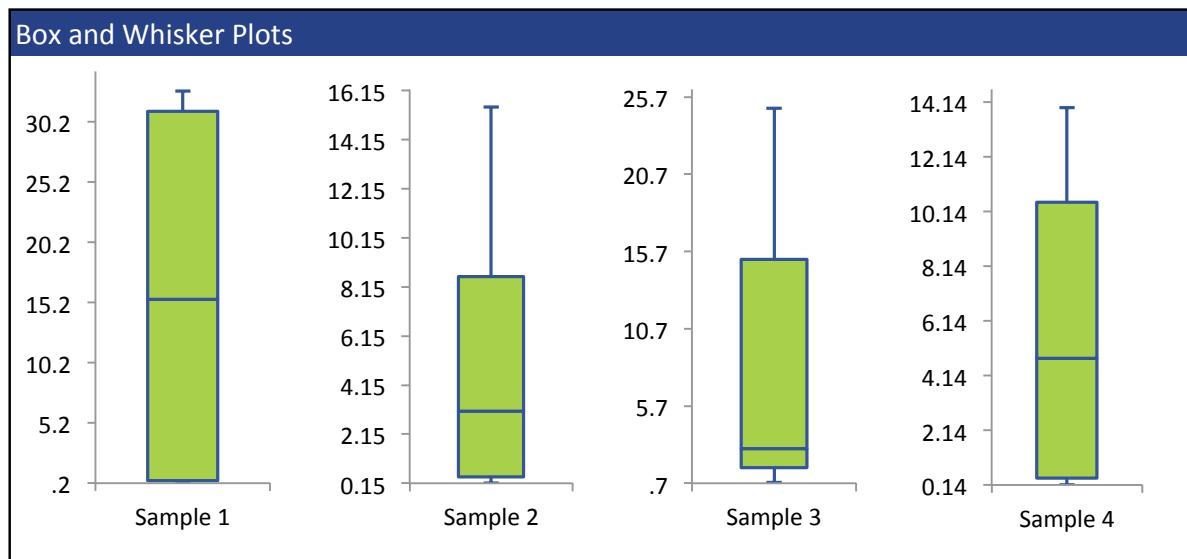
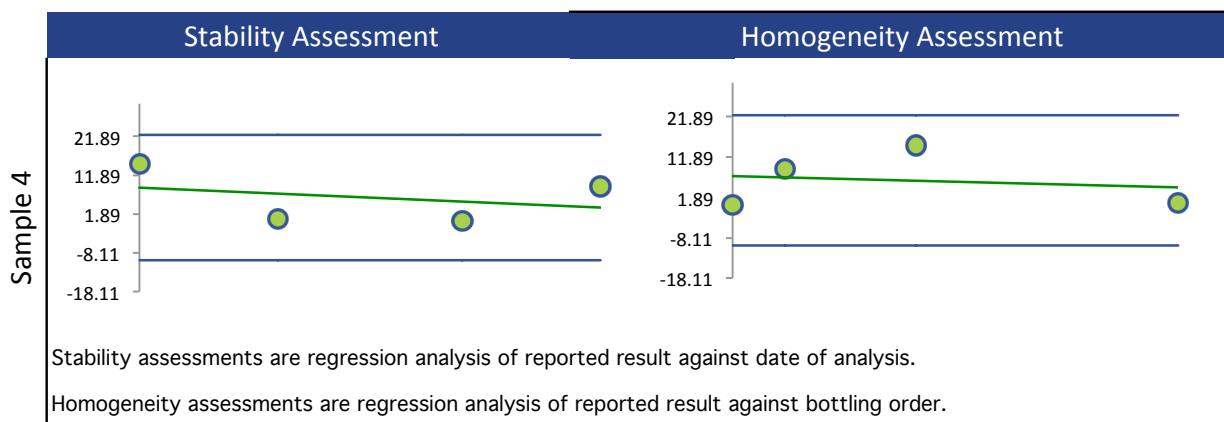
Annex A Summary by Analyte

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Annex A Summary by Analyte

BENDIOCARB



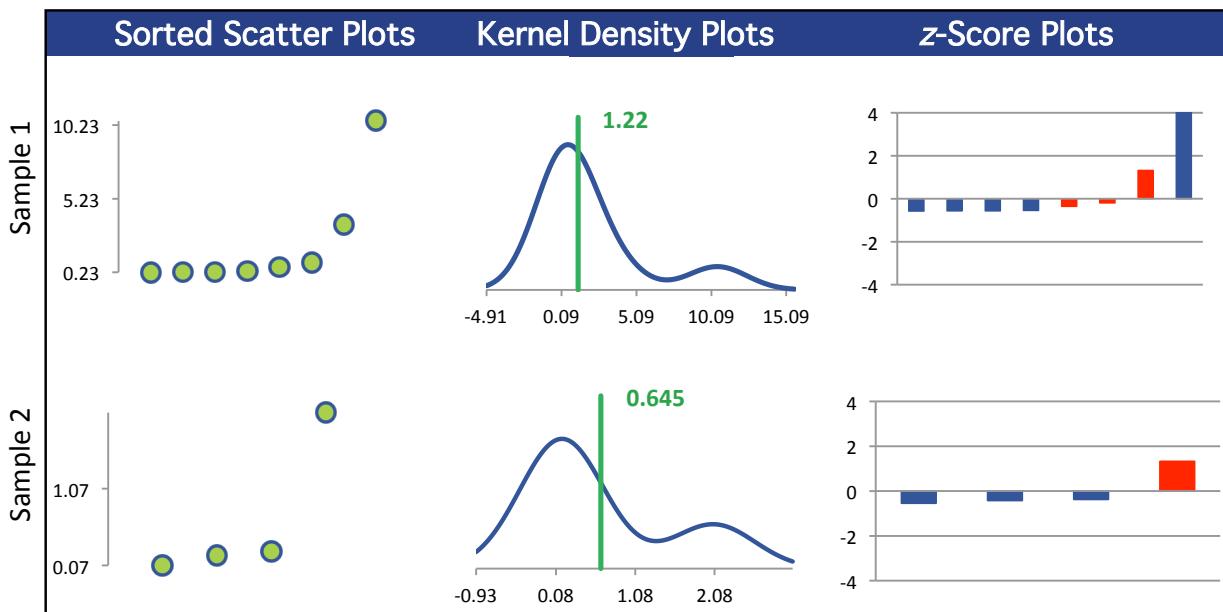
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Summary Statistics	Excluded	Excluded	Excluded	Excluded
Statistic	C22-1	C22-2	C22-3	C22-4
N	8	4	5	0
Median $\mu\text{g/L}$	0.444	0.226	0.400	
Robust Mean $\mu\text{g/L}$	1.22	0.645	1.95	
U $\mu\text{g/L}$	0.756	0.669	1.77	
Robust Standard Deviation $\mu\text{g/L}$	1.71	1.07	3.17	
Regression Standard Deviation $\mu\text{g/L}$	0.439	0.267	0.660	
Stability Flag				
Homogeneity Flag				
Standard Deviation Used (SDPA) $\mu\text{g/L}$	1.71	1.07	3.17	
Outliers	0	0	0	2
$ z > 3.0$	1	0	0	0
$2 < z < 3$	0	0	1	0

Methods Used

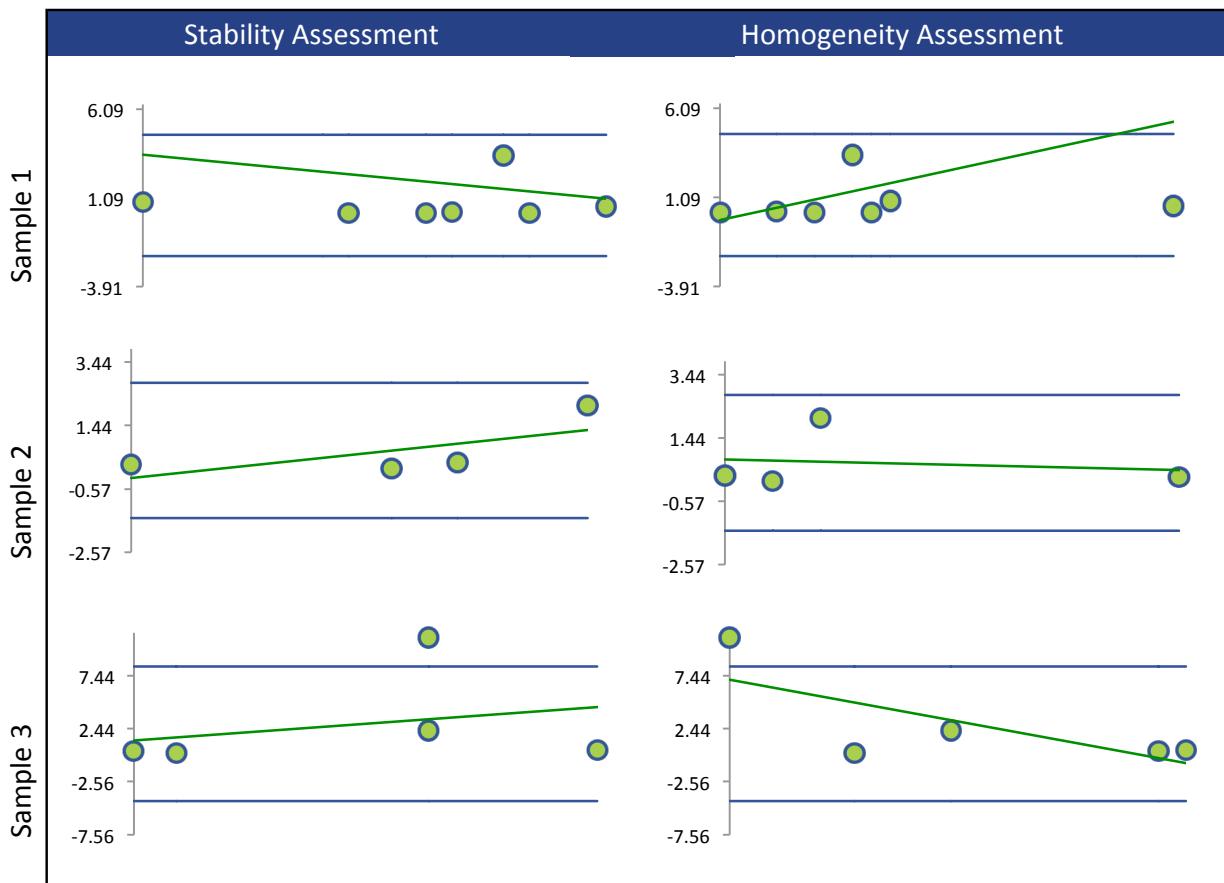
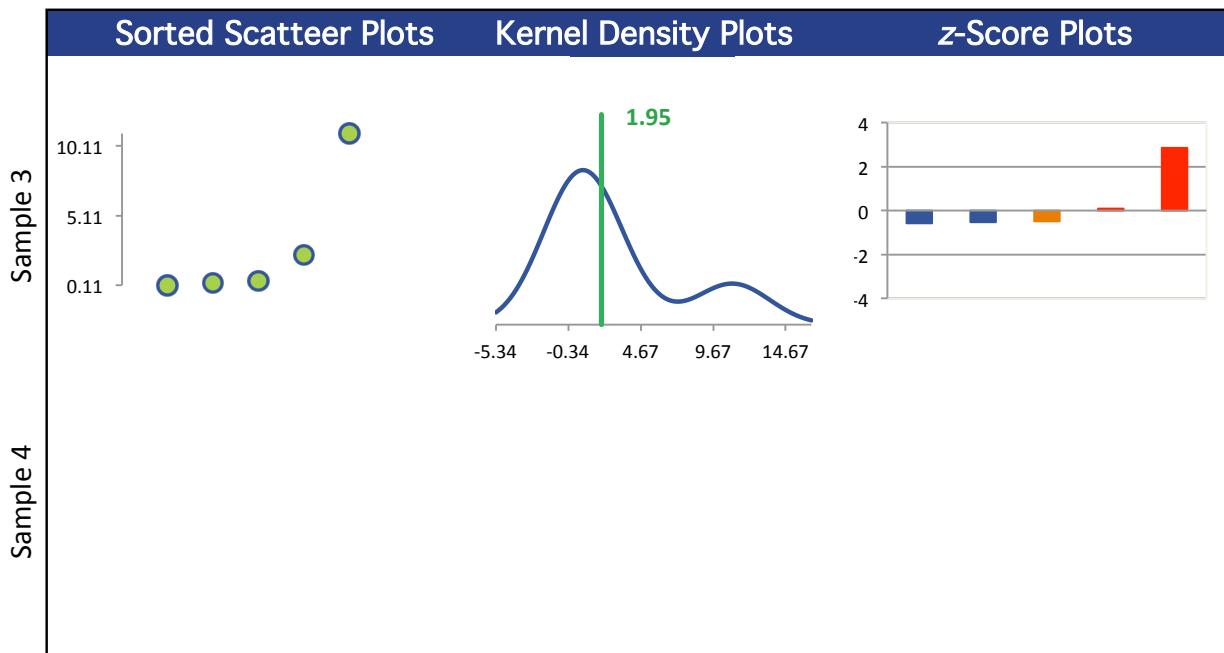
Method	C22-1	C22-2	C22-3	C22-4
LC-MS/MS (Blue)	5	3	2	0
GC/MS (Red)	2	0	2	0
HPLC (Green)	1	1	1	0

All summary stats and the plots below are based on the data excluding any flagged outliers



Annex A Summary by Analyte

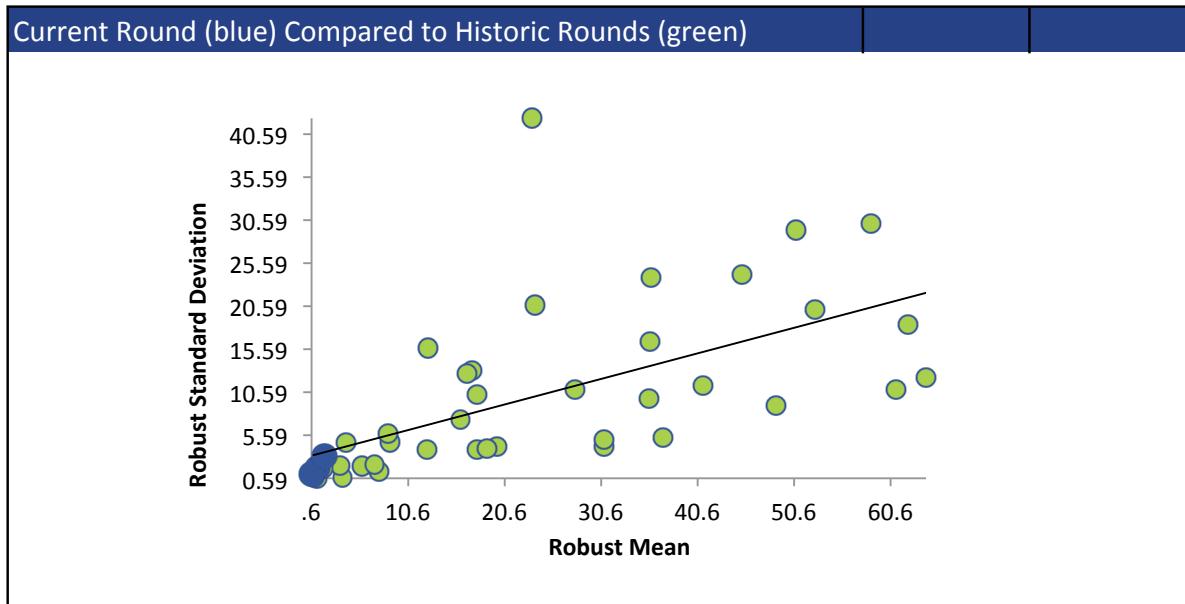
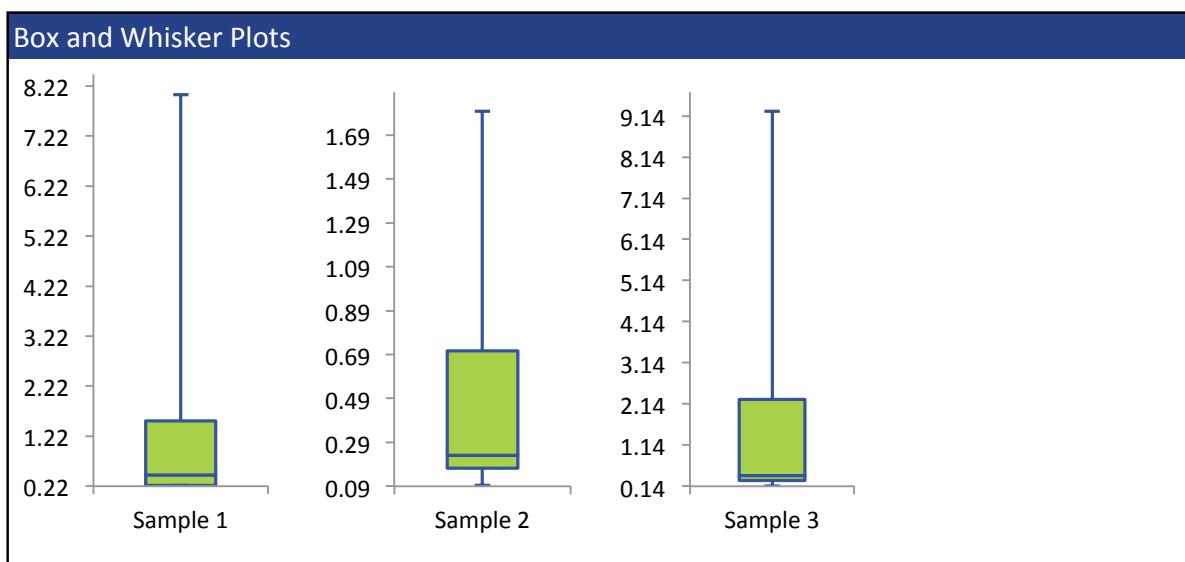
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Annex A Summary by Analyte

CARBARYL

	Stability Assessment	Homogeneity Assessment
Sample 4	Stability assessments are regression analysis of reported result against date of analysis.	Homogeneity assessments are regression analysis of reported result against bottling order.



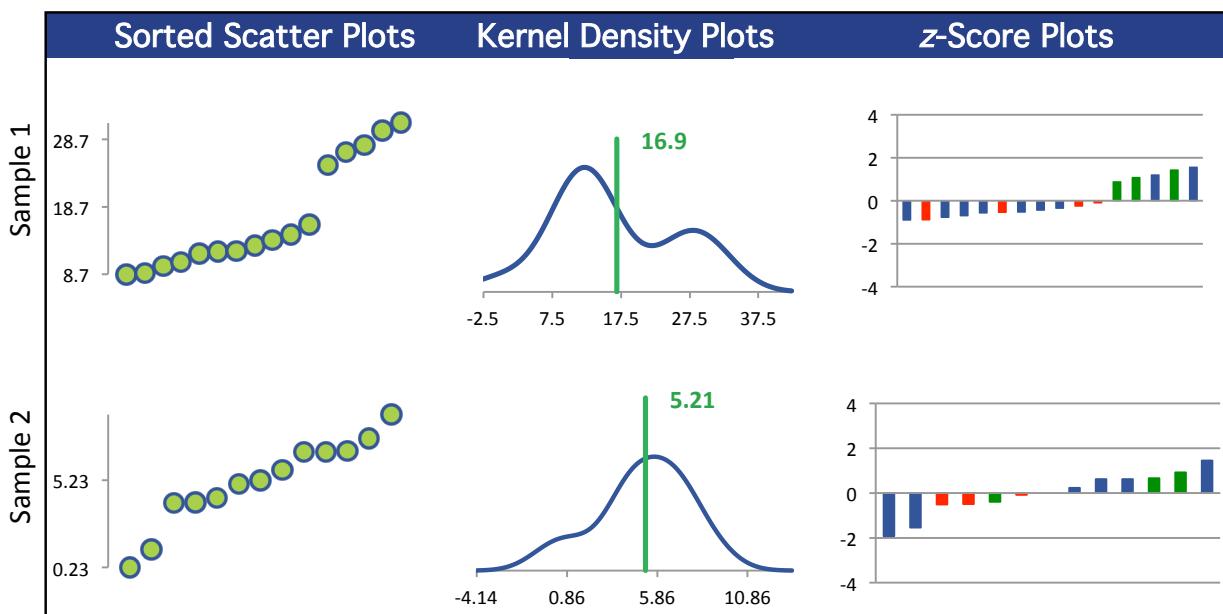
CARBOFURAN**Summary Statistics**

Statistic	C22-1	C22-2	C22-3	C22-4
N	16	13	12	14
Median $\mu\text{g/L}$	13.3	5.20	1.90	7.78
Robust Mean $\mu\text{g/L}$	16.9	5.21	1.85	9.16
$\text{U } \mu\text{g/L}$	2.85	0.898	0.451	2.32
Robust Standard Deviation $\mu\text{g/L}$	9.13	2.59	1.25	6.93
Regression Standard Deviation $\mu\text{g/L}$	5.24	1.61	0.571	2.83
Stability Flag				
Homogeneity Flag				
Standard Deviation Used (SDPA) $\mu\text{g/L}$	9.13	2.59	1.25	6.93
Outliers	0	0	0	0
$ \text{z} > 3.0$	0	0	0	0
$2 < \text{z} < 3$	0	0	1	0

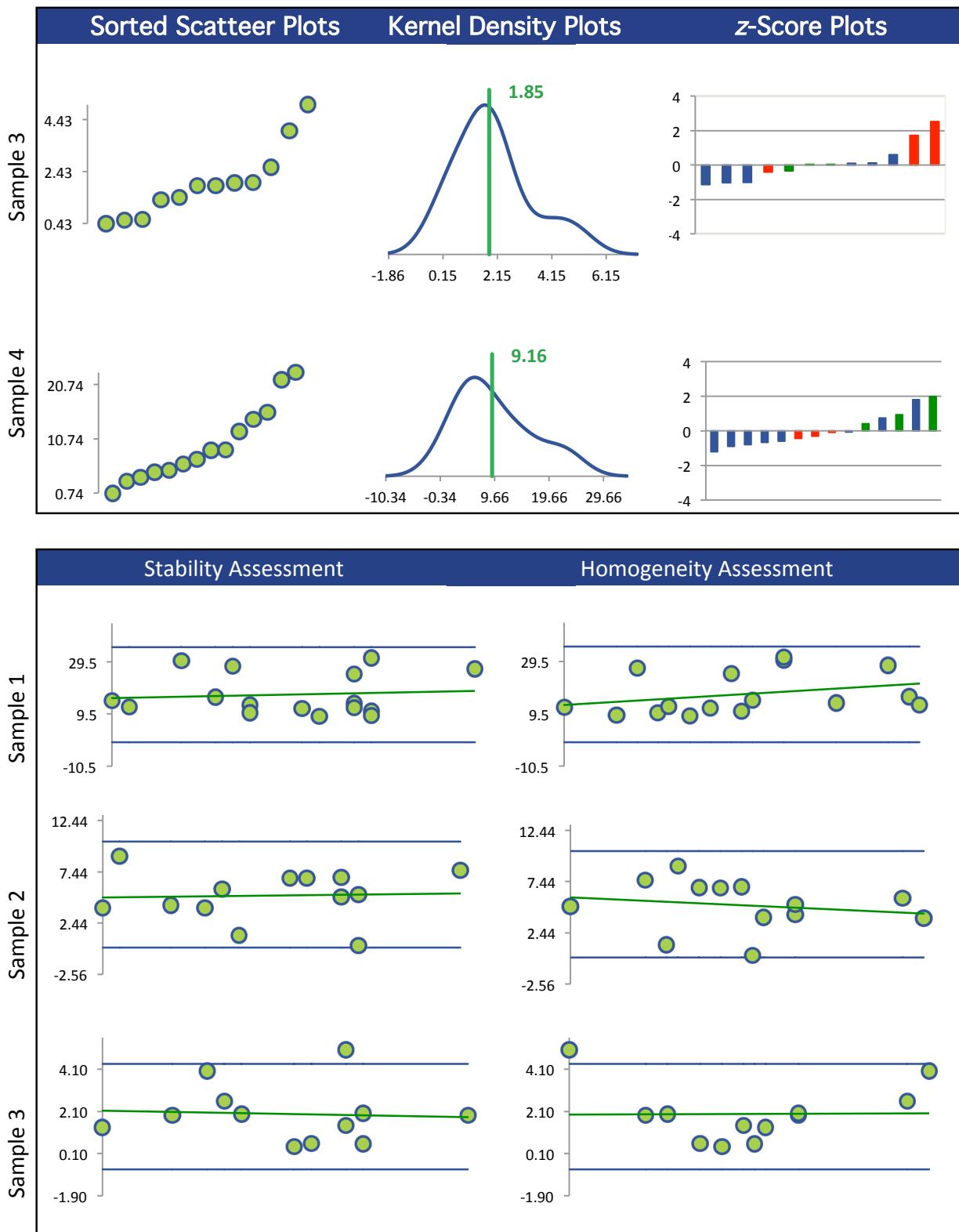
Methods Used

Method	C22-1	C22-2	C22-3	C22-4
LC-MS/MS (Blue)	9	7	6	8
GC/MS (Red)	4	3	3	3
HPLC (Green)	3	3	3	3

