

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

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

CHEMTECH PROJEC	CT NUMBER: bf111424			
SequenceID :	bf111424	NA	NO	YES
1. Chromatograms Lal	peled/Compounds Identified. (Field samples and Method	d 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 Free series	quency - Performed every 24 hours for 600 series and 12	hours for 8000		_
analysis and continu	- Initial Calibration performed within 30 days before sar ning calibration performed within 24 hours of sample and 2 hours for 8000 series			<b>√</b>
5. GC/MS Calibration	Met:			✓
a. Initial calibration If not met, list those co	Meet Criteria ompounds and their recoveries which fall outside the acc	eptable range.		<b>√</b>
	ration(CCC) Meet Criteria ompounds and their recoveries which fall outside the acc	eptable range.		✓
<ol> <li>Blank Contaminatic</li> <li>a. B/N Fraction</li> </ol>	on - If yes, list compounds and concentrations in each bla	.nk:	<b>√</b>	

d. Acid Fraction

7. Surrogate Recover If not met, list those of	ies Meet Criteria	<u>✓</u>
a. B/N Fraction	Typically only two phenolic surrogates failed in the sample P4762-02 proving matrix interference, which can also l abnormal chromatogram. Hence this analysis will be final.	be witnessed by the
d. Acid Fraction		
1	ix Spike Duplicate Recoveries Meet Criteria compounds and their recoveries which fall outside the acceptable range.	<u>✓</u>
a. B/N Fraction	Recovery and RPD failed for some compound in P4788-01MS/MSD and P4807-05MS/MSD due to matrix interferent action is required.	nce. No corrective
d. Acid Fraction		
9. Internal Standard A Comments:	Area/Retention Time Shift Meet Criteria	✓
chromatogram. He	failed in the sample P4791-02 due to the oily matrix of the sample which can be observed by the abnormatic ence no corrective action is required.	l
Internal standard t 10. Extraction Holdir confirms the matr If not met, list numbe	failed even in the rerun samples of P4796-05, P4796-01, P4795-01, P4811-01 and P4816-01 Hence this ng Time.Met ix interference. er of days exceeded for each sample:	<b>_</b>
11. Analysis Holding	Time Met	4
If not met, list numbe	er of days exceeded for each sample:	•

ADDITIONAL COMMENTS:

Recovery failed for some compound in PB164911BS/BSD but their hit is not present in any of the associated samples. The data will be used for hardcopies. The sample P4793-01 had to be analyzed with 50X dilution due to the oily nature of the sample, and even in this diluted sample surrogates failed proving matrix interference which can also be witnessed by the abnormal chromatogram. Hence no corrective action is required.



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