

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

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

CHEMTECH PROJEC	T NUMBER: bp050225			
SequenceID :	bp050225	NA	NO	YES
1. Chromatograms Lab	eled/Compounds Identified. (Field samples and Method Bla	unks)		✓
<b>e</b> ,	cifications. DFTPP Meet Criteria Criteria ARE DIFFERENT CRITERIA FOR NY ASP CLP, CLP AN	ND NJ)		_✓
3. GC/MS Tuning Freq series	uency - Performed every 24 hours for 600 series and 12 hour	rs for 8000		<b>√</b>
analysis and continu	- Initial Calibration performed within 30 days before sample ing calibration performed within 24 hours of sample analysis 2 hours for 8000 series			✓
5. GC/MS Calibration	Met:			✓
a. Initial calibration If not met, list those co	Meet Criteria mpounds and their recoveries which fall outside the acceptat	ble range. ———		✓
If not met, list those co	ration(CCC) Meet Criteria mpounds and their recoveries which fall outside the acceptate und #77 which is not present in parameter list of the associat	-	<u>√</u>	
-	n - If yes, list compounds and concentrations in each blank:			

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			
8. Matrix Spike/Matrix Spike Duplicate Recoveries Meet Criteria If not met, list those compounds and their recoveries which fall outside the acceptable range.		<u>✓</u>	
a. B/N Fraction Recovery and RPD failed for some compound in Q1916-01MS/MSD and Q1928-01MS/MSD due to action is required.	matrix interfere	nce. No correc	<u>tive</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:		✓	

# ADDITIONAL COMMENTS:

Recovery of only two compounds are biased high in PB167823BS but their Hit is not present in any associated samples. The data will be used for hardcopies.



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