

Data Path : Z:\SVOASRV\HPCHEM1\BNA G\DATA\BG081120\
 Data File : BG046218.D
 Acq On : 11 Aug 2020 15:25
 Operator : CG/JU
 Sample : L3624-02
 Misc :
 ALS Vial : 7 Sample Multiplier: 1

Instrument :
 BNA_G
 ClientSampleId :
 SS-TP-1

Integration Parameters: rteint.p
 Integrator: RTE
 Smoothing : ON Filtering: 5
 Sampling : 1 Min Area: 3 % of largest Peak
 Start Thrs: 0.2 Max Peaks: 100
 Stop Thrs : 0 Peak Location: TOP

If leading or trailing edge < 100 prefer < Baseline drop else tangent >
 Peak separation: 5

Method : Z:\SVOASRV\HPCHEM1\BNA_G\METHODS\8270-BG073020.M
 Title : ASP BNA STANDARDS FOR 5 POINT CALIBRATION

Signal : TIC

peak #	R.T. min	first scan	max scan	last scan	PK TY	peak height	corr. area	corr. % max.	% of total
1	3.155	6	11	22	rVB	263053	365237	8.16%	1.781%
2	5.065	331	336	346	rVB	30606	48436	1.08%	0.236%
3	5.535	409	416	427	rBV	728345	1136240	25.38%	5.540%
4	7.115	677	685	695	rBV	726763	1246537	27.85%	6.078%
5	7.950	819	827	836	rBV	287201	472544	10.56%	2.304%
6	9.101	1014	1023	1032	rBV	473472	834685	18.65%	4.070%
7	10.746	1296	1303	1312	rVB	379998	662927	14.81%	3.232%
8	13.202	1713	1721	1732	rBV	1387111	2162446	48.31%	10.544%
9	14.025	1856	1861	1867	rBV	63658	99660	2.23%	0.486%
10	14.577	1947	1955	1963	rBV2	604828	991020	22.14%	4.832%
11	16.058	2201	2207	2220	rBV	1340902	2155616	48.15%	10.511%
12	17.315	2415	2421	2431	rBV	933440	1420976	31.74%	6.929%
13	19.930	2860	2866	2880	rBV	3441619	4476419	100.00%	21.827%
14	21.228	3083	3087	3097	rBV	468679	662547	14.80%	3.231%
15	21.563	3137	3144	3154	rBV	1073657	1734283	38.74%	8.456%
16	22.380	3279	3283	3291	rVB6	41194	81384	1.82%	0.397%
17	24.671	3662	3673	3682	rBV2	631875	1786525	39.91%	8.711%
18	25.934	3883	3888	3904	rVB8	52145	171199	3.82%	0.835%

