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

MATRIX: Water

CHEMTECH PROJECT NUMBER: P3845

METHOD: 8270-Modified/3510 NO NA YES 1. Chromatograms Labeled/Compounds Identified. (Field samples and Method Blanks) 2. GC/MS Tuning Specifications. DFTPP Meet Criteria. (NOTE THAT THERE ARE DIFFERENT CRITERIA FOR NY ASP CLP, CLP AND NJ) GC/MS Tuning Frequency - Performed every 24 hours for 600 series and 12 hours for 3. 8000 Series. 4. GC/MS Calibration - Initial Calibration performed within 30 days before sample analysis and continuing calibration performed within 24 hours of sample analysis for 600 series and 12 hours for 8000 series. 5. GC/MS Calibration Requirements. The Initial Calibration met the requirements. The Continuous Calibration met the requirements. 6. Blank Contamination - If yes, list compounds and concentrations in each blank: 7. Surrogate Recoveries Meet Criteria If not met, list those compounds and their recoveries which fall outside the acceptable ranges. The Surrogate recoveries met the acceptable criteria except for PT-ACIDS-WP [2,4,6-Tribromophenol - 156%, 2-Methylnaphthalene-d10 - 0%, Fluoranthene-d10 - 0%, Nitrobenzene-d5 - 124%, Phenol-d6 - 115%], PT-ACIDS-WPDL [2,4,6-Tribromophenol - 144%, 2-Fluorophenol - 107%, 2-Methylnaphthalene-d10 - 0%, Fluoranthene-d10 - 0%, Nitrobenzene-d5 - 131%, Phenol-d6 - 120%], PT-ACIDS-WPDL2 [2-Methylnaphthalene-d10 - 0%, Fluoranthene-d10 - 0%, Phenol-d6 - 103%], PB163341BS [2-Fluorophenol - 102% and Phenol-d6 - 101%], This sample was extracted for full scan analysis and above mention surrogates were not part of full scan

extraction therefore no corrective action is required.

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

NA

NO

YES

8.	Matrix Spike/Matrix Spike Duplicate Recoveries Meet Criteria		•
	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 .The RPI {PB163341BSD} with File ID: BN034053.D met criteria except for 4 methylphenol[21%], due to difference in results of BS and BSD.		
9.	Internal Standard Area/Retention Time Shift Meet Criteria		•
	Comments:		
10.	Extraction Holding Time Met		✓
	If not met, list number of days exceeded for each sample:		
11.	Analysis Holding Time Met		✓
	If not met, list number of days exceeded for each sample:		
	The Holding Times were met for all analysis.		
ADDIT	TONAL COMMENTS:		
Sample	s PT-ACIDS-WP, PT-ACIDS-WPDL were diluted due to high concentr	ations.	
The Fo	rm 6 is not included in the data package because the Initial Calibration w	as performed using 7 points.	
Please 1	use %D calculated based on Avg RF and CCRF for all compounds using	Average Response Factor when	the
%RSD	value for a compound is <15% for the Initial Calibration curve and use 9	6D calculated based on Amount	added
and Cal	culated amount for all compounds using Linear Regression when the %l	RSD value for a compound is >	15% for
he Initi	ial Calibration curve for SW-846 analysis.		
QA RE	VIEW	Date	