

NEW JERSEY LAB ID#:20012 : NEW YORK LAB ID#: 11376

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

CHEMTECH PROJECT N	JMBER: bM092324			
SequenceID :	bM092324	NA	NO	YES
1. Chromatograms Labeled	Compounds Identified. (Field samples and Method I	Blanks)		✓
2. GC/MS Tuning Specific (NOTE THAT THERE AR	AND NJ)		_	
3. GC/MS Tuning Frequen series	y - Performed every 24 hours for 600 series and 12 h	iours for 8000		√
	al Calibration performed within 30 days before samp alibration performed within 24 hours of sample analy rs for 8000 series	•		✓
5. GC/MS Calibration Met				✓
a. Initial calibration Me If not met, list those compo	t Criteria and their recoveries which fall outside the accept	ptable range.		√
-	(CCC) Meet Criteria ands and their recoveries which fall outside the accep oppound #77 which is not present in parameter list o		<u> </u>	
-	yes, list compounds and concentrations in each blan	·	<u> </u>	

d. Acid Fraction

7. Surrogate Recoveries Meet CriteriaIf not met, list those compounds and their recoveries which fall outside the acceptable ranges.a. B/N Fraction	 <u>✓</u>	
 d. Acid Fraction 8. Matrix Spike/Matrix Spike Duplicate Recoveries Meet Criteria If not met, list those compounds and their recoveries which fall outside the acceptable range. a. B/N Fraction <u>Recovery and RPD fail for some compound in P4135-01MS/MSD due to matrix interference.</u> 	 <u> </u>	
d. Acid Fraction9. 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:	 <u> </u>	

ADDITIONAL COMMENTS:

One surrogate Terphenyl-d14 in PB163525BL and PB163606BL was marginally biased high. The data will be used for the hard copies. Recovery fail high for compound #81,90,92 in PB163606BS, however as they are passing in the ccc properly, the data will be used for the hard copies. The samples P4142-01 & P4146-01 analyzed with 5X each for concentrated matrix.



YES

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