

# **Test Group Summary Report**

## **C16 Volatile Organic Compounds in Water**

### **June 2024 PT Round**

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**Issued: July 30, 2024**

## Table of Contents

<b>1.0</b>	<b>The Proficiency Testing Report .....</b>	<b>1</b>
<b>2.0</b>	<b>Definitions.....</b>	<b>1</b>
<b>3.0</b>	<b>Scoring System .....</b>	<b>1</b>
<b>3.1</b>	<b>Homogeneity and Stability Assessment .....</b>	<b>2</b>
<b>3.2</b>	<b>The z score .....</b>	<b>2</b>
<b>3.2</b>	<b>Composite(PT) Score .....</b>	<b>2</b>
<b>3.3</b>	<b>Identifying Bias.....</b>	<b>2</b>
<b>3.4</b>	<b>Deviations from Evaluation Procedure .....</b>	<b>3</b>
<b>4.0</b>	<b>PT Round Specific Data Summary.....</b>	<b>3</b>
<b>4.1</b>	<b>Summary statistics.....</b>	<b>3</b>
<b>4.2</b>	<b>z- Score Plots.....</b>	<b>3</b>
<b>4.3</b>	<b>kernel density plots .....</b>	<b>3</b>
<b>4.4</b>	<b>stability and homogeneity Plots .....</b>	<b>3</b>
<b>4.5</b>	<b>Box-and-Whisker Plots.....</b>	<b>3</b>
<b>4.6</b>	<b>Historic Comparison Plot .....</b>	<b>3</b>
	<b>Annex A Summary by Analyte .....</b>	<b>4</b>

## 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><math>\pm u</math>:</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](http://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:

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

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 $\leq 2$	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<sup>1</sup>

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<sup>th</sup> and 95<sup>th</sup> 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.

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<sup>1</sup> For some reports, the colour coding for methods is not being displayed properly.

## Annex A Summary by Analyte

### 1,1,1-TRICHLOROETHANE

#### Summary Statistics

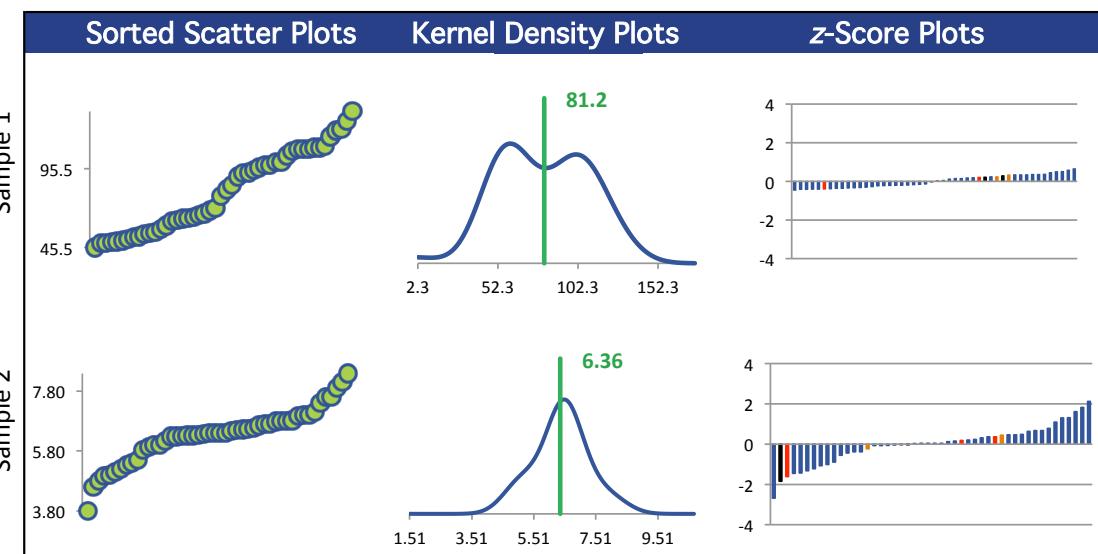
#### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	48	48	48	48
Median $\mu\text{g/L}$	80.2	6.40	61.7	31.4
Robust Mean $\mu\text{g/L}$	81.2	6.36	62.4	31.3
U $\mu\text{g/L}$	5.20	0.154	1.19	0.599
Robust Standard Deviation $\mu\text{g/L}$	28.8	0.855	6.60	3.32
Regression Standard Deviation $\mu\text{g/L}$	12.2	0.954	9.36	4.69
Stability Flag				
Homogeneity Flag	Homogeneity			
Standard Deviation Used (SDPA) $\mu\text{g/L}$	77.9	0.954	9.36	4.69
Outliers	0	0	0	0
$ z  > 3.0$	0	0	0	0
$2 <  z  < 3$	0	2	2	1

#### Methods Used

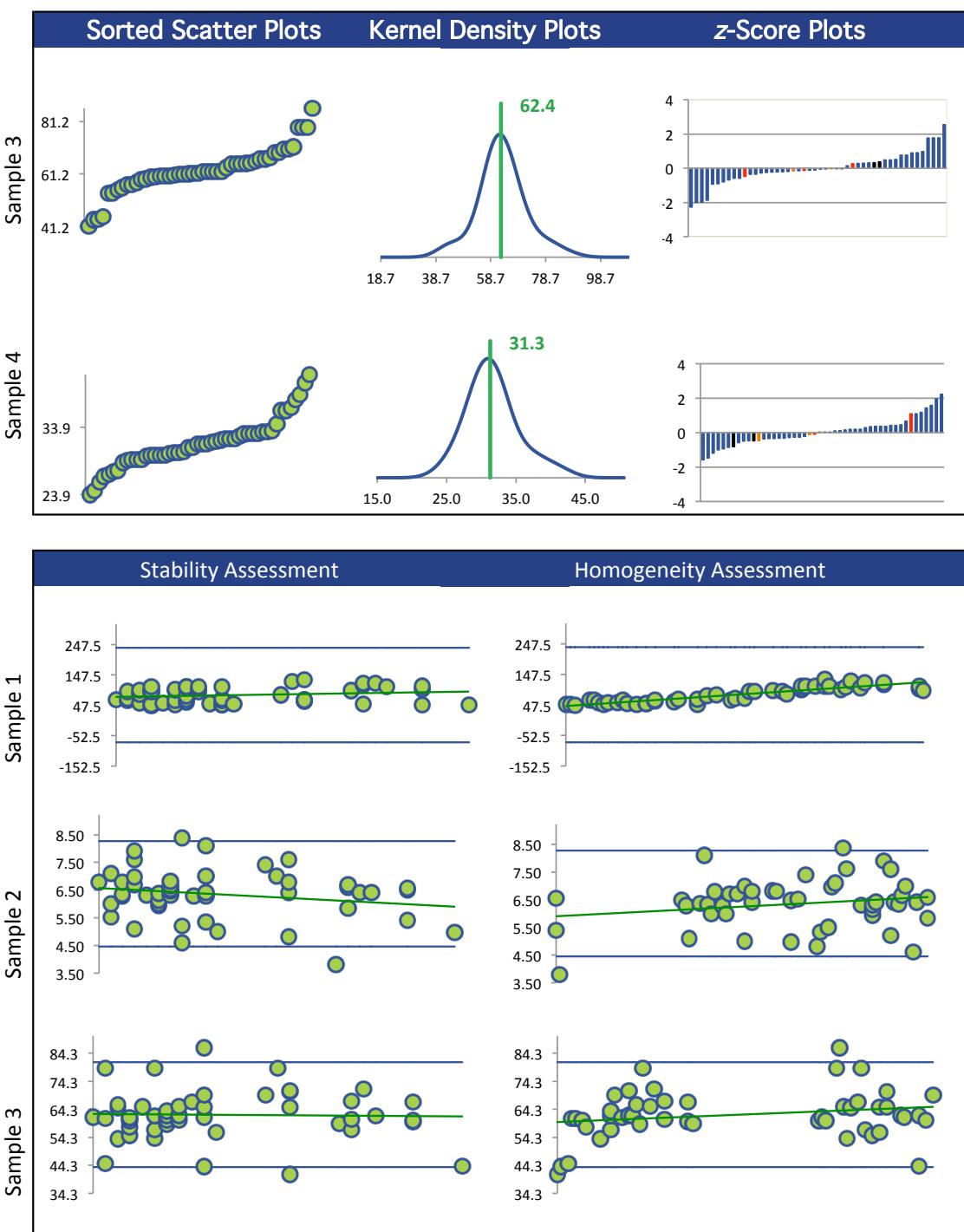
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	26	26	26	26
GC/MS - HEADSPACE (Red)	19	19	19	19
GC/FID - PURGE AND TRAP (Green)	1	1	1	1
GC/MS (Orange)	2	2	2	2

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



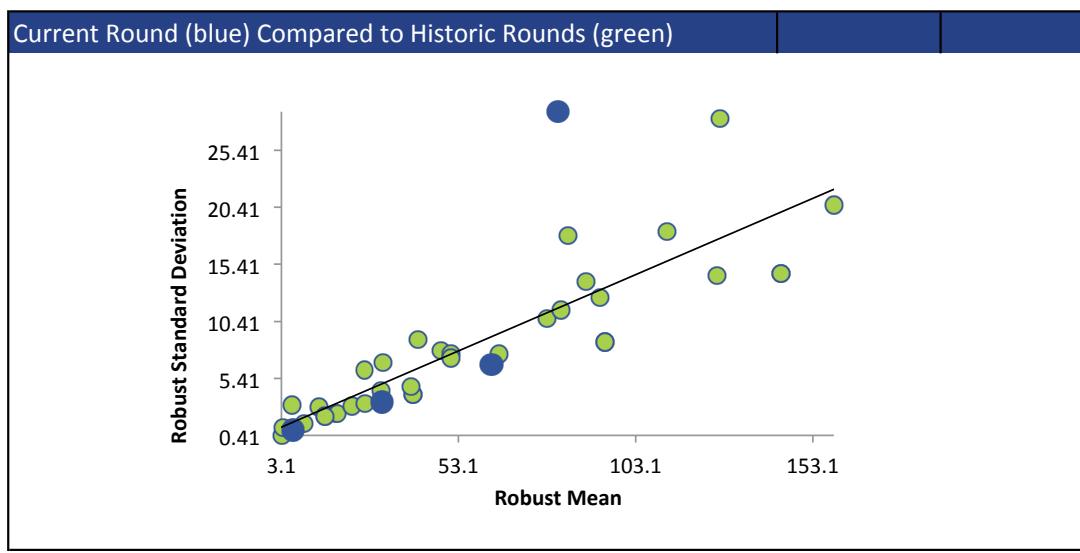
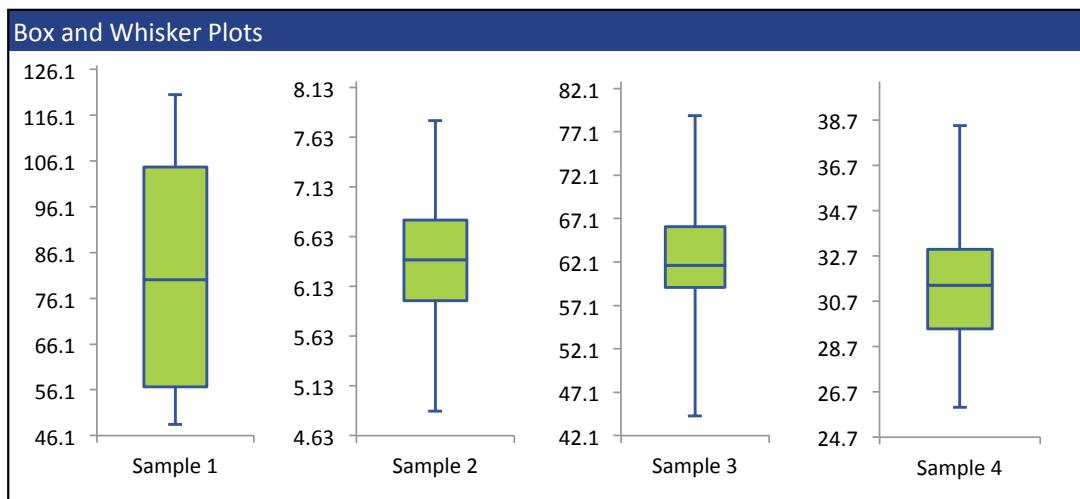
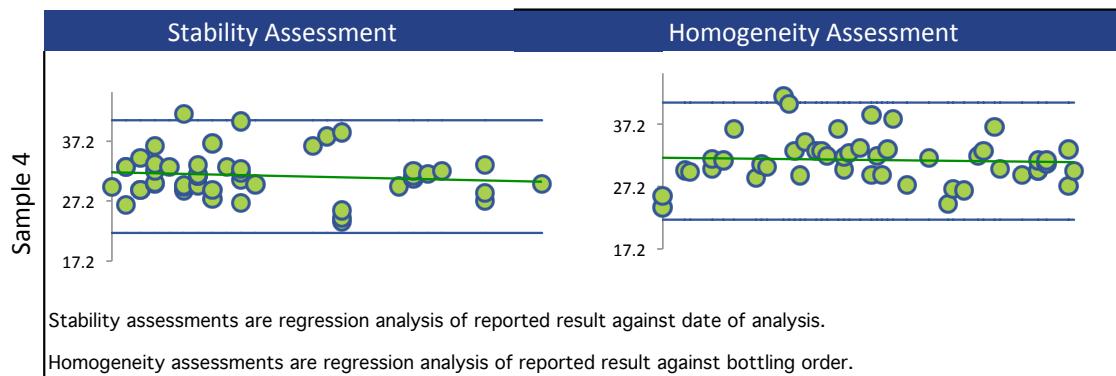
Annex A Summary by Analyte

1,1,1-TRICHLOROETHANE



## Annex A Summary by Analyte

### 1,1,1-TRICHLOROETHANE



## Annex A Summary by Analyte

### 1,1,2,2-TETRACHLOROETHANE

#### Summary Statistics

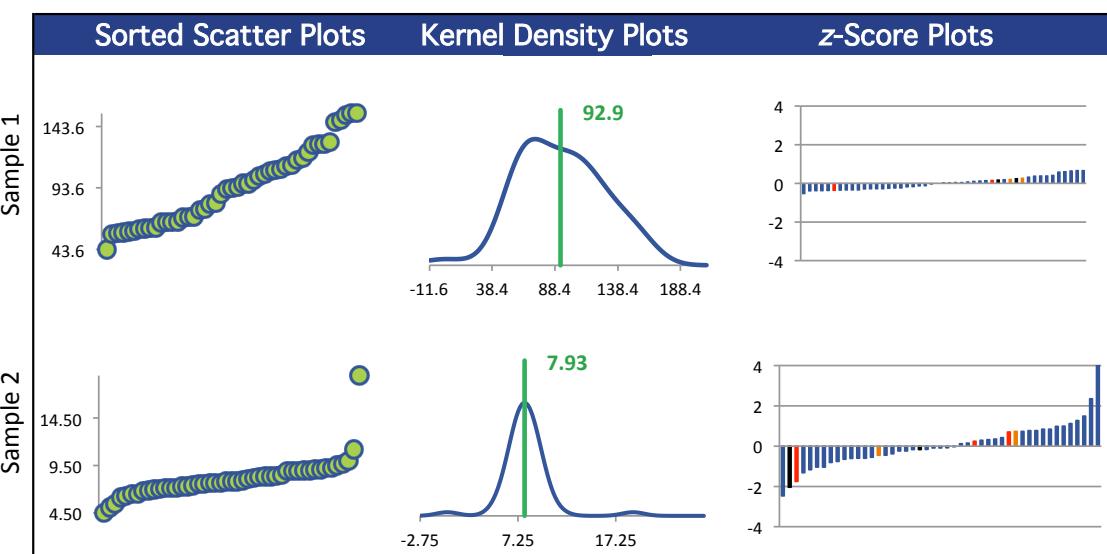
#### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	47	47	47	47
Median $\mu\text{g/L}$	93.4	7.80	72.0	38.0
Robust Mean $\mu\text{g/L}$	92.9	7.93	73.7	38.3
U $\mu\text{g/L}$	6.09	0.232	1.97	1.08
Robust Standard Deviation $\mu\text{g/L}$	33.4	1.27	10.8	5.91
Regression Standard Deviation $\mu\text{g/L}$	16.3	1.39	12.9	6.71
Stability Flag				
Homogeneity Flag	Homogeneity		Homogeneity	
Standard Deviation Used (SDPA) $\mu\text{g/L}$	91.8	1.39	17.8	6.71
Outliers	0	0	0	0
$ z  > 3.0$	0	1	1	1
$2 <  z  < 3$	0	3	0	2

#### Methods Used

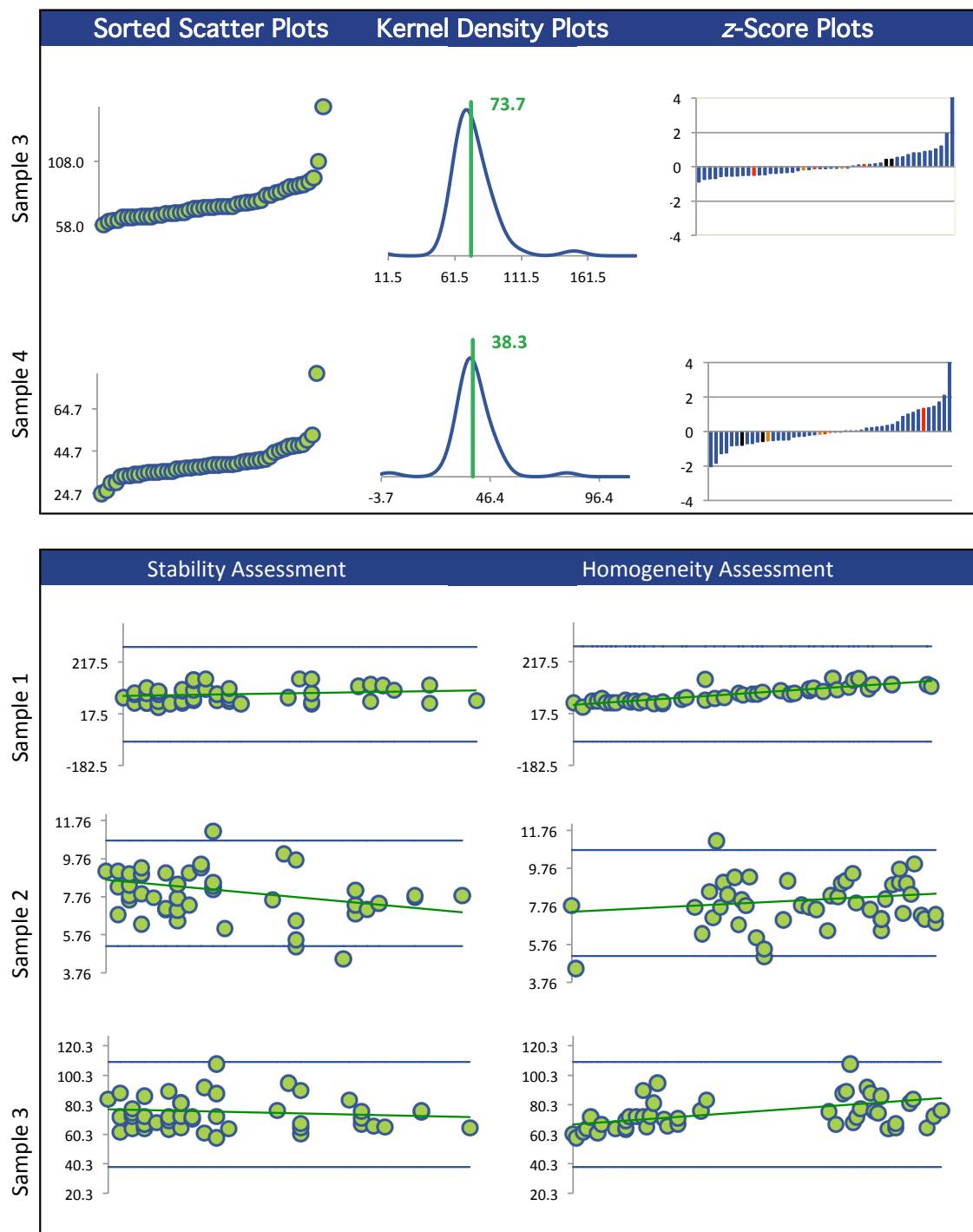
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	27	27	27	27
GC/MS - HEADSPACE (Red)	16	16	16	16
GC/FID - PURGE AND TRAP (Green)	1	1	1	1
GC/MS/MS - HEADSPACE (Orange)	1	1	1	1
GC/MS (Black)	2	2	2	2

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



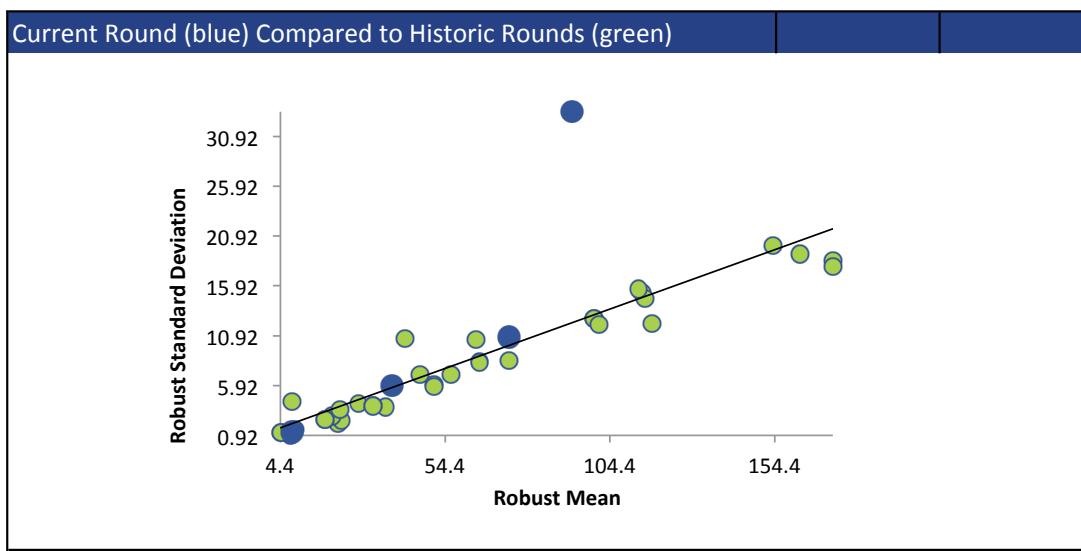
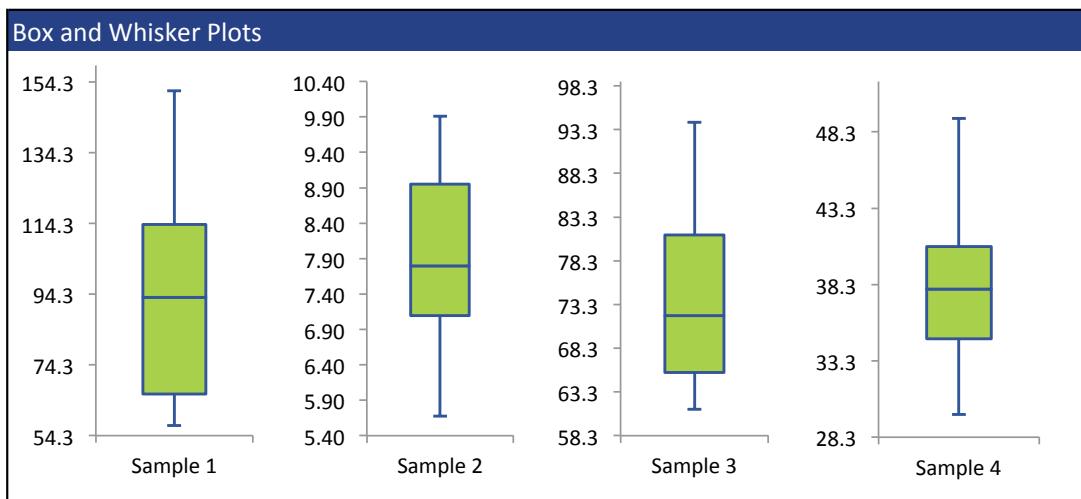
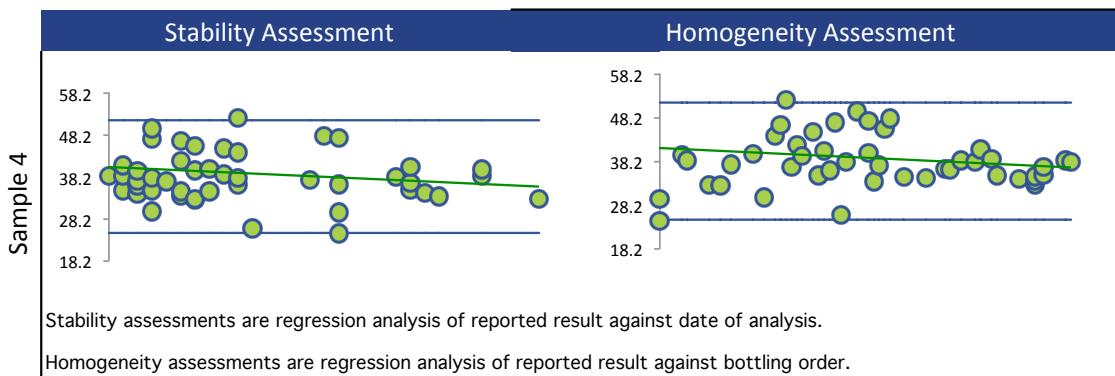
Annex A Summary by Analyte

1,1,2,2-TETRACHLOROETHANE



## Annex A Summary by Analyte

### 1,1,2,2-TETRACHLOROETHANE



## Annex A Summary by Analyte

### 1,1,2-TRICHLOROETHANE

#### Summary Statistics

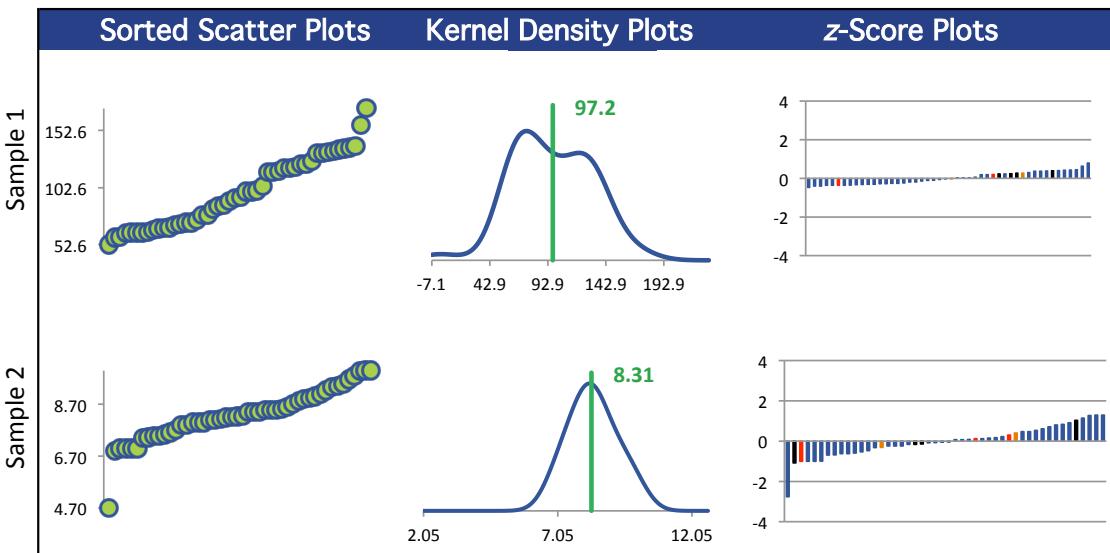
#### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	48	48	48	48
Median $\mu\text{g/L}$	93.8	8.25	76.9	39.8
Robust Mean $\mu\text{g/L}$	97.2	8.31	76.9	39.4
$U \mu\text{g/L}$	6.03	0.172	1.49	0.736
Robust Standard Deviation $\mu\text{g/L}$	33.4	0.952	8.28	4.08
Regression Standard Deviation $\mu\text{g/L}$	14.6	1.25	11.5	5.90
Stability Flag				
Homogeneity Flag	Homogeneity	Homogeneity		
Standard Deviation Used ( $SDPA \mu\text{g/L}$ )	93.6	1.31	11.5	5.90
Outliers	0	0	0	0
$ z  > 3.0$	0	0	0	0
$2 <  z  < 3$	0	1	2	2

#### Methods Used

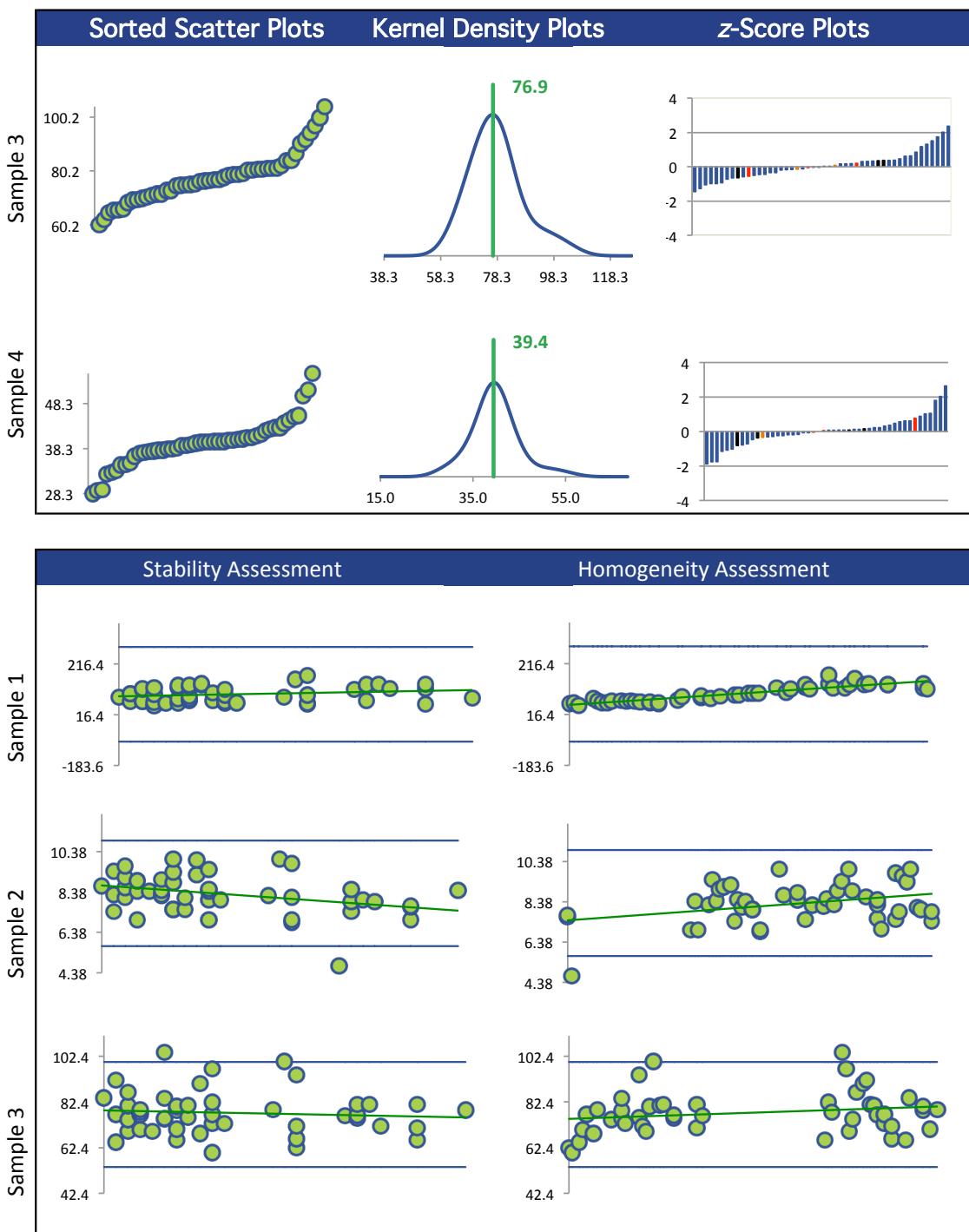
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	26	26	26	26
GC/MS - HEADSPACE (Red)	19	19	19	19
GC/FID - PURGE AND TRAP (Green)	1	1	1	1
GC/MS (Orange)	2	2	2	2

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



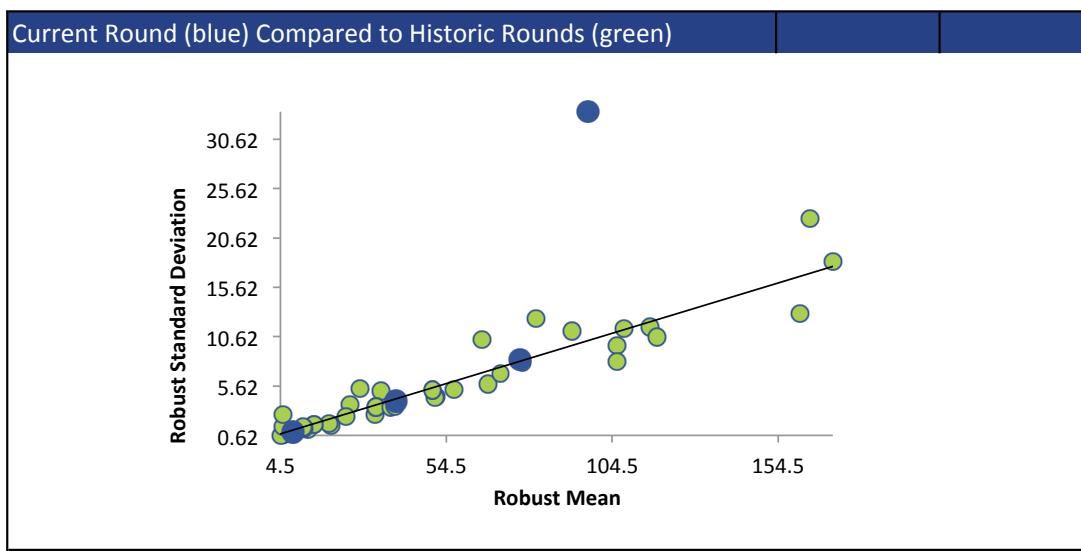
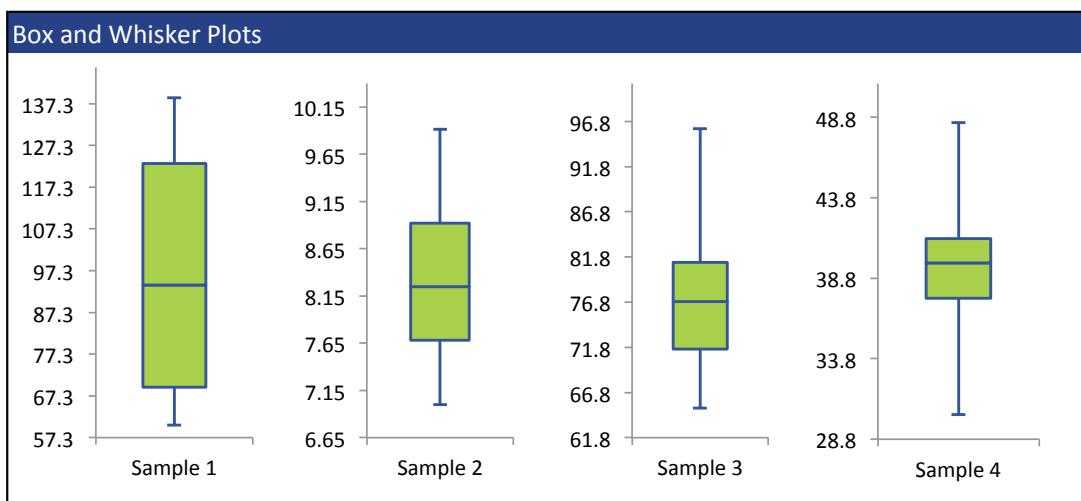
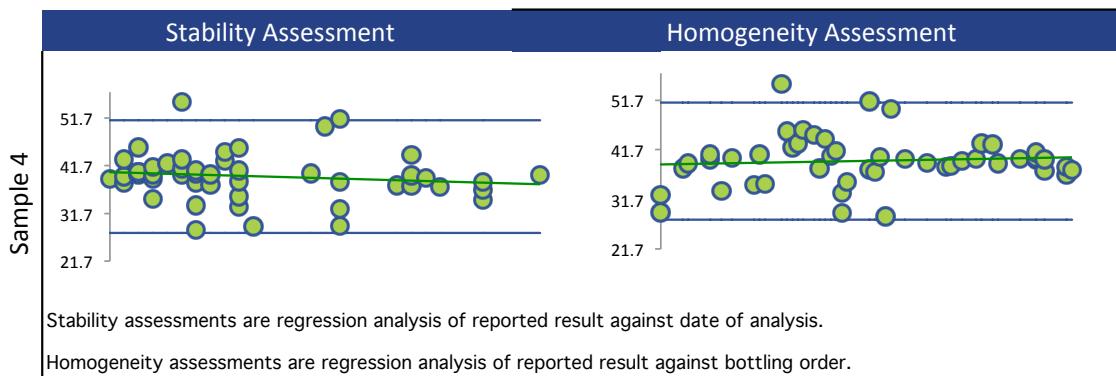
Annex A Summary by Analyte

1,1,2-TRICHLOROETHANE



## Annex A Summary by Analyte

### 1,1,2-TRICHLOROETHANE



## Annex A Summary by Analyte

### 1,1-DICHLOROETHANE

#### Summary Statistics

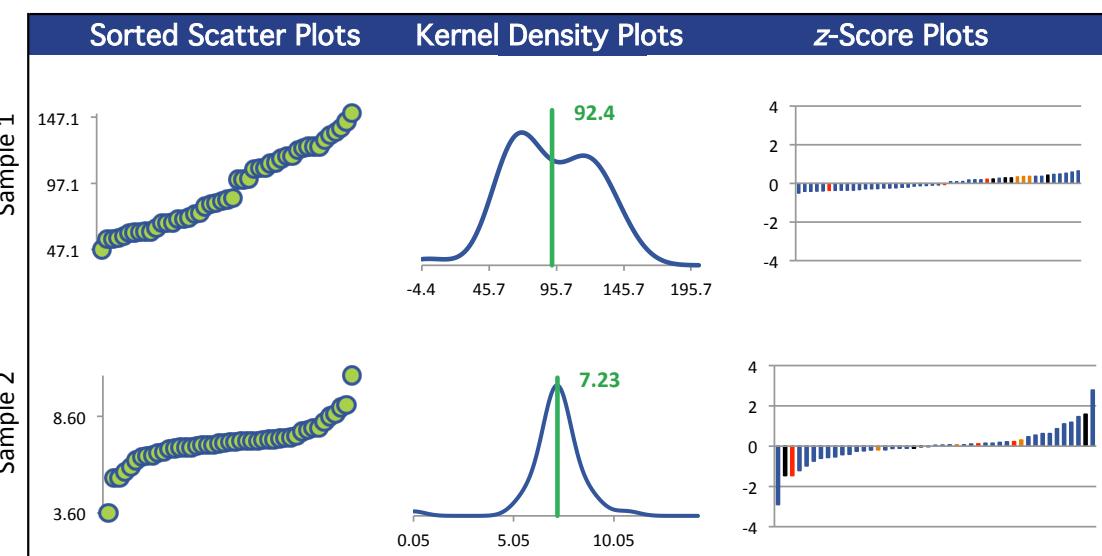
#### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	47	45	47	47
Median $\mu\text{g/L}$	84.9	7.28	71.0	36.1
Robust Mean $\mu\text{g/L}$	92.4	7.23	70.9	36.1
$U \mu\text{g/L}$	6.02	0.141	1.51	0.613
Robust Standard Deviation $\mu\text{g/L}$	33.0	0.758	8.27	3.36
Regression Standard Deviation $\mu\text{g/L}$	13.9	1.08	10.6	5.42
Stability Flag				
Homogeneity Flag	Homogeneity	Homogeneity		
Standard Deviation Used ( $SDPA \mu\text{g/L}$ )	88.2	1.25	10.6	5.42
Outliers	0	0	0	0
$ z  > 3.0$	0	0	0	0
$2 <  z  < 3$	0	2	3	1

#### Methods Used

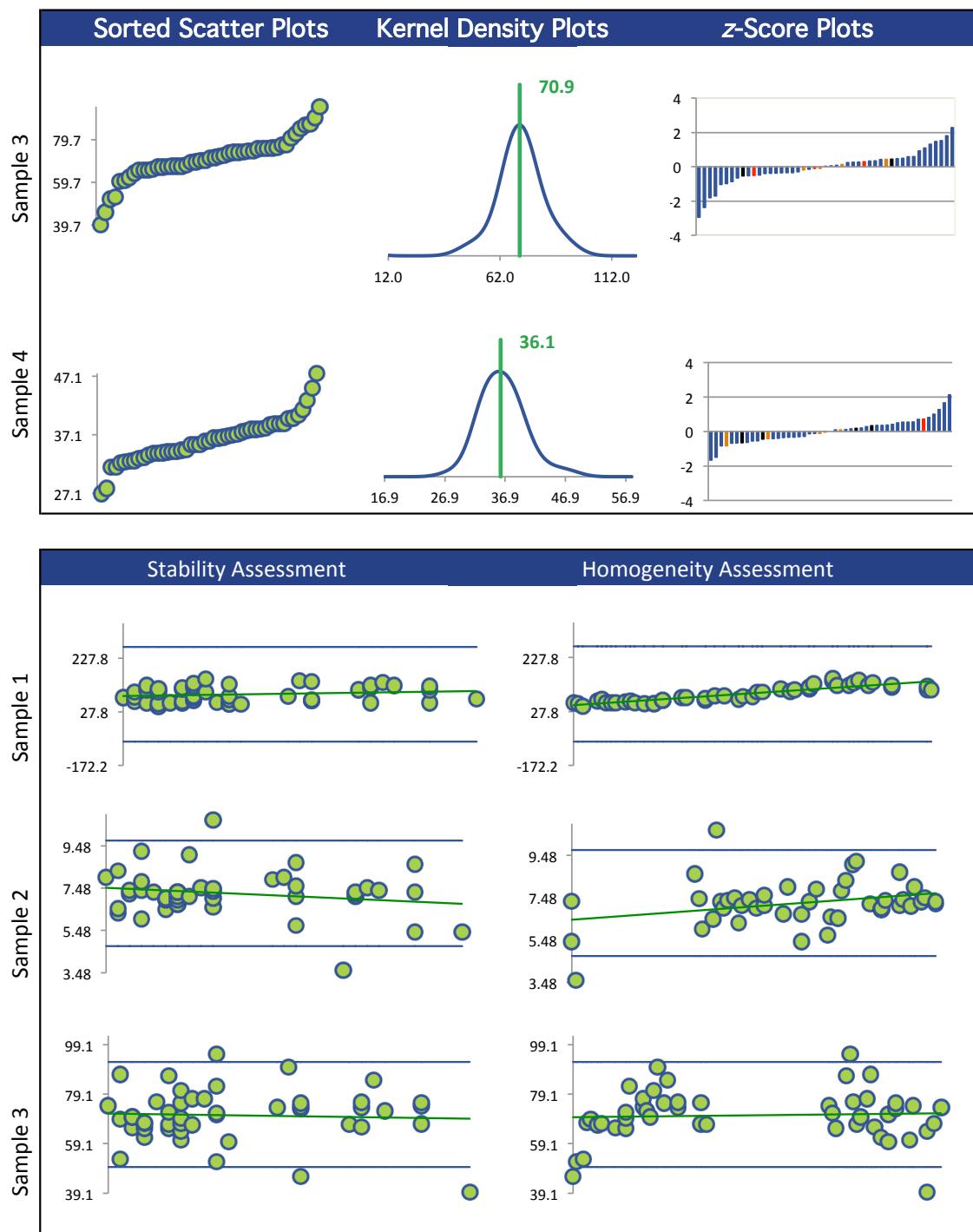
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	25	24	25	25
GC/MS - HEADSPACE (Red)	19	19	19	19
GC/FID - PURGE AND TRAP (Green)	1	1	1	1
GC/MS (Orange)	2	1	2	2

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



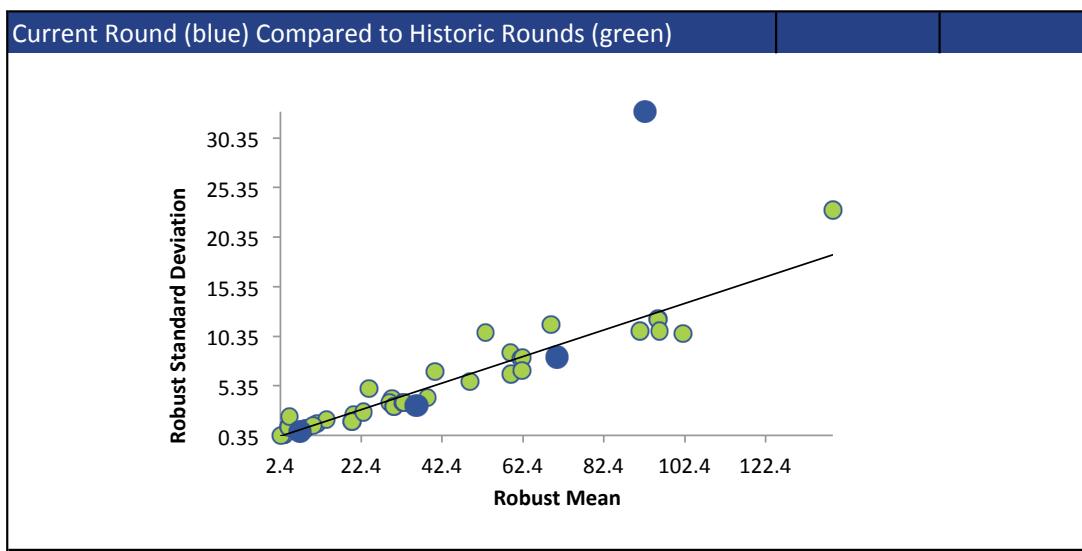
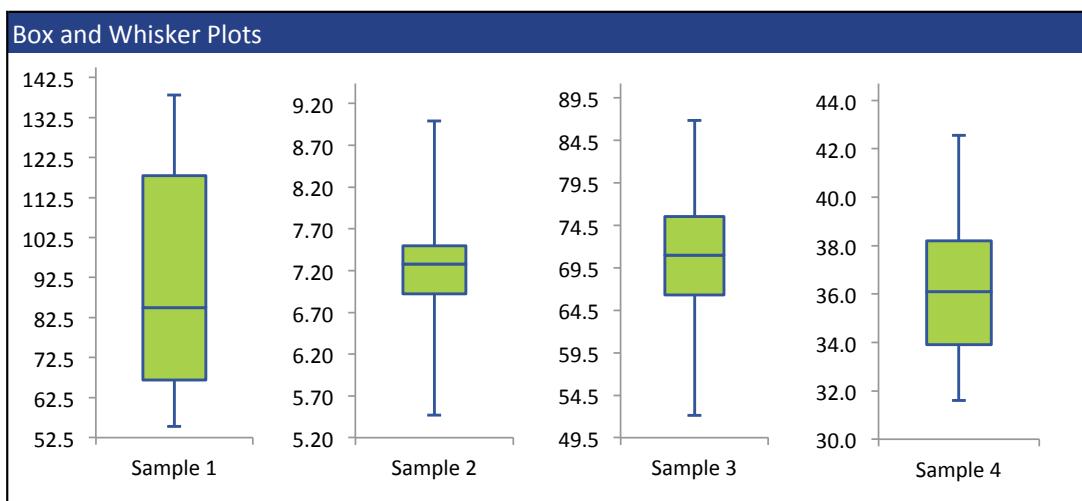
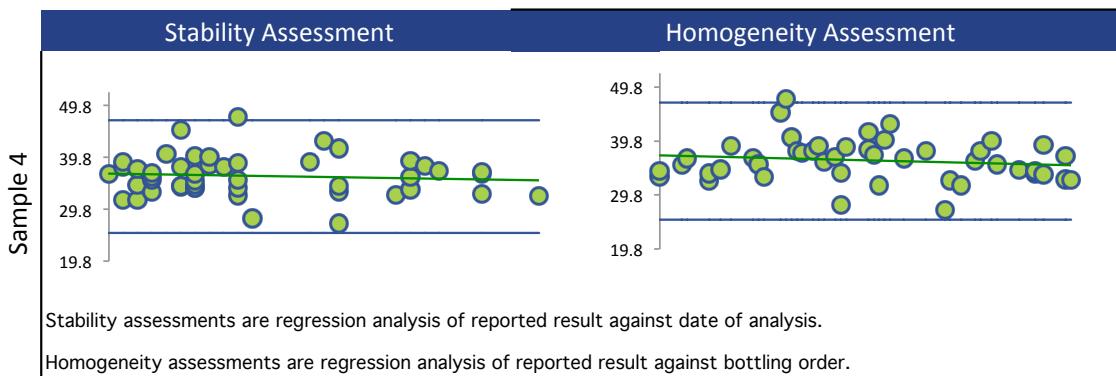
Annex A Summary by Analyte

1,1-DICHLOROETHANE



## Annex A Summary by Analyte

### 1,1-DICHLOROETHANE



## Annex A Summary by Analyte

### 1,1-DICHLOROETHYLENE

#### Summary Statistics

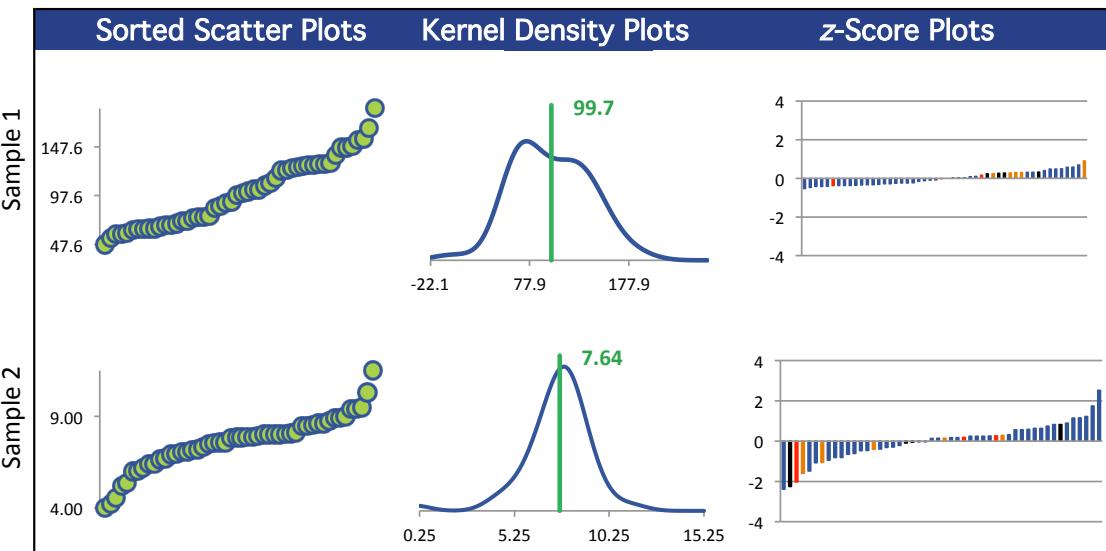
#### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	50	50	50	50
Median $\mu\text{g/L}$	99.2	7.86	75.7	38.1
Robust Mean $\mu\text{g/L}$	99.7	7.64	76.2	39.0
U $\mu\text{g/L}$	6.70	0.225	1.84	0.962
Robust Standard Deviation $\mu\text{g/L}$	37.9	1.27	10.4	5.44
Regression Standard Deviation $\mu\text{g/L}$	19.9	1.53	15.2	7.80
Stability Flag				
Homogeneity Flag	Homogeneity			
Standard Deviation Used (SDPA) $\mu\text{g/L}$	96.0	1.53	15.2	7.80
Outliers	0	0	0	0
$ z  > 3.0$	0	0	0	0
$2 <  z  < 3$	0	4	2	2

