

Info

Modified Sequence AcD3-AAFTK<AcD3K*>NQK<AcD3K*>DPGVLDK-COOH

Ion score 55.32

Protein P50543

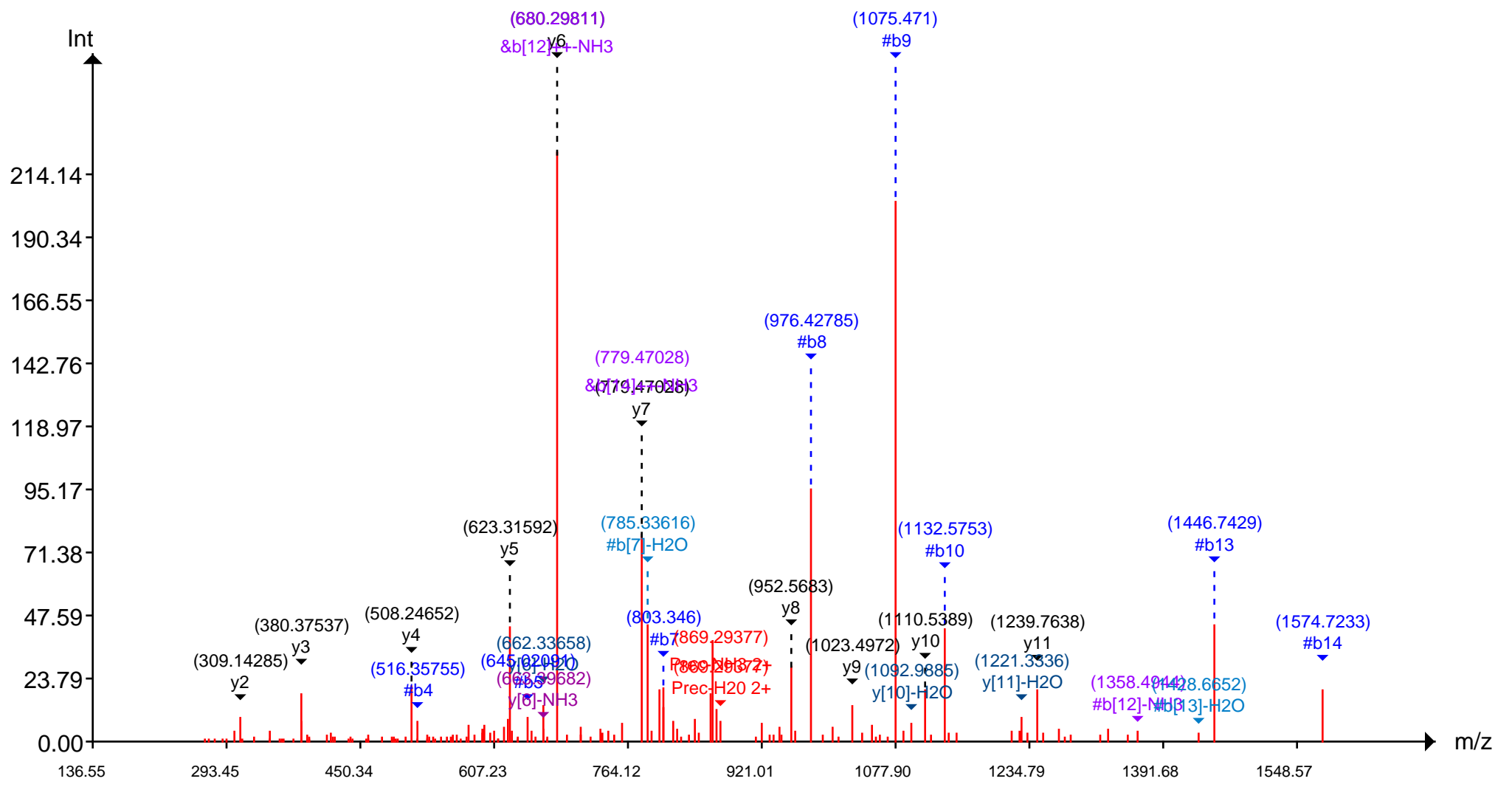
Confidence level 99.00%

Precursor m/z 897.98991

Identity Threshold 47.57

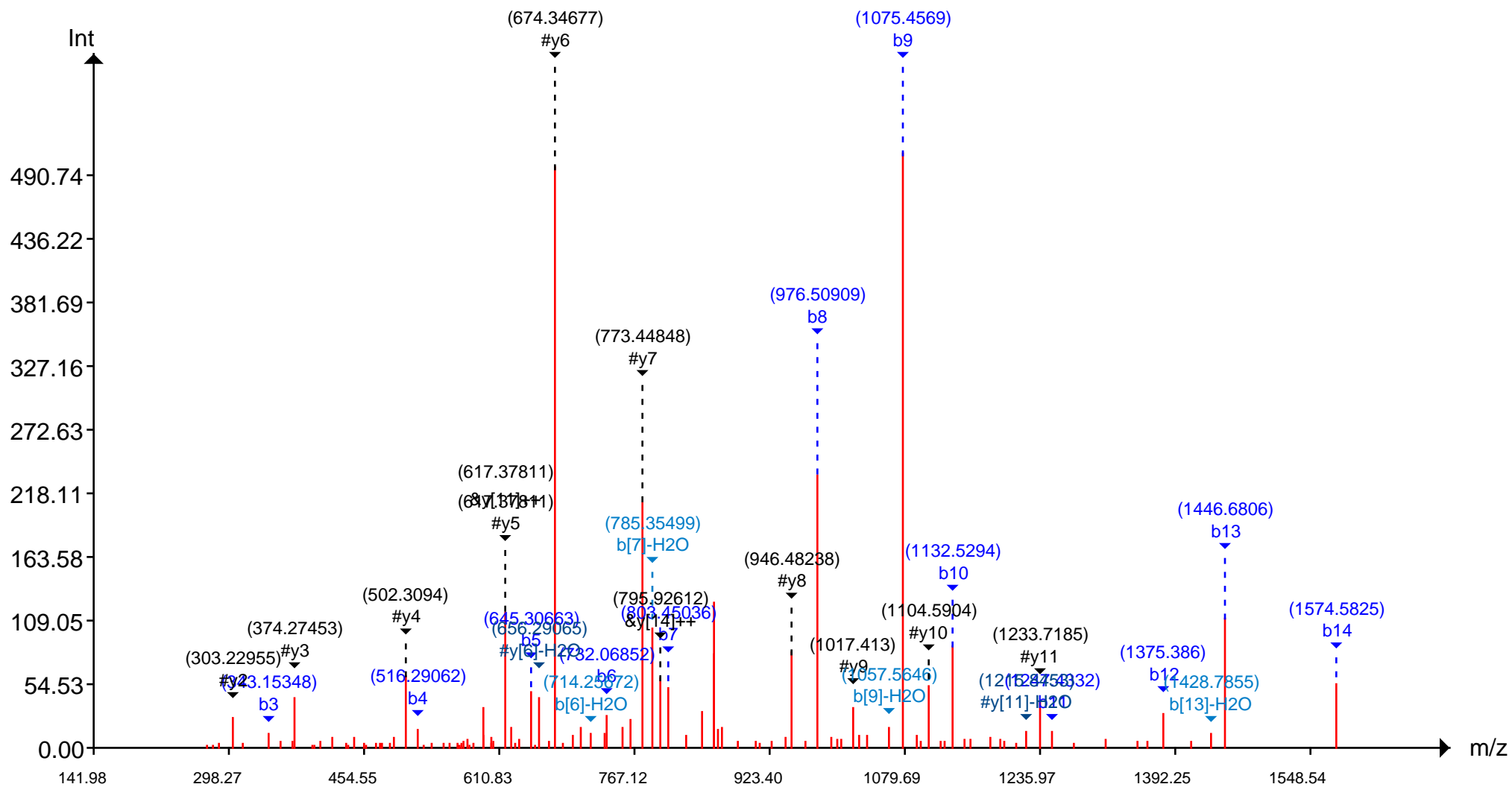
Charge 2+

Mass error 0.003064



Info

Modified Sequence	AcD3-IAIK<AcD3K*>ESAK<AcD3K*>VGDQAQR<C13>-COOH	Ion score	81.8
Protein	Q99JY9	Confidence level	99.00%
Precursor m/z	878.00749	Identity Threshold	45.12
Charge	2+	Mass error	0.002457



Info

Modified Sequence AcD3-IAIK<AcD3K*>ESAK<AcD3K*>VGDQAQR-COOH

Ion score 103.98

Protein Q99JY9

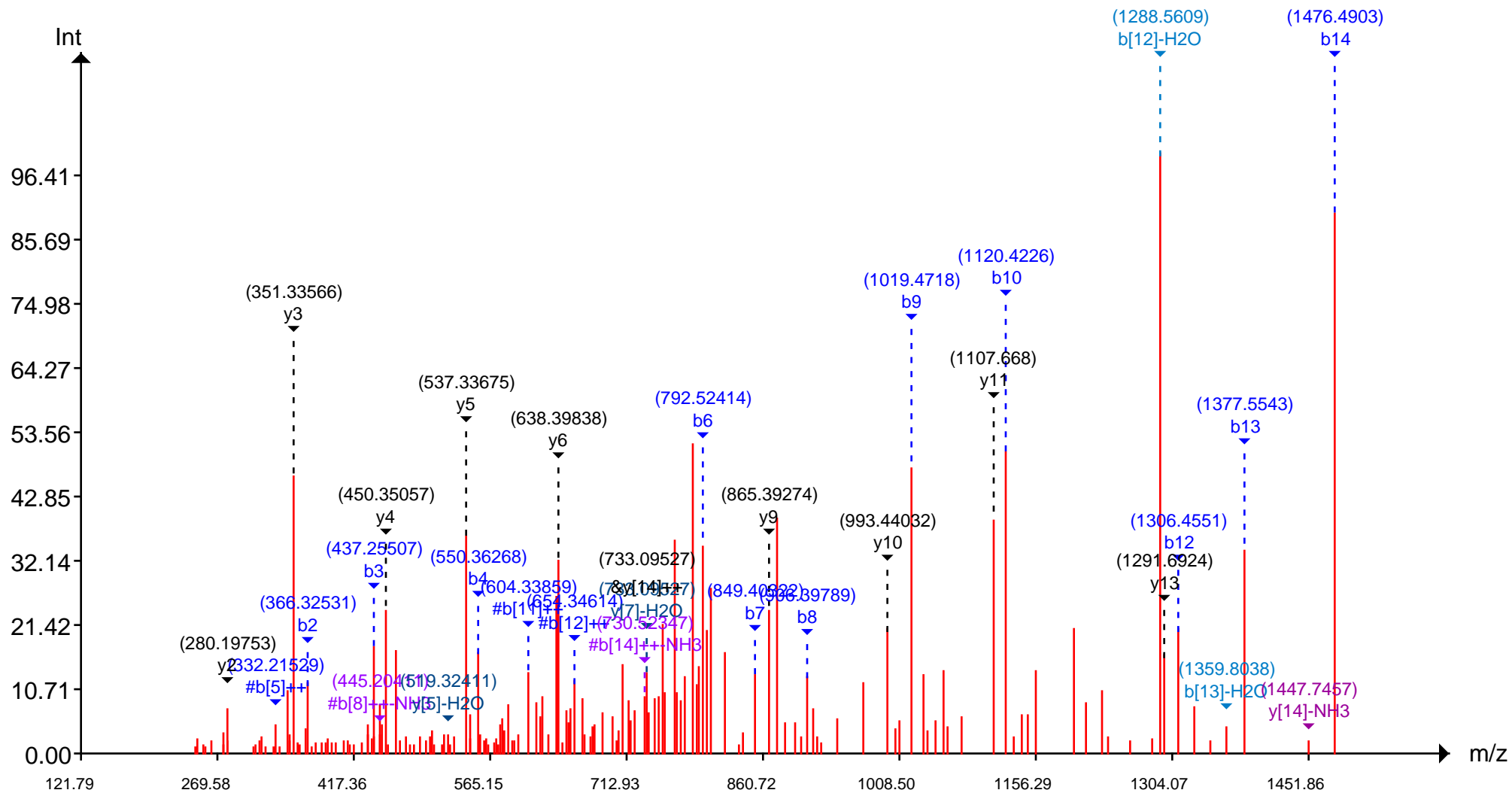
Confidence level 99.00%

Precursor m/z 874.99705

Identity Threshold 46.20

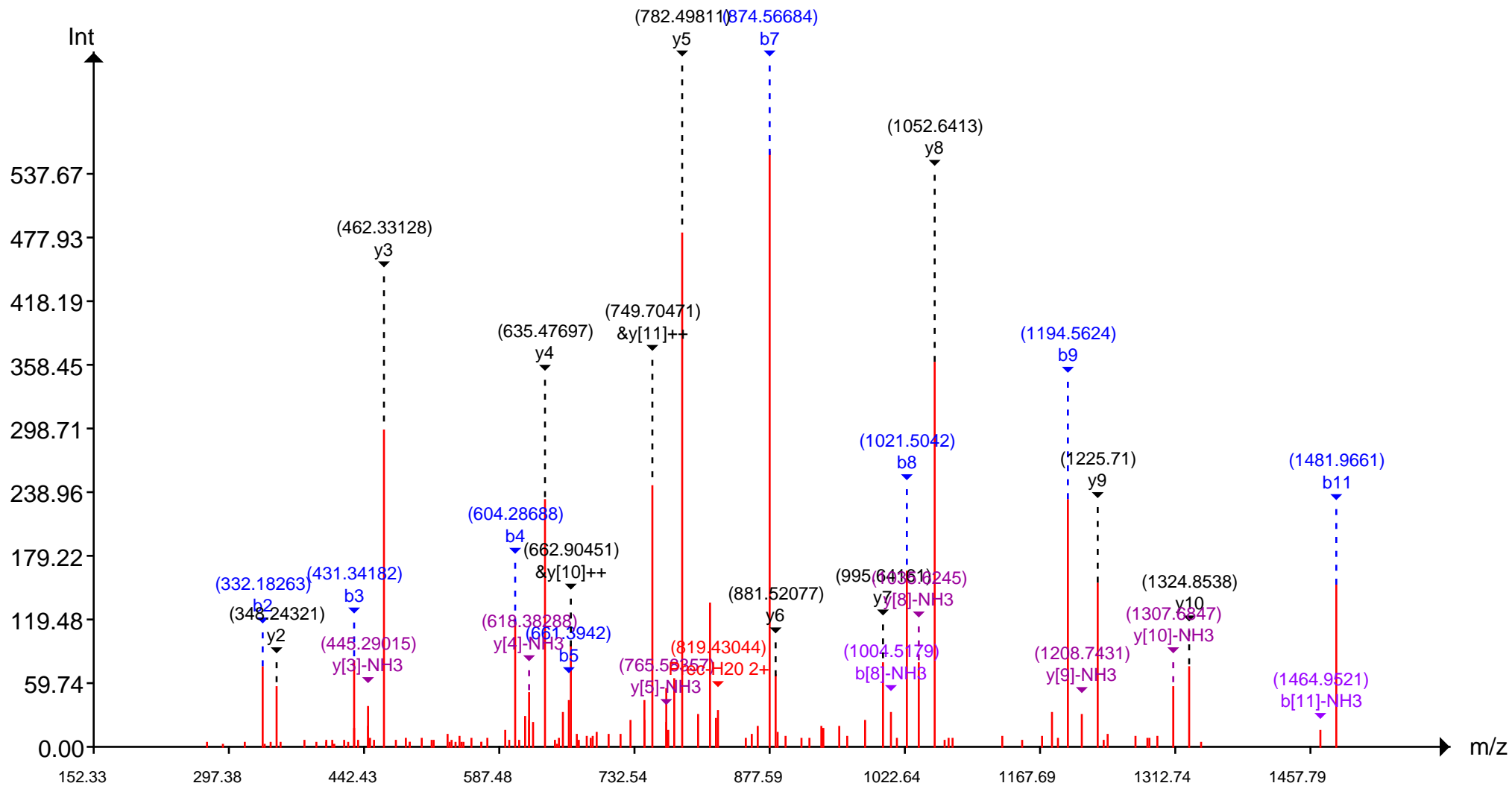
Charge 2+

Mass error 0.001719



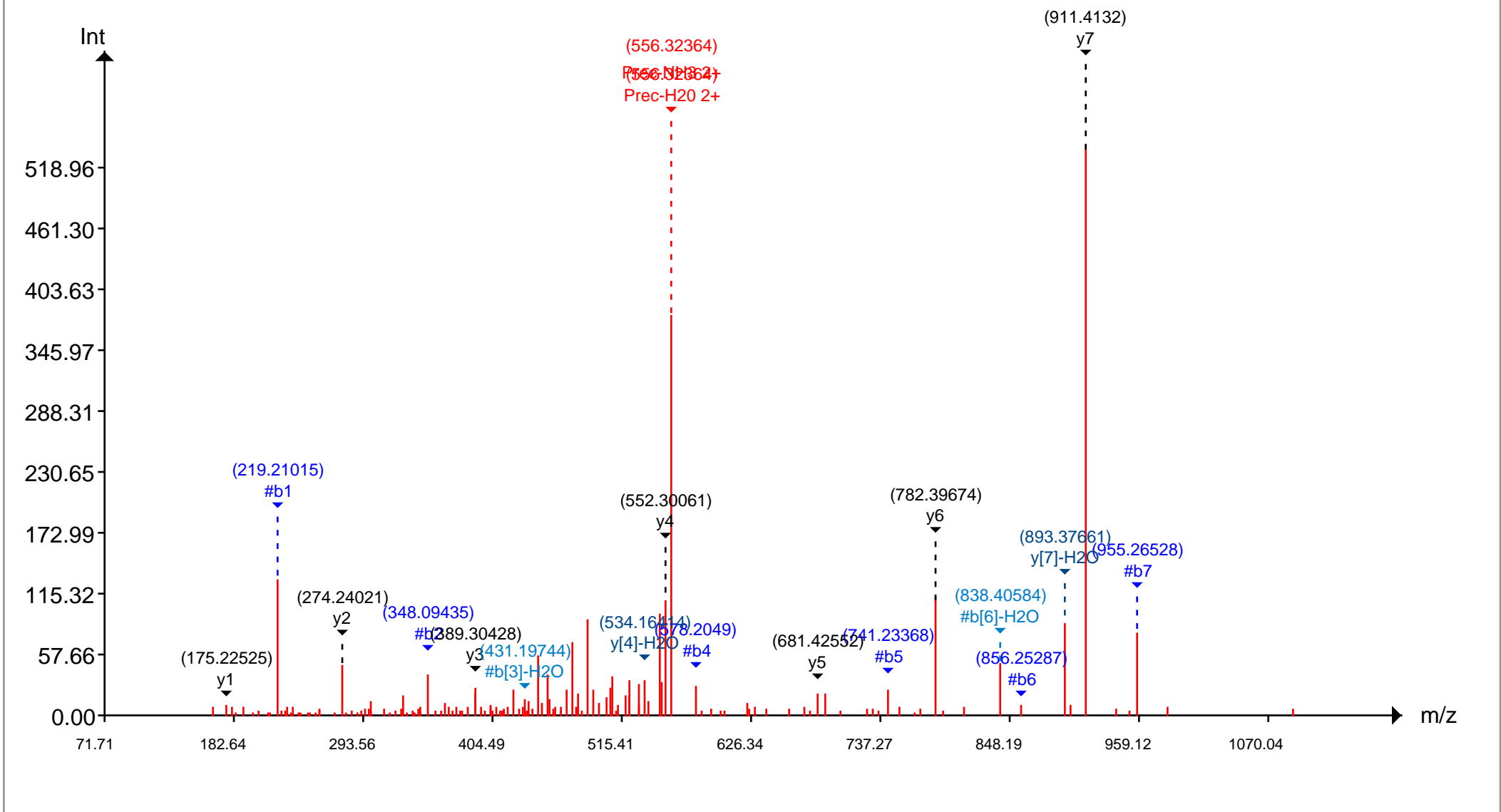
Info

Modified Sequence	AcD3-FK<AcD3K*>AINQGGLTSVAVR<C13>-COOH	Ion score	79.68
Protein	Q9QUM9	Confidence level	99.00%
Precursor m/z	828.98633	Identity Threshold	44.65
Charge	2+	Mass error	8.02E-4



Info

Modified Sequence	AcD3-LK<AcD3K*>VK<AcD3K*>GNVFK<AcD3K*>NK<AcD3K*>R-COOH	Ion score	65.15
Protein	P84099	Confidence level	99.00%
Precursor m/z	828.52588	Identity Threshold	37.11
Charge	2+	Mass error	0.002082



Info

Modified Sequence AcD3-K<AcD3K*>ETEYDVR-COOH

Ion score 49.46

Protein Q61176

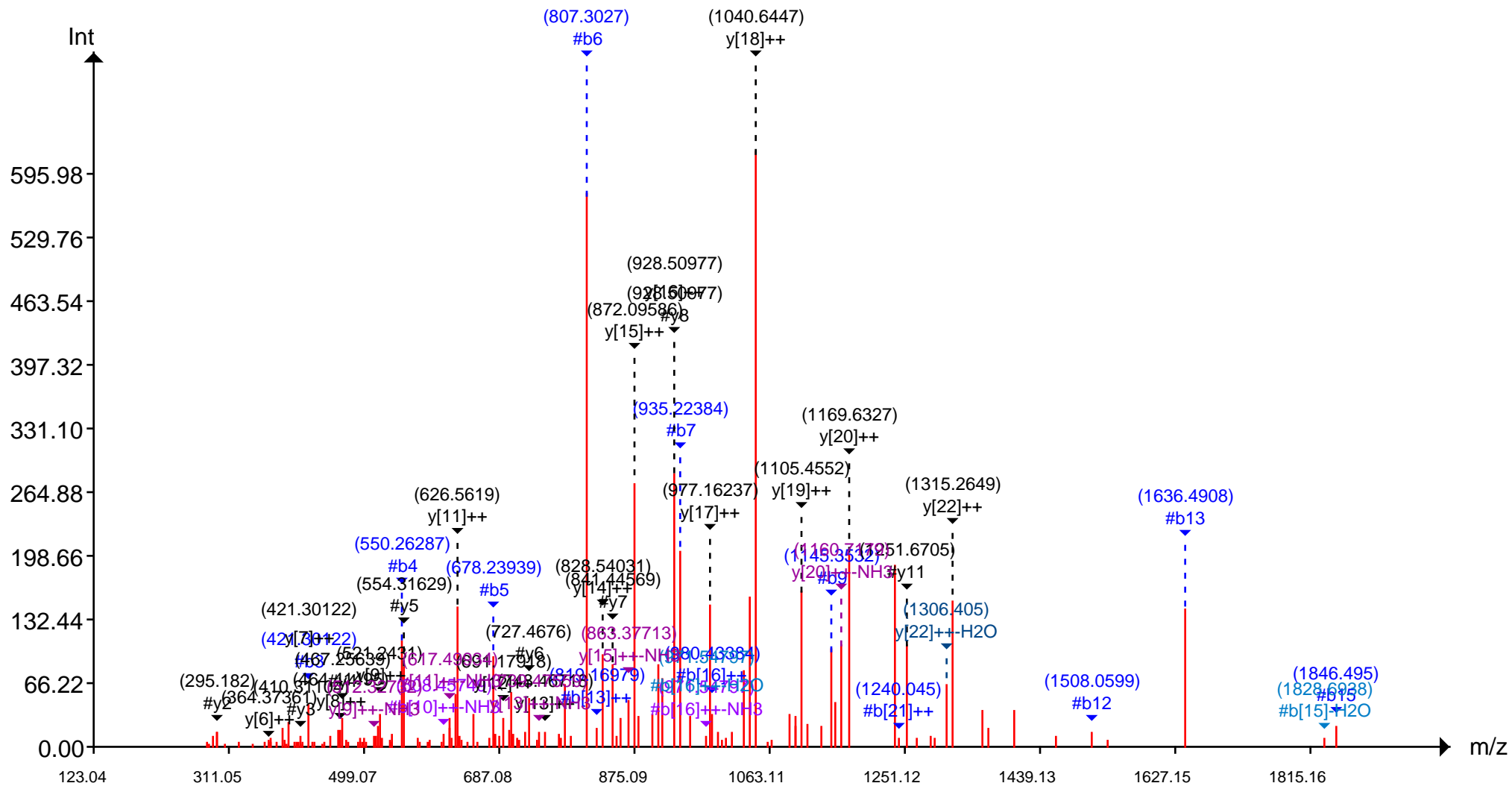
Confidence level 99.00%

Precursor m/z 565.28596

Identity Threshold 47.45

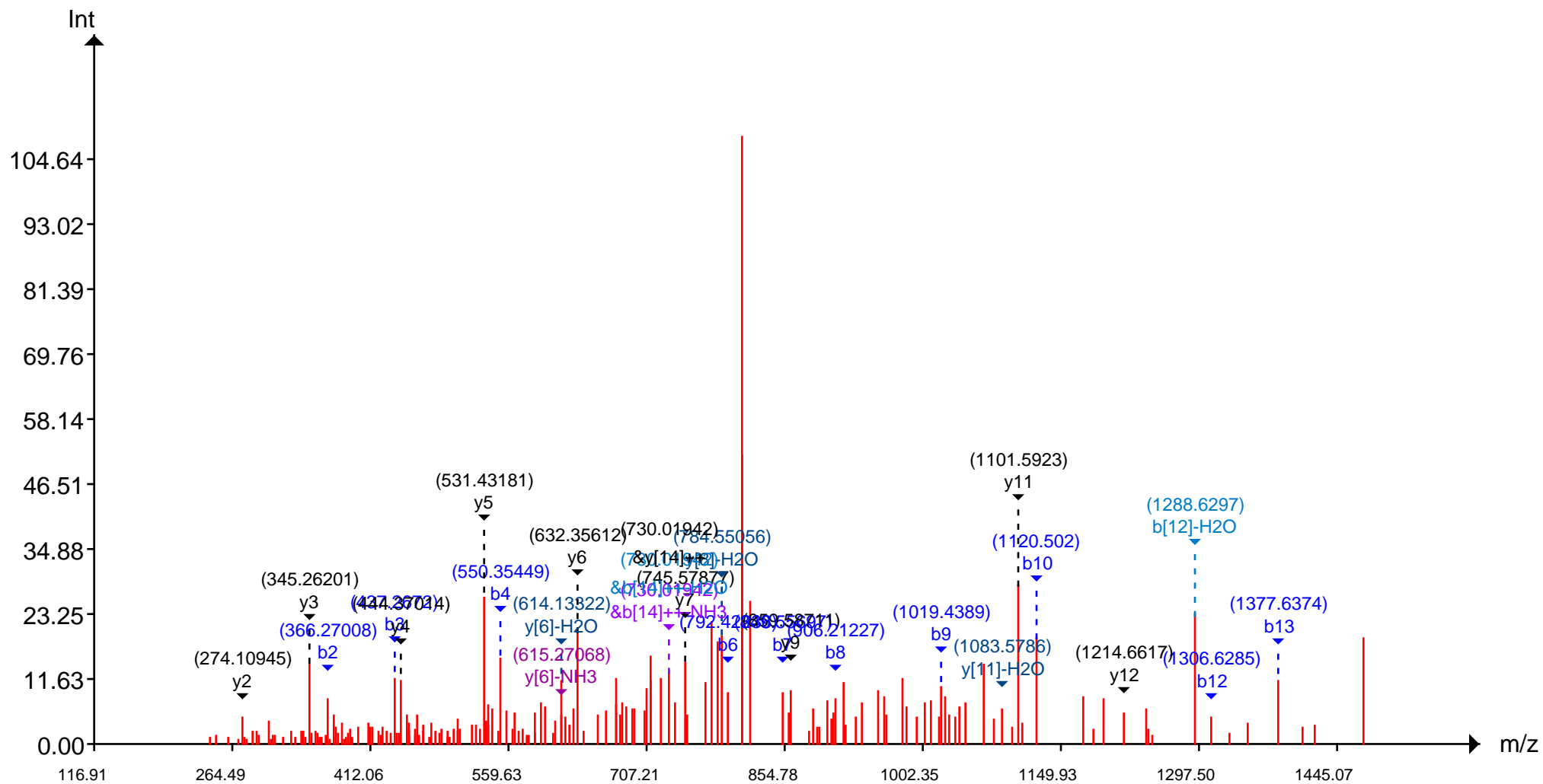
Charge 2+

Mass error 4.07E-4



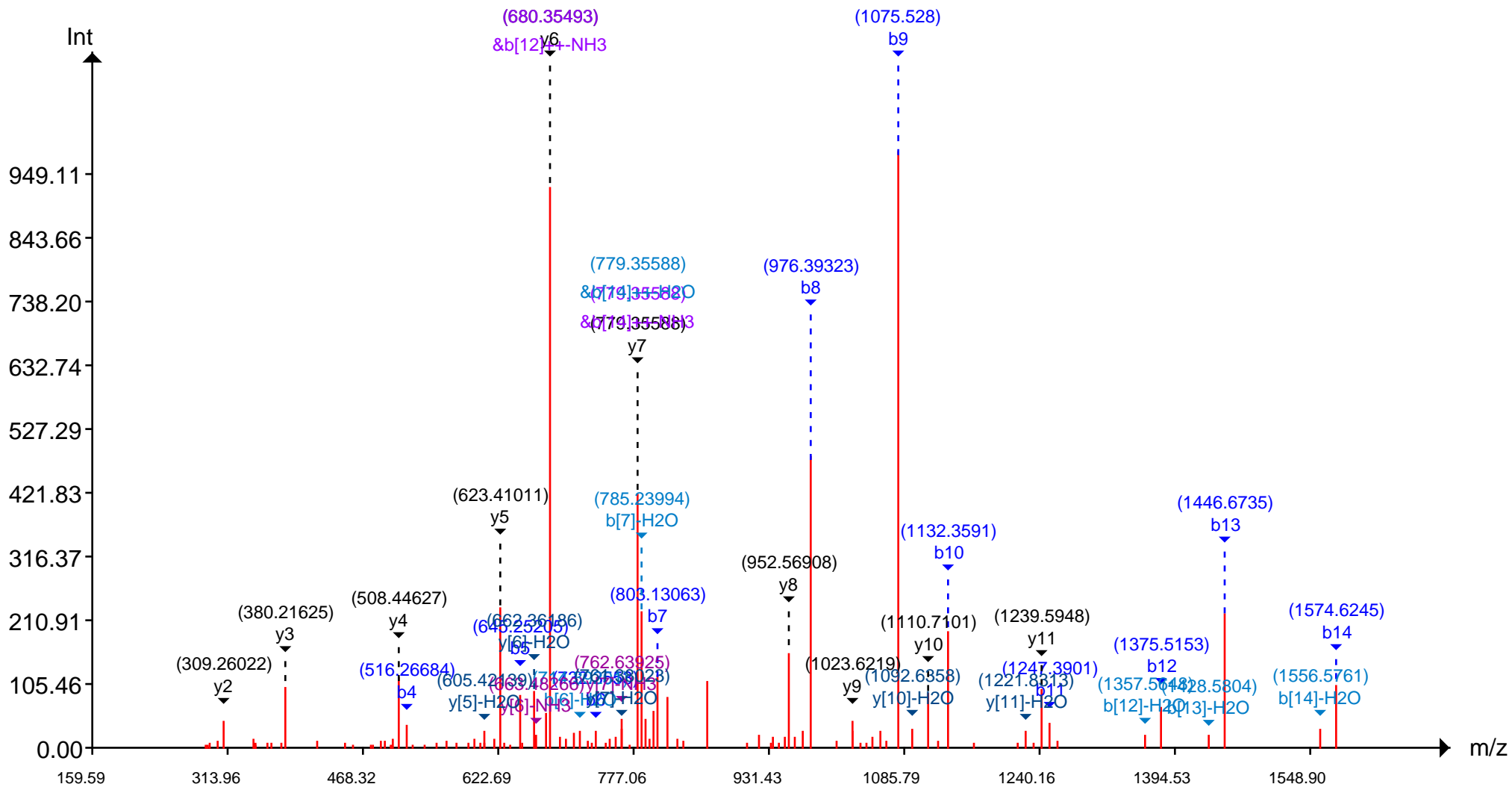
Info

Modified Sequence	AcD3-LVYEQEQPLSC<Cmm*>DEPILSNK<AcD3K*>SGDNR<C13>-COOH	Ion score	79.24
Protein	P70296	Confidence level	99.00%
Precursor m/z	963.13895	Identity Threshold	46.88
Charge	3+	Mass error	0.003085



Info

Modified Sequence	AcD3-FK<AcD3K*>AINQGGLTSVAVR-COOH	Ion score	46.14
Protein	Q9QUM9	Confidence level	99.00%
Precursor m/z	825.97616	Identity Threshold	45.56
Charge	2+	Mass error	6.04E-4



Info

Modified Sequence AcD3-IAIK<AcD3K*>ESAK<AcD3K*>VGDQAQR<C13>-COOH

Ion score 103.9

Protein Q99JY9

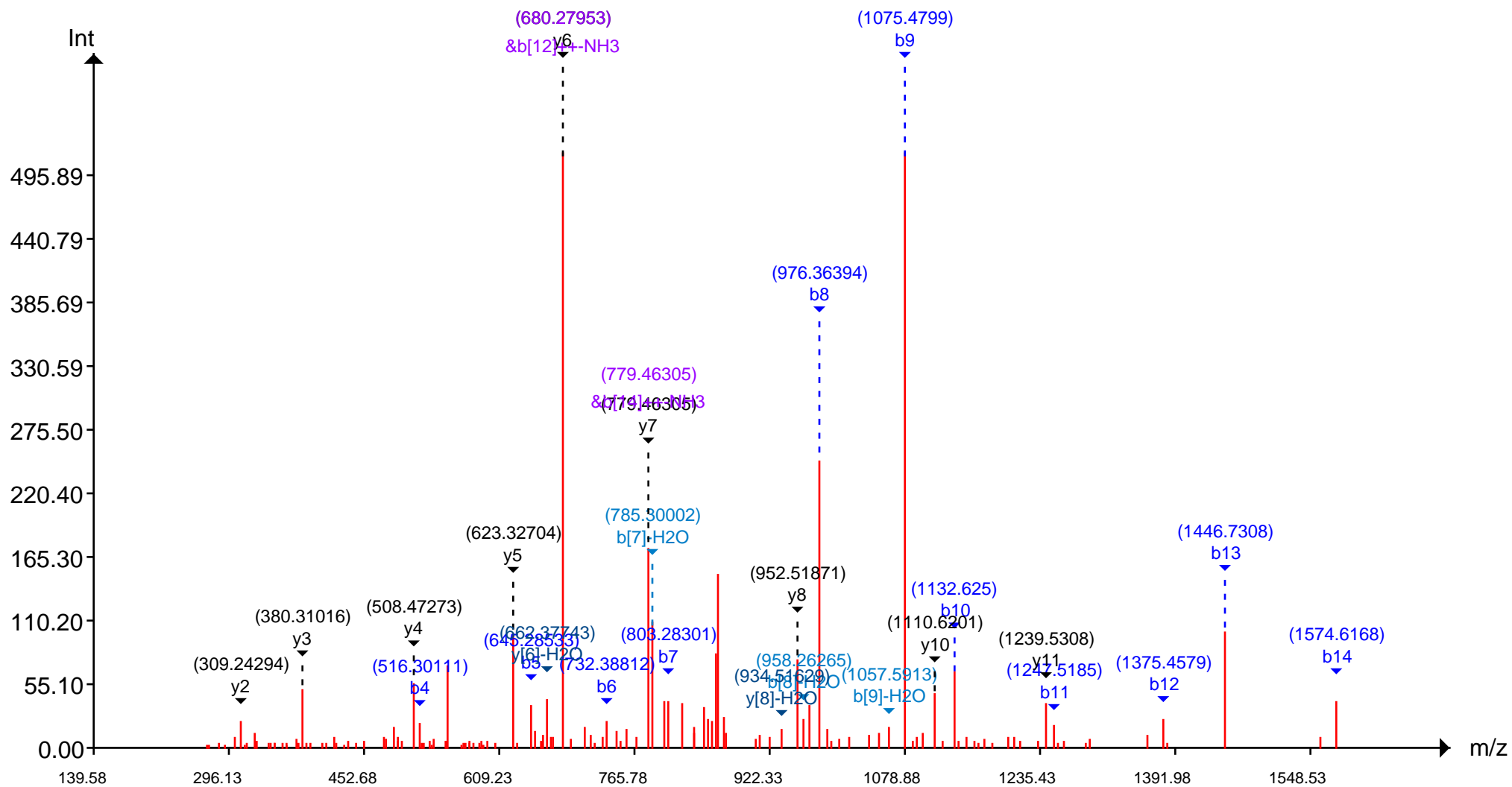
Confidence level 99.00%

Precursor m/z 878.00745

Identity Threshold 45.14

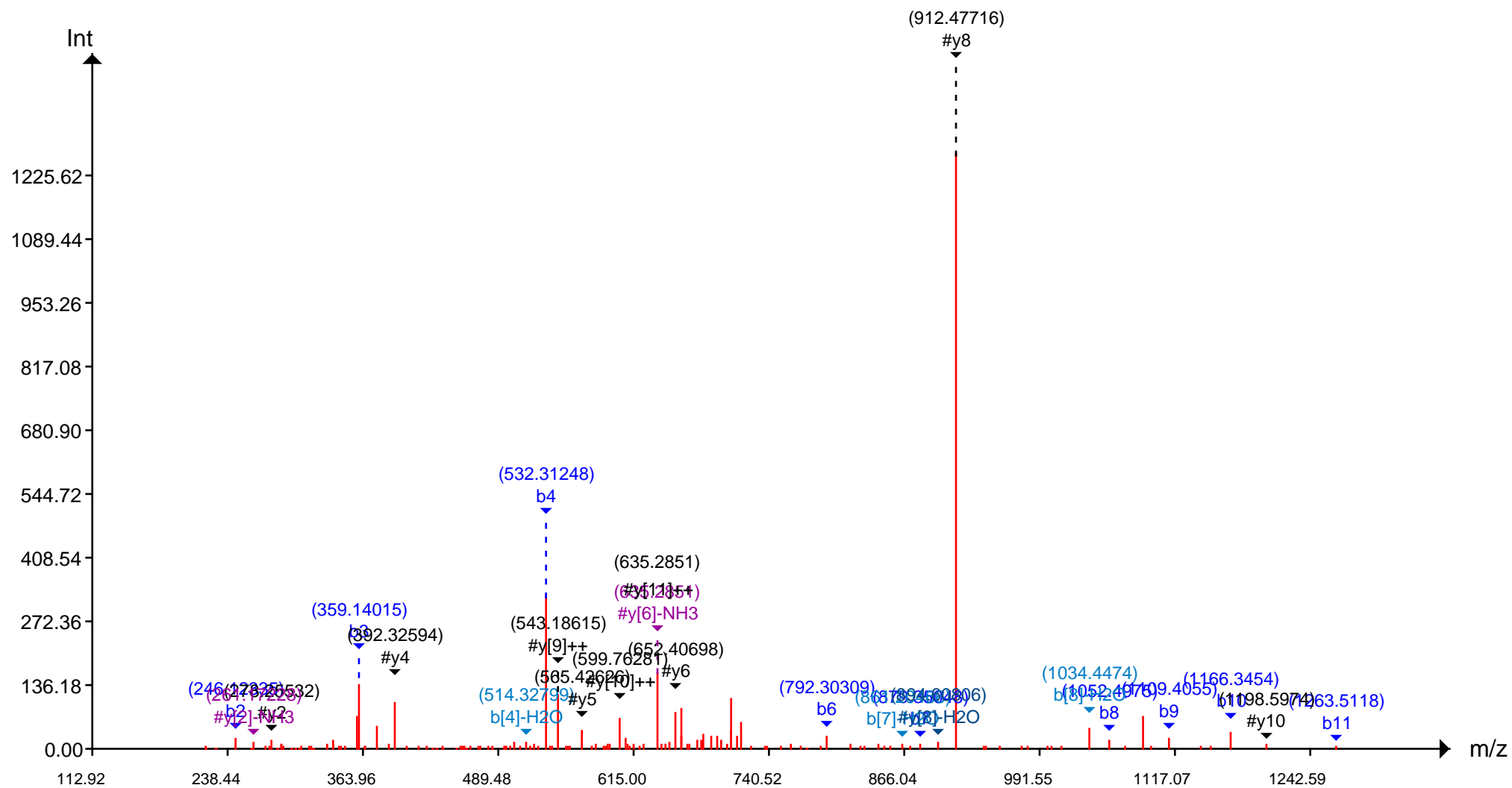
Charge 2+

Mass error 0.002377



Info

Modified Sequence	AcD3-IAIK<AcD3K*>ESAK<AcD3K*>VGDQAQR<C13>-COOH	Ion score	72.66
Protein	Q99JY9	Confidence level	99.00%
Precursor m/z	878.0076	Identity Threshold	45.10
Charge	2+	Mass error	0.002677



Info

Modified Sequence AcD3-EALK<AcD3K*>PYSK<AcD3K*>GGPR<C13>-COOH

Ion score 59.1

Protein P06745

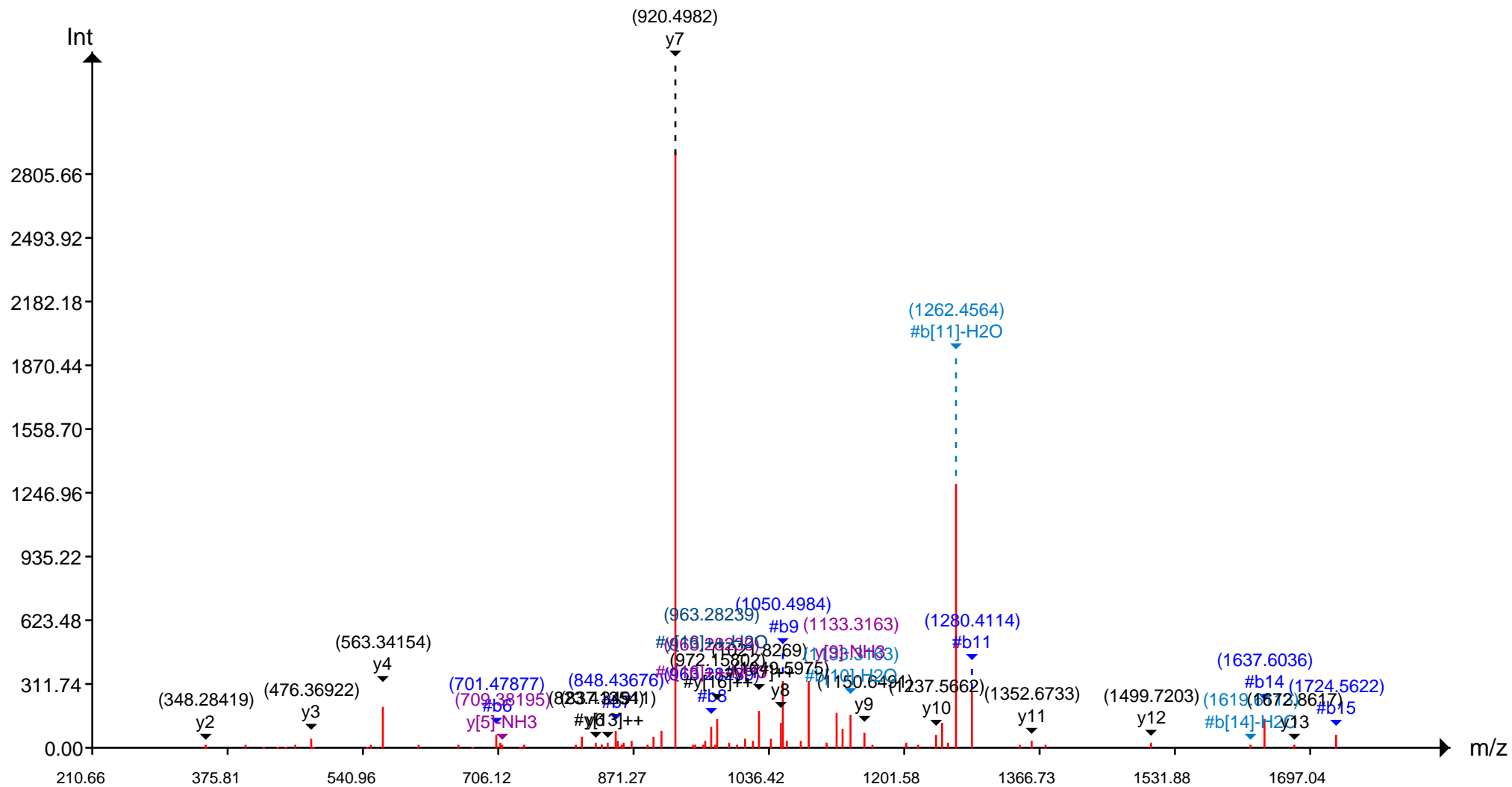
Confidence level 99.00%

Precursor m/z 722.41652

Identity Threshold 46.07

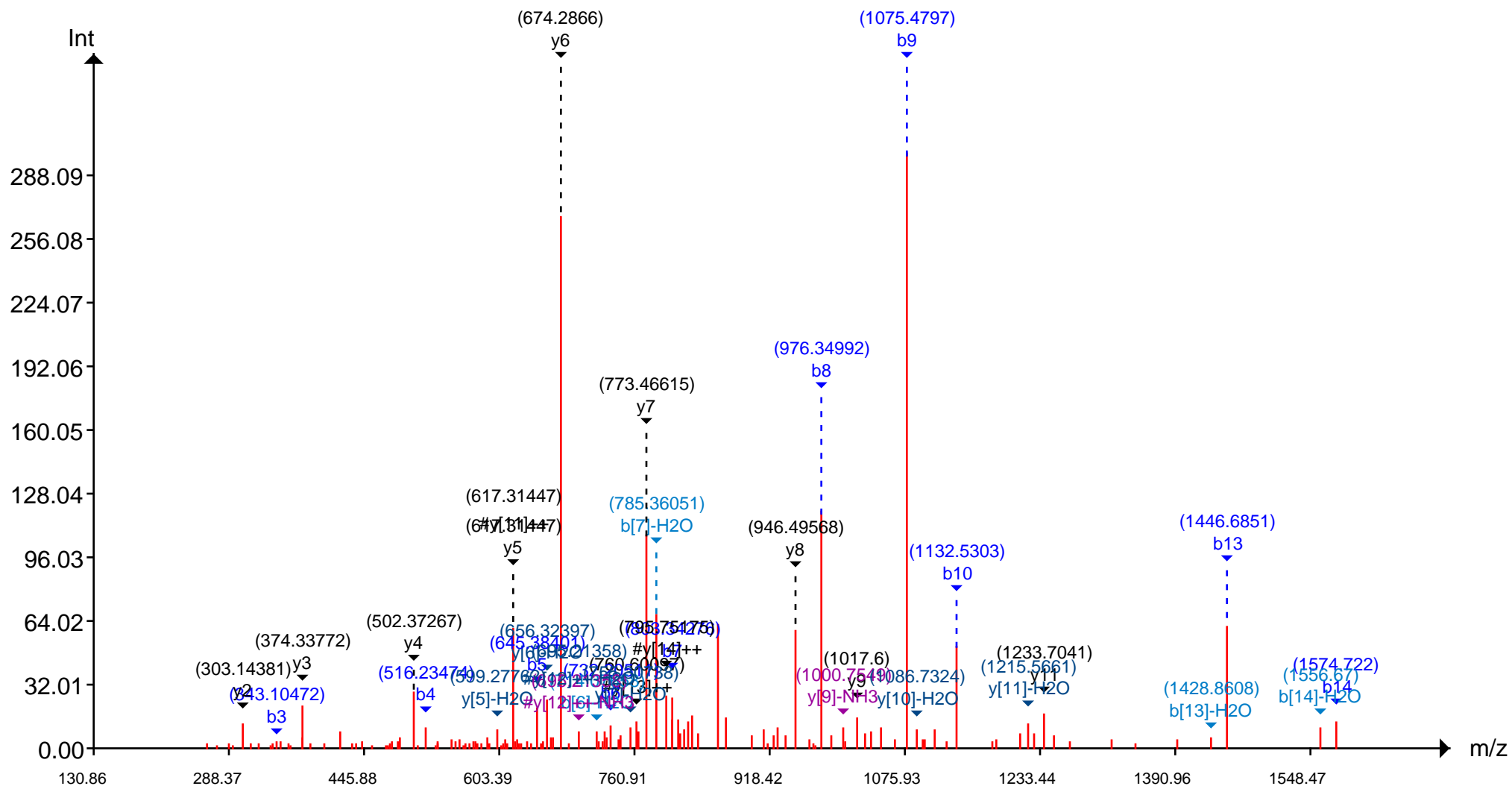
Charge 2+

Mass error 9.98E-4



Info

Modified Sequence	AcD3-IVGVNK<AcD3K*>M<Mox*>DSTEPPYSQK<AcD3K*>R-COOH	Ion score	71.83
Protein	P10126	Confidence level	99.00%
Precursor m/z	1100.5685	Identity Threshold	47.97
Charge	2+	Mass error	0.003277



Info

Modified Sequence AcD3-IAIK<AcD3K*>ESAK<AcD3K*>VGDQAQR-COOH

Ion score 83.3

Protein Q99JY9

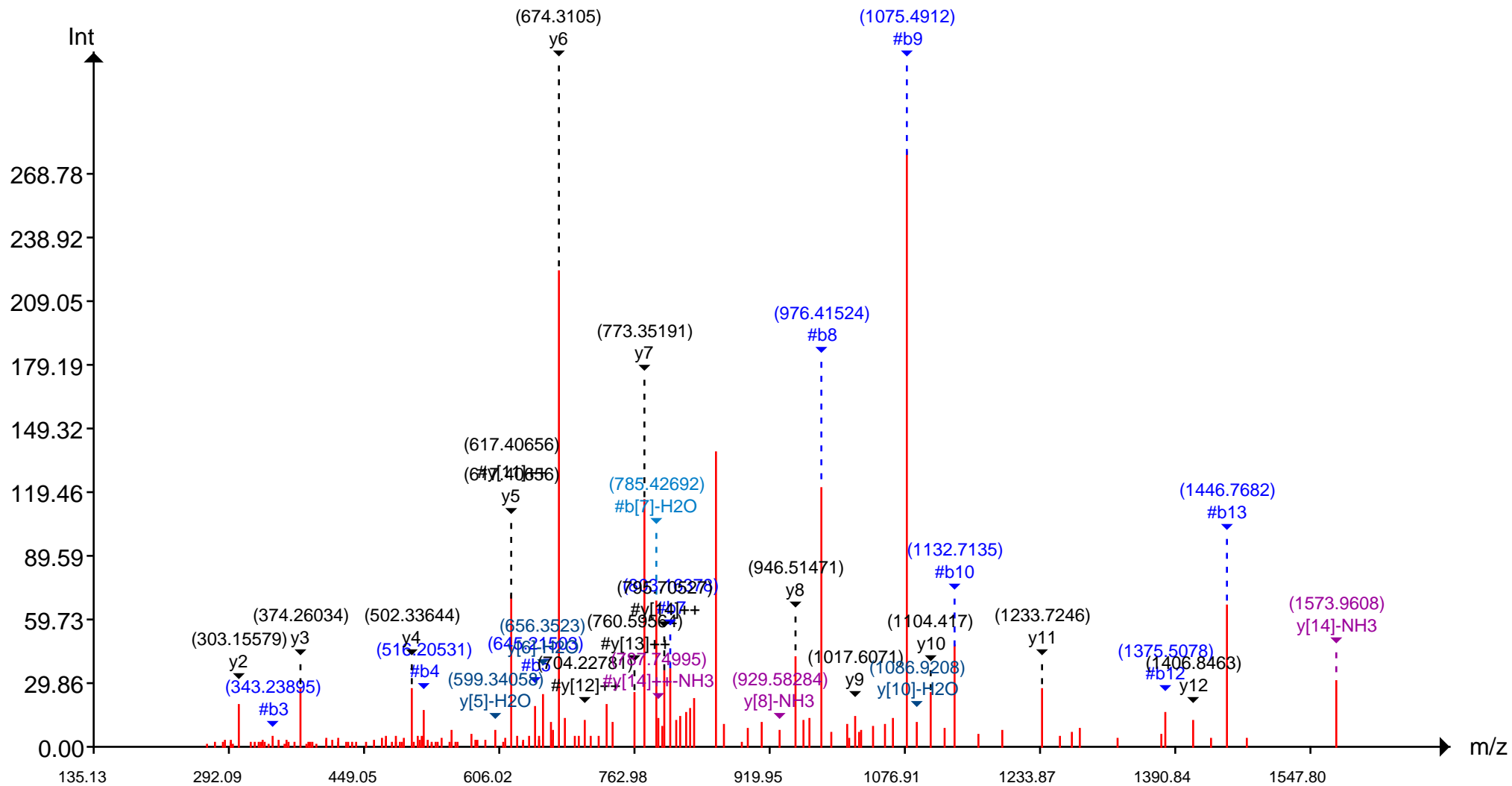
Confidence level 99.00%

Precursor m/z 874.99746

Identity Threshold 46.10

Charge 2+

Mass error 0.002539



Info

Modified Sequence AcD3-IAIK<AcD3K*>ESAK<AcD3K*>VGDQAQR-COOH

Ion score 80.63

Protein Q99JY9

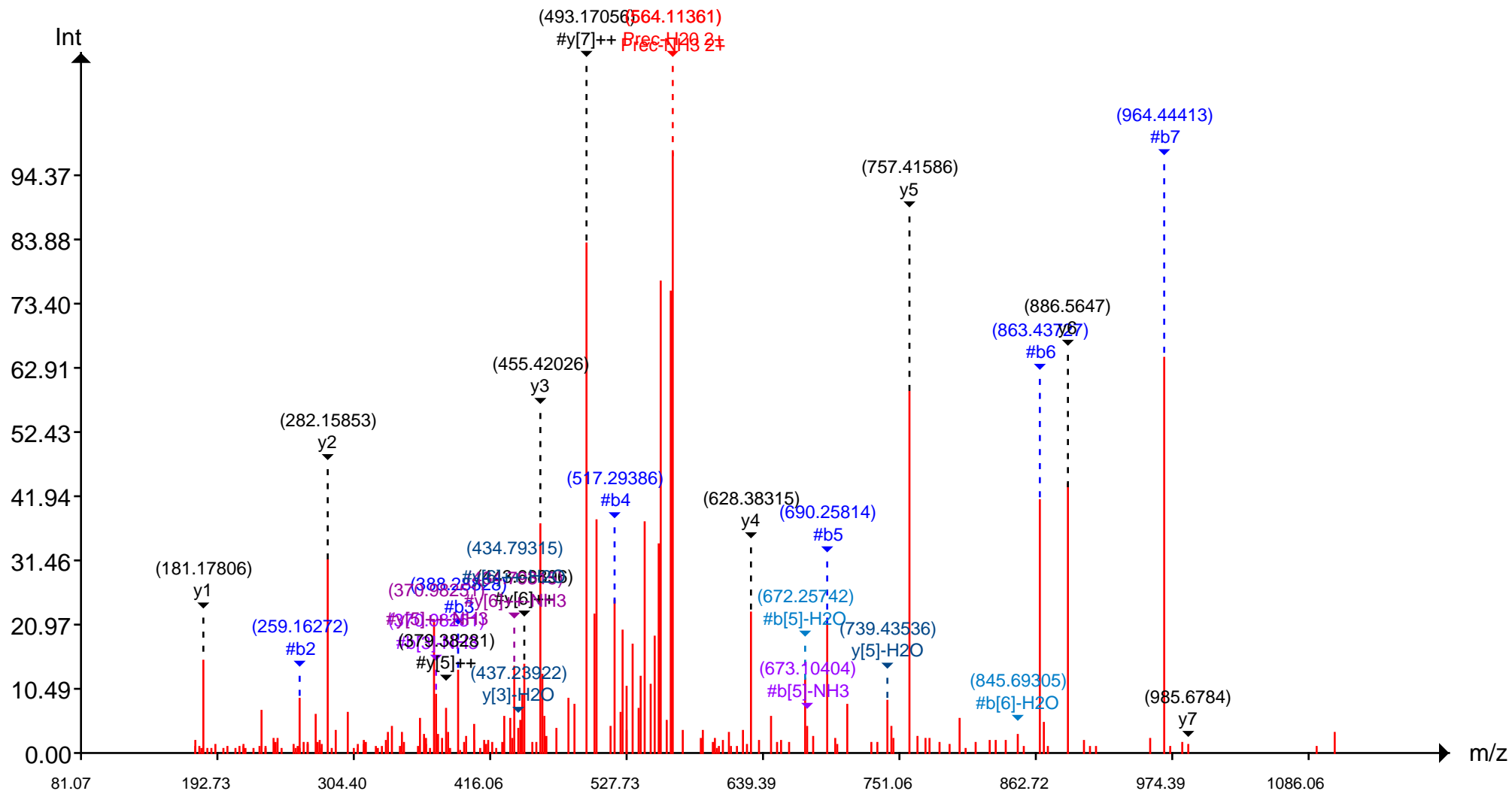
Confidence level 99.00%

Precursor m/z 874.9976

Identity Threshold 46.08

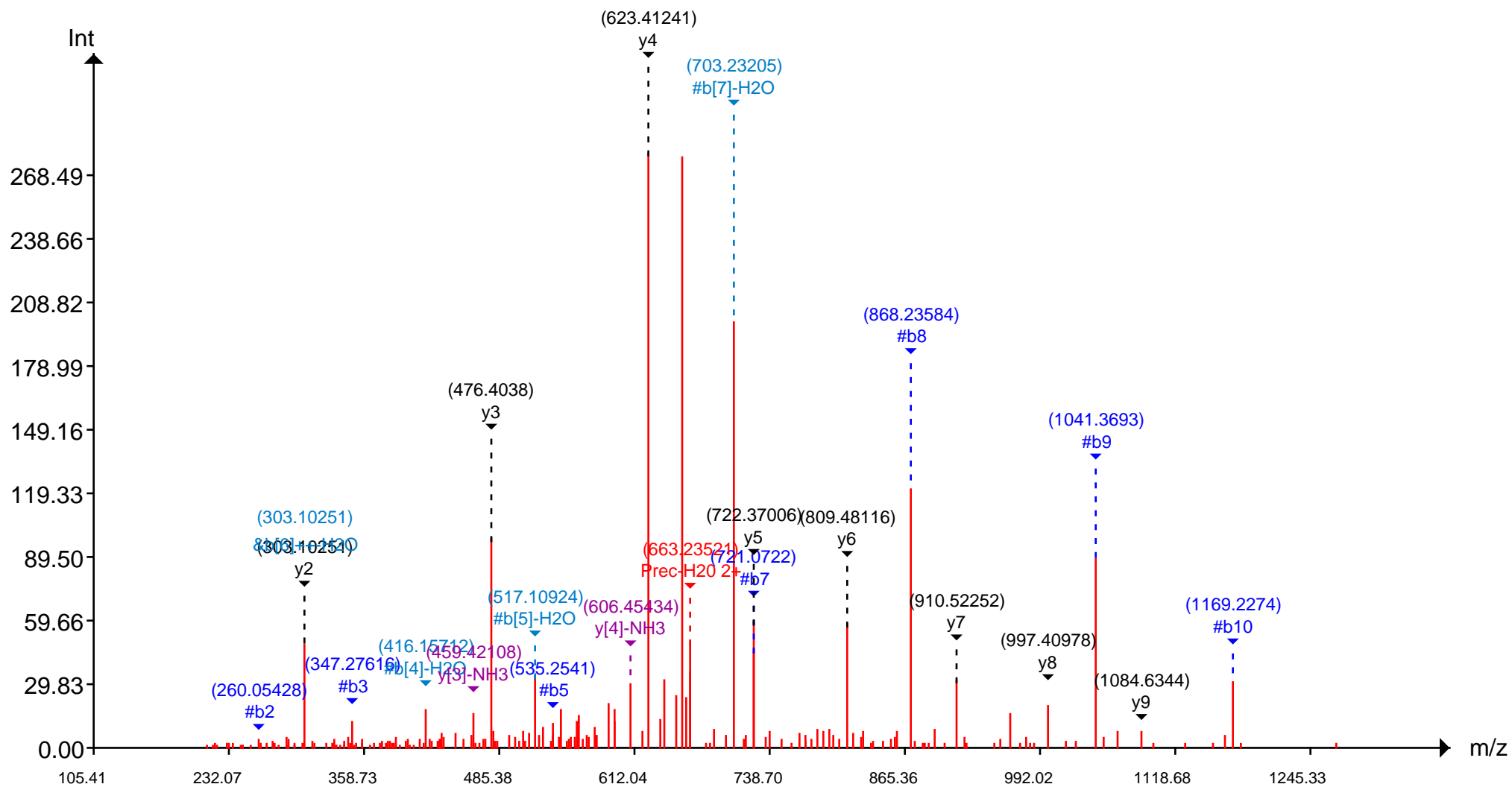
Charge 2+

Mass error 0.002819



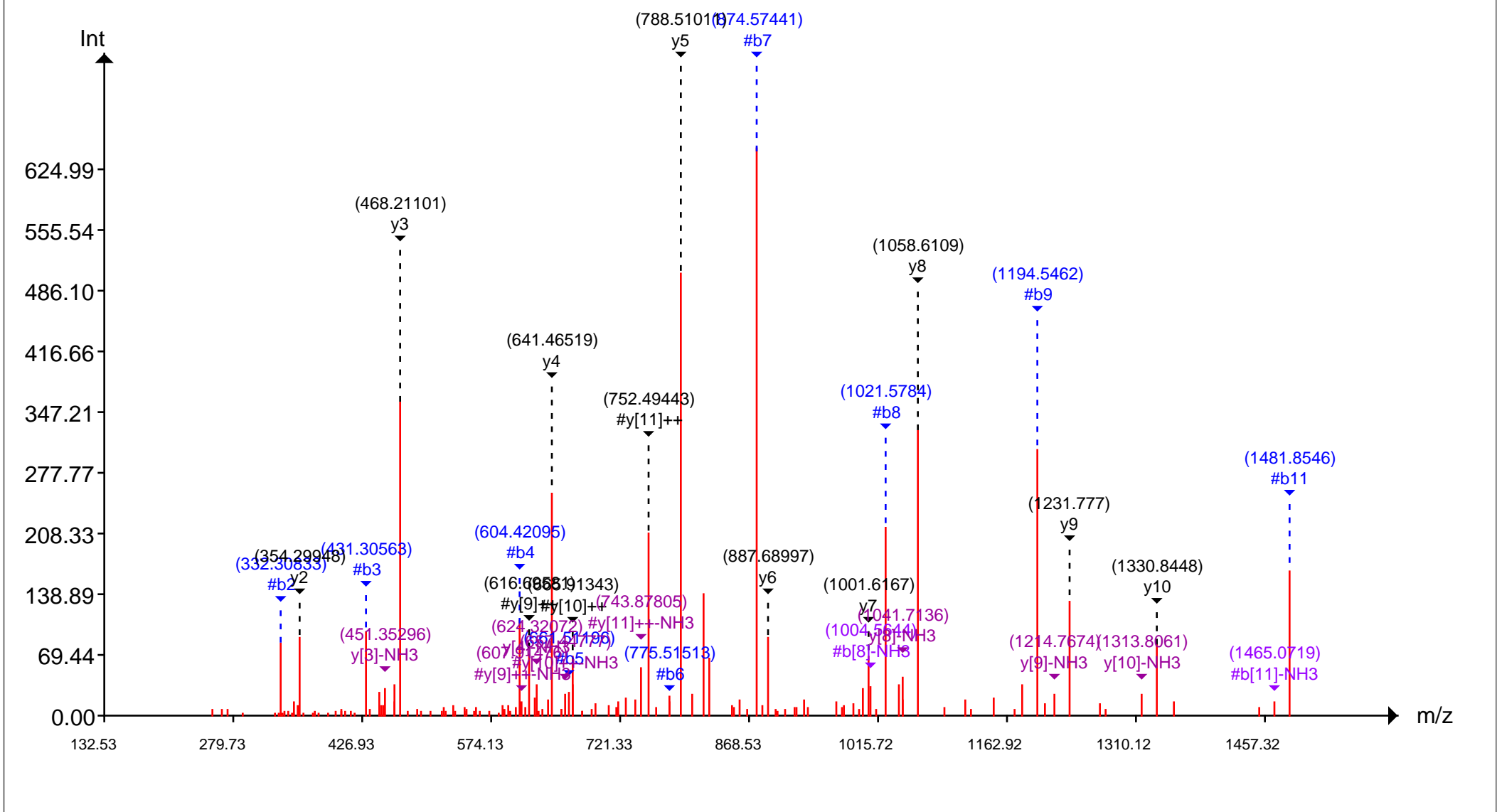
Info

Modified Sequence	AcD3-NVEEK<AcD3K*>K<AcD3K*>TR<C13>-COOH	Ion score	54.6
Protein	P97855, P97379	Confidence level	99.00%
Precursor m/z	572.83455	Identity Threshold	46.70
Charge	2+	Mass error	4.34E-4



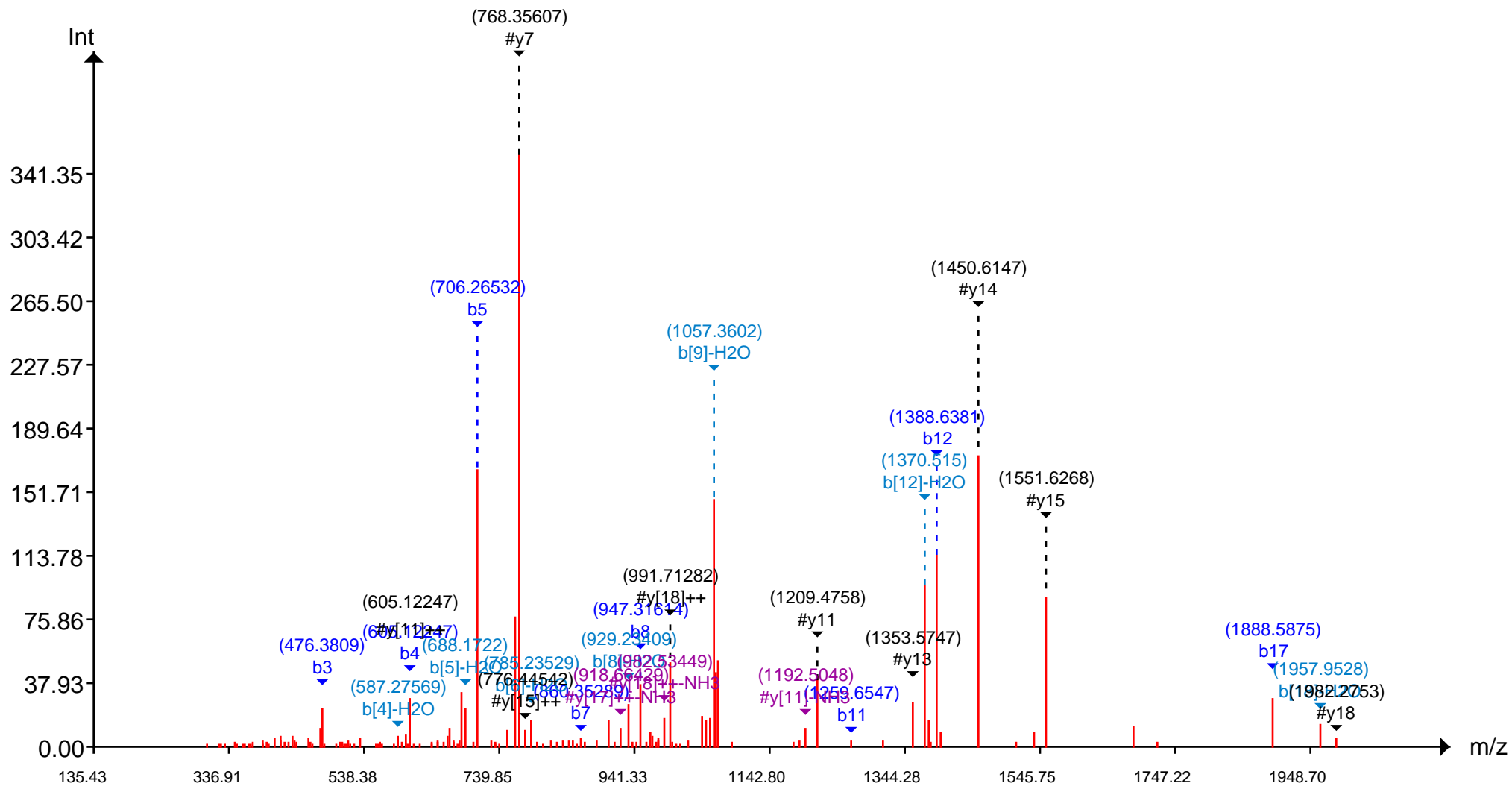
Info

Modified Sequence	AcD3-DVSSTSVFK<AcD3K*>QR-COOH	Ion score	56.71
Protein	Q8R1F1	Confidence level	99.00%
Precursor m/z	672.35791	Identity Threshold	48.09
Charge	2+	Mass error	0.001302



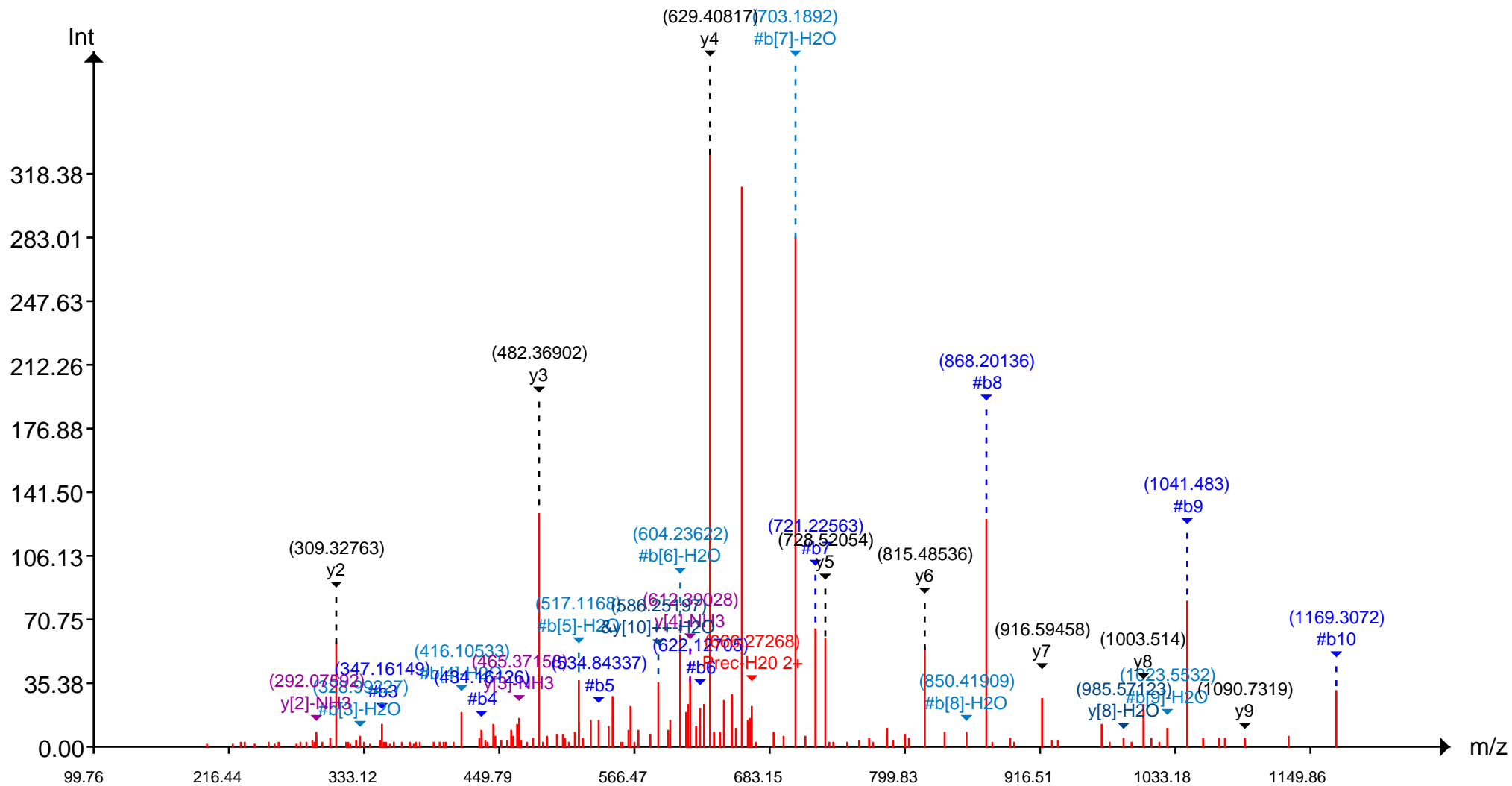
Info

Modified Sequence	AcD3-LK<AcD3K*>VK<AcD3K*>GNVFK<AcD3K*>NK<AcD3K*>R<C13>-COOH	Ion score	83.71
Protein	P84099	Confidence level	99.00%
Precursor m/z	831.53565	Identity Threshold	34.88
Charge	2+	Mass error	0.001481



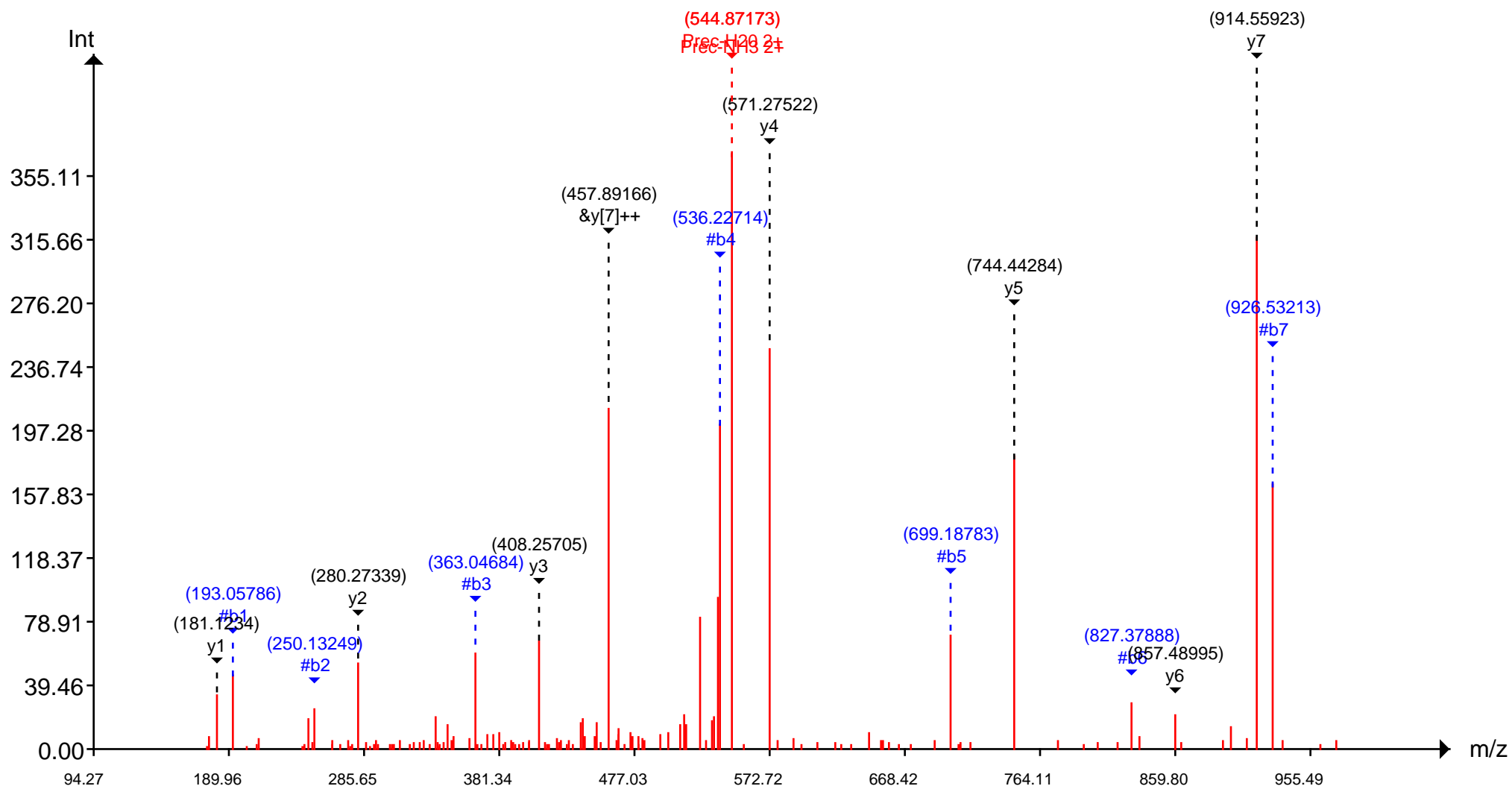
Info

Modified Sequence	AcD3-EQK<AcD3K*>ETPGSQPSEPC<Cmm*>SGVSR<C13>-COOH	Ion score	51.38
Protein	P27546	Confidence level	99.00%
Precursor m/z	1078.5121	Identity Threshold	46.15
Charge	2+	Mass error	0.003071



