

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

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

| CHEMTECH PROJEC | CT NUMBER: bf013025 | | | |
|---|--|---------------|----------|----------|
| SequenceID : | bf013025 | NA | NO | YES |
| 1. Chromatograms Lal | eled/Compounds Identified. (Field samples and Method E | Blanks) | | ✓ |
| e 1 | cifications. DFTPP Meet Criteria Criteria ARE DIFFERENT CRITERIA FOR NY ASP CLP, CLP A | AND NJ) | | _ |
| 3. GC/MS Tuning Free series | uency - Performed every 24 hours for 600 series and 12 ho | ours for 8000 | | _√ |
| analysis and continu | - Initial Calibration performed within 30 days before samp ing calibration performed within 24 hours of sample analy 2 hours for 8000 series | | | ✓ |
| 5. GC/MS Calibration | Met: | | | ✓ |
| a. Initial calibration If not met, list those co | Meet Criteria mpounds and their recoveries which fall outside the accept | table range. | | √ |
| | ration(CCC) Meet Criteria mpounds and their recoveries which fall outside the accept | table range. | | ✓ |
| Blank Contaminatic a. B/N Fraction | n - If yes, list compounds and concentrations in each blank | c | <u>√</u> | |

d. Acid Fraction

| 7. Surrogate Recoveries Meet CriteriaIf not met, list those compounds and their recoveries which fall outside the acceptable ranges.a. B/N Fraction | | <u> </u> |
|---|--------------|----------|
| 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. a. B/N Fraction <u>The Recovery Fail for Some Compounds in Q1209-05MS/MSD due to Matrix interference</u> | <u> </u> | |
| d. Acid Fraction | | |
| 9. Internal Standard Area/Retention Time Shift Meet Criteria Comments: | | _ |
| 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:

In PB166335BS Compound #41 Failing High which Hit not present in any associated samples.



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