

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

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

CHEMTECH PROJE	CT NUMBER:	bf030625			
SequenceID :	bf030625		NA	NO	YES
1. Chromatograms La	beled/Compounds Ide	entified. (Field samples and 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 Fre series	equency - Performed e	very 24 hours for 600 series and 12 hours for 8000			<b>√</b>
analysis and contin		performed within 30 days before sample rmed within 24 hours of sample analysis es			✓
5. GC/MS Calibration	n Met:				✓
a. Initial calibration If not met, list those c		ecoveries which fall outside the acceptable range.			✓
	oration(CCC) Meet C ompounds and their r	riteria ecoveries which fall outside the acceptable range.		<u>√</u>	
•		h and compounds #23,26,32,54,56,57,62,65,77 are bill with Hit of this compound they will be analyzed aga	6		
6. Blank Contamination	on - If yes, list compo	unds and concentrations in each blank:		✓	
a. B/N Fraction					

d. Acid Fraction

7. Surrogate Recove If not met, list those	ries Meet Criteria compounds and their recoveries which fall outside the acceptable ranges.			✓
a. B/N Fraction				
d. Acid Fraction				
•	rix Spike Duplicate Recoveries Meet Criteria		✓	
If not met, list those	compounds and their recoveries which fall outside the acceptable range.			
a. B/N Fraction	The Recovery is biased low in 3,3-Dichlorobenzidine and Benzidine and biased high in Atrazine in the to matrix interference. Hence, no corrective action is required.	<u>e samples Q14</u>	92-01MS/MSD c	<u>lue</u>
d. Acid Fraction				
9. Internal Standard Comments:	Area/Retention Time Shift Meet Criteria			<b>√</b>
10. Extraction Holdi If not met, list numb	ng Time Met er of days exceeded for each sample:			<b>√</b>
11. Analysis Holding If not met, list numb	g Time Met er of days exceeded for each sample:		<u> </u>	

# ADDITIONAL COMMENTS:

The Recovery is biased high for very few compounds in PB166984BS and PB166982BS/BSD. These compounds are not present in any associated samples. This data will be used for hardcopies.



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