All summary stats and the plots below are based on the data excluding any flagged outliers

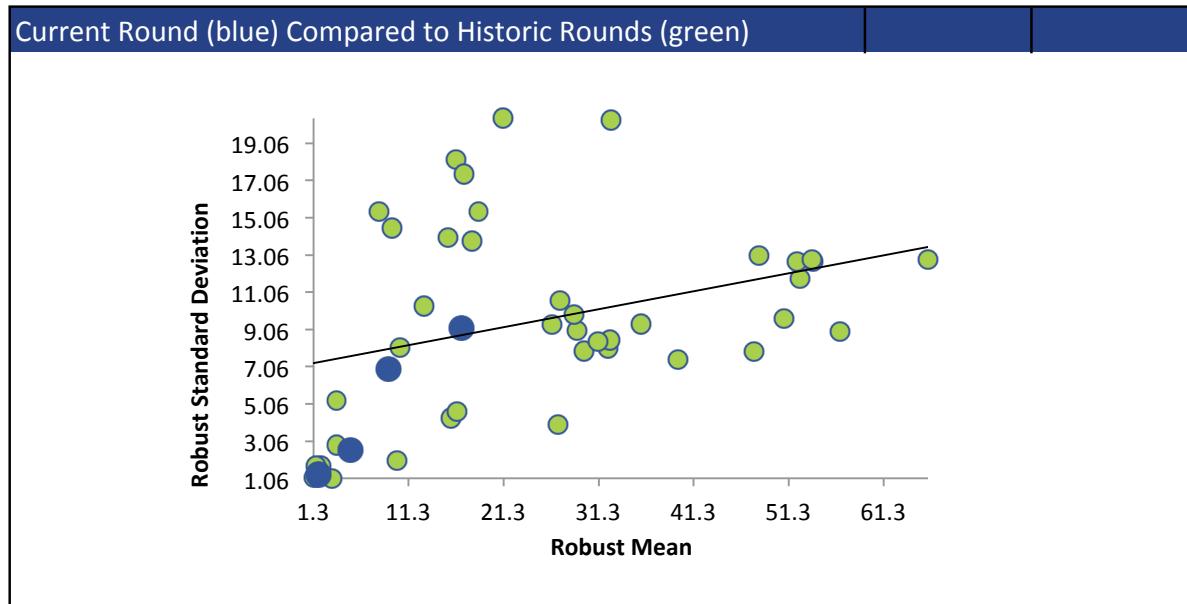
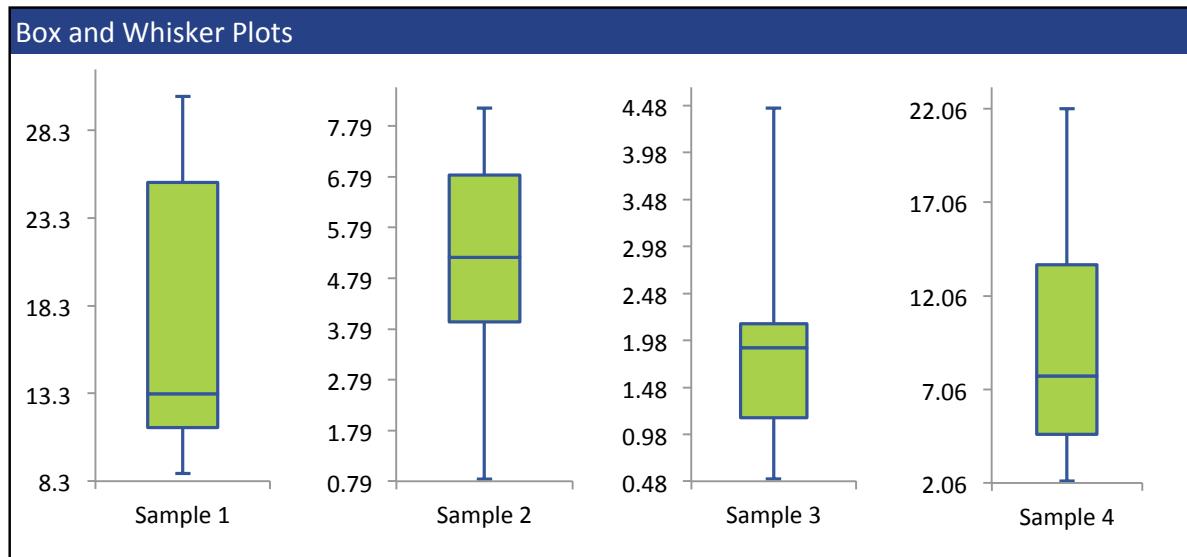
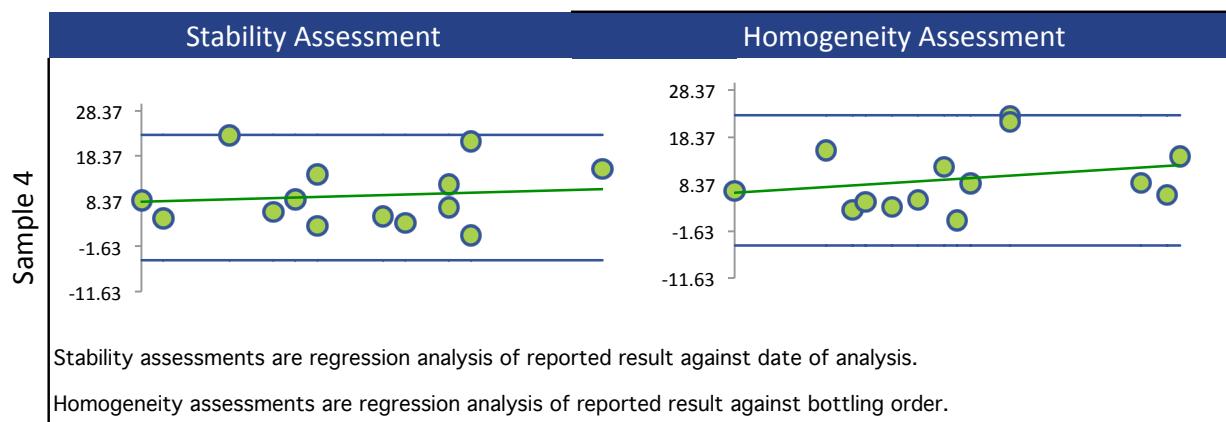


CARBOFURAN



Annex A Summary by Analyte

CARBOFURAN



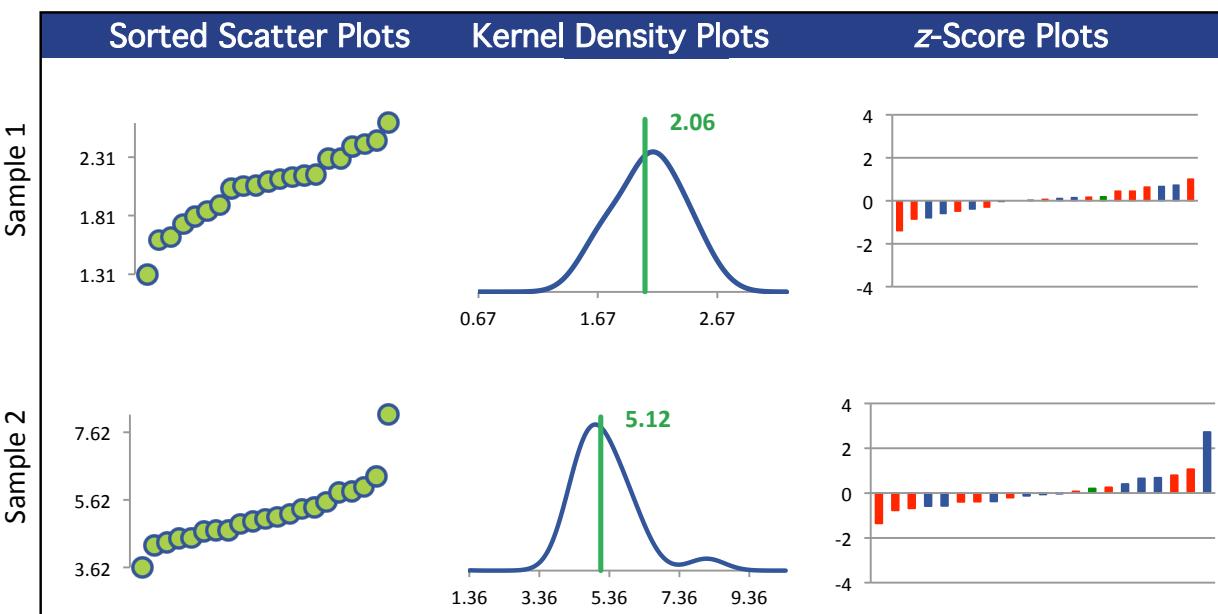
CHLORPYRIFOS ETHYL**Summary Statistics**

Statistic	C22-1	C22-2	C22-3	C22-4
N	21	21	21	21
Median $\mu\text{g/L}$	2.10	5.05	7.58	3.76
Robust Mean $\mu\text{g/L}$	2.06	5.12	7.49	3.87
U $\mu\text{g/L}$	0.0889	0.203	0.281	0.125
Robust Standard Deviation $\mu\text{g/L}$	0.326	0.745	1.03	0.458
Regression Standard Deviation $\mu\text{g/L}$	0.535	1.11	1.55	0.874
Stability Flag				
Homogeneity Flag				
Standard Deviation Used (SDPA) $\mu\text{g/L}$	0.535	1.11	1.55	0.874
Outliers	0	0	0	0
$ z > 3.0$	0	0	0	1
$2 < z < 3$	0	1	0	0

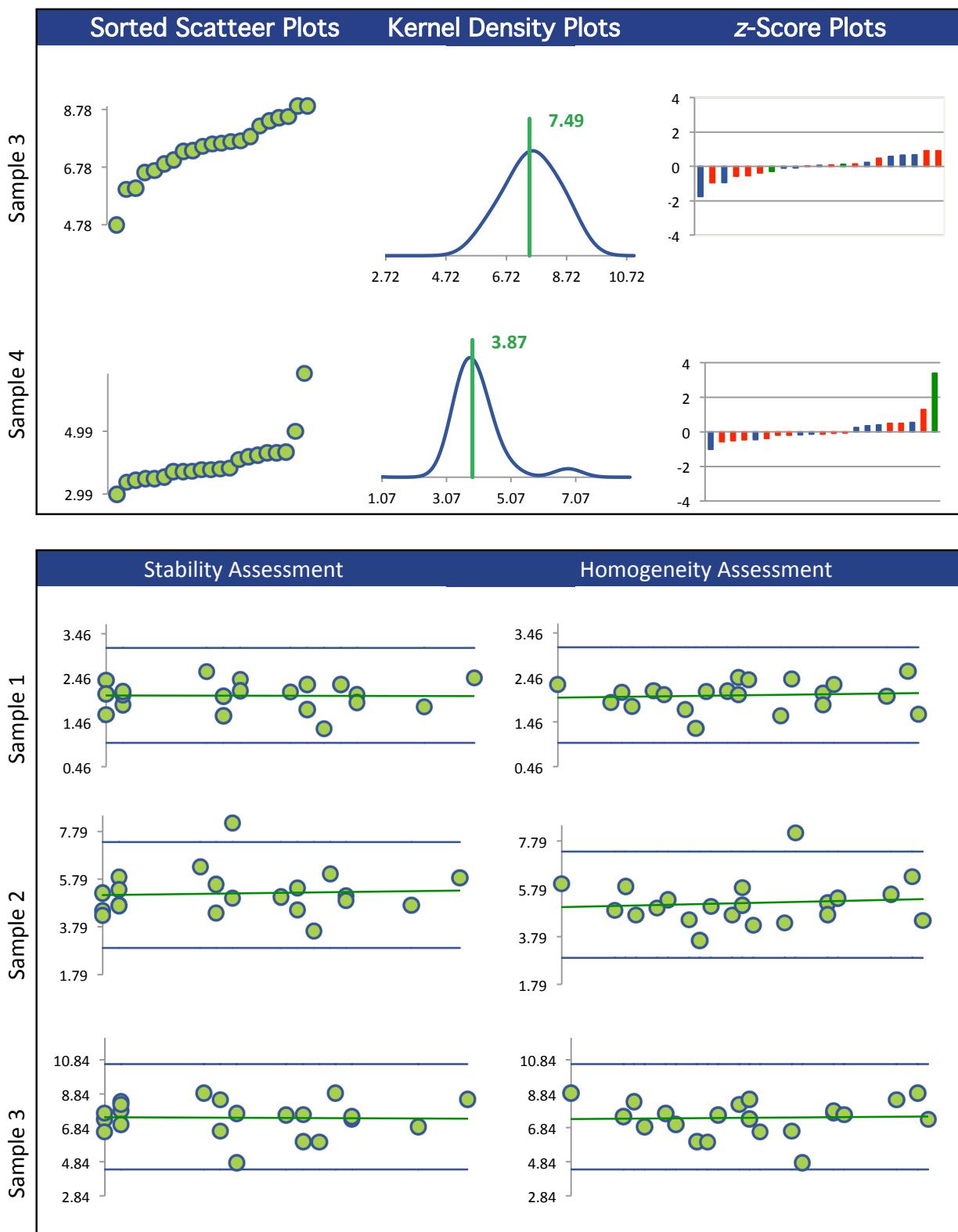
Methods Used

Method	C22-1	C22-2	C22-3	C22-4
GC/MS (Blue)	10	10	10	10
LC-MS/MS (Red)	8	8	8	8
GC/MS/MS (Green)	3	3	3	3

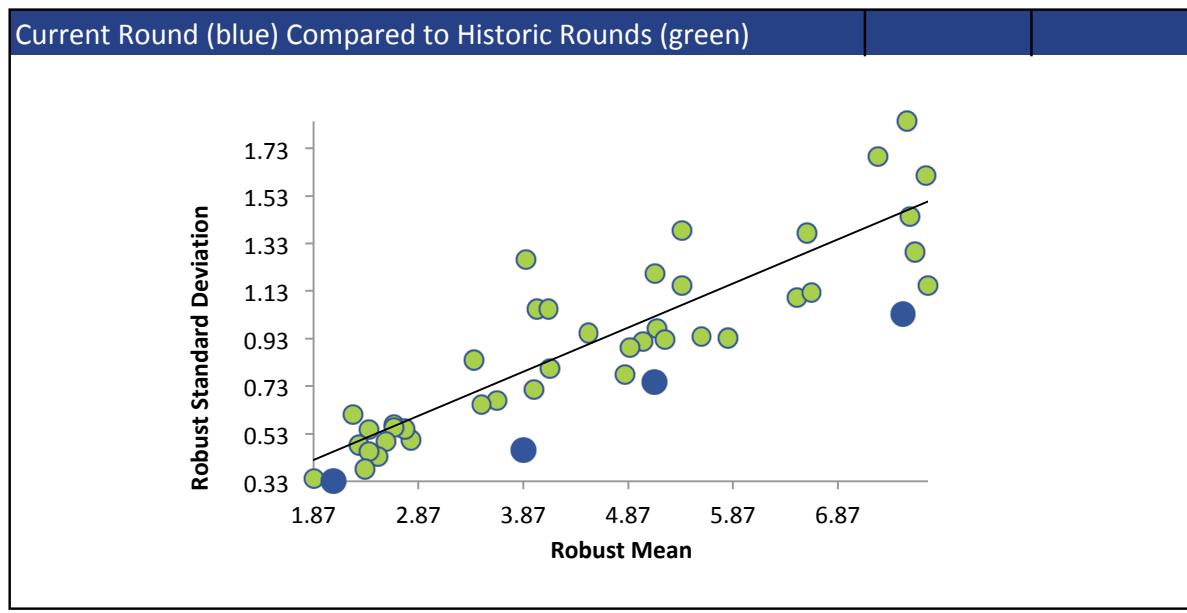
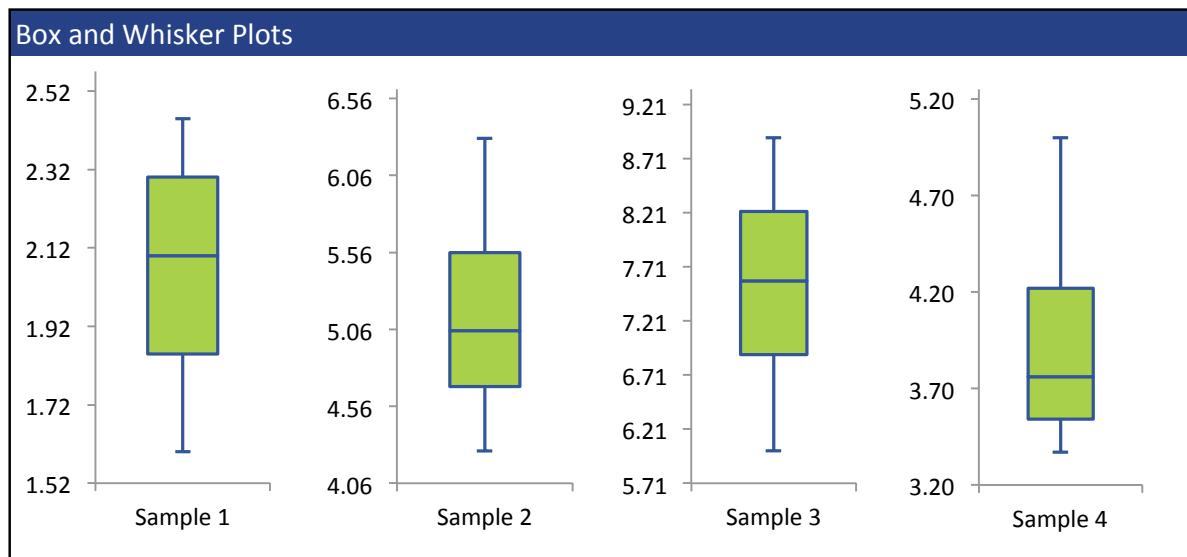
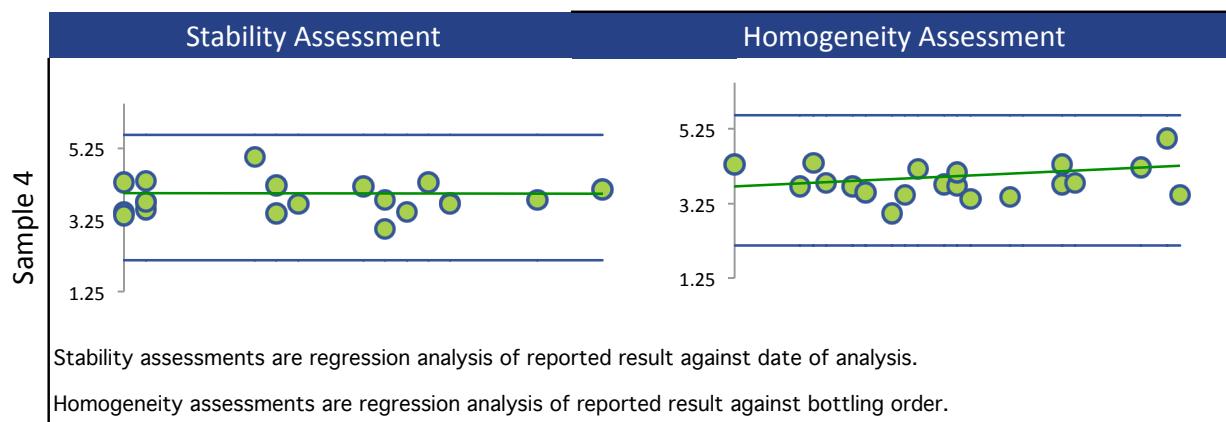
All summary stats and the plots below are based on the data excluding any flagged outliers



CHLORPYRIFOS ETHYL



CHLORPYRIFOS ETHYL



CYANAZINE

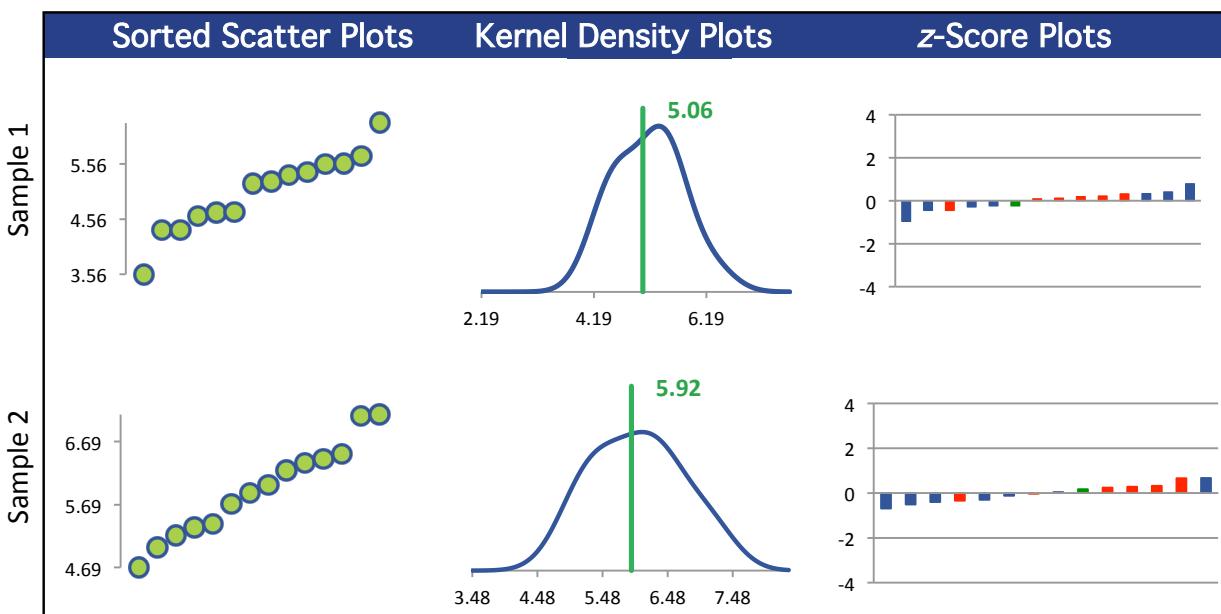
Summary Statistics

Statistic	C22-1	C22-2	C22-3	C22-4
N	14	14	14	14
Median $\mu\text{g/L}$	5.22	5.94	8.14	2.93
Robust Mean $\mu\text{g/L}$	5.06	5.92	8.02	2.89
$\text{U } \mu\text{g/L}$	0.230	0.283	0.384	0.0802
Robust Standard Deviation $\mu\text{g/L}$	0.687	0.848	1.15	0.240
Regression Standard Deviation $\mu\text{g/L}$	1.56	1.79	2.37	0.958
Stability Flag				
Homogeneity Flag				
Standard Deviation Used (SDPA) $\mu\text{g/L}$	1.56	1.79	2.37	0.958
Outliers	0	0	0	0
$ \text{z} > 3.0$	0	0	0	0
$2 < \text{z} < 3$	0	0	0	0

