

Test Group Summary Report

C06A OC Pesticides in Water

June 2025 PT Round

Issued: August 25, 2025

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1.0 The Proficiency Testing Report

The Proficiency Testing Report consists of two parts.

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

2.0 Definitions

The PTC Proficiency Testing Reports contain some terms that new participants may not be familiar with:

- PT Code:* The PT Code is unique to each analyte that a participant is registered for.
- Lab Info:* If a participant is accredited by CALA, this three-digit number in the Laboratory Information is the appendix number that the accredited method is assigned to.
- N:* The number of participants results that were used to calculate the summary statistics. This excludes qualified data (e.g., <) and any results that were flagged as outliers.
- Assigned Value:* The Assigned Value is the Robust Mean of the reported results, outliers excluded. This is often referred to as the "target" value.
- Reported Value:* The result reported by the participant.
- SDPTA:* The Standard Deviation of Proficiency Testing Assessment; this value is used to determine the acceptance limits for the PT evaluation. The calculations examine the Robust Standard Deviation (from consensus STD) and Regression Standard Deviation (STD from historical studies or TNI limits); and, if any, Homogeneity and Stability flags to determine the SDPTA (whichever is higher).
- z-Score:* A value assigned to each reported result that is a measure of the degree to which it deviates from the Assigned Value.
- PT Score:* The composite score of the four results reported for each analyte. It is normalized to a score out of 100.
- Bias:* A flag assigned if bias is detected using the re-scaled z-score procedure.
- U:* The Uncertainty of the assigned value.

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 *Statistical methods for use in proficiency testing by interlaboratory comparison*.

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.

$$RSZ = \frac{\square z}{\sqrt{N}} \quad \text{where } \begin{aligned} z &= \text{the z- score} \\ N &= \text{the number of samples} \end{aligned}$$

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.

Annex A Summary by Analyte

a - Chlordane

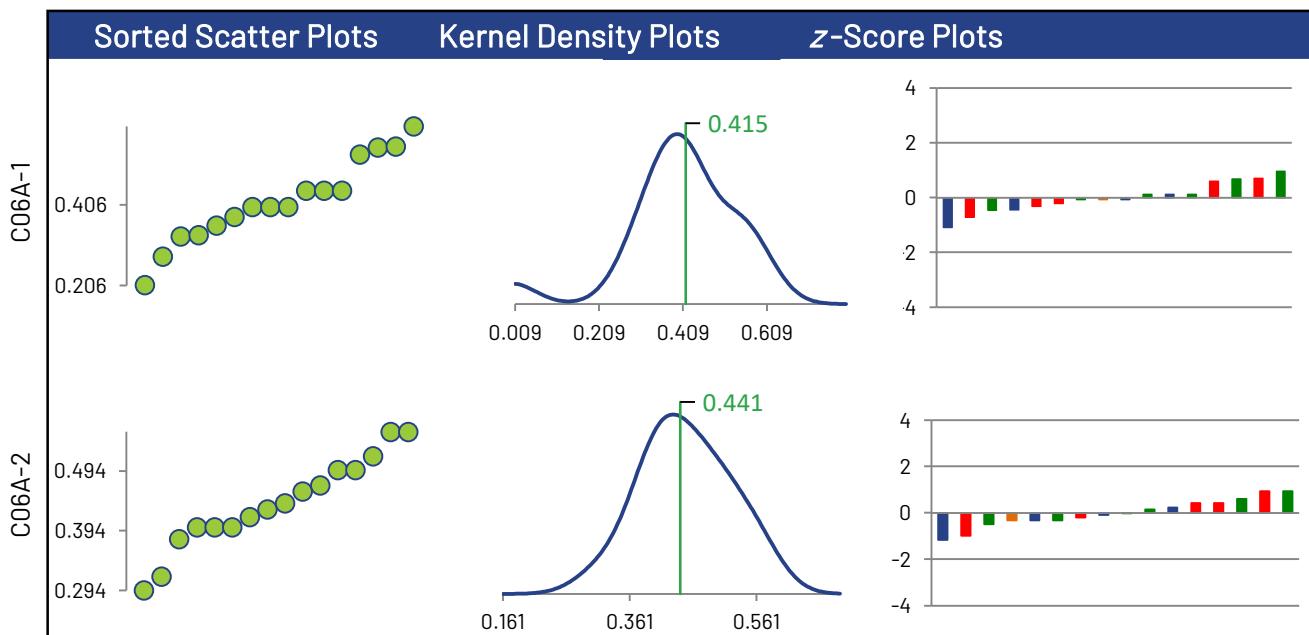
Summary Statistics

Statistic	C06A-1	C06A-2	C06A-3	C06A-4
No. of results used for summary statistics	16	16	18	18
Median µg/L	0.400	0.435	3.67	3.22
Robust Mean µg/L	0.415	0.441	3.51	3.15
Uncertainty of the Assigned Value µg/L	0.0353	0.0263	0.386	0.227
Robust Standard Deviation µg/L	0.113	0.0840	1.31	0.772
Regression Standard Deviation µg/L	0.0788	0.0838	0.667	0.598
Stability Flag	Stability	Stability		Stability
Homogeneity Flag				
Standard Deviation Used (SDPA) µg/L	0.191	0.127	1.31	1.39
Flagged Outliers	1	1	0	0
$ z > 3.0$	0	0	0	0
$2 < z < 3$	0	0	1	0
Failure Rate %*		5.6		

Methods Used

Method	C06A-1	C06A-2	C06A-3	C06A-4
GC/MS (Blue)	4	4	5	5
GC/MS/MS (Red)	5	5	5	5
GC/ECD (Green)	6	6	7	7
GCXGC/ECD (Orange)	1	1	1	1

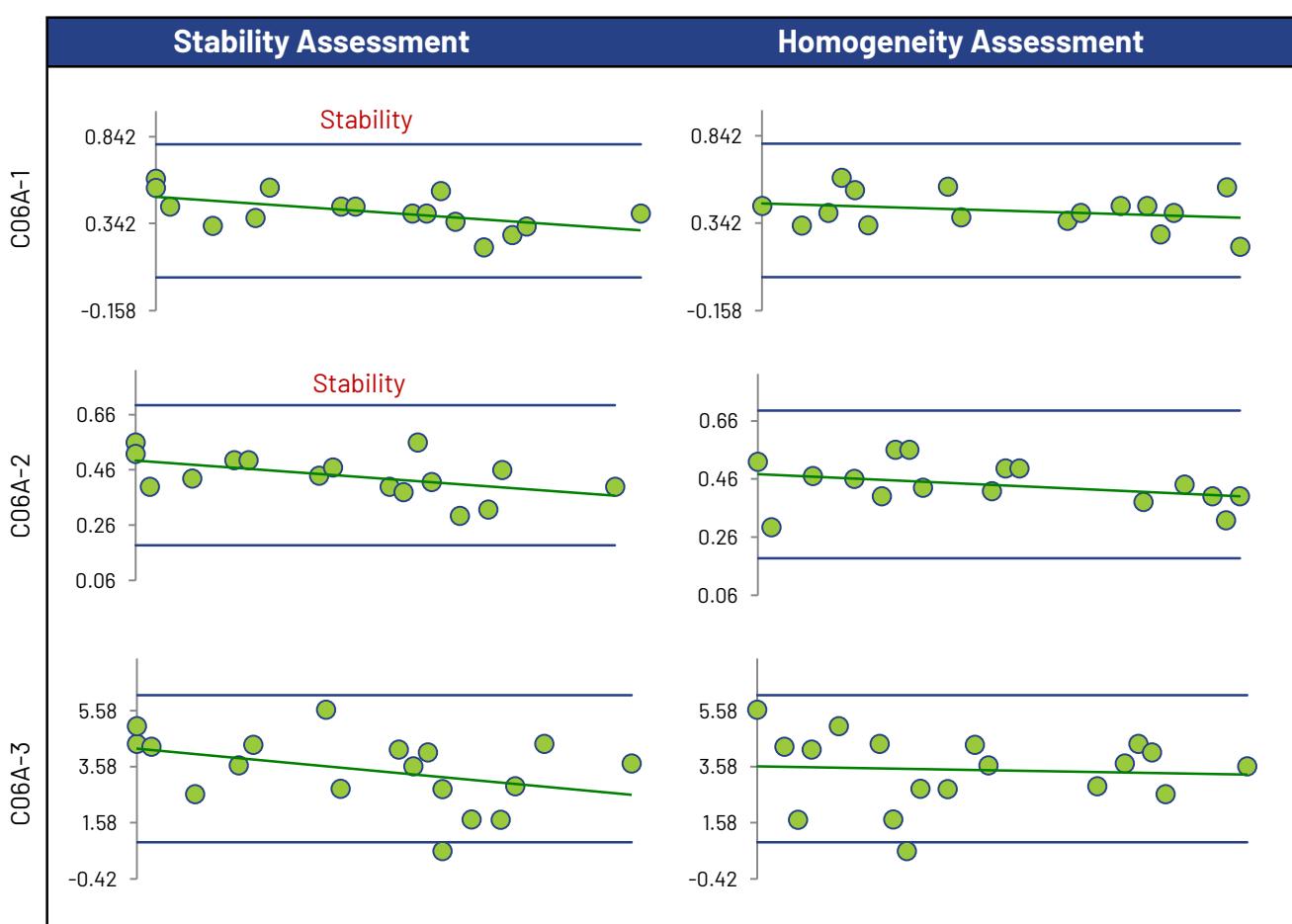
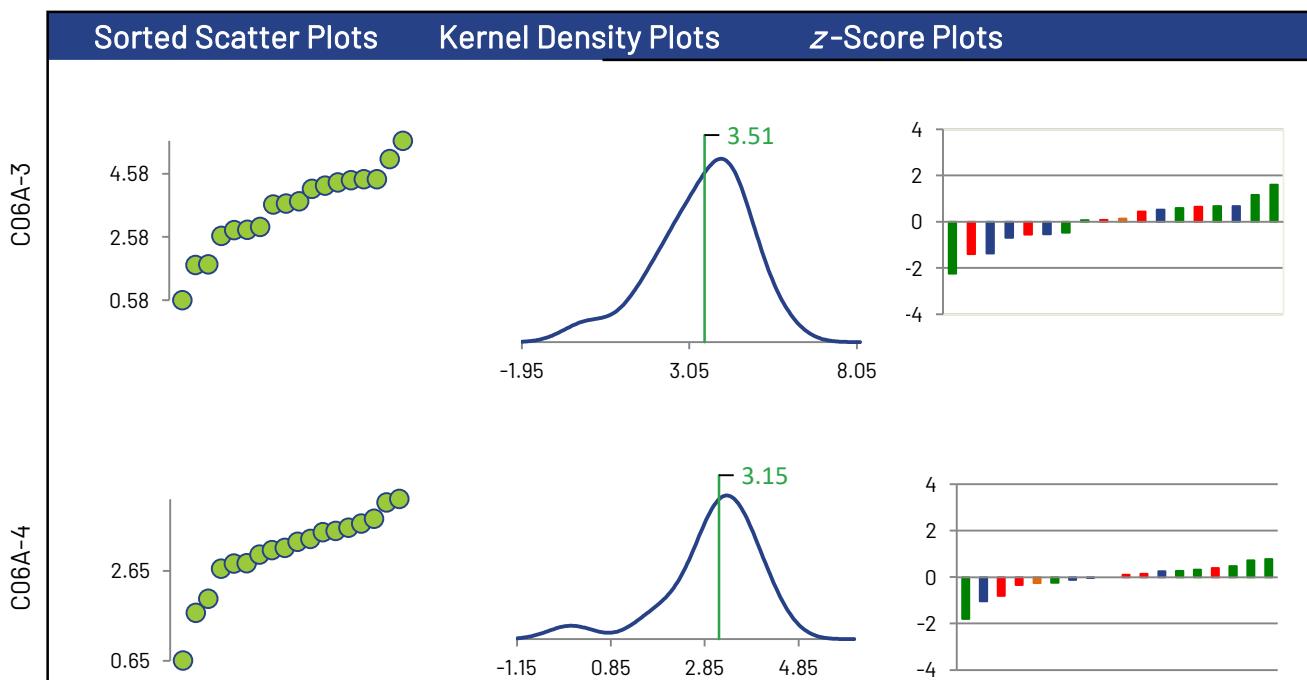
All summary stats, except the failure rate, and the plots below are based on the data excluding any flagged outliers



* Displayed failure rate does not account for the RDL modified z-Scores or inaccurately reported non-detects.

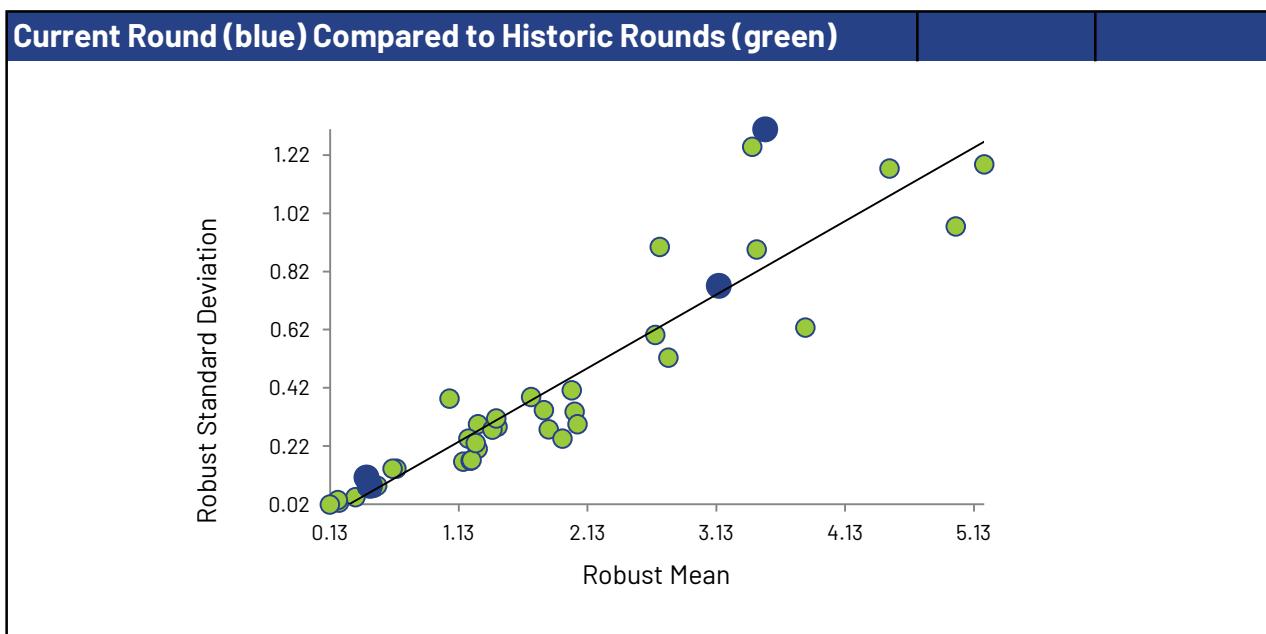
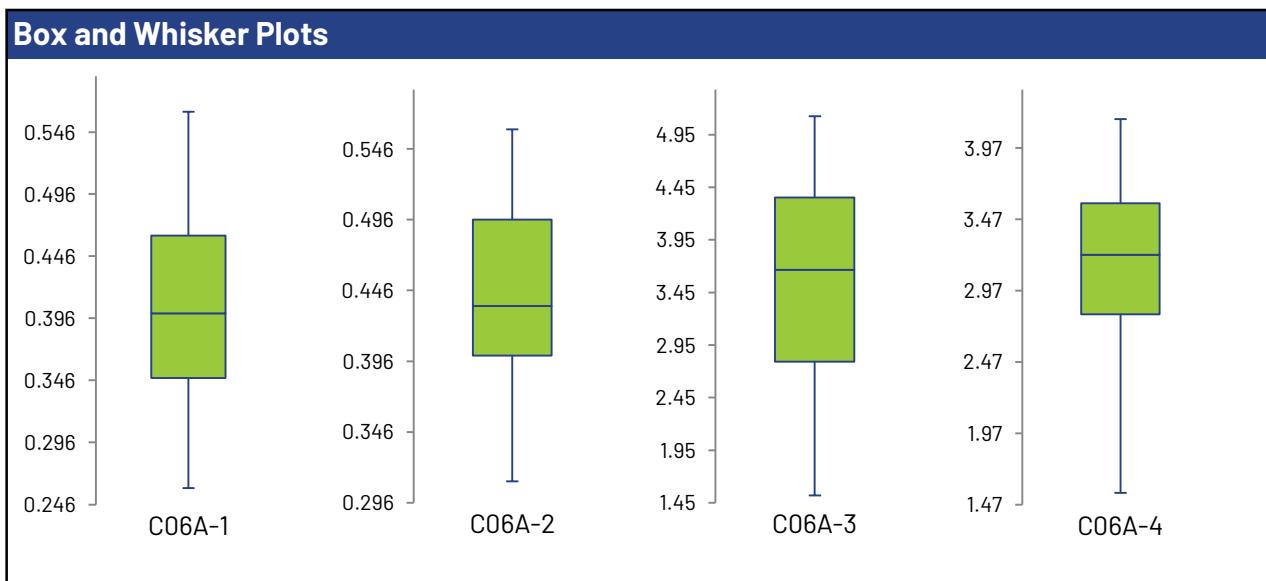
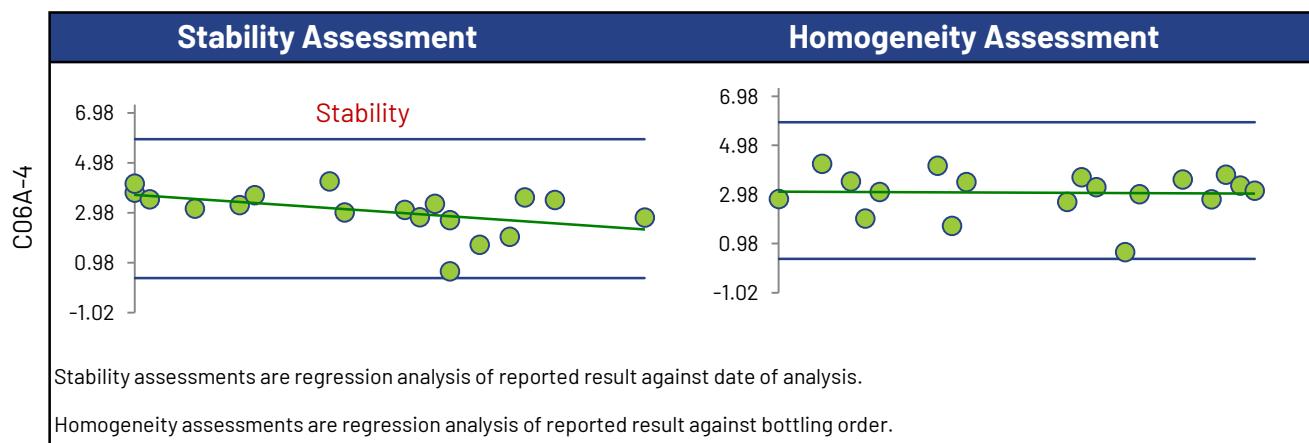
Annex A Summary by Analyte

a - Chlordane



Annex A Summary by Analyte

a - Chlordane



Annex A Summary by Analyte

A -BHC

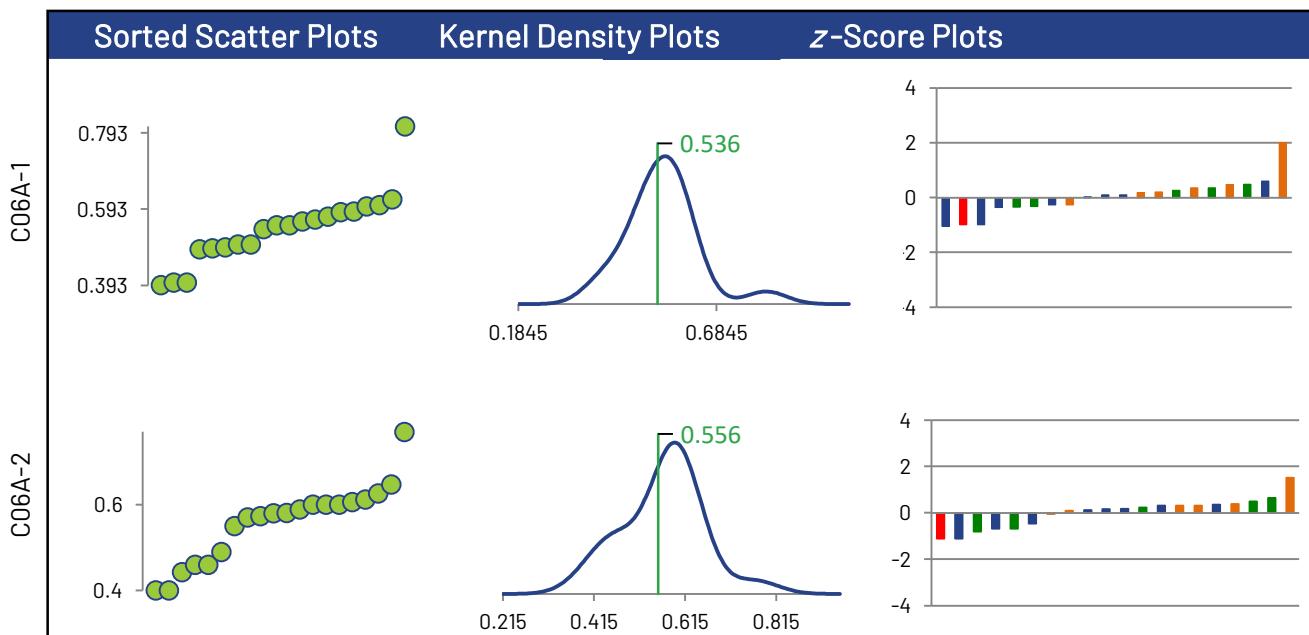
Summary Statistics

Statistic	C06A-1	C06A-2	C06A-3	C06A-4
No. of results used for summary statistics	20	20	20	20
Median µg/L	0.550	0.581	0.557	0.587
Robust Mean µg/L	0.536	0.556	0.553	0.569
Uncertainty of the Assigned Value µg/L	0.0210	0.0248	0.0204	0.0221
Robust Standard Deviation µg/L	0.0750	0.0886	0.0730	0.0790
Regression Standard Deviation µg/L	0.137	0.142	0.142	0.146
Stability Flag				
Homogeneity Flag				
Standard Deviation Used (SDPA) µg/L	0.137	0.142	0.142	0.146
Flagged Outliers	0	0	0	0
$ z > 3.0$	0	0	0	0
$2 < z < 3$	0	0	0	0
Failure Rate %*	0.0			

Methods Used

Method	C06A-1	C06A-2	C06A-3	C06A-4
GC/ECD (Blue)	8	8	8	8
GCXGC/ECD (Red)	1	1	1	1
GC/MS/MS (Green)	5	5	5	5
GC/MS (Orange)	6	6	6	6

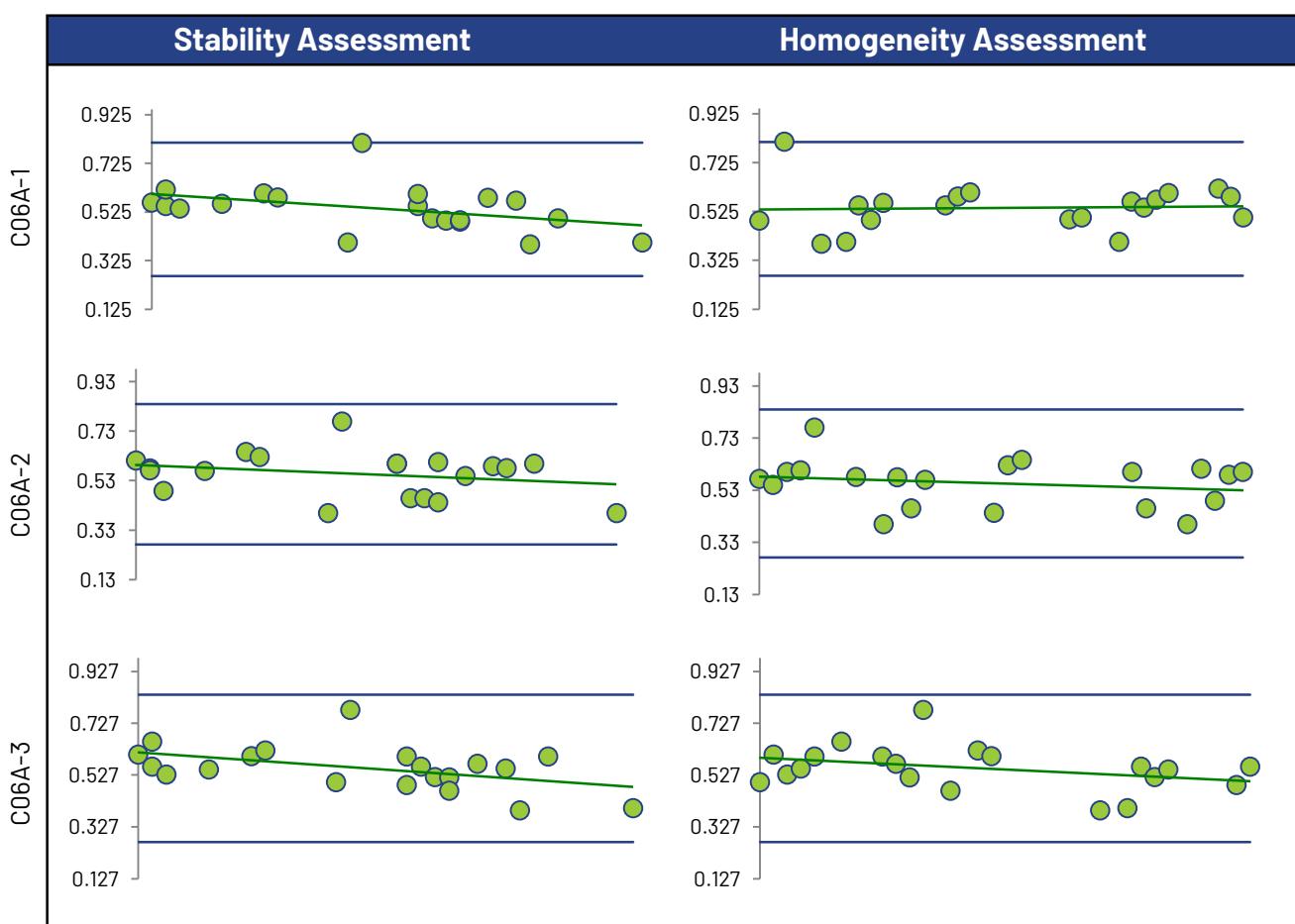
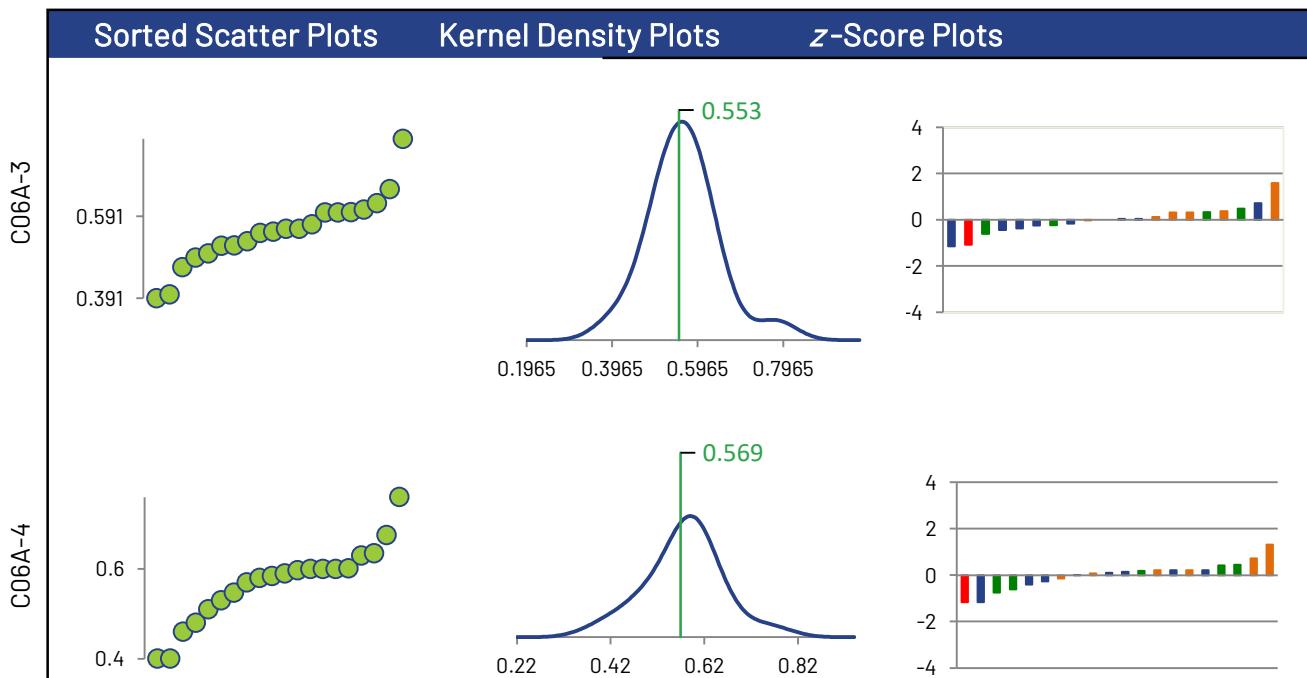
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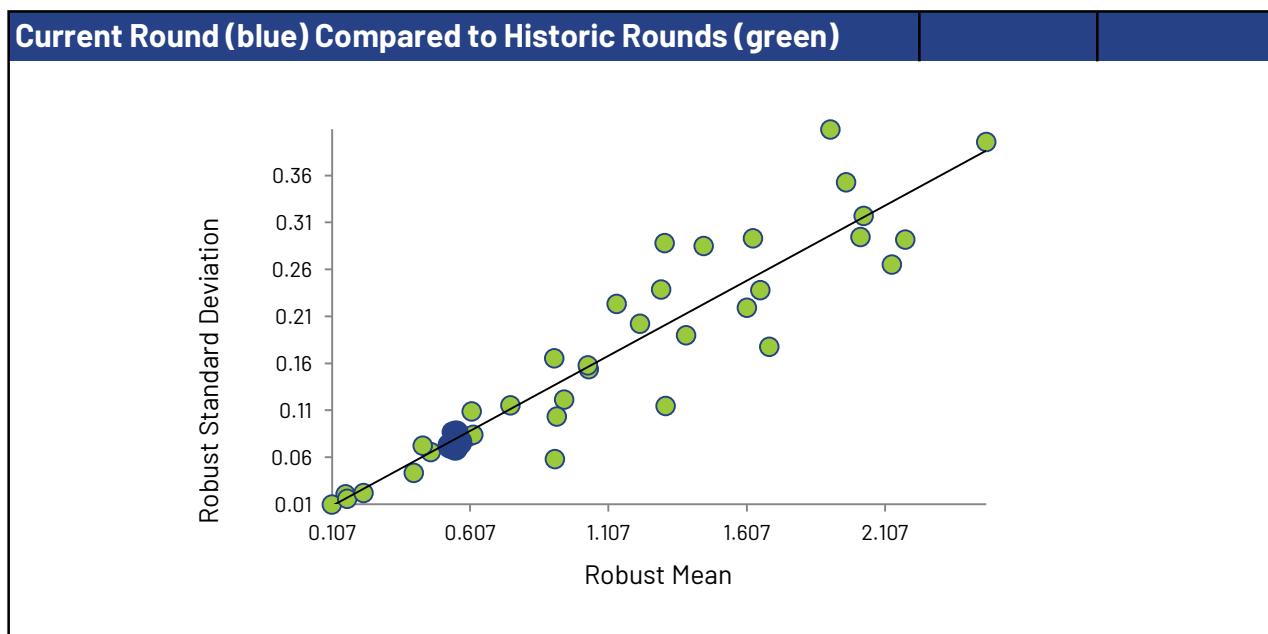
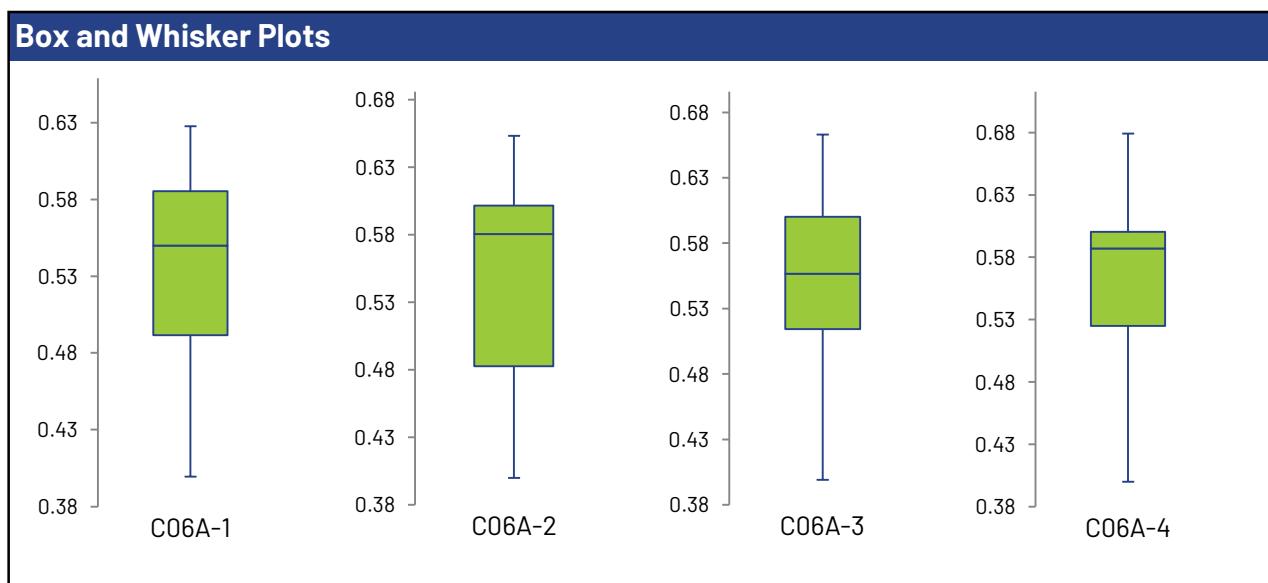
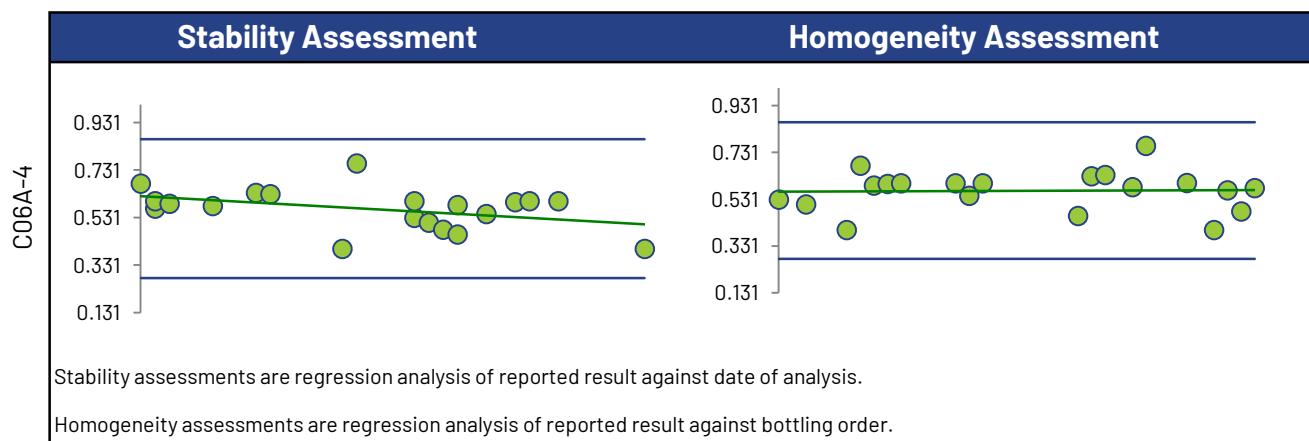
Annex A Summary by Analyte

