

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

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

| CHEMTECH PROJEC   | CT NUMBER: bf061825  |                |          |            |
|---|--|----------------|----------|------------|
| SequenceID :  | bf061825   | NA             | NO       | YES        |
| 1. Chromatograms Lal  | beled/Compounds Identified. (Field samples and Method Bla  | anks)          |          | ✓          |
| <b>e</b> 1  | ccifications. DFTPP Meet Criteria Criteria<br>E ARE DIFFERENT CRITERIA FOR NY ASP CLP, CLP AN  | ND NJ)         |          | _ <b>√</b> |
| 3. GC/MS Tuning Free<br>series                                  | quency - Performed every 24 hours for 600 series and 12 hou  | urs for 8000   |          | _ ✓        |
| analysis and continu  | <ul> <li>Initial Calibration performed within 30 days before sample<br/>ing calibration performed within 24 hours of sample analysis</li> <li>hours for 8000 series</li> </ul> |                |          | ✓          |
| 5. GC/MS Calibration  | Met:   |                |          | ✓          |
| a. Initial calibration<br>If not met, list those co             | Meet Criteria<br>ompounds and their recoveries which fall outside the acceptal   | ble range. ——— |          | ✓          |
|   | ration(CCC) Meet Criteria ompounds and their recoveries which fall outside the acceptal  | ble range.     |          | ✓          |
| <ol> <li>Blank Contaminatic</li> <li>a. B/N Fraction</li> </ol> | n - If yes, list compounds and concentrations in each blank:   |                | <u>_</u> |            |

d. Acid Fraction

| <ul> <li>7. Surrogate Recoveries Meet Criteria</li> <li>If not met, list those compounds and their recoveries which fall outside the acceptable ranges.</li> <li>a. B/N Fraction</li> </ul> | <br>✓        |
|---|--------------|
| <ul> <li>d. Acid Fraction</li> <li>8. Matrix Spike/Matrix Spike Duplicate Recoveries Meet Criteria</li> </ul>   | 1            |
| a. B/N Fraction       Recovery and RPD failed in the Q2339-04MS/MSD due to matrix interface. No corrective action is required.  | <br><u> </u> |
| <ul> <li>d. Acid Fraction</li> <li>9. Internal Standard Area/Retention Time Shift Meet Criteria</li> </ul>  | <br><u> </u> |
| Comments:<br>10. Extraction Holding Time Met  |              |
| If not met, list number of days exceeded for each sample:   | <br><u> </u> |
| 11. Analysis Holding Time Met<br>If not met, list number of days exceeded for each sample:  | <br><b>√</b> |

# ADDITIONAL COMMENTS:

Recovery of some compounds are biased high in PB168505BS due to matrix interference. The data will be used for hardcopies.



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