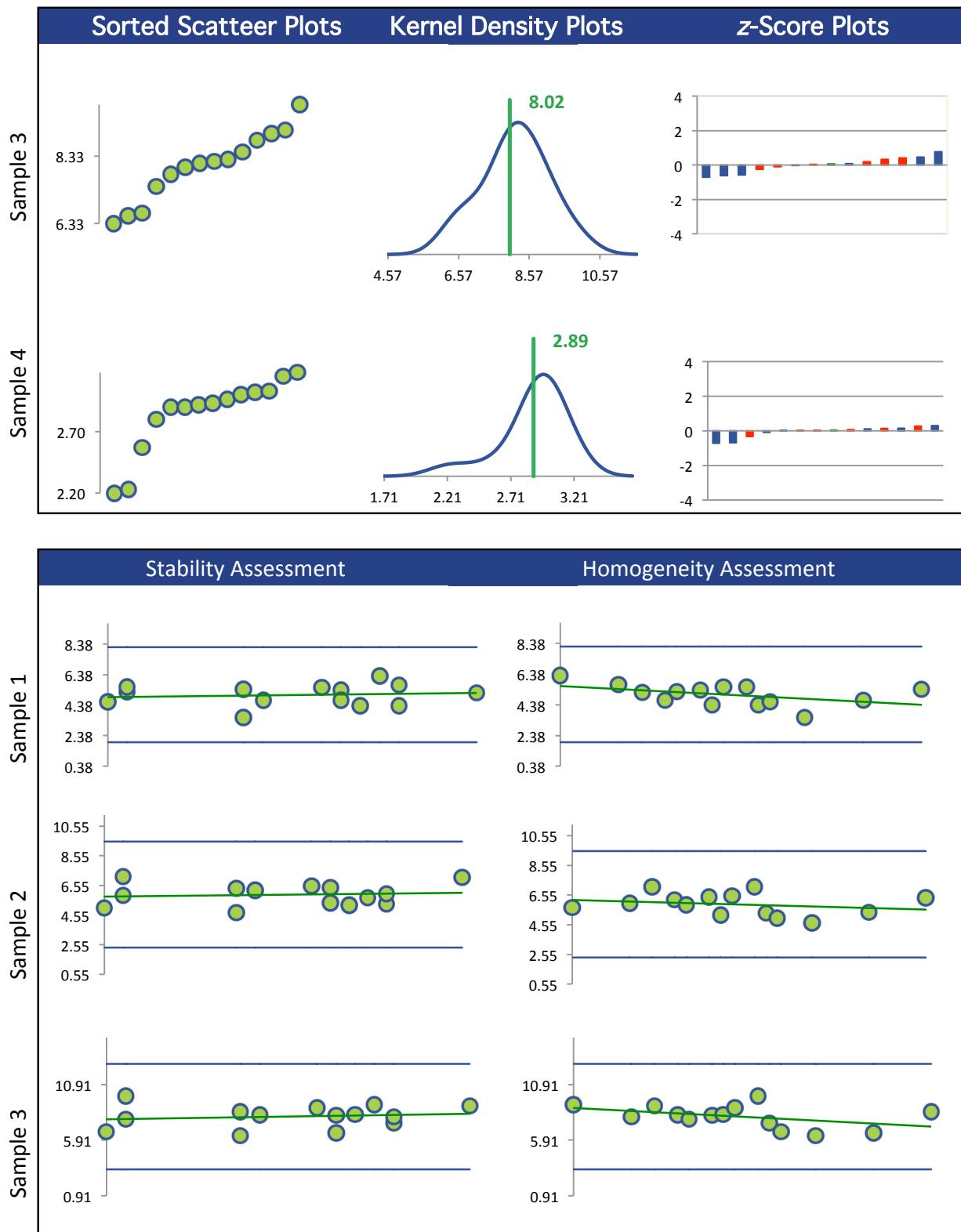
Methods Used

Method	C22-1	C22-2	C22-3	C22-4
GC/MS (Blue)	7	7	7	7
LC-MS/MS (Red)	5	5	5	5
GC/MS/MS (Green)	1	1	1	1
HPLC (Orange)	1	1	1	1

All summary stats and the plots below are based on the data excluding any flagged outliers

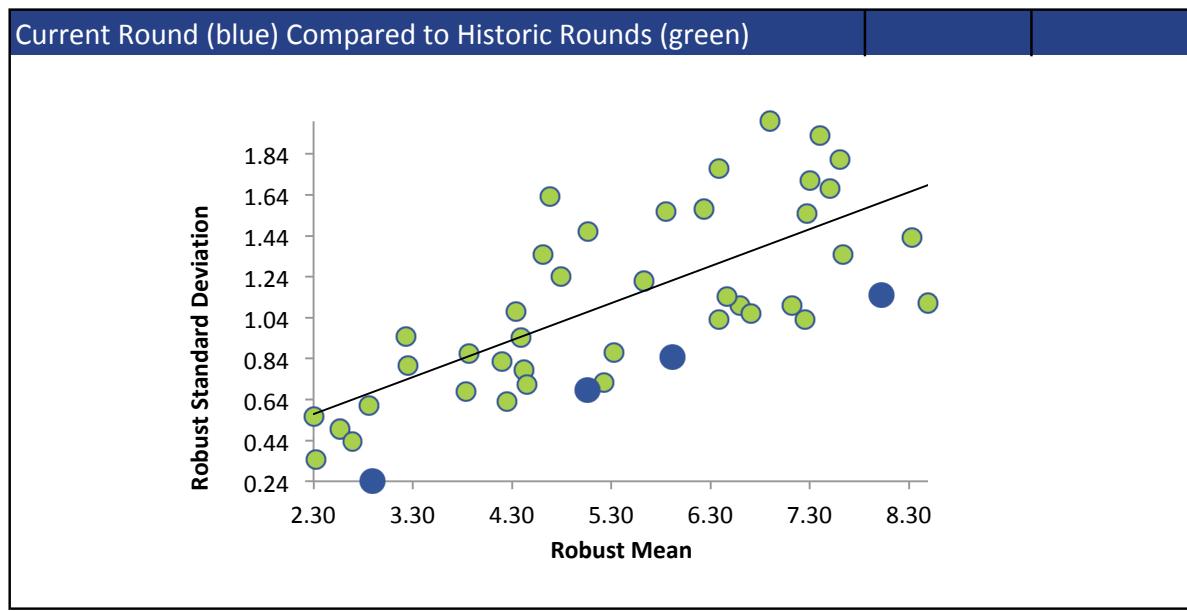
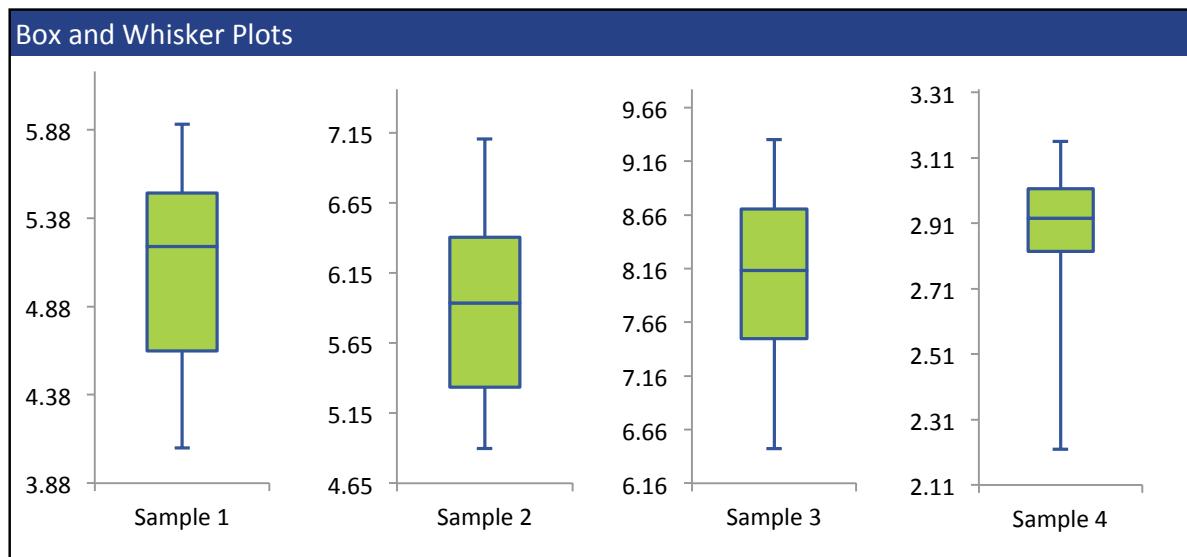
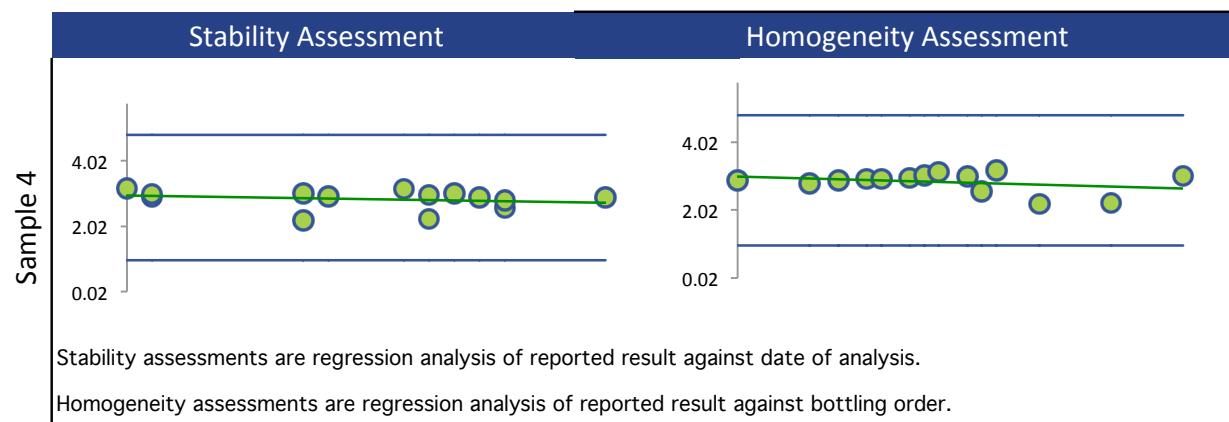


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Annex A Summary by Analyte

CYANAZINE



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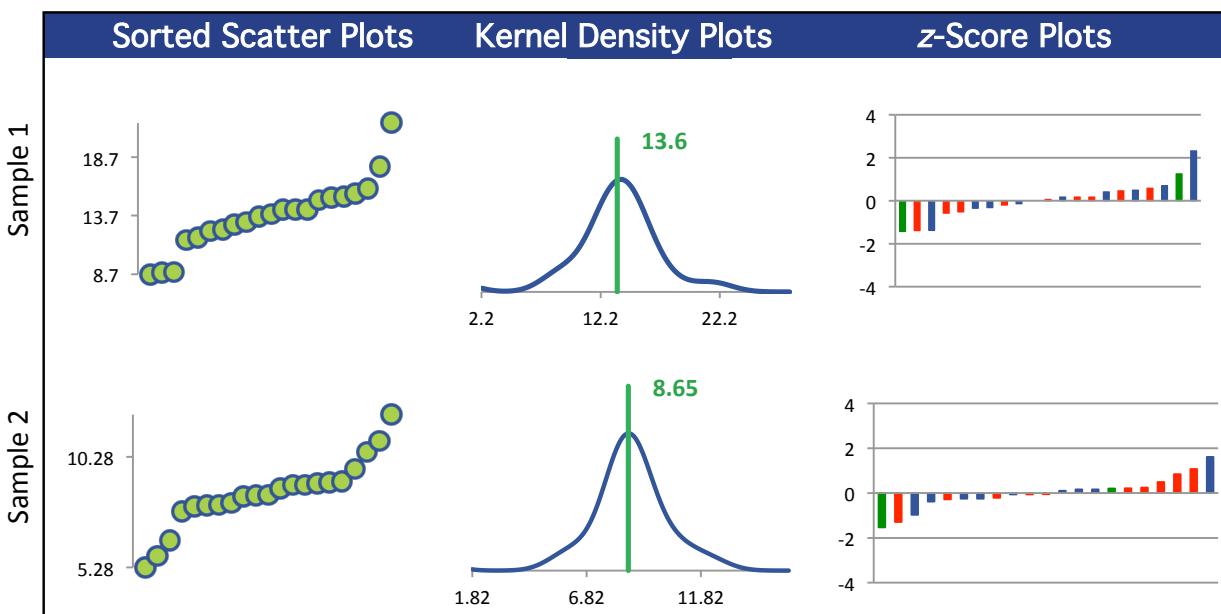
Summary Statistics

Statistic	C22-1	C22-2	C22-3	C22-4
N	21	21	21	19
Median $\mu\text{g/L}$	13.8	8.55	7.65	1.30
Robust Mean $\mu\text{g/L}$	13.6	8.65	7.44	1.30
U $\mu\text{g/L}$	0.706	0.360	0.352	0.0665
Robust Standard Deviation $\mu\text{g/L}$	2.59	1.32	1.29	0.232
Regression Standard Deviation $\mu\text{g/L}$	3.45	2.20	1.89	0.331
Stability Flag				
Homogeneity Flag				Homogeneity
Standard Deviation Used (SDPA) $\mu\text{g/L}$	3.45	2.20	1.89	0.363
Outliers	0	0	0	1
$ z > 3.0$	0	0	1	0
$2 < z < 3$	1	0	0	0

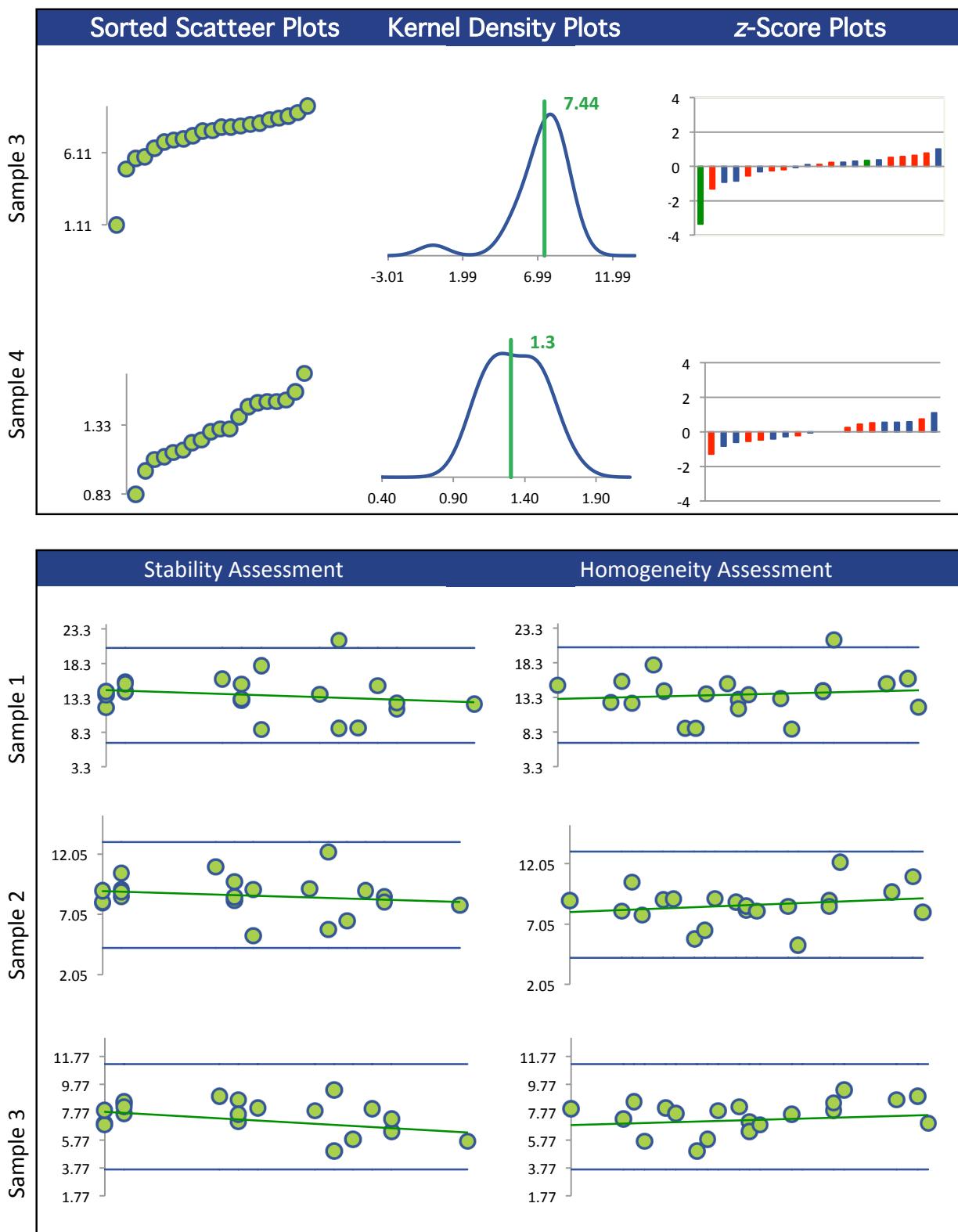
Methods Used

Method	C22-1	C22-2	C22-3	C22-4
GC/MS/MS (Blue)	2	2	2	1
LC-MS/MS (Red)	9	9	9	8
GC/MS (Green)	9	9	9	9
GC/ECD (Orange)	1	1	1	1

All summary stats and the plots below are based on the data excluding any flagged outliers

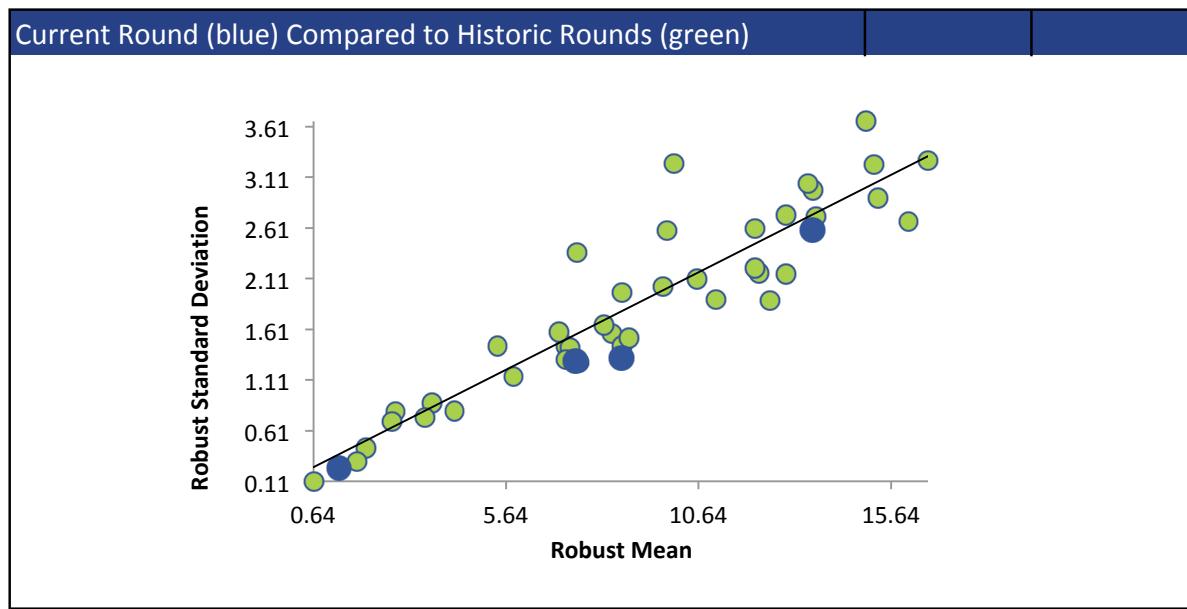
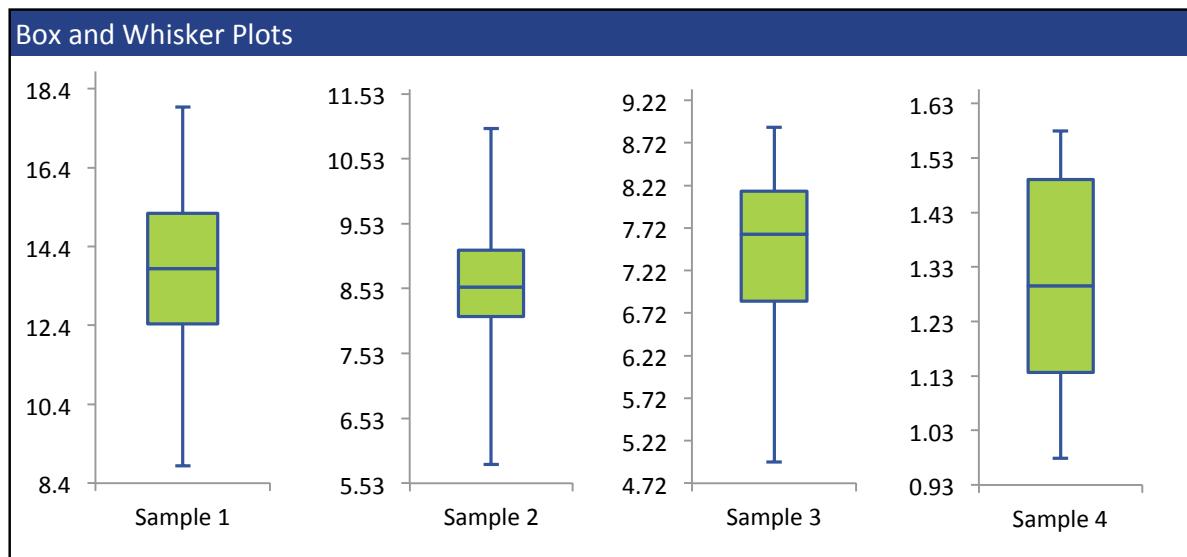
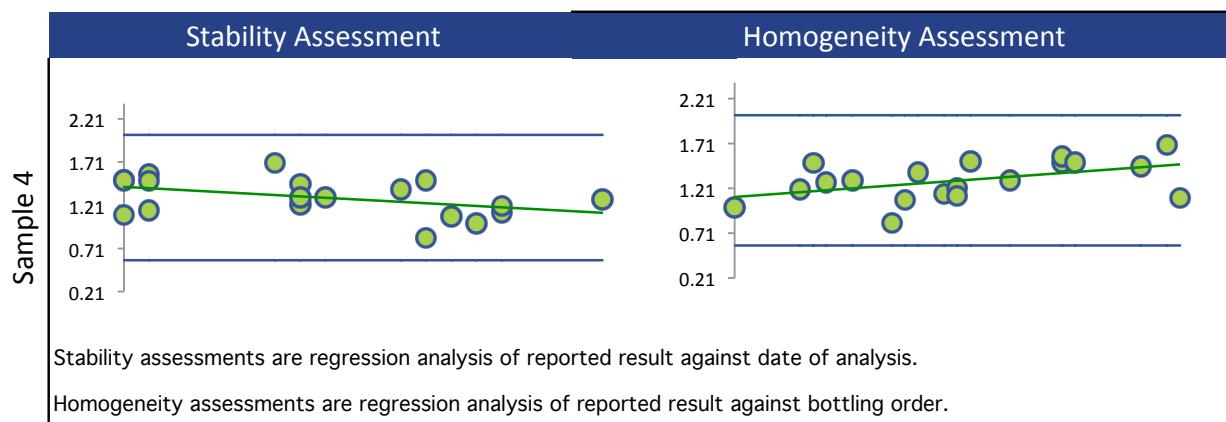


DIAZINON



Annex A Summary by Analyte

DIAZINON



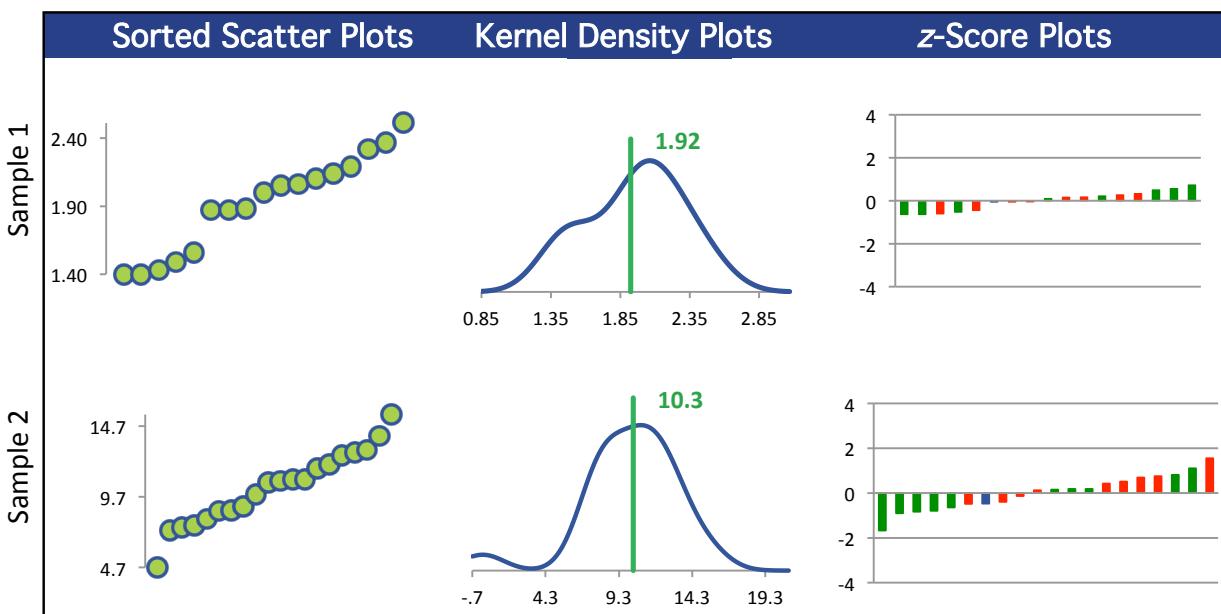
DIMETHOATE**Summary Statistics**

Statistic	C22-1	C22-2	C22-3	C22-4
N	17	20	20	20
Median $\mu\text{g/L}$	2.00	10.8	8.91	4.43
Robust Mean $\mu\text{g/L}$	1.92	10.3	8.56	4.16
U $\mu\text{g/L}$	0.122	0.749	0.805	0.313
Robust Standard Deviation $\mu\text{g/L}$	0.402	2.68	2.88	1.12
Regression Standard Deviation $\mu\text{g/L}$	0.807	3.35	2.81	1.48
Stability Flag				
Homogeneity Flag				
Standard Deviation Used (SDPA) $\mu\text{g/L}$	0.807	3.35	2.88	1.48
Outliers	0	0	0	0
$ z > 3.0$	0	0	0	1
$2 < z < 3$	0	0	0	0