#### Methods Used

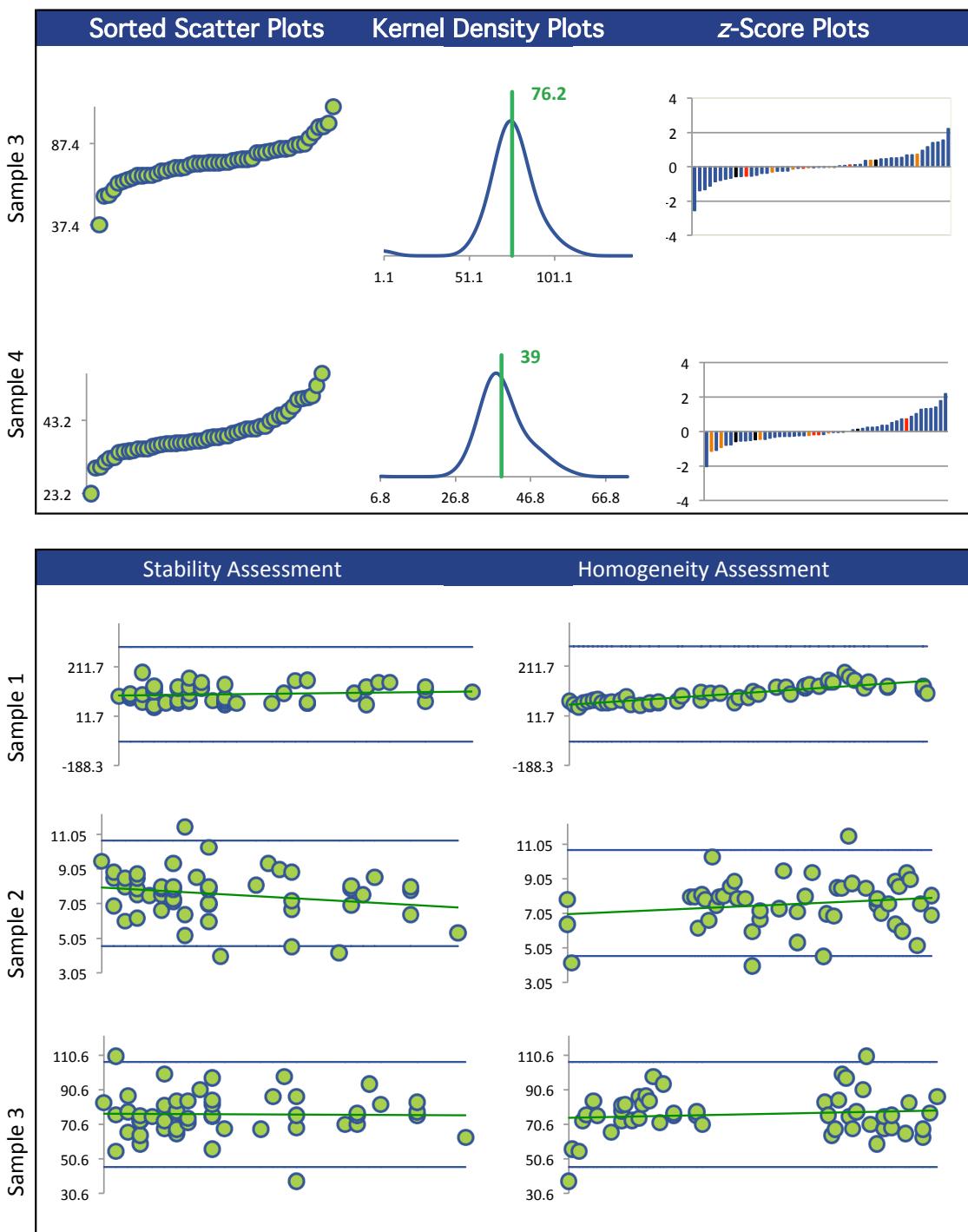
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	27	27	27	27
GC/MS - HEADSPACE (Red)	19	19	19	19
GC/FID - PURGE AND TRAP (Green)	1	1	1	1
GC/MS/MS - HEADSPACE (Orange)	1	1	1	1
GC/MS (Black)	2	2	2	2

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



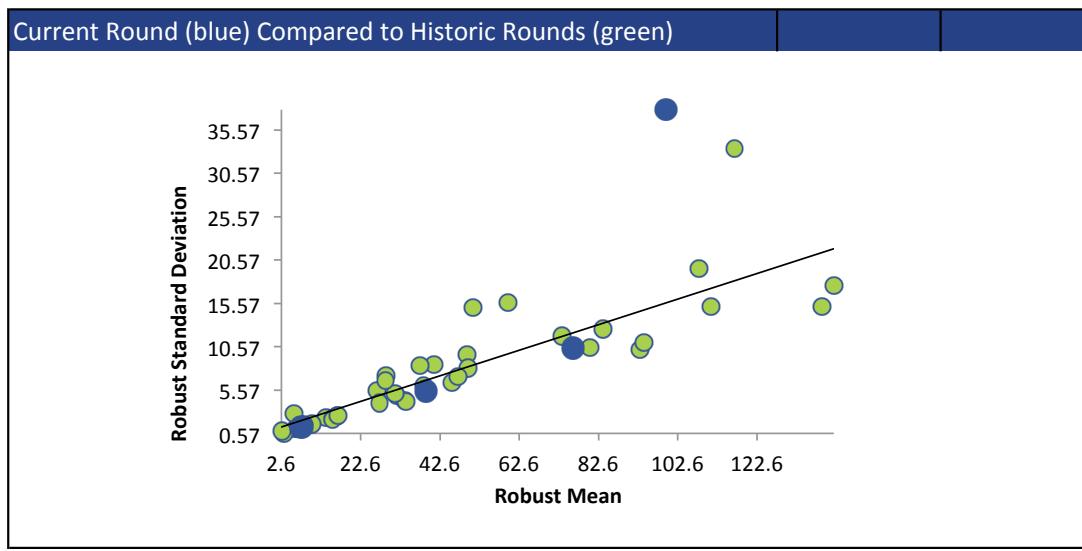
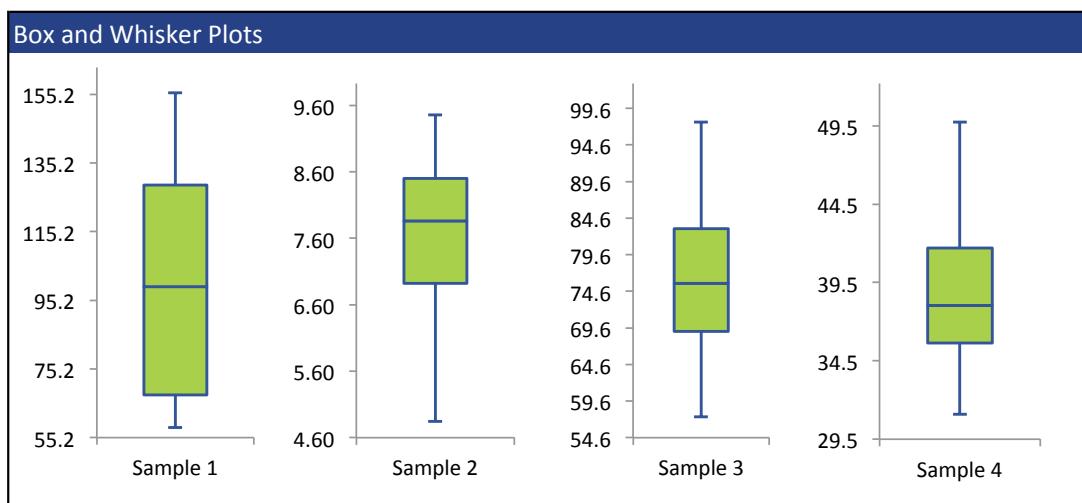
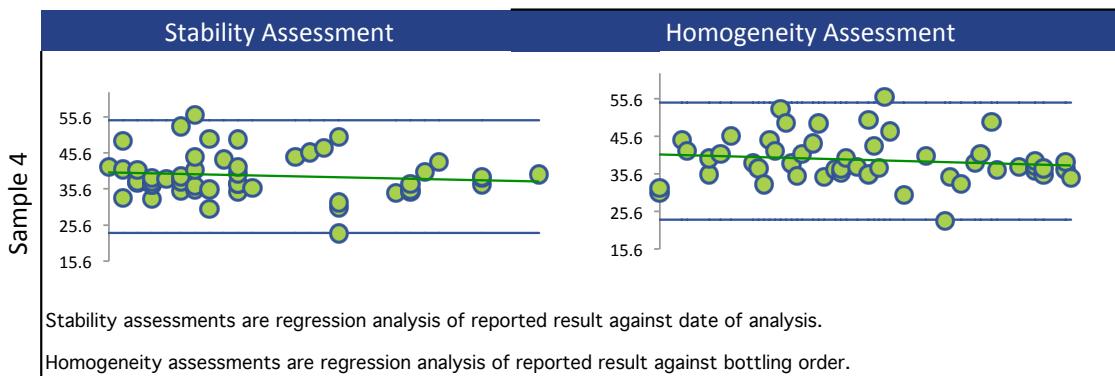
Annex A Summary by Analyte

1,1-DICHLOROETHYLENE



## Annex A Summary by Analyte

### 1,1-DICHLOROETHYLENE



## Annex A Summary by Analyte

### 1,2-DICHLOROBENZENE

#### Summary Statistics

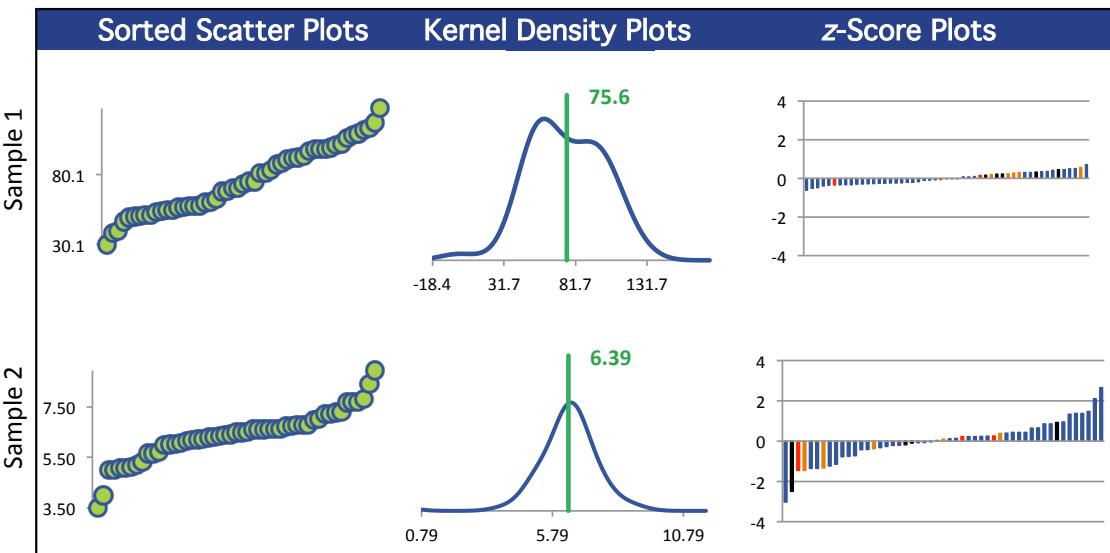
#### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	51	51	51	51
Median $\mu\text{g/L}$	73.1	6.48	60.1	30.4
Robust Mean $\mu\text{g/L}$	75.6	6.39	59.9	31.0
U $\mu\text{g/L}$	4.60	0.161	1.19	0.555
Robust Standard Deviation $\mu\text{g/L}$	26.3	0.920	6.79	3.17
Regression Standard Deviation $\mu\text{g/L}$	11.3	0.958	8.98	4.65
Stability Flag				
Homogeneity Flag	Homogeneity			
Standard Deviation Used (SDPA) $\mu\text{g/L}$	72.8	0.958	8.98	4.65
Outliers	0	0	0	0
$ z  > 3.0$	0	1	0	1
$2 <  z  < 3$	0	3	2	3

#### Methods Used

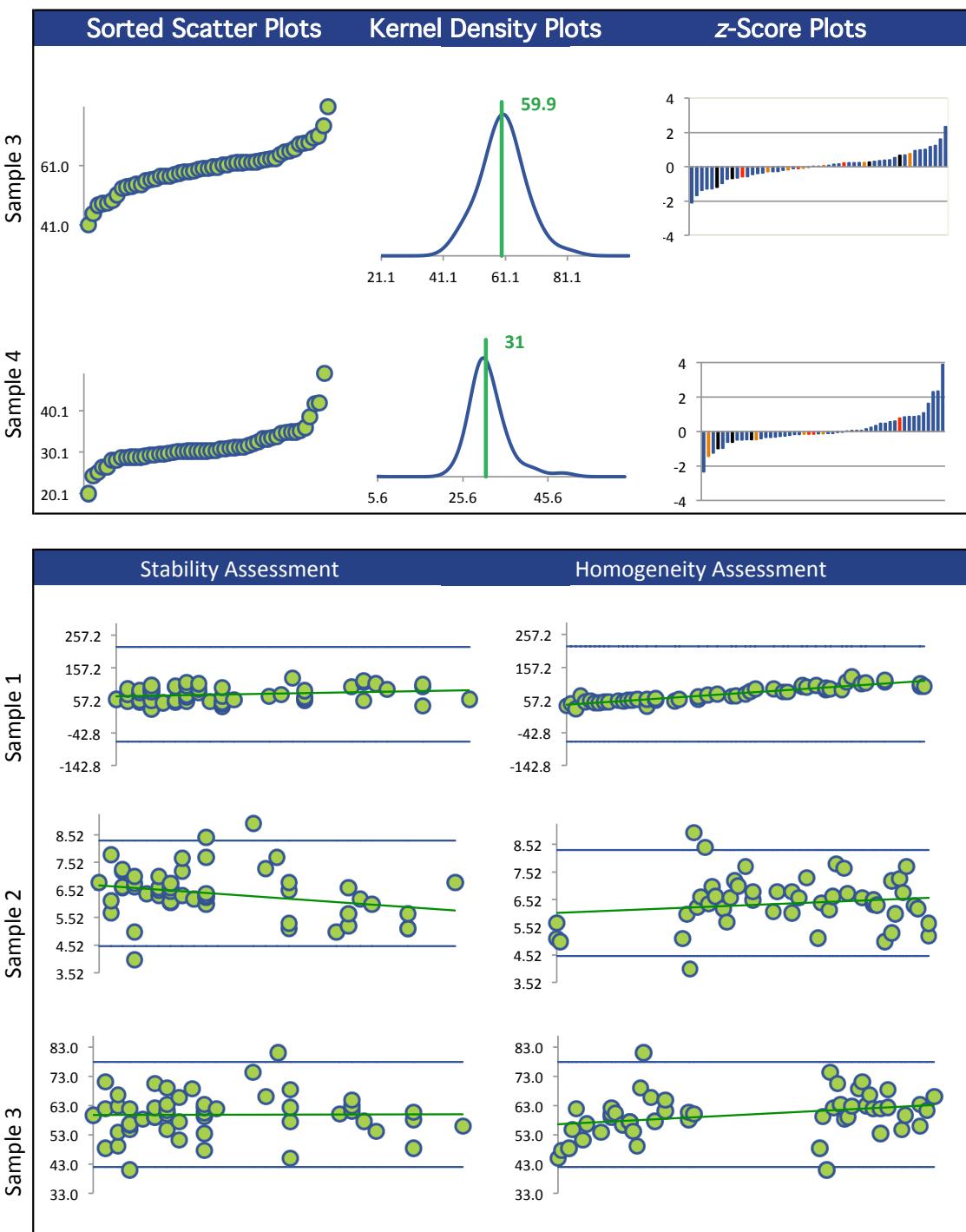
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	28	28	28	28
GC/MS - HEADSPACE (Red)	19	19	19	19
GC/MS/MS - HEADSPACE (Green)	1	1	1	1
GC/FID - PURGE AND TRAP (Orange)	1	1	1	1
GC/MS (Black)	2	2	2	2

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



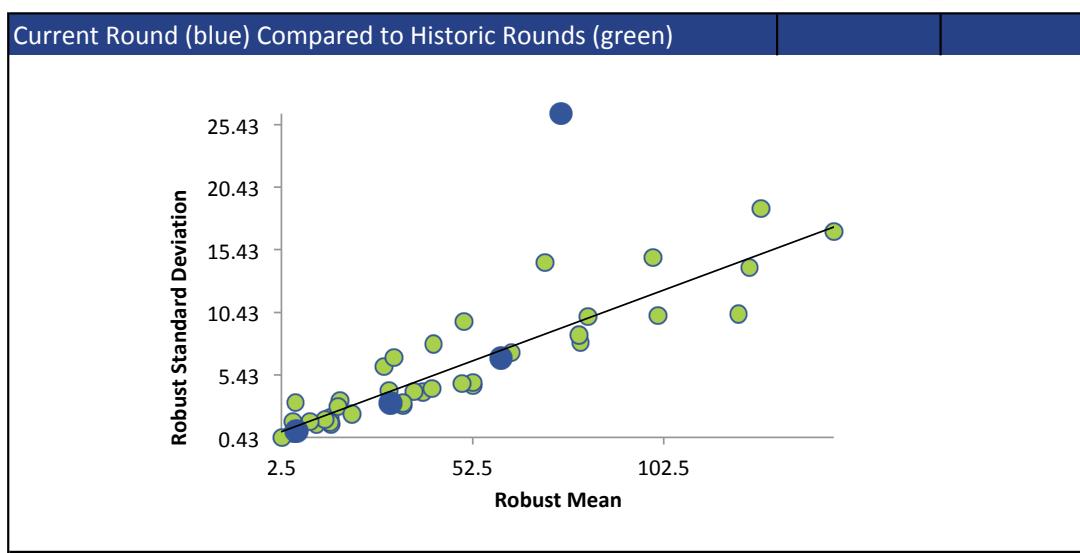
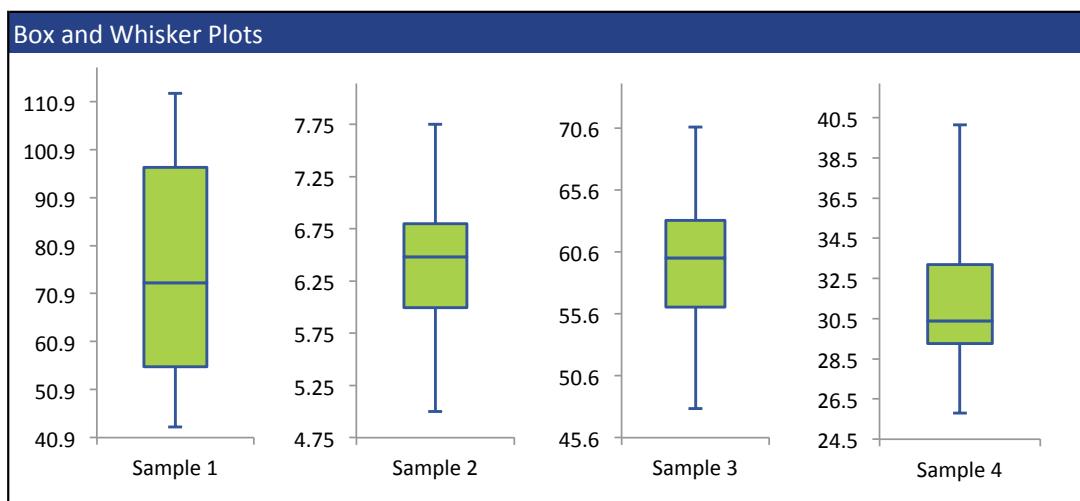
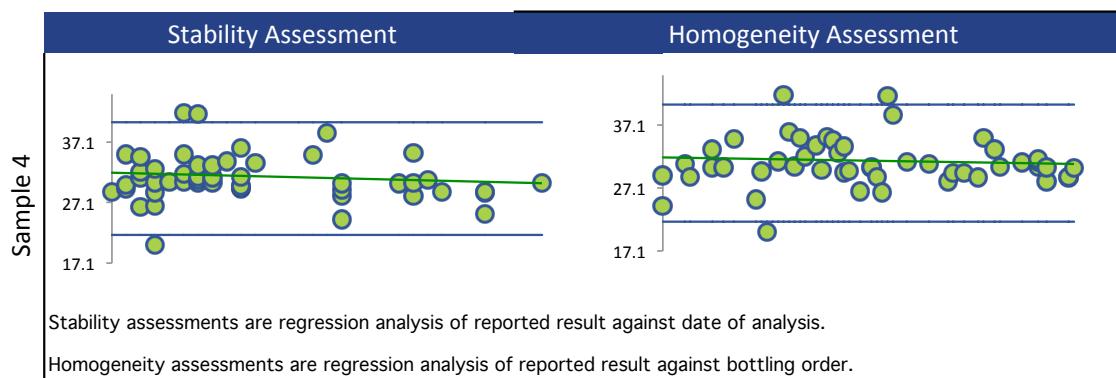
Annex A Summary by Analyte

1,2-DICHLOROBENZENE



## Annex A Summary by Analyte

### 1,2-DICHLOROBENZENE



## Annex A Summary by Analyte

### 1,2-DICHLOROETHANE

#### Summary Statistics

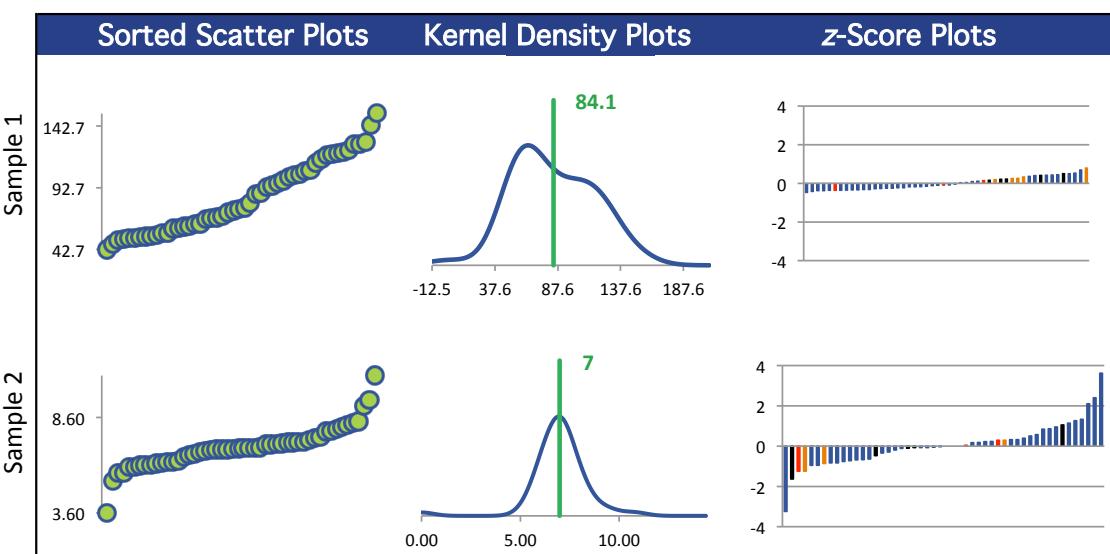
#### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	50	50	50	50
Median $\mu\text{g/L}$	76.0	7.00	66.8	34.1
Robust Mean $\mu\text{g/L}$	84.1	7.00	67.2	34.4
$U \mu\text{g/L}$	5.66	0.156	1.45	0.606
Robust Standard Deviation $\mu\text{g/L}$	32.0	0.885	8.20	3.43
Regression Standard Deviation $\mu\text{g/L}$	12.6	1.05	10.1	5.16
Stability Flag				
Homogeneity Flag	Homogeneity			
Standard Deviation Used ( $SDPA \mu\text{g/L}$ )	85.7	1.05	10.1	5.16
Outliers	0	0	0	0
$ z  > 3.0$	0	2	0	1
$2 <  z  < 3$	0	2	2	1

#### Methods Used

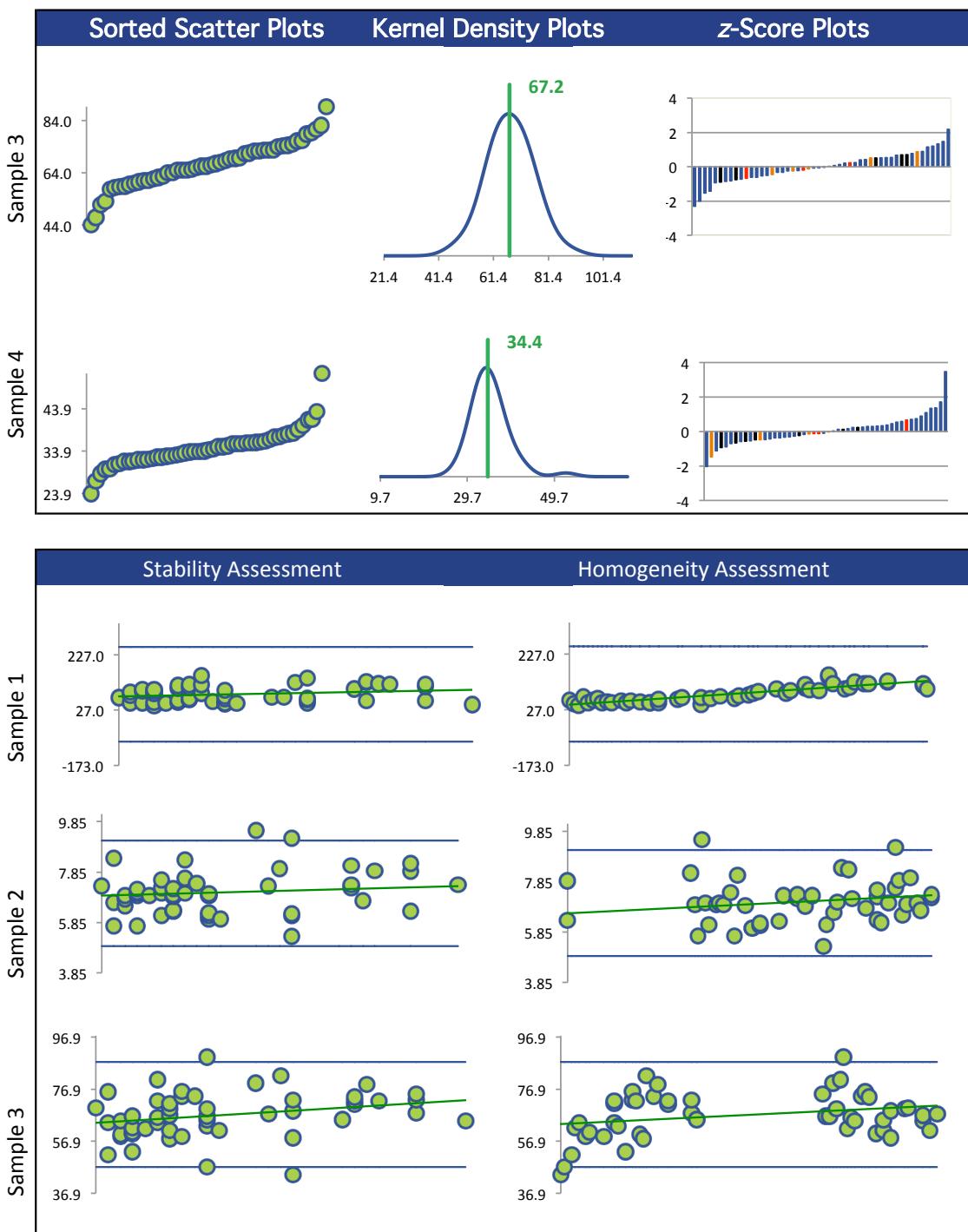
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	27	27	27	27
GC/MS - HEADSPACE (Red)	19	19	19	19
GC/MS/MS - HEADSPACE (Green)	1	1	1	1
GC/FID - PURGE AND TRAP (Orange)	1	1	1	1
GC/MS (Black)	2	2	2	2

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



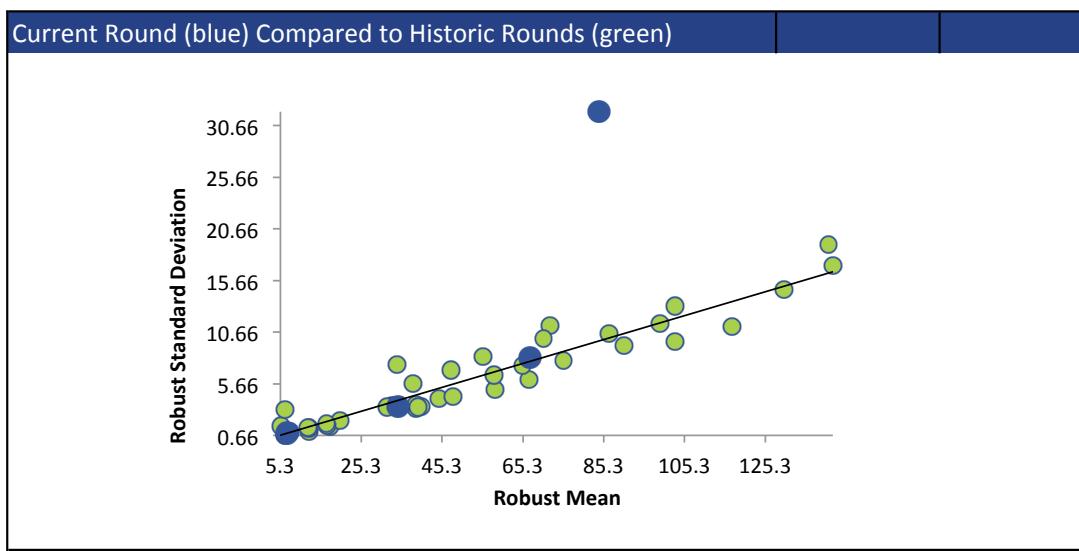
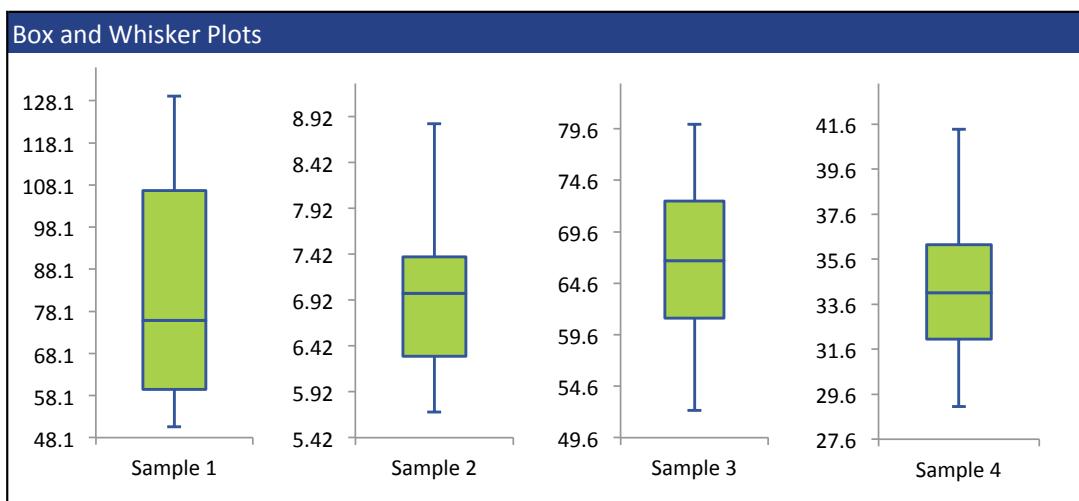
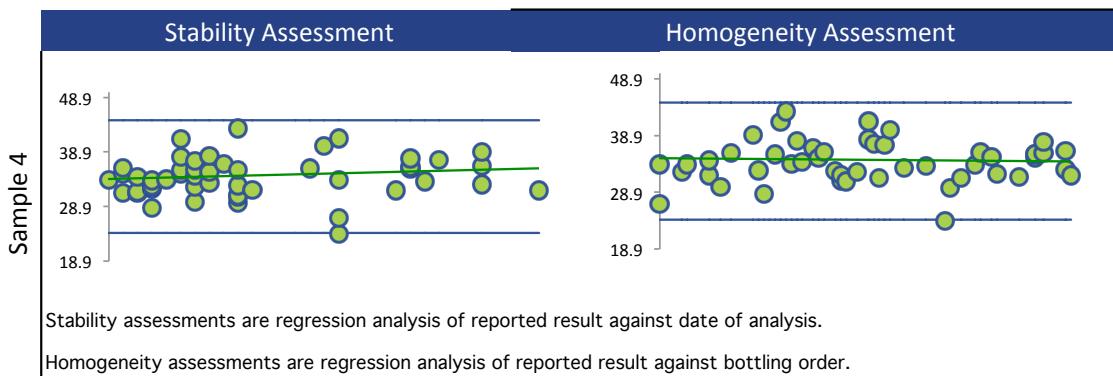
Annex A Summary by Analyte

1,2-DICHLOROETHANE



## Annex A Summary by Analyte

### 1,2-DICHLOROETHANE



## Annex A Summary by Analyte

### 1,2-DICHLOROPROPANE

#### Summary Statistics

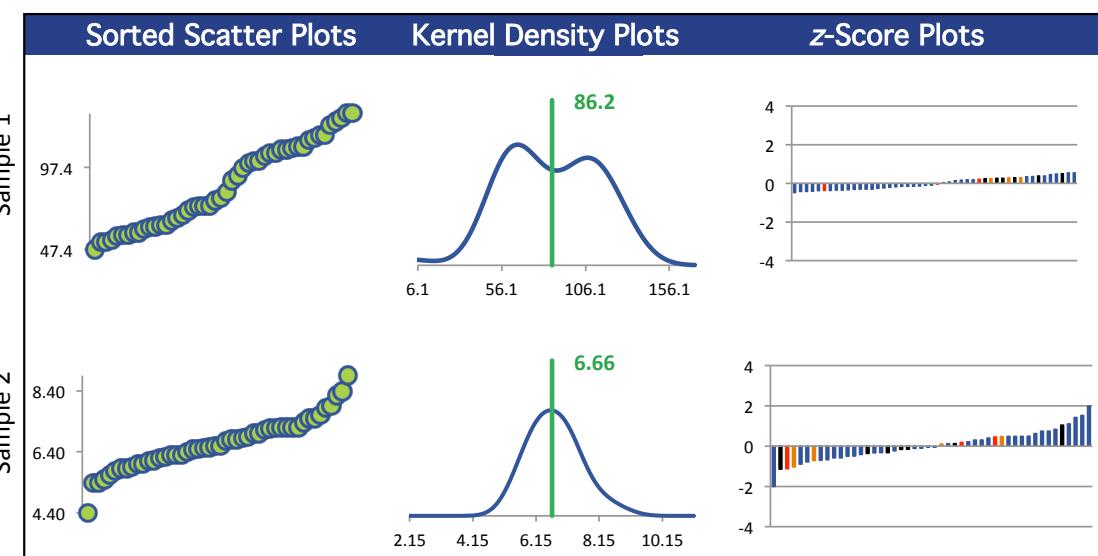
#### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	48	48	48	48
Median $\mu\text{g/L}$	80.3	6.60	65.0	33.0
Robust Mean $\mu\text{g/L}$	86.2	6.66	65.3	33.3
$U \mu\text{g/L}$	5.25	0.147	0.877	0.516
Robust Standard Deviation $\mu\text{g/L}$	29.1	0.817	4.86	2.86
Regression Standard Deviation $\mu\text{g/L}$	12.9	1.00	9.79	5.00
Stability Flag	Stability			
Homogeneity Flag	Homogeneity			
Standard Deviation Used ( $SDPA$ ) $\mu\text{g/L}$	79.3	1.13	9.79	5.00
Outliers	0	0	0	0
$ z  > 3.0$	0	0	0	0
$2 <  z  < 3$	0	0	2	1

#### Methods Used

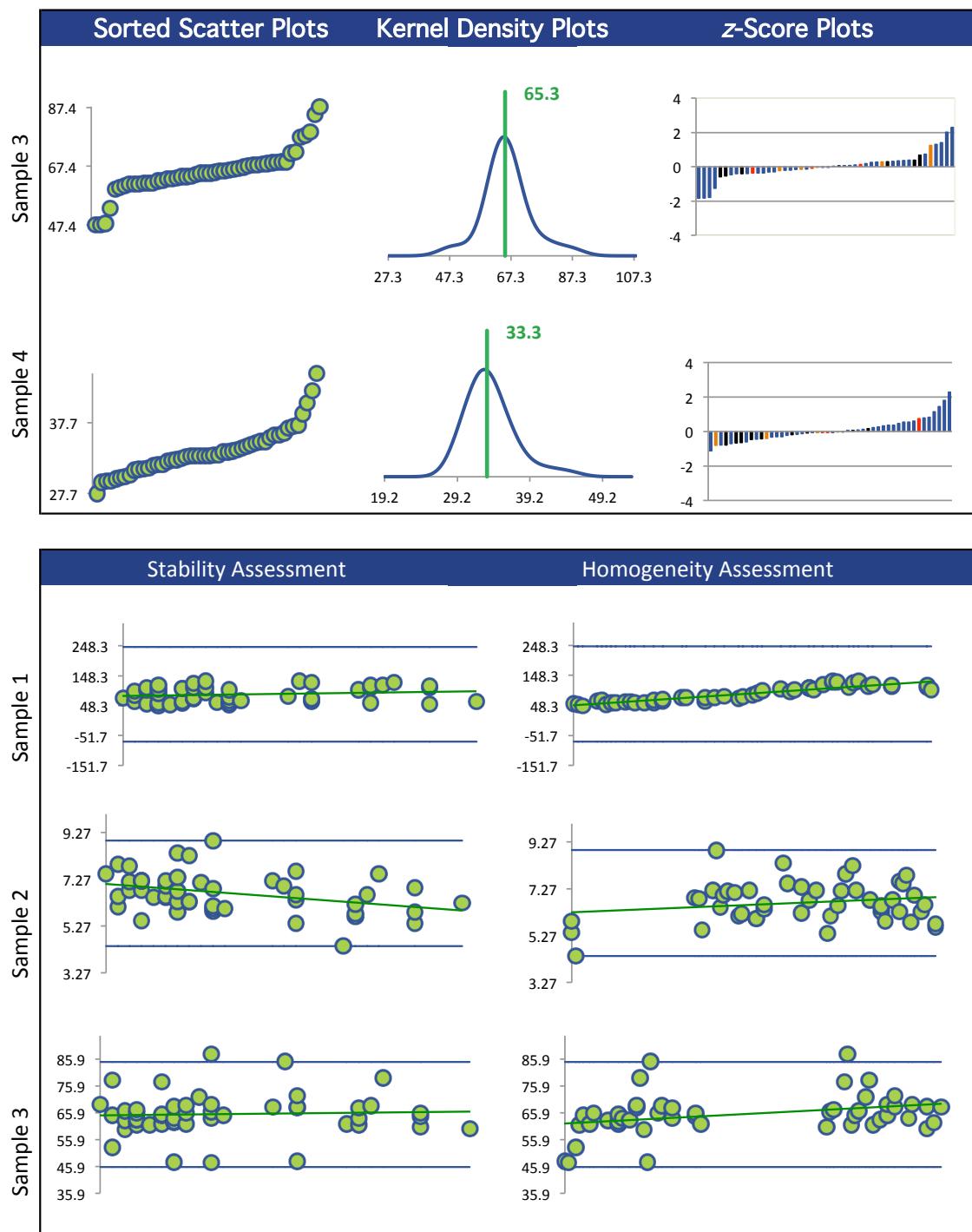
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	26	26	26	26
GC/MS - HEADSPACE (Red)	19	19	19	19
GC/FID - PURGE AND TRAP (Green)	1	1	1	1
GC/MS (Orange)	2	2	2	2

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



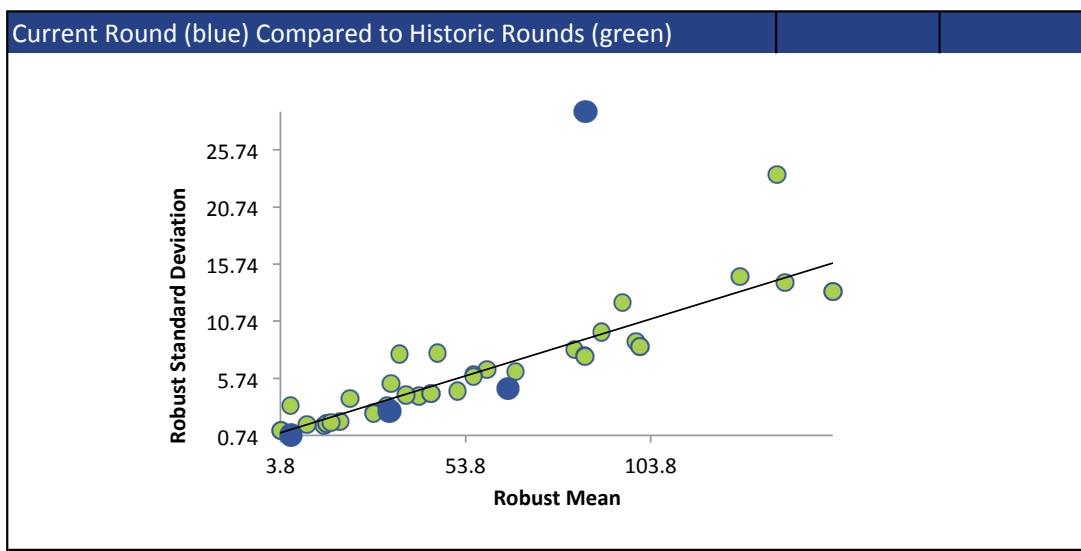
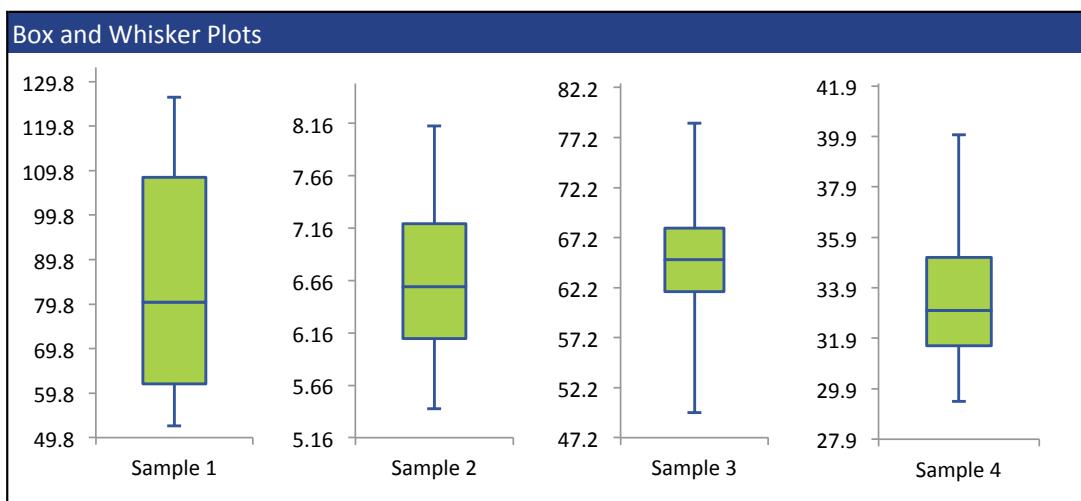
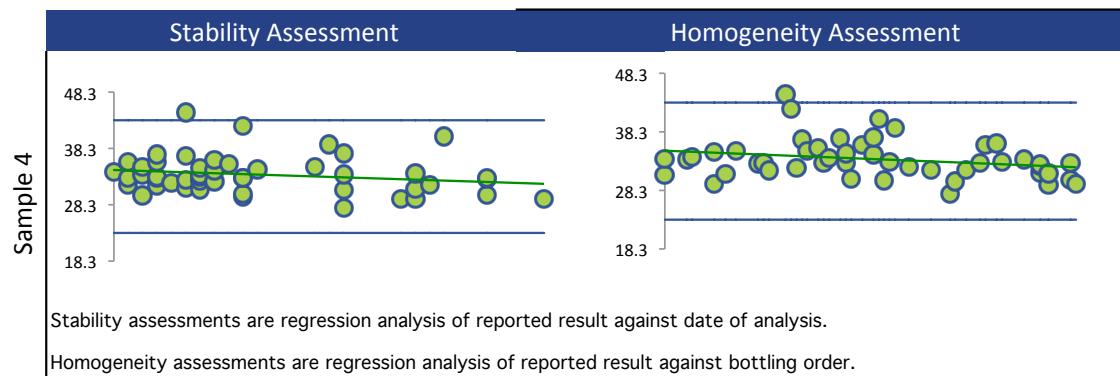
Annex A Summary by Analyte

1,2-DICHLOROPROPANE



## Annex A Summary by Analyte

### 1,2-DICHLOROPROPANE



## Annex A Summary by Analyte

### 1,3-DICHLOROBENZENE

#### Summary Statistics

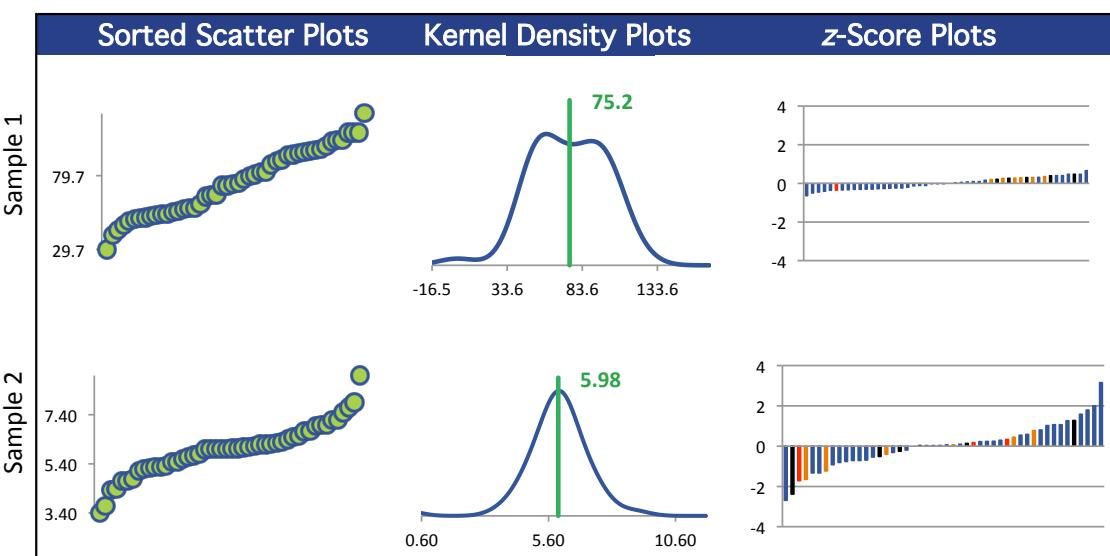
#### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	48	48	48	48
Median $\mu\text{g/L}$	74.6	6.03	58.2	29.3
Robust Mean $\mu\text{g/L}$	75.2	5.98	57.4	29.7
$U \mu\text{g/L}$	4.51	0.173	1.48	0.637
Robust Standard Deviation $\mu\text{g/L}$	25.0	0.959	8.21	3.53
Regression Standard Deviation $\mu\text{g/L}$	11.3	0.897	8.61	4.45
Stability Flag				
Homogeneity Flag	Homogeneity			
Standard Deviation Used ( $SDPA \mu\text{g/L}$ )	69.8	0.959	8.61	4.45
Outliers	0	0	0	0
$ z  > 3.0$	0	1	0	0
$2 <  z  < 3$	0	2	1	3

#### Methods Used

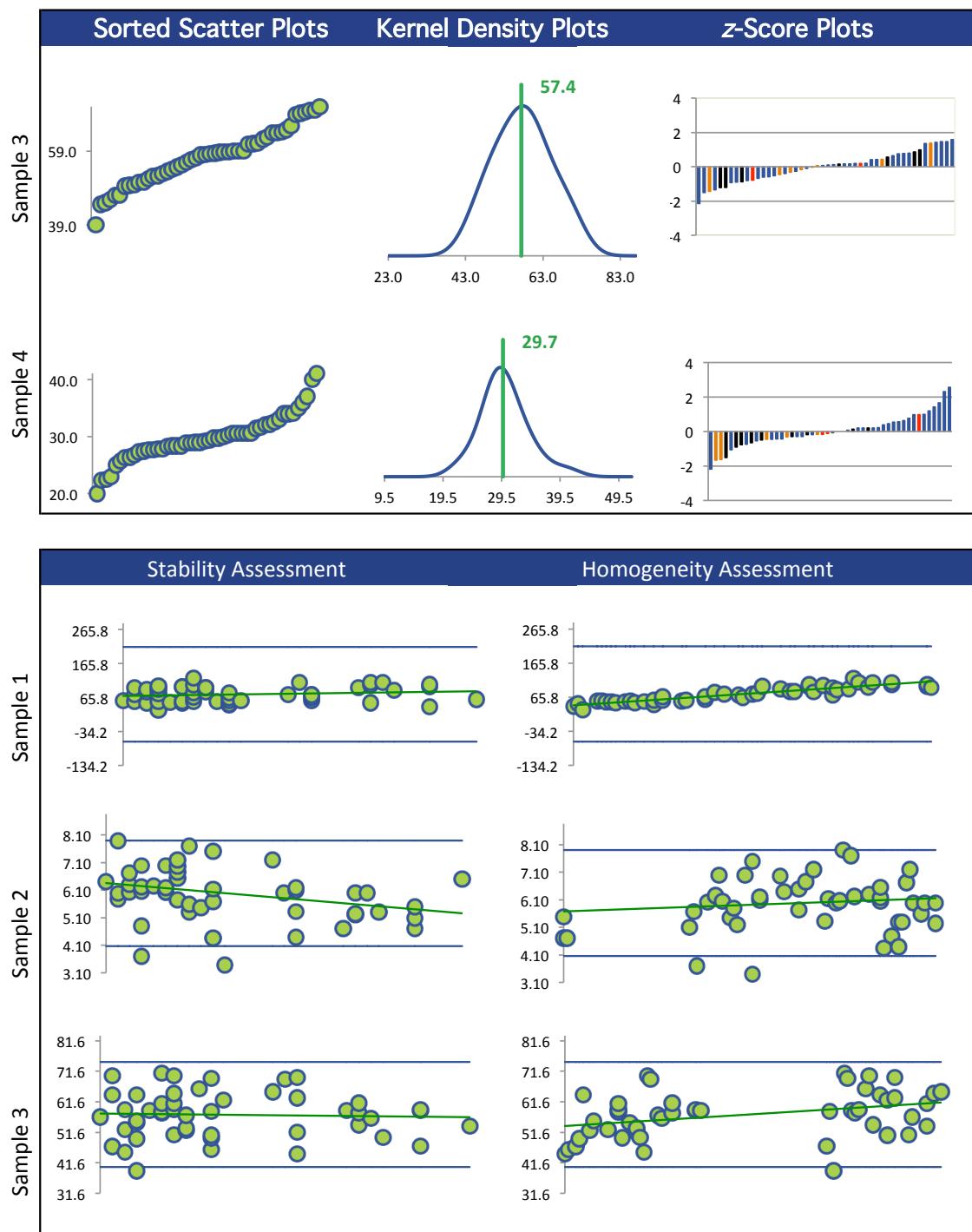
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	25	25	25	25
GC/MS - HEADSPACE (Red)	19	19	19	19
GC/FID - PURGE AND TRAP (Green)	1	1	1	1
GC/MS (Orange)	3	3	3	3

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



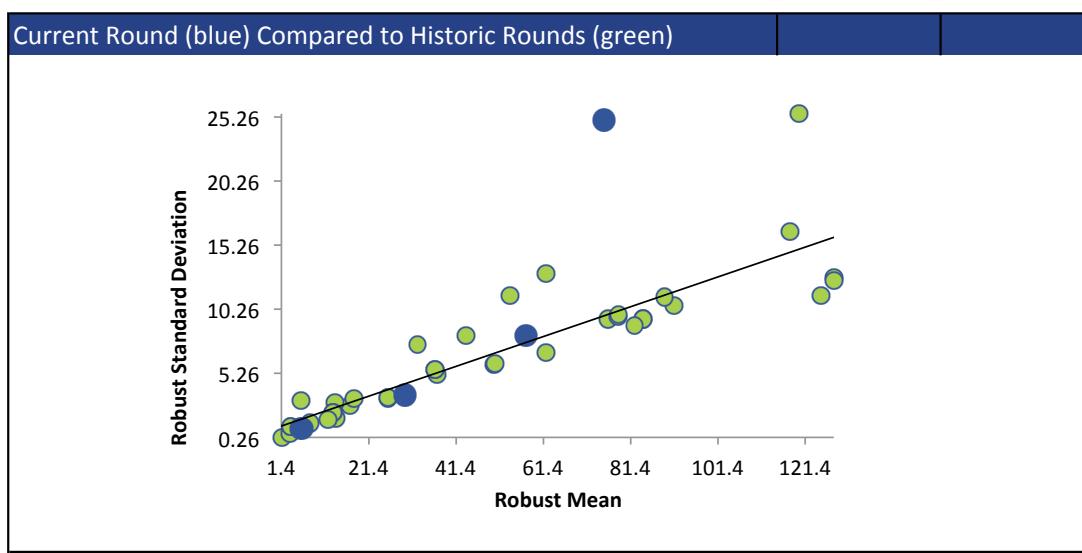
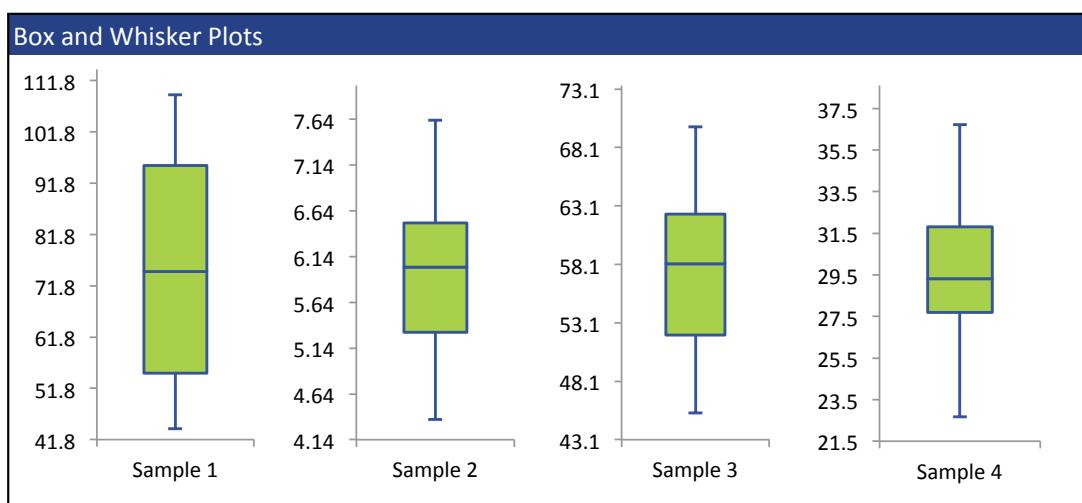
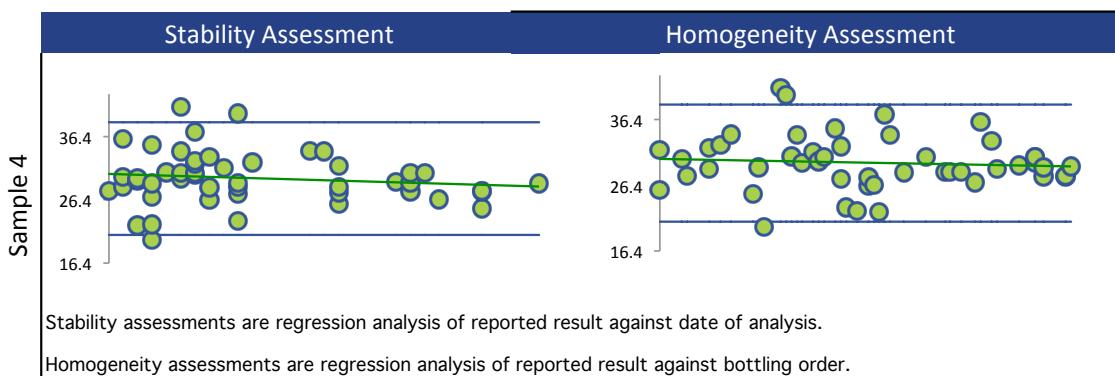
## Annex A Summary by Analyte