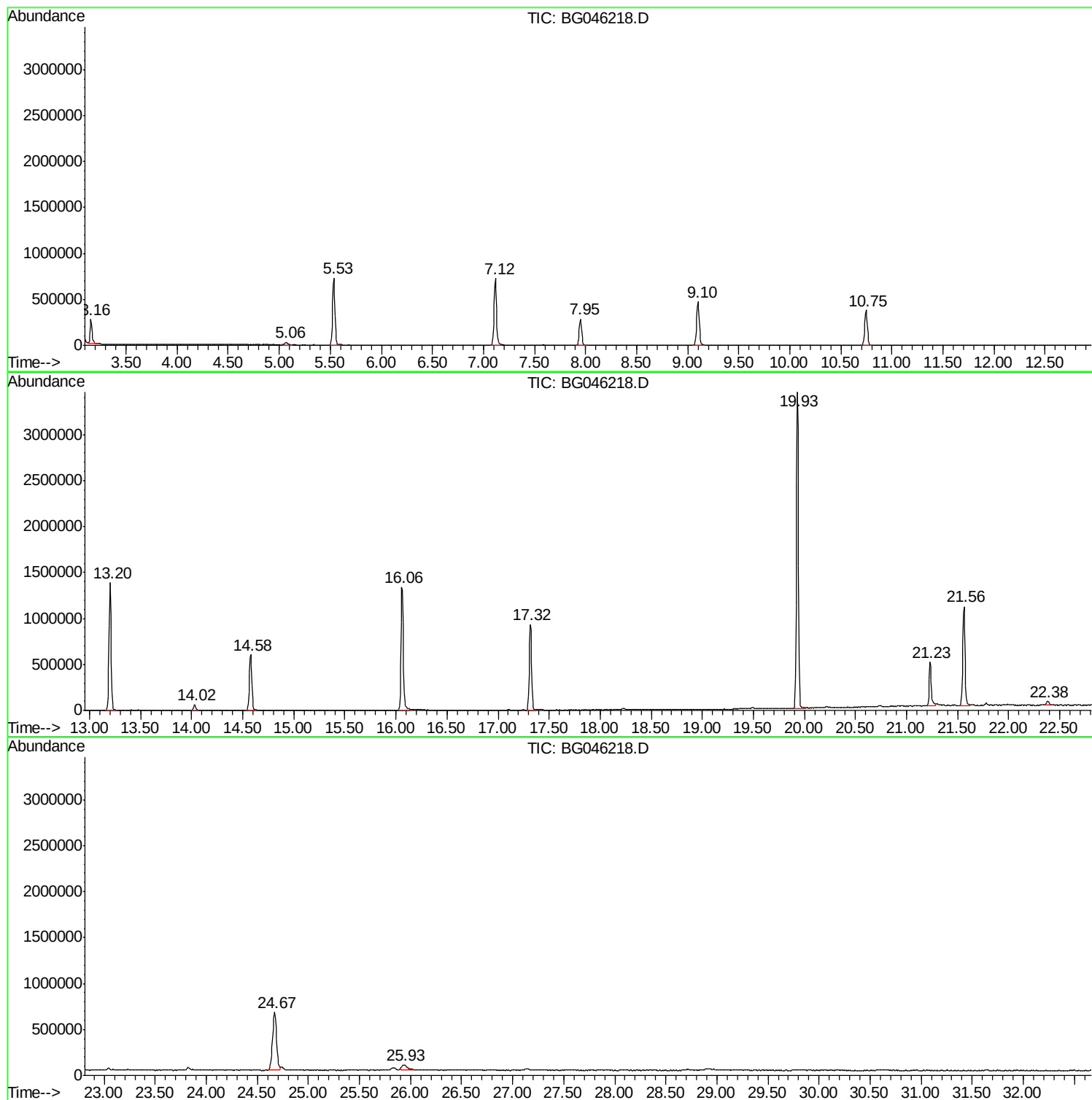
Sum of corrected areas: 20508681

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Quant Method : Z:\SVOASRV\HPCHEM1\BNA G\METHODS\8270-BG073020.M
Quant Title : ASP BNA STANDARDS FOR 5 POINT CALIBRATION