Methods Used

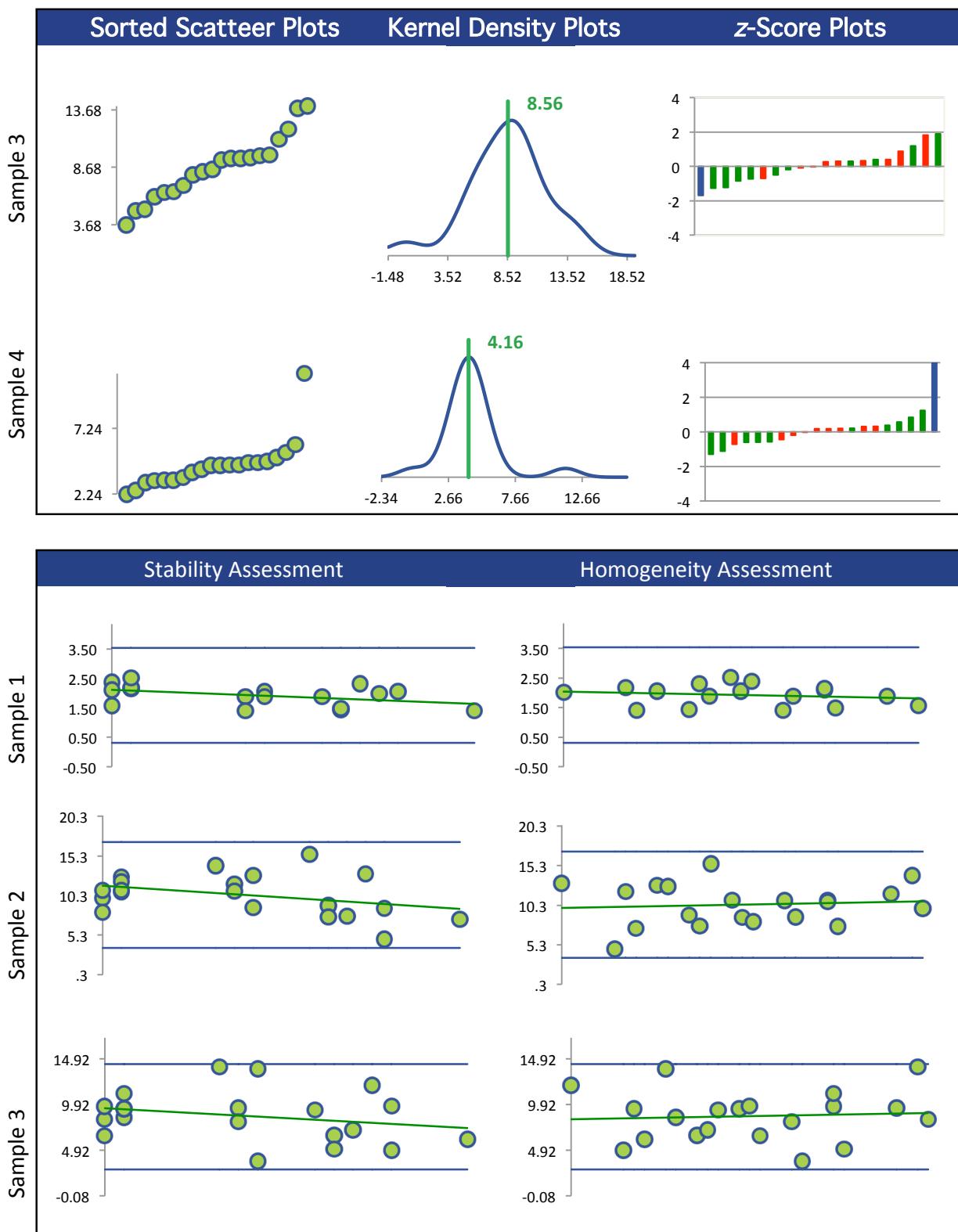
Method	C22-1	C22-2	C22-3	C22-4
GC/MS (Blue)	8	10	10	10
LC-MS/MS (Red)	8	9	9	9
GC/MS/MS (Green)	1	1	1	1

All summary stats and the plots below are based on the data excluding any flagged outliers



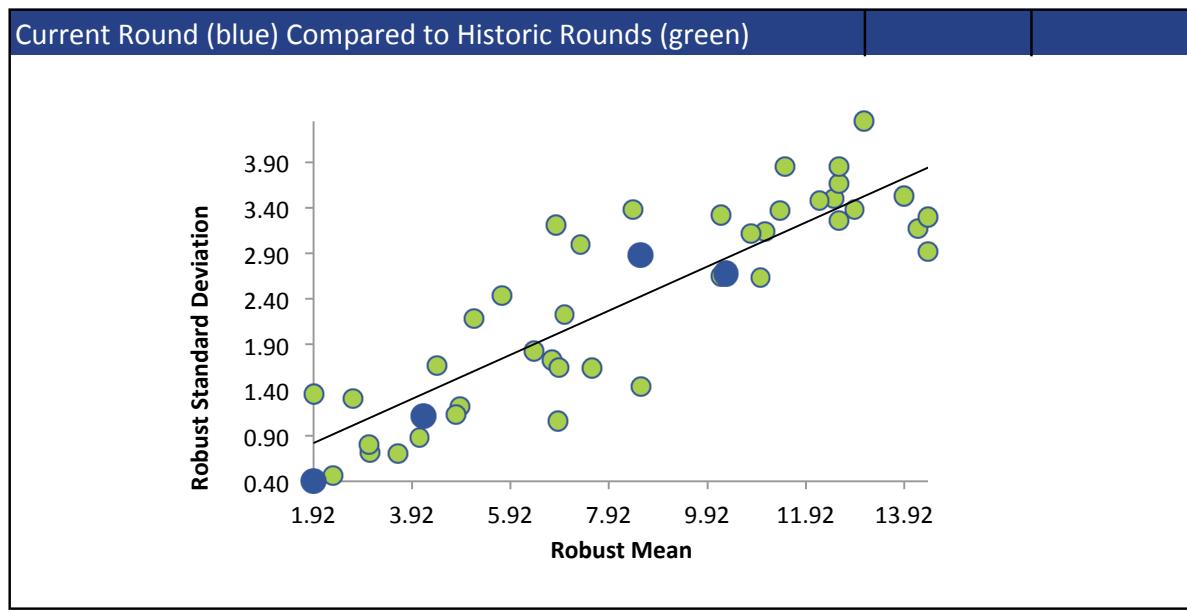
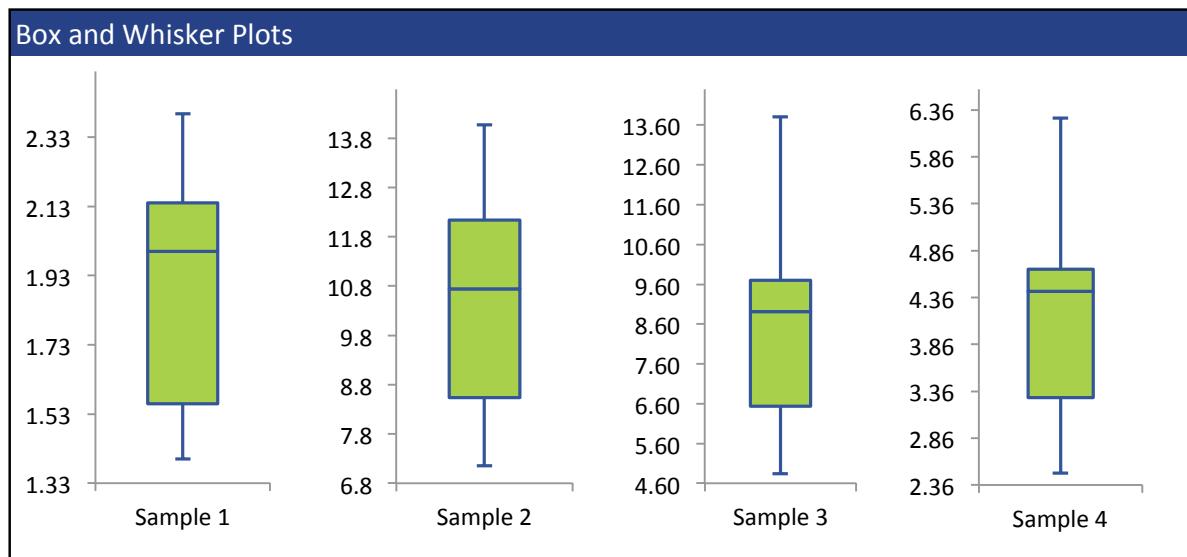
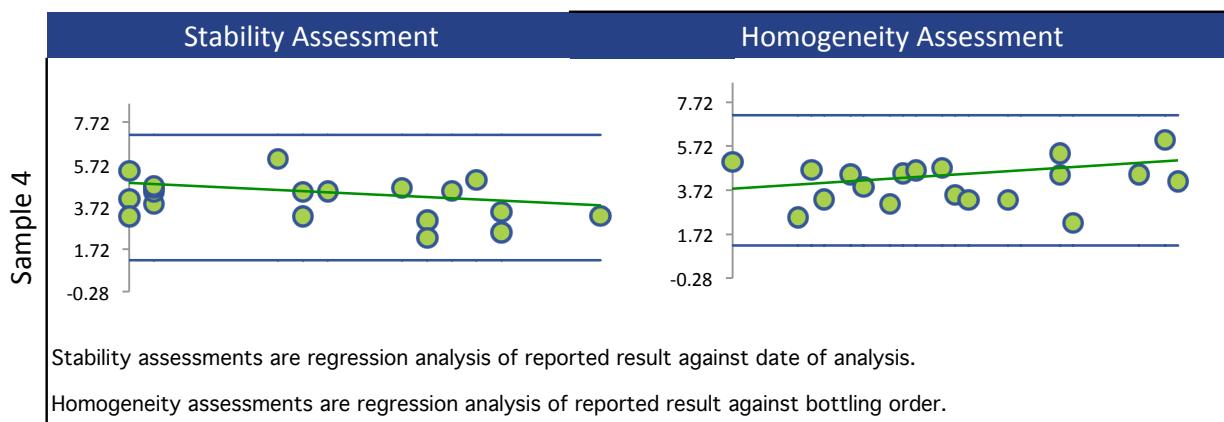
Annex A Summary by Analyte

DIMETHOATE



Annex A Summary by Analyte

DIMETHOATE



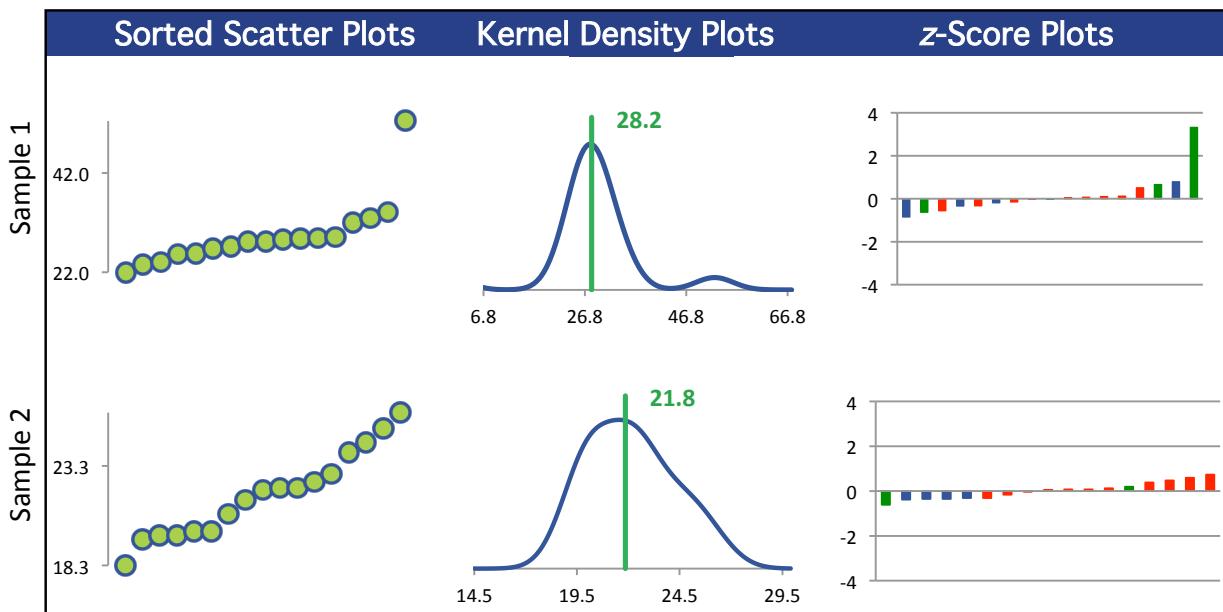
DIURON**Summary Statistics**

Statistic	C22-1	C22-2	C22-3	C22-4
N	17	17	17	17
Median $\mu\text{g/L}$	28.1	22.1	33.5	41.0
Robust Mean $\mu\text{g/L}$	28.2	21.8	33.2	41.1
$\text{U } \mu\text{g/L}$	1.20	0.725	1.22	1.36
Robust Standard Deviation $\mu\text{g/L}$	3.97	2.39	4.01	4.50
Regression Standard Deviation $\mu\text{g/L}$	7.30	5.78	8.48	10.4
Stability Flag				
Homogeneity Flag				
Standard Deviation Used (SDPA) $\mu\text{g/L}$	7.30	5.78	8.48	10.4
Outliers	0	0	0	0
$ \text{z} > 3.0$	1	0	0	0
$2 < \text{z} < 3$	0	0	0	0

Methods Used

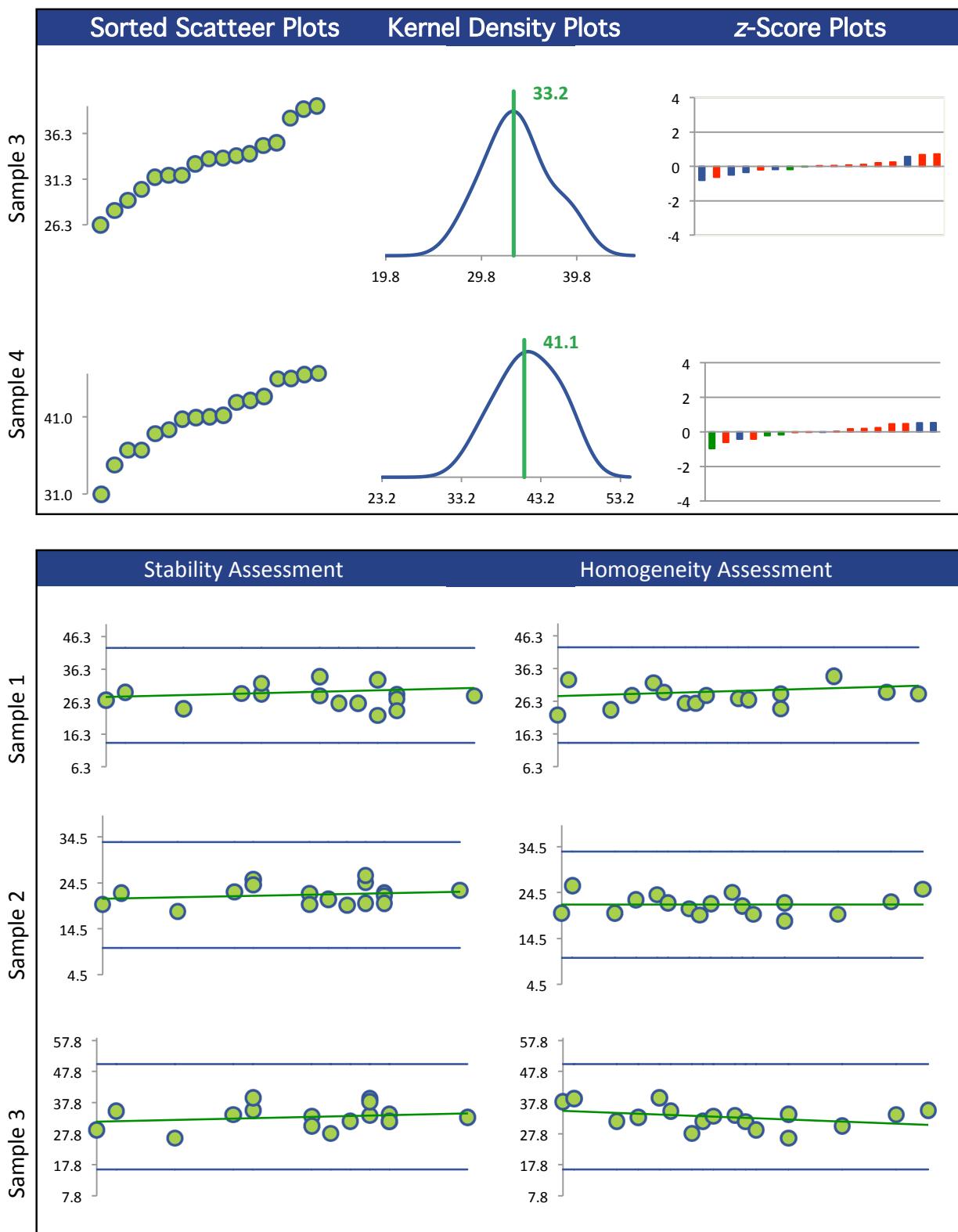
Method	C22-1	C22-2	C22-3	C22-4
GC/MS (Blue)	4	4	4	4
HPLC (Red)	5	5	5	5
LC-MS/MS (Green)	8	8	8	8

All summary stats and the plots below are based on the data excluding any flagged outliers



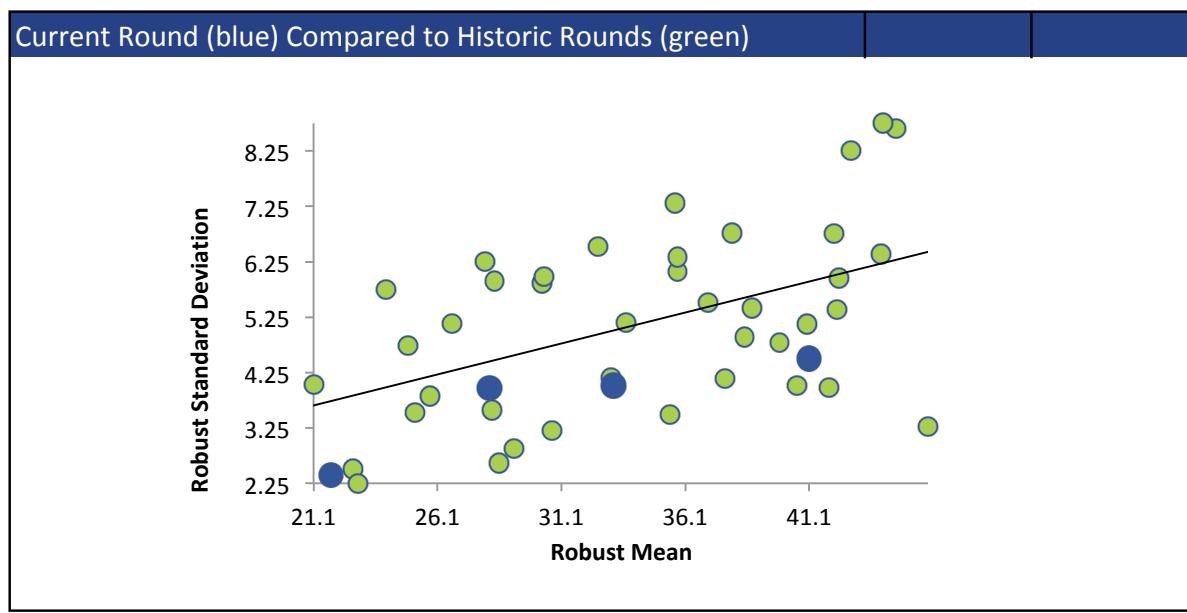
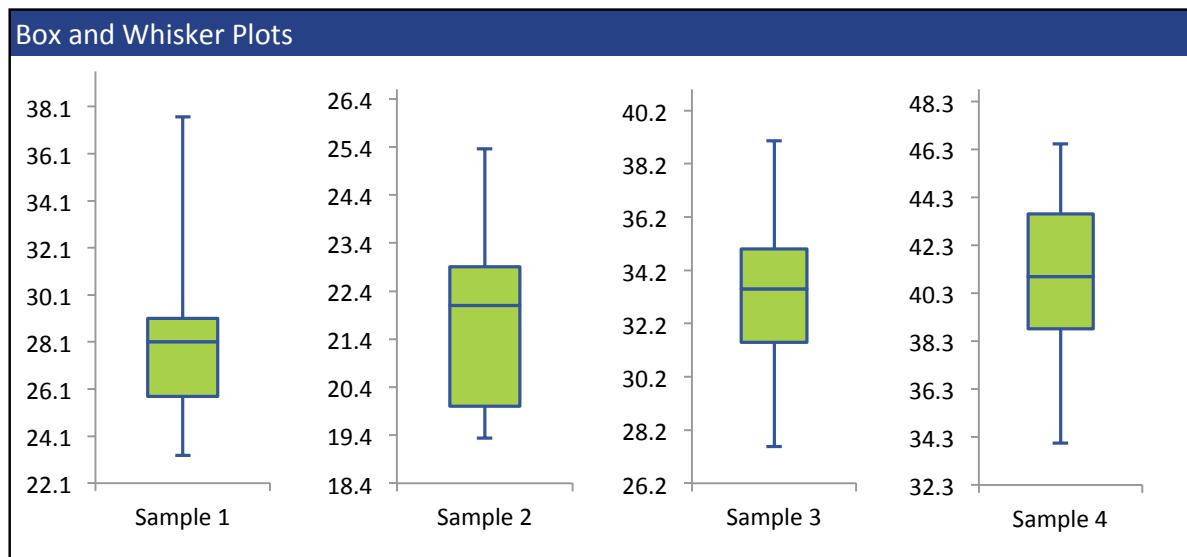
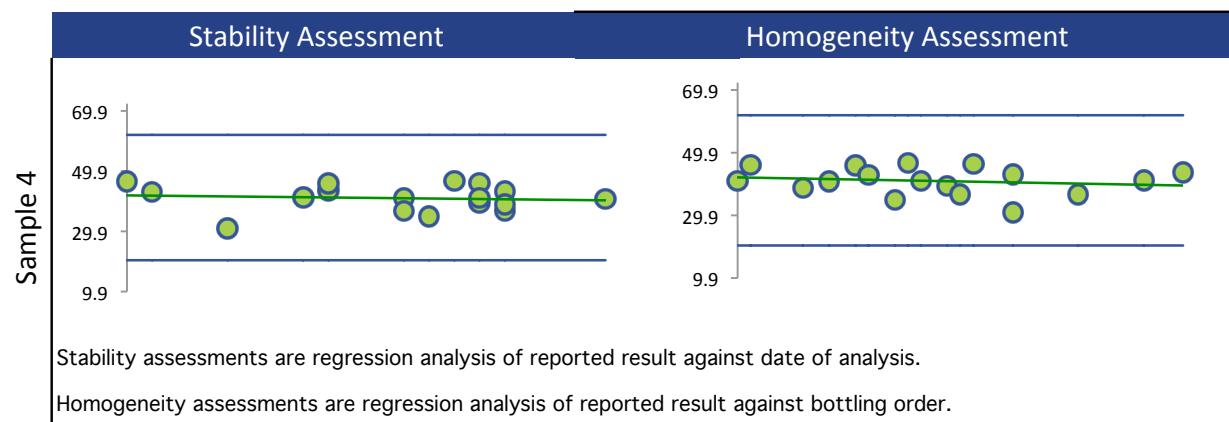
Annex A Summary by Analyte