A -BHC



Annex A Summary by Analyte

A -BHC



Annex A Summary by Analyte

Aldrin

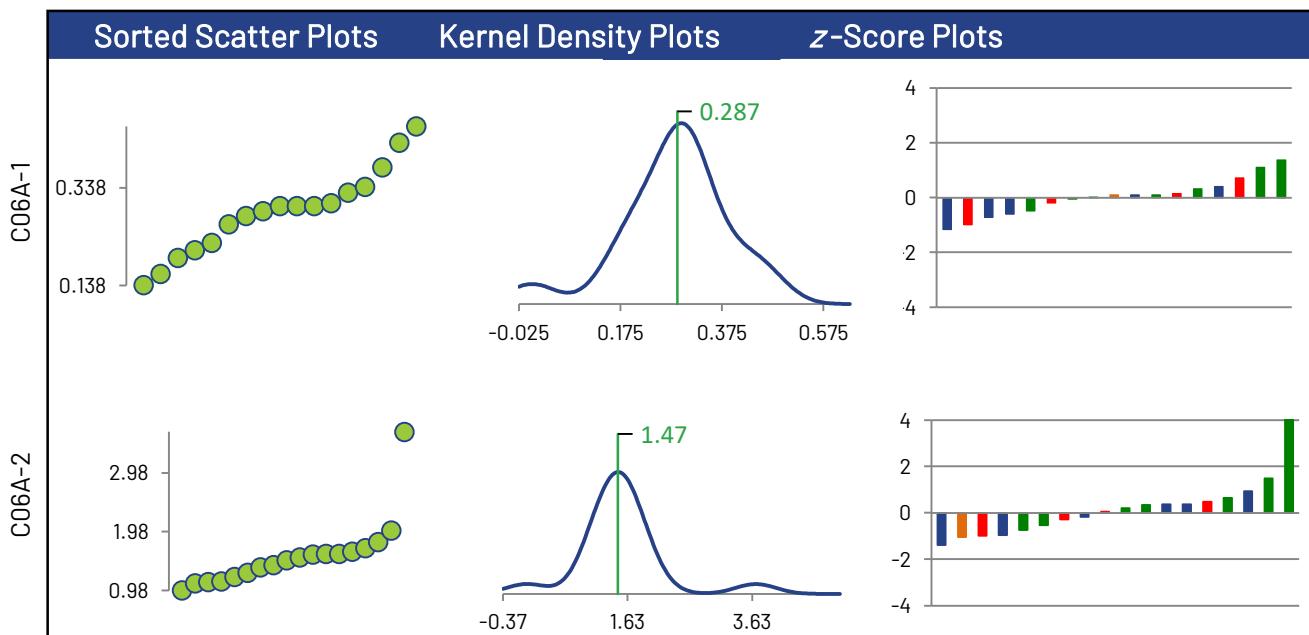
Summary Statistics

Statistic	C06A-1	C06A-2	C06A-3	C06A-4
No. of results used for summary statistics	17	18	18	17
Median µg/L	0.300	1.52	3.47	0.476
Robust Mean µg/L	0.287	1.47	3.33	0.481
Uncertainty of the Assigned Value µg/L	0.0281	0.0990	0.342	0.0327
Robust Standard Deviation µg/L	0.0927	0.336	1.16	0.108
Regression Standard Deviation µg/L	0.0923	0.356	0.768	0.135
Stability Flag				
Homogeneity Flag	Homogeneity			
Standard Deviation Used (SDPA) µg/L	0.129	0.356	1.16	0.135
Flagged Outliers	0	0	0	0
$ z > 3.0$	0	1	0	0
$2 < z < 3$	0	0	0	0
Failure Rate %*		5.6		

Methods Used

Method	C06A-1	C06A-2	C06A-3	C06A-4
GC/MS (Blue)	5	6	6	5
GC/MS/MS (Red)	4	4	4	4
GC/ECD (Green)	7	7	7	7
GCXGC/ECD (Orange)	1	1	1	1

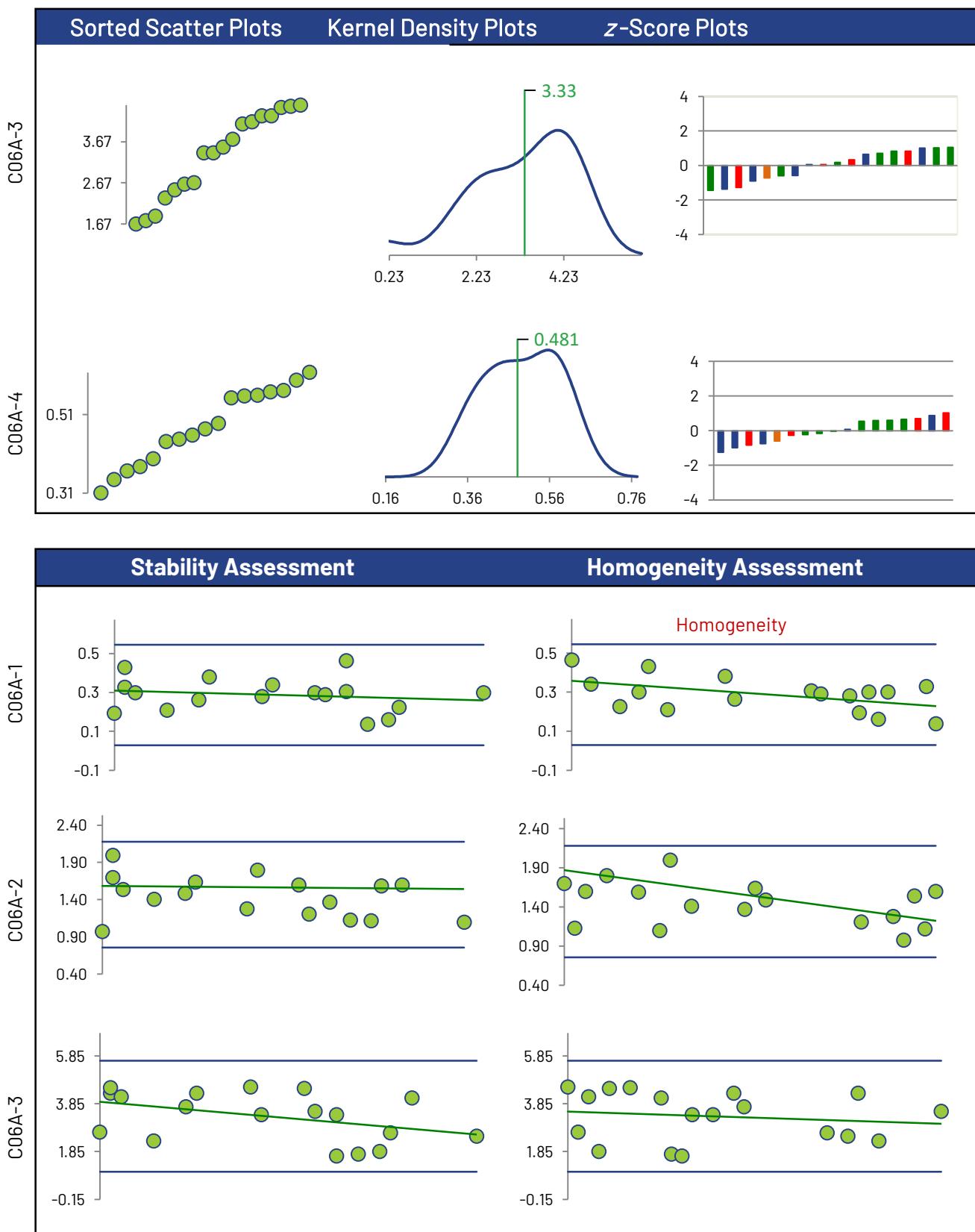
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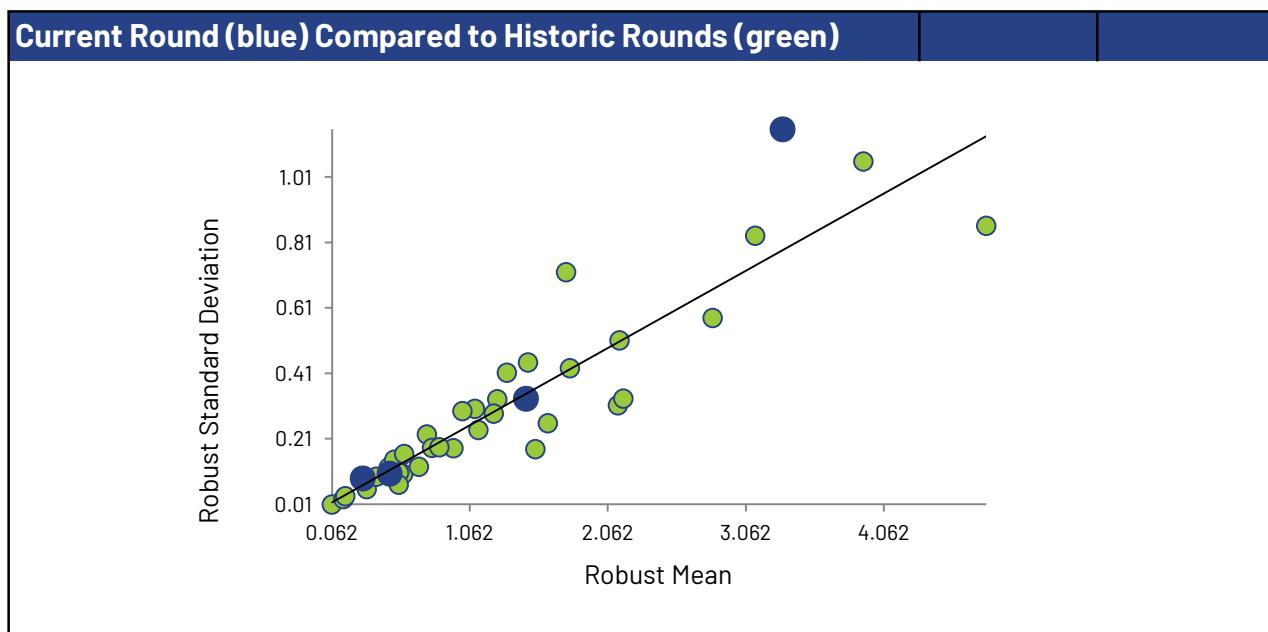
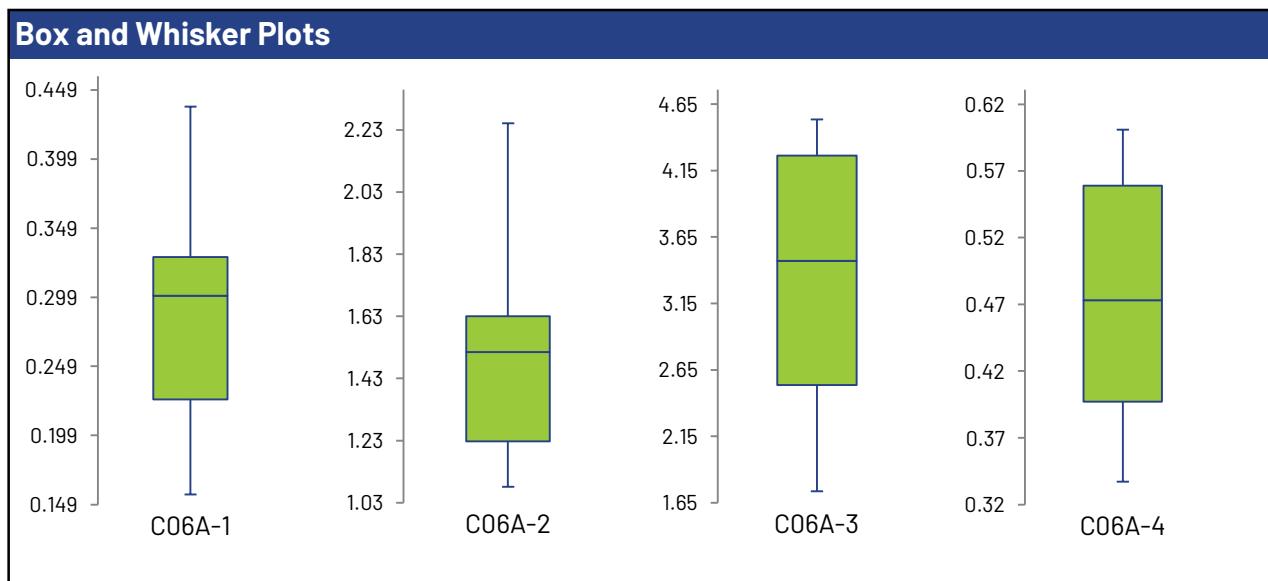
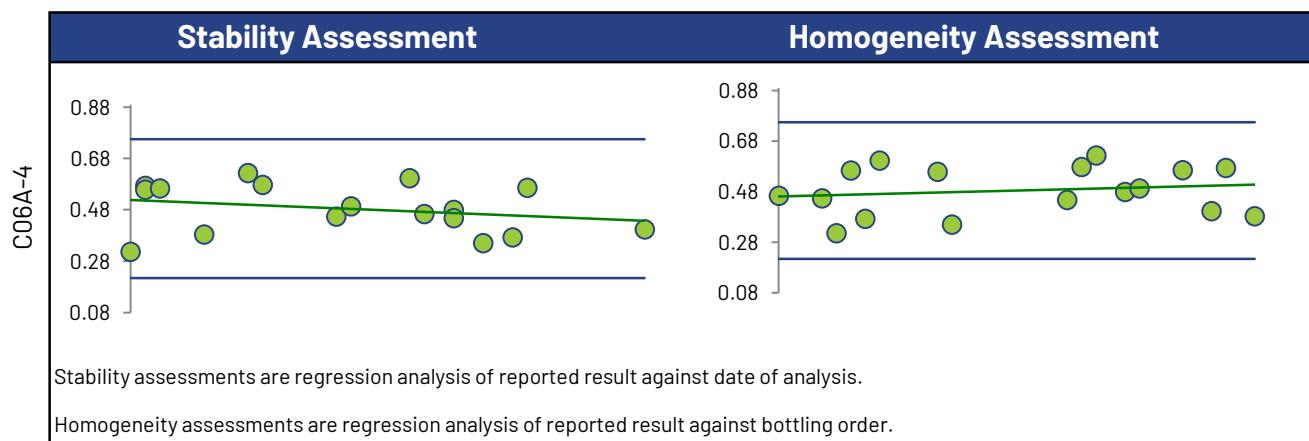
Annex A Summary by Analyte

Aldrin



Annex A Summary by Analyte

Aldrin



Annex A Summary by Analyte

Dieldrin

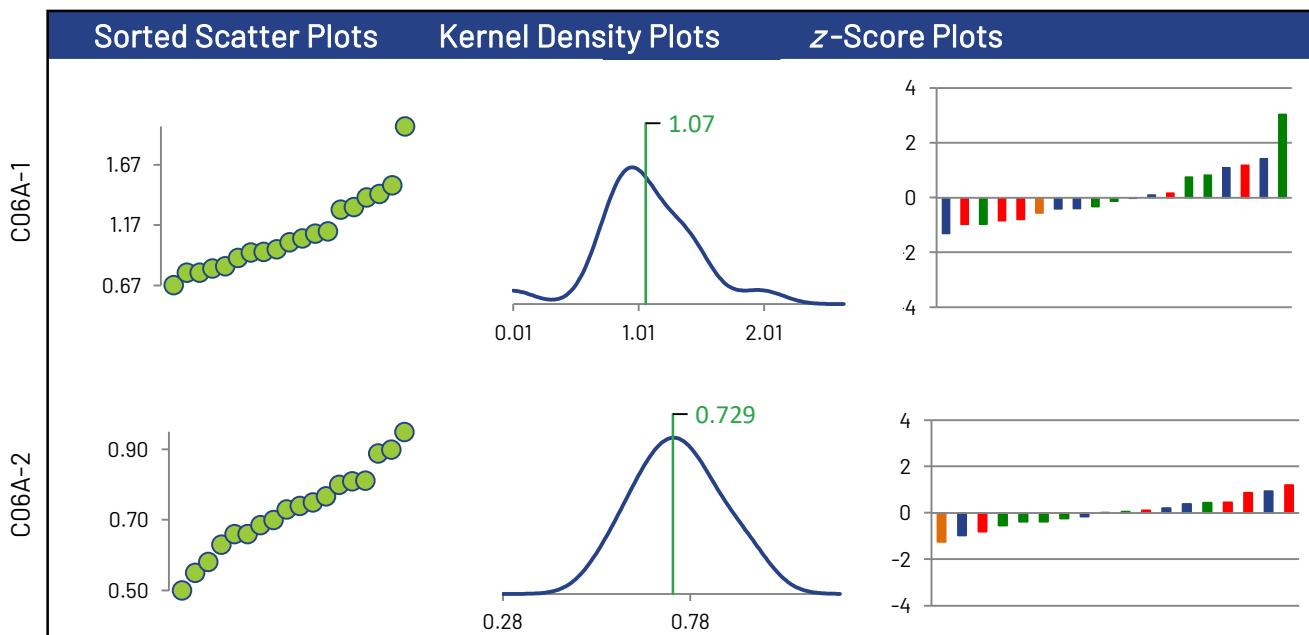
Summary Statistics

Statistic	C06A-1	C06A-2	C06A-3	C06A-4
No. of results used for summary statistics	19	18	19	19
Median µg/L	1.03	0.735	3.44	3.03
Robust Mean µg/L	1.07	0.729	3.35	2.79
Uncertainty of the Assigned Value µg/L	0.0866	0.0389	0.280	0.172
Robust Standard Deviation µg/L	0.302	0.132	0.976	0.599
Regression Standard Deviation µg/L	0.233	0.163	0.694	0.582
Stability Flag		Stability		Stability
Homogeneity Flag				
Standard Deviation Used (SDPA) µg/L	0.302	0.184	0.976	1.04
Flagged Outliers	0	1	0	0
z >3.0	1	0	0	0
2< z <3	0	0	1	0
Failure Rate %*		5.3		

Methods Used

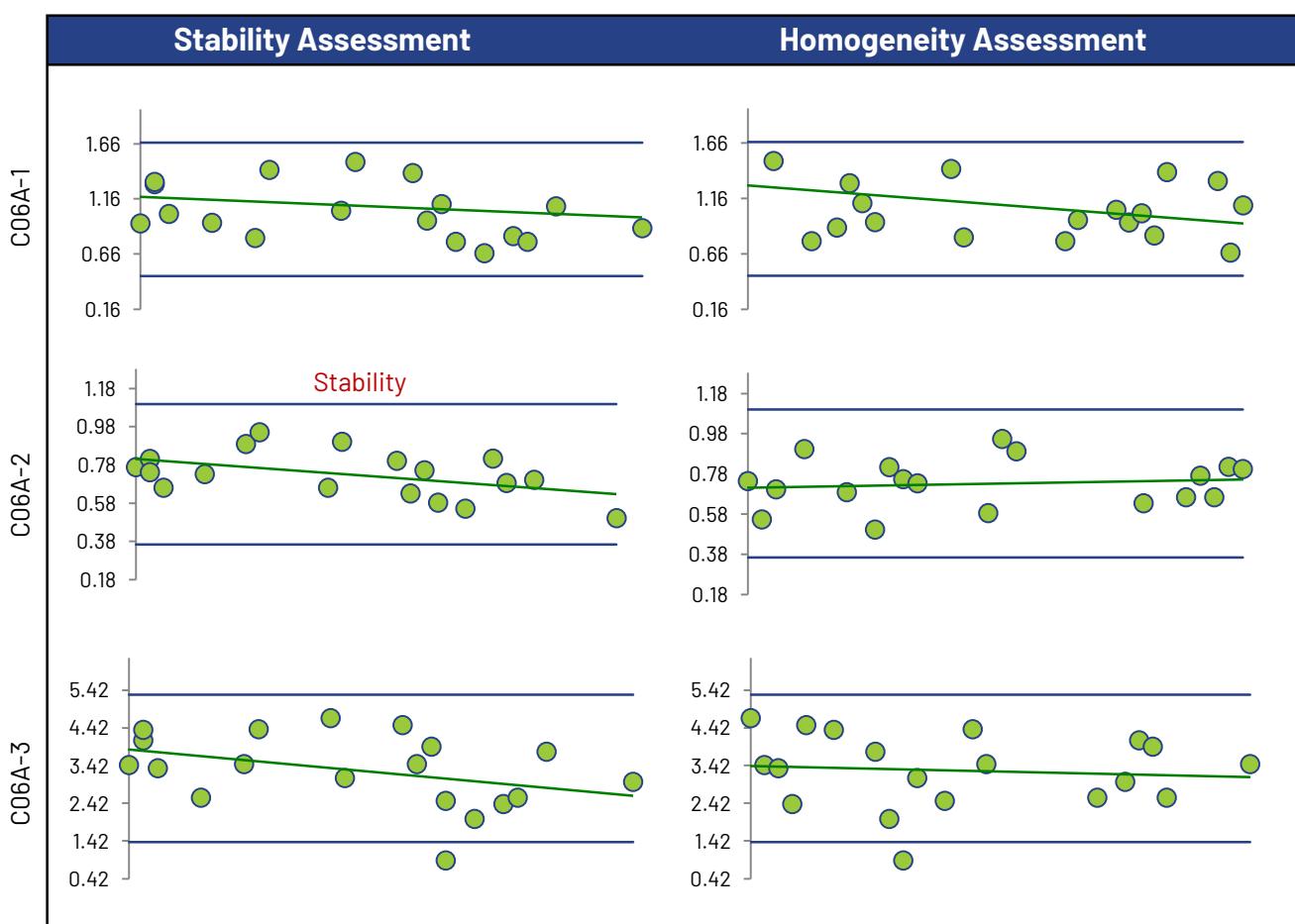
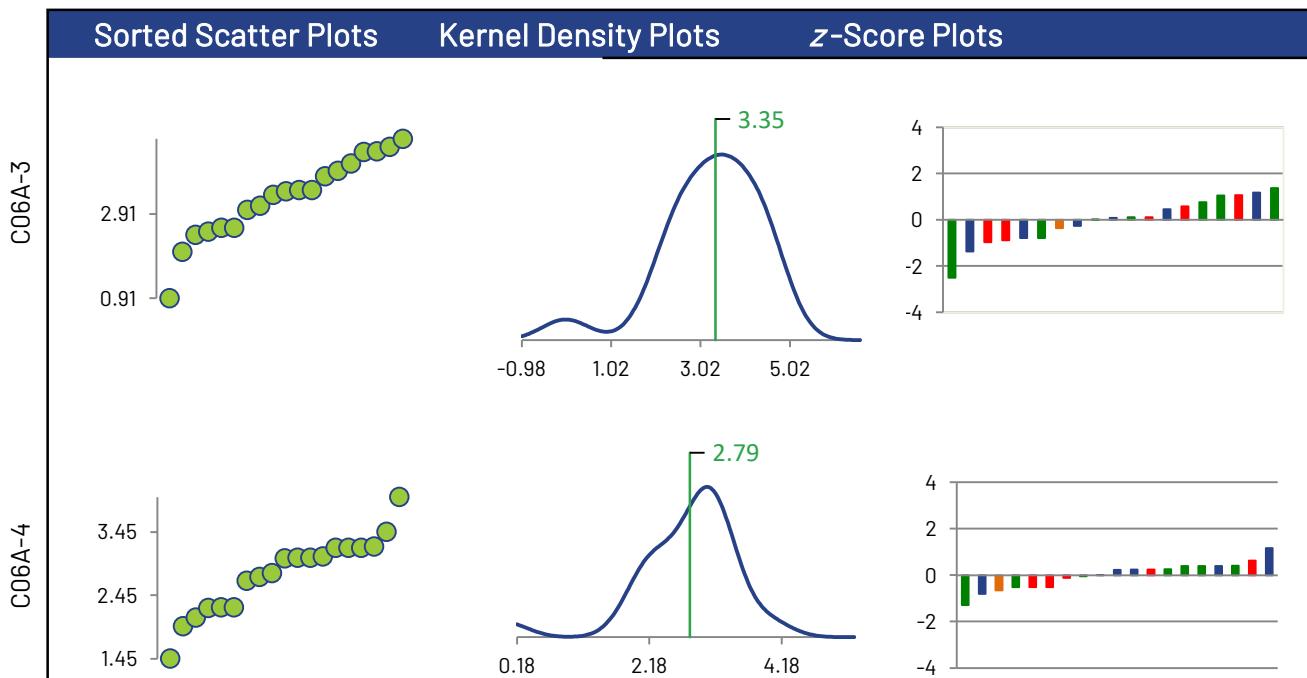
Method	C06A-1	C06A-2	C06A-3	C06A-4
GC/MS(Blue)	6	6	6	6
GC/MS/MS(Red)	5	5	5	5
GC/ECD(Green)	7	6	7	7
GCXGC/ECD(Orange)	1	1	1	1