TIC Library : C:\Database\NIST11.L
TIC Integration Parameters: LSCINT.P



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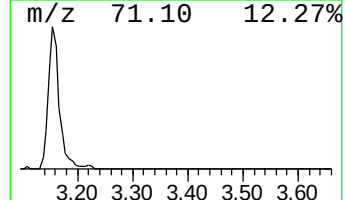
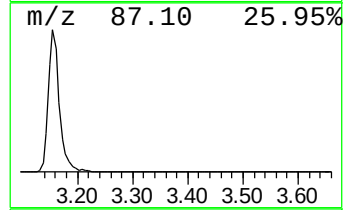
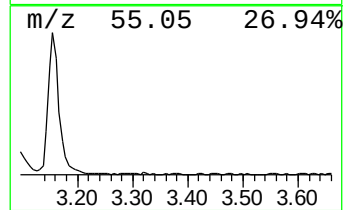
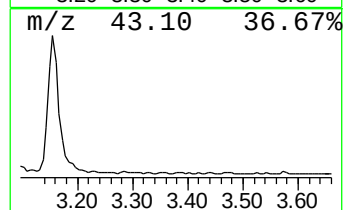
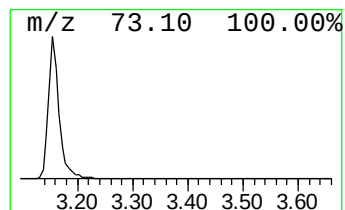
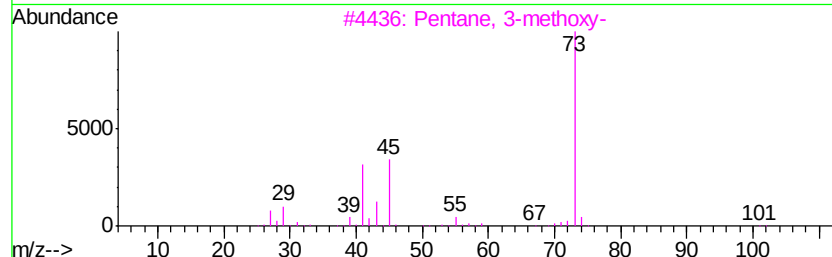
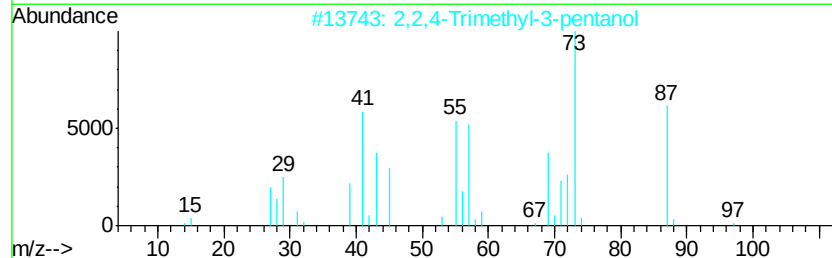
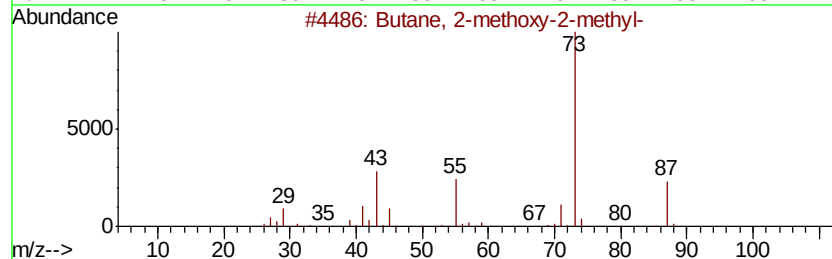
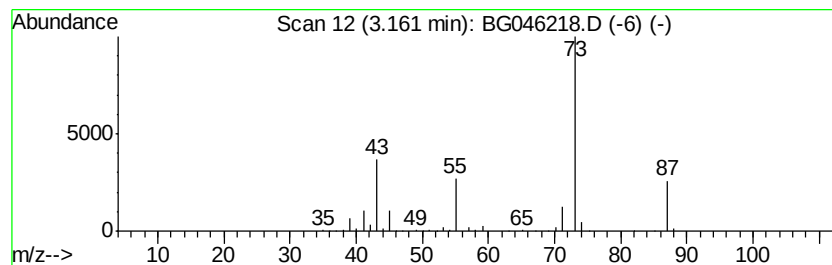
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TIC Library : C:\Database\NIST11.L
TIC Integration Parameters: LSCINT.P

Peak Number 1 Butane, 2-methoxy-2-methyl- Concentration Rank 1

R.T.	EstConc	Area	Relative to ISTD	R.T.
3.16	15.46 ng	365237	1,4-Dichlorobenzene-d4	7.95

Hit#	of	Tentative ID	MW	MolForm	CAS#	Qual
1	5	Butane, 2-methoxy-2-methyl-	102	C6H14O	000994-05-8	83
2		2,2,4-Trimethyl-3-pentanol	130	C8H18O	005162-48-1	43
3		Pentane, 3-methoxy-	102	C6H14O	036839-67-5	17
4		1,3-Dioxolane, 2-methyl-	88	C4H8O2	000497-26-7	9
5		Silane, tetramethyl-	88	C4H12Si	000075-76-3	9



