

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

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

CHEMTECH PROJEC	CT NUMBER:	BP052325			
SequenceID :	BP052325		NA	NO	YES
1. Chromatograms Lal	peled/Compounds Ider	tified. (Field samples and Method Blanks)			✓
2. GC/MS Tuning Spo (NOTE THAT THERE		eet Criteria Criteria RITERIA FOR NY ASP CLP, CLP AND NJ	I)		<b>√</b>
3. GC/MS Tuning Free series	quency - Performed ev	ery 24 hours for 600 series and 12 hours for 8	8000		✓
analysis and continu		erformed within 30 days before sample ned within 24 hours of sample analysis			<b>√</b>
5. GC/MS Calibration	Met:				✓
a. Initial calibration If not met, list those co		coveries which fall outside the acceptable ran	1ge		✓
•		and Compounds #41 & 56 are biased high, it a properly passing CCC.	f any samples are found with hit o	of these	
	pration(CCC) Meet Cri compounds and their re	teria coveries which fall outside the acceptable ran	nge.	<u> </u>	
<ol> <li>Blank Contaminatic</li> <li>a. B/N Fraction</li> </ol>	n - If yes, list compou	nds and concentrations in each blank:		✓	

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			
<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>	
Internal standard fail in Q2095-08 which is not associated for required compound.			
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:

Rahul



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