DIURON



Annex A Summary by Analyte

DIURON



MALATHION

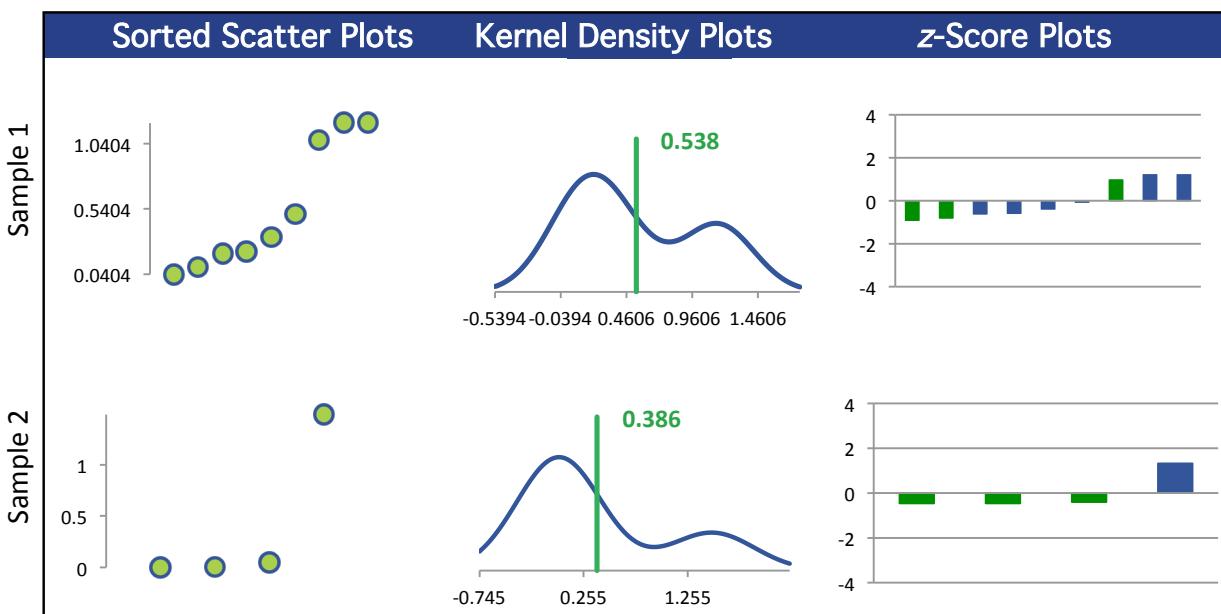
Summary Statistics

Statistic	C22-1	C22-2	C22-3	C22-4
N	9	4	4	4
Median $\mu\text{g/L}$	0.324	0.0271	0.0606	0.0270
Robust Mean $\mu\text{g/L}$	0.538	0.386	0.608	0.243
$\text{U } \mu\text{g/L}$	0.228	0.522	0.806	0.320
Robust Standard Deviation $\mu\text{g/L}$	0.548	0.835	1.29	0.512
Regression Standard Deviation $\mu\text{g/L}$	0.366	0.336	0.380	0.308
Stability Flag				
Homogeneity Flag				
Standard Deviation Used (SDPA) $\mu\text{g/L}$	0.548	0.835	1.29	0.512
Outliers	0	0	0	0
$ z > 3.0$	0	0	0	0
$2 < z < 3$	0	0	0	0

Methods Used

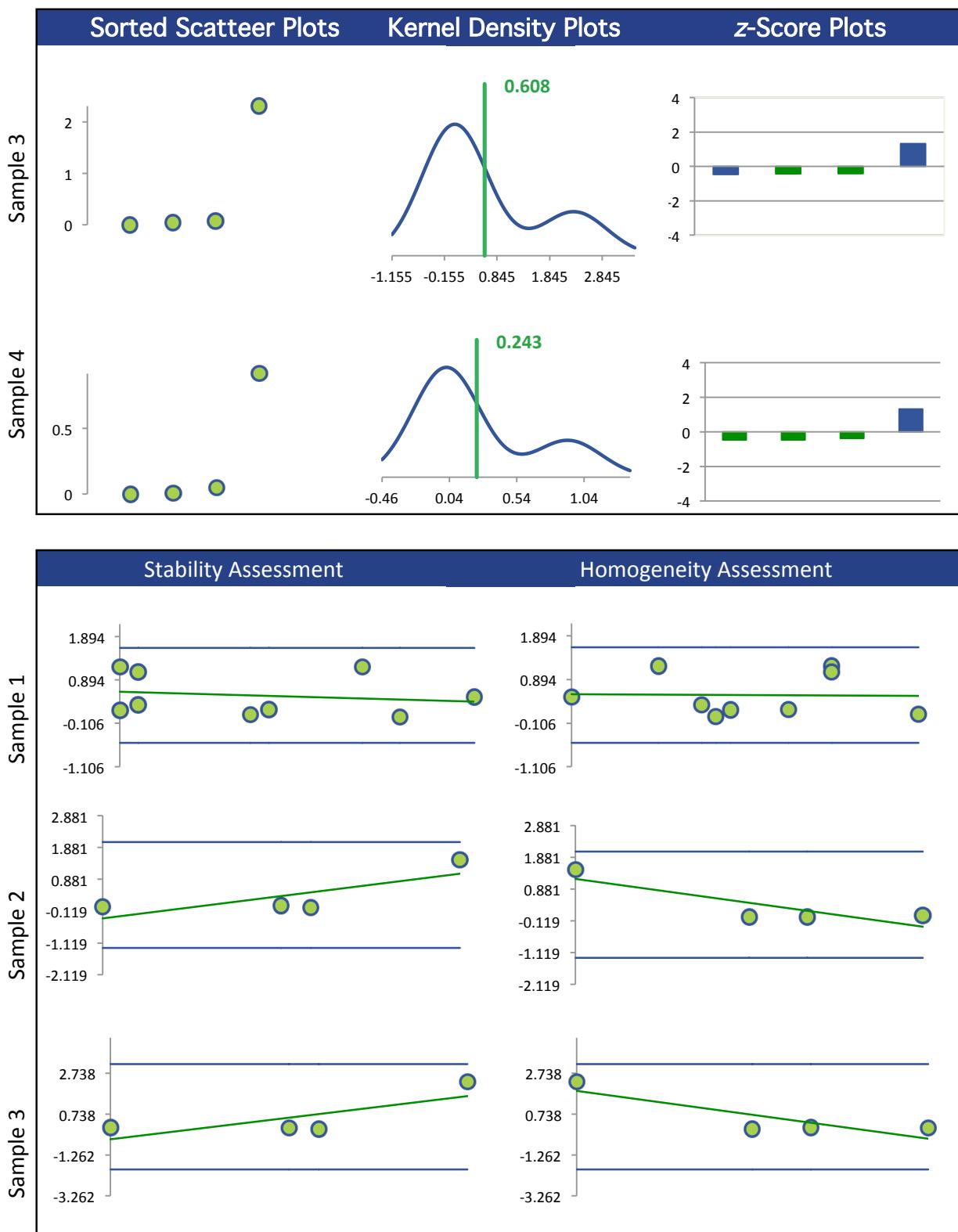
Method	C22-1	C22-2	C22-3	C22-4
LC-MS/MS (Blue)	3	2	2	2
GC/MS (Red)	5	1	1	1
GC/MS/MS (Green)	1	1	1	1

All summary stats and the plots below are based on the data excluding any flagged outliers



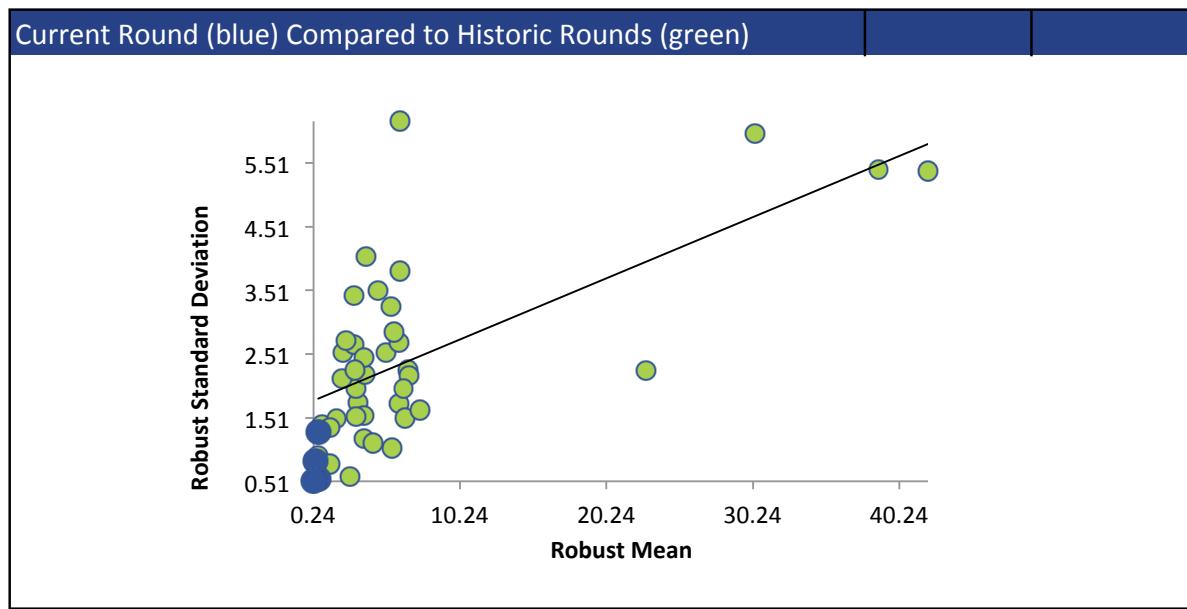
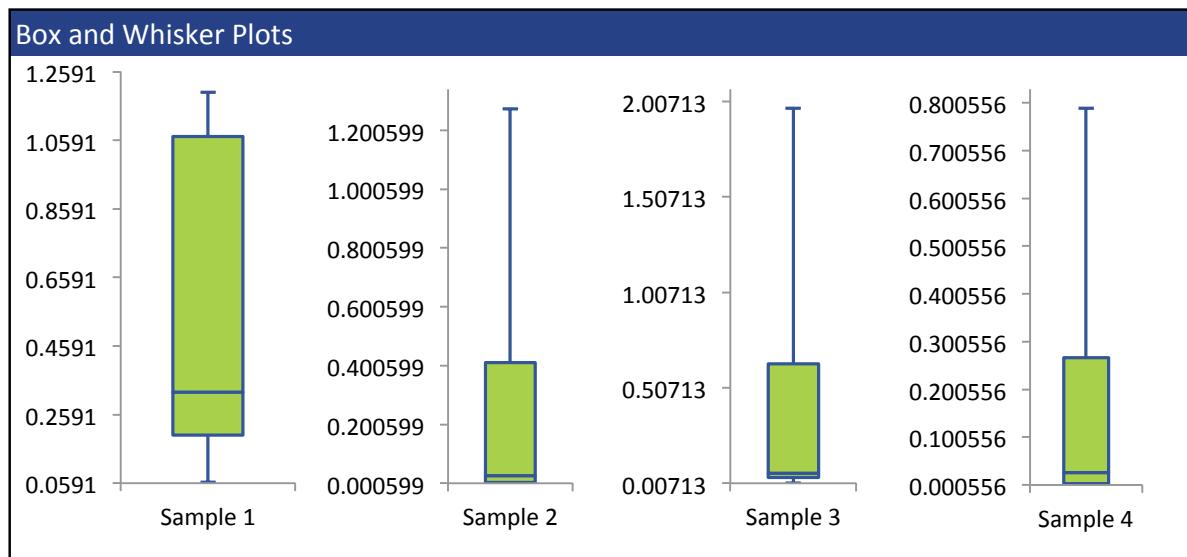
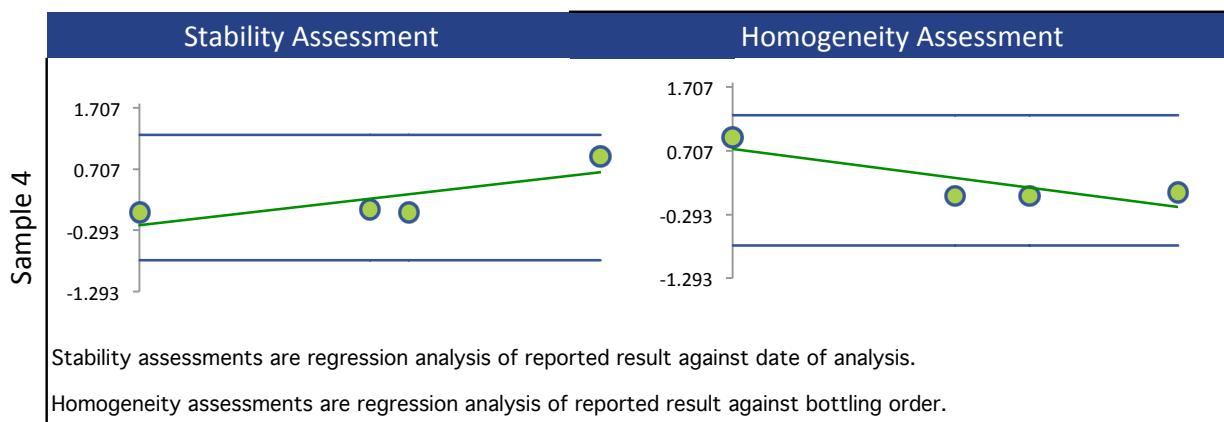
Annex A Summary by Analyte

MALATHION



Annex A Summary by Analyte

MALATHION



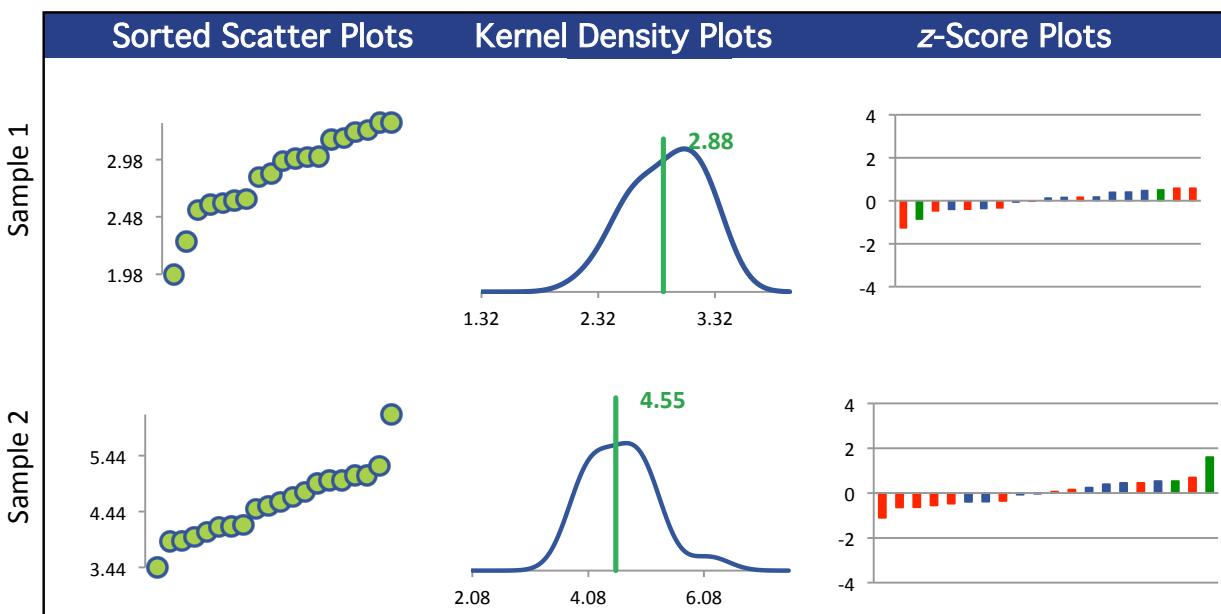
METOLACHLOR**Summary Statistics**

Statistic	C22-1	C22-2	C22-3	C22-4
N	19	20	20	20
Median $\mu\text{g/L}$	2.97	4.58	7.01	7.86
Robust Mean $\mu\text{g/L}$	2.88	4.55	6.96	7.95
$\text{U } \mu\text{g/L}$	0.103	0.163	0.237	0.219
Robust Standard Deviation $\mu\text{g/L}$	0.358	0.584	0.849	0.784
Regression Standard Deviation $\mu\text{g/L}$	0.707	1.01	1.44	1.61
Stability Flag				
Homogeneity Flag				
Standard Deviation Used (SDPA) $\mu\text{g/L}$	0.707	1.01	1.44	1.61
Outliers	0	0	0	0
$ z > 3.0$	0	0	0	0
$2 < z < 3$	0	0	0	1

Methods Used

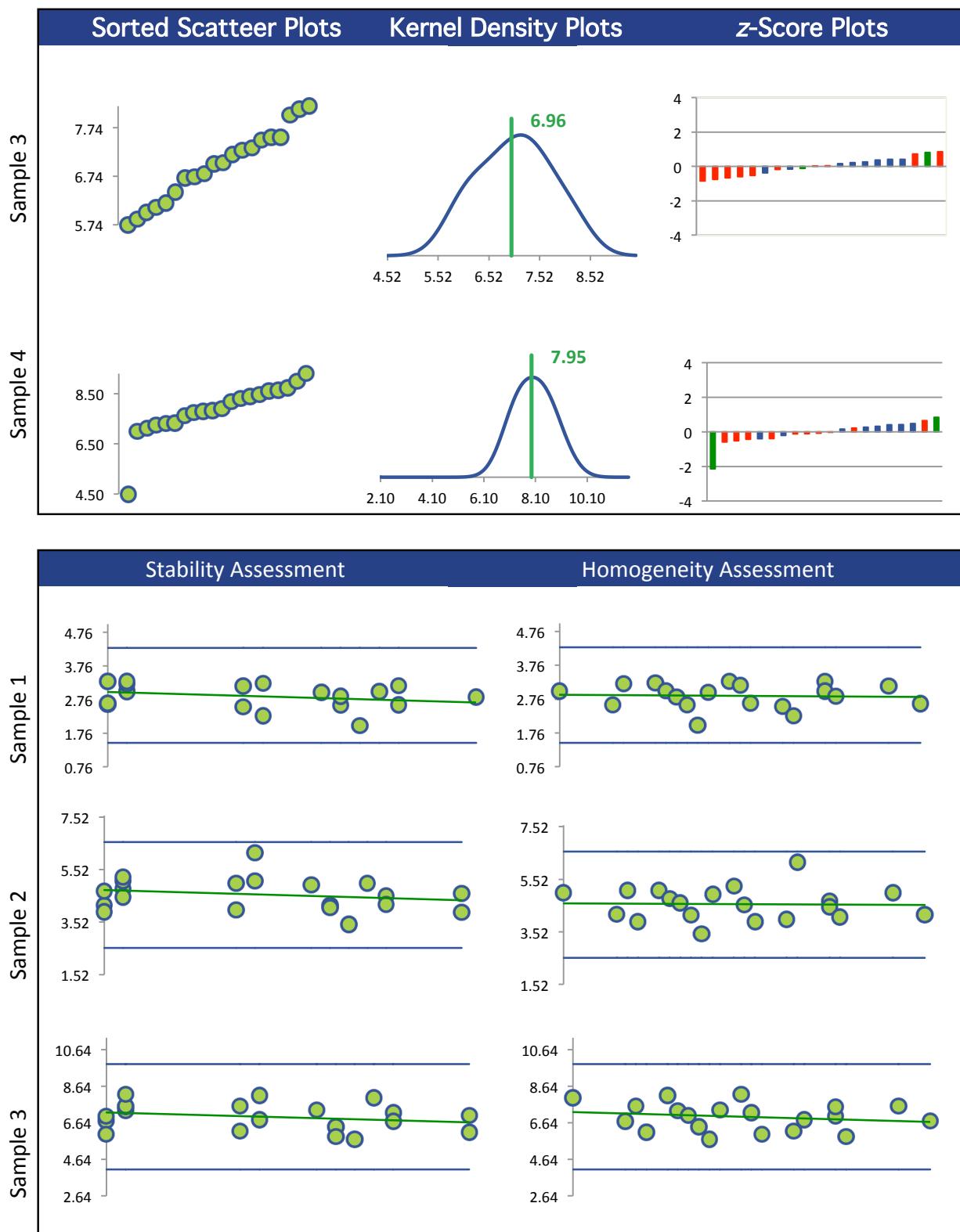
Method	C22-1	C22-2	C22-3	C22-4
GC/MS (Blue)	8	9	9	9
GC/MS/MS (Red)	2	2	2	2
LC-MS/MS (Green)	8	8	8	8
LC/MS (Orange)	1	1	1	1

All summary stats and the plots below are based on the data excluding any flagged outliers



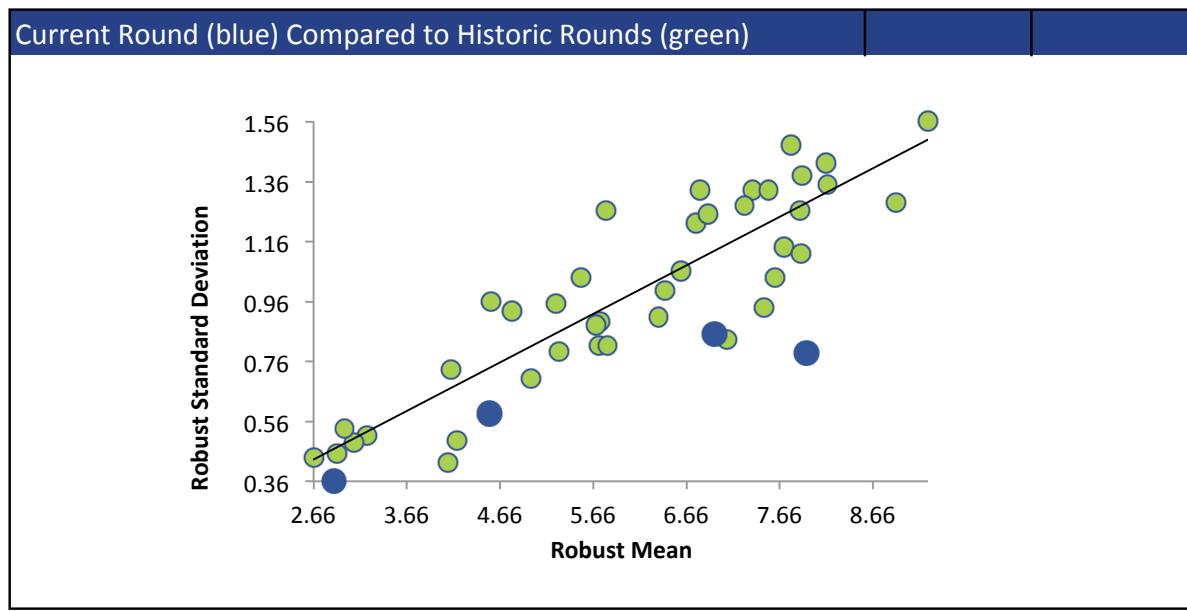
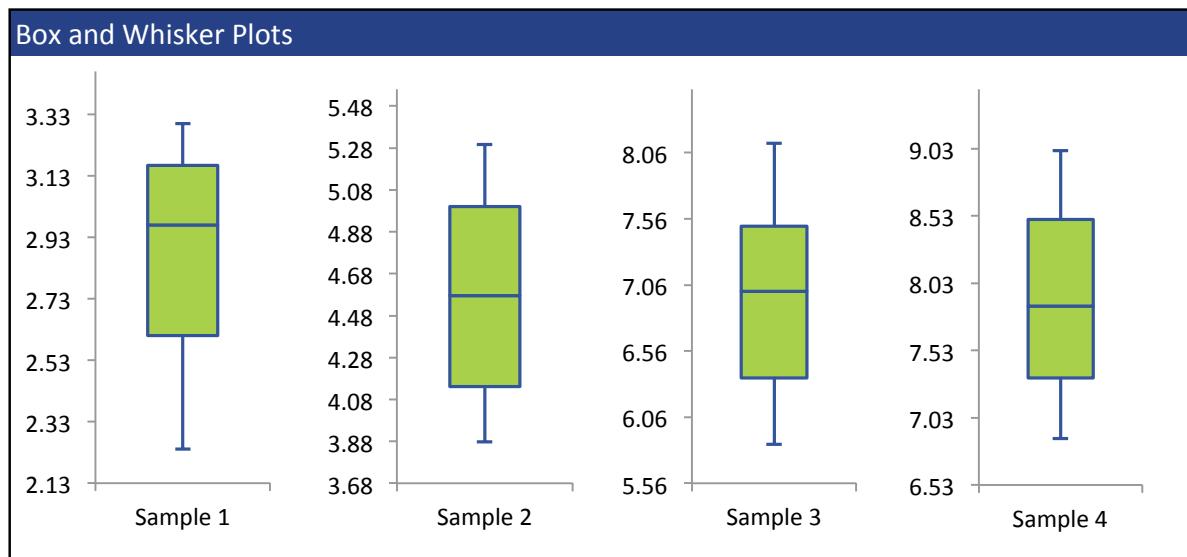
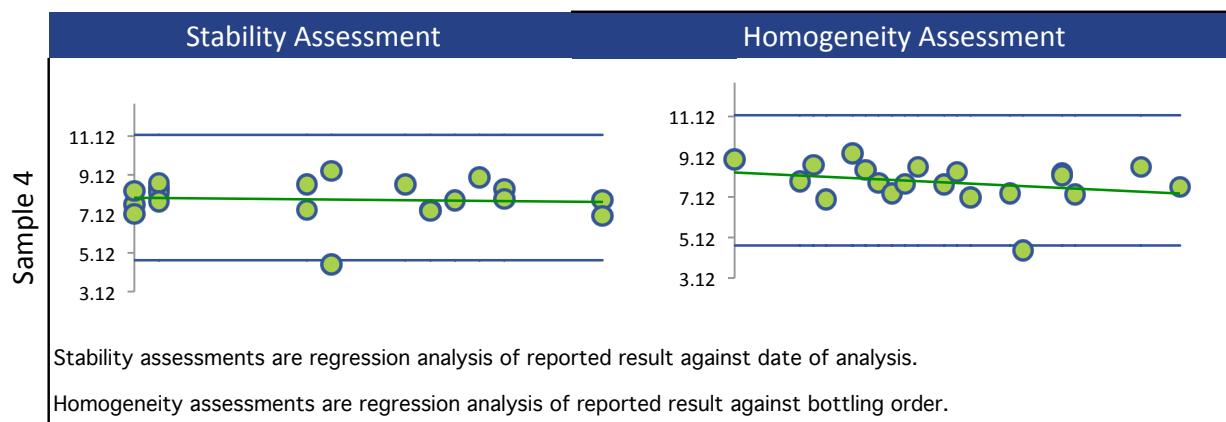
Annex A Summary by Analyte