### 1,3-DICHLOROBENZENE



## Annex A Summary by Analyte

### 1,3-DICHLOROBENZENE



## Annex A Summary by Analyte

### 1,4-DICHLOROBENZENE

#### Summary Statistics

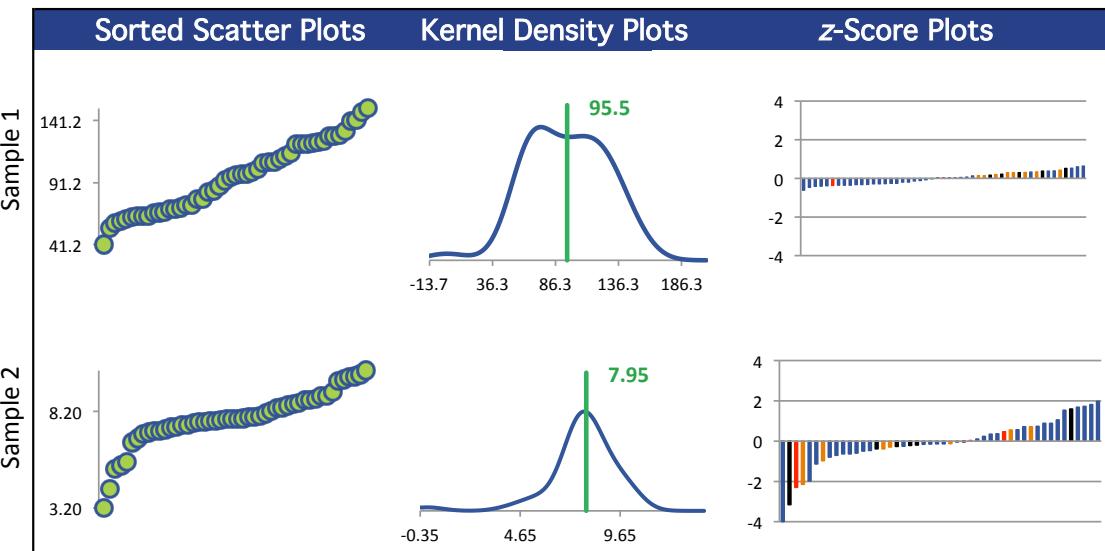
#### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	49	48	49	49
Median $\mu\text{g/L}$	97.2	7.80	75.7	38.6
Robust Mean $\mu\text{g/L}$	95.5	7.95	73.9	39.1
U $\mu\text{g/L}$	5.68	0.207	1.69	0.846
Robust Standard Deviation $\mu\text{g/L}$	31.8	1.15	9.47	4.74
Regression Standard Deviation $\mu\text{g/L}$	14.3	1.19	11.1	5.86
Stability Flag				
Homogeneity Flag	Homogeneity			
Standard Deviation Used (SDPA) $\mu\text{g/L}$	87.4	1.19	11.1	5.86
Outliers	0	0	0	0
$ z  > 3.0$	0	2	0	0
$2 <  z  < 3$	0	2	0	2

#### Methods Used

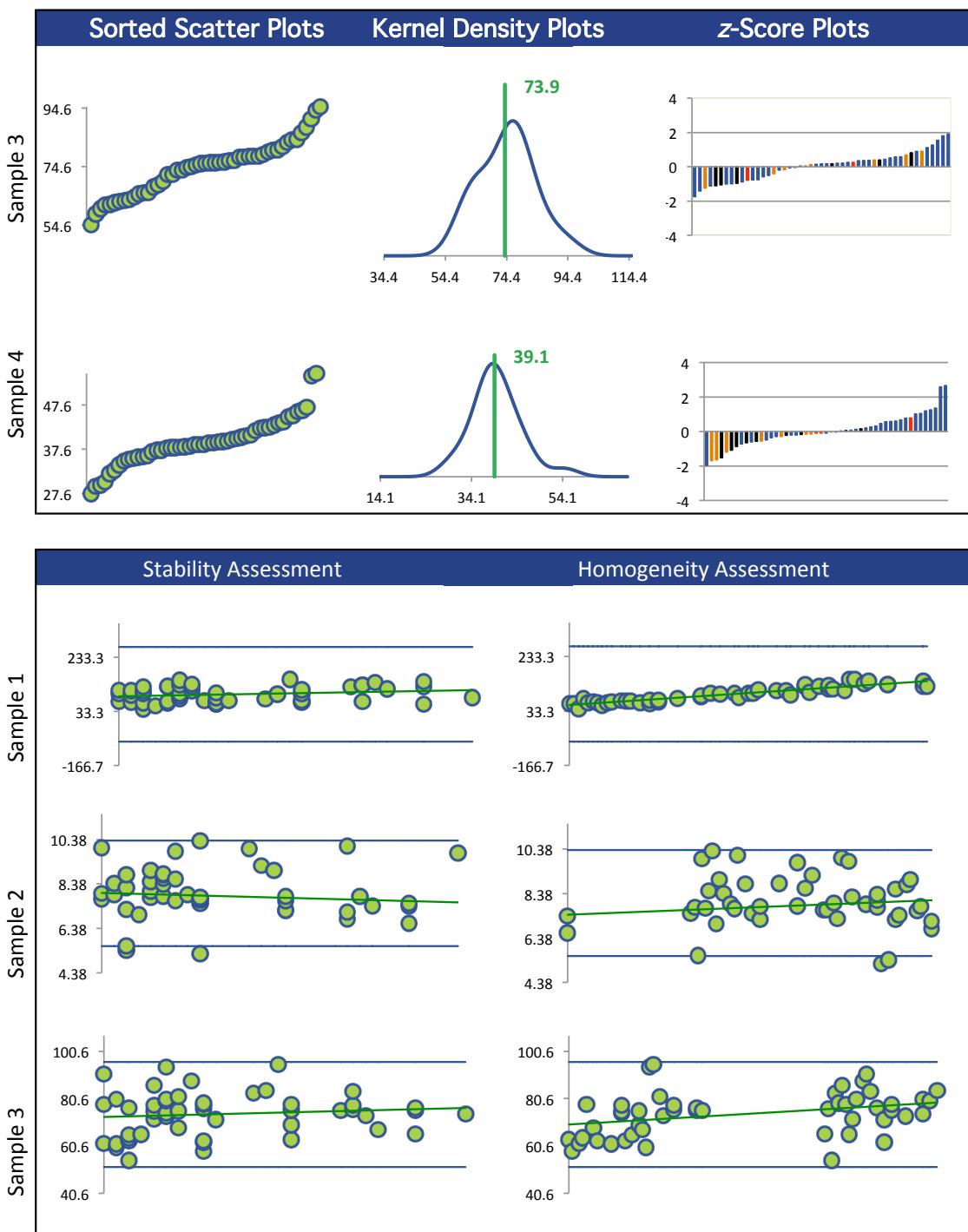
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	26	26	26	26
GC/MS - HEADSPACE (Red)	19	19	19	19
GC/MS/MS - HEADSPACE (Green)	1	1	1	1
GC/FID - PURGE AND TRAP (Orange)	1	1	1	1
GC/MS (Black)	2	1	2	2

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



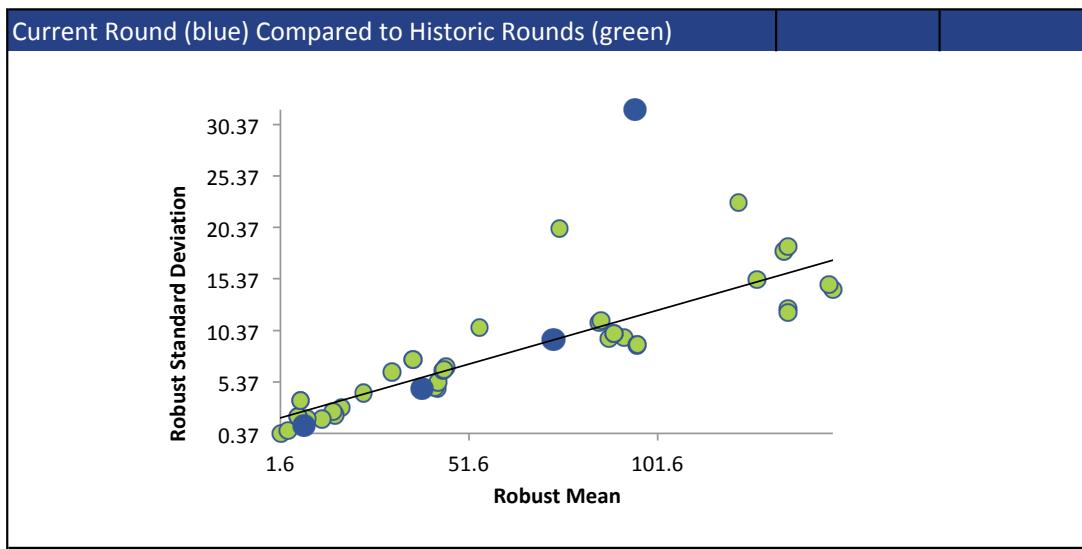
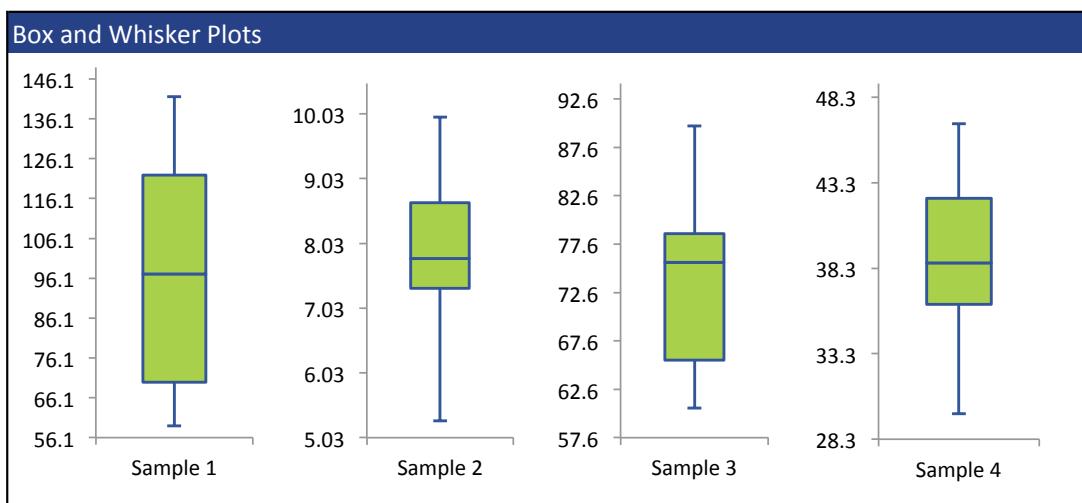
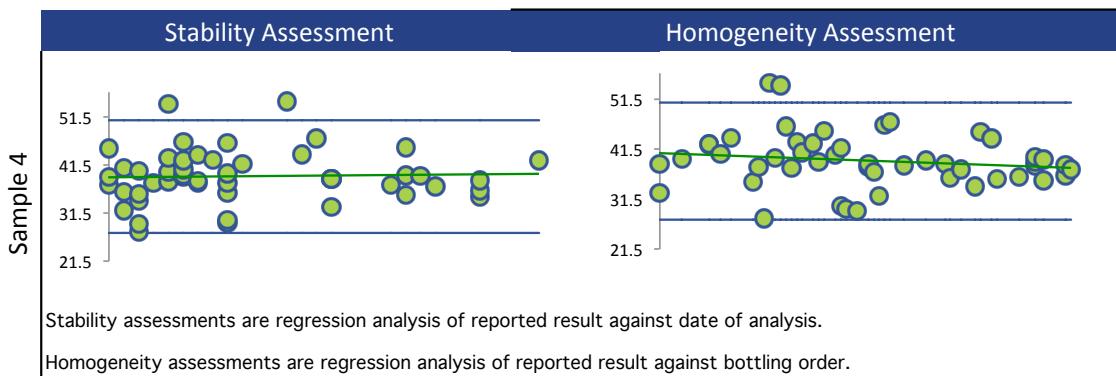
Annex A Summary by Analyte

1,4-DICHLOROBENZENE



## Annex A Summary by Analyte

### 1,4-DICHLOROBENZENE



## Annex A Summary by Analyte

### ACETONE (2-PROPANONE)

#### Summary Statistics

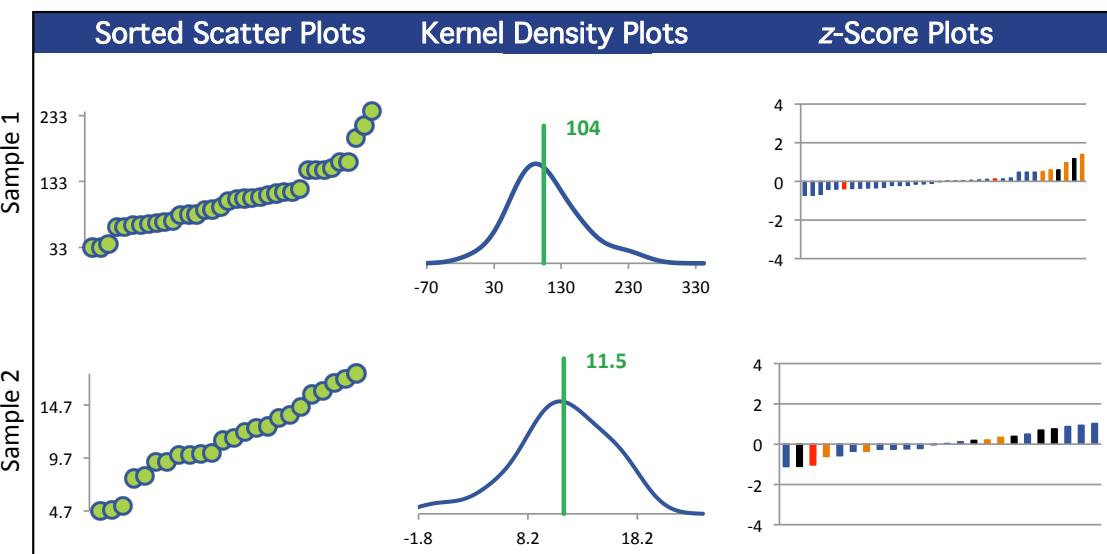
#### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	36	24	36	36
Median $\mu\text{g/L}$	105	11.5	82.5	46.1
Robust Mean $\mu\text{g/L}$	104	11.5	84.1	44.5
U $\mu\text{g/L}$	9.46	1.08	4.71	2.56
Robust Standard Deviation $\mu\text{g/L}$	45.4	4.25	22.6	12.3
Regression Standard Deviation $\mu\text{g/L}$	26.2	4.73	21.7	12.4
Stability Flag	Stability			
Homogeneity Flag	Homogeneity			
Standard Deviation Used (SDPA) $\mu\text{g/L}$	96.7	6.10	22.6	12.4
Outliers	0	0	0	0
$ z  > 3.0$	0	0	1	0
$2 <  z  < 3$	0	0	2	1

#### Methods Used

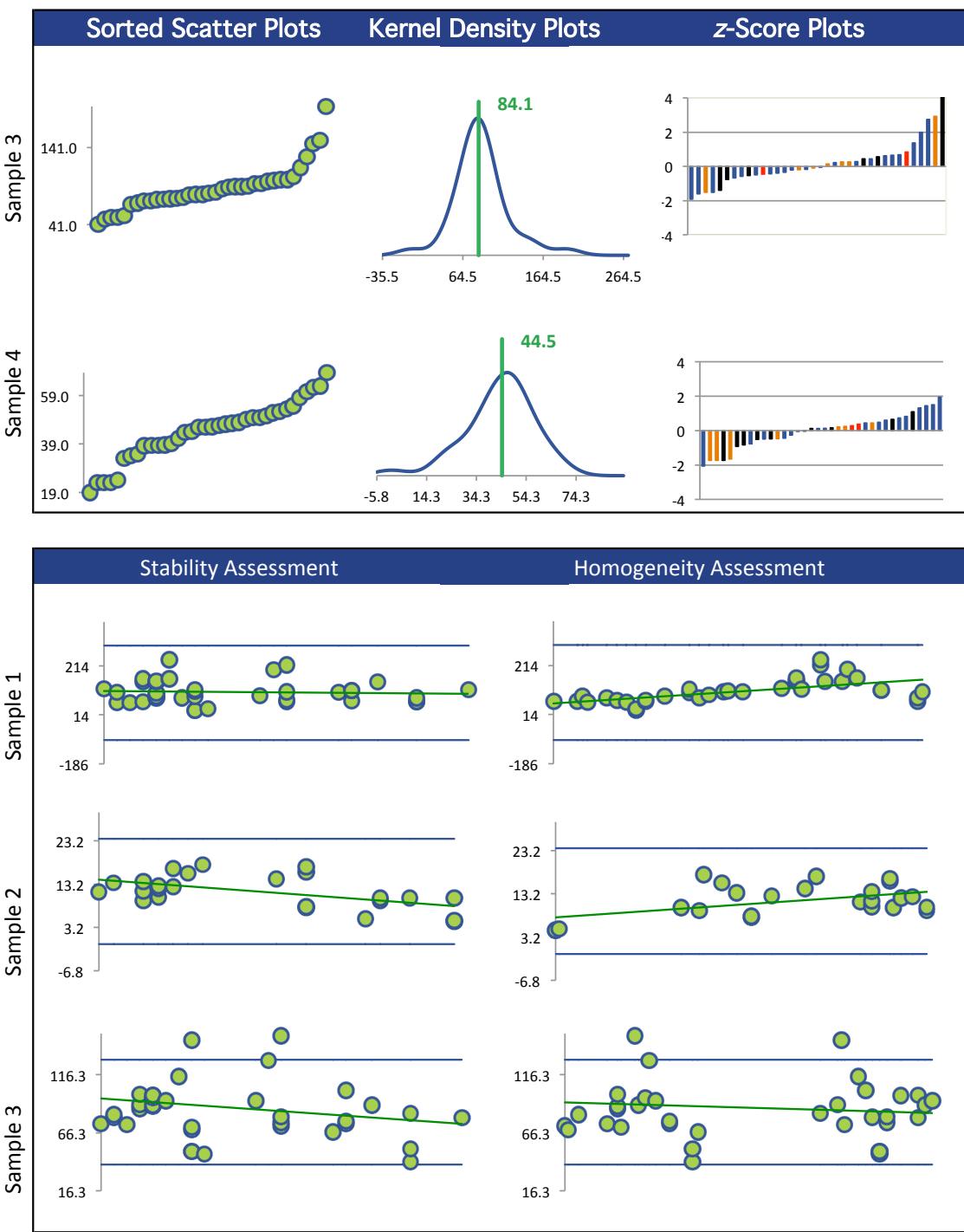
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - HEADSPACE (Blue)	14	6	14	14
GC/MS - PURGE AND TRAP (Red)	20	16	20	20
GC/FID - PURGE AND TRAP (Green)	1	1	1	1
GC/MS (Orange)	1	1	1	1

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



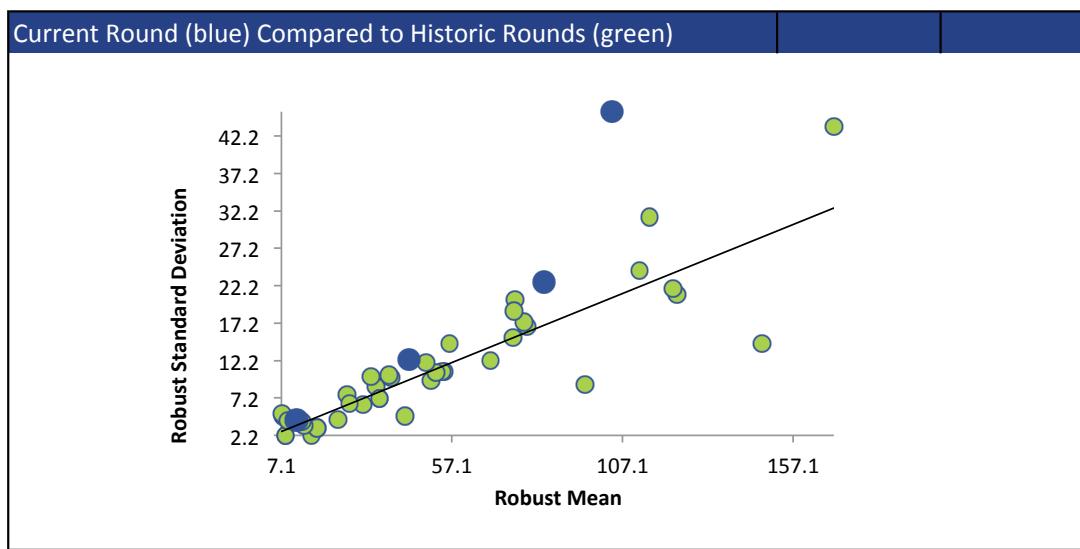
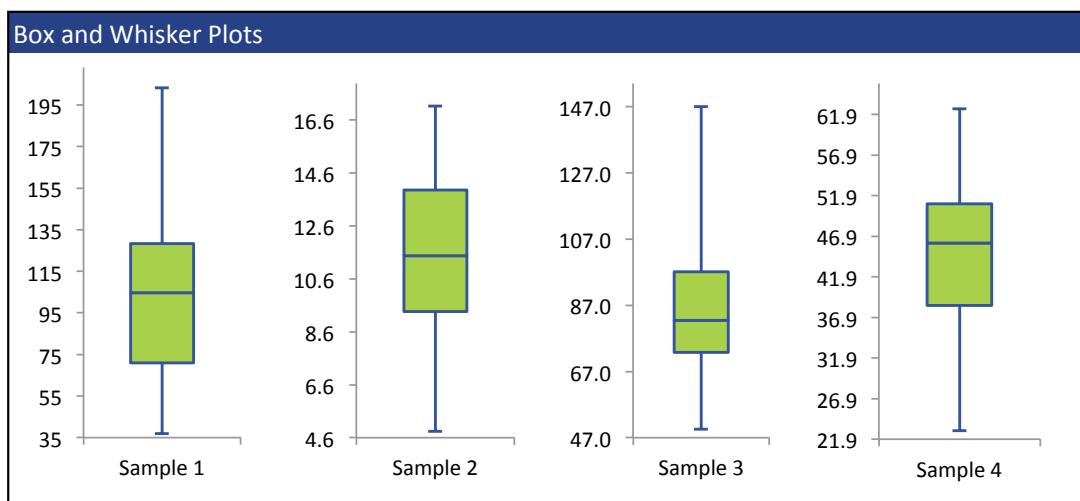
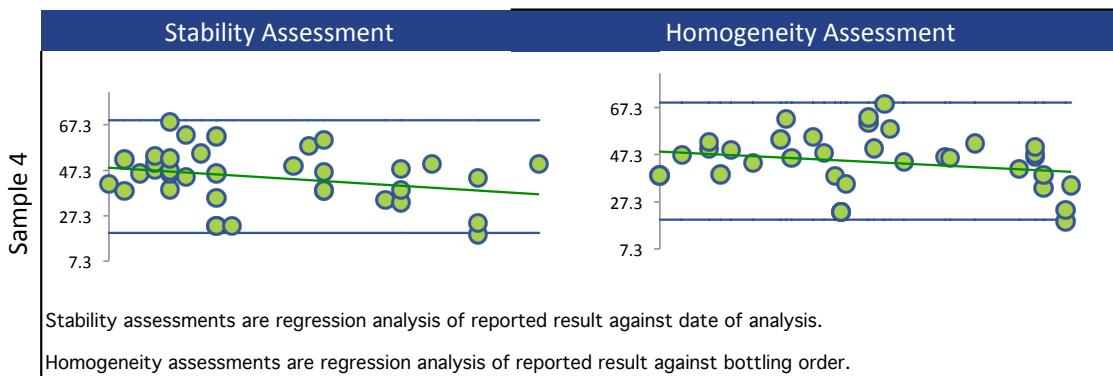
Annex A Summary by Analyte

ACETONE (2-PROPANONE)



## Annex A Summary by Analyte

### ACETONE (2-PROPANONE)



## Annex A Summary by Analyte

### BENZENE

#### Summary Statistics

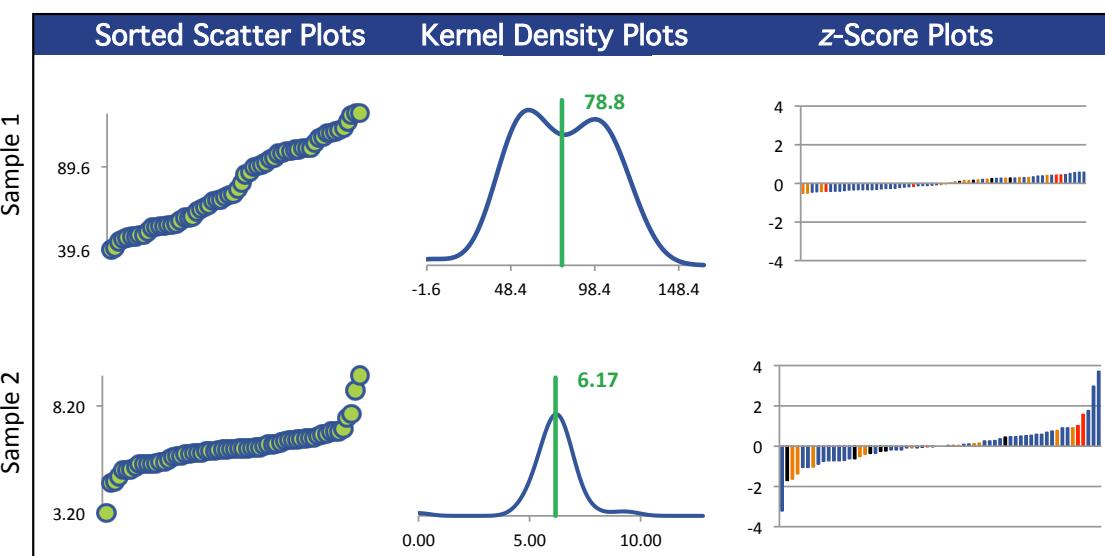
#### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	62	62	62	62
Median $\mu\text{g/L}$	74.9	6.17	60.0	30.6
Robust Mean $\mu\text{g/L}$	78.8	6.17	60.6	30.9
$U \mu\text{g/L}$	4.46	0.108	1.02	0.491
Robust Standard Deviation $\mu\text{g/L}$	28.1	0.678	6.43	3.09
Regression Standard Deviation $\mu\text{g/L}$	11.8	0.925	9.08	4.64
Stability Flag				
Homogeneity Flag	Homogeneity			
Standard Deviation Used ( $SDPA \mu\text{g/L}$ )	75.0	0.925	9.08	4.64
Outliers	0	0	0	0
$ z  > 3.0$	0	2	0	1
$2 <  z  < 3$	0	1	2	0

#### Methods Used

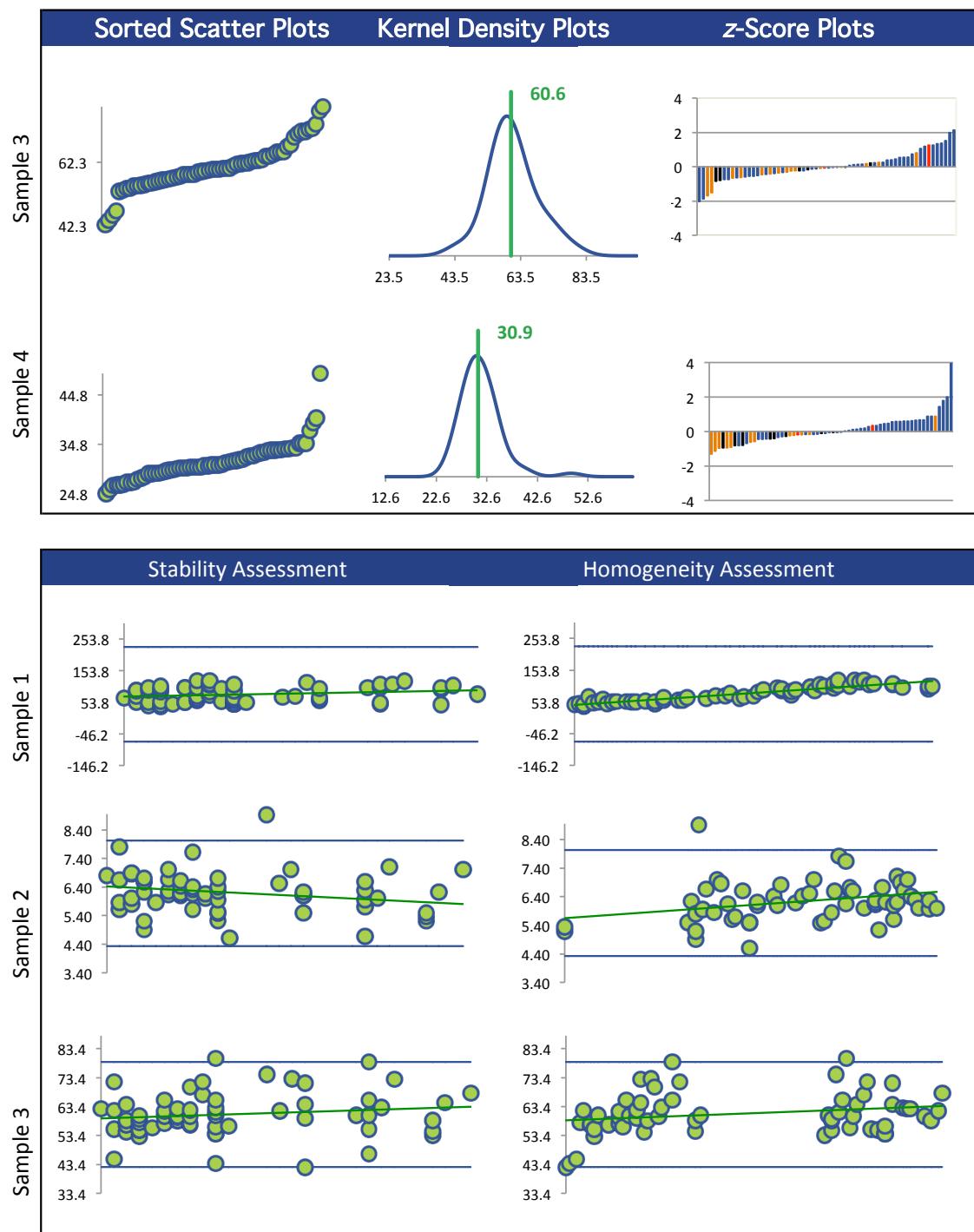
Method	C16-1	C16-2	C16-3	C16-4
GC/MS (Blue)	6	6	6	6
GC/MS - PURGE AND TRAP (Red)	31	31	31	31
GC/MS - HEADSPACE (Green)	22	22	22	22
GC/FID - HEADSPACE (Orange)	1	1	1	1
GC/MS/MS - HEADSPACE (Black)	1	1	1	1
GC/FID - PURGE AND TRAP (Yellow)	1	1	1	1

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



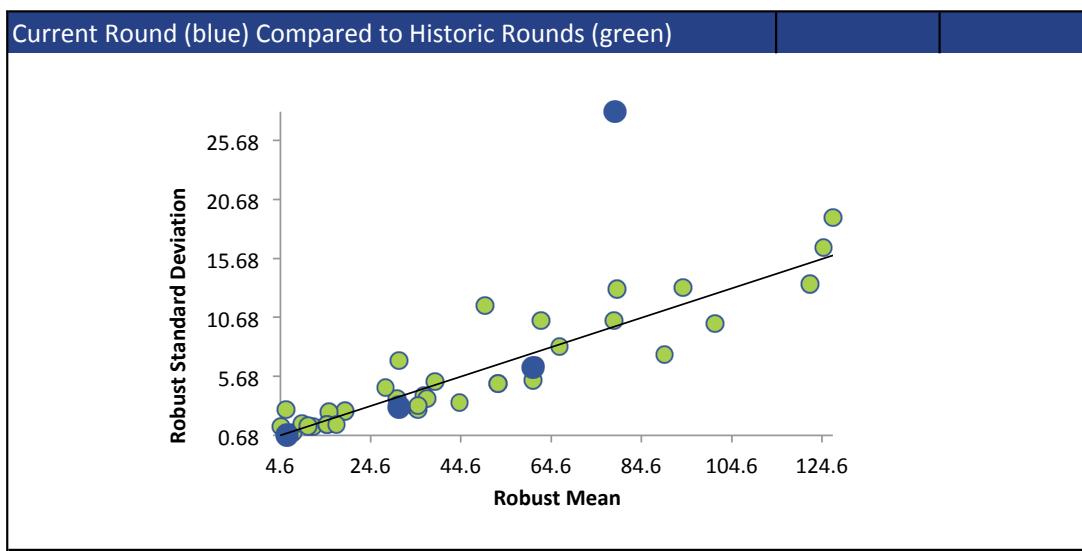
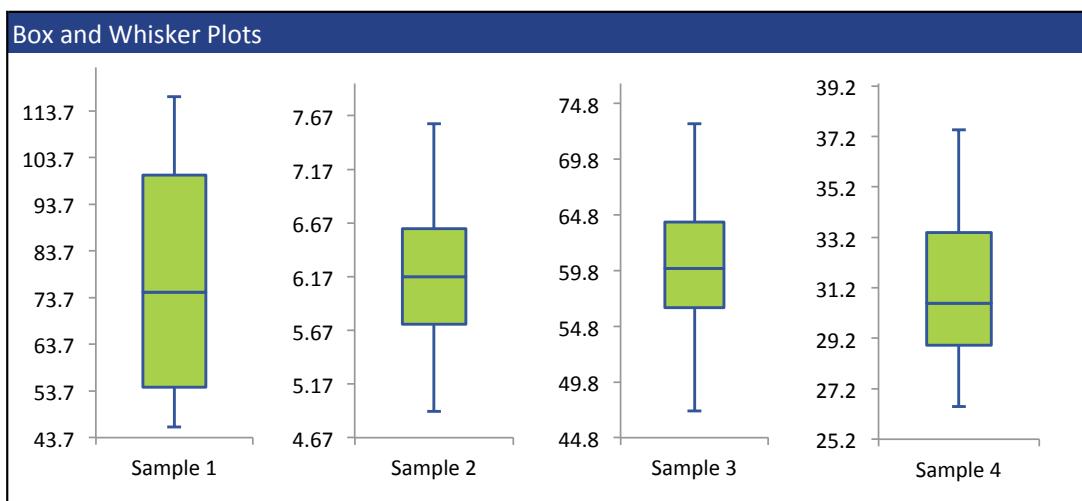
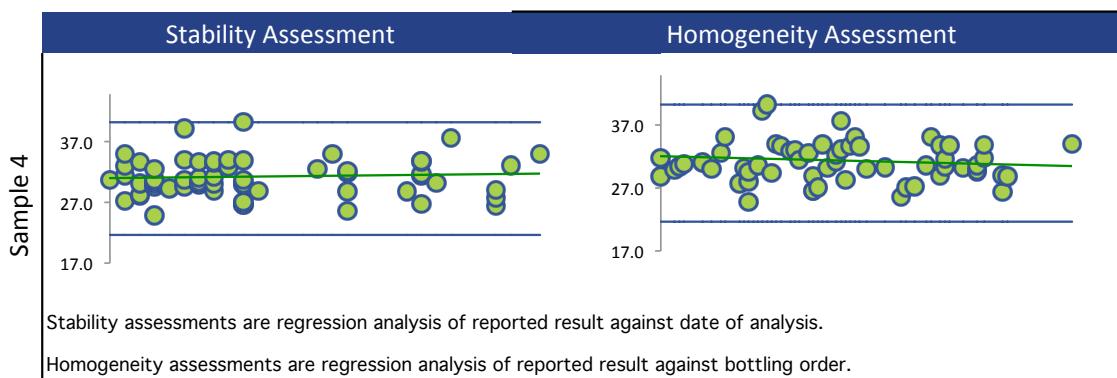
Annex A Summary by Analyte

BENZENE



## Annex A Summary by Analyte

### BENZENE



Annex A Summary by Analyte

## BROMODICHLOROMETHANE

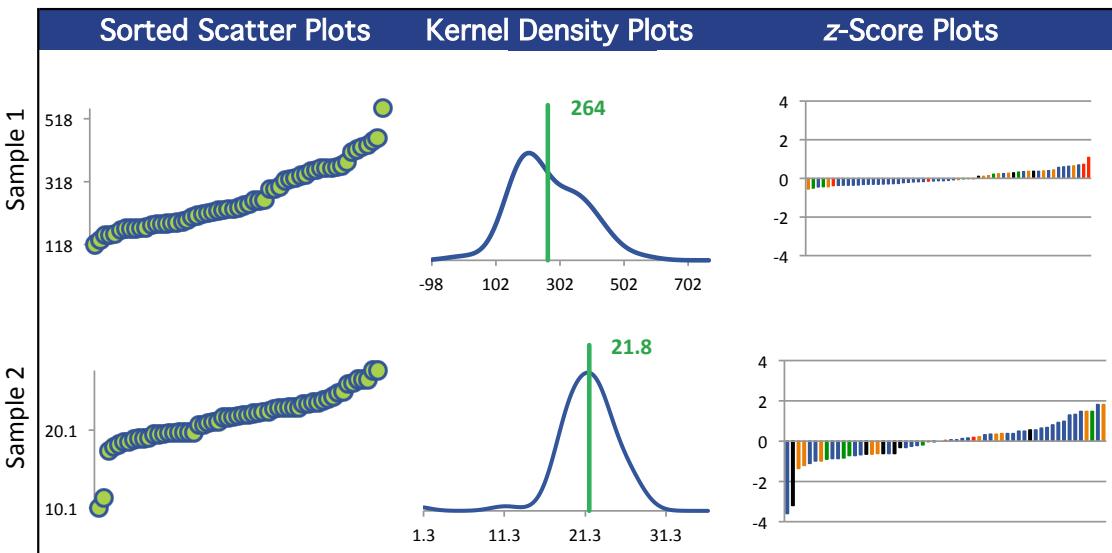
### Summary Statistics

Statistic	C16-1	C16-2	C16-3	C16-4
N	57	57	57	57
Median $\mu\text{g/L}$	235	21.9	211	106
Robust Mean $\mu\text{g/L}$	264	21.8	212	108
$U \mu\text{g/L}$	17.4	0.480	4.29	2.47
Robust Standard Deviation $\mu\text{g/L}$	105	2.90	25.9	14.9
Regression Standard Deviation $\mu\text{g/L}$	39.6	3.26	31.8	16.2
Stability Flag				
Homogeneity Flag	Homogeneity			
Standard Deviation Used ( $SDPA \mu\text{g/L}$ )	262	3.26	31.8	16.2
Outliers	0	0	0	0
$ z  > 3.0$	0	2	2	3
$2 <  z  < 3$	0	0	2	4

### Methods Used

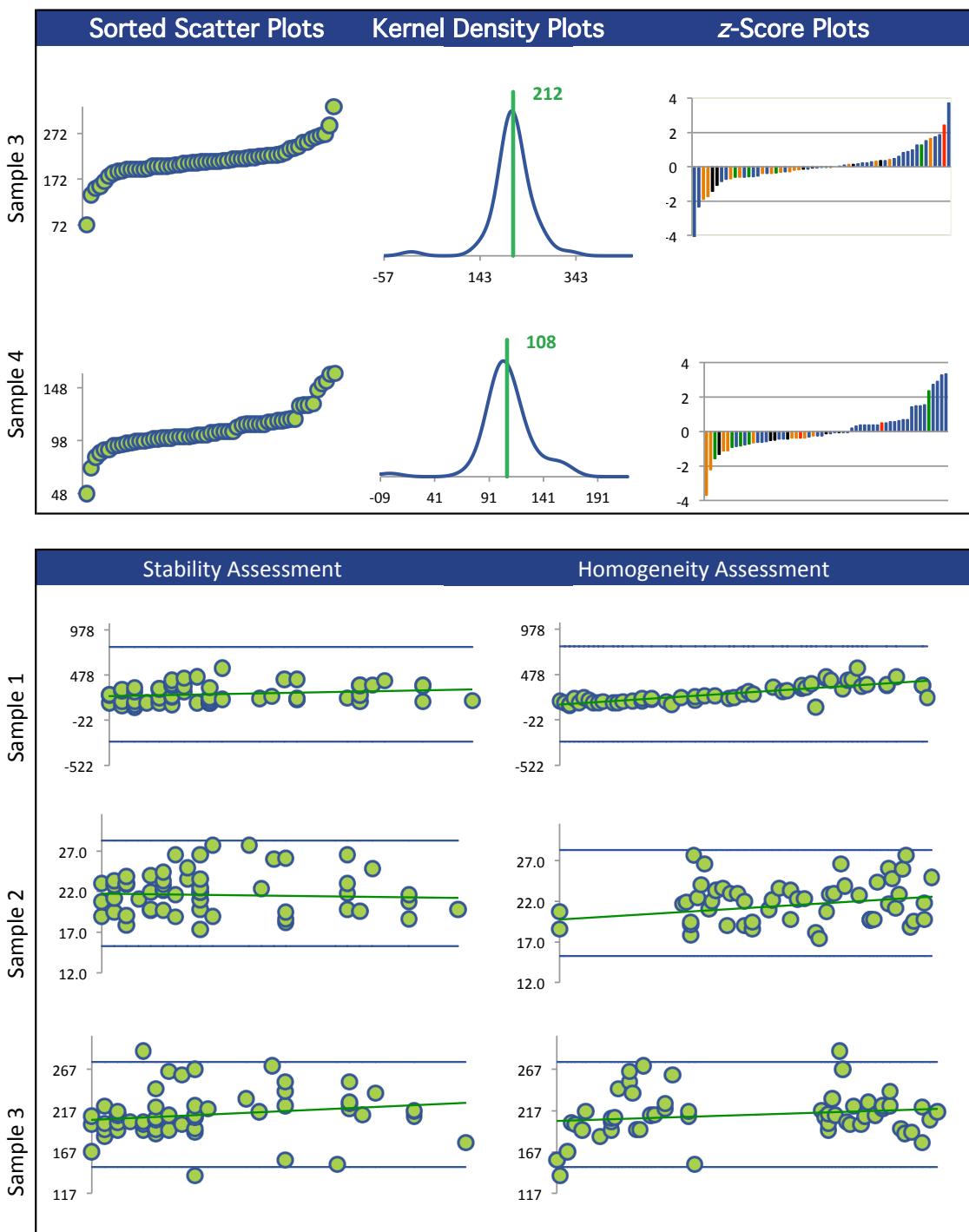
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	28	28	28	28
GC/MS (Red)	5	5	5	5
GC/MS - HEADSPACE (Green)	22	22	22	22
GC/MS/MS - HEADSPACE (Orange)	1	1	1	1
GC/FID - PURGE AND TRAP (Black)	1	1	1	1

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



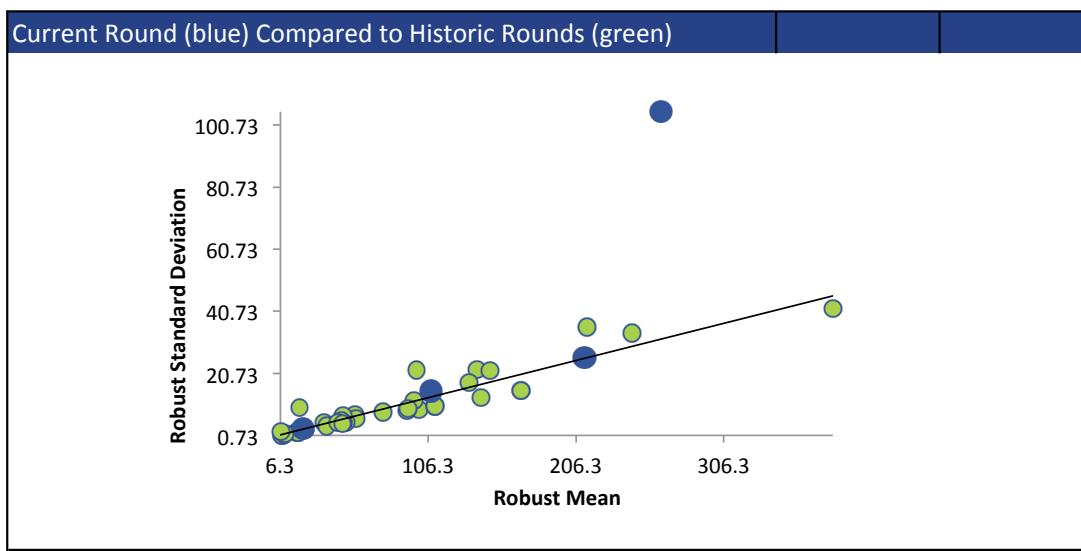
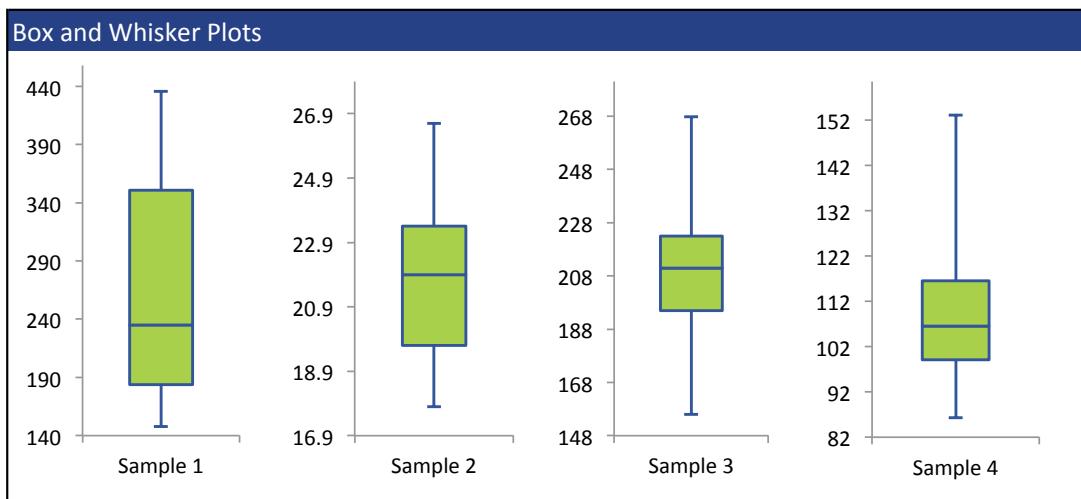
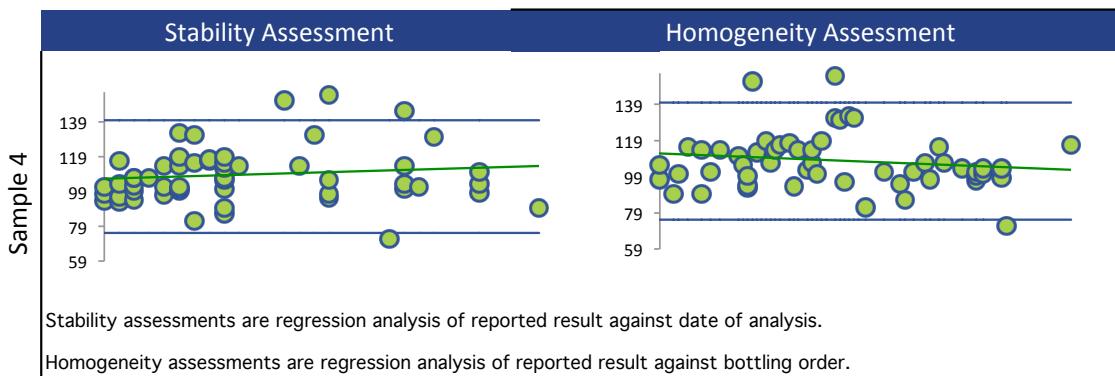
Annex A Summary by Analyte

BROMODICHLOROMETHANE



## Annex A Summary by Analyte

### BROMODICHLOROMETHANE



## Annex A Summary by Analyte

### BROMOFORM

#### Summary Statistics

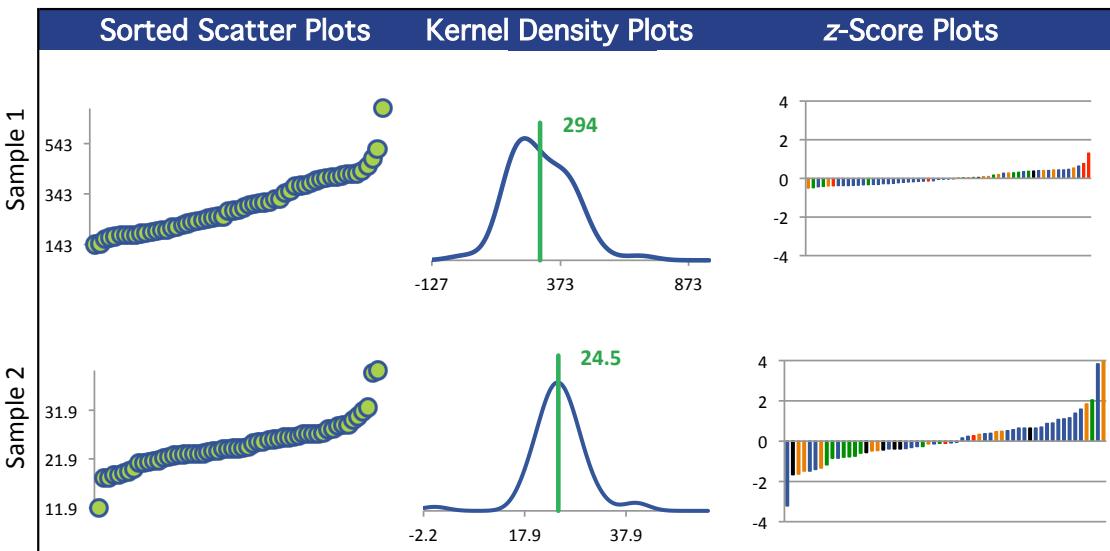
#### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	57	57	57	57
Median $\mu\text{g/L}$	281	24.1	235	120
Robust Mean $\mu\text{g/L}$	294	24.5	239	122
$U \mu\text{g/L}$	18.5	0.649	5.50	3.28
Robust Standard Deviation $\mu\text{g/L}$	112	3.92	33.2	19.8
Regression Standard Deviation $\mu\text{g/L}$	44.1	3.68	35.8	18.2
Stability Flag				
Homogeneity Flag	Homogeneity			
Standard Deviation Used (SDPA) $\mu\text{g/L}$	298	3.92	35.8	19.8
Outliers	0	0	0	0
$ z  > 3.0$	0	3	2	1
$2 <  z  < 3$	0	1	2	4

