

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

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

CHEMTECH PROJE	CT NUMBER:	bf042225			
SequenceID :	bf042225		NA	NO	YES
1. Chromatograms La	beled/Compounds Identi	fied. (Field samples and Method Blanks)	<u> </u>		
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 every 24 hours for 600 series and 12 hours for 8000 series					
analysis and contin	•	ormed within 30 days before sample ed within 24 hours of sample analysis	<u>√</u>		
5. GC/MS Calibration	Met:		<u>_</u>		
a. Initial calibration If not met, list those c		veries which fall outside the acceptable range.	<u> </u>		
	oration(CCC) Meet Criter ompounds and their reco	ria veries which fall outside the acceptable range.	<u></u>		
Compound #85 is properly passing C	-	if any samples are found with hit of this compo	ound they will be re-analyzed	again with a	
6. Blank Contamination	on - If yes, list compound	ls and concentrations in each blank:	<u>✓</u>		
a. B/N Fraction					

d. Acid Fraction

<ul><li>7. Surrogate Recoveries Meet Criteria</li><li>If not met, list those compounds and their recoveries which fall outside the acceptable ranges.</li><li>a. B/N Fraction</li></ul>	<u>✓</u>	 
d. Acid Fraction		
<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:	<b>√</b>	 
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:

A very few compounds are biased high in the PB167672BS/BSD. The data will be used for hardcopies.



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