

d. Acid Fraction

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

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

CHEMTECH PROJECT NU	MBER: <u>be1031</u>	124				
SequenceID:	be103124	_		NA	NO	YES
1. Chromatograms Labeled/0	Compounds Identified. (Fig.	eld samples and Method Blanks)				_
2. GC/MS Tuning Specifica (NOTE THAT THERE ARE		a Criteria FOR NY ASP CLP, CLP AND NJ)				_
3. GC/MS Tuning Frequency series	- Performed every 24 hour	rs for 600 series and 12 hours for 8000				✓
4. GC/MS Calibration - Initi analysis and continuing ca for 600 series and 12 hour	libration performed within	ithin 30 days before sample 24 hours of sample analysis				_
5. GC/MS Calibration Met:						✓
a. Initial calibration Meet If not met, list those compou		hich fall outside the acceptable range.				_
b. Continuous Calibration If not met, list those compou		hich fall outside the acceptable range.			_	
Compounds #6,77 are bia with a properly passing C		any samples are found with hit of these	compounds they wi	ll be re-anal	yzed	
6. Blank Contamination - If	ves, list compounds and cor	ncentrations in each blank:			<u> </u>	
a. B/N Fraction						

7. Surrogate Recover If not met, list those	ries Meet Criteria compounds and their recoveries which fall outside the acceptable ranges.					
a. B/N Fraction	Surrogates failed in the sample P4578-02, due to the limited volume of this water sample it will be re-analyzed to confirm the matrix interference and will not be re-extracted.					
d. Acid Fraction						
	rix Spike Duplicate Recoveries Meet Criteria compounds and their recoveries which fall outside the acceptable range.					
a. B/N Fraction	Recovery of certain compounds were affected even in the diluted sample of both P4640-01MS and Mence no corrective action is required.	MSD proving matrix interfer	rence.			
d. Acid Fraction						
9. Internal Standard A	Area/Retention Time Shift Meet Criteria					
Internal standard	failed in P4616-01MSD.					
10. Extraction Holdin If not met, list number	ng Time Met er of days exceeded for each sample:					
11. Analysis Holding If not met, list number	g Time Met er of days exceeded for each sample:					
The samples P463	MMENTS: some compound in PB164533BS/BSD which hit are not present in any associated samples 39-03, P4640-01 and its associated MS/MSD had to be analyzed with 5X dilution each due s difficult to inject otherwise. Hence no corrective action is required.					

<u>krunal</u>
Analyst

Date



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