#### Methods Used

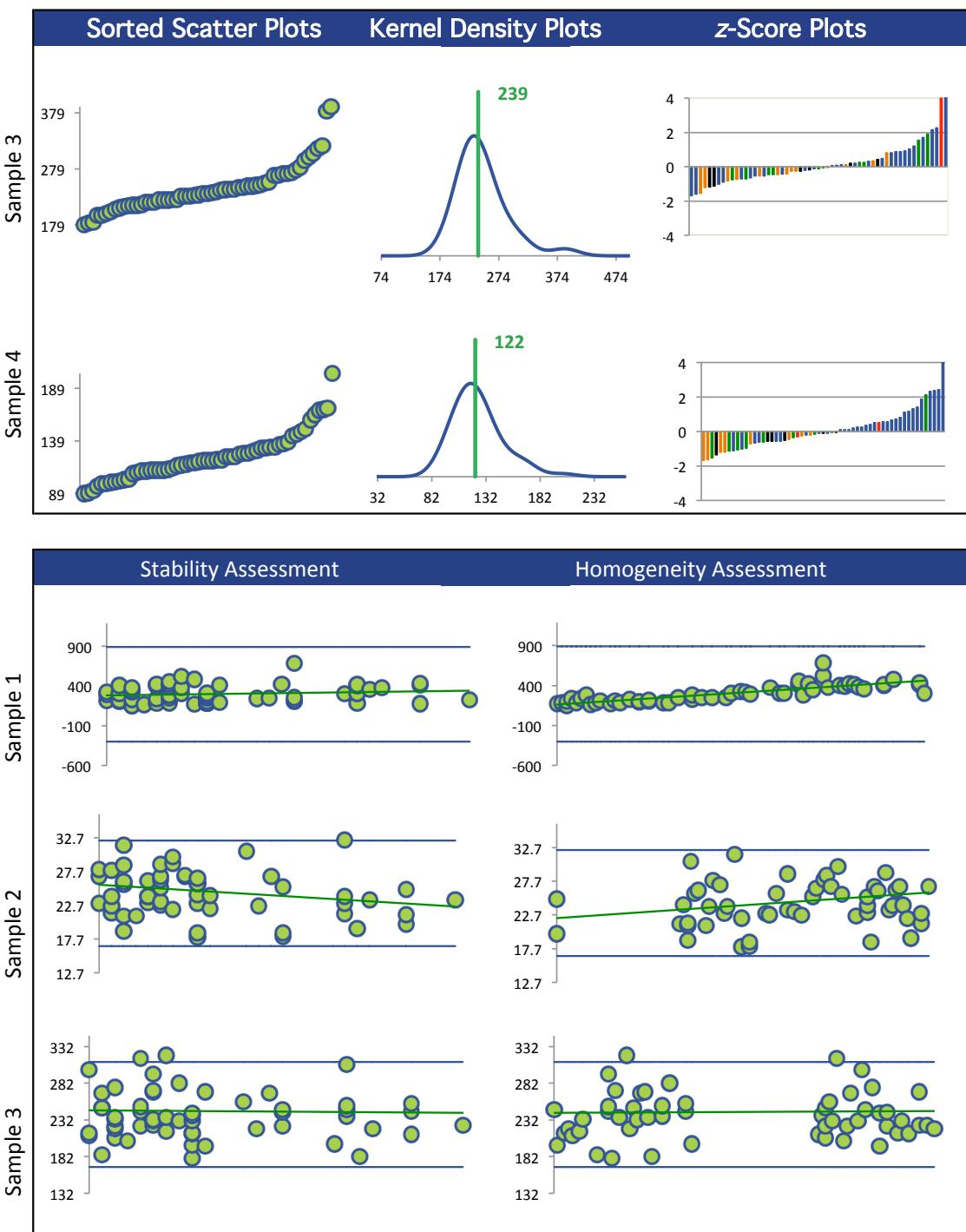
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	28	28	28	28
GC/MS (Red)	5	5	5	5
GC/MS - HEADSPACE (Green)	22	22	22	22
GC/MS/MS - HEADSPACE (Orange)	1	1	1	1
GC/FID - PURGE AND TRAP (Black)	1	1	1	1

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



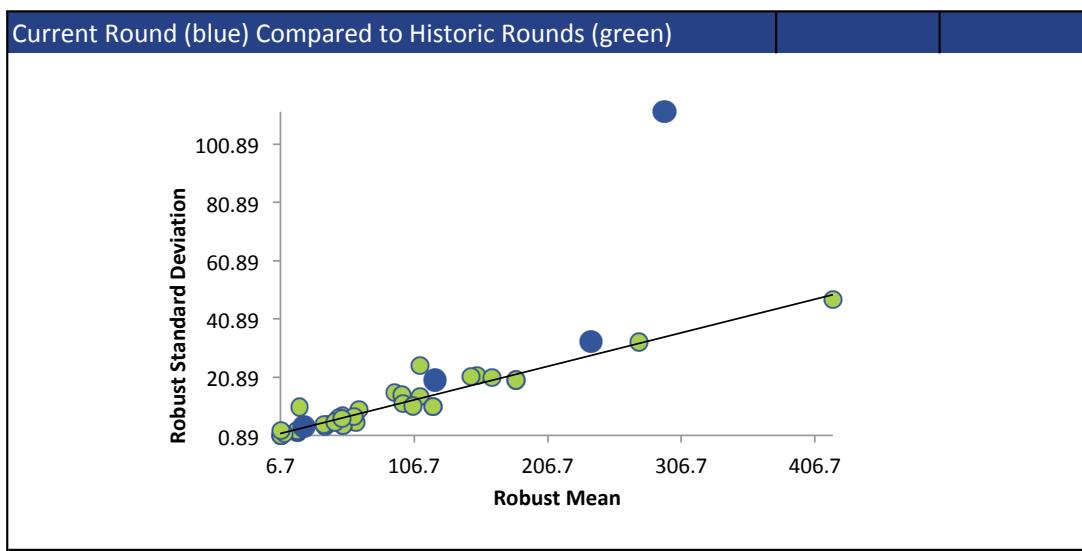
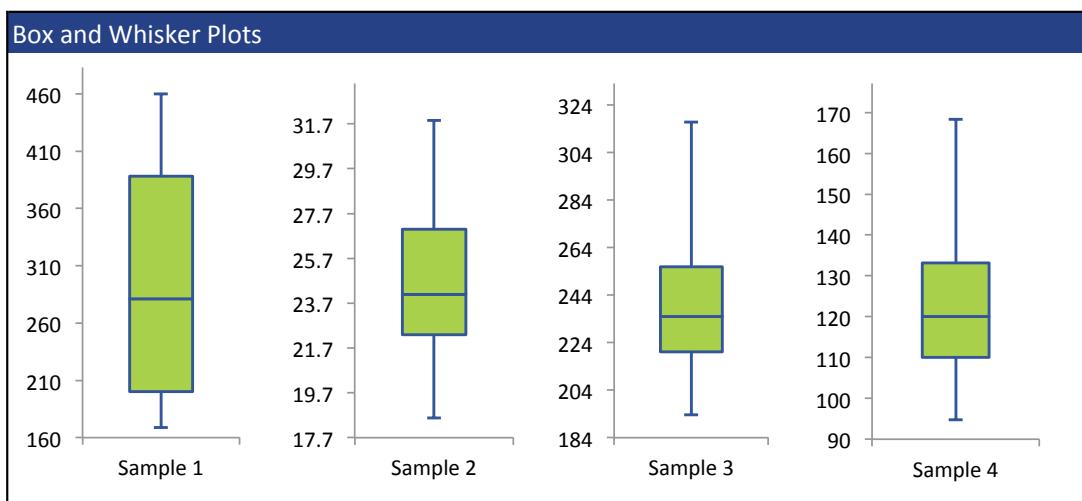
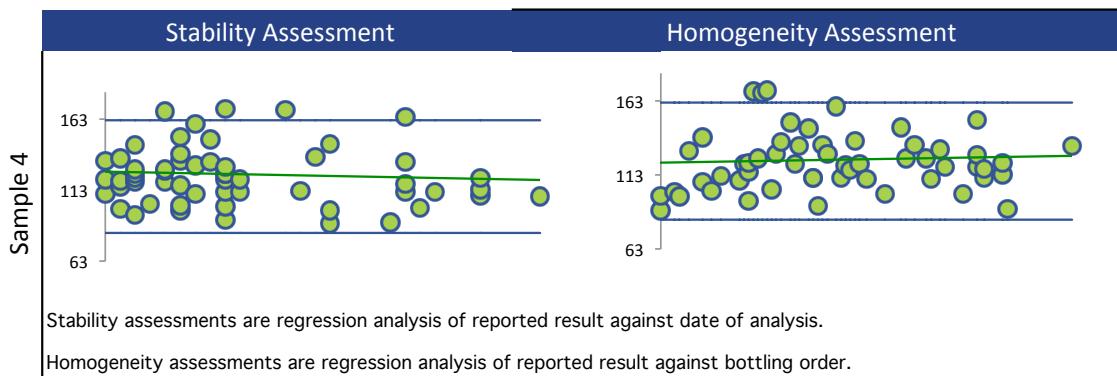
Annex A Summary by Analyte

BROMOFORM



## Annex A Summary by Analyte

### BROMOFORM



Annex A Summary by Analyte

## CARBON TETRACHLORIDE

### Summary Statistics

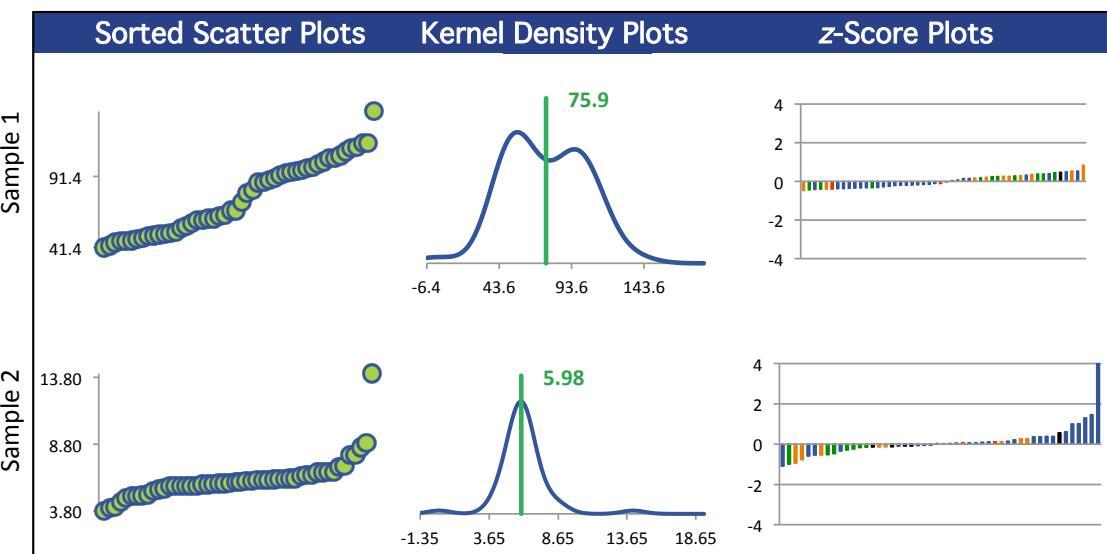
### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	50	50	50	50
Median $\mu\text{g/L}$	70.3	6.00	58.5	29.4
Robust Mean $\mu\text{g/L}$	75.9	5.98	59.2	29.8
$U \mu\text{g/L}$	4.90	0.153	1.32	0.728
Robust Standard Deviation $\mu\text{g/L}$	27.7	0.864	7.48	4.12
Regression Standard Deviation $\mu\text{g/L}$	11.4	0.897	8.88	4.47
Stability Flag		Stability		
Homogeneity Flag	Homogeneity			
Standard Deviation Used ( $SDPA \mu\text{g/L}$ )	72.6	2.00	8.88	4.47
Outliers	0	0	0	0
$ z  > 3.0$	0	1	0	2
$2 <  z  < 3$	0	0	6	1

### Methods Used

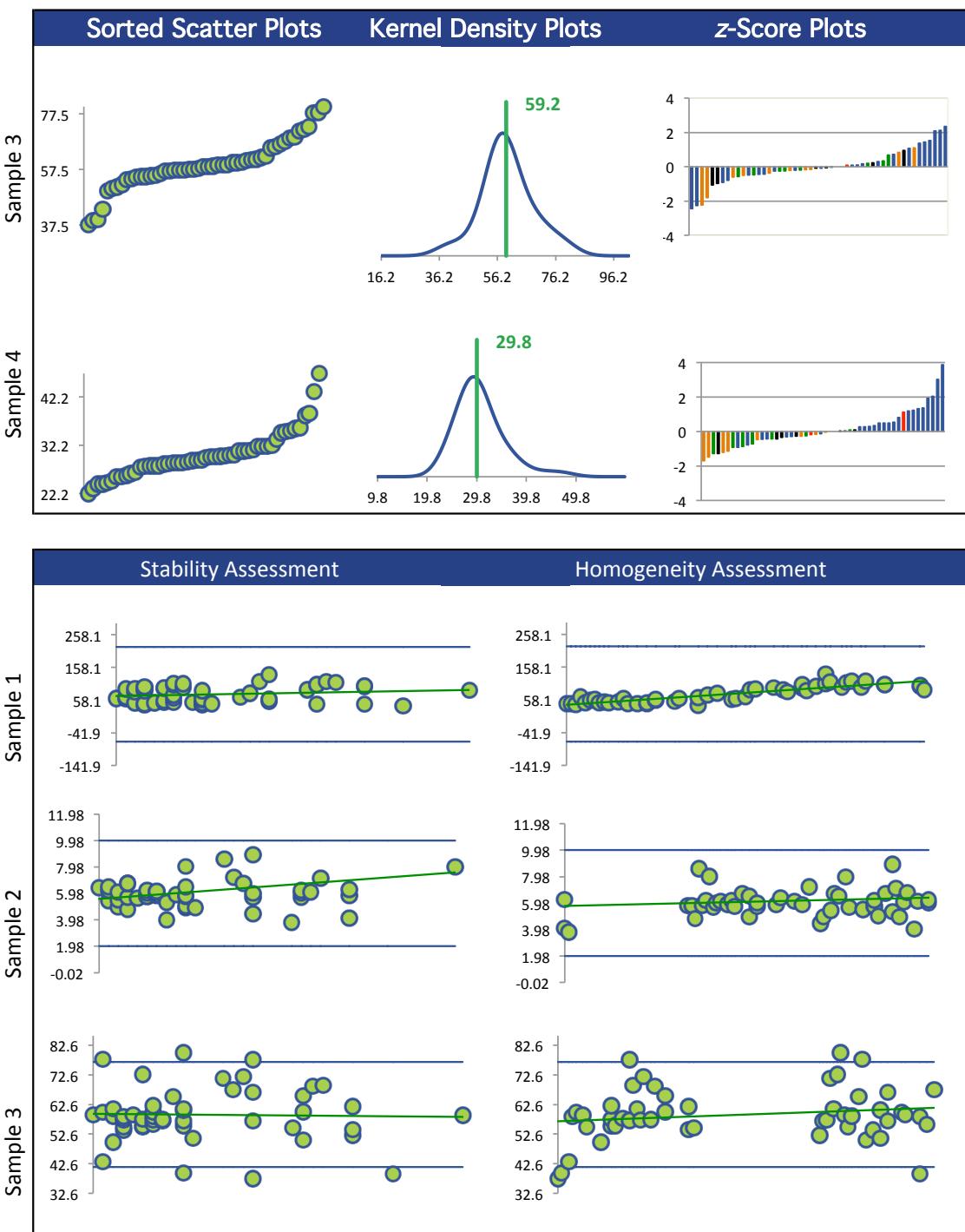
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	27	27	27	27
GC/MS - HEADSPACE (Red)	19	19	19	19
GC/MS/MS - HEADSPACE (Green)	1	1	1	1
GC/FID - PURGE AND TRAP (Orange)	1	1	1	1
GC/MS (Black)	2	2	2	2

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



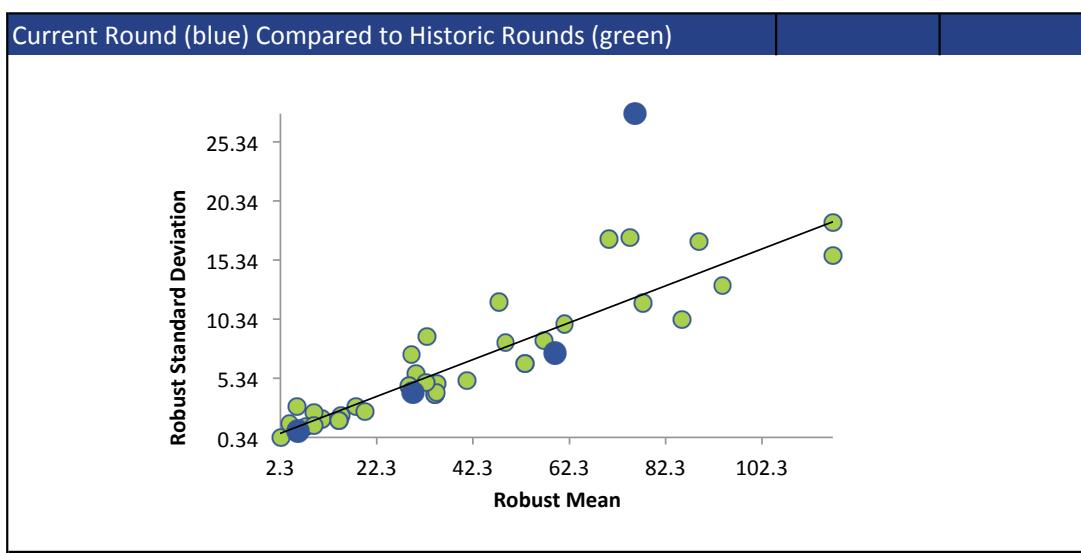
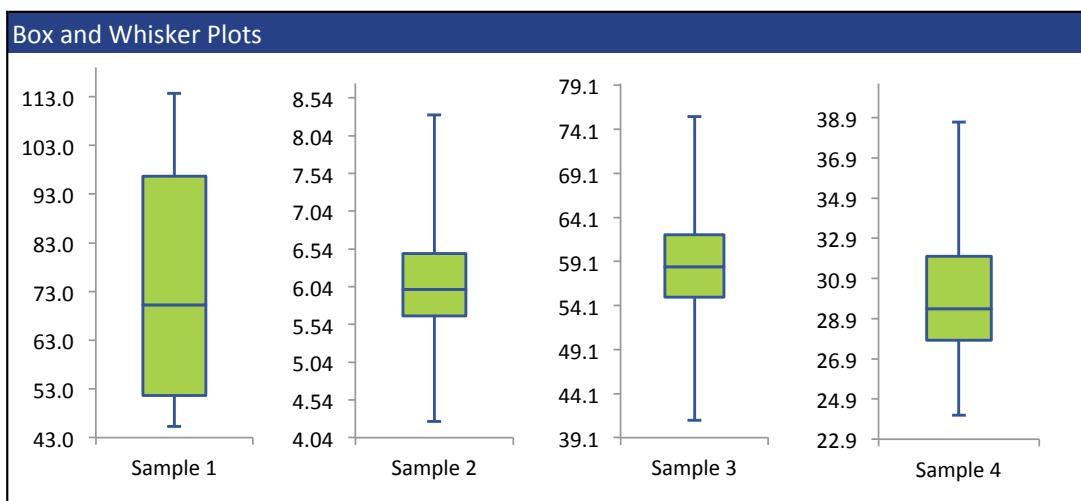
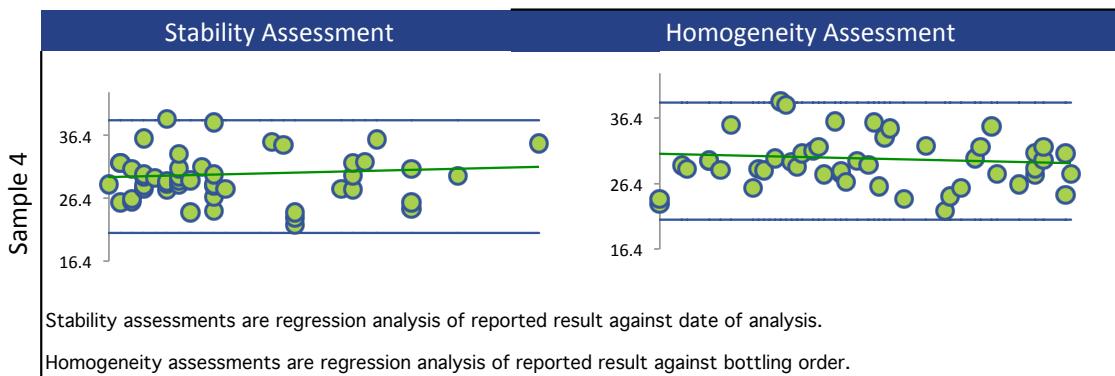
Annex A Summary by Analyte

CARBON TETRACHLORIDE



## Annex A Summary by Analyte

### CARBON TETRACHLORIDE



Annex A Summary by Analyte

## CHLOROBENZENE

### Summary Statistics

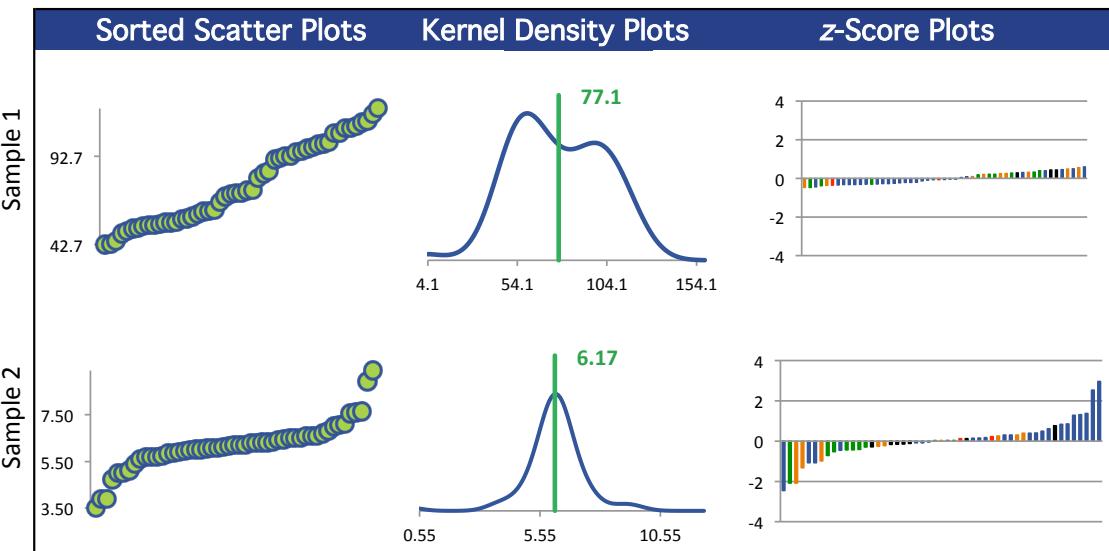
### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	51	51	51	51
Median $\mu\text{g/L}$	71.9	6.20	58.5	30.2
Robust Mean $\mu\text{g/L}$	77.1	6.17	59.6	30.4
$U \mu\text{g/L}$	4.59	0.128	1.06	0.497
Robust Standard Deviation $\mu\text{g/L}$	26.2	0.733	6.03	2.84
Regression Standard Deviation $\mu\text{g/L}$	11.6	0.926	8.93	4.56
Stability Flag	Stability			
Homogeneity Flag	Homogeneity			
Standard Deviation Used ( $SDPA \mu\text{g/L}$ )	71.7	1.09	8.93	4.56
Outliers	0	0	0	0
$ z  > 3.0$	0	0	0	1
$2 <  z  < 3$	0	5	1	2

### Methods Used

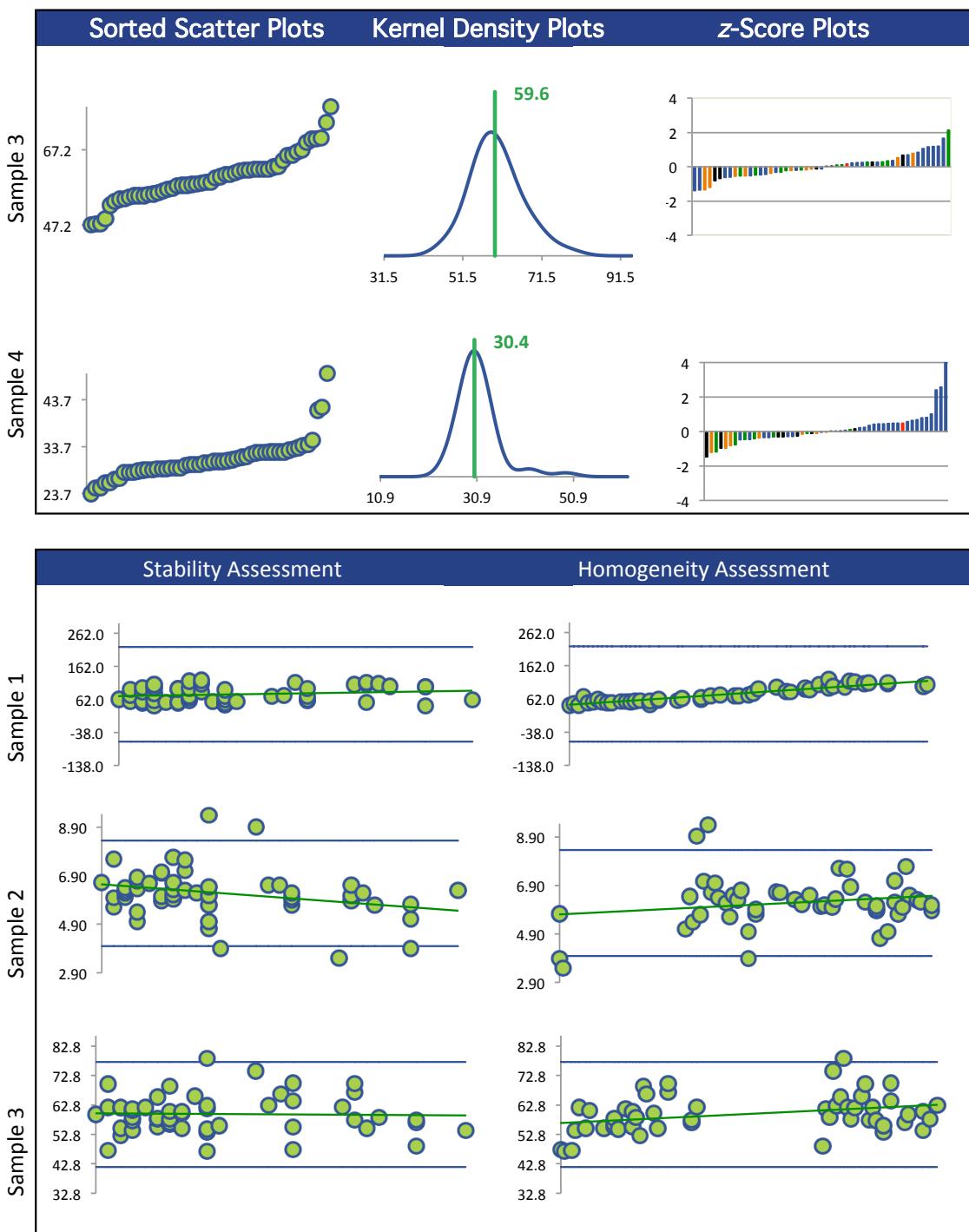
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	28	28	28	28
GC/MS - HEADSPACE (Red)	19	19	19	19
GC/MS/MS - HEADSPACE (Green)	1	1	1	1
GC/FID - PURGE AND TRAP (Orange)	1	1	1	1
GC/MS (Black)	2	2	2	2

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



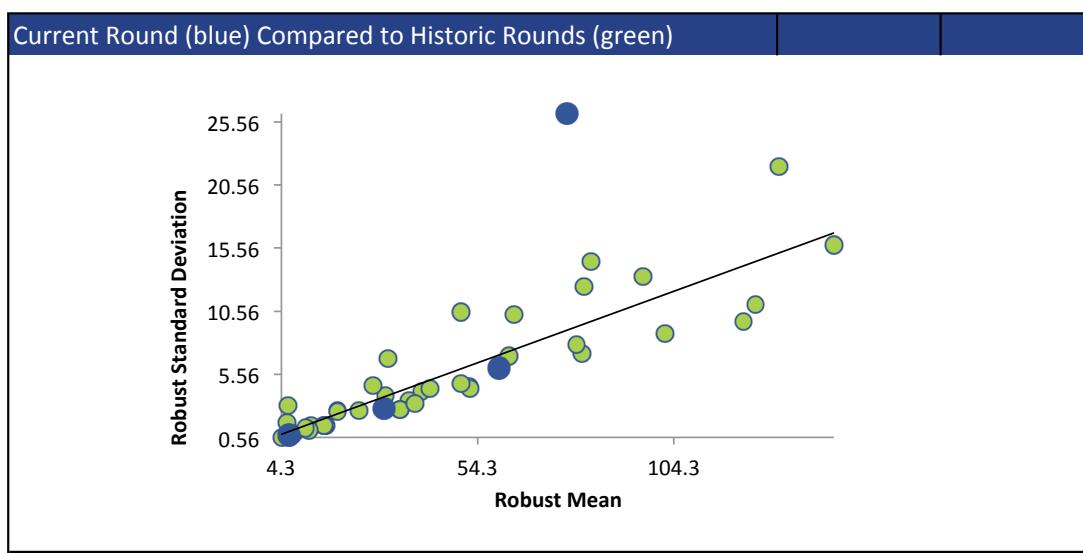
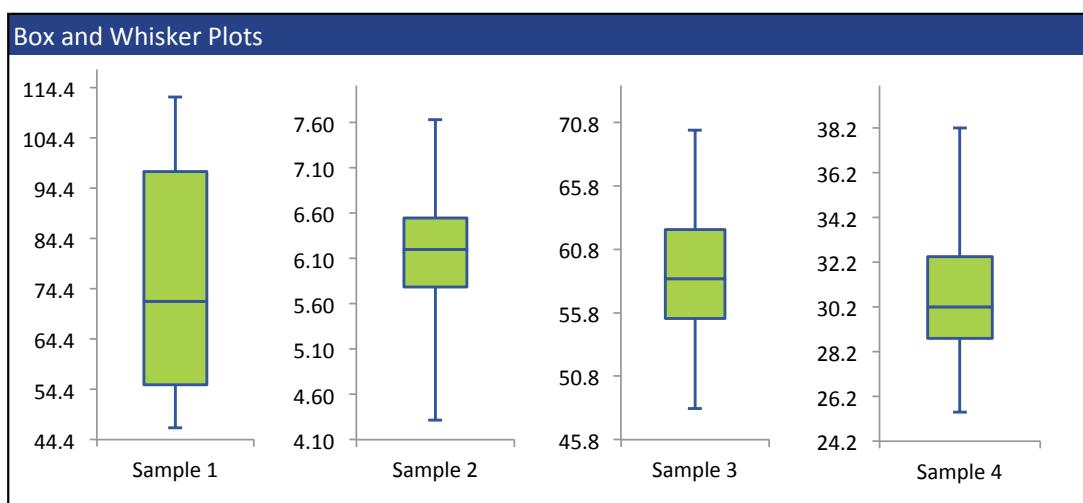
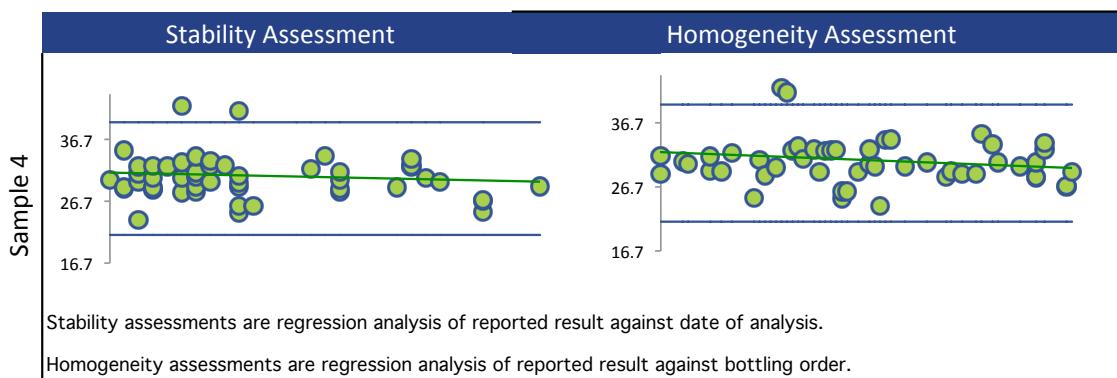
Annex A Summary by Analyte

CHLOROBENZENE



## Annex A Summary by Analyte

### CHLOROBENZENE



Annex A Summary by Analyte

## CHLORODIBROMOMETHANE

### Summary Statistics

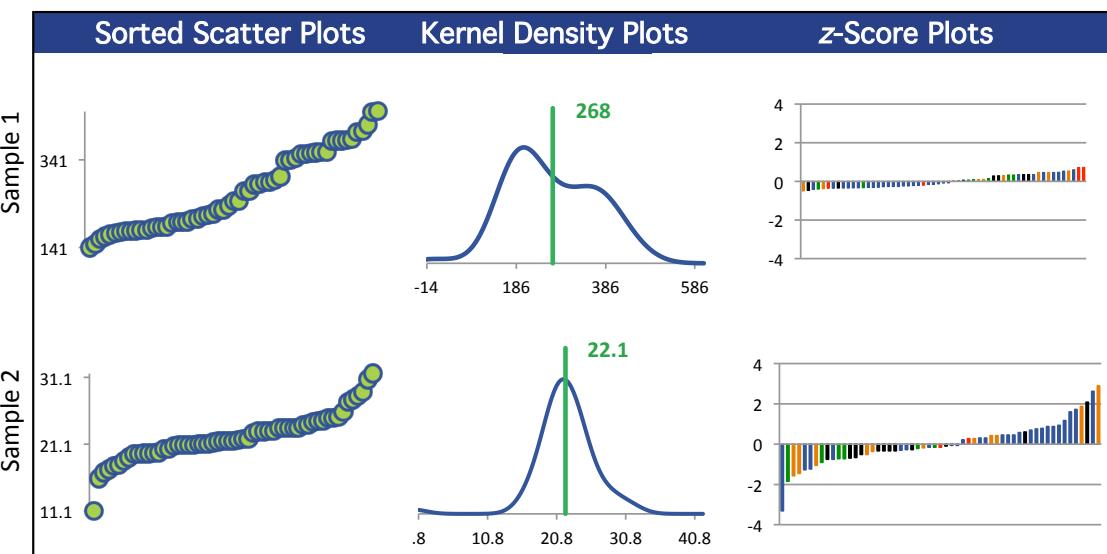
### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	57	57	57	56
Median $\mu\text{g/L}$	246	21.6	214	110
Robust Mean $\mu\text{g/L}$	268	22.1	215	110
$U \mu\text{g/L}$	16.3	0.497	4.19	2.09
Robust Standard Deviation $\mu\text{g/L}$	98.6	3.00	25.3	12.5
Regression Standard Deviation $\mu\text{g/L}$	40.1	3.32	32.2	16.4
Stability Flag				
Homogeneity Flag	Homogeneity			
Standard Deviation Used ( $SDPA \mu\text{g/L}$ )	255	3.32	32.2	16.4
Outliers	0	0	0	1
$ z  > 3.0$	0	1	0	1
$2 <  z  < 3$	0	3	4	3

### Methods Used

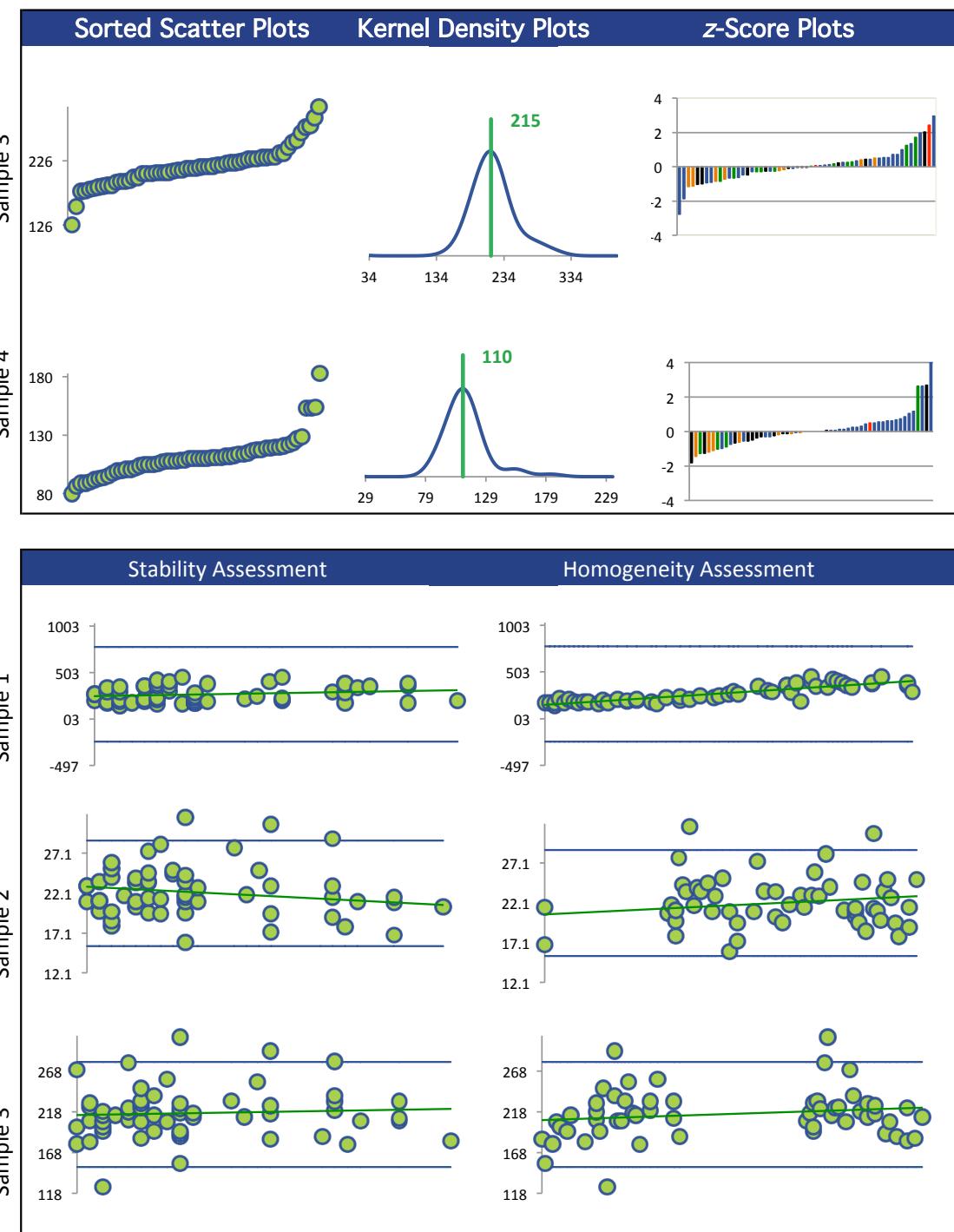
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	28	28	28	27
GC/MS (Red)	5	5	5	5
GC/MS - HEADSPACE (Green)	22	22	22	22
GC/MS/MS - HEADSPACE (Orange)	1	1	1	1
GC/FID - PURGE AND TRAP (Black)	1	1	1	1

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



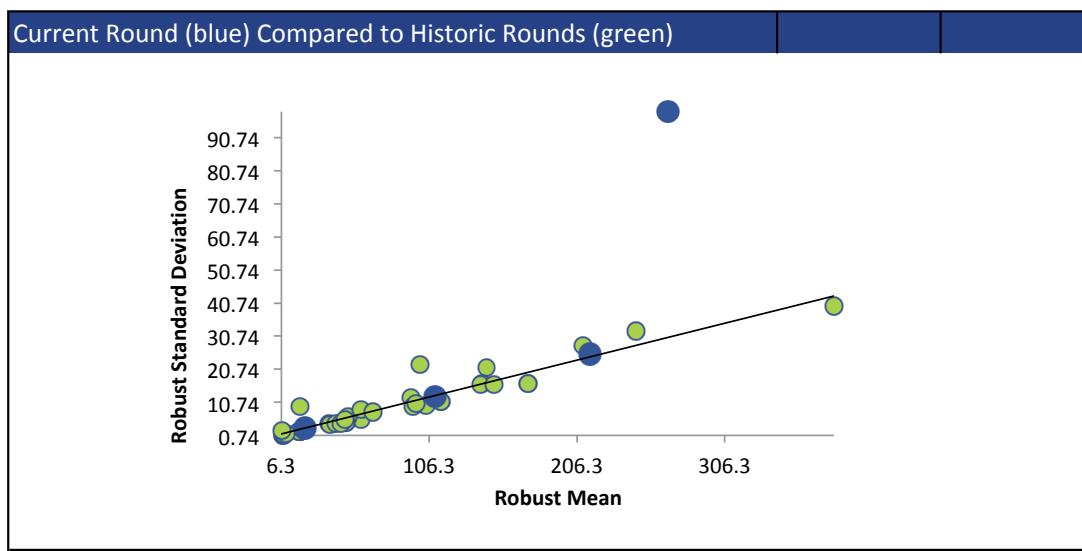
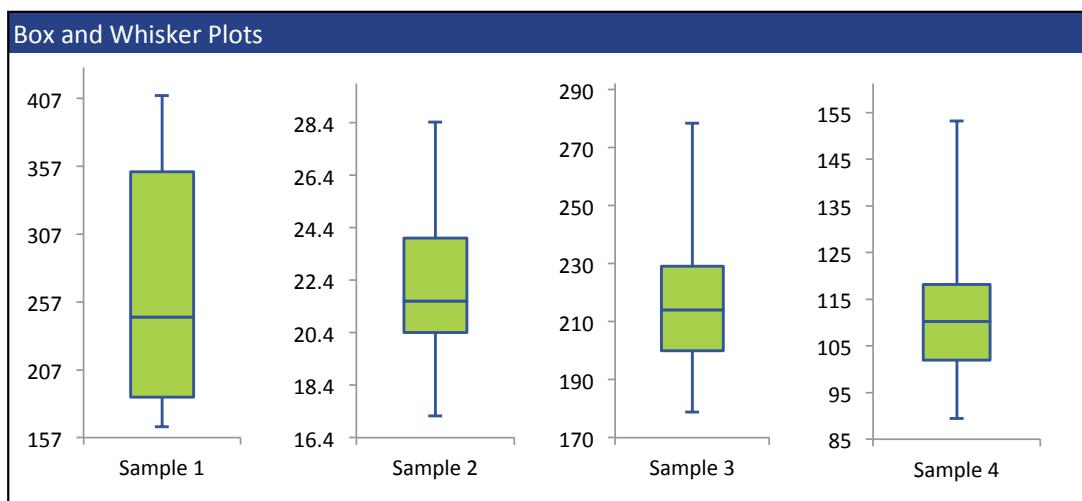
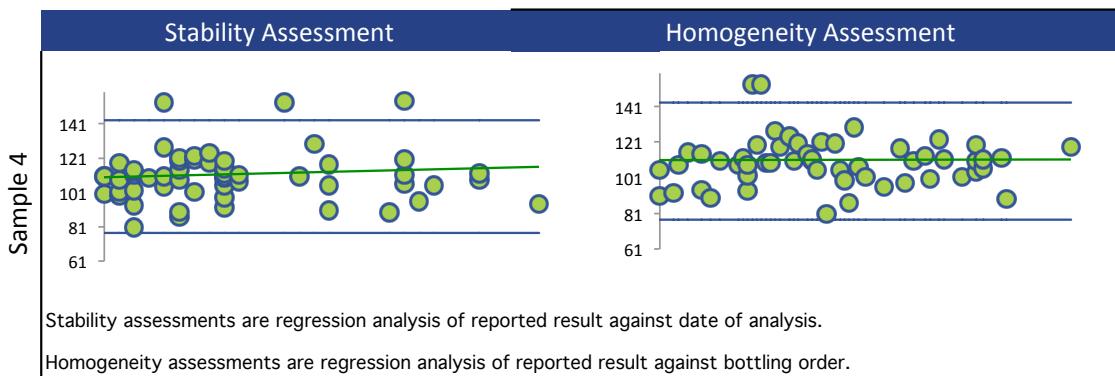
Annex A Summary by Analyte

## CHLORODIBROMOMETHANE



## Annex A Summary by Analyte

### CHLORODIBROMOMETHANE



## Annex A Summary by Analyte

### CHLOROFORM

#### Summary Statistics

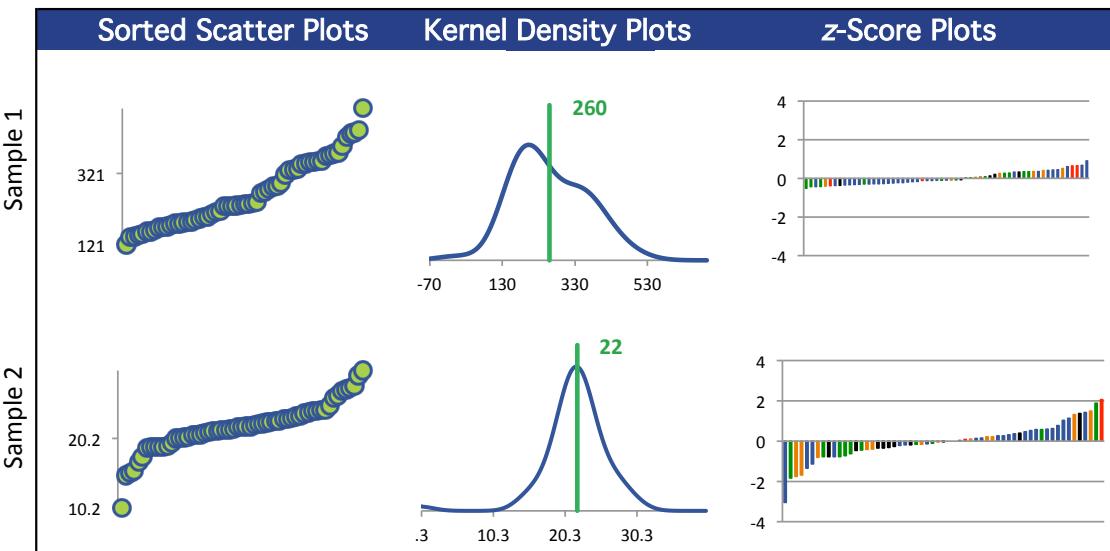
#### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	59	59	59	59
Median $\mu\text{g/L}$	234	21.9	211	108
Robust Mean $\mu\text{g/L}$	260	22.0	210	107
U $\mu\text{g/L}$	16.3	0.483	4.78	2.16
Robust Standard Deviation $\mu\text{g/L}$	100	2.97	29.4	13.3
Regression Standard Deviation $\mu\text{g/L}$	39.0	3.29	31.5	16.1
Stability Flag				
Homogeneity Flag	Homogeneity	Homogeneity		
Standard Deviation Used (SDPA) $\mu\text{g/L}$	266	3.88	31.5	16.1
Outliers	0	0	0	0
$ z  > 3.0$	0	1	1	3
$2 <  z  < 3$	0	1	5	1

#### Methods Used

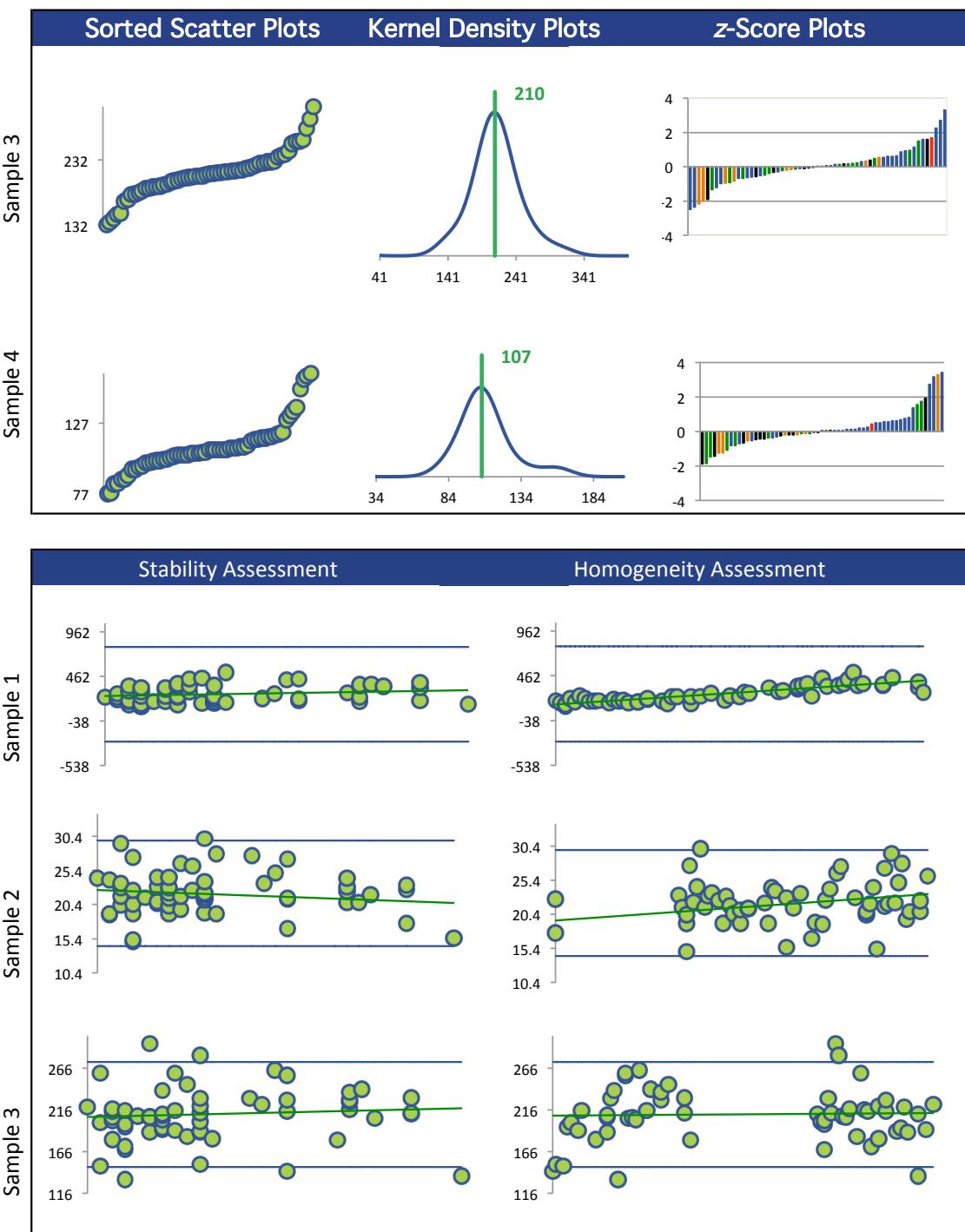
Method	C16-1	C16-2	C16-3	C16-4
GC/MS (Blue)	5	5	5	5
GC/MS - PURGE AND TRAP (Red)	30	30	30	30
GC/MS - HEADSPACE (Green)	22	22	22	22
GC/MS/MS - HEADSPACE (Orange)	1	1	1	1
GC/FID - PURGE AND TRAP (Black)	1	1	1	1

