

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 Id	ntified. (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)					✓
3. GC/MS Tuning Fre series	equency - Performed e	very 24 hours for 600 series and 12 hours for	8000		√
analysis and contin		erformed within 30 days before sample rmed within 24 hours of sample analysis			<u> </u>
5. GC/MS Calibration	n Met:				✓
a. Initial calibratio If not met, list those c		ecoveries which fall outside the acceptable rar	nge		√
	bration(CCC) Meet C compounds and their r	iteria ecoveries which fall outside the acceptable rar	nge.	∕	
	marginally high and o present in any associa	ompound #23,26,32,54,56,57,62,65,77 are fai ed samples.	il high side in SSTDCCC (BF1-	41858.D).	
6. Blank Contaminati	on - If yes, list compo	unds and concentrations in each blank:		√	
a. B/N Fraction					

d. Acid Fraction

7. Surrogate Recoveries Meet CriteriaIf not met, list those compounds and their recoveries which fall outside the acceptable ranges.a. B/N Fraction	—		✓
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. a. B/N Fraction <u>The Recovery fail for some compound in Q1492-01MS/MSD due to matrix interference.</u> 		<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:		<u>✓</u>	

ADDITIONAL COMMENTS:

The Recovery fail for some compound in PB166984BS and PB166982BS/BSD. Which hit are not present in any associated samples.



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