All summary stats, except the failure rate, and the plots below are based on the data excluding any flagged outliers



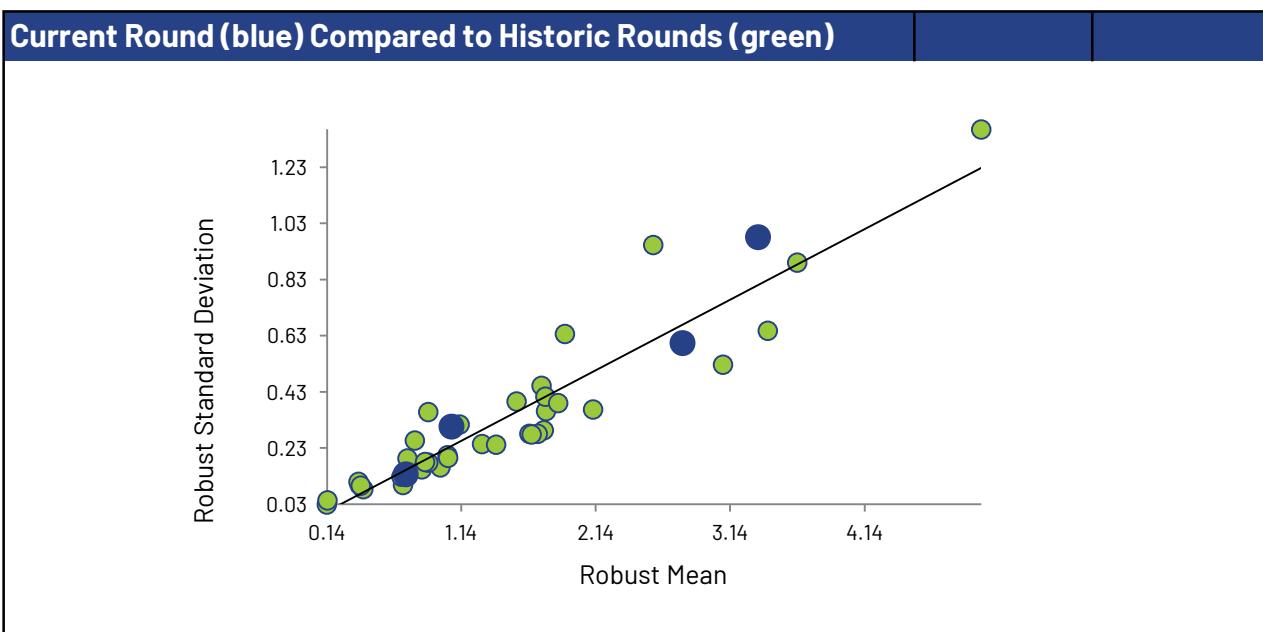
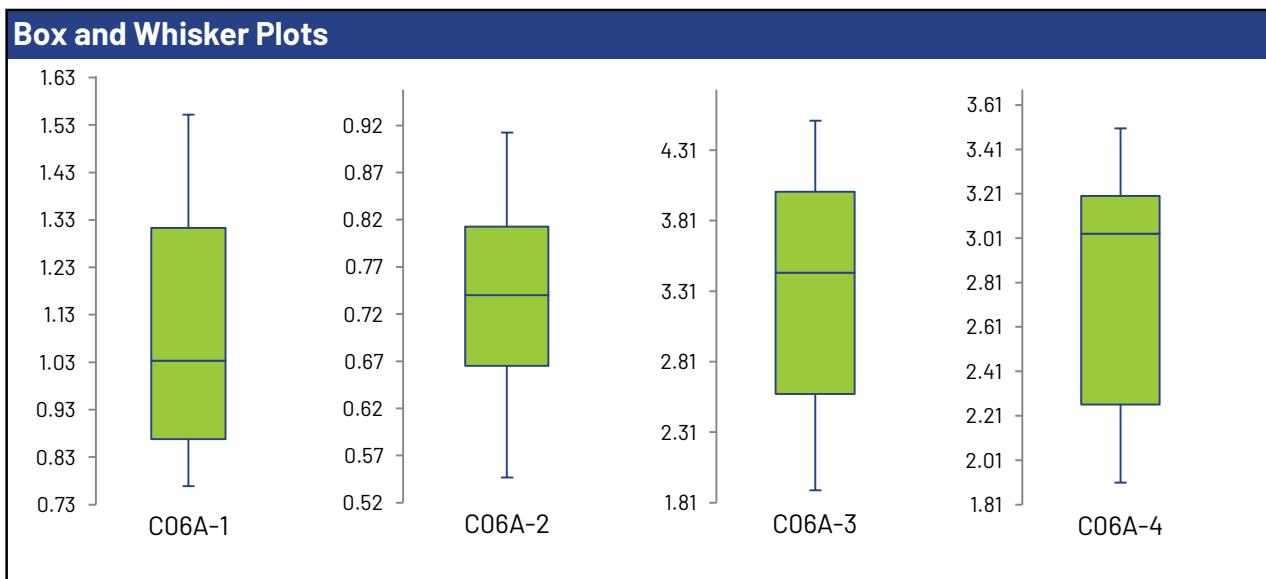
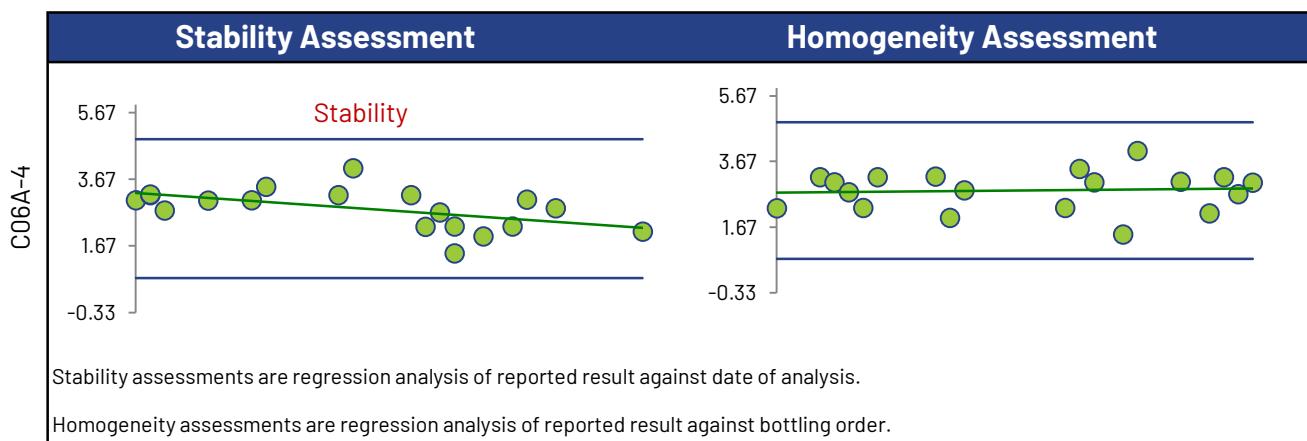
* Displayed failure rate does not account for the RDL modified z-Scores or inaccurately reported non-detects.

Dieldrin



Annex A Summary by Analyte

Dieldrin



Endosulfan I

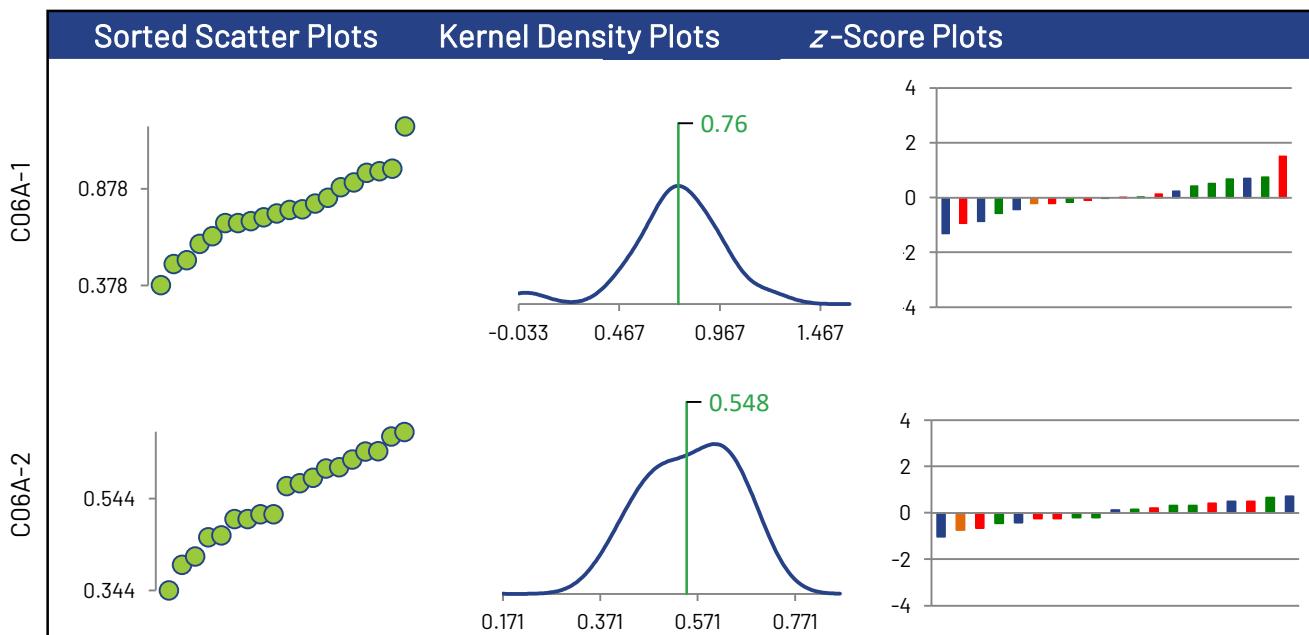
Summary Statistics

Statistic	C06A-1	C06A-2	C06A-3	C06A-4
No. of results used for summary statistics	20	19	20	20
Median µg/L	0.759	0.572	2.02	1.04
Robust Mean µg/L	0.760	0.548	2.03	1.06
Uncertainty of the Assigned Value µg/L	0.0531	0.0310	0.141	0.0352
Robust Standard Deviation µg/L	0.190	0.108	0.506	0.126
Regression Standard Deviation µg/L	0.199	0.148	0.507	0.271
Stability Flag	Stability	Stability		
Homogeneity Flag				
Standard Deviation Used (SDPA) µg/L	0.291	0.200	0.507	0.271
Flagged Outliers	0	1	0	0
$ z > 3.0$	0	0	0	0
$2 < z < 3$	0	0	1	0
Failure Rate %*		5.0		

Methods Used

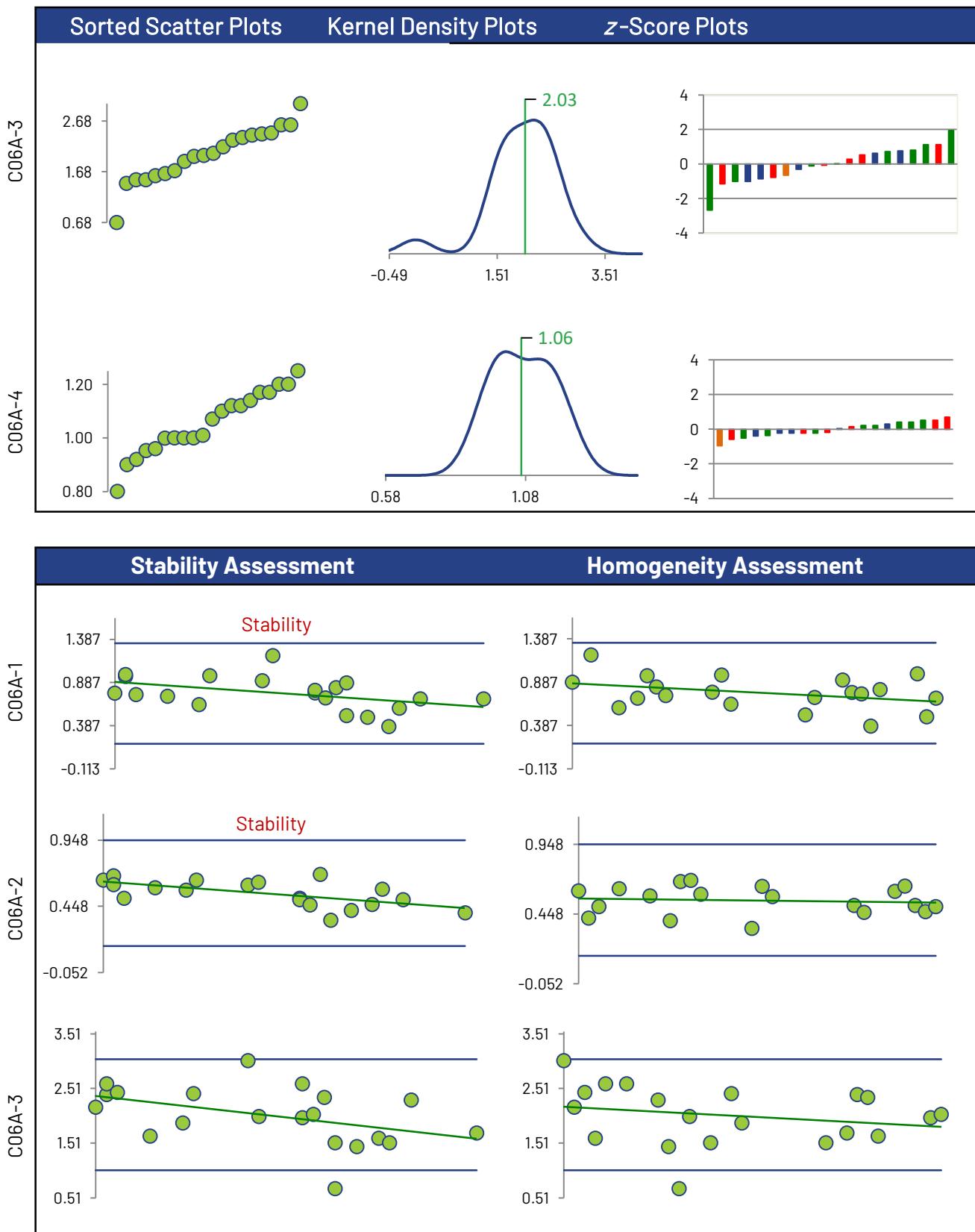
Method	C06A-1	C06A-2	C06A-3	C06A-4
GC/MS/MS (Blue)	5	5	5	5
GC/MS (Red)	6	6	6	6
GC/ECD (Green)	8	7	8	8
GCXGC/ECD (Orange)	1	1	1	1

All summary stats, except the failure rate, and the plots below are based on the data excluding any flagged outliers



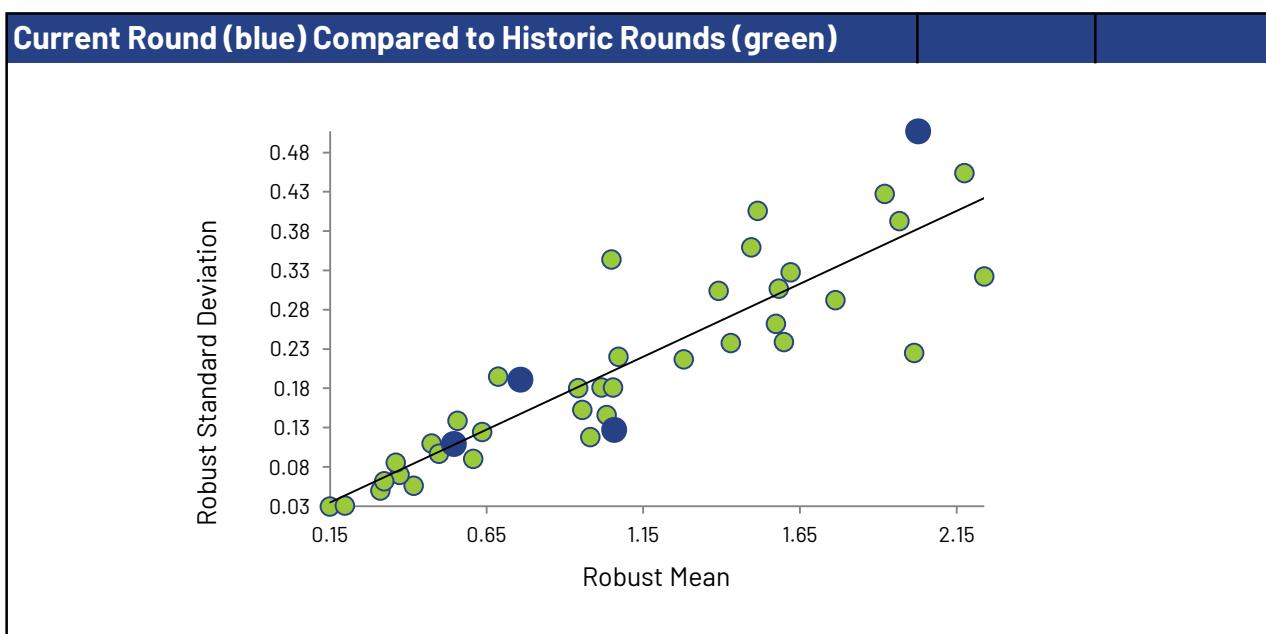
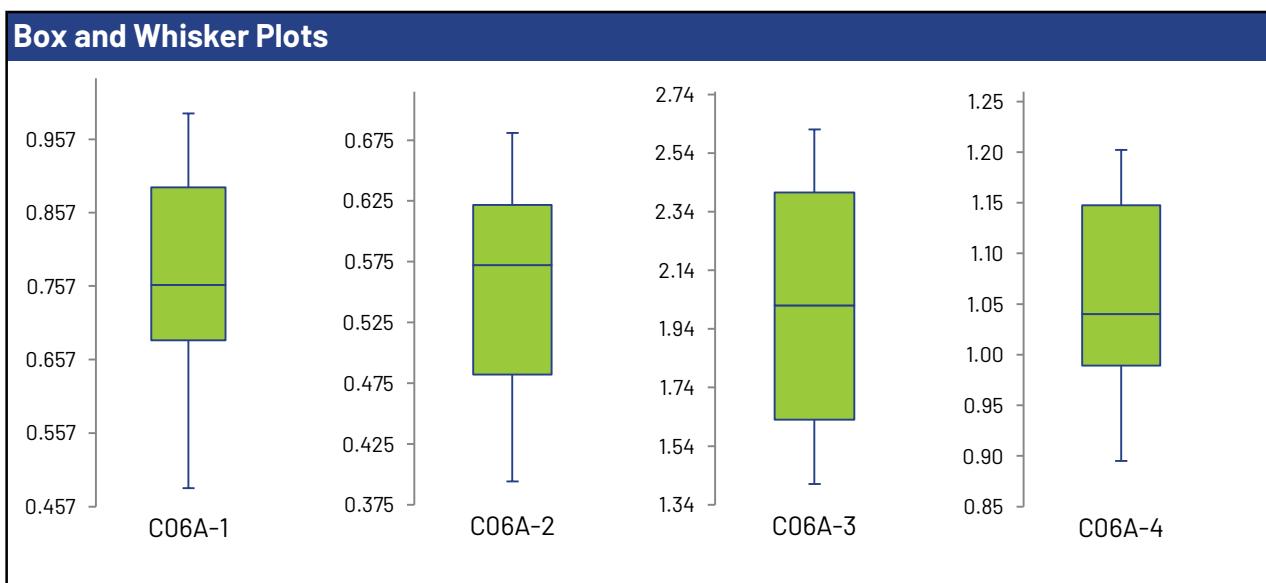
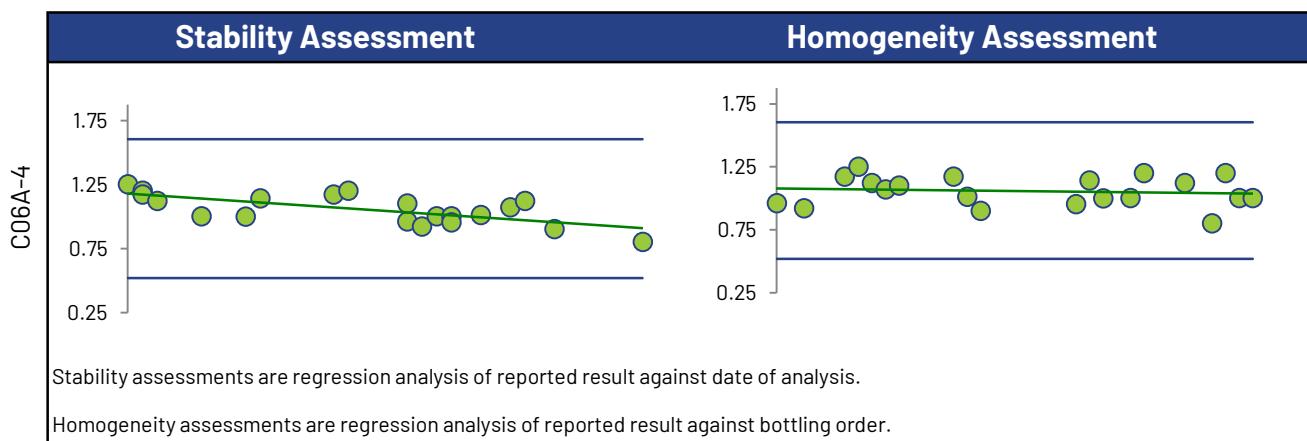
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Endosulfan I



Annex A Summary by Analyte

Endosulfan I



Annex A Summary by Analyte

Endosulfan II

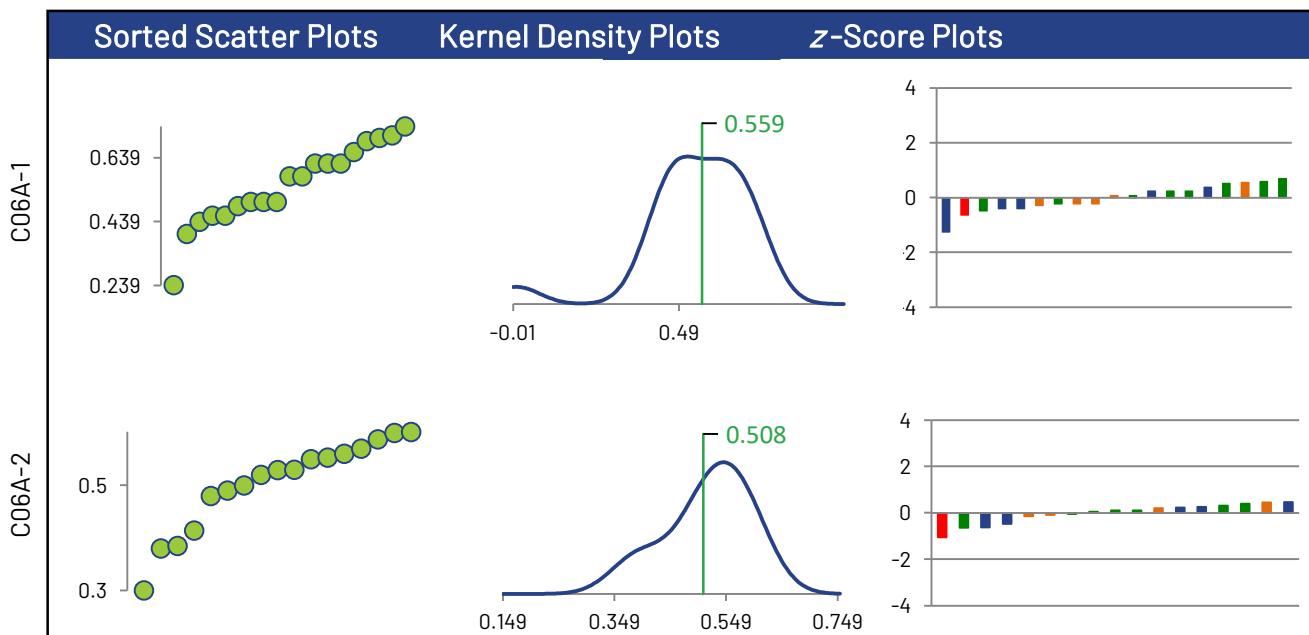
Summary Statistics

Statistic	C06A-1	C06A-2	C06A-3	C06A-4
No. of results used for summary statistics	19	17	19	19
Median µg/L	0.580	0.529	1.85	0.910
Robust Mean µg/L	0.559	0.508	1.89	0.866
Uncertainty of the Assigned Value µg/L	0.0364	0.0263	0.163	0.0465
Robust Standard Deviation µg/L	0.127	0.0868	0.569	0.162
Regression Standard Deviation µg/L	0.159	0.149	0.429	0.221
Stability Flag	Stability	Stability	Stability	Stability
Homogeneity Flag				
Standard Deviation Used (SDPA) µg/L	0.254	0.199	0.970	0.314
Flagged Outliers	0	1	0	0
$ z > 3.0$	0	0	0	0
$2 < z < 3$	0	0	0	0
Failure Rate %*		5.3		

Methods Used

Method	C06A-1	C06A-2	C06A-3	C06A-4
GC/MS/MS (Blue)	5	5	5	5
GCXGC/ECD (Red)	1	1	1	1
GC/ECD (Green)	8	7	8	8
GC/MS (Orange)	5	4	5	5