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



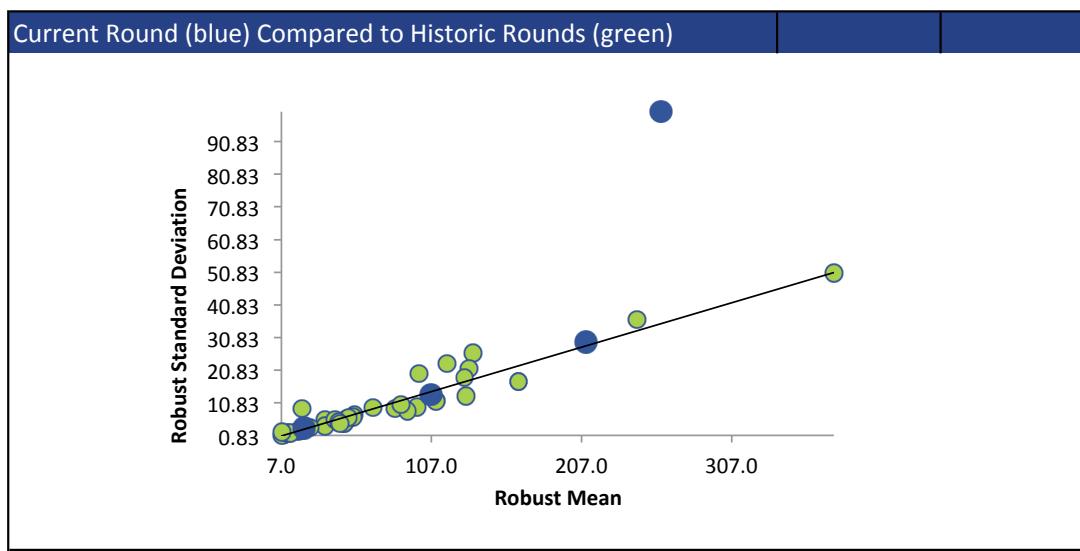
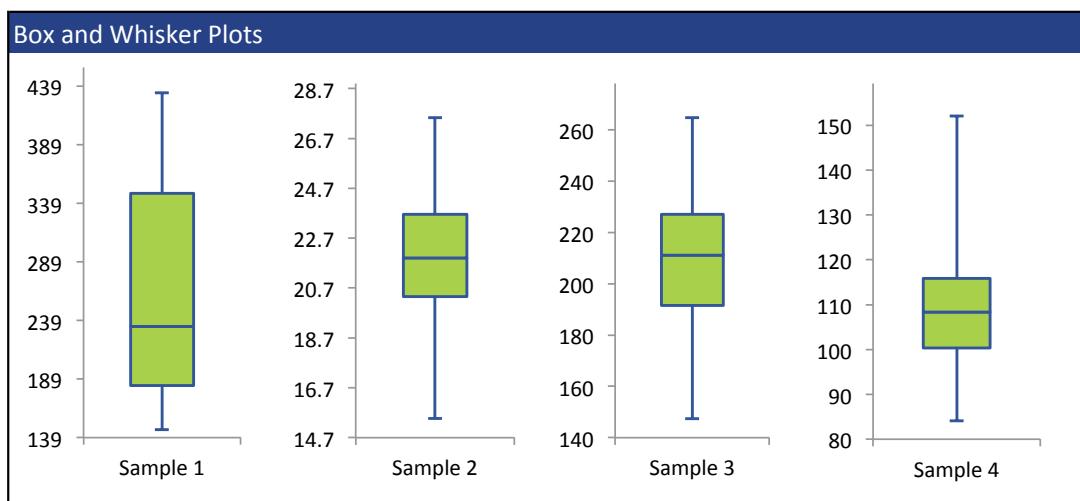
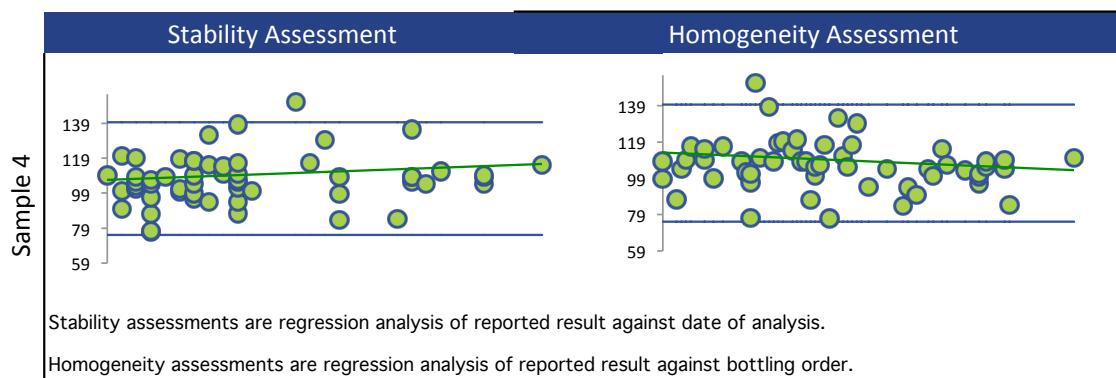
Annex A Summary by Analyte

CHLOROFORM



## Annex A Summary by Analyte

### CHLOROFORM



Annex A Summary by Analyte

## CIS-1,2-DICHLOROETHYLENE

### Summary Statistics

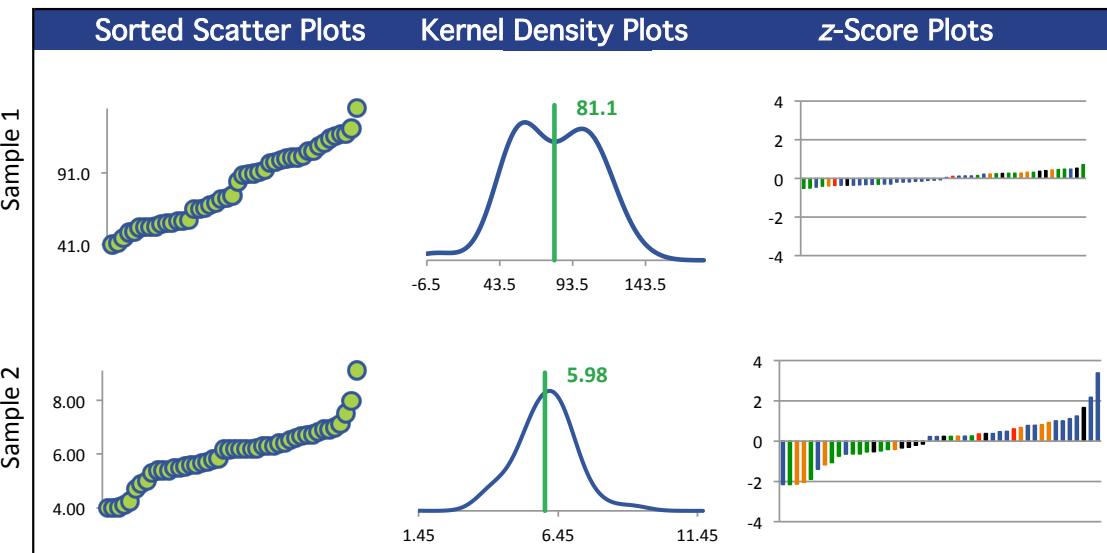
### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	46	46	45	46
Median $\mu\text{g/L}$	80.0	6.19	61.0	30.6
Robust Mean $\mu\text{g/L}$	81.1	5.98	61.4	30.5
$U \mu\text{g/L}$	5.33	0.170	1.38	0.599
Robust Standard Deviation $\mu\text{g/L}$	28.9	0.923	7.43	3.25
Regression Standard Deviation $\mu\text{g/L}$	12.2	0.897	9.21	4.58
Stability Flag				Stability
Homogeneity Flag	Homogeneity			
Standard Deviation Used ( $SDPA \mu\text{g/L}$ )	77.4	0.923	9.21	4.76
Outliers	0	0	1	0
$ z  > 3.0$	0	1	0	0
$2 <  z  < 3$	0	5	2	4

### Methods Used

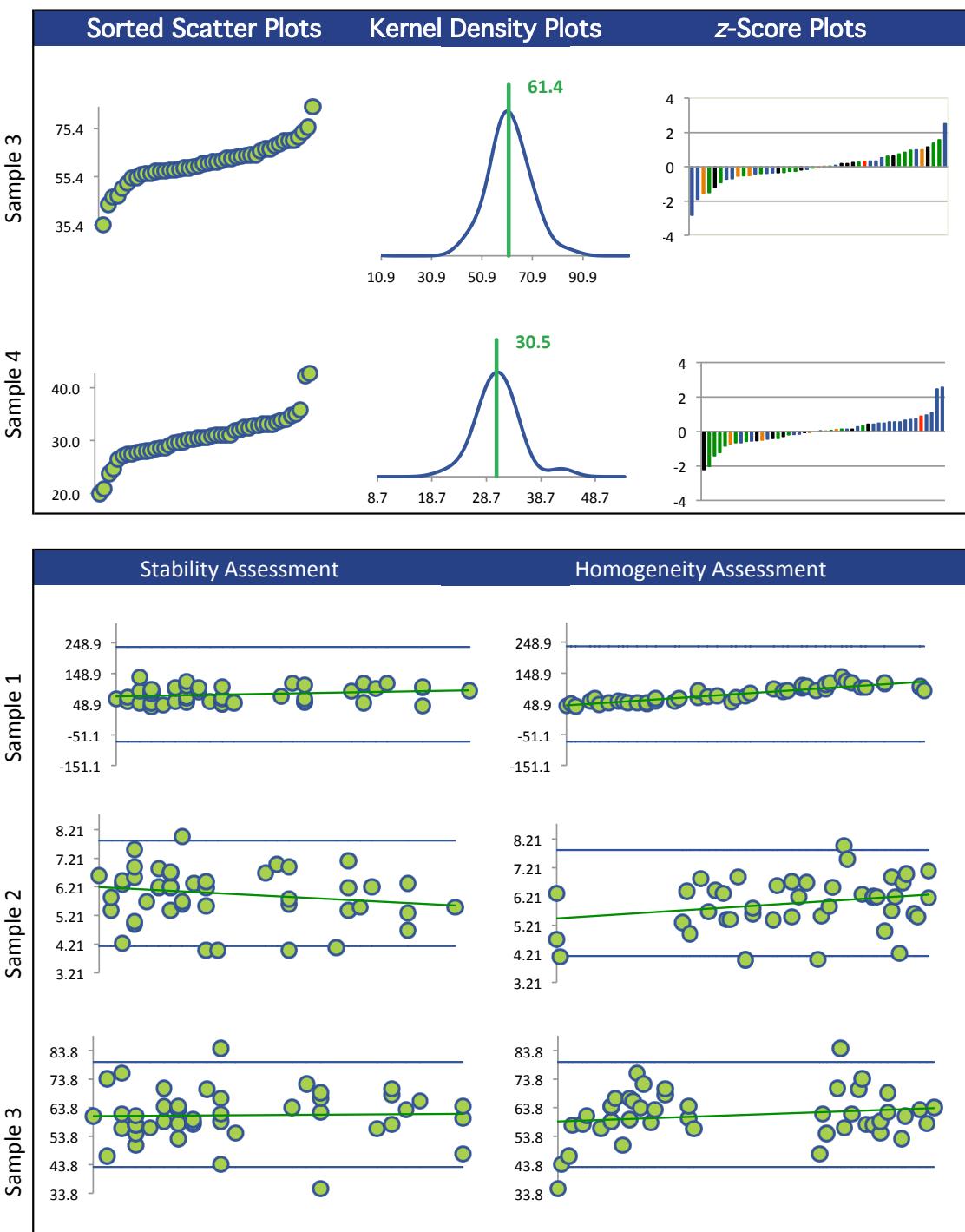
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	24	24	23	24
GC/MS (Red)	3	3	3	3
GC/MS - HEADSPACE (Green)	18	18	18	18
GC/FID - PURGE AND TRAP (Orange)	1	1	1	1

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



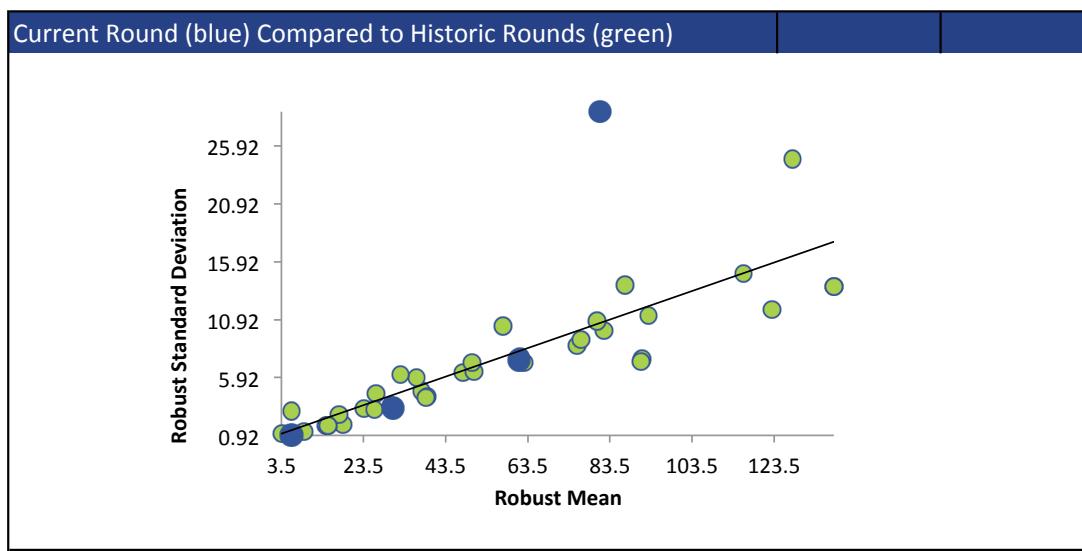
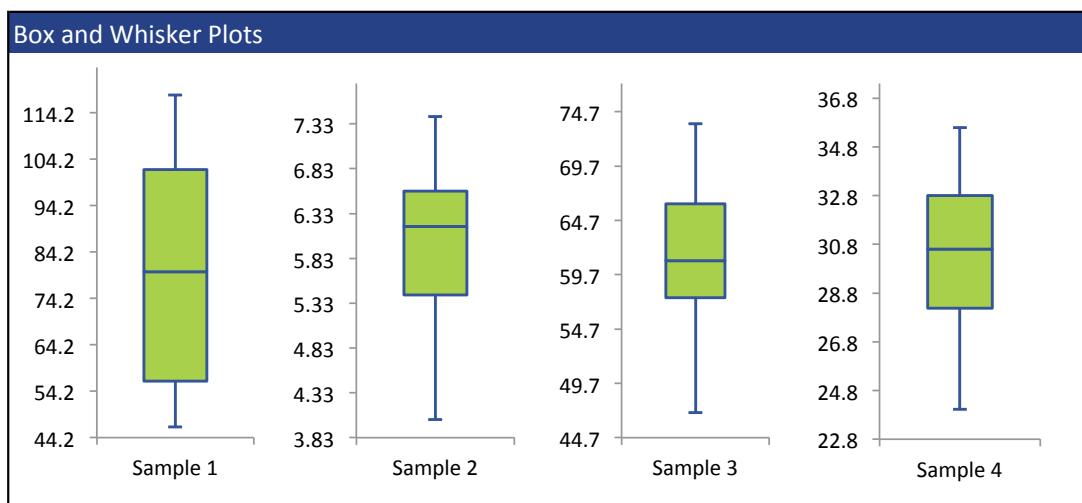
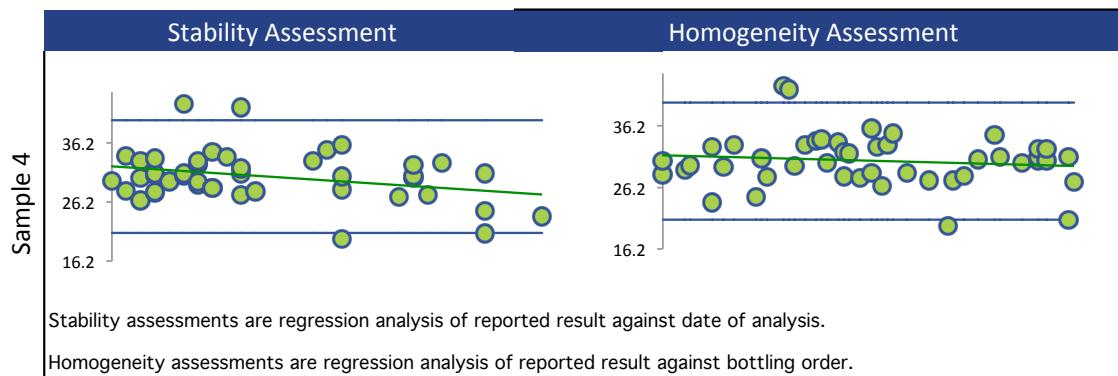
Annex A Summary by Analyte

CIS-1,2-DICHLOROETHYLENE



## Annex A Summary by Analyte

### CIS-1,2-DICHLOROETHYLENE



Annex A Summary by Analyte

## CIS-1,3-DICHLOROPROPENE

### Summary Statistics

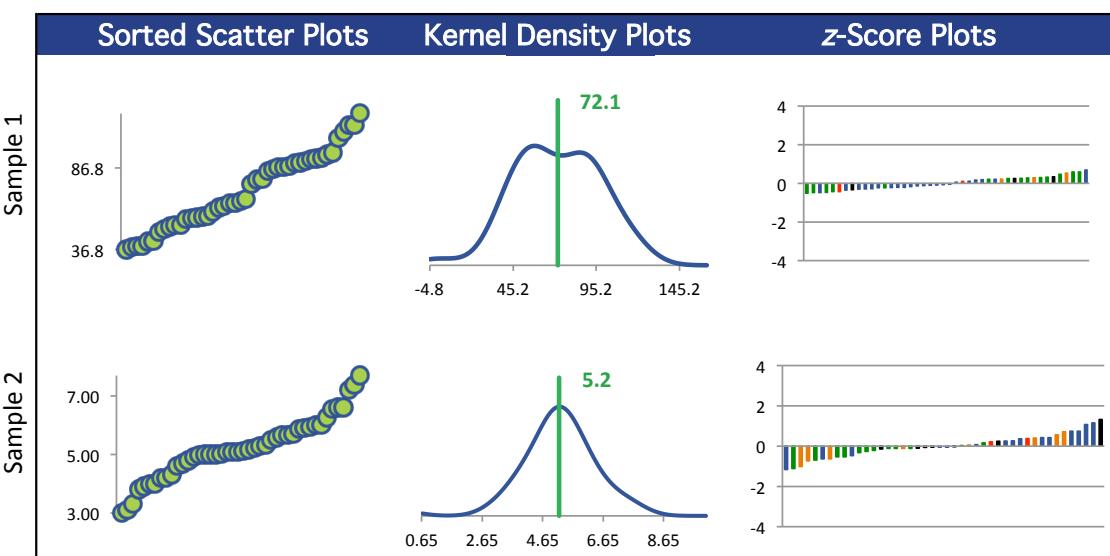
### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	44	44	44	44
Median $\mu\text{g/L}$	67.1	5.11	52.4	26.8
Robust Mean $\mu\text{g/L}$	72.1	5.20	52.9	27.0
$U \mu\text{g/L}$	4.82	0.194	1.67	0.767
Robust Standard Deviation $\mu\text{g/L}$	25.6	1.03	8.85	4.07
Regression Standard Deviation $\mu\text{g/L}$	10.8	0.781	7.94	4.06
Stability Flag		Stability		Stability
Homogeneity Flag	Homogeneity			
Standard Deviation Used ( $SDPA \mu\text{g/L}$ )	68.4	1.89	8.85	5.77
Outliers	0	0	0	0
$ z  > 3.0$	0	0	1	0
$2 <  z  < 3$	0	0	3	1

### Methods Used

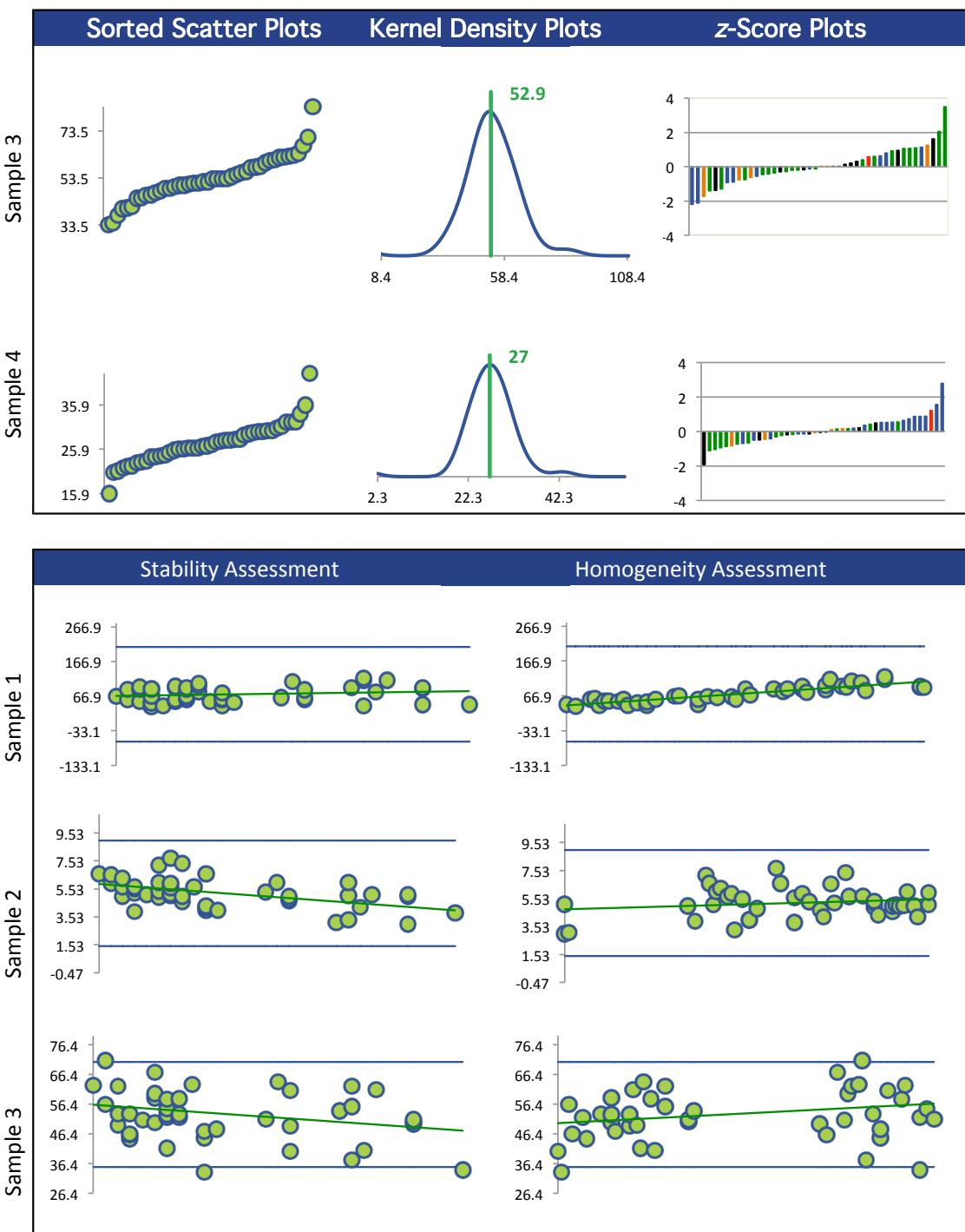
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	23	23	23	23
GC/MS - HEADSPACE (Red)	18	18	18	18
GC/MS (Green)	3	3	3	3

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



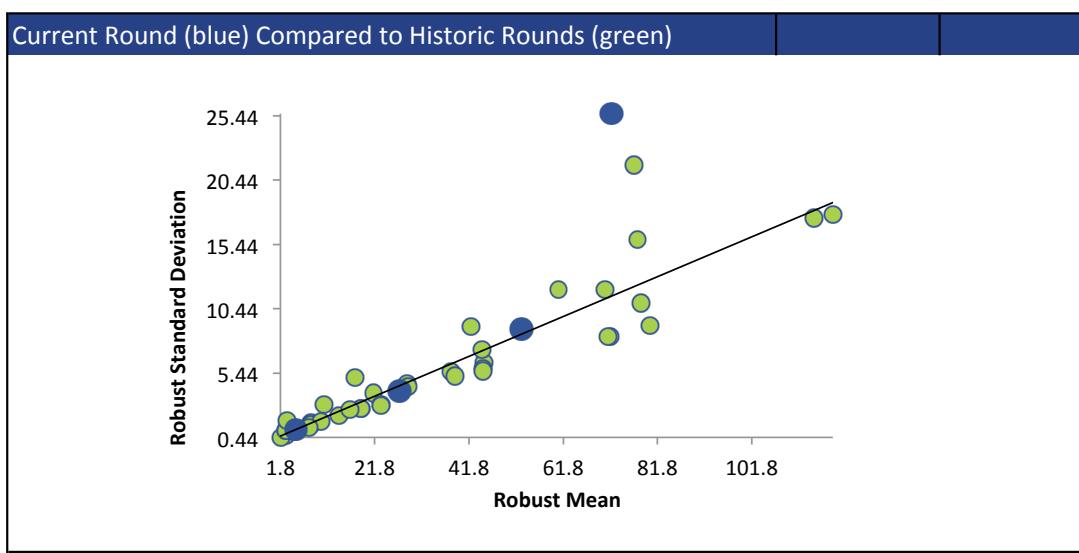
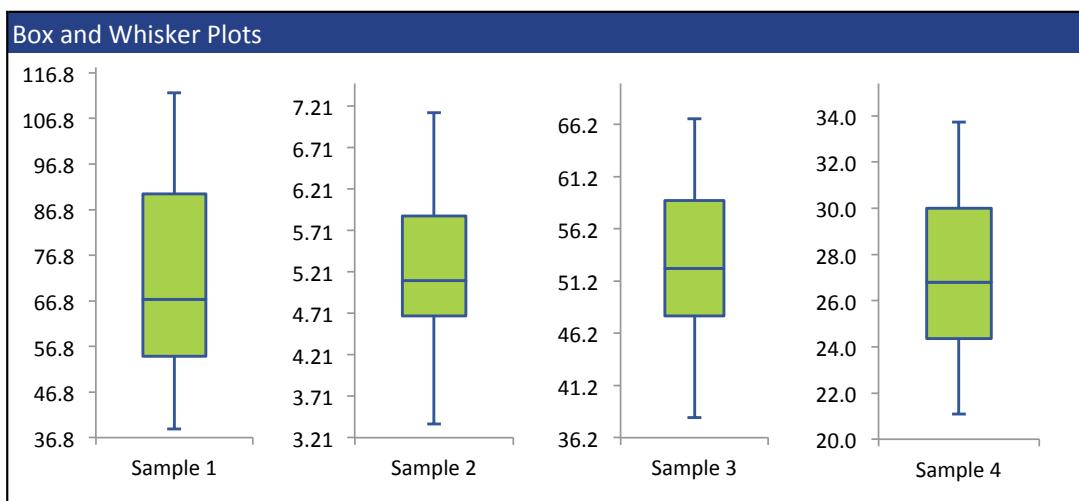
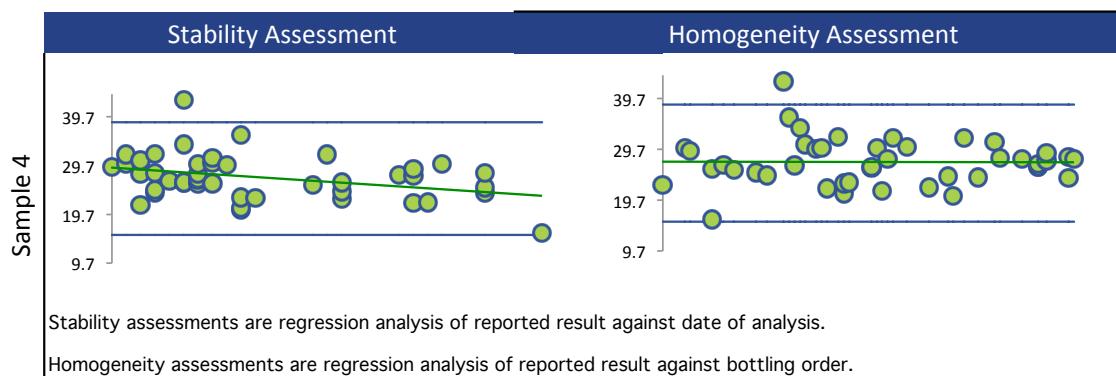
Annex A Summary by Analyte

CIS-1,3-DICHLOROPROPENE



## Annex A Summary by Analyte

### CIS-1,3-DICHLOROPROPENE



## Annex A Summary by Analyte

### DICHLOROMETHANE

#### Summary Statistics

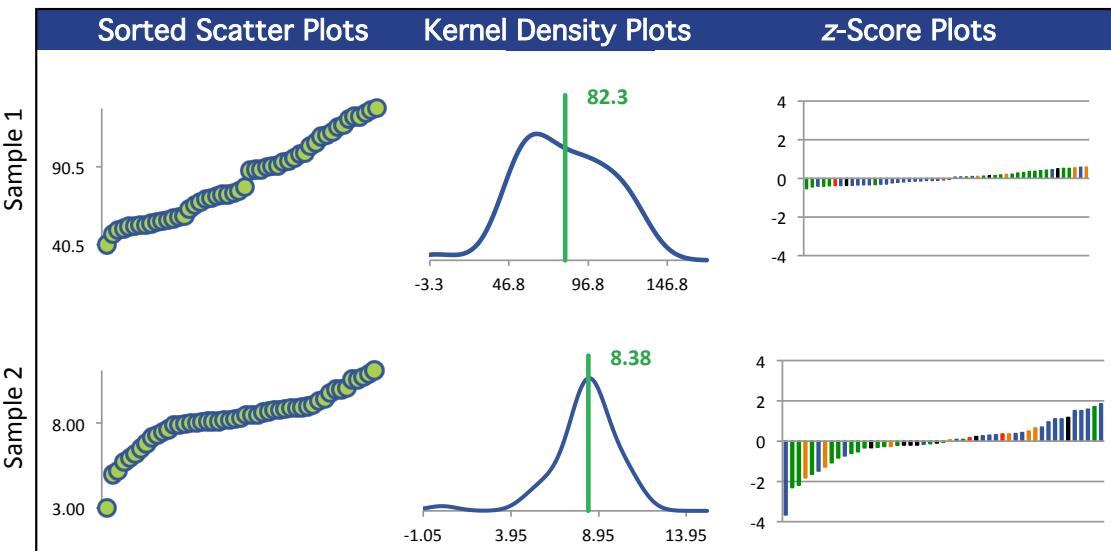
#### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	50	49	50	50
Median $\mu\text{g/L}$	76.2	8.30	63.3	33.1
Robust Mean $\mu\text{g/L}$	82.3	8.38	64.0	33.4
$U \mu\text{g/L}$	5.16	0.257	1.52	0.764
Robust Standard Deviation $\mu\text{g/L}$	29.2	1.44	8.59	4.32
Regression Standard Deviation $\mu\text{g/L}$	14.4	1.47	11.2	5.85
Stability Flag			Stability	
Homogeneity Flag	Homogeneity			
Standard Deviation Used ( $SDPA \mu\text{g/L}$ )	77.7	1.47	14.3	5.85
Outliers	0	0	0	0
$ z  > 3.0$	0	1	0	0
$2 <  z  < 3$	0	2	0	3

#### Methods Used

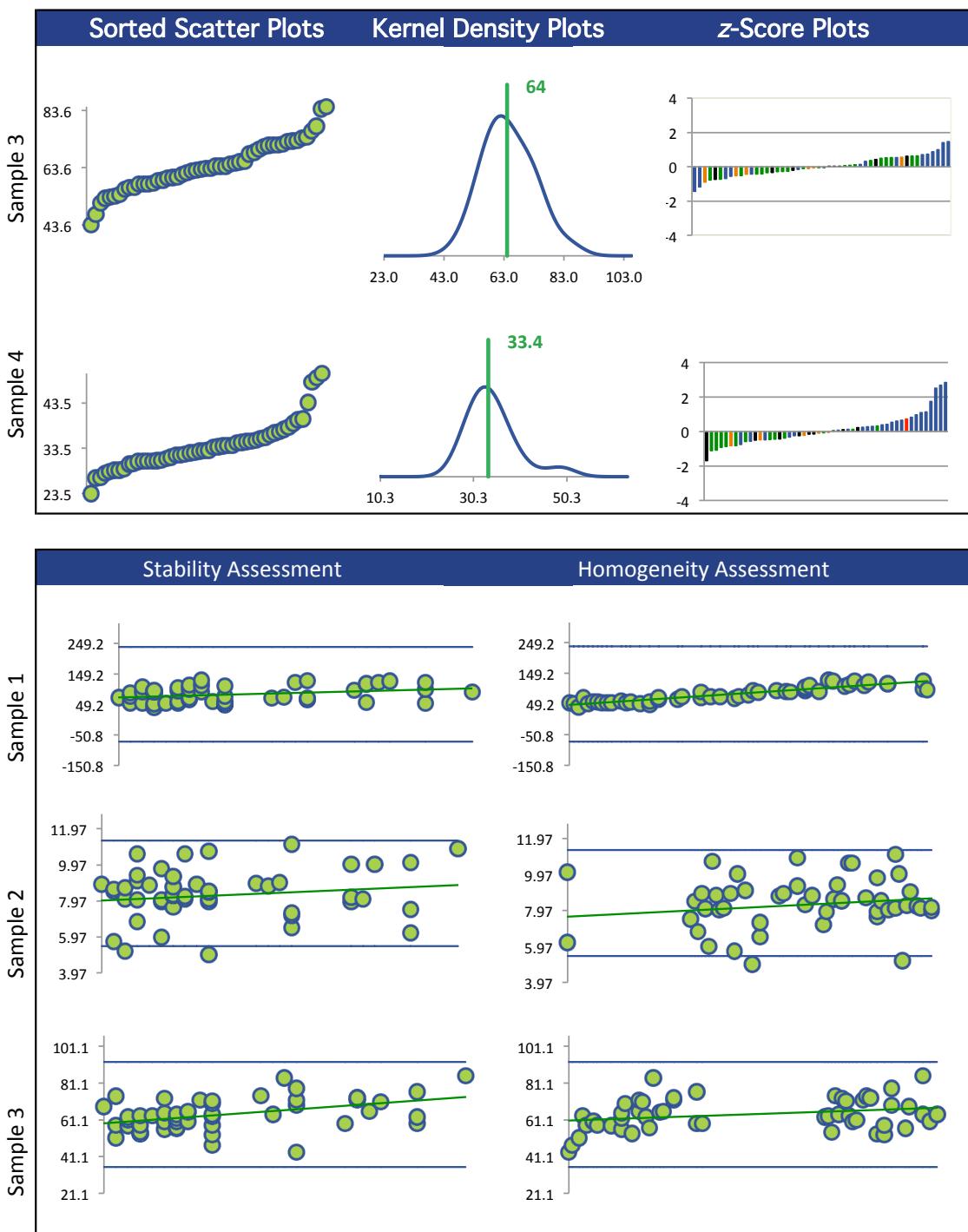
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	27	26	27	27
GC/MS - HEADSPACE (Red)	19	19	19	19
GC/MS/MS - HEADSPACE (Green)	1	1	1	1
GC/FID - PURGE AND TRAP (Orange)	1	1	1	1
GC/MS (Black)	2	2	2	2

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



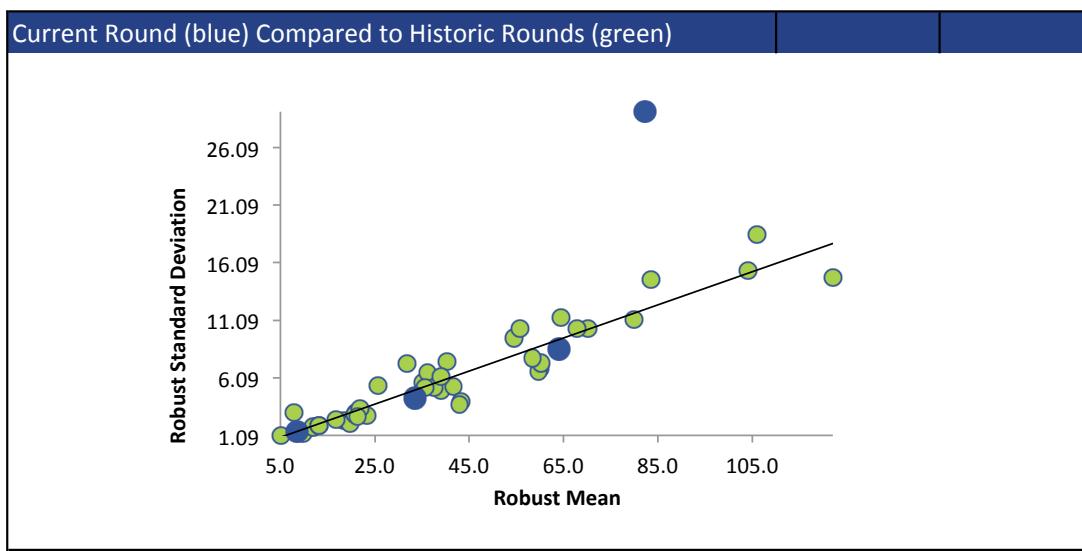
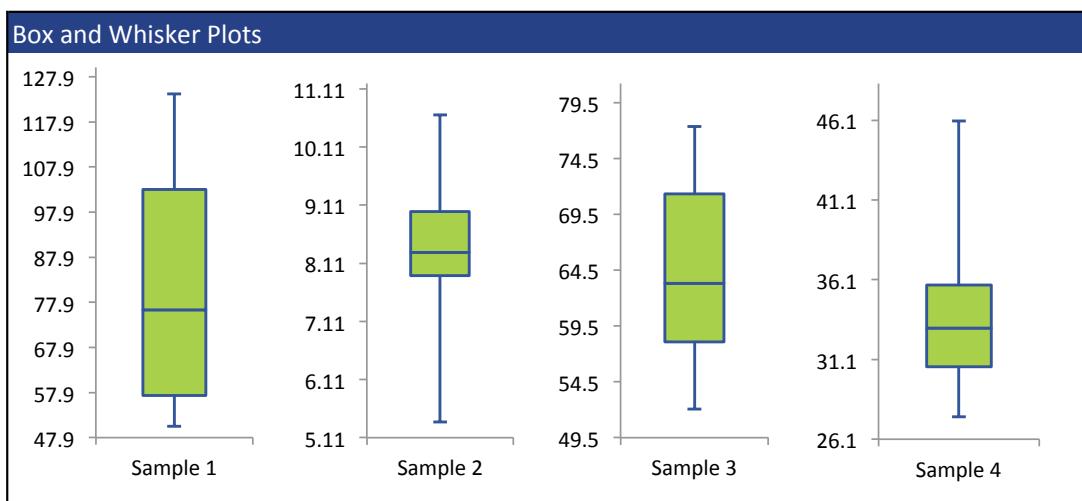
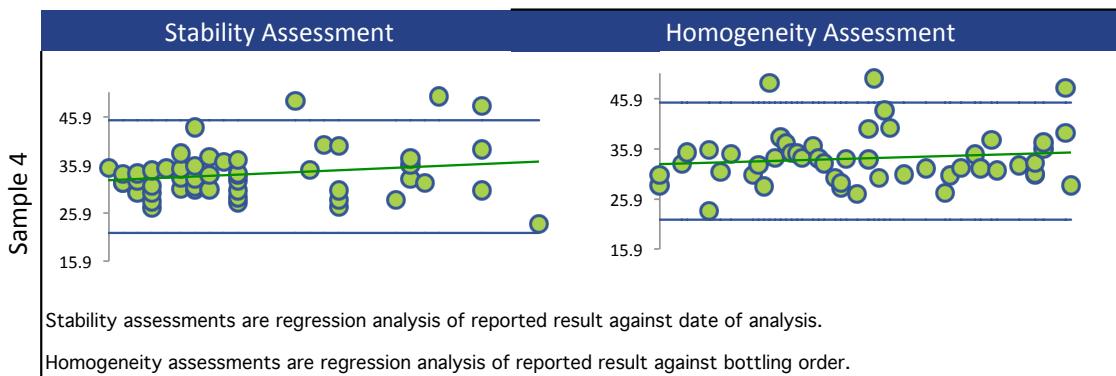
Annex A Summary by Analyte

DICHLOROMETHANE



## Annex A Summary by Analyte

### DICHLOROMETHANE



Annex A Summary by Analyte

## ETHYLBENZENE

### Summary Statistics

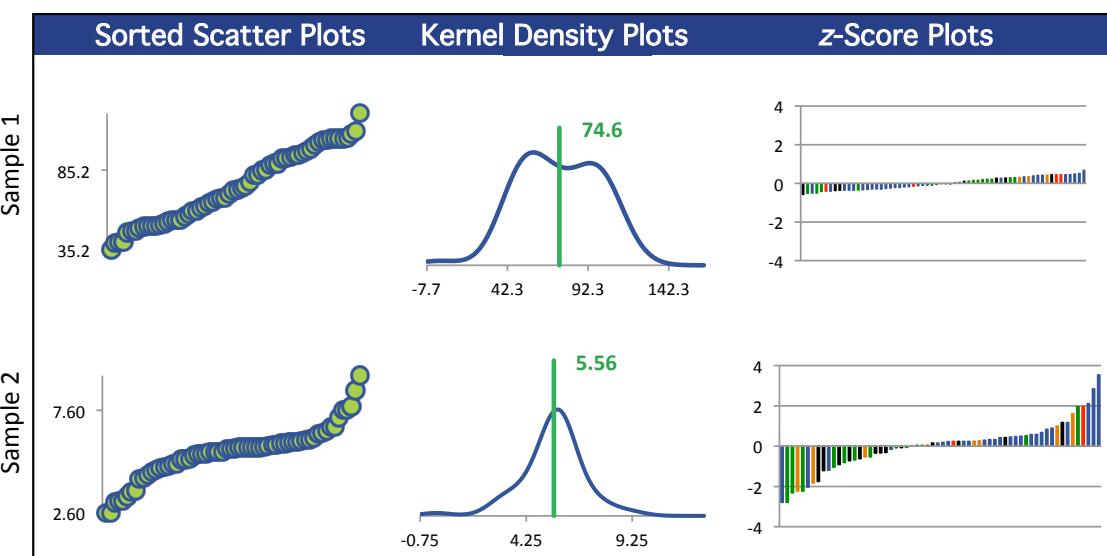
### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	62	62	62	62
Median $\mu\text{g/L}$	72.8	5.74	56.9	29.0
Robust Mean $\mu\text{g/L}$	74.6	5.56	56.7	28.9
$U \mu\text{g/L}$	4.08	0.168	1.40	0.594
Robust Standard Deviation $\mu\text{g/L}$	25.7	1.06	8.80	3.74
Regression Standard Deviation $\mu\text{g/L}$	11.2	0.835	8.50	4.34
Stability Flag				
Homogeneity Flag	Homogeneity			
Standard Deviation Used (SDPA) $\mu\text{g/L}$	67.9	1.06	8.80	4.34
Outliers	0	0	0	0
$ z  > 3.0$	0	1	0	2
$2 <  z  < 3$	0	8	1	3

### Methods Used

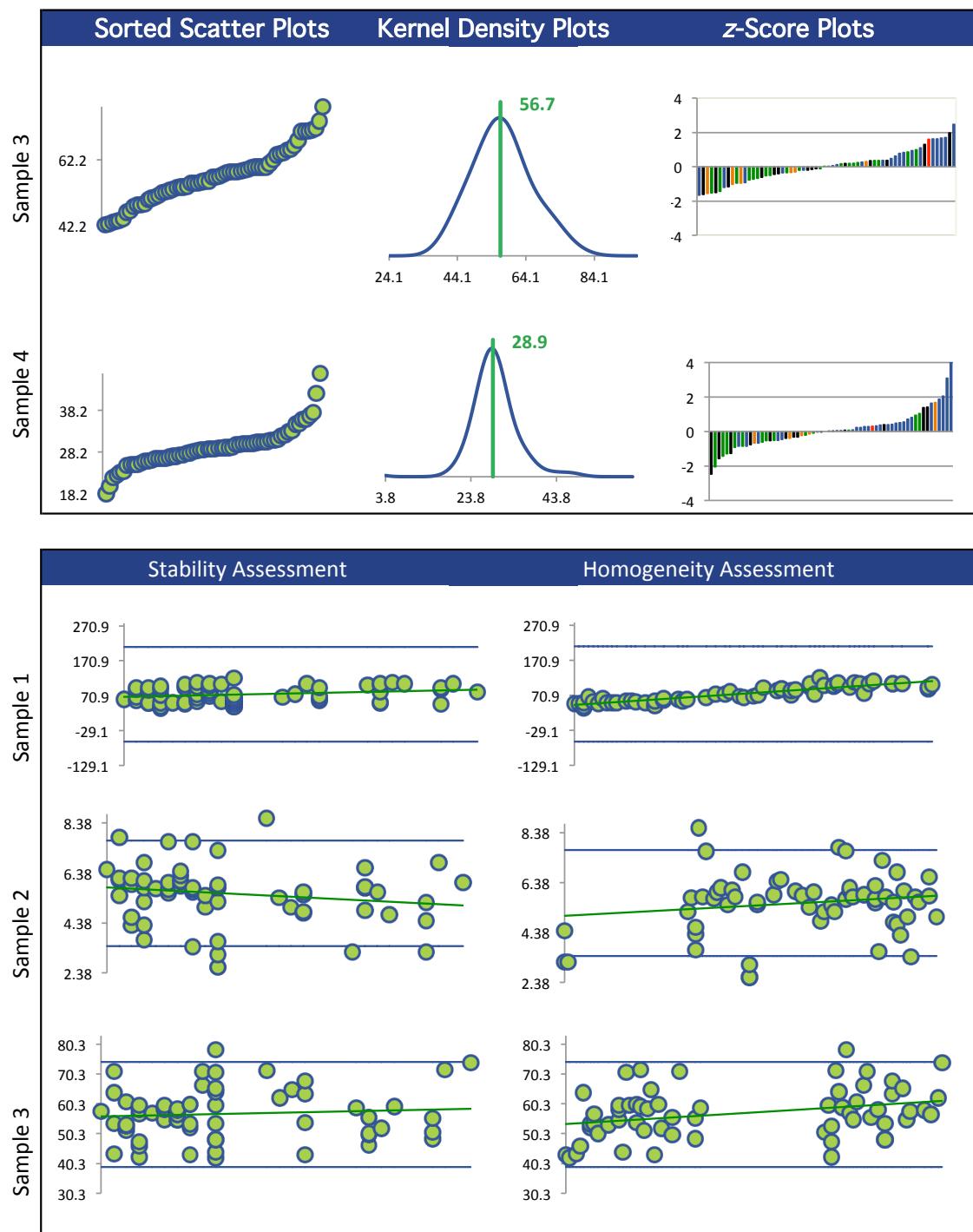
Method	C16-1	C16-2	C16-3	C16-4
GC/MS (Blue)	6	6	6	6
GC/MS - PURGE AND TRAP (Red)	31	31	31	31
GC/MS - HEADSPACE (Green)	22	22	22	22
GC/FID - HEADSPACE (Orange)	1	1	1	1
GC/MS/MS - HEADSPACE (Black)	1	1	1	1
GC/FID - PURGE AND TRAP (Yellow)	1	1	1	1

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



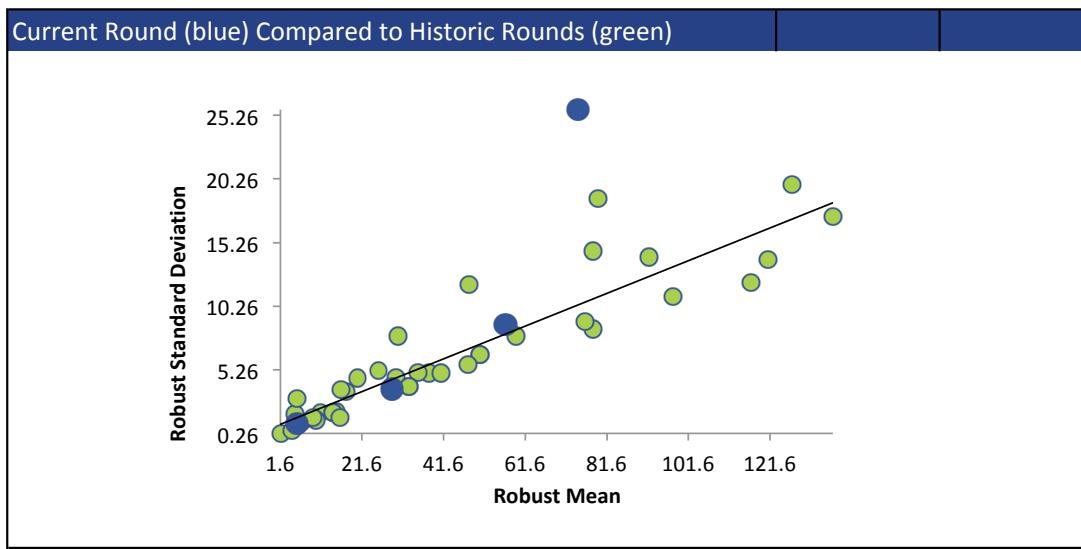
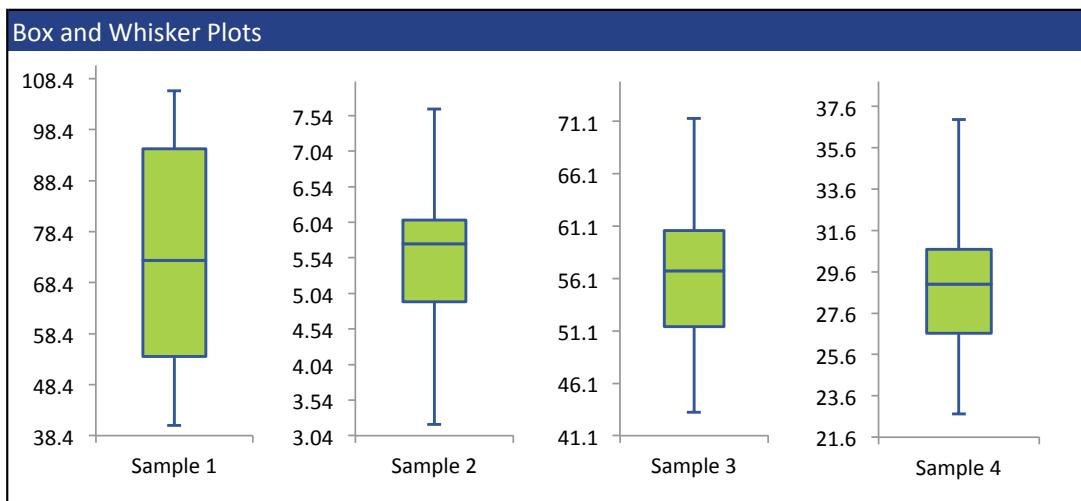
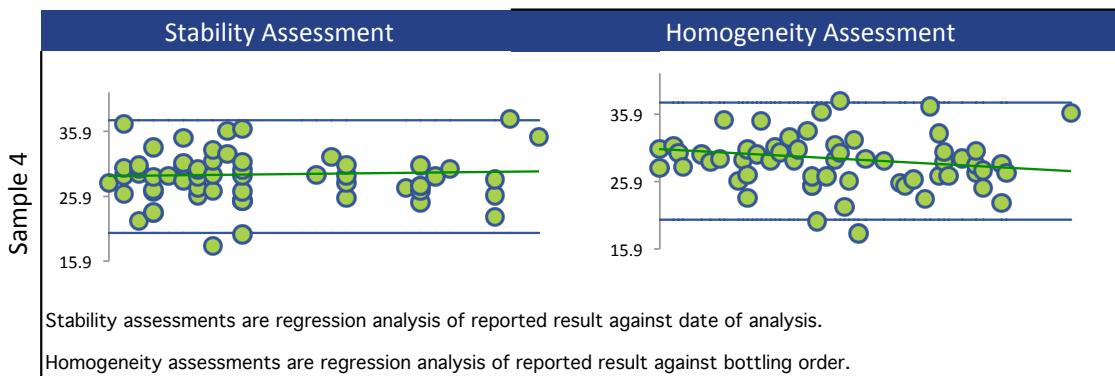
Annex A Summary by Analyte

ETHYLBENZENE



## Annex A Summary by Analyte

### ETHYLBENZENE



Annex A Summary by Analyte

## ETHYLENE DIBROMIDE

### Summary Statistics

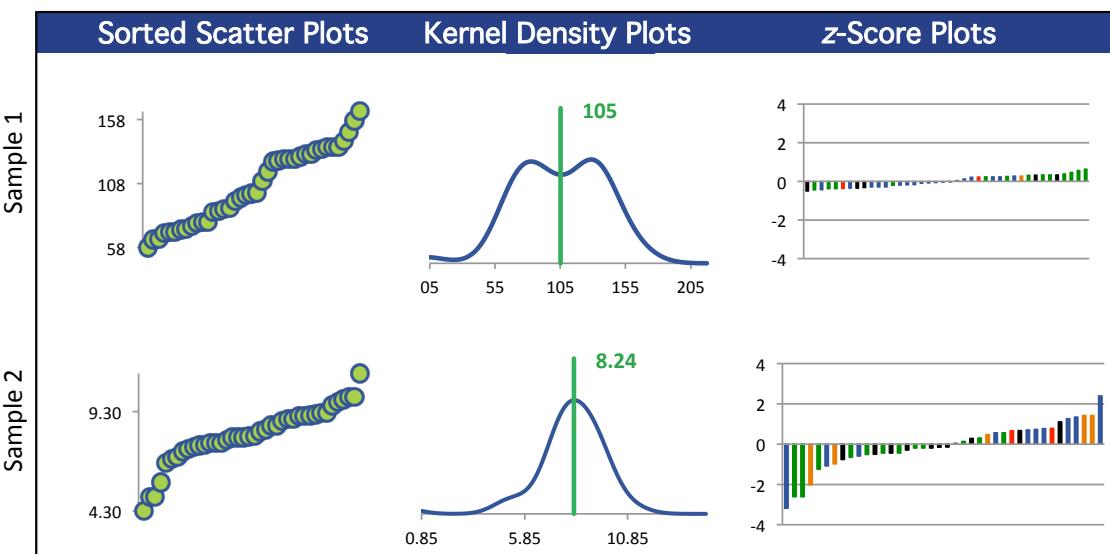
### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	40	40	40	40
Median $\mu\text{g/L}$	101	8.06	76.9	40.0
Robust Mean $\mu\text{g/L}$	105	8.24	78.3	40.6
U $\mu\text{g/L}$	6.56	0.245	1.80	0.876
Robust Standard Deviation $\mu\text{g/L}$	33.2	1.24	9.12	4.43
Regression Standard Deviation $\mu\text{g/L}$	15.8	1.24	11.7	6.09
Stability Flag				
Homogeneity Flag	Homogeneity			
Standard Deviation Used (SDPA) $\mu\text{g/L}$	91.8	1.24	11.7	6.09
Outliers	0	0	0	0
$ z  > 3.0$	0	1	0	0
$2 <  z  < 3$	0	4	1	1

