

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

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

CHEMTECH PROJE	CT NUMBER:	bf080924			
SequenceID :	bf080924		NA	NO	YES
1. Chromatograms La	beled/Compounds Ide	ntified. (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 Fre series	equency - Performed e	very 24 hours for 600 series and 12 hours for 8000			_
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 Met:					✓
a. Initial calibration If not met, list those c		ecoveries which fall outside the acceptable range.			✓
b. Continuous Calil If not met, list those c		<u>✓</u>			
2		n CCC. And Hexachlorocyclopentadiene is biased hig ll be reanalyzed with a properly passing CCC.	gh in the CCC if any samples	are	
6. Blank Contamination - If yes, list compounds and concentrations in each blank:					
a. B/N Fraction					

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 and RPD failed for some compound in P3460-01MS/MSD and P3509-01MS/MSD due to matrix interference action is required.	ence. No correc	<u>etive</u>
d. Acid Fraction		
9. Internal Standard Area/Retention Time Shift Meet Criteria Comments:	<u>√</u>	
Internal standard fail in P3507-01 and P3394-02. I.S failed in the sample P3509-01 and its associated MD/MSD due to matrix interference. No corrective action is required.		
10. Extraction Holding Time Met If not met, list number of days exceeded for each sample:		<u>√</u>
11. Analysis Holding Time Met		
If not met, list number of days exceeded for each sample:	<u> </u>	

ADDITIONAL COMMENTS:

One surrogate (Terphenyl-d14) is marginally biased high in PB162358BS. The data will be used for hardcopies. Recovery of compound #74,80 failed biased high in PB162358BS which Hit are not present in any associated samples. Recovery of compound #15,74,80 failed biased high in PB162423BS which Hit are not present in any associated samples.

Recovery of compound #15,74,80,89 failed biased high in PB162489BS which Hit are not present in any associated samples. The sample P3507-01 had to be analyzed with 5X dilution due to dirty and viscous matrix with was difficult to inject otherwise. And internal standard failed even in this diluted sample proving matrix interference. Hence no corrective action is required.



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