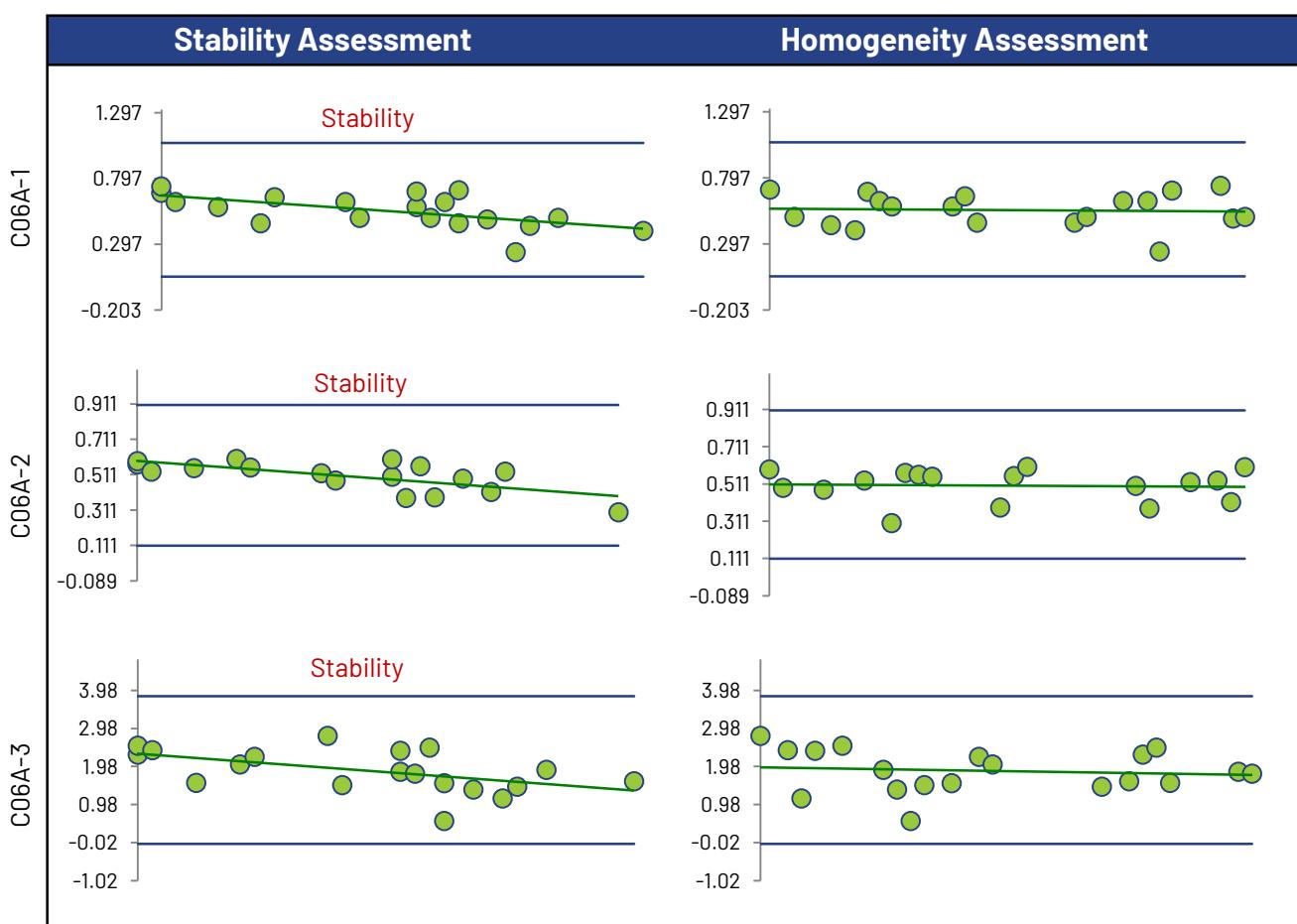
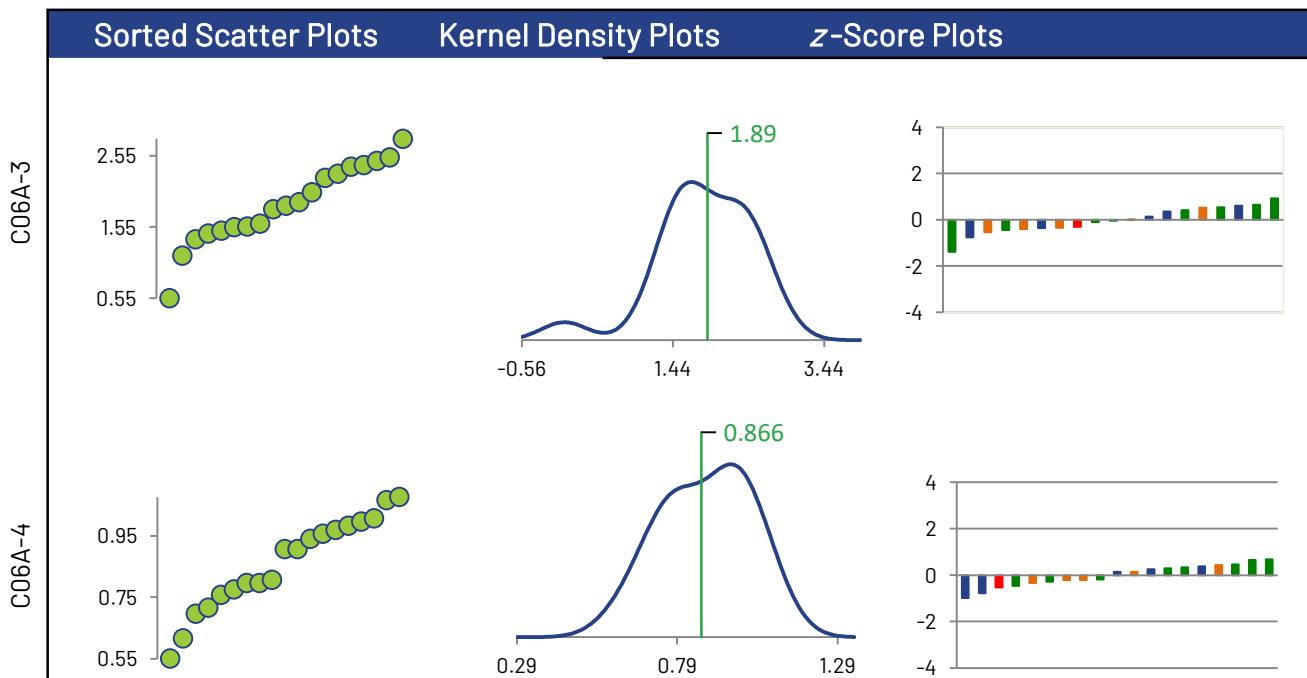
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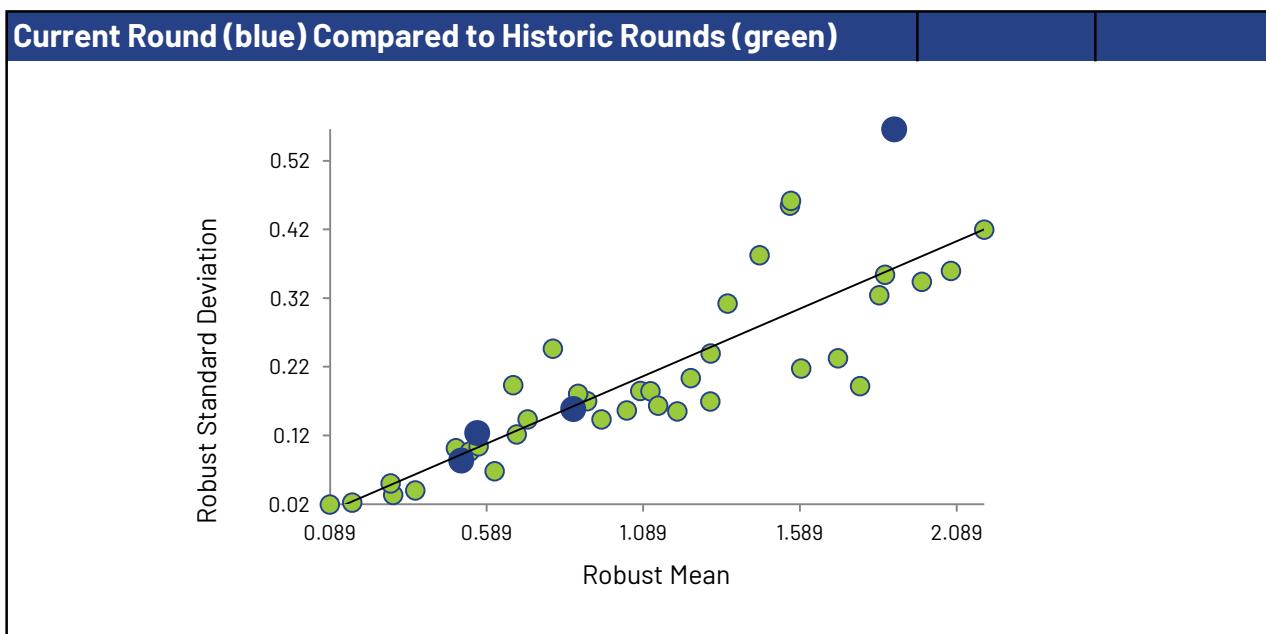
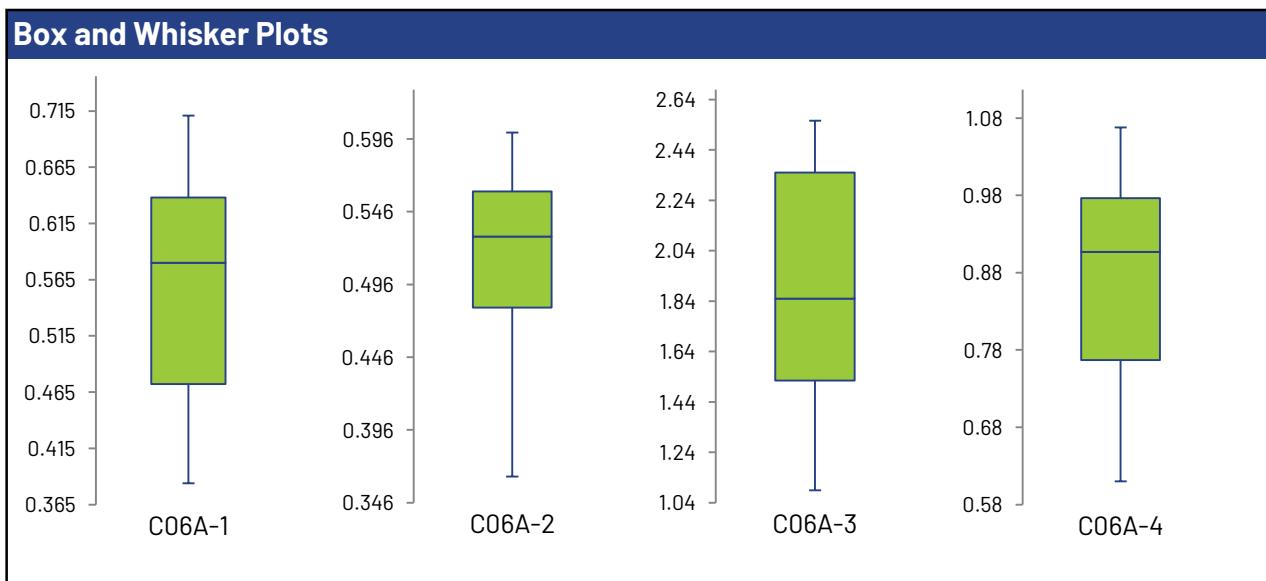
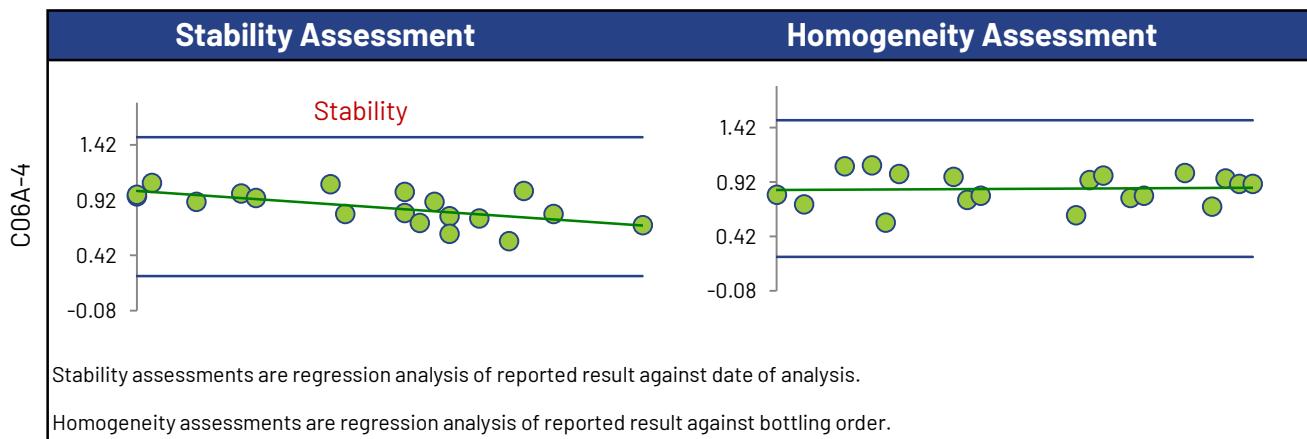
Annex A Summary by Analyte

Endosulfan II



Annex A Summary by Analyte

Endosulfan II



Annex A Summary by Analyte

Endrin

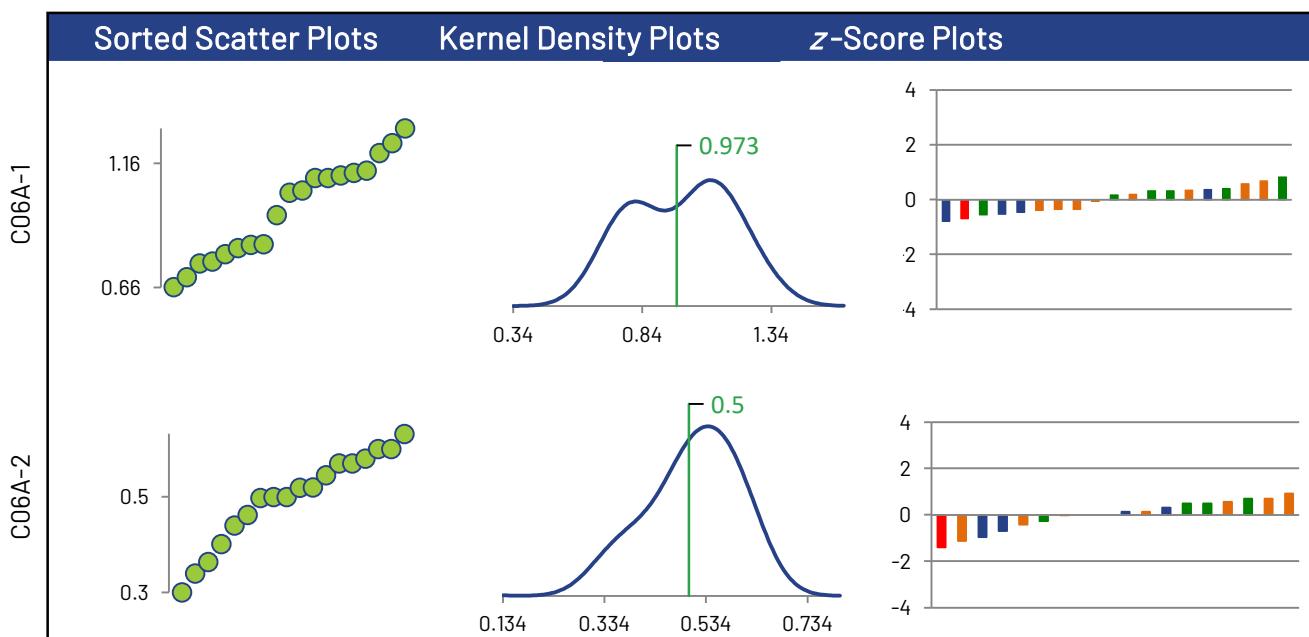
Summary Statistics

Statistic	C06A-1	C06A-2	C06A-3	C06A-4
No. of results used for summary statistics	19	18	19	19
Median µg/L	1.04	0.510	2.04	1.04
Robust Mean µg/L	0.973	0.500	1.97	1.00
Uncertainty of the Assigned Value µg/L	0.0639	0.0301	0.181	0.0602
Robust Standard Deviation µg/L	0.223	0.102	0.632	0.210
Regression Standard Deviation µg/L	0.246	0.143	0.461	0.252
Stability Flag	Stability			
Homogeneity Flag				
Standard Deviation Used (SDPA) µg/L	0.394	0.143	0.632	0.252
Flagged Outliers	0	1	0	0
$ z > 3.0$	0	0	1	0
$2 < z < 3$	0	0	1	1
Failure Rate %*		5.3		

Methods Used

Method	C06A-1	C06A-2	C06A-3	C06A-4
GC/MS/MS (Blue)	4	4	4	4
GCXGC/ECD (Red)	1	1	1	1
GC/MS (Green)	6	6	6	6
GC/ECD (Orange)	8	7	8	8

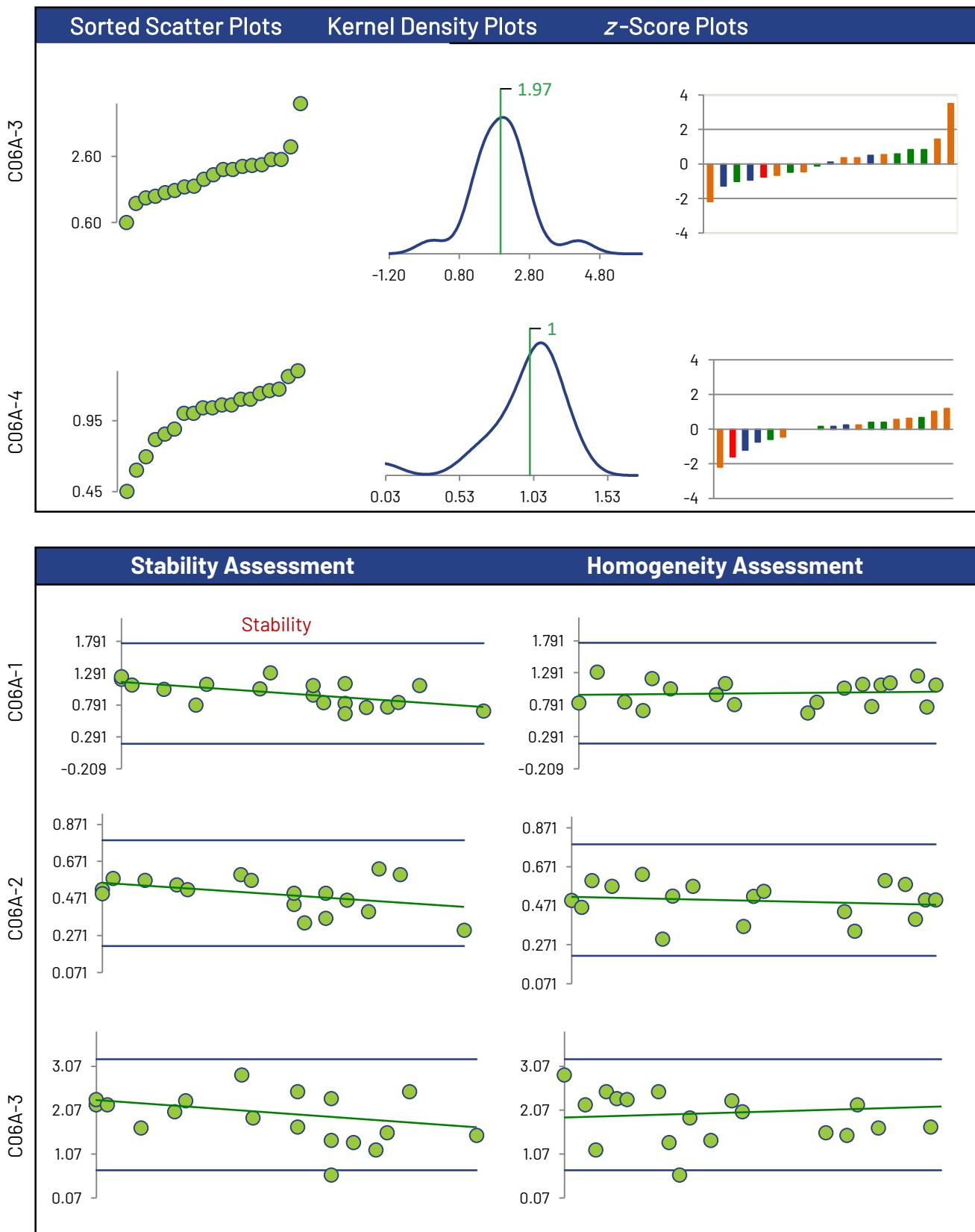
All summary stats, except the failure rate, and the plots below are based on the data excluding any flagged outliers



* Displayed failure rate does not account for the RDL modified z -Scores or inaccurately reported non-detects.

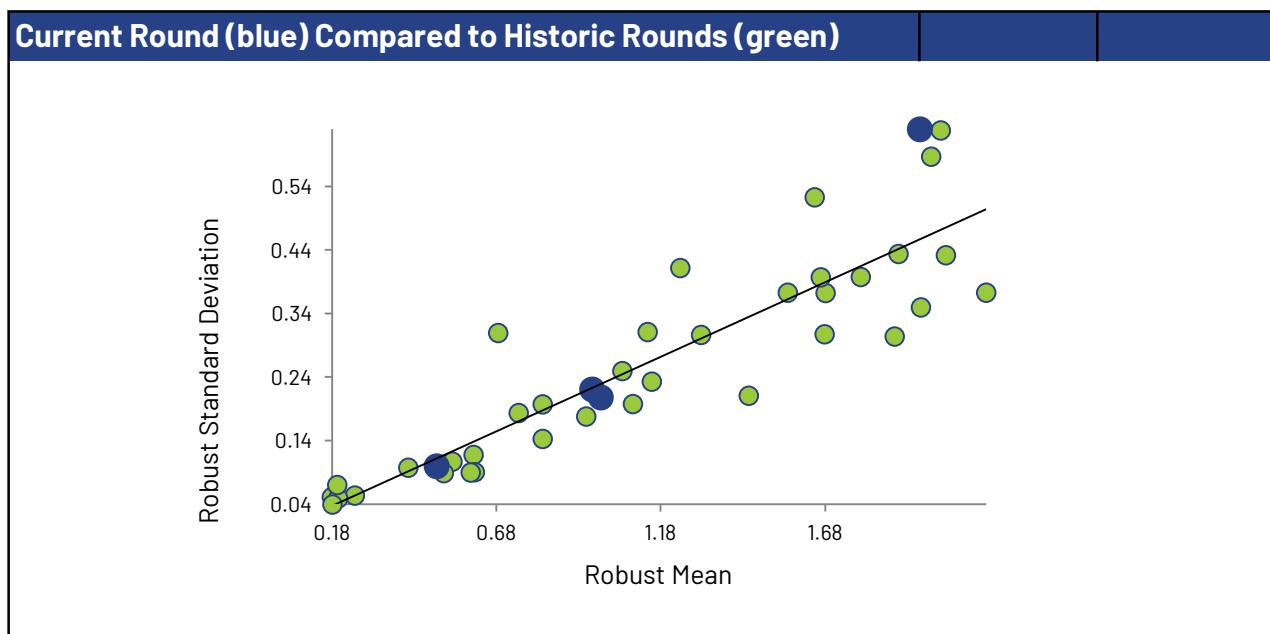
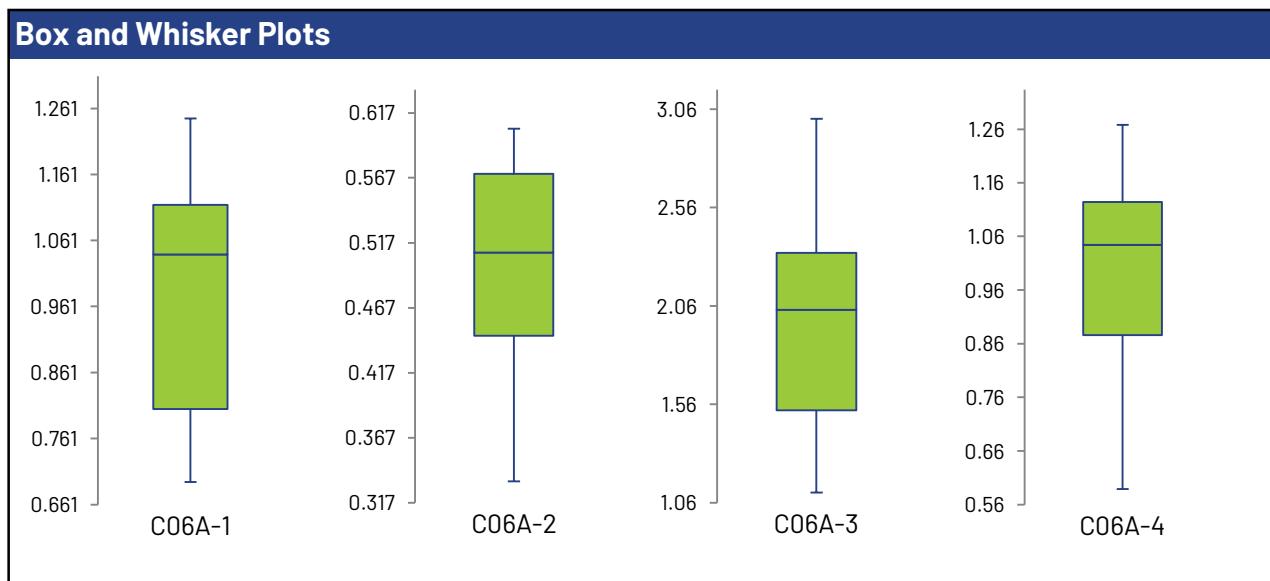
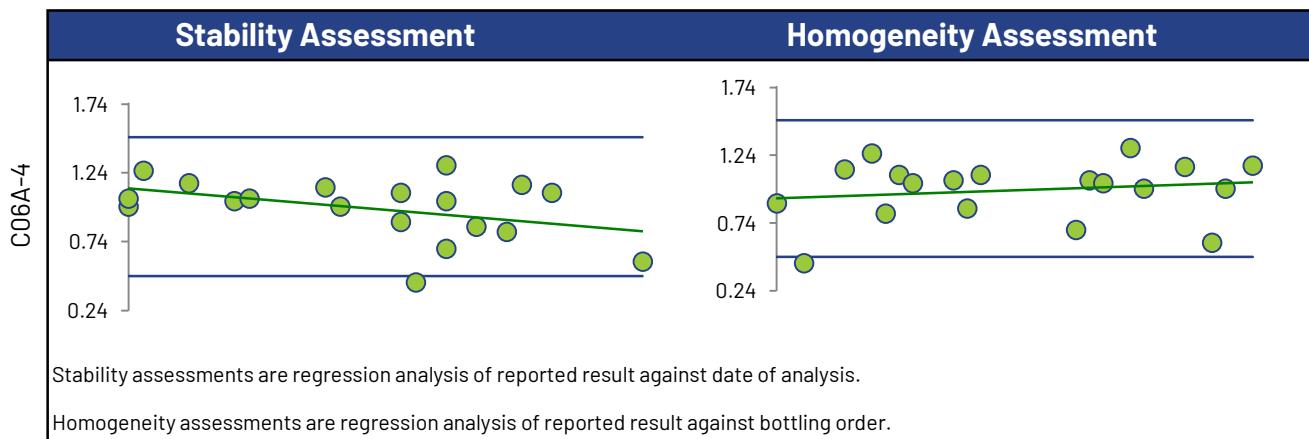
Annex A Summary by Analyte

Endrin



Annex A Summary by Analyte

Endrin



Annex A Summary by Analyte

g - Chlordane

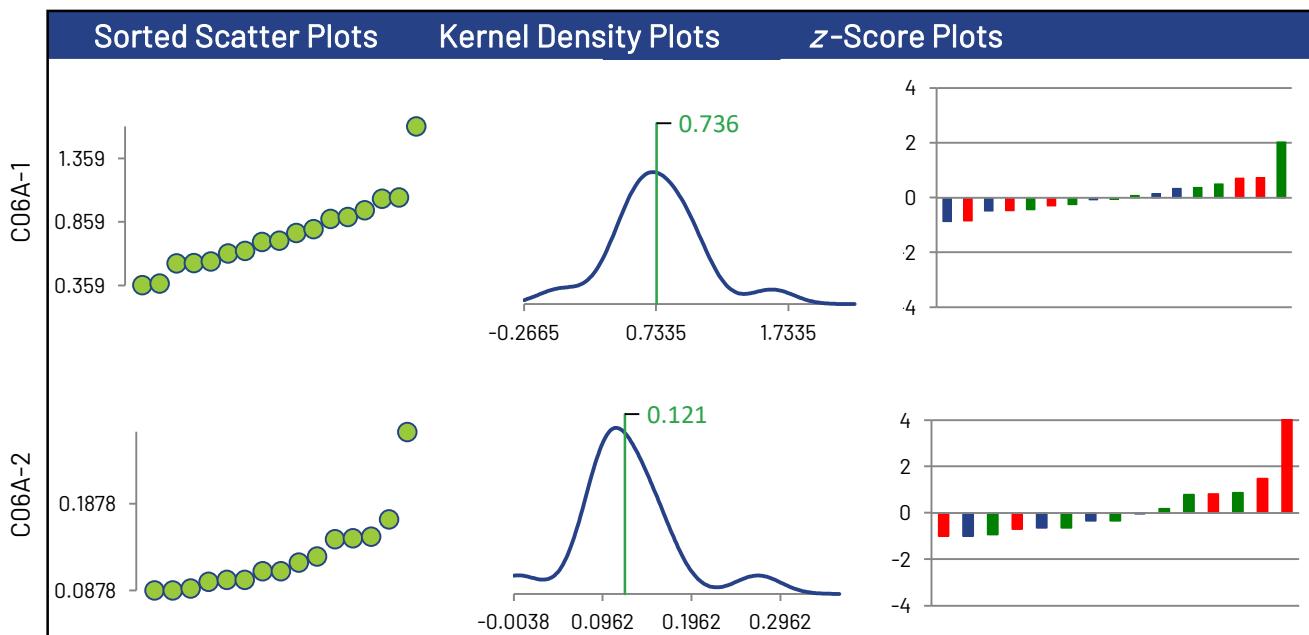
Summary Statistics

Statistic	C06A-1	C06A-2	C06A-3	C06A-4
No. of results used for summary statistics	17	15	17	17
Median µg/L	0.710	0.110	2.62	2.02
Robust Mean µg/L	0.736	0.121	2.63	1.98
Uncertainty of the Assigned Value µg/L	0.0800	0.0107	0.327	0.146
Robust Standard Deviation µg/L	0.264	0.0332	1.08	0.483
Regression Standard Deviation µg/L	0.128	0.0229	0.453	0.341
Stability Flag				
Homogeneity Flag	Homogeneity			
Standard Deviation Used (SDPA) µg/L	0.431	0.0332	1.08	0.483
Flagged Outliers	0	1	0	0
$ z > 3.0$	0	1	0	0
$2 < z < 3$	1	0	1	0
Failure Rate %*		5.9		

Methods Used

Method	C06A-1	C06A-2	C06A-3	C06A-4
GC/MS (Blue)	5	4	5	5
GC/MS/MS (Red)	5	5	5	5
GC/ECD (Green)	7	6	7	7

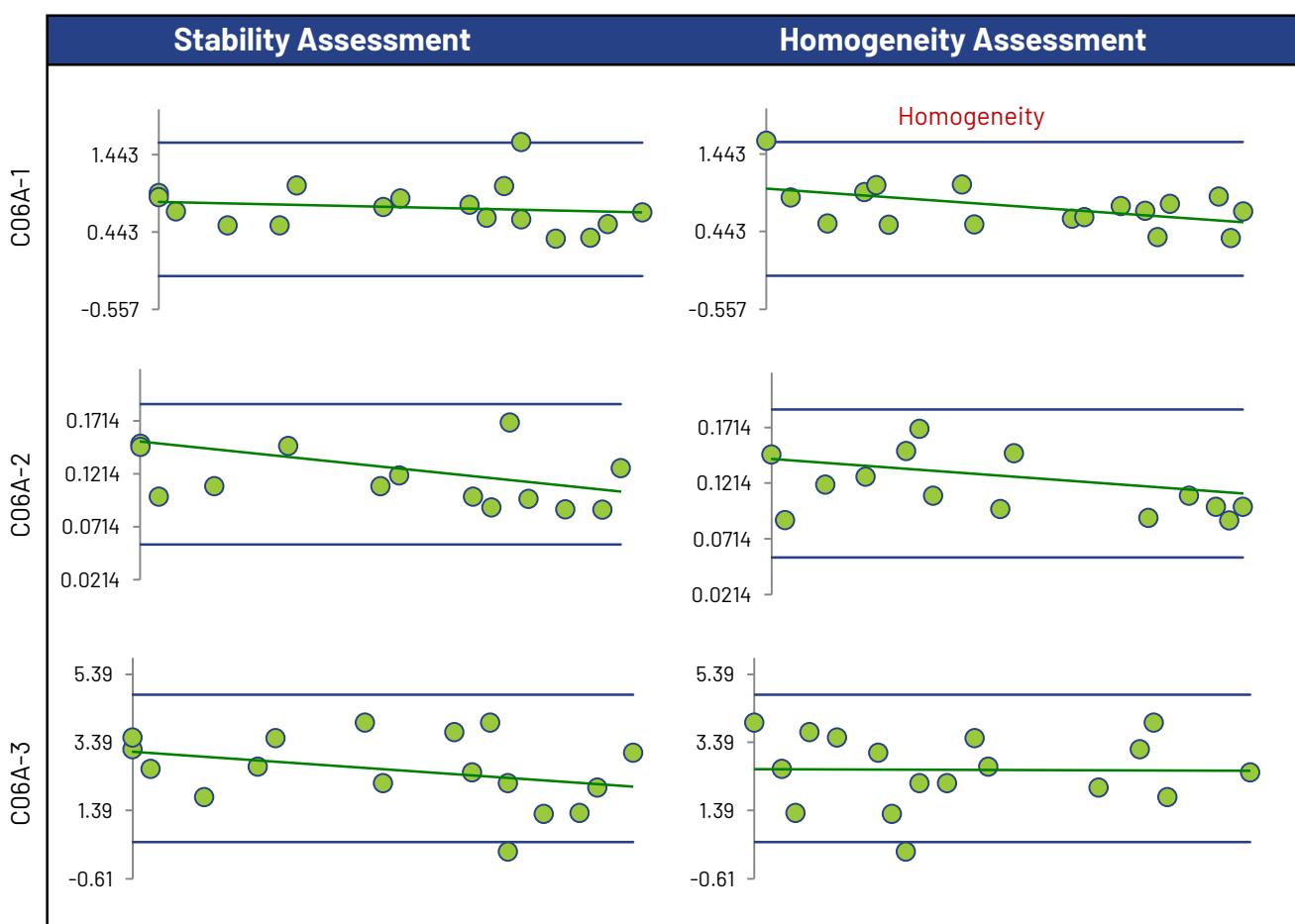
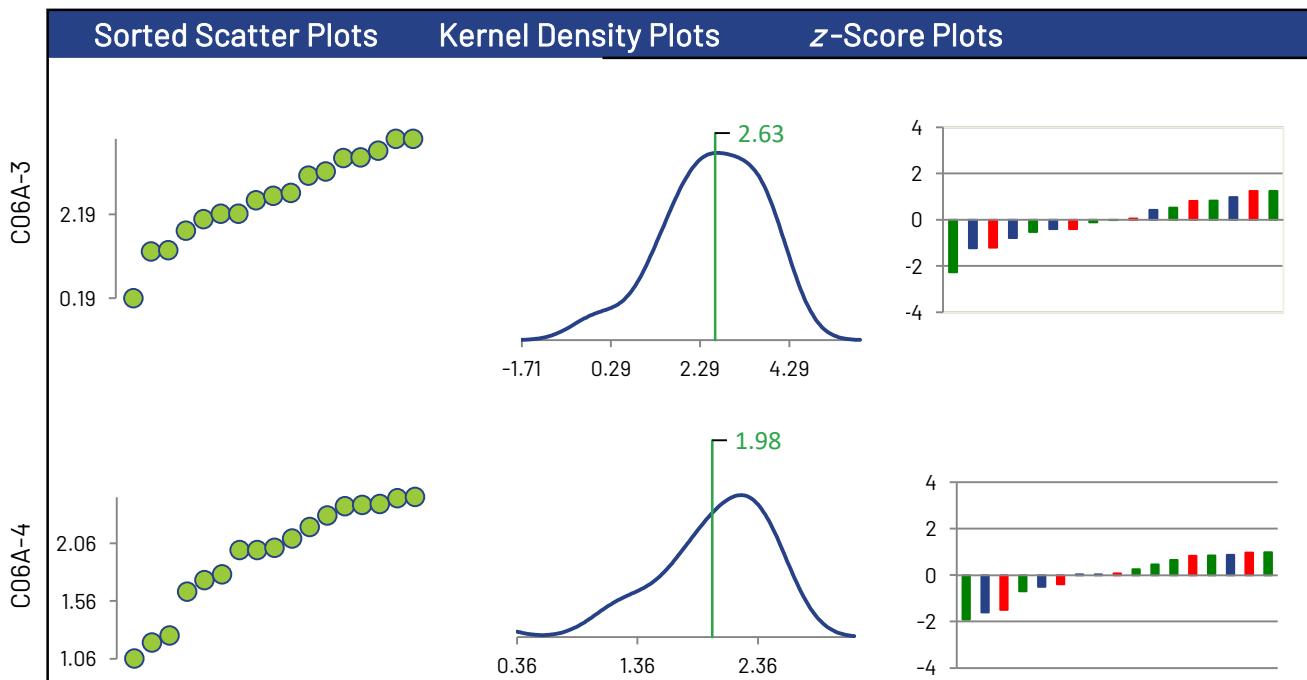
All summary stats, except the failure rate, and the plots below are based on the data excluding any flagged outliers



* Displayed failure rate does not account for the RDL modified z-Scores or inaccurately reported non-detects.