### Methods Used

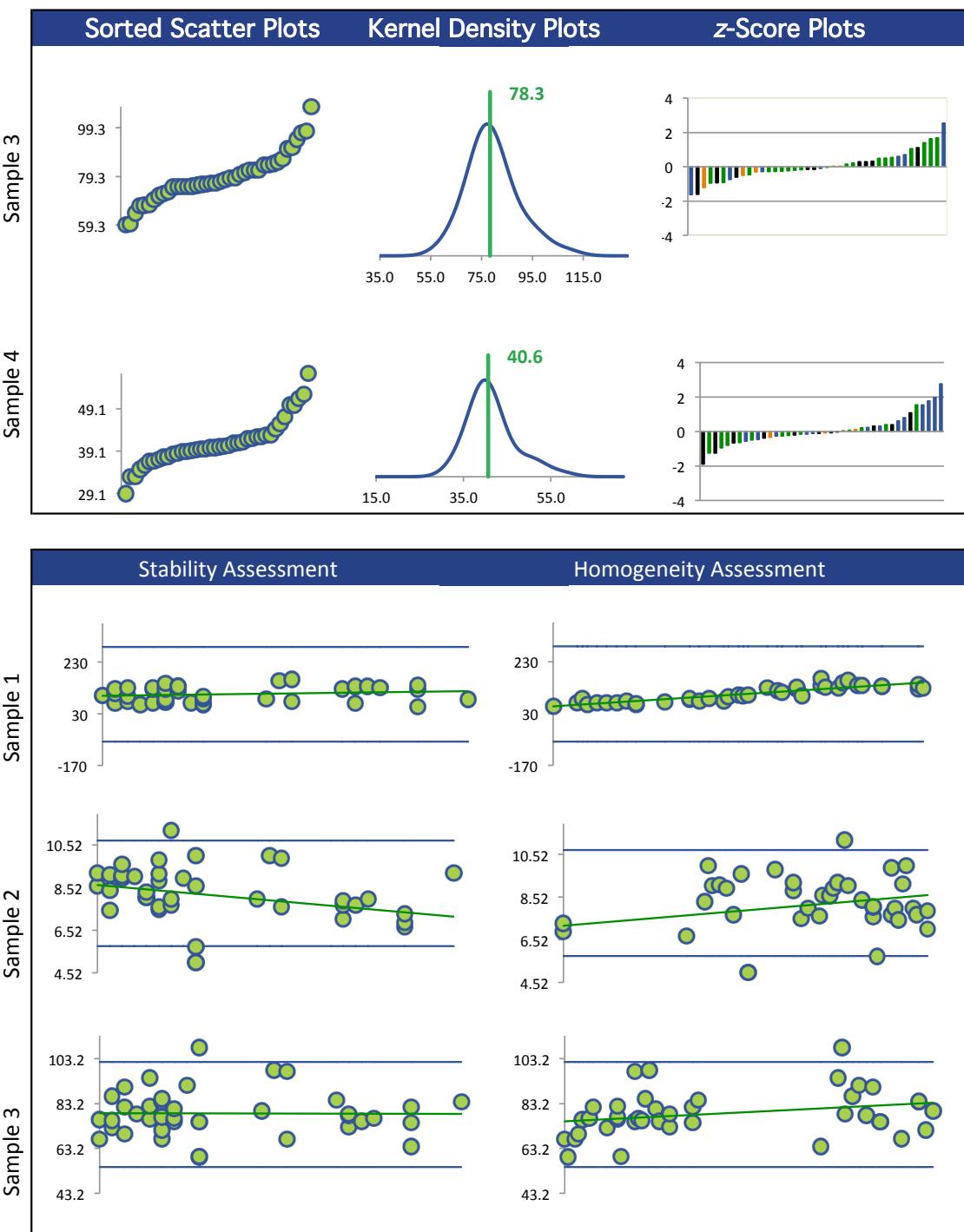
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	22	22	22	22
GC/MS - HEADSPACE (Red)	16	16	16	16
GC/MS (Green)	2	2	2	2

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



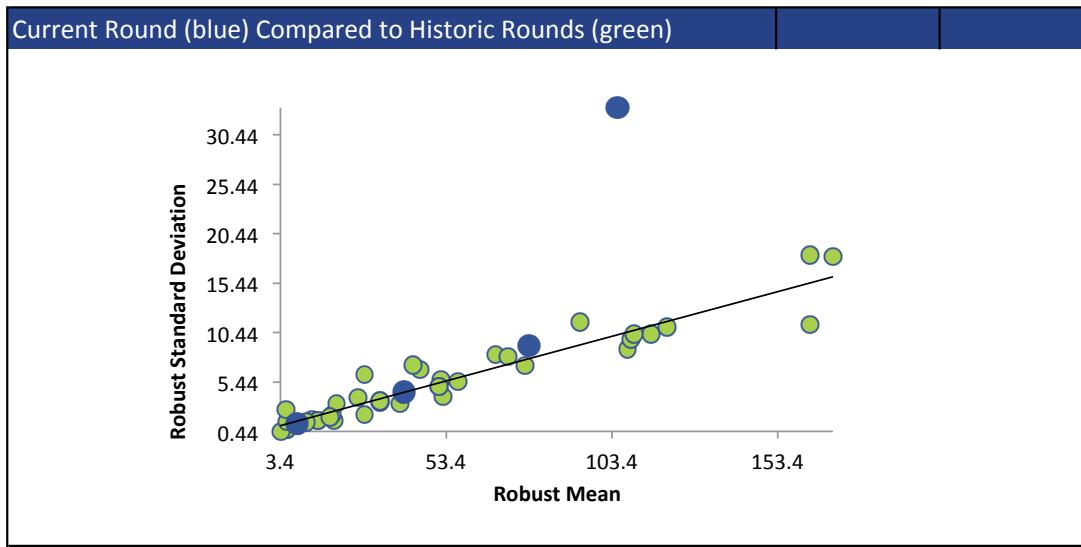
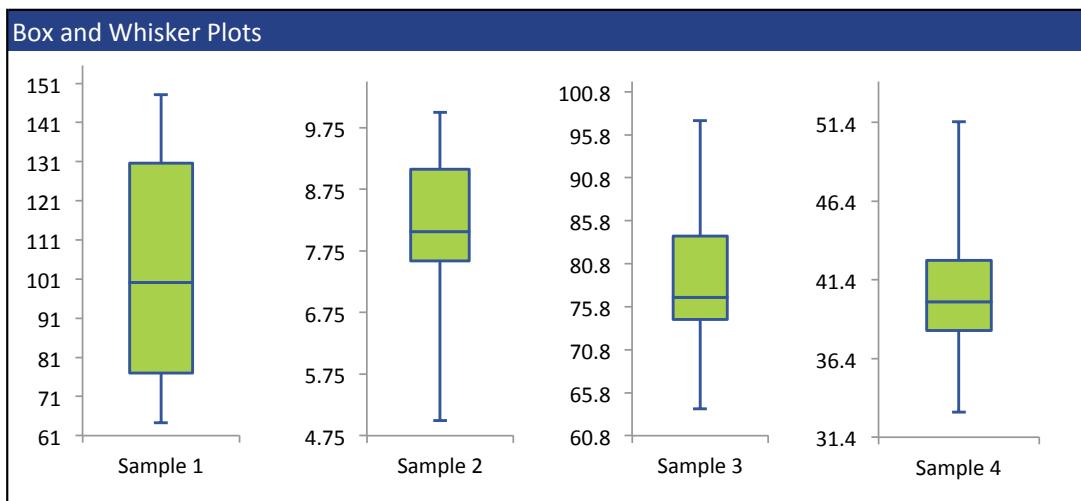
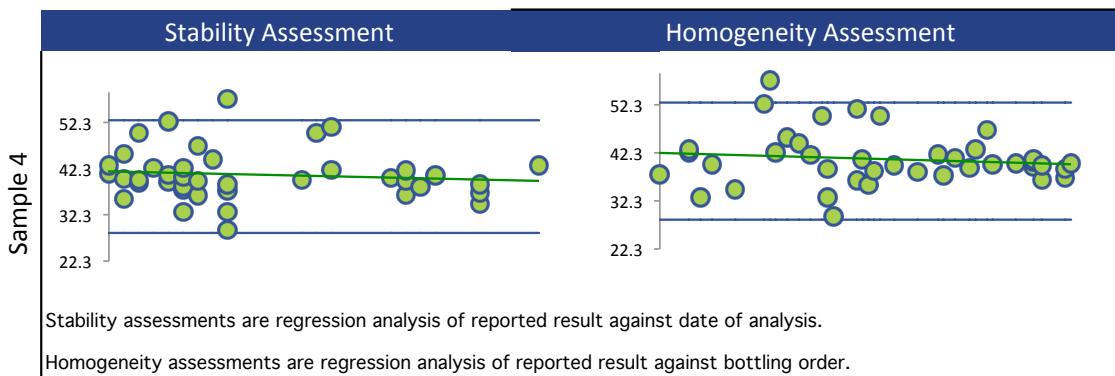
Annex A Summary by Analyte

ETHYLENE DIBROMIDE



## Annex A Summary by Analyte

### ETHYLENE DIBROMIDE



## Annex A Summary by Analyte

### M,P-XYLENE

#### Summary Statistics

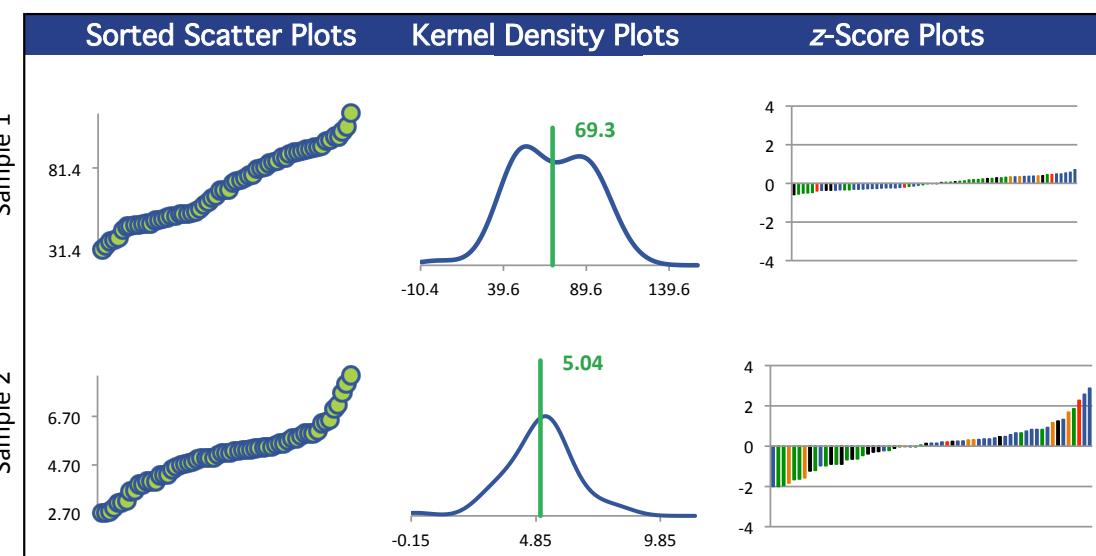
#### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	62	61	62	62
Median $\mu\text{g/L}$	68.2	5.20	53.7	27.5
Robust Mean $\mu\text{g/L}$	69.3	5.04	53.6	27.1
$U \mu\text{g/L}$	3.89	0.187	1.37	0.657
Robust Standard Deviation $\mu\text{g/L}$	24.5	1.17	8.65	4.14
Regression Standard Deviation $\mu\text{g/L}$	10.4	0.757	8.04	4.06
Stability Flag				
Homogeneity Flag	Homogeneity			
Standard Deviation Used ( $SDPA \mu\text{g/L}$ )	64.7	1.17	8.65	4.14
Outliers	0	0	0	0
$ z  > 3.0$	0	0	0	1
$2 <  z  < 3$	0	3	3	6

#### Methods Used

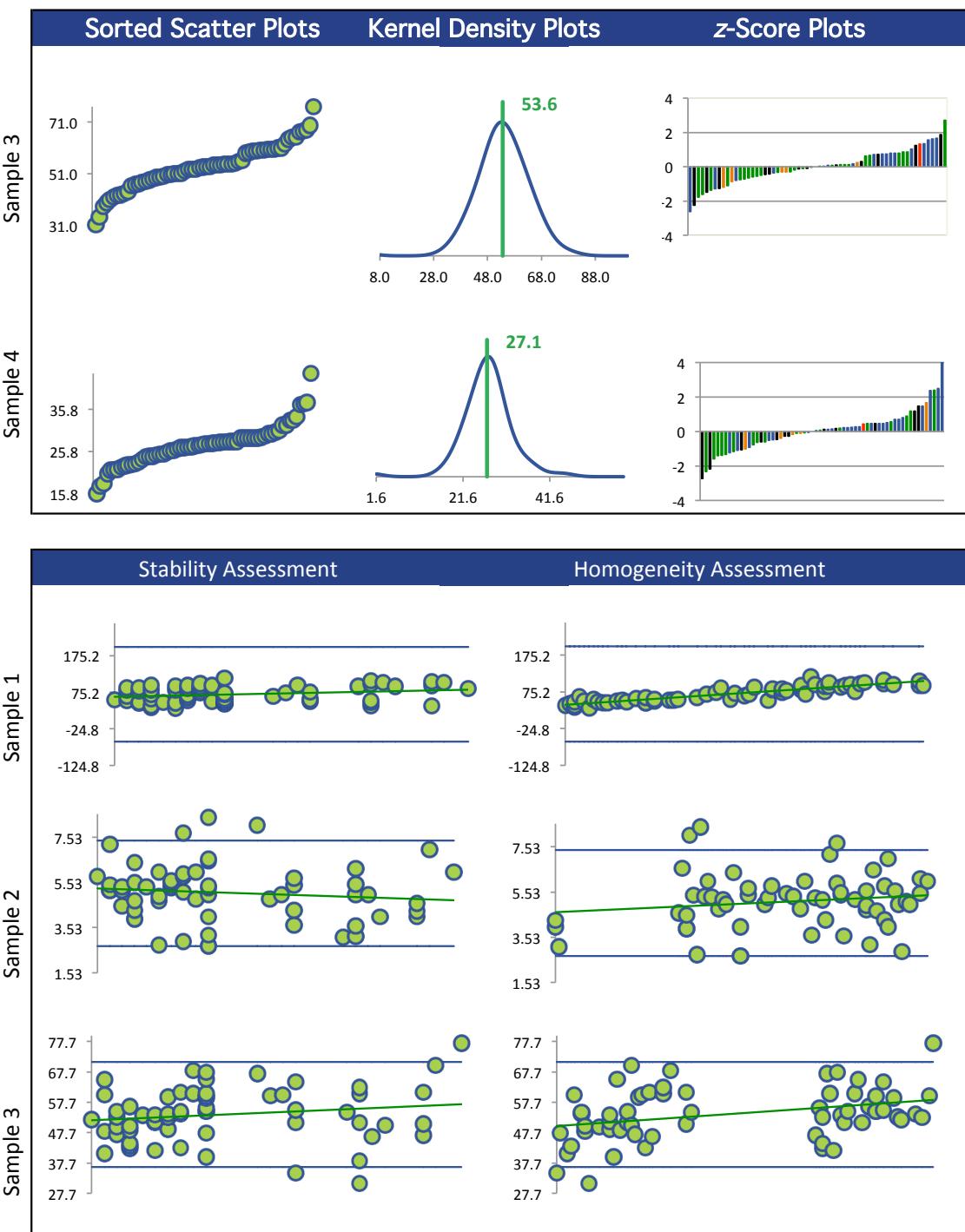
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	31	31	31	31
GC/MS (Red)	6	5	6	6
GC/MS - HEADSPACE (Green)	22	22	22	22
GC/FID - HEADSPACE (Orange)	1	1	1	1
GC/MS/MS - HEADSPACE (Black)	1	1	1	1
GC/FID - PURGE AND TRAP (Yellow)	1	1	1	1

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



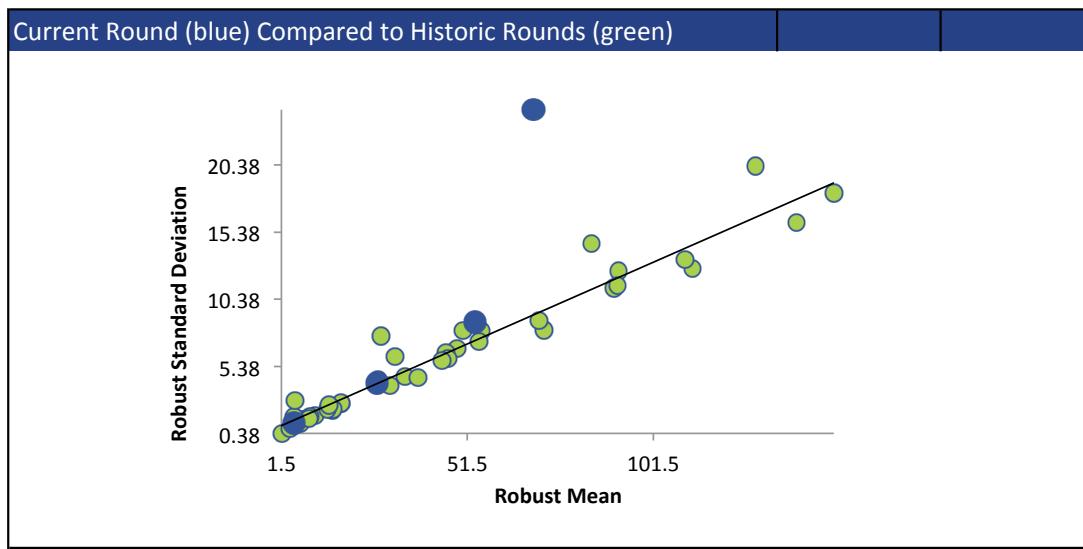
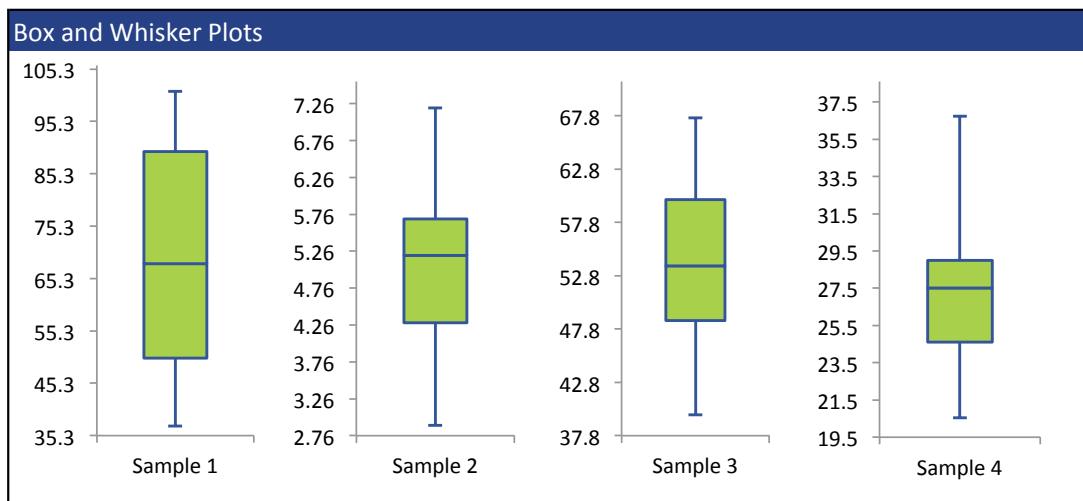
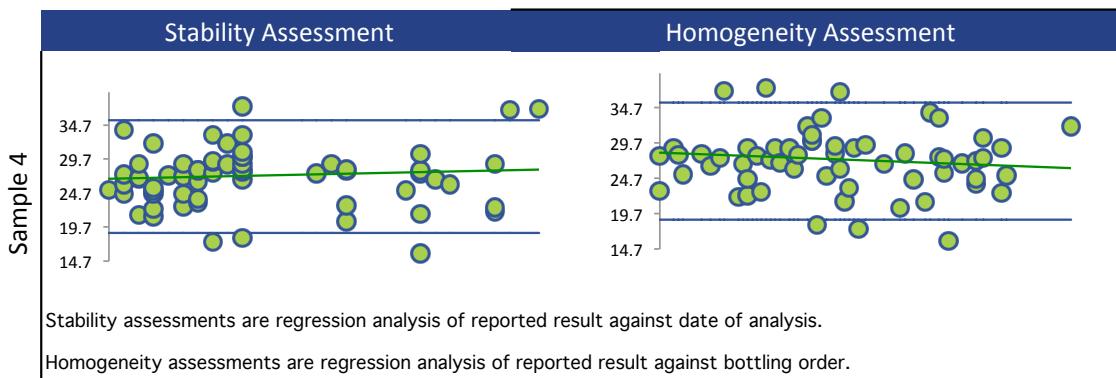
## Annex A Summary by Analyte

### M,P-XYLENE



## Annex A Summary by Analyte

### M,P-XYLENE



## Annex A Summary by Analyte

### METHYL ETHYL KETONE

#### Summary Statistics

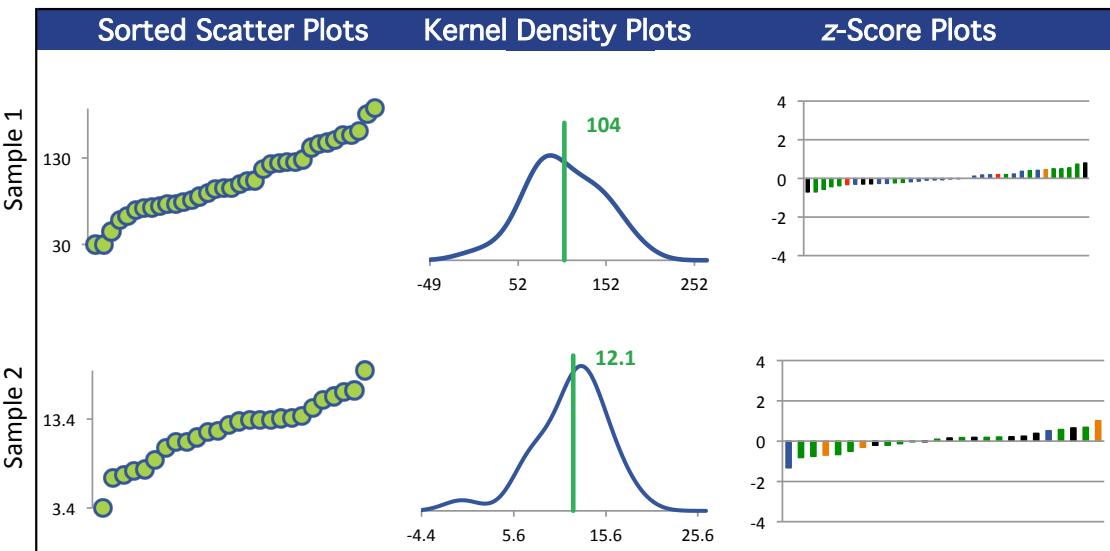
#### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	36	26	36	36
Median $\mu\text{g/L}$	97.9	12.9	84.6	43.0
Robust Mean $\mu\text{g/L}$	104	12.1	82.7	43.0
$U \mu\text{g/L}$	8.90	0.890	3.75	2.17
Robust Standard Deviation $\mu\text{g/L}$	42.7	3.63	18.0	10.4
Regression Standard Deviation $\mu\text{g/L}$	23.4	2.72	18.6	9.68
Stability Flag				
Homogeneity Flag	Homogeneity	Homogeneity		
Standard Deviation Used ( $SDPA \mu\text{g/L}$ )	104	6.63	18.6	10.4
Outliers	0	0	0	0
$ z  > 3.0$	0	0	1	0
$2 <  z  < 3$	0	0	0	0

#### Methods Used

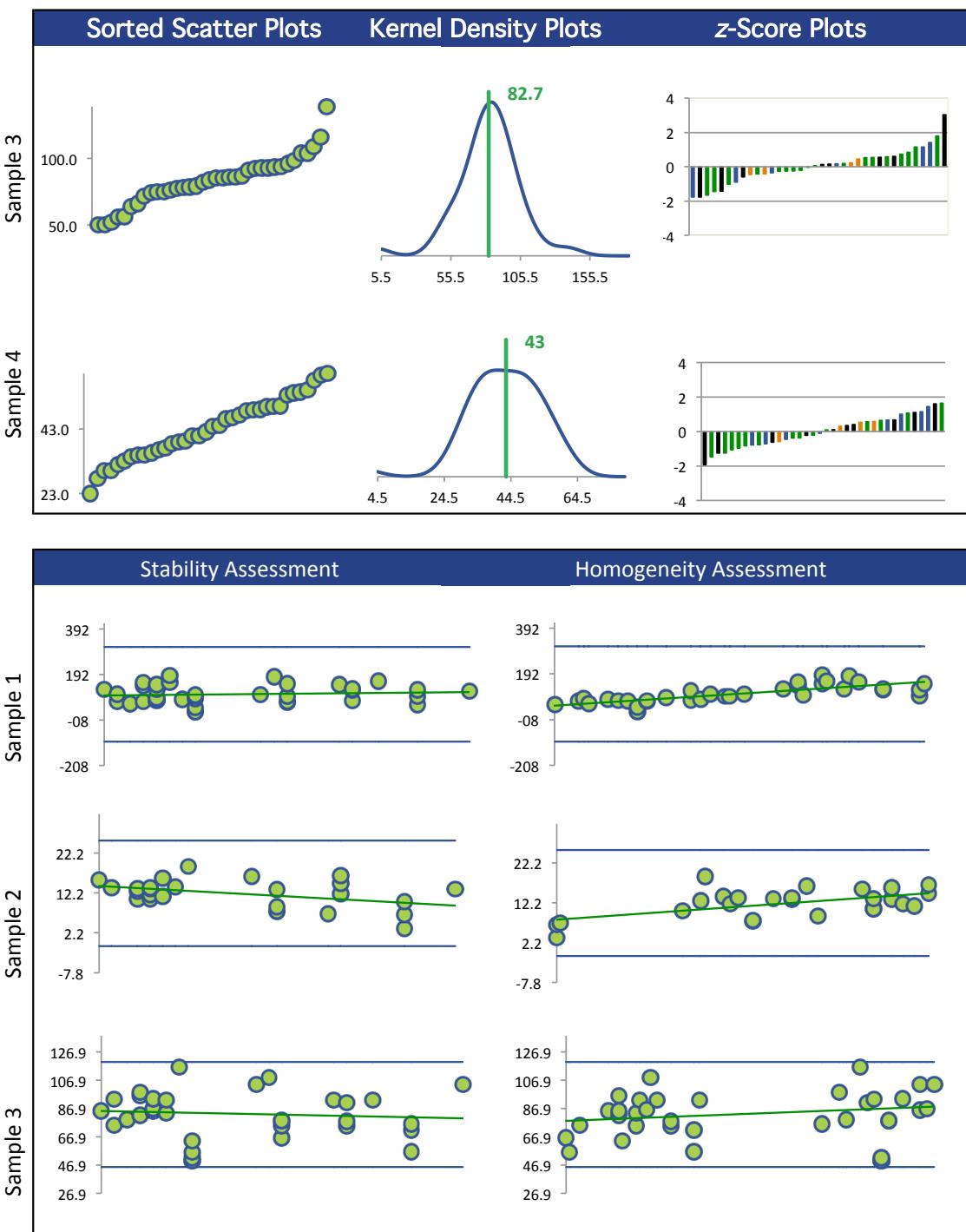
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - HEADSPACE (Blue)	14	7	14	14
GC/MS - PURGE AND TRAP (Red)	20	17	20	20
GC/FID - PURGE AND TRAP (Green)	1	1	1	1
GC/MS (Orange)	1	1	1	1

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



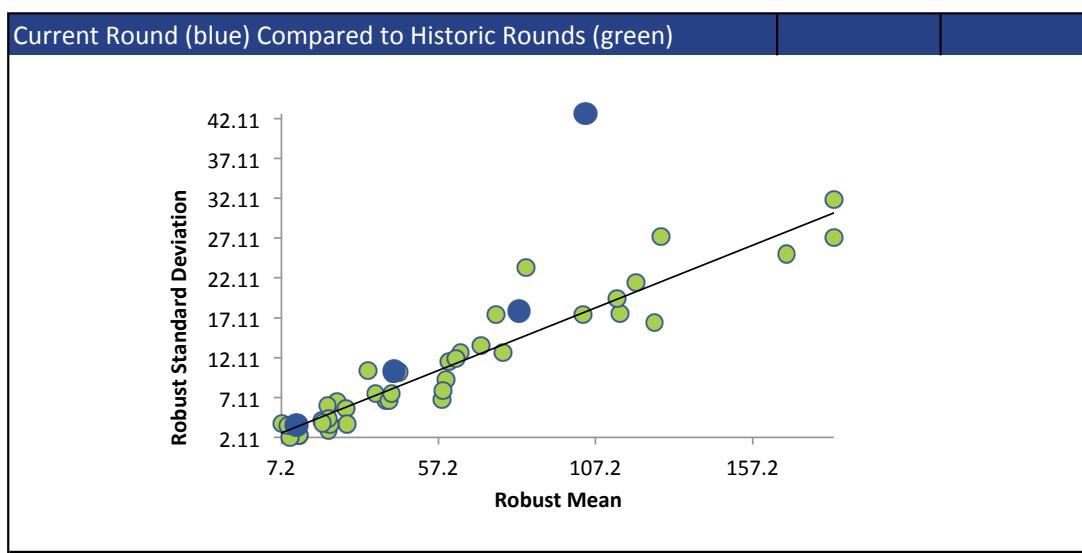
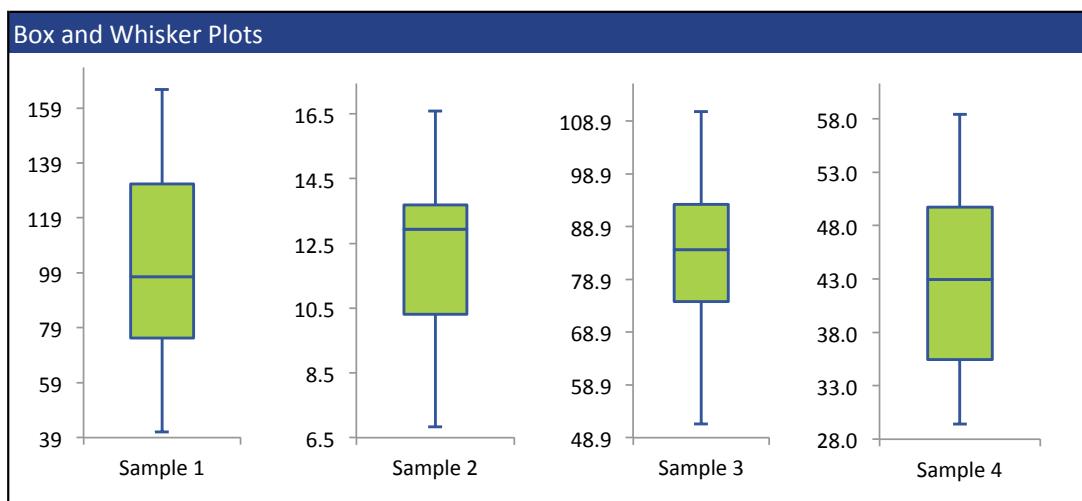
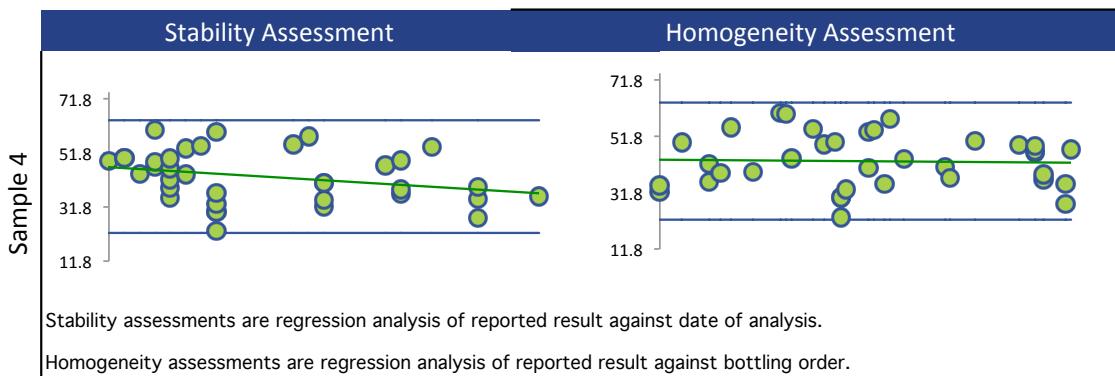
Annex A Summary by Analyte

METHYL ETHYL KETONE



## Annex A Summary by Analyte

### METHYL ETHYL KETONE



Annex A Summary by Analyte

### METHYL ISOBUTYL KETONE (MIBK)

#### Summary Statistics

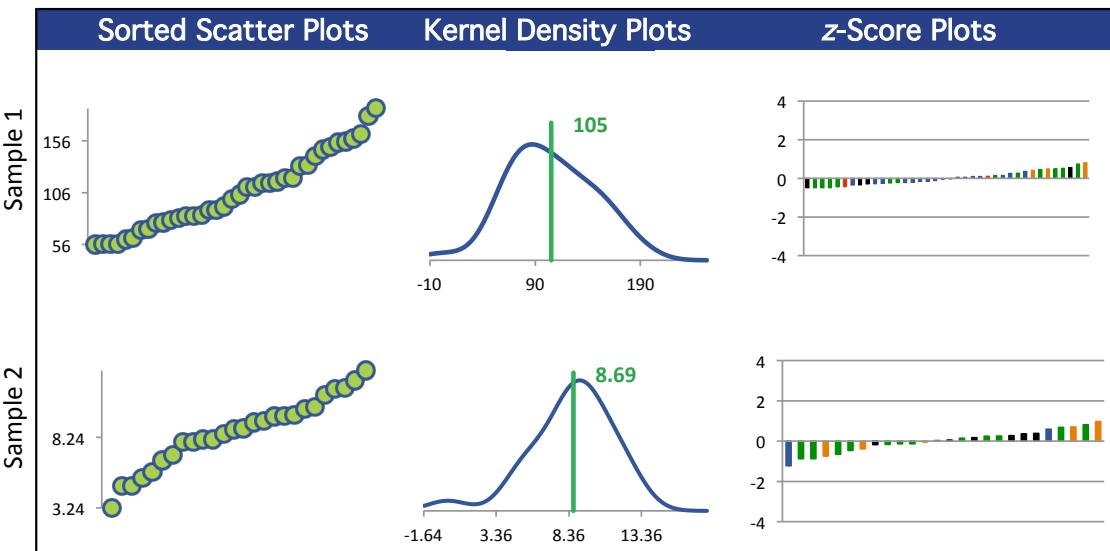
#### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	38	26	38	38
Median $\mu\text{g/L}$	102	8.85	77.7	39.1
Robust Mean $\mu\text{g/L}$	105	8.69	78.5	40.1
$U \mu\text{g/L}$	8.15	0.657	3.41	1.73
Robust Standard Deviation $\mu\text{g/L}$	40.2	2.68	16.8	8.55
Regression Standard Deviation $\mu\text{g/L}$	15.8	1.30	11.8	6.01
Stability Flag				
Homogeneity Flag	Homogeneity	Homogeneity		
Standard Deviation Used ( $SDPA$ ) $\mu\text{g/L}$	101	4.45	16.8	8.55
Outliers	0	0	0	0
$ z  > 3.0$	0	0	2	0
$2 <  z  < 3$	0	0	0	2

#### Methods Used

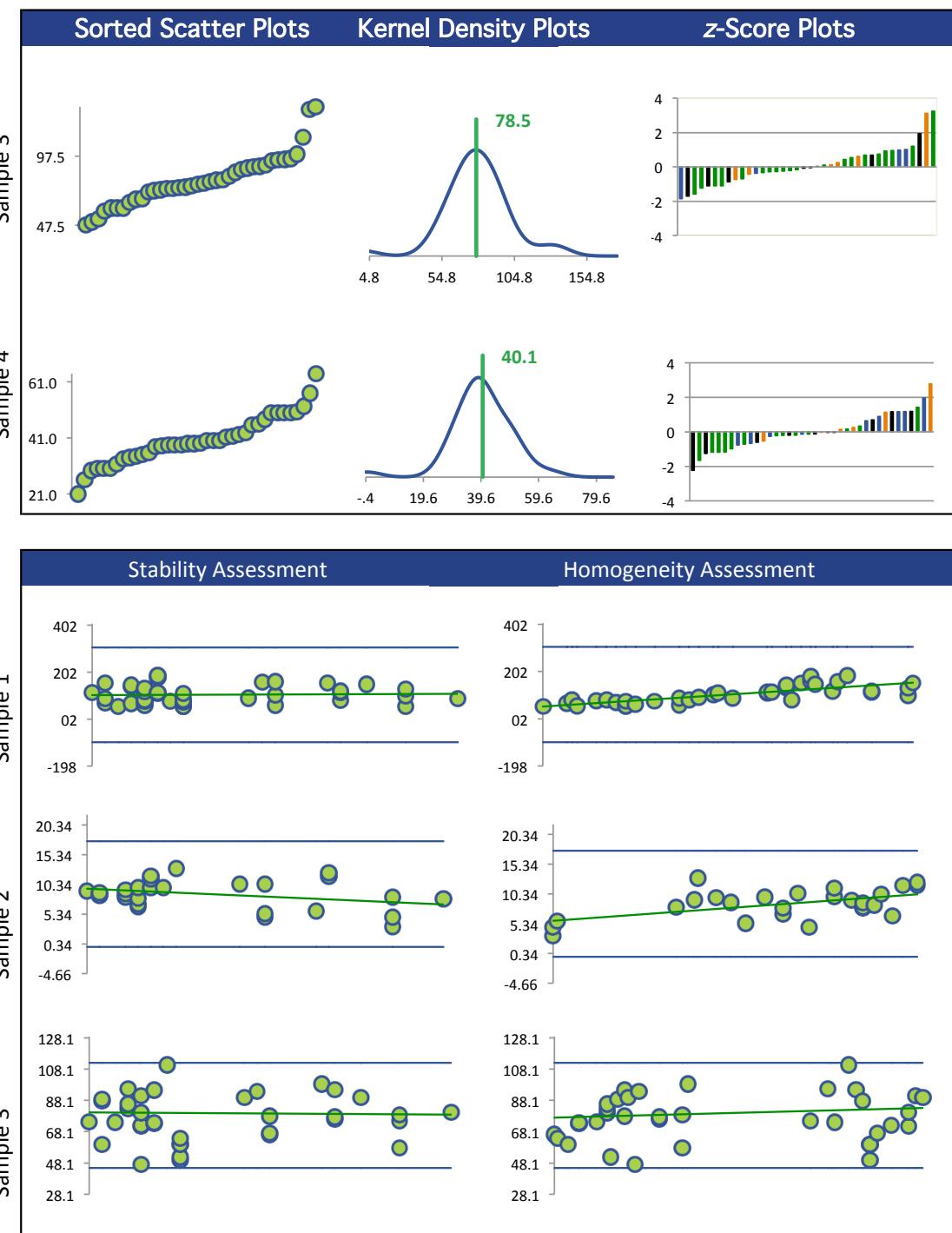
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	22	17	22	22
GC/MS - HEADSPACE (Red)	13	7	13	13
GC/FID (Green)	1	0	1	1
GC/MS (Orange)	2	2	2	2

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



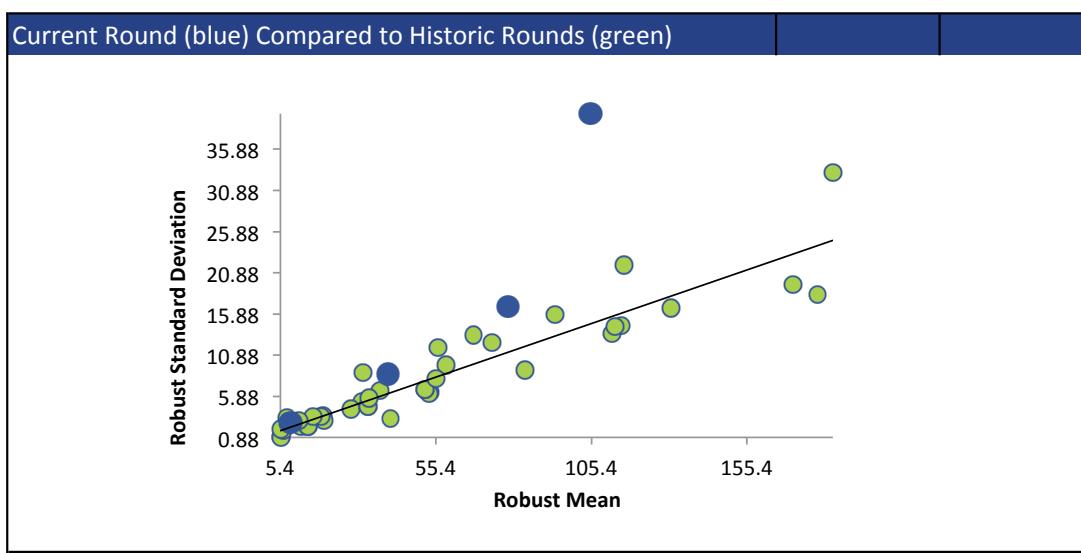
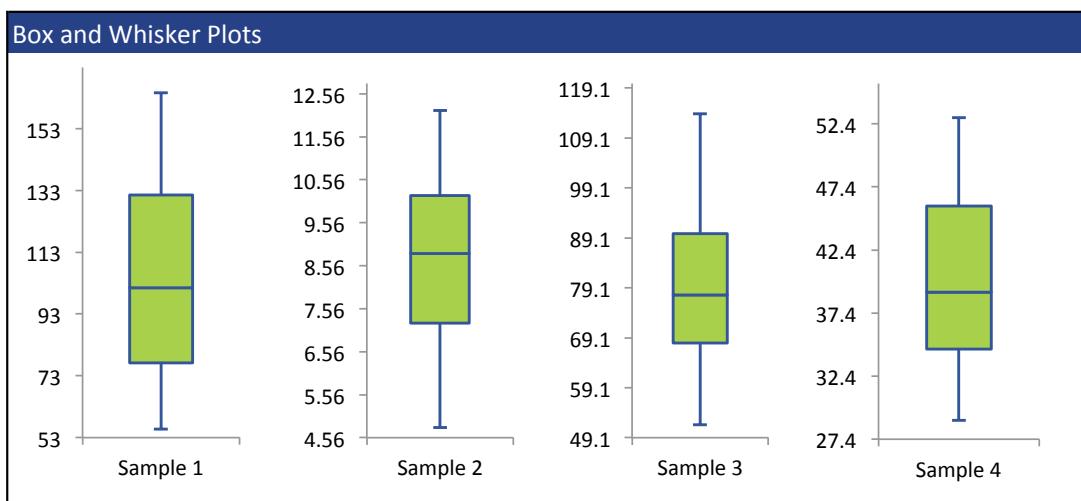
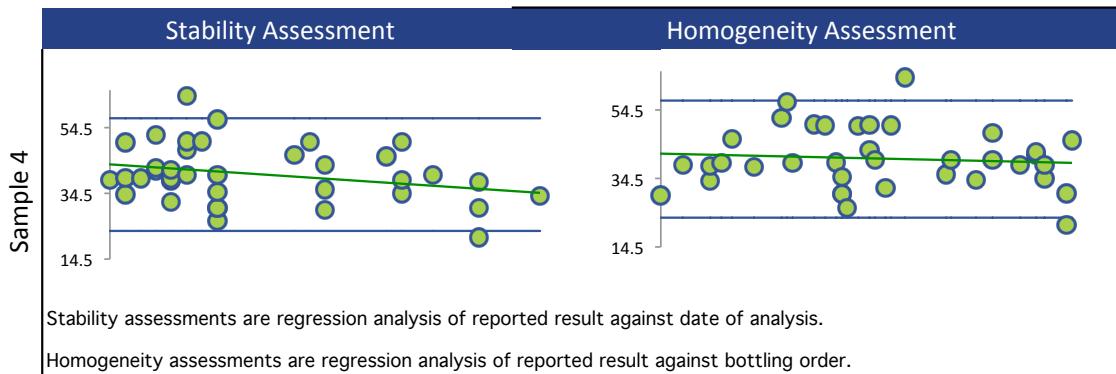
Annex A Summary by Analyte

METHYL ISOBUTYL KETONE (MIBK)



## Annex A Summary by Analyte

### METHYL ISOBUTYL KETONE (MIBK)



Annex A Summary by Analyte

## METHYL T-BUTYL ETHER

### Summary Statistics

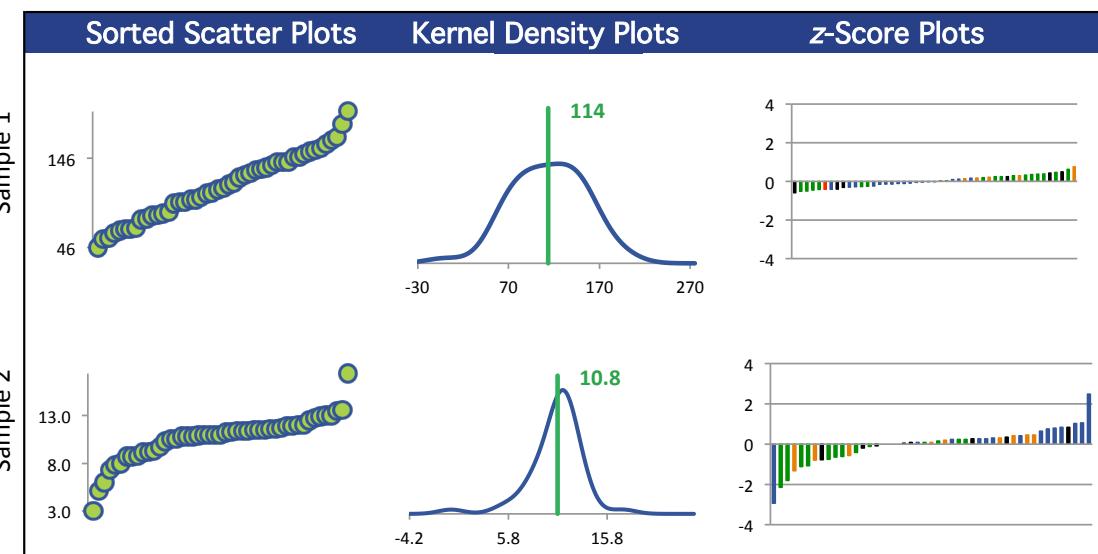
### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	47	47	47	47
Median $\mu\text{g/L}$	113	11.0	82.0	42.7
Robust Mean $\mu\text{g/L}$	114	10.8	81.6	43.1
$U \mu\text{g/L}$	7.40	0.334	2.28	1.23
Robust Standard Deviation $\mu\text{g/L}$	40.6	1.83	12.5	6.73
Regression Standard Deviation $\mu\text{g/L}$	22.9	2.17	16.3	8.61
Stability Flag				
Homogeneity Flag	Homogeneity	Homogeneity		
Standard Deviation Used ( $SDPA \mu\text{g/L}$ )	112	2.66	16.3	8.61
Outliers	0	0	0	0
$ z  > 3.0$	0	0	0	0
$2 <  z  < 3$	0	3	2	1

### Methods Used

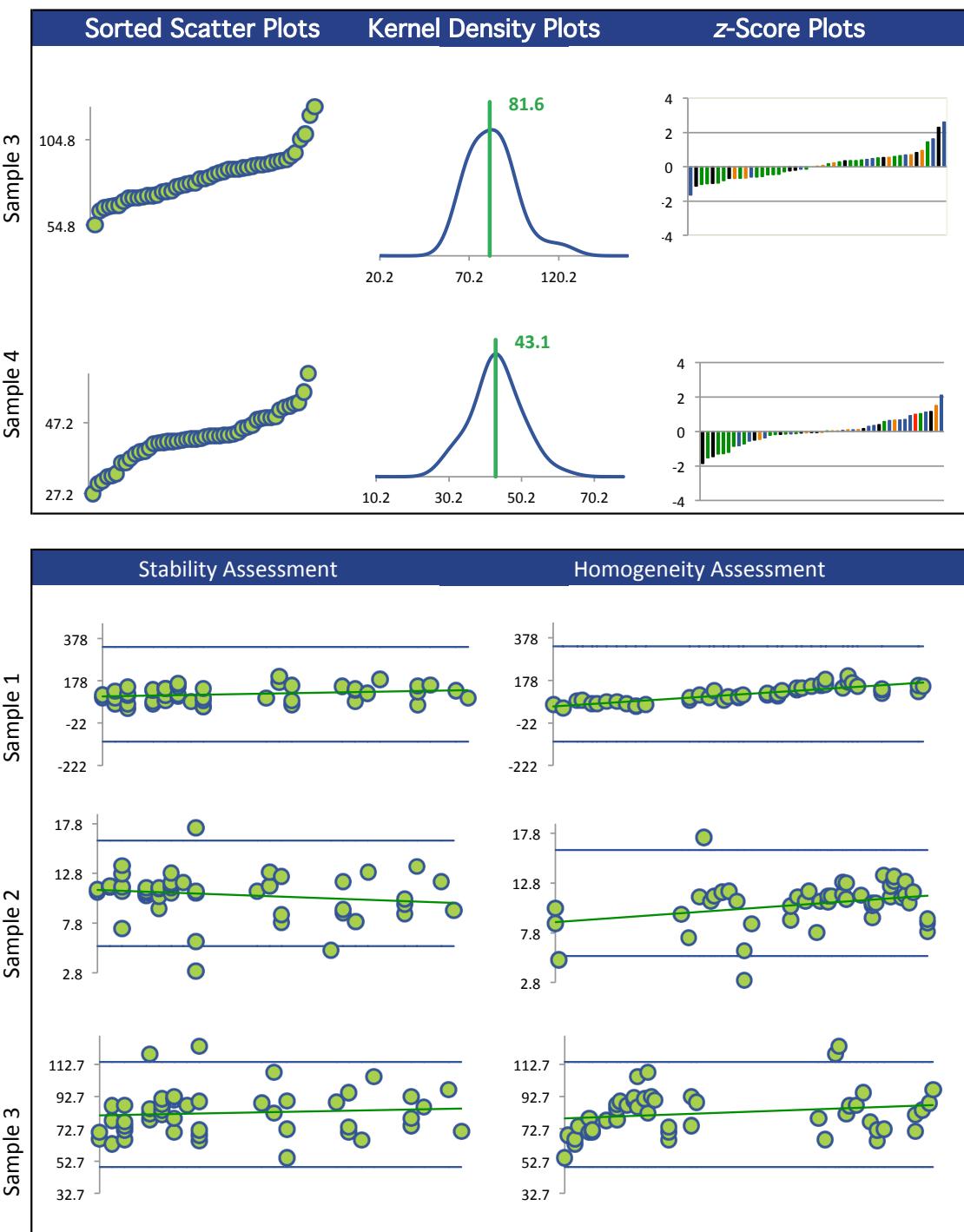
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	24	24	24	24
GC/MS - HEADSPACE (Red)	20	20	20	20
GC/MS (Green)	3	3	3	3