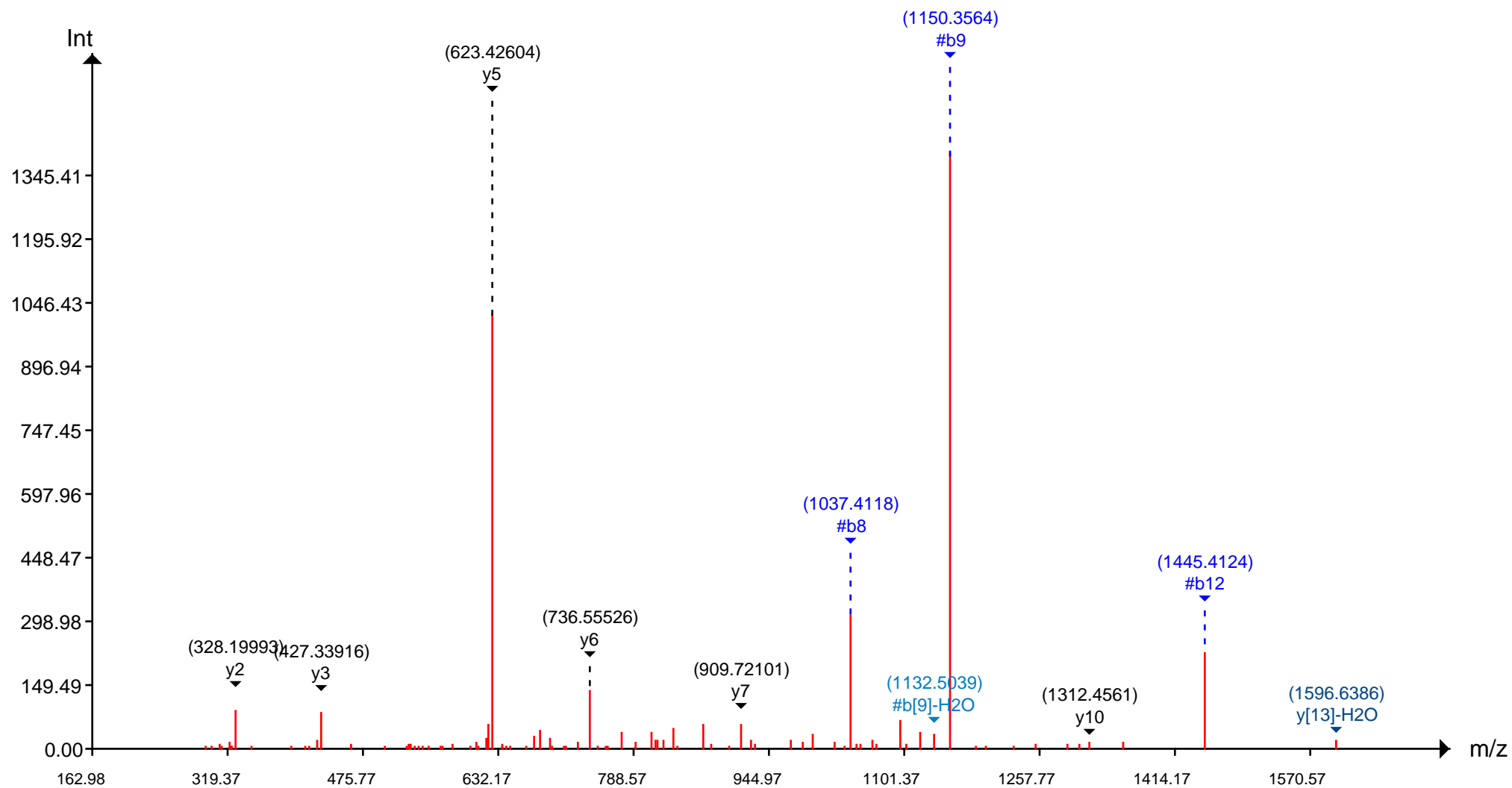
Info

Modified Sequence	AcD3-DVSSTSVFK<AcD3K*>QR<C13>-COOH	Ion score	57.14
Protein	Q8R1F1	Confidence level	99.00%
Precursor m/z	675.36802	Identity Threshold	47.97
Charge	2+	Mass error	0.00138



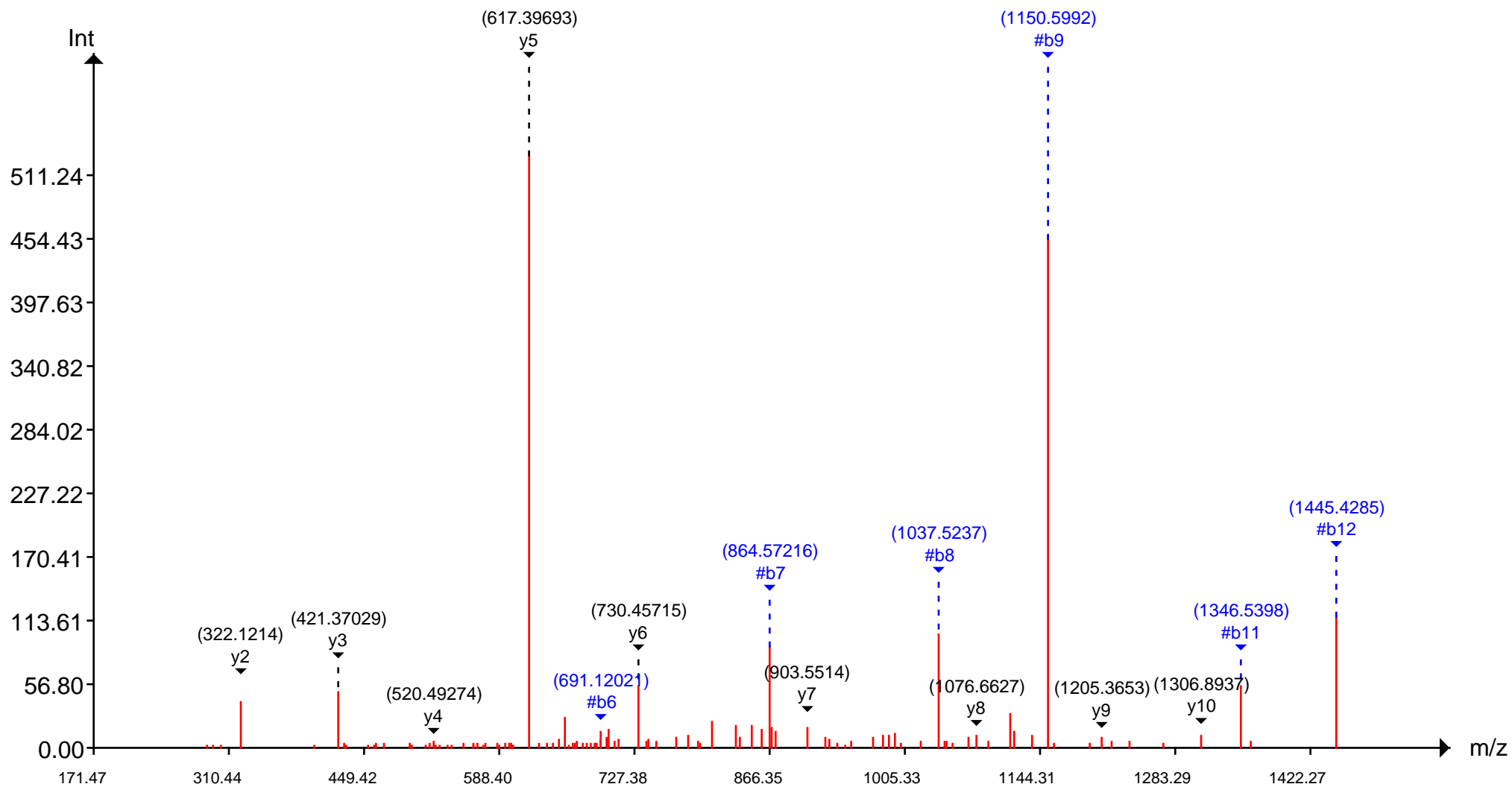
Info

Modified Sequence	AcD3-M<Mox*>GLK<AcD3K*>YQVR<C13>-COOH	Ion score	62.55
Protein	P70677	Confidence level	99.00%
Precursor m/z	553.81551	Identity Threshold	47.70
Charge	2+	Mass error	-3.38E-4



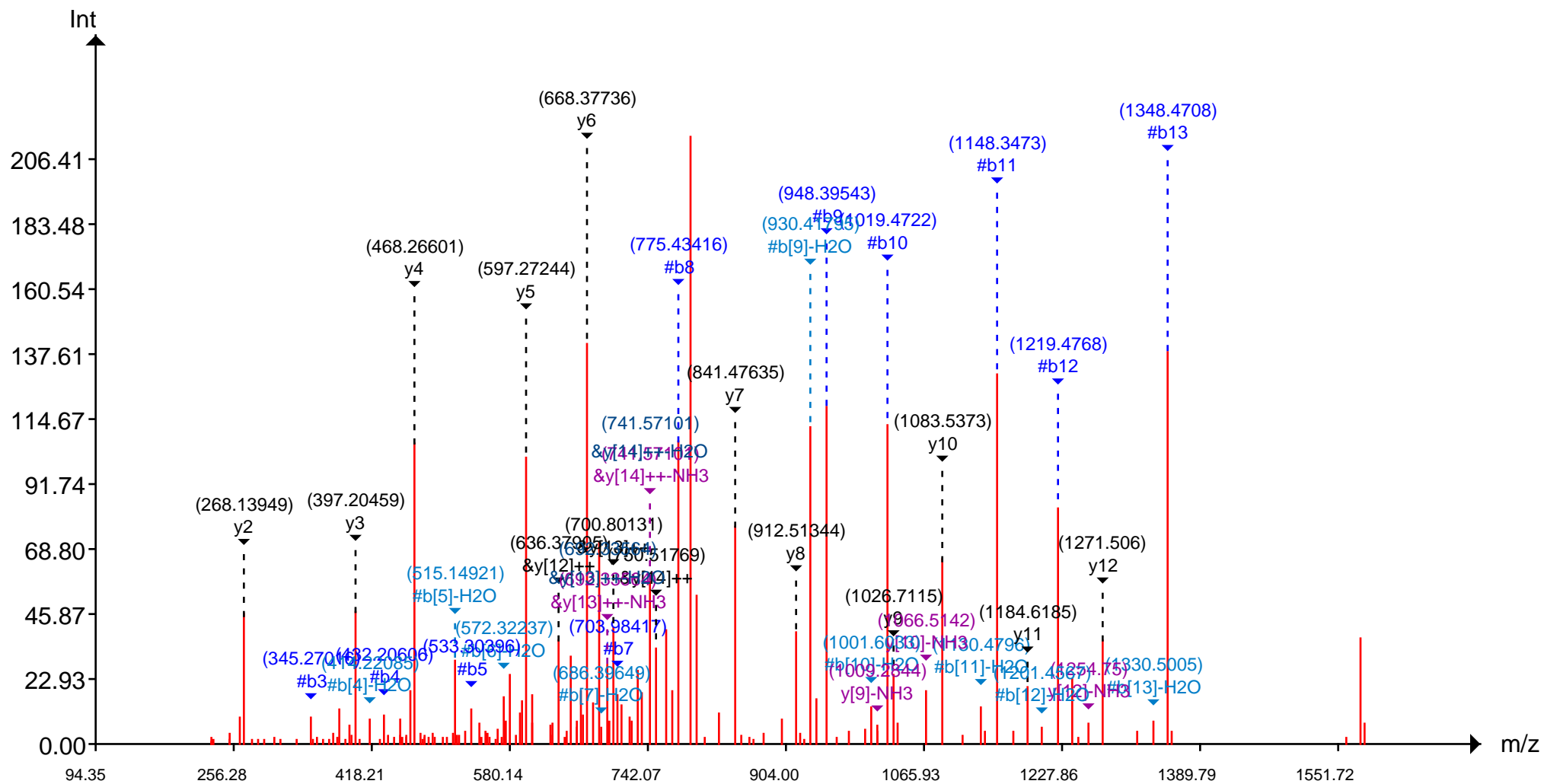
Info

Modified Sequence	AcD3-IAAC<Cmm*>TEK<AcD3K*>K<AcD3K*>IPVVFR<C13>-COOH	Ion score	45.75
Protein	Q9R0P3	Confidence level	99.00%
Precursor m/z	887.02409	Identity Threshold	43.64
Charge	2+	Mass error	0.002424



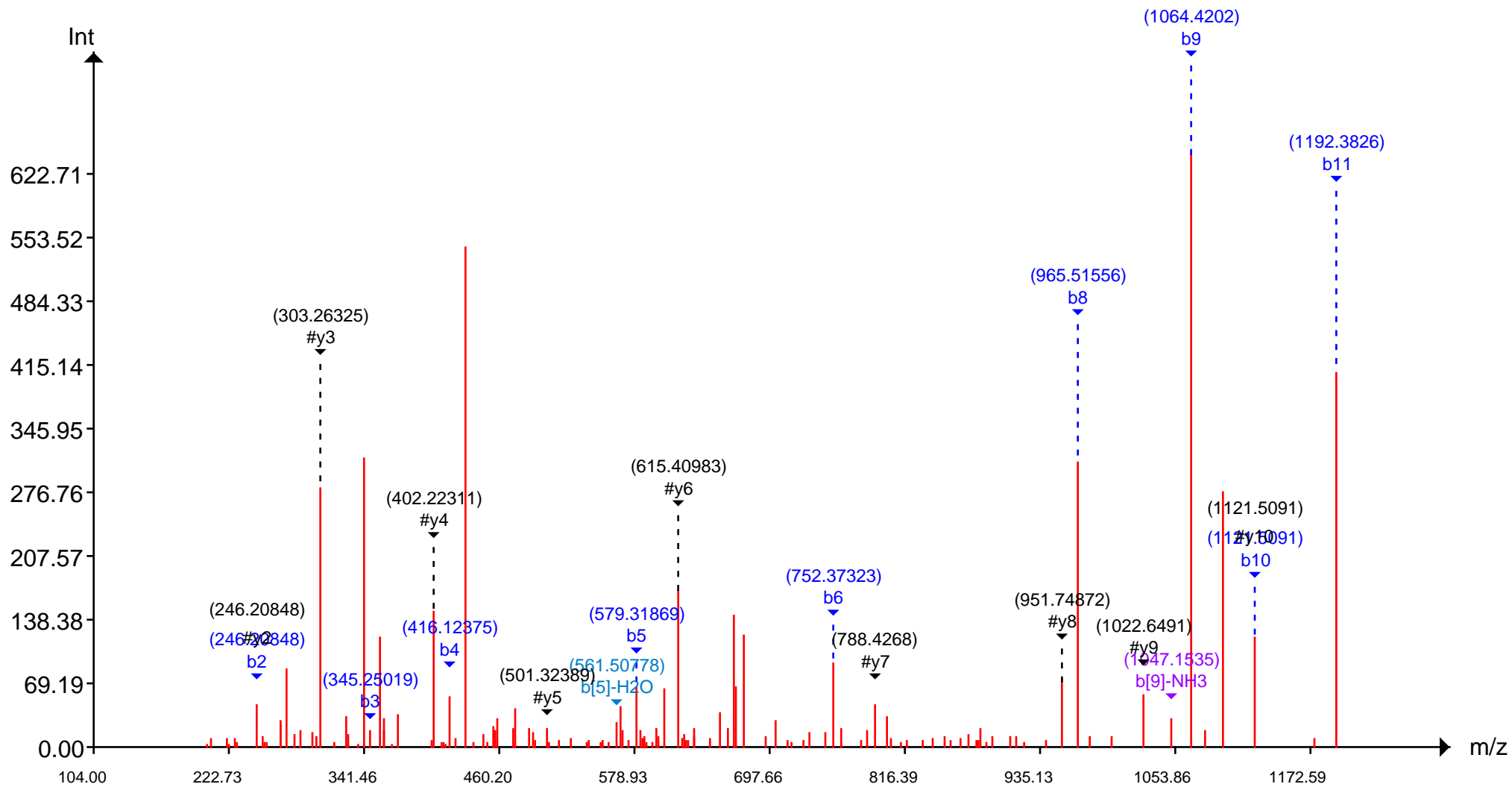
Info

Modified Sequence	AcD3-IAAC<Cmm*>TEK<AcD3K*>K<AcD3K*>IPVVFR-COOH	Ion score	70.0
Protein	Q9R0P3	Confidence level	99.00%
Precursor m/z	884.0141	Identity Threshold	44.73
Charge	2+	Mass error	0.002585



Info

Modified Sequence	AcD3-AVESTGNAK<AcD3K*>AEAESR<C13>-COOH	Ion score	79.75
Protein	Q9EQK5	Confidence level	99.00%
Precursor m/z	808.41107	Identity Threshold	47.95
Charge	2+	Mass error	0.001283



Info

Modified Sequence AcD3-LSVAYK<AcD3K*>NVVGAR-COOH

Ion score 56.81

Protein Q9CQV8, P68510, P61982, P63101

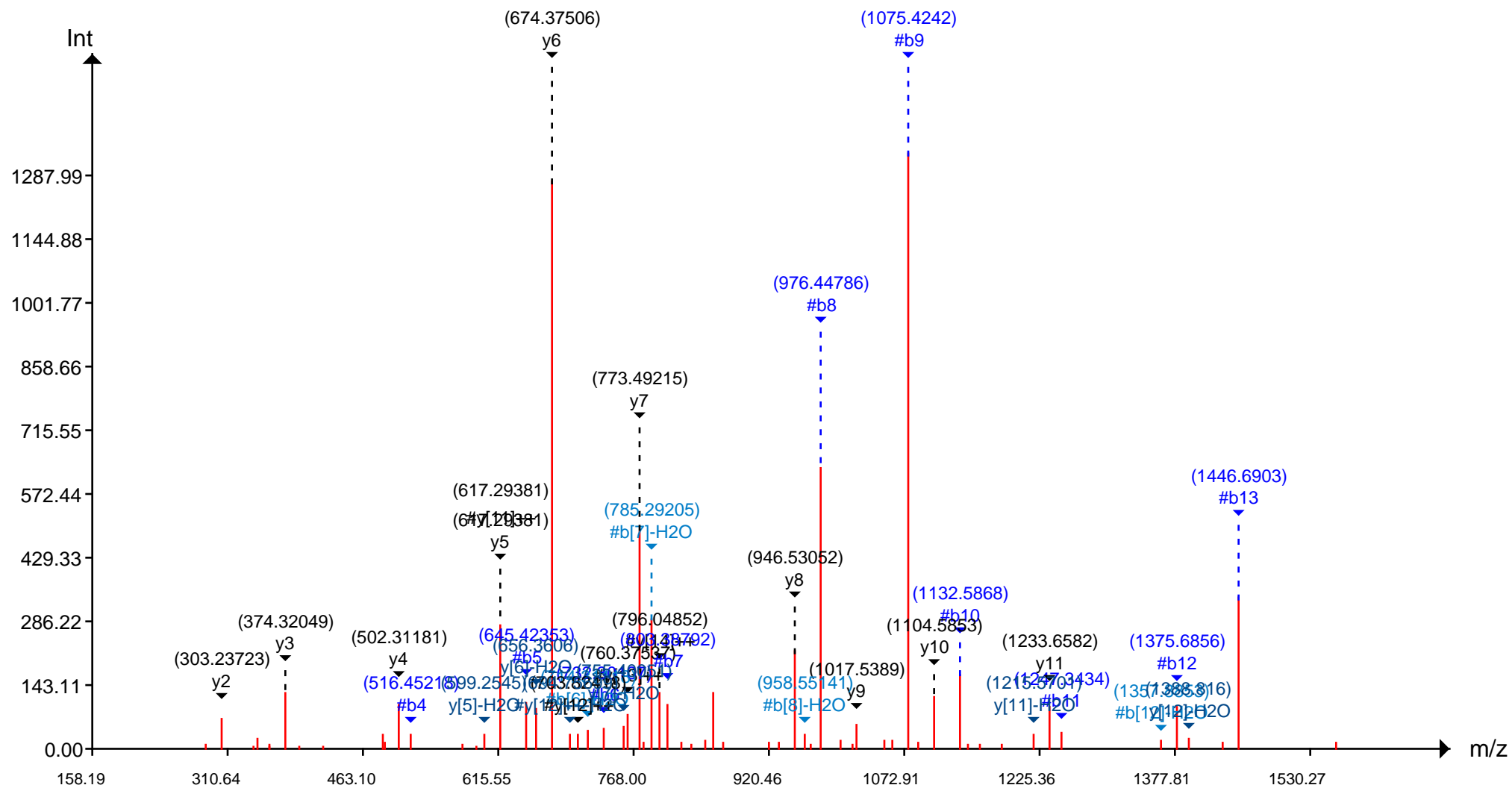
Confidence level 99.00%

Precursor m/z 683.90198

Identity Threshold 45.44

Charge 2+

Mass error 7.12E-4



Info

Modified Sequence AcD3-IAIK<AcD3K*>ESAK<AcD3K*>VGDQAQR-COOH

Ion score 95.24

Protein Q99JY9

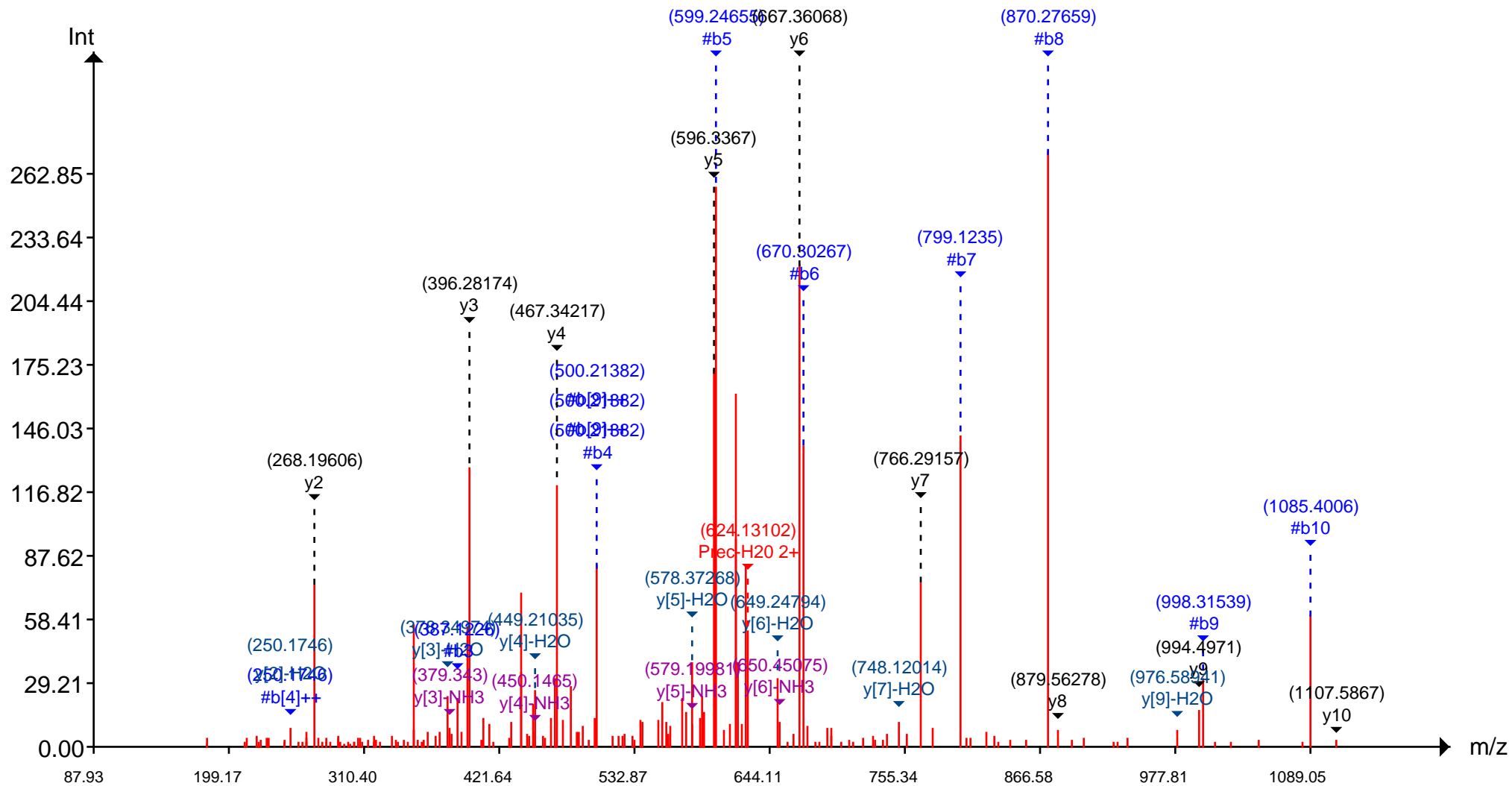
Confidence level 99.00%

Precursor m/z 874.99725

Identity Threshold 46.20

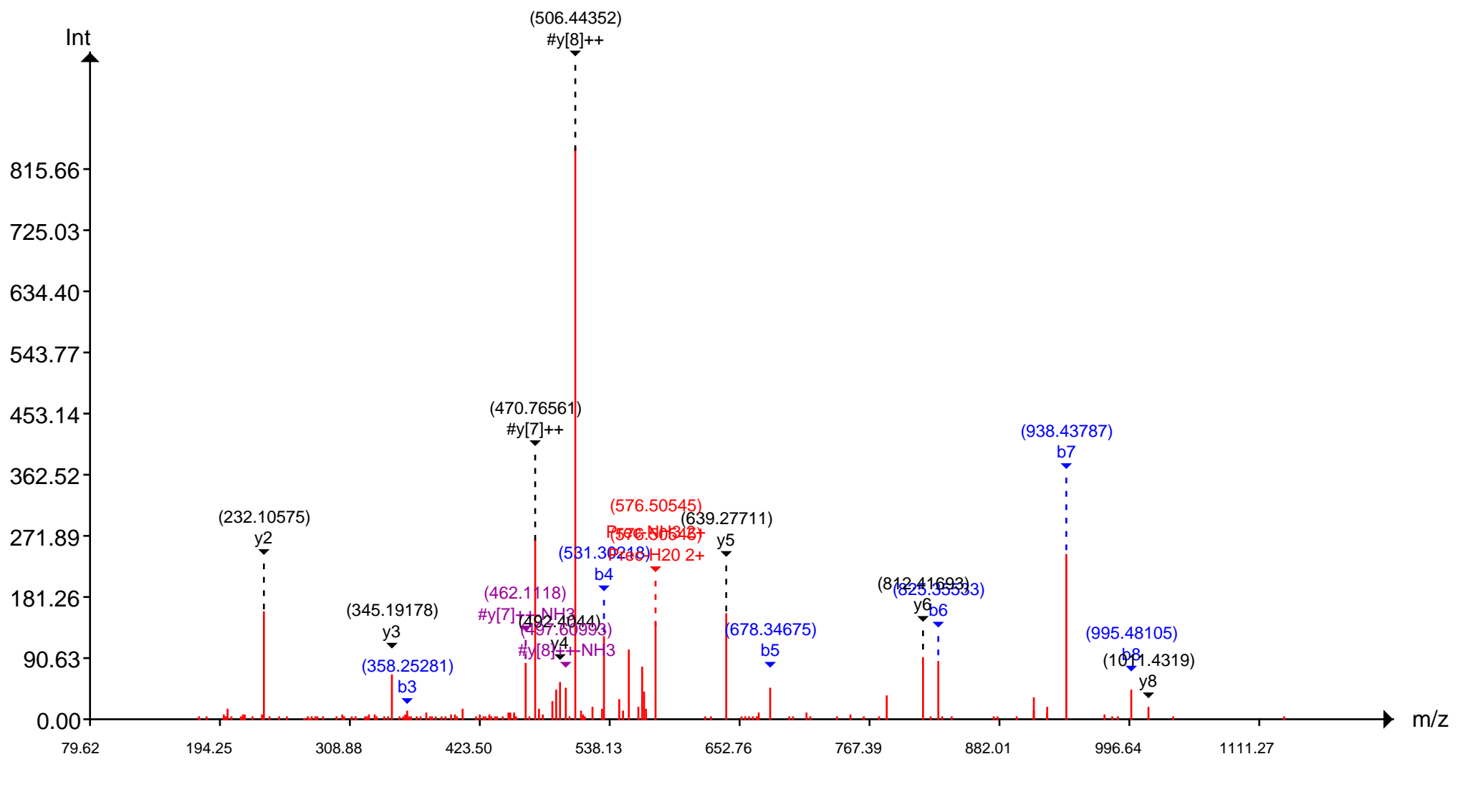
Charge 2+

Mass error 0.002119

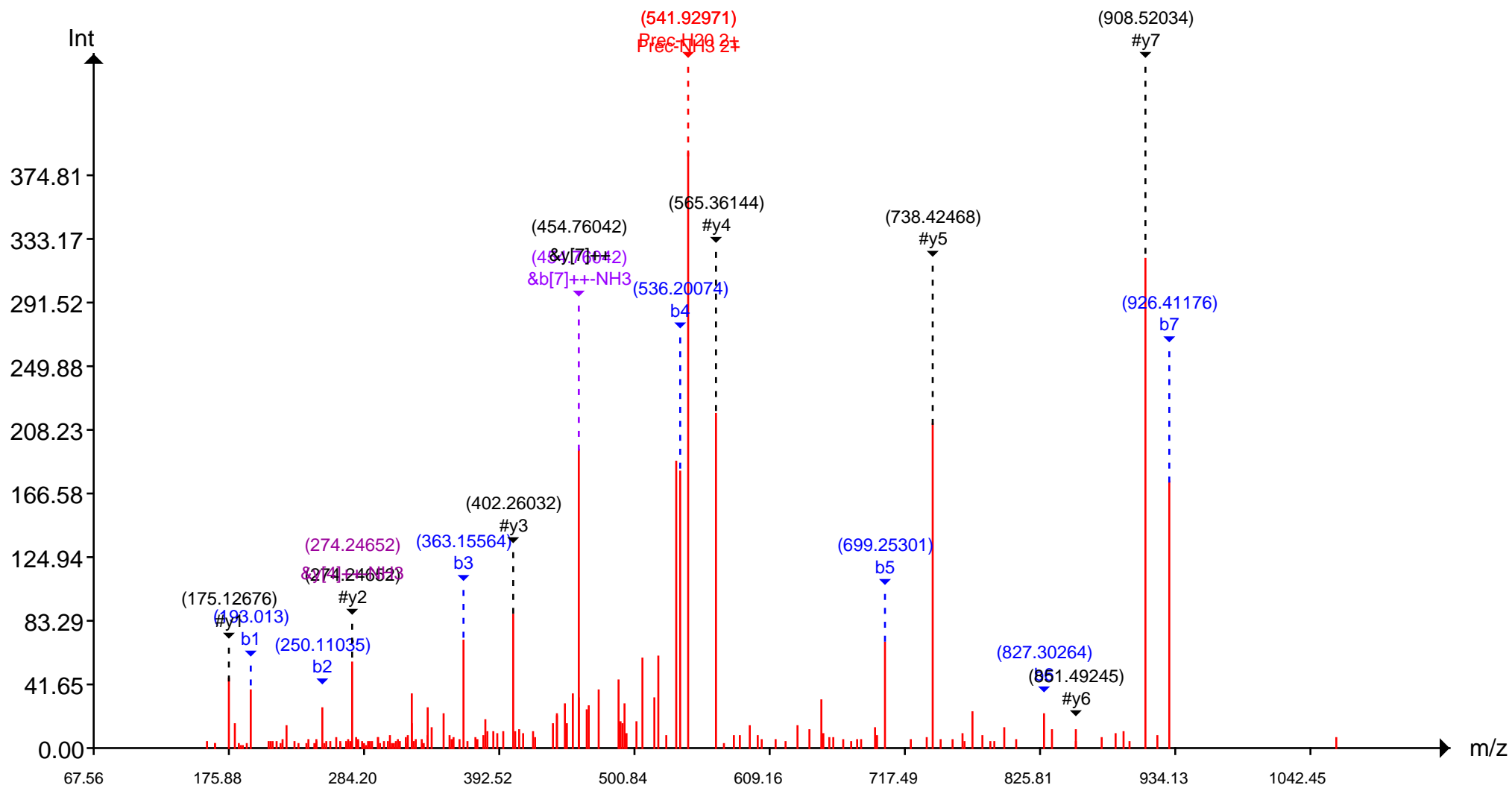


Info

Modified Sequence	AcD3-LLDLVAEAQSR<C13>-COOH	Ion score	75.61
Protein	Q3U1Z5	Confidence level	99.00%
Precursor m/z	633.36569	Identity Threshold	46.31
Charge	2+	Mass error	6.72E-4



Info			
Modified Sequence	AcD3-LAQQ<AcD3K*>M<Mox*>M<Mox*>IGR-COOH	Ion score	49.22
Protein	P52480, P53657	Confidence level	99.00%
Precursor m/z	585.31818	Identity Threshold	47.86
Charge	2+	Mass error	3.18E-4



Info

Modified Sequence AcD3-M<Mox*>GLK<AcD3K*>YQVR-COOH

Protein P70677

Precursor m/z 550.80555

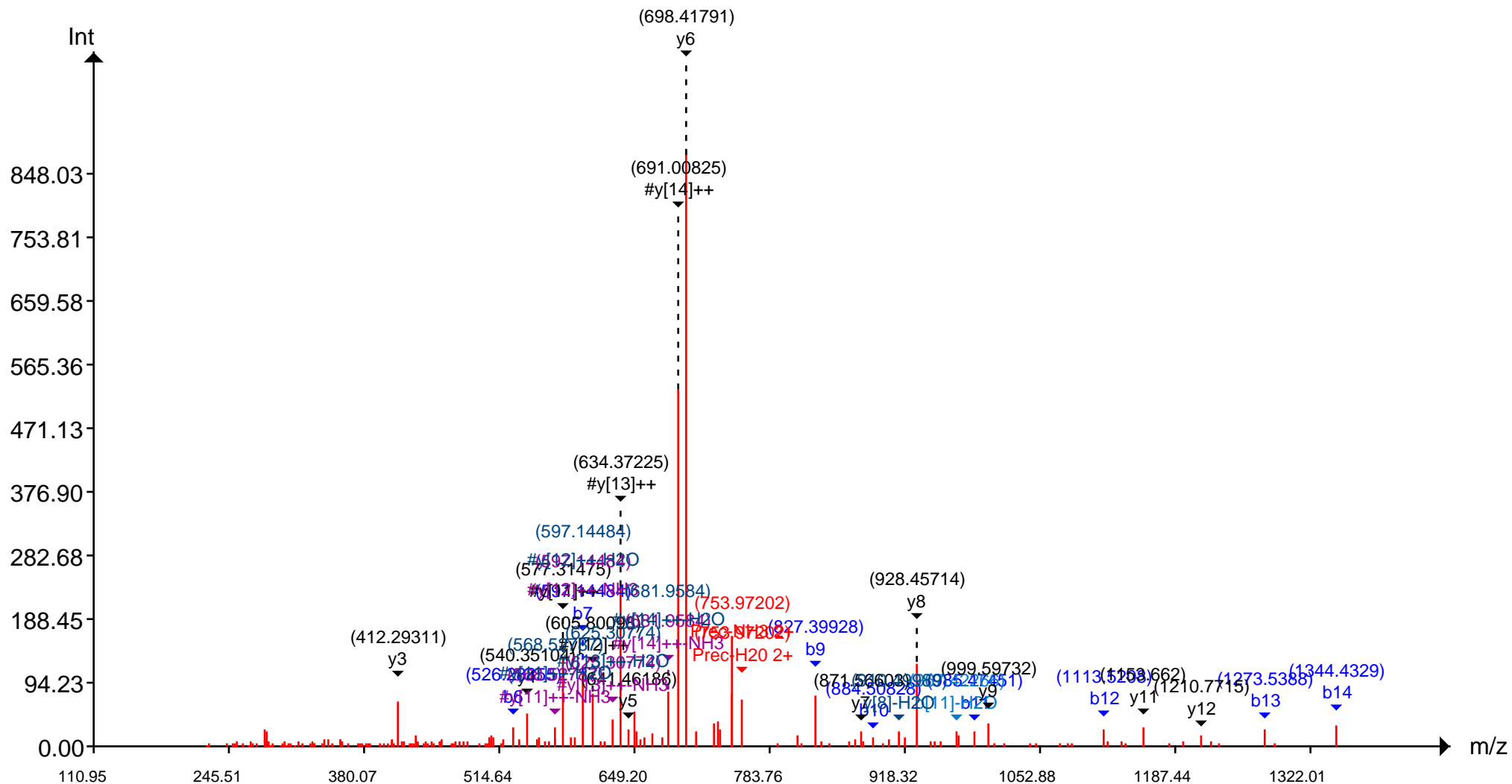
Charge 2+

Ion score 51.17

Confidence level 99.00%

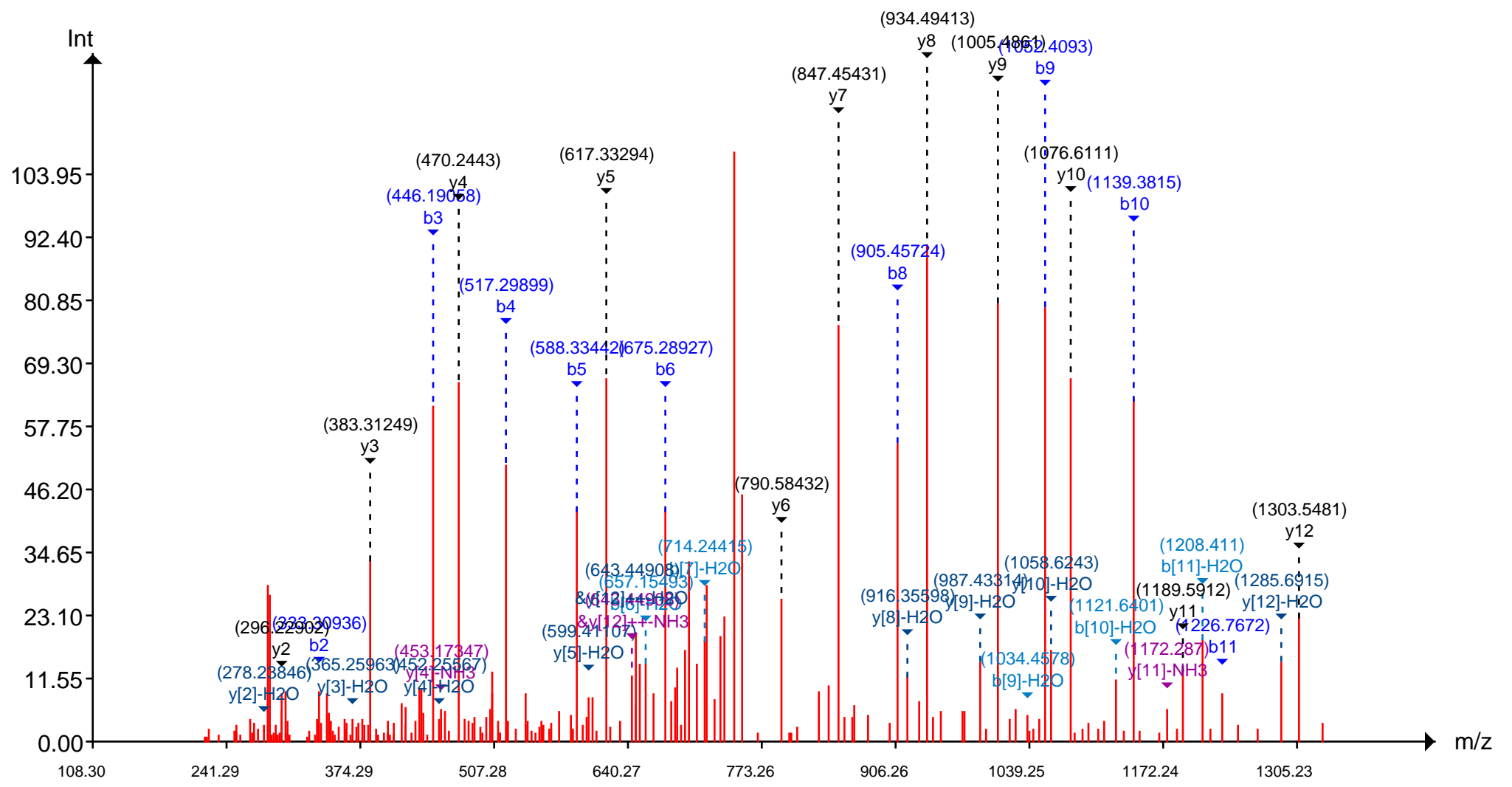
Identity Threshold 47.53

Mass error -1.16E-4

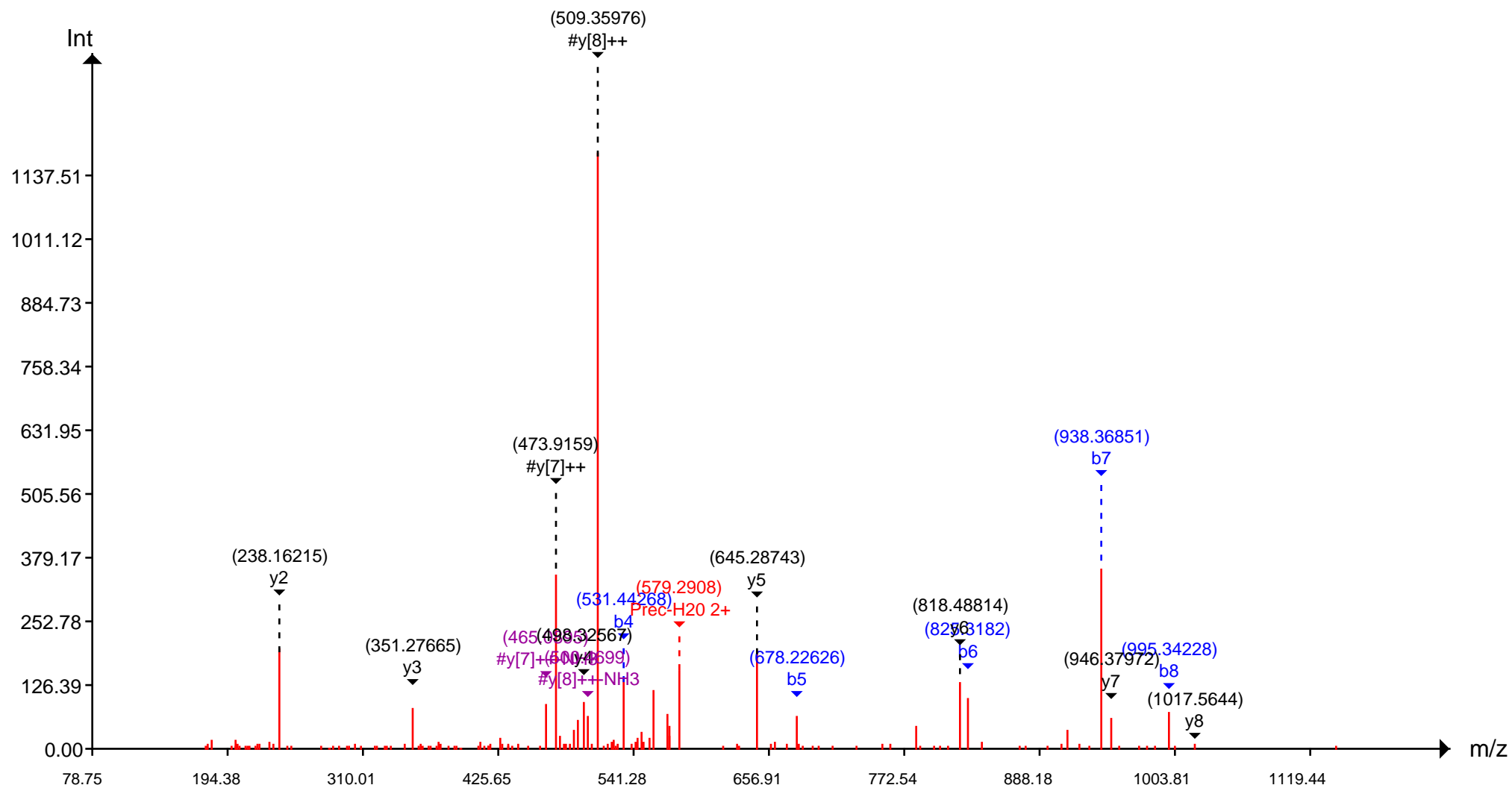


Info

Modified Sequence	AcD3-VLGGPGAGK<AcD3K*>GTQC<Cmm*>AR<C13>-COOH	Ion score	51.52
Protein	Q9DBP5	Confidence level	99.00%
Precursor m/z	762.91215	Identity Threshold	47.95
Charge	2+	Mass error	4.68E-4



Info			
Modified Sequence	AcD3-K<AcD3K*>NIAASGK<AcD3K*>FSSDR<C13>-COOH	Ion score	83.49
Protein	Q9ET01	Confidence level	99.00%
Precursor m/z	761.42044	Identity Threshold	47.62
Charge	2+	Mass error	0.002292



Info

Modified Sequence AcD3-LAQK<AcD3K*>M<Mox*>M<Mox*>IGR<C13>-COOH

Ion score 54.04

Protein P52480, P53657

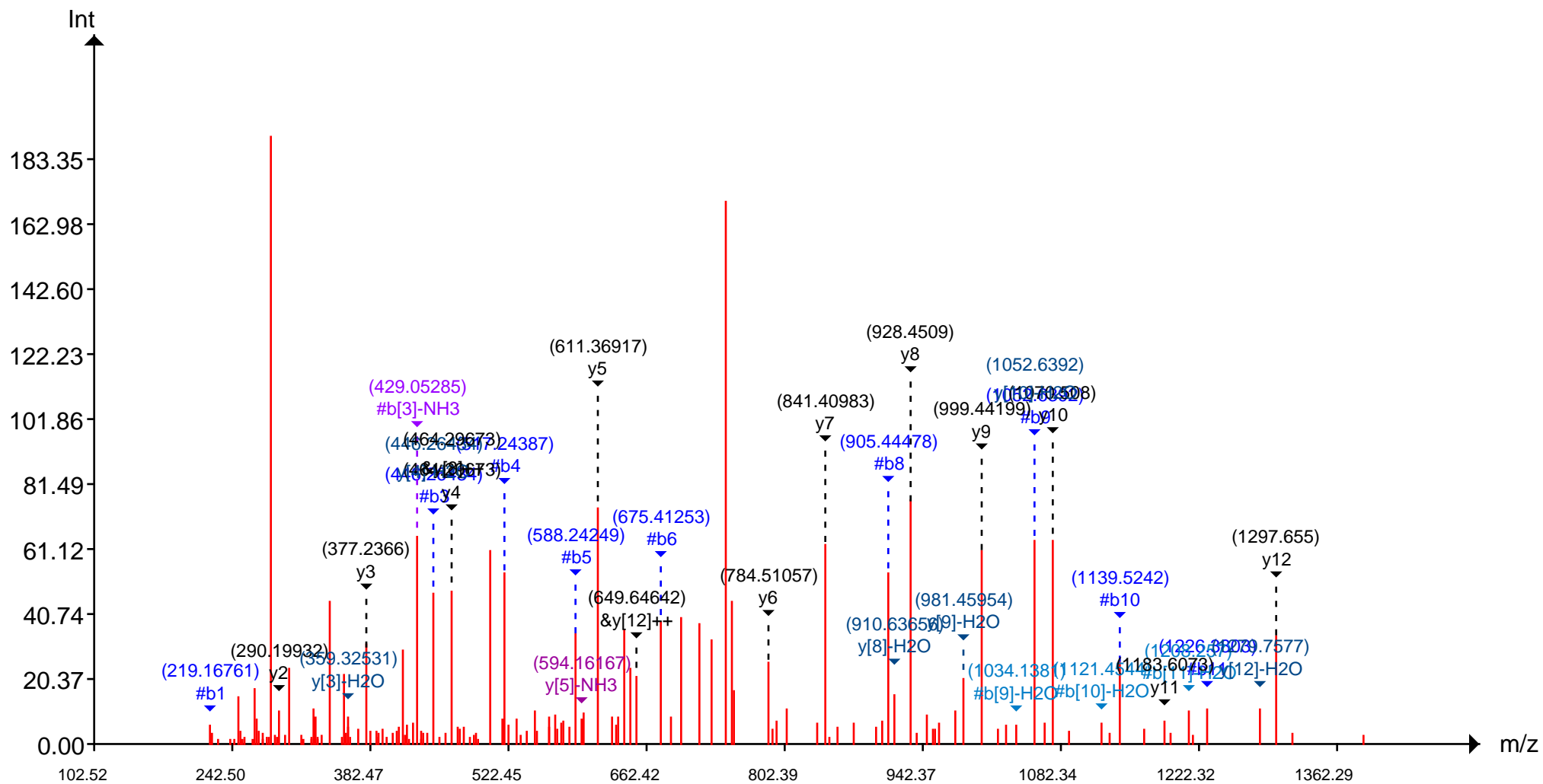
Confidence level 99.00%

Precursor m/z 588.32796

Identity Threshold 47.60

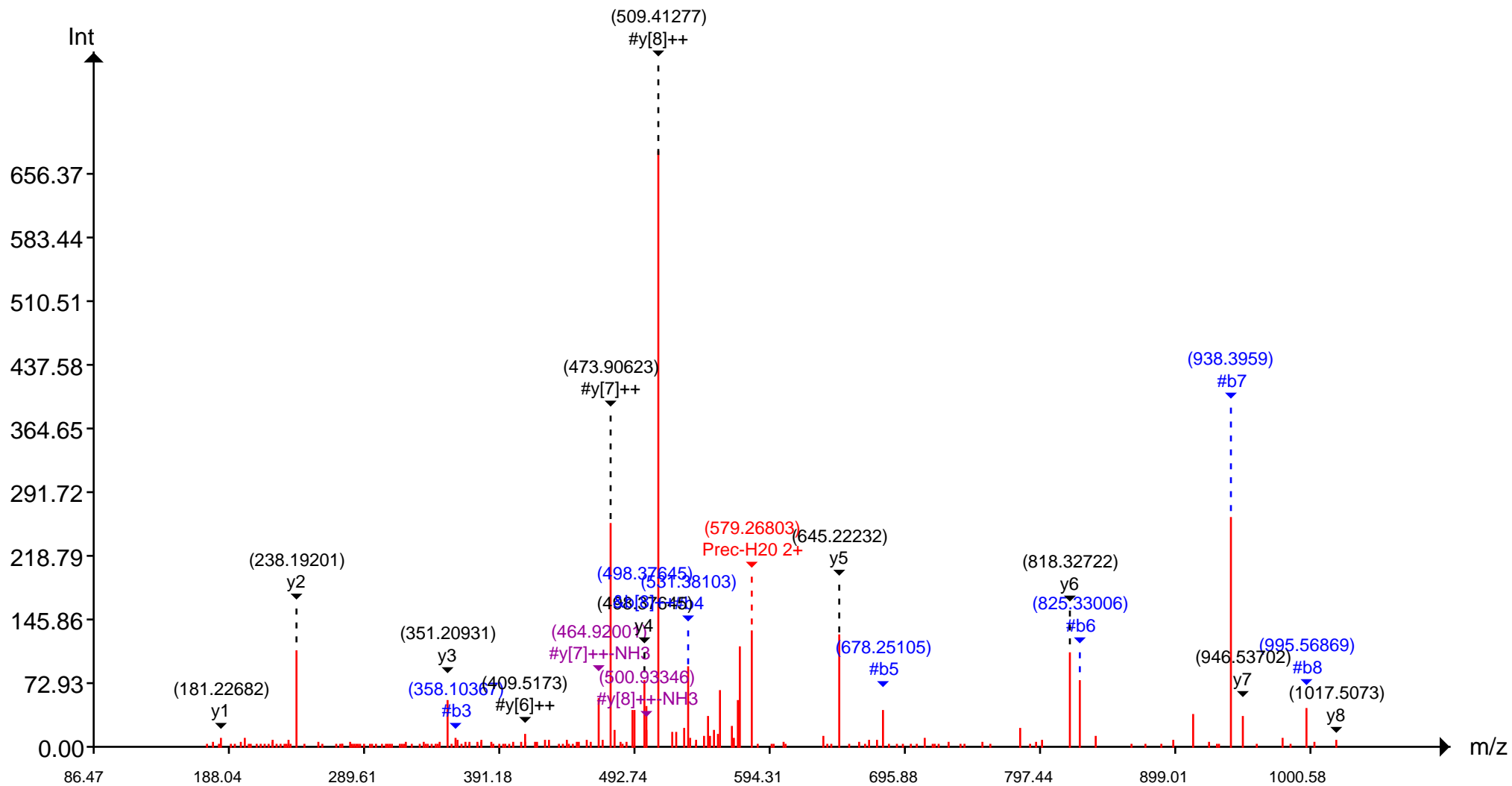
Charge 2+

Mass error -2.64E-4



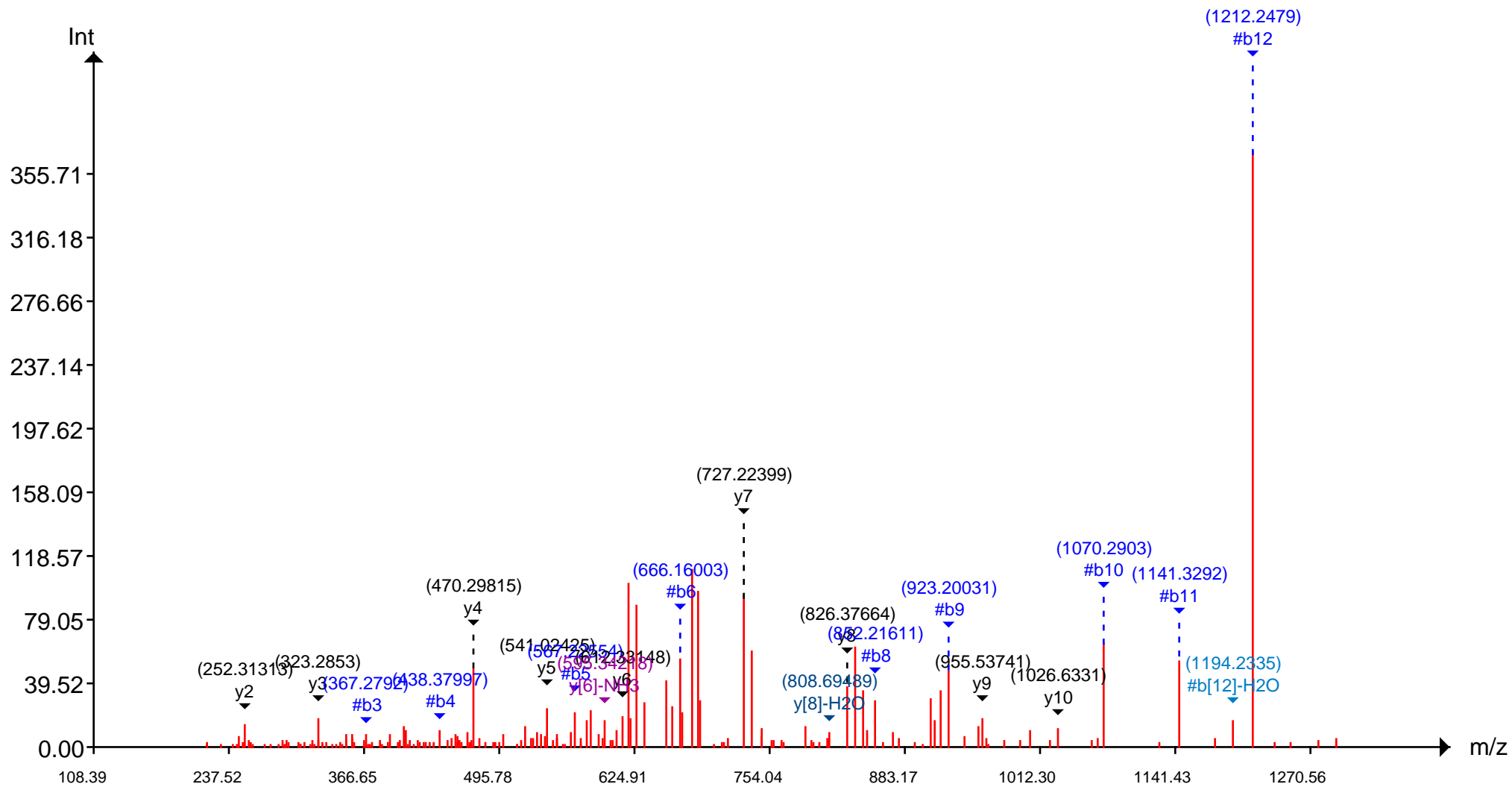
Info

Modified Sequence	AcD3-K<AcD3K*>NIAASGK<AcD3K*>FSSDR-COOH	Ion score	69.17
Protein	Q9ET01	Confidence level	99.00%
Precursor m/z	758.41017	Identity Threshold	47.84
Charge	2+	Mass error	0.001894



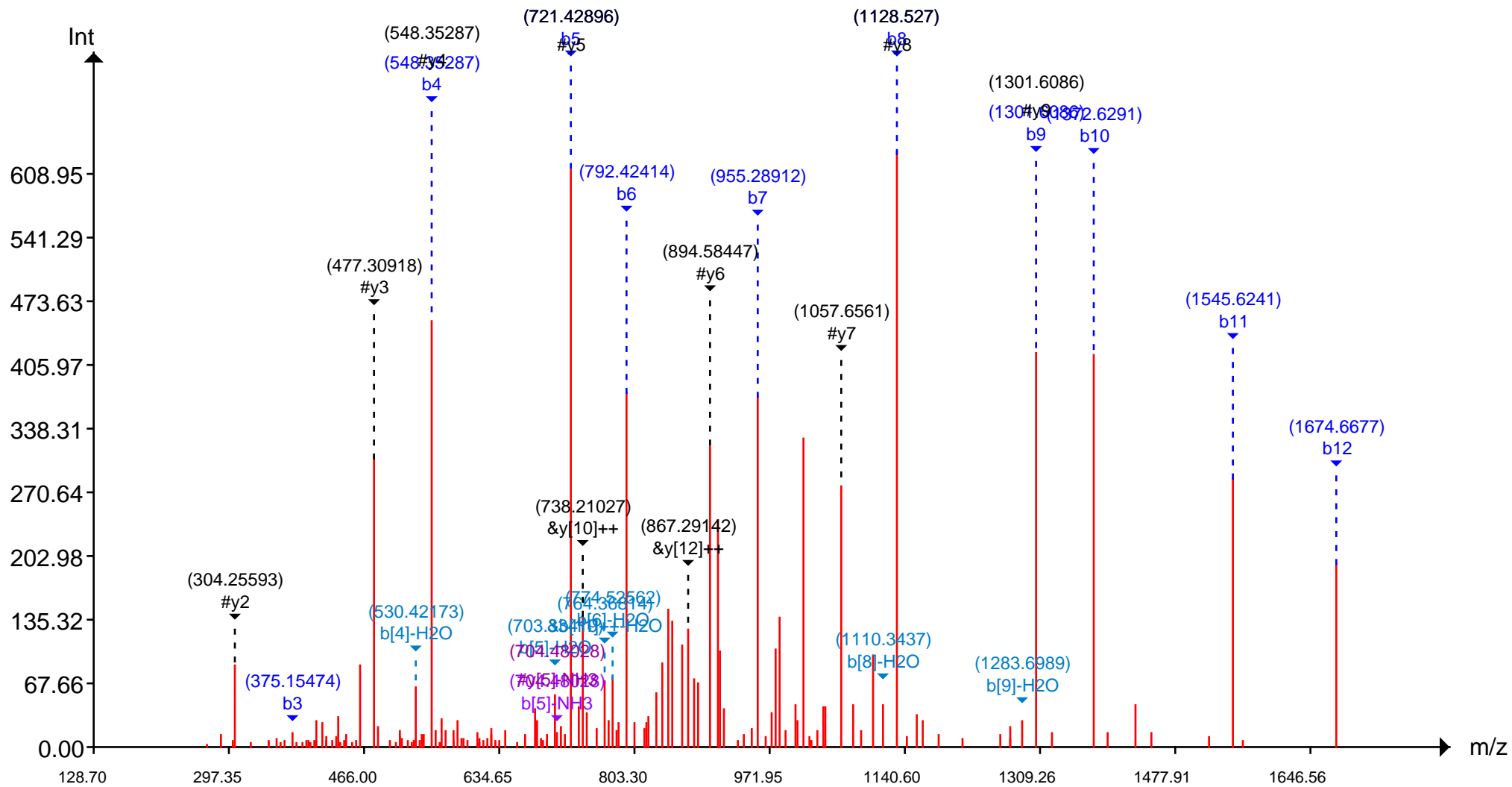
Info

Modified Sequence	AcD3-LAQK<AcD3K*>M<Mox*>M<Mox*>IGR<C13>-COOH	Ion score	48.01
Protein	P52480, P53657	Confidence level	99.00%
Precursor m/z	588.32822	Identity Threshold	47.63
Charge	2+	Mass error	2.56E-4



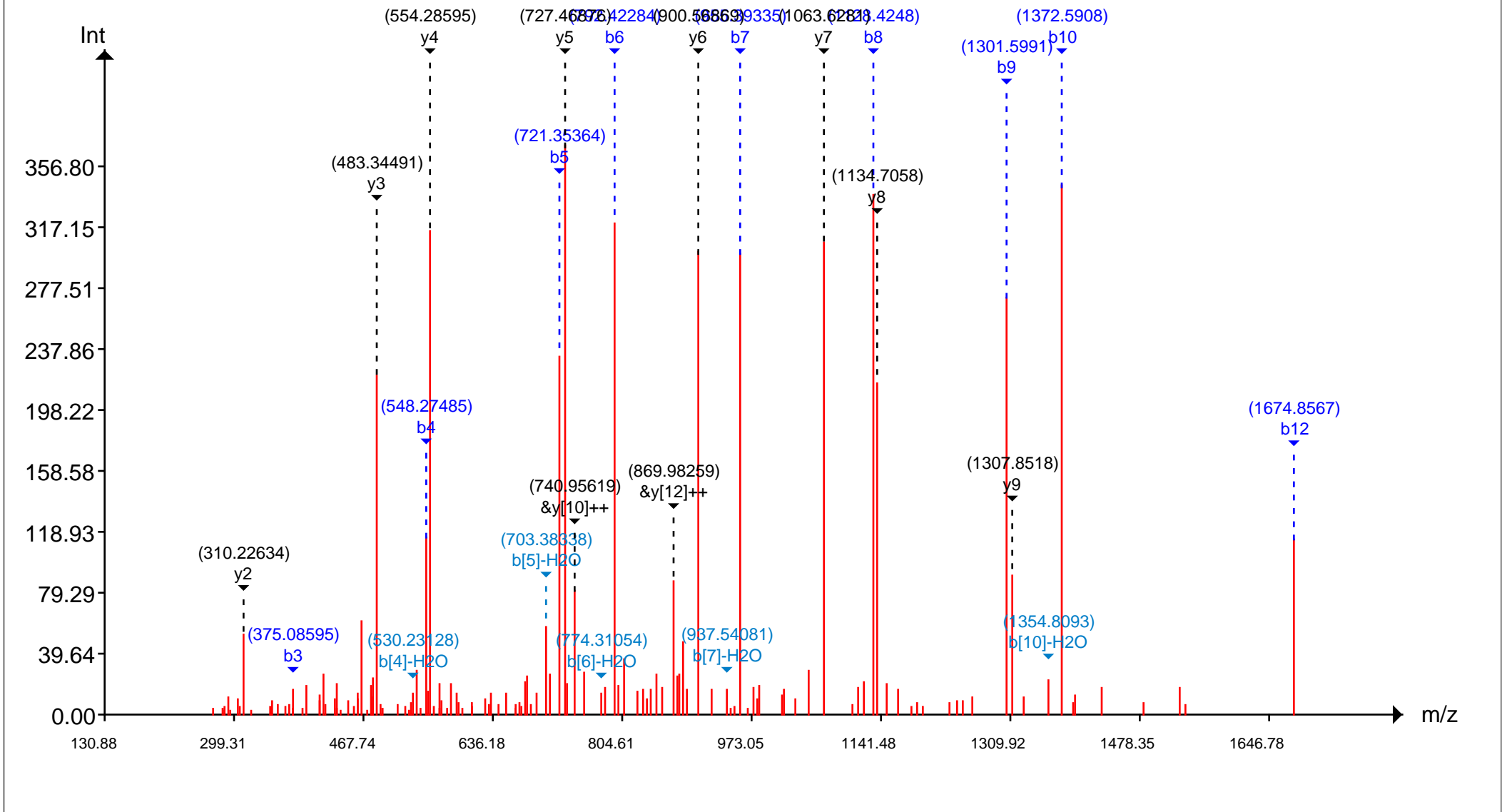
Info

Modified Sequence	AcD3-SSM<Mox*>AEVDAAM<Mox*>AAR<C13>-COOH	Ion score	57.76
Protein	O88569	Confidence level	99.00%
Precursor m/z	696.8174	Identity Threshold	44.66
Charge	2+	Mass error	6.95E-4



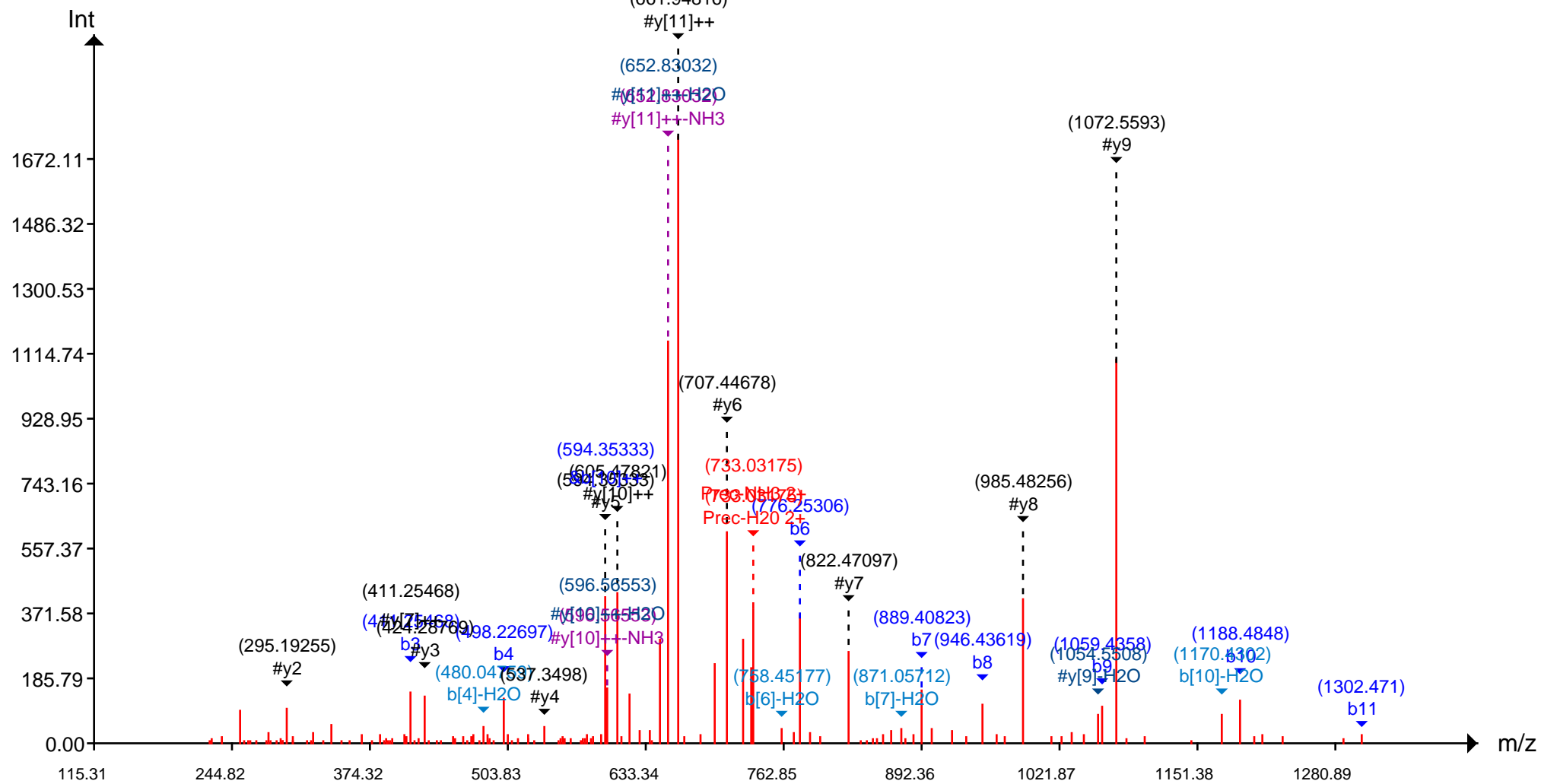
Info

Modified Sequence	AcD3-AEEK<AcD3K*>K<AcD3K*>AYK<AcD3K*>K<AcD3K*>AK<AcD3K*>ER-COOH	Ion score	76.3
Protein	Q810U5	Confidence level	99.00%
Precursor m/z	925.04064	Identity Threshold	44.12
Charge	2+	Mass error	0.001466



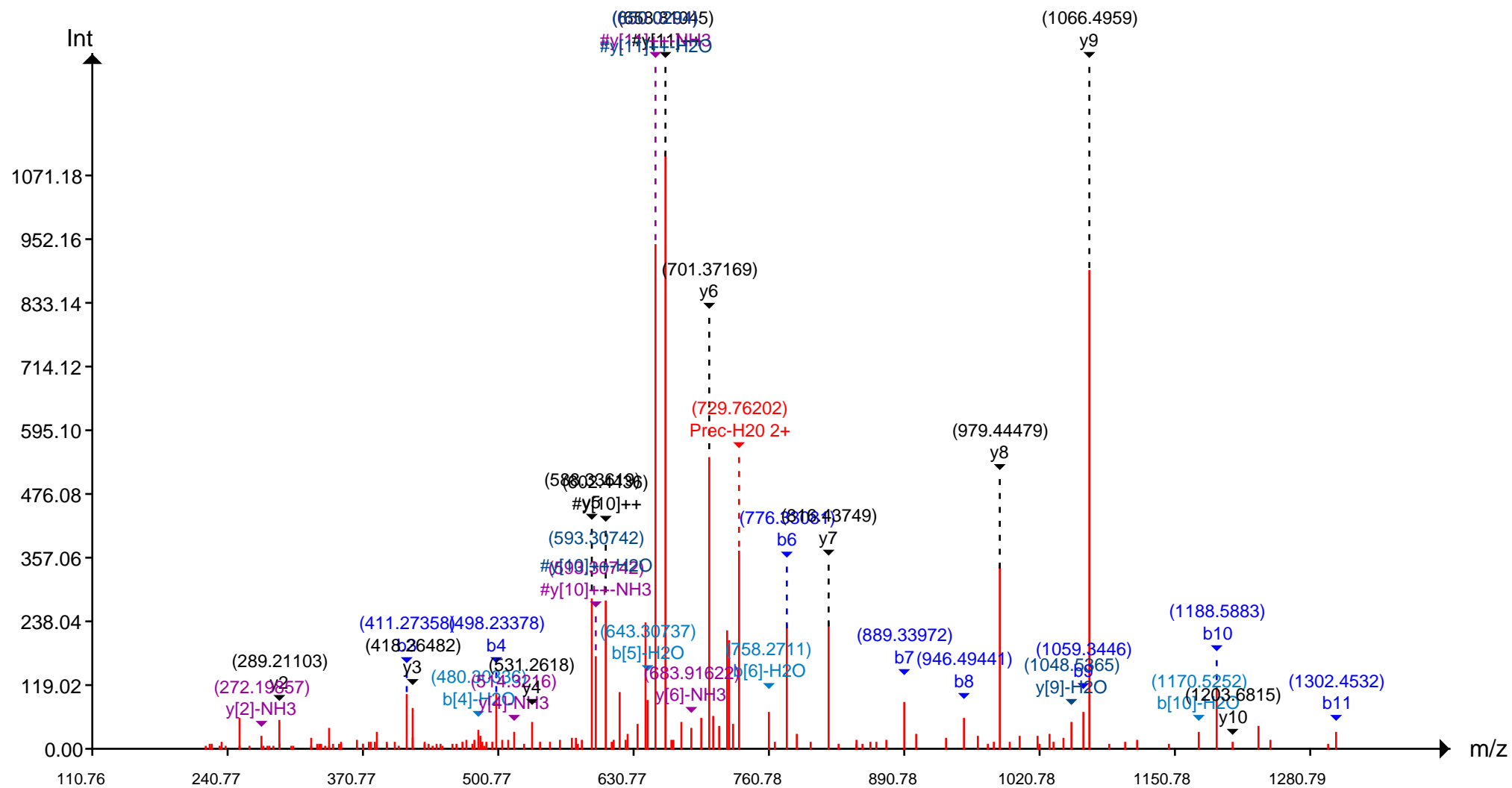
Info

Modified Sequen...	AcD3-AEEK<AcD3K*>K<AcD3K*>AYK<AcD3K*>K<AcD3K*>AK<AcD3K*>ER<C13>-COOH	Ion score	56.41
Prote...	Q810U5	Confidence level	99.00%
Precursor m...	928.05087	Identity Threshold	42.88
Char...	2+	Mass error	0.001785



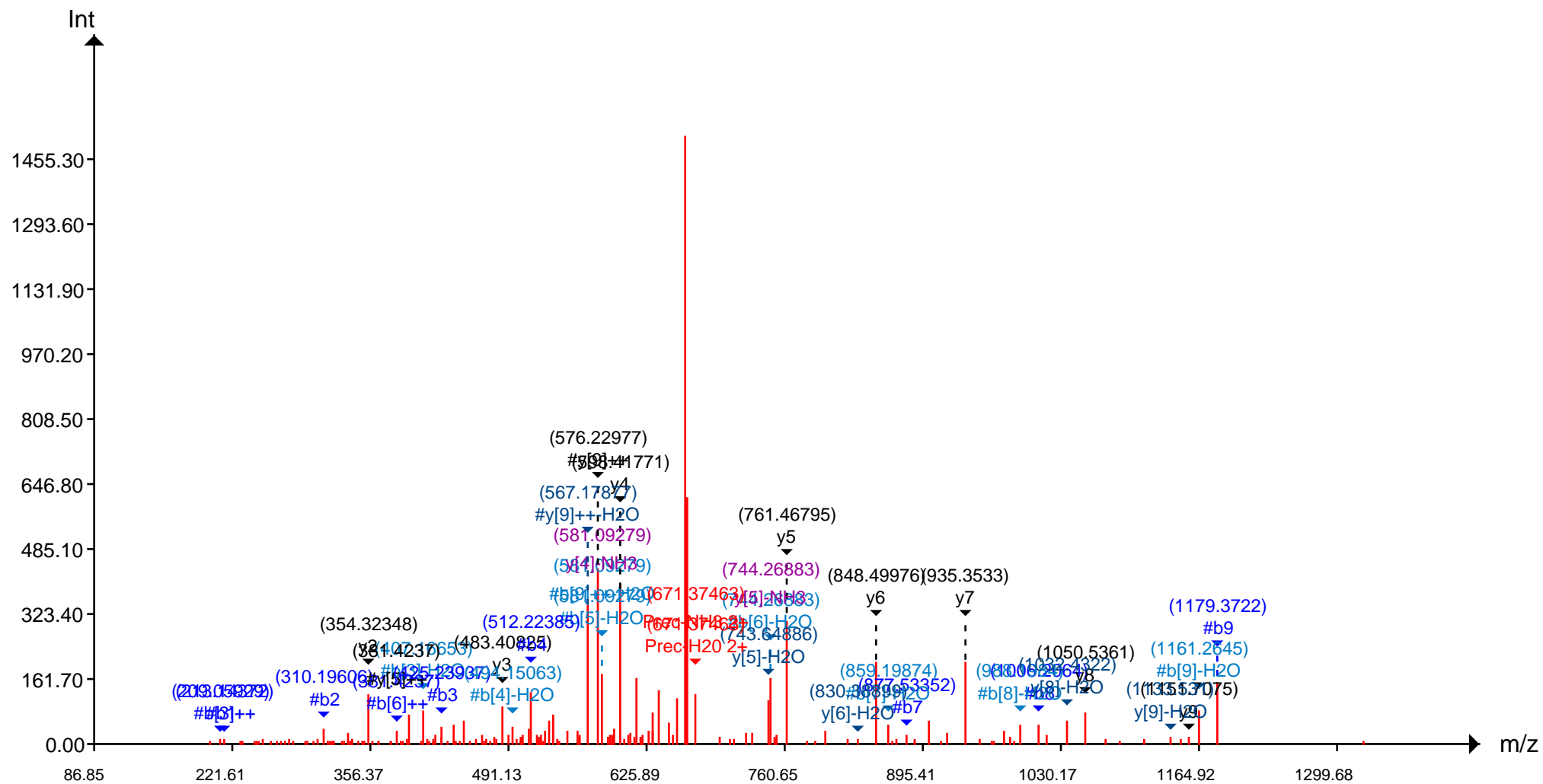
Info

Modified Sequence	AcD3-DLHSYDLGIENR<C13>-COOH	Ion score	65.73
Protein	O88844	Confidence level	99.00%
Precursor m/z	741.87226	Identity Threshold	47.50
Charge	2+	Mass error	0.001452



