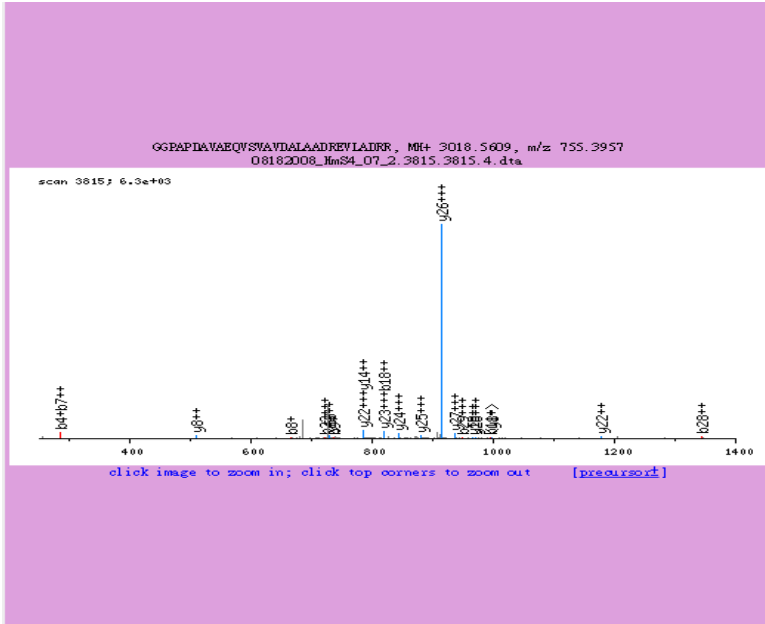


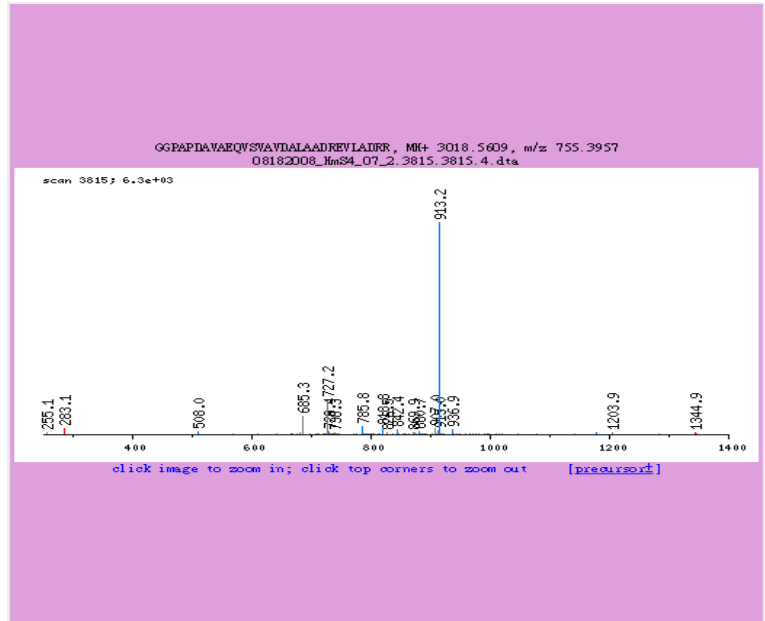
Supplementary Figure 1. Tandem mass spectra of the *H. marismortui* proteins identified with only a single unique peptide. The gene locus name, peptide sequence, peptide probability, precursor charge, calculated neutral peptide mass, precursor neutral mass, mass difference, Xcorr, and deltaCN are indicated in the top right corner. *Top left panel:* MS/MS spectrum showing the matched b- and y-ions; *Bottom left panel:* MS/MS spectrum showing the m/z of the matched b- and y-ions. *Bottom right panel:* Table of highlighted matched b- and y-ions.

rrnAC2681

GGPAPDAVAEQVSVAVDALAADREVLADRR

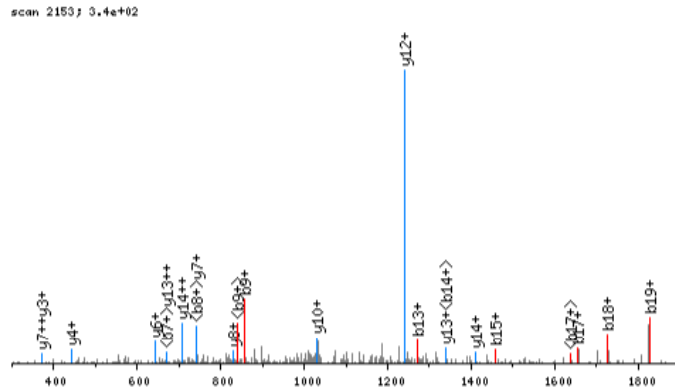


Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	Deltacn
0.9998	4	3017.6	3018.6	1.0181	1.981	0.182



b ⁺	b ²⁺	b ³⁺	#	AA	#	y ⁺	y ²⁺	y ³⁺
58.0293	29.5186	20.0150	1	G	30			
115.0508	58.0293	39.0221	2	G	29	2961.5394	1481.2736	987.8517
212.1035	106.5557	71.3731	3	P	28	2904.5179	1452.7629	968.8445
283.1406	142.0742	95.0521	4	A	27	2807.4652	1404.2365	936.4936
380.1934	190.6006	127.4030	5	P	26	2736.4280	1368.7179	912.8146
495.2203	248.1141	165.7453	6	D	25	2639.3753	1320.1916	880.4636
566.2575	283.6326	189.4244	7	A	24	2524.3483	1262.6781	842.1213
665.3259	333.1668	222.4472	8	V	23	2433.3112	1227.1595	818.4423
736.3630	368.6854	246.1262	9	A	22	2354.2428	1177.6253	785.4195
865.4056	433.2067	289.1404	10	E	21	2283.2057	1142.1068	761.7404
993.4641	497.2360	331.8266	11	Q	20	2154.1631	1077.5855	718.7263
1092.5326	546.7702	364.8494	12	V	19	2026.1045	1013.5562	676.0401
1179.5646	590.2862	393.8601	13	S	18	1927.0361	964.0220	643.0173
1278.6330	639.8204	426.8829	14	V	17	1840.0041	920.5060	614.0066
1349.6701	675.3390	450.5619	15	A	16	1740.9357	870.9717	580.9838
1448.7385	724.8732	483.5847	16	V	15	1669.8986	835.4532	557.3047
1563.7655	782.3867	521.9270	17	D	14	1570.8301	785.9190	524.2819
1634.8026	817.9052	545.6061	18	A	13	1455.8032	728.4055	485.9396
1747.8867	874.4472	583.3008	19	L	12	1384.7661	692.8870	462.2606
1818.9238	909.9658	606.9798	20	A	11	1271.6820	636.3449	424.5659
1889.9609	945.4844	630.6588	21	A	10	1200.6449	600.8264	400.8869
2004.9878	1002.9978	669.0012	22	D	9	1129.6078	565.3078	377.2078
2161.0889	1081.0484	721.0349	23	R	8	1014.5809	507.7943	338.8655
2290.1315	1145.5697	764.0491	24	E	7	858.4797	429.7438	286.8318
2389.1999	1195.1039	797.0719	25	V	6	729.4371	365.2225	243.8176
2502.2840	1251.6459	834.7666	26	L	5	630.3687	315.6883	210.7948
2573.3211	1287.1645	858.4456	27	A	4	517.2847	259.1462	173.1001
2688.3481	1344.6779	896.7879	28	D	3	446.2476	223.6277	149.4211
2844.4492	1422.7285	948.8216	29	R	2	331.2206	166.1142	111.0788
			30	R	1	175.1195	88.0637	59.0451

SVVGDDAVPDAESVTAVPR, MH+ 2098.0304, m/z 1049.5188
 08182008_Im8S3_12_4_21S3_21S3_2.dta

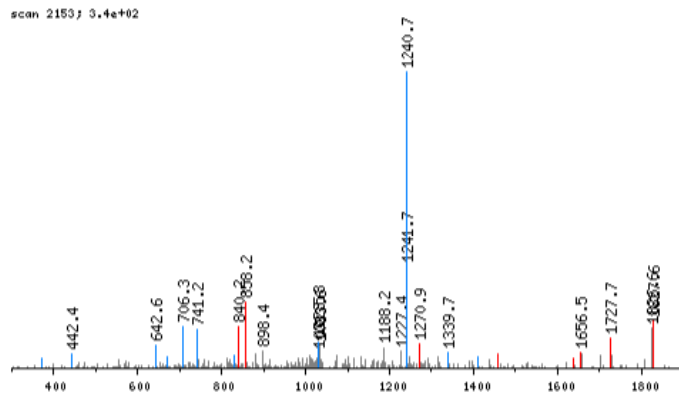


click image to zoom in; click top corners to zoom out [precursor±]

rrnAC1938 SVVGDDAVPDAESVTAVPR

Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	Deltacn
1	2	2097	2098	1.005	4.522	0.482

SVVGDDAVPDAESVTAVPR, MH+ 2098.0304, m/z 1049.5188
 08182008_Im8S3_12_4_21S3_21S3_2.dta

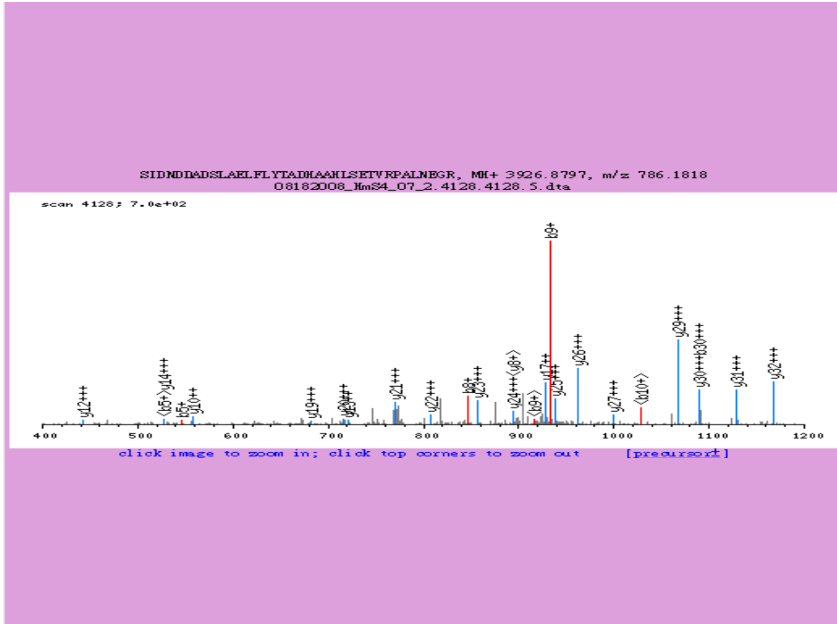


click image to zoom in; click top corners to zoom out [precursor±]

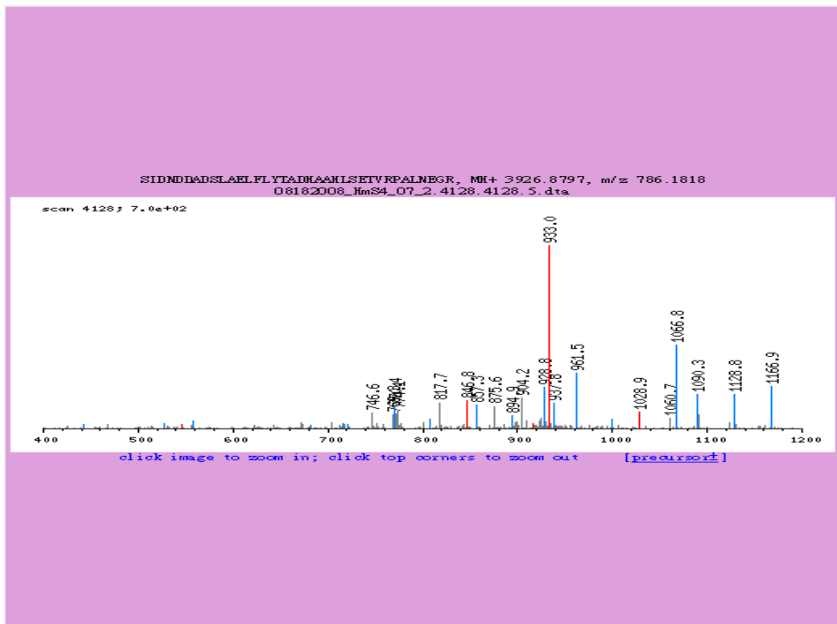
b ⁺	#	AA	#	y ⁺	y ²⁺
88.0399	1	S	21		
187.1083	2	V	20	2010.9984	1006.0031
286.1767	3	V	19	1911.9300	956.4689
343.1981	4	G	18	1812.8616	906.9347
458.2251	5	D	17	1755.8401	878.4240
573.2520	6	D	16	1640.8132	820.9105
688.2790	7	D	15	1525.7862	763.3970
759.3161	8	A	14	1410.7593	705.8835
858.3845	9	V	13	1339.7222	670.3650
955.4373	10	P	12	1240.6537	620.8308
1070.4642	11	D	11	1143.6010	572.3044
1141.5013	12	A	10	1028.5740	514.7909
1270.5439	13	E	9	957.5369	479.2724
1357.5759	14	S	8	828.4943	414.7511
1456.6444	15	V	7	741.4623	371.2351
1557.6920	16	T	6	642.3939	321.7009
1656.7605	17	V	5	541.3462	271.1770
1727.7976	18	A	4	442.2778	221.6428
1826.8660	19	V	3	371.2407	166.1243
1923.9187	20	P	2	272.1723	136.5900
	21	R	1	175.1195	88.0637

rrnAC3286

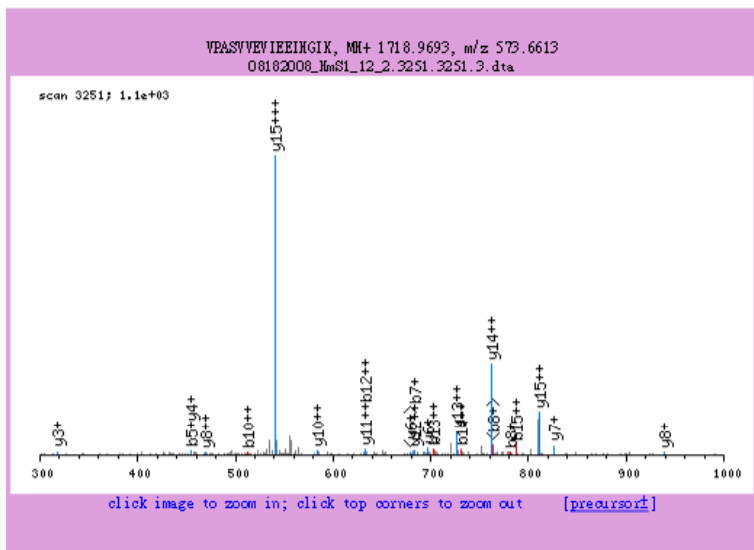
SIDNDDADSLAELFLYTADHAAHLSETVR PALNEGR



Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	Deltaen
0.9999	5	3925.9	3925.9	0.0201	3.367	0.133