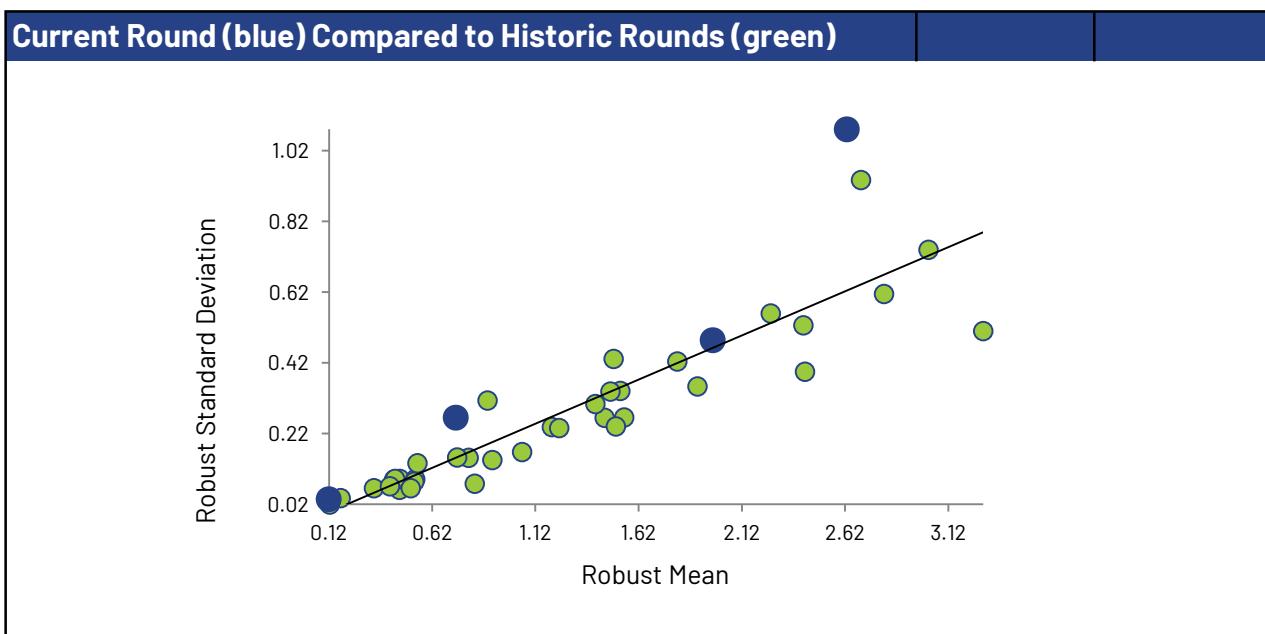
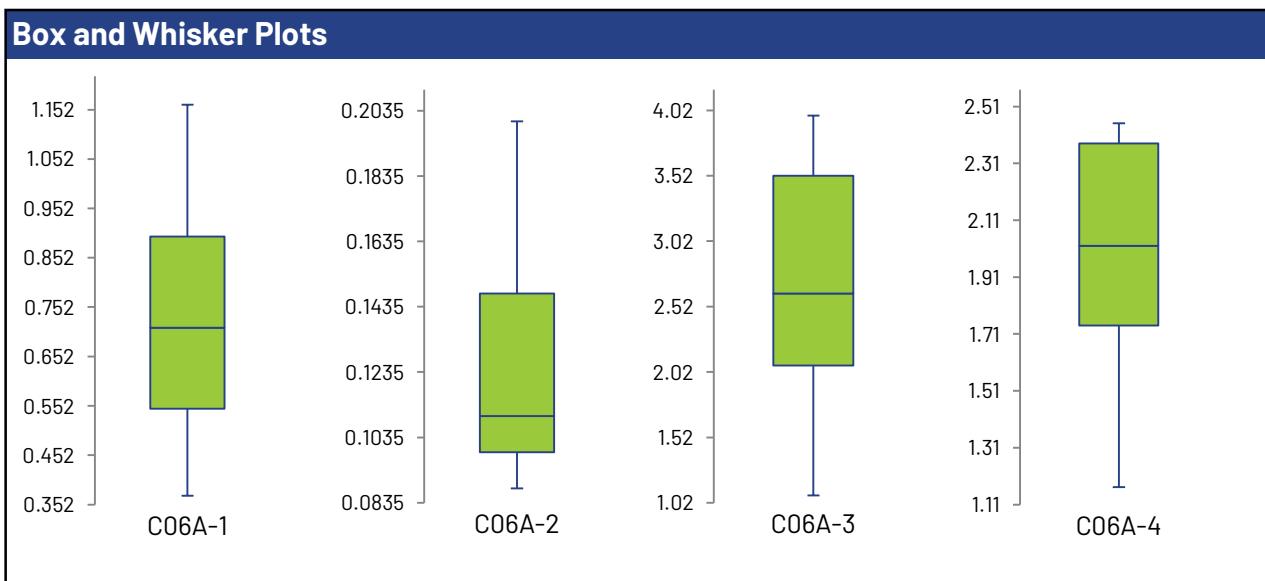
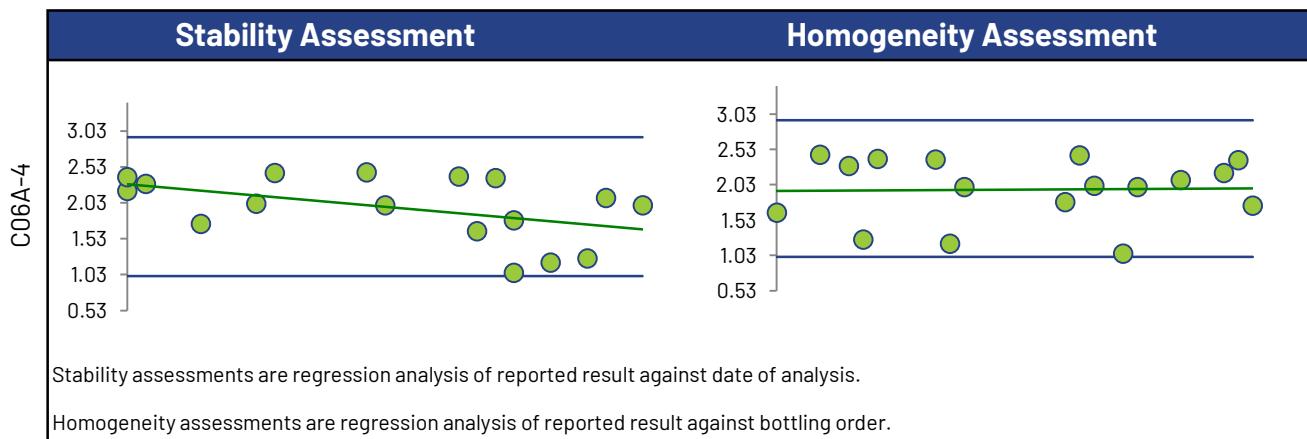
Annex A Summary by Analyte

g - Chlordane



Annex A Summary by Analyte

g - Chlordane



Heptachlor

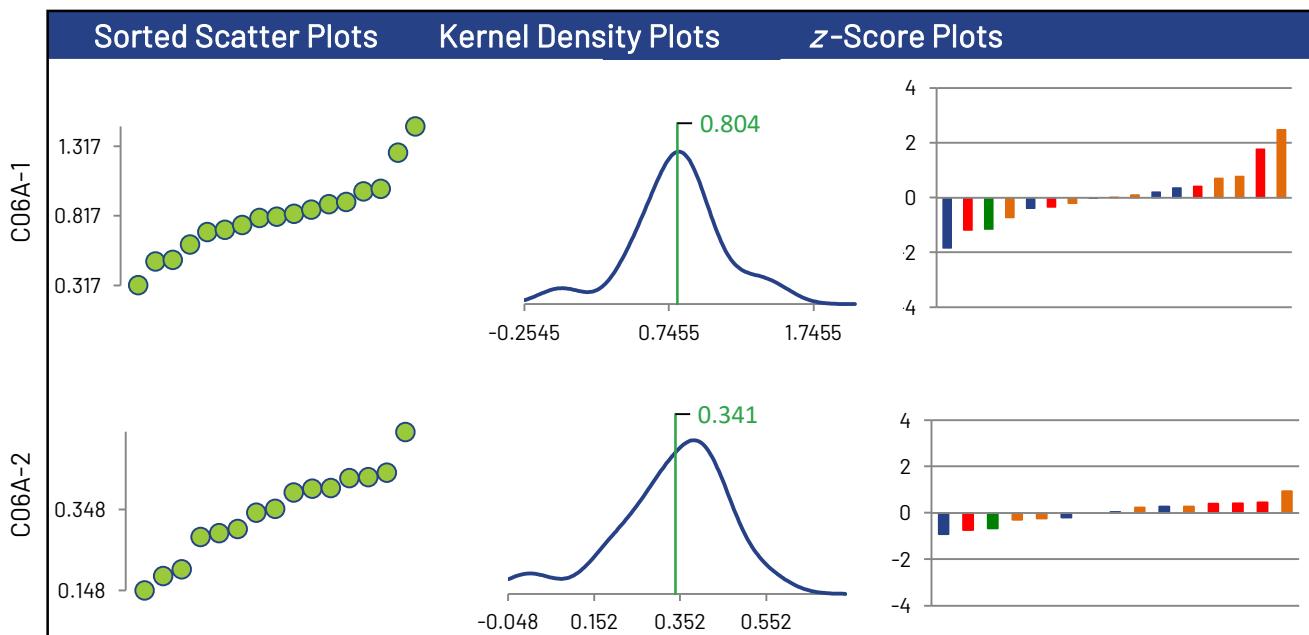
Summary Statistics

Statistic	C06A-1	C06A-2	C06A-3	C06A-4
No. of results used for summary statistics	17	15	17	17
Median µg/L	0.810	0.350	3.90	1.20
Robust Mean µg/L	0.804	0.341	3.47	1.20
Uncertainty of the Assigned Value µg/L	0.0755	0.0374	0.376	0.101
Robust Standard Deviation µg/L	0.249	0.116	1.24	0.332
Regression Standard Deviation µg/L	0.264	0.112	1.14	0.394
Stability Flag		Stability	Stability	
Homogeneity Flag				
Standard Deviation Used (SDPA) µg/L	0.264	0.211	2.21	0.394
Flagged Outliers	0	1	0	0
z >3.0	0	0	0	0
2< z <3	1	0	0	1
Failure Rate %*		5.9		

Methods Used

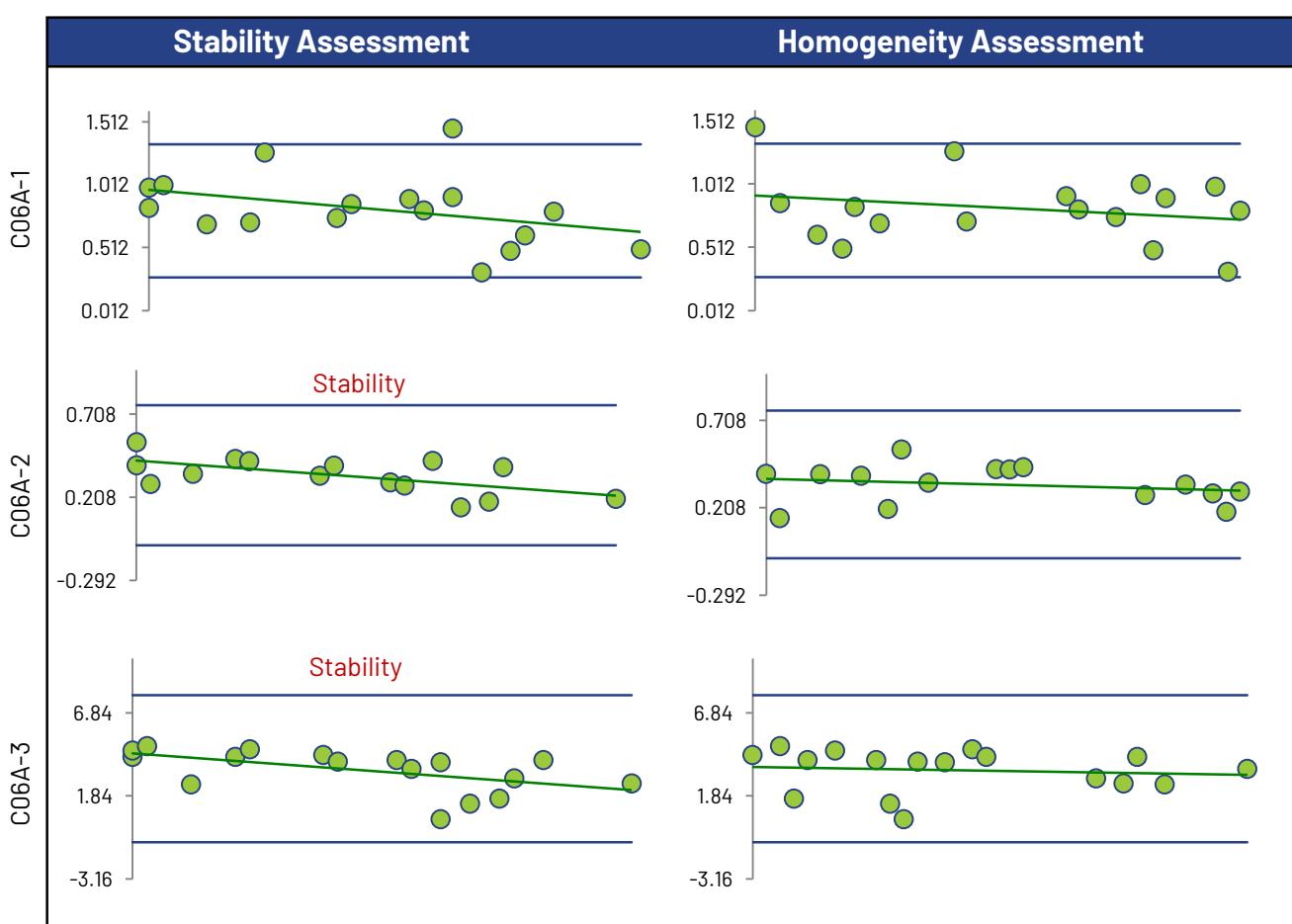
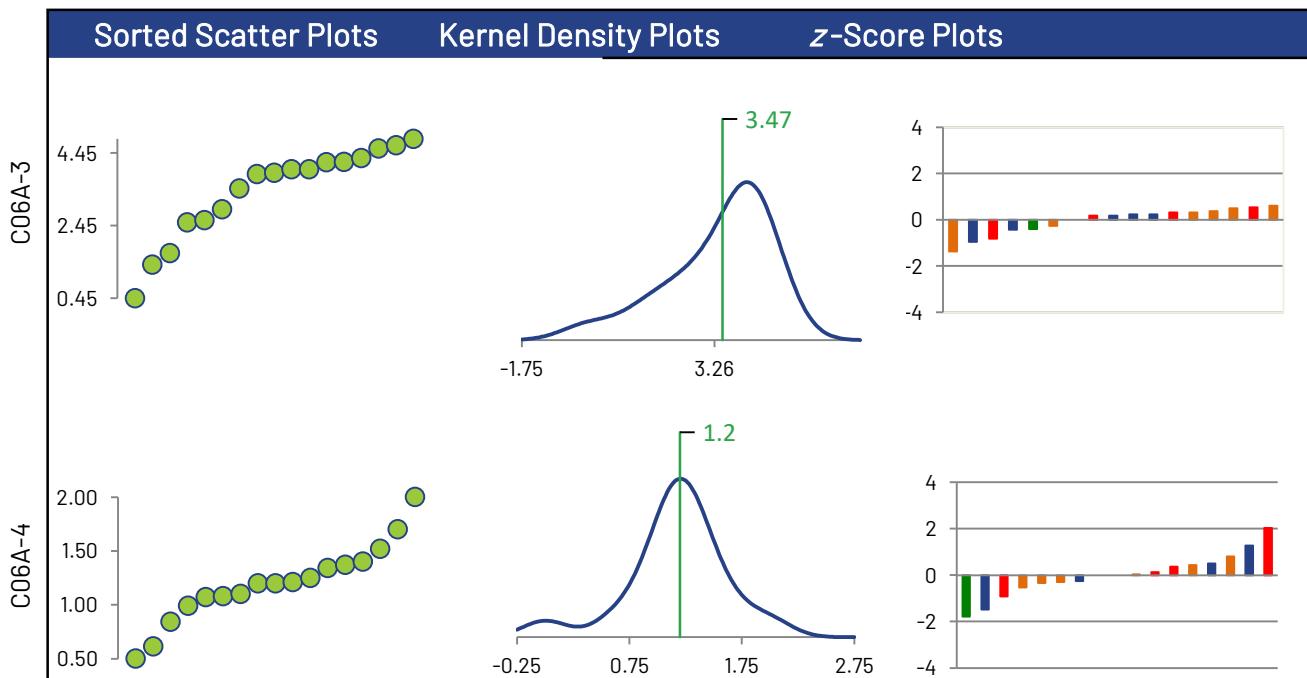
Method	C06A-1	C06A-2	C06A-3	C06A-4
GC/MS(Blue)	5	4	5	5
GC/MS/MS(Red)	4	4	4	4
GCXGC/ECD(Green)	1	1	1	1
GC/ECD(Orange)	7	6	7	7

All summary stats, except the failure rate, and the plots below are based on the data excluding any flagged outliers



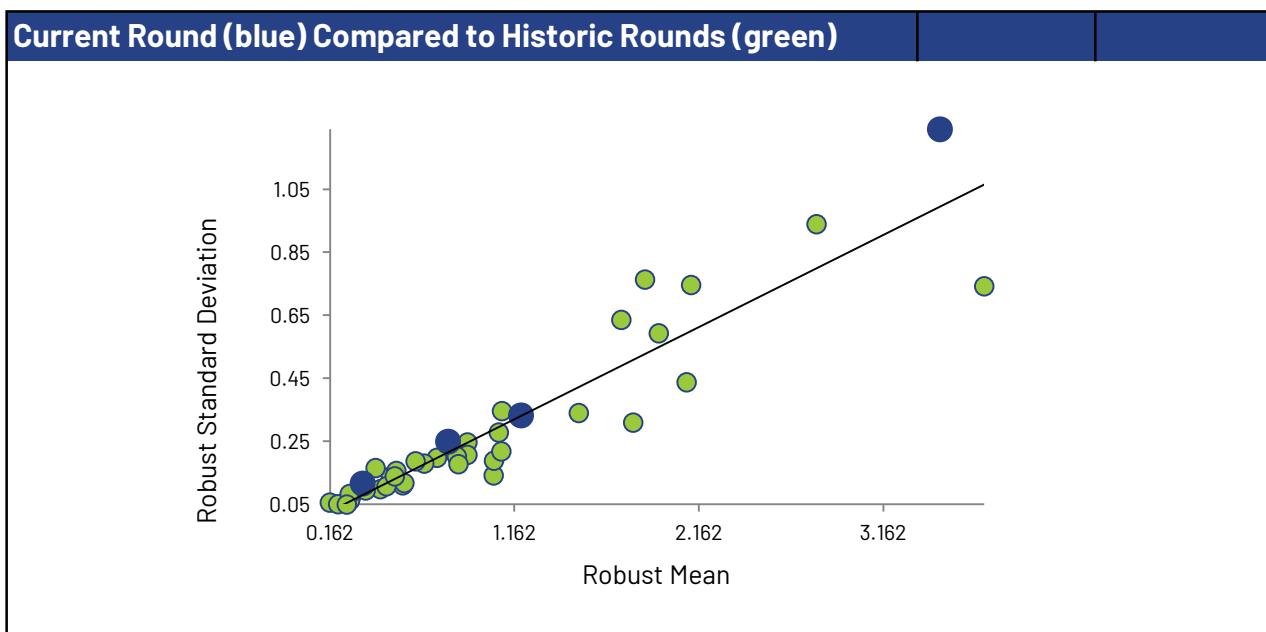
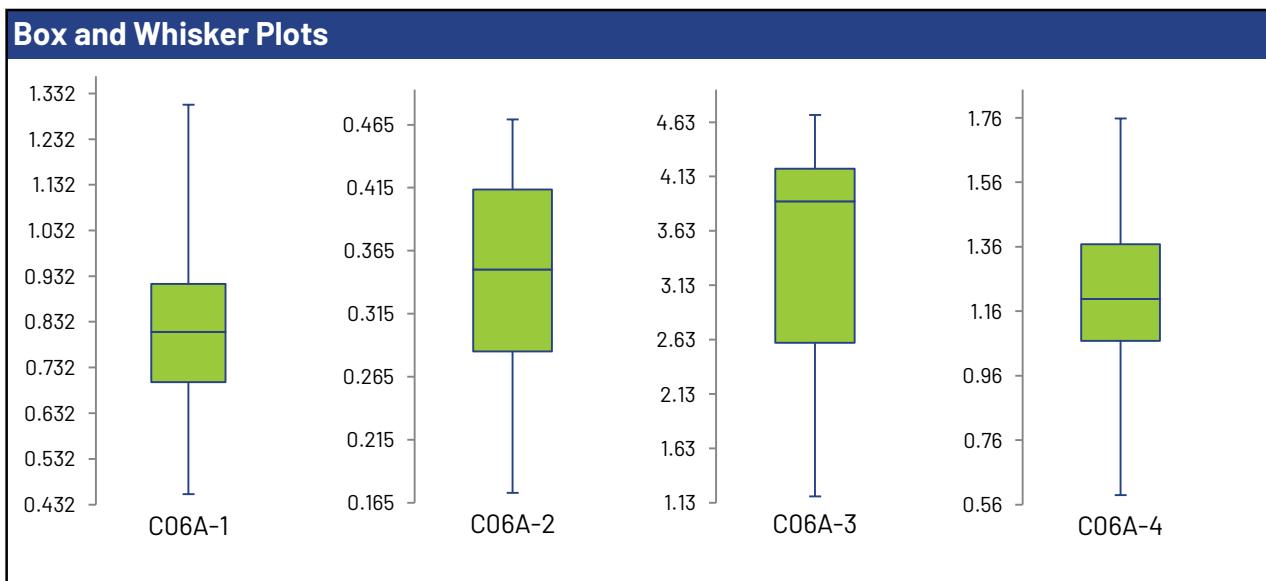
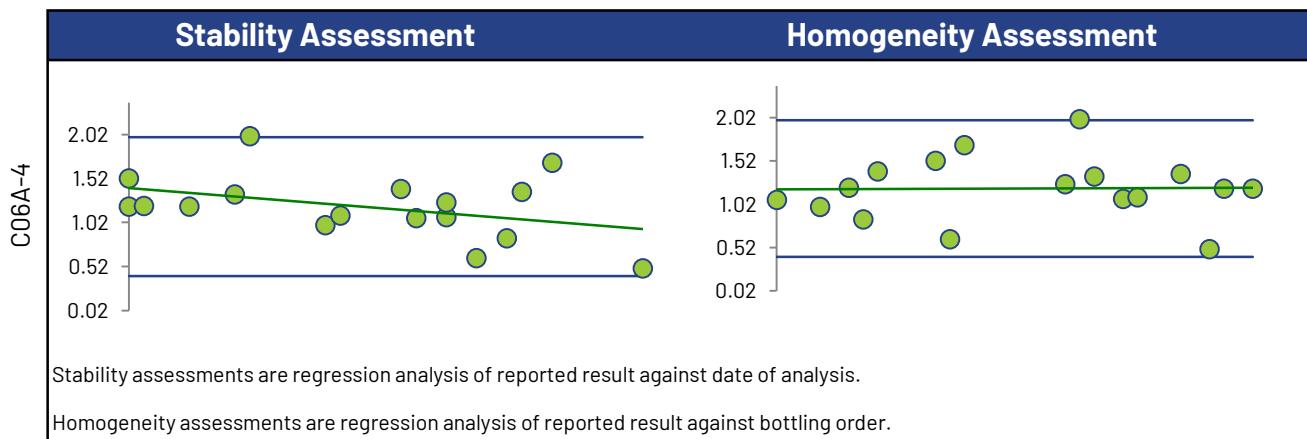
* Displayed failure rate does not account for the RDL modified z-Scores or inaccurately reported non-detects.

