

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

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

CHEMTECH PROJE	CT NUMBER: bp041625				
SequenceID :	bp041625		NA	NO	YES
1. Chromatograms La	beled/Compounds Identified. (Field samples and I	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	quency - Performed every 24 hours for 600 series	and 12 hours for 8000			_✓
analysis and contin	- Initial Calibration performed within 30 days bef ung calibration performed within 24 hours of sam 2 hours for 8000 series				✓
5. GC/MS Calibration	Met:				✓
a. Initial calibration If not met, list those c	n Meet Criteria compounds and their recoveries which fall outside t	he acceptable range.			✓
	oration(CCC) Meet Criteria compounds and their recoveries which fall outside t	the acceptable range.			
Compound #9 and with a properly pa	# 69 are biased high in the CCC, if any samples as asing CCC.	re found with hit of these compounds	they will be re-	analyzed	
6. Blank Contamination	on - If yes, list compounds and concentrations in ea	ach blank:		√	
a. B/N Fraction					

d. Acid Fraction

7. Surrogate Recoveries Meet Criteria If 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 CriteriaIf not met, list those compounds and their recoveries which fall outside the acceptable range.a. B/N Fraction	<u>✓</u>		
d. Acid Fraction			
9. Internal Standard Area/Retention Time Shift Meet Criteria Comments:		<u>✓</u>	
Internal standard failed in the sample Q1791-10 due to limited volume availability of this water sample it can n re-extracted, Hence it will be re-analyzed.	ot be		
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 a very few compound are biased high in PB167599BS/BSD and PB167606BS but their hit is not present in any associated samples. The data will be used for hardcopies.



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