

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 PROJECT NU	MBER:	BN022025				
SequenceID:	BN022025			NA	NO	YES
1. Chromatograms Labeled/C	Compounds Identif	ied. (Field samples and Method Blan	ıks)			_
2. GC/MS Tuning Specificat (NOTE THAT THERE ARE		t Criteria Criteria TERIA FOR NY ASP CLP, CLP ANI	D NJ)			<b>✓</b>
3. GC/MS Tuning Frequency series	- Performed ever	y 24 hours for 600 series and 12 hours	s for 8000			<b>✓</b>
	libration performe	ormed within 30 days before sample d within 24 hours of sample analysis				<u> </u>
5. GC/MS Calibration Met:						_
a. Initial calibration Meet If not met, list those compound		veries which fall outside the acceptable	le range.			<b>✓</b>
b. Continuous Calibration( If not met, list those compound	` '	ia veries which fall outside the acceptable	le range.			_
6. Blank Contamination - If y a. B/N Fraction	yes, list compound	s and concentrations in each blank:			<u> </u>	

7. C	in Mark Circle			,
7. Surrogate Recover If not met, list those	compounds and their recoveries which fall outside the acceptable ranges.		<del></del>	<u> </u>
a. B/N Fraction				
d. Acid Fraction				
•	rix Spike Duplicate Recoveries Meet Criteria compounds and their recoveries which fall outside the acceptable range.		<u> </u>	
<ul><li>a. B/N Fraction</li><li>d. Acid Fraction</li></ul>	Being a poor recovery compound recovery of 1,4-Dioxane is marginally biased low in the Q1380-10MS. compounds are biased high in the Q1380-11MSD but are not present in the parameters list of the sample. required.			
9. Internal Standard A	Area/Retention Time Shift Meet Criteria			✓
10. Extraction Holdin	ng Time Met er of days exceeded for each sample:			✓
11. Analysis Holding If not met, list numbe	g Time Met er of days exceeded for each sample:		<u> </u>	
be re-extracted ar	MMENTS: Methylnaphthalene-d10 is biased low in the sample Q1380-12. Due to the limited volume of this dollar however reanalysis of this sample will result in the same surrogate failure so this analysis will be compound is biased high in PB166758BS. Which hit are not present in any associated sample	ll be final.	rill not	

Rahul	<u>-</u>	02/20/2025
Analyst		Date



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