

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

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

CHEMTECH PROJECT	NUMBER:	bf081424			
SequenceID :	bf081424		NA	NO	YES
1. Chromatograms Labe	led/Compounds Ide	entified. (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 Frequ	ency - Performed e	very 24 hours for 600 series and 12 hours for 8000 serie	es		1
	ng calibration perfo	performed within 30 days before sample rmed within 24 hours of sample analysis		_	✓
5. GC/MS Calibration M	let:				✓
a. Initial calibration M If not met, list those com		ecoveries which fall outside the acceptable range.			✓
	pounds and their r	iteria ecoveries which fall outside the acceptable range. C but not present in parameter list of the associated sam		<u> </u>	
-	-	unds and concentrations in each blank:	-	√	

d. Acid Fraction

7. Surrogate RecoverIf not met, list thosea. B/N Fraction	ries Meet Criteria		
d. Acid Fraction			
•	rix Spike Duplicate Recoveries Meet Criteria	<u>✓</u>	
a. B/N Fraction	Recovery and RPD failed for some compound in P3466-02MS, P3466-03MSD, P3475-03MS, P3475-04MSD of interference. No corrective action is required.	due to matrix	
d. Acid Fraction			
9. Internal Standard . Comments:	Area/Retention Time Shift Meet Criteria	√	
	fail in P3548-06, P3548-12, P3475-01, P3526-01. One I.S (Acenaphthene-d10) failed in the sample us matrix hence this result will be final.	P3549-01 due	
10. Extraction Holdi If not met, list numb	ng Time Met er of days exceeded for each sample:		<u>√</u>
11. Analysis Holding If not met, list numb	g Time Met		

ADDITIONAL COMMENTS:

Recovery of some compound fail high side in PB162577BS which Hit are not present in any associated samples. Surrogates failed in the sample P3543-01 due to the presence of nontargeted hydrocarbons which can be witnessed by the chromatogram. Hence this analysis will be final. I.S failed in the sample P3475-02 and its associated MS/MSD (P3475-03MS, P3475-04MSD) due to matrix interference. Hence no corrective action is required.



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