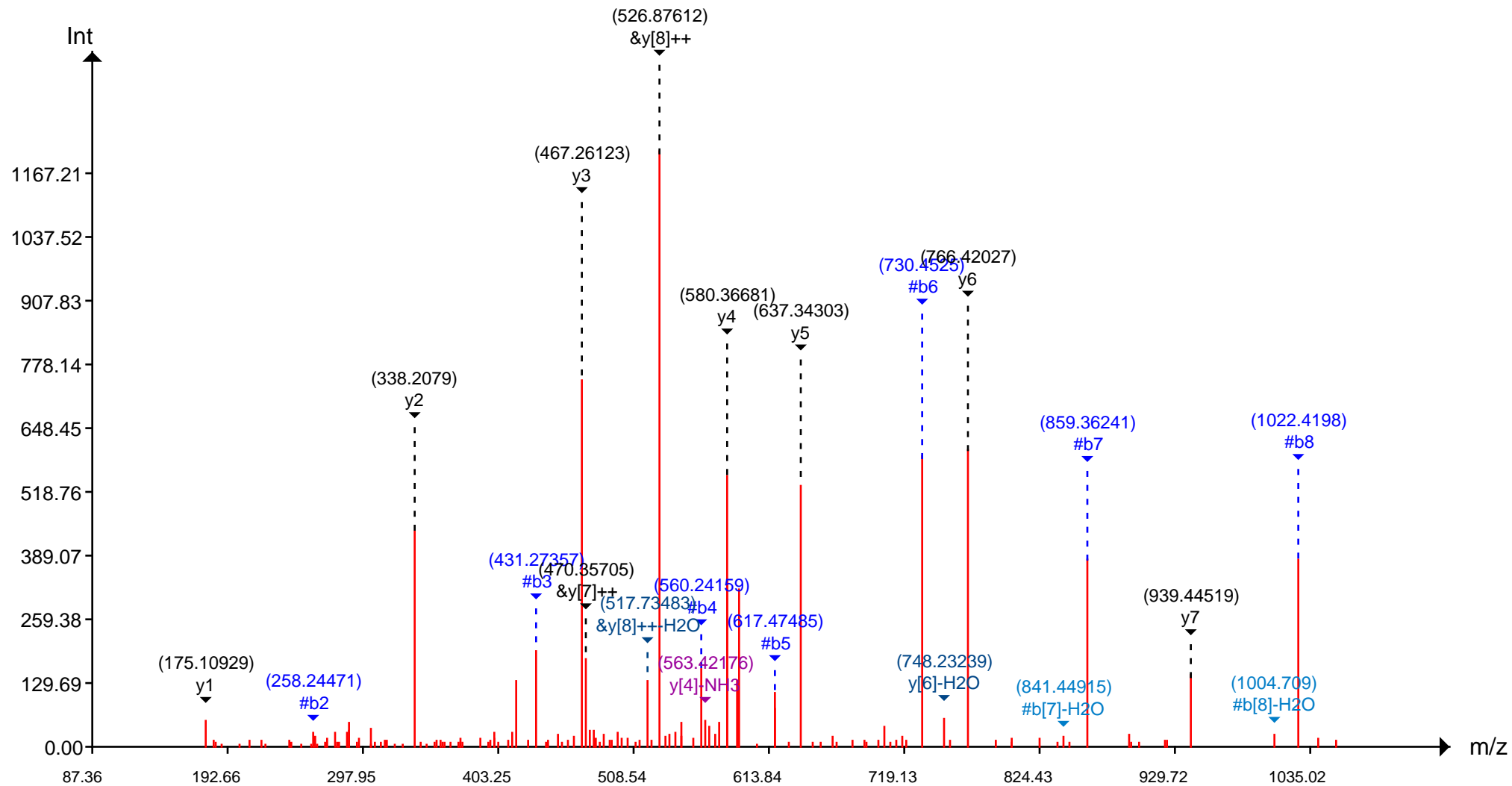
Info

Modified Sequence	AcD3-DLHSYDLGIENR-COOH	Ion score	71.07
Protein	O88844	Confidence level	99.00%
Precursor m/z	738.86197	Identity Threshold	47.09
Charge	2+	Mass error	0.001014



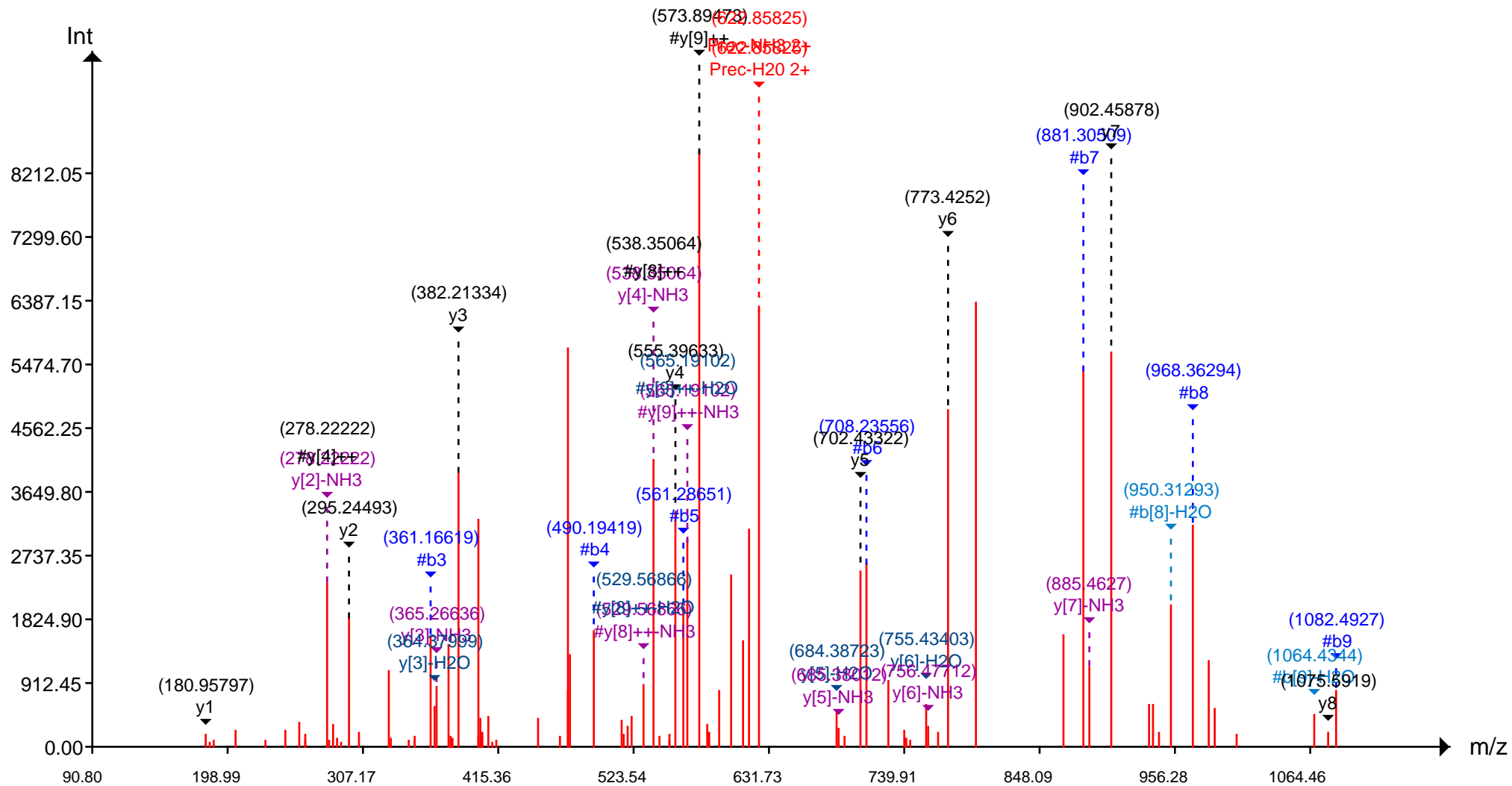
Info

Modified Sequence	AcD3-YTDSSYDEK<AcD3K*>R<C13>-COOH	Ion score	57.31
Protein	P10649	Confidence level	99.00%
Precursor m/z	680.31789	Identity Threshold	45.41
Charge	2+	Mass error	8.06E-4



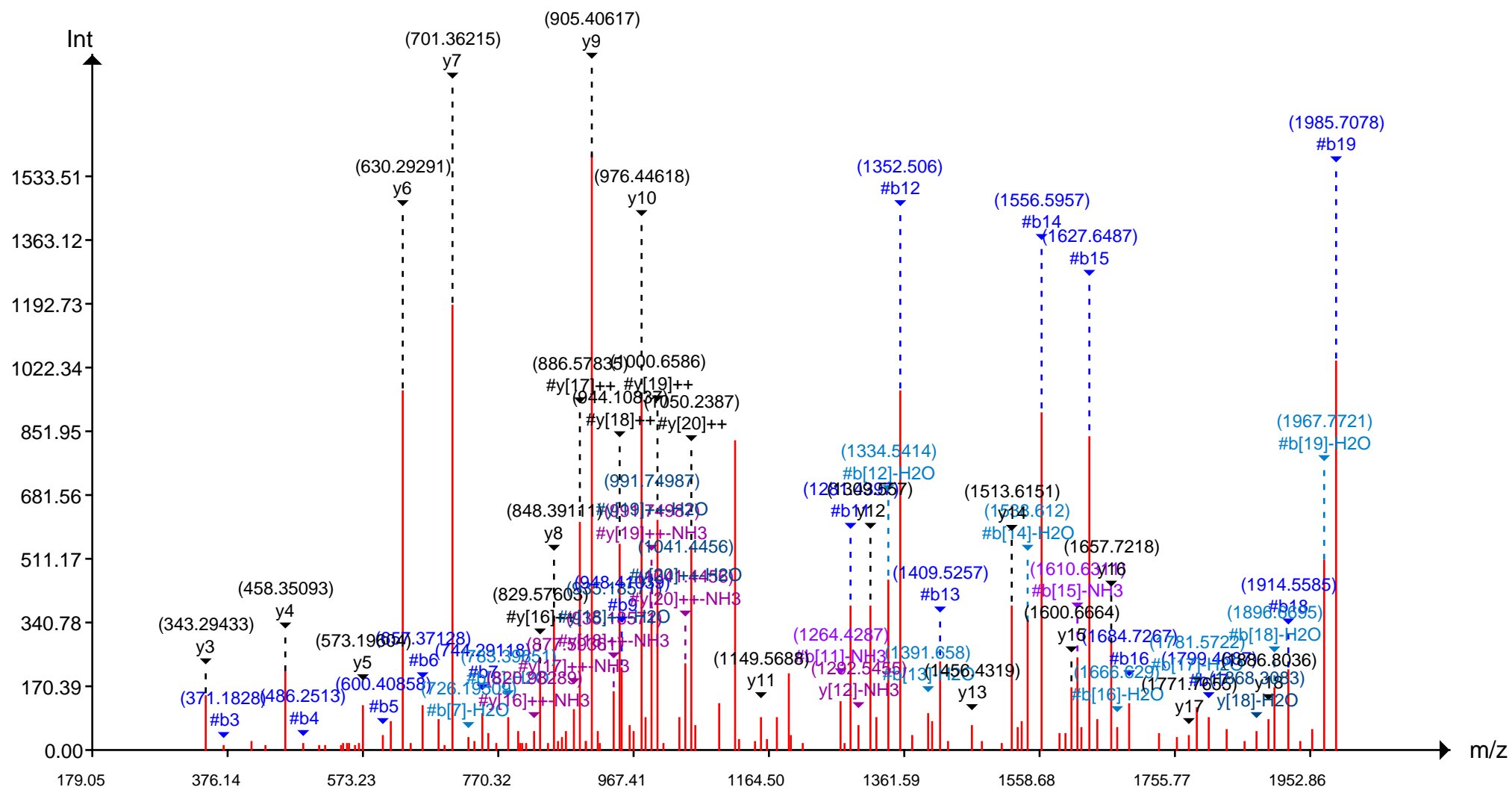
Info

Modified Sequence	AcD3-VLK<AcD3K*>EGIEYR-COOH	Ion score	64.19
Protein	Q61599	Confidence level	99.00%
Precursor m/z	598.84334	Identity Threshold	46.69
Charge	2+	Mass error	2.07E-4



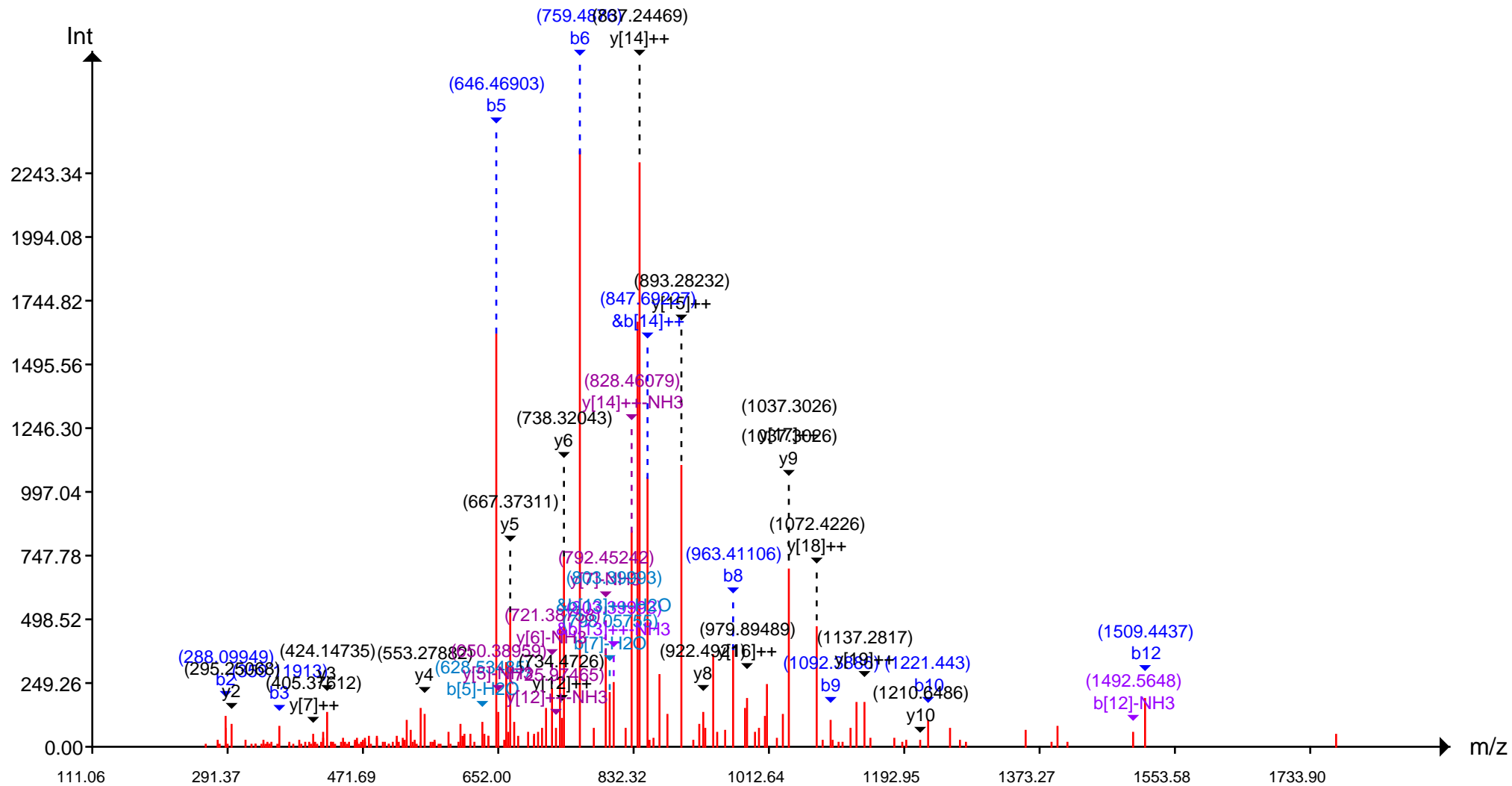
Info

Modified Sequence	AcD3-AAK<AcD3K*>EAM<Mox*>K<AcD3K*>SNR<C13>-COOH	Ion score	49.04
Protein	Q8CIN4	Confidence level	99.00%
Precursor m/z	631.84366	Identity Threshold	47.94
Charge	2+	Mass error	-0.001411



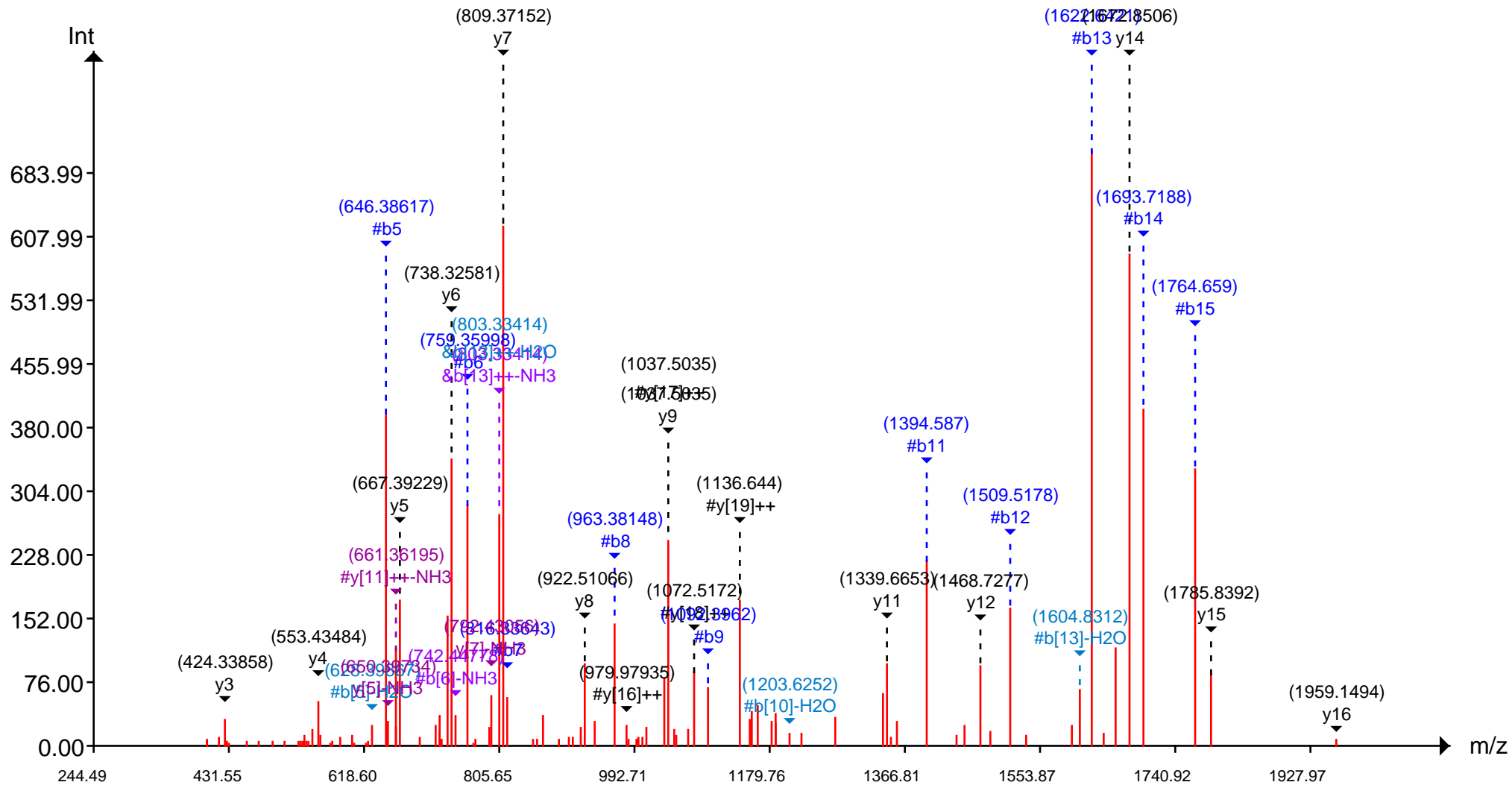
Info

Modified Sequence	AcD3-LVIDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR-COOH	Ion score	138.23
Protein	P63260	Confidence level	99.00%
Precursor m/z	1129.0295	Identity Threshold	45.52
Charge	2+	Mass error	0.002257



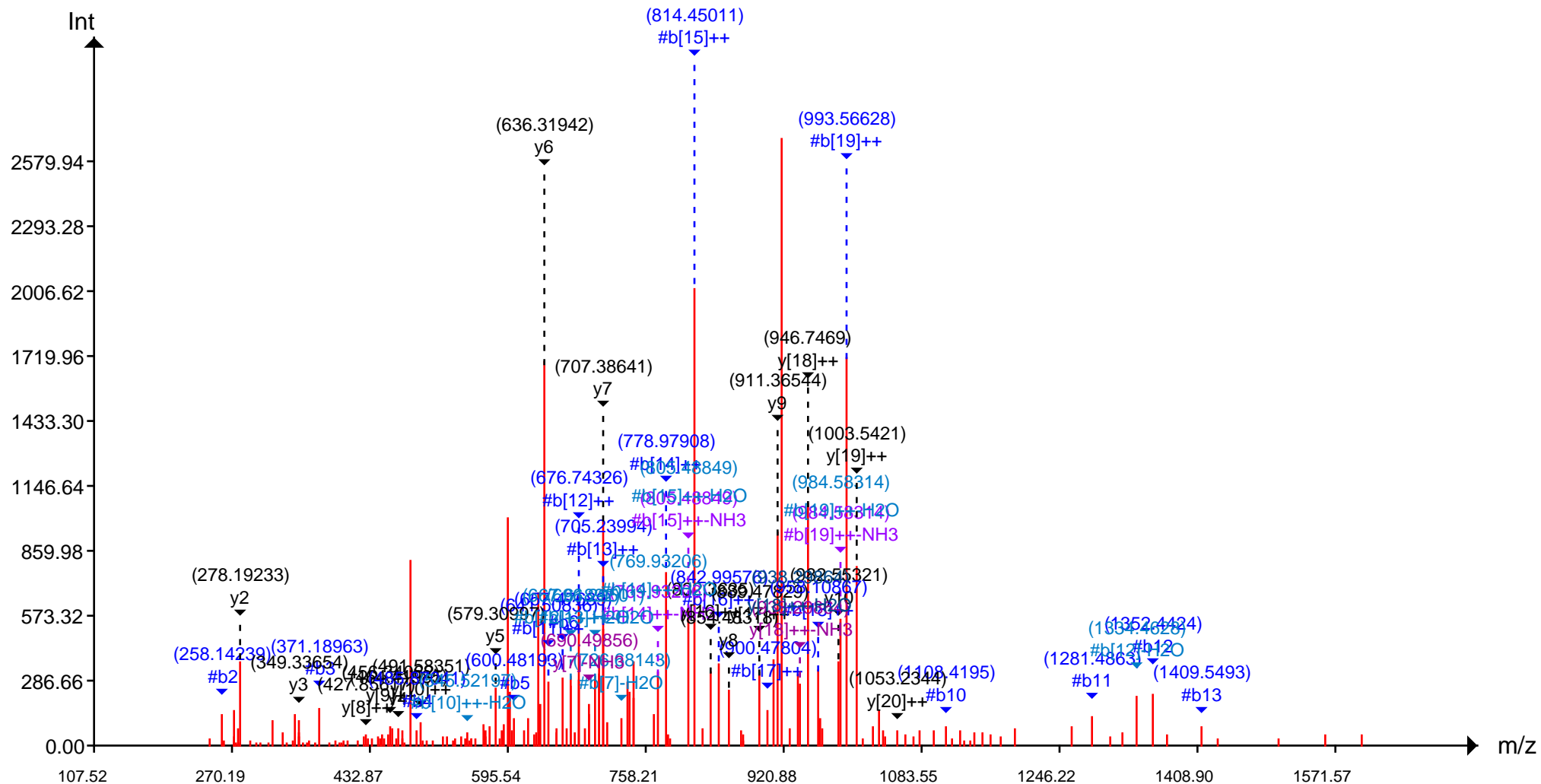
Info

Modified Sequence	AcD3-LEANK<AcD3K*>IGFEEK<AcD3K*>DIAANEENR<C13>-COOH	Ion score	58.29
Protein	Q9JJU8	Confidence level	99.00%
Precursor m/z	811.08532	Identity Threshold	47.91
Charge	3+	Mass error	0.002168



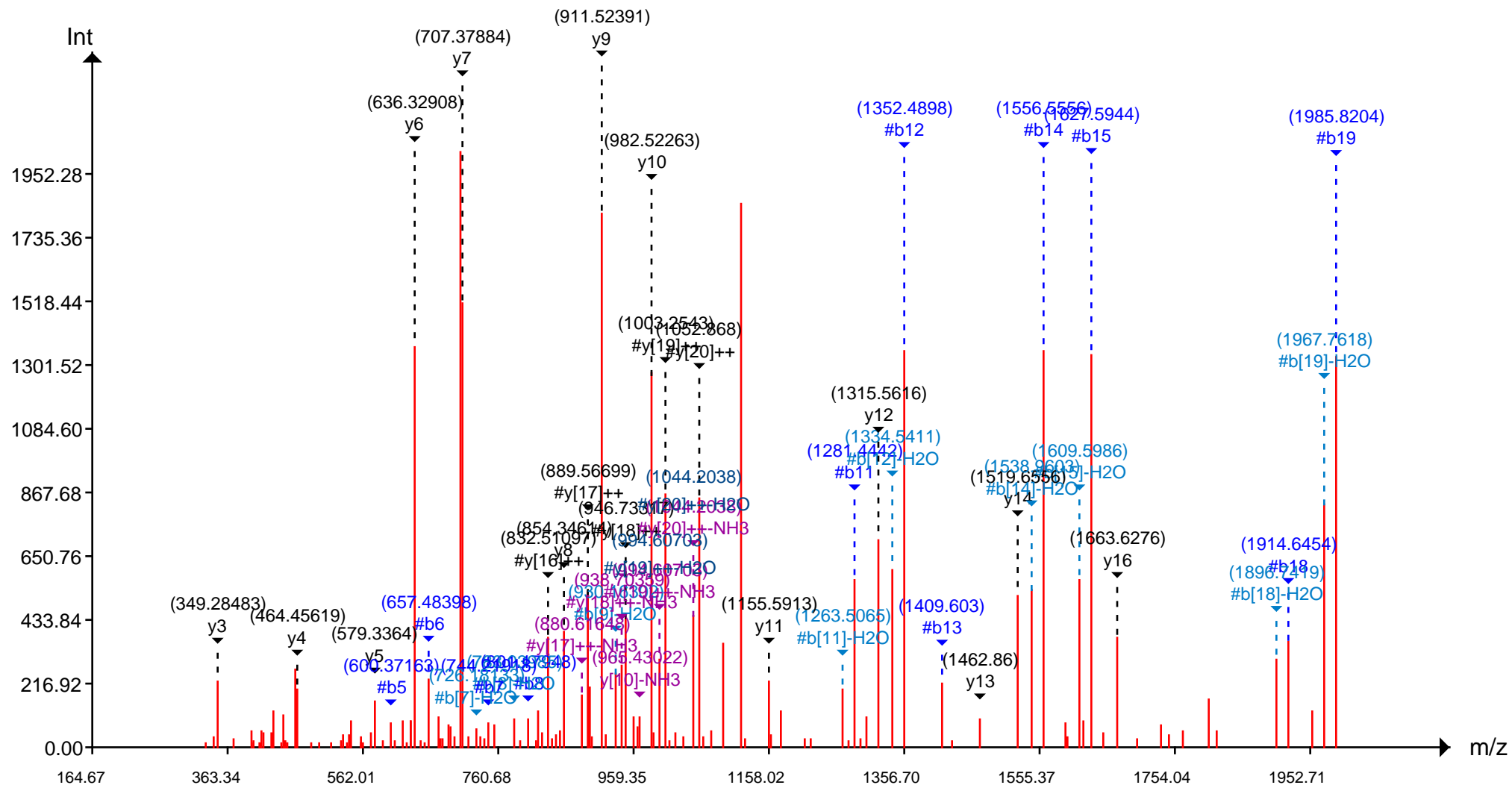
Info

Modified Sequence	AcD3-LEANK<AcD3K*>IGFEEK<AcD3K*>DIAANEENR<C13>-COOH	Ion score	99.45
Protein	Q9JJU8	Confidence level	99.00%
Precursor m/z	1216.1245	Identity Threshold	47.91
Charge	2+	Mass error	0.002484



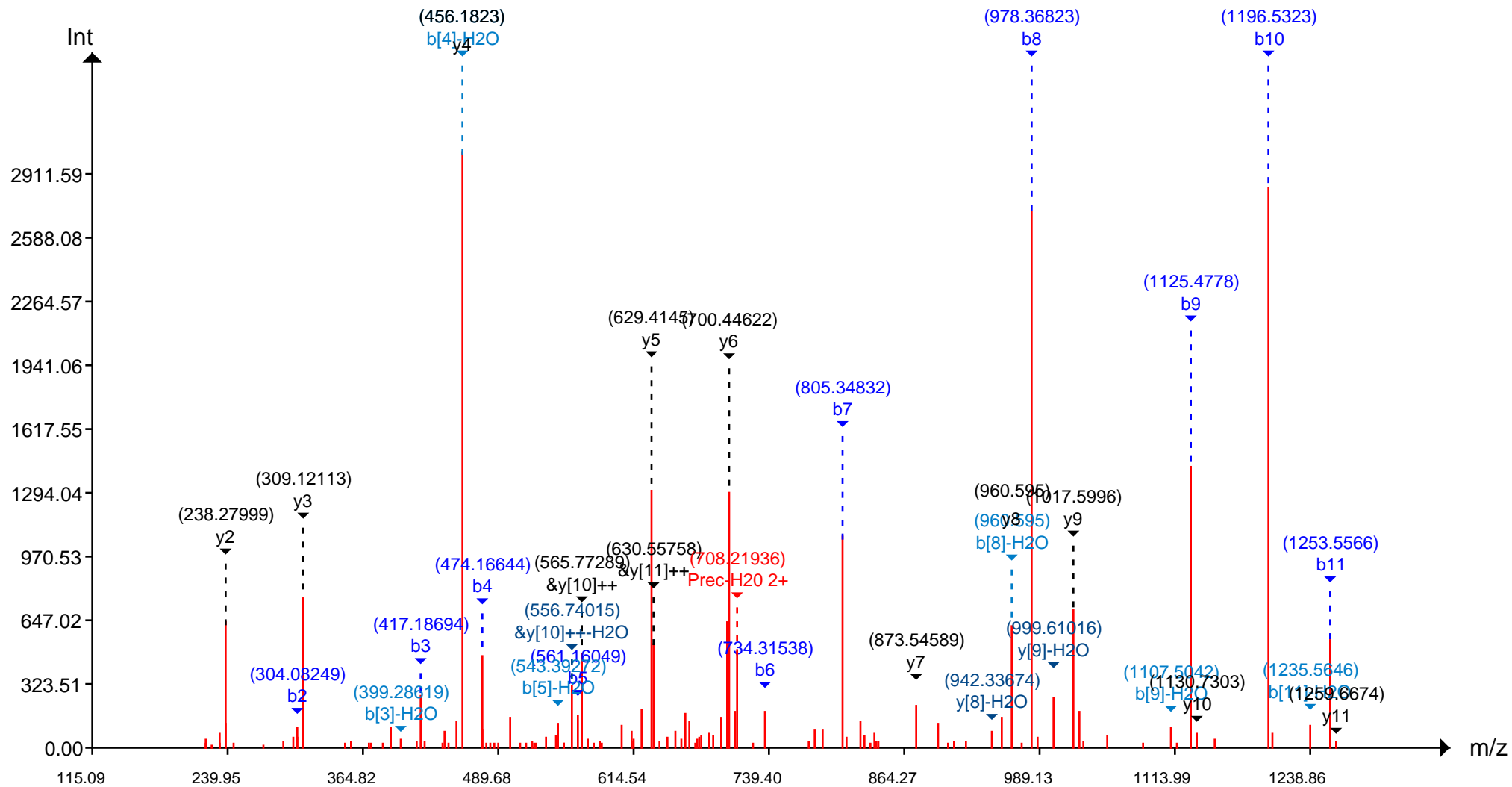
Info

Modified Sequence	AcD3-LVIDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH	Ion score	51.69
Protein	P63260	Confidence level	99.00%
Precursor m/z	755.02825	Identity Threshold	46.30
Charge	3+	Mass error	5.9E-4



Info

Modified Sequence	AcD3-LVIDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH	Ion score	99.25
Protein	P63260	Confidence level	99.00%
Precursor m/z	1132.039	Identity Threshold	46.38
Charge	2+	Mass error	0.001116



Info

Modified Sequence AcD3-EELGSK<AcD3K*>AK<AcD3K*>FAGR<C13>-COOH

Ion score 89.02

Protein P17182

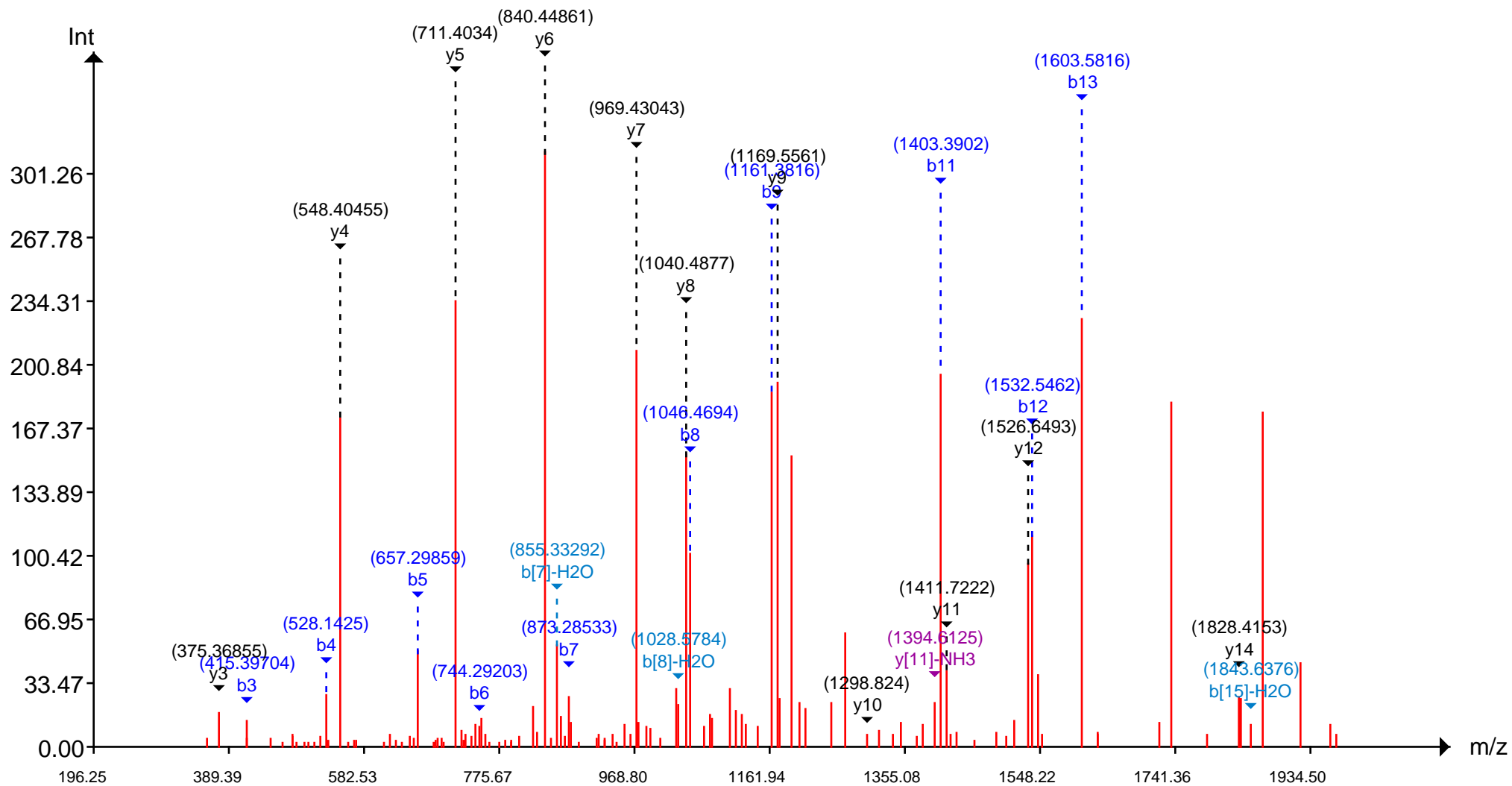
Confidence level 99.00%

Precursor m/z 717.40618

Identity Threshold 46.98

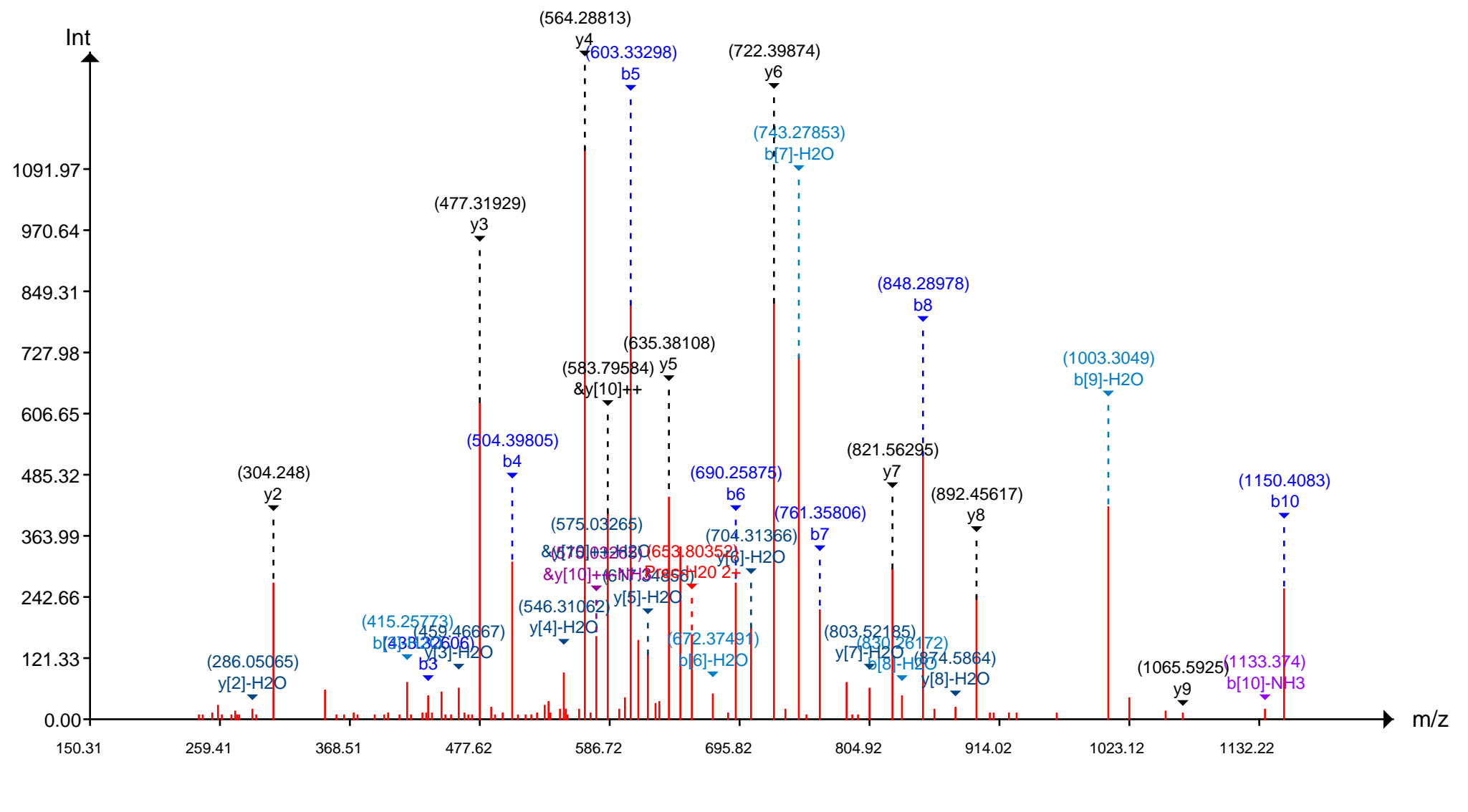
Charge 2+

Mass error 0.001055

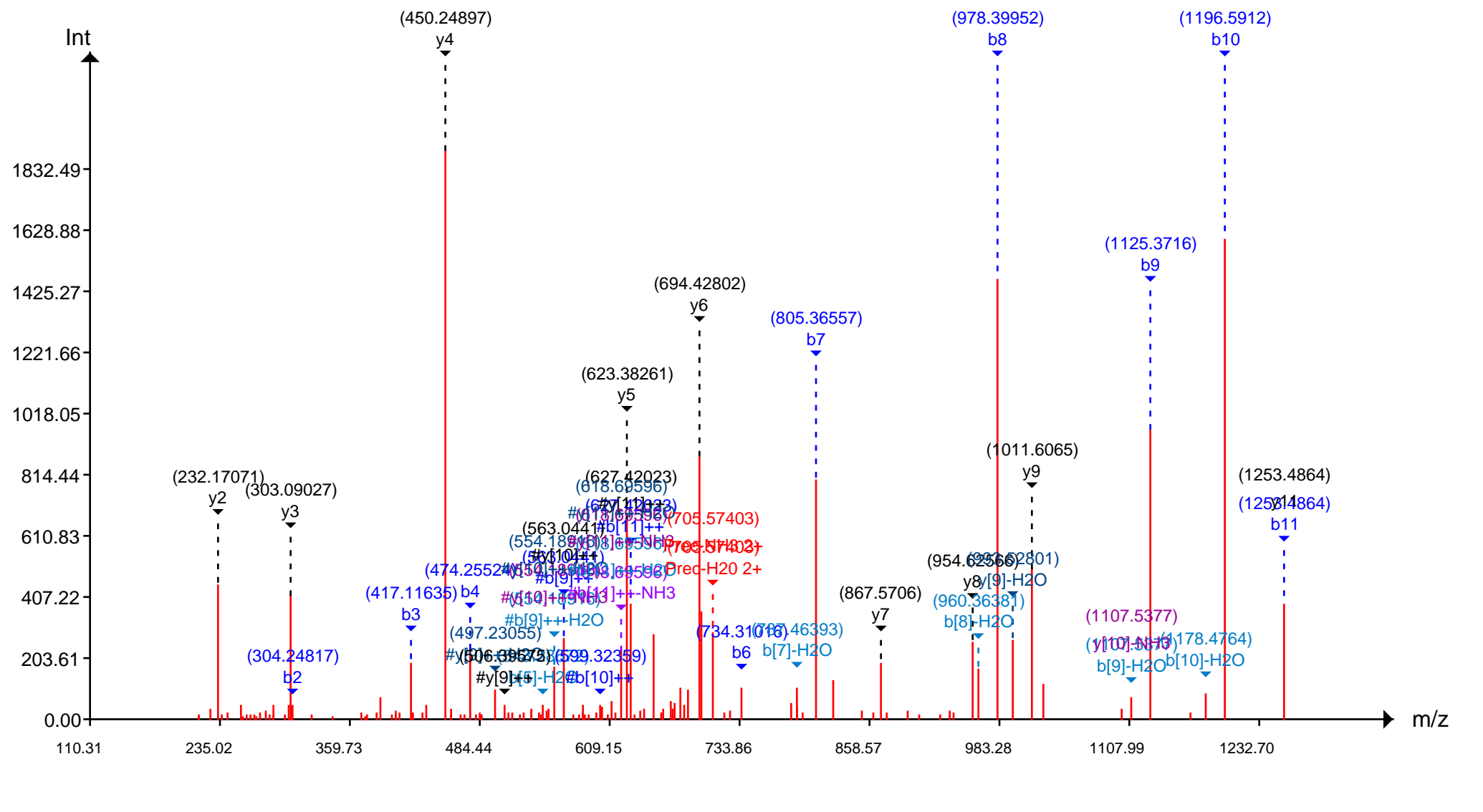


Info

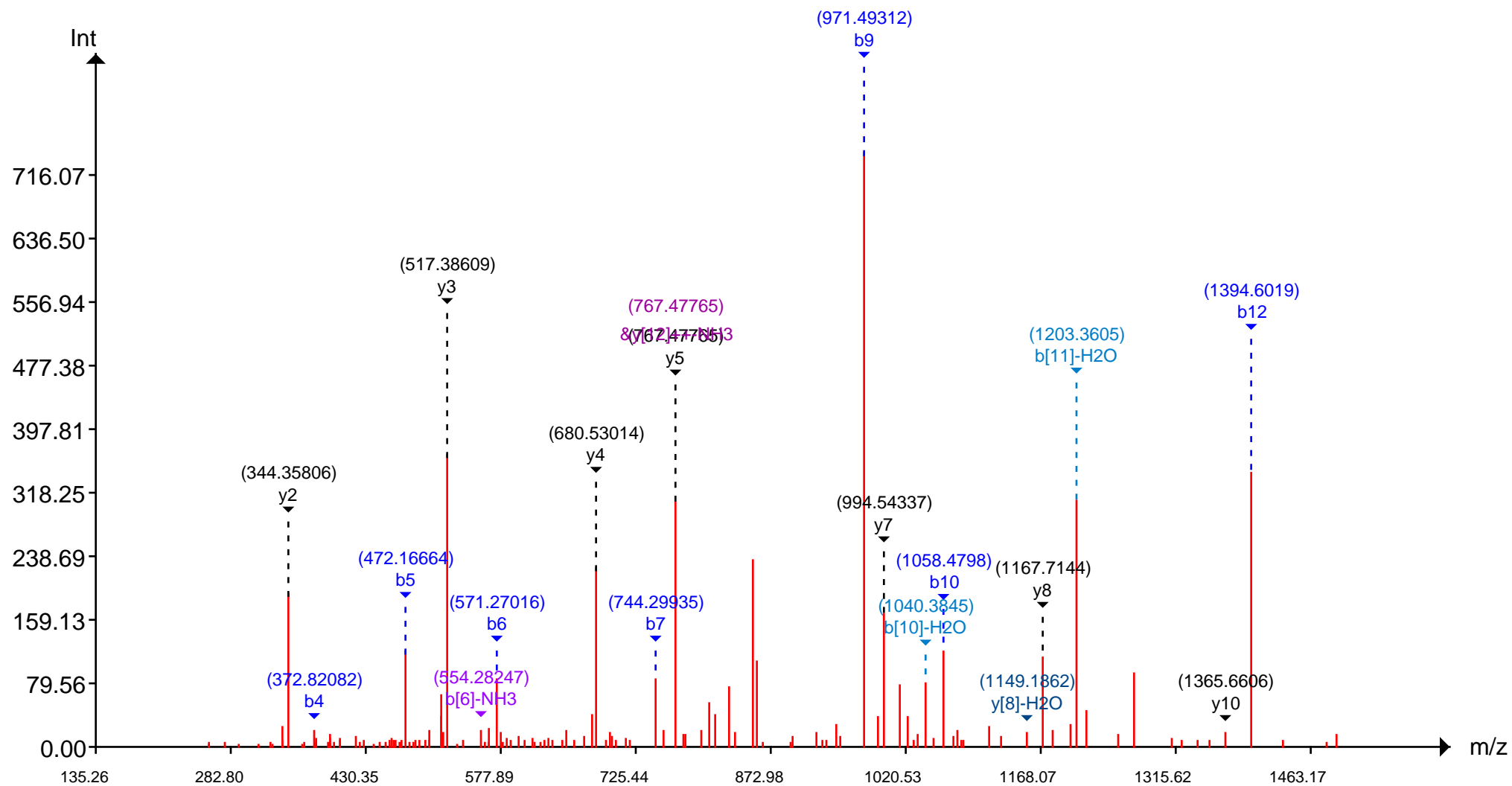
Modified Sequence	AcD3-QQILESEK<AcD3K*>DLEEAEEYK<AcD3K*>EAR-COOH	Ion score	66.04
Protein	P48428	Confidence level	99.00%
Precursor m/z	1286.6361	Identity Threshold	47.31
Charge	2+	Mass error	0.00391



Info			
Modified Sequence	AcD3-ITK<AcD3K*>AVSASK<AcD3K*>ER-COOH	Ion score	73.1
Protein	P43276	Confidence level	99.00%
Precursor m/z	662.8931	Identity Threshold	44.95
Charge	2+	Mass error	8.5E-4

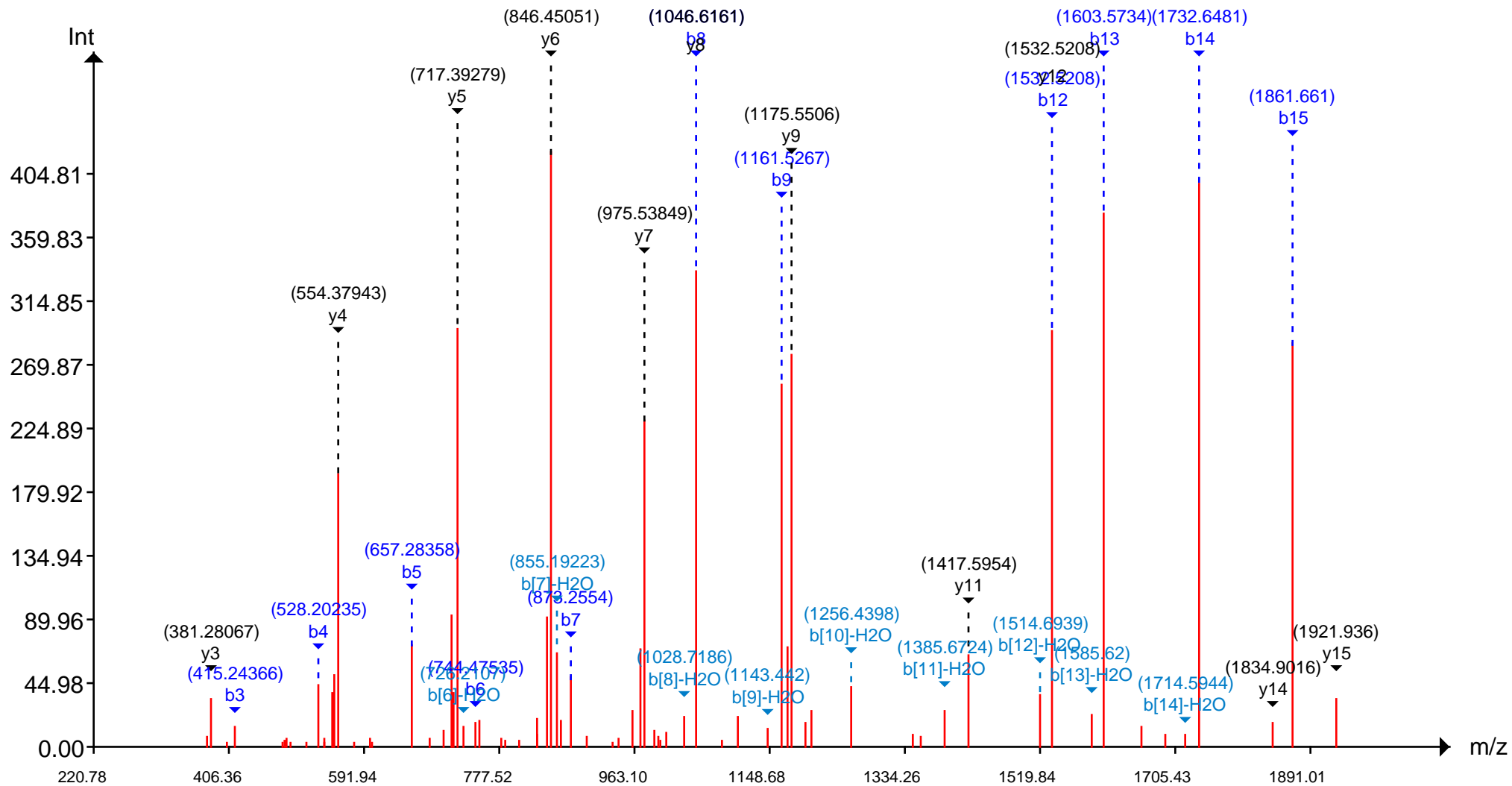


Info			
Modified Sequence	AcD3-EELGSK<AcD3K*>AK<AcD3K*>FAGR-COOH	Ion score	70.4
Protein	P17182	Confidence level	99.00%
Precursor m/z	714.39625	Identity Threshold	47.38
Charge	2+	Mass error	0.001337



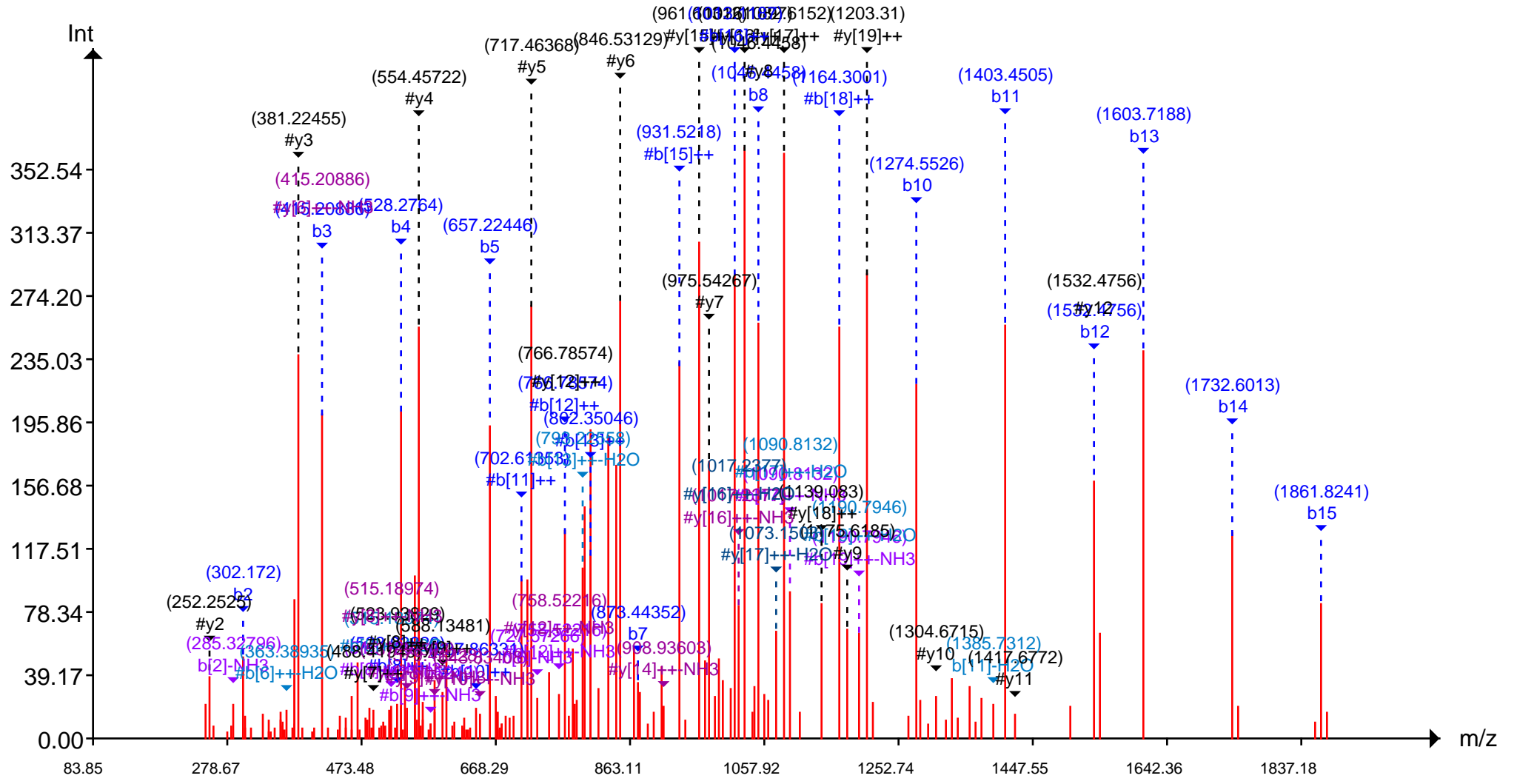
Info

Modified Sequence	AcD3-AAANVVK<AcD3K*>NISYK<AcD3K*>YR<C13>-COOH	Ion score	45.54
Protein	P28076, O35522	Confidence level	99.00%
Precursor m/z	869.50158	Identity Threshold	45.39
Charge	2+	Mass error	0.001944

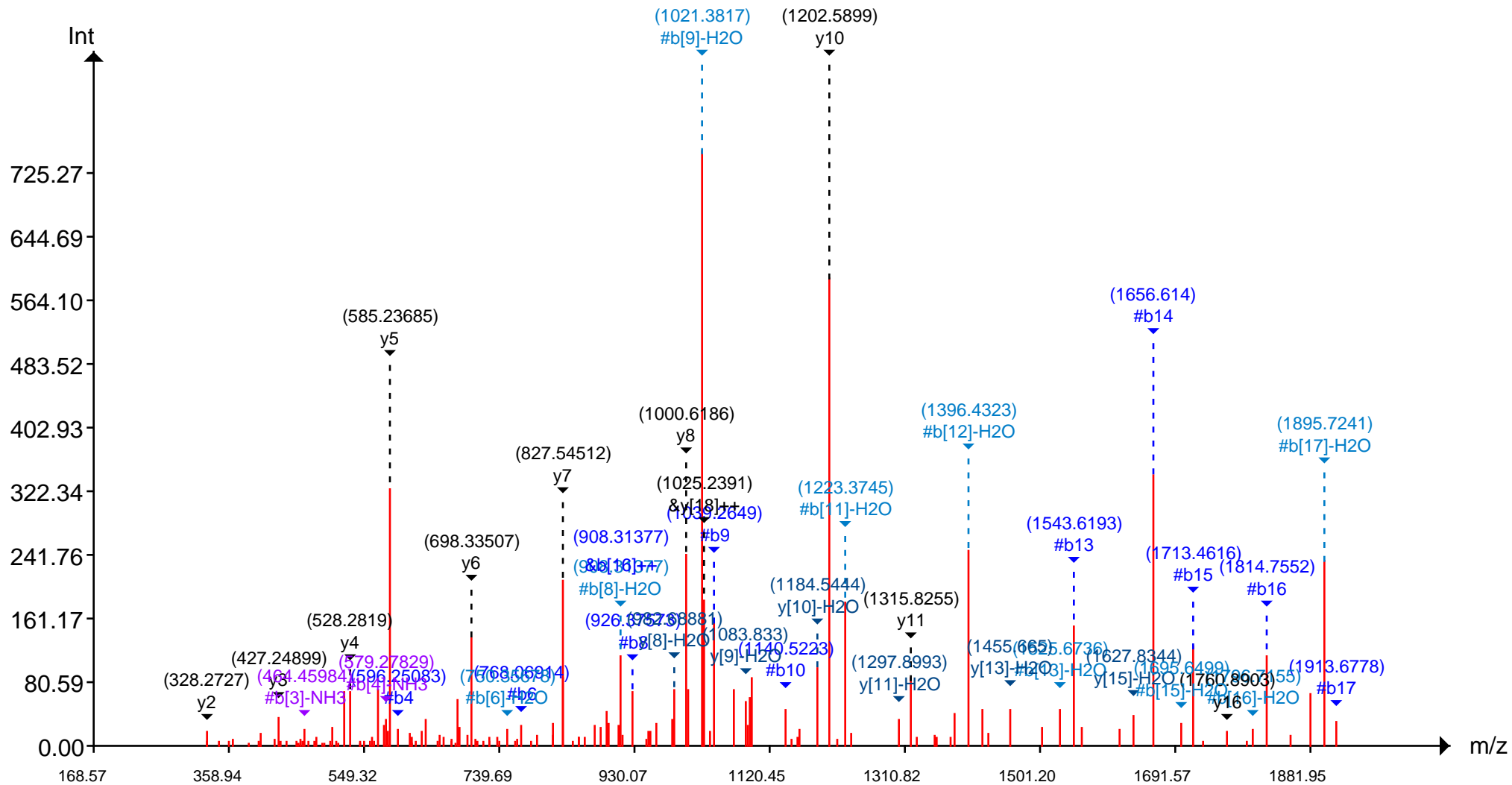


Info

Modified Sequence	AcD3-QQILESEK<AcD3K*>DLEEAEYK<AcD3K*>EAR<C13>-COOH	Ion score	78.26
Protein	P48428	Confidence level	99.00%
Precursor m/z	1289.6462	Identity Threshold	47.57
Charge	2+	Mass error	0.003968

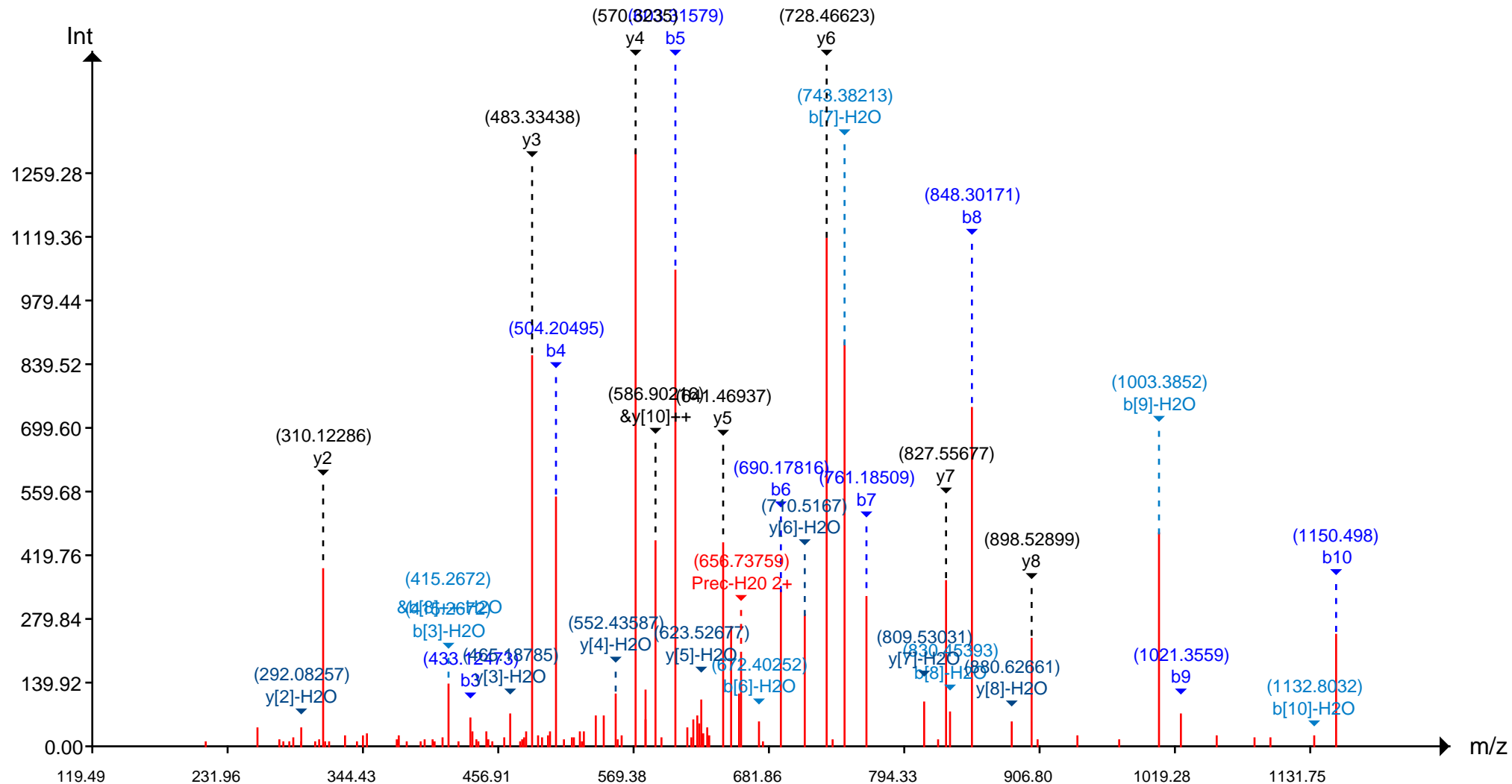


Info			
Modified Sequence	AcD3-QQILESEK<AcD3K*>DLEEAEYK<AcD3K*>EAR<C13>-COOH	Ion score	79.25
Protein	P48428	Confidence level	99.00%
Precursor m/z	860.09977	Identity Threshold	47.56
Charge	3+	Mass error	0.003602



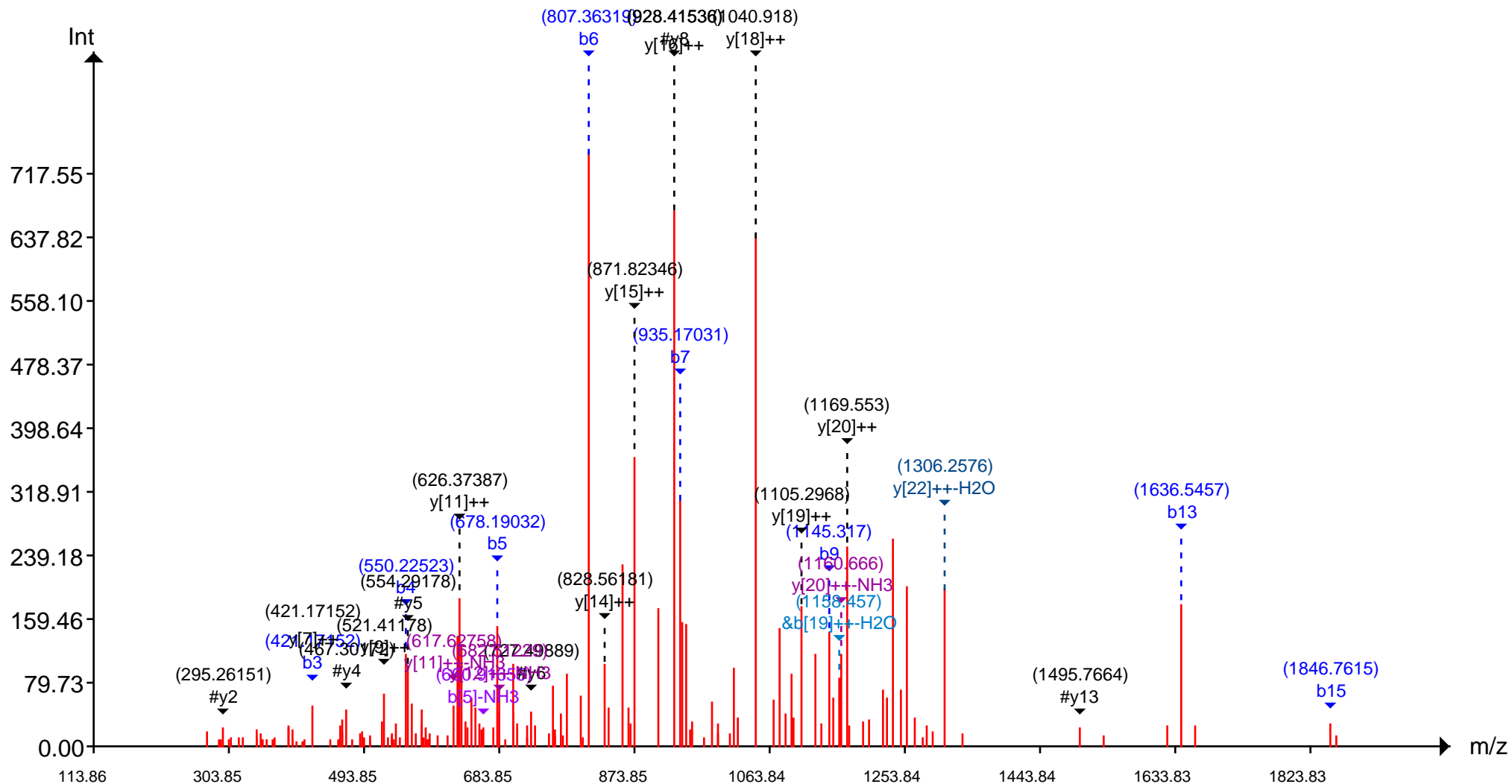
Info

Modified Sequence	AcD3-FDK<AcD3K*>DGDGTITTK<AcD3K*>ELGTVM<MoX*>R<C13>-COOH	Ion score	63.16
Protein	P62204	Confidence level	99.00%
Precursor m/z	1121.0728	Identity Threshold	47.96
Charge	2+	Mass error	0.002203



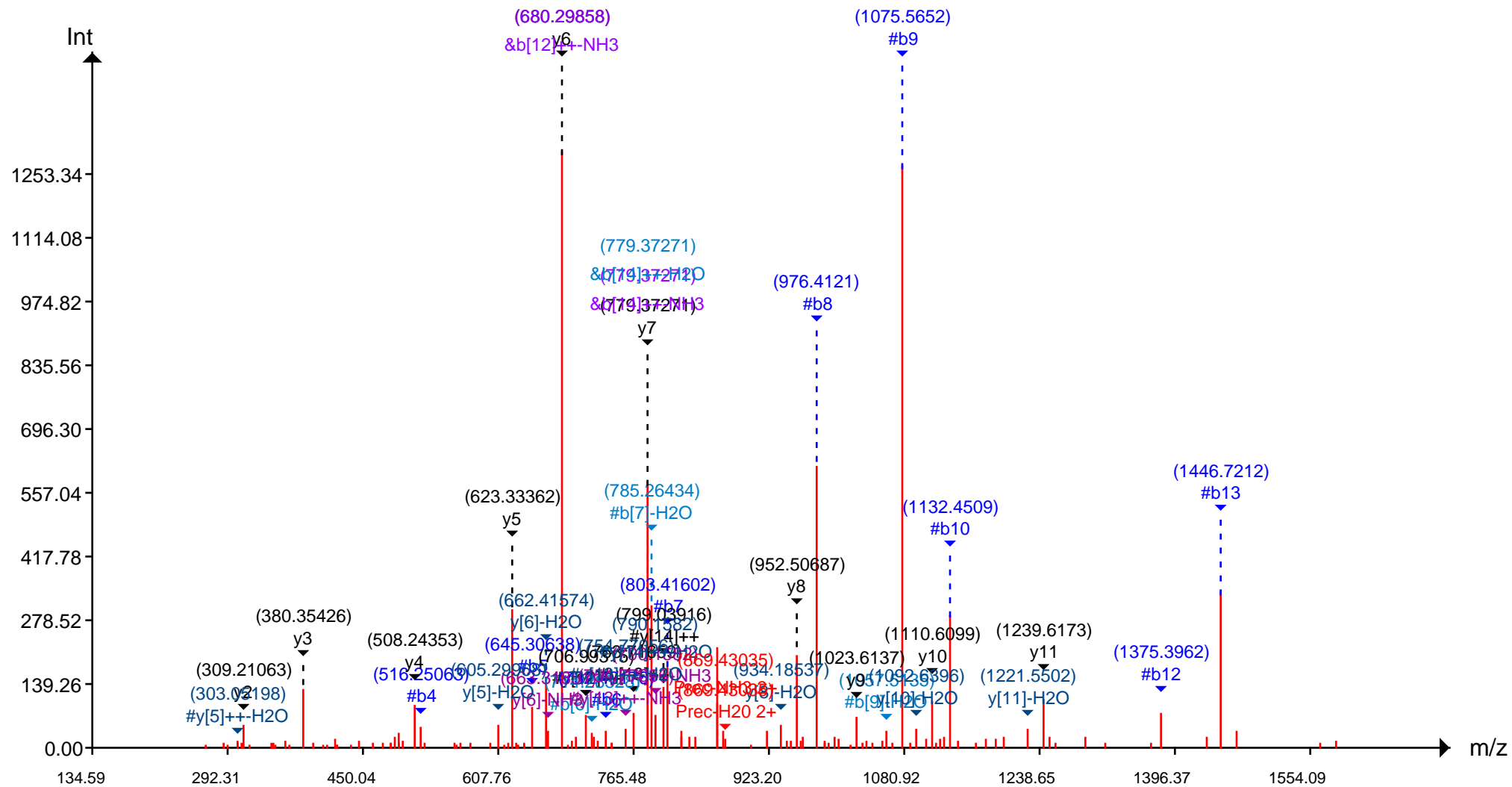
Info

Modified Sequence	AcD3-ITK<AcD3K*>AVSASK<AcD3K*>ER<C13>-COOH	Ion score	55.5
Protein	P43276	Confidence level	99.00%
Precursor m/z	665.90253	Identity Threshold	44.00
Charge	2+	Mass error	-4.31E-4



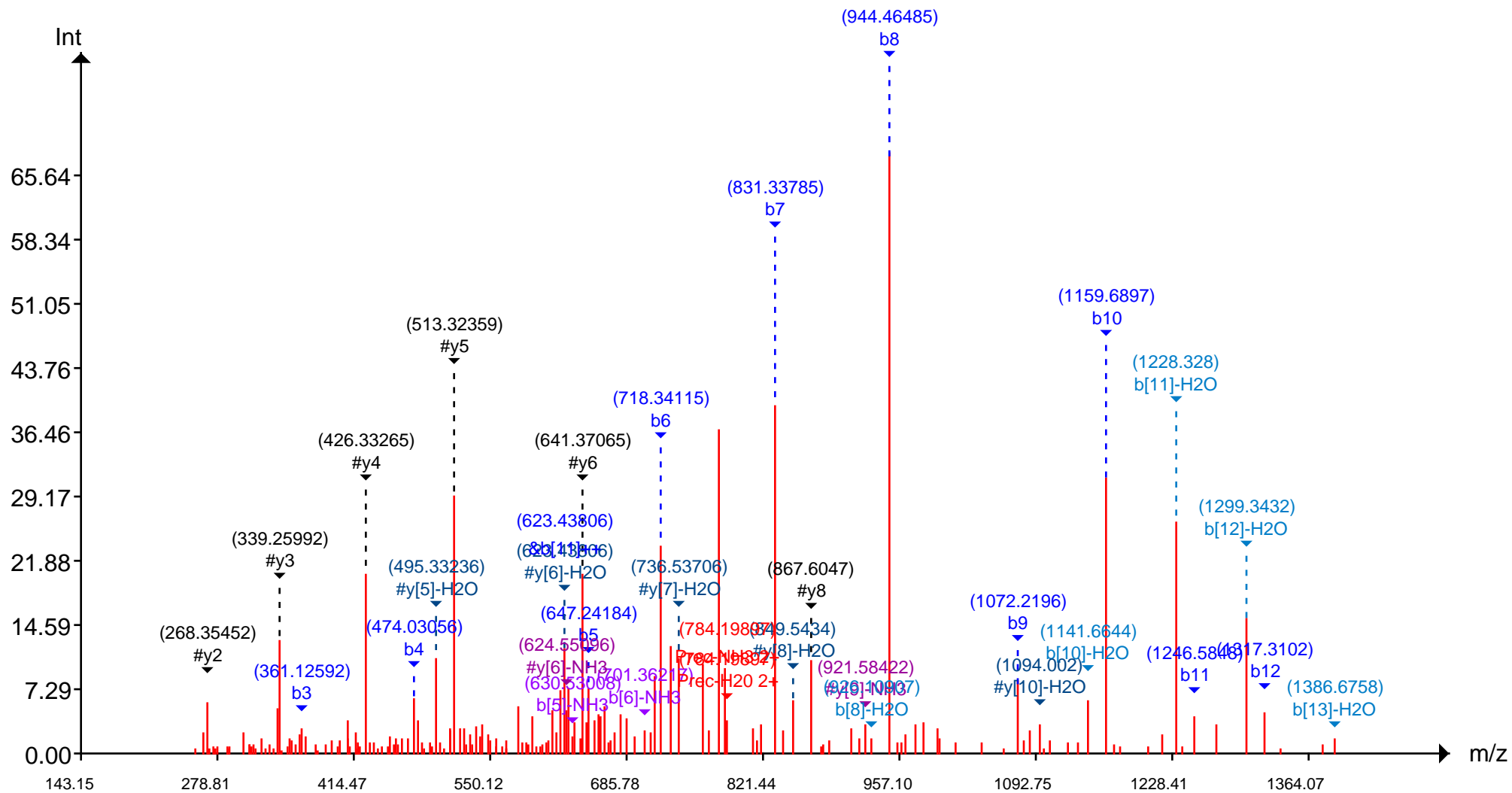
Info

Modified Sequence	AcD3-LVYEQEQPLSC<Cmm*>DEPILSNK<AcD3K*>SGDNR<C13>-COOH	Ion score	55.27
Protein	P70296	Confidence level	99.00%
Precursor m/z	963.13831	Identity Threshold	46.84
Charge	3+	Mass error	0.001165



