

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

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

CHEMTECH PROJECT NUMBER:	bm040925			
SequenceID : bm040925		NA	NO	YES
1. Chromatograms Labeled/Compounds Iden	ified. (Field samples and Method Blanks)			✓
2. GC/MS Tuning Specifications. DFTPP Meet Criteria Criteria (NOTE THAT THERE ARE DIFFERENT CRITERIA FOR NY ASP CLP, CLP AND NJ)				✓
3. GC/MS Tuning Frequency - Performed ever series	ery 24 hours for 600 series and 12 hours for 8000			_ ✓
 GC/MS Calibration - Initial Calibration per analysis and continuing calibration perform for 600 series and 12 hours for 8000 series 				√
5. GC/MS Calibration Met:				✓
a. Initial calibration Meet Criteria If not met, list those compounds and their rec	overies which fall outside the acceptable range.			√
b. Continuous Calibration(CCC) Meet Crit If not met, list those compounds and their rec	eria overies which fall outside the acceptable range.		<u>✓</u>	
The compound #77 is biased high in the C	CC. But not present in the parameters list of the samples analyzed.			
 Blank Contamination - If yes, list compour a. B/N Fraction 	ds and concentrations in each blank:		<u>✓</u>	

d. Acid Fraction

7. Surrogate Recoveries Meet Criteria ✓ If not met, list those compounds and their recoveries which fall outside the acceptable ranges.	
a. B/N Fraction	
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 Recovery failed for some compounds in Q1744-01MS/MSD due to matrix interference. Hence, no corrective action is required 	
d. Acid Fraction	
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 ✓	

ADDITIONAL COMMENTS:

Recovery failed for some compounds in PB167521BS/BSD. This data will be used for hardcopies. The sample Q1739-01 had to be analyzed with 20X dilution due to oily, viscous and dirty matrix. Due to this reason it was extracted in 5ml volume. Even in this diluted sample, surrogates failed due to the presence of non-targeted hydrocarbons which can be witnessed by the abnormal chromatogram. Hence, this analysis will be final.



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