

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

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

CHEMTECH PROJE	CT NUMBER:	bn102424			
SequenceID :	bn102424		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)					<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		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>√</b>
	bration(CCC) Meet Cr compounds and their re	iteria coveries which fall outside the acceptable range.			✓
Some compound in samples.	n the SSTDCCC(BN0	34701.D) are biased low but they are not present ir	n parameter list of the associated		
6. Blank Contamination	on - If yes, list compo	inds and concentrations in each 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	Surrogates Failed in the sample P4438-03 and P4440-01, due to the limited availability of these water re-analyzed to confirm the matrix interference and will not be re-extracted.	samples they w	<u>ill be</u>			
d. Acid Fraction						
	x Spike Duplicate Recoveries Meet Criteria ompounds and their recoveries which fall outside the acceptable range.	<u>✓</u>	·			
a. B/N Fraction						
d. Acid Fraction						
9. Internal Standard Ar Comments:	rea/Retention Time Shift Meet Criteria		<u> </u>			
Internal Standard F	ailed in PB164229BL,PB164229BS/BSD,PB164230BS which are not associated for required	d compound.				
10. Extraction Holding If not met, list number	g Time Met of days exceeded for each sample:			✓		
11. Analysis Holding If not met, list number	Time Met of days exceeded for each sample:		✓			

# ADDITIONAL COMMENTS:

Recovery of 1,4-Dioxane is marginally biased low in PB164229BS. The data will be used for hardcopies. Recovery failed for some compound in PB164229BS/BSD and PB164230BS which are not present in parameter list of the associated samples. The sample P4432-01 had to be analyzed with 20X dilution due to oily nature of the sample which can also be observed from the abnormal chromatogram. Hence no corrective action is required.



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