

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

GC/MS VOA CONFORMANCE/NON-CONFORMANCE SUMMARY

CHEMTECH PROJECT NUMBER: vn071825

SequenceID : vn071825

	NA	NO	YES
1. Chromatograms Labeled/Compounds Identified. (Field samples and Method Blanks)	_____	_____	<u>✓</u>
2. GC/MS Tuning Specifications. BFB Meet Criteria (NOTE THAT THERE ARE DIFFERENT CRITERIA FOR NY ASP CLP, CLP AND NJ)	_____	_____	<u>✓</u>
3. GC/MS Tuning Frequency - Performed every 24 hours for 600 series and 12 hours for 8000 series	_____	_____	<u>✓</u>
4. GC/MS Calibration - Initial Calibration performed before sample analysis and continuing calibration performed within 24 hours of sample analysis for 600 series and 12 hours for 8000 series	_____	_____	<u>✓</u>
5. GC/MS Calibration Requirements	_____	_____	<u>✓</u>
5a. Initial Calibration Meet Criteria If not met, list those compounds and their recoveries which fall outside the acceptable ranges.	<u>✓</u>	_____	_____
5b. Continuous Calibration(CCC) Meet Criteria If not met, list those compounds and their recoveries which fall outside the acceptable ranges.	_____	_____	<u>✓</u>
6. Blank Contamination - If yes, list compounds and concentrations in each blank:	_____	<u>✓</u>	_____
7. Surrogate Recoveries Meet Criteria If not met, list those compounds and their recoveries which fall outside the acceptable ranges.	_____	<u>✓</u>	_____
8a. 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>	_____	_____
8b. Blank Spike/Blank spike Duplicate Recoveries Meet Criteria If not met, list those compounds and their recoveries which fall outside the acceptable range. carbon disulfide recovery failed low	_____	<u>✓</u>	_____
9. Internal Standard Area/Retention Time Shift Meet Criteria	_____	_____	<u>✓</u>
Comments:			

MMDadoda

Analyst

07/21/2025

Date

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NA NO YES

10. Analysis Holding Time Met

If not met, list number of days exceeded for each sample:

____✓

ADDITIONAL COMMENTS:

Q2585-01 sample analyzed at straight 5x dilution as per previous history on the sample.

Q2594-01 sample is dark yellow and foamy in nature ,lab has analyzed staright 5x dilution on the sample

Sequence QC(BS,BSD) failing for low recovery,no sample has been reported for any positive hit for the analyte,sdg 2630 and 2585 do not required that analyte and 2594 lab has analyzed at elevated RL (5X DILUTION) and sample has no any positive hit of any analyte,but as mentioned earlier samples is dark yellow in color and foamy lab has not analyzed any straight analysis for the samples ,lab will use all analysis as final analysis

MMDadoda

Analyst

07/21/2025

Date



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