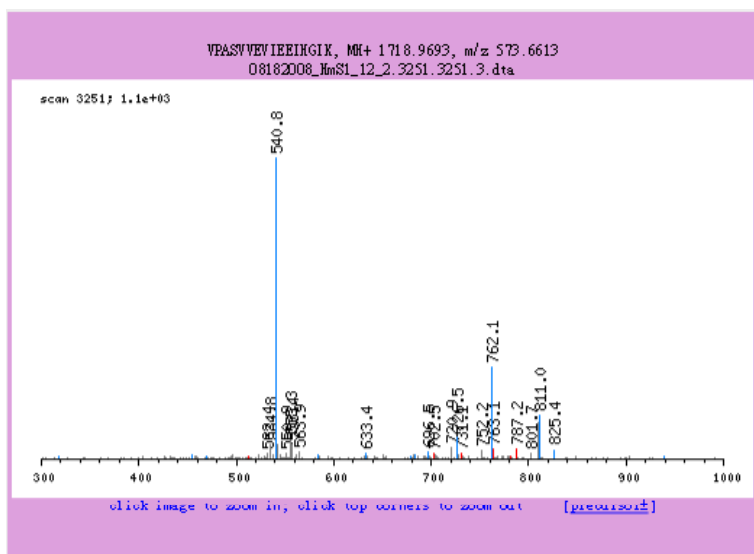
b ⁺	b ²⁺	b ³⁺	#	AA	#	y ⁺	y ²⁺	y ³⁺
88.0999	44.5298	30.0185	1	S	96			
201.1239	101.0659	67.7132	2	I	95	3839.8477	1920.4277	1280.6211
316.1509	158.5793	106.0555	3	D	94	3726.7636	1863.8857	1242.9264
430.1938	215.6008	144.0698	4	N	93	3611.7966	1806.3722	1204.5841
545.2207	273.1143	182.4121	5	D	92	3497.6937	1749.3508	1166.5698
660.2477	330.6277	220.7544	6	D	91	3382.6668	1691.8373	1128.2275
731.2848	366.1463	244.4335	7	A	90	3267.6398	1634.3238	1089.8852
846.3117	423.6598	282.7758	8	D	89	3196.6027	1598.8053	1066.2061
933.3438	467.1758	311.7865	9	S	88	3081.5758	1541.2918	1027.8638
1046.4278	523.7178	349.4812	10	L	87	2994.5437	1497.7758	998.8531
1117.4649	559.2364	373.1602	11	A	86	2881.4597	1441.2338	961.1584
1246.5075	623.7577	416.1744	12	E	85	2810.4226	1405.7152	937.4794
1359.5916	680.2997	453.8691	13	L	84	2681.3800	1341.1939	894.4652
1506.6600	753.8339	502.8919	14	F	83	2568.2959	1284.6519	856.7705
1619.7441	810.3759	540.5866	15	L	82	2421.2275	1211.1177	807.7477
1782.8074	891.9076	594.9410	16	Y	81	2308.1434	1154.5756	770.0530
1883.8551	942.4315	628.6236	17	T	80	2145.0801	1073.0440	715.6986
1954.8922	977.9500	652.3026	18	A	79	2044.0324	1022.5201	682.0160
2069.9191	1035.4635	690.6449	19	D	78	1972.9953	987.0016	658.3370
2206.9780	1103.9929	736.3312	20	H	77	1857.9684	929.4881	619.9947
2278.0152	1139.5115	760.0103	21	A	76	1720.9095	860.9586	574.3084
2349.0523	1175.0301	783.6893	22	A	75	1649.8723	825.4401	530.6293
2486.1112	1243.5595	829.3756	23	H	74	1578.8352	789.9215	526.9503
2599.1953	1300.1015	867.0703	24	L	73	1441.7763	721.3921	481.2640
2686.2273	1343.6176	896.0810	25	S	72	1328.6922	664.8500	443.5693
2815.2699	1408.1388	939.0952	26	E	71	1241.6602	621.3340	414.5586
2916.3176	1458.6627	972.7777	27	T	70	1112.6176	556.8127	371.5444
3015.3860	1508.1969	1005.8005	28	V	69	1011.5700	506.2889	337.8619
3171.4871	1586.2475	1057.8342	29	R	68	912.5015	456.7547	304.8391
3268.5398	1634.7738	1090.1852	30	P	67	756.4004	378.7041	252.8054
3339.5770	1670.2924	1113.8642	31	A	66	659.3477	330.1777	220.4544
3452.6610	1726.8344	1151.5589	32	L	65	588.3105	294.6592	196.7754
3566.7039	1783.8559	1189.5732	33	N	64	475.2265	238.1172	159.0807
3695.7465	1848.3772	1232.5874	34	E	63	361.1896	181.0957	121.0664
3752.7660	1876.8879	1251.5946	35	G	62	232.1410	116.5744	78.0522
			36	R	61	175.1195	88.0637	59.0451



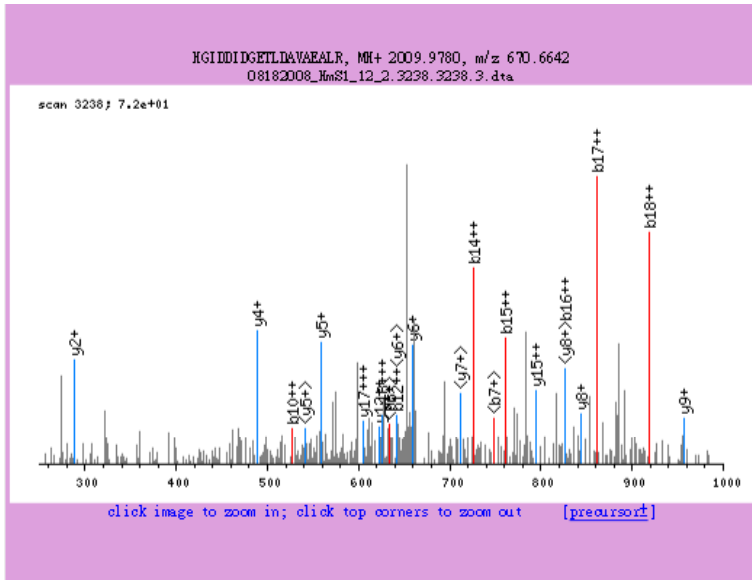
rrnAC1913

VPASVVEVIEEIHGK

Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	DeltaCn
1	3	1718	1718	0.0029	3.243	0.292

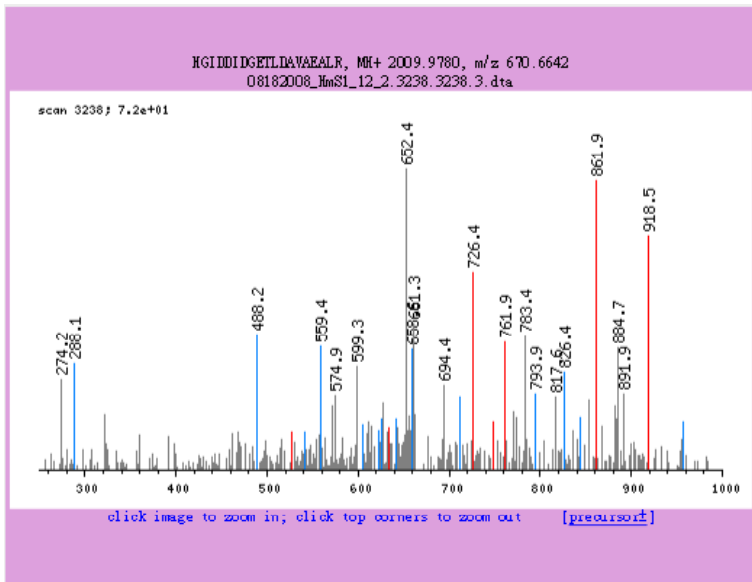


b ⁺	b ²⁺	# AA	#	y ⁺	y ²⁺	y ³⁺
100.0762	50.5420	1	V	16		
197.1290	99.0684	2	P	15	1619.9008	810.4543 540.6388
268.1661	134.5870	3	A	14	1522.8481	761.9280 508.2879
355.1981	178.1030	4	S	13	1451.8110	726.4094 484.6089
454.2666	227.6372	5	V	12	1364.7789	682.8934 455.5982
553.3330	277.1714	6	V	11	1265.7105	633.3592 422.5754
682.3776	341.6927	7	E	10	1166.6421	583.8250 389.5526
781.4460	391.2269	8	V	9	1037.5995	519.3037 346.5384
894.5300	447.7689	9	I	8	938.5311 469.7695	313.5156
1023.5726	512.2902	10	E	7	825.4470	413.2274 275.8209
1152.6152	576.8115	11	E	6	696.4044	348.7061 232.8067
1265.6993	633.3536	12	I	5	567.3619	284.1848 189.7925
1402.7582	701.8830	13	H	4	454.2778	227.6428 152.0978
1459.7797	730.3937	14	G	3	317.2189	159.1134 106.4115
1572.8637	786.9358	15	I	2	280.1974	130.6026 87.4044
		16	K	1	147.1134	74.0606 49.7097



rrnAC0470 HGIDDIDGETLDAVAEALR

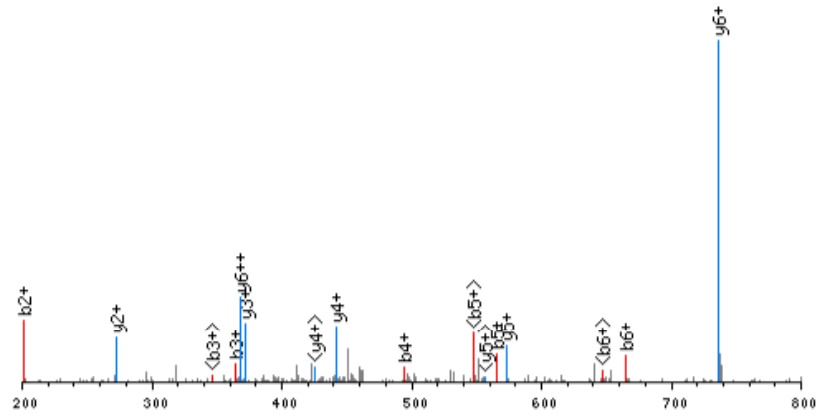
Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	Deltacn
0.9878	3	2009	2009	-0.0119	2.381	0.103



b ⁺	b ²⁺	#	AA	#	y ⁺	y ²⁺	y ³⁺
138.0667	69.5373	1	H	19			
195.0882	98.0480	2	G	18	1872.9191	936.9635	624.9782
308.1723	154.5900	3	I	17	1815.8976	908.4527	605.9711
423.1992	212.1035	4	D	16	1702.8135	851.9107	568.2764
538.2262	269.6170	5	D	15	1587.7866	794.3972	529.9341
651.3102	326.1590	6	I	14	1472.7597	736.8837	491.5918
766.3372	383.6725	7	D	13	1359.6756	680.3417	453.8971
823.3586	412.1832	8	G	12	1244.6487	622.8282	415.5548
952.4012	476.7045	9	E	11	1187.6272	594.3175	396.5476
1053.4489	527.2284	10	T	10	1058.5846	529.7962	353.5334
1166.5330	583.7704	11	L	9	957.5369	479.2724	319.8509
1281.5599	641.2839	12	D	8	844.4529	422.7303	282.1562
1332.5970	676.8024	13	A	7	729.4259	365.2169	243.8139
1451.6654	726.3366	14	V	6	658.3888	329.6983	220.1348
1522.7025	761.8552	15	A	5	559.3204	280.1641	187.1120
1651.7451	826.3765	16	E	4	488.2833	244.6455	163.4330
1722.7822	861.8950	17	A	3	359.2407	180.1243	120.4188
1835.8663	918.4371	18	L	2	288.2036	144.6057	96.7397
		19	R	1	175.1195	88.0637	59.0451

TVYMAVPR, MI+ 936.4977, m/z 468.7525
08182008_ImS3_07_3.1746.1746.2.dta

scan 1746; 3.7e+02



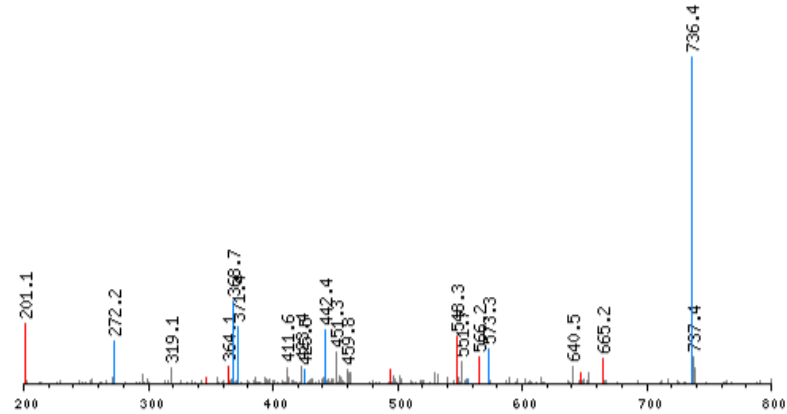
click image to zoom in; click top corners to zoom out [\[precursor†\]](#)

rrnAC3378 TVYMAVPR

Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	Deltaen
0.9991	2	935.49	935.49	0.0019	1.935	0.143

TVYMAVPR, MI+ 936.4977, m/z 468.7525
08182008_ImS3_07_3.1746.1746.2.dta

scan 1746; 3.7e+02



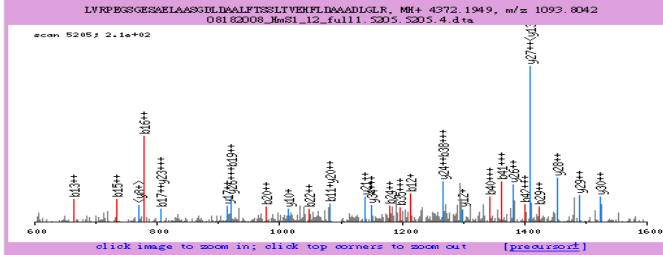
click image to zoom in; click top corners to zoom out [\[precursor†\]](#)

b ⁺	b ²⁺	# AA	#	y ⁺	y ²⁺
102.0555	51.5317	1	T	8	
201.1239	101.0639	2	V	7	835.4500
364.1872	182.5975	3	Y	6	736.3816
495.2277	248.1178	4	M	5	573.3183
566.2648	283.6363	5	A	4	442.2778
665.3333	333.1705	6	V	3	371.2407
762.3860	381.6969	7	P	2	272.1723
		8	R	1	175.1195
					88.0637

