

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

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

CHEMTECH PROJE	CT NUMBER:	bn082324				
SequenceID :	bn082324			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)						√
3. GC/MS Tuning Fre series	equency - Performed e	very 24 hours for 600 series and 12 hours for 8000	-			_
analysis and contin	•	erformed within 30 days before sample rend within 24 hours of sample analysis	-			✓
5. GC/MS Calibration	n Met:		_			✓
a. Initial calibration If not met, list those c		ecoveries which fall outside the acceptable range.	-			✓
	bration(CCC) Meet Co ompounds and their re	iteria coveries which fall outside the acceptable range.	-		<u> </u>	
Compound #20 is of the associated s		bound #34 is biased High in the SSTDCCC(BN0335	599.D) but are not in	the parame	eters list	
6. Blank Contamination	on - If yes, list compo	unds 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	h fall outside the acceptable ranges.			✓
a. B/N Fraction				
d. Acid Fraction				
8. Matrix Spike/Matrix Spike Duplicate Recoveries Meet C If not met, list those compounds and their recoveries which			<u>✓</u>	
a. B/N Fraction <u>The Recovery and RPD Fail for Some Corequired.</u>	ompounds in P3641-02MSD,P3707-03MSD due to Matrix i	nterference. No	corrective actio	<u>n is</u>
d. Acid Fraction				
9. Internal Standard Area/Retention Time Shift Meet Criter Comments:	ia			<u> </u>
10. Extraction Holding Time Met If not met, list number of days exceeded for each sample:				<u> </u>
 Analysis Holding Time Met If not met, list number of days exceeded for each sample: 			√	

ADDITIONAL COMMENTS:

In PB162859BS Compound #34 is biased High but is not part of associated Sample parameter list. In PB162937BS Compound #24 is biased low but is not part of associated Sample parameter list.

Analyst



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