

Data Path : Z:\HPCHEM1\BNA_F\DATA\BF062215\
 Data File : BF080125.D
 Acq On : 23 Jun 2015 8:12
 Operator : TP/IZ
 Sample : G2715-01 5X
 Misc :
 ALS Vial : 31 Sample Multiplier: 1

Instrument :
 BNA_F
 ClientSampleId :
 76-INFIELD(0-2)-1

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

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

Method : Z:\HPCHEM1\BNA_F\METHODS\8270-BF061715.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	5.533	212	214	219	rVB	53319	58758	10.15%	1.234%
2	5.933	246	249	251	rBV	193073	219804	37.96%	4.615%
3	6.916	333	335	338	rBV	225471	229355	39.61%	4.815%
4	7.087	348	350	352	rBV	307058	252556	43.62%	5.302%
5	7.327	368	371	373	rBV	189950	249484	43.09%	5.238%
6	7.476	382	384	387	rVB	179770	183651	31.72%	3.856%
7	7.876	417	419	421	rBV	179499	142381	24.59%	2.989%
8	8.607	481	483	486	rBV	286148	338495	58.46%	7.107%
9	9.682	575	577	580	rBV	209974	282488	48.79%	5.931%
10	10.070	610	611	613	rBV	13682	14352	2.48%	0.301%
11	10.379	636	638	641	rBV	327396	411823	71.13%	8.646%
12	11.088	698	700	704	rVB	20890	19730	3.41%	0.414%
13	11.179	706	708	710	rBV	151348	167550	28.94%	3.518%
14	11.888	767	770	772	rBV	560646	483248	83.46%	10.146%
15	13.408	898	903	906	rBV	5482	8576	1.48%	0.180%
16	13.545	913	915	918	rBV	340729	341556	58.99%	7.171%
17	14.082	958	962	964	rBV	11316	13999	2.42%	0.294%
18	14.185	968	971	975	rVV4	3241	9648	1.67%	0.203%
19	14.585	1004	1006	1008	rBV2	19930	32738	5.65%	0.687%
20	14.756	1017	1021	1022	rBV4	2930	7981	1.38%	0.168%
21	14.791	1022	1024	1027	rBV	482975	579004	100.00%	12.156%
22	14.916	1033	1035	1039	rBV4	7354	11135	1.92%	0.234%
23	15.134	1052	1054	1057	rBV3	3642	6366	1.10%	0.134%
24	15.385	1073	1076	1080	rBV3	26668	51262	8.85%	1.076%
25	15.957	1124	1126	1128	rBV3	6563	12198	2.11%	0.256%
26	16.208	1145	1148	1152	rVB	22575	34673	5.99%	0.728%
27	16.642	1183	1186	1189	rBV	406956	576847	99.63%	12.111%
28	17.248	1237	1239	1243	rVB2	12873	23363	4.04%	0.491%