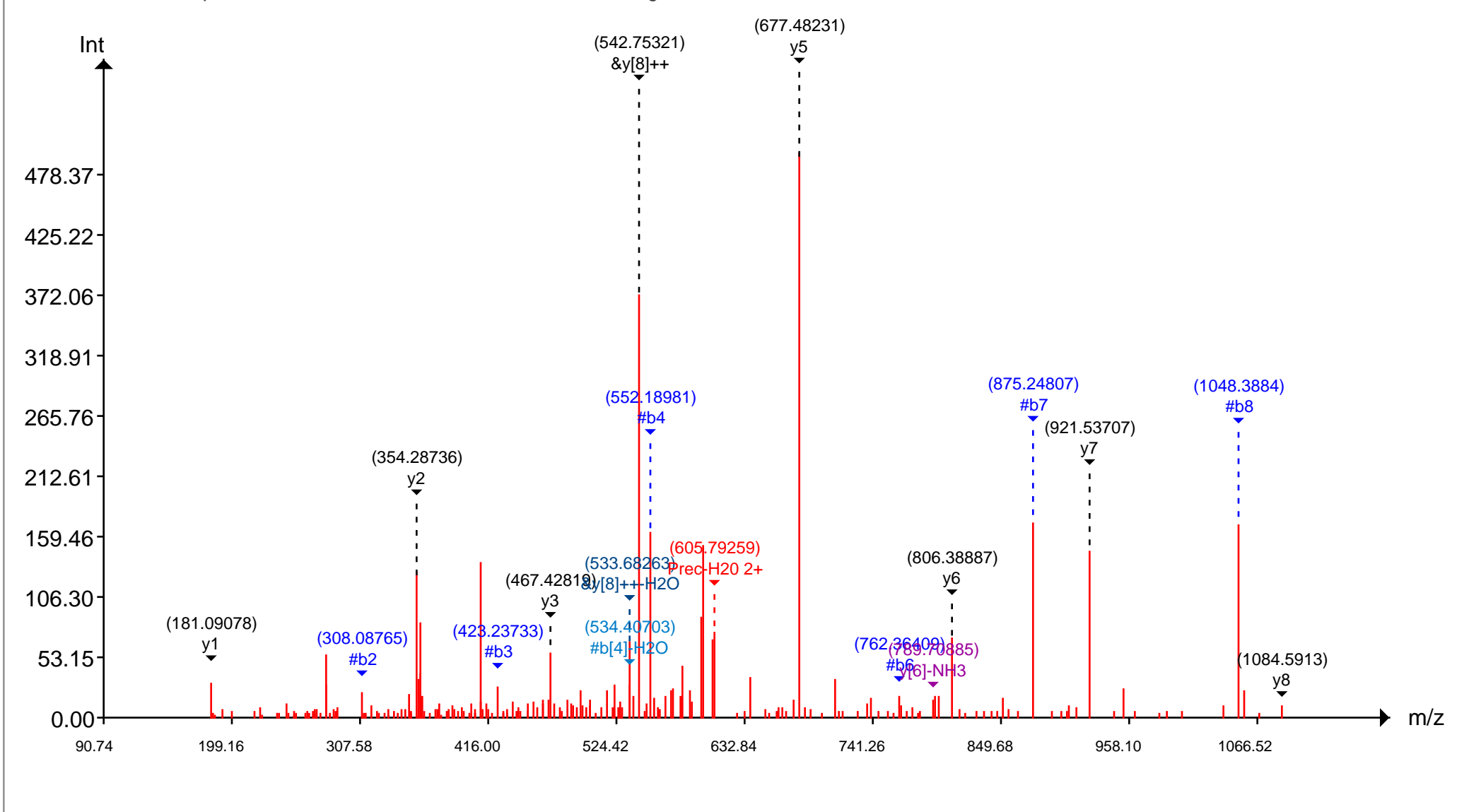
Info

Modified Sequence	AcD3-IAIK<AcD3K*>ESAK<AcD3K*>VGDQAQR<C13>-COOH	Ion score	75.97
Protein	Q99JY9	Confidence level	99.00%
Precursor m/z	878.00766	Identity Threshold	45.10
Charge	2+	Mass error	0.002797



Info

Modified Sequence	AcD3-DAELK<AcD3K*>ALLQSSASR<C13>-COOH	Ion score	83.06
Protein	P50580	Confidence level	99.00%
Precursor m/z	792.94463	Identity Threshold	47.04
Charge	2+	Mass error	0.001448



Info

Modified Sequence AcD3-VYDEPLLK<AcD3K*>R<C13>-COOH

Ion score 50.63

Protein A2A8Z1

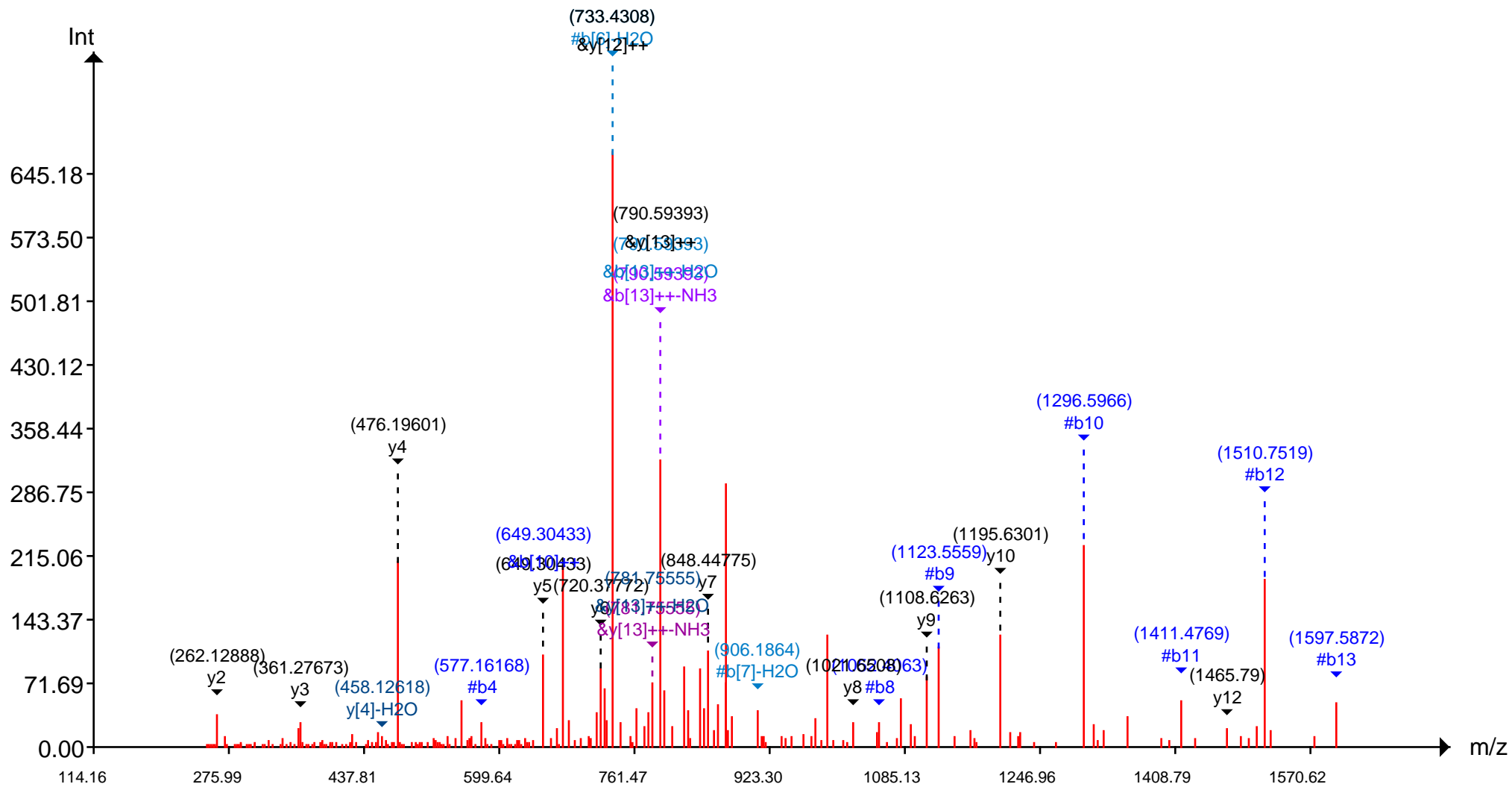
Confidence level 99.00%

Precursor m/z 614.86094

Identity Threshold 46.15

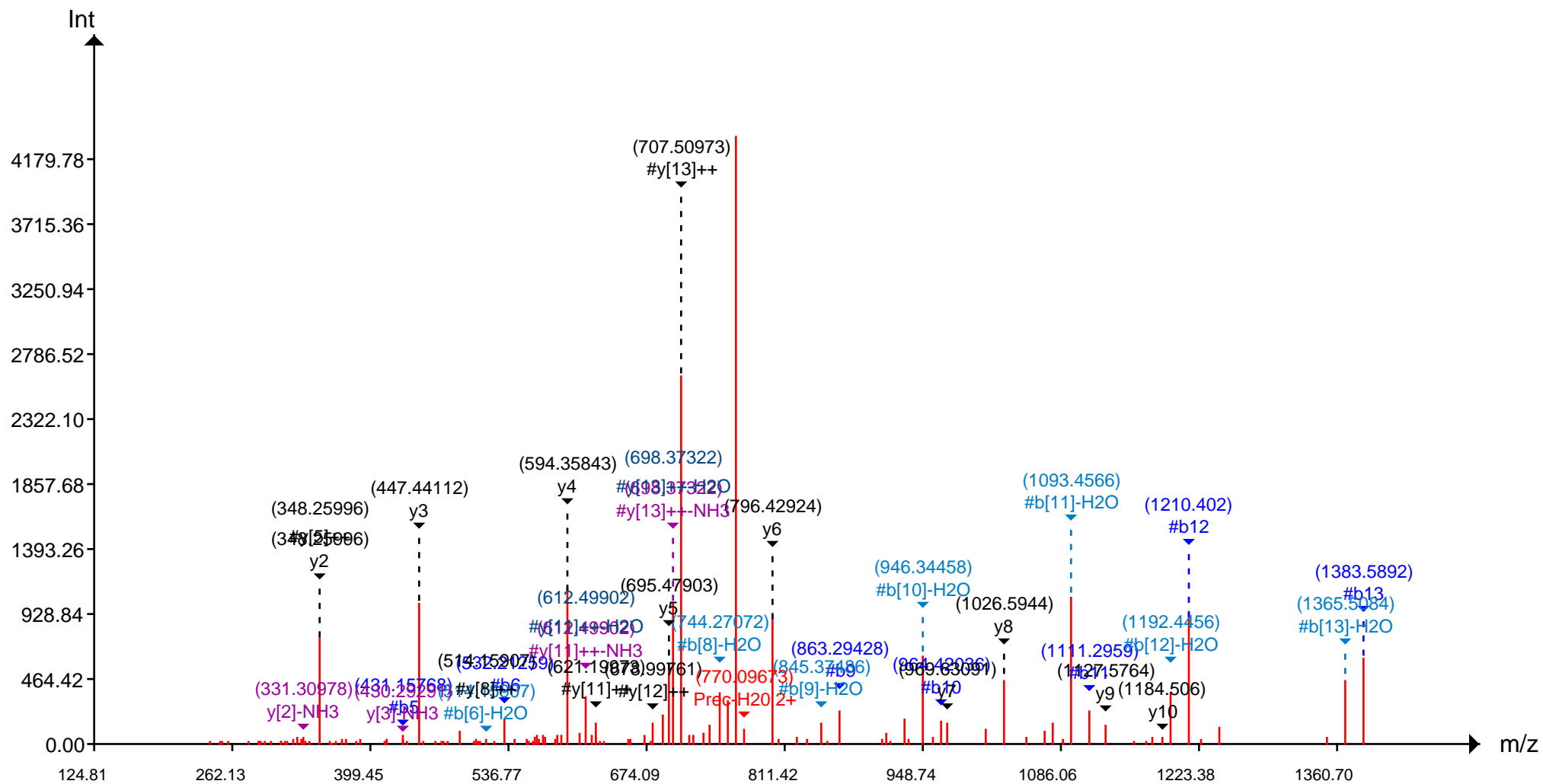
Charge 2+

Mass error -3.9E-4



Info

Modified Sequence	AcD3-FNPK<AcD3K*>SSK<AcD3K*>QAK<AcD3K*>DVSR-COOH	Ion score	48.68
Protein	O08585	Confidence level	99.00%
Precursor m/z	886.49105	Identity Threshold	47.22
Charge	2+	Mass error	0.00217



Info

Modified Sequence AcD3-VGDGGTgk<AcD3K*>TTFVK<AcD3K*>R-COOH

Ion score 62.85

Protein P62827

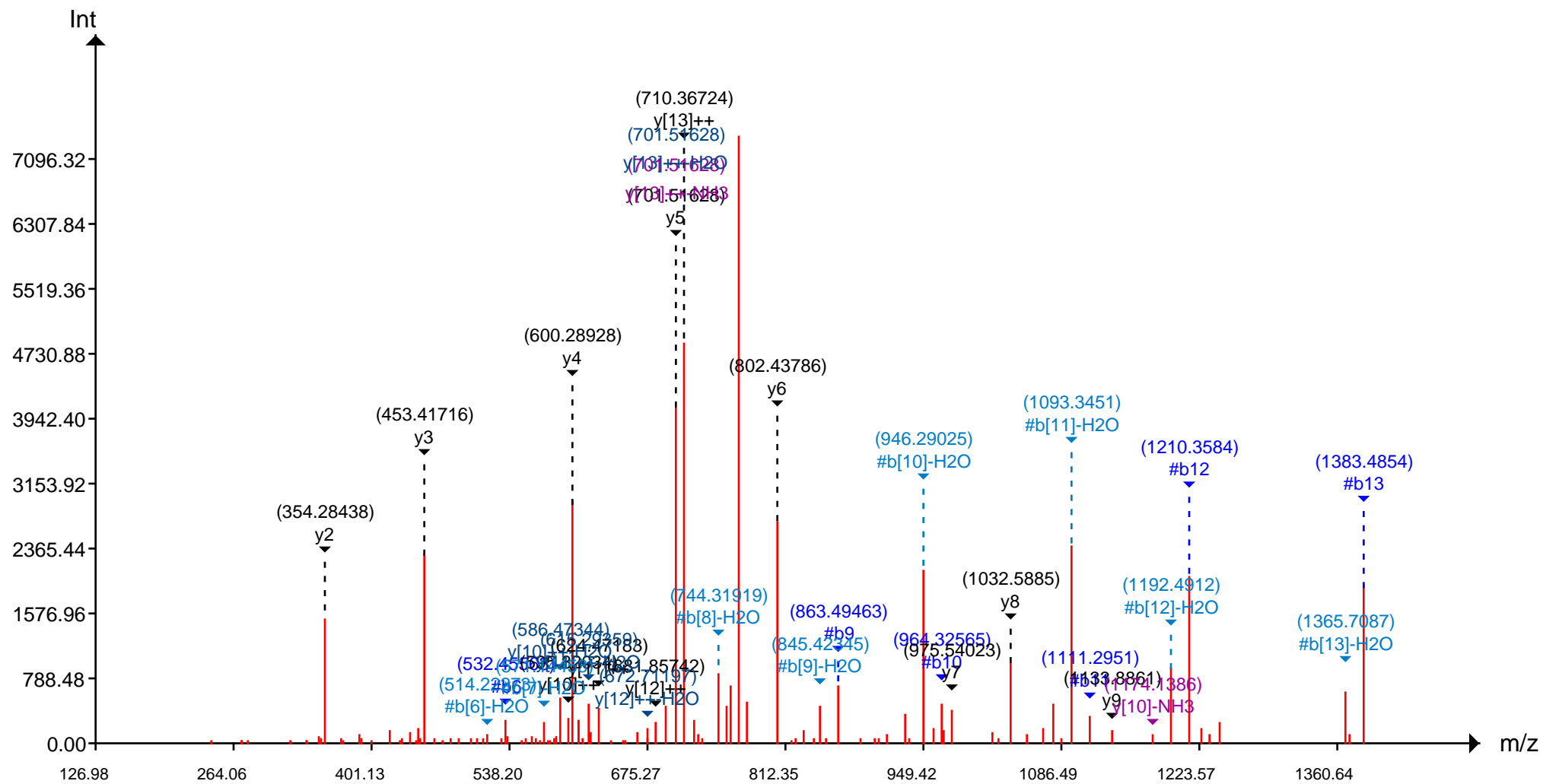
Confidence level 99.00%

Precursor m/z 779.43334

Identity Threshold 47.33

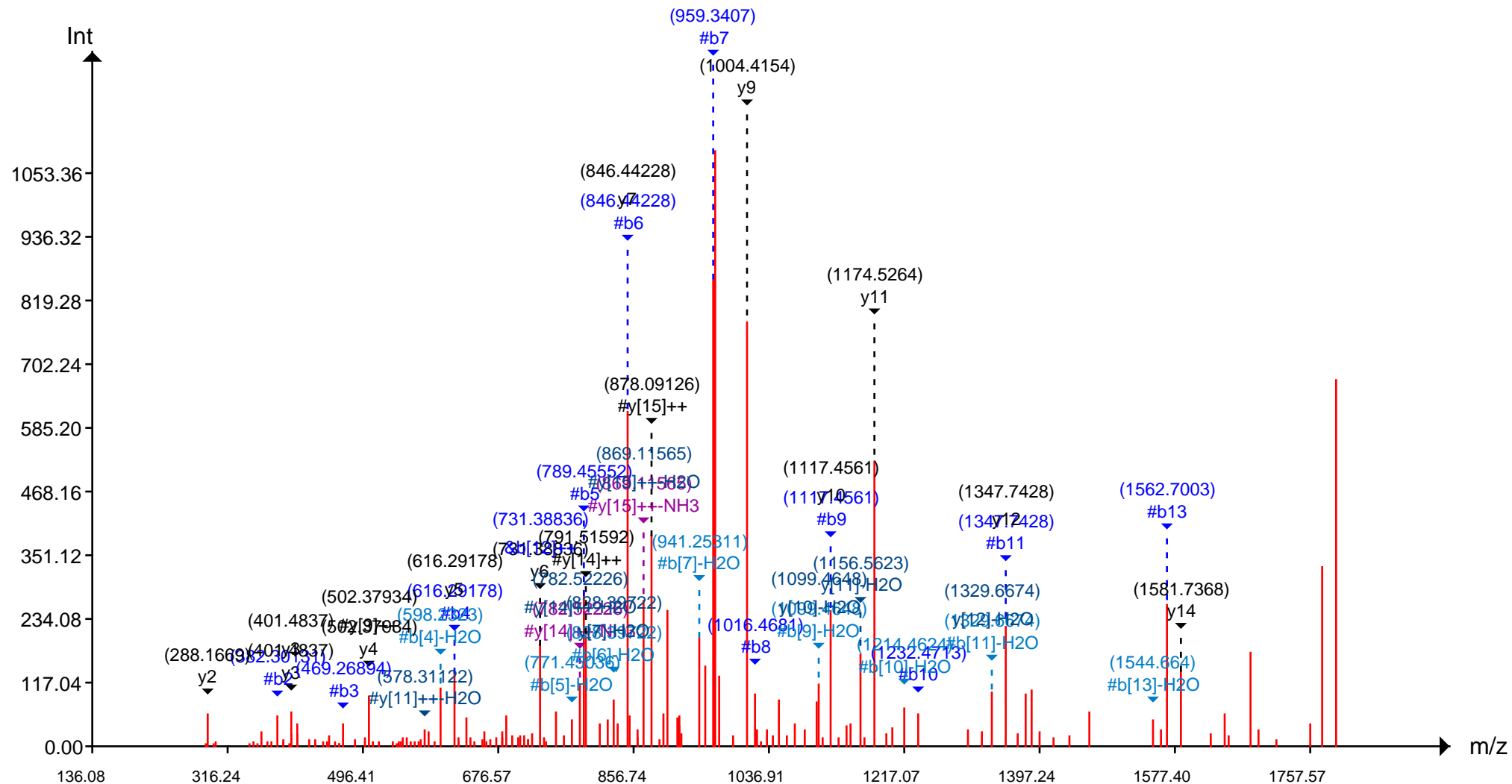
Charge 2+

Mass error 0.001222



Info

Modified Sequence	AcD3-VGDGGTGK<AcD3K*>TTFVK<AcD3K*>R<C13>-COOH	Ion score	50.84
Protein	P62827	Confidence level	99.00%
Precursor m/z	782.44285	Identity Threshold	46.93
Charge	2+	Mass error	1.0E-4



Info

Modified Sequence AcD3-YK<AcD3K*>SM<Mox*>K<AcD3K*>GLGTDDNTLIR-COOH

Ion score 84.92

Protein P97429

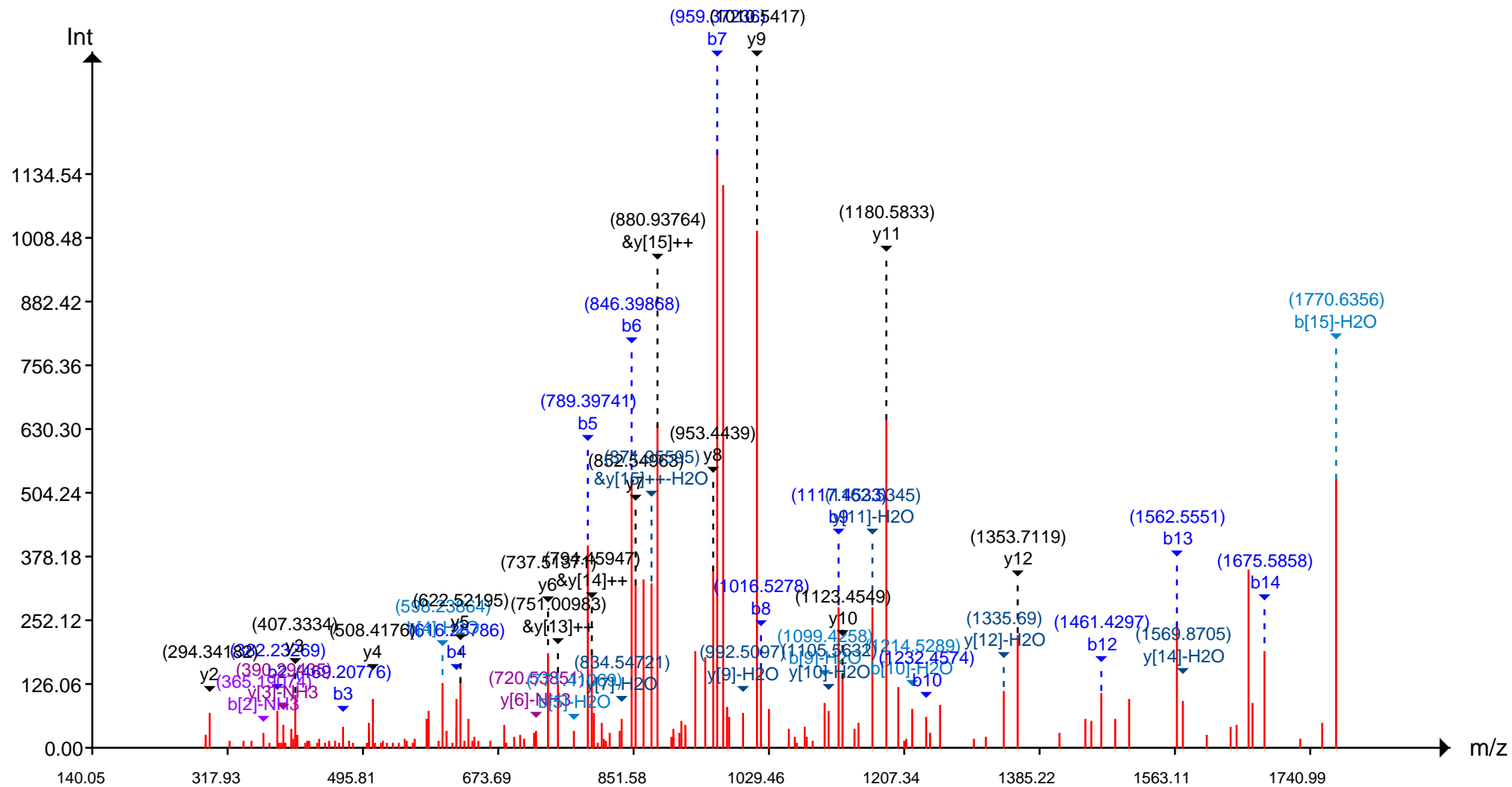
Confidence level 99.00%

Precursor m/z 982.01251

Identity Threshold 47.94

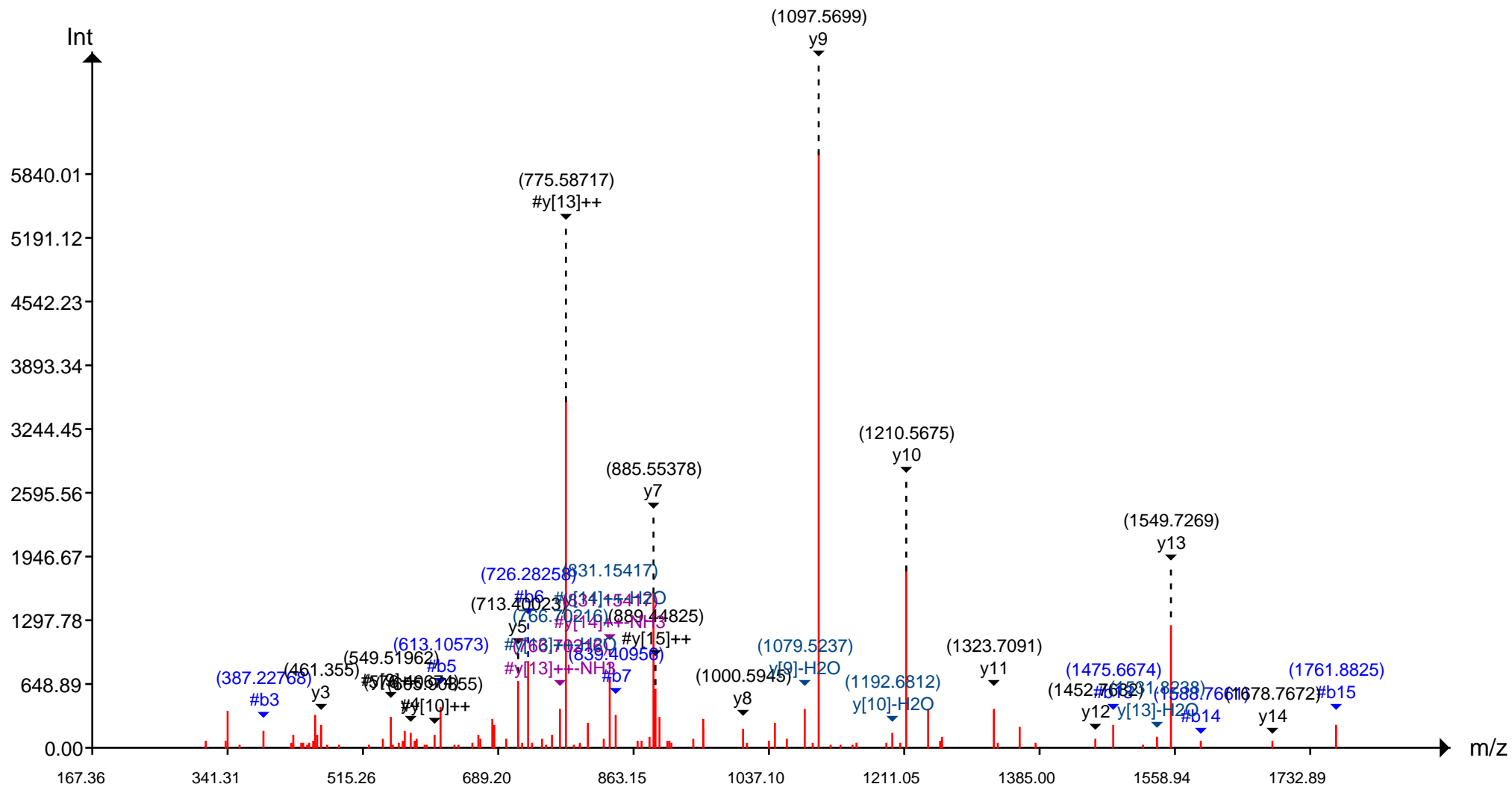
Charge 2+

Mass error 0.00264



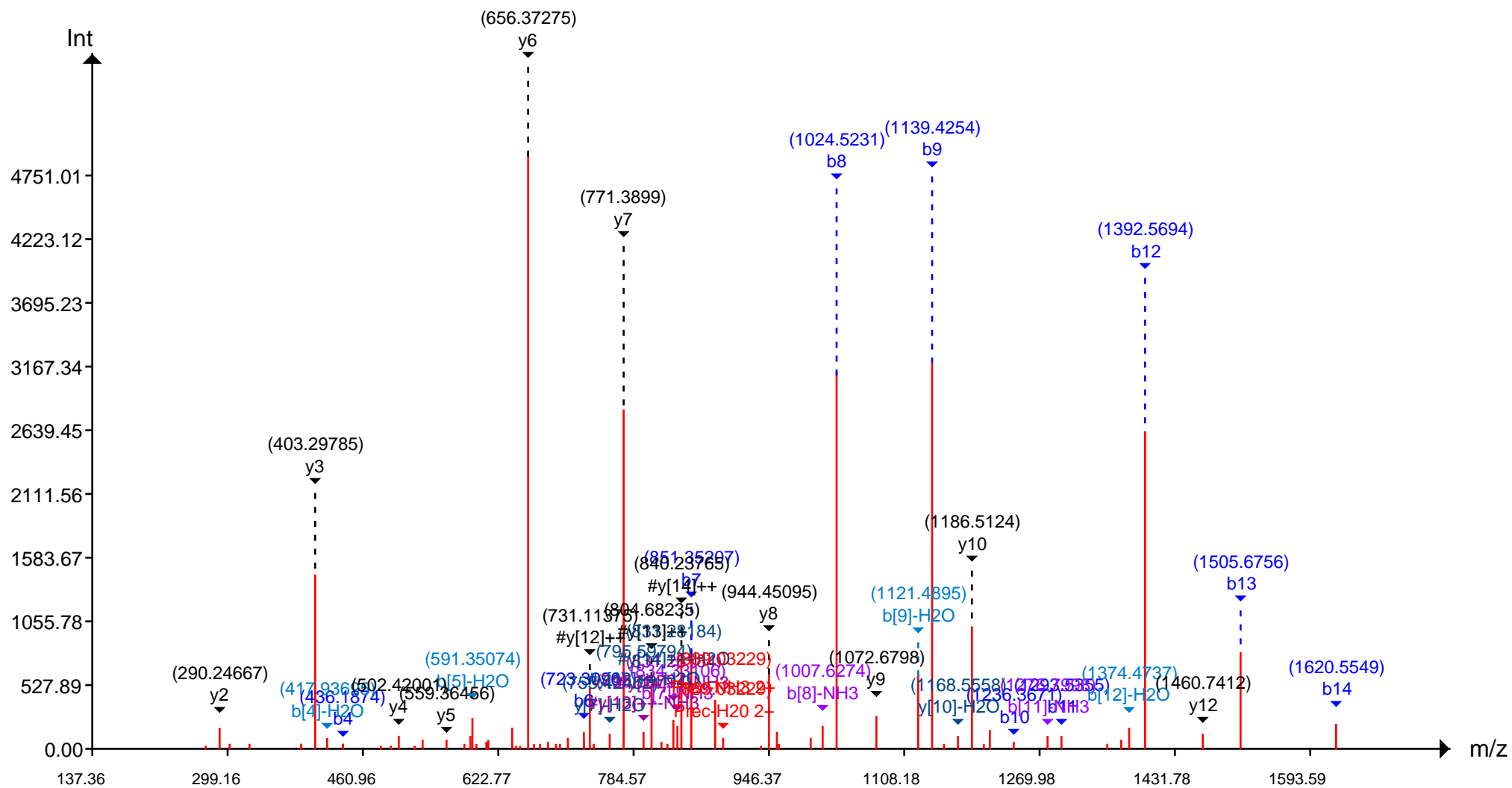
Info

Modified Sequence	AcD3-YK<AcD3K*>SM<Mox*>K<AcD3K*>GLGTTDNTLIR<C13>-COOH	Ion score	81.29
Protein	P97429	Confidence level	99.00%
Precursor m/z	985.02256	Identity Threshold	47.82
Charge	2+	Mass error	0.002599



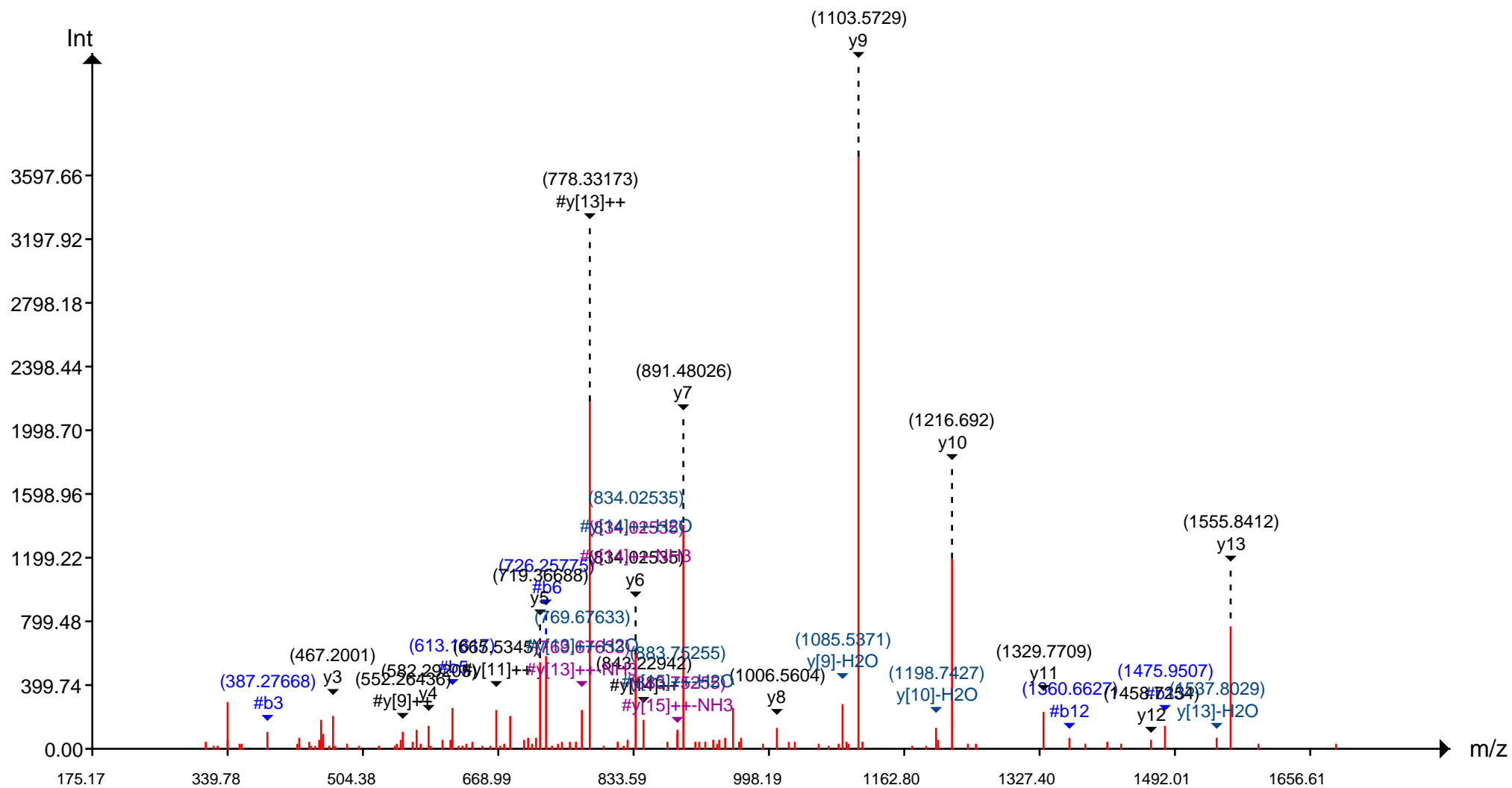
Info

Modified Sequence	AcD3-IVEPEILPDGDHDLK<AcD3K*>R-COOH	Ion score	76.01
Protein	P05064, P05063	Confidence level	99.00%
Precursor m/z	968.51973	Identity Threshold	47.79
Charge	2+	Mass error	0.002905



Info

Modified Sequence	AcD3-AAFTK<AcD3K*>NQK<AcD3K*>DPGVLDK-COOH	Ion score	94.4
Protein	P50543	Confidence level	99.00%
Precursor m/z	897.98929	Identity Threshold	47.71
Charge	2+	Mass error	0.001824



Info

Modified Sequence AcD3-IVEPEILPDGDHDLK<AcD3K*>R<C13>-COOH

Ion score 81.82

Protein P05064, P05063

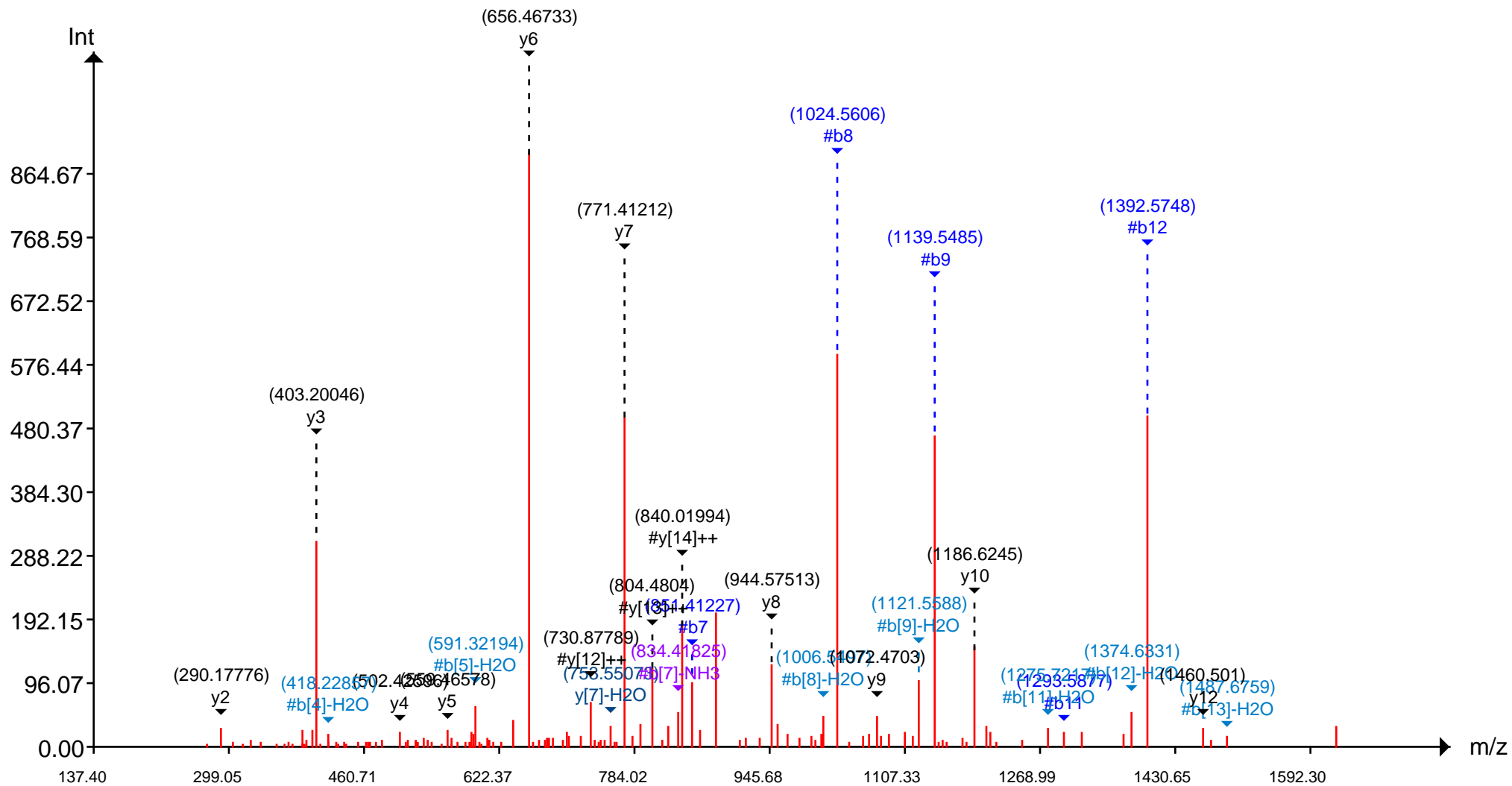
Confidence level 99.00%

Precursor m/z 971.52929

Identity Threshold 47.46

Charge 2+

Mass error 0.001883



Info

Modified Sequence AcD3-AAFTK<AcD3K*>NQK<AcD3K*>DPGVLDK-COOH

Ion score 74.18

Protein P50543

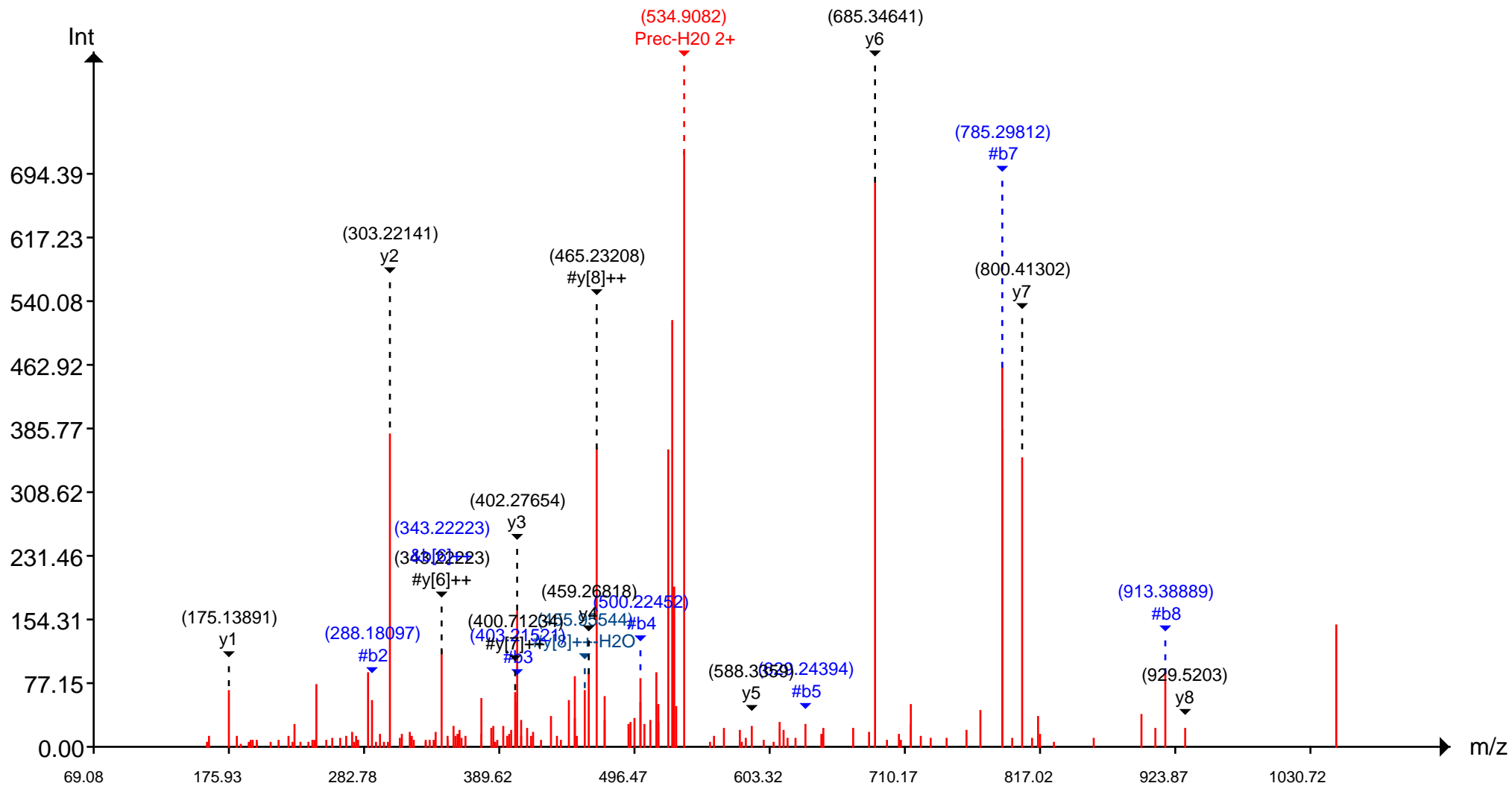
Confidence level 99.00%

Precursor m/z 897.98942

Identity Threshold 47.66

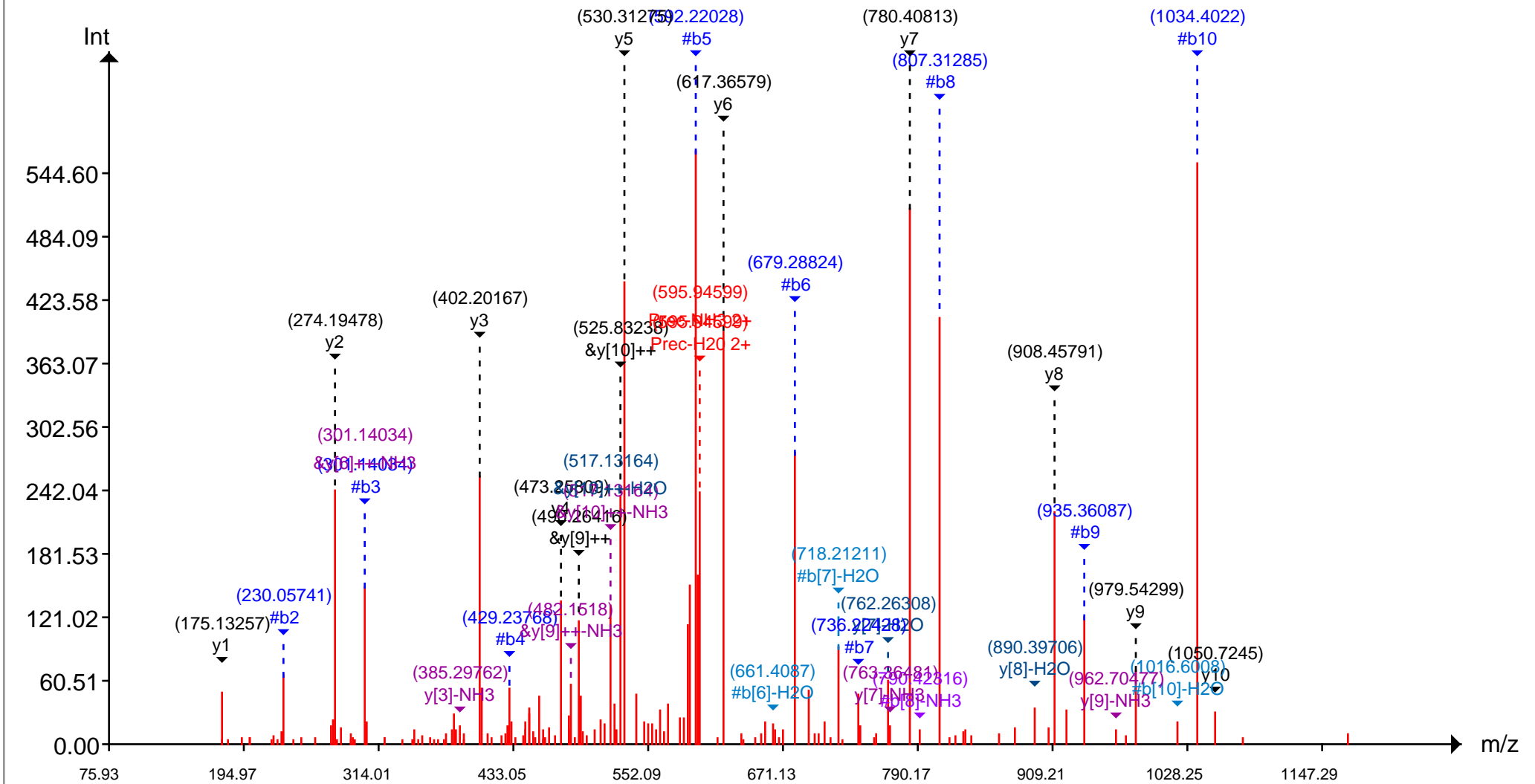
Charge 2+

Mass error 0.002084



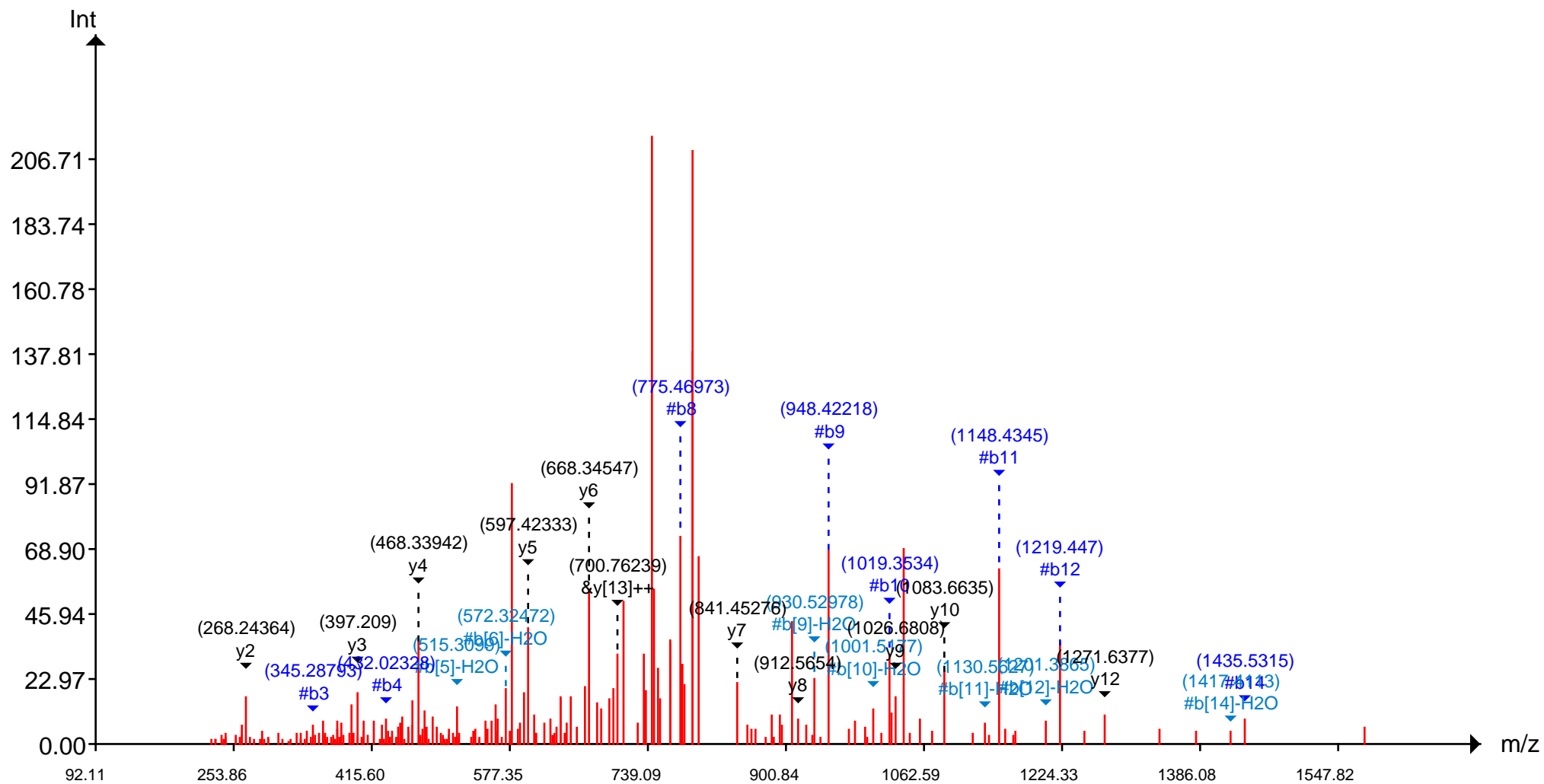
Info

Modified Sequence	AcD3-LEDPEGVQR-COOH	Ion score	47.96
Protein	Q9CQE5	Confidence level	99.00%
Precursor m/z	544.27656	Identity Threshold	47.33
Charge	2+	Mass error	1.01E-4



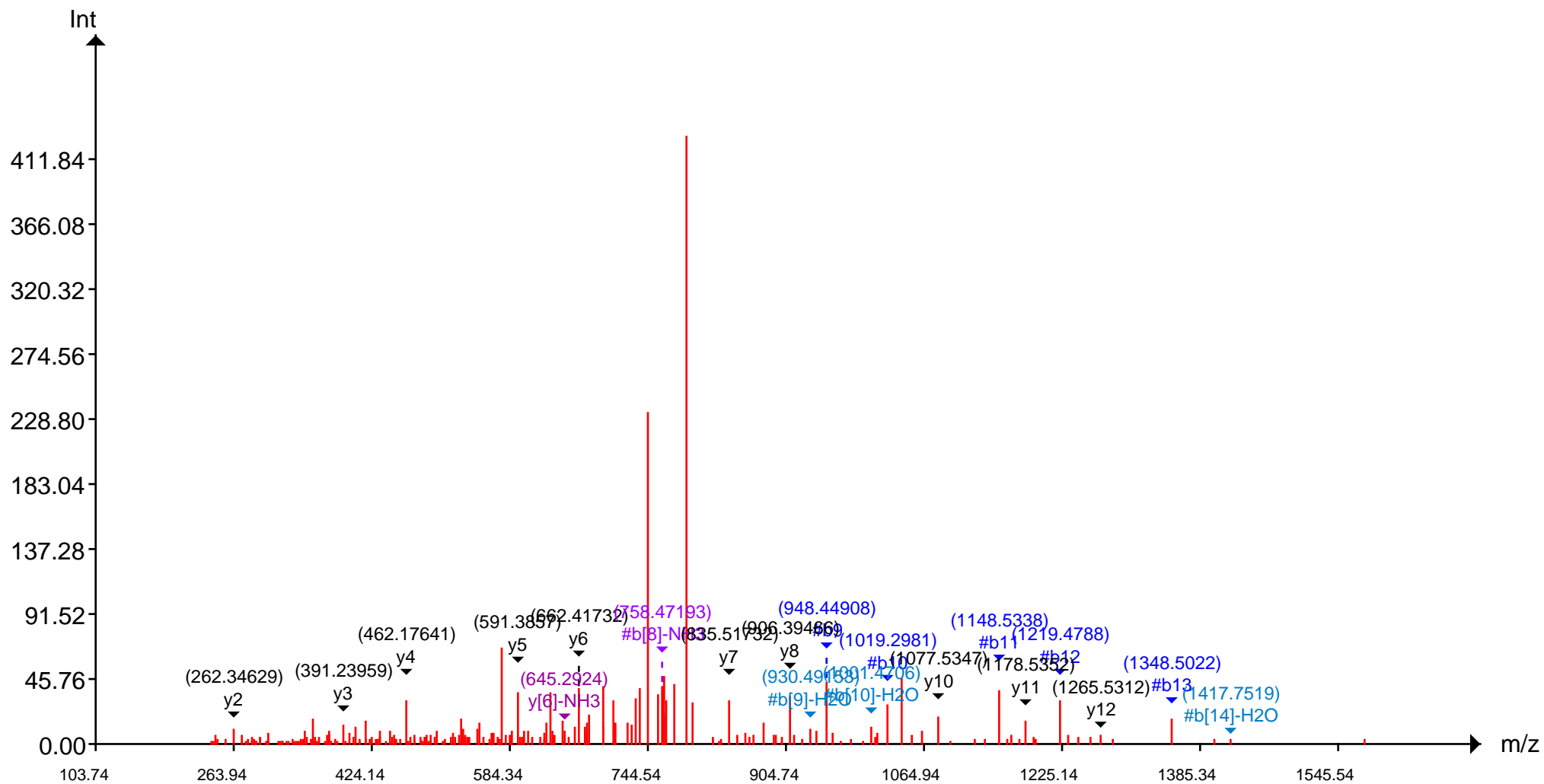
Info

Modified Sequence	AcD3-IAAQYSGAQVR-COOH	Ion score	96.24
Protein	Q9D8N0	Confidence level	99.00%
Precursor m/z	604.82681	Identity Threshold	48.20
Charge	2+	Mass error	2.28E-4



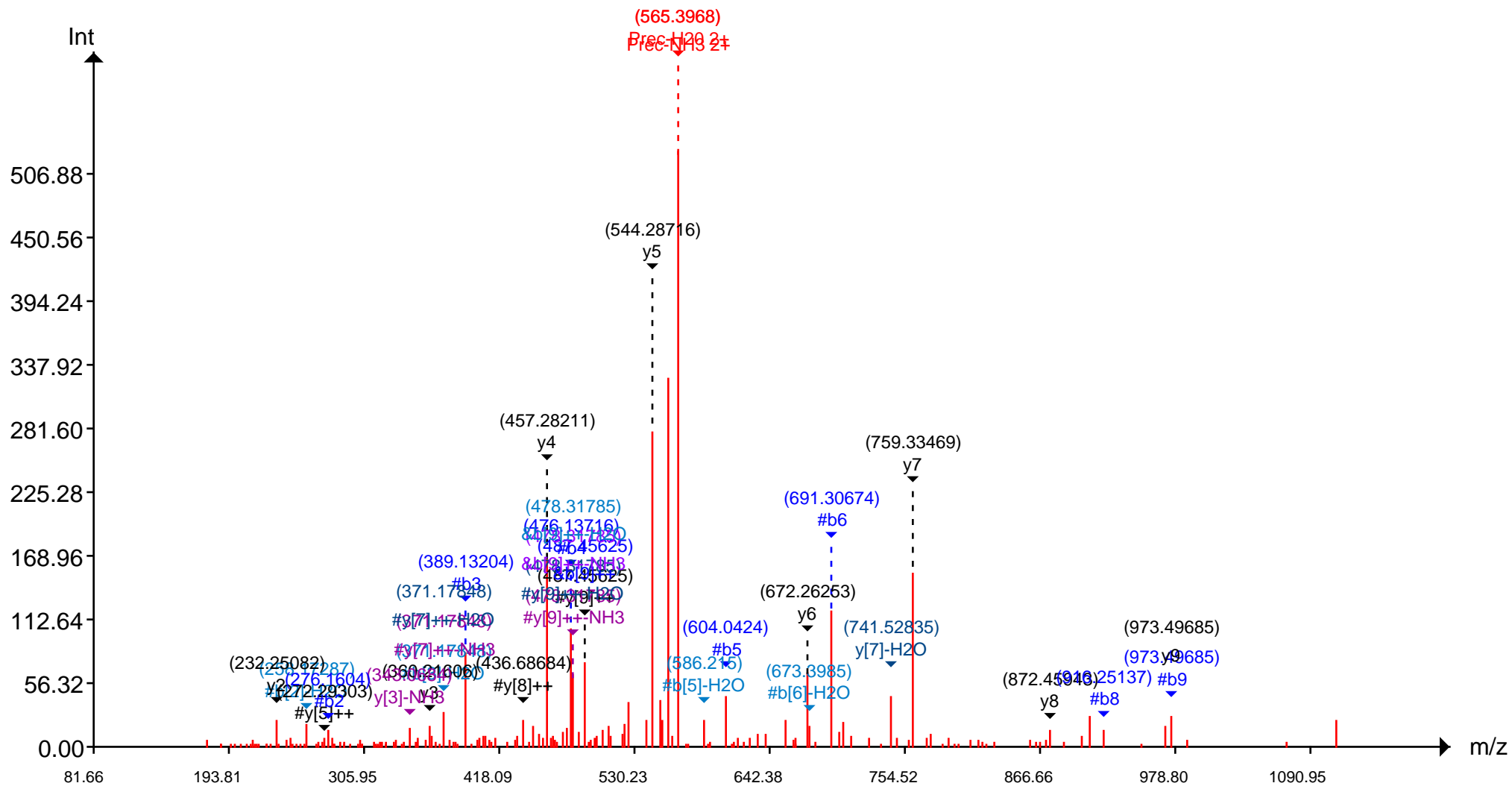
Info

Modified Sequence	AcD3-AVESTGNAK<AcD3K*>AEAESR<C13>-COOH	Ion score	50.51
Protein	Q9EQK5	Confidence level	99.00%
Precursor m/z	808.4108	Identity Threshold	47.89
Charge	2+	Mass error	7.43E-4



Info

Modified Sequence	AcD3-AVESTGNAK<AcD3K*>AEAESR-COOH	Ion score	51.16
Protein	Q9EQK5	Confidence level	99.00%
Precursor m/z	805.40093	Identity Threshold	47.59
Charge	2+	Mass error	0.001145



Info

Modified Sequence AcD3-ETLSQSPQGR-COOH

Protein P60335

Precursor m/z 574.29194

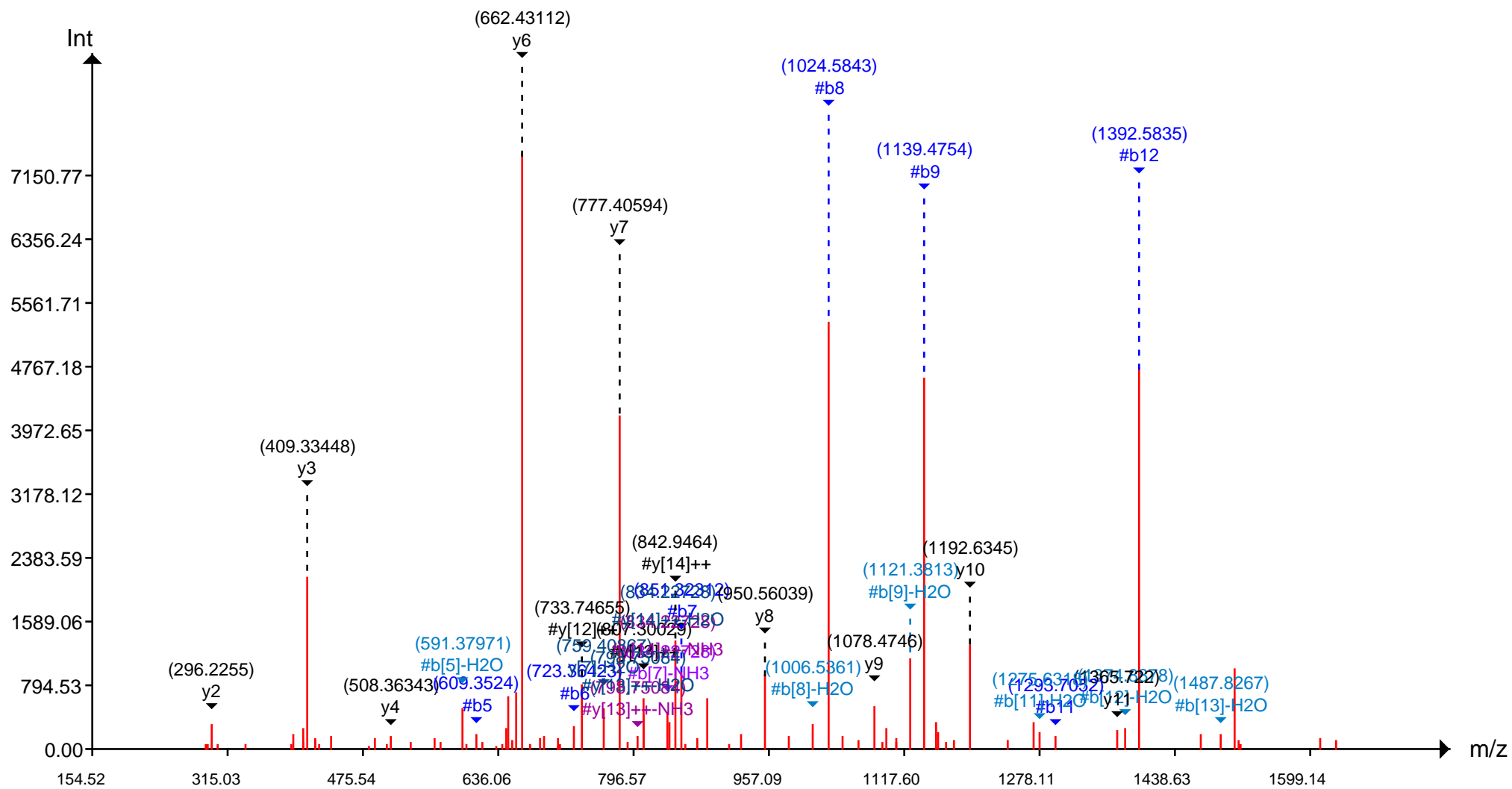
Charge 2+

Ion score 65.67

Confidence level 99.00%

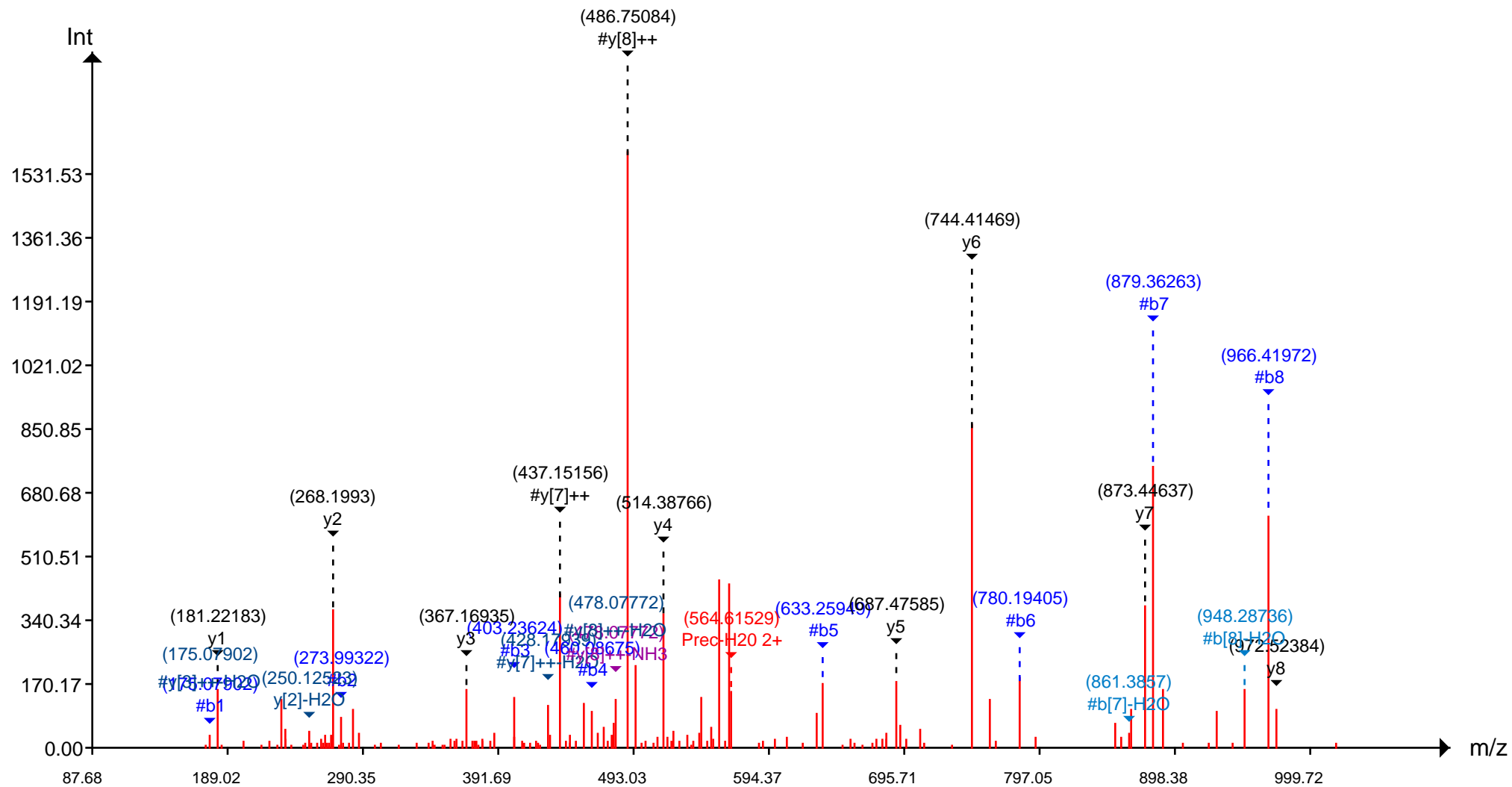
Identity Threshold 47.55

Mass error -0.001503



Info

Modified Sequence	AcD3-AAFTK<AcD3K*>NQK<AcD3K*>DPGVLDK<C13>-COOH	Ion score	62.34
Protein	P50543	Confidence level	99.00%
Precursor m/z	900.99919	Identity Threshold	47.12
Charge	2+	Mass error	0.001482



Info

Modified Sequence AcD3-EVEGK<AcD3K*>M<Mox*>VSR<C13>-COOH

Ion score 54.49

Protein P63028

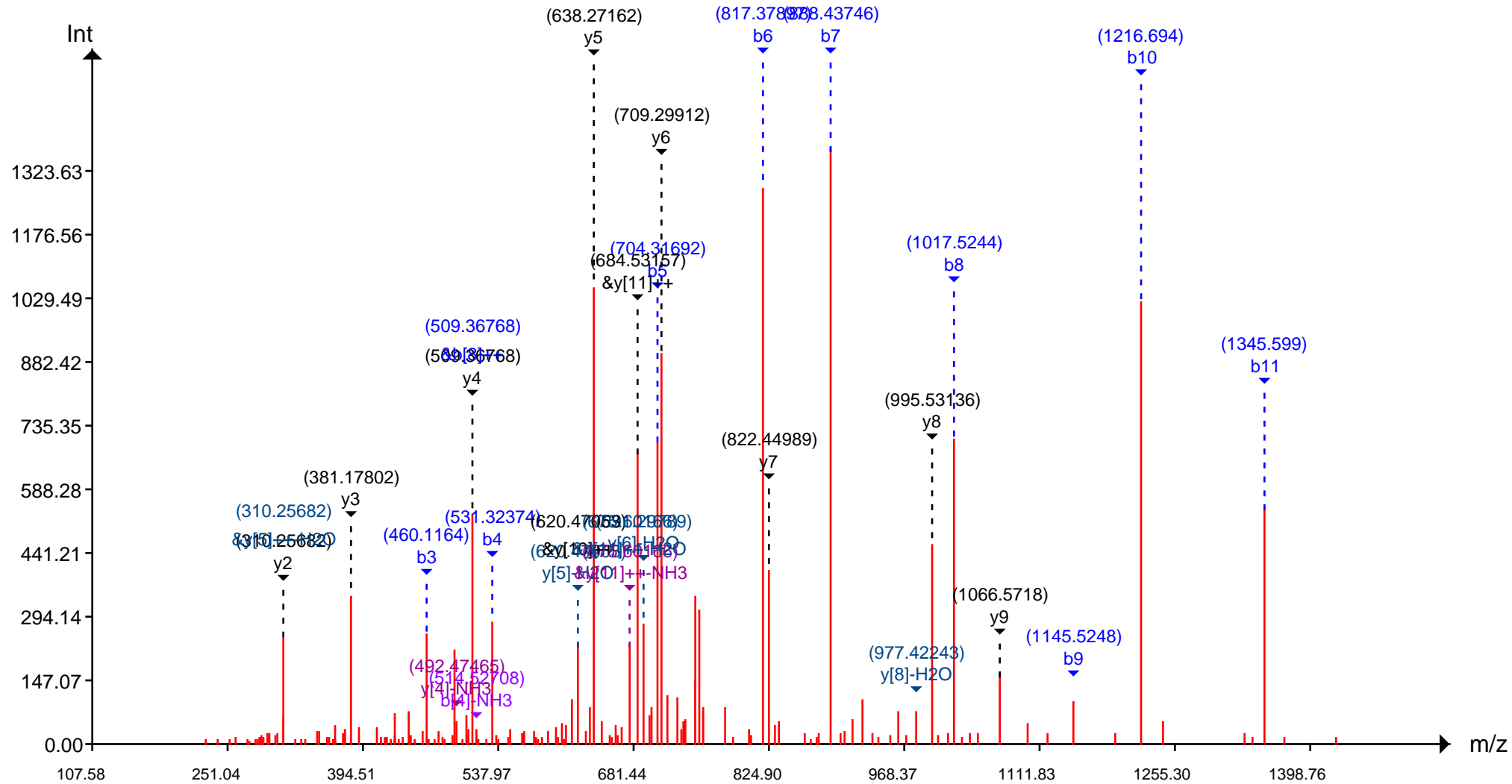
Confidence level 99.00%

Precursor m/z 573.80532

Identity Threshold 47.96

Charge 2+

Mass error -3.78E-4



Info

Modified Sequence AcD3-IQK<AcD3K*>AK<AcD3K*>LAEQAER<C13>-COOH

Ion score 66.6

Protein O70456, P68254

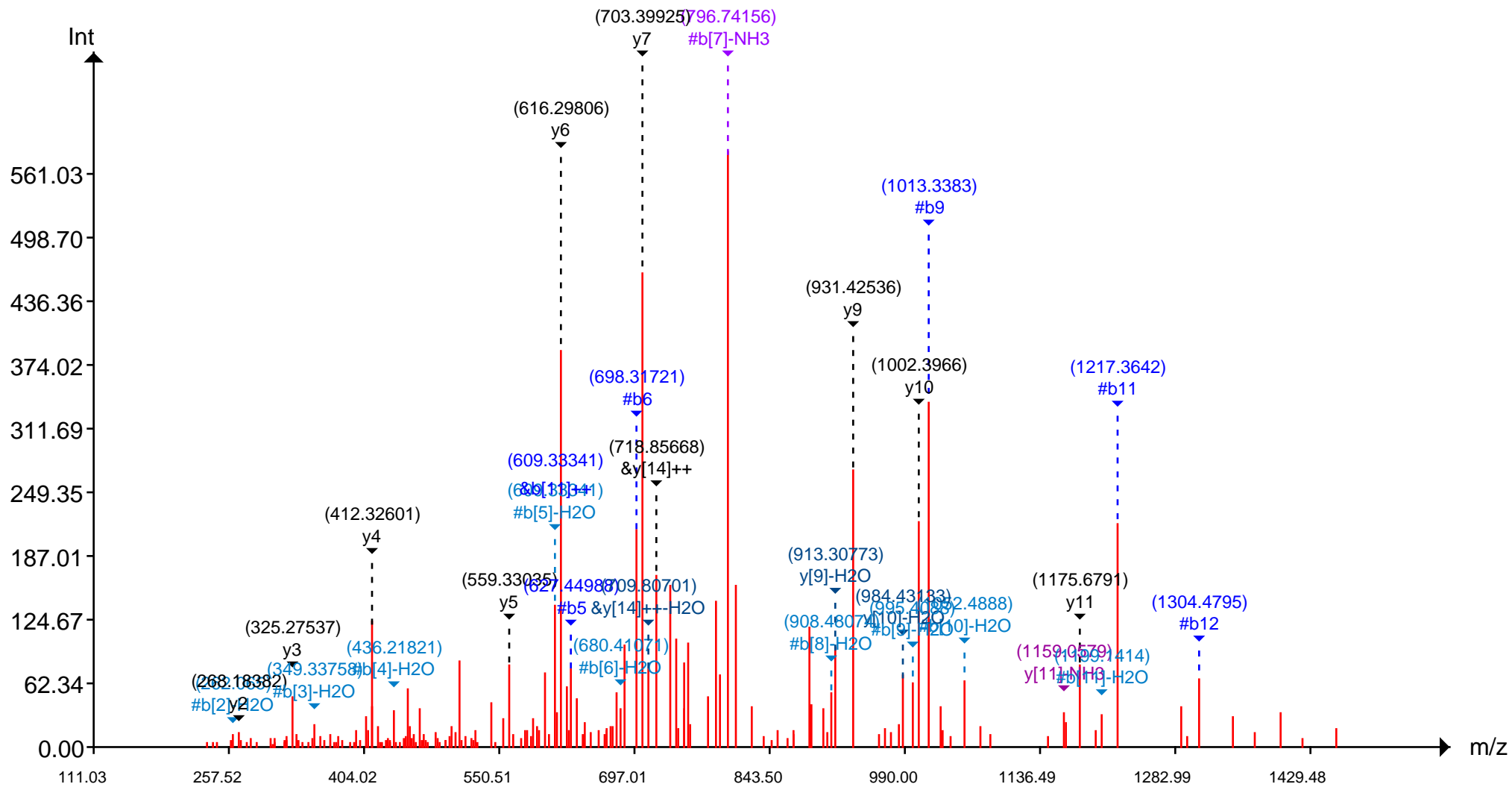
Confidence level 99.00%

Precursor m/z 763.45394

Identity Threshold 44.19

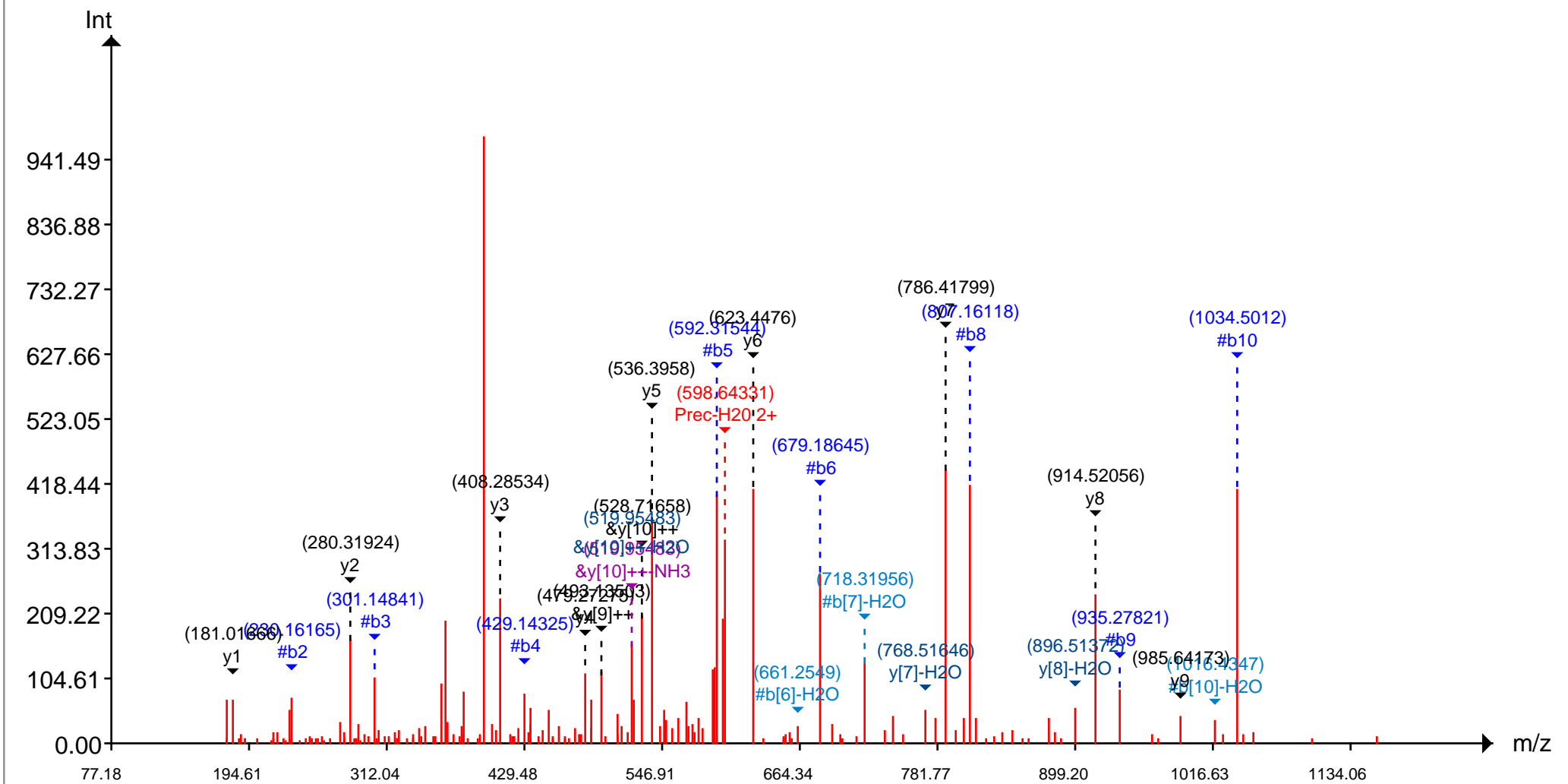
Charge 2+

Mass error 0.001635



Info

Modified Sequence	AcD3-FSSSK<AcD3K*>ADLSGM<Mox*>SGSR<C13>-COOH	Ion score	54.18
Protein	Q9D154, Q5SV42	Confidence level	99.00%
Precursor m/z	814.89387	Identity Threshold	46.66
Charge	2+	Mass error	5.9E-4