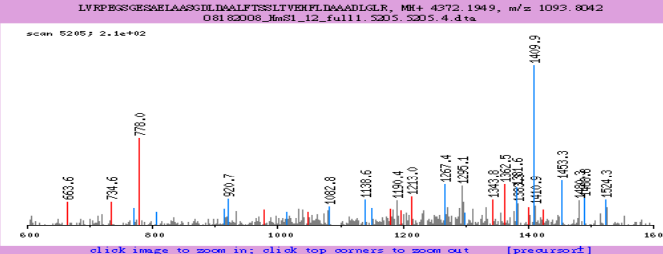
rrnAC3088

LVRPEGSGESAEALAASGDLDAALFTSSL

TVEHFLDAAADLGLR



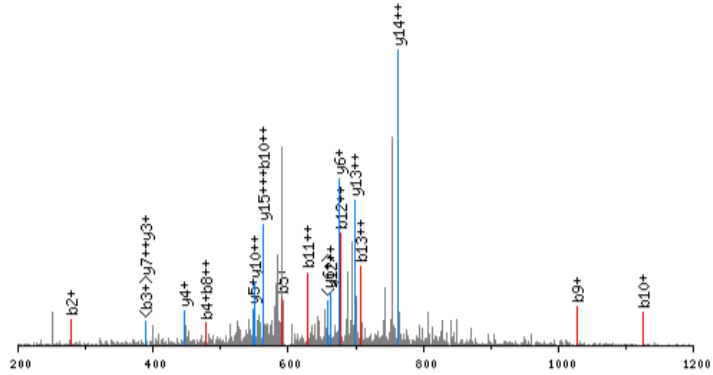
Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	DeltaCn
0.999	4	4371.2	4372.2	1.0103	3.706	0.046



b ⁺	b ²⁺	b ³⁺	#	AA	#	y ⁺	y ²⁺	y ³⁺				
114	0919	57	3499	38	R	25	4259	1108	2130	0593	1430	3755
213	1839	107	0841	71	T	23	4160	0424	2080	5251	1387	3527
369	2614	185	1946	123	R	41	4003	9413	2002	4746	1335	3190
466	3142	239	6610	156	P	40	3906	8885	1953	9482	1302	9681
595	3568	298	1823	199	E	39	3777	8459	1889	4269	1259	9539
652	3782	326	6930	218	G	38	3720	8245	1860	9161	1240	9467
739	4103	370	2090	247	S	37	3633	7924	1817	4001	1211	9360
796	4317	398	7198	266	G	36	3576	7710	1788	8894	1192	9289
925	4743	463	2411	309	S	35	3447	7284	1724	3681	1149	9147
1012	5063	506	7571	338	S	34	3360	6963	1680	8521	1120	9040
1089	5435	542	2756	361	A	33	3289	6593	1645	3335	1097	8923
1212	5861	606	7069	404	R	32	3160	6166	1580	8122	1054	8718
1325	6301	663	3390	442	L	31	3047	5726	1524	2702	1016	8561
1396	7072	698	8575	466	A	30	2976	4955	1488	7516	992	8370
1467	7443	734	3761	489	A	29	2905	4583	1453	2331	969	8180
1554	7764	777	8921	518	S	28	2818	4263	1409	7171	940	8023
1611	7978	806	4028	537	G	27	2761	4049	1381	2063	921	7866
1726	8248	863	9163	576	D	26	2646	3779	1323	6929	882	7979
1839	9088	920	4583	613	L	25	2533	2938	1267	1508	845	7832
1954	9358	977	9718	652	D	24	2418	2669	1209	6374	806	7609
2025	9729	1013	4904	675	A	23	2347	2298	1174	1188	783	7481
2097	0100	1049	0089	699	A	22	2216	1827	1138	6092	759	7028
2210	0941	1105	5510	737	R	21	2016	0402	1008	5240	672	6853
2357	1625	1179	0852	786	F	20	1914	9925	958	0002	639	0027
2458	2102	1229	6990	820	T	19	1827	9605	914	4842	609	9920
2545	2422	1273	1230	849	S	18	1740	9285	870	9681	580	9814
2632	2742	1316	6410	878	S	17	1627	8444	814	4261	543	2867
2745	3583	1373	1831	915	L	16	1526	7967	763	9023	509	6041
2846	4060	1423	7069	949	T	15	1427	7283	714	3681	476	5813
2945	4744	1473	2411	982	V	14	1327	6601	649	8468	433	5671
3074	5170	1537	7624	1025	E	13	1216	6260	581	3173	397	8806
3211	5759	1606	2919	1071	R	12	1114	5584	907	7831	338	8500
3358	6443	1679	8261	1120	F	11	901	4743	451	2411	301	1633
3471	7284	1736	3681	1157	L	10	814	4261	543	2867	509	6041
3586	7553	1793	8816	1196	D	9	786	4474	393	7276	262	8210
3657	7924	1829	4001	1219	A	8	715	4103	358	2090	239	1420
3728	8295	1864	9187	1243	A	7	644	3731	322	6905	215	4629
3799	8667	1900	4372	1267	A	6	573	3360	287	1719	191	7839
3914	8936	1957	9507	1305	D	5	458	3091	229	6585	153	4416
4027	9777	2014	4927	1343	L	4	345	2250	173	1164	115	7469
4084	9991	2043	0035	1362	G	3	288	2036	144	6057	96	7397
4198	0632	2099	5455	1400	L	2	175	1195	88	0637	59	0451
				43	R	1						

NYEAILDHAVETGVDR, MH+ 1801.8721, m/z 601.2955
20081001M4_516_690.02121.02121.3.dta

scan 2121; 9.2e+02



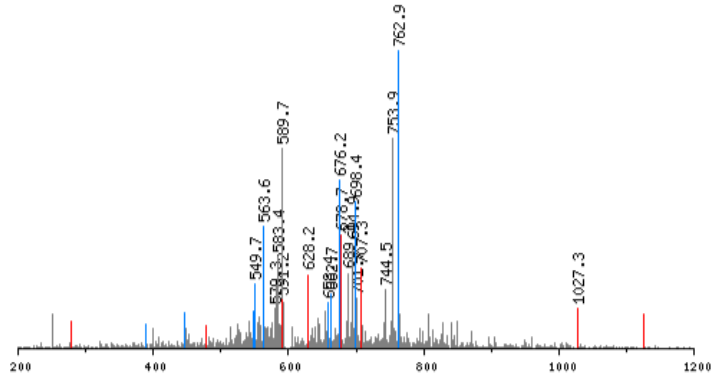
click image to zoom in; click top corners to zoom out [precursor]

rrnAC2998 NYEAILDHAVETGVDR

Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	DeltaCn
1	3	1800.9	1800.9	0.002192	2.972	0.349

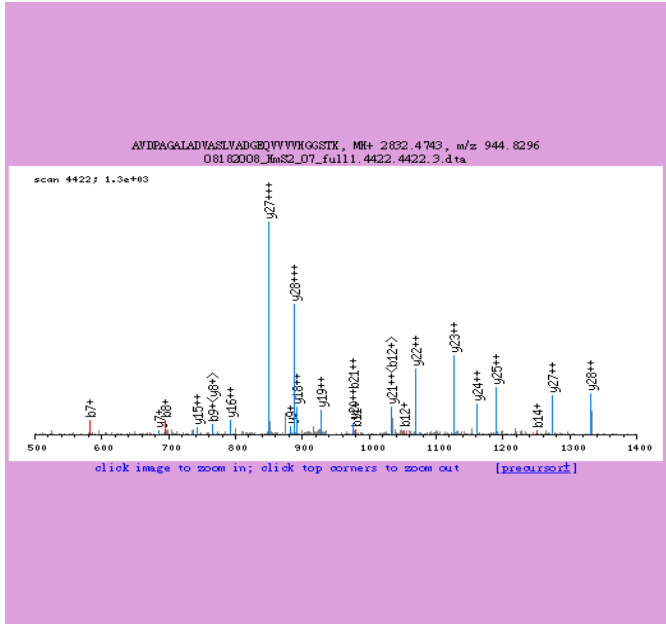
NYEAILDHAVETGVDR, MH+ 1801.8721, m/z 601.2955
20081001M4_516_690.02121.02121.3.dta

scan 2121; 9.2e+02



click image to zoom in; click top corners to zoom out [precursor]

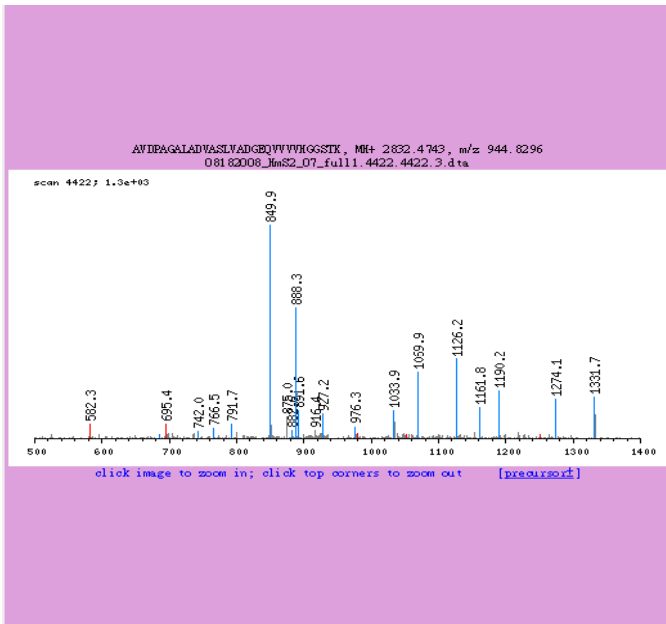
b ⁺	b ²⁺	#	AA	#	y ⁺	y ²⁺	y ³⁺
115.0308	58.0293	1	N	16			
278.1141	139.5610	2	Y	15	1687.8291	844.4185	563.2816
407.1567	204.0822	3	E	14	1524.7658	762.8868	508.9272
478.1938	239.6008	4	A	13	1395.7232	698.3655	465.9130
591.2779	296.1428	5	I	12	1324.6861	662.8470	442.2339
704.3619	352.6849	6	L	11	1211.6020	606.3049	404.5392
819.3889	410.1983	7	D	10	1098.5180	549.7629	366.8445
956.4478	478.7278	8	H	9	983.4910	492.2494	328.5022
1027.4849	514.2464	9	A	8	846.4321	423.7200	282.8159
1126.5533	563.7806	10	V	7	775.3950	388.2014	259.1369
1255.5959	628.3019	11	E	6	676.3266	338.6672	226.1141
1356.6436	678.8257	12	T	5	547.2840	274.1459	183.0999
1413.6630	707.3364	13	G	4	446.2363	223.6221	149.4173
1512.7334	756.8706	14	V	3	389.2149	195.1113	130.4102
1627.7804	814.3841	15	D	2	290.1464	145.5771	97.3874
		16	R	1	175.1195	88.0637	59.0451



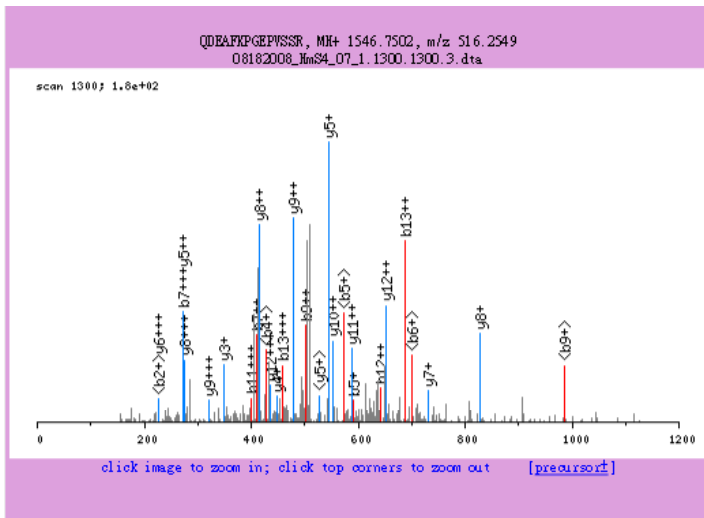
rrnAC2676

AVDPAGALADVASLVADGEQVVVVHGGSTK

Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	DeltaCn
1	3	2831.5	2831.5	0.0199	4.67	0.521



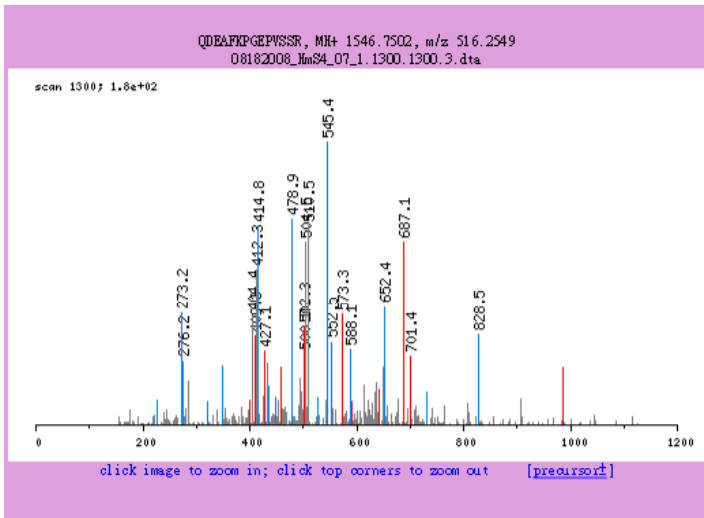
b ⁺	b ²⁺	#	AA	#	y ⁺	y ²⁺	y ³⁺
72.0449	36.5264	1	A	30			
171.1134	86.0606	2	V	29	2761.4372	1381.2225	921.1510
286.1403	143.5741	3	D	28	2662.3688	1331.6883	888.1281
383.1931	192.1004	4	P	27	2547.3419	1274.1748	849.7858
454.2302	227.6190	5	A	26	2450.2891	1225.6485	817.4349
511.2516	256.1297	6	G	25	2379.2520	1190.1299	793.7559
582.2888	291.6483	7	A	24	2322.2305	1161.6192	774.7487
695.3728	348.1903	8	L	23	2251.1934	1126.1006	751.0697
766.4099	383.7089	9	A	22	2138.1093	1069.5586	713.3750
881.4369	441.2223	10	D	21	2067.0722	1034.0400	689.6960
980.5053	490.7566	11	V	20	1952.0453	976.5266	651.3536
1051.5424	526.2751	12	A	19	1852.9769	926.9923	618.3308
1138.5744	569.7911	13	S	18	1781.9398	891.4738	594.6518
1251.6585	626.3332	14	L	17	1694.9077	847.9578	565.6411
1350.7269	675.8674	15	V	16	1581.8237	791.4157	527.9464
1421.7640	711.3859	16	A	15	1482.7552	741.8815	494.9236
1536.7910	768.8994	17	D	14	1411.7181	706.3630	471.2446
1593.8124	797.4101	18	G	13	1296.6912	648.8495	432.9023
1722.8530	861.9314	19	E	12	1239.6697	620.3988	413.8951
1850.9136	925.9607	20	Q	11	1110.6271	555.8175	370.8809
1949.9820	975.4949	21	V	10	982.5686	491.7882	328.1947
2049.0504	1025.0291	22	V	9	883.5001	442.2540	295.1719
2148.1188	1074.5633	23	V	8	784.4317	392.7198	262.1491
2247.1873	1124.0975	24	V	7	685.3633	343.1856	229.1263
2384.2462	1192.6270	25	I	6	586.2949	293.6514	196.1035
2441.2676	1221.1377	26	G	5	449.2360	225.1219	150.4172
2498.2891	1249.6485	27	G	4	392.2145	196.6112	131.4101
2585.3211	1293.1645	28	S	3	335.1931	168.1004	112.4029
2686.3688	1343.6883	29	T	2	248.1610	124.5844	83.3922
		30	K	1	147.1134	74.0606	49.7097



rrnAC2630

QDEAFKPGEPVSSR

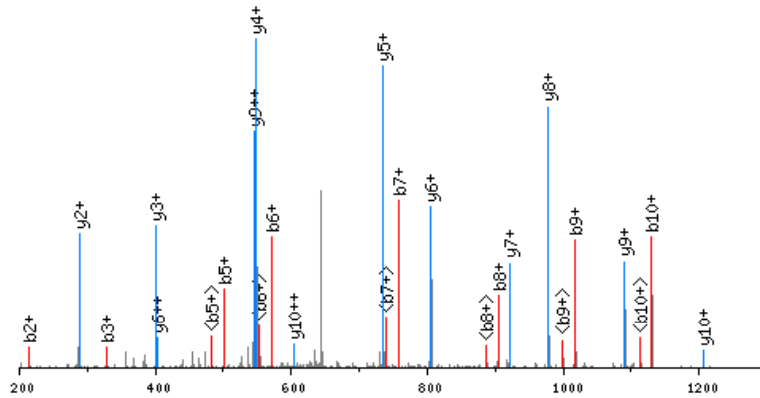
Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	Deltacn
0.9994	3	1545.7	1545.7	0.0015	2.625	0.161



b ⁺	b ²⁺	b ³⁺	#	AA	#	y ⁺	y ²⁺	y ³⁺
129.0664	65.0371	43.6940	1	Q	14			
244.0933	122.5506	82.0963	2	D	13	1418.6916	709.8497	473.5691
373.1359	187.0719	125.0305	3	E	12	1303.6646	652.3962	435.2268
444.1731	222.5904	148.7296	4	A	11	1174.6220	587.8149	392.2126
591.2415	296.1246	197.7524	5	F	10	1103.5849	552.2964	368.5335
719.3364	360.1721	240.4507	6	K	9	956.5165	478.7622	319.5107
816.3892	408.6985	272.8016	7	P	8	828.4216	414.7147	276.8124
873.4107	437.2092	291.8088	8	G	7	731.3688	366.1883	244.4615
1002.4533	501.7305	334.8230	9	E	6	674.3473	337.6776	225.4543
1099.5060	550.2569	367.1739	10	P	5	545.3047	273.1563	182.4401
1198.5744	599.7911	400.1967	11	V	4	448.2520	224.6299	150.0892
1285.6065	643.3071	429.2074	12	S	3	349.1836	175.0957	117.0664
1372.6385	686.8232	458.2180	13	S	2	262.1515	131.5797	88.0557
			14	R	1	175.1195	88.0637	59.0451

