

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

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

CHEMTECH PROJEC	CT NUMBER: bf052325			
SequenceID :	bf052325	NA	NO	YES
1. Chromatograms Lal	peled/Compounds Identified. (Field samples and Method Blan	nks)		✓
<b>e</b> 1	ecifications. DFTPP Meet Criteria Criteria E ARE DIFFERENT CRITERIA FOR NY ASP CLP, CLP AN	ND NJ)		_ <b>√</b>
3. GC/MS Tuning Free series	quency - 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 ompounds and their recoveries which fall outside the acceptab	ole range.		<b>√</b>
	ration(CCC) Meet Criteria ompounds and their recoveries which fall outside the acceptab	ole range.		✓
<ol> <li>Blank Contaminatic</li> <li>a. B/N Fraction</li> </ol>	n - If yes, list compounds and concentrations in each blank:		<u> </u>	

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>Recovery and RPD fail for some compound in Q2097-03MSD, Q2095-04MS/MSD, Q2109-01MS/M</u></li> </ul>	SD due to mati	rix interference	
d. Acid Fraction			
9. Internal Standard Area/Retention Time Shift Meet Criteria Comments:		✓	
Internal standard fail in Q2102-01,03.			
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:		_ <b>√</b>	

# ADDITIONAL COMMENTS:

Recovery of 3,3-Dichlorobenzidine is marginally biased low in the PB168126BS. The data will be used for hardcopies.



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