Info

Modified Sequence AcD3-IAAQYSGAQVR<C13>-COOH

Ion score 63.12

Protein Q9D8N0

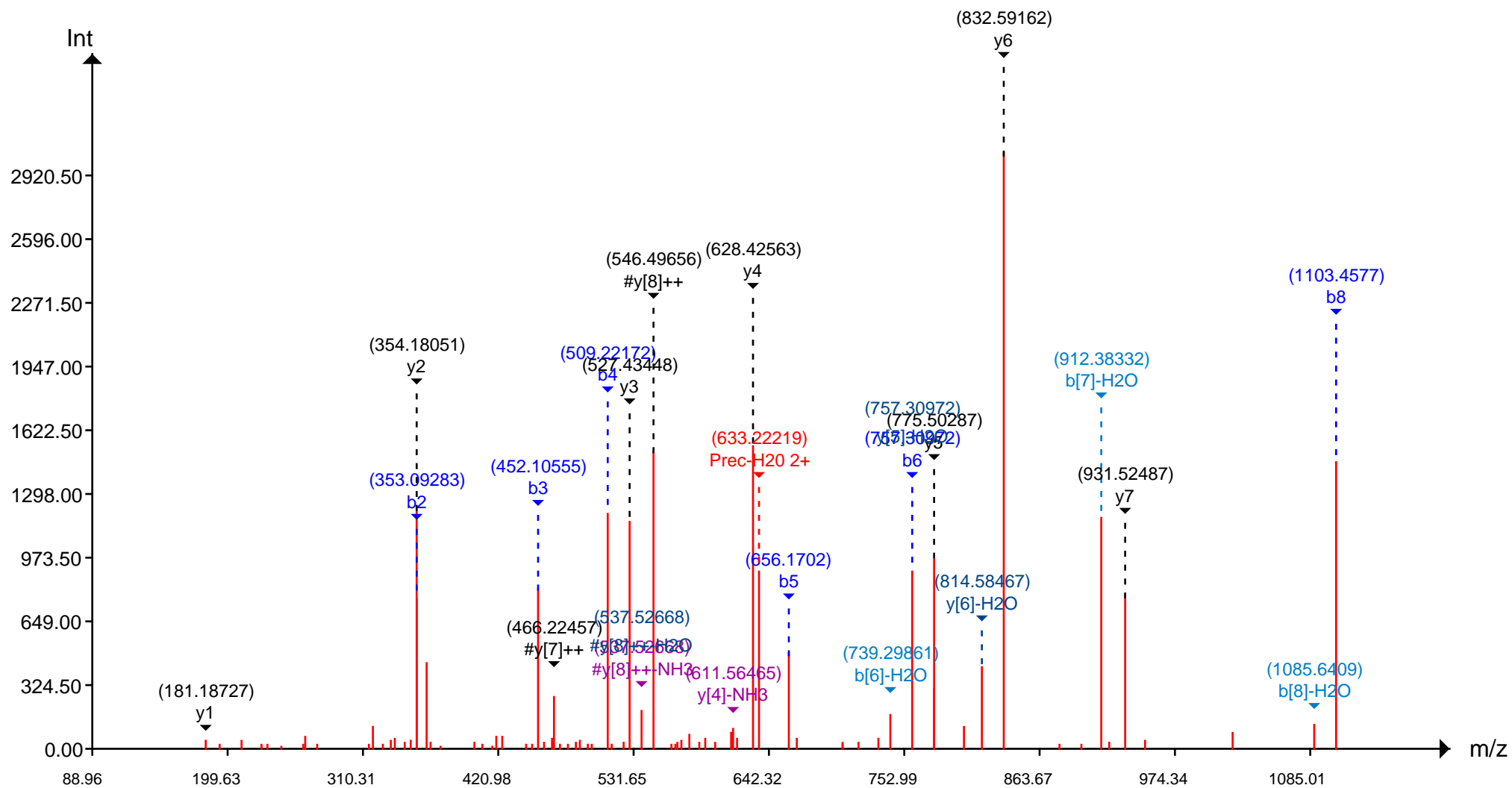
Confidence level 99.00%

Precursor m/z 607.83679

Identity Threshold 47.89

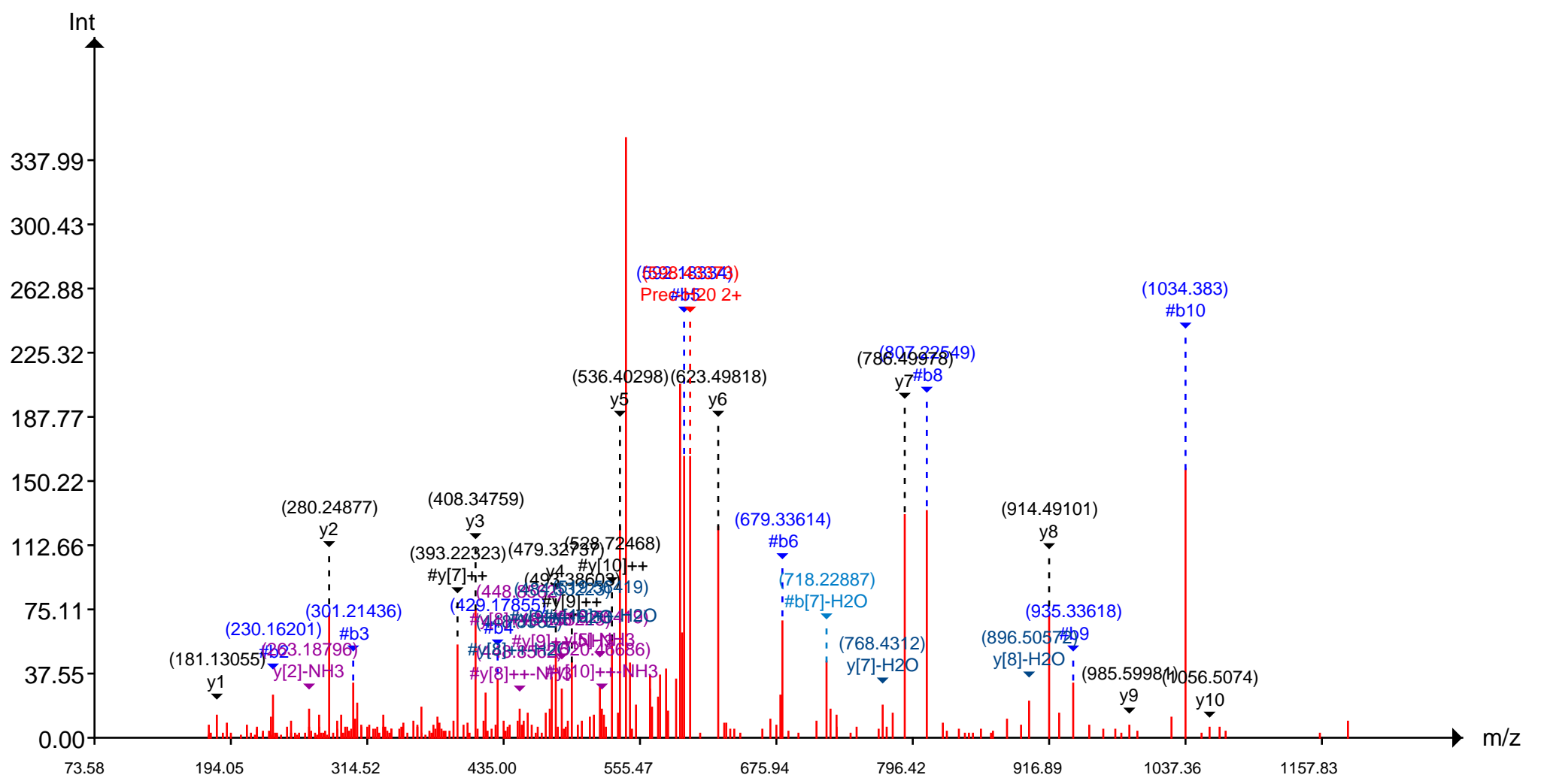
Charge 2+

Mass error 4.7E-5

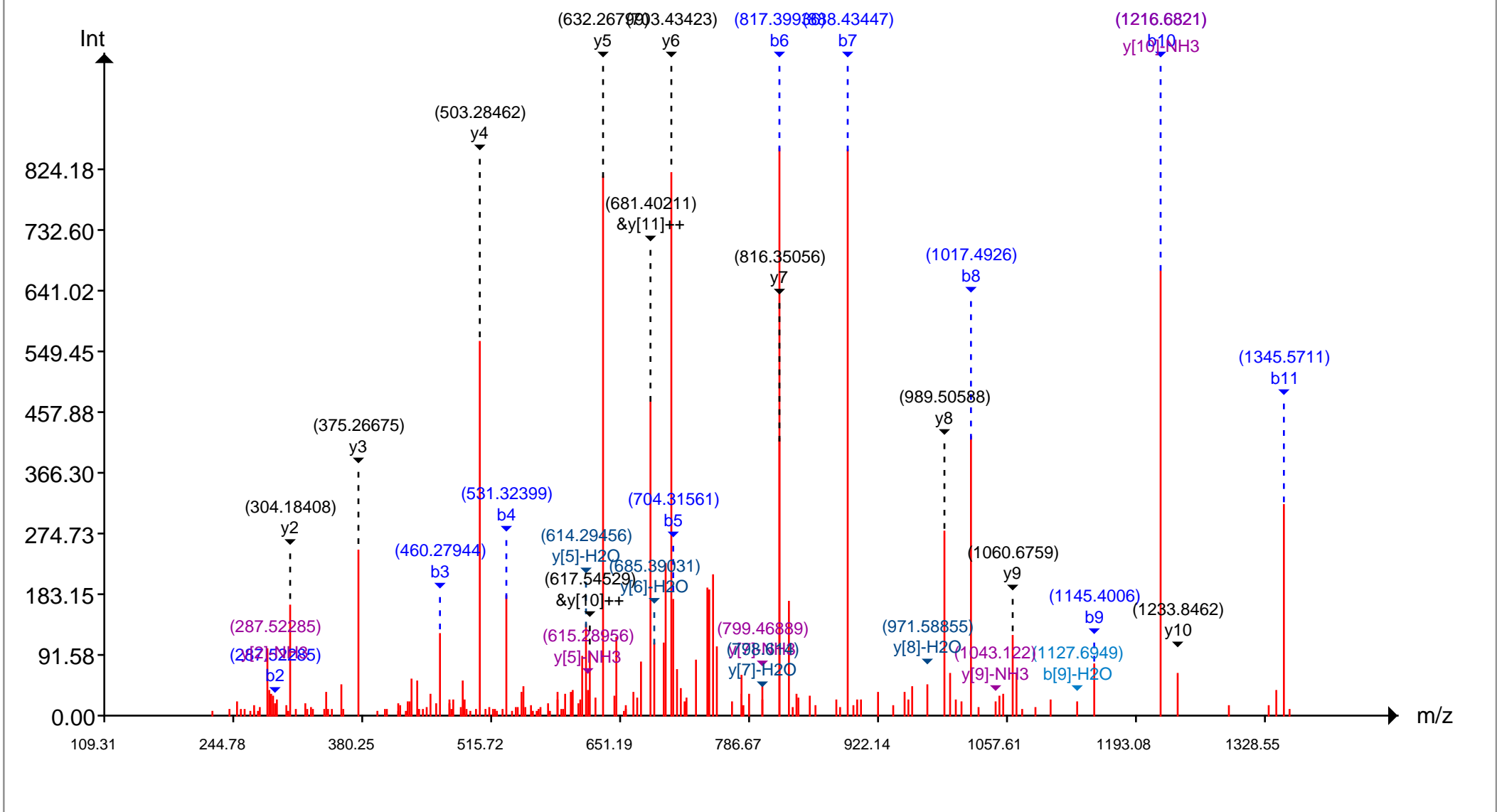


Info

Modified Sequence	AcD3-FC<Cmm*>VGFTK<AcD3K*>K<AcD3K*>R<C13>-COOH	Ion score	53.86
Protein	P97351	Confidence level	99.00%
Precursor m/z	642.36542	Identity Threshold	47.19
Charge	2+	Mass error	0.001307



Info			
Modified Sequence	AcD3-IAAQYSGAQVR<C13>-COOH	Ion score	76.45
Protein	Q9D8N0	Confidence level	99.00%
Precursor m/z	607.83659	Identity Threshold	47.85
Charge	2+	Mass error	-3.53E-4



Info

Modified Sequence AcD3-IQK<AcD3K*>AK<AcD3K*>LAEQAER-COOH

Ion score 80.82

Protein O70456, P68254

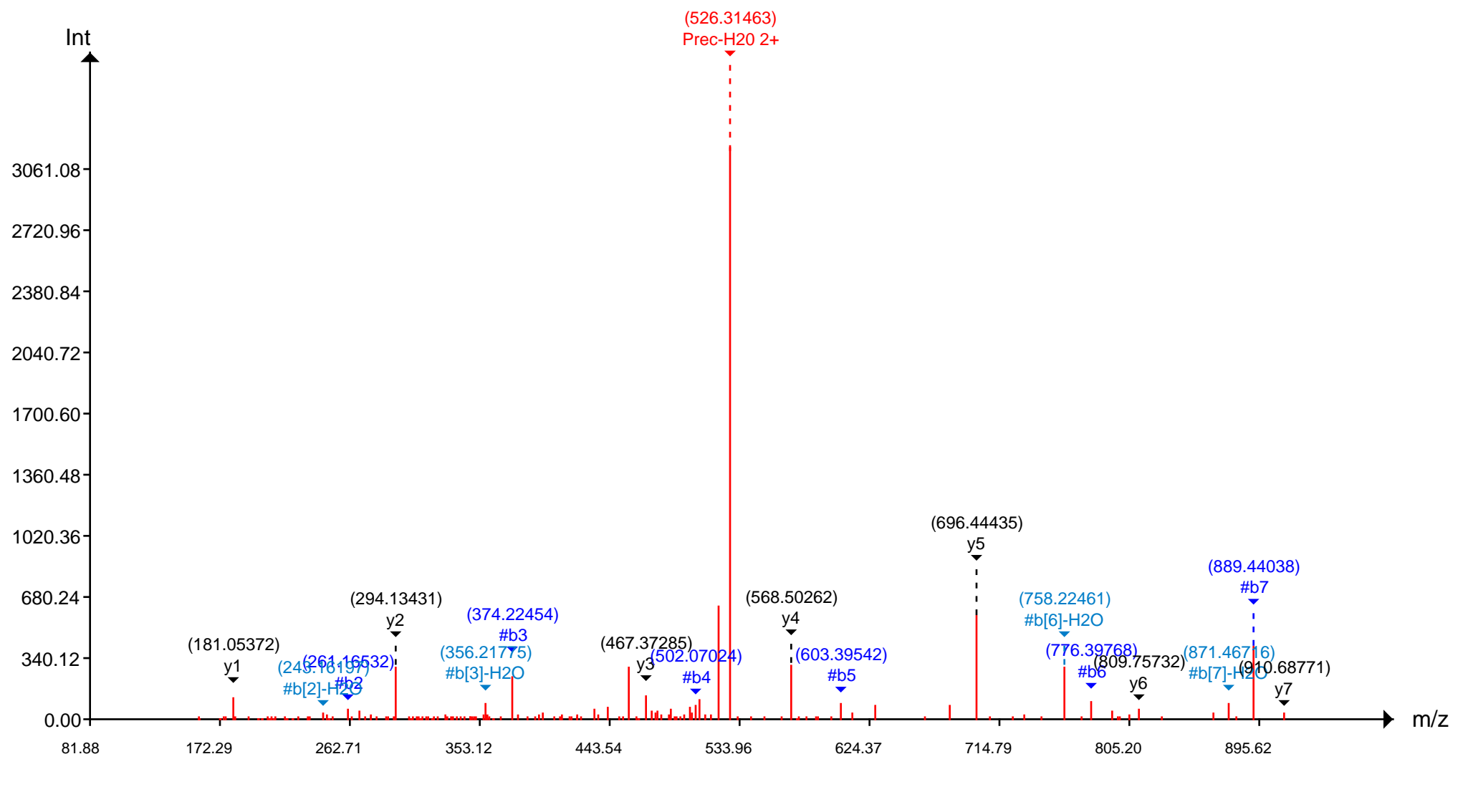
Confidence level 99.00%

Precursor m/z 760.44347

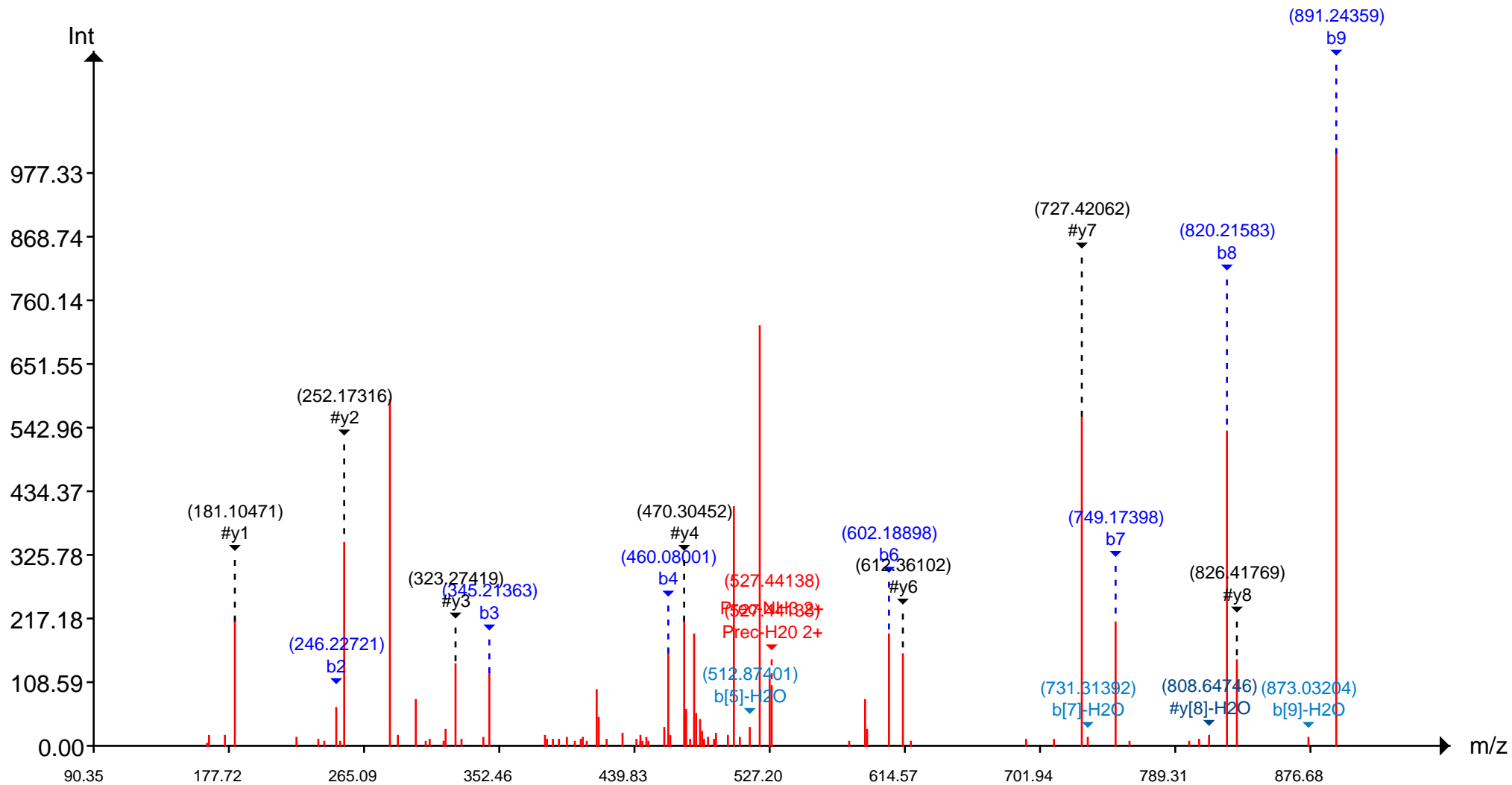
Identity Threshold 45.46

Charge 2+

Mass error 8.36E-4



Info			
Modified Sequence	AcD3-NTLQTK<AcD3K*>LR<C13>-COOH	Ion score	49.22
Protein	Q7TPR4, Q9JI91, O88990, P57780	Confidence level	99.00%
Precursor m/z	535.33213	Identity Threshold	43.98
Charge	2+	Mass error	-8.35E-4



Info

Modified Sequence AcD3-AEVDAAAM<Mox*>AAR<C13>-COOH

Protein O88569

Precursor m/z 536.26715

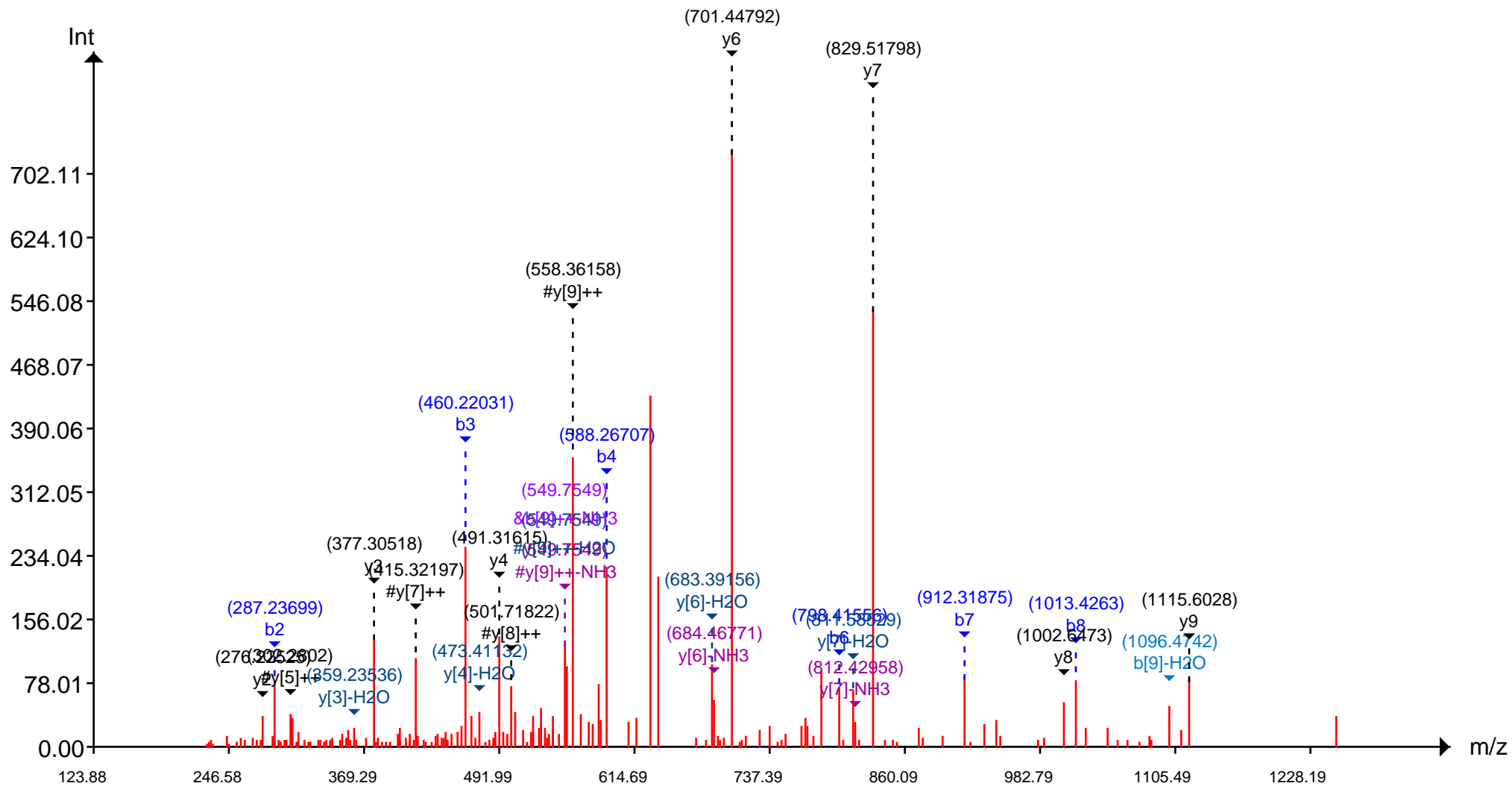
Charge 2+

Ion score 61.4

Confidence level 99.00%

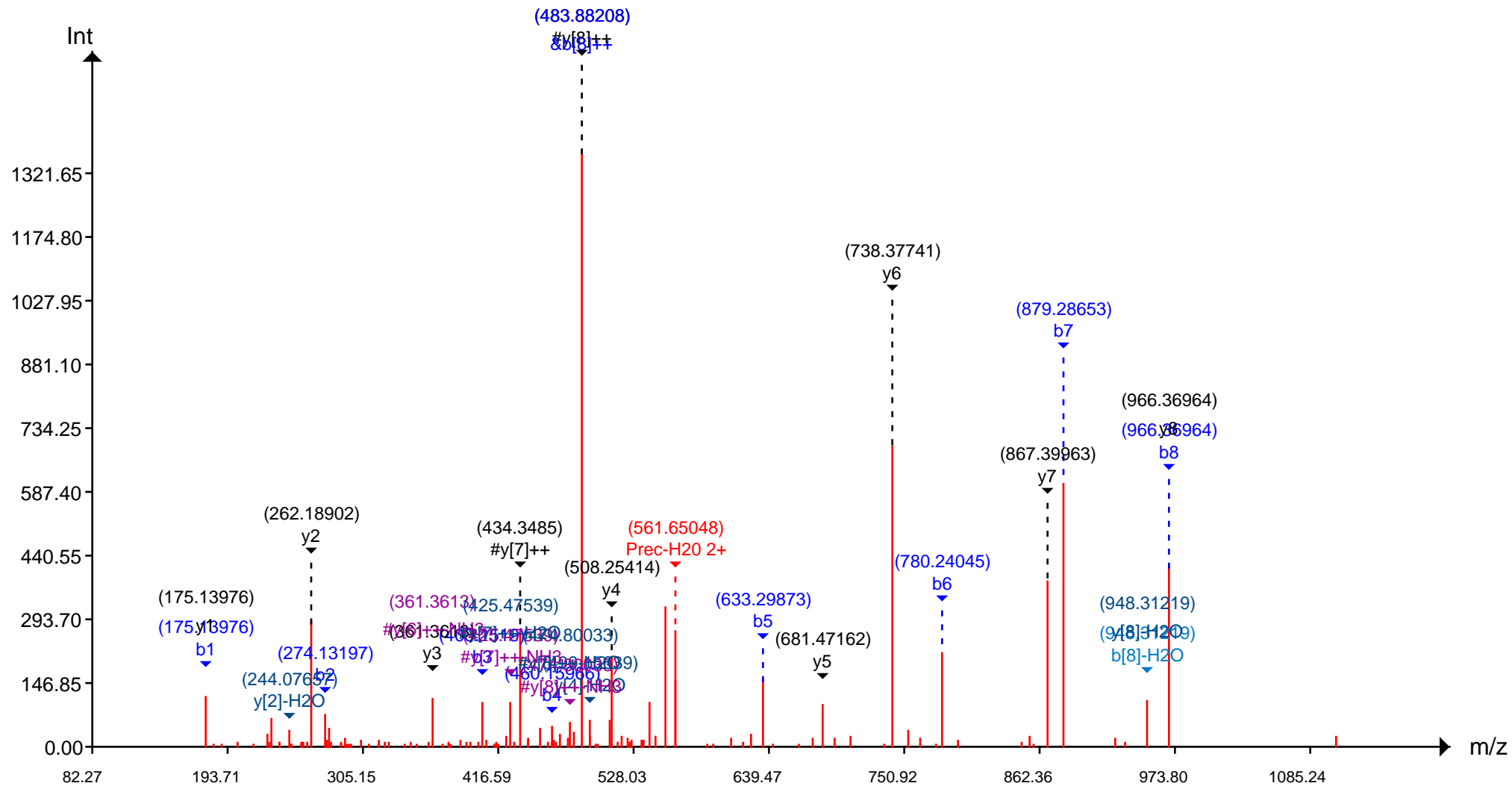
Identity Threshold 46.77

Mass error -3.48E-4



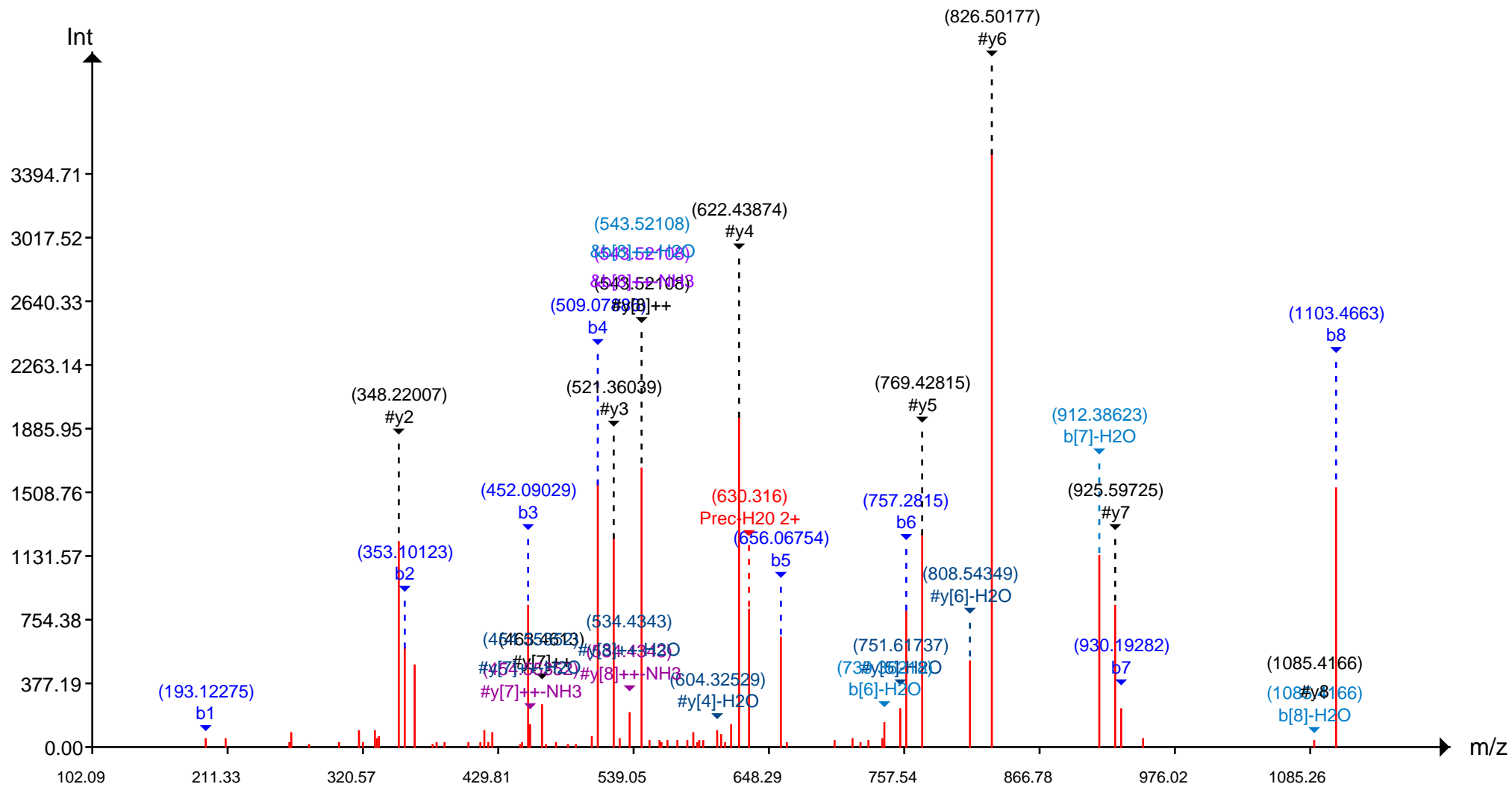
Info

Modified Sequence	AcD3-QLK<AcD3K*>QPLNTR-COOH	Ion score	60.18
Protein	P35235	Confidence level	99.00%
Precursor m/z	644.87812	Identity Threshold	46.07
Charge	2+	Mass error	-5.7E-5



Info

Modified Sequence	AcD3-EVEGK<AcD3K*>M<Mox*>VSR-COOH	Ion score	66.89
Protein	P63028	Confidence level	99.00%
Precursor m/z	570.79548	Identity Threshold	47.78
Charge	2+	Mass error	8.4E-5



Info

Modified Sequence AcD3-FC<Cmm*>VGFTK<AcD3K*>K<AcD3K*>R-COOH

Ion score 74.06

Protein P97351

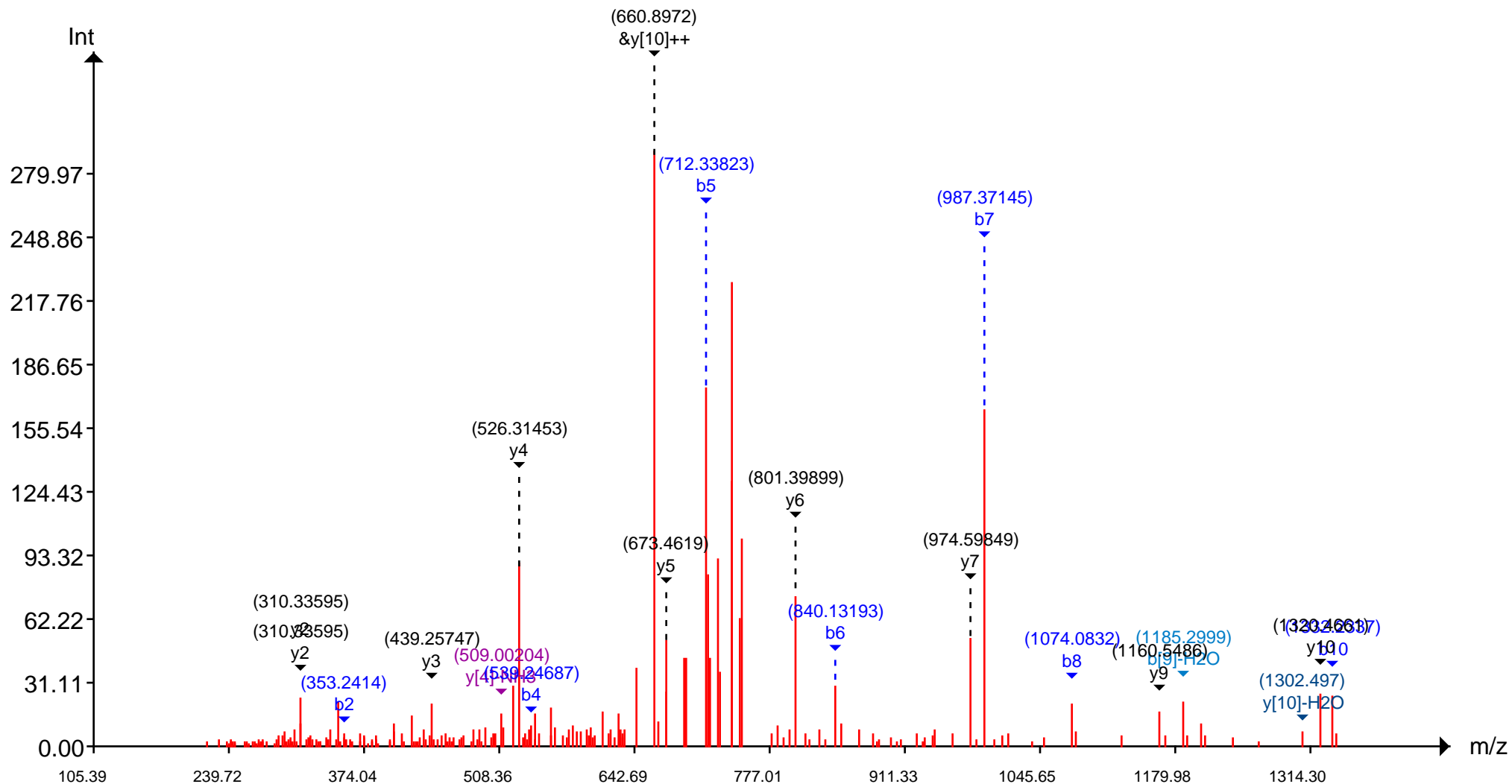
Confidence level 99.00%

Precursor m/z 639.35558

Identity Threshold 47.69

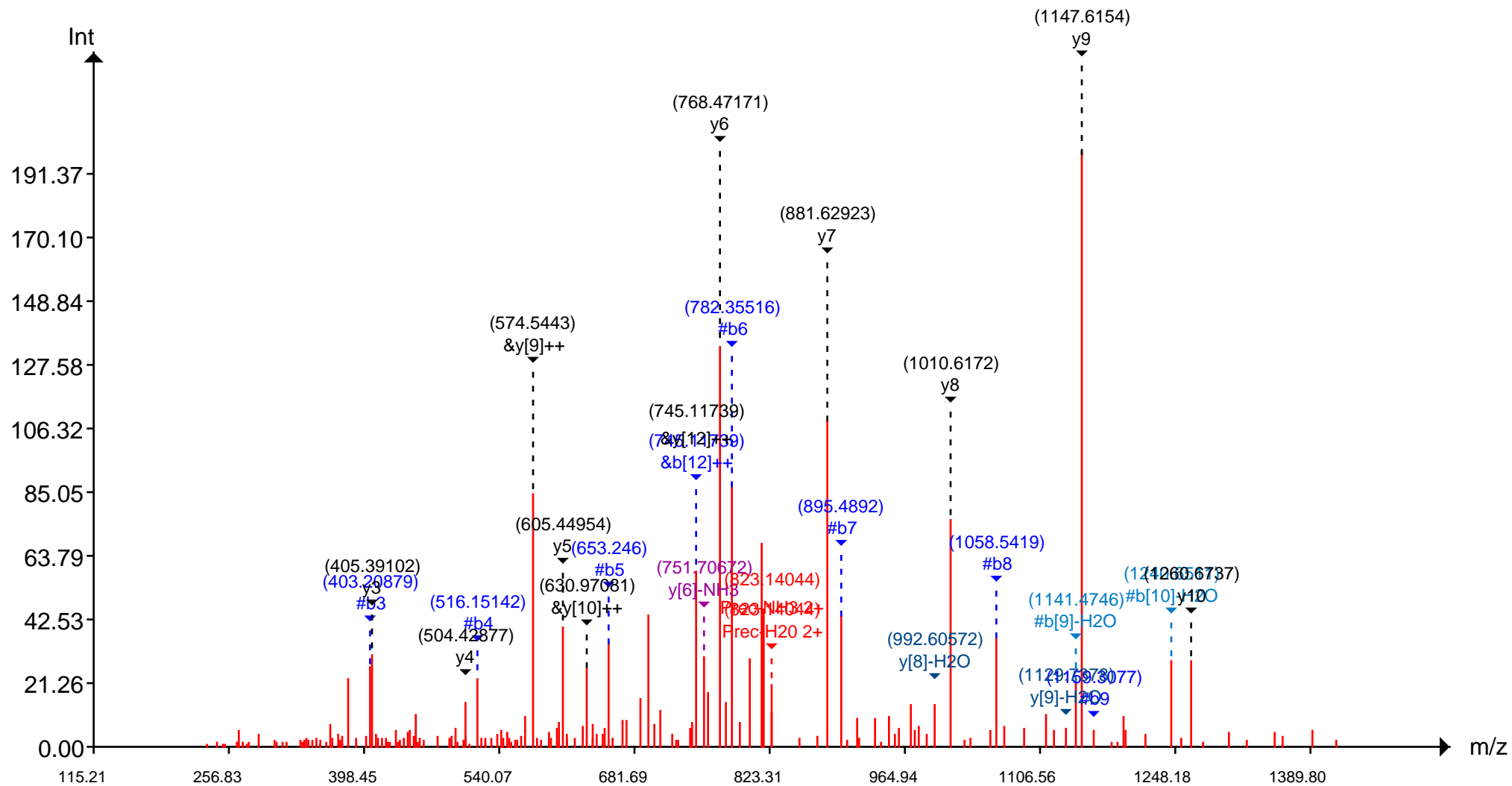
Charge 2+

Mass error 0.001769



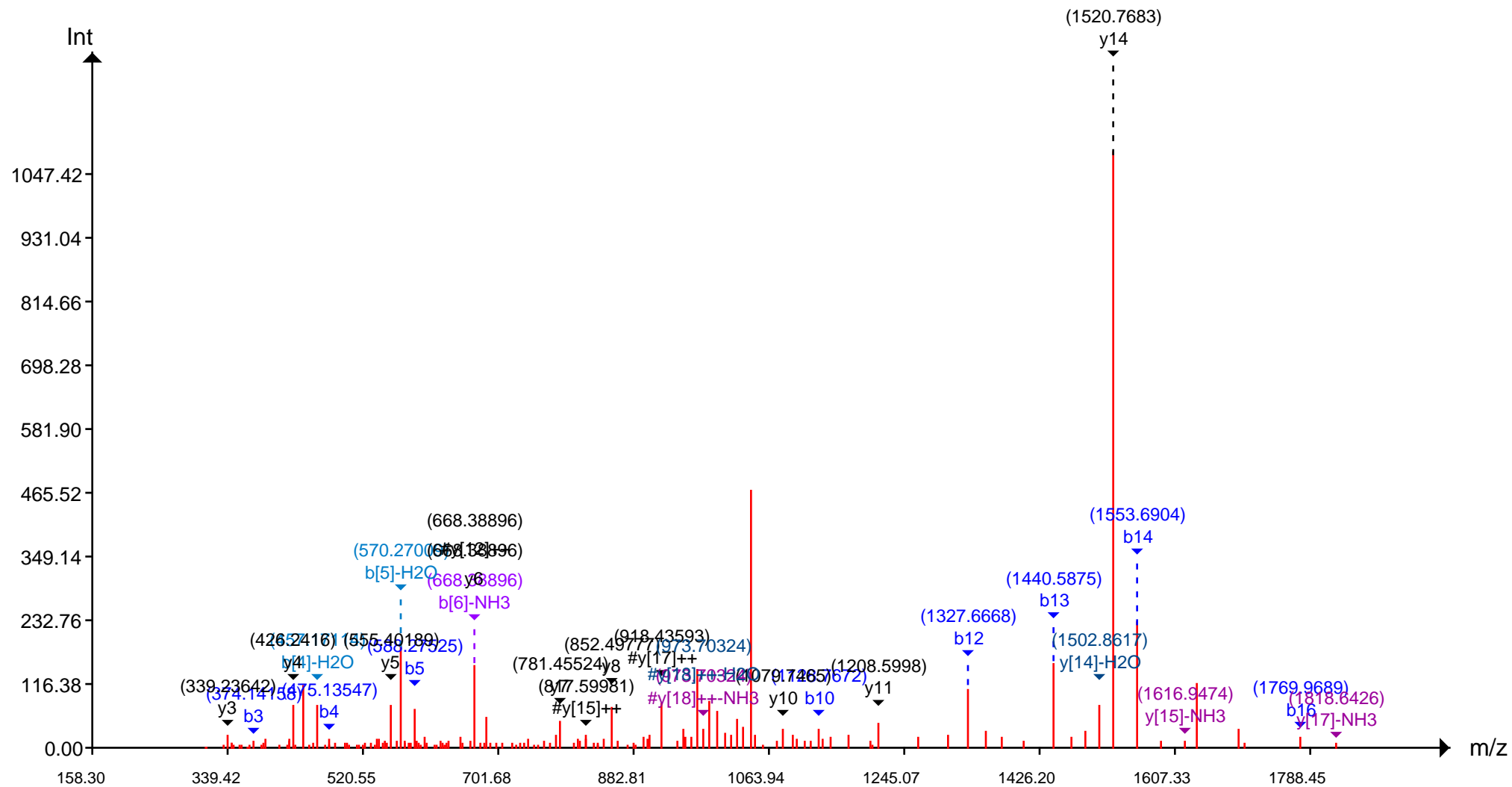
Info

Modified Sequence	AcD3-M<Mox*>C<Cmm*>GEK<AcD3K*>QFSEER<C13>-COOH	Ion score	48.82
Protein	P39749	Confidence level	99.00%
Precursor m/z	756.83753	Identity Threshold	43.60
Charge	2+	Mass error	6.42E-4



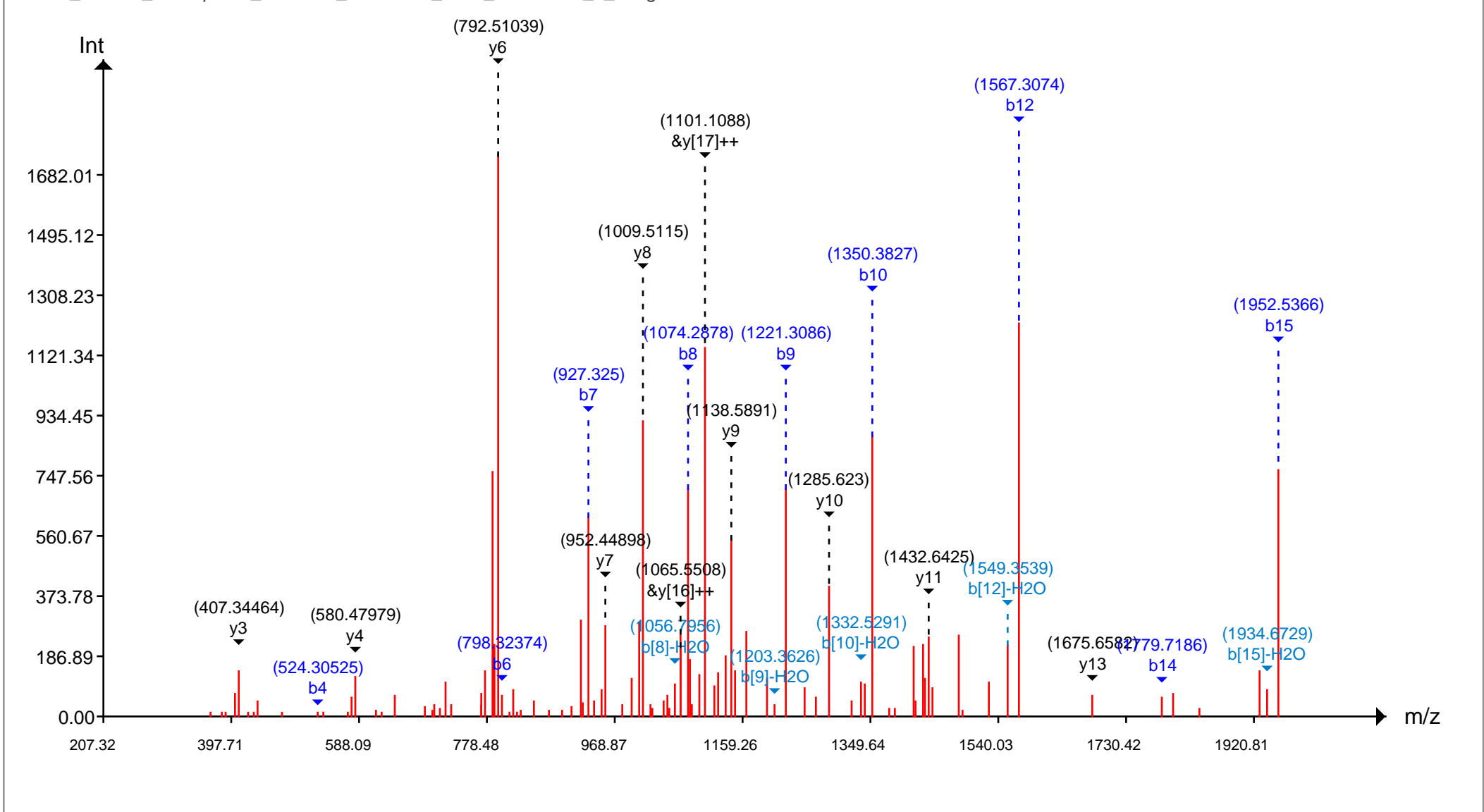
Info

Modified Sequence	AcD3-EDLIHEIYTVGK<AcD3K*>R-COOH	Ion score	58.98
Protein	P14148	Confidence level	99.00%
Precursor m/z	831.95289	Identity Threshold	47.76
Charge	2+	Mass error	0.001702



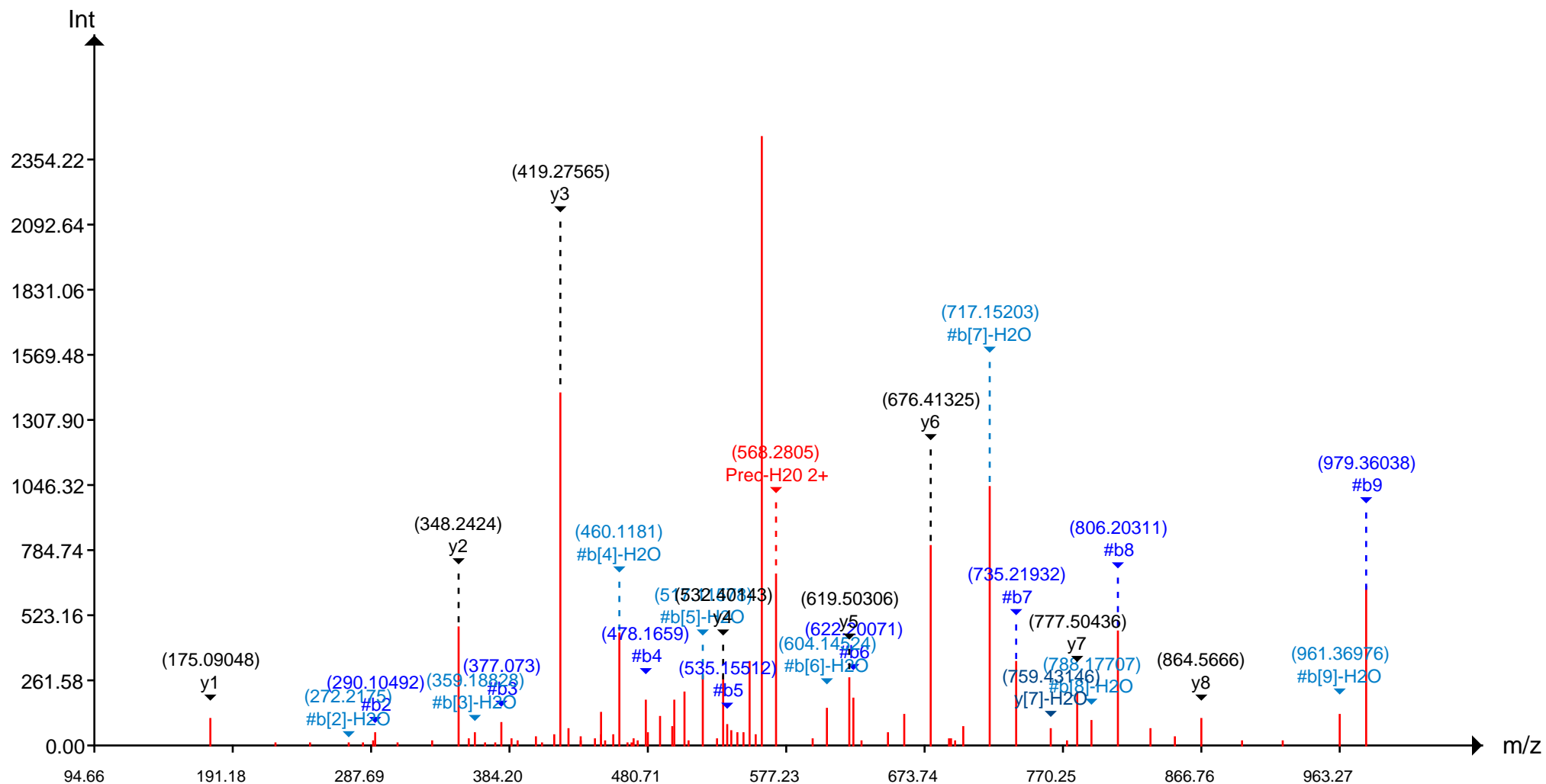
Info

Modified Sequence	AcD3-VQTTLPSQEVQALLESTGR<C13>-COOH	Ion score	47.47
Protein	Q9WU84	Confidence level	99.00%
Precursor m/z	1054.5739	Identity Threshold	47.21
Charge	2+	Mass error	0.00367



Info

Modified Sequence	AcD3-IAM<Mox*>M<Mox*>C<Cmm*>NEFFEGC<Cmm*>PDK<AcD3K*>EPR<C13>-COOH	Ion score	48.01
Protein	P07091	Confidence level	99.00%
Precursor m/z	1180.0086	Identity Threshold	40.40
Charge	2+	Mass error	0.002343



Info

Modified Sequence AcD3-DESTGSIK<AcD3K*>R-COOH

Ion score 63.15

Protein P05064

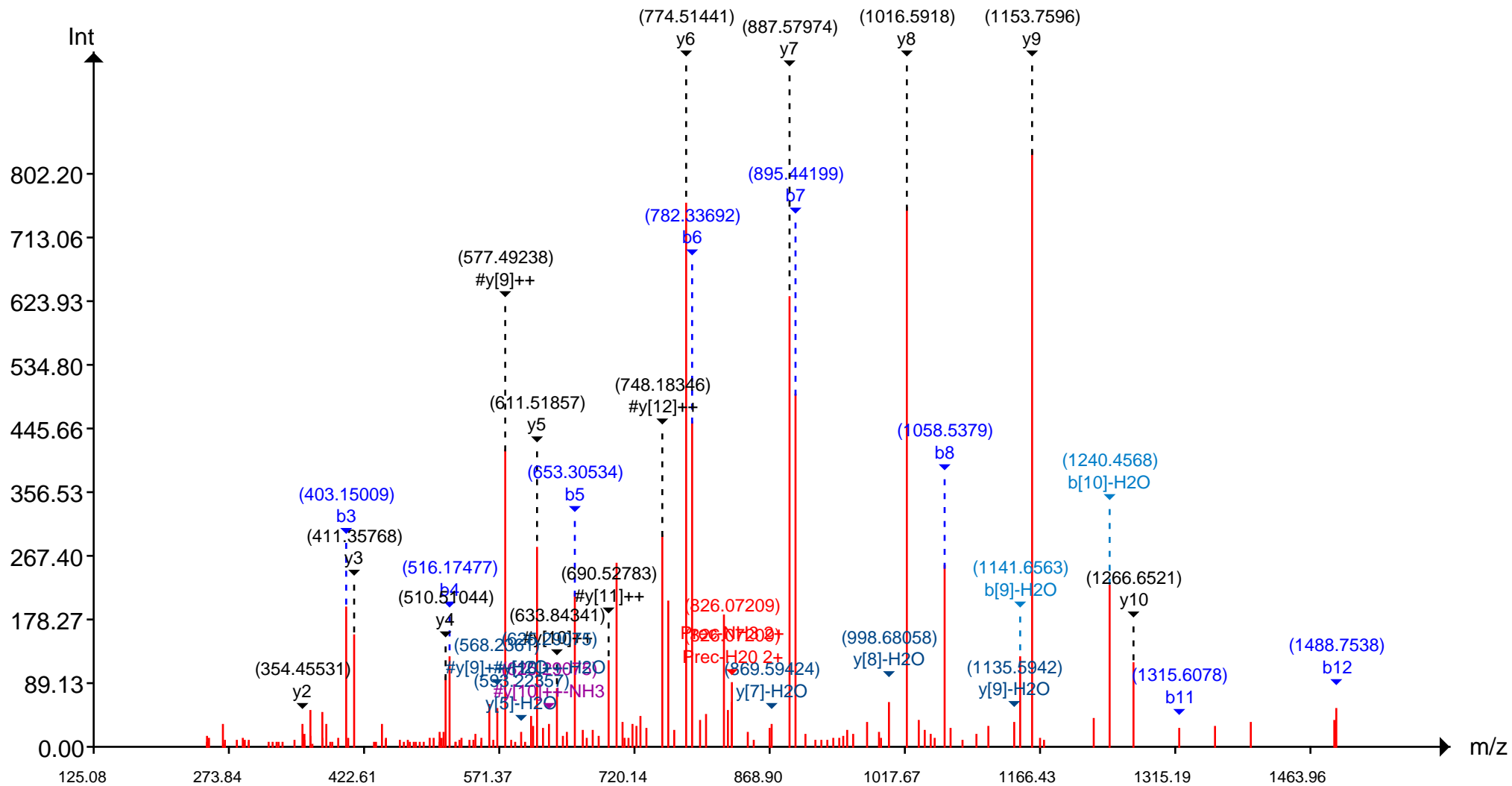
Confidence level 99.00%

Precursor m/z 577.30195

Identity Threshold 47.87

Charge 2+

Mass error 2.3E-5



Info

Modified Sequence AcD3-EDLIHEIYTVGK<AcD3K*>R<C13>-COOH

Ion score 68.56

Protein P14148

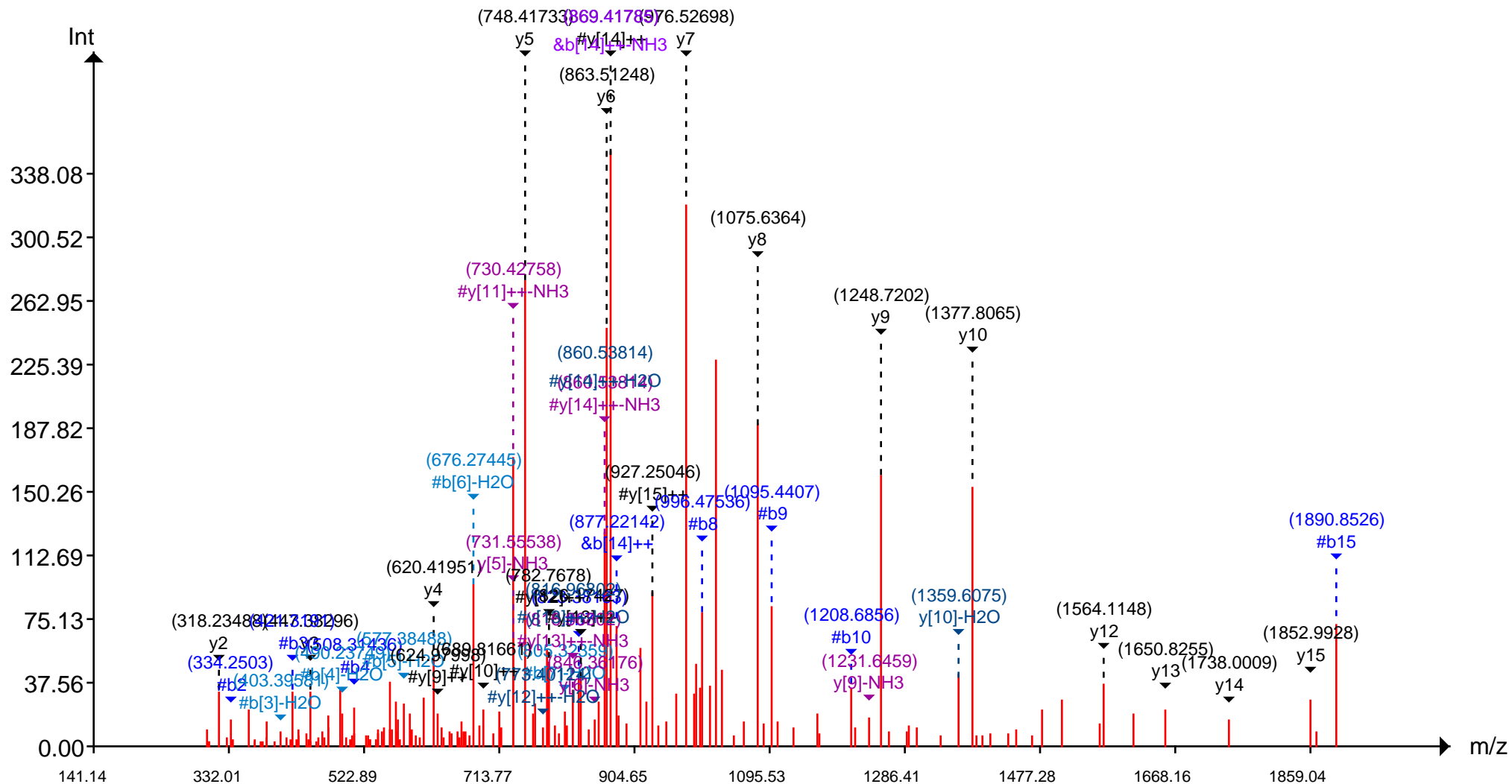
Confidence level 99.00%

Precursor m/z 834.96311

Identity Threshold 47.20

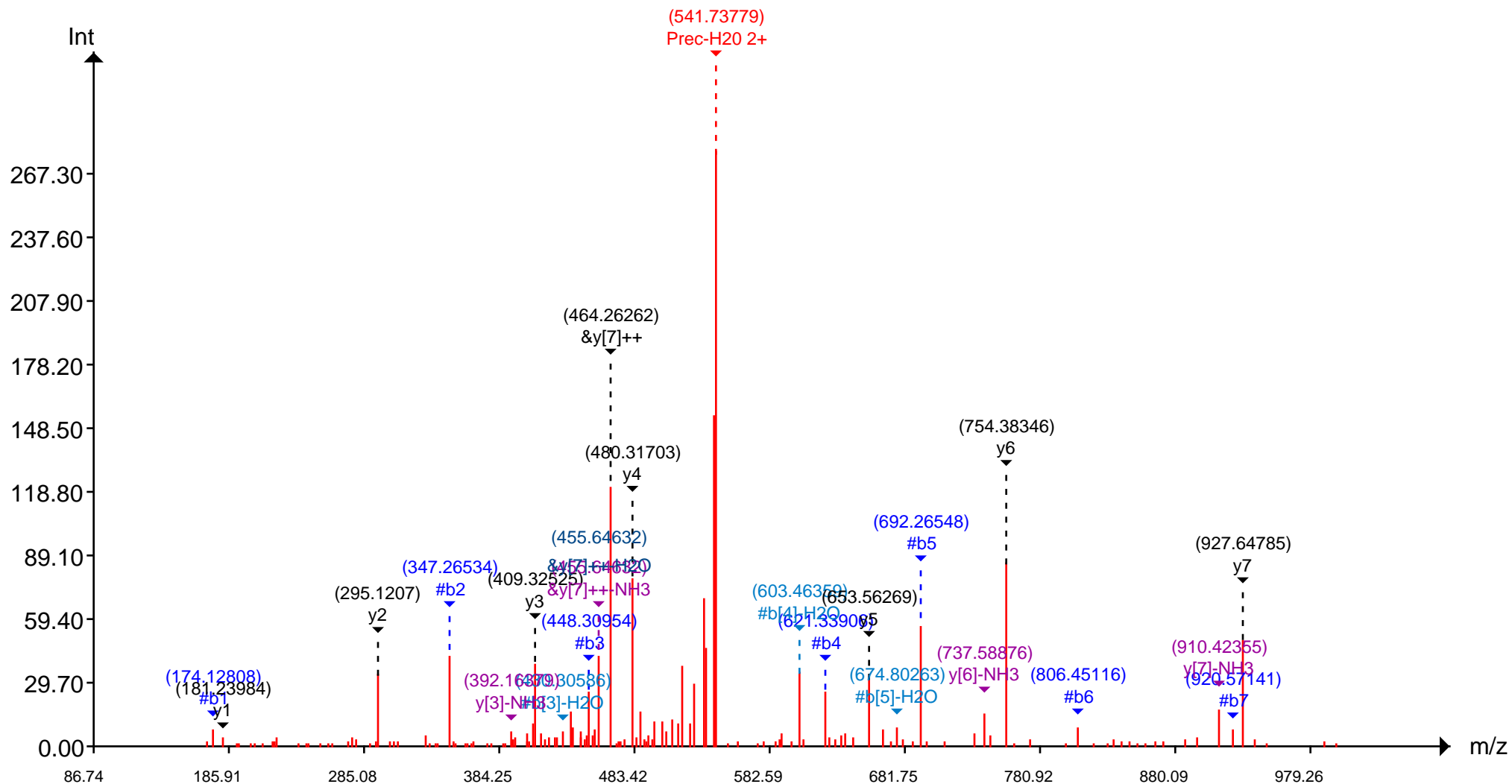
Charge 2+

Mass error 0.0020



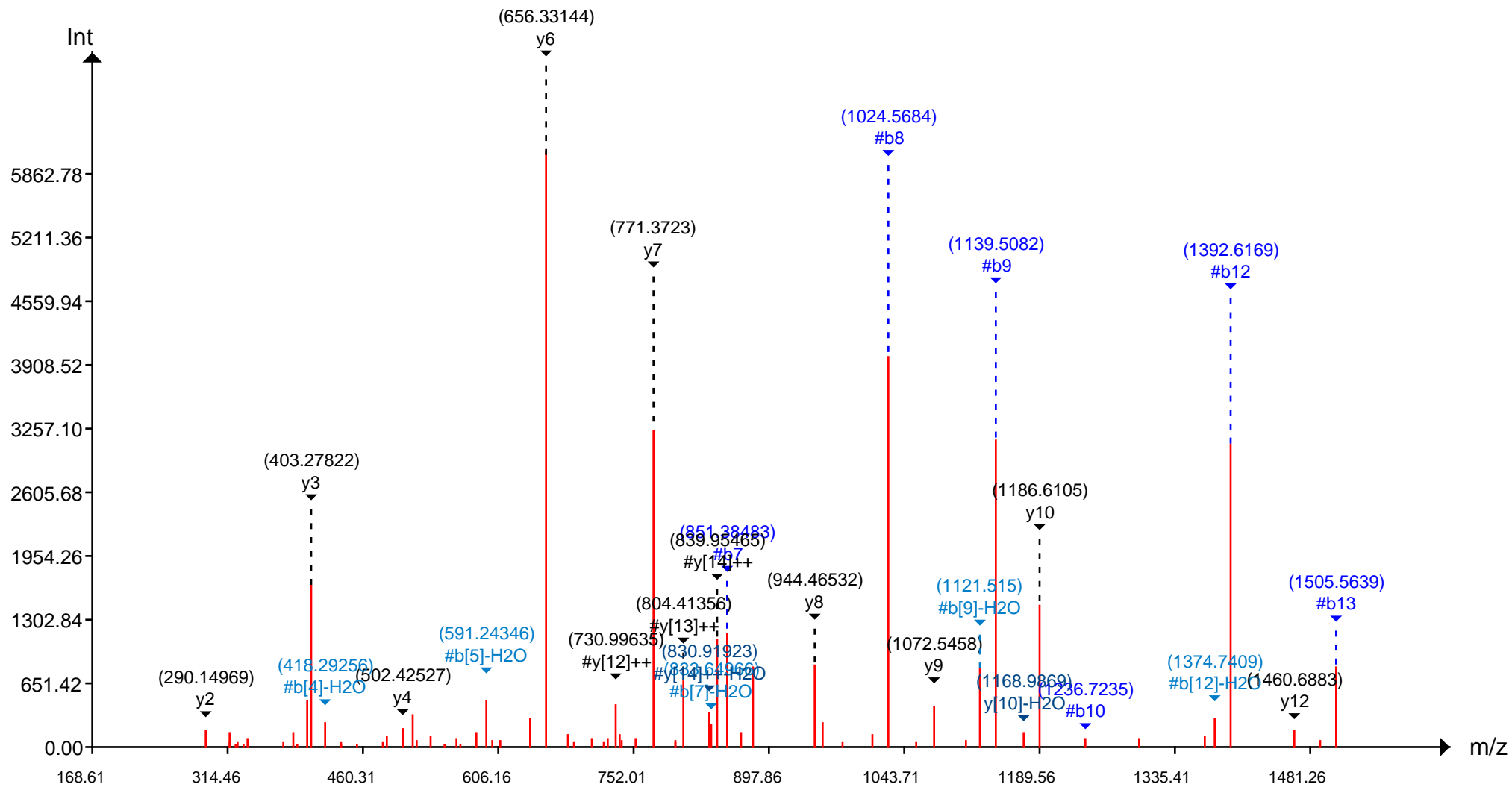
Info

Modified Sequence	AcD3-K<AcD3K*>DSSSVEK<AcD3K*>VLDQK<AcD3K*>EHR<C13>-COOH	Ion score	107.56
Protein	Q99020	Confidence level	99.00%
Precursor m/z	1036.0628	Identity Threshold	47.46
Charge	2+	Mass error	0.003229



Info

Modified Sequence	AcD3-QK<AcD3K*>TK<AcD3K*>ANNR<C13>-COOH	Ion score	52.96
Protein	O08585	Confidence level	99.00%
Precursor m/z	550.82645	Identity Threshold	45.71
Charge	2+	Mass error	-7.82E-4