Heptachlor



Annex A Summary by Analyte

Heptachlor



Heptachlor epoxide

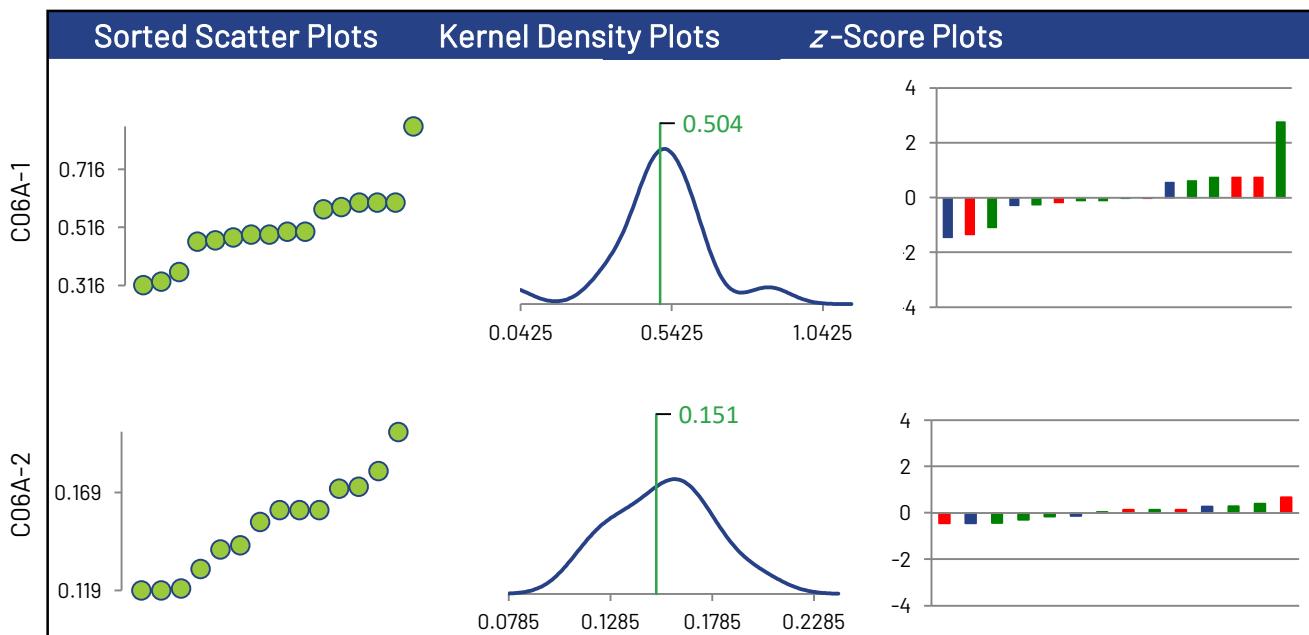
Summary Statistics

Statistic	C06A-1	C06A-2	C06A-3	C06A-4
No. of results used for summary statistics	16	14	16	16
Median µg/L	0.495	0.157	1.70	1.00
Robust Mean µg/L	0.504	0.151	1.66	0.956
Uncertainty of the Assigned Value µg/L	0.0356	0.00892	0.141	0.0444
Robust Standard Deviation µg/L	0.114	0.0267	0.451	0.142
Regression Standard Deviation µg/L	0.130	0.0719	0.319	0.204
Stability Flag				Stability
Homogeneity Flag				
Standard Deviation Used (SDPA) µg/L	0.130	0.0719	0.451	0.223
Flagged Outliers	0	1	0	0
z >3.0	0	0	1	0
2< z <3	1	0	0	0
Failure Rate %*			6.3	

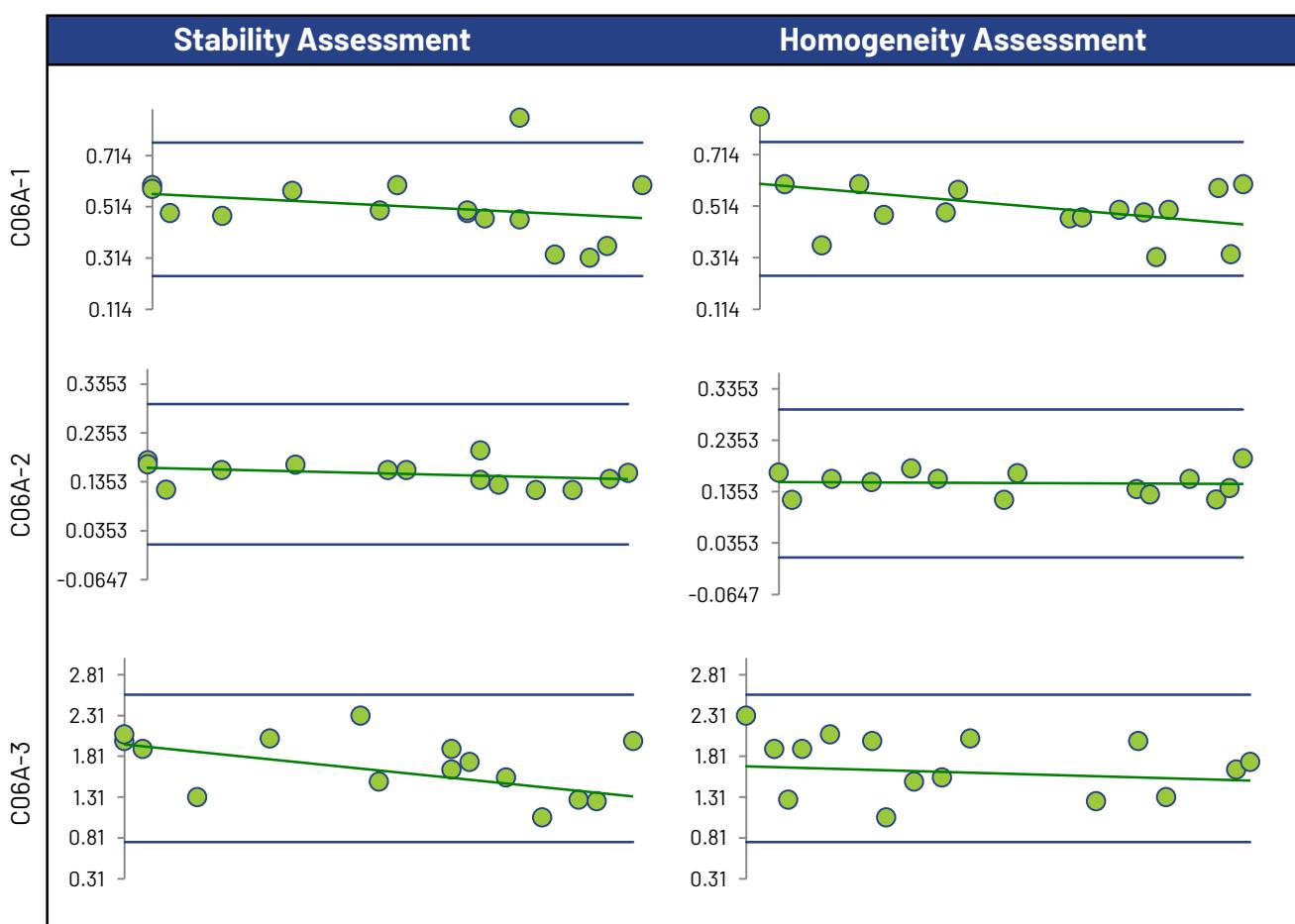
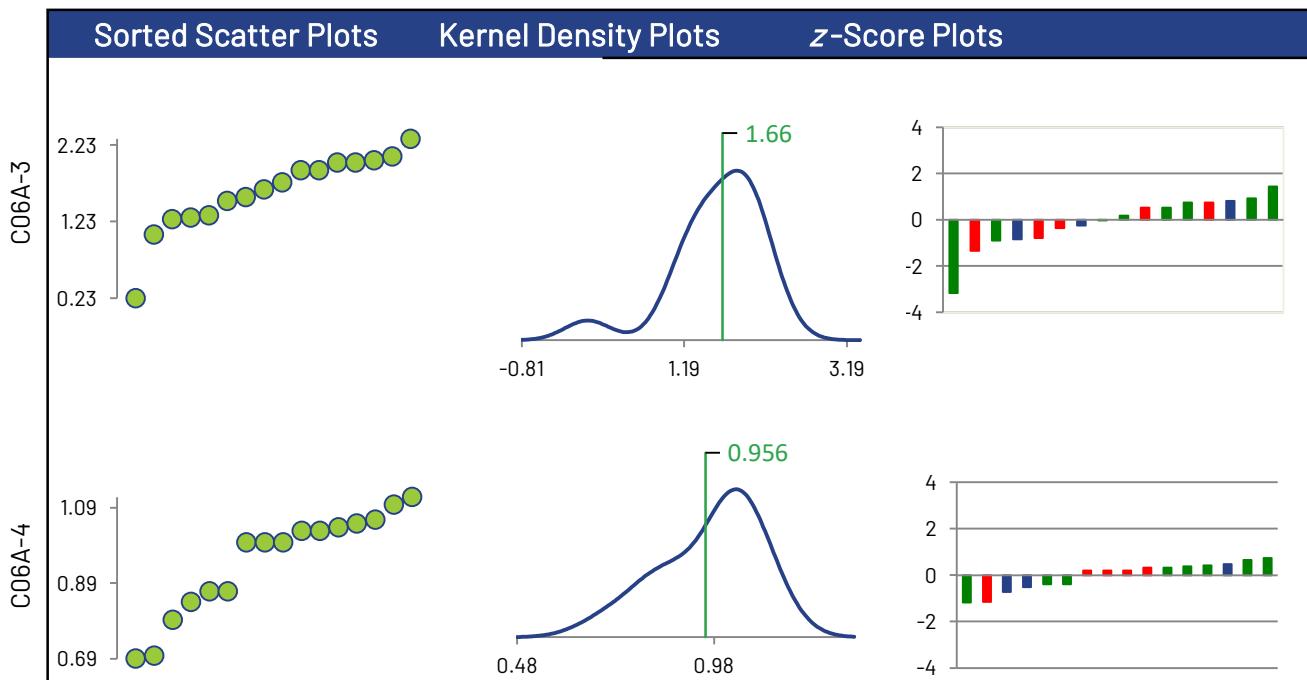
Methods Used

Method	C06A-1	C06A-2	C06A-3	C06A-4
GC/MS/MS (Blue)	3	3	3	3
GC/MS (Red)	5	4	5	5
GC/ECD (Green)	8	7	8	8

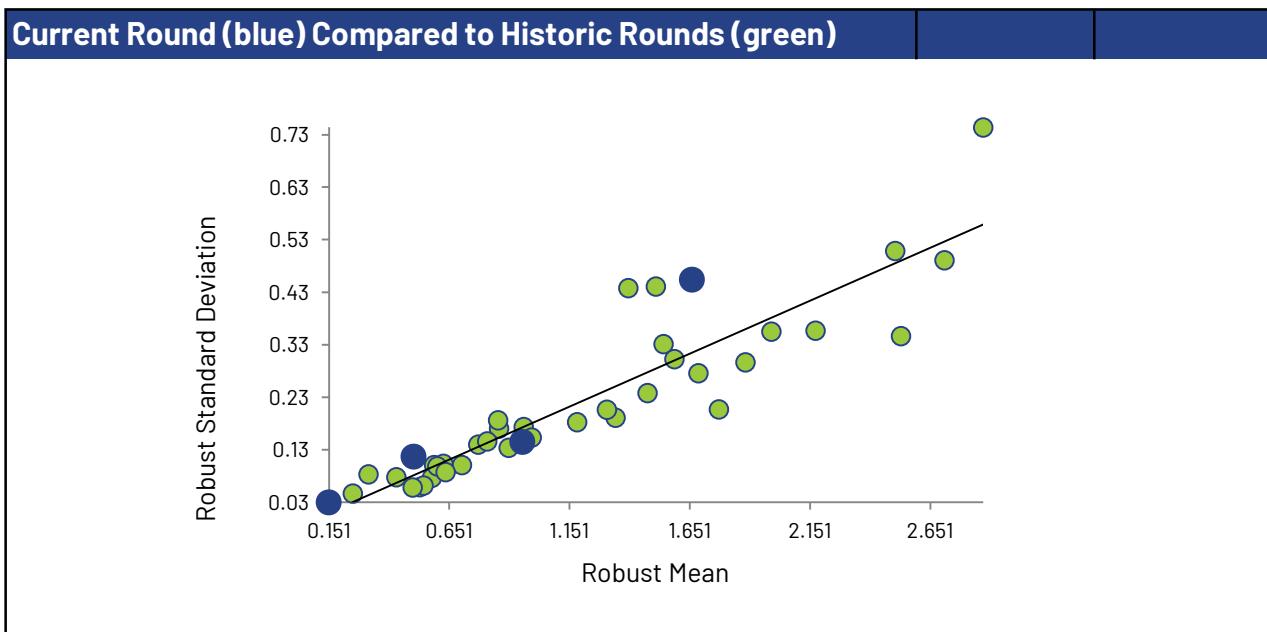
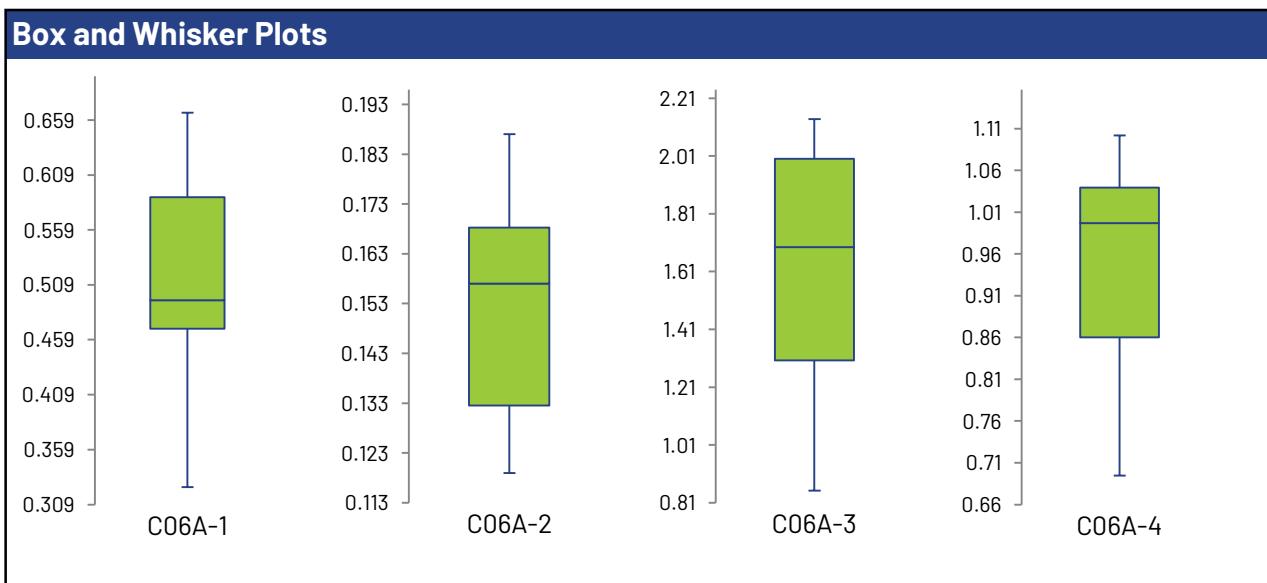
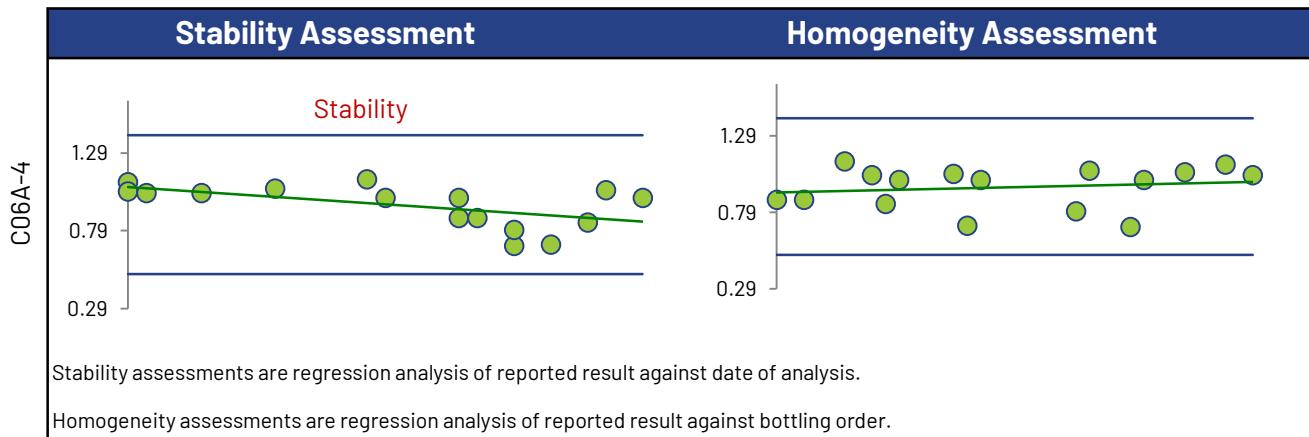
All summary stats, except the failure rate, and the plots below are based on the data excluding any flagged outliers



* Displayed failure rate does not account for the RDL modified z-Scores or inaccurately reported non-detects.

Heptachlor epoxide

Heptachlor epoxide



Lindane (gamma-BHC)

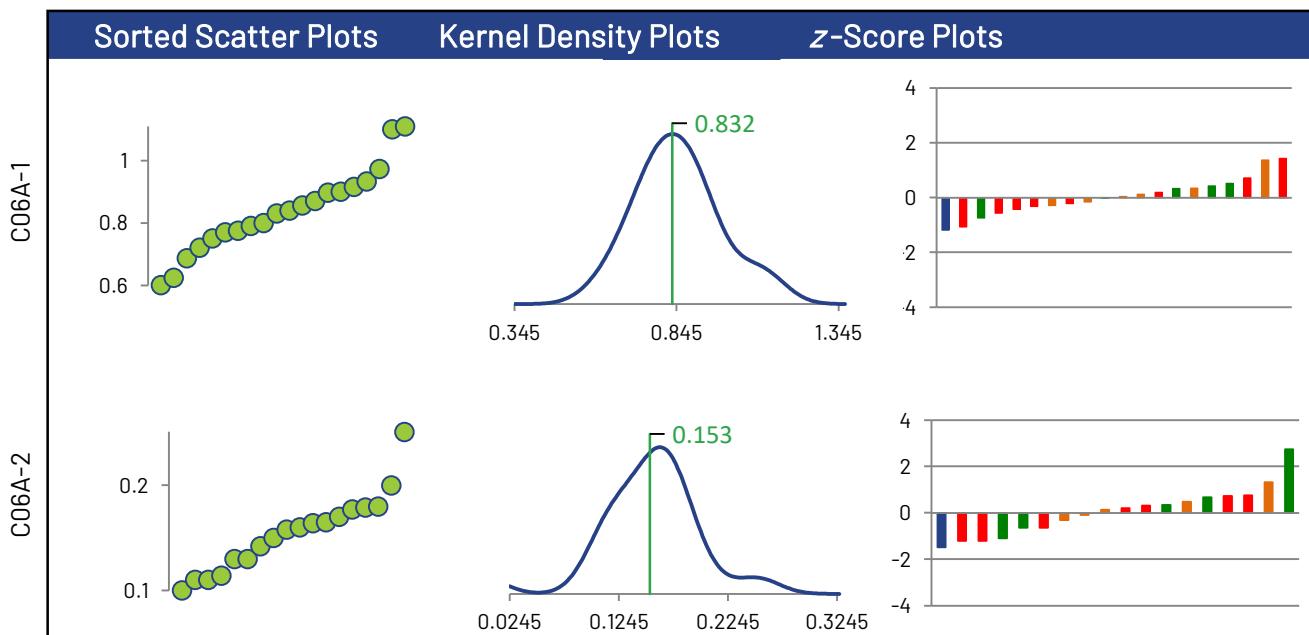
Summary Statistics

Statistic	C06A-1	C06A-2	C06A-3	C06A-4
No. of results used for summary statistics	20	18	20	20
Median µg/L	0.835	0.159	2.43	1.29
Robust Mean µg/L	0.832	0.153	2.31	1.27
Uncertainty of the Assigned Value µg/L	0.0363	0.0105	0.122	0.0598
Robust Standard Deviation µg/L	0.130	0.0357	0.437	0.214
Regression Standard Deviation µg/L	0.195	0.0357	0.541	0.297
Stability Flag				Stability
Homogeneity Flag				
Standard Deviation Used (SDPA) µg/L	0.195	0.0357	0.541	0.330
Flagged Outliers	0	1	0	0
z >3.0	0	0	1	0
2< z <3	0	1	0	0
Failure Rate %*			5.0	

Methods Used

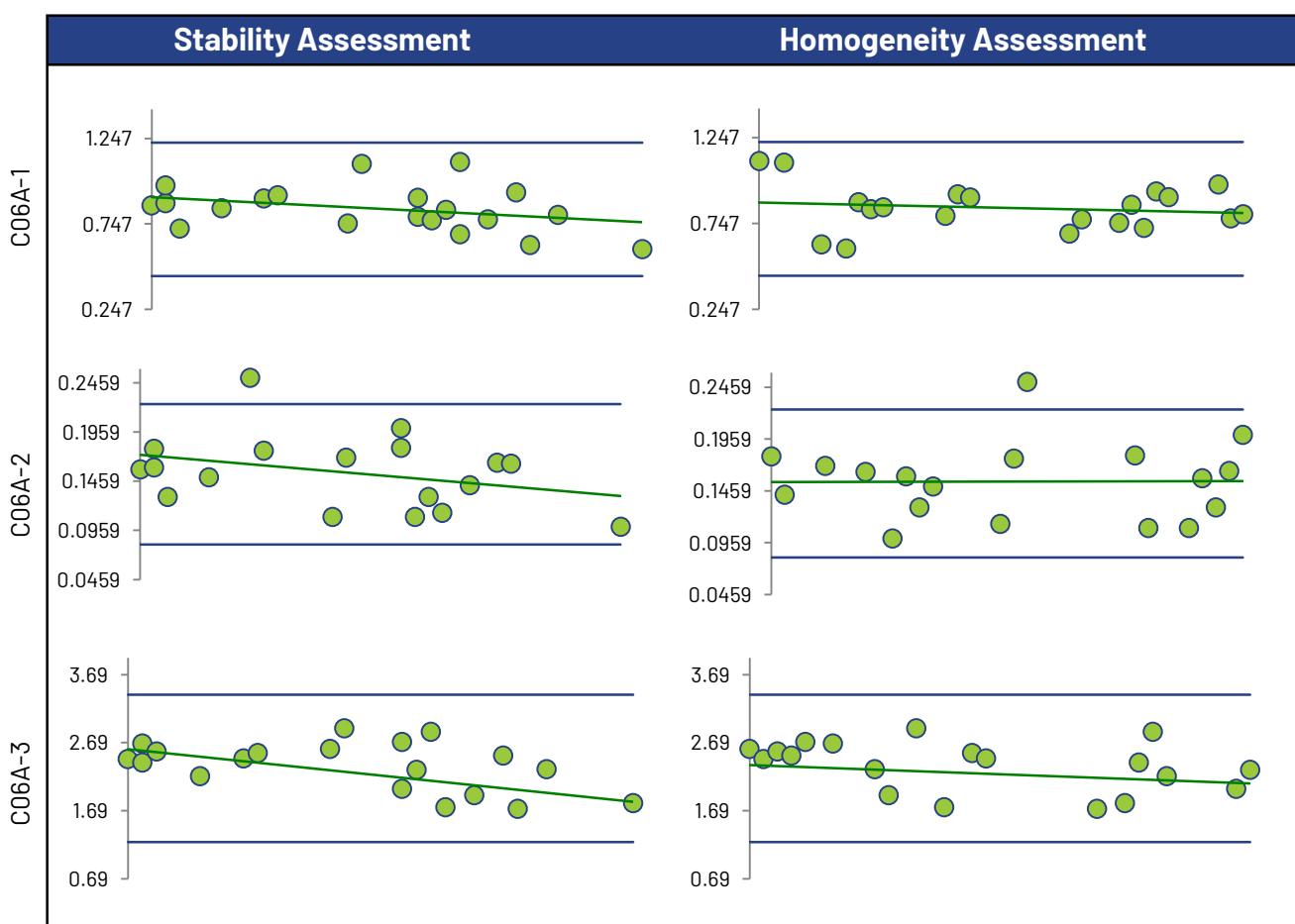
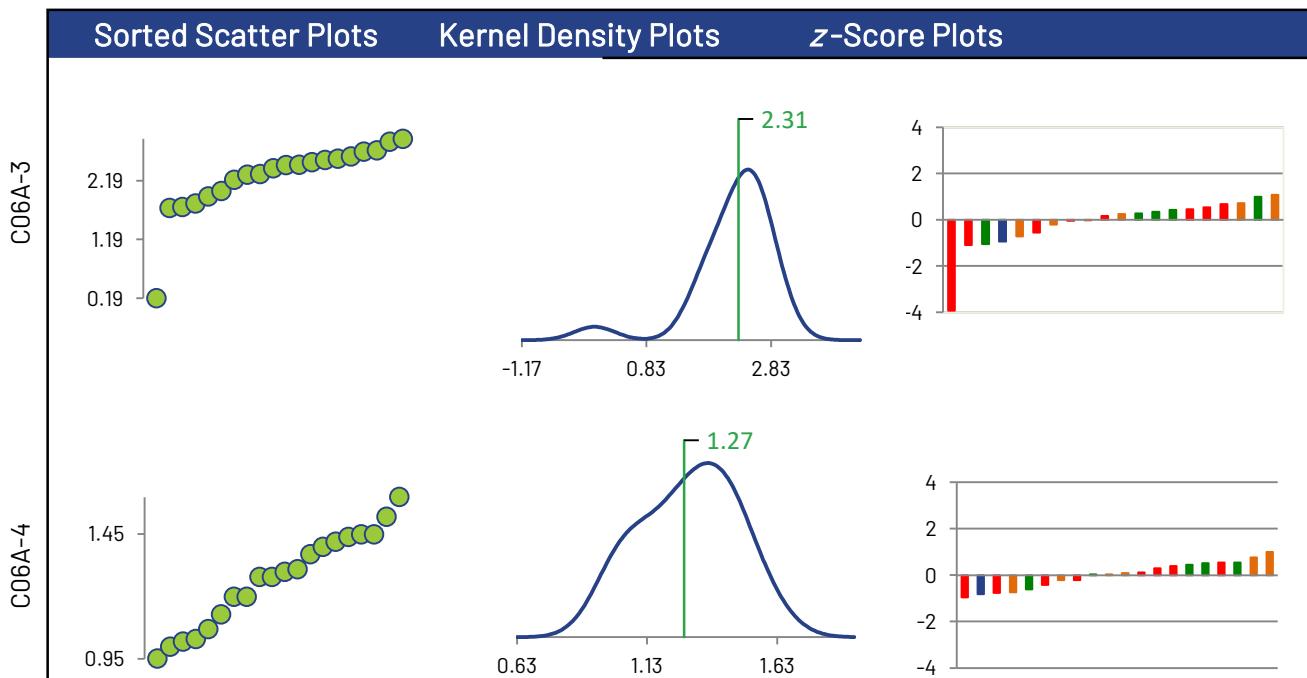
Method	C06A-1	C06A-2	C06A-3	C06A-4
GCXGC/ECD (Blue)	1	1	1	1
GC/ECD (Red)	8	7	8	8
GC/MS/MS (Green)	5	5	5	5
GC/MS (Orange)	6	5	6	6

All summary stats, except the failure rate, and the plots below are based on the data excluding any flagged outliers



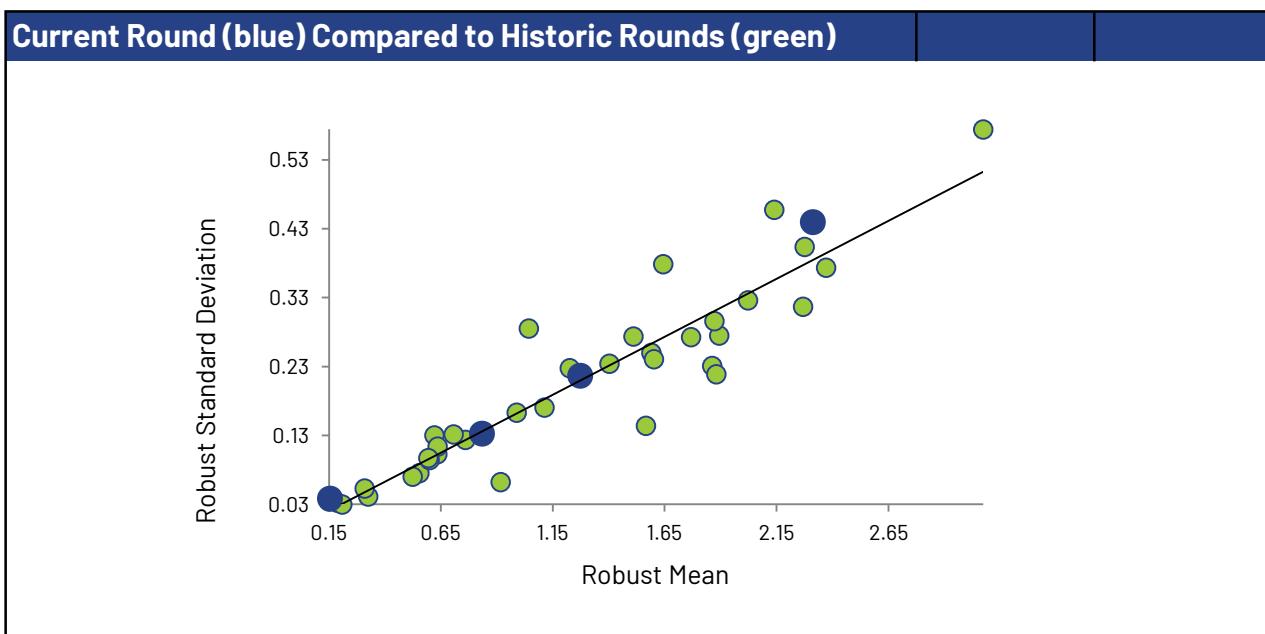
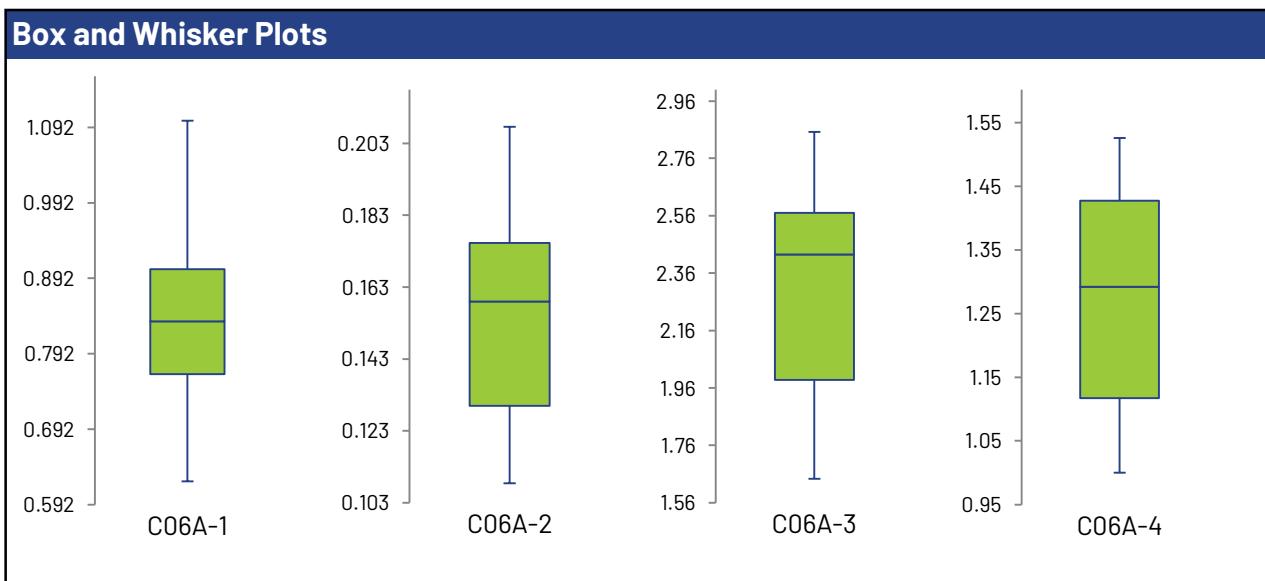
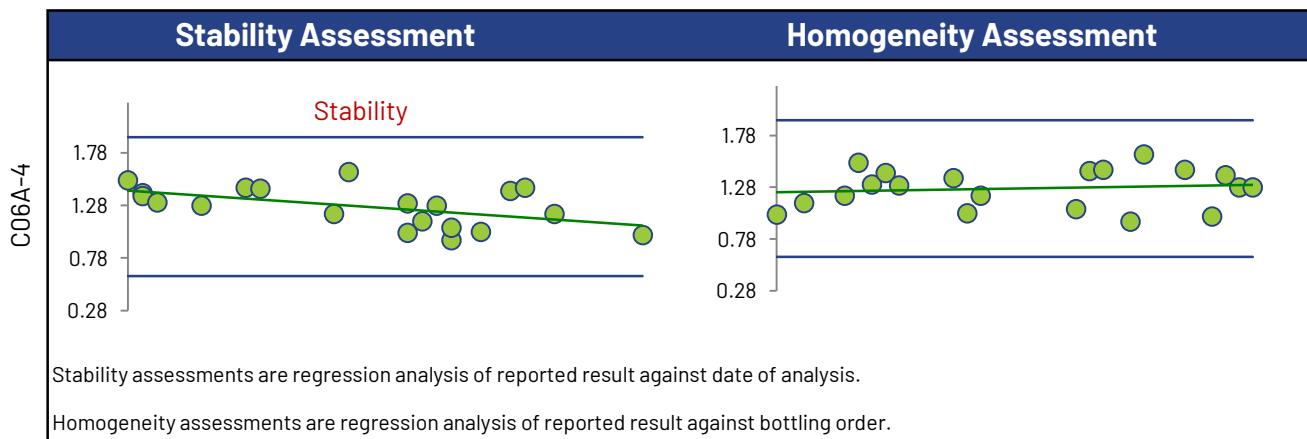
* Displayed failure rate does not account for the RDL modified z-Scores or inaccurately reported non-detects.