VDLGDAWFLLR, MH+ 1304.7003, m/z 652.8538
08182008_Xms2_07_2.3643.3643.2.dta

scan 3643; 2.7e+03



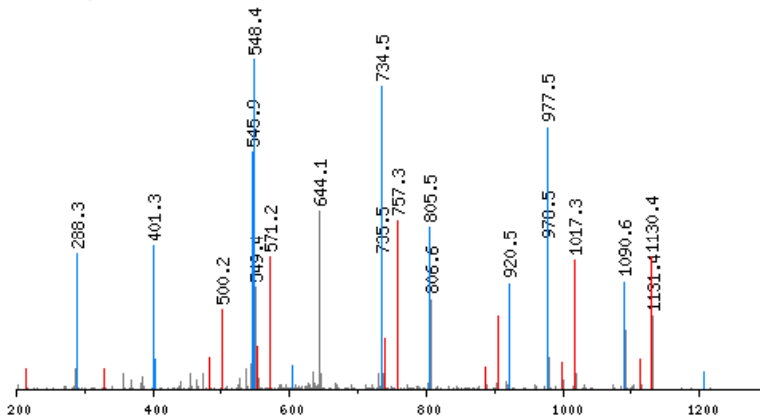
click image to zoom in; click top corners to zoom out [\[precursor!\]](#)

rrnAC2574 VDLGDAWFLLR

Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	DeltaCn
1	2	1303.7	1303.7	0.0009	4.365	0.432

VDLGDAWFLLR, MH+ 1304.7003, m/z 652.8538
08182008_Xms2_07_2.3643.3643.2.dta

scan 3643; 2.7e+03

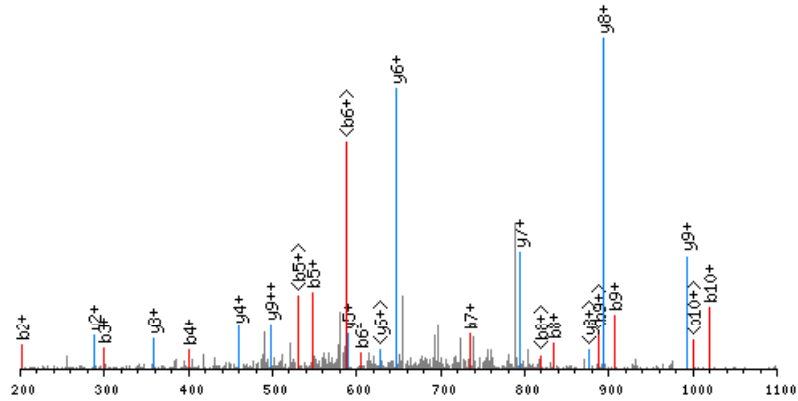


click image to zoom in; click top corners to zoom out [\[precursor!\]](#)

b ⁺	#	AA	#	y ⁺	y ²⁺
100.0762	1	V	11		
215.1032	2	D	10	1205.6319	603.3199
328.1872	3	L	9	1090.6049	545.8064
385.2087	4	G	8	977.5209	489.2644
500.2357	5	D	7	920.4994	460.7536
571.2728	6	A	6	805.4725	403.2401
757.3521	7	W	5	734.4354	367.7216
904.4205	8	F	4	548.3560	274.6819
1017.5046	9	L	3	401.2876	201.1477
1130.5886	10	L	2	288.2036	144.6057
	11	R	1	175.1195	88.0637

SLVTFGETALR, MH+ 1193.6530, m/z 597.3301
08182008_HmS1_12_2.2603.2603.2.dta

scan 2603: 6.4e+02



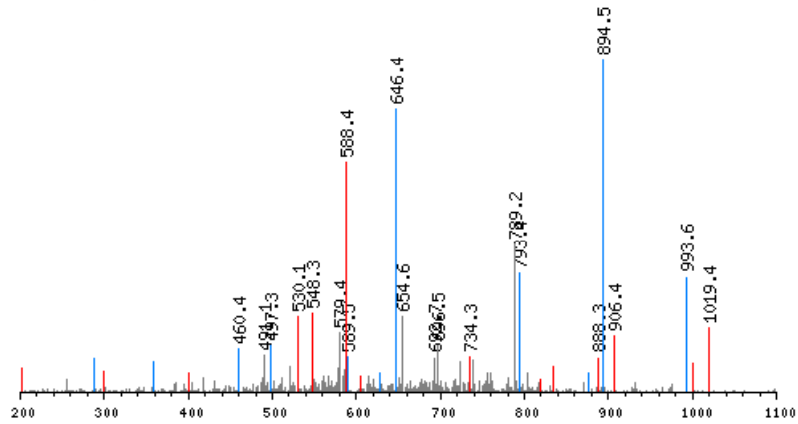
click image to zoom in; click top corners to zoom out [\[precursor\]](#)

rrnAC2551 SLVTFGETALR

Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	DeltaCn
0.9998	2	1192.6	1192.6	0.0014	2.845	0.442

SLVTFGETALR, MH+ 1193.6530, m/z 597.3301
08182008_HmS1_12_2.2603.2603.2.dta

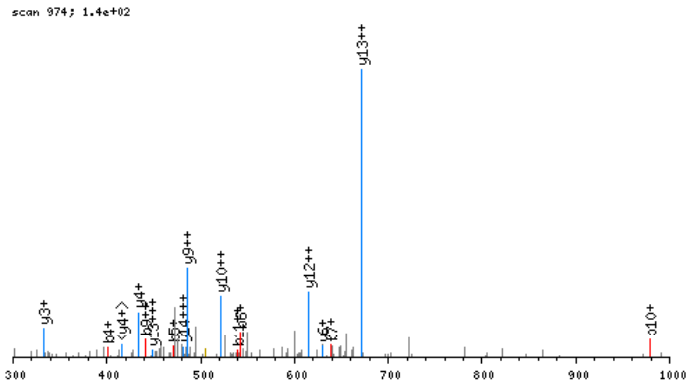
scan 2603: 6.4e+02



click image to zoom in; click top corners to zoom out [\[precursor\]](#)

b ⁺	#	AA	#	y ⁺	y ²⁺
88.0399	1	S	11		
201.1239	2	L	10	1106.6210	553.8144
300.1923	3	V	9	993.5369	497.2724
401.2400	4	T	8	894.4685	447.7382
548.3084	5	F	7	793.4208	397.2143
605.3299	6	G	6	646.3524	323.6801
734.3725	7	E	5	589.3310	295.1694
835.4202	8	T	4	460.2884	230.6481
906.4573	9	A	3	359.2407	180.1243
1019.5413	10	L	2	288.2036	144.6057
	11	R	1	175.1195	88.0637

AVNDAAPKDPVTGTR, MH+ 1511.7818, m/z 504.5988
08182008_HmS2_07_1_0974.0974.3.dta

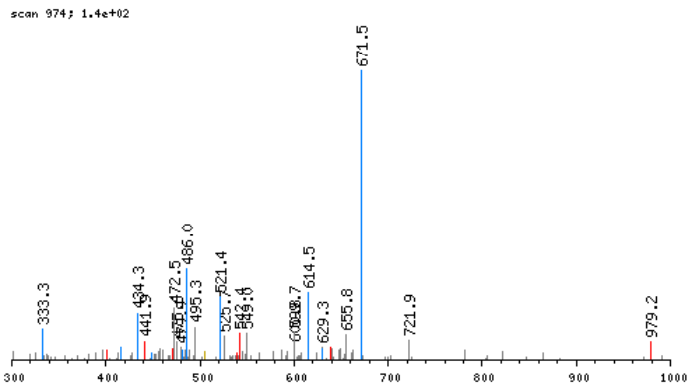


click image to zoom in; click top corners to zoom out [precursor]

rrnAC2470 AVNDAAPKDPVTGTR

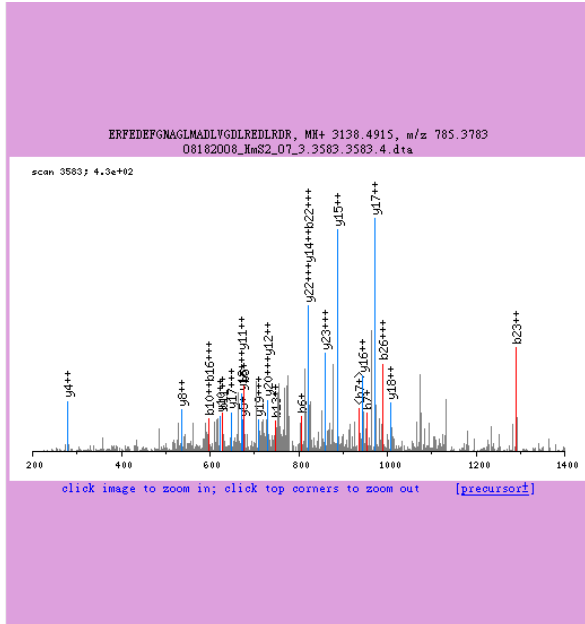
Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	Deltacn
0.9998	3	1510.8	1510.8	0.0043	2.088	0.297

AVNDAAPKDPVTGTR, MH+ 1511.7818, m/z 504.5988
08182008_HmS2_07_1_0974.0974.3.dta



click image to zoom in; click top corners to zoom out [precursor]

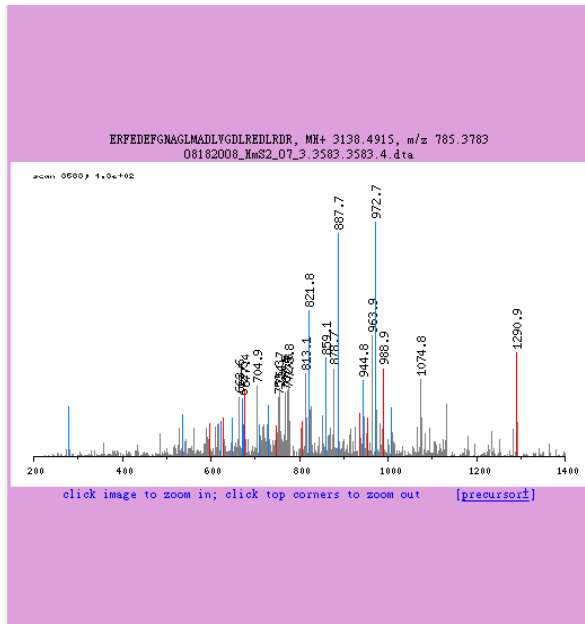
b ⁺	b ²⁺	#	AA	#	y ⁺	y ²⁺	y ³⁺
72.0449	36.5264	1	A	15			
171.1134	86.0606	2	V	14	1440.7447	720.8763	480.9201
285.1563	143.0821	3	N	13	1341.6763	671.3420	447.8973
400.1832	200.5955	4	D	12	1227.6333	614.3206	409.8830
471.2203	236.1141	5	A	11	1112.6064	556.8071	371.5407
542.2575	271.6326	6	A	10	1041.5693	521.2886	347.8616
639.3102	320.1590	7	P	9	970.5322	485.7700	324.1826
767.4052	384.2065	8	K	8	873.4794	437.2436	291.8317
882.4321	441.7200	9	D	7	745.3844	373.1961	249.1334
979.4849	490.2464	10	P	6	630.3575	315.6827	210.7910
1078.5533	539.7806	11	V	5	533.3047	267.1563	178.4401
1179.6010	590.3044	12	T	4	434.2363	217.6221	145.4173
1236.6224	618.8151	13	G	3	333.1886	167.0982	111.7348
1337.6701	669.3390	14	T	2	276.1672	138.5875	92.7276
		15	R	1	175.1195	88.0637	59.0451



rrnAC1709

ERFEDEFGNAGLMADLVGDLREDLRDR

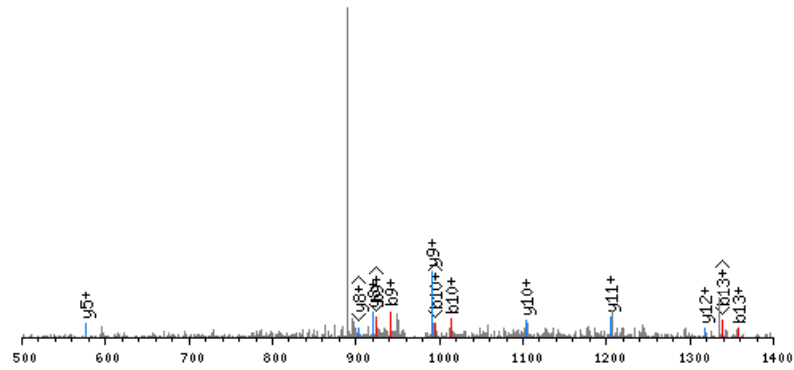
Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	DeltaCn
0.9955	4	3137.5	3137.5	0.0067	3.713	0.108



b ⁺	b ²⁺	b ³⁺	#	AA	#	y ⁺	y ²⁺	y ³⁺
130.0504	65.5291	44.0220	1	E	27			
286.1515	143.5797	96.0557	2	R	26	3009.4489	1505.2283	1003.8215
433.2199	217.1139	145.0785	3	F	25	2853.3477	1427.1778	951.7878
562.2625	281.6352	188.0927	4	E	24	2706.2793	1353.6436	902.7650
677.2895	339.1487	226.4350	5	D	23	2577.2367	1289.1223	859.7508
806.3321	403.6699	269.4492	6	E	22	2462.2098	1231.6088	821.4085
933.4003	477.2042	316.4720	7	F	21	2333.1672	1167.0673	776.3943
1010.4220	505.7149	337.4792	8	G	20	2186.0988	1093.5533	729.3715
1124.4649	562.7364	375.4935	9	N	19	2129.0773	1065.0426	710.3643
1195.5020	598.2549	399.1725	10	A	18	2015.0344	1008.0211	672.3500
1252.5235	626.7656	418.1797	11	G	17	1943.9973	972.5026	648.6710
1365.6075	683.3077	455.8744	12	L	16	1886.9758	943.9918	629.6638
1496.6480	748.8279	499.5546	13	M	15	1773.8918	887.4498	591.9691
1567.6851	784.3465	523.2336	14	A	14	1642.8513	821.9295	548.2890
1682.7121	841.8599	561.5759	15	D	13	1571.8142	786.4110	524.6099
1795.7961	898.4020	599.2706	16	L	12	1456.7872	728.8975	486.2676
1894.8645	947.9362	632.2934	17	V	11	1343.7032	672.3555	448.5729
1951.8860	976.4469	651.3006	18	G	10	1244.6347	622.8213	415.5501
2066.9129	1033.9604	689.6429	19	D	9	1187.6133	594.3105	396.5430
2179.9970	1090.5024	727.3716	20	L	8	1072.5863	536.7971	358.2007
2336.0981	1168.5530	779.3713	21	R	7	959.5023	480.2550	320.5060
2465.1407	1233.0743	822.3855	22	E	6	803.4012	402.2045	268.4723
2580.1677	1290.5877	860.7278	23	D	5	674.3586	337.6832	225.4581
2693.2517	1347.1298	898.4225	24	L	4	559.3316	280.1697	187.1158
2849.3528	1425.1803	950.4562	25	R	3	446.2476	223.6277	149.4211
2964.3798	1482.6938	988.7985	26	D	2	290.1464	145.5771	97.3874
			27	R	1	175.1195	88.0637	59.0451