Info

Modified Sequence AcD3-AAFTK<AcD3K*>NQK<AcD3K*>DPGVLDK-COOH

Ion score 74.25

Protein P50543

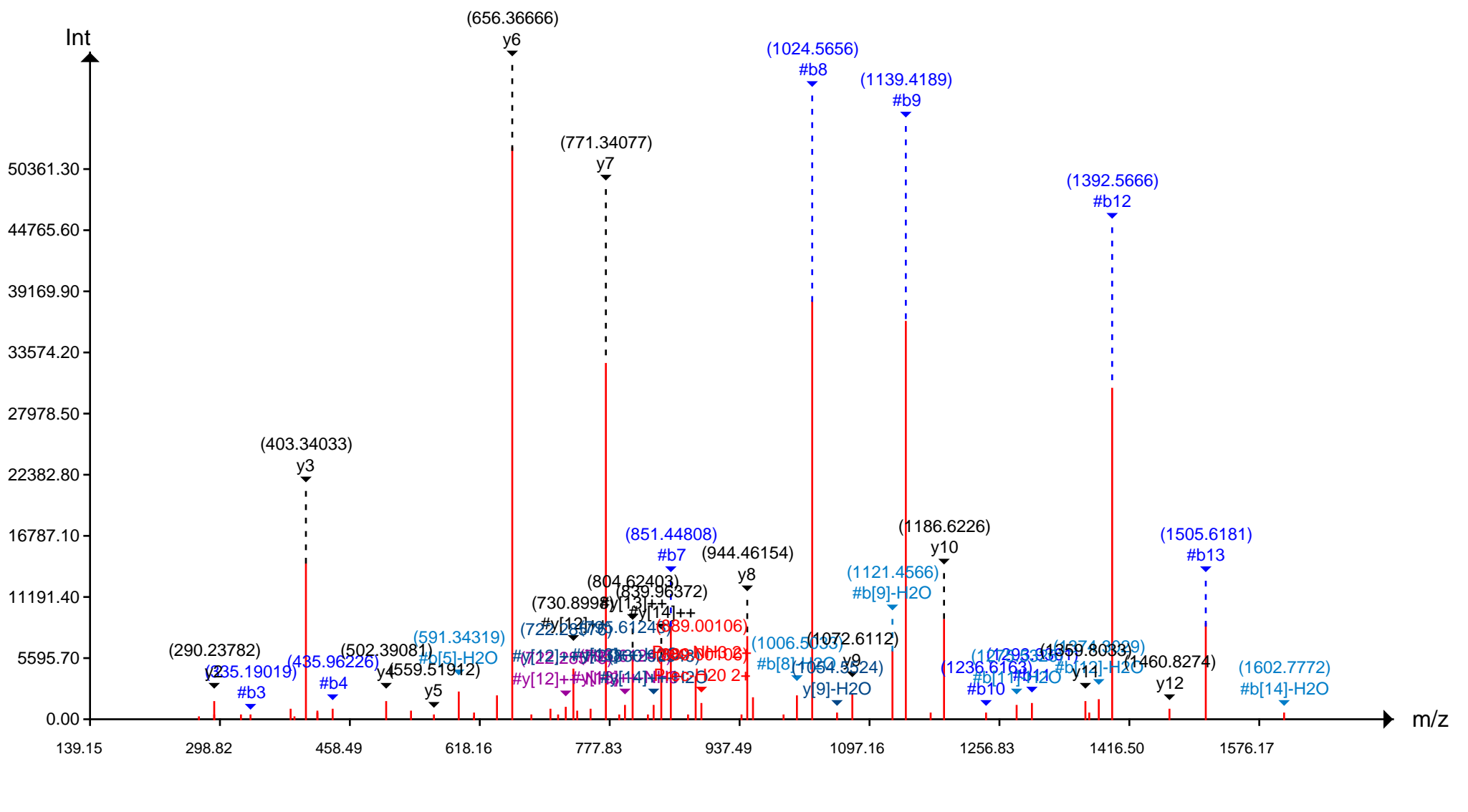
Confidence level 99.00%

Precursor m/z 897.98943

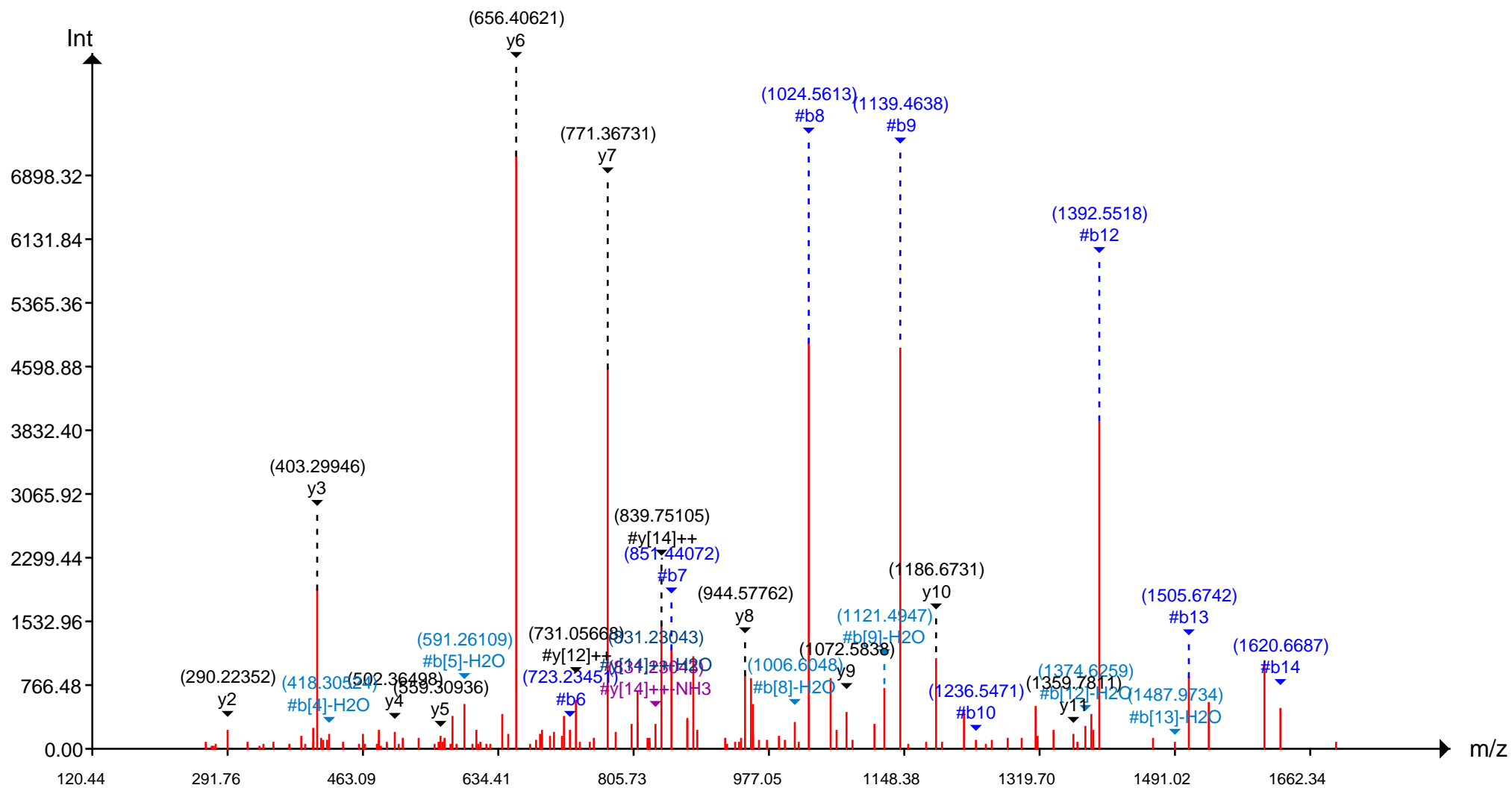
Identity Threshold 47.66

Charge 2+

Mass error 0.002104



Info			
Modified Sequence	AcD3-AAFTK<AcD3K*>NQK<AcD3K*>DPGVLDR-COOH	Ion score	107.92
Protein	P50543	Confidence level	99.00%
Precursor m/z	897.98974	Identity Threshold	47.62
Charge	2+	Mass error	0.002724



Info

Modified Sequence AcD3-AAFTK<AcD3K*>NQK<AcD3K*>DPGVLDK-COOH

Ion score 67.25

Protein P50543

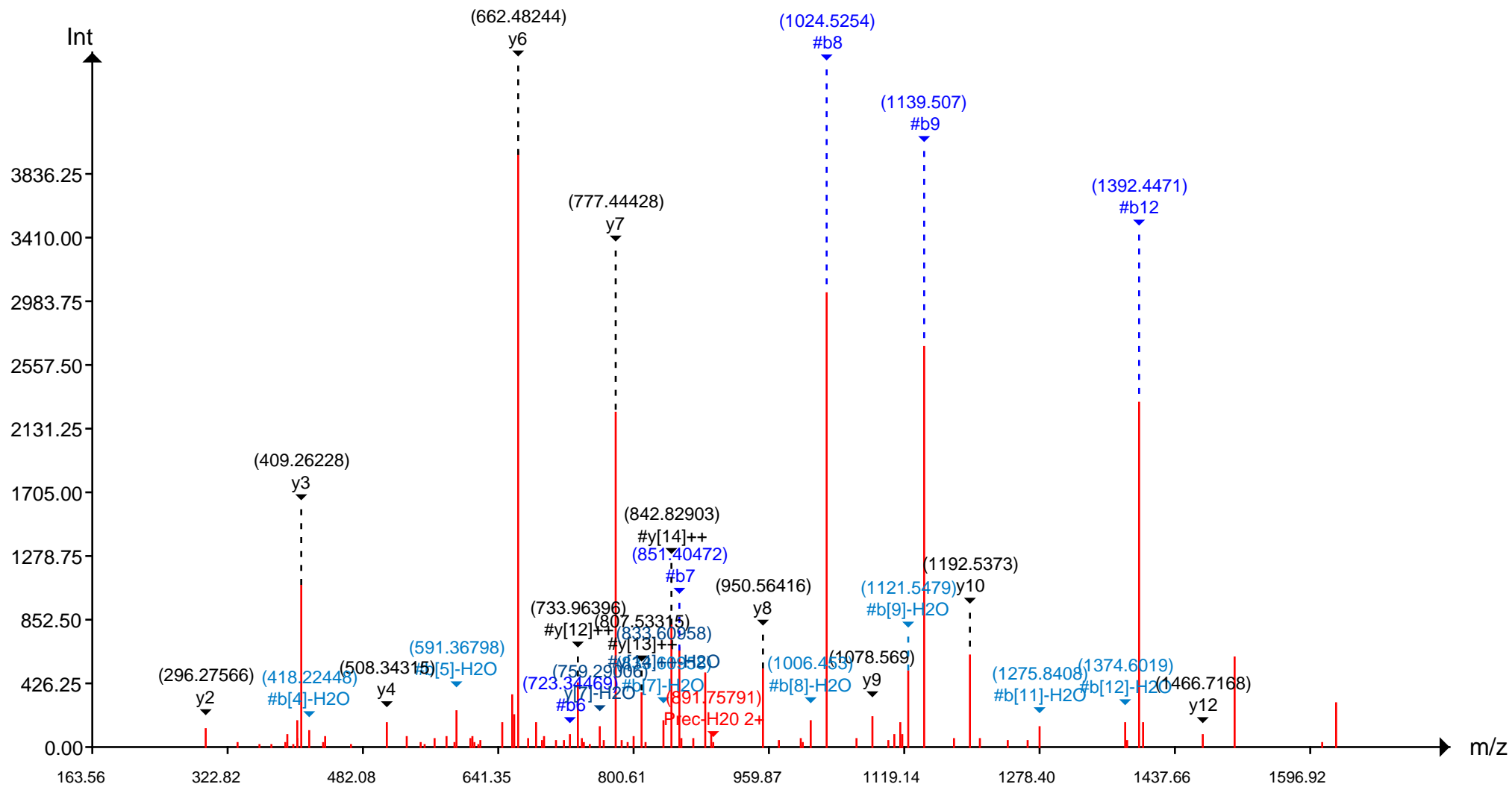
Confidence level 99.00%

Precursor m/z 897.99012

Identity Threshold 47.62

Charge 2+

Mass error 0.003484



Info

Modified Sequence AcD3-AAFTK<AcD3K*>NQK<AcD3K*>DPGVLDK<C13>-COOH

Ion score 66.75

Protein P50543

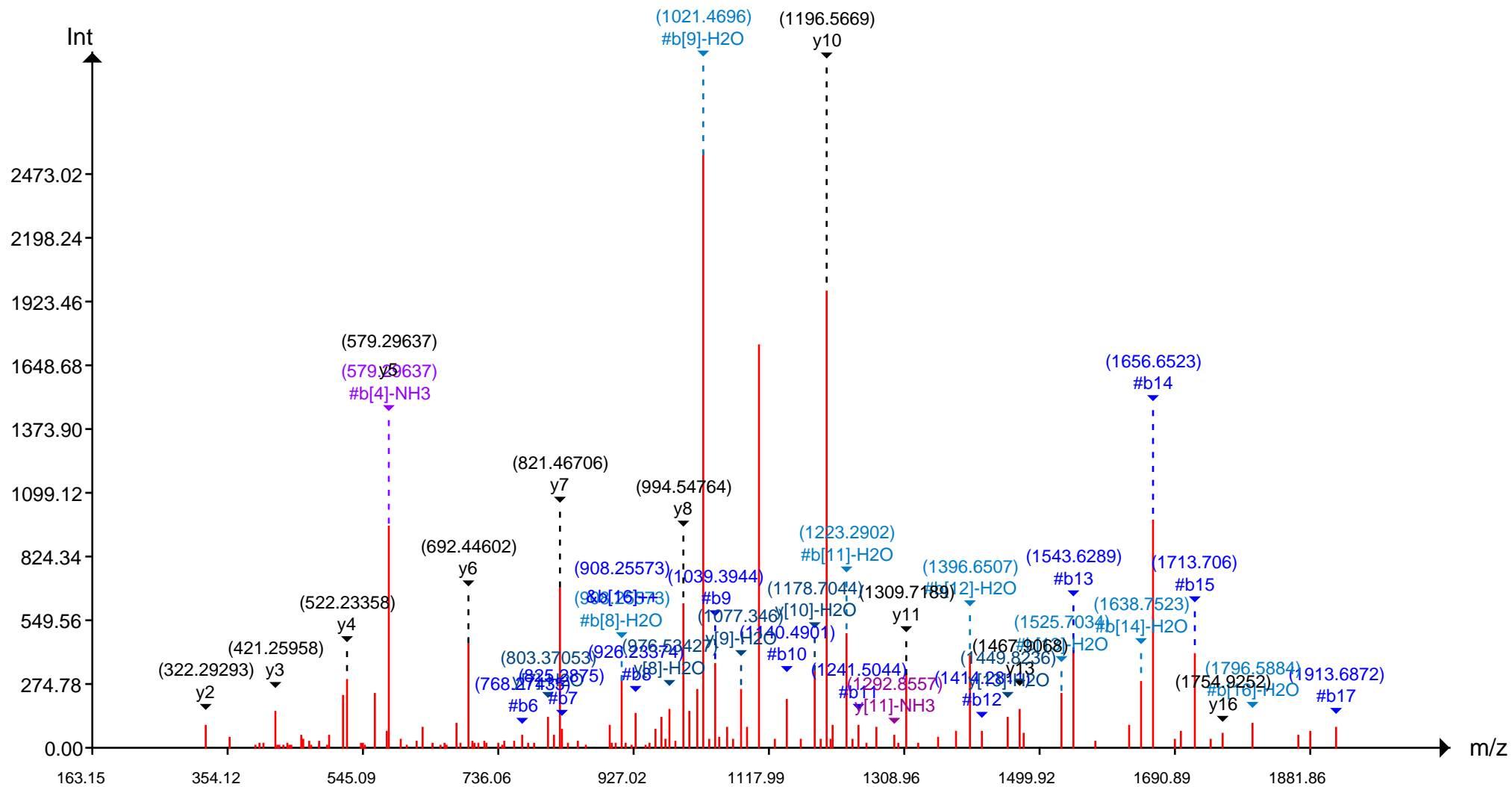
Confidence level 99.00%

Precursor m/z 900.99662

Identity Threshold 47.32

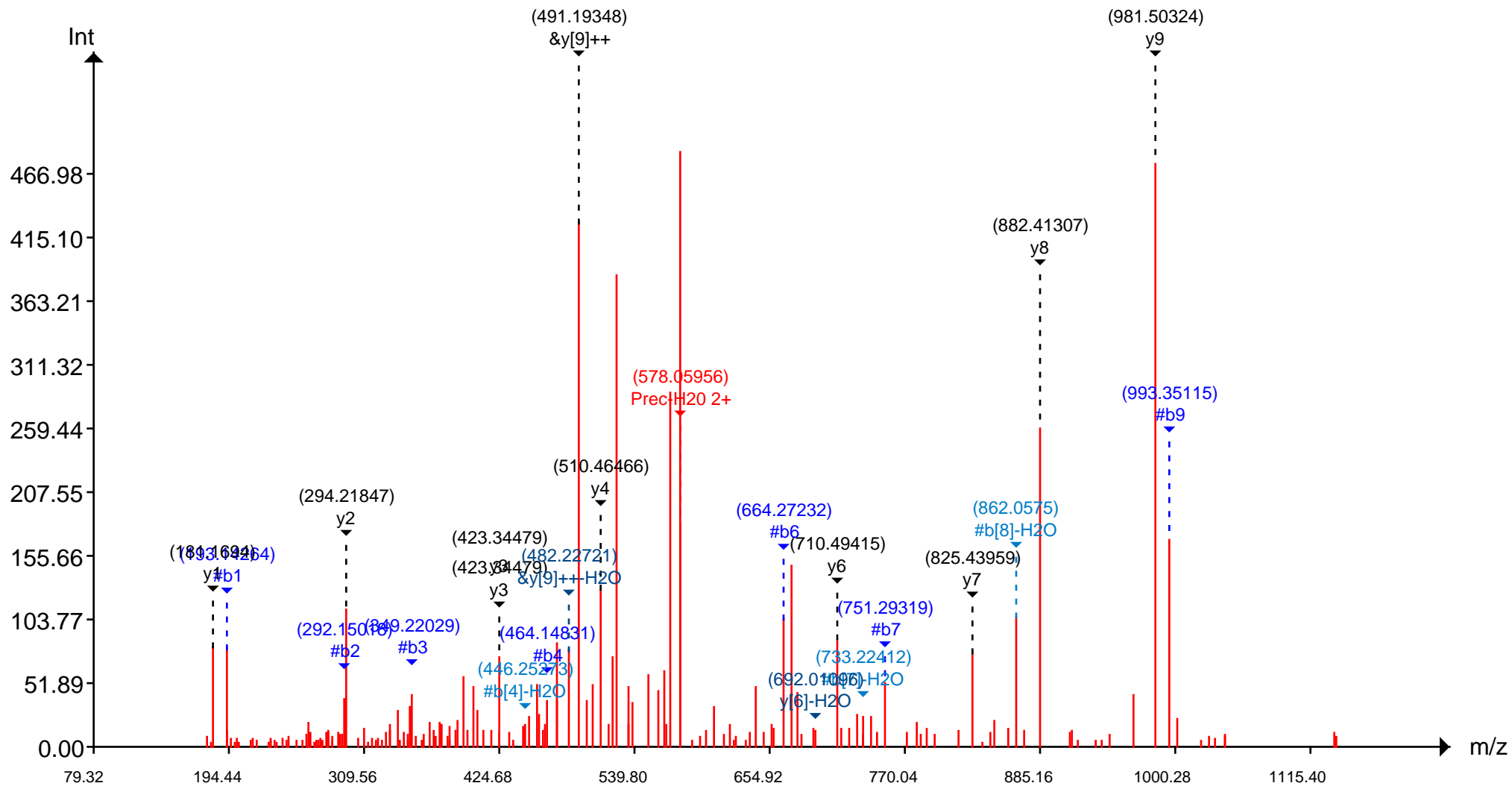
Charge 2+

Mass error -0.003658



Info

Modified Sequence	AcD3-FDK<AcD3K*>DGDGTITTK<AcD3K*>ELGTVM<MoX*>R-COOH	Ion score	99.89
Protein	P62204	Confidence level	99.00%
Precursor m/z	1118.0627	Identity Threshold	47.77
Charge	2+	Mass error	0.002144



Info

Modified Sequence AcD3-M<Mox*>VGDEASELR<C13>-COOH

Protein P61161

Precursor m/z 587.28303

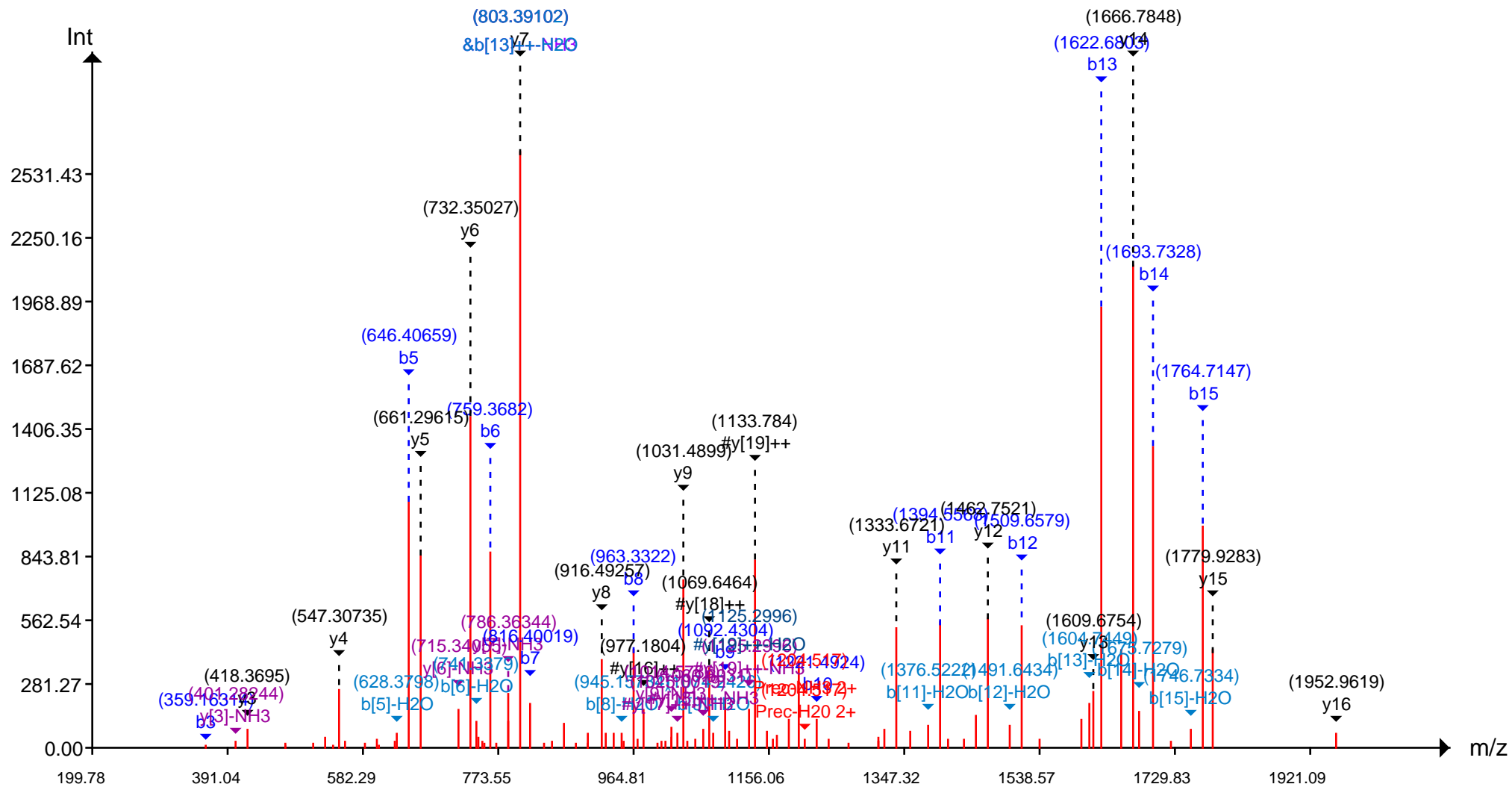
Charge 2+

Ion score 49.0

Confidence level 99.00%

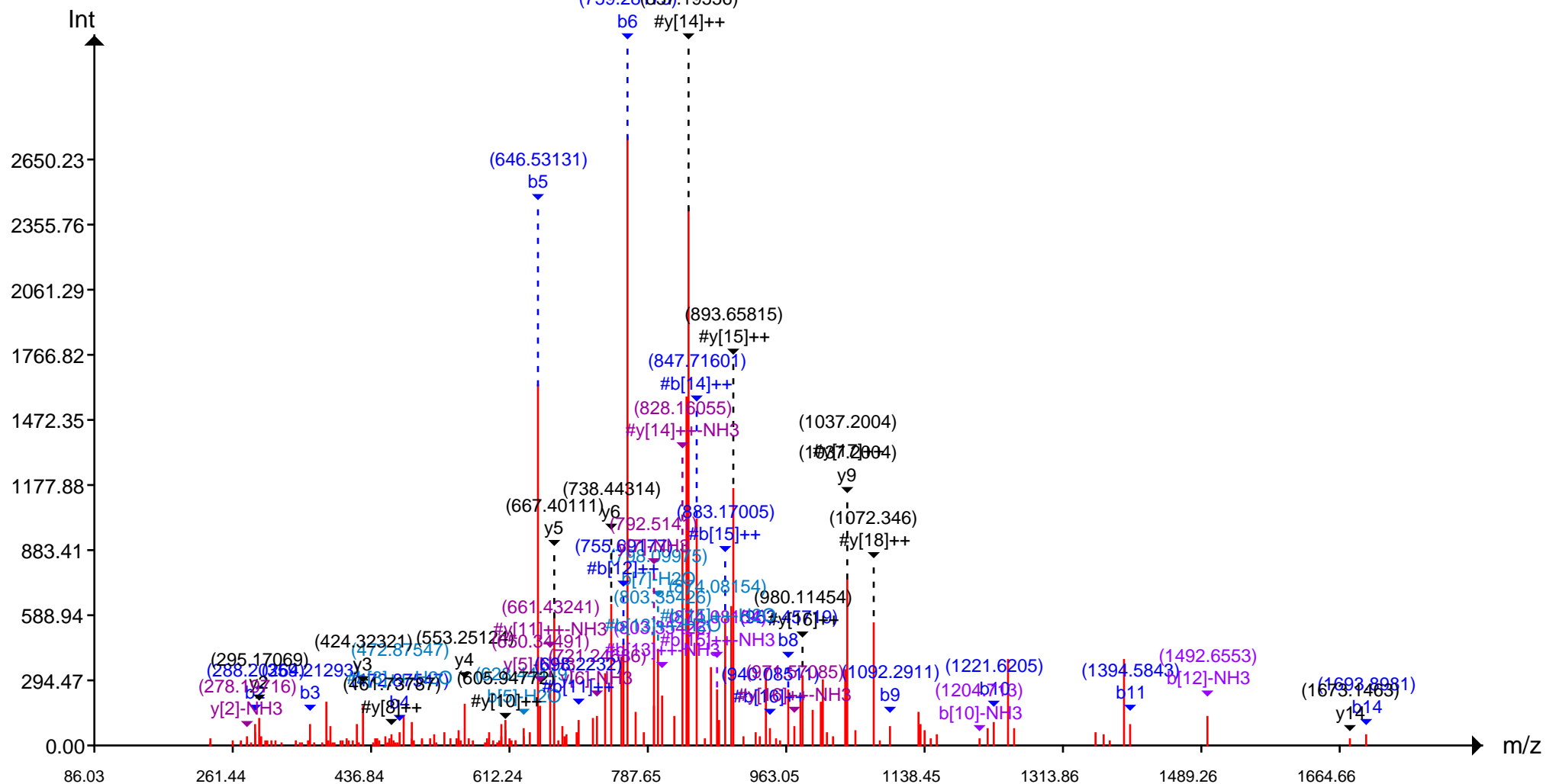
Identity Threshold 46.33

Mass error -2.96E-4



Info

Modified Sequence	AcD3-LEANK<AcD3K*>IGFEEK<AcD3K*>DIAANEENR-COOH	Ion score	115.31
Protein	Q9JJU8	Confidence level	99.00%
Precursor m/z	1213.1145	Identity Threshold	47.87
Charge	2+	Mass error	0.002625



Info

Modified Sequence AcD3-LEANK<AcD3K*>IGFEEK<AcD3K*>DIAANEENR<C13>-COOH

Ion score 51.52

Protein Q9JJU8

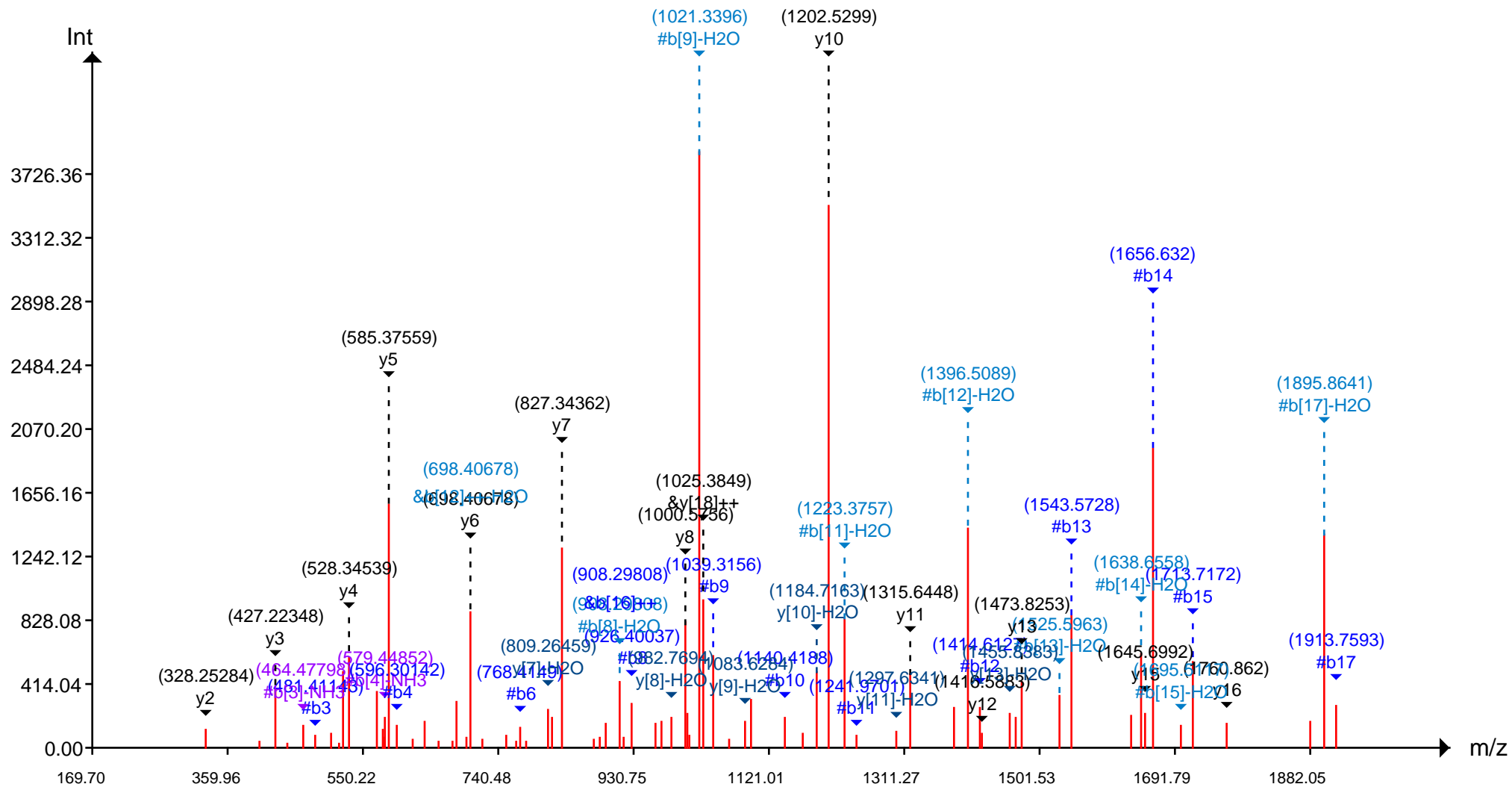
Confidence level 99.00%

Precursor m/z 811.08537

Identity Threshold 47.92

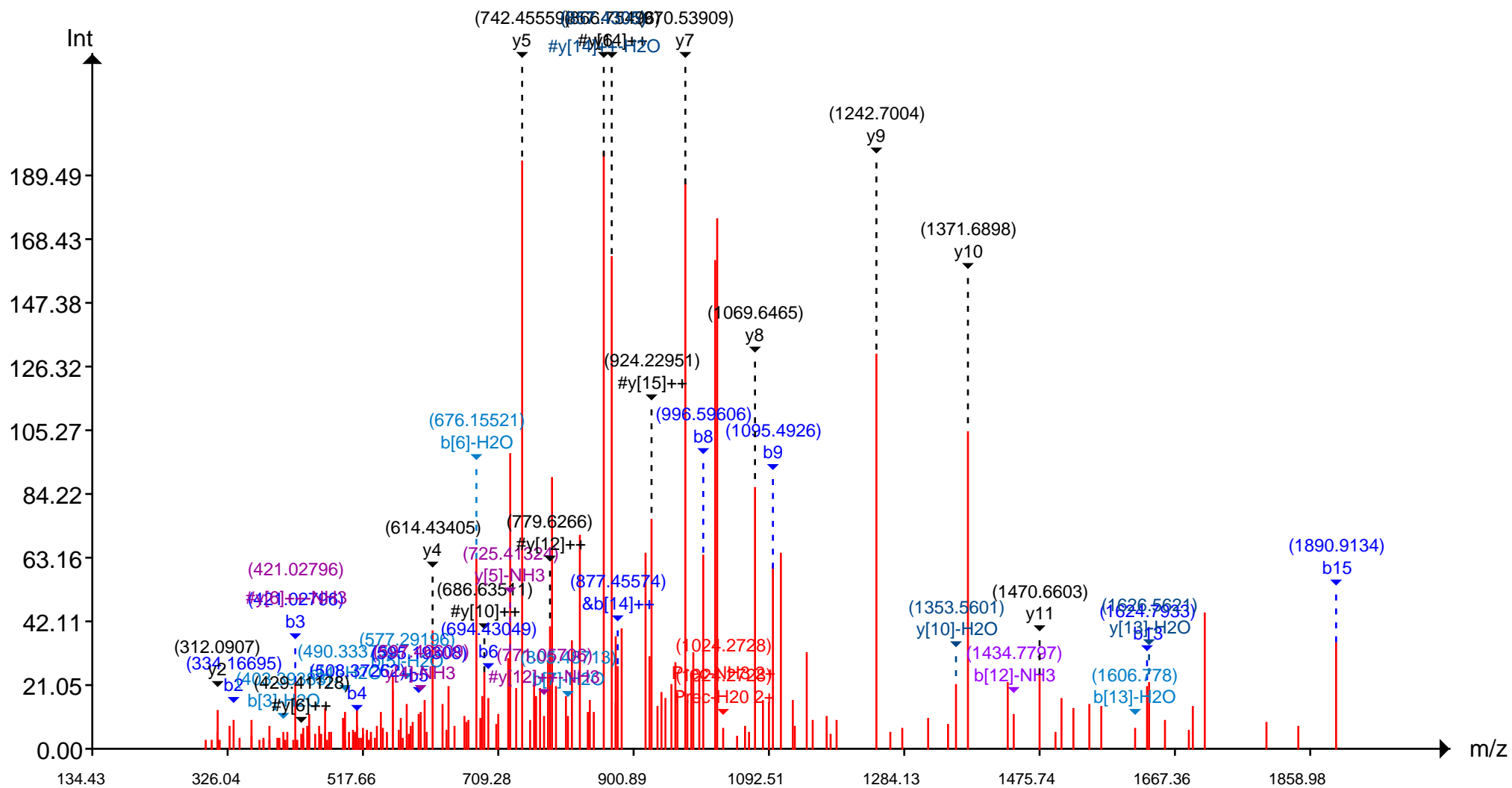
Charge 3+

Mass error 0.002318



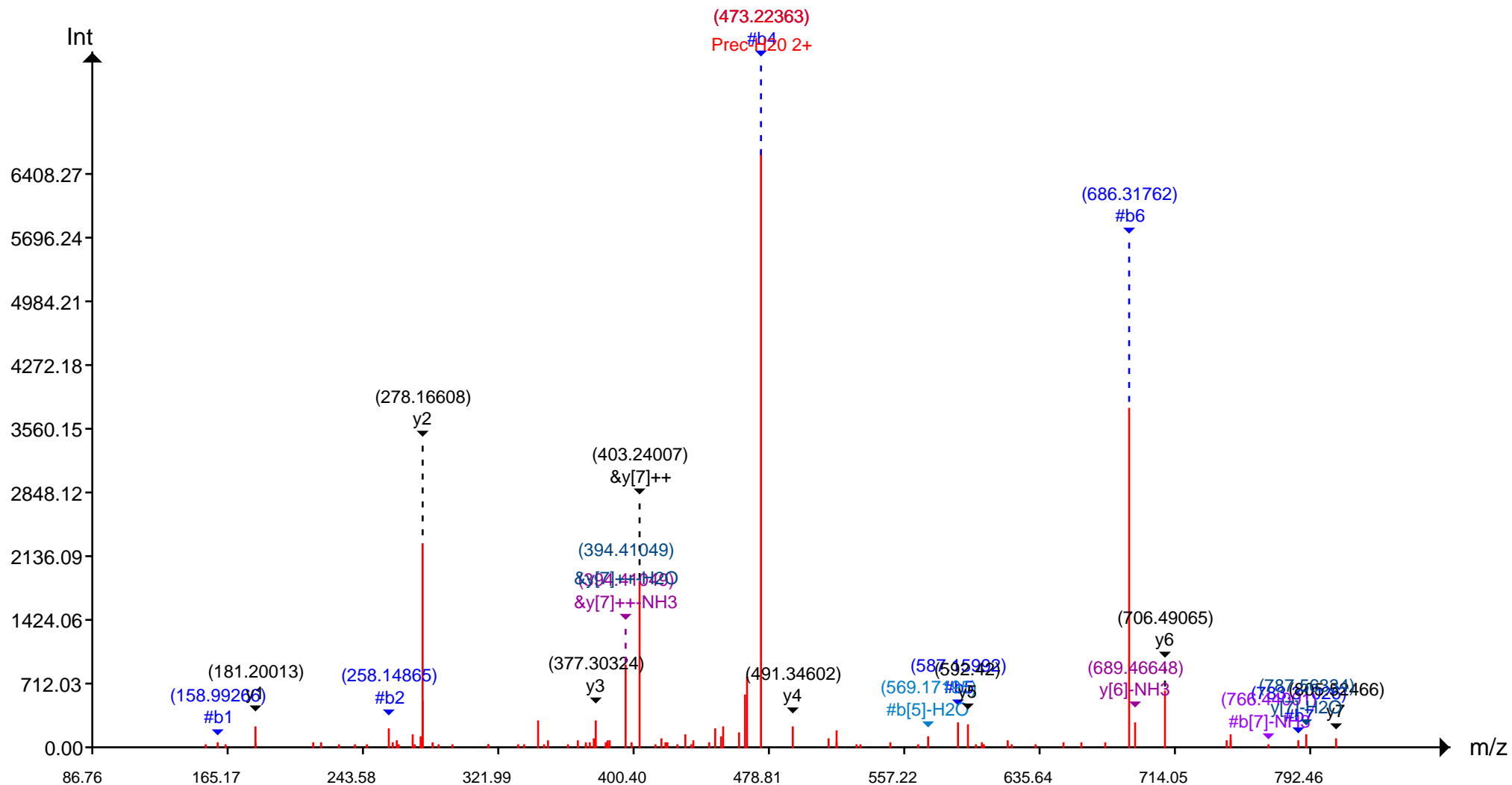
Info

Modified Sequence	AcD3-FDK<AcD3K*>DGDGTITTK<AcD3K*>ELGTVM<MoX*>R<C13>-COOH	Ion score	99.28
Protein	P62204	Confidence level	99.00%
Precursor m/z	1121.0724	Identity Threshold	47.93
Charge	2+	Mass error	0.001403



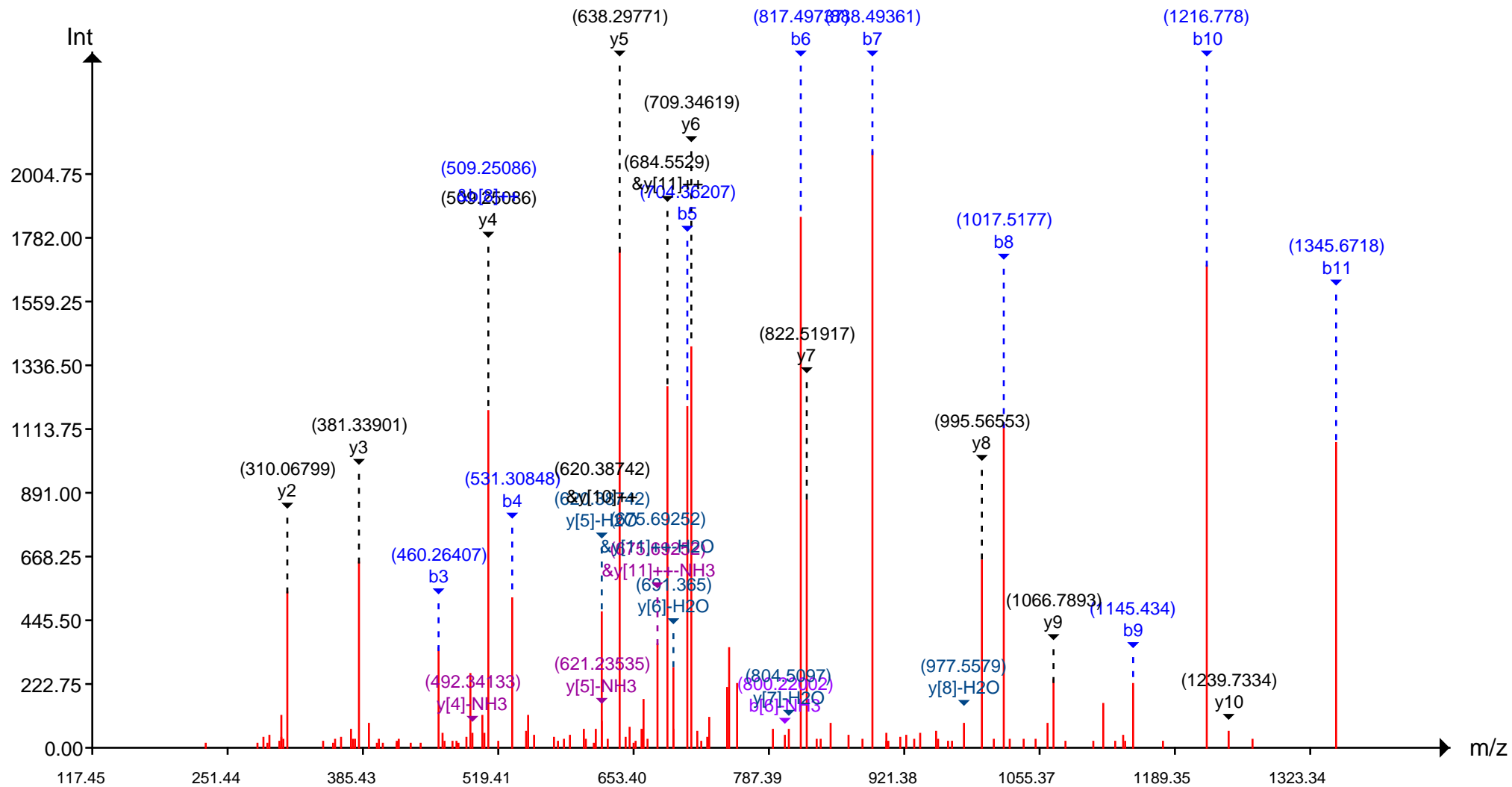
Info

Modified Sequence	AcD3-K<AcD3K*>DSSSVEK<AcD3K*>VLDQK<AcD3K*>EHR-COOH	Ion score	51.38
Protein	Q99020	Confidence level	99.00%
Precursor m/z	1033.0527	Identity Threshold	47.74
Charge	2+	Mass error	0.003171



Info

Modified Sequence	AcD3-IVNTNVPR<C13>-COOH	Ion score	61.12
Protein	P34884	Confidence level	99.00%
Precursor m/z	482.28997	Identity Threshold	45.74
Charge	2+	Mass error	-0.003002



Info

Modified Sequence AcD3-IQK<AcD3K*>AK<AcD3K*>LAEQAER<C13>-COOH

Ion score 81.11

Protein O70456, P68254

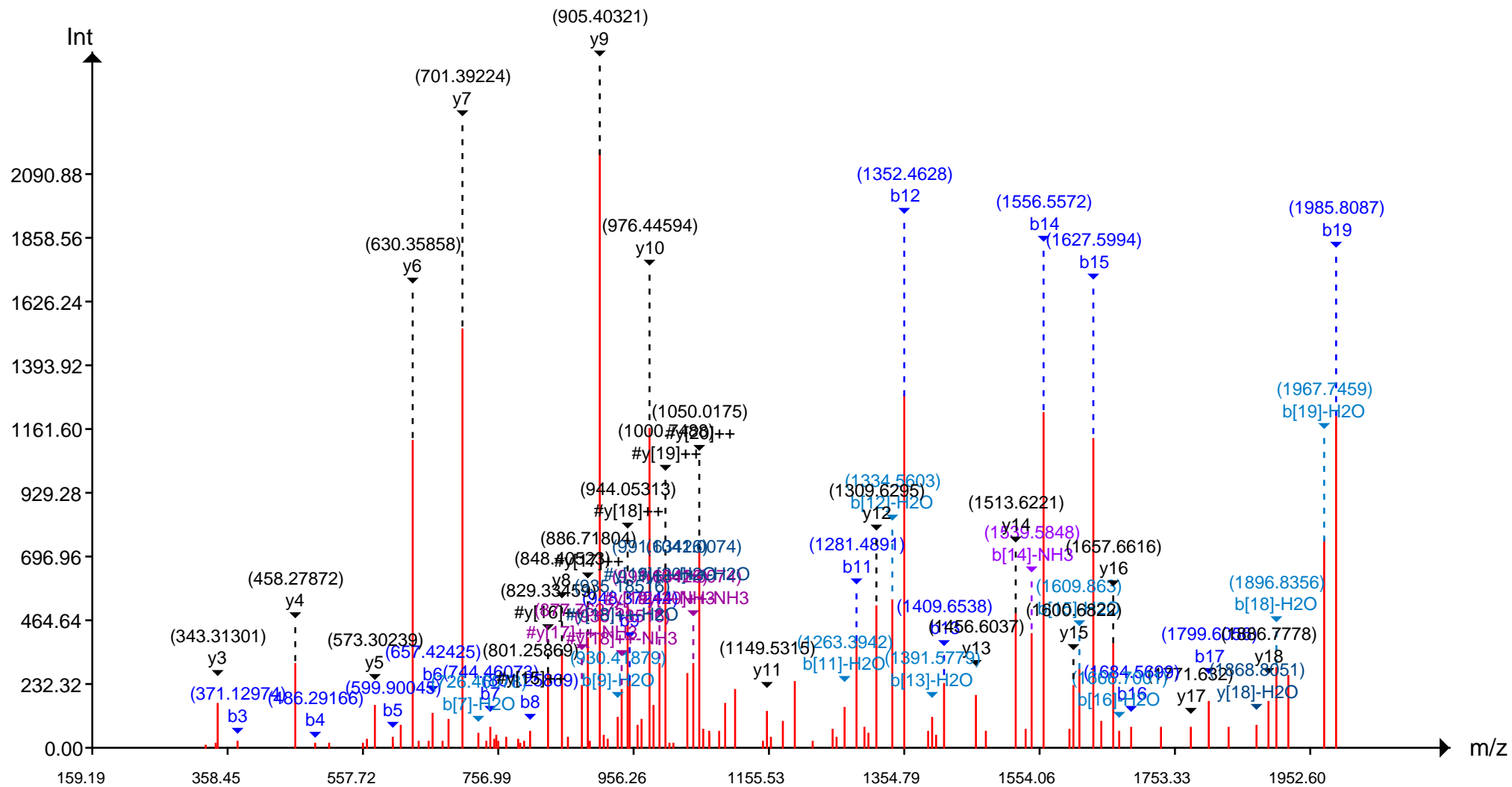
Confidence level 99.00%

Precursor m/z 763.45384

Identity Threshold 44.20

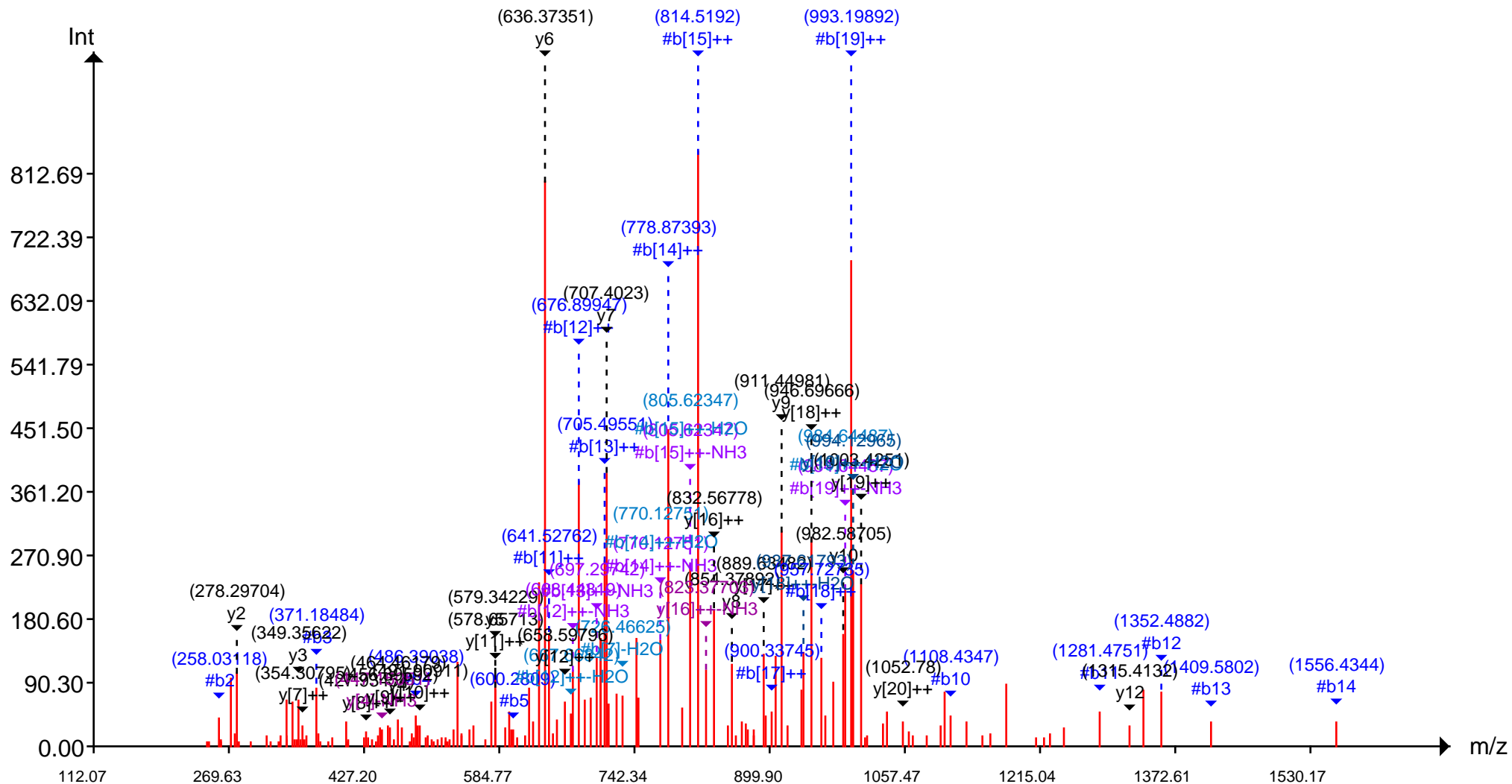
Charge 2+

Mass error 0.001435



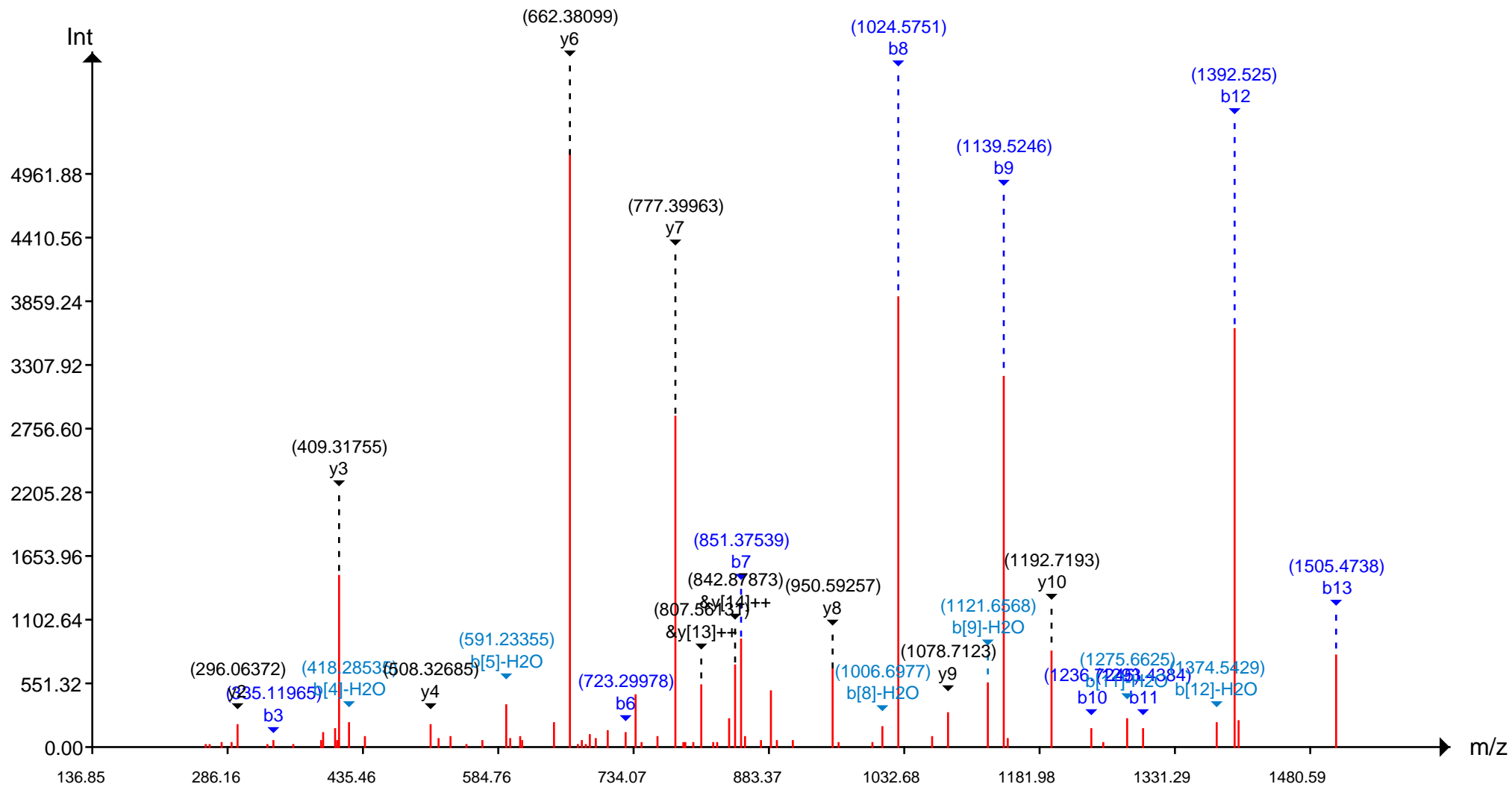
Info

Modified Sequence	AcD3-LVIDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR-COOH	Ion score	141.17
Protein	P63260	Confidence level	99.00%
Precursor m/z	1129.0298	Identity Threshold	45.59
Charge	2+	Mass error	0.002857



Info

Modified Sequence	AcD3-LVIDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH	Ion score	75.23
Protein	P63260	Confidence level	99.00%
Precursor m/z	755.0287	Identity Threshold	46.36
Charge	3+	Mass error	0.00194



Info

Modified Sequence AcD3-AAFTK<AcD3K*>NQK<AcD3K*>DPGVLDK<C13>-COOH

Ion score 77.42

Protein P50543

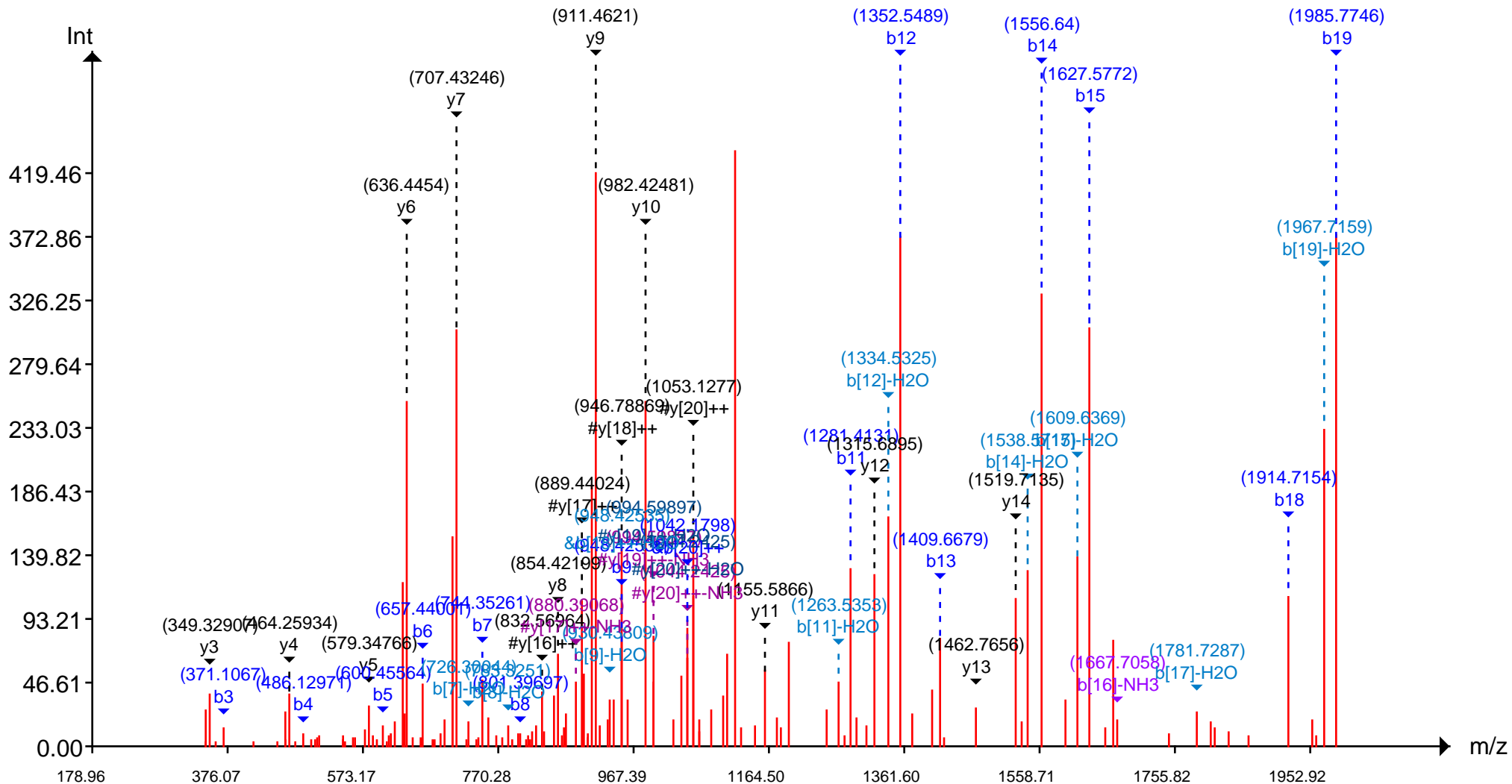
Confidence level 99.00%

Precursor m/z 900.99928

Identity Threshold 47.12

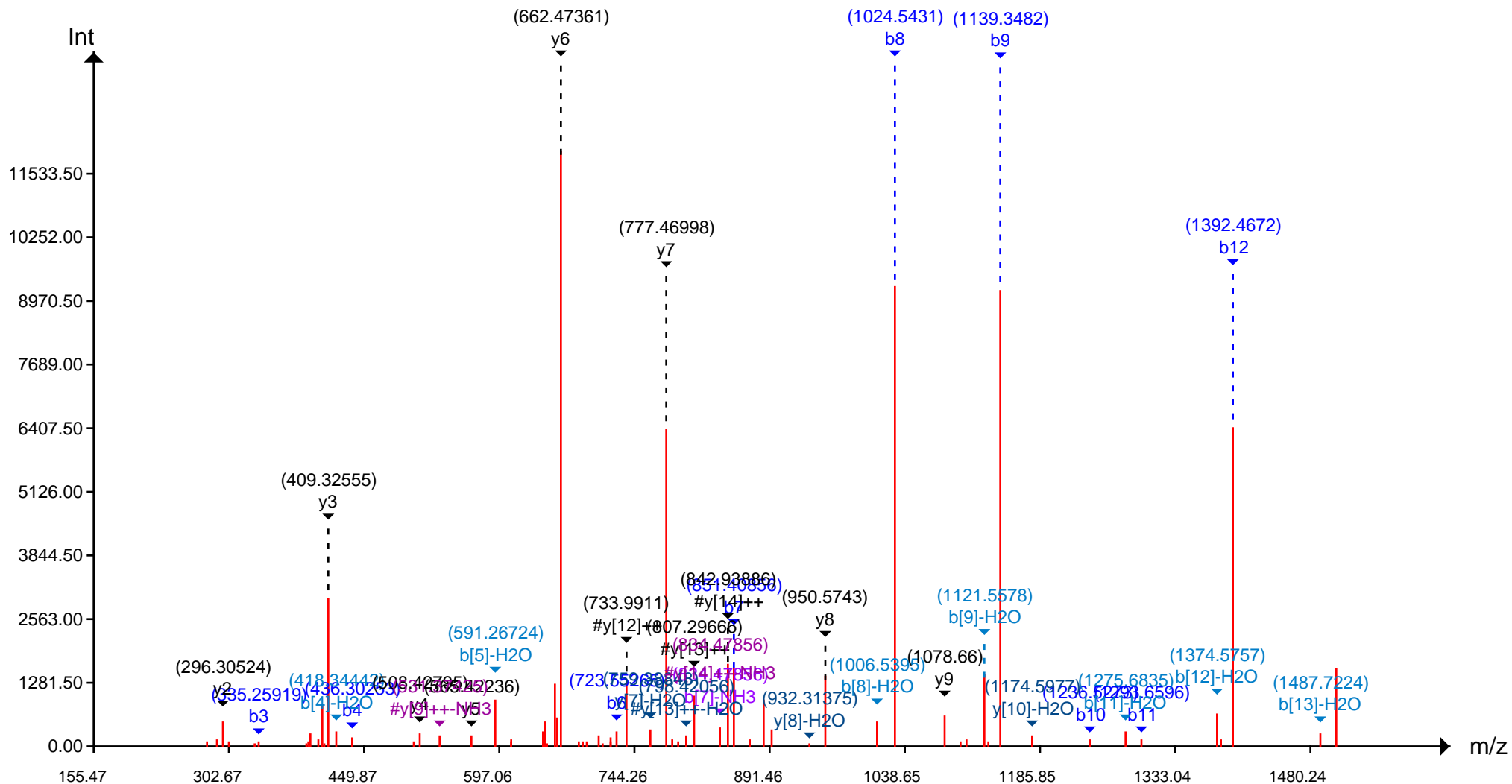
Charge 2+

Mass error 0.001662



Info

Modified Sequence	AcD3-LVIDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH	Ion score	80.56
Protein	P63260	Confidence level	99.00%
Precursor m/z	1132.0399	Identity Threshold	46.36
Charge	2+	Mass error	0.002916



Info

Modified Sequence AcD3-AAFTK<AcD3K*>NQK<AcD3K*>DPGVLDK<C13>-COOH

Ion score 66.61

Protein P50543

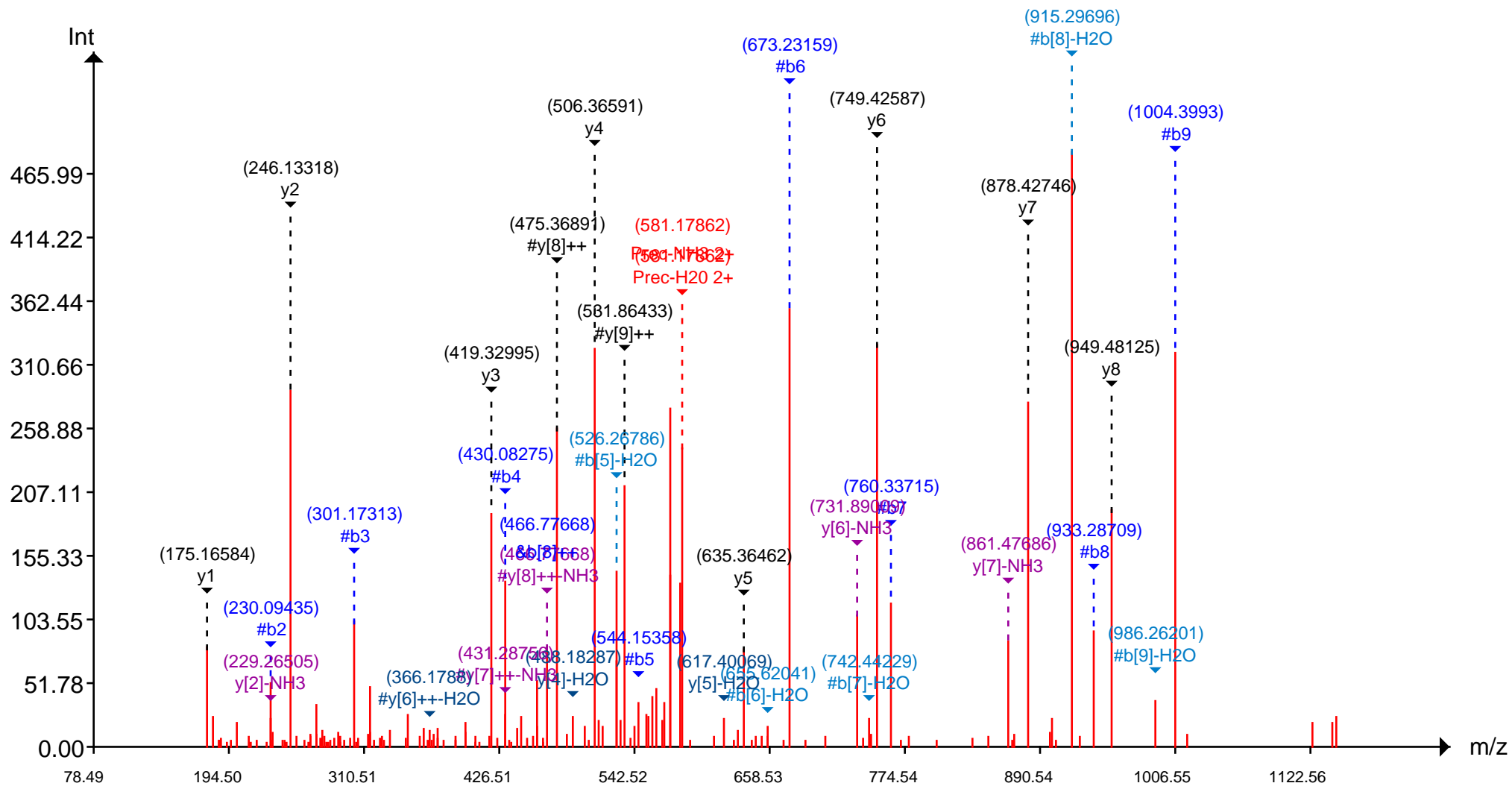
Confidence level 99.00%

Precursor m/z 900.99891

Identity Threshold 47.09

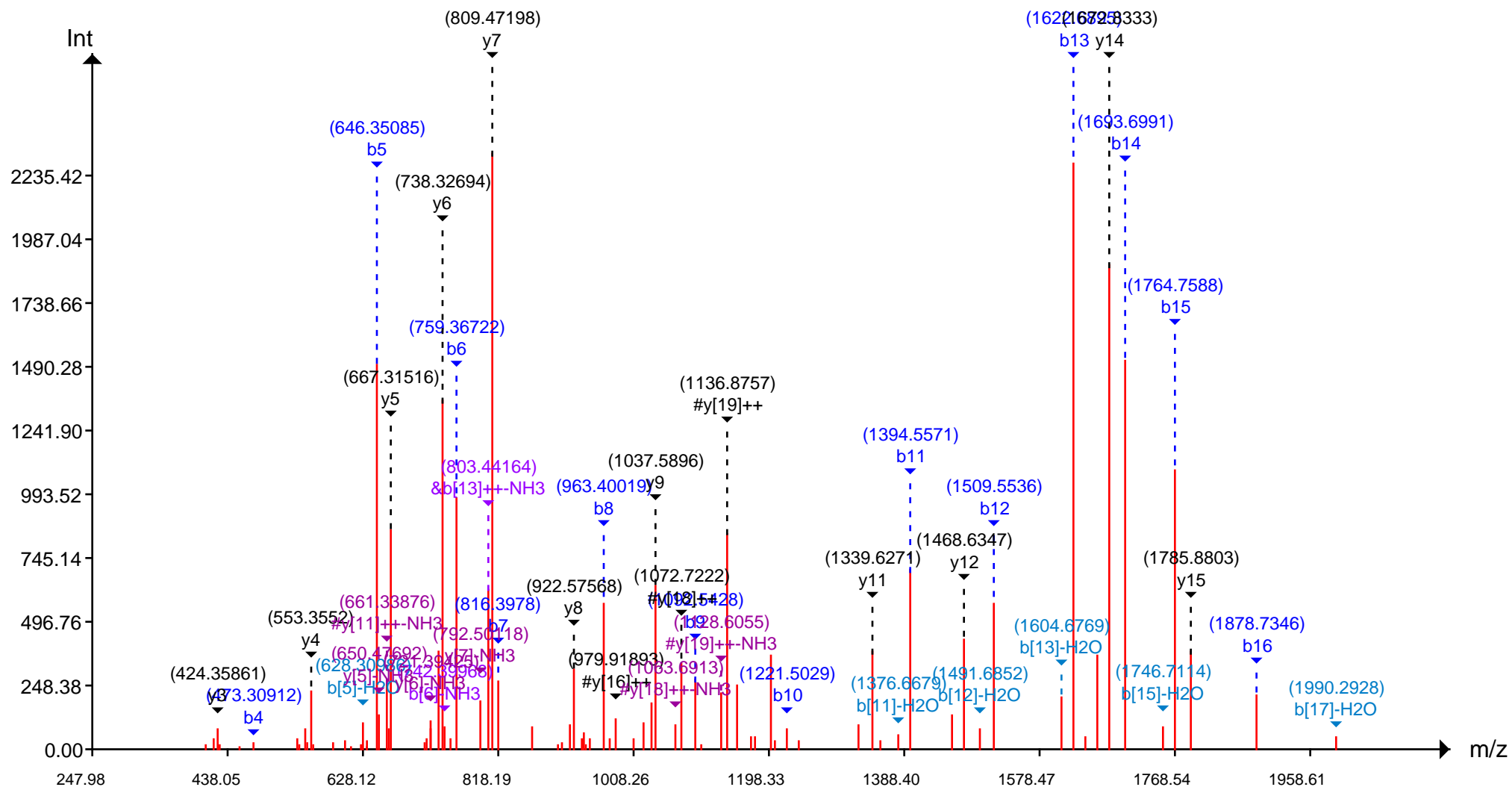
Charge 2+

Mass error 9.22E-4



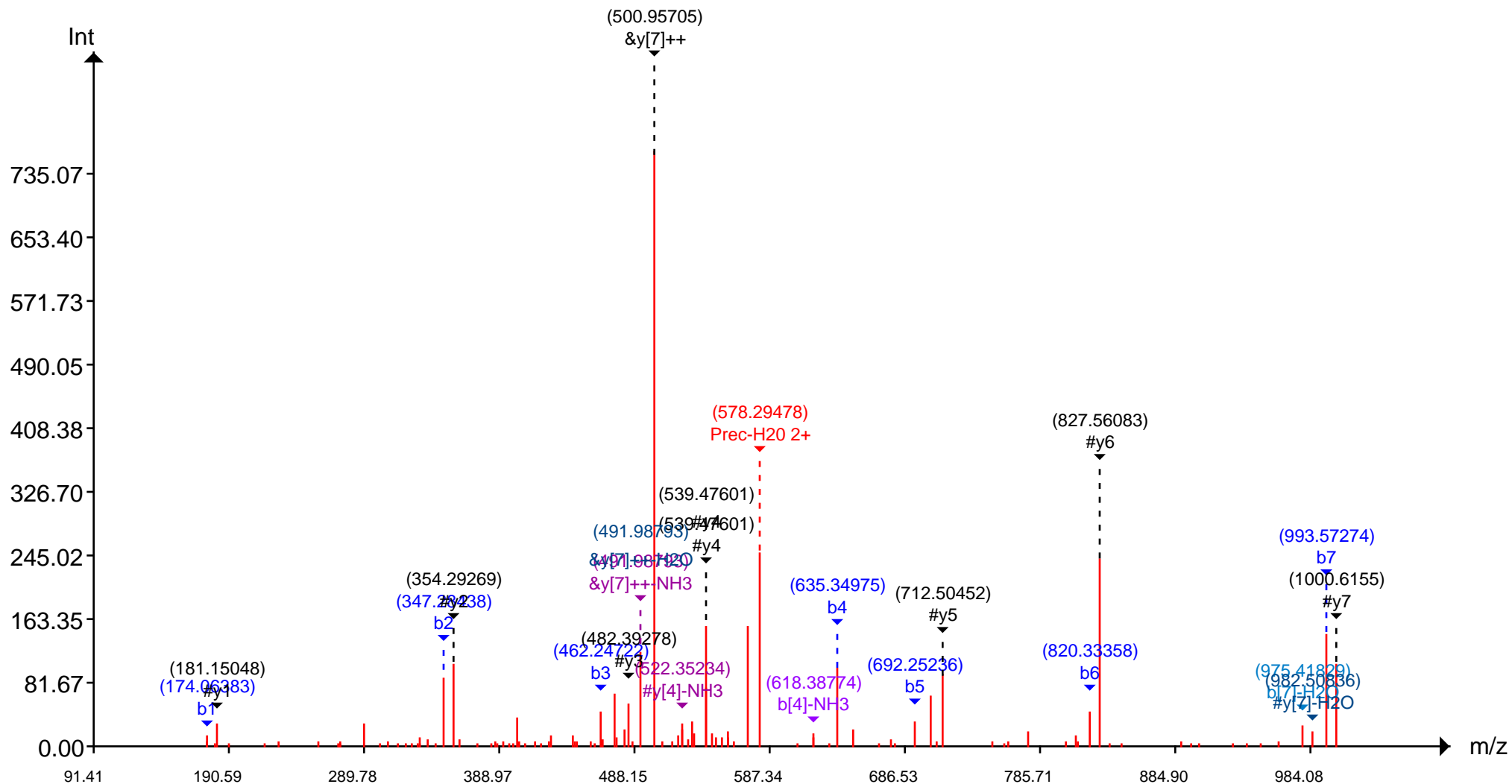
Info

Modified Sequence	AcD3-AIAENESK<AcD3K*>AR-COOH	Ion score	57.98
Protein	Q923D4	Confidence level	99.00%
Precursor m/z	589.81776	Identity Threshold	48.18
Charge	2+	Mass error	4.2E-5



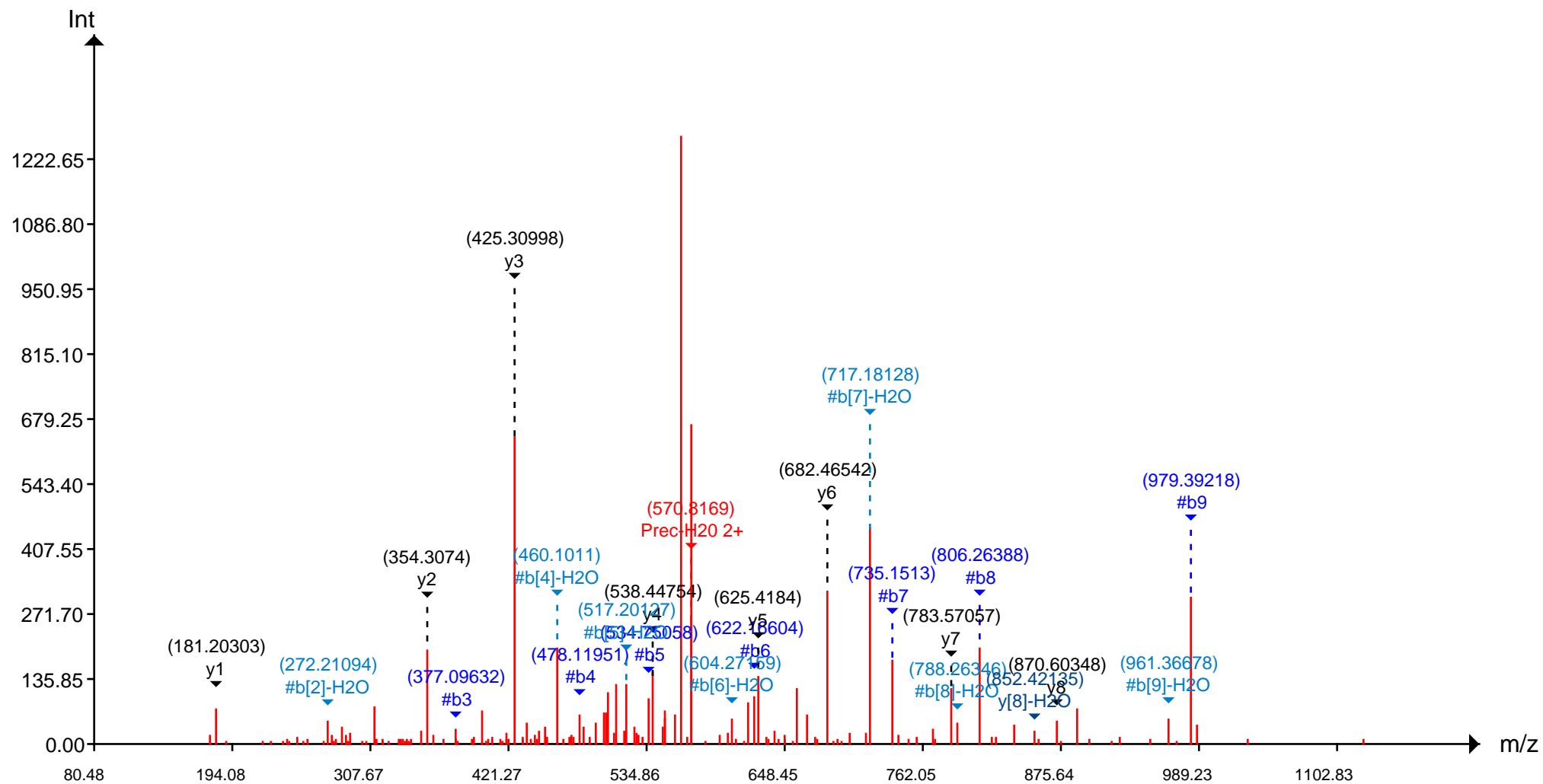
Info

Modified Sequence	AcD3-LEANK<AcD3K*>IGFEEK<AcD3K*>DIAANEENR<C13>-COOH	Ion score	118.56
Protein	Q9JJU8	Confidence level	99.00%
Precursor m/z	1216.1247	Identity Threshold	47.92
Charge	2+	Mass error	0.002884