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



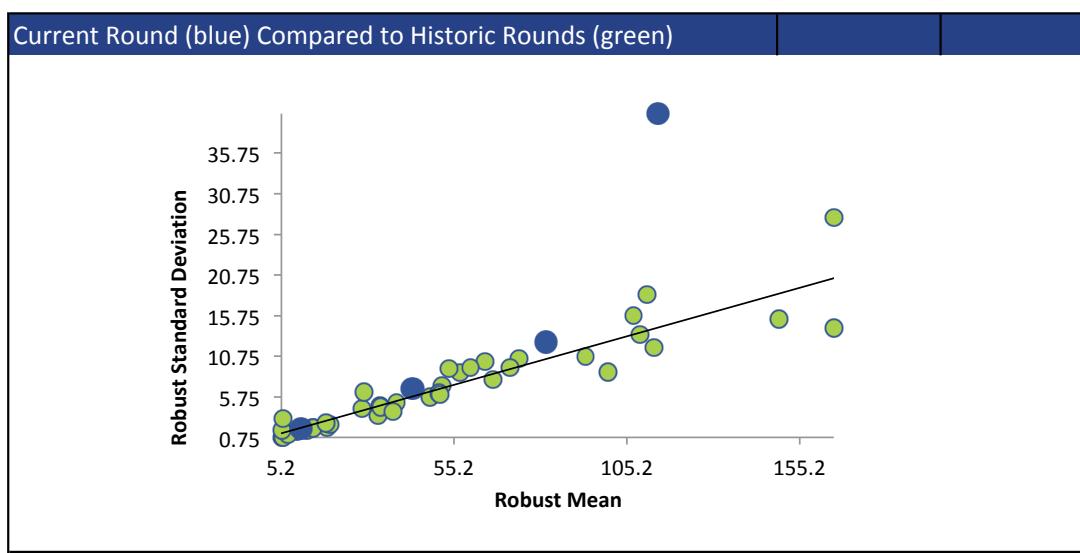
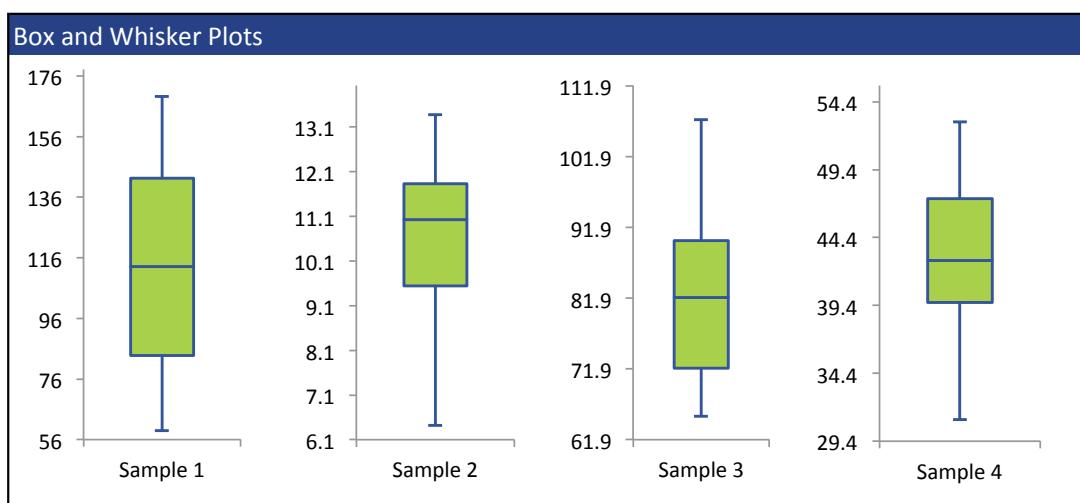
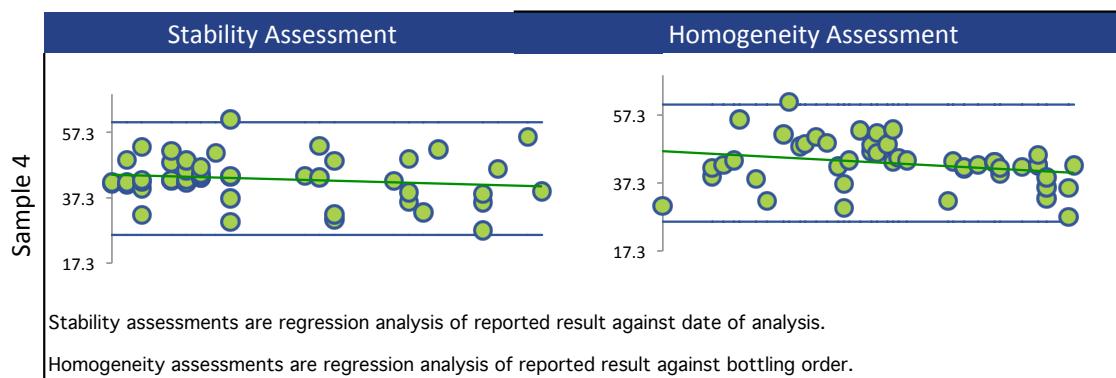
Annex A Summary by Analyte

METHYL T-BUTYL ETHER



## Annex A Summary by Analyte

## METHYL T-BUTYL ETHER



## Annex A Summary by Analyte

### O-XYLENE

#### Summary Statistics

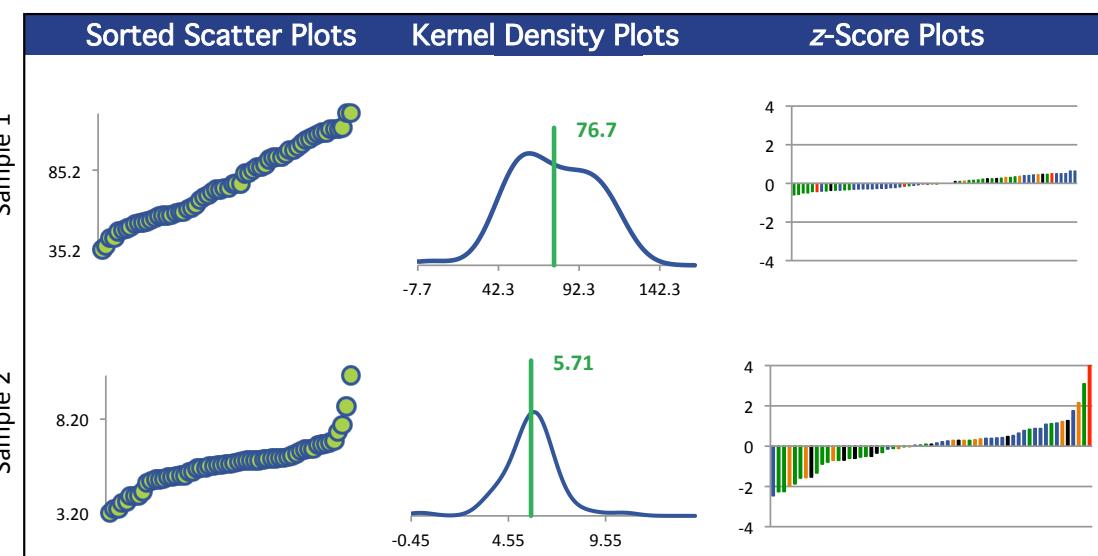
#### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	62	59	62	62
Median $\mu\text{g/L}$	74.0	5.80	59.8	30.3
Robust Mean $\mu\text{g/L}$	76.7	5.71	59.3	30.2
$U \mu\text{g/L}$	4.21	0.166	1.47	0.722
Robust Standard Deviation $\mu\text{g/L}$	26.5	1.02	9.25	4.55
Regression Standard Deviation $\mu\text{g/L}$	11.5	0.857	8.90	4.53
Stability Flag				
Homogeneity Flag	Homogeneity			
Standard Deviation Used ( $SDPA \mu\text{g/L}$ )	69.0	1.02	9.25	4.55
Outliers	0	0	0	0
$ z  > 3.0$	0	2	0	1
$2 <  z  < 3$	0	4	2	3

#### Methods Used

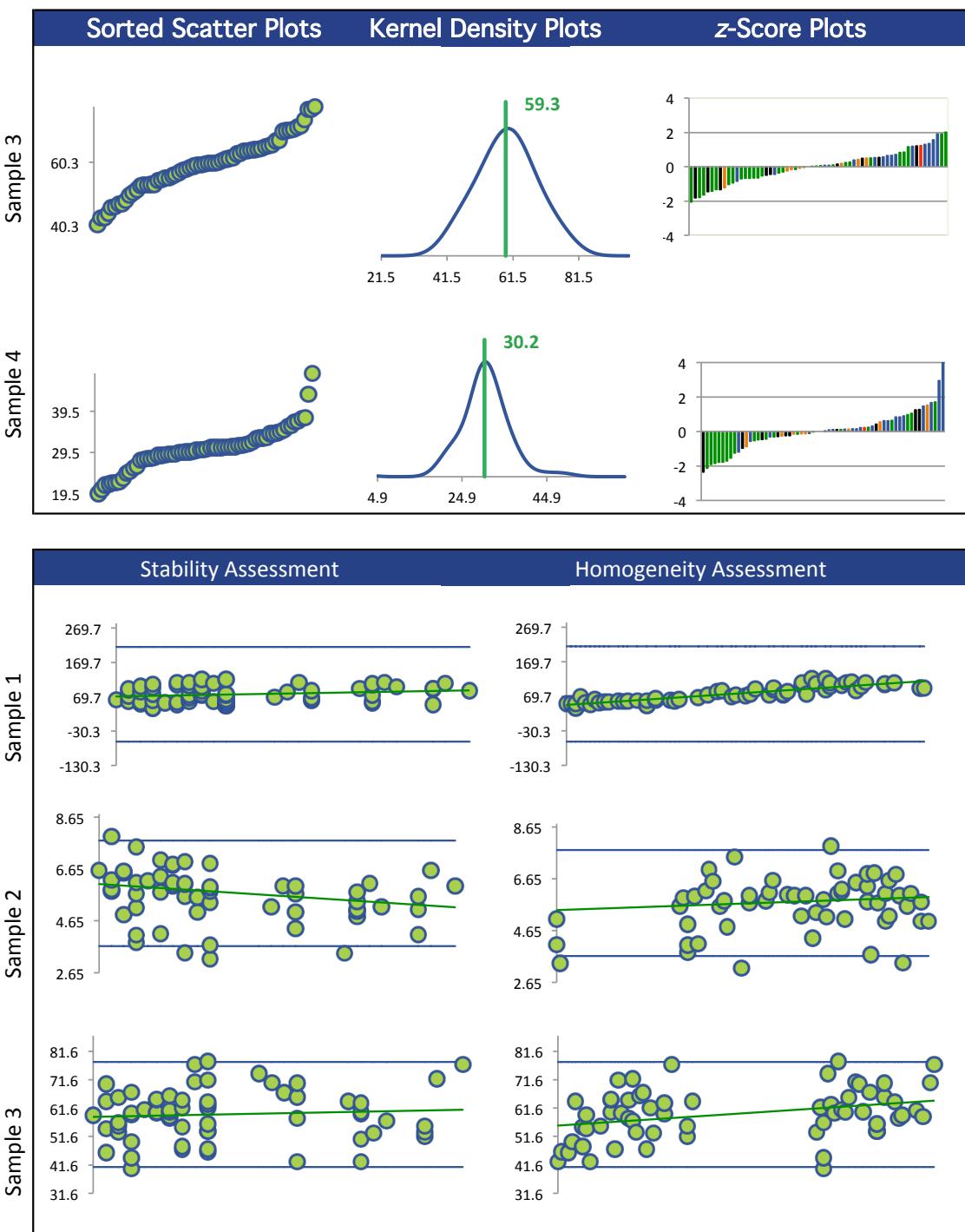
Method	C16-1	C16-2	C16-3	C16-4
GC/MS (Blue)	6	5	6	6
GC/MS - PURGE AND TRAP (Red)	31	31	31	31
GC/MS - HEADSPACE (Green)	22	21	22	22
GC/FID - HEADSPACE (Orange)	1	0	1	1
GC/MS/MS - HEADSPACE (Black)	1	1	1	1
GC/FID - PURGE AND TRAP (Yellow)	1	1	1	1

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



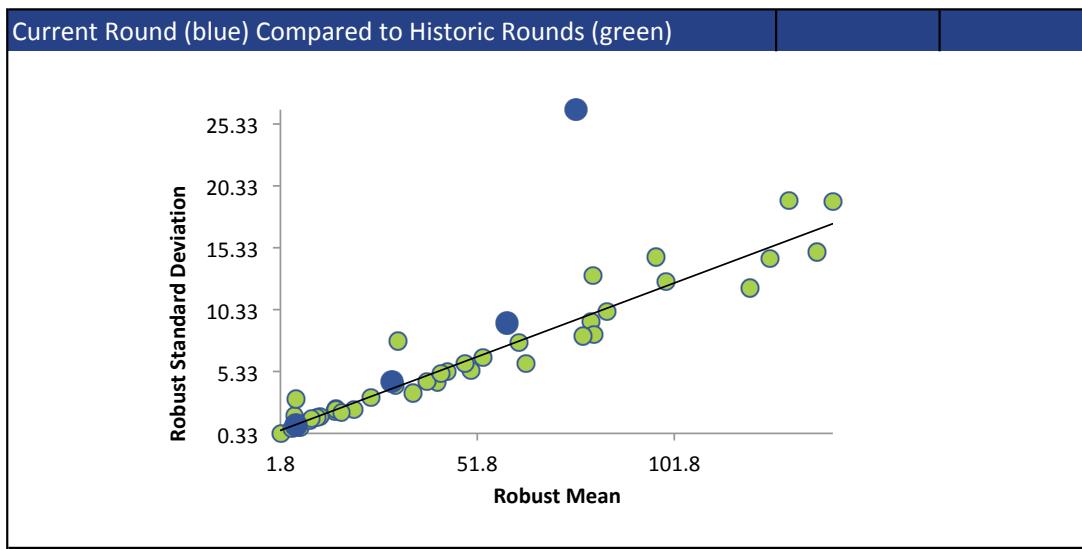
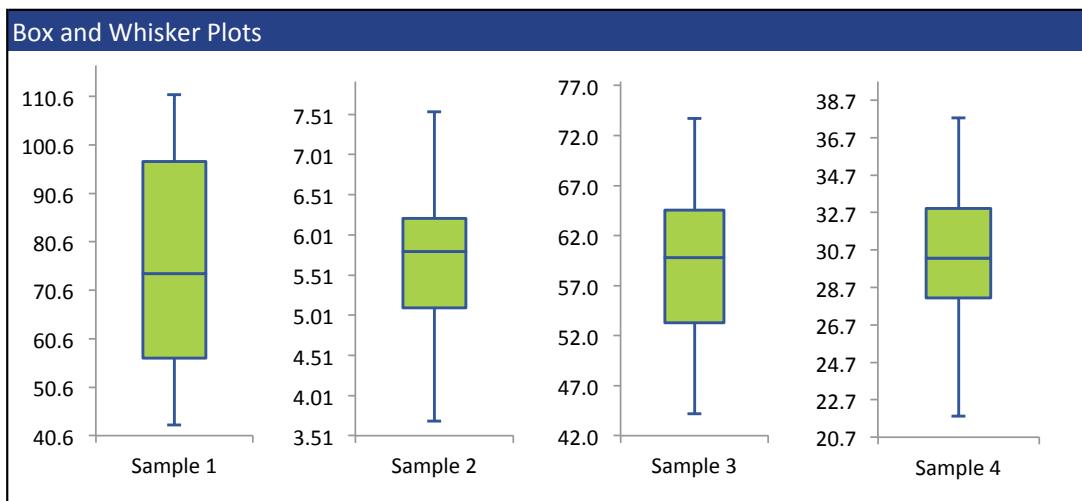
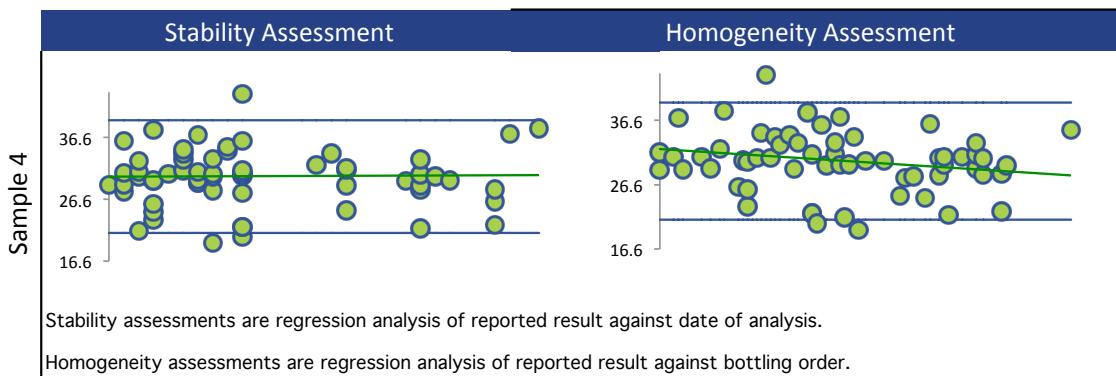
## Annex A Summary by Analyte

### O-XYLENE



## Annex A Summary by Analyte

### O-XYLENE



## Annex A Summary by Analyte

### STYRENE

#### Summary Statistics

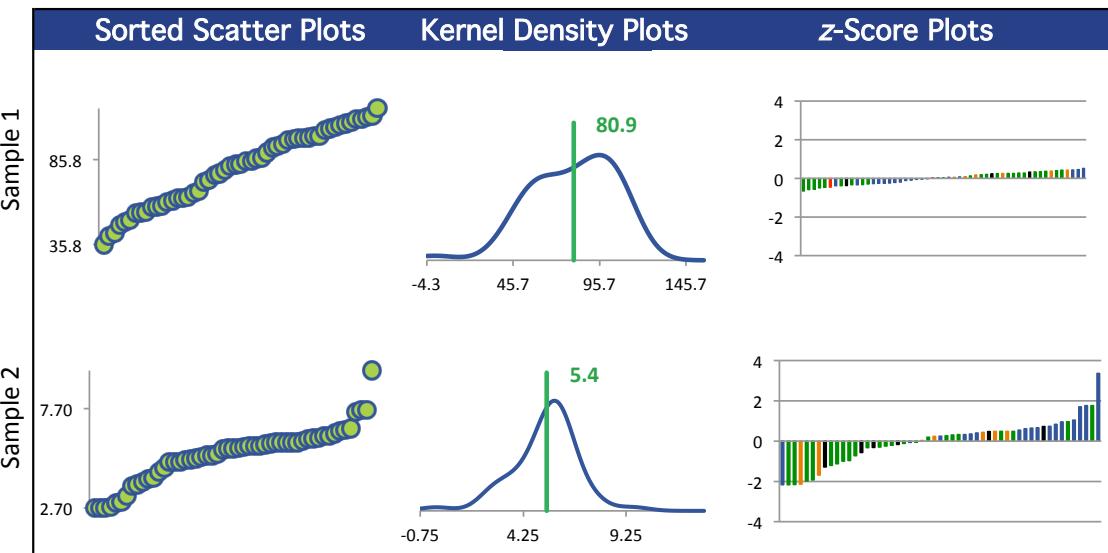
#### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	53	53	53	53
Median $\mu\text{g/L}$	83.1	5.70	58.1	29.8
Robust Mean $\mu\text{g/L}$	80.9	5.40	58.5	29.5
$U \mu\text{g/L}$	4.22	0.215	1.56	0.735
Robust Standard Deviation $\mu\text{g/L}$	24.6	1.25	9.08	4.28
Regression Standard Deviation $\mu\text{g/L}$	12.1	0.810	8.77	4.42
Stability Flag				
Homogeneity Flag	Homogeneity			
Standard Deviation Used ( $SDPA \mu\text{g/L}$ )	67.9	1.25	9.08	4.42
Outliers	0	0	0	0
$ z  > 3.0$	0	1	0	1
$2 <  z  < 3$	0	4	1	2

#### Methods Used

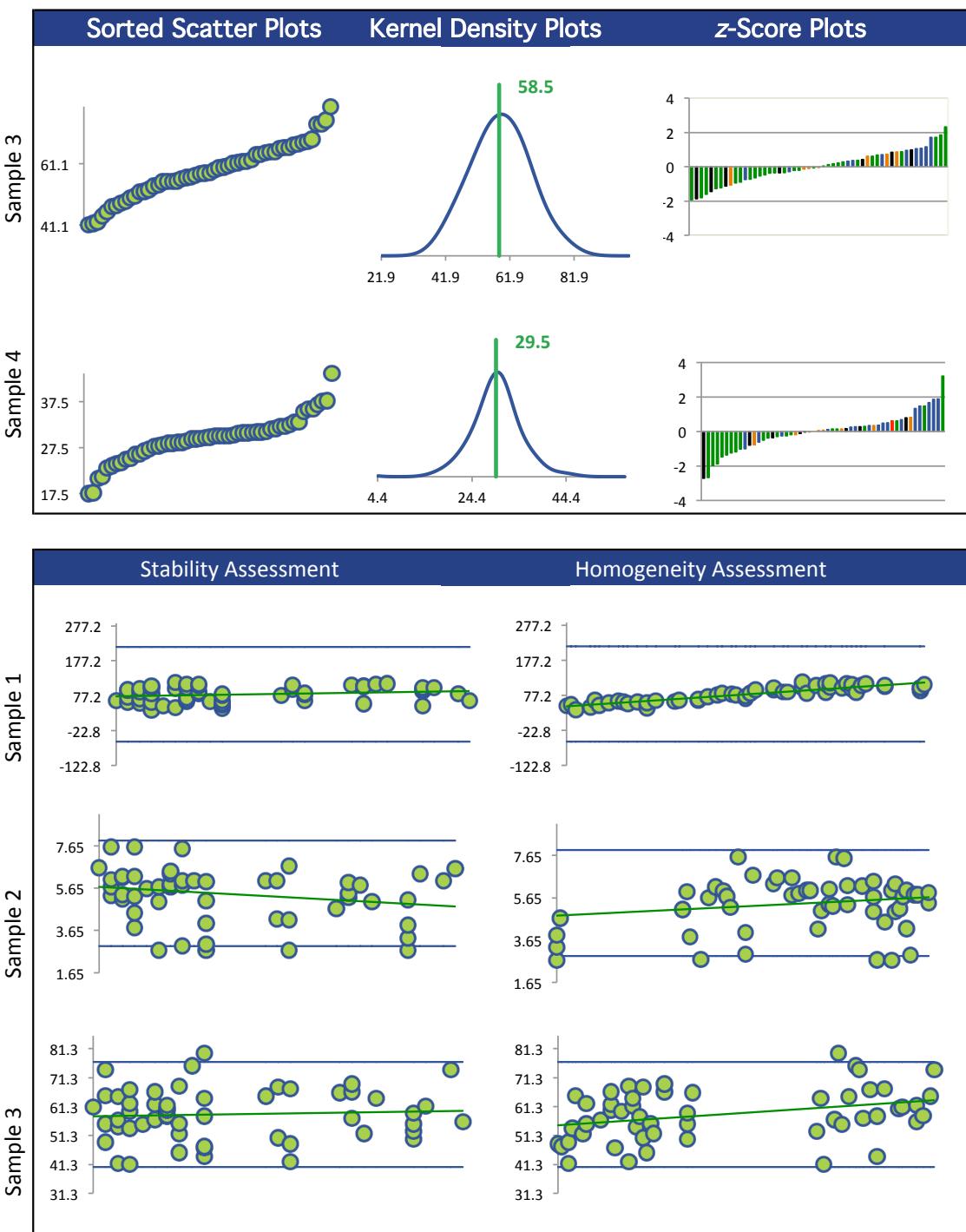
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	25	25	25	25
GC/MS - HEADSPACE (Red)	22	22	22	22
GC/MS (Green)	5	5	5	5
GC/PID (Orange)	1	1	1	1

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



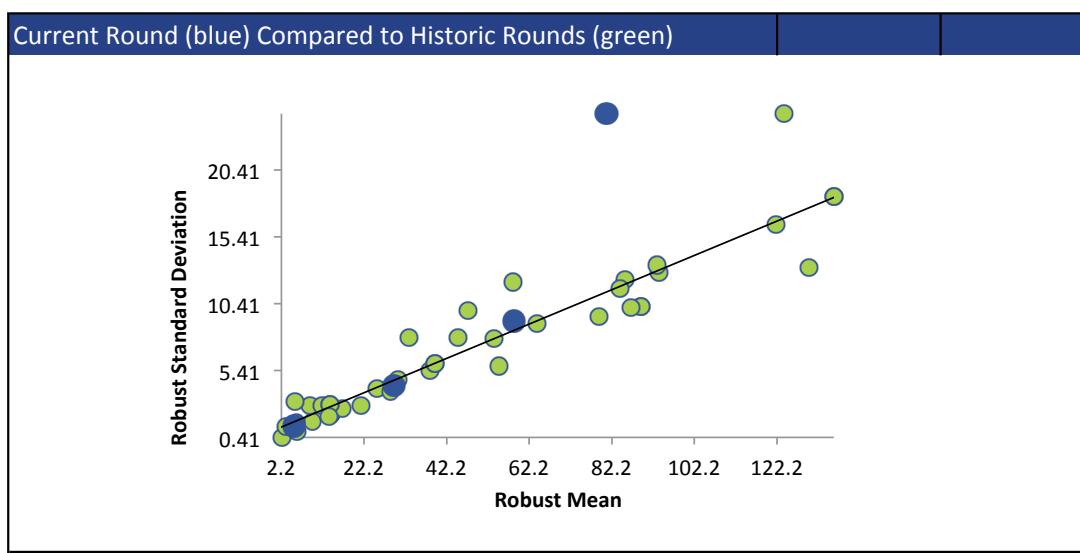
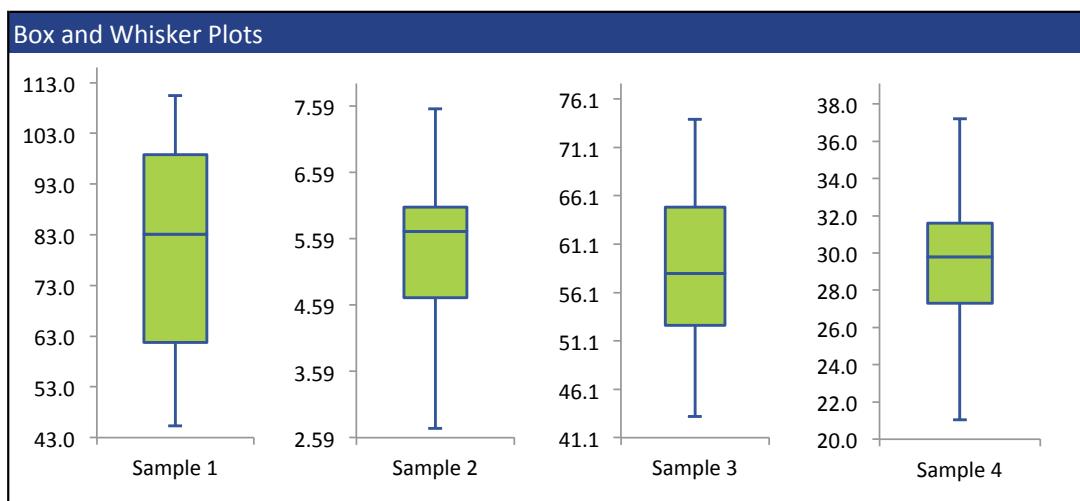
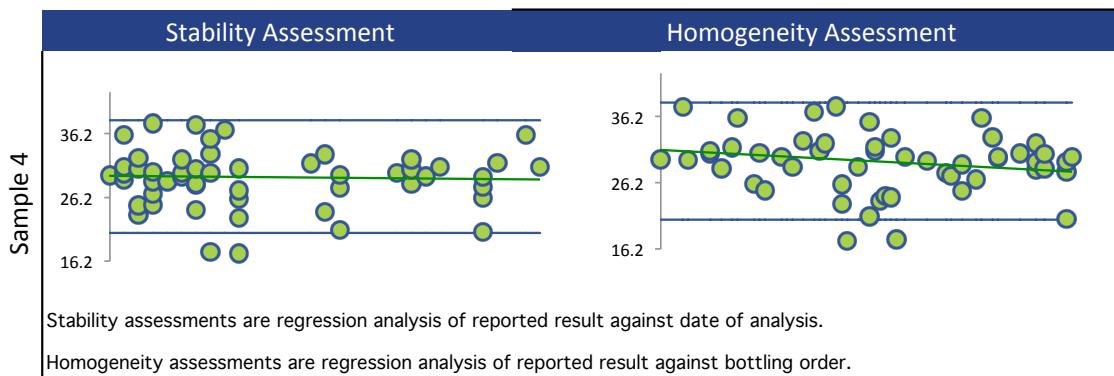
Annex A Summary by Analyte

STYRENE



## Annex A Summary by Analyte

### STYRENE



Annex A Summary by Analyte

## TETRACHLOROETHYLENE

### Summary Statistics

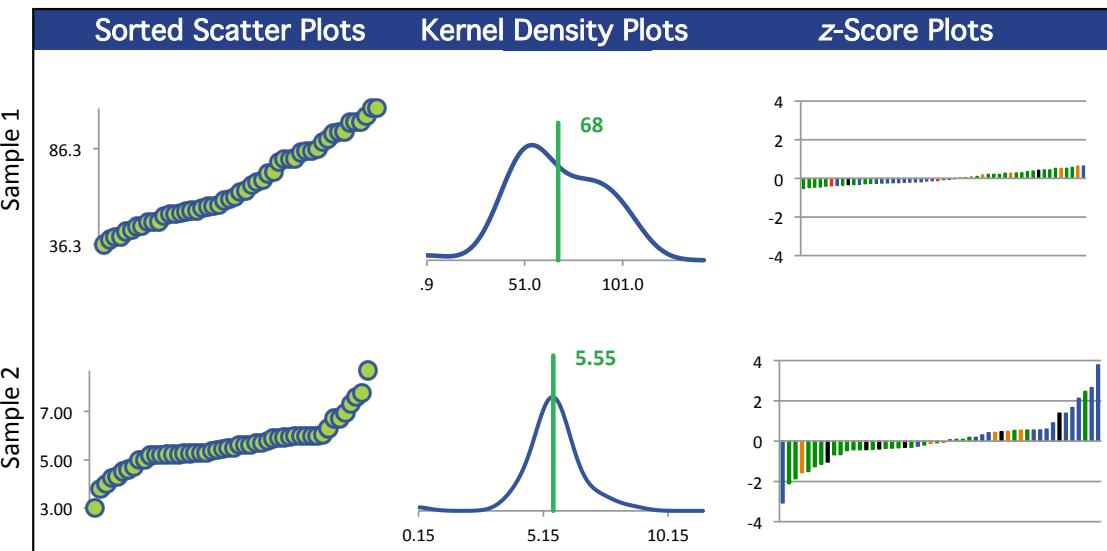
### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	51	50	51	51
Median $\mu\text{g/L}$	63.6	5.50	52.0	26.5
Robust Mean $\mu\text{g/L}$	68.0	5.55	53.1	26.9
$U \mu\text{g/L}$	4.04	0.139	1.51	0.586
Robust Standard Deviation $\mu\text{g/L}$	23.1	0.784	8.63	3.35
Regression Standard Deviation $\mu\text{g/L}$	10.2	0.833	7.96	4.03
Stability Flag				
Homogeneity Flag	Homogeneity			
Standard Deviation Used ( $SDPA \mu\text{g/L}$ )	61.0	0.833	8.63	4.03
Outliers	0	0	0	0
$ z  > 3.0$	0	2	0	2
$2 <  z  < 3$	0	4	0	4

### Methods Used

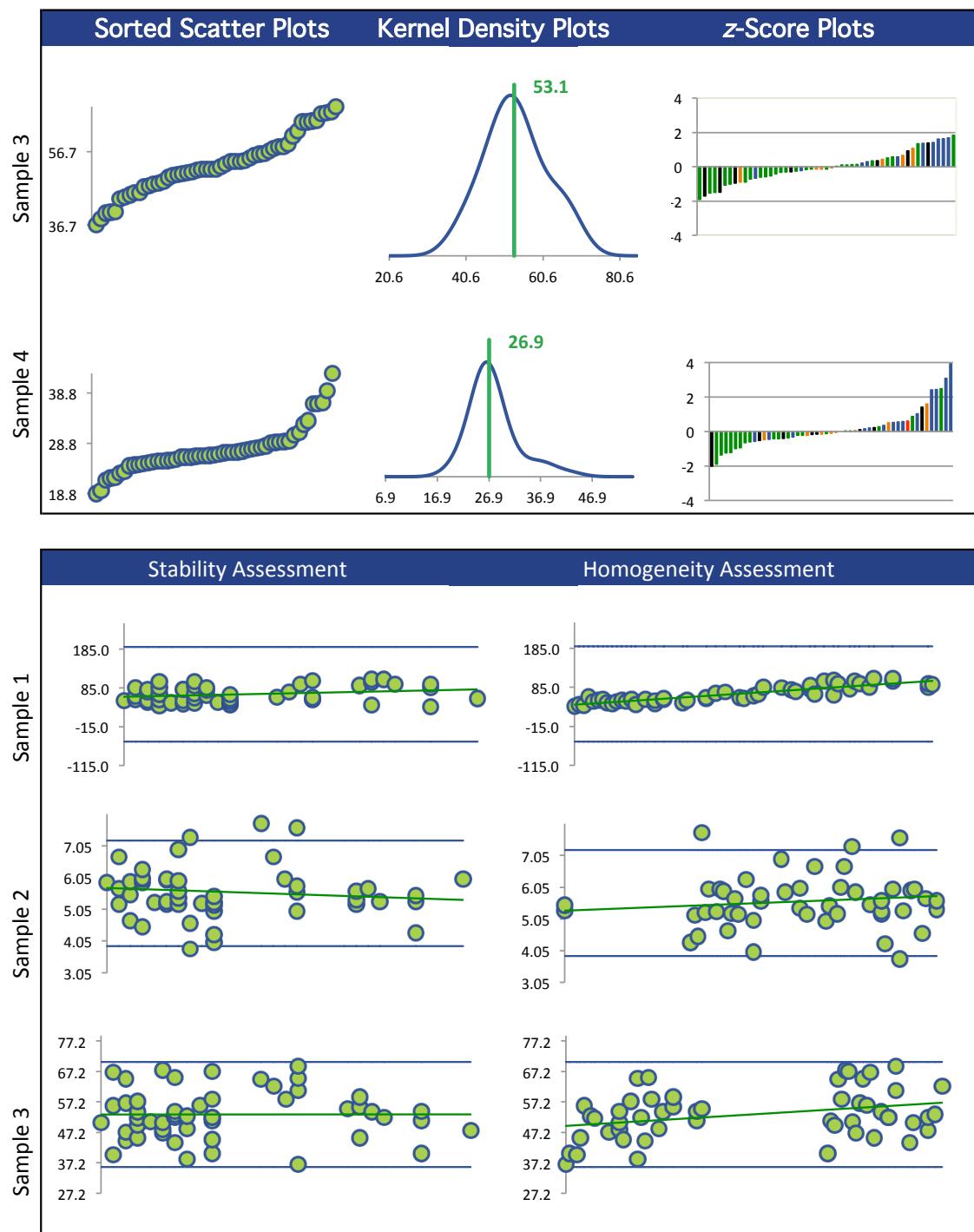
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	28	28	28	28
GC/MS - HEADSPACE (Red)	19	19	19	19
GC/MS/MS - HEADSPACE (Green)	1	1	1	1
GC/FID - PURGE AND TRAP (Orange)	1	1	1	1
GC/MS (Black)	2	1	2	2

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



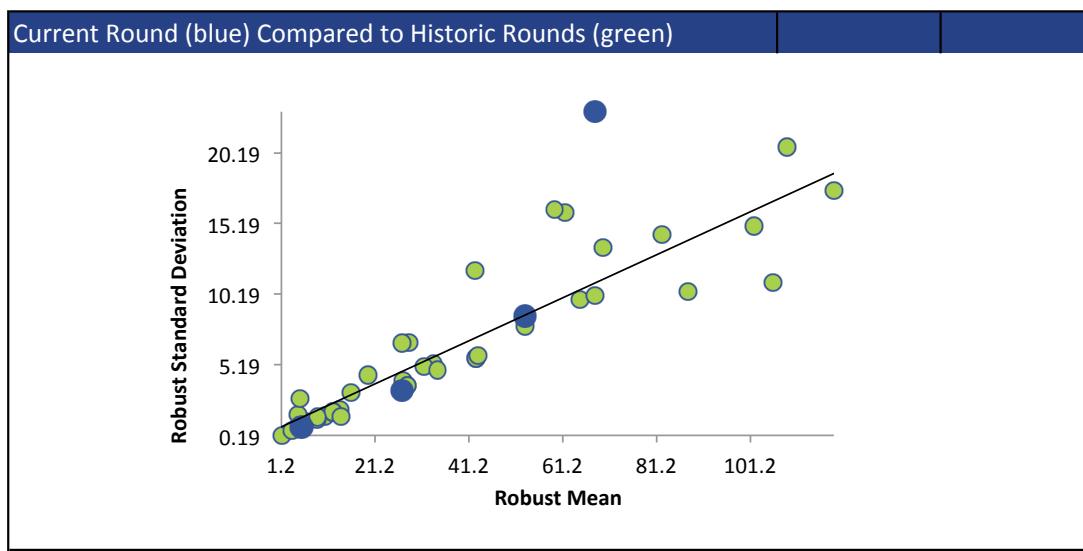
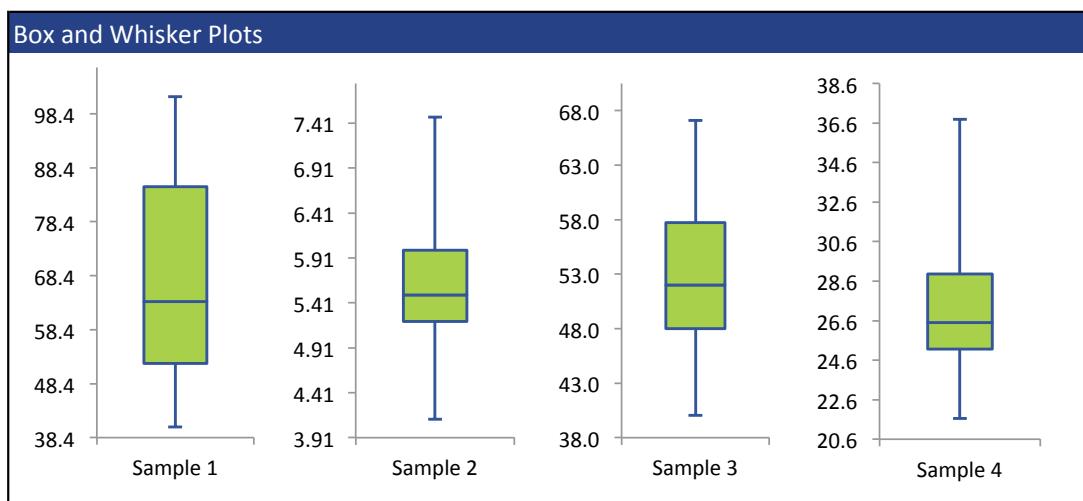
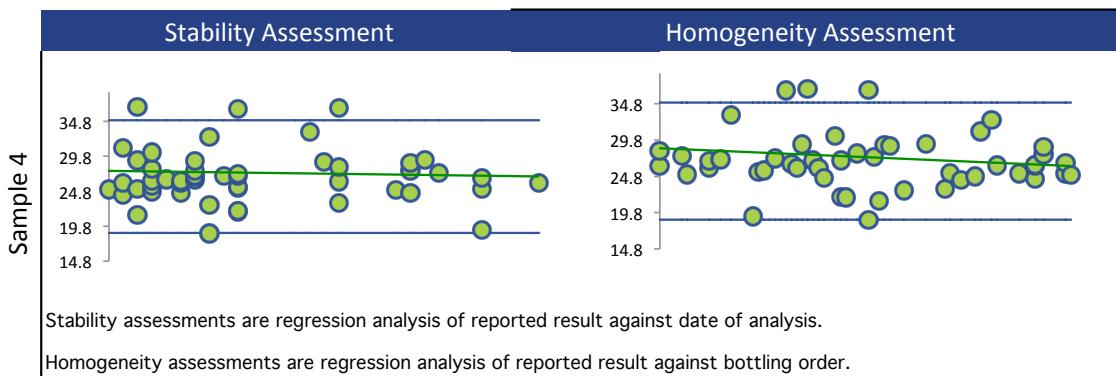
Annex A Summary by Analyte

TETRACHLOROETHYLENE



## Annex A Summary by Analyte

### TETRACHLOROETHYLENE



## Annex A Summary by Analyte

### TOLUENE

#### Summary Statistics

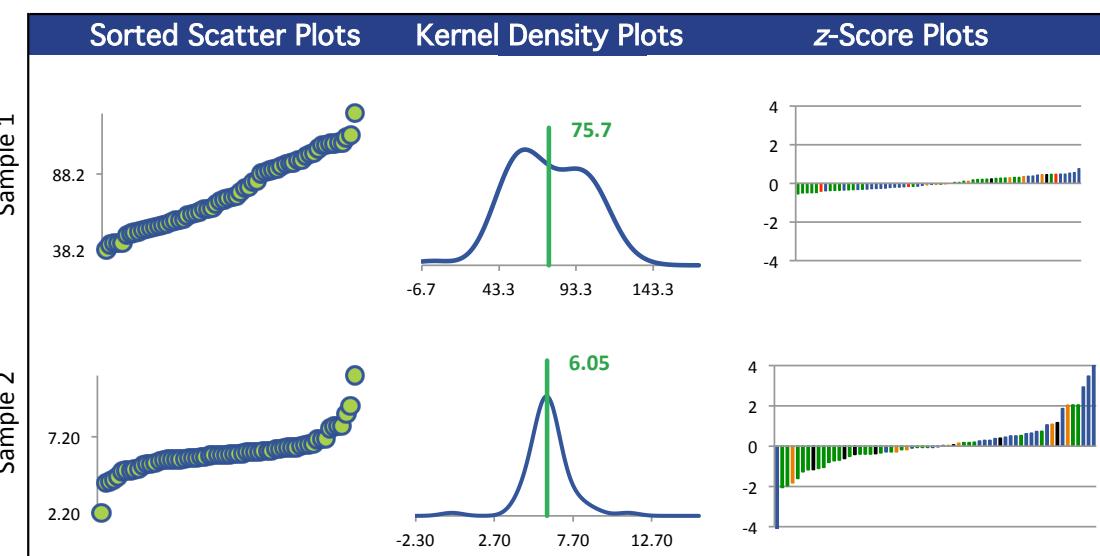
#### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	62	62	62	61
Median $\mu\text{g/L}$	72.2	6.01	58.4	29.7
Robust Mean $\mu\text{g/L}$	75.7	6.05	58.5	29.9
$U \mu\text{g/L}$	4.11	0.132	1.27	0.624
Robust Standard Deviation $\mu\text{g/L}$	25.9	0.829	7.99	3.90
Regression Standard Deviation $\mu\text{g/L}$	11.4	0.908	8.77	4.48
Stability Flag				
Homogeneity Flag	Homogeneity			
Standard Deviation Used (SDPA) $\mu\text{g/L}$	68.8	0.908	8.77	4.48
Outliers	0	0	0	1
$ z  > 3.0$	0	3	0	1
$2 <  z  < 3$	0	5	1	2

#### Methods Used

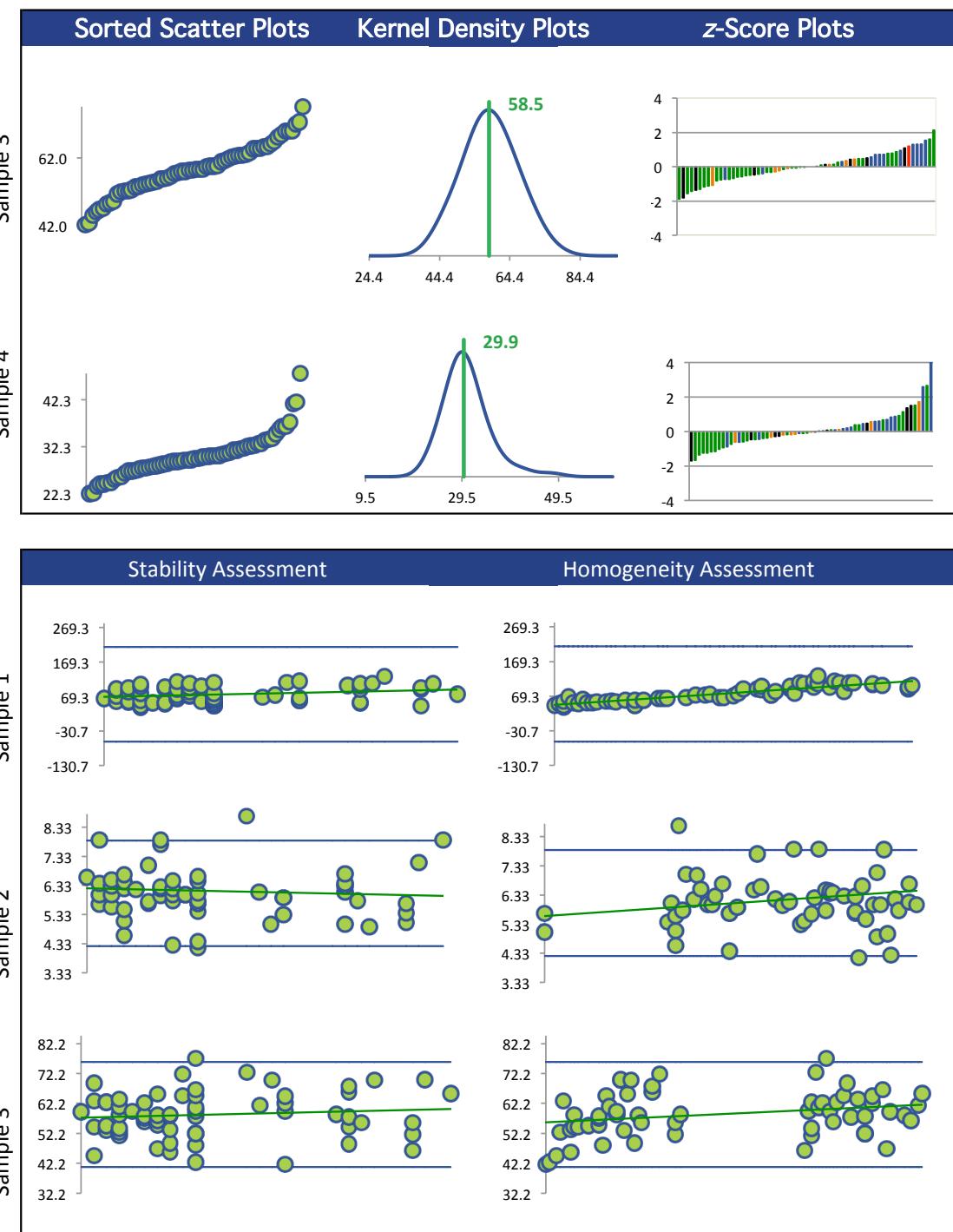
Method	C16-1	C16-2	C16-3	C16-4
GC/MS (Blue)	6	6	6	6
GC/MS - PURGE AND TRAP (Red)	31	31	31	30
GC/MS - HEADSPACE (Green)	22	22	22	22
GC/FID - HEADSPACE (Orange)	1	1	1	1
GC/MS/MS - HEADSPACE (Black)	1	1	1	1
GC/FID - PURGE AND TRAP (Yellow)	1	1	1	1

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



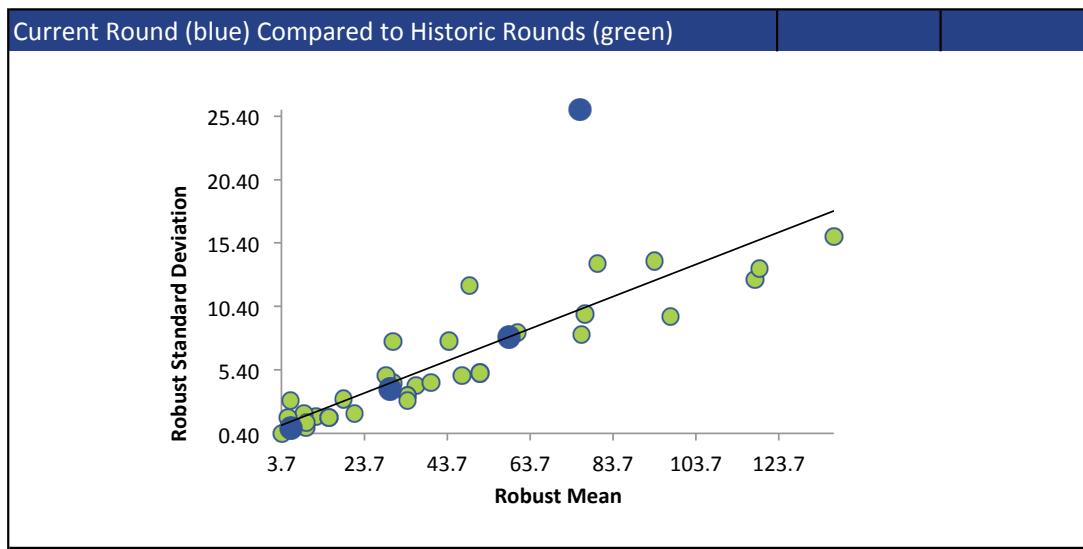
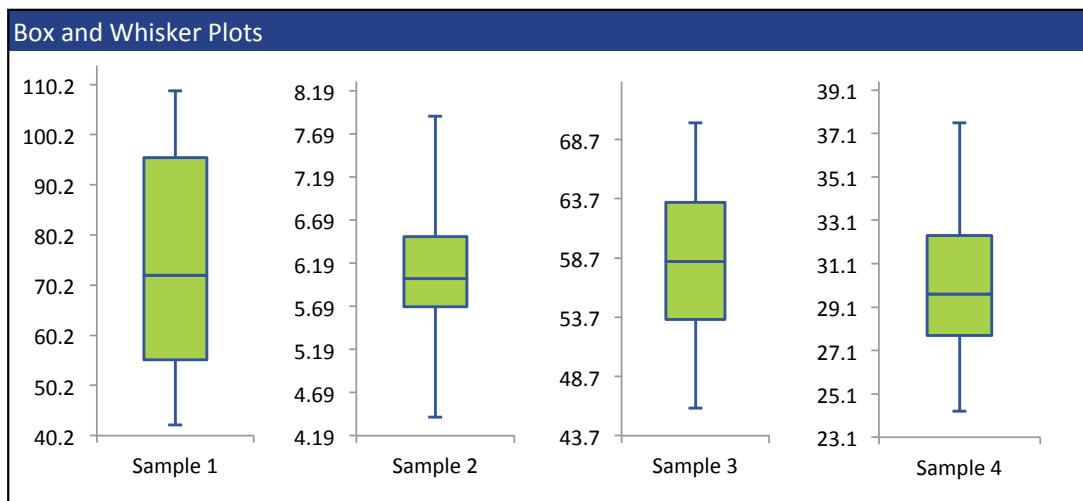
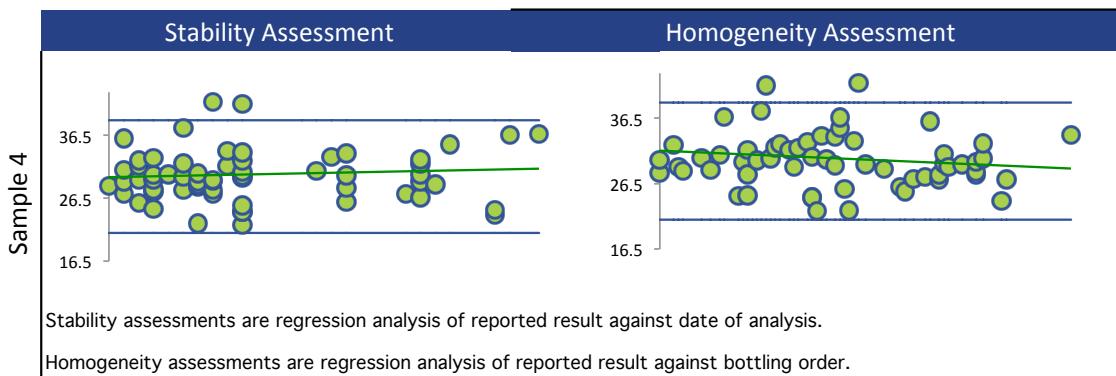
Annex A Summary by Analyte

