

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

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

CHEMTECH PROJE	CT NUMBER:	bn110424			
SequenceID :	bn110424		NA	NO	YES
1. Chromatograms La	beled/Compounds Ide	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			_		
analysis and contin	•	erformed within 30 days before sample med within 24 hours of sample analysis s			✓
5. GC/MS Calibration	n Met:				✓
a. Initial calibration If not met, list those c		coveries which fall outside the acceptable range.			✓
b. Continuous Calibration(CCC) Meet Criteria If not met, list those compounds and their recoveries which fall outside the acceptable range.				<u> </u>	
•	in the SSTDCCC(BN ne associated samples.)34837.D) and SSTDCCC(BN034850.D) are bias	sed low but they are not present in		
6. Blank Contamination - If yes, list compounds and concentrations in each blank:					
a. B/N Fraction					

d. Acid Fraction

7. Surrogate Recoveries M	leet Criteria		1	
If not met, list those comp	ounds and their recoveries which fall outside the acceptable ranges.			
a. B/N Fraction Sur	rogate Failed in the sample P4665-02, due to the limited volume of this water sample it will be re-	-analyzed and n	ot re-extracted.	
d. Acid Fraction				
	ike Duplicate Recoveries Meet Criteria ounds and their recoveries which fall outside the acceptable range.	<u> </u>		
a. B/N Fraction				
d. Acid Fraction				
9. Internal Standard Area/I Comments:	Retention Time Shift Meet Criteria		<u> </u>	
Internal Standards Fail compound.	ed in P4652-02,03,05,06,P4600-05, PB164594BS/BSD which are not associated for red	quired		
10. Extraction Holding Tir If not met, list number of c	ne Met lays exceeded for each sample:			✓
11. Analysis Holding Time If not met, list number of c	e Met lays exceeded for each sample:		√	

ADDITIONAL COMMENTS:

Recovery fail for some compound in PB164594BS/BSD which are not present in parameter list of the associated samples. The data will be used for hardcopies.



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