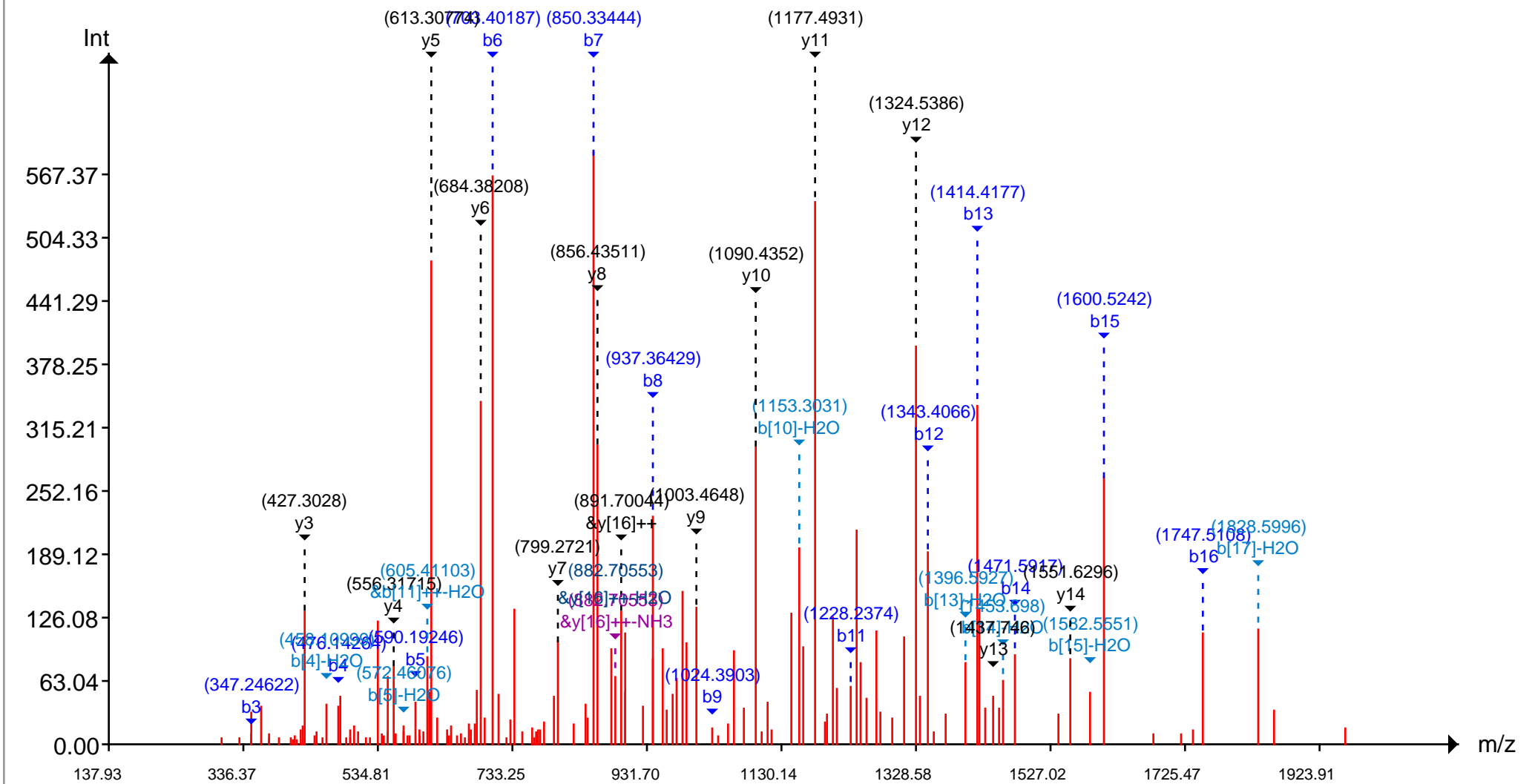
Info

Modified Sequence	AcD3-QK<AcD3K*>DK<AcD3K*>GQK<AcD3K*>R<C13>-COOH	Ion score	46.43
Protein	Q9CT10	Confidence level	99.00%
Precursor m/z	587.357	Identity Threshold	44.42
Charge	2+	Mass error	-3.96E-4



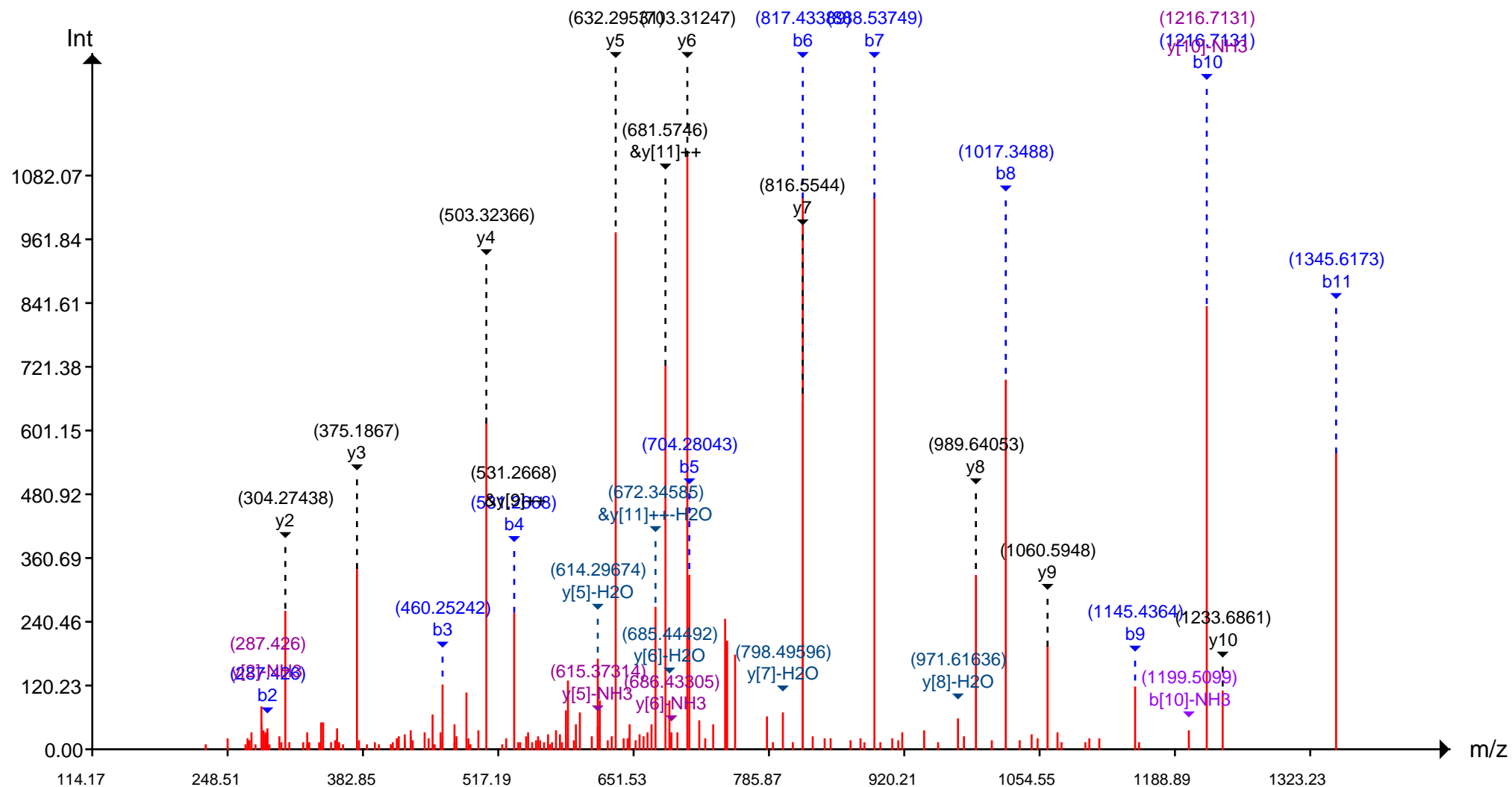
Info

Modified Sequence	AcD3-DESTGSIK<AcD3K*>R<C13>-COOH	Ion score	50.41
Protein	P05064	Confidence level	99.00%
Precursor m/z	580.31202	Identity Threshold	47.73
Charge	2+	Mass error	2.1E-5



Info

Modified Sequence	AcD3-TVTENIFSSM<Mox*>GDAGEM<Mox*>VR<C13>-COOH	Ion score	75.7
Protein	P26039	Confidence level	99.00%
Precursor m/z	1013.966	Identity Threshold	44.67
Charge	2+	Mass error	0.001615



Info

Modified Sequence AcD3-IQK<AcD3K*>AK<AcD3K*>LAEQAER-COOH

Ion score 88.32

Protein O70456, P68254

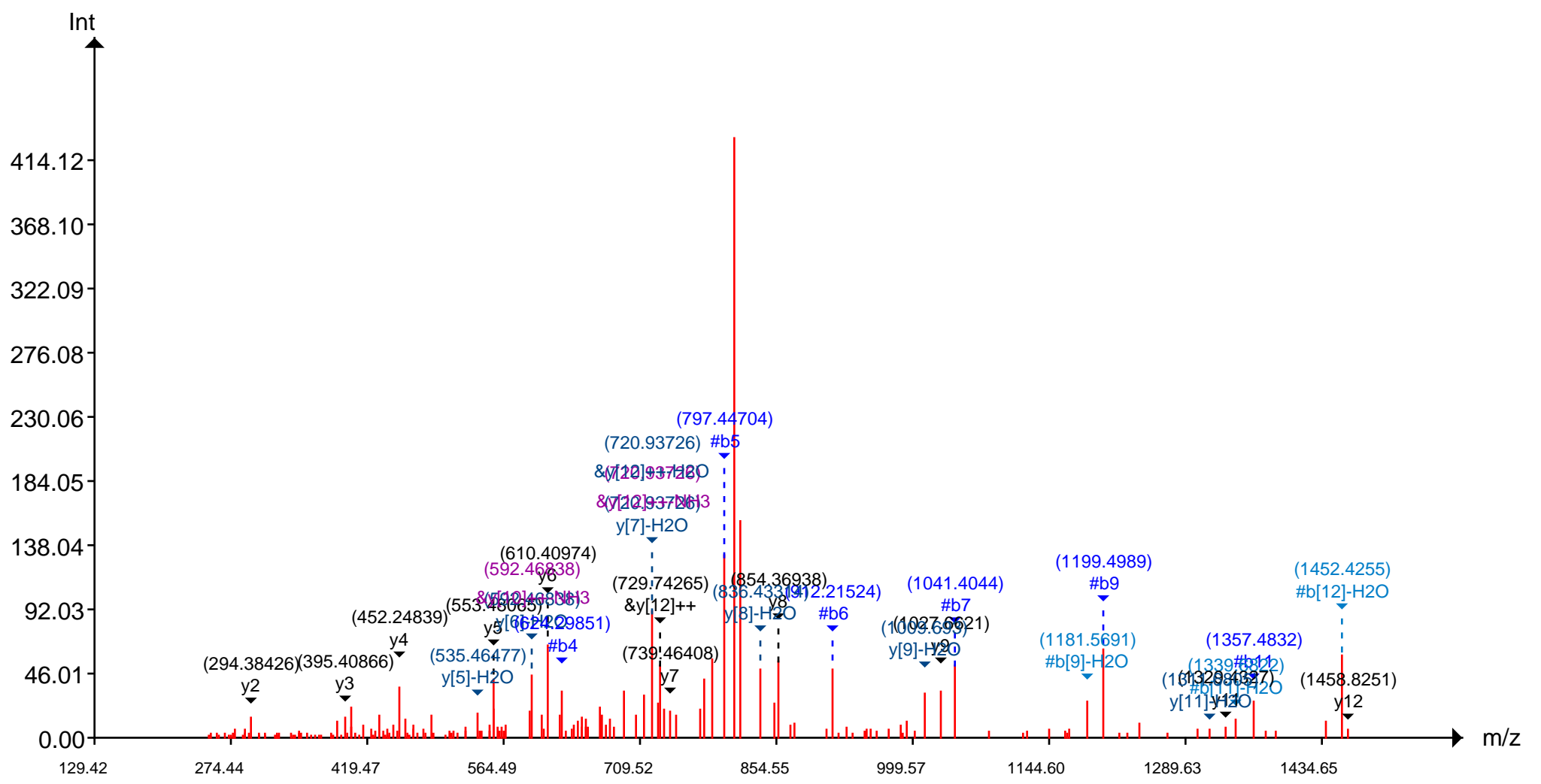
Confidence level 99.00%

Precursor m/z 760.44385

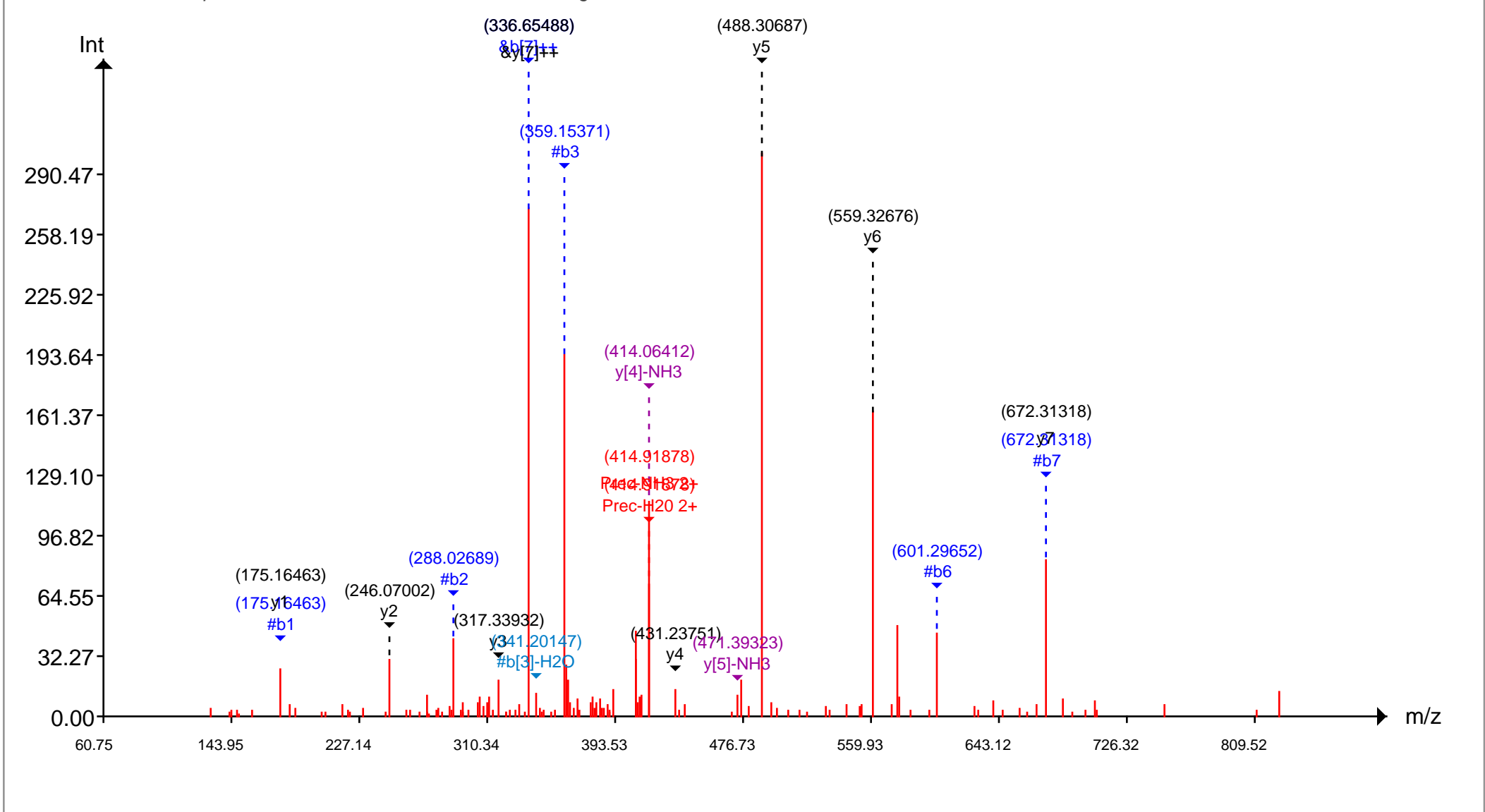
Identity Threshold 45.41

Charge 2+

Mass error 0.001596

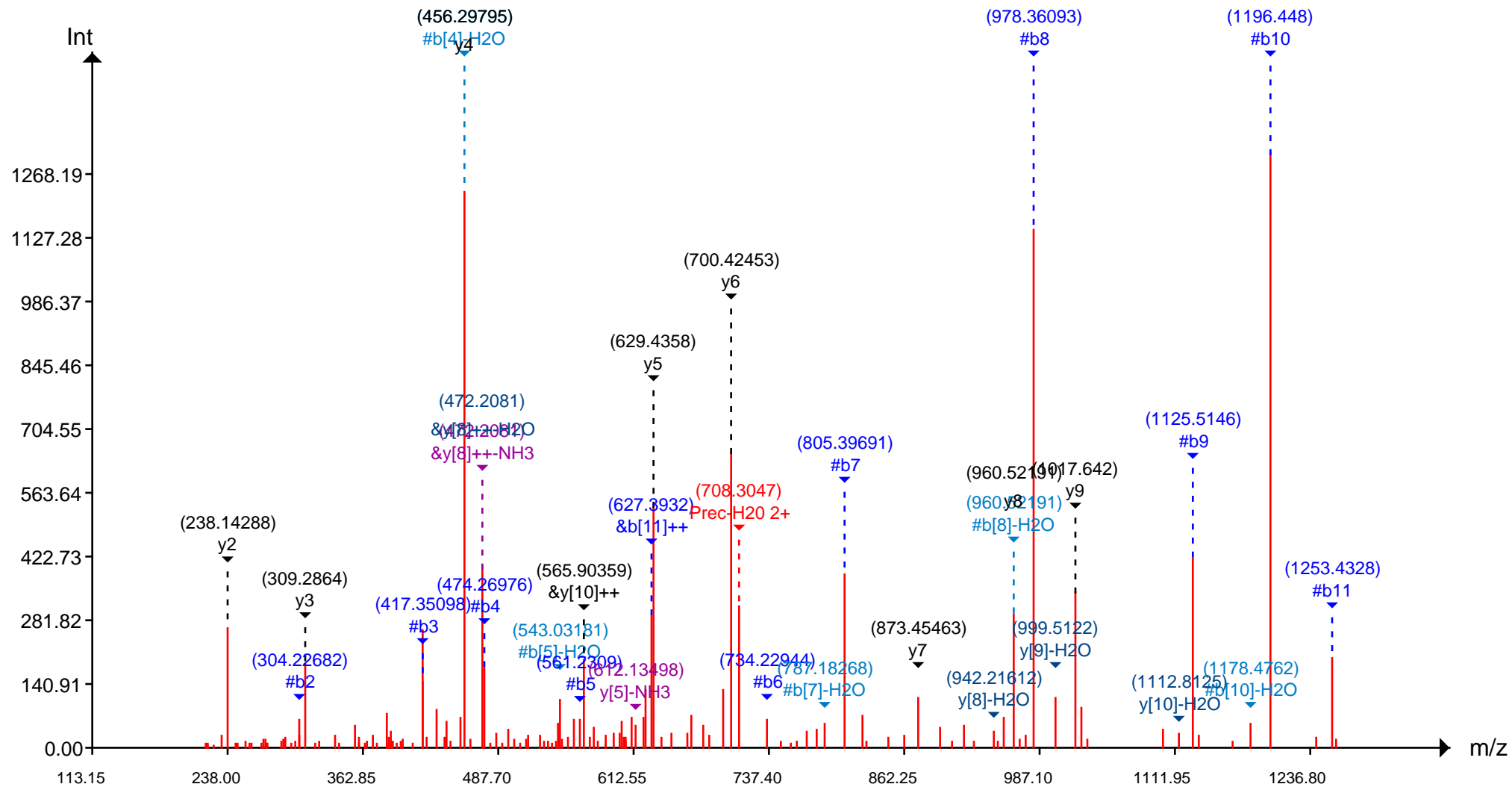


Info			
Modified Sequence	AcD3-M<Mox*>EEK<AcD3K*>K<AcD3K*>DEGTGTLR<C13>-COOH	Ion score	57.99
Protein	P26039	Confidence level	99.00%
Precursor m/z	825.91908	Identity Threshold	47.99
Charge	2+	Mass error	0.00122



Info

Modif...	AcD3-ELAGNAAR-COOH	Ion score
	Q8CGP5, Q8CGP6, Q8CGP7, P22752, Q6GSS7, Q64522, Q64523, Q8BFU2, Q8R1M2, Q8CCK0, P27661, Q9QZQ8	Confidence level
	423.72891	Identity Threshold
	2+	Mass error



Info

Modified Sequence AcD3-EELGSK<AcD3K*>AK<AcD3K*>FAGR<C13>-COOH

Ion score 64.79

Protein P17182

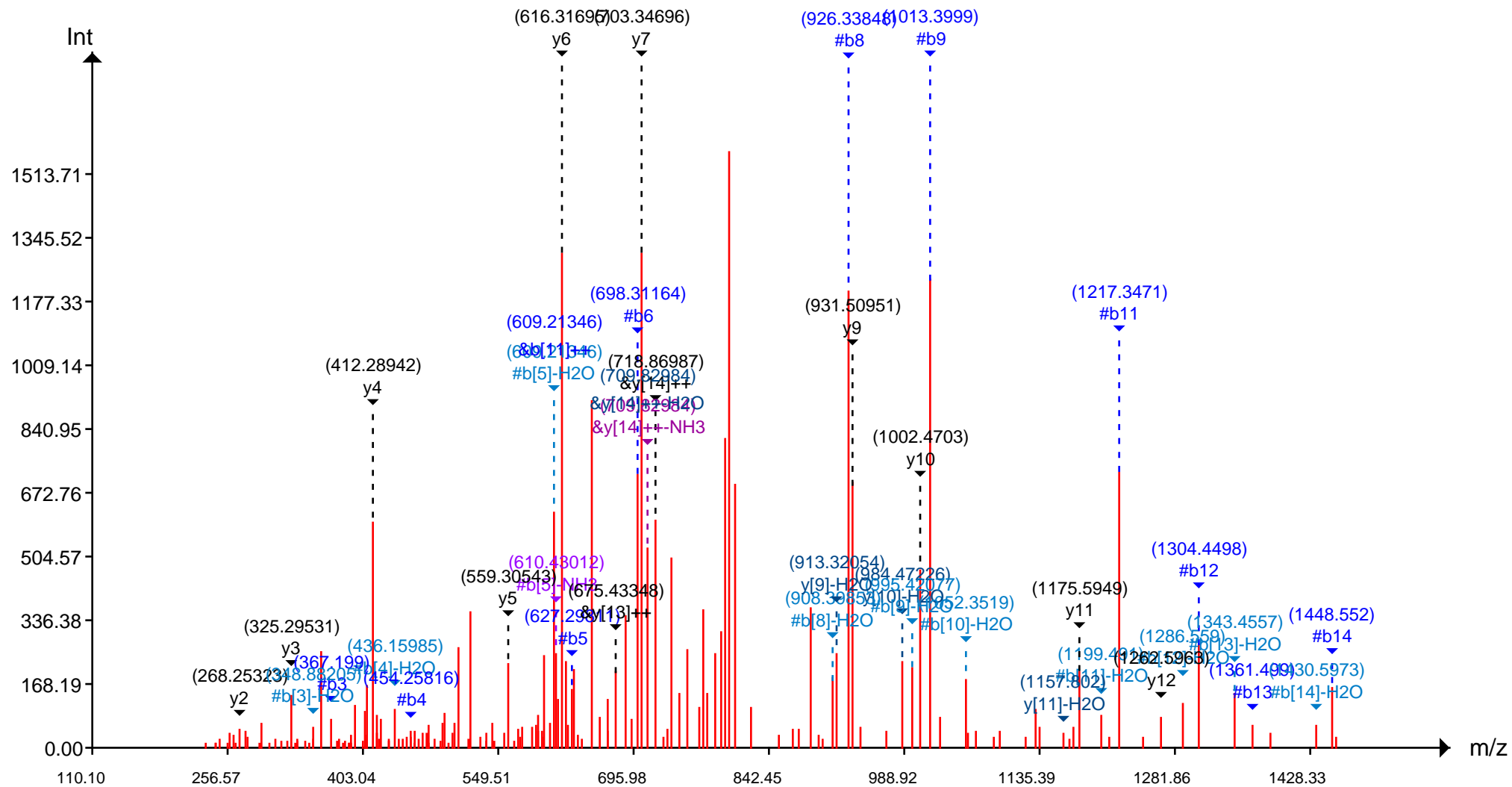
Confidence level 99.00%

Precursor m/z 717.40611

Identity Threshold 47.06

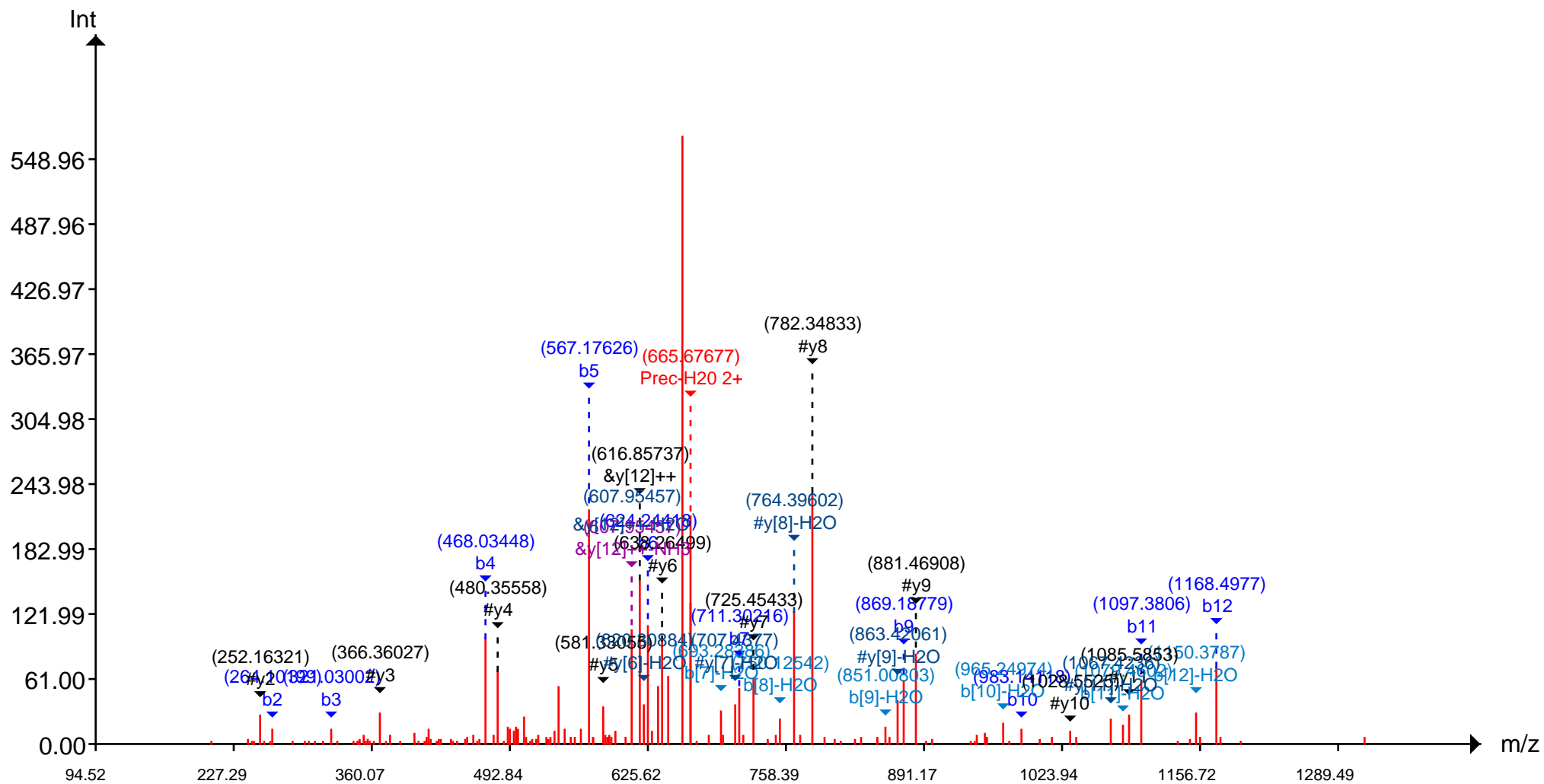
Charge 2+

Mass error 9.15E-4



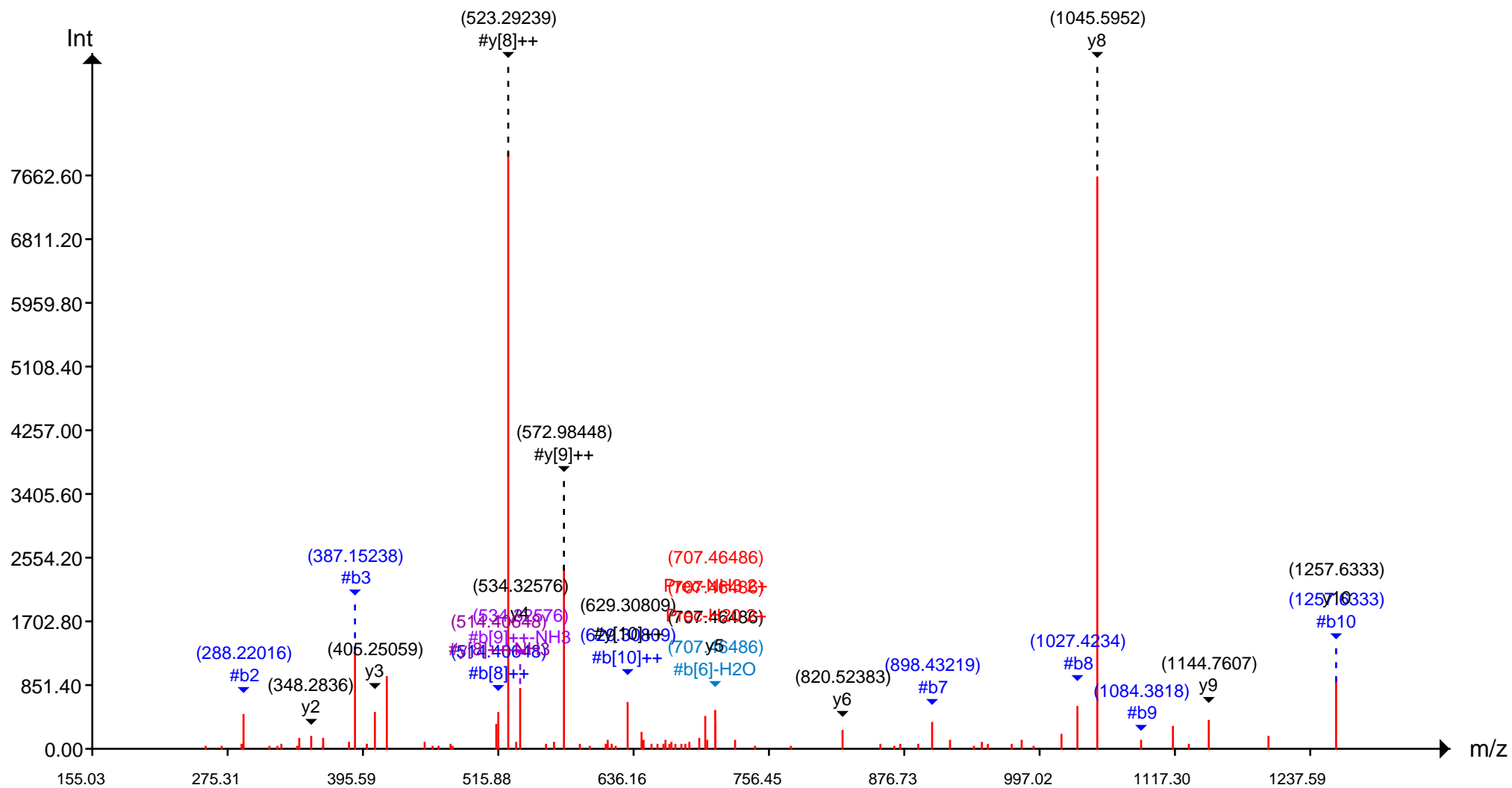
Info

Modified Sequence	AcD3-FSSSK<AcD3K*>ADLSGM<Mox*>SGSR<C13>-COOH	Ion score	62.99
Protein	Q9D154, Q5SV42	Confidence level	99.00%
Precursor m/z	814.89411	Identity Threshold	46.74
Charge	2+	Mass error	0.00107



Info

Modified Sequence	AcD3-AM<Mox*>GM<Mox*>VGS GTNNAR<C13>-COOH	Ion score	72.78
Protein	Q8VDM4	Confidence level	99.00%
Precursor m/z	674.80901	Identity Threshold	44.71
Charge	2+	Mass error	-0.001131



Info

Modified Sequence AcD3-EIVPQIK<AcD3K*>EGK<AcD3K*>R-COOH

Ion score 56.65

Protein Q9DBJ1

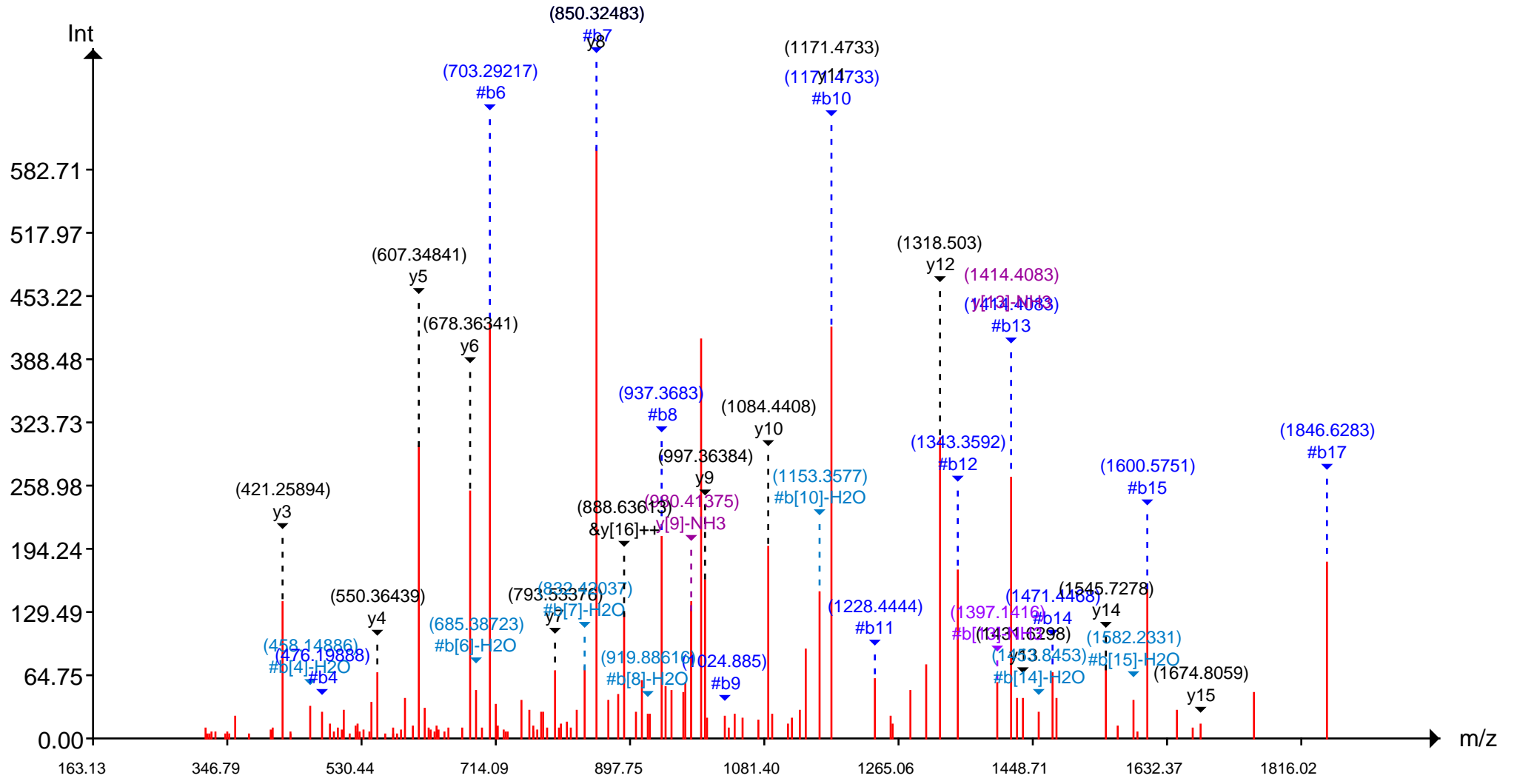
Confidence level 99.00%

Precursor m/z 716.42989

Identity Threshold 44.04

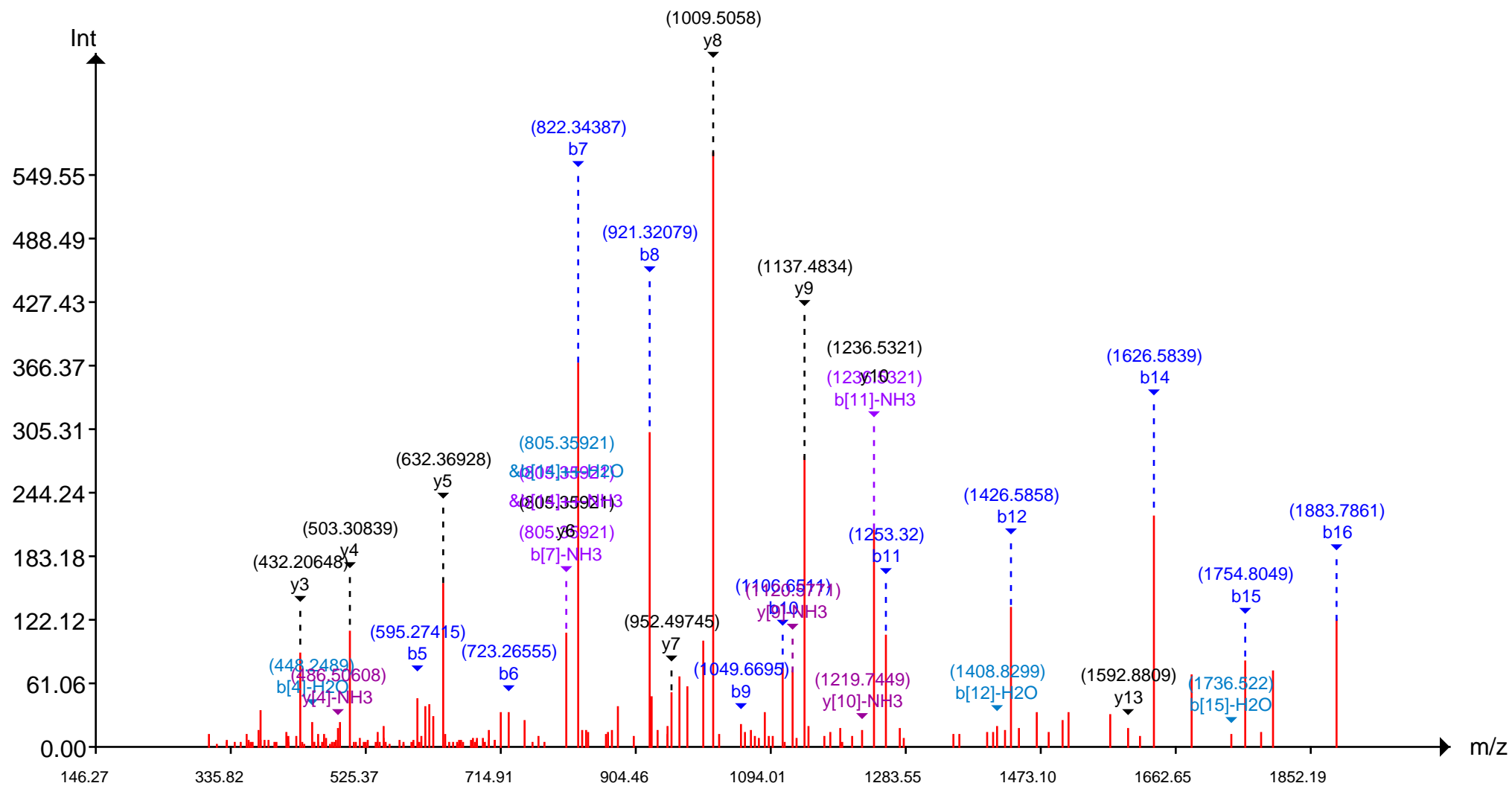
Charge 2+

Mass error 9.29E-4



Info

Modified Sequence	AcD3-TVTENIFSSM<Mox*>GDAGEM<Mox*>VR-COOH	Ion score	97.95
Protein	P26039	Confidence level	99.00%
Precursor m/z	1010.9558	Identity Threshold	43.54
Charge	2+	Mass error	0.001357



Info

Modified Sequence AcD3-YVSAEQVVQGM<Mox*>K<AcD3K*>EAQER-COOH

Ion score 66.72

Protein Q9CQU0

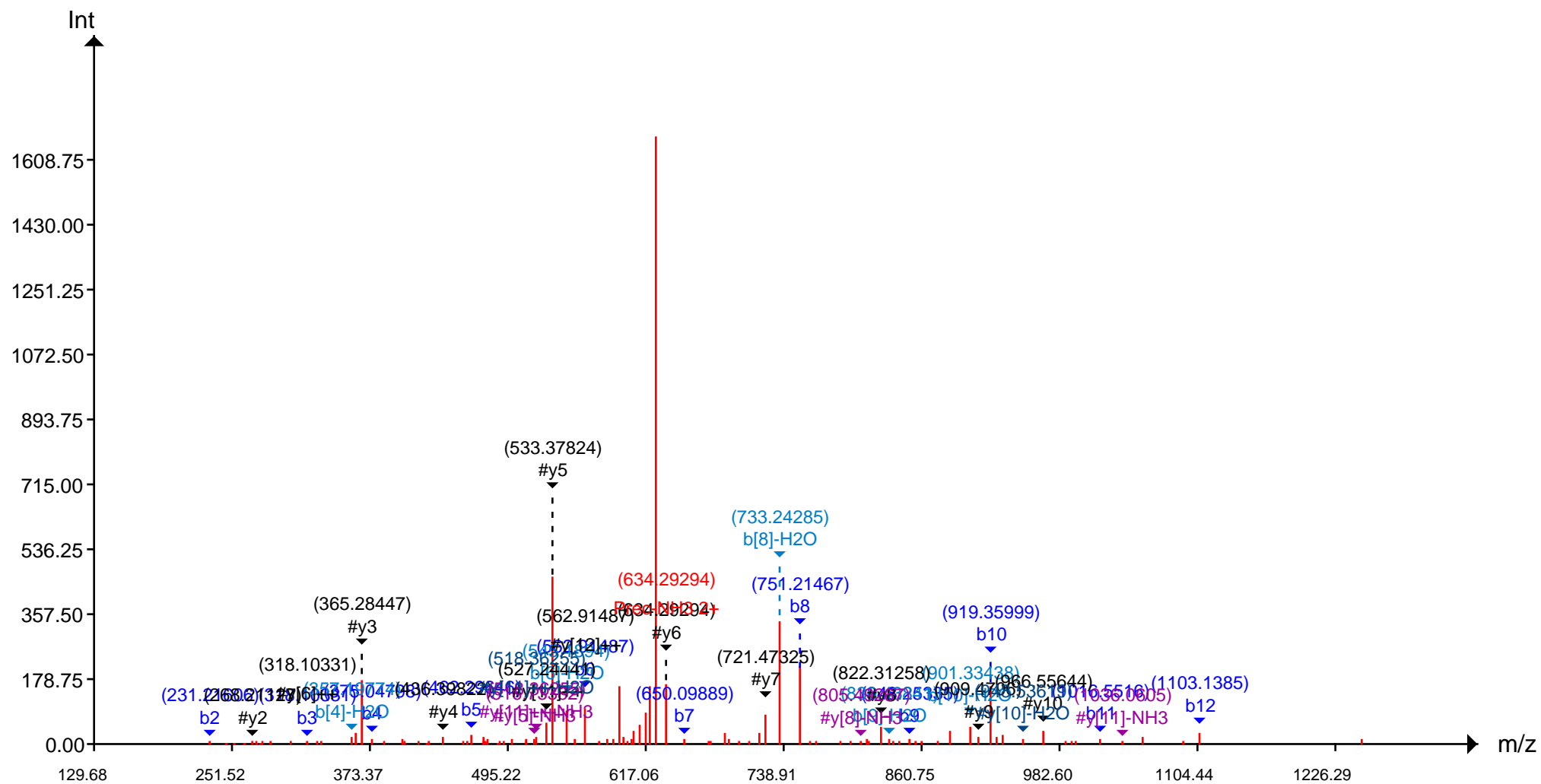
Confidence level 99.00%

Precursor m/z 1029.5089

Identity Threshold 47.49

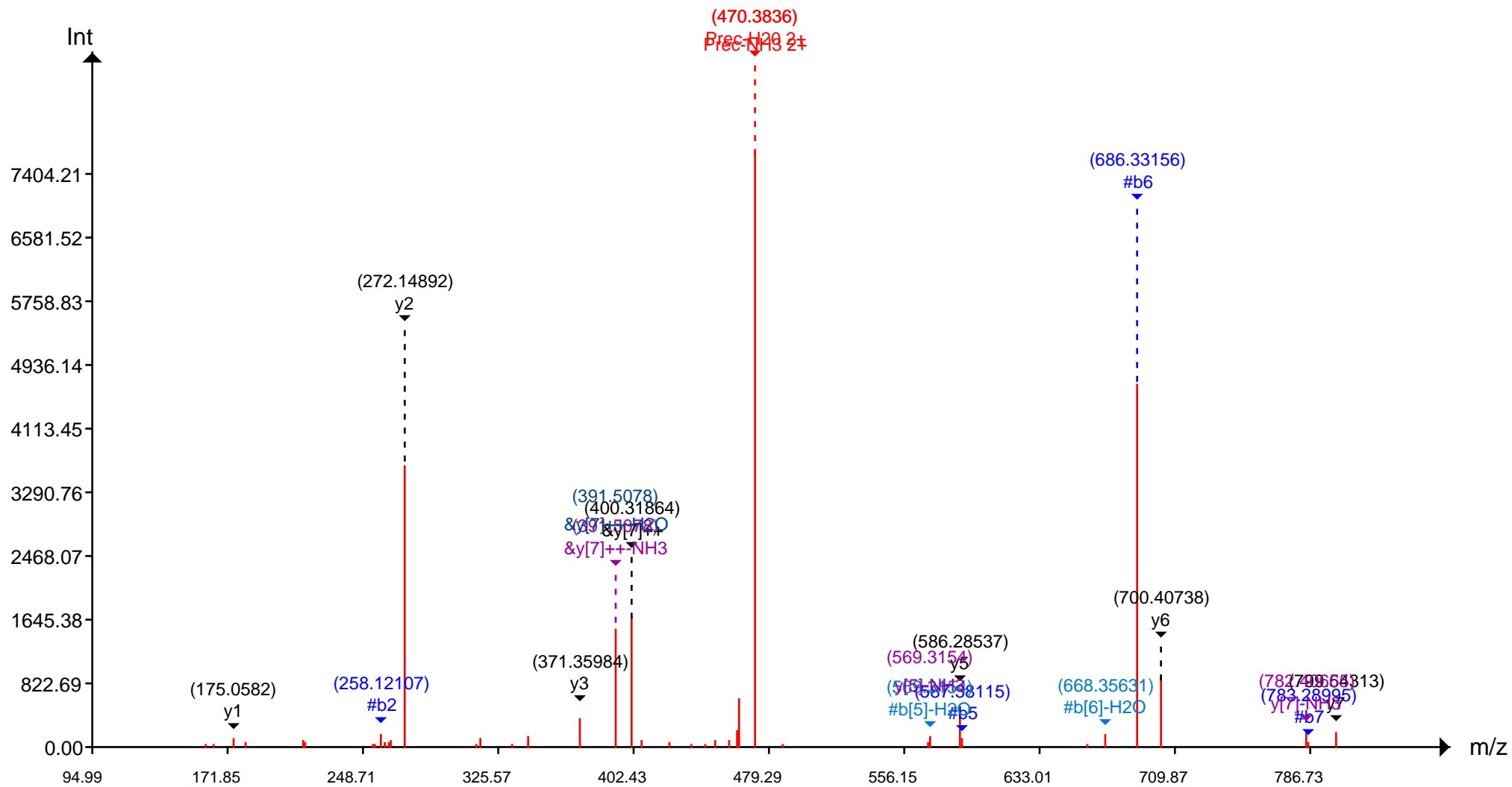
Charge 2+

Mass error 0.002638



Info

Modified Sequence	AcD3-NASGSTSTPAPSR<C13>-COOH	Ion score	75.78
Protein	Q91V92	Confidence level	99.00%
Precursor m/z	642.32153	Identity Threshold	47.35
Charge	2+	Mass error	-3.07E-4



Info

Modified Sequence AcD3-IVNTNVPR-COOH

Ion score 51.93

Protein P34884

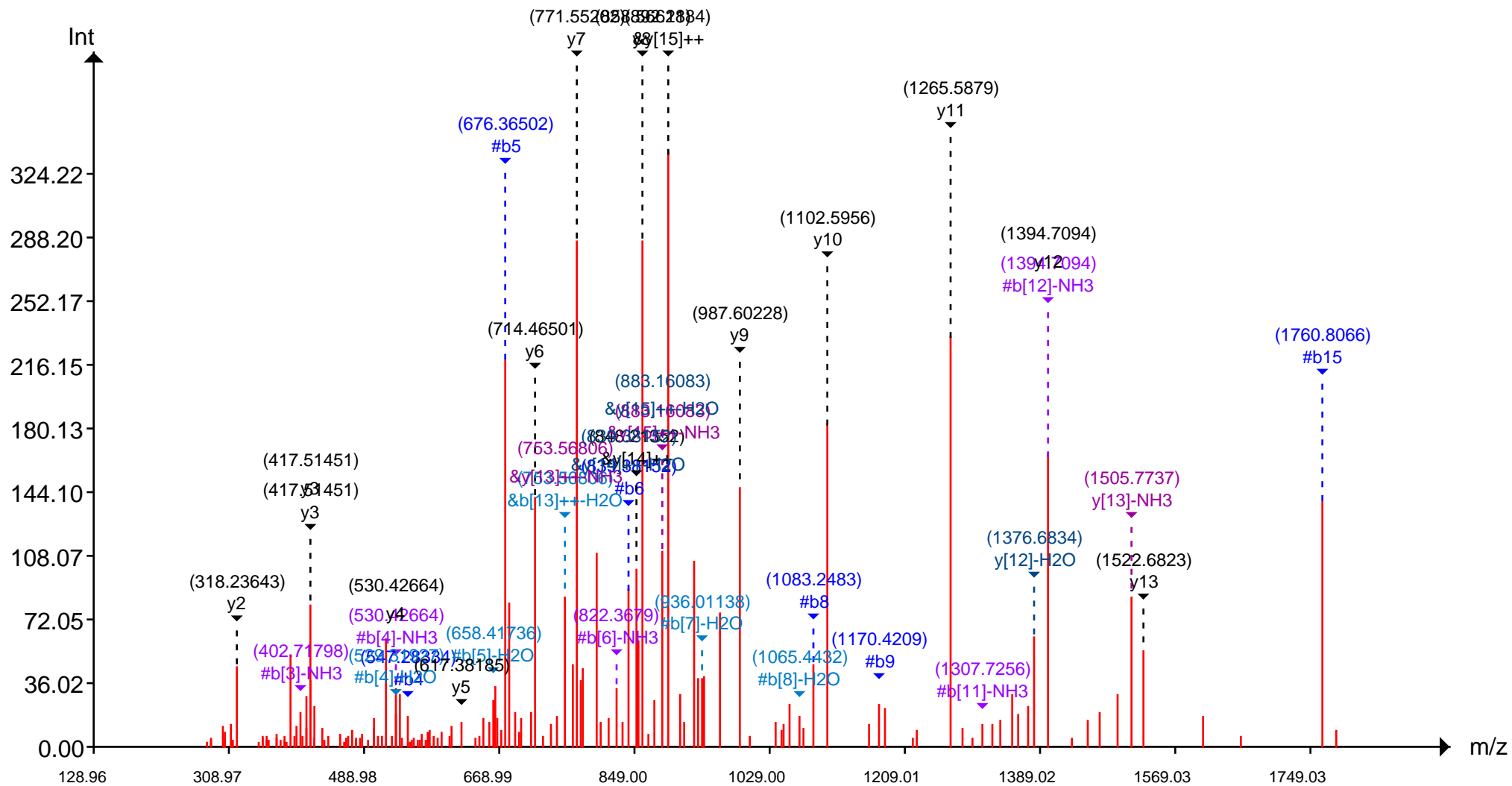
Confidence level 99.00%

Precursor m/z 479.28149

Identity Threshold 46.47

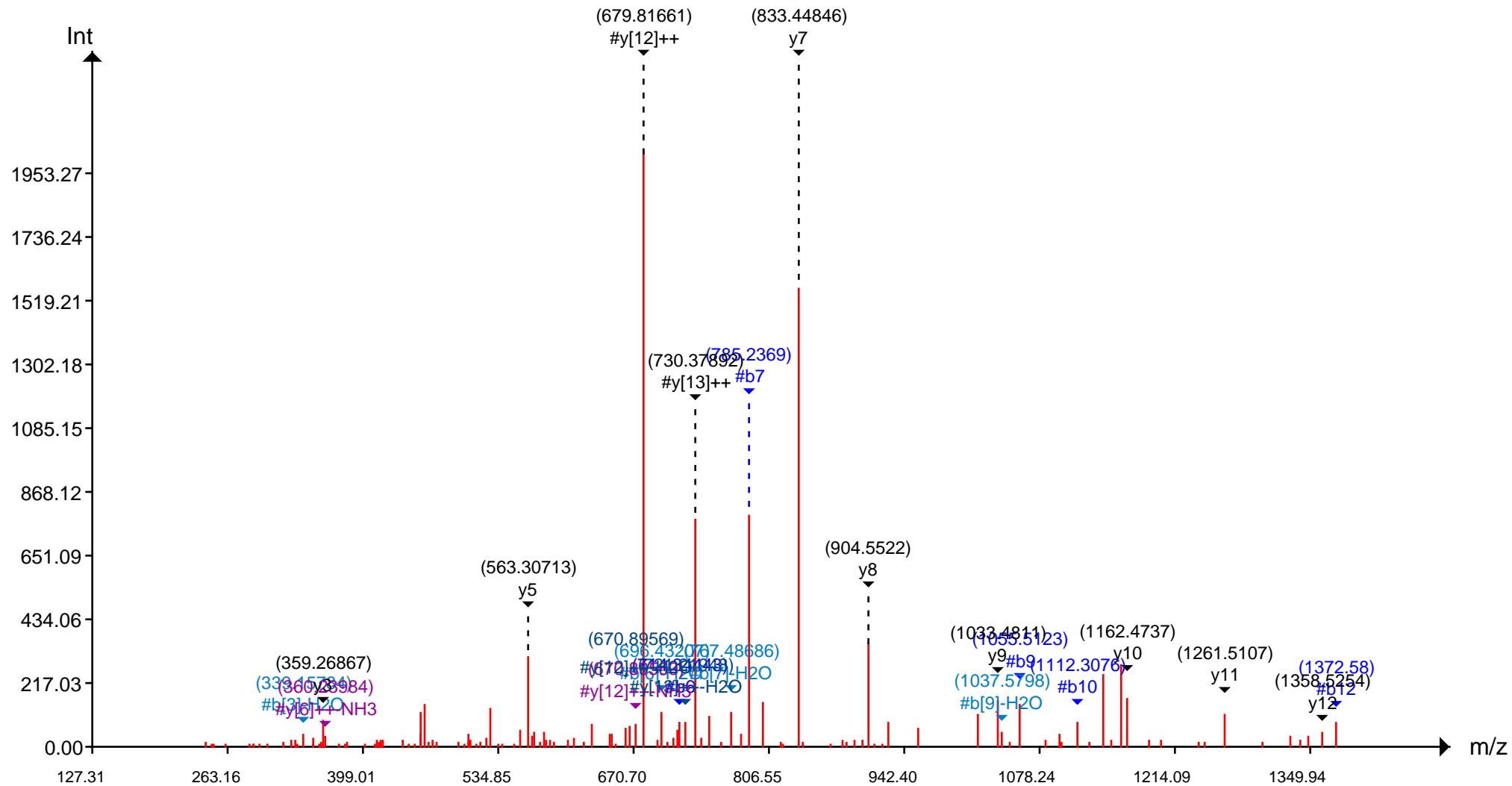
Charge 2+

Mass error 1.8E-4



Info

Modified Sequence	AcD3-ISK<AcD3K*>QEYDESGPSIVHR<C13>-COOH	Ion score	74.69
Protein	P60710, P63260	Confidence level	99.00%
Precursor m/z	971.00085	Identity Threshold	48.20
Charge	2+	Mass error	0.001796



Info

Modified Sequence AcD3-LTPVEEAPK<AcD3K*>GM<Mox*>LAR-COOH

Ion score 54.87

Protein Q61599

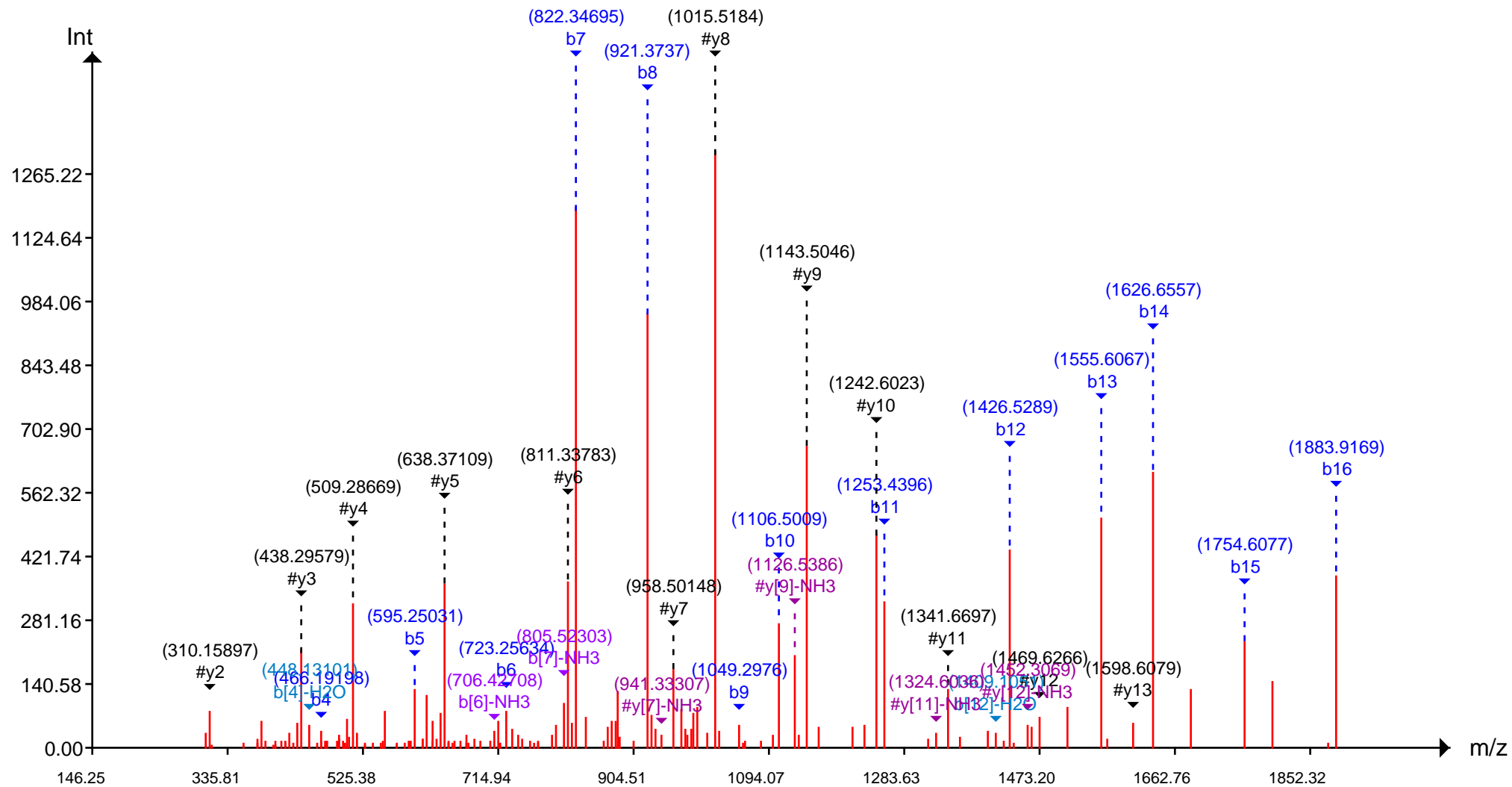
Confidence level 99.00%

Precursor m/z 809.44475

Identity Threshold 47.68

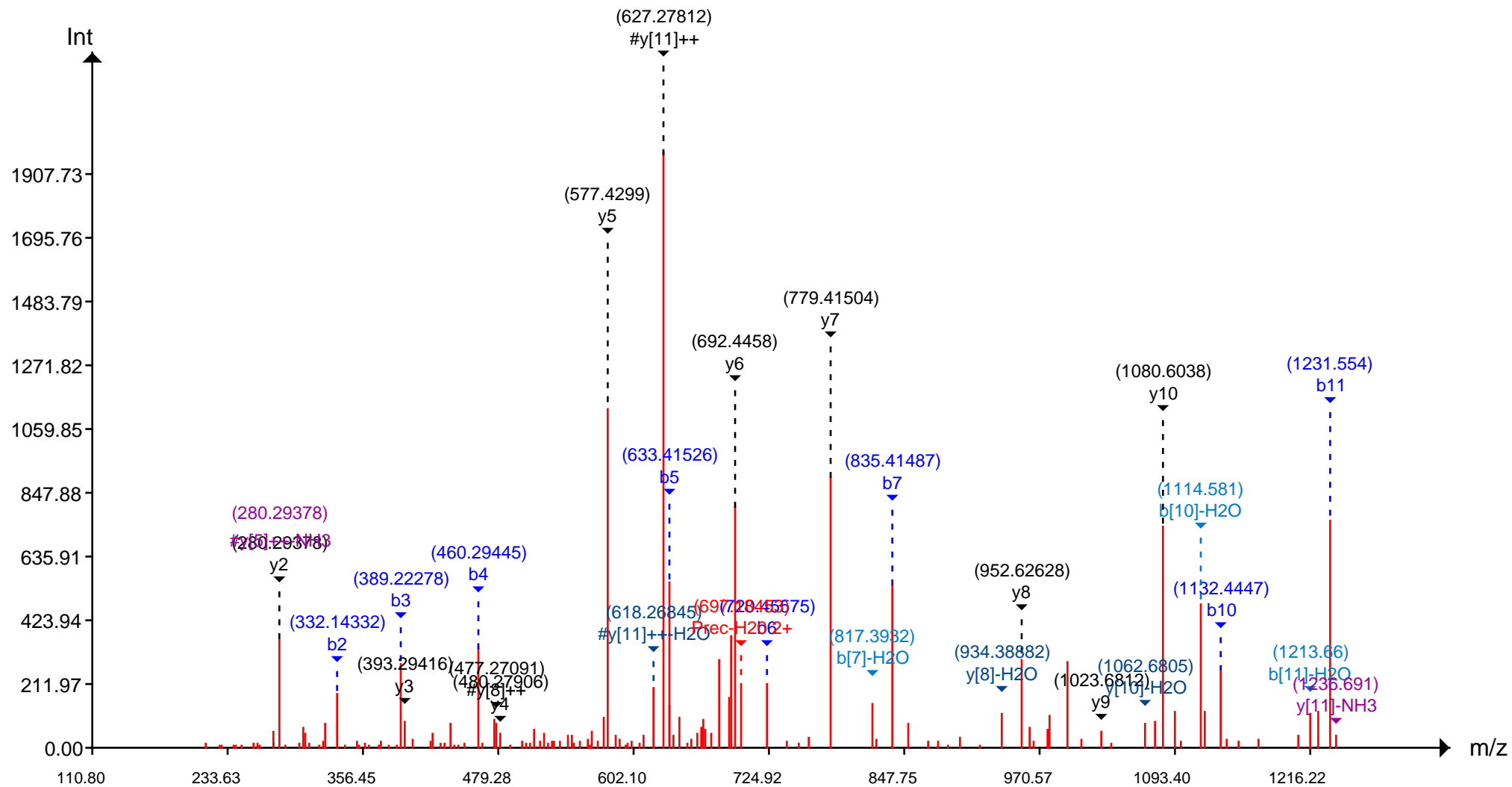
Charge 2+

Mass error 0.003519



Info

Modified Sequence	AcD3-YVSAEQVVQGM<Mox*>K<AcD3K*>EAQER<C13>-COOH	Ion score	96.6
Protein	Q9CQU0	Confidence level	99.00%
Precursor m/z	1032.5191	Identity Threshold	47.81
Charge	2+	Mass error	0.002896



Info

Modified Sequence AcD3-IK<AcD3K*>GAK<AcD3K*>SDPSIVR<C13>-COOH

Ion score 56.23

Protein P62748

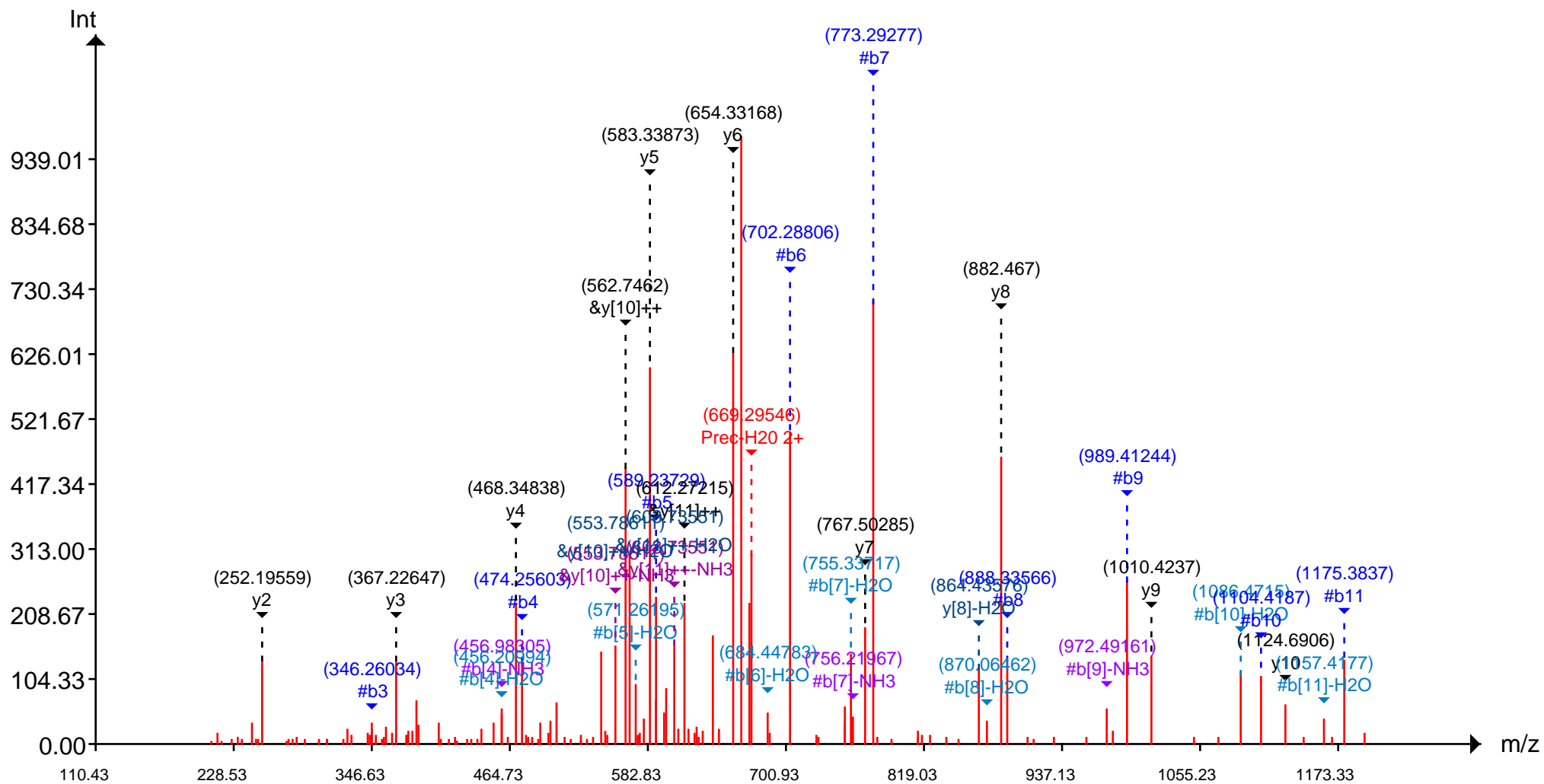
Confidence level 99.00%

Precursor m/z 706.43205

Identity Threshold 42.59

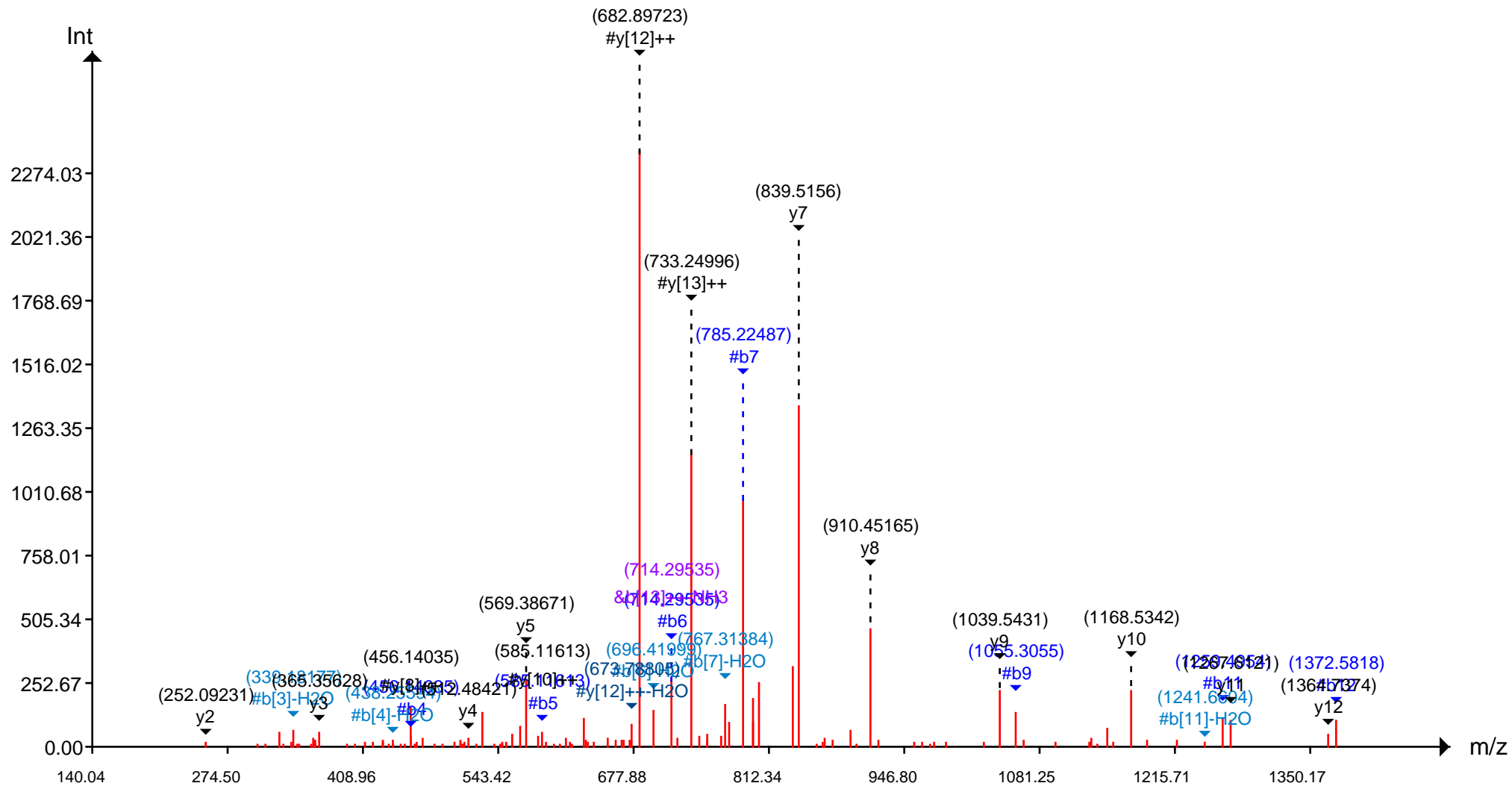
Charge 2+

Mass error 7.47E-4



Info

Modified Sequence	AcD3-SVNQLDADTDAR<C13>-COOH	Ion score	85.7
Protein	P10107	Confidence level	99.00%
Precursor m/z	678.3325	Identity Threshold	47.09
Charge	2+	Mass error	5.0E-4



