

d. Acid Fraction

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

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

CHEMTECH PROJEC	T NUMBER:	bf123024			
SequenceID:	bf123024		NA	NO	YES
1. Chromatograms Labo	eled/Compounds Ide	entified. (Field samples and Method Blanks)			_
2. GC/MS Tuning Spec (NOTE THAT THERE		Meet Criteria Criteria CRITERIA FOR NY ASP CLP, CLP AND NJ)			✓
3. GC/MS Tuning Frequencies	uency - Performed e	very 24 hours for 600 series and 12 hours for 8000			<b>✓</b>
	ng calibration perfo	performed within 30 days before sample rmed within 24 hours of sample analysis es			_
5. GC/MS Calibration N	Met:				✓
a. Initial calibration		ecoveries which fall outside the acceptable range.			<b>✓</b>
b. Continuous Calibra If not met, list those con	, ,	riteria ecoveries which fall outside the acceptable range.		_	
Compounds #54,56, will be analyzed aga		ed high in CCC (BF141031.D) if any samples are founding CCC.	with Hit of this compound	d they	
6. Blank Contamination	- If yes, list compo	unds and concentrations in each blank:			
a. B/N Fraction					

7. Surrogate Recoveries Meet Criteria If not met, list those compounds and their recoveries which fall outside the acceptable ranges.	_
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 The Recovery and RPD Fail for Some Compounds in P5391-01MS/MSD due to Matrix interference. Hence, no corrective action is required.	
d. Acid Fraction	
9. Internal Standard Area/Retention Time Shift Meet Criteria ✓	
Internal Standard Failed in the samples P5386-01,03 due to the presence of nontargeted hydrocarbons which can be observed by the abnormal chromatogram. Hence, no corrective action is required. Even in the diluted sample P5318-01 internal standard failed due to concentrated matrix and the presence of nontargeted hydrocarbons which can be observed by the Extraction Holding Time Met abnormal chromatogram. Hence, no corrective action required. 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:	
ADDITIONAL COMMENTS:	

One surrogate Nitrobenzene-d5 was marginally biased high in both PB165872BS and PB165900BL. One surrogate 2,4,6-Tribromophenol was marginally biased high in PB165872BS. The data will be used for the hard copies. Recovery for some compounds were biased high in PB165875\BSD, PB165894BS and PB165872BS. This data will be used for hardcopies. The samples P5390-01 and P5318-01 had to be analyzed with 2X dilution each due to viscous matrix.



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