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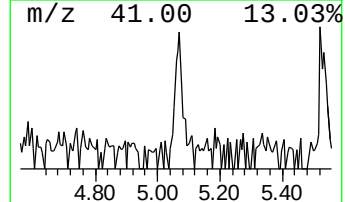
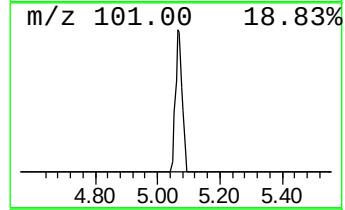
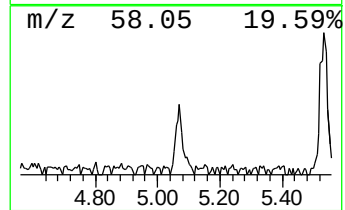
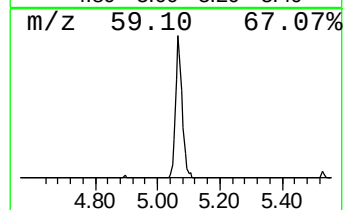
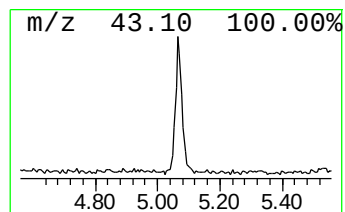
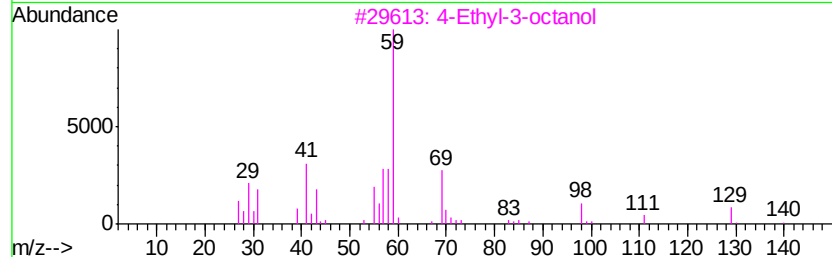
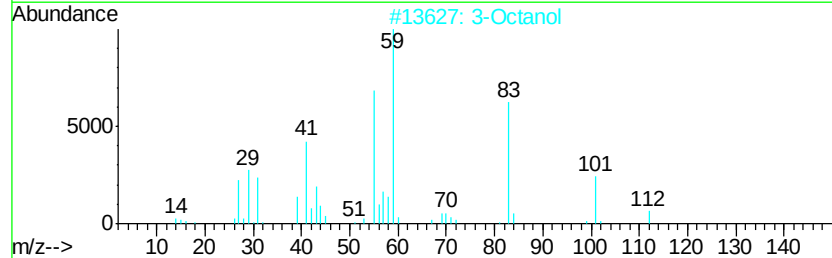
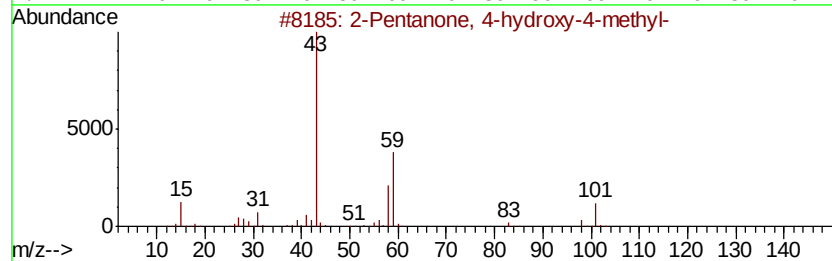
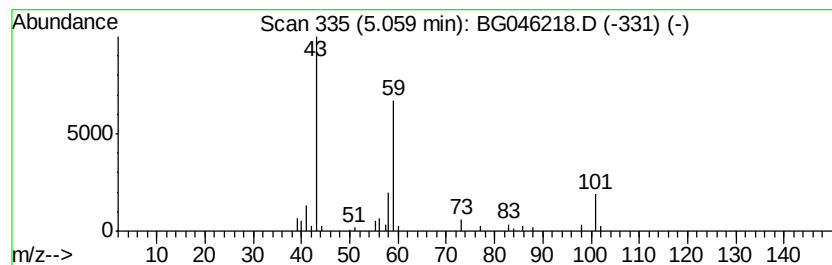
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 Peak Number 2 2-Pentanone, 4-hydroxy-4-me... Concentration Rank 3

R.T.	EstConc	Area	Relative to ISTD	R.T.
5.06	2.05 ng	48436	1,4-Dichlorobenzene-d4	7.95

Hit#	of	5	Tentative ID	MW	MolForm	CAS#	Qual
1	2-Pentanone, 4-hydroxy-4-methyl-	116	C6H12O2	000123-42-2	59		
2	3-Octanol	130	C8H18O	000589-98-0	28		
3	4-Ethyl-3-octanol	158	C10H22O	019781-26-1	28		
4	3-Pentanol	88	C5H12O	000584-02-1	25		
5	2-Hexanol, 2-methyl-	116	C7H16O	000625-23-0	25		