METOLACHLOR



Annex A Summary by Analyte

METOLACHLOR



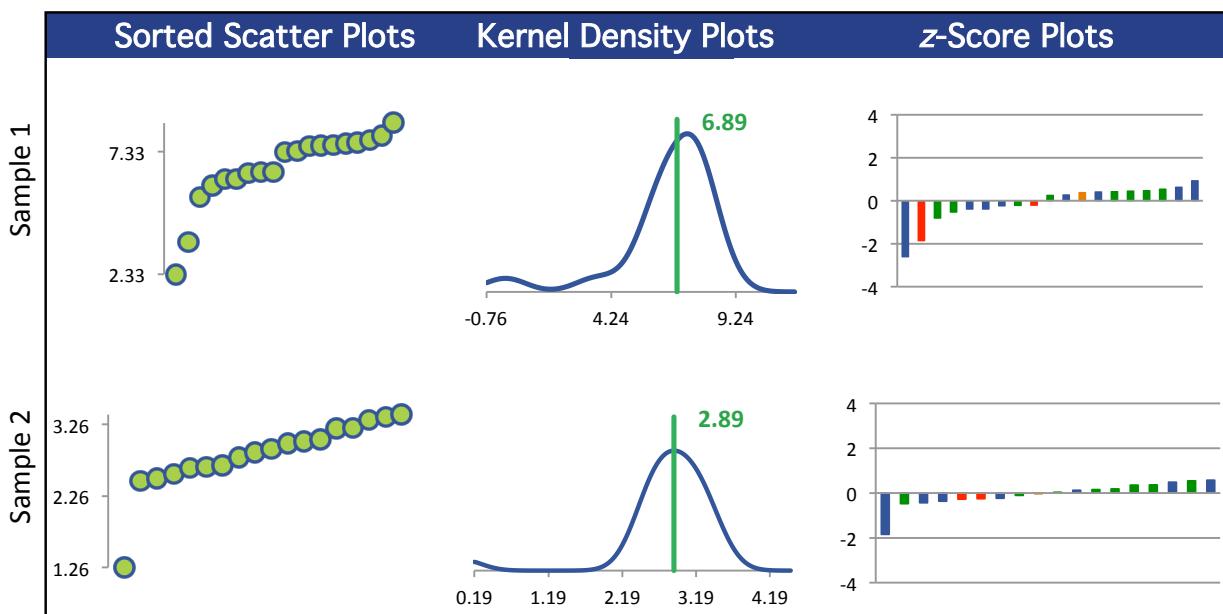
METRIBUZIN**Summary Statistics**

Statistic	C22-1	C22-2	C22-3	C22-4
N	19	18	19	20
Median $\mu\text{g/L}$	7.32	2.90	4.06	6.05
Robust Mean $\mu\text{g/L}$	6.89	2.89	3.96	6.08
$\text{U } \mu\text{g/L}$	0.324	0.109	0.145	0.200
Robust Standard Deviation $\mu\text{g/L}$	1.13	0.370	0.504	0.716
Regression Standard Deviation $\mu\text{g/L}$	1.75	0.888	1.12	1.58
Stability Flag				
Homogeneity Flag				
Standard Deviation Used (SDPA) $\mu\text{g/L}$	1.75	0.888	1.12	1.58
Outliers	0	0	0	0
$ \text{z} > 3.0$	0	0	0	0
$2 < \text{z} < 3$	1	0	0	2

Methods Used

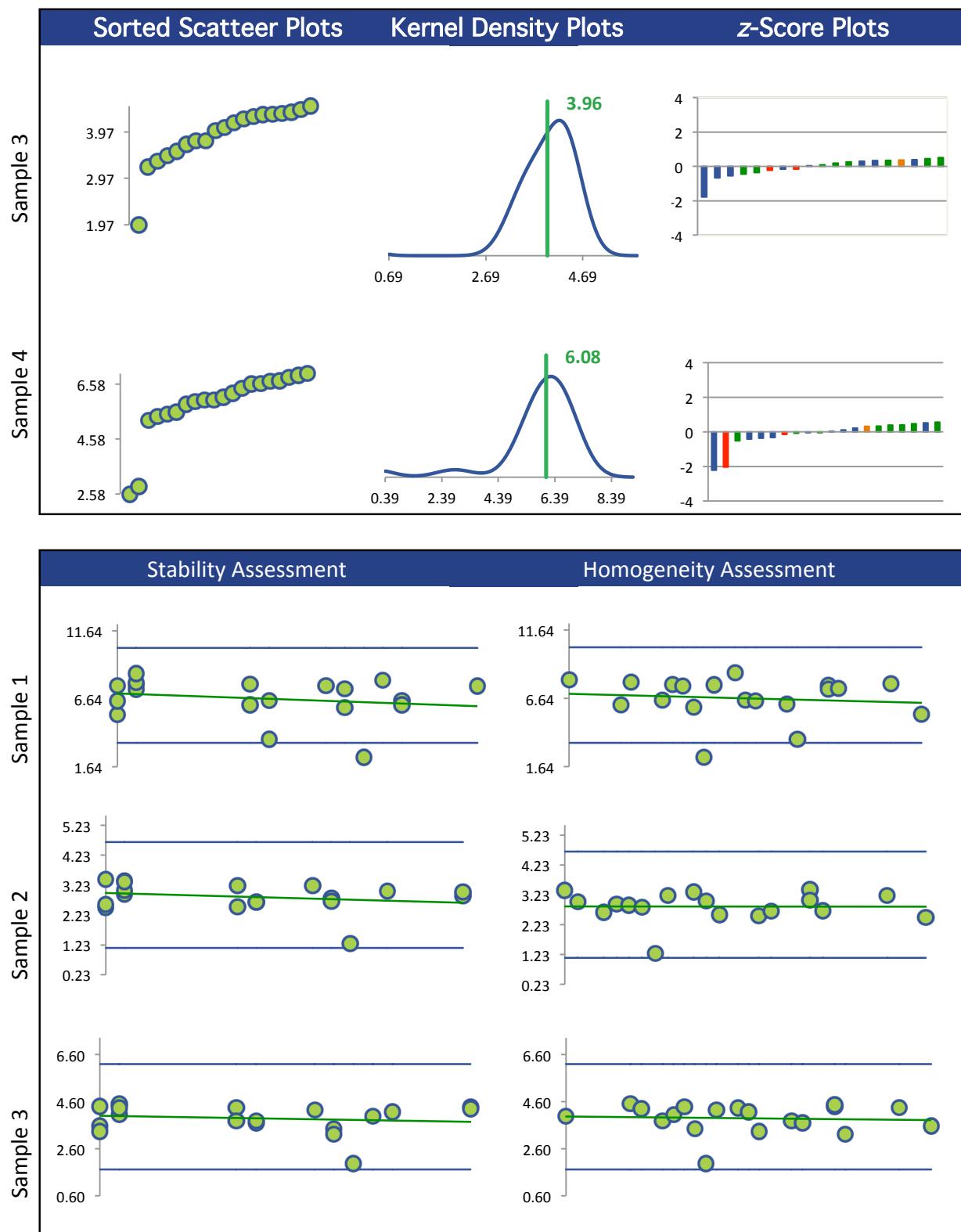
Method	C22-1	C22-2	C22-3	C22-4
GC/MS (Blue)	8	7	8	9
GC/MS/MS (Red)	2	2	2	2
LC-MS/MS (Green)	8	8	8	8
LC/MS (Orange)	1	1	1	1

All summary stats and the plots below are based on the data excluding any flagged outliers



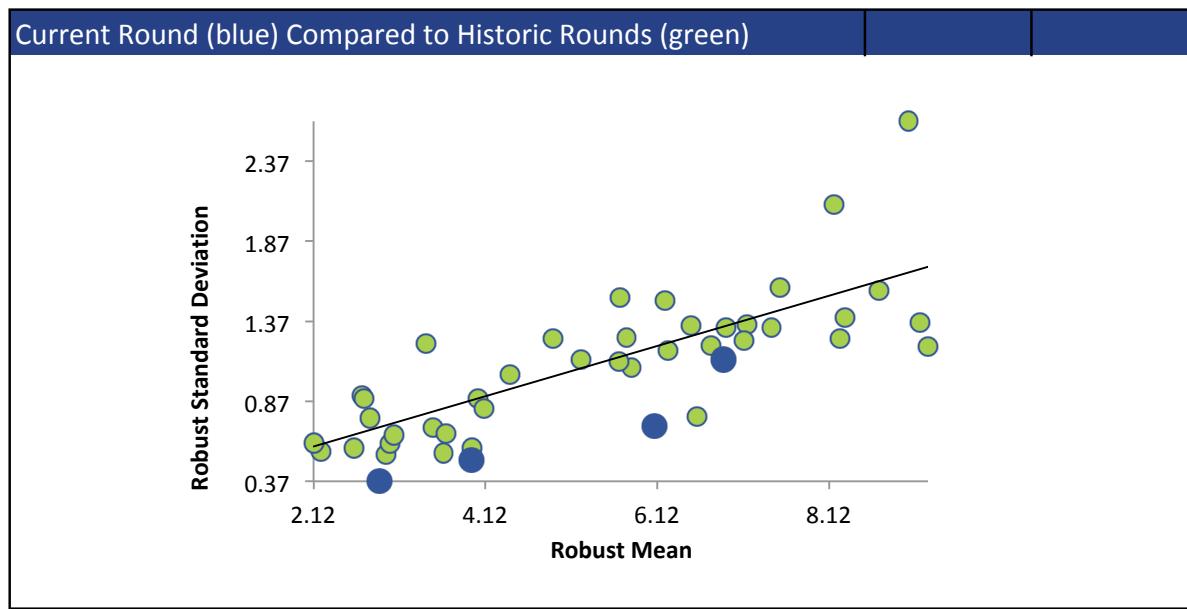
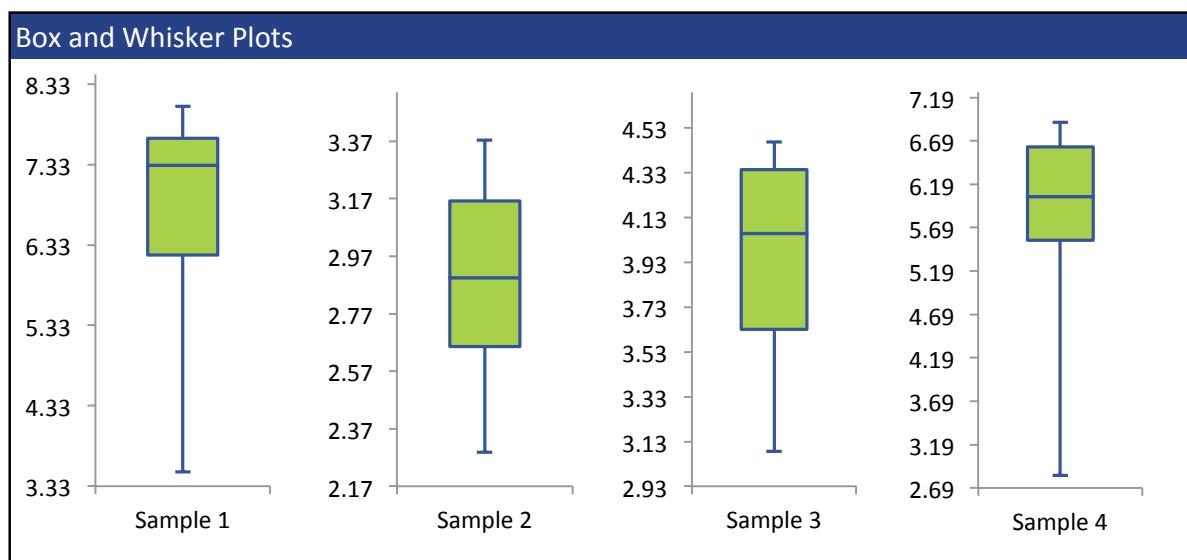
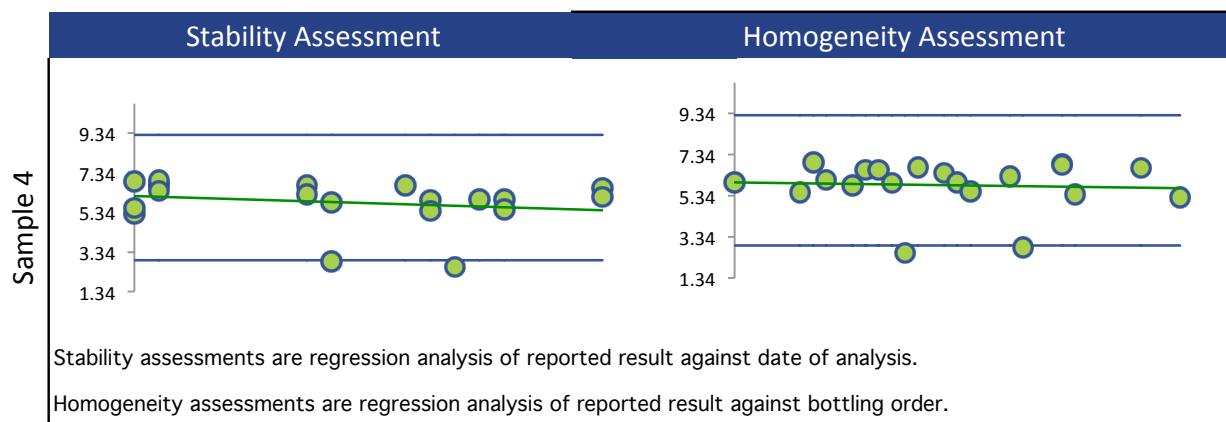
Annex A Summary by Analyte

METRIBUZIN



Annex A Summary by Analyte

METRIBUZIN



PARATHION (ETHYL)

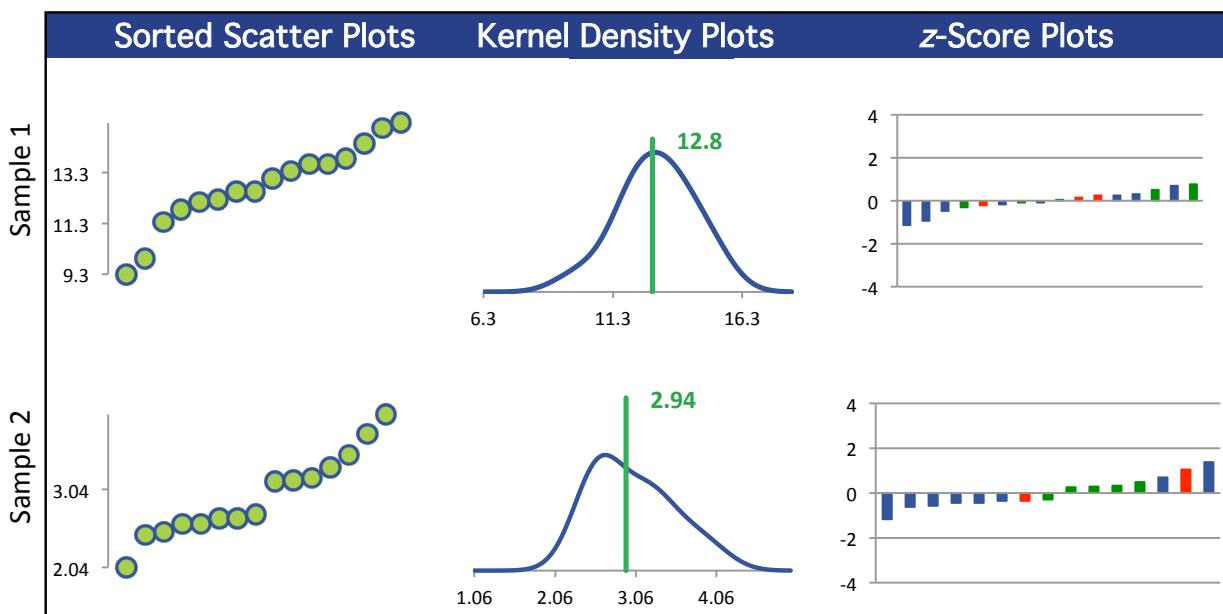
Summary Statistics

Statistic	C22-1	C22-2	C22-3	C22-4
N	16	15	16	16
Median $\mu\text{g/L}$	12.8	2.72	11.2	6.04
Robust Mean $\mu\text{g/L}$	12.8	2.94	11.4	6.04
$\text{U } \mu\text{g/L}$	0.522	0.185	0.534	0.290
Robust Standard Deviation $\mu\text{g/L}$	1.67	0.572	1.71	0.928
Regression Standard Deviation $\mu\text{g/L}$	3.08	0.768	2.75	1.50
Stability Flag				
Homogeneity Flag				
Standard Deviation Used (SDPA) $\mu\text{g/L}$	3.08	0.768	2.75	1.50
Outliers	0	1	0	0
$ \text{z} > 3.0$	0	0	0	0
$2 < \text{z} < 3$	0	0	0	1

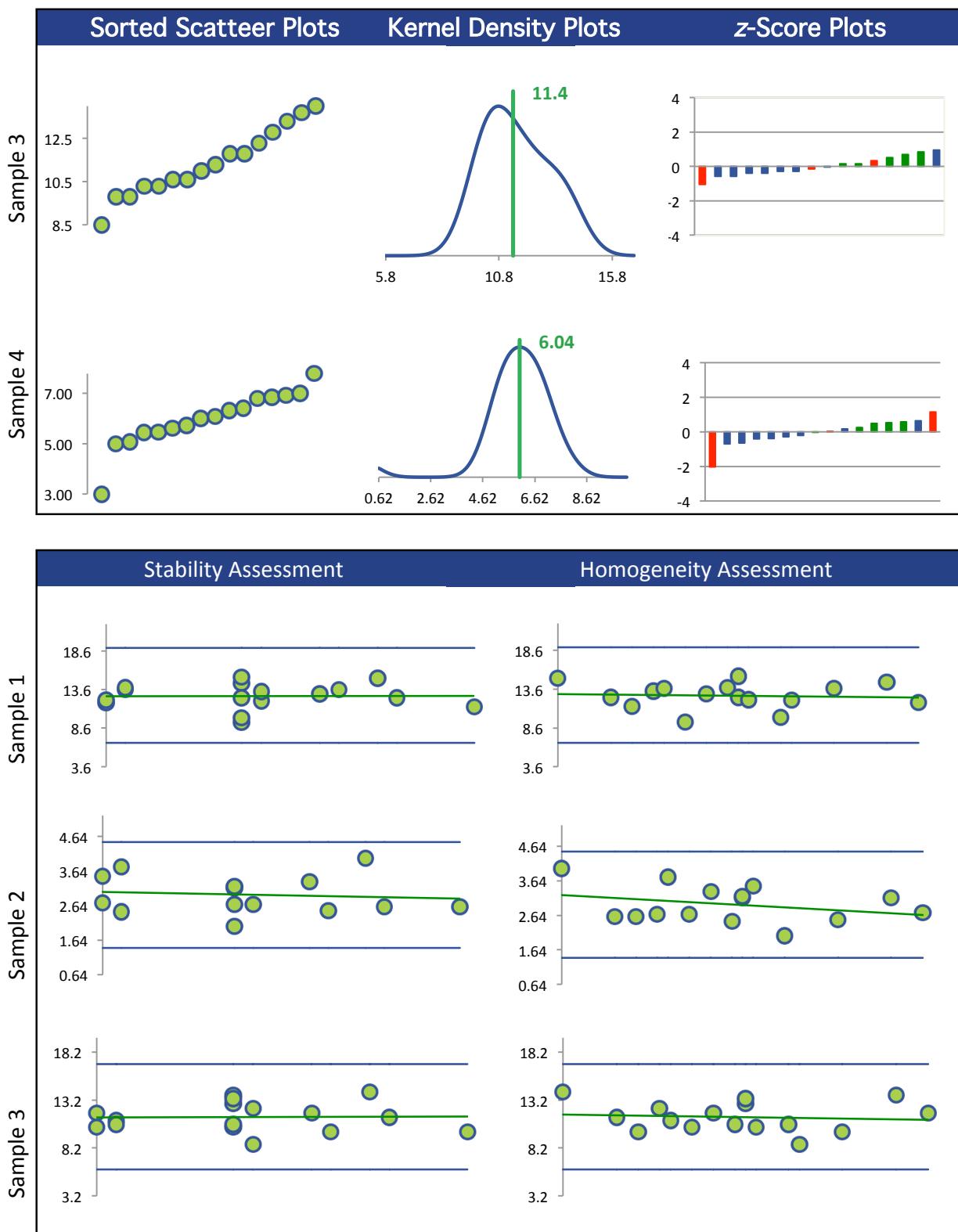
Methods Used

Method	C22-1	C22-2	C22-3	C22-4
GC/MS (Blue)	8	8	8	8
LC-MS/MS (Red)	5	5	5	5
GC/MS/MS (Green)	3	2	3	3

All summary stats and the plots below are based on the data excluding any flagged outliers

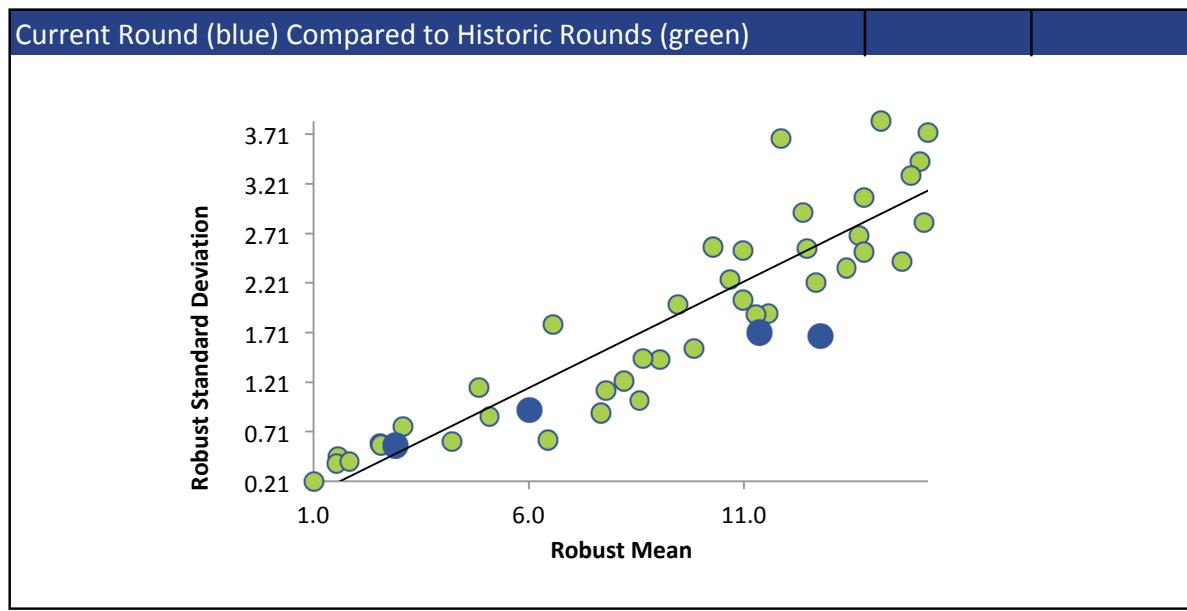
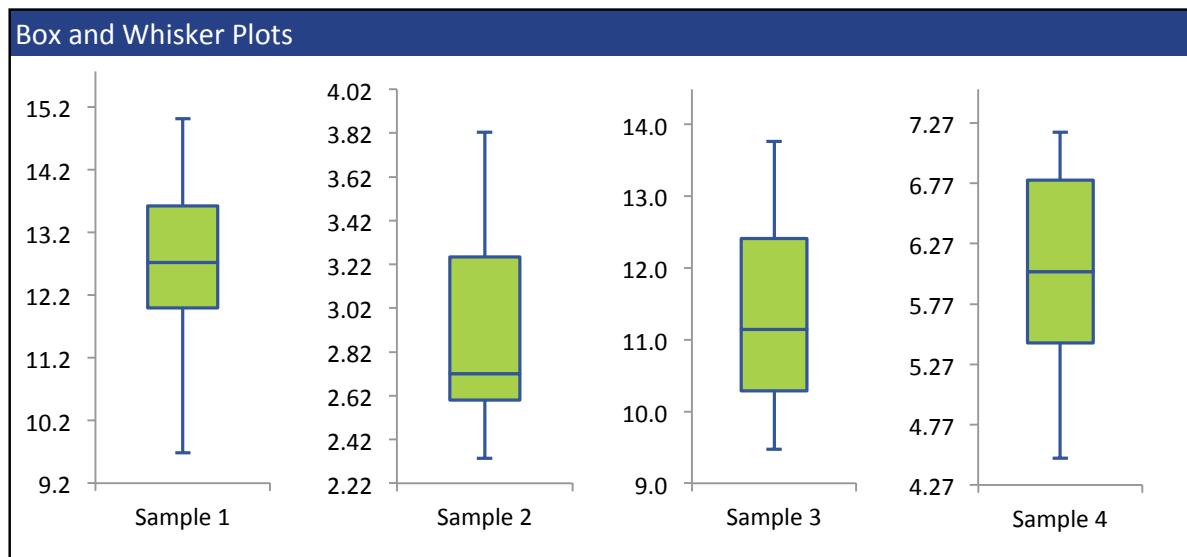
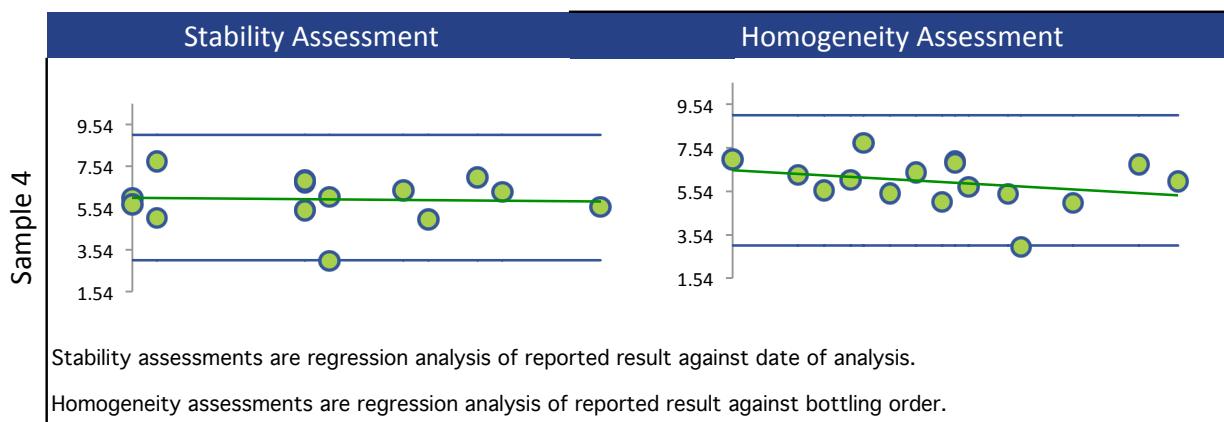


