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

GC/MS VOA CONFORMANCE/NON-CONFORMANCE SUMMARY

CHEM	TECH PROJECT NUMBER: Q1664 M	ATRIX: Solid			
METH	OD: 8260D				
			NA	NO	YES
1.	Chromatograms Labeled/Compounds Identified. (Field samples a	nd Method Blanks)	1 11 1	110	\checkmark
2.	GC/MS Tuning Specifications BFB Meet Criteria (NOTE THAT THERE ARE DIFFERENT CR ASP CLP, CLP AND NJ)	ITERIA FOR NY			✓
3.	GC/MS Tuning Frequency - Performed every 24 hours for 600 ser 8000 Series.	ies and 12 hours for			\checkmark
4.	GC/MS Calibration - Initial Calibration performed before sample a continuing calibration performed within 24 hours of sample analys 12 hours for 8000 series.				✓
5.	GC/MS Calibration Requirements.			\checkmark	
	The Initial Calibration met the requirements.				
	The Continuous Calibration File ID VN086140.D met the requirer Hexanone and 4-Methyl-2-Pentanone, are failing high but no positi samples therefore no corrective action taken.				
6.	Blank Contamination - If yes, list compounds and concentrations i	n each blank:		\checkmark	
7.	Surrogate Recoveries Meet Criteria				\checkmark
	If not met, list those compounds and their recoveries which fall ou ranges.	tside the acceptable			
8.	Matrix Spike/Matrix Spike Duplicate Recoveries Meet Criteria			\checkmark	
	If not met, list those compounds and their recoveries which fall ou range.	tside the acceptable			
	The MS {Q1664-05MS} with File ID: VN086163.D recoveries me for all compounds except for 1,1,2,2-Tetrachloroethane[33%],due interference.				
	The MSD {Q1664-06MSD} with File ID: VN086164.D recoverie requirements except for 1,1,2,2-Tetrachloroethane[35%] and Tetrachloroethene[161%],due to matrix interference.	s met the acceptable			
	The RPD for {Q1664-06MSD} with File ID: VN086164.D met cr Bromomethane[32%],due to difference in results of MS and MSD				
	The Blank Spike recoveries met the acceptable requirements .				

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GC/MS VOA CONFORMANCE/NON-CONFORMANCE SUMMARY (CONTINUED)

		NA	NO	YES
9.	Internal Standard Area/Retention Time Shift Meet Criteria			\checkmark
	Comments:			
10.	Analysis Holding Time Met			\checkmark
	If not met, list number of days exceeded for each sample:			

ADDITIONAL COMMENTS:

Trip Blank was not provided with this set of samples.

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

Date