Info

Modified Sequence AcD3-LTPVEEAPK<AcD3K*>GM<Mox*>LAR<C13>-COOH

Ion score 61.85

Protein Q61599

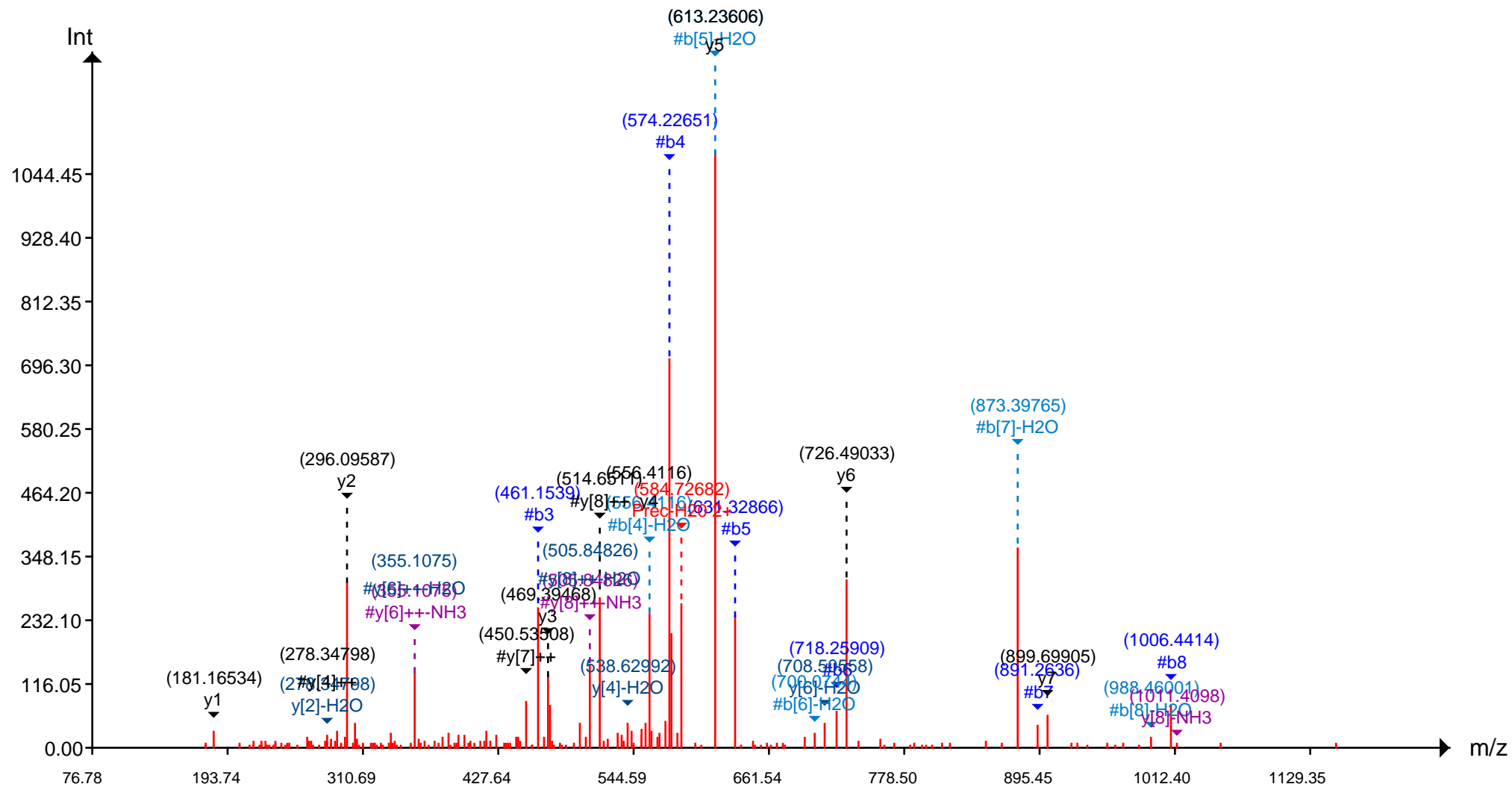
Confidence level 99.00%

Precursor m/z 812.45434

Identity Threshold 47.16

Charge 2+

Mass error 0.002557



Info

Modified Sequence AcD3-IEK<AcD3K*>LGSK<AcD3K*>DR<C13>-COOH

Ion score 56.25

Protein Q921Y0

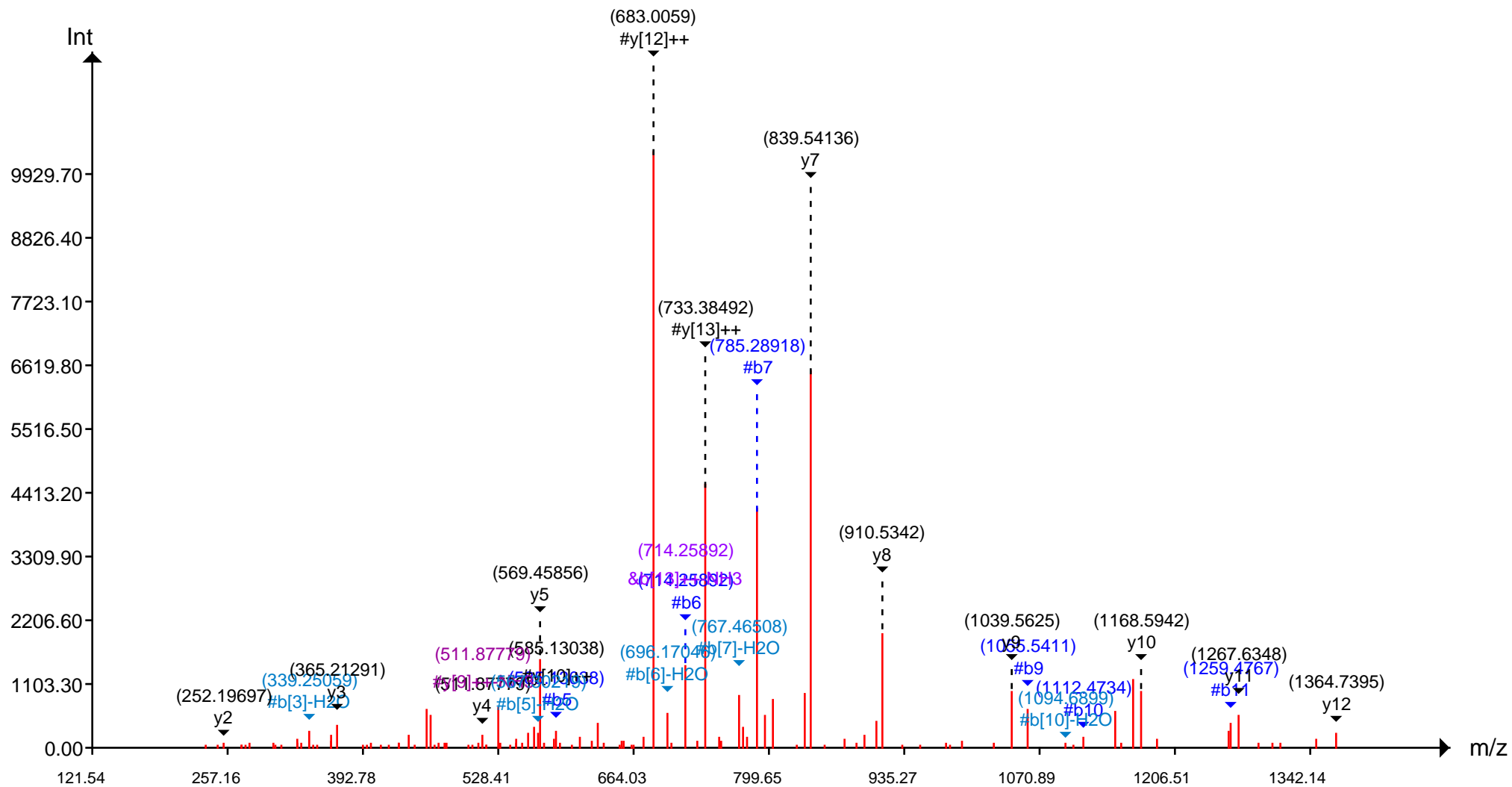
Confidence level 99.00%

Precursor m/z 593.85779

Identity Threshold 45.01

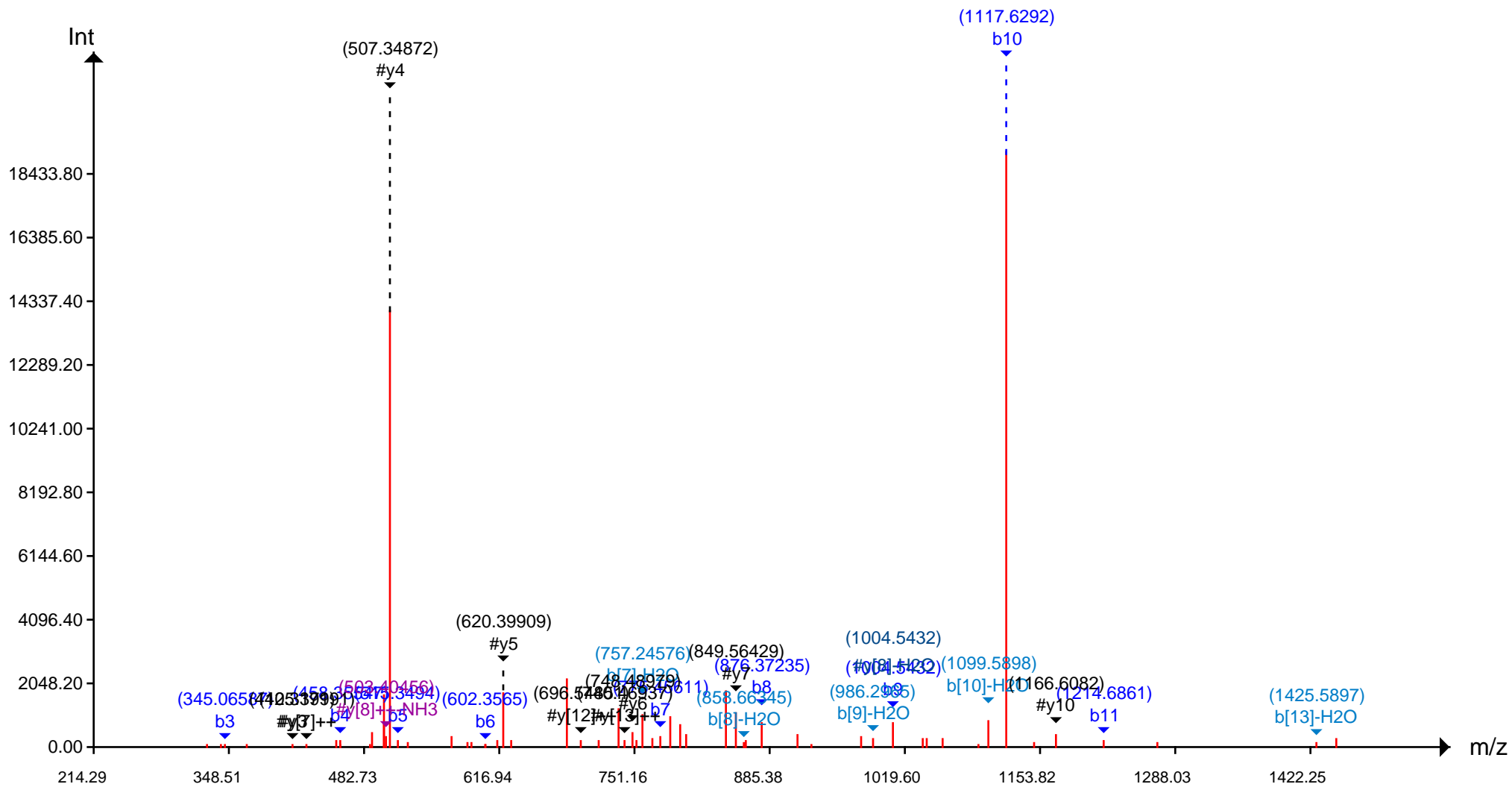
Charge 2+

Mass error -3.7E-5



Info

Modified Sequence	AcD3-LTPVEEAPK<AcD3K*>GM<Mox*>LAR<C13>-COOH	Ion score	59.61
Protein	Q61599	Confidence level	99.00%
Precursor m/z	812.45485	Identity Threshold	47.01
Charge	2+	Mass error	0.003577



Info

Modified Sequence AcD3-VSLIGSK<AcD3K*>TQIPTQR<C13>-COOH

Ion score 45.81

Protein Q9R1P4

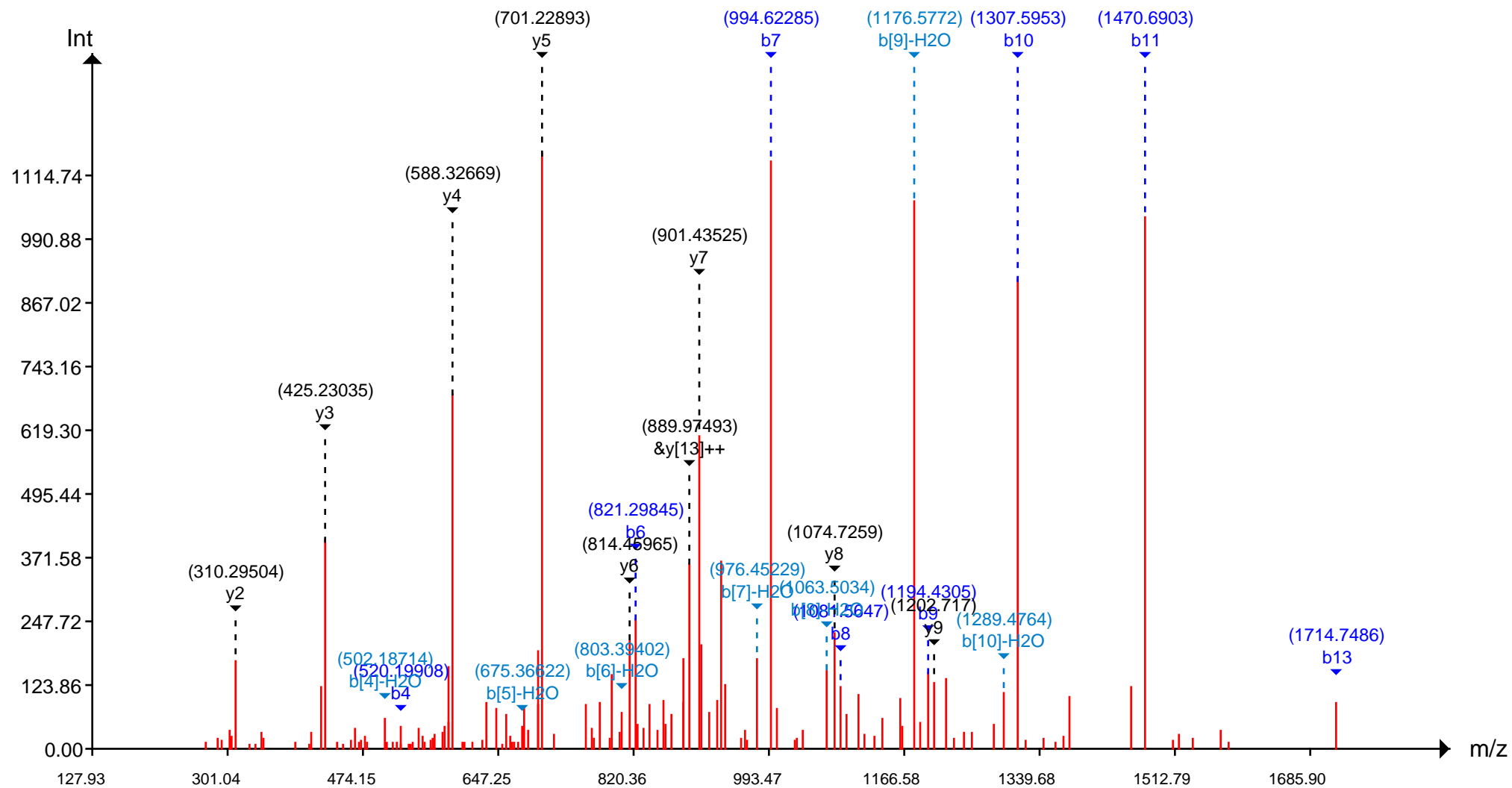
Confidence level 99.00%

Precursor m/z 812.48644

Identity Threshold 43.30

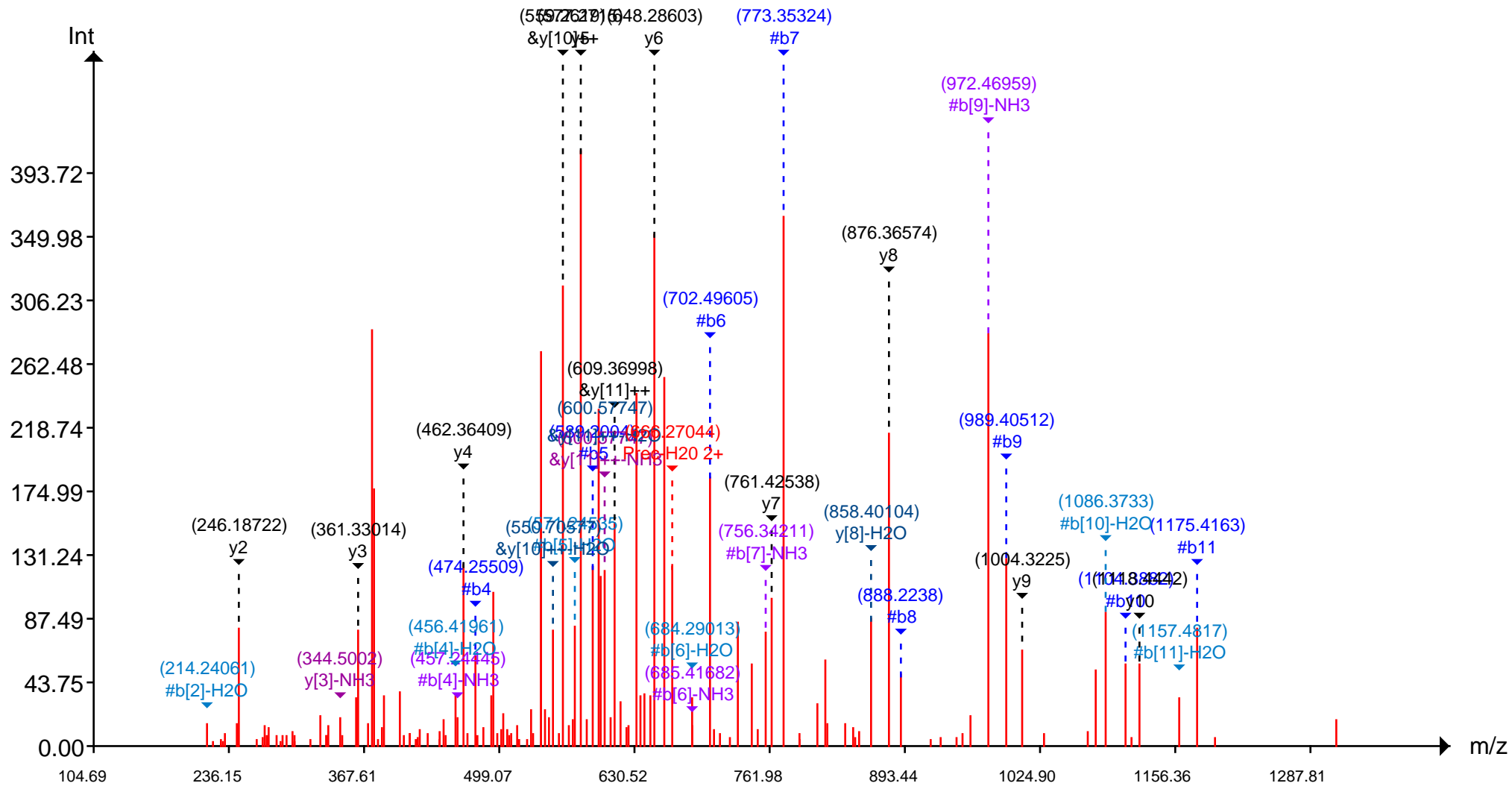
Charge 2+

Mass error 0.001358



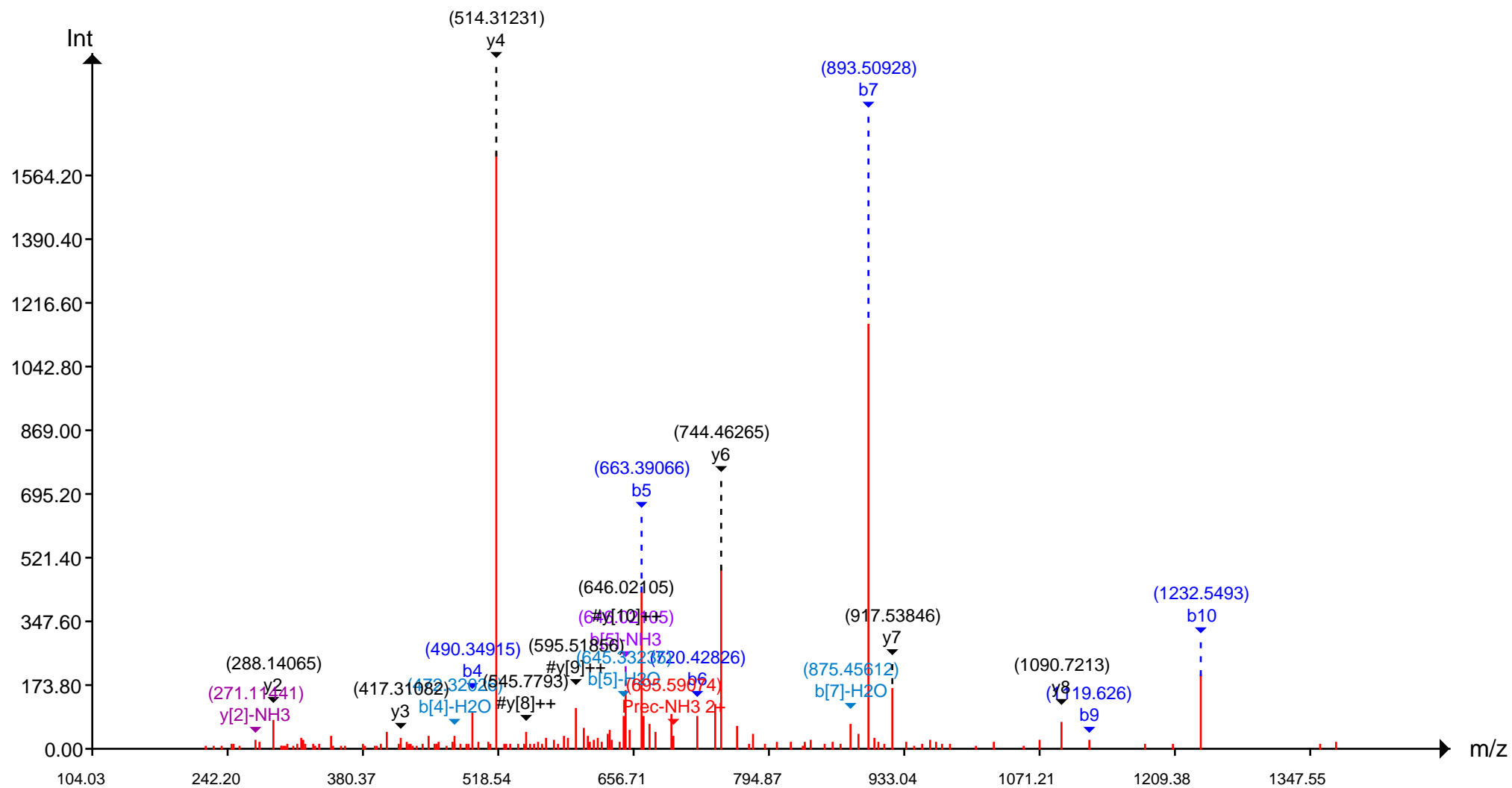
Info

Modified Sequence	AcD3-ATEK<AcD3K*>K<AcD3K*>QK<AcD3K*>SILYDER<C13>-COOH	Ion score	57.34
Protein	P49722	Confidence level	99.00%
Precursor m/z	948.03468	Identity Threshold	46.31
Charge	2+	Mass error	0.00157



Info

Modified Sequence	AcD3-SVNQLDADTDAR-COOH	Ion score	70.85
Protein	P10107	Confidence level	99.00%
Precursor m/z	675.32262	Identity Threshold	46.16
Charge	2+	Mass error	8.81E-4



Info

Modified Sequence AcD3-ATVK<AcD3K*>K<AcD3K*>GK<AcD3K*>PELR-COOH

Ion score 40.38

Protein P62830

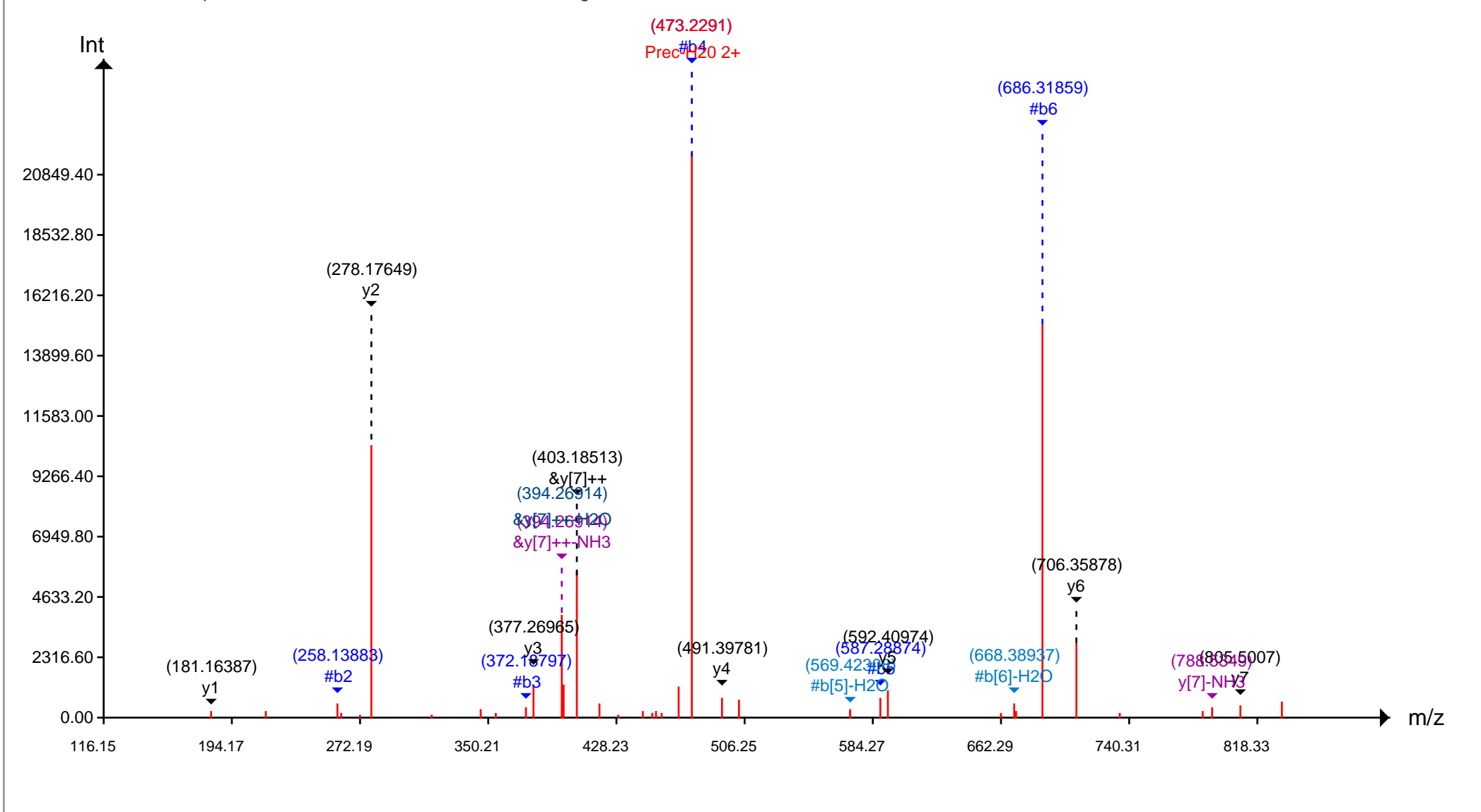
Confidence level 99.00%

Precursor m/z 703.94144

Identity Threshold 39.92

Charge 2+

Mass error 1.03E-4



Info

Modified Sequence AcD3-IVNTNVPR<C13>-COOH

Ion score 67.34

Protein P34884

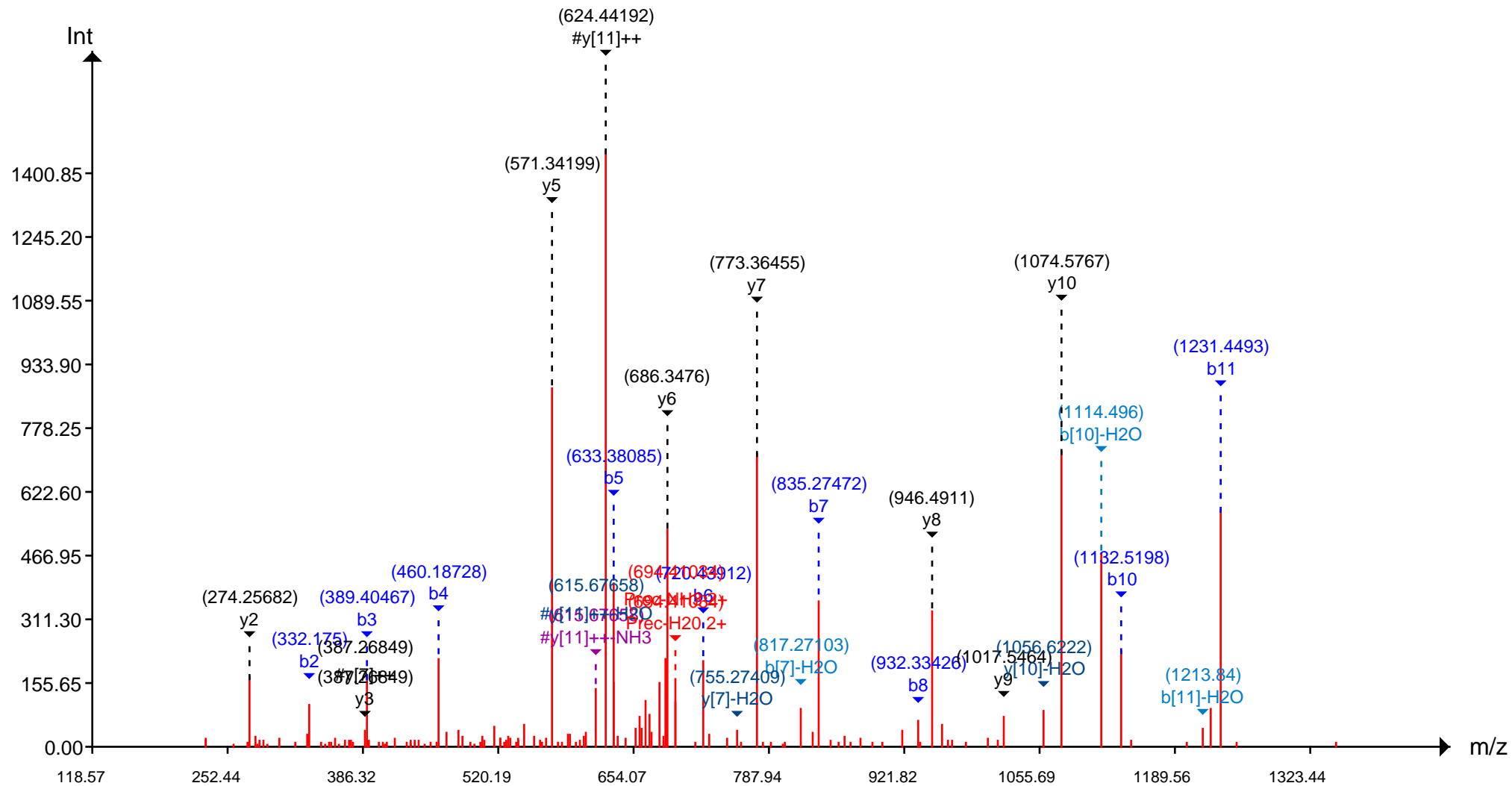
Confidence level 99.00%

Precursor m/z 482.2911

Identity Threshold 45.65

Charge 2+

Mass error -7.42E-4



Info

Modified Sequence AcD3-IK<AcD3K*>GAK<AcD3K*>SDPSIVR-COOH

Ion score 63.47

Protein P62748

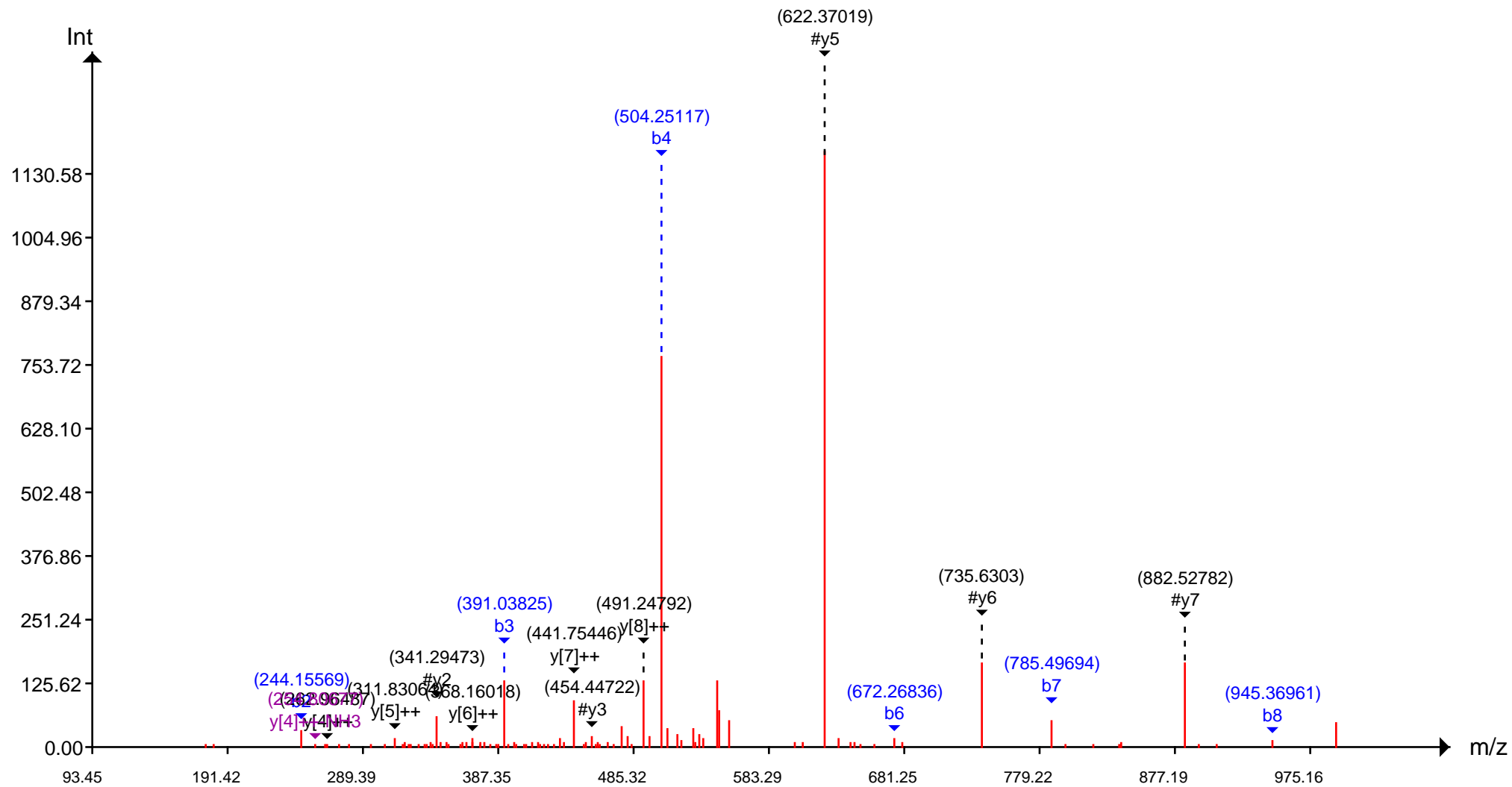
Confidence level 99.00%

Precursor m/z 703.42212

Identity Threshold 43.98

Charge 2+

Mass error 0.001029



Info

Modified Sequence AcD3-VVFLPALC<Cmm*>R<C13>-COOH

Ion score 50.57

Protein P12970

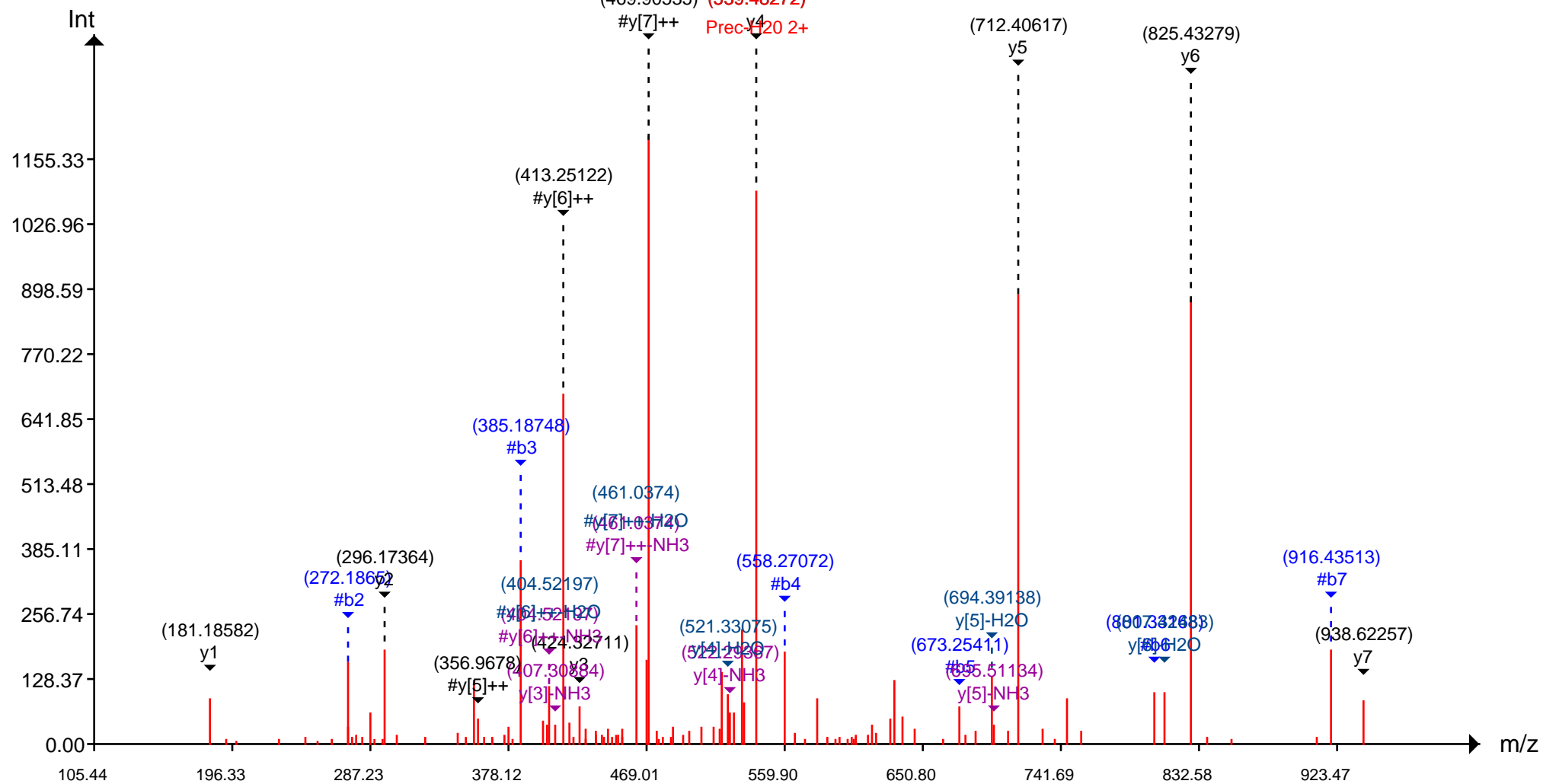
Confidence level 99.00%

Precursor m/z 563.33507

Identity Threshold 45.32

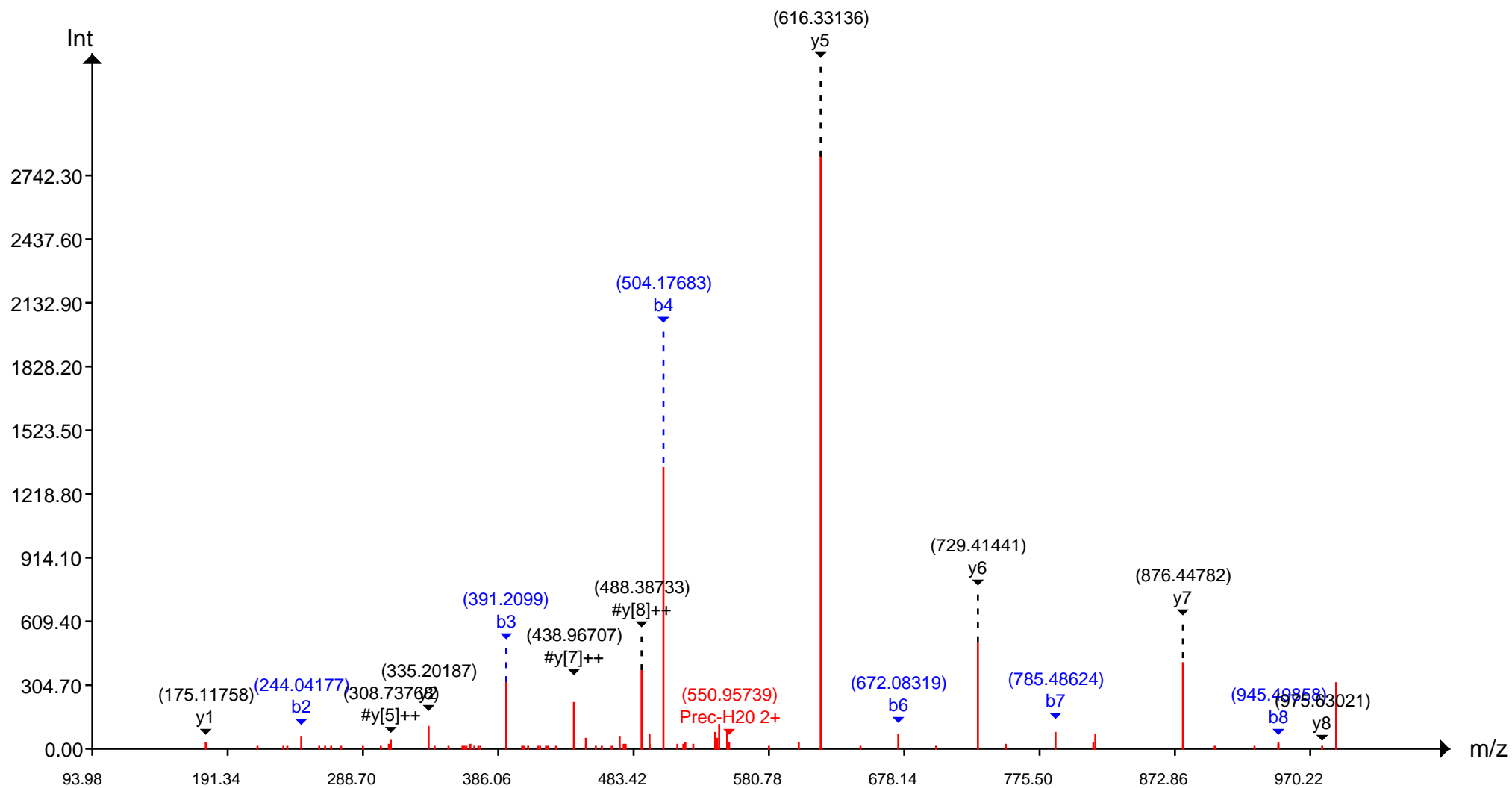
Charge 2+

Mass error 4.82E-4



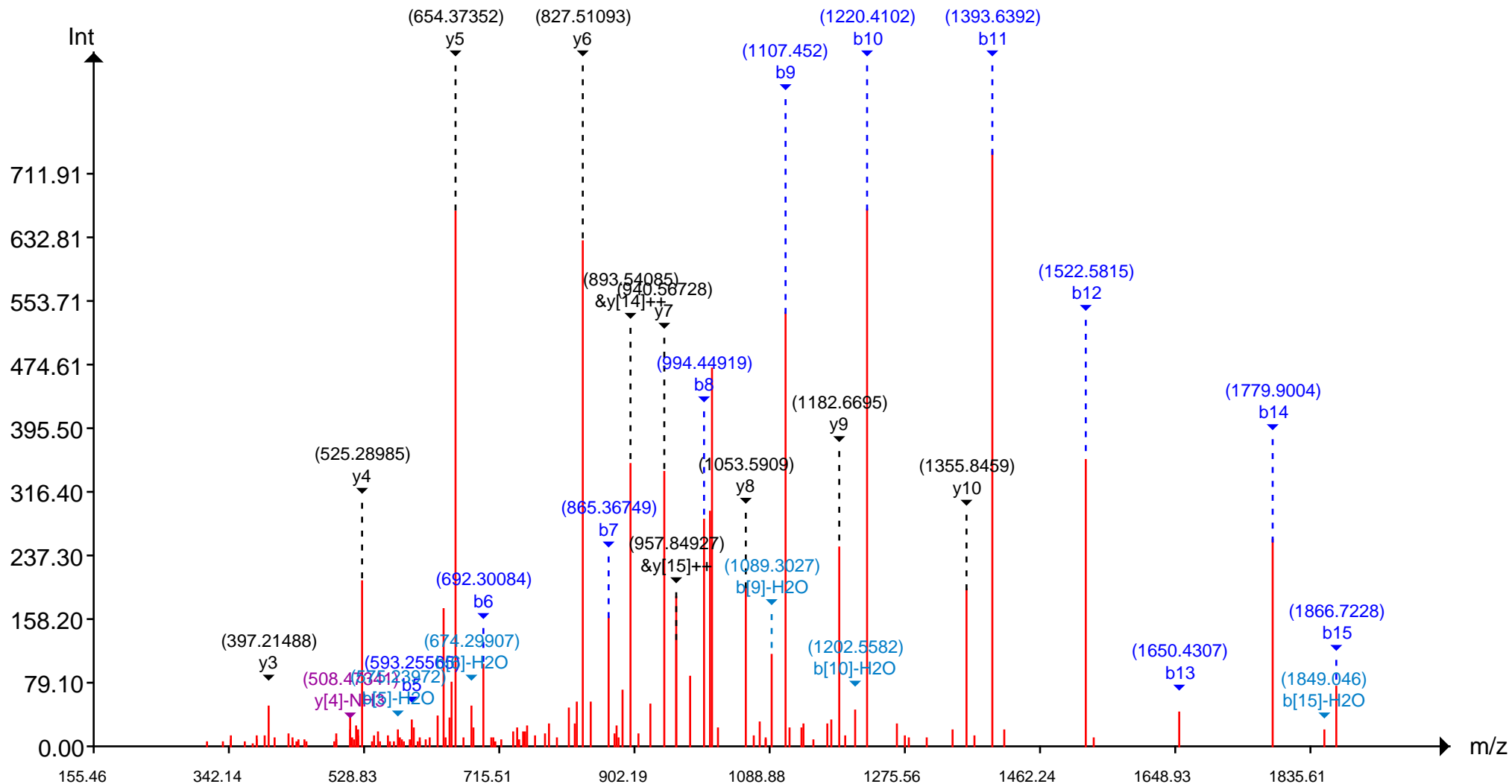
Info

Modified Sequence	AcD3-LIIK<AcD3K*>DQDR<C13>-COOH	Ion score	56.05
Protein	P80316	Confidence level	99.00%
Precursor m/z	548.83273	Identity Threshold	44.82
Charge	2+	Mass error	7.0E-4



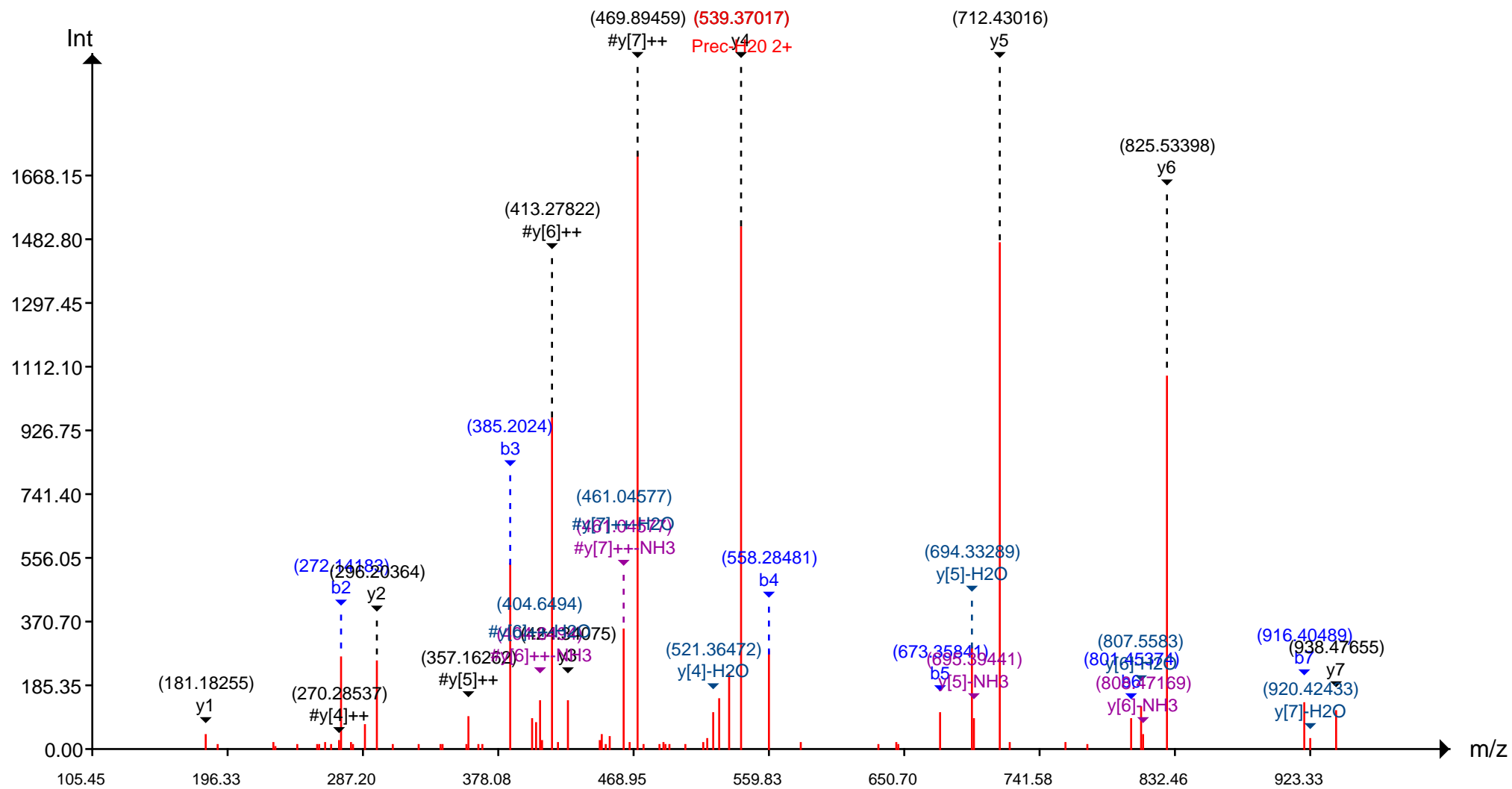
Info

Modified Sequence	AcD3-VVFLPALC<Cmm*>R-COOH	Ion score	58.9
Protein	P12970	Confidence level	99.00%
Precursor m/z	560.32408	Identity Threshold	46.55
Charge	2+	Mass error	-0.001356



Info

Modified Sequence	AcD3-SESPM<Mox*>VK<AcD3K*>EILK<AcD3K*>EQESR<C13>-COOH	Ion score	76.31
Protein	Q99LX0	Confidence level	99.00%
Precursor m/z	1024.0383	Identity Threshold	48.04
Charge	2+	Mass error	0.002417



Info

Modified Sequence AcD3-LIIK<AcD3K*>DQDR<C13>-COOH

Ion score 60.88

Protein P80316

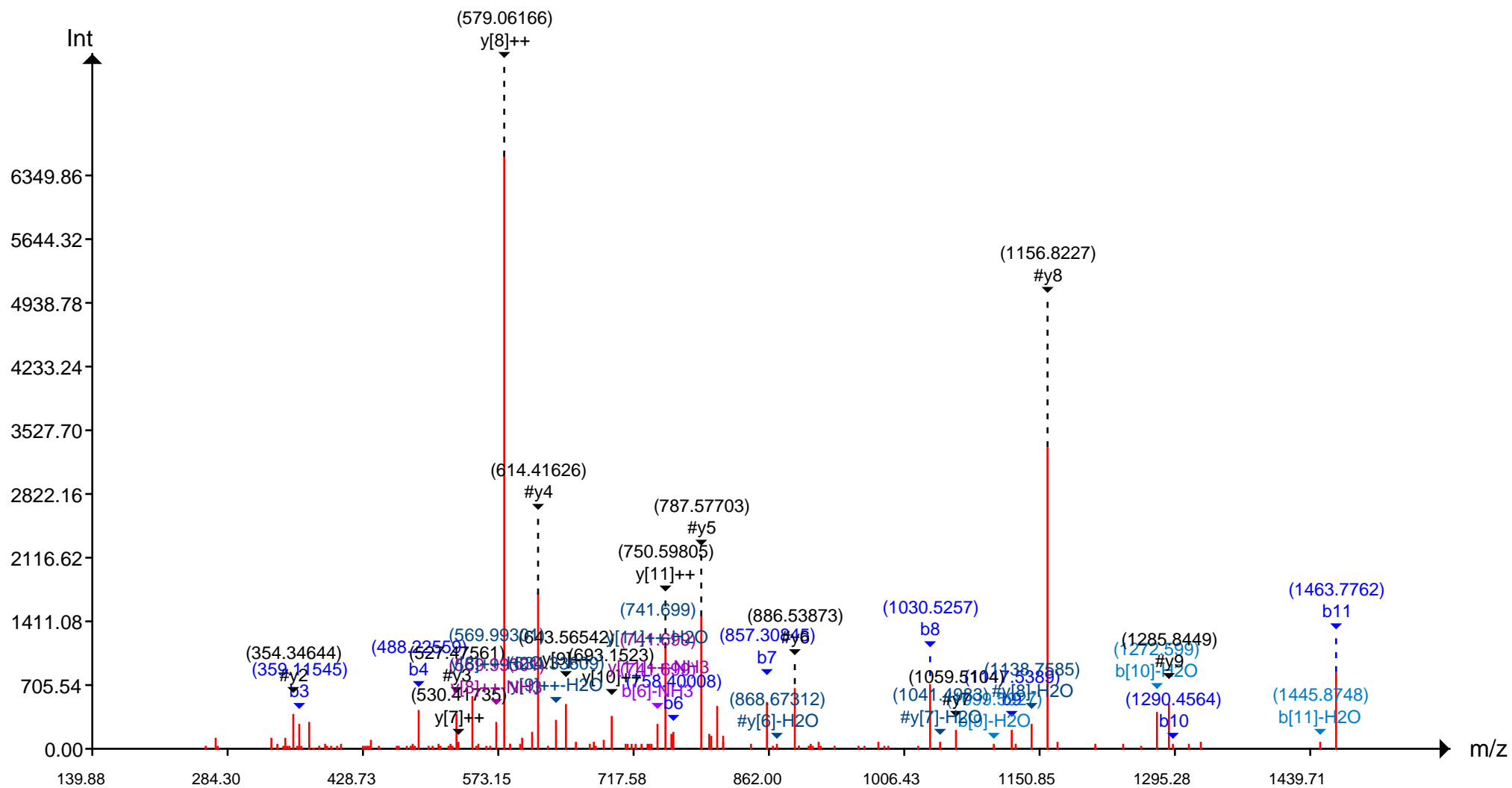
Confidence level 99.00%

Precursor m/z 548.83231

Identity Threshold 44.61

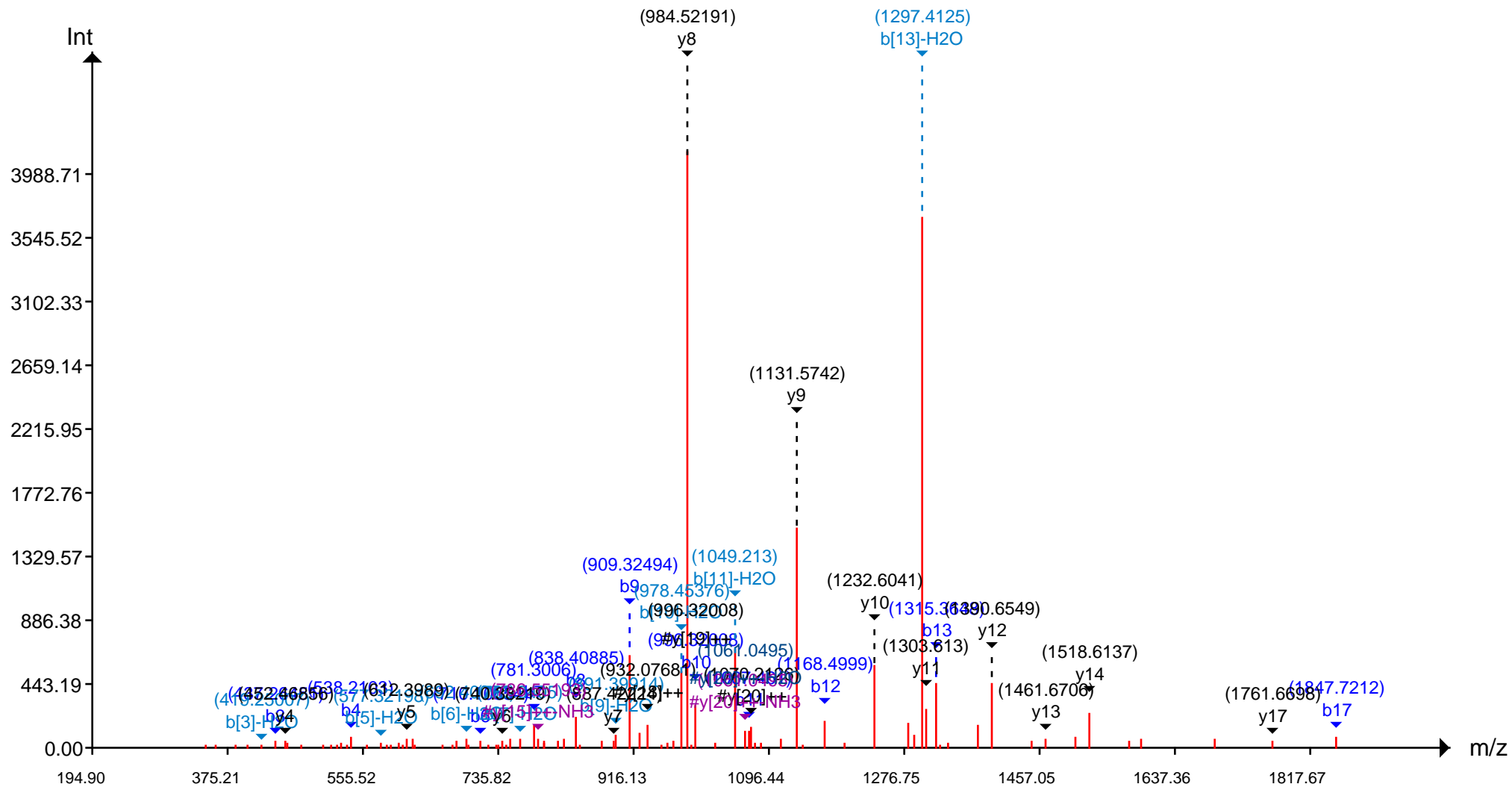
Charge 2+

Mass error -1.4E-4



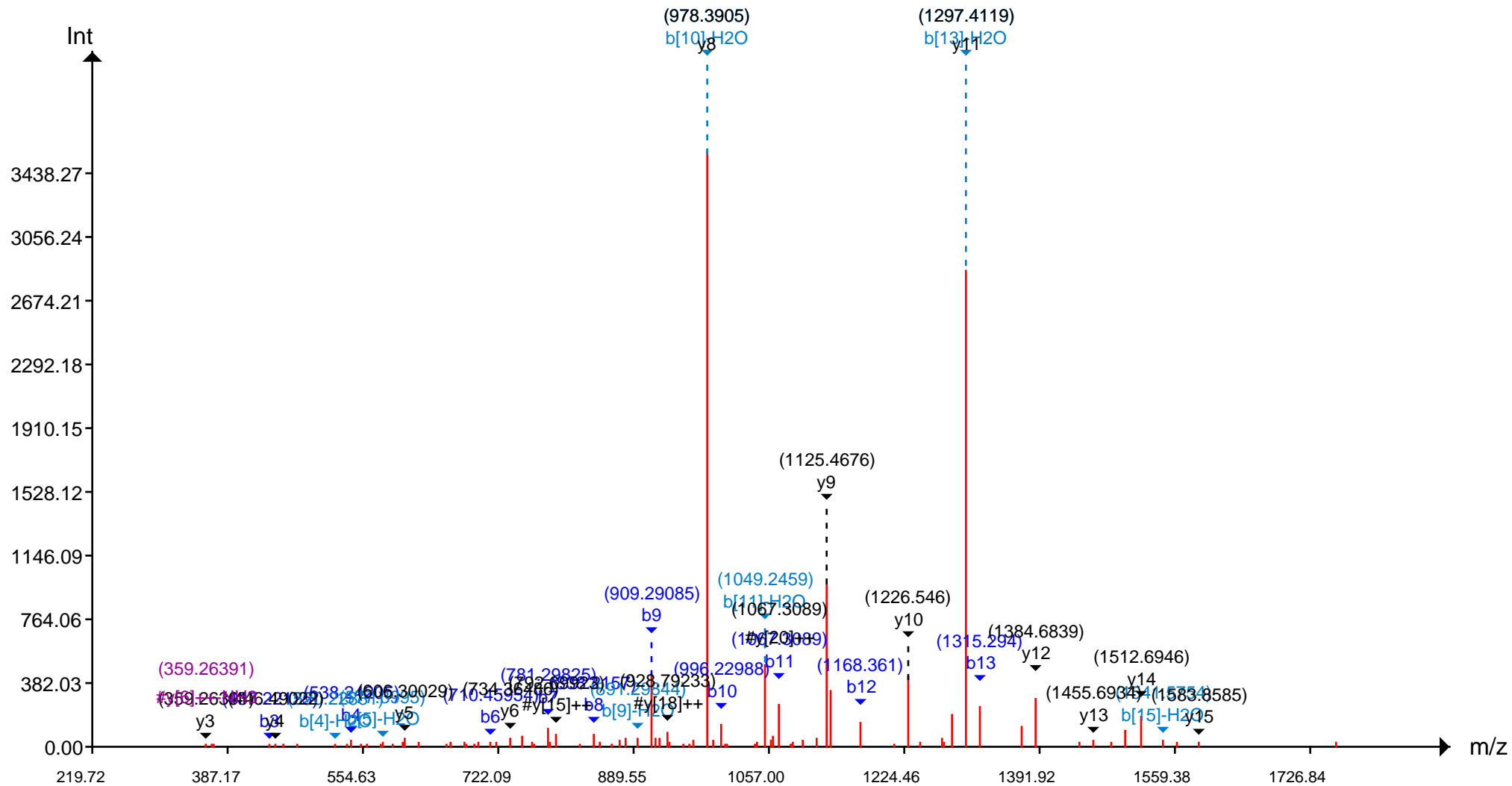
Info

Modified Sequence	AcD3-VDVEPK<AcD3K*>VK<AcD3K*>SK<AcD3K*>K<AcD3K*>R<C13>-COOH	Ion score	55.81
Protein	P62315	Confidence level	99.00%
Precursor m/z	822.51699	Identity Threshold	38.37
Charge	2+	Mass error	0.001209



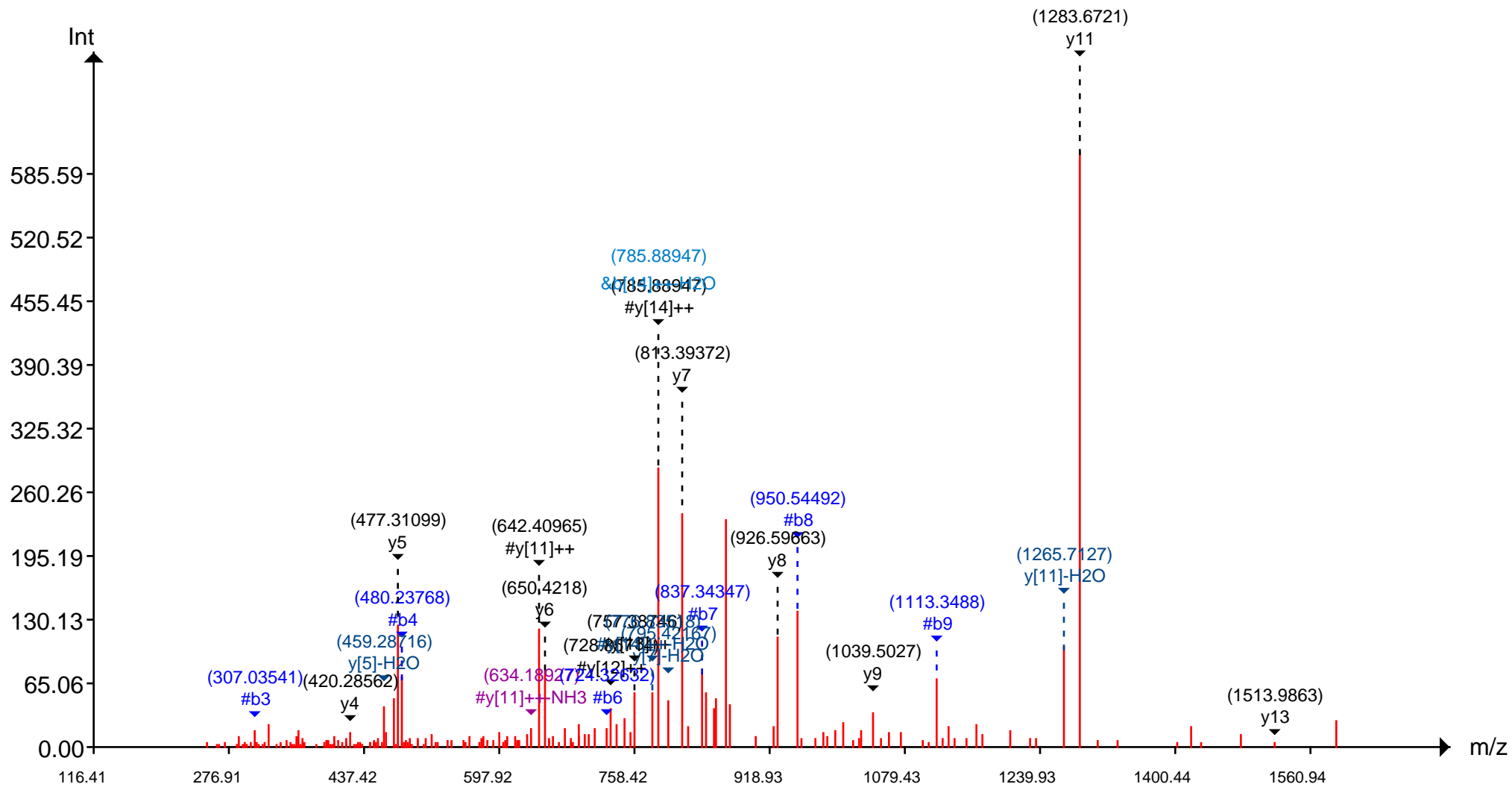
Info

Modified Sequence	AcD3-DFETGDAGASATFPM<Mox*>QC<Cmm*>SALR<C13>-COOH	Ion score	90.47
Protein	P63242	Confidence level	99.00%
Precursor m/z	1150.012	Identity Threshold	42.62
Charge	2+	Mass error	0.002673



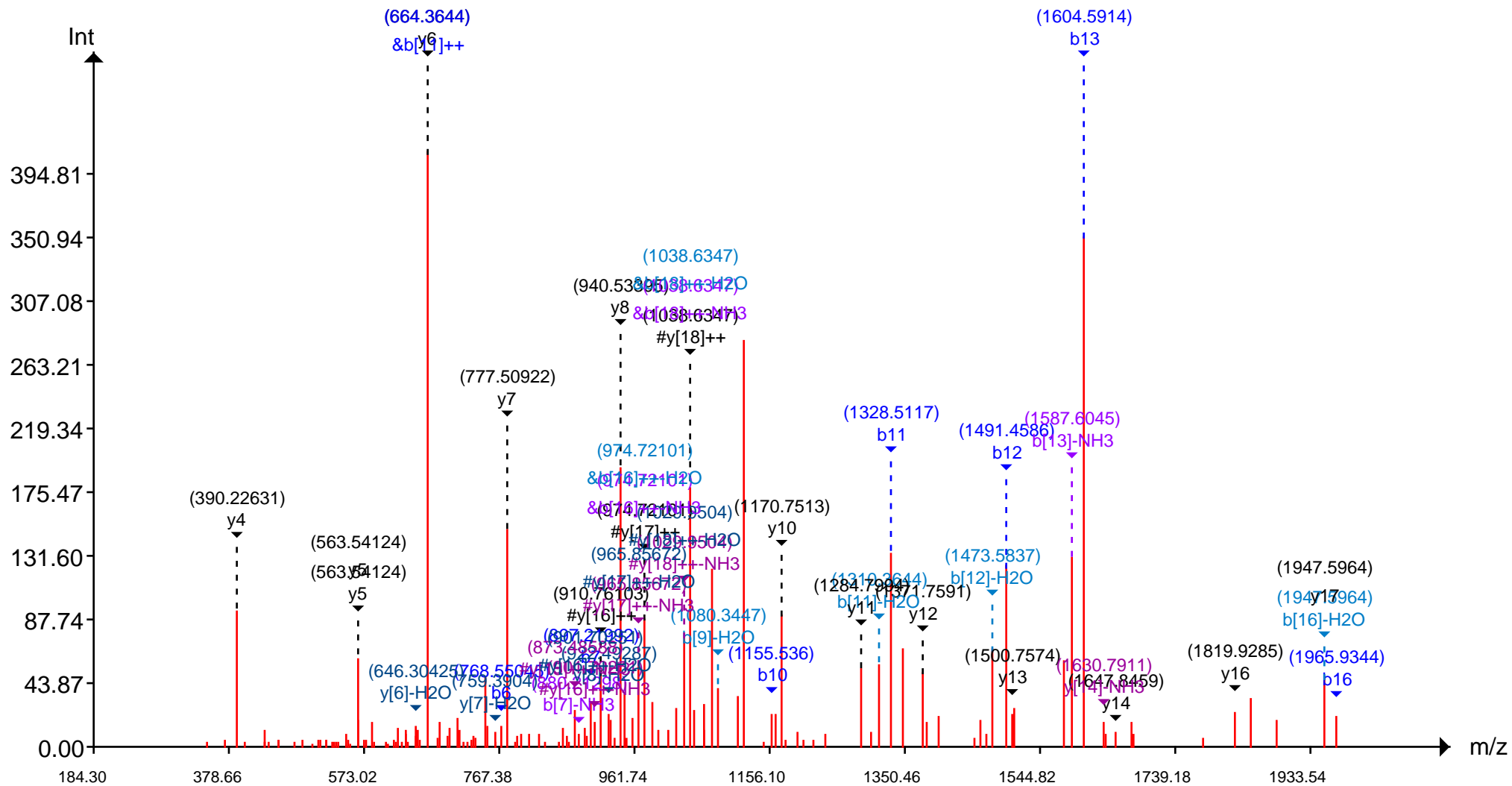
Info

Modified Sequence	AcD3-DFETGDAGASATFPM<Mox*>QC<Cmm*>SALR-COOH	Ion score	112.25
Protein	P63242	Confidence level	99.00%
Precursor m/z	1147.0024	Identity Threshold	41.59
Charge	2+	Mass error	0.003615



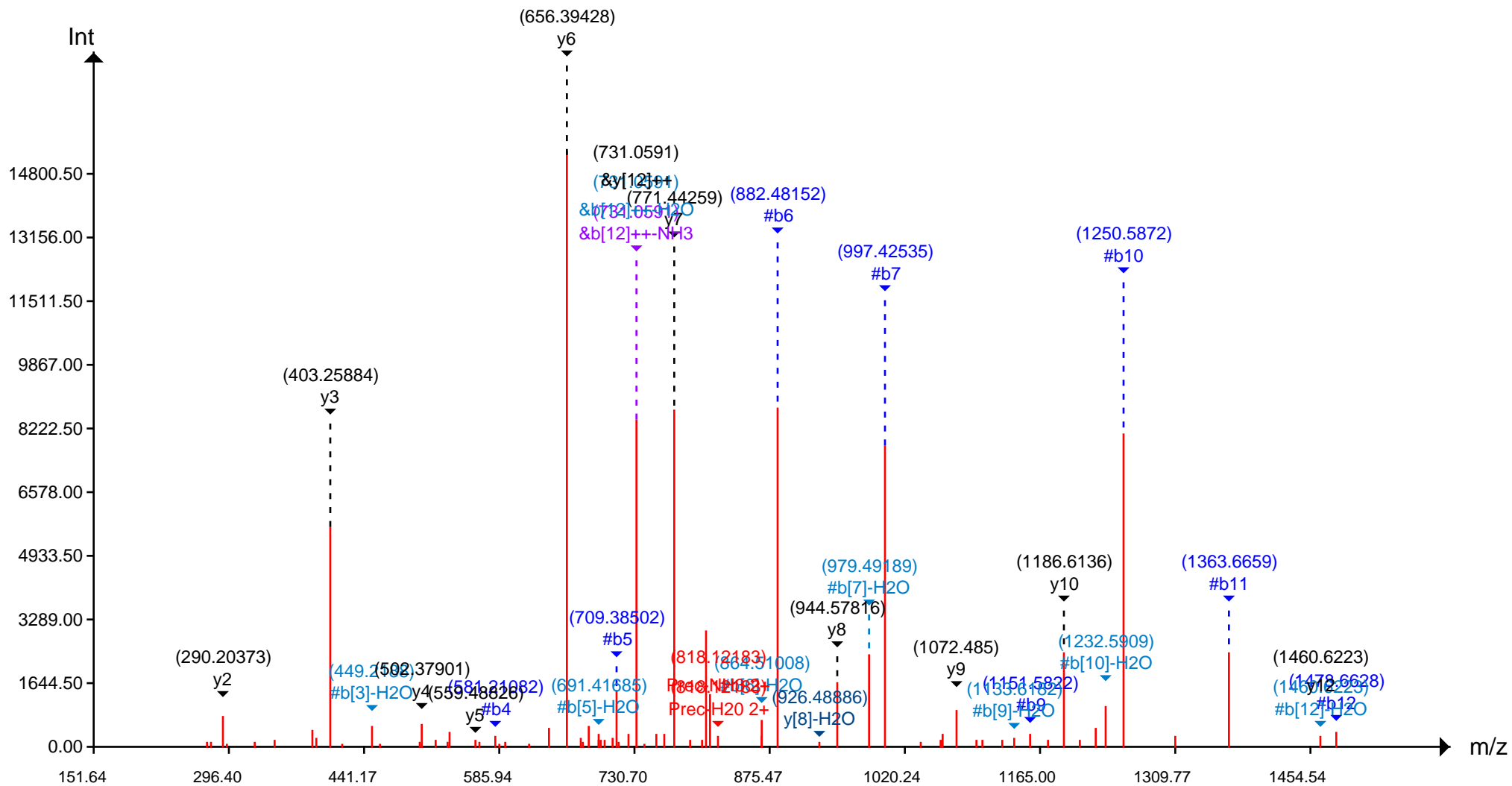
Info

Modified Sequence	AcD3-FGGK<AcD3K*>PM<Mox*>IITYK<AcD3K*>GGTSR-COOH	Ion score	54.41
Protein	P13020	Confidence level	99.00%
Precursor m/z	881.97972	Identity Threshold	47.74
Charge	2+	Mass error	0.001498



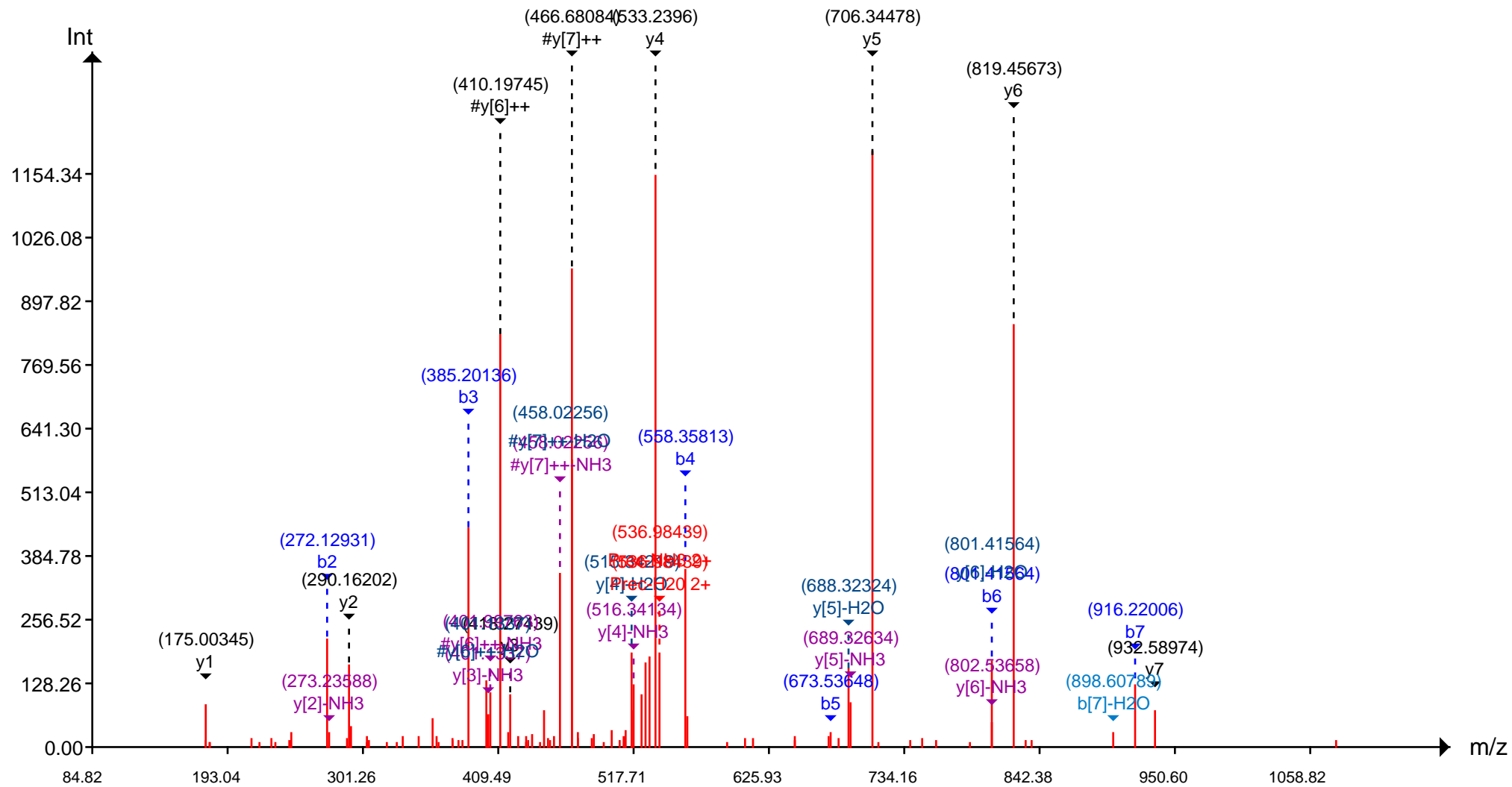
