

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

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

CHEMTECH PROJECT	NUMBER: Bf032525			
SequenceID :	Bf032525	NA	NO	YES
1. Chromatograms Labele	l/Compounds Identified. (Field samples and Method Bla	unks)		
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 Freque series	cy - Performed every 24 hours for 600 series and 12 hou	rs for 8000		
	itial Calibration performed within 30 days before sample calibration performed within 24 hours of sample analysis urs for 8000 series			
5. GC/MS Calibration Me	:	<u>✓</u>		
a. Initial calibration Meet Criteria If not met, list those compounds and their recoveries which fall outside the acceptable range.				
Compound #77 is bias	d low in the CCC but not present in the parameters list o	f any associated samples.		
b. Continuous Calibrati If not met, list those comp	on(CCC) Meet Criteria ounds and their recoveries which fall outside the acceptal	→ble range.		
<ul><li>6. Blank Contamination -</li><li>a. B/N Fraction</li></ul>	f yes, list compounds 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.	<u>✓</u>	 
<ul><li>a. B/N Fraction</li><li>d. Acid Fraction</li></ul>		
<ul><li>8. Matrix Spike/Matrix Spike Duplicate Recoveries Meet Criteria</li><li>If not met, list those compounds and their recoveries which fall outside the acceptable range.</li><li>a. B/N Fraction</li></ul>	<u> </u>	 
d. Acid Fraction		
9. Internal Standard Area/Retention Time Shift Meet Criteria Comments:	<u>✓</u>	 
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 the PB167274BL/BS, PB167261BL, PB167193TB, PB167230BL, PB167230BSD, PB167254BL, PB167254BS and also 2,4,6-Tribromophenol is biased high in PB167274BS. Recovers of a very few compounds are slightly out of QC limits for PB167274BS, PB167261BS, PB167230BS/BSD. The data will be used for hardcopies.



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