

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

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

CHEMTECH PROJECT	NUMBER: bm082124			
SequenceID :	bm082124	NA	NO	YES
1. Chromatograms Label	d/Compounds Identified. (Field samples and Method	od 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 Freque	ncy - Performed every 24 hours for 600 series and 1	2 hours for 8000 series		1
	nitial Calibration performed within 30 days before says calibration performed within 24 hours of sample arours for 8000 series	•		
5. GC/MS Calibration M	et:			<b>√</b>
a. Initial calibration M If not met, list those com	eet Criteria bounds and their recoveries which fall outside the ac	cceptable range.		✓
	on(CCC) Meet Criteria bounds and their recoveries which fall outside the ac 9 are biased high in the CCC.	cceptable range.	✓	
-	If yes, list compounds and concentrations in each b	lank:	<u> </u>	

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 Criteria If not met, list those compounds and their recoveries which fall outside the acceptable range.		<u> </u>	
a. B/N Fraction Recovery failed for some compound in P3617-08MS/MSD due to matrix interference. No corrective a	ction is requir	ed.	
d. Acid Fraction			
9. Internal Standard Area/Retention Time Shift Meet Criteria Comments:			<u> </u>
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:		<u>✓</u>	
ADDITIONAL COMMENTS:			

Terphenyl-d14 was marginally biased high in PB162815BS and 2-Fluorobiphenyl & Terphenyl-d14 were marginally biased high in PB162782BL. The data will be used for the hard copies.

Recovery of some compound are biased high in PB162815BS and PB162782BS but there Hit are not present in any associated samples.

Rahul



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