Lindane (gamma-BHC)



Annex A Summary by Analyte

Lindane (gamma-BHC)



Annex A Summary by Analyte

Mirex

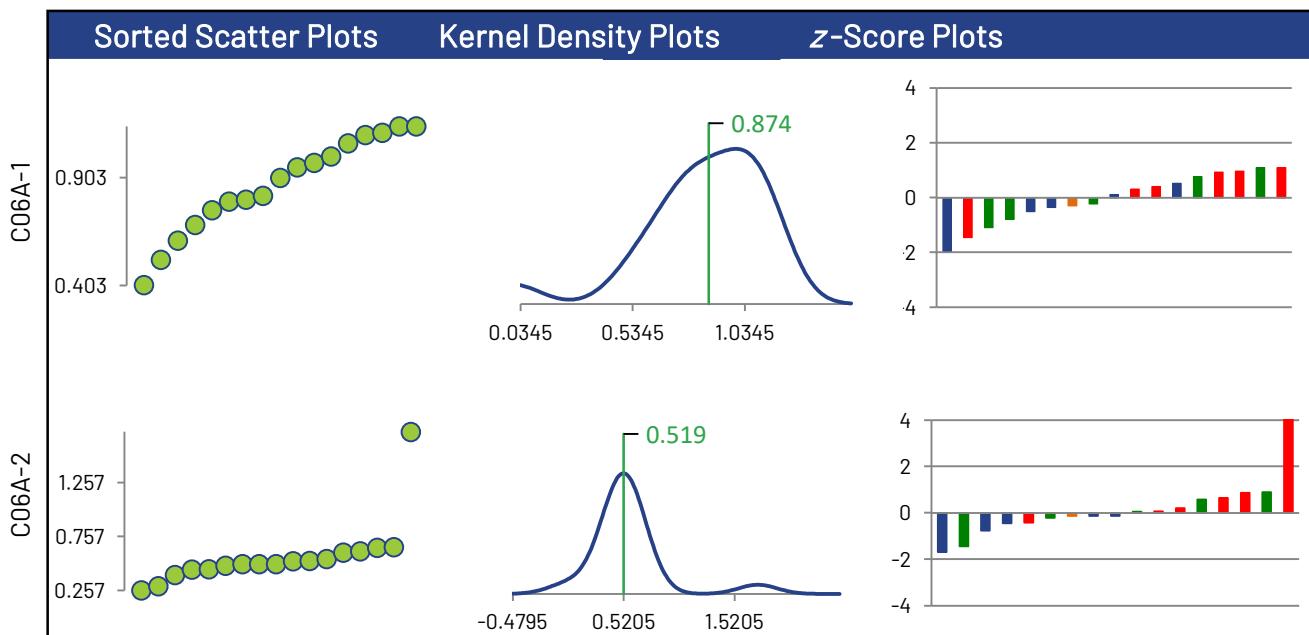
Summary Statistics

Statistic	C06A-1	C06A-2	C06A-3	C06A-4
No. of results used for summary statistics	17	17	18	18
Median µg/L	0.900	0.500	1.81	1.50
Robust Mean µg/L	0.874	0.519	1.66	1.49
Uncertainty of the Assigned Value µg/L	0.0722	0.0367	0.140	0.0893
Robust Standard Deviation µg/L	0.238	0.121	0.476	0.303
Regression Standard Deviation µg/L	0.243	0.156	0.436	0.395
Stability Flag				Stability
Homogeneity Flag				
Standard Deviation Used (SDPA) µg/L	0.243	0.156	0.476	0.525
Flagged Outliers	0	0	0	0
$ z > 3.0$	0	1	0	0
$2 < z < 3$	0	0	1	0
Failure Rate %*			5.6	

Methods Used

Method	C06A-1	C06A-2	C06A-3	C06A-4
GC/MS (Blue)	5	5	5	5
GC/ECD (Red)	6	6	7	7
GC/MS/MS (Green)	5	5	5	5
GCXGC/ECD (Orange)	1	1	1	1

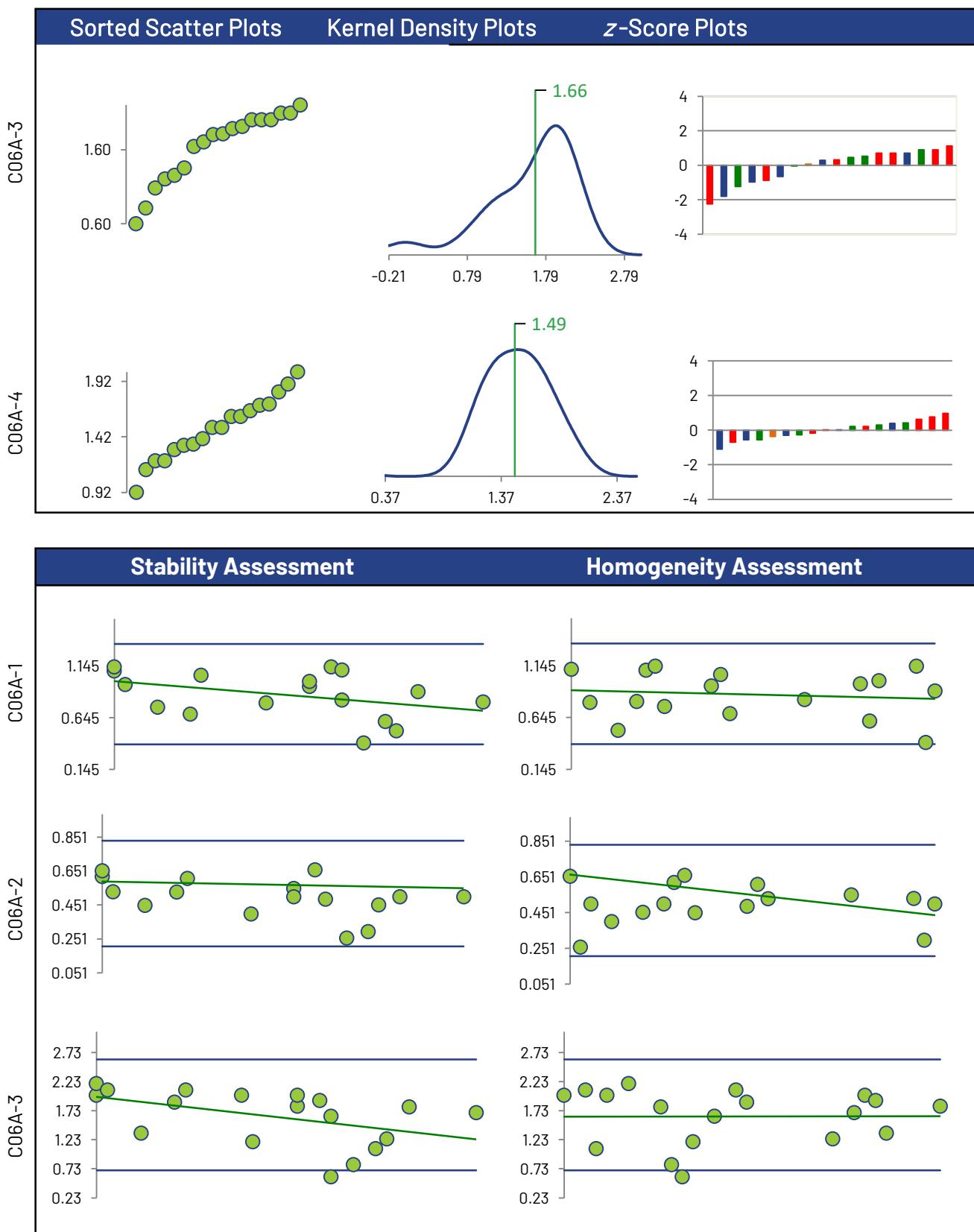
All summary stats, except the failure rate, and the plots below are based on the data excluding any flagged outliers



* Displayed failure rate does not account for the RDL modified z-Scores or inaccurately reported non-detects.

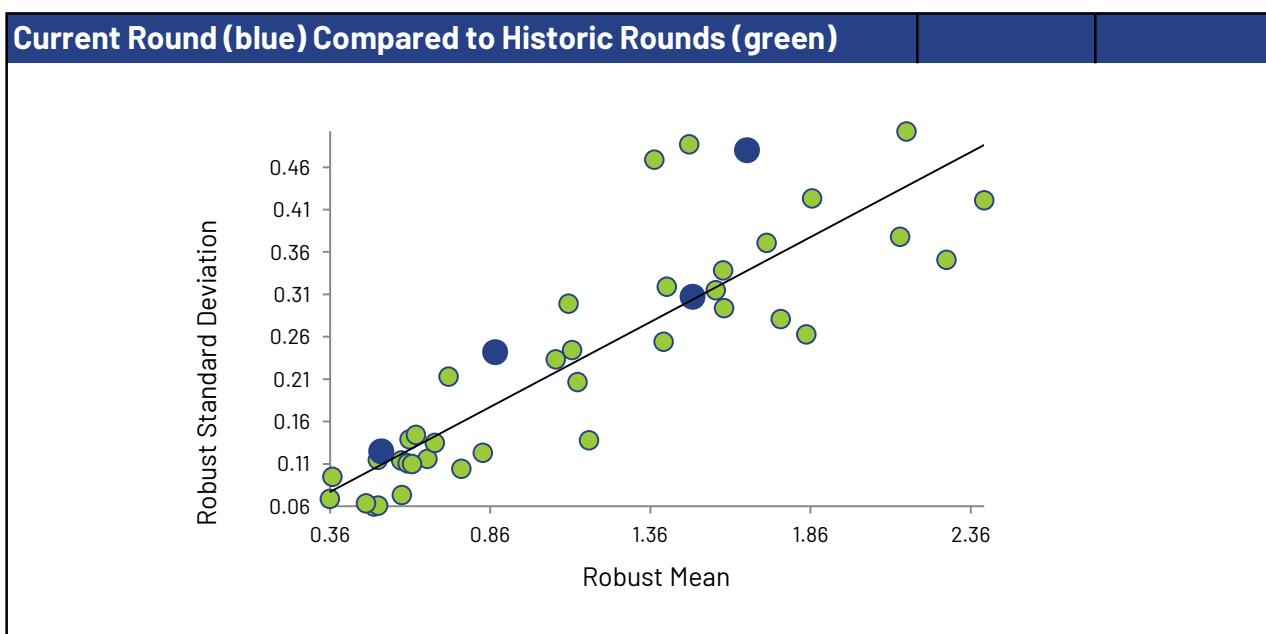
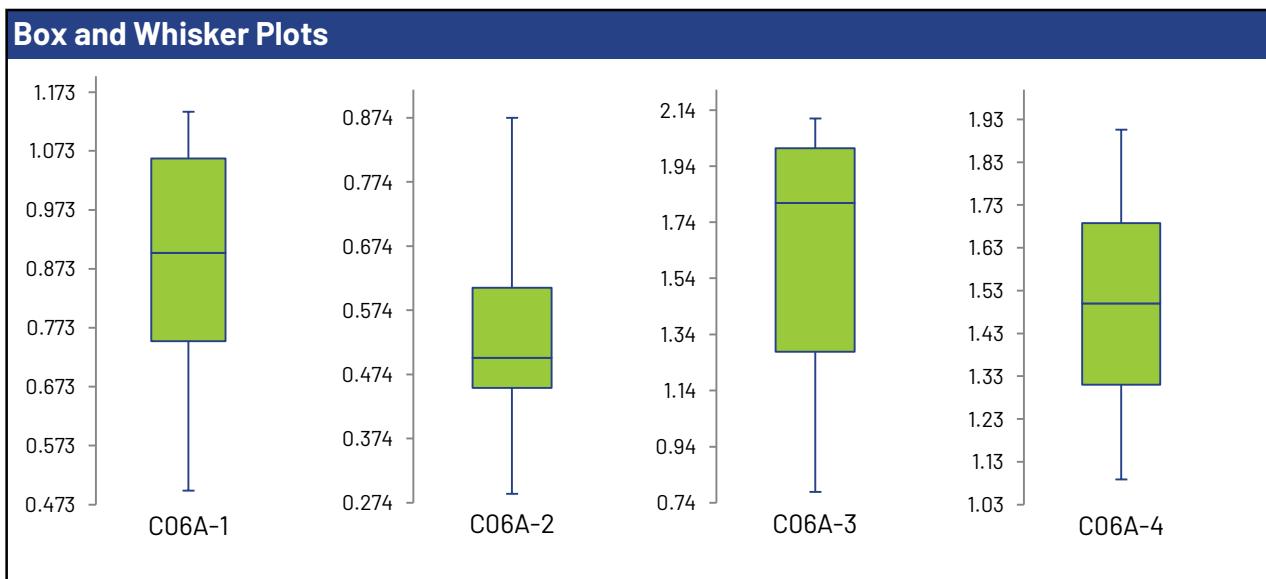
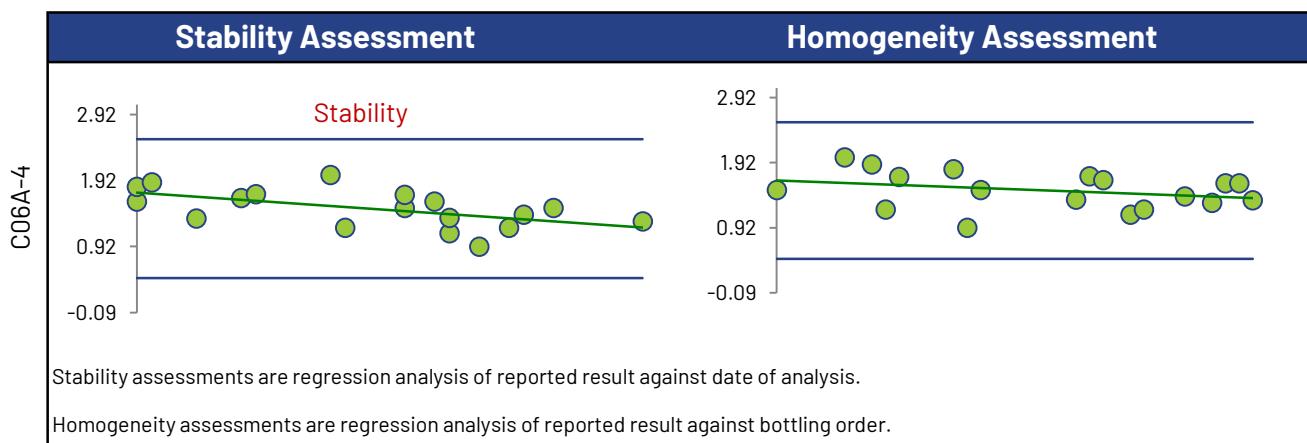
Annex A Summary by Analyte

Mirex



Annex A Summary by Analyte

Mirex



Annex A Summary by Analyte

o,p'-DDT

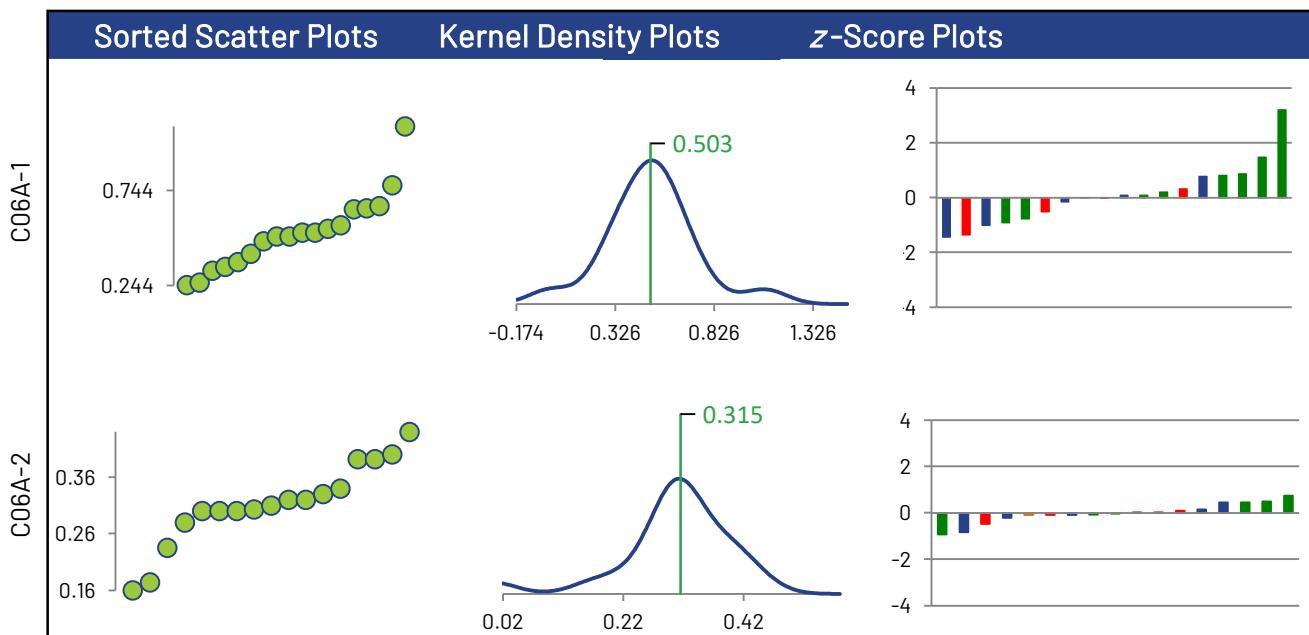
Summary Statistics

Statistic	C06A-1	C06A-2	C06A-3	C06A-4
No. of results used for summary statistics	18	17	19	19
Median µg/L	0.510	0.310	1.87	1.03
Robust Mean µg/L	0.503	0.315	1.88	1.06
Uncertainty of the Assigned Value µg/L	0.0530	0.0214	0.195	0.0837
Robust Standard Deviation µg/L	0.180	0.0706	0.680	0.292
Regression Standard Deviation µg/L	0.134	0.0954	0.415	0.248
Stability Flag		Stability		Stability
Homogeneity Flag				
Standard Deviation Used (SDPA) µg/L	0.180	0.169	0.680	0.588
Flagged Outliers	0	1	0	0
$ z > 3.0$	1	0	0	0
$2 < z < 3$	0	0	1	0
Failure Rate %*		5.3		

Methods Used

Method	C06A-1	C06A-2	C06A-3	C06A-4
GC/MS/MS (Blue)	5	5	5	5
GC/MS (Red)	4	4	5	5
GC/ECD (Green)	8	7	8	8
GCXGC/ECD (Orange)	1	1	1	1

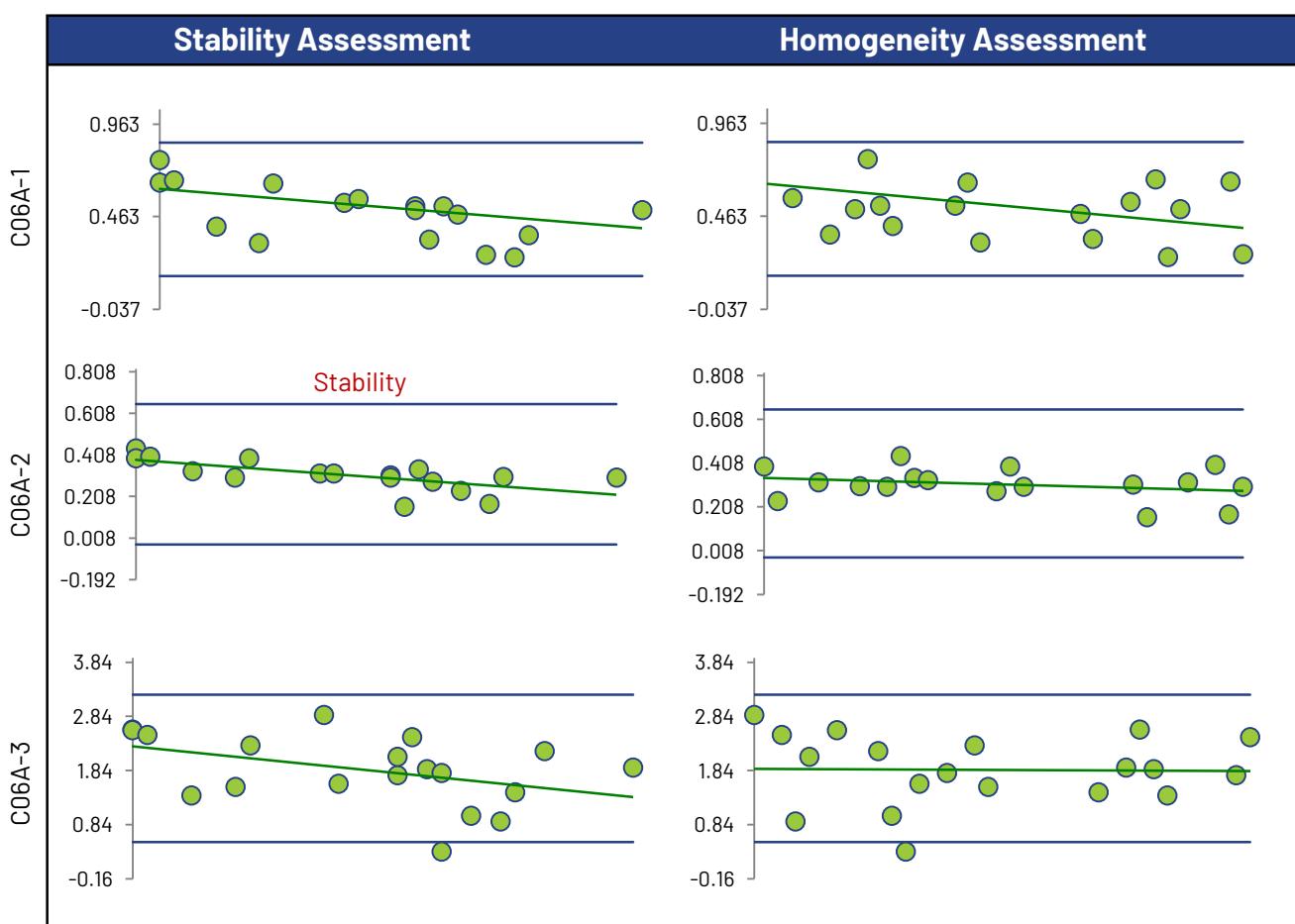
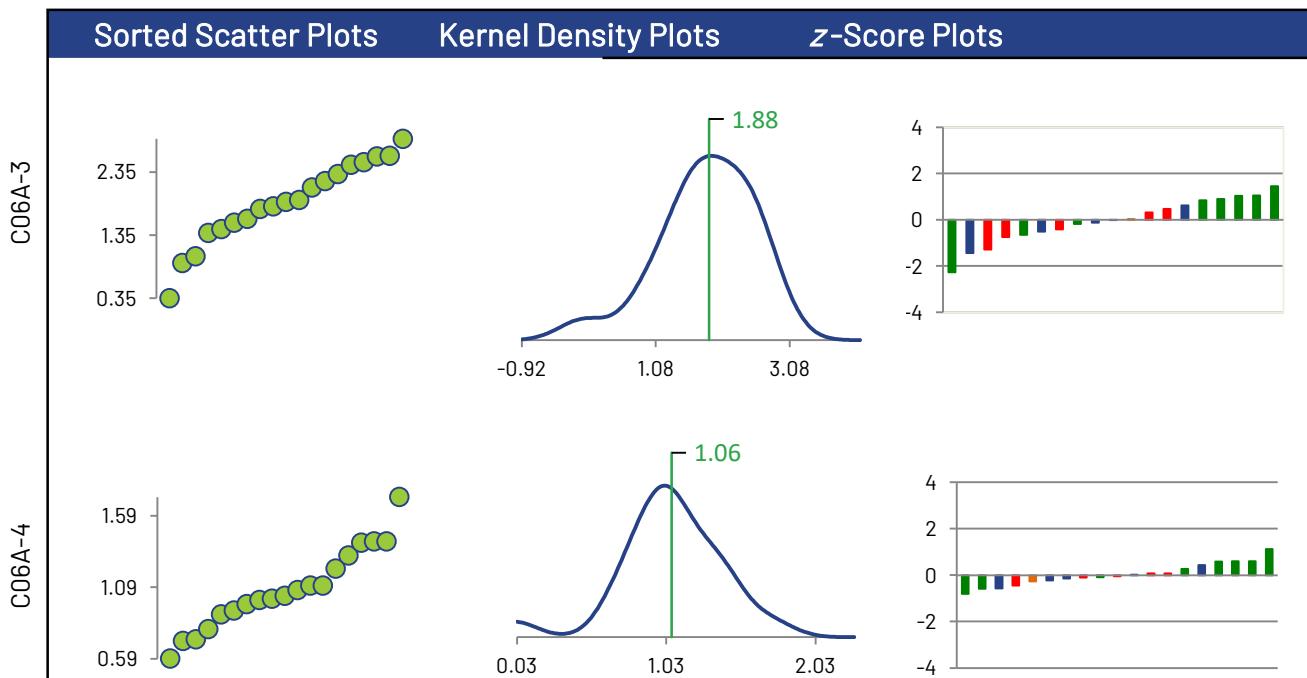
All summary stats, except the failure rate, and the plots below are based on the data excluding any flagged outliers



* Displayed failure rate does not account for the RDL modified z-Scores or inaccurately reported non-detects.