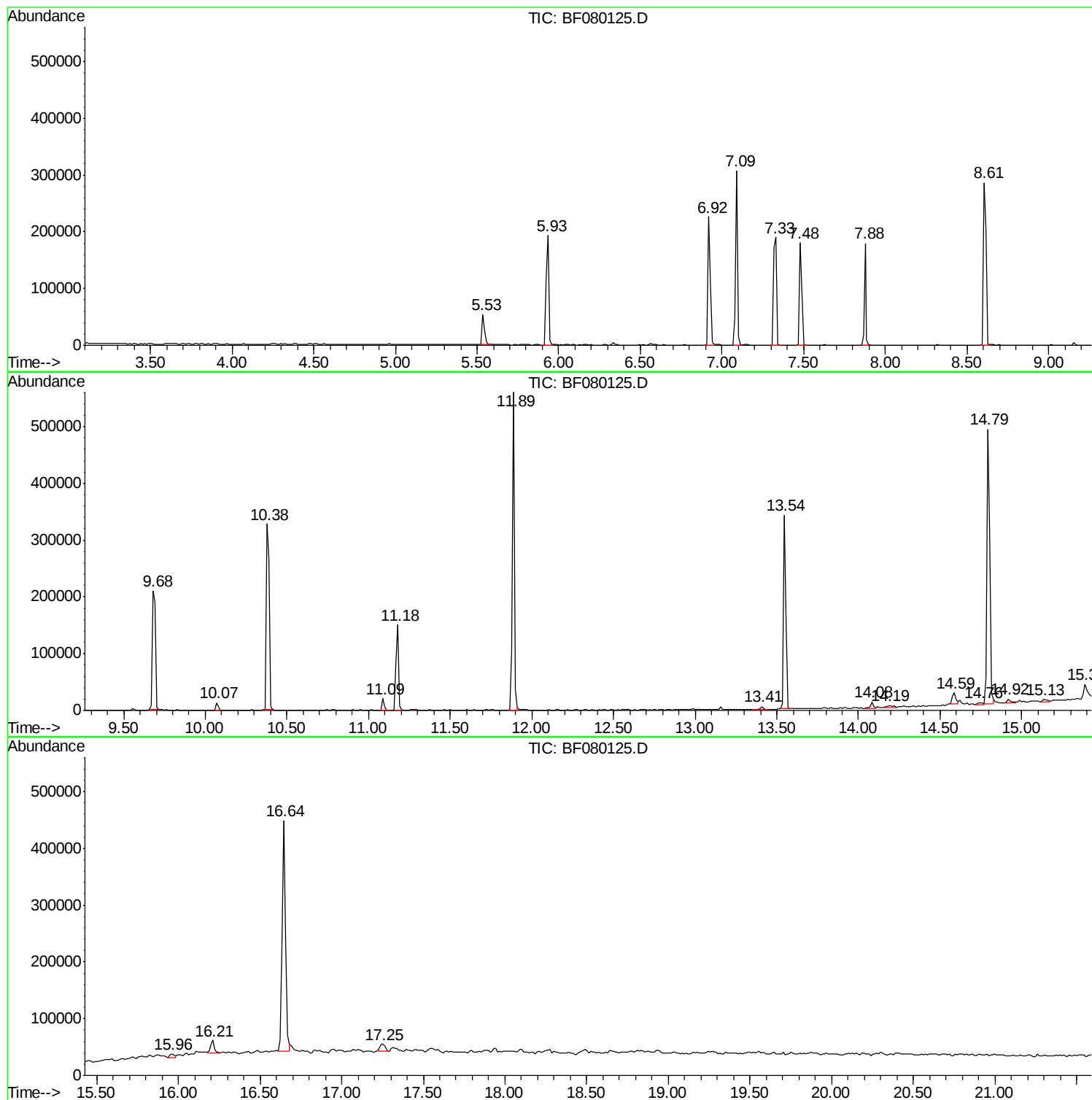
Sum of corrected areas: 4763021

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Quant Method : Z:\HPCHEM1\BNA_F\METHODS\8270-BF061715.M
Quant Title : ASP BNA STANDARDS FOR 5 POINT CALIBRATION

