ALLIANCE 284 Sheffield Street, Mountainside New Jersey 07092 NEW JERSEY LAB ID#: 20012: NEW YORK LAB ID#: 11376

GC/MS SEMI-VOLATILE ANALYSIS CONFORMANCE/NON-CONFORMANCE SUMMARY

CHEM	TECH PROJECT NUMBER: P4866 MA	TRIX: Water			
METH	OD: 8270-Modified/3510				
1.	Chromatograms Labeled/Compounds Identified. (Field samples and I	Method Blanks)	NA	NO	YES ✔
2.	GC/MS Tuning Specifications. DFTPP Meet Criteria. (NOTE THAT THERE ARE DIFFERENT CRITERIA FOR NY ASF CLP AND NJ)	CLP,			✓
3.	GC/MS Tuning Frequency - Performed every 24 hours for 600 series 8000 Series.	and 12 hours for			✓
4.	GC/MS Calibration - Initial Calibration performed within 30 days bef analysis and continuing calibration performed within 24 hours of same 600 series and 12 hours for 8000 series.				✓
5.	GC/MS Calibration Requirements.			\checkmark	
	The Initial Calibration met the requirements .				
	The Continuous Calibration File ID BN035219.D met the requiremen Fluorobiphenyl,Fluoranthene-d10,Nitrobenzene-d5 and Terphenyl-d1 our target compound, therefore no corrective action taken.				
	The Continuous Calibration File ID BN035235.D met the requiremen Fluorobiphenyl and Terphenyl-d14 which are not our target compound corrective action taken.				
6.	Blank Contamination - If yes, list compounds and concentrations in ea	ach blank:		\checkmark	
7.	Surrogate Recoveries Meet Criteria			\checkmark	
	If not met, list those compounds and their recoveries which fall outsid ranges.	e the acceptable			
	The Surrogate recoveries met the acceptable criteria except for RW7-SP200-20241114 [Nitrobenzene-d5 - 47%, Terphenyl-d14 - 661 RW7-SP201-20241114 [2-Fluorobiphenyl - 24%, Nitrobenzene-d5 - 5 d14 - 573%], PB165012BL [2-Fluorobiphenyl - 42%, Terphenyl-d14 - 323%], PB165012BS [2-Fluorobiphenyl - 22%, Terphenyl-d14 - 556%], PB165012BSD [2-Fluorobiphenyl - 19% and Terphenyl-d14 - 567%] was not associated withclient parameter list, therefore no corrective ac	54%, Terphenyl- ,failing surrogates			

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<u>GC/MS SEMI-VOLATILE ANALYSIS CONFORMANCE/NON-CONFORMANCE SUMMARY</u> (<u>CONTINUED</u>)

		NA	NO	YES
8.	Matrix Spike/Matrix Spike Duplicate Recoveries Meet Criteria			\checkmark
	If not met, list those compounds and their recoveries which fall outside the acceptable range.			
	The Blank Spike met requirements for all samples . The Blank Spike Duplicate met requirements for all samples .			
9.	Internal Standard Area/Retention Time Shift Meet Criteria		\checkmark	
	Comments: The Internal Standards Areas met the acceptable requirements except for RW7-SP200-20241114, RW7-SP201-20241114, failing Internal Standards were not associated with client parameter list, therefore no corrective action taken.			
10.	Extraction Holding Time Met			\checkmark
	If not met, list number of days exceeded for each sample:			
11.	Analysis Holding Time Met			\checkmark
	If not met, list number of days exceeded for each sample:			

ADDITIONAL COMMENTS:

Samples RW7-SP200-20241114 was diluted at 2X. This sample initially analyzed straight in sequence BN112124 but END CCAL is missing therefore based on that result lab analyzed this sample directly with 2X while Straight analysis given as screening data in Miscellaneous section.

The laboratory certifies that the all-electronic diskette deliverable exactly match the data summary forms (i.e. Form Is)." The Form 6 is not included in the data package because the Initial Calibration was performed using 7 points.

Please use %D calculated based on Avg RF and CCRF for all compounds using Average Response Factor when the %RSD value for a compound is 20% for the Initial Calibration curve and use %D calculated based on Amount added and Calculated amount for all compounds using Linear Regression when the %RSD value for a compound is > 20% for the Initial Calibration curve for SW-846 analysis.

QA REVIEW