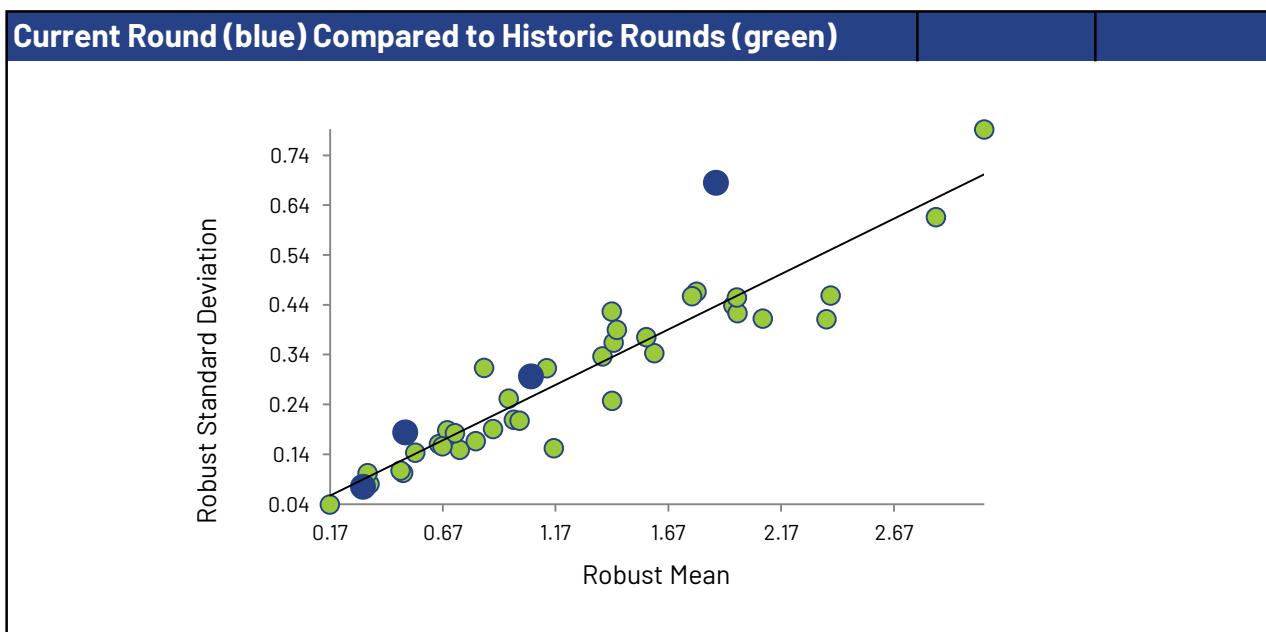
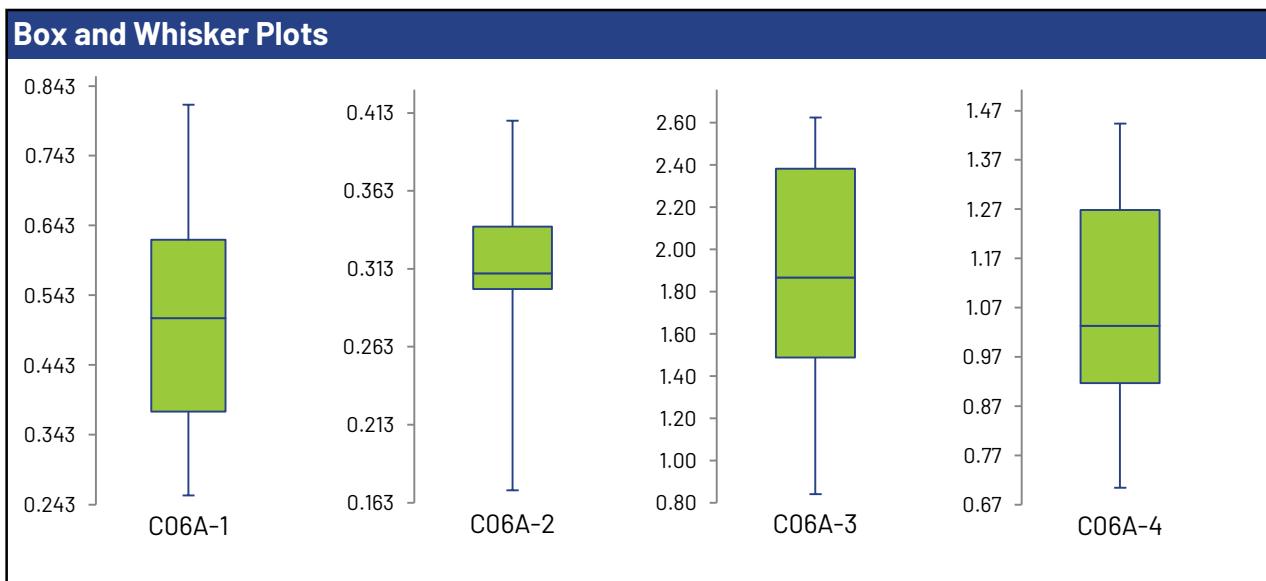
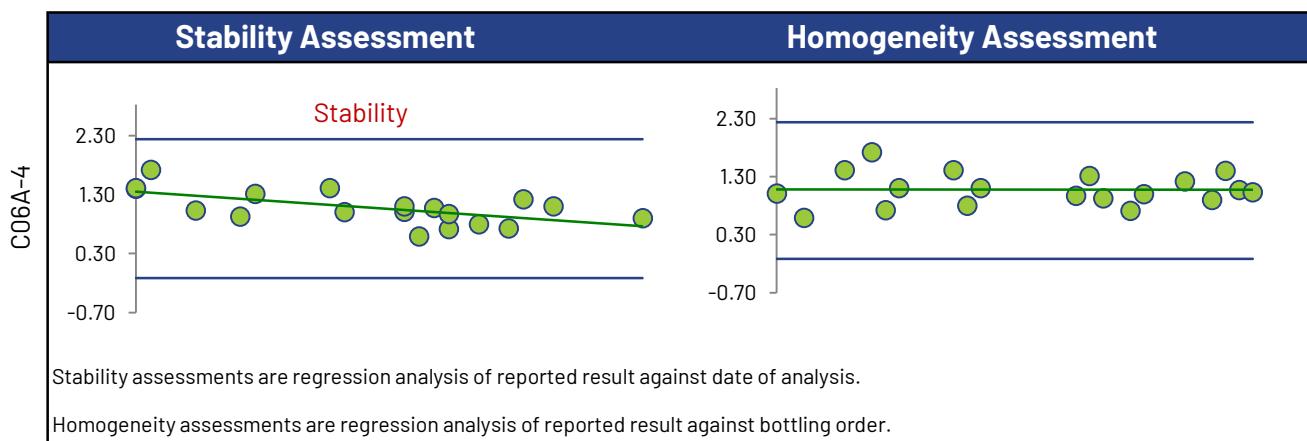
Annex A Summary by Analyte

o,p'-DDT



Annex A Summary by Analyte

o,p'-DDT



Annex A Summary by Analyte

p,p' - DDT

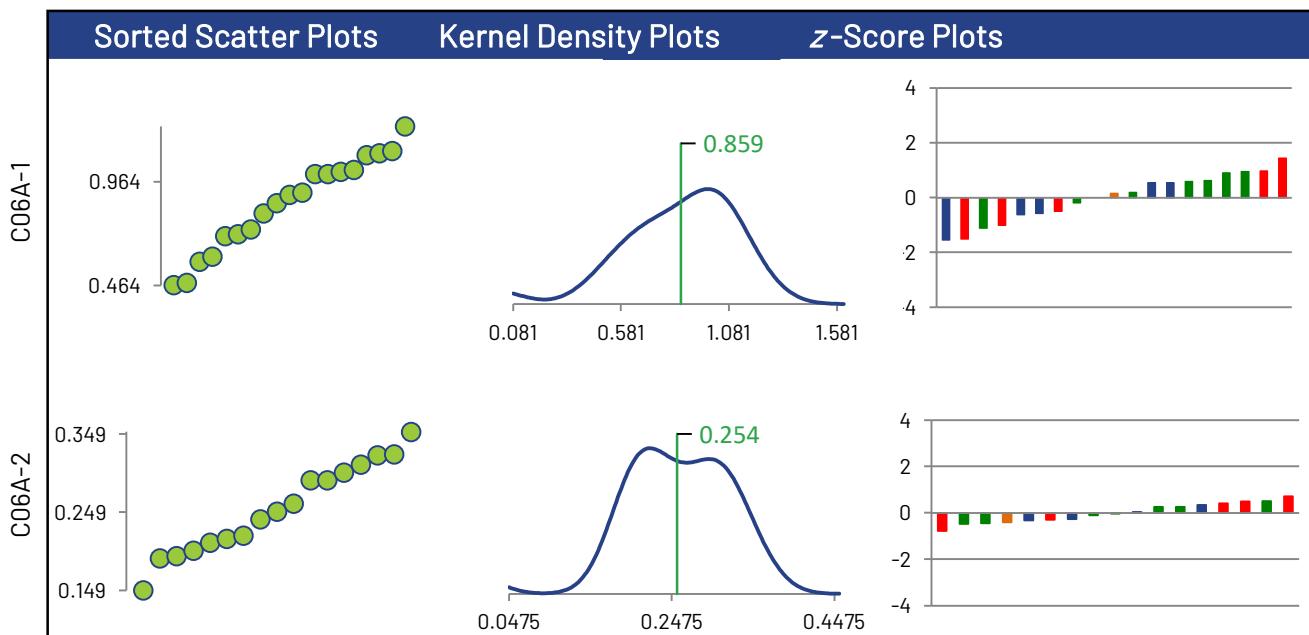
Summary Statistics

Statistic	C06A-1	C06A-2	C06A-3	C06A-4
No. of results used for summary statistics	19	17	19	19
Median µg/L	0.900	0.250	1.97	1.39
Robust Mean µg/L	0.859	0.254	1.95	1.39
Uncertainty of the Assigned Value µg/L	0.0731	0.0195	0.191	0.0952
Robust Standard Deviation µg/L	0.255	0.0644	0.667	0.332
Regression Standard Deviation µg/L	0.230	0.0834	0.496	0.359
Stability Flag		Stability		Stability
Homogeneity Flag				
Standard Deviation Used (SDPA) µg/L	0.255	0.136	0.667	0.596
Flagged Outliers	0	1	0	0
z >3.0	0	0	0	0
2< z <3	0	0	1	0
Failure Rate %*		5.3		

Methods Used

Method	C06A-1	C06A-2	C06A-3	C06A-4
GC/MS (Blue)	5	4	5	5
GC/MS/MS (Red)	5	5	5	5
GC/ECD (Green)	8	7	8	8
GCXGC/ECD (Orange)	1	1	1	1

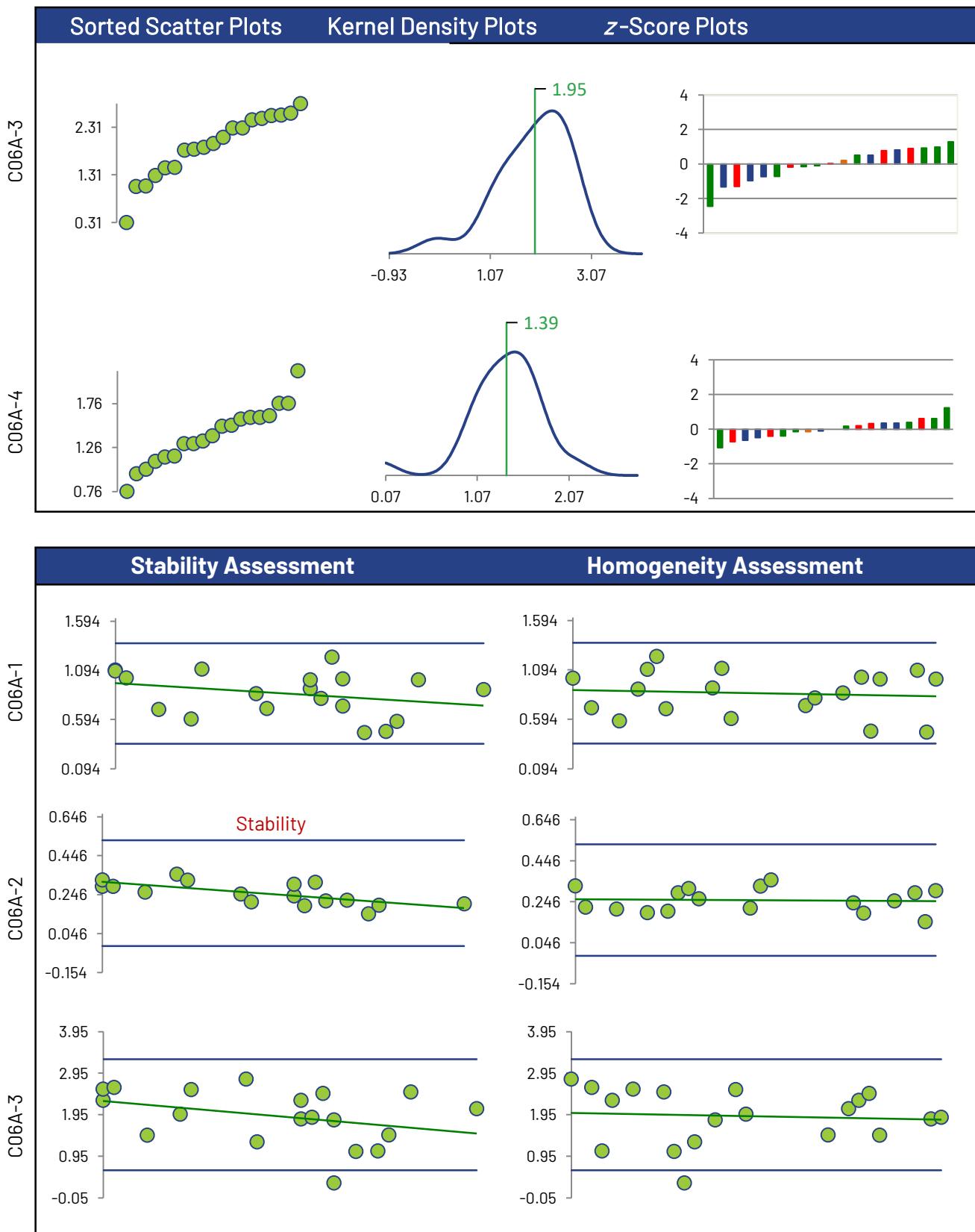
All summary stats, except the failure rate, and the plots below are based on the data excluding any flagged outliers



* Displayed failure rate does not account for the RDL modified z-Scores or inaccurately reported non-detects.

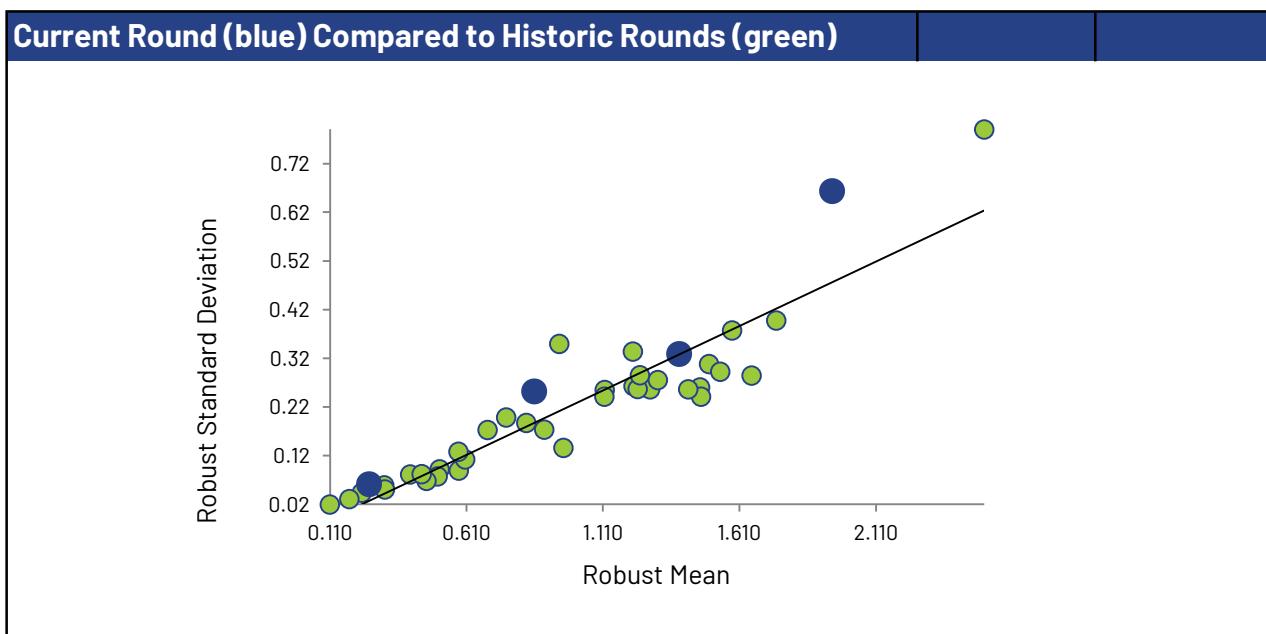
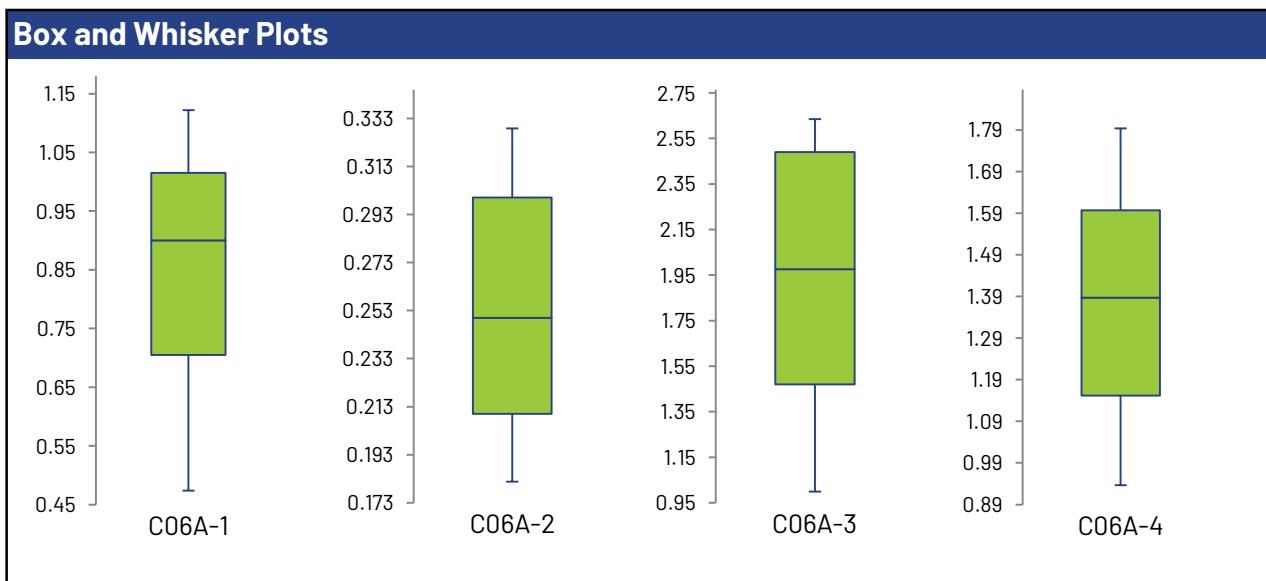
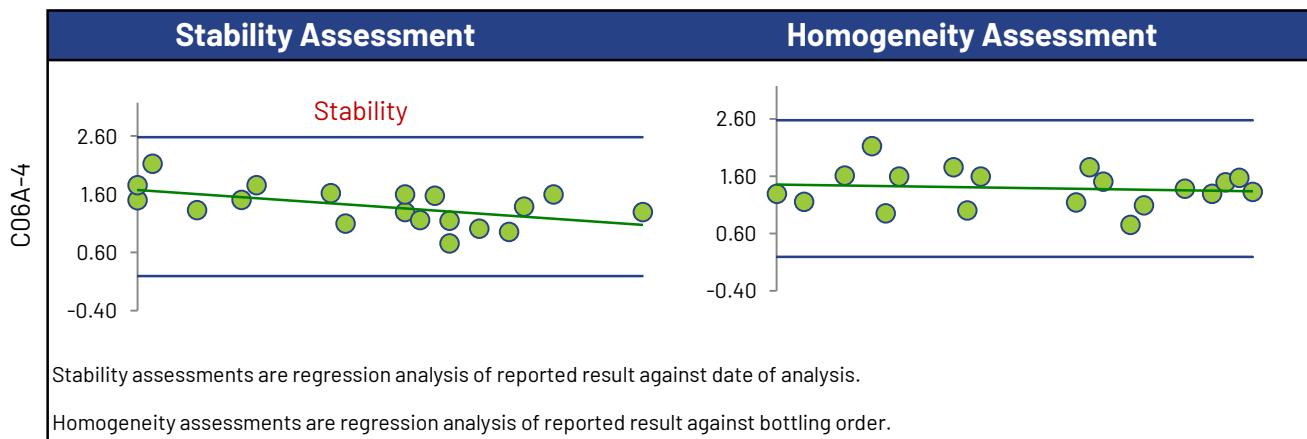
Annex A Summary by Analyte

p,p'-DDT



Annex A Summary by Analyte

p,p'-DDT



Annex A Summary by Analyte

p,p' Methoxychlor

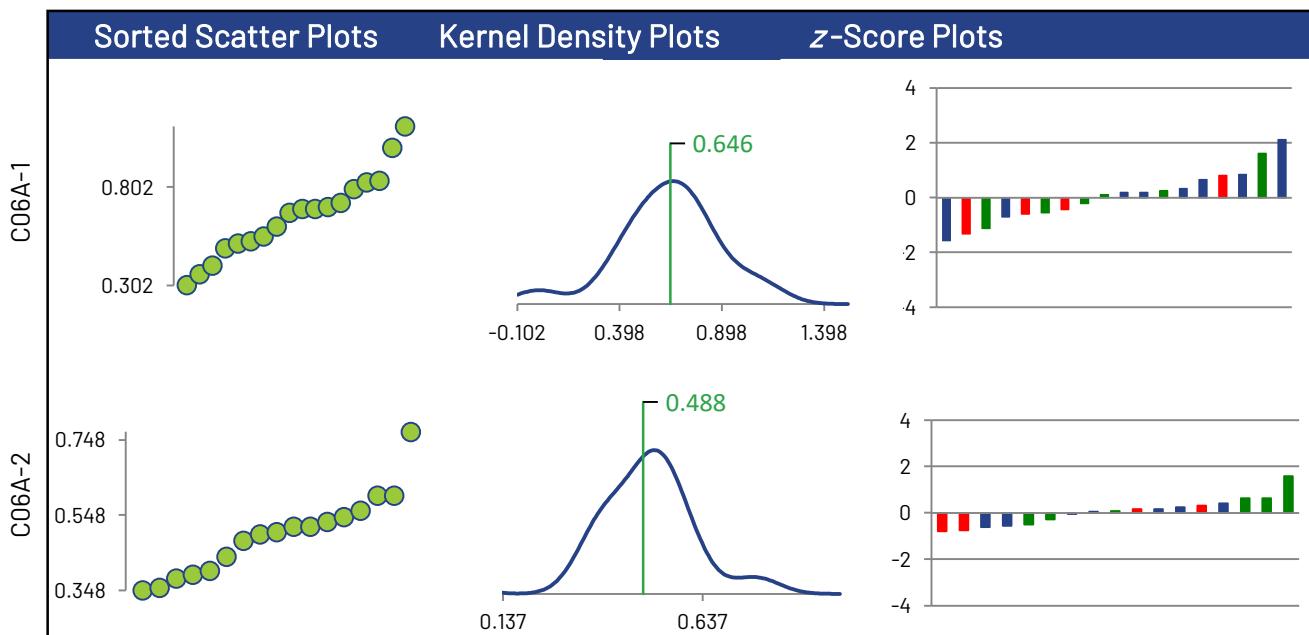
Summary Statistics

Statistic	C06A-1	C06A-2	C06A-3	C06A-4
No. of results used for summary statistics	18	17	18	18
Median µg/L	0.680	0.503	2.16	1.19
Robust Mean µg/L	0.646	0.488	2.17	1.24
Uncertainty of the Assigned Value µg/L	0.0645	0.0309	0.134	0.0725
Robust Standard Deviation µg/L	0.219	0.102	0.455	0.246
Regression Standard Deviation µg/L	0.184	0.152	0.503	0.308
Stability Flag			Stability	
Homogeneity Flag		Homogeneity		Homogeneity
Standard Deviation Used (SDPA) µg/L	0.219	0.177	0.806	0.394
Flagged Outliers	0	1	0	0
z >3.0	0	0	0	0
2< z <3	1	0	0	0
Failure Rate %*		5.6		

Methods Used

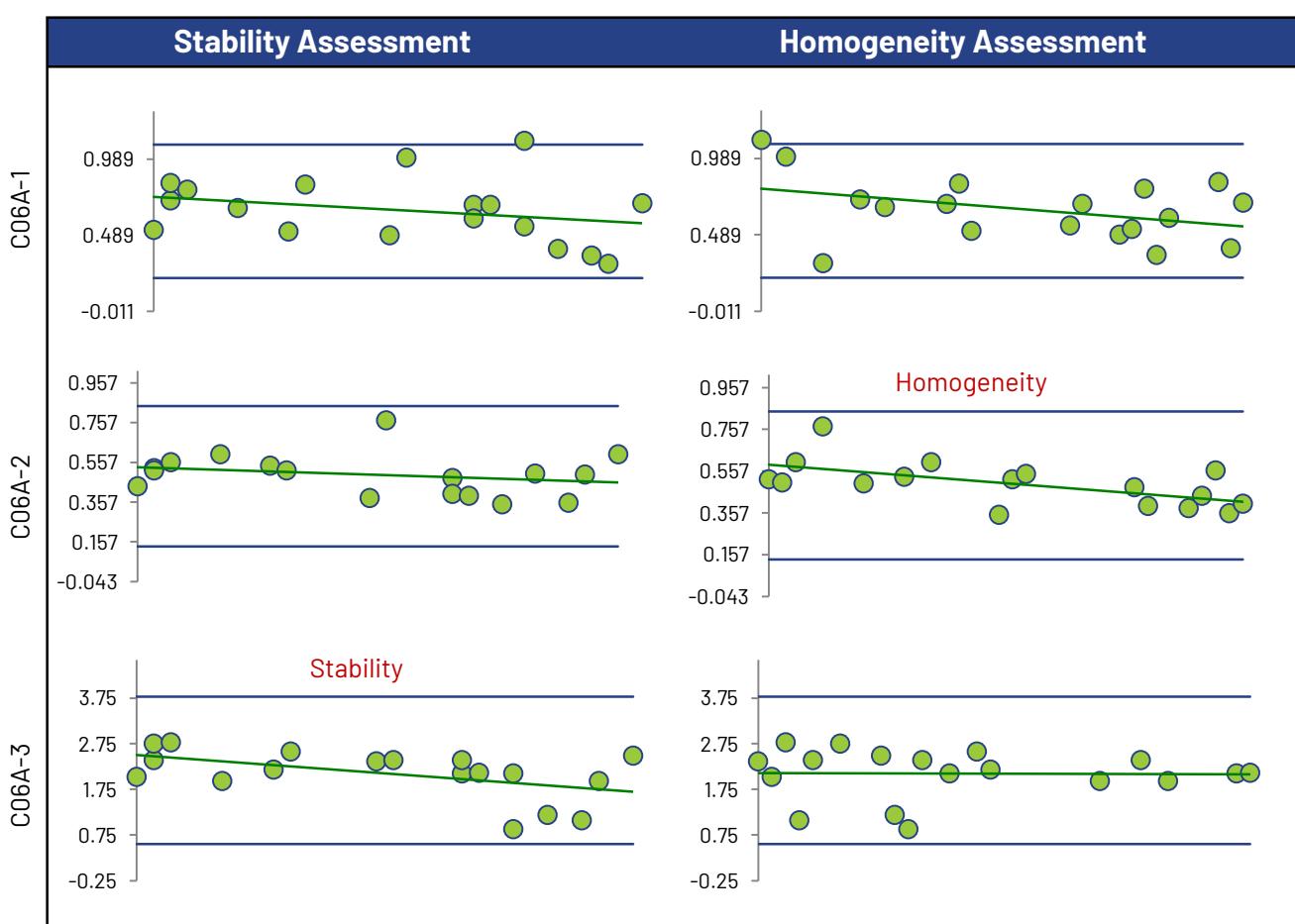
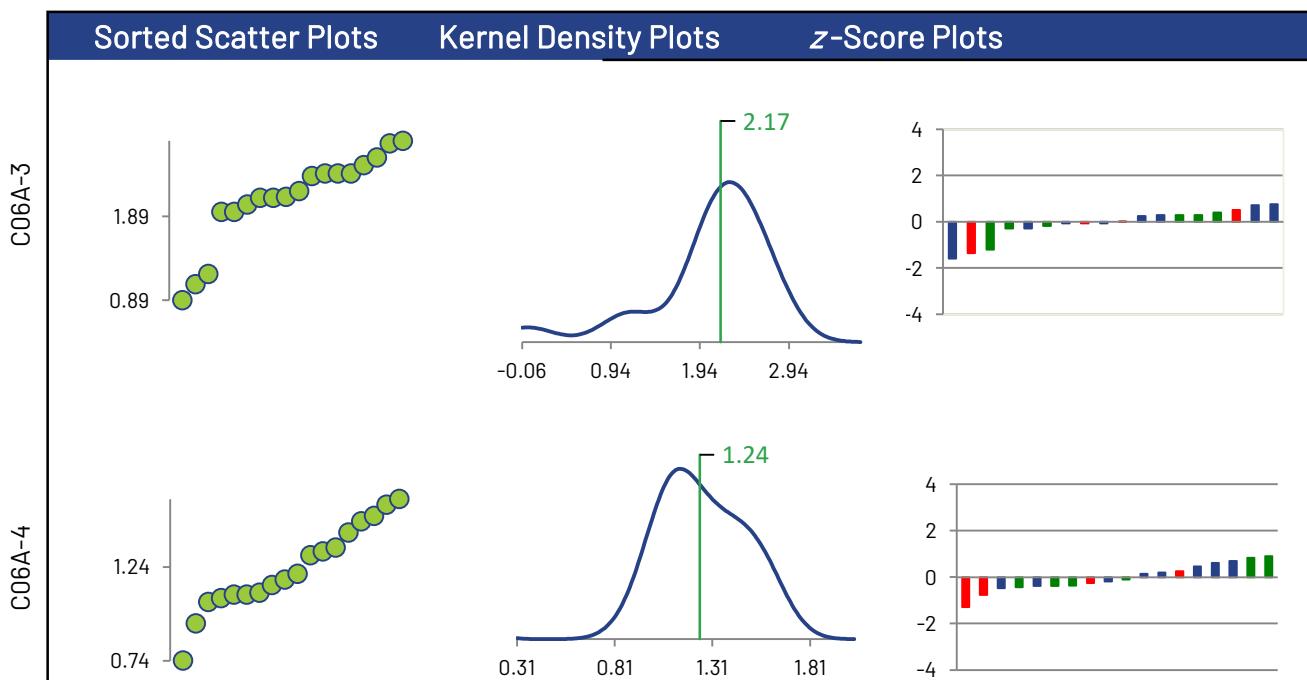
Method	C06A-1	C06A-2	C06A-3	C06A-4
GC/ECD (Blue)	8	7	8	8
GC/MS/MS (Red)	4	4	4	4
GC/MS (Green)	6	6	6	6

All summary stats, except the failure rate, and the plots below are based on the data excluding any flagged outliers



* Displayed failure rate does not account for the RDL modified z-Scores or inaccurately reported non-detects.

p,p' Methoxychlor



Annex A Summary by Analyte

p,p' Methoxychlor