TIC Library : C:\DATABASE\NIST02.L
TIC Integration Parameters: LSCINT.P



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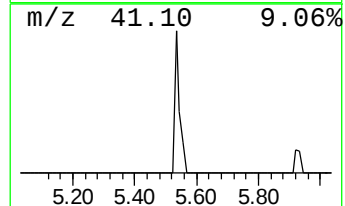
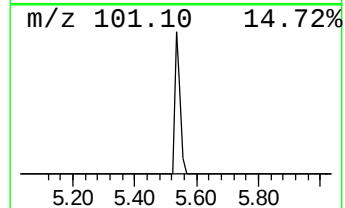
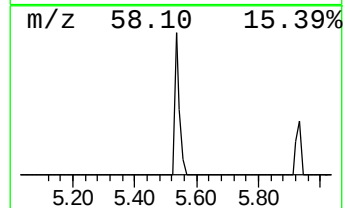
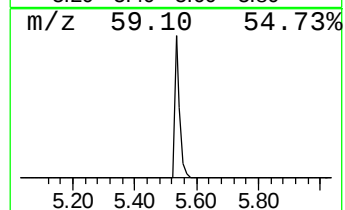
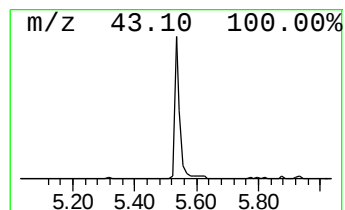
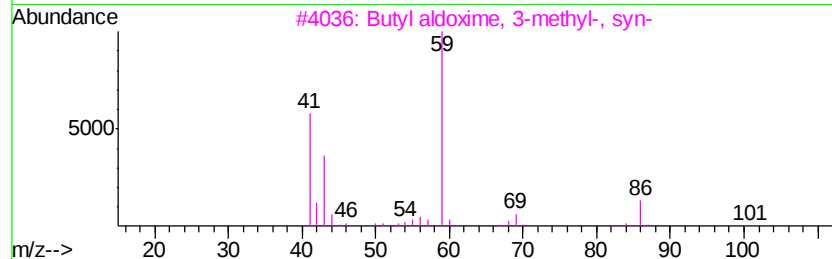
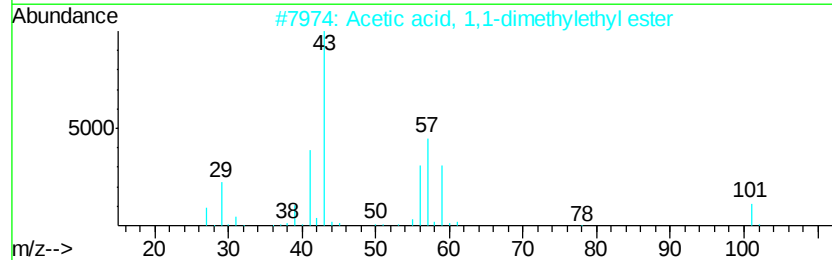
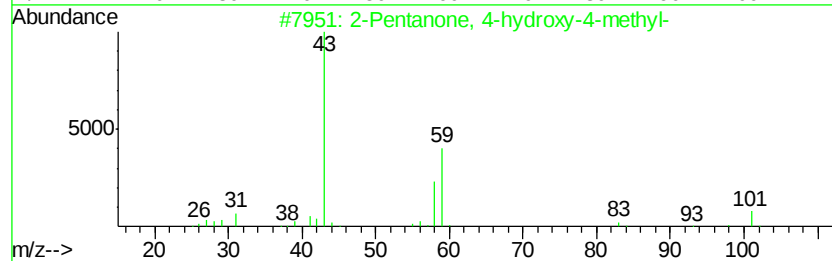
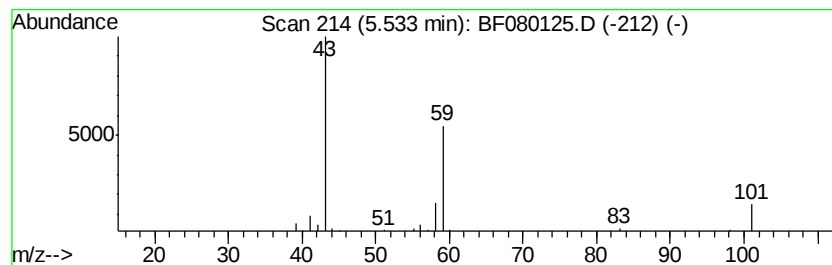
Quant Method : Z:\HPCHEM1\BNA_F\METHODS\8270-BF061715.M
Quant Title : ASP BNA STANDARDS FOR 5 POINT CALIBRATION

TIC Library : C:\DATABASE\NIST02.L
TIC Integration Parameters: LSCINT.P

Peak Number 1 2-Pentanone, 4-hydroxy-4-me... Concentration Rank 3

R.T.	EstConc	Area	Relative to ISTD	R.T.
5.53	4.71 ng	58758	1,4-Dichlorobenzene-d4	7.33

Hit#	of	5	Tentative ID	MW	MolForm	CAS#	Qual
1			2-Pentanone, 4-hydroxy-4-methyl-	116	C6H12O2	000123-42-2	56
2			Acetic acid, 1,1-dimethylethyl e...	116	C6H12O2	000540-88-5	28
3			Butyl aldoxime, 3-methyl-, syn-	101	C5H11NO	005780-40-5	25
4			Diacetamide	101	C4H7NO2	000625-77-4	9
5			4-Penten-2-one, 4-methyl-	98	C6H10O	003744-02-3	9