PARATHION (ETHYL)



Annex A Summary by Analyte

PARATHION (ETHYL)



PHORATE

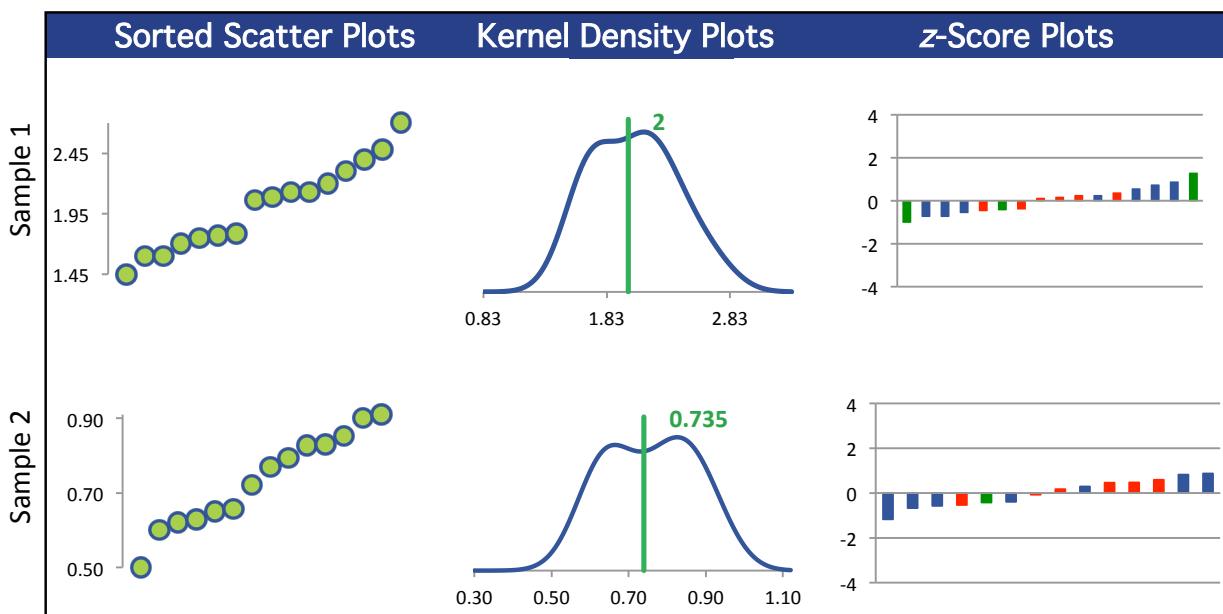
Summary Statistics

Statistic	C22-1	C22-2	C22-3	C22-4
N	16	14	16	15
Median $\mu\text{g/L}$	2.08	0.745	3.28	2.67
Robust Mean $\mu\text{g/L}$	2.00	0.735	3.19	2.73
$\text{U } \mu\text{g/L}$	0.122	0.0461	0.160	0.133
Robust Standard Deviation $\mu\text{g/L}$	0.391	0.138	0.513	0.413
Regression Standard Deviation $\mu\text{g/L}$	0.549	0.201	0.875	0.749
Stability Flag				
Homogeneity Flag				
Standard Deviation Used (SDPA) $\mu\text{g/L}$	0.549	0.201	0.875	0.749
Outliers	0	1	0	1
$ \text{z} > 3.0$	0	0	0	0
$2 < \text{z} < 3$	0	0	0	0

Methods Used

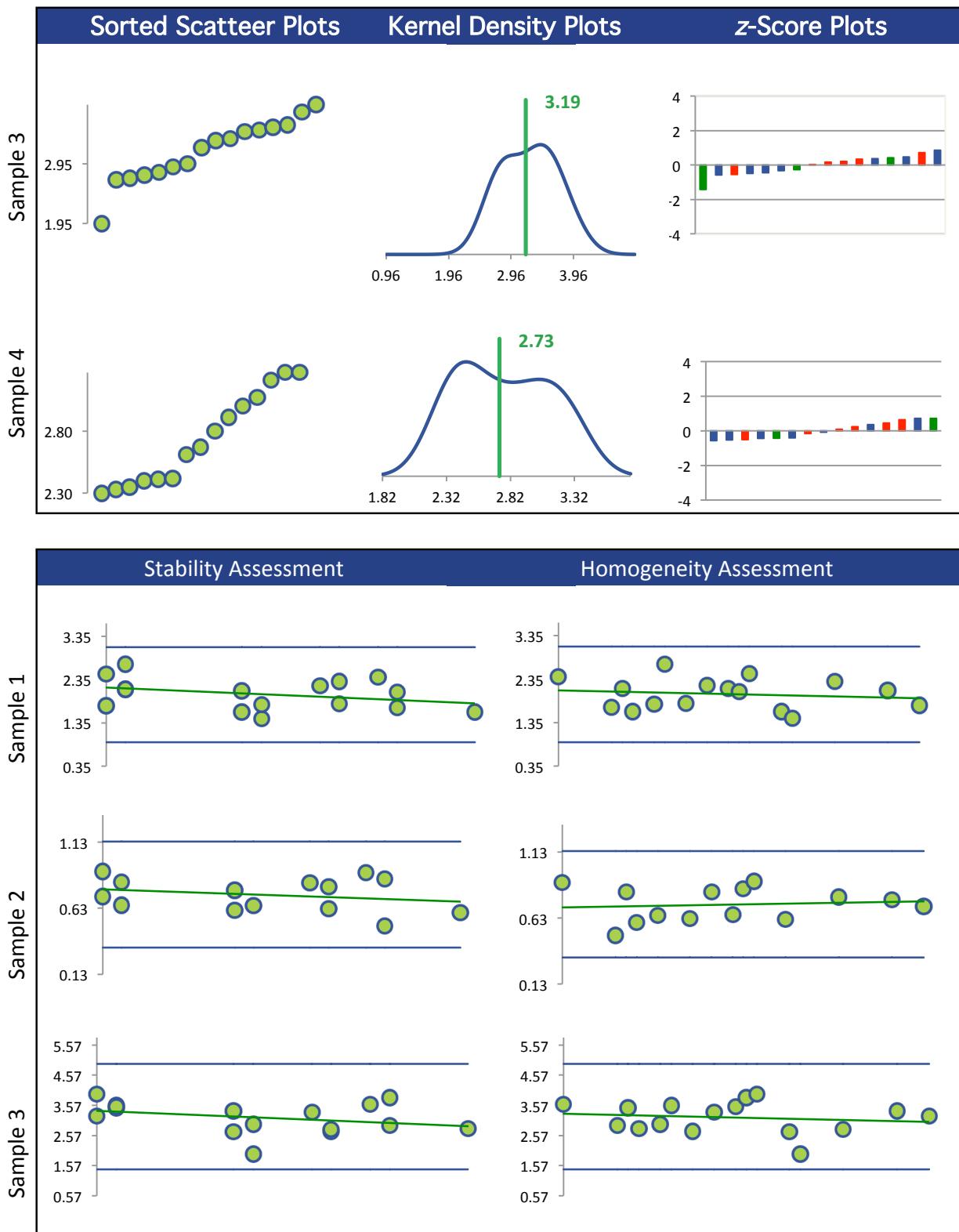
Method	C22-1	C22-2	C22-3	C22-4
GC/MS/MS (Blue)	3	1	3	2
GC/MS (Red)	7	7	7	7
LC-MS/MS (Green)	6	6	6	6

All summary stats and the plots below are based on the data excluding any flagged outliers



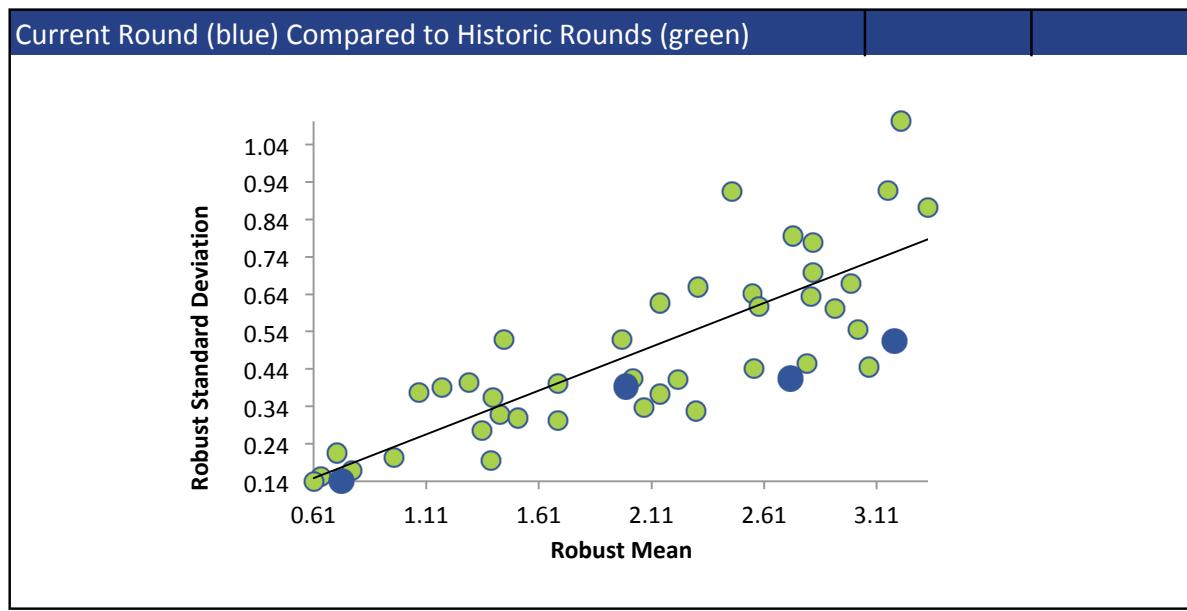
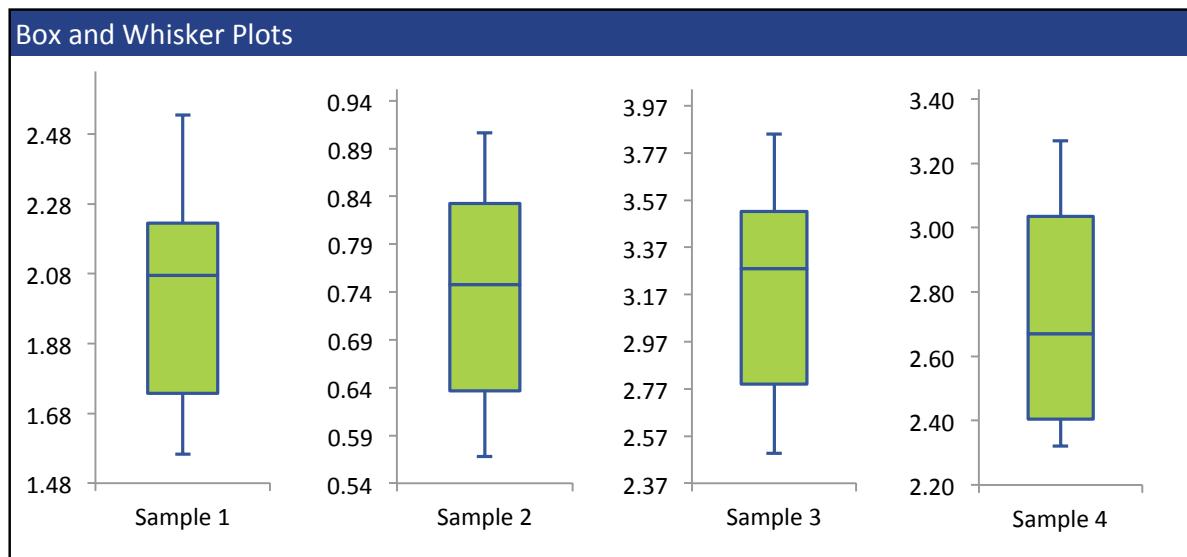
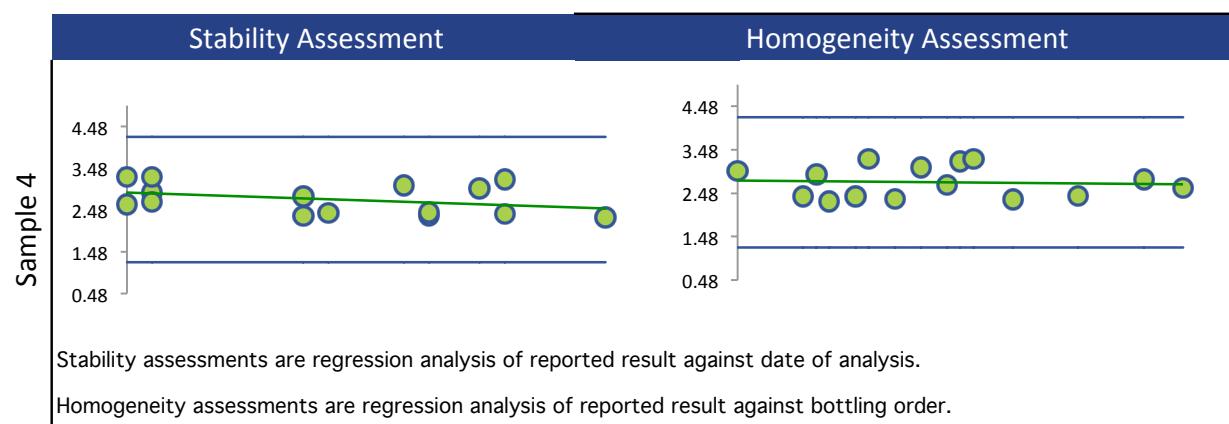
Annex A Summary by Analyte

PHORATE



Annex A Summary by Analyte

PHORATE



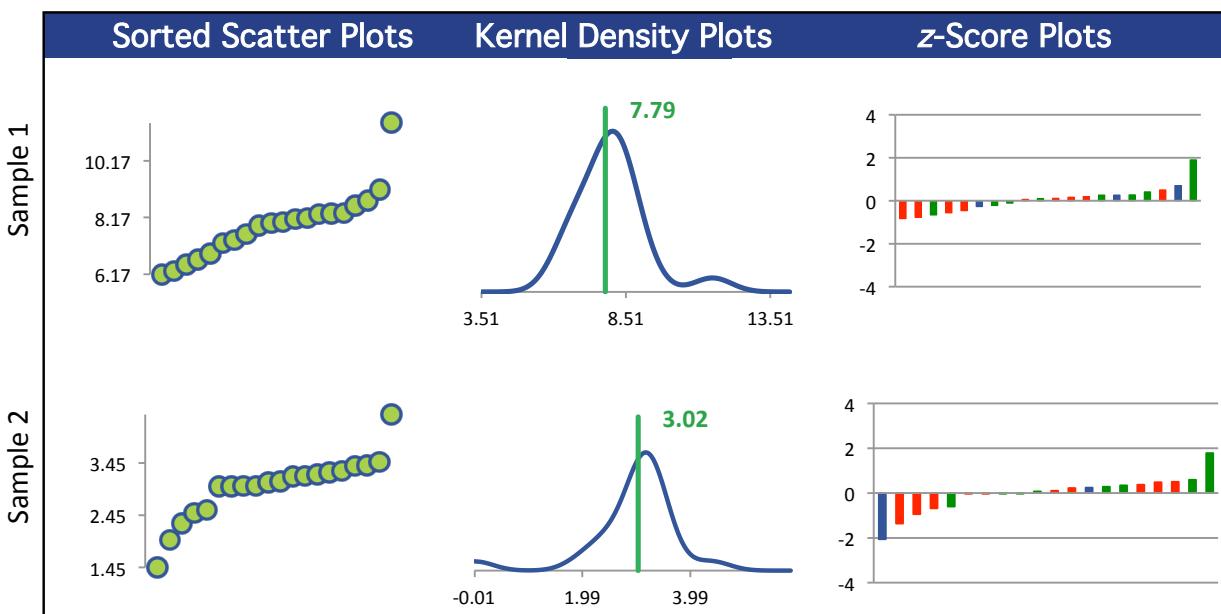
SIMAZINE**Summary Statistics**

Statistic	C22-1	C22-2	C22-3	C22-4
N	20	20	20	20
Median $\mu\text{g/L}$	7.99	3.09	1.80	5.80
Robust Mean $\mu\text{g/L}$	7.79	3.02	1.78	5.63
U $\mu\text{g/L}$	0.288	0.129	0.0696	0.221
Robust Standard Deviation $\mu\text{g/L}$	1.03	0.461	0.249	0.790
Regression Standard Deviation $\mu\text{g/L}$	1.95	0.763	0.454	1.41
Stability Flag				
Homogeneity Flag				
Standard Deviation Used (SDPA) $\mu\text{g/L}$	1.95	0.763	0.454	1.41
Outliers	0	0	0	0
$ z > 3.0$	0	0	1	0
$2 < z < 3$	0	1	0	1

Methods Used

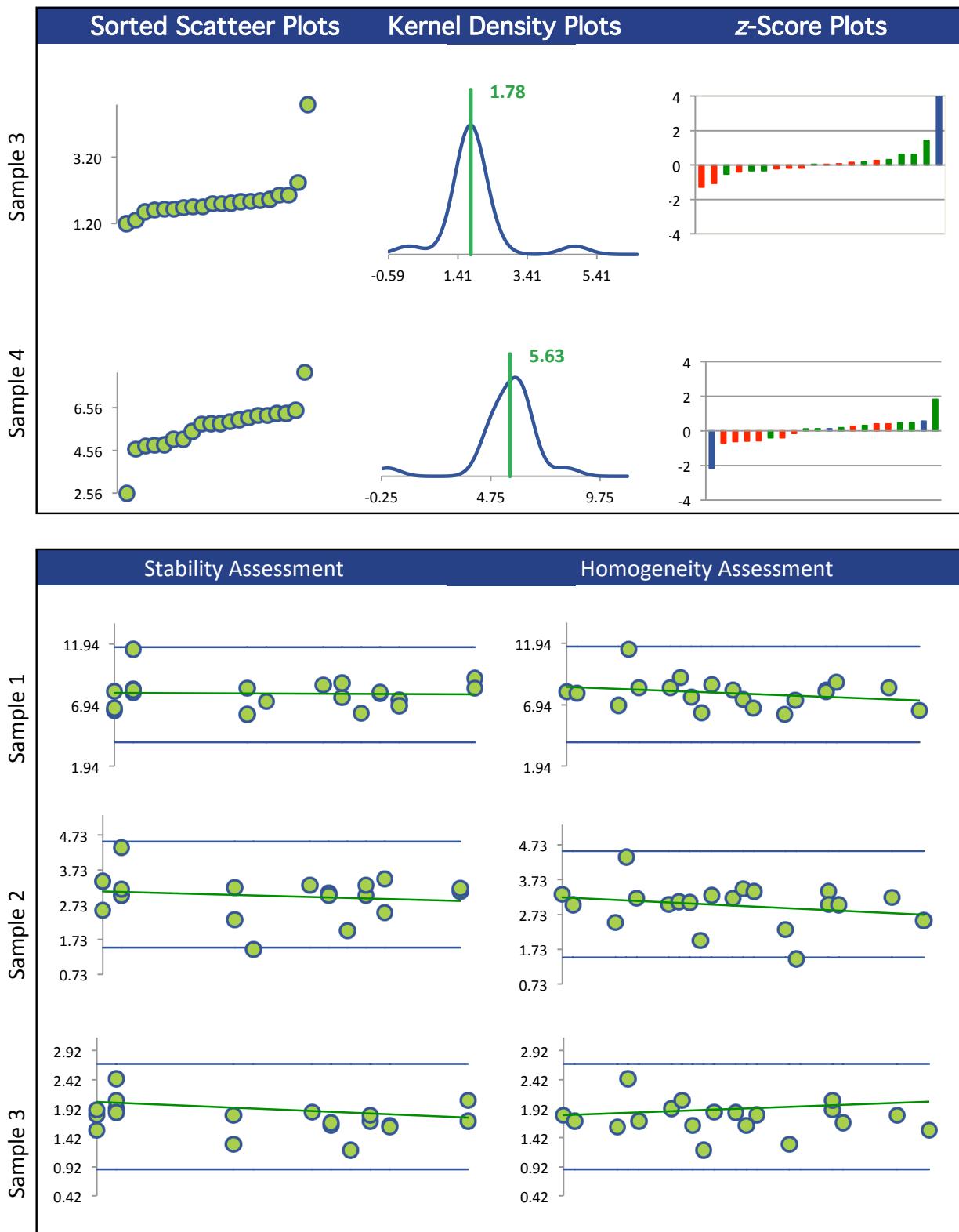
Method	C22-1	C22-2	C22-3	C22-4
GC/MS (Blue)	8	8	8	8
LC-MS/MS (Red)	8	8	8	8
GC/MS/MS (Green)	1	1	1	1
HPLC (Orange)	2	2	2	2
LC/MS (Black)	1	1	1	1

All summary stats and the plots below are based on the data excluding any flagged outliers



