

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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