VTDADLLTIADDEVTDAR, MH+ 1932.9402, m/z 966.9737
08182008_HmS2_07_3_3310.3310.2.dta

scan 3310; 7.9e+02



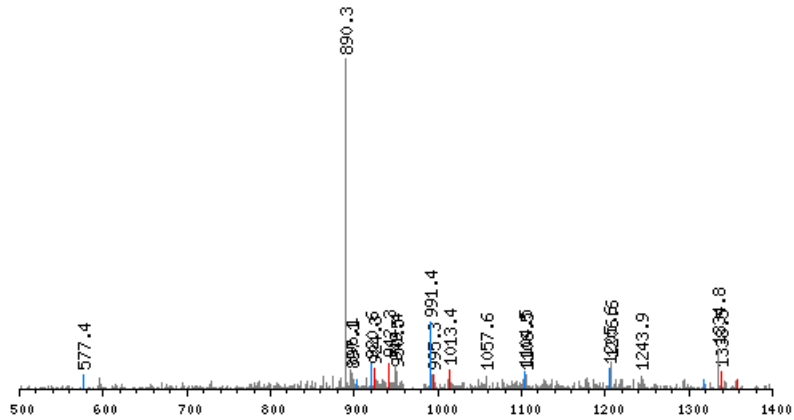
click image to zoom in; click top corners to zoom out [\[precursor\]](#)

rrnAC1690 VTDADLLTIADDEVTDAR

Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	Deltaen
1	2	1931.9	1931.9	0.0059	2.769	0.358

VTDADLLTIADDEVTDAR, MH+ 1932.9402, m/z 966.9737
08182008_HmS2_07_3_3310.3310.2.dta

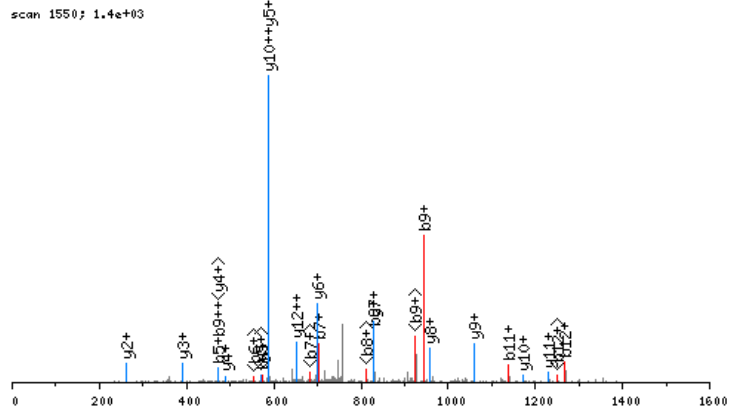
scan 3310; 7.9e+02



click image to zoom in; click top corners to zoom out [\[precursor\]](#)

b ⁺	#	AA	#	y ⁺	y ²⁺
100.0762	1	V	18		
201.1239	2	T	17	1833.8718	917.4398
316.1509	3	D	16	1732.8241	866.9160
387.1880	4	A	15	1617.7972	809.4025
502.2149	5	D	14	1546.7601	773.8839
615.2990	6	L	13	1431.7331	716.3705
728.3830	7	L	12	1318.6490	659.8284
829.4307	8	T	11	1205.5650	603.2864
942.5148	9	I	10	1104.5173	552.7626
1013.5519	10	A	9	991.4332	496.2205
1128.5788	11	D	8	920.3961	460.7020
1257.6214	12	E	7	805.3692	403.1885
1356.6899	13	V	6	676.3266	338.6672
1457.7375	14	T	5	577.2582	289.1330
1572.7645	15	D	4	476.2105	238.6092
1687.7914	16	D	3	361.1836	181.0957
1758.8285	17	A	2	246.1566	123.5822
	18	R	1	175.1195	88.0637

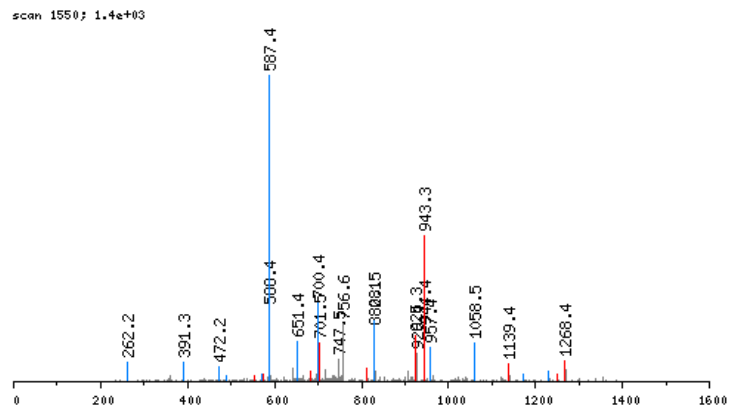
LDAGDTEQLPVESR, MH+ 1529.7447, m/z 765.3760
08182008_HmS2_07_3.1550.1550.2.dta



rrnAC1479 LDAGDTEQLPVESR

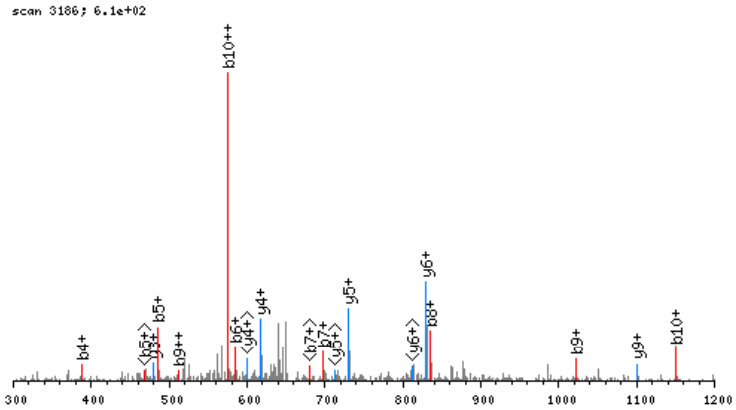
Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	Deltacn
1	2	1528.7	1528.7	0.001	4.362	0.429

LDAGDTEQLPVESR, MH+ 1529.7447, m/z 765.3760
08182008_HmS2_07_3.1550.1550.2.dta



b ⁺	b ²⁺	# AA	#	y ⁺	y ²⁺
114.0919	57.5499	1	I	14	
229.1188	115.0633	2	D	13	1416.6607
300.1559	150.5819	3	A	12	1301.6337
357.1774	179.0926	4	G	11	1230.5966
472.2044	236.6061	5	D	10	1173.5752
573.2520	287.1299	6	T	9	1058.5482
702.2946	351.6512	7	E	8	957.5005
830.3532	415.6805	8	Q	7	828.4579
943.4373	472.2225	9	L	6	700.3994
1040.4900	520.7489	10	P	5	587.3153
1139.5584	570.2831	11	V	4	490.2625
1268.6010	634.8044	12	E	3	391.1941
1355.6331	678.3204	13	S	2	262.1515
		14	R	1	175.1195

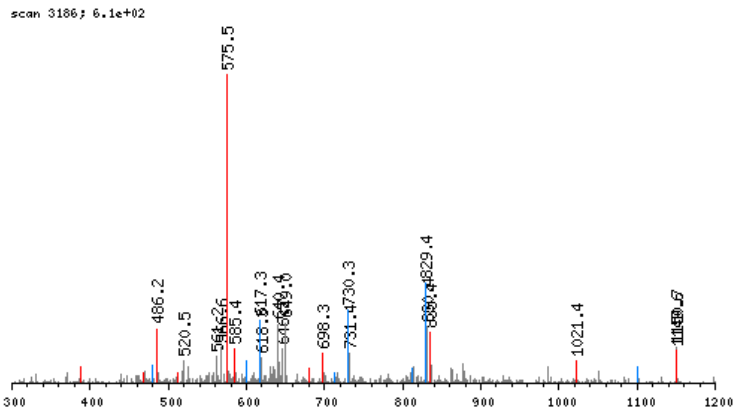
VDDGVVLHWQF, MK+ 1314.6483, m/z 657.8278
08182008_NmS1_12_2_3186.3186.2.dta



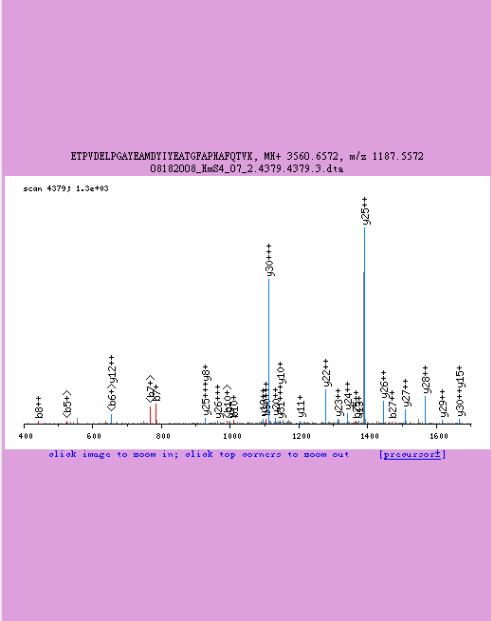
rrnAC1131 VDDGVVLHWQF

Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	Deltaen
1	2	1313.6	1313.6	0.0034	3.478	0.365

VDDGVVLHWQF, MK+ 1314.6483, m/z 657.8278
08182008_NmS1_12_2_3186.3186.2.dta



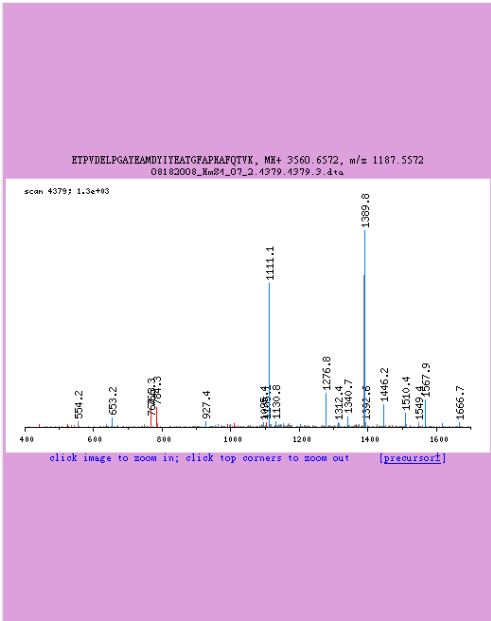
b ⁺	b ²⁺	#	AA	#	y ⁺	y ²⁺
100.0762	50.5420	1	V	11		
215.1032	108.0555	2	D	10	1215.5798	608.2938
330.1301	165.5690	3	D	9	1100.5529	550.7804
387.1516	194.0797	4	G	8	985.5260	493.2669
486.2200	243.6139	5	V	7	928.5045	464.7562
585.2884	293.1481	6	V	6	829.4361	415.2220
698.3725	349.6902	7	L	5	730.3677	365.6877
835.4314	418.2196	8	H	4	617.2836	309.1457
1021.5107	511.2593	9	W	3	480.2247	240.6163
1149.5693	575.2886	10	Q	2	294.1454	147.5766
		11	F	1	166.0868	83.5473



rrnAC0917

ETPVDELPGAYEAMDYIYEATGFAPHAFTVK

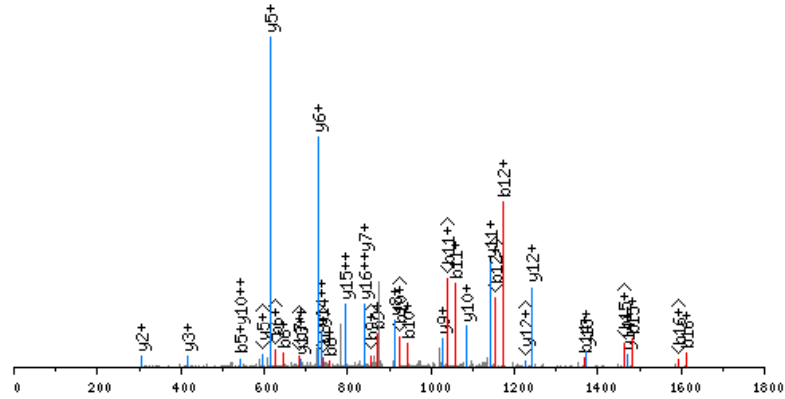
Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	Deltacn
1	3	3559.6	3560.6	0.9899	3.586	0.363



b*	b ²⁺	b ³⁺	# AA	#	y*	y ²⁺	y ³⁺
130.0504	65.5291	44.0220	1	E	32		
231.0981	116.0530	77.7046	2	T	31	3431.6146	1716.3112
328.1509	164.5793	110.0555	3	P	30	3330.5669	1665.7874
427.2193	214.1135	143.0783	4	Y	29	3233.5141	1617.2610
542.2462	271.6270	181.4206	5	D	28	3134.4457	1567.7268
671.2888	336.1483	224.4348	6	E	27	3019.4188	1510.2133
784.3729	392.6903	262.1295	7	L	26	2890.3762	1445.6920
881.4256	441.2167	294.4804	8	P	25	2777.2921	1389.1500
938.4471	469.7275	313.4876	9	G	24	2680.2394	1340.6236
1009.4842	505.2460	337.1666	10	A	23	2623.2179	1312.1129
1172.5475	586.7777	391.5211	11	Y	22	2552.1808	1276.5943
1301.5901	651.2990	434.5353	12	E	21	2389.1175	1195.0626
1372.6273	686.8175	458.2143	13	A	20	2260.0749	1130.5413
1503.6677	752.3378	501.8945	14	M	19	2189.0378	1095.0228
1618.6947	809.8513	540.2368	15	D	18	2057.9973	1029.5025
1781.7580	891.3829	594.5912	16	Y	17	1942.9703	971.9891
1894.8421	947.9249	632.2859	17	I	16	1779.9070	890.4574
2057.9054	1029.4566	686.6404	18	Y	15	1666.8229	833.9154
2186.9480	1093.9779	729.6545	19	E	14	1503.7596	752.3837
2257.9851	1129.4965	753.3336	20	A	13	1374.7170	687.8624
2359.0328	1180.0203	787.0161	21	T	12	1303.6799	652.3439
2416.0542	1208.5310	806.0233	22	G	11	1202.6322	601.8200
2563.1227	1282.0652	855.0461	23	F	10	1145.6108	573.3093
2634.1598	1317.5838	878.7251	24	A	9	998.5423	499.7751
2731.2125	1366.1102	911.0761	25	P	8	927.5052	464.2565
2868.2715	1434.6396	956.7624	26	H	7	830.4525	415.7901
2939.3086	1470.1582	980.4414	27	A	6	693.3936	347.2007
3086.3770	1543.6924	1029.4642	28	F	5	622.3564	311.6821
3214.4356	1607.7217	1072.1504	29	Q	4	475.2880	238.1479
3315.4832	1658.2455	1105.8330	30	T	3	347.2294	174.1186
3414.5517	1707.7797	1138.8558	31	Y	2	246.1818	123.5948
			32	K	1	147.1134	74.0606