TOLUENE



## Annex A Summary by Analyte

### TOLUENE



Annex A Summary by Analyte

## TRANS-1,2-DICHLOROETHYLENE

### Summary Statistics

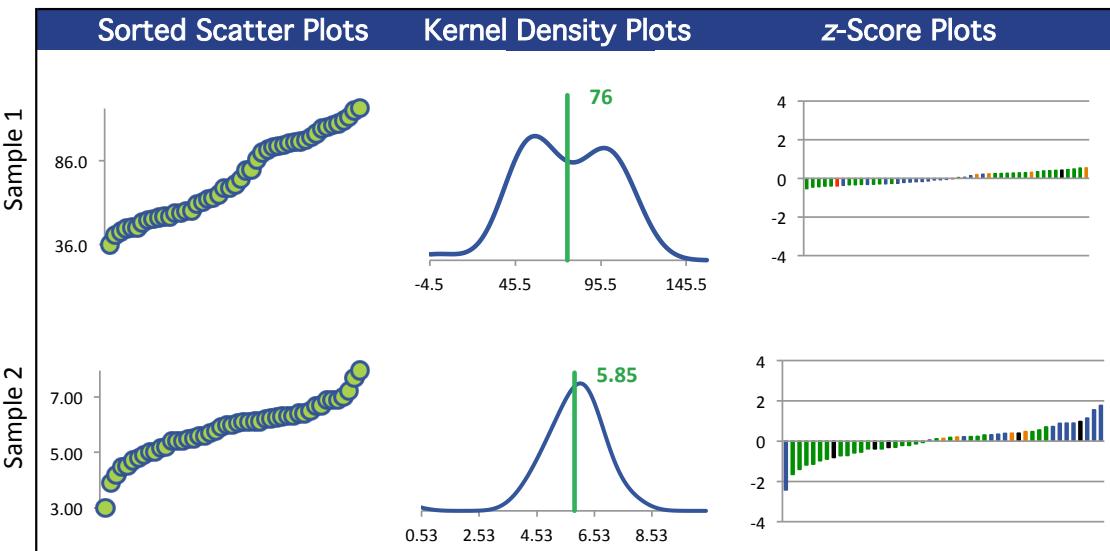
### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	47	47	47	47
Median $\mu\text{g/L}$	71.7	6.00	58.7	30.0
Robust Mean $\mu\text{g/L}$	76.0	5.85	58.7	29.4
$U \mu\text{g/L}$	5.01	0.167	1.39	0.676
Robust Standard Deviation $\mu\text{g/L}$	27.5	0.917	7.62	3.71
Regression Standard Deviation $\mu\text{g/L}$	11.4	0.878	8.81	4.42
Stability Flag				
Homogeneity Flag	Homogeneity	Homogeneity		
Standard Deviation Used ( $SDPA \mu\text{g/L}$ )	74.2	1.18	8.81	4.42
Outliers	0	0	0	0
$ z  > 3.0$	0	0	0	0
$2 <  z  < 3$	0	1	1	1

### Methods Used

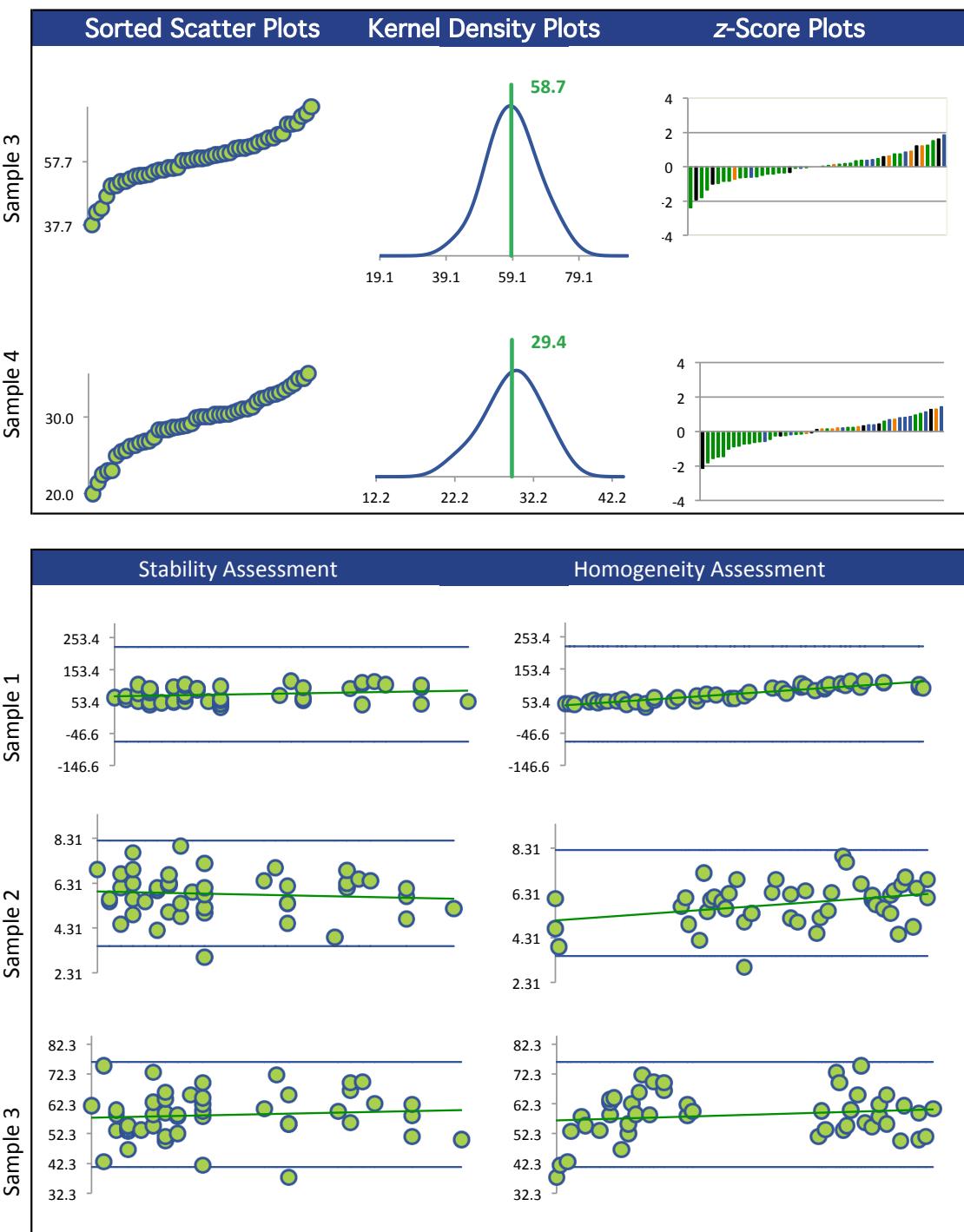
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	24	24	24	24
GC/MS (Red)	3	3	3	3
GC/MS - HEADSPACE (Green)	19	19	19	19
GC/FID - PURGE AND TRAP (Orange)	1	1	1	1

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



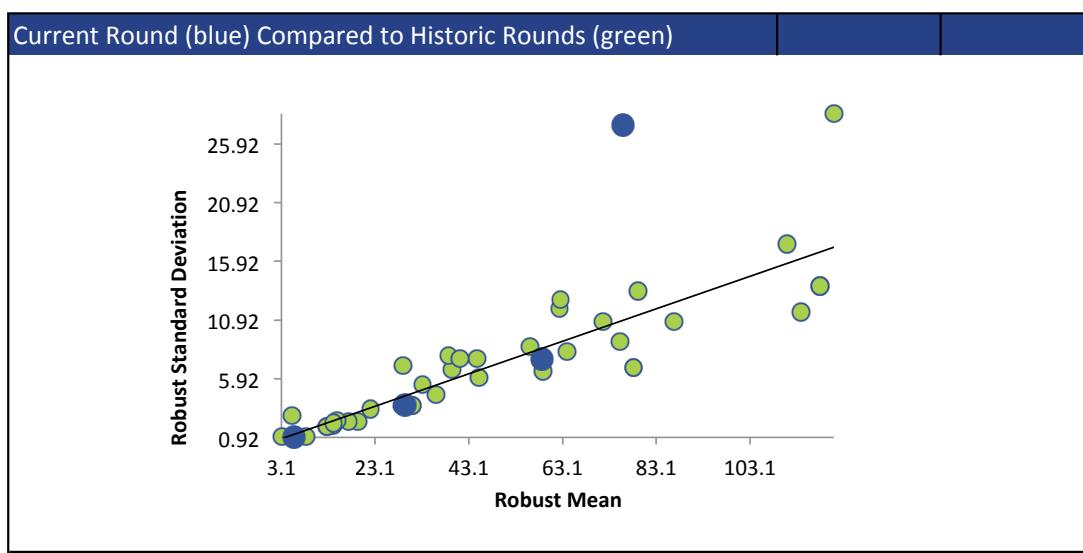
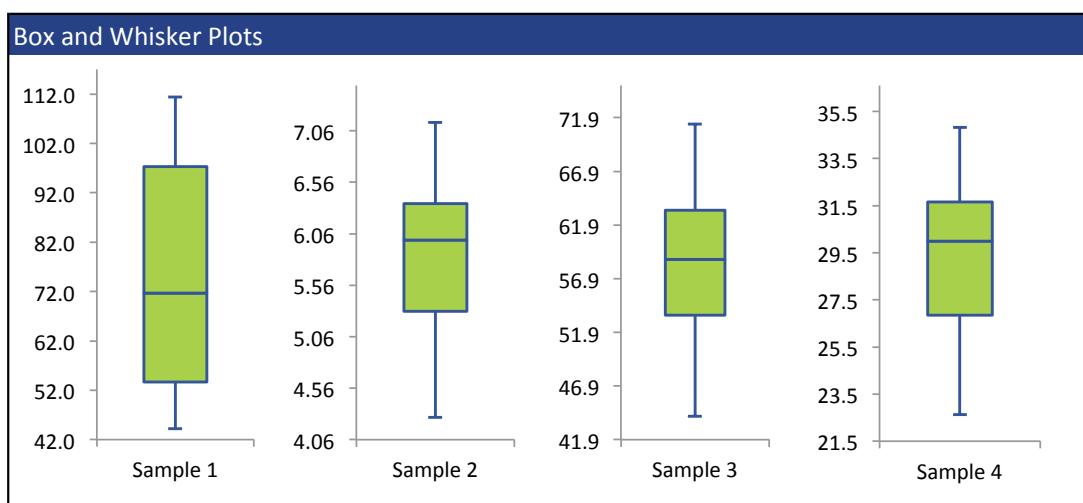
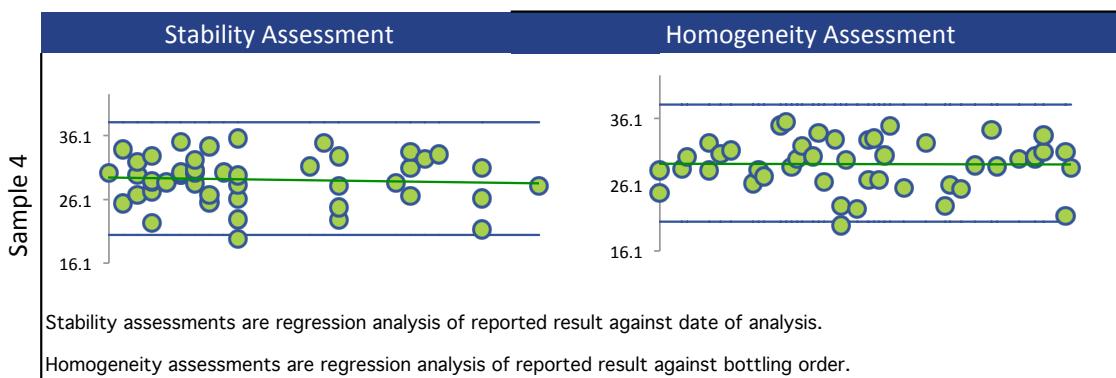
Annex A Summary by Analyte

TRANS-1,2-DICHLOROETHYLENE



## Annex A Summary by Analyte

### TRANS-1,2-DICHLOROETHYLENE



Annex A Summary by Analyte

## TRANS-1,3-DICHLOROPROPENE

### Summary Statistics

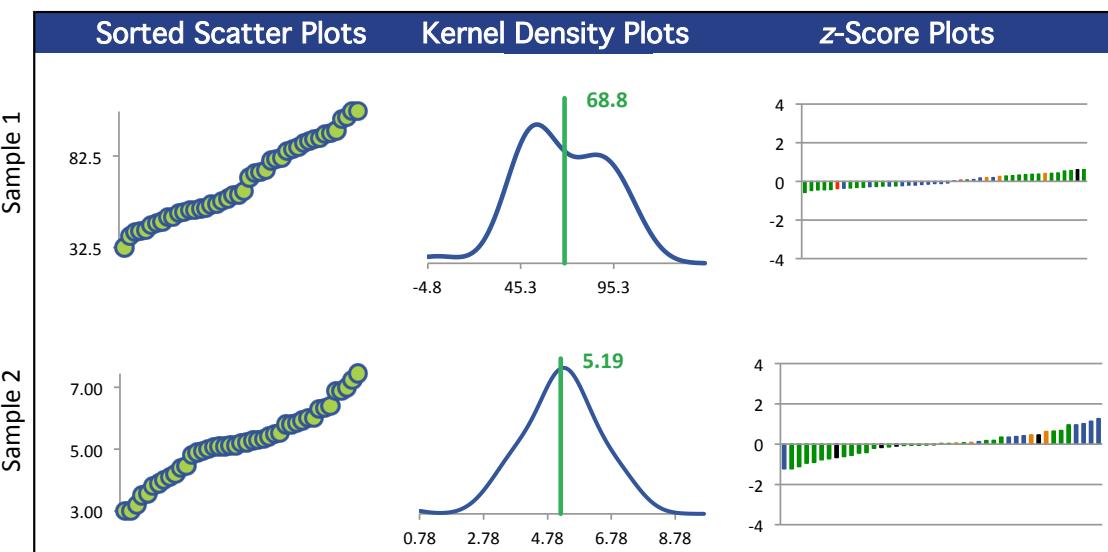
### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	44	43	44	44
Median $\mu\text{g/L}$	62.3	5.20	51.5	26.4
Robust Mean $\mu\text{g/L}$	68.8	5.19	52.2	25.9
U $\mu\text{g/L}$	4.58	0.227	1.62	0.872
Robust Standard Deviation $\mu\text{g/L}$	24.3	1.19	8.61	4.63
Regression Standard Deviation $\mu\text{g/L}$	10.3	0.779	7.83	3.88
Stability Flag		Stability		Stability
Homogeneity Flag	Homogeneity			
Standard Deviation Used (SDPA) $\mu\text{g/L}$	62.9	1.80	8.61	5.44
Outliers	0	0	0	0
$ z  > 3.0$	0	0	1	0
$2 <  z  < 3$	0	0	3	1

### Methods Used

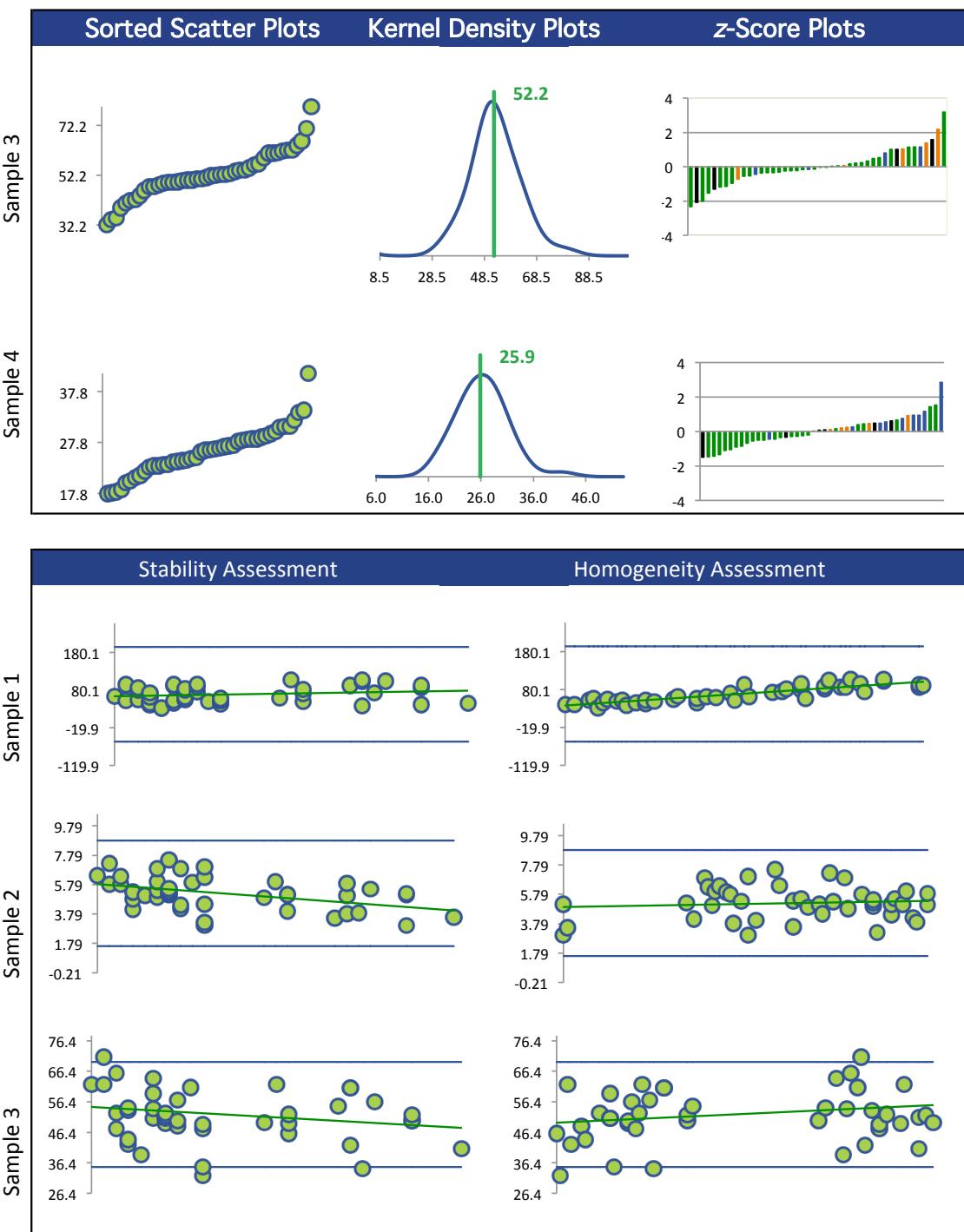
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - HEADSPACE (Blue)	18	18	18	18
GC/MS - PURGE AND TRAP (Red)	23	23	23	23
GC/MS (Green)	3	2	3	3

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



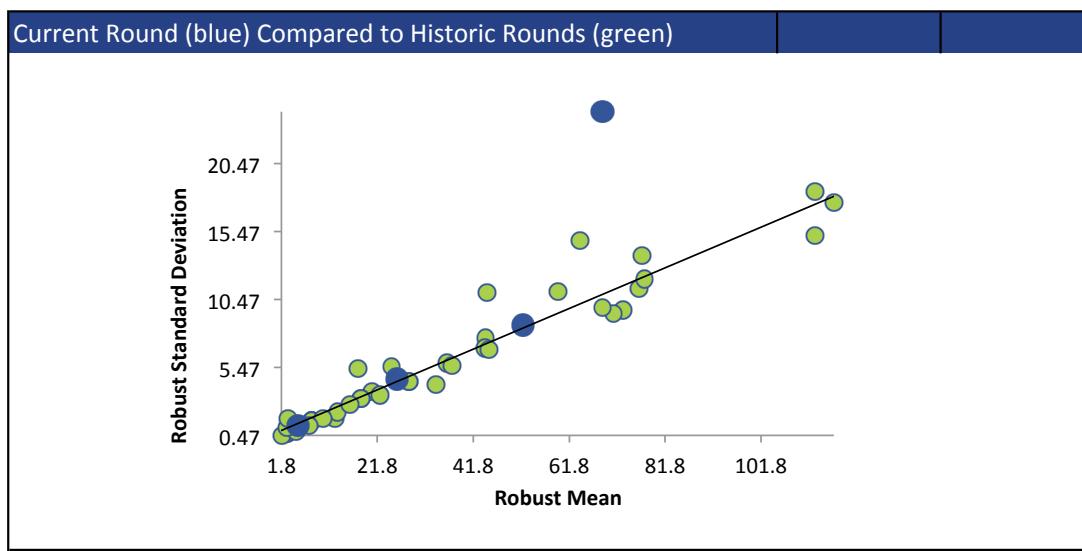
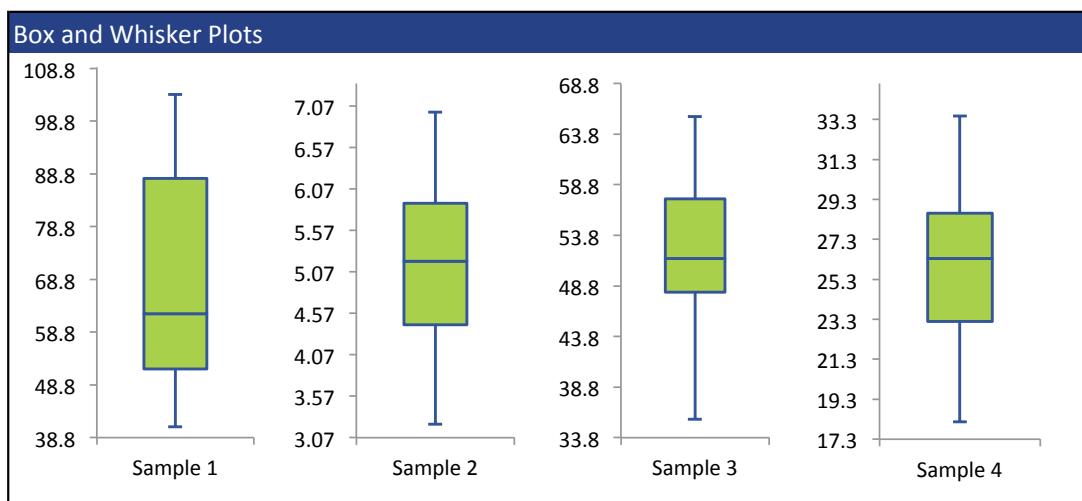
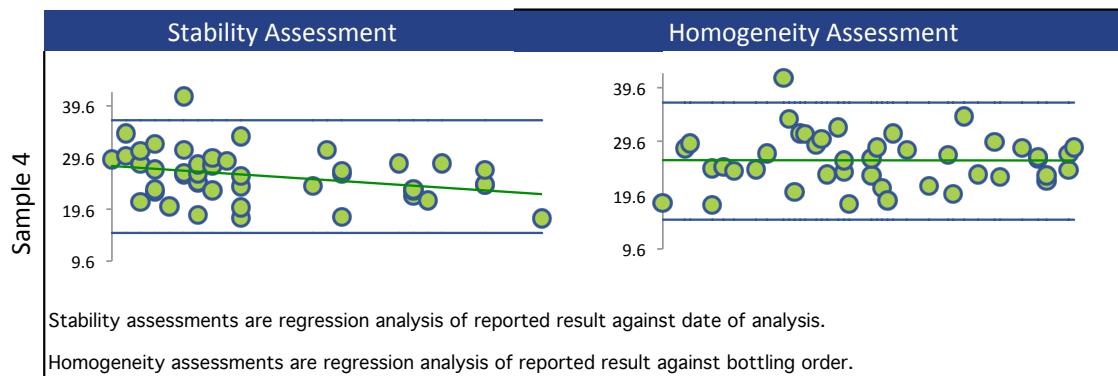
Annex A Summary by Analyte

TRANS-1,3-DICHLOROPROPENE



## Annex A Summary by Analyte

### TRANS-1,3-DICHLOROPROPENE



Annex A Summary by Analyte

## TRICHLOROETHYLENE

### Summary Statistics

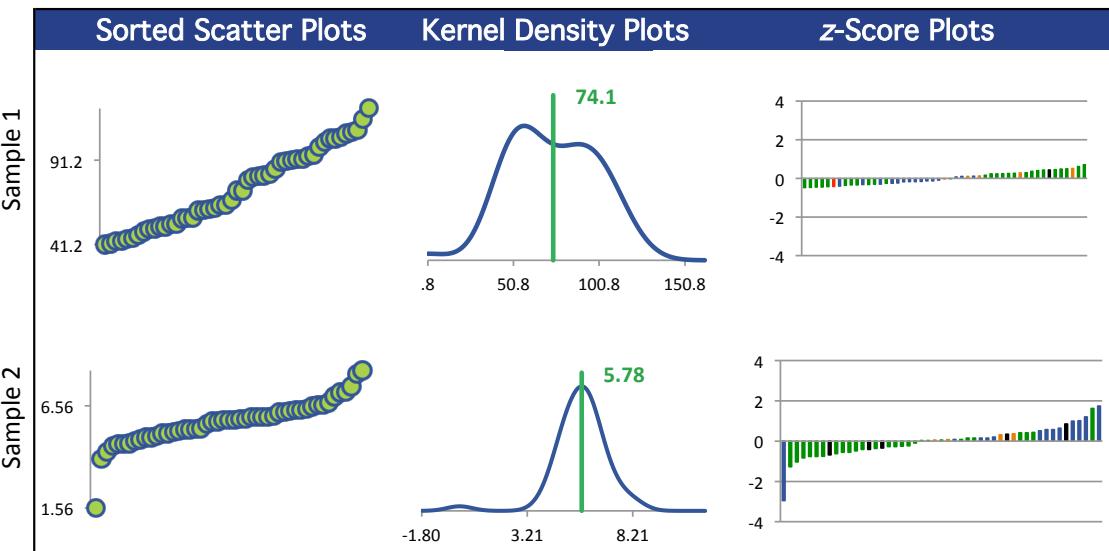
### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	49	49	49	49
Median $\mu\text{g/L}$	73.1	5.85	57.6	29.0
Robust Mean $\mu\text{g/L}$	74.1	5.78	57.3	29.2
$U \mu\text{g/L}$	4.64	0.164	1.47	0.664
Robust Standard Deviation $\mu\text{g/L}$	26.0	0.917	8.25	3.72
Regression Standard Deviation $\mu\text{g/L}$	13.0	1.01	10.0	5.10
Stability Flag	Stability			
Homogeneity Flag	Homogeneity			
Standard Deviation Used ( $SDPA$ ) $\mu\text{g/L}$	67.6	1.43	10.0	5.10
Outliers	0	0	0	0
$ z  > 3.0$	0	0	0	1
$2 <  z  < 3$	0	1	0	1

### Methods Used

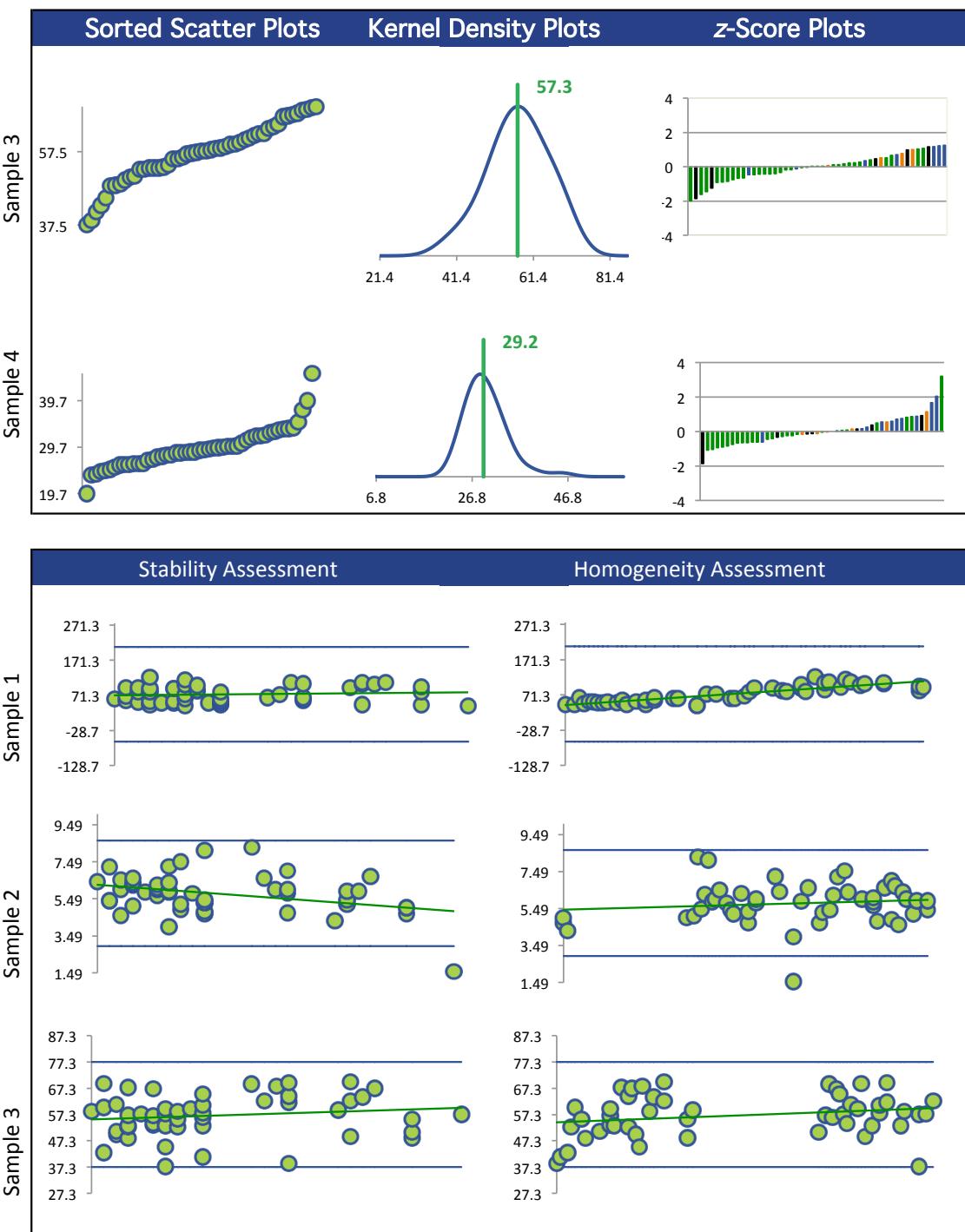
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	27	27	27	27
GC/MS - HEADSPACE (Red)	18	18	18	18
GC/MS/MS - HEADSPACE (Green)	1	1	1	1
GC/FID - PURGE AND TRAP (Orange)	1	1	1	1
GC/MS (Black)	2	2	2	2

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



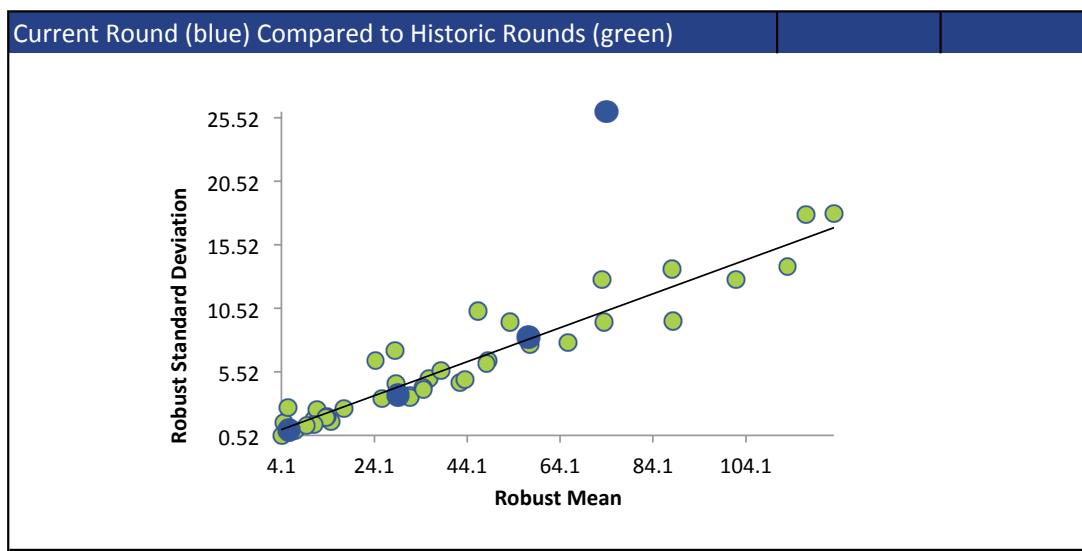
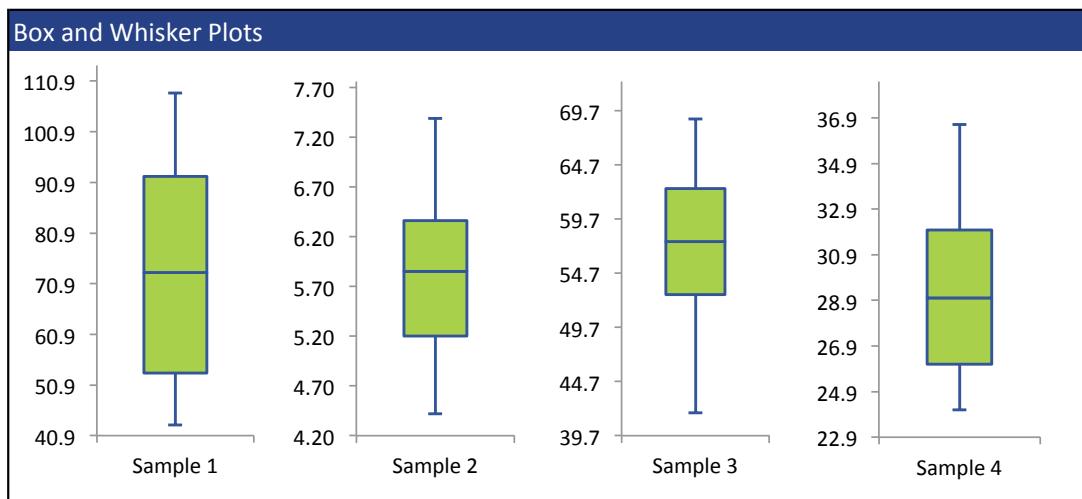
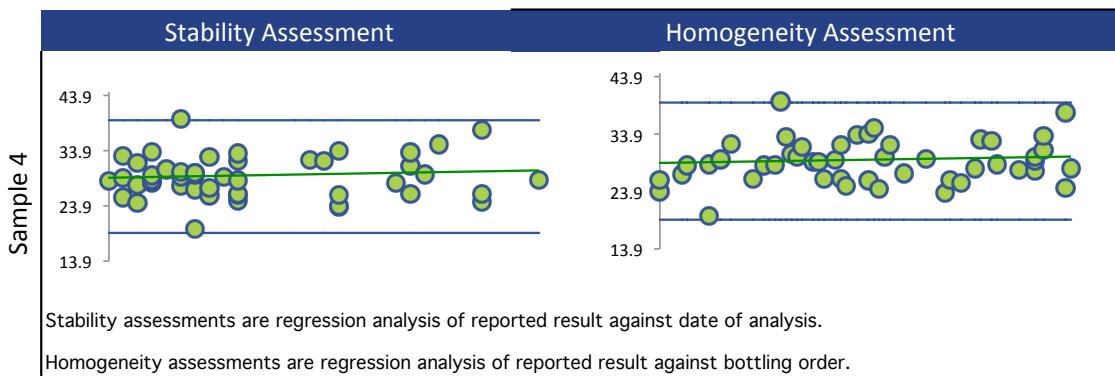
Annex A Summary by Analyte

TRICHLOROETHYLENE



## Annex A Summary by Analyte

### TRICHLOROETHYLENE



Annex A Summary by Analyte

## TRICHLOROFLUOROMETHANE

### Summary Statistics

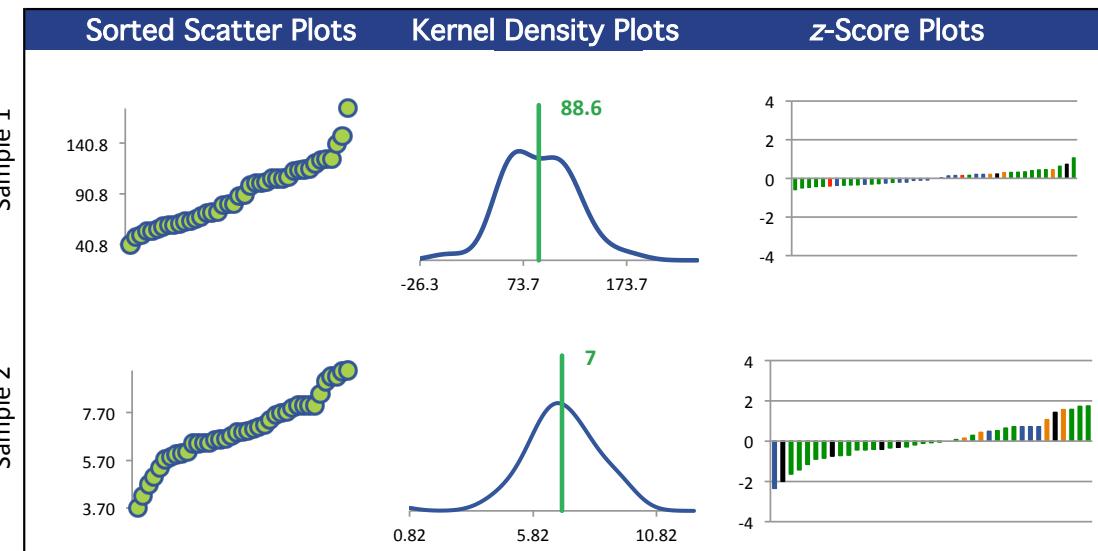
### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	41	39	41	41
Median $\mu\text{g/L}$	88.5	6.90	68.5	34.4
Robust Mean $\mu\text{g/L}$	88.6	7.00	69.0	34.9
$U \mu\text{g/L}$	6.23	0.282	2.03	1.06
Robust Standard Deviation $\mu\text{g/L}$	31.9	1.41	10.4	5.42
Regression Standard Deviation $\mu\text{g/L}$	17.7	1.40	13.8	6.98
Stability Flag				
Homogeneity Flag	Homogeneity			
Standard Deviation Used ( $SDPA \mu\text{g/L}$ )	81.3	1.41	13.8	6.98
Outliers	0	0	0	0
$ z  > 3.0$	0	0	0	0
$2 <  z  < 3$	0	1	3	1

### Methods Used

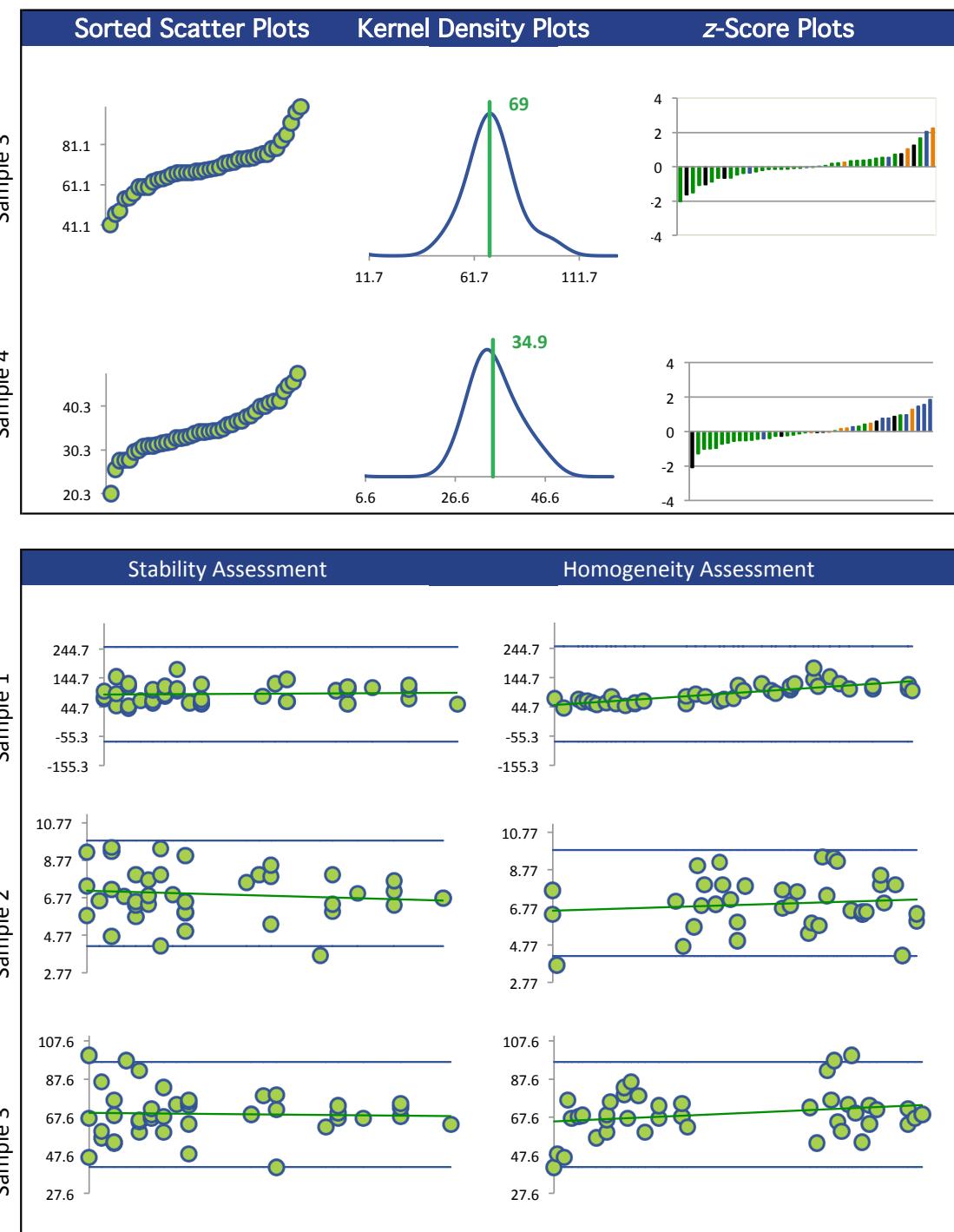
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	21	21	21	21
GC/MS - HEADSPACE (Red)	18	17	18	18
GC/MS (Green)	2	1	2	2

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



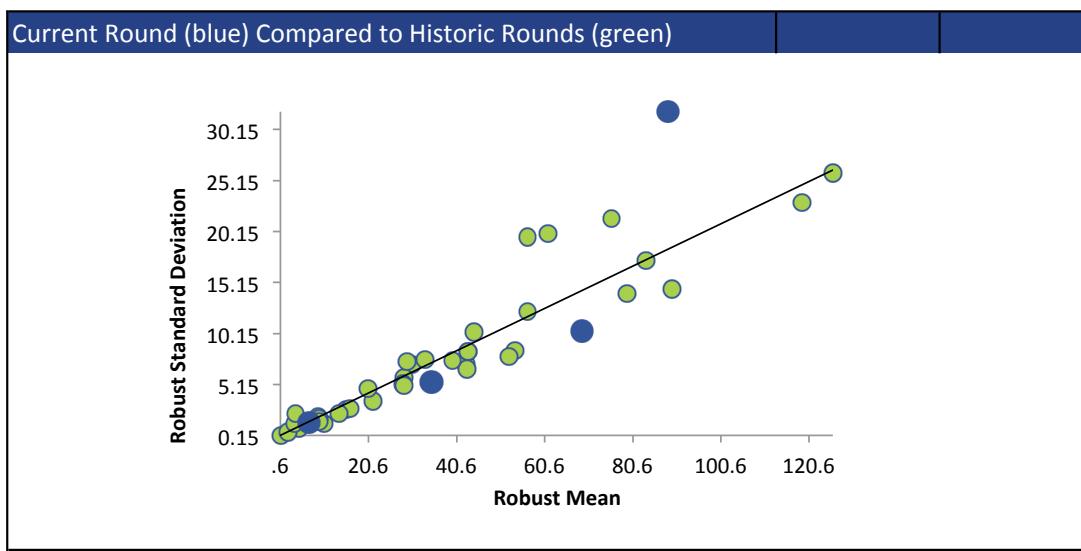
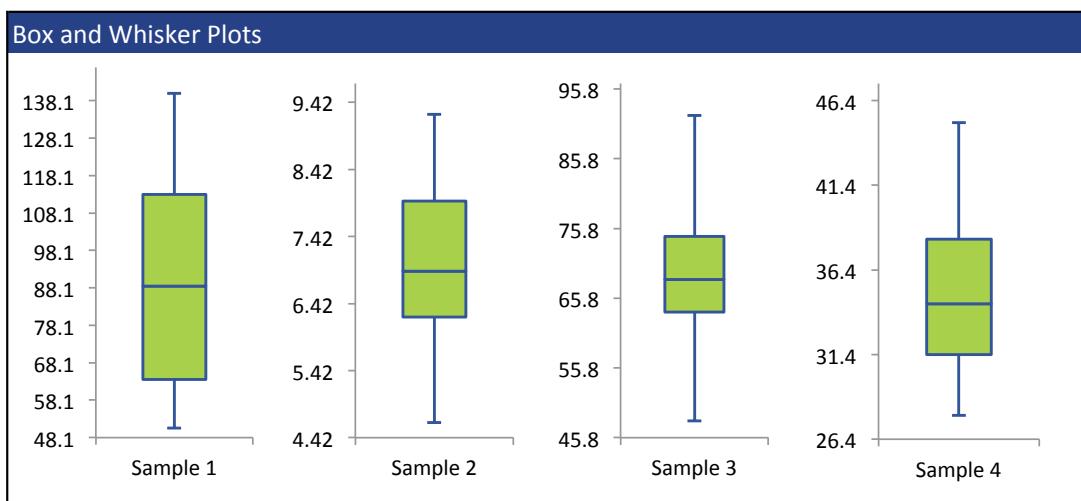
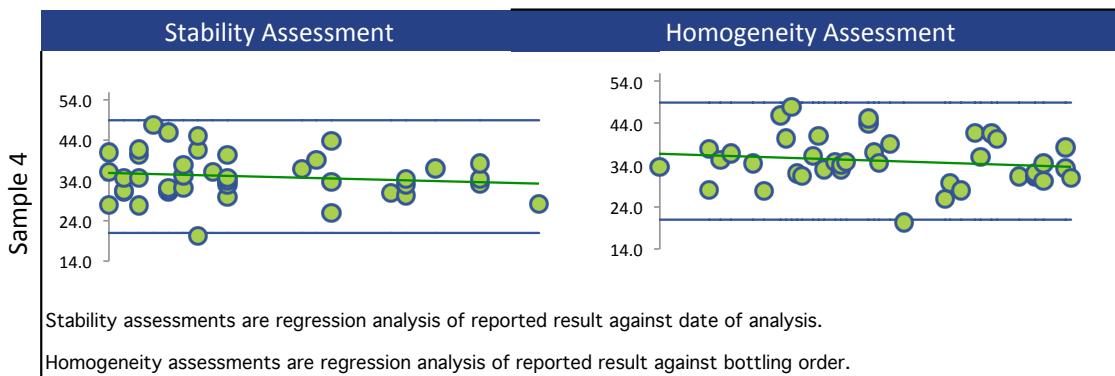
Annex A Summary by Analyte

TRICHLOROFLUOROMETHANE



## Annex A Summary by Analyte

### TRICHLOROFLUOROMETHANE



Annex A Summary by Analyte

## VINYL CHLORIDE

### Summary Statistics

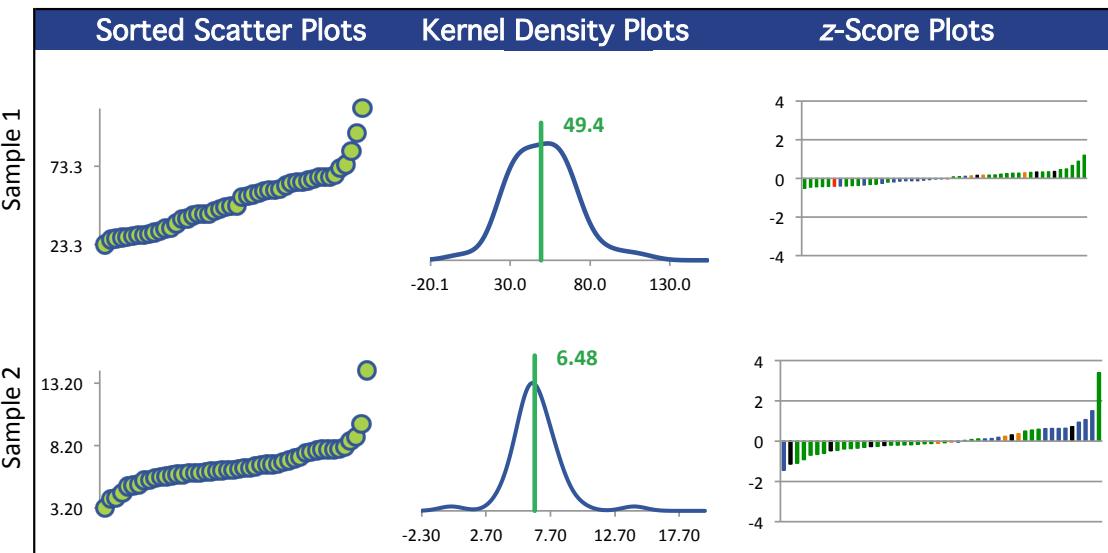
### Excluded

Statistic	C16-1	C16-2	C16-3	C16-4
N	48	48	48	48
Median $\mu\text{g/L}$	48.1	6.30	40.1	17.1
Robust Mean $\mu\text{g/L}$	49.4	6.48	40.5	17.7
$U \mu\text{g/L}$	3.28	0.229	1.45	0.552
Robust Standard Deviation $\mu\text{g/L}$	18.2	1.27	8.03	3.06
Regression Standard Deviation $\mu\text{g/L}$	11.1	1.46	9.12	3.99
Stability Flag				
Homogeneity Flag	Homogeneity	Homogeneity		
Standard Deviation Used ( $SDPA \mu\text{g/L}$ )	50.4	2.28	9.12	3.99
Outliers	0	0	0	0
$ z  > 3.0$	0	1	1	0
$2 <  z  < 3$	0	0	3	3

### Methods Used

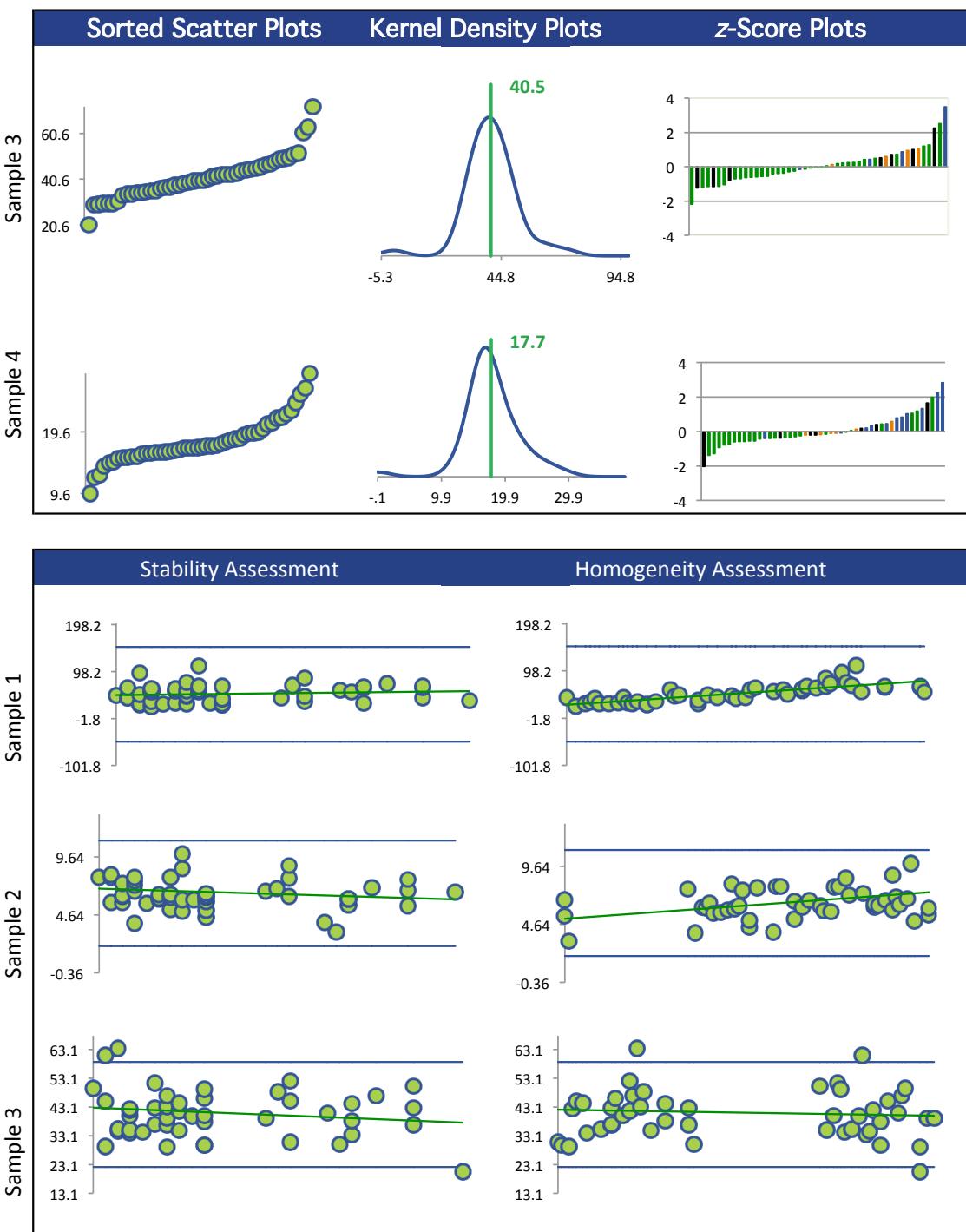
Method	C16-1	C16-2	C16-3	C16-4
GC/MS - PURGE AND TRAP (Blue)	27	27	27	27
GC/MS - HEADSPACE (Red)	18	18	18	18
GC/MS/MS - HEADSPACE (Green)	1	1	1	1
GC/MS (Orange)	2	2	2	2

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



Annex A Summary by Analyte

VINYL CHLORIDE



## Annex A Summary by Analyte

### VINYL CHLORIDE