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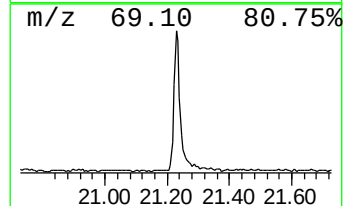
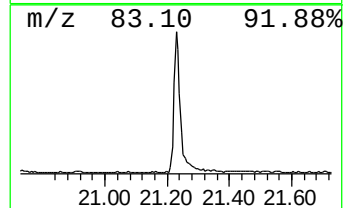
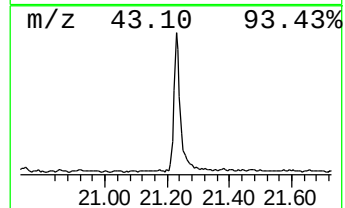
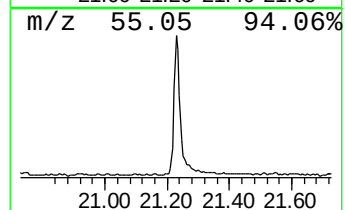
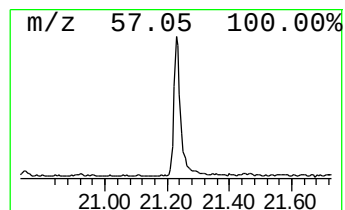
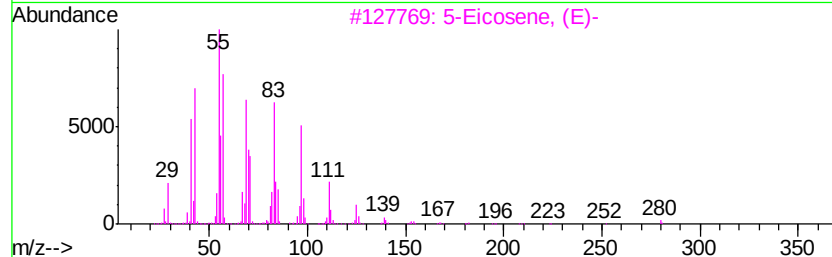
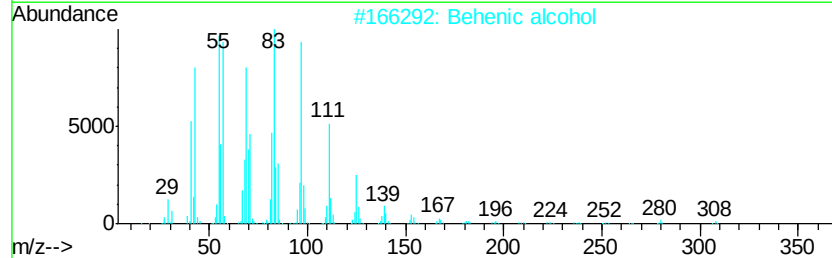
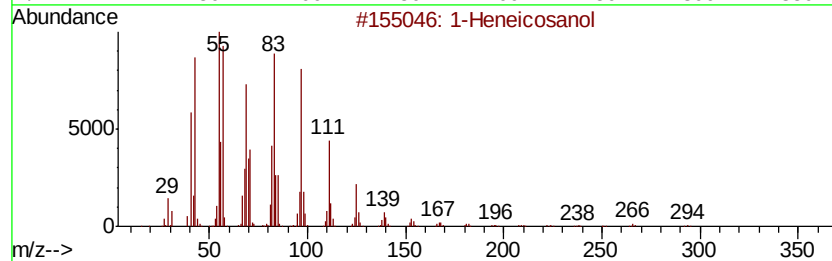
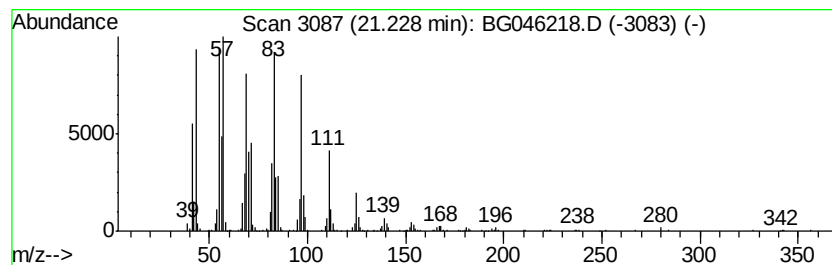
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Peak Number 3 1-Heneicosanol Concentration Rank 2

R.T.	EstConc	Area	Relative to ISTD	R.T.
21.23	7.64 ng	662547	Chrysene-d12	21.56

Hit#	of	Tentative ID	MW	MolForm	CAS#	Qual
1	5	1-Heneicosanol	312	C21H44O	015594-90-8	95
2		Behenic alcohol	326	C22H46O	000661-19-8	95
3		5-Eicosene, (E)-	280	C20H40	074685-30-6	95
4		n-Nonadecanol-1	284	C19H40O	001454-84-8	94
5		3-Eicosene, (E)-	280	C20H40	074685-33-9	93



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TIC Top Hit name	RT	EstConc	Units	Response	--Internal Standard--			
					#	RT	Resp	Conc
Butane, 2-methoxy...	3.16	15.5	ng	365237	1	7.95	472544	20.0
2-Pentanone, 4-hy...	5.06	2.0	ng	48436	1	7.95	472544	20.0
1-Heneicosanol	21.23	7.6	ng	662547	5	21.56	1734280	20.0