

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

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

CHEMTECH PROJECT	NUMBER:	bf120924				
SequenceID :	bf120924			NA	NO	YES
1. Chromatograms Label	ed/Compounds Ide	ntified. (Field samples an	d Method Blanks)			✓
2. GC/MS Tuning Specifications. DFTPP Meet Criteria Criteria (NOTE THAT THERE ARE DIFFERENT CRITERIA FOR NY ASP CLP, CLP AND NJ)						<b>_</b>
3. GC/MS Tuning Freque series	ency - Performed e	very 24 hours for 600 series	es and 12 hours for 8000			✓
	g calibration perfor	erformed within 30 days b med within 24 hours of sa s				✓
5. GC/MS Calibration M	et:					✓
a. Initial calibration M If not met, list those com		coveries which fall outsid	e the acceptable range.			<b>√</b>
	pounds and their re	coveries which fall outsid	le the acceptable range. 0783.D) but not present in an	ny samples so CCC is ok	✓	
6. Blank Contamination	- If yes, list compo	inds and concentrations ir	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 <u>The Recovery and RPD Fail for Some Compounds in P5095-04MS/MSD due to Matrix integration</u></li> </ul>	terference	<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:		✓	

ADDITIONAL COMMENTS:

Recovery is biased high for some compounds in PB1653432BS/BSD and PB165435BS. The data will be used for hardcopies.

Hitesh



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