

284 Sheffield Street, Mountainside, New Jersey 07092, Phone : 908 789 8900, Fax : 908 789 8922

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

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

CHEMTECH PROJECT NU	MBER: <u>BM02282</u>	25			
SequenceID:	BM022825		NA	NO	YES
1. Chromatograms Labeled/	Compounds Identified. (Field	samples and Method Blanks)			_
	tions. DFTPP Meet Criteria C DIFFERENT CRITERIA FO	Criteria OR NY ASP CLP, CLP AND NJ)			_
3. GC/MS Tuning Frequency series	- Performed every 24 hours f	for 600 series and 12 hours for 8000			_
	al Calibration performed within libration performed within 24 s for 8000 series				_
5. GC/MS Calibration Met:					✓
a. Initial calibration Meet If not met, list those compou		h fall outside the acceptable range.			_
Compounds #26,32,41,54 will be analyzed again w		d high in the CCC. If any samples are found v	with Hit of this compou	and they	
b. Continuous Calibration If not met, list those compou	· · · · · · · · · · · · · · · · · · ·	h fall outside the acceptable range.			
6. Blank Contamination - If a. B/N Fraction	ves, list compounds and conce	entrations in each blank:		<u> </u>	

d. Acid Fraction

7. Surrogate Recoveries Meet Criteria If not met, list those compounds and their recoveries whi	ich fall outside the acceptable ranges.		<u> </u>	
a. B/N Fraction	. •			
d. Acid Fraction Typically two phenolic surrogates fai chromatogram. Therefore this analys	iled in the sample Q1456-02 due to matrix interference which is will be final.	ch can be observed b	by the abnormal	
8. Matrix Spike/Matrix Spike Duplicate Recoveries Mee			<u>✓</u>	
If not met, list those compounds and their recoveries whi	ich fall outside the acceptable range.			
a. B/N Fraction Recovery failed for some compounds in	in Q1458-01MS/MSD due to matrix interference. Hence, no	corrective action is	required.	
d. Acid Fraction				
9. Internal Standard Area/Retention Time Shift Meet Crit Comments:	teria			<b>✓</b>
10. Extraction Holding Time Met If not met, list number of days exceeded for each sample	e:			_
11. Analysis Holding Time Met If not met, list number of days exceeded for each sample	»:		<u> </u>	
	in PB166930BL. One surrogate Terphenyl-d14 is sligh			

One surrogate Terphenyl-d14 is slightly biased high in PB166930BL. One surrogate Terphenyl-d14 is slightly biased high and also recovery of few compounds were biased high in PB166930BS. The data will be used for hardcopies. The sample Q1457-01 had to be analyzed with 5X dilution due to dirty and viscous matrix. Hence, this analysis is final.

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	Rahul	02/28/2025	



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