TVLTQTGGDALNPTLER, MH+ 1785.9347, m/z 893.4710
20081001M4_685_968.01566.01566.2.dta

scan 1566: 1.6e+03



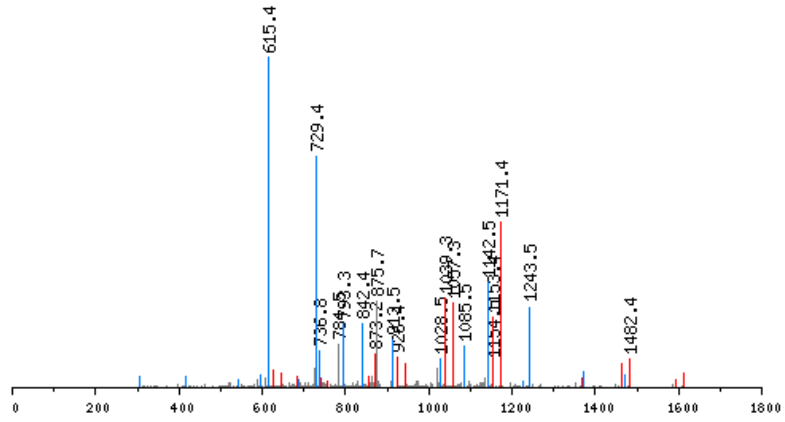
click image to zoom in; click top corners to zoom out [\[precursor!\]](#)

rrnAC0845 TVLTQTGGDALNPTLER

Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	Deltaen
1	2	1784.9	1784.9	0.000152	4.429	0.512

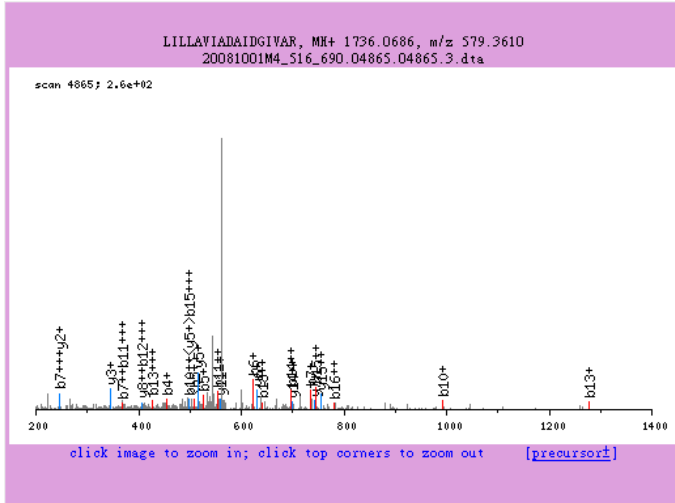
TVLTQTGGDALNPTLER, MH+ 1785.9347, m/z 893.4710
20081001M4_685_968.01566.01566.2.dta

scan 1566: 1.6e+03



click image to zoom in; click top corners to zoom out [\[precursor!\]](#)

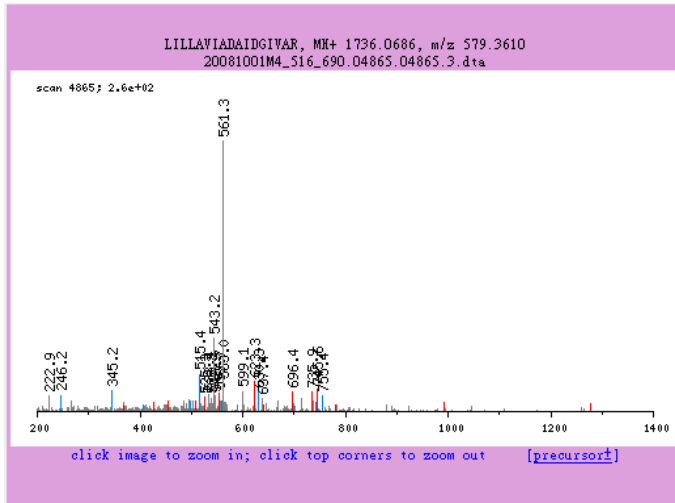
b ⁺	#	AA	#	y ⁺	y ²⁺
102.0555	1	T	17		
201.1239	2	V	16	1684.8870	842.9474
314.2080	3	L	15	1585.8186	793.4132
415.2557	4	T	14	1472.7345	736.8712
543.3142	5	Q	13	1371.6868	686.3473
644.3619	6	T	12	1243.6283	622.3180
701.3834	7	G	11	1142.5806	571.7942
758.4048	8	G	10	1085.5591	543.2835
873.4318	9	D	9	1028.5376	514.7727
944.4689	10	A	8	913.5107	457.2593
1057.5530	11	L	7	842.4736	421.7407
1171.5959	12	N	6	729.3895	365.1987
1268.6487	13	P	5	615.3466	308.1772
1369.6963	14	T	4	518.2938	259.6508
1482.7804	15	L	3	417.2462	209.1270
1611.8230	16	E	2	304.1621	152.5850
	17	R	1	175.1195	88.0637



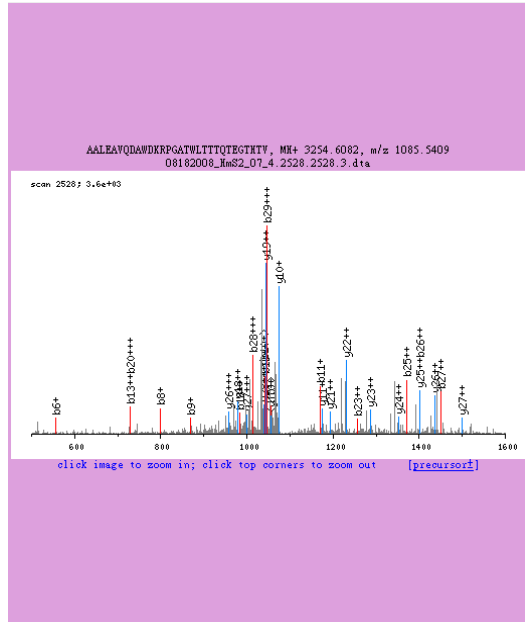
rrnAC0788

LILLAVIADAIDGIVAR

Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	Deltacn
0.9985	3	1735.1	1735.1	-0.0004	2.871	0.143



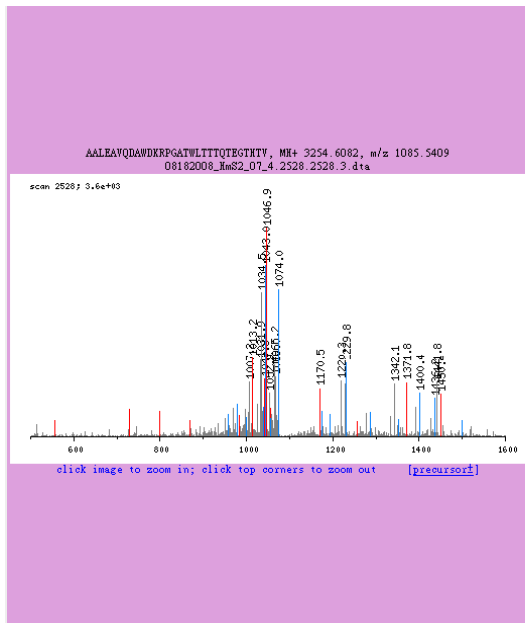
b ⁺	b ²⁺	b ³⁺	#	AA	#	y ⁺	y ²⁺	y ³⁺
114.0919	57.5499	38.7025	1	L	17			
227.1760	114.0919	76.3972	2	I	16	1622.9845	811.9962	541.6667
340.2600	170.6339	114.0919	3	L	15	1509.9005	755.4541	503.9720
453.3441	227.1760	151.7866	4	L	14	1396.8164	698.9121	466.2773
524.3812	262.6945	175.4656	5	A	13	1283.7323	642.3701	428.5827
623.4496	312.2287	208.4884	6	V	12	1212.6952	606.8515	404.9036
736.5337	368.7707	246.1831	7	I	11	1113.6268	557.3173	371.8808
807.5708	404.2893	269.8621	8	A	10	1000.5427	500.7753	334.1861
922.5977	461.8028	308.2045	9	D	9	929.5056	465.2567	310.5071
993.6348	497.3213	331.8835	10	A	8	814.4787	407.7433	272.1648
1106.7189	553.8634	369.5782	11	I	7	743.4416	372.2247	248.4857
1221.7459	611.3768	407.9205	12	D	6	630.3575	315.6827	210.7910
1278.7673	639.8876	426.9277	13	G	5	515.3306	258.1692	172.4487
1391.8514	696.4296	464.6223	14	I	4	458.3091	229.6585	153.4416
1490.9198	745.9638	497.6451	15	V	3	345.2250	173.1164	115.7469
1561.9569	781.4824	521.3242	16	A	2	246.1566	123.5822	82.7241
			17	R	1	175.1195	88.0637	59.0451



rrnAC0753

AALEAVQDAWDKRPGATWLTTTQTEGTHTV

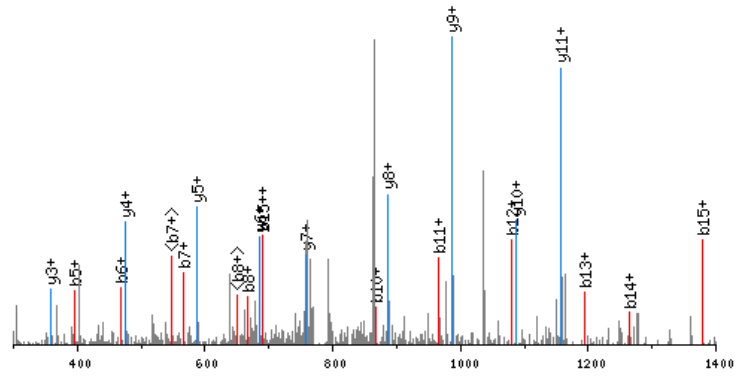
Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	Deltacn
1	3	3253.6	3254.6	1.0179	5.036	0.349



b ⁺	b ²⁺	b ³⁺	#	AA	#	y ⁺	y ²⁺	y ³⁺
72.0449	96.5264	24.6869	1	A	30			
143.0821	72.0449	48.3659	2	A	29	3183.5711	1592.2895	1061.8622
256.1661	128.5870	86.0606	3	L	28	3112.5340	1556.7709	1038.1832
385.2087	193.1083	129.0748	4	E	27	2999.4499	1500.2289	1000.4885
456.2458	228.6268	152.7538	5	A	26	2870.4073	1435.7076	957.4743
555.3142	278.1610	185.7766	6	V	25	2799.3702	1400.1890	939.1953
683.3728	342.1903	228.4628	7	Q	24	2700.3018	1350.6548	900.7725
798.3998	399.7038	266.8031	8	D	23	2572.2432	1286.6255	858.0863
869.4369	435.2223	290.4842	9	A	22	2457.2163	1229.1120	819.7440
1055.5162	528.2620	352.5106	10	W	21	2386.1791	1193.5935	796.0649
1170.5431	585.7755	390.8529	11	D	20	2200.0998	1100.5538	734.0385
1298.6381	649.8230	433.5512	12	K	19	2085.0729	1043.0404	695.6962
1454.7392	727.8735	485.5850	13	R	18	1956.9779	978.9929	652.9979
1551.7920	776.3999	517.9359	14	P	17	1800.8768	900.9423	600.9642
1608.8134	804.9106	536.9430	15	G	16	1703.8241	852.4159	568.6132
1679.8505	840.4292	560.6221	16	A	15	1646.8026	823.9052	549.6061
1780.8982	890.9530	594.3046	17	T	14	1575.7655	788.3867	525.9270
1966.9775	983.9927	656.3311	18	W	13	1474.7178	737.8628	492.2445
2080.0616	1040.5347	694.0257	19	L	12	1288.6385	644.8232	430.2180
2181.1093	1091.0586	727.7083	20	T	11	1175.5544	588.2811	392.5234
2282.1570	1141.5824	761.3909	21	T	10	1074.5067	537.7573	358.8408
2383.2046	1192.1062	795.0734	22	T	9	973.4591	487.2334	325.1582
2511.2632	1256.1355	837.7596	23	Q	8	872.4114	436.7096	291.4757
2612.3109	1306.6594	871.4422	24	T	7	744.3528	372.6803	248.7895
2741.3535	1371.1807	914.4564	25	E	6	643.3051	322.1565	215.1069
2798.3749	1399.6914	933.4635	26	G	5	514.2625	257.6352	172.0927
2899.4226	1450.2152	967.1461	27	T	4	457.2411	229.1244	153.0856
3036.4815	1518.7447	1012.8324	28	I	3	356.1934	178.6006	119.4030
3137.5292	1569.2685	1046.5150	29	T	2	219.1345	110.0712	73.7167
			30	V	1	118.0868	59.5473	40.0342

GPAGIAVTEAVLDAIR, MH+ 1552.8699, m/z 776.9386
08182008_MmS3_12_full11.3715.3715.2.dta

scan 3715; 2.4e+02



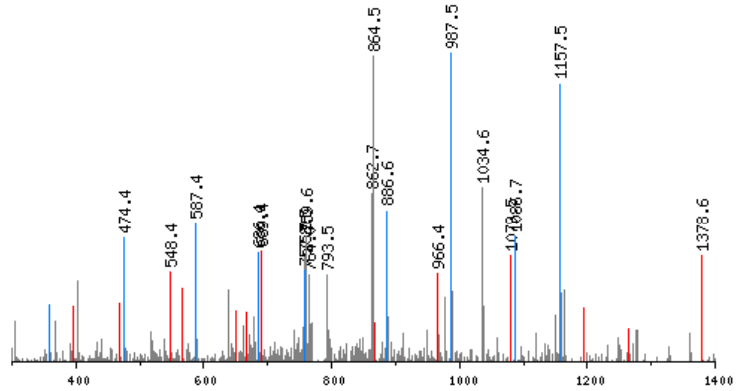
click image to zoom in; click top corners to zoom out [\[precursor!\]](#)

rrnAC0547 GPAGIAVTEAVLDAIR

Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	DeltaC
1	2	1551.9	1551.8	-0.0285	4.075	0.258

