

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

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

CHEMTECH PROJEC	T NUMBER:	bf062525			
SequenceID :	bf062525		NA	NO	YES
1. Chromatograms Lab	eled/Compounds Ide	entified. (Field samples and Method Blanks)	✓		
2. GC/MS Tuning Spe (NOTE THAT THERE	<u> </u>				
3. GC/MS Tuning Frequency - Performed every 24 hours for 600 series and 12 hours for 8000 series					
	ing calibration perfo	performed within 30 days before sample rmed within 24 hours of sample analysis es	<u> </u>		
5. GC/MS Calibration	Met:		<u> </u>		
a. Initial calibration If not met, list those co		ecoveries which fall outside the acceptable range.	<u> </u>		
b. Continuous Calibr If not met, list those co		riteria ecoveries which fall outside the acceptable range.			
-		, but not present in the parameters list of the samples analy re found with hit of these compounds they will be re-analy			

 \checkmark

6. Blank Contamination - If yes, list compounds 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 fall outside the acceptable ranges.	<u>✓</u>			
a. B/N Fraction					
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 PB168592BSDRPD failed due to result diffraction between BS and BSD. Q2386-02MS/MSD Recovery failed due to matrix interference, and RPD failed due to result difference between MS and MSD. 					
d. Acid Fraction	ecovery failed due to matrix interference, and KPD failed due to result	anterence between MS and MSD.			
9. Internal Standard Area/Retention Time Comments:	Shift Meet Criteria	<u> </u>			
10. Extraction Holding Time Met If not met, list number of days exceeded f	or each sample:	_ ✓			
11. Analysis Holding Time Met If not met, list number of days exceeded f	or each sample:	<u>✓</u>			

ADDITIONAL COMMENTS:

Only one compound 3,3-Dichlorobenzidine is marginally biased low in the PB168592BS/BSD, PB168601BS. The data will be used for hardcopies.



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