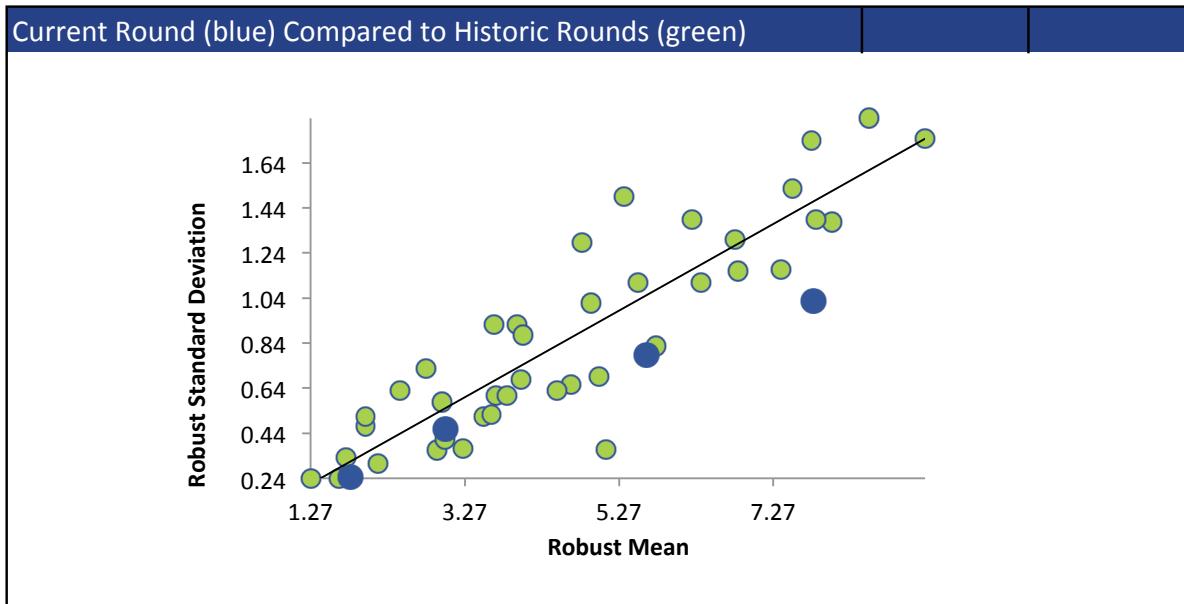
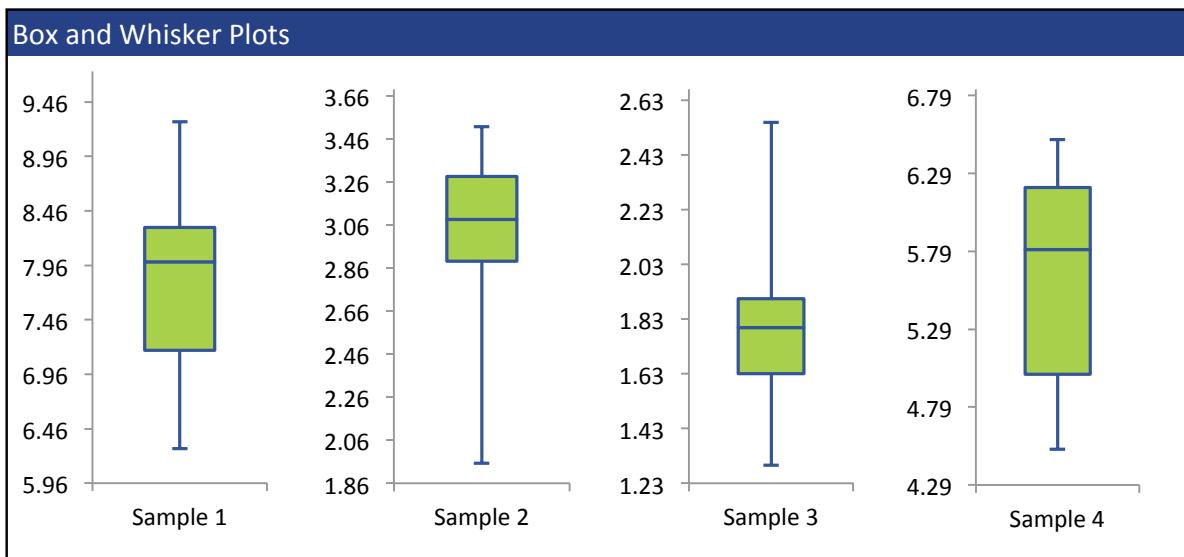
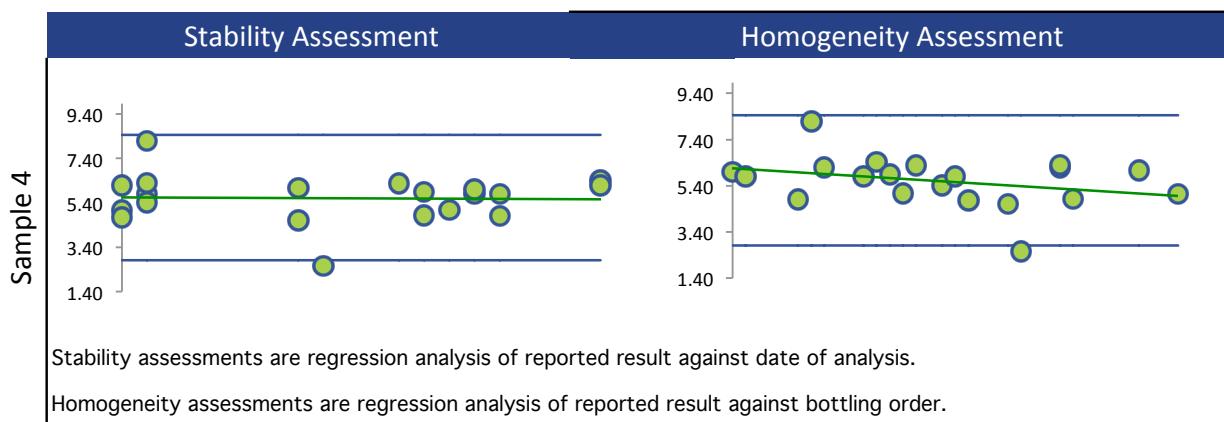
Annex A Summary by Analyte

SIMAZINE



Annex A Summary by Analyte

SIMAZINE



TERBUFOS

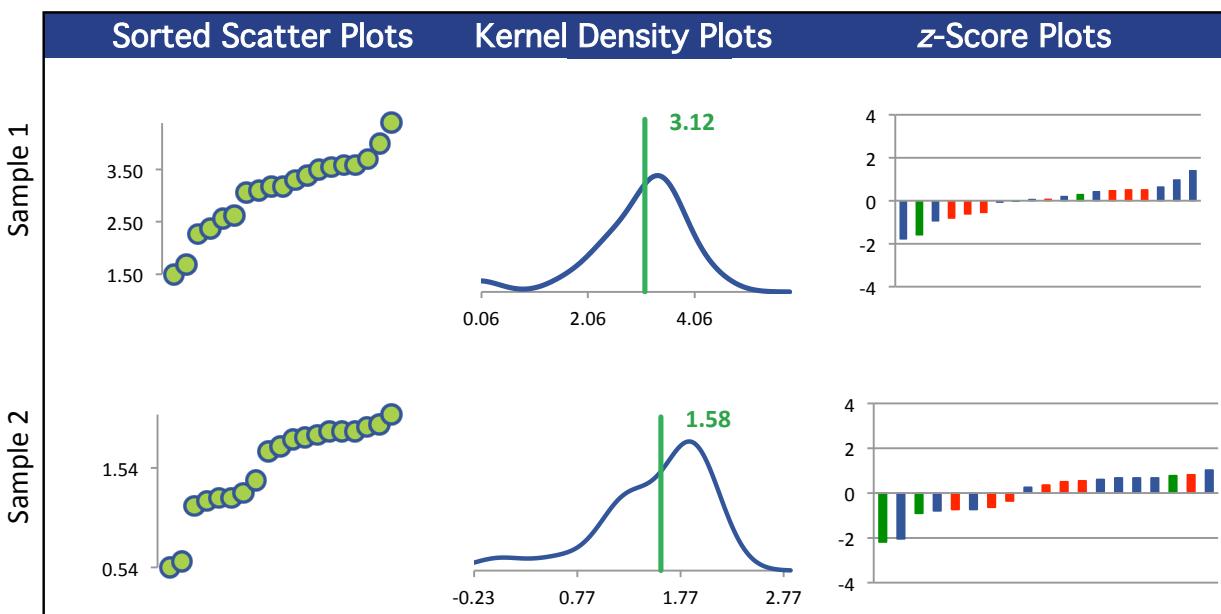
Summary Statistics

Statistic	C22-1	C22-2	C22-3	C22-4
N	19	19	18	19
Median $\mu\text{g/L}$	3.18	1.75	1.00	2.13
Robust Mean $\mu\text{g/L}$	3.12	1.58	0.966	1.95
U $\mu\text{g/L}$	0.208	0.124	0.0807	0.133
Robust Standard Deviation $\mu\text{g/L}$	0.726	0.433	0.274	0.465
Regression Standard Deviation $\mu\text{g/L}$	0.908	0.480	0.308	0.583
Stability Flag				
Homogeneity Flag				
Standard Deviation Used (SDPA) $\mu\text{g/L}$	0.908	0.480	0.308	0.583
Outliers	0	0	0	0
$ z > 3.0$	0	0	1	0
$2 < z < 3$	0	2	0	0

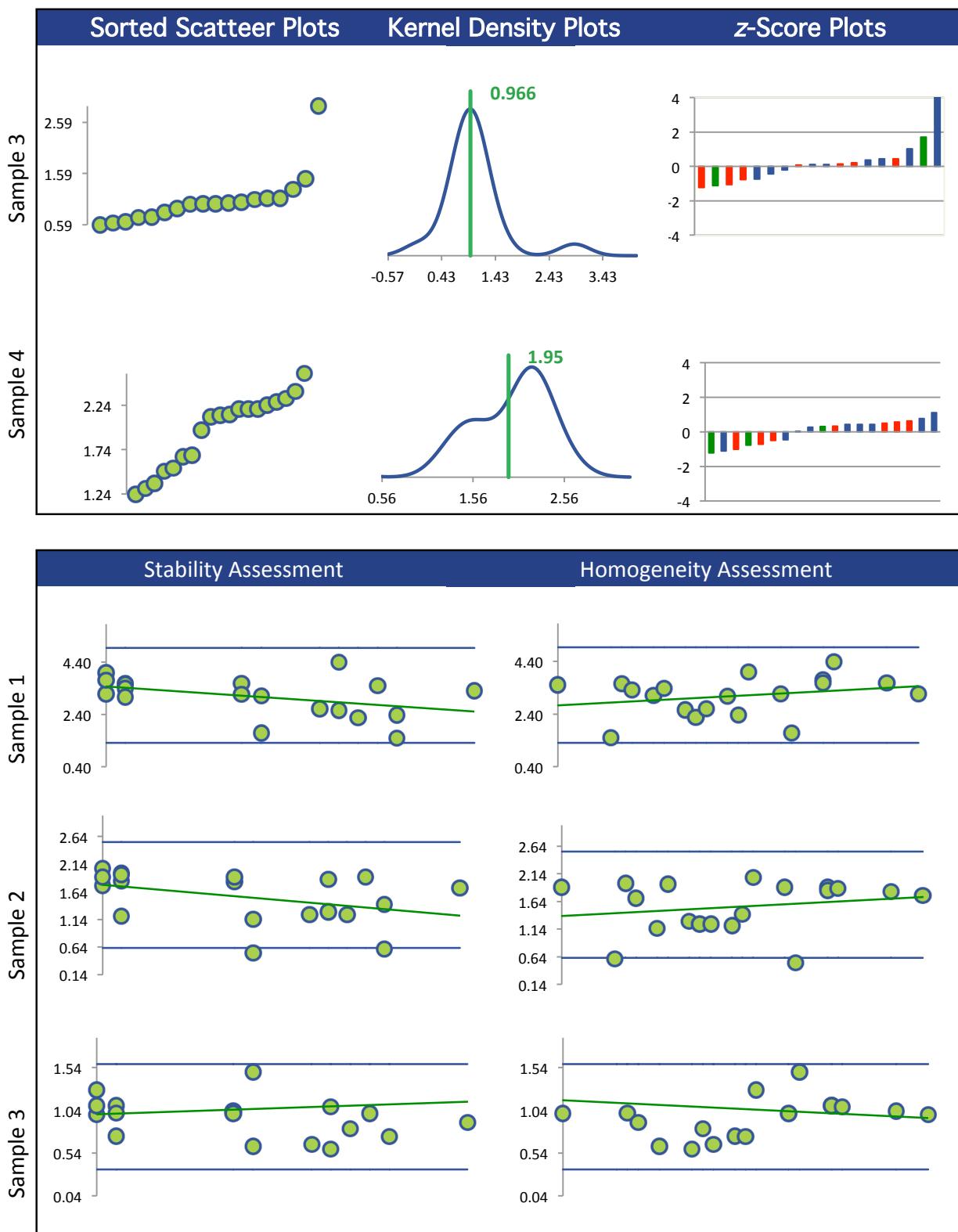
Methods Used

Method	C22-1	C22-2	C22-3	C22-4
GC/MS (Blue)	9	9	9	9
GC/MS/MS (Red)	3	3	2	3
LC-MS/MS (Green)	7	7	7	7

All summary stats and the plots below are based on the data excluding any flagged outliers

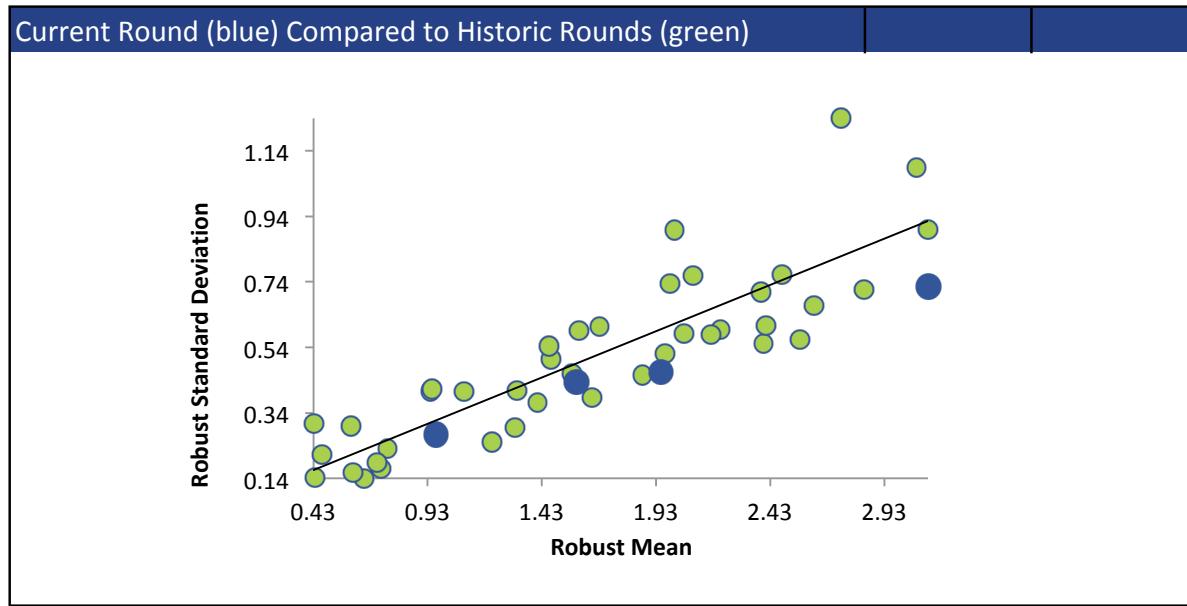
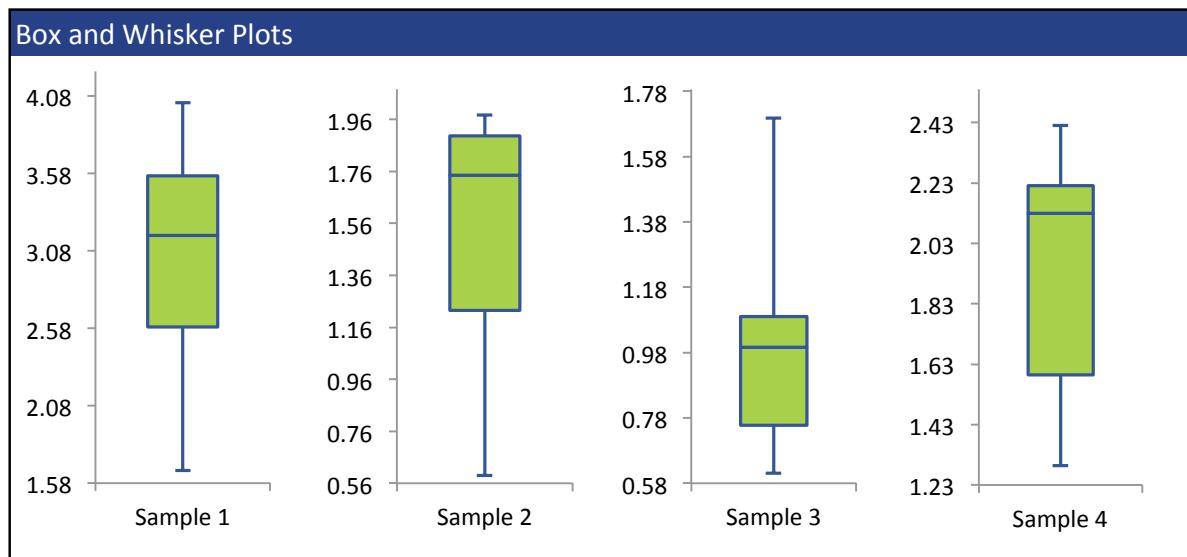
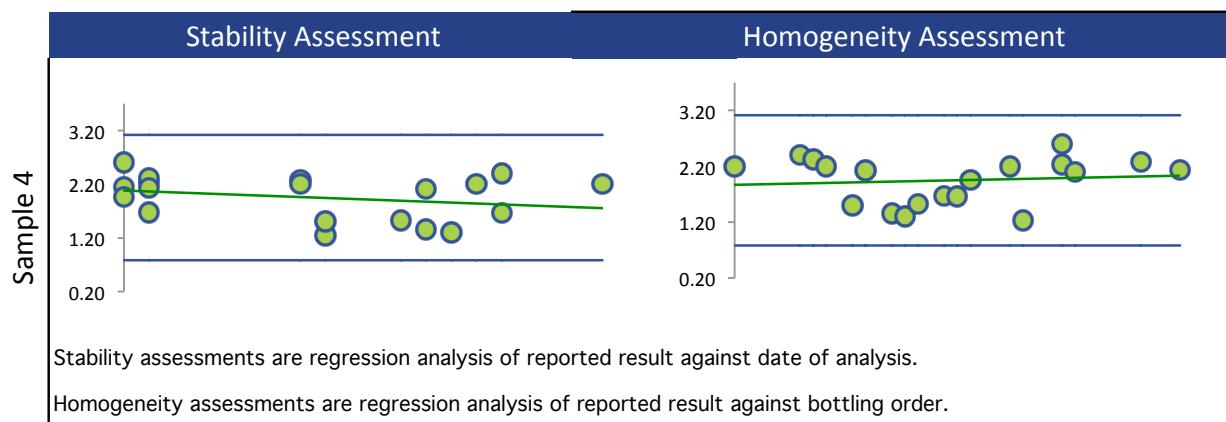


TERBUFOS



Annex A Summary by Analyte

TERBUFOS



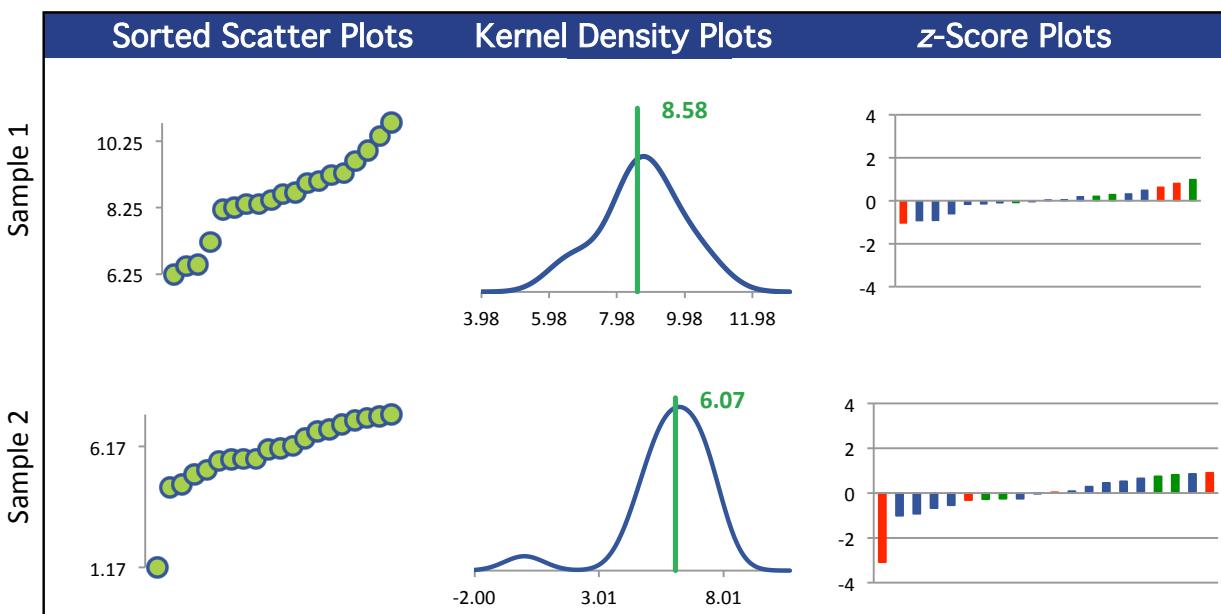
TRIFLURALIN**Summary Statistics**

Statistic	C22-1	C22-2	C22-3	C22-4
N	19	20	20	20
Median $\mu\text{g/L}$	8.67	6.08	1.66	3.05
Robust Mean $\mu\text{g/L}$	8.58	6.07	1.65	3.03
U $\mu\text{g/L}$	0.401	0.324	0.0813	0.172
Robust Standard Deviation $\mu\text{g/L}$	1.40	1.16	0.291	0.615
Regression Standard Deviation $\mu\text{g/L}$	2.24	1.59	0.442	0.800
Stability Flag				
Homogeneity Flag				
Standard Deviation Used (SDPA) $\mu\text{g/L}$	2.24	1.59	0.442	0.800
Outliers	0	0	0	0
$ z > 3.0$	0	1	0	1
$2 < z < 3$	0	0	0	0

Methods Used

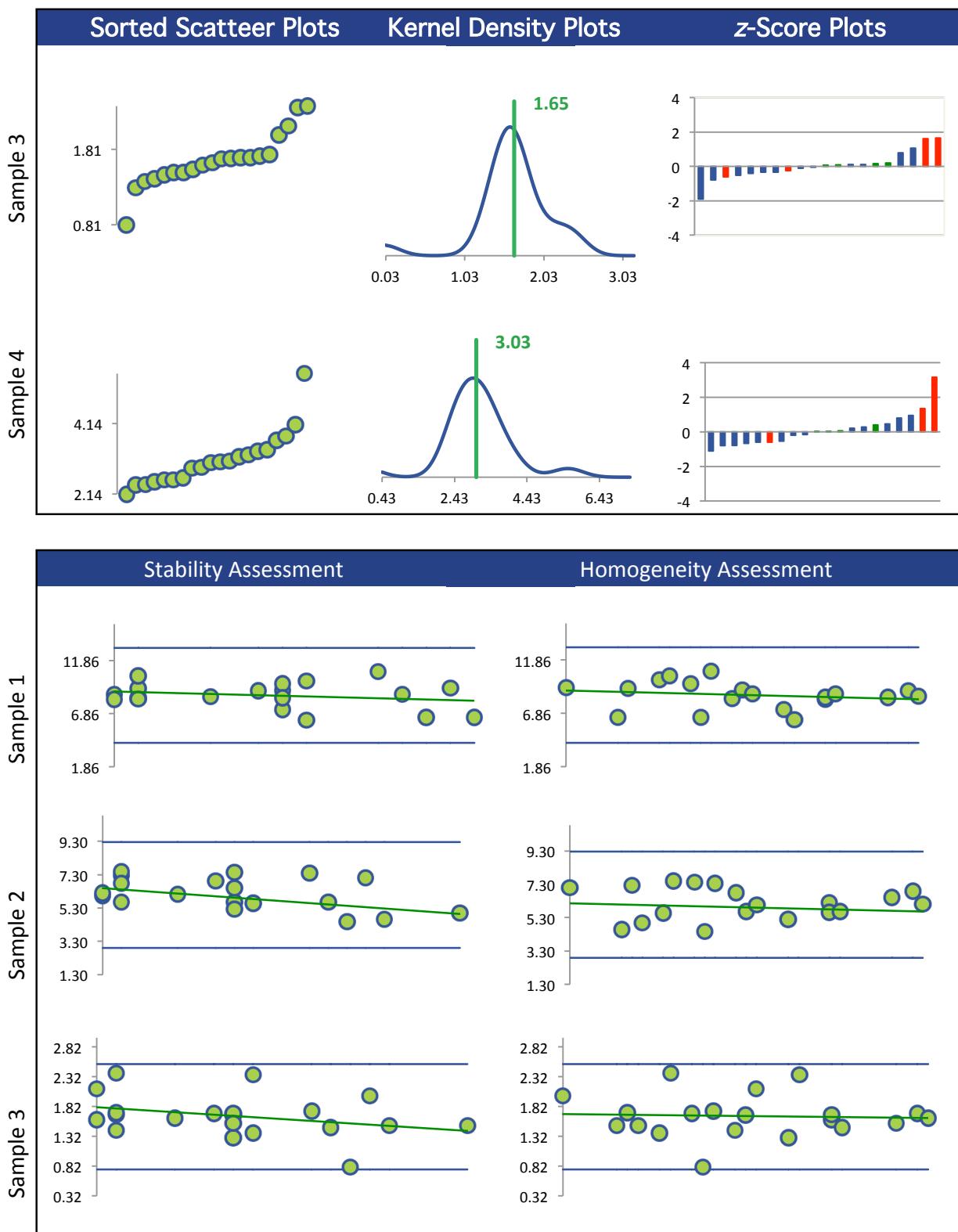
Method	C22-1	C22-2	C22-3	C22-4
GC/MS/MS (Blue)	3	3	3	3
GC/MS (Red)	10	11	11	11
GC/ECD (Green)	2	2	2	2
LC-MS/MS (Orange)	4	4	4	4

All summary stats and the plots below are based on the data excluding any flagged outliers



Annex A Summary by Analyte

TRIFLURALIN



Annex A Summary by Analyte

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