GPAGIAVTEAVLDAIR, MH+ 1552.8699, m/z 776.9386
08182008_MmS3_12_full11.3715.3715.2.dta

scan 3715; 2.4e+02

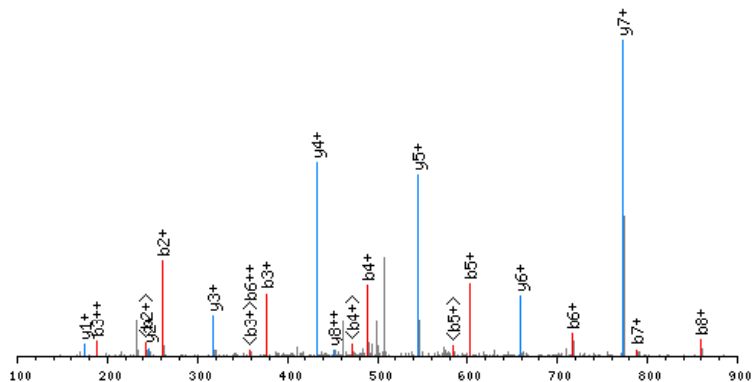


click image to zoom in; click top corners to zoom out [\[precursor!\]](#)

b ⁺	b ²⁺	#	AA	#	y ⁺	y ²⁺
58.0293	29.5186	1	G	16		
155.0821	78.0449	2	P	15	1495.8484	748.4281
226.1192	113.5635	3	A	14	1398.7957	699.9017
283.1406	142.0742	4	G	13	1327.7585	664.3832
396.2247	198.6163	5	I	12	1270.7371	635.8725
467.2618	234.1348	6	A	11	1157.6530	579.3304
566.3302	283.6690	7	V	10	1086.6159	543.8119
667.3779	334.1929	8	T	9	987.5475	494.2777
796.4205	398.7142	9	E	8	886.4998	443.7538
867.4576	434.2327	10	A	7	757.4572	379.2325
966.5260	483.7669	11	V	6	686.4201	343.7140
1079.6101	540.3090	12	L	5	587.3517	294.1798
1194.6370	597.8224	13	D	4	474.2676	237.6377
1265.6741	633.3410	14	A	3	359.2407	180.1243
1378.7582	689.8830	15	I	2	288.2036	144.6057
		16	R	1	175.1195	88.0637

MEDLLDAAR, MH+ 1033.4988, m/z 517.2530
08182008_HmS3_12_1.2251.2251.2.dta

scan 2251; 8.3e+02



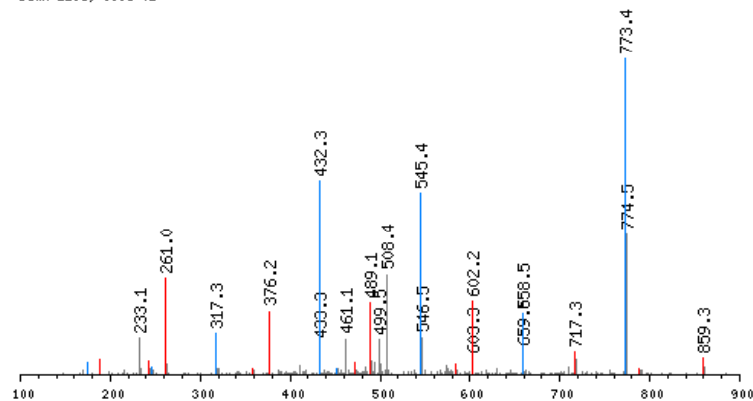
click image to zoom in; click top corners to zoom out [\[precursor!\]](#)

rrnAC0446 MEDLLDAAR

Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	Deltacn
0.9996	2	1032.5	1032.5	0.0004	3.1	0.294

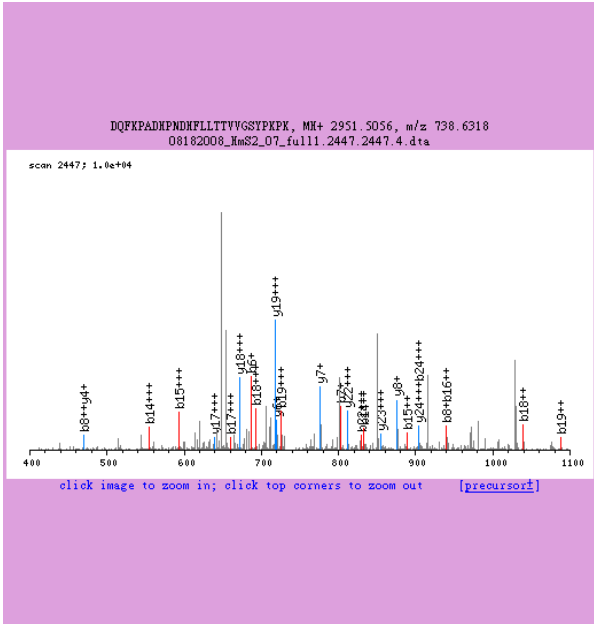
MEDLLDAAR, MH+ 1033.4988, m/z 517.2530
08182008_HmS3_12_1.2251.2251.2.dta

scan 2251; 8.3e+02



click image to zoom in; click top corners to zoom out [\[precursor!\]](#)

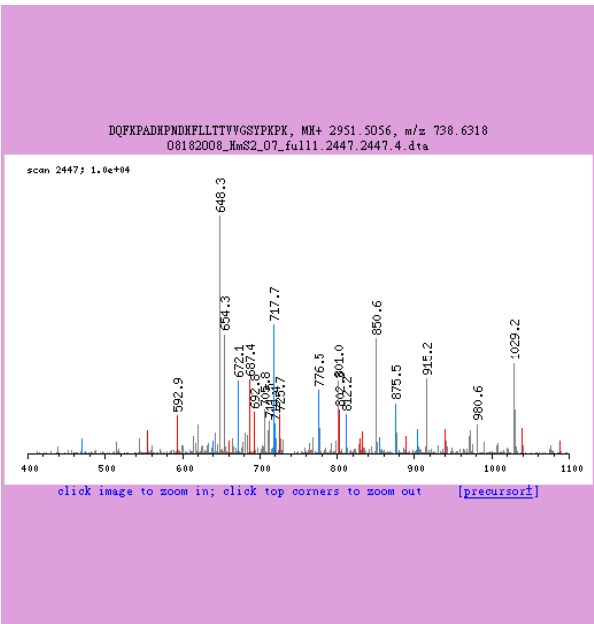
b ⁺	b ²⁺	#	AA	#	y ⁺	y ²⁺
132.0483	66.5281	1	M	9		
261.0909	131.0494	2	E	8	902.4583	451.7331
376.1178	188.5628	3	D	7	773.4157	387.2118
489.2019	245.1049	4	I	6	658.3888	329.6983
602.2860	301.6469	5	L	5	545.3047	273.1563
717.3129	359.1604	6	D	4	432.2207	216.6142
788.3500	394.6789	7	A	3	317.1937	159.1008
859.3871	430.1975	8	A	2	246.1566	123.5822
		9	R	1	175.1195	88.0637



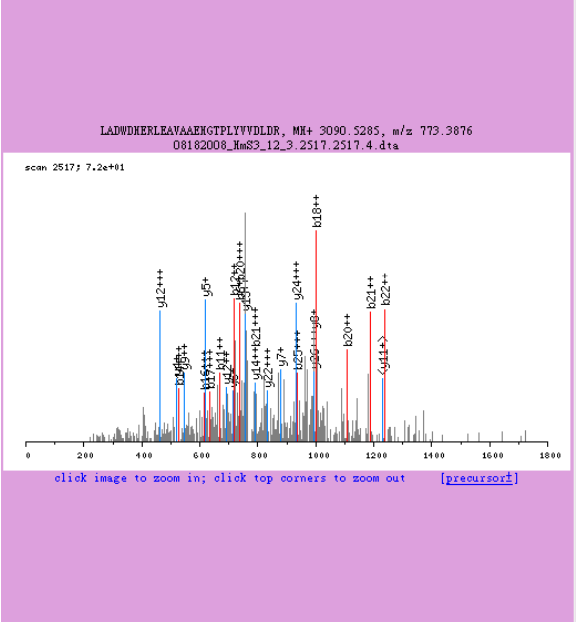
rrnAC0254

DQFKPADHPNDHLLTTVVGSPYKPK

Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	Deltacn
0.9999	4	2950.5	2951.5	0.9933	3.425	0.314



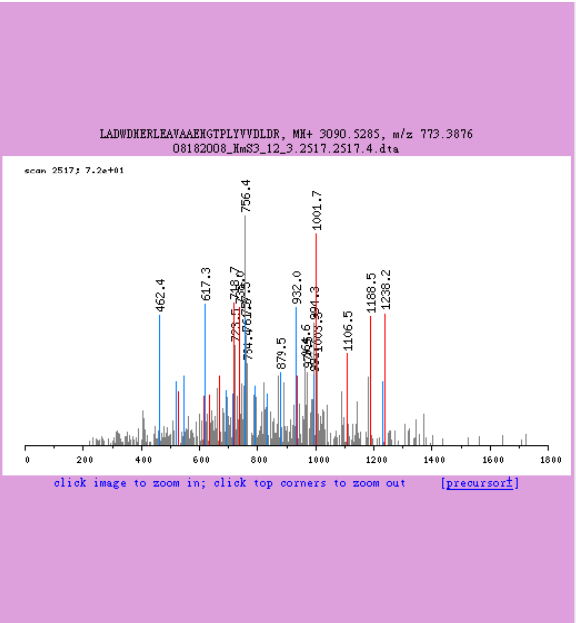
b ⁺	b ²⁺	b ³⁺	#	AA	#	y ⁺	y ²⁺	y ³⁺
116.0348	58.5213	39.3501	1	D	26			
244.0933	122.5506	82.0363	2	Q	25	2836.4786	1418.7432	946.1648
391.1618	196.0848	131.0591	3	F	24	2708.4200	1354.7139	903.4786
519.2567	260.1323	173.7575	4	X	23	2561.3516	1281.1797	854.4558
616.3095	308.6587	206.1084	5	P	22	2433.2567	1217.1322	811.7574
687.3466	344.1772	229.7874	6	A	21	2336.2039	1168.6059	779.4065
802.3735	401.6907	268.1297	7	D	20	2265.1668	1133.0873	755.7275
939.4325	470.2201	313.8160	8	H	19	2150.1398	1075.5738	717.3852
1036.4852	518.7465	346.1670	9	P	18	2013.0809	1007.0444	671.6989
1150.5281	575.7680	384.1813	10	M	17	1916.0282	958.5180	639.3479
1265.5551	633.2815	422.5236	11	D	16	1801.9852	901.4965	601.3336
1402.6140	701.8109	468.2099	12	H	15	1686.9583	843.9831	562.9913
1549.6824	775.3451	517.2327	13	F	14	1549.8994	775.4536	517.3050
1662.7665	831.8872	554.9274	14	I	13	1402.8310	701.9194	468.2822
1775.8505	888.4292	592.6221	15	L	12	1289.7469	645.2774	430.5875
1876.8982	938.9530	626.3046	16	T	11	1176.6628	588.8353	392.8928
1977.9459	989.4769	659.9872	17	T	10	1075.6152	538.3115	359.2103
2077.0143	1039.0111	693.0100	18	V	9	974.5675	487.7877	325.5277
2176.0827	1088.5453	726.0328	19	V	8	875.4991	438.2535	292.5049
2233.1042	1117.0560	745.0399	20	G	7	776.4307	388.7192	259.4821
2320.1362	1160.5720	774.0506	21	S	6	719.4092	360.2085	240.4750
2483.1995	1242.1037	828.4051	22	Y	5	632.3772	316.6925	211.4643
2580.2523	1290.6301	860.7560	23	P	4	469.3138	235.1608	157.1098
2708.3473	1354.6776	903.4543	24	X	3	372.2611	186.6345	124.7589
2805.4000	1403.2039	935.8052	25	P	2	244.1661	122.5870	82.0606
			26	X	1	147.1134	74.0606	49.7097



rrnAC0198

LADWDHERLEAVAAEHGTPLYVVDLDR

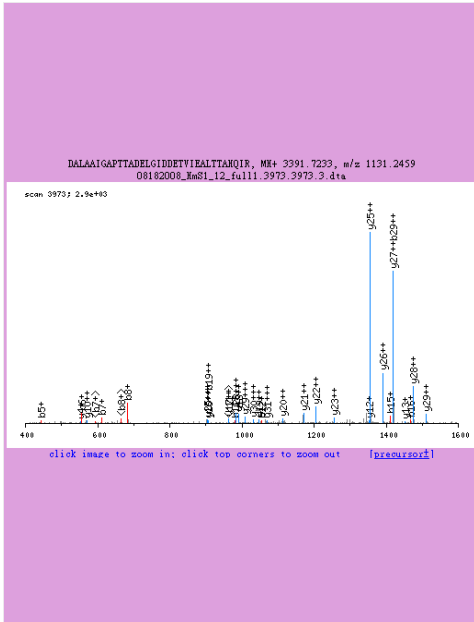
Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	Deltacn
1	4	3089.5	3091.5	2.0104	3.64	0.228



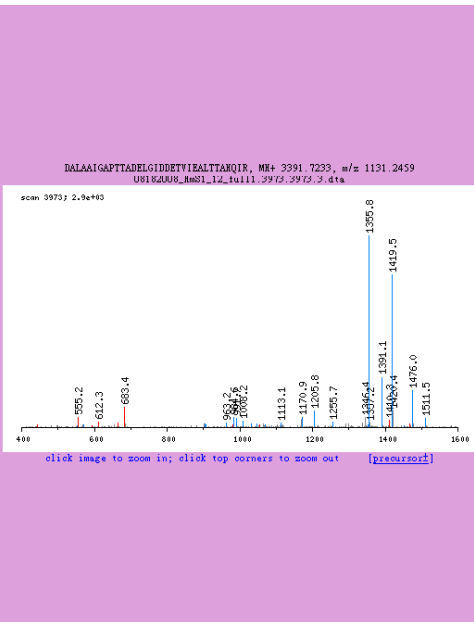
b ⁺	b ²⁺	b ³⁺	#	AA	#	y ⁺	y ²⁺	y ³⁺
114.0919	57.5499	38.7025	1	L	27			
185.1290	93.0684	62.3816	2	A	26	2977.4444	1489.2261	993.1534
300.1559	150.5819	100.7239	3	D	25	2906.4073	1453.7076	969.4743
486.2353	243.6215	162.7503	4	W	24	2791.3804	1396.1941	931.1320
601.2622	301.1350	201.0926	5	D	23	2605.3011	1303.1544	869.1056
738.3211	369.6645	246.7789	6	I	22	2490.2741	1245.6410	830.7633
867.3637	434.1858	289.7931	7	E	21	2353.2152	1177.1115	785.0769
1023.4648	512.2363	341.8268	8	R	20	2224.1726	1112.5902	742.0628
1136.5489	568.7784	379.5215	9	L	19	2068.0715	1034.5397	690.0290
1265.5915	633.2997	422.5357	10	E	18	1954.9874	977.9976	652.3344
1336.6286	668.8182	446.2147	11	A	17	1825.9448	913.4763	609.3202
1435.6970	718.3524	479.2376	12	V	16	1754.9077	877.9578	585.6411
1506.7341	753.8710	502.9166	13	A	15	1655.8393	828.4236	552.6183
1577.7712	789.3895	526.5956	14	A	14	1584.8022	792.9050	528.3393
1706.8138	853.9108	569.6098	15	E	13	1513.7651	757.3865	505.2602
1843.8727	922.4403	615.2961	16	I	12	1384.7225	692.8652	462.2460
1900.8942	950.9510	634.3033	17	G	11	1247.6636	624.3357	416.5597
2001.9419	1001.4749	667.9858	18	T	10	1190.6421	595.8250	397.5526
2098.9946	1050.0012	700.3368	19	P	9	1089.5944	545.3011	363.8700
2212.0787	1106.5433	738.0315	20	L	8	992.5417	496.7747	331.5191
2375.1420	1188.0749	792.3859	21	Y	7	879.4576	440.2327	293.8244
2474.2104	1237.6091	825.4087	22	V	6	716.3943	358.7011	239.4700
2573.2789	1287.1433	858.4315	23	V	5	617.3259	309.1668	206.4472
2688.3058	1344.6568	896.7738	24	D	4	518.2575	259.6326	173.4244
2801.3899	1401.1988	934.4685	25	L	3	403.2305	202.1192	135.0821
2916.4168	1458.7123	972.8108	26	D	2	290.1464	145.5771	97.3874
			27	R	1	175.1195	88.0637	59.0451

