

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

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

CHEMTECH PROJE	CT NUMBER:	bf022625				
SequenceID :	bf022625			NA	NO	YES
1. Chromatograms La	beled/Compounds Id	ntified. (Field samples and Method Blar	nks)			✓
2. GC/MS Tuning Specifications. DFTPP Meet Criteria Criteria (NOTE THAT THERE ARE DIFFERENT CRITERIA FOR NY ASP CLP, CLP AND NJ) —						<u> </u>
3. GC/MS Tuning Fre series	quency - Performed e	very 24 hours for 600 series and 12 hour	rs for 8000			<b>_</b>
analysis and contin		erformed within 30 days before sample rmed within 24 hours of sample analysis es				✓
5. GC/MS Calibration	Met:					✓
a. Initial calibration If not met, list those c		ecoveries which fall outside the acceptab	vle range.			<b>√</b>
b. Continuous Calibration(CCC) Meet Criteria If not met, list those compounds and their recoveries which fall outside the acceptable range.					<u>✓</u>	
	5 biased high in the S h proper passing CC	STDCCC(BF141775.D). If any samples 2.	are found with Hit of this con	mpound they	y will be	
6. Blank Contamination - If yes, list compounds and concentrations in each blank:					✓	
a. B/N Fraction						

d. Acid Fraction

7. Surrogate Recoveries Meet Criteria ✓						
a. B/N Fraction <u>Terphenyl-d14 surrogate is marginally biased high in PB166853BS. The data</u>	a will be used for hardcopies.					
d. Acid Fraction						
<ul><li>8. Matrix Spike/Matrix Spike Duplicate Recoveries Meet Criteria</li><li>If not met, list those compounds and their recoveries which fall outside the acceptable range</li><li>a. B/N Fraction</li></ul>	а	—	<u> </u>			
d. Acid Fraction						
9. Internal Standard Area/Retention Time Shift Meet Criteria Comments:			<u> </u>			
10. Extraction Holding Time Met If not met, list number of days exceeded for each sample:			<b>√</b>			
11. Analysis Holding Time Met If not met, list number of days exceeded for each sample:		✓				

#### ADDITIONAL COMMENTS:

The recovery is biased high for compound #80 in PB166853BS and the recovery is biased high for compound #84 in PB166875BS/BSD. The recovery is biased low for compound #88 in PB166875BS. This data will be used for hardcopies. The sample Q1428-01 had to be analyzed with 2X dilution due to dirty viscous and concentrated matrix. Hence this analysis will be will be final.



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