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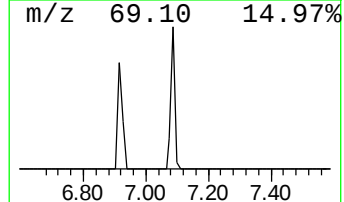
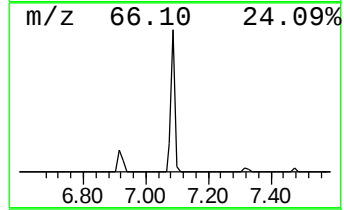
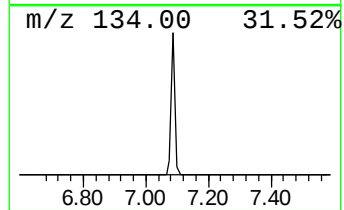
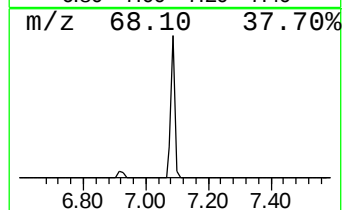
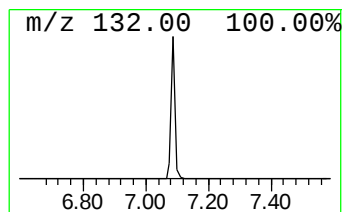
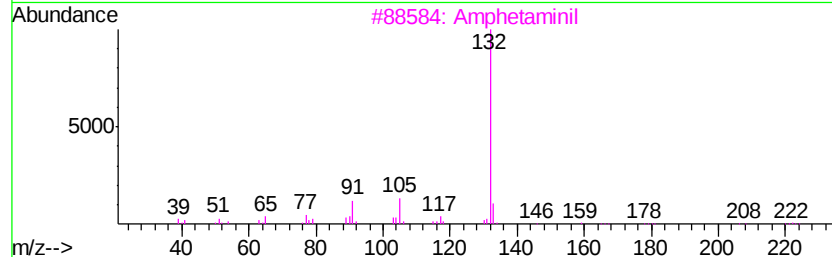
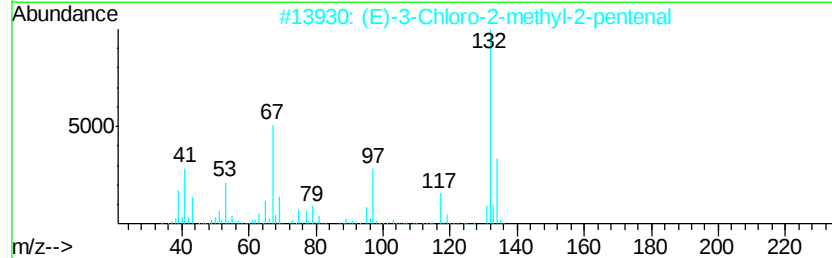
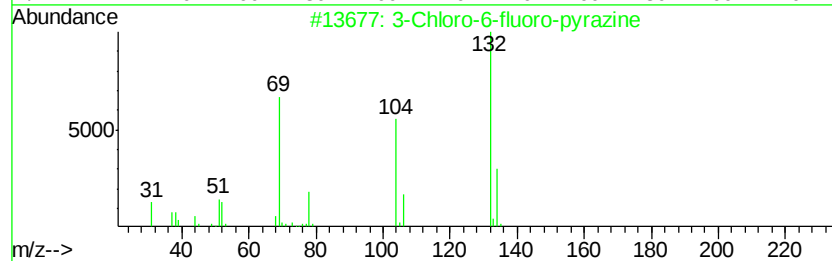
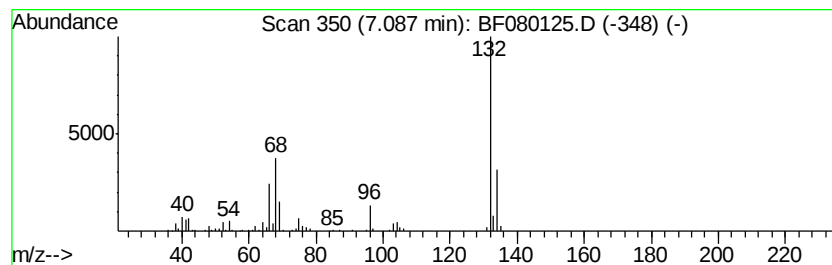
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Peak Number 2 unknown7.09 Concentration Rank 1

R.T.	EstConc	Area	Relative to ISTD	R.T.
7.09	20.25 ng	252556	1,4-Dichlorobenzene-d4	7.33

Hit#	of	Tentative ID	MW	MolForm	CAS#	Qual
1	5	3-Chloro-6-fluoro-pyrazine	132	C4H2ClFN2	1000146-10-7	35
2		(E)-3-Chloro-2-methyl-2-pentenal	132	C6H9ClO	031357-76-3	35
3		Amphetaminil	250	C17H18N2	017590-01-1	12
4		5-Aminoindole	132	C8H8N2	005192-03-0	12
5		4-Methylpyrrolo[1,2-a]pyrazine	132	C8H8N2	064608-60-2	9



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TIC Top Hit name	RT	EstConc	Units	Response	--Internal Standard--			
					#	RT	Resp	Conc
2-Pentanone, 4-hy...	5.53	4.7	ng	58758	1	7.33	249484	20.0
unknown7.09	7.09	20.3	ng	252556	1	7.33	249484	20.0