rrnAC0175

DALAAIGAPTTADELGIDDETVEALTTAHQIR



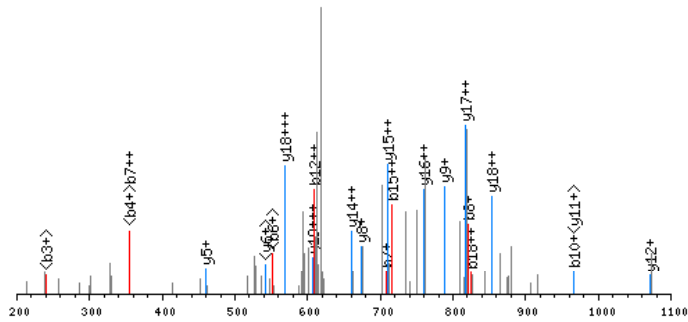
Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	Deltaen
1	3	3390.7	3392.7	2.0229	3.441	0.341



b ⁺	b ²⁺	#	AA	#	y ⁺	y ²⁺	y ³⁺
116.0948	58.5213	1	D	33			
187.0719	94.0399	2	A	32	3276.6963	1638.8521	1092.9040
300.1559	150.5819	3	L	31	3205.6592	1603.3295	1069.2250
371.1931	186.1004	4	A	30	3092.5752	1546.7915	1031.5303
442.2302	221.6190	5	A	29	3021.5380	1511.2729	1007.8512
555.3142	278.1610	6	I	28	2950.5009	1475.7544	984.1722
612.3357	306.6718	7	G	27	2837.4169	1419.2129	946.4775
683.3728	342.1903	8	A	26	2780.3954	1390.7016	927.4704
780.4256	390.7167	9	P	25	2709.3583	1355.1831	903.7913
881.4733	441.2405	10	T	24	2612.3055	1306.6567	871.4404
982.5209	491.7644	11	T	23	2511.2579	1256.1328	837.7578
1053.5580	527.2829	12	A	22	2410.2102	1205.6090	804.0753
1168.5850	584.7964	13	D	21	2339.1731	1170.0904	780.3962
1297.6276	649.3177	14	E	20	2224.1461	1112.5770	742.0539
1410.7116	705.8597	15	L	19	2095.1035	1048.0557	699.0397
1467.7331	734.3705	16	G	18	1982.0195	991.5136	661.3450
1580.8172	790.9125	17	I	17	1924.9980	963.0029	642.3379
1695.8441	848.4260	18	D	16	1811.9139	906.4609	604.6432
1810.8711	905.9394	19	D	15	1696.8870	848.9474	566.3009
1939.9137	970.4607	20	E	14	1581.8600	791.4339	527.9586
2040.9613	1020.9846	21	T	13	1452.8175	726.9126	484.9444
2140.0297	1070.5188	22	V	12	1351.7698	676.3888	451.2618
2253.1138	1127.0608	23	I	11	1252.7014	626.8546	418.2390
2382.1564	1191.5821	24	E	10	1139.6173	570.3126	380.5443
2453.1935	1227.1007	25	A	9	1010.5747	505.7913	337.5301
2566.2776	1283.6427	26	L	8	939.5376	470.2727	313.8511
2667.3253	1334.1665	27	T	7	826.4535	413.7307	276.1564
2768.3729	1384.6904	28	T	6	725.4058	363.2068	242.4738
2839.4101	1420.2089	29	A	5	624.3582	312.6830	208.7913
2976.4690	1488.7384	30	H	4	553.3211	277.1644	185.1122
3104.5275	1552.7671	31	Q	3	416.2621	208.6350	139.4259
3217.6116	1609.3097	32	I	2	288.2036	144.6057	96.7397
		33	R	1	175.1195	88.0637	59.0451

ANADVHHISGHDGGTGASPK, MH+ 1889.9106, m/z 630.6417
08182008_HmS2_07_2_0973.0973.3.dta

scan 973; 2.3e+01



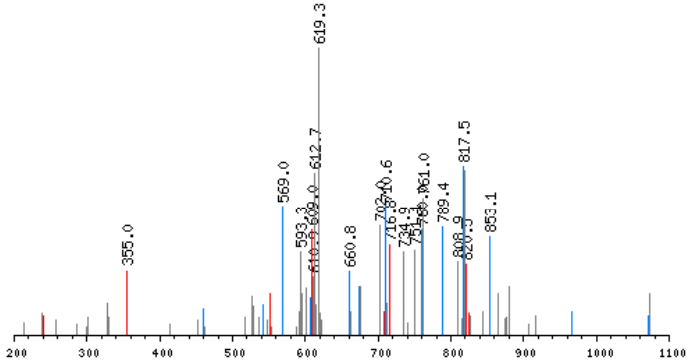
click image to zoom in; click top corners to zoom out [precursor]

rrnAC0169 ANADVHHISGHDGGTGASPK

Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	Deltaen
1	3	1888.9	1889.9	1.0033	2.662	0.283

ANADVHHISGHDGGTGASPK, MH+ 1889.9106, m/z 630.6417
08182008_HmS2_07_2_0973.0973.3.dta

scan 973; 2.3e+01

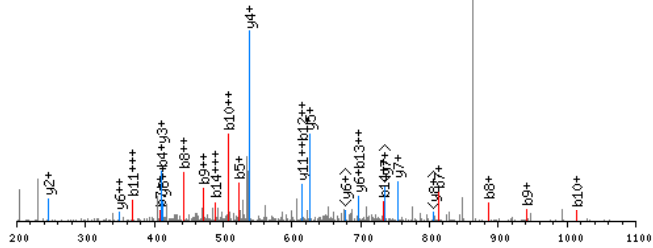


click image to zoom in; click top corners to zoom out [precursor]

b ⁺	b ²⁺	#	AA	#	y ⁺	y ²⁺	y ³⁺
72.0449	36.5264	1	A	20			
186.0879	93.5478	2	N	19	1818.8735	909.9406	606.9630
257.1250	129.0664	3	A	18	1704.8305	852.9192	568.9487
372.1519	186.5799	4	D	17	1633.7934	817.4006	545.2697
471.2203	236.1141	5	Y	16	1518.7665	759.8872	506.9274
570.2888	285.6483	6	V	15	1419.6981	710.3529	473.9046
707.3477	354.1777	7	H	14	1320.6297	660.8187	440.8818
820.4317	410.7198	8	I	13	1183.5707	592.2893	395.1955
907.4638	454.2358	9	S	12	1070.4867	535.7473	357.5008
964.4852	482.7465	10	G	11	983.4546	492.2312	328.4901
1101.5441	551.2760	11	H	10	926.4332	463.7205	309.4829
1216.5711	608.7894	12	D	9	789.3743	395.1910	263.7966
1273.5925	637.3002	13	G	8	674.3473	337.6776	225.4543
1330.6140	665.8109	14	G	7	617.3259	309.1668	206.4472
1431.6617	716.3348	15	T	6	560.3044	280.6561	187.4400
1488.6831	744.8455	16	G	5	459.2567	230.1323	153.7575
1559.7203	780.3640	17	A	4	402.2353	201.6215	134.7503
1646.7523	823.8801	18	S	3	331.1981	166.1030	111.0713
1743.8050	872.4064	19	P	2	244.1661	122.5870	82.0606
		20	K	1	147.1134	74.0606	49.7097

VAHTLYEAGASEYAR, MH+ 1637.7924, m/z 546.6023
08182008_ImmS2_07_4.1573.1573.3.dta

scan 1573; 9.8e+02



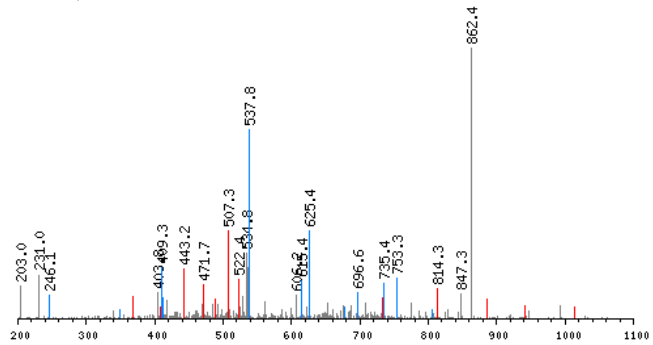
click image to zoom in; click top corners to zoom out [\[precursor±\]](#)

pNG7330 VAHTLYEAGASEYAR

Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	Deltaen
1	3	1636.8	1636.8	-0.0064	3.073	0.371

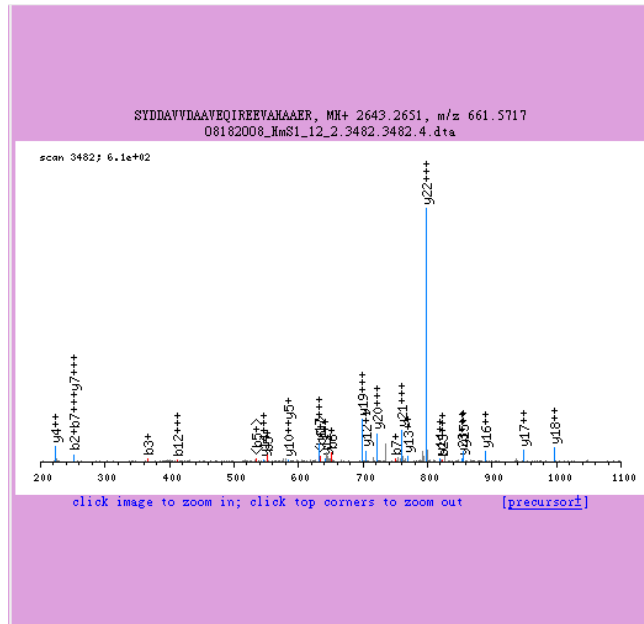
VAHTLYEAGASEYAR, MH+ 1637.7924, m/z 546.6023
08182008_ImmS2_07_4.1573.1573.3.dta

scan 1573; 9.8e+02



click image to zoom in; click top corners to zoom out [\[precursor±\]](#)

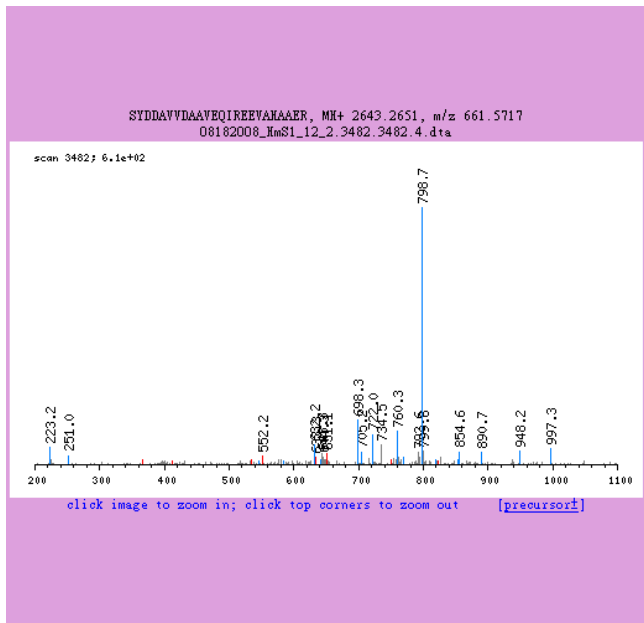
b ⁺	b ²⁺	b ³⁺	#	AA	#	y ⁺	y ²⁺	y ³⁺
100.0762	50.5420	34.0306	1	V	15			
171.1134	86.0606	57.7097	2	A	14	1538.7239	769.8659	513.5799
308.1723	154.5900	103.3960	3	I	13	1467.6868	734.3473	489.9008
409.2199	205.1139	137.0785	4	T	12	1330.6279	665.8179	444.2145
522.3040	261.6559	174.7732	5	L	11	1229.5802	615.2940	410.5320
685.3673	343.1876	229.1277	6	Y	10	1116.4962	558.7520	372.8373
814.4099	407.7089	272.1419	7	E	9	953.4328	477.2203	318.4828
885.4470	443.2274	295.8209	8	A	8	824.3903	412.6990	275.4686
942.4685	471.7382	314.8281	9	G	7	753.3531	377.1805	251.7896
1013.5056	507.2567	338.5071	10	A	6	696.3317	348.6698	232.7824
1100.5376	550.7727	367.5178	11	S	5	625.2946	313.1512	209.1034
1229.5802	615.2940	410.5320	12	E	4	538.2625	269.6352	180.0927
1392.6436	696.8257	464.8864	13	Y	3	409.2199	205.1139	137.0785
1463.6807	732.3443	488.5654	14	A	2	246.1566	123.5822	82.7241
			15	R	1	175.1195	88.0637	59.0451



pNG7248

SYDDAVVDAAVEQIREEVAAHAAER

Peptide prob.	Precursor charge	Calc neutral pep mass	Precursor neutral mass	Mass difference	Xcorr	Deltacn
0.999	4	2642.3	2642.3	0.0077	2.584	0.176



b ⁺	b ²⁺	b ³⁺	#	AA	#	y ⁺	y ²⁺	y ³⁺
88.0399	44.5238	30.0185	1	S	24			
251.1032	126.0555	84.3729	2	Y	23	2556.2330	1278.6204	852.7496
366.1301	183.5690	122.7153	3	D	22	2393.1697	1197.0888	798.3951
481.1571	241.0824	161.0576	4	D	21	2278.1428	1139.5753	760.0528
552.1942	276.6010	184.7366	5	A	20	2163.1158	1082.0618	721.7105
651.2626	326.1352	217.7594	6	Y	19	2092.0787	1046.5433	698.0315
750.3310	375.6694	250.7822	7	Y	18	1993.0103	997.0091	665.0086
865.3580	433.1829	289.1245	8	D	17	1893.9419	947.4749	631.9858
936.3951	468.7014	312.8036	9	A	16	1778.9149	889.9614	593.6435
1007.4322	504.2200	336.4826	10	A	15	1707.8778	854.4428	569.9645
1106.5006	553.7542	369.5054	11	Y	14	1636.8407	818.9243	546.2855
1235.5432	618.2755	412.5196	12	E	13	1537.7723	769.3901	513.2626
1363.6018	682.3048	455.2058	13	Q	12	1408.7297	704.8688	470.2485
1476.6858	738.8468	492.9005	14	I	11	1280.6711	640.8395	427.5623
1632.7869	816.8974	544.9342	15	R	10	1167.5871	584.2974	389.8676
1761.8295	881.4187	587.9484	16	E	9	1011.4859	506.2469	337.8339
1890.8721	945.9400	630.9626	17	E	8	882.4434	441.7256	294.8197
1989.9405	995.4742	663.9854	18	V	7	753.4008	377.2043	251.8055
2060.9777	1030.9927	687.6644	19	A	6	654.3323	327.6701	218.7827
2198.0366	1099.5222	733.3507	20	H	5	583.2952	292.1515	195.1036
2269.0737	1135.0408	757.0298	21	A	4	446.2363	223.6221	149.4173
2340.1108	1170.5593	780.7088	22	A	3	375.1992	188.1035	125.7383
2469.1534	1235.0806	823.7230	23	E	2	304.1621	152.5850	102.0592
			24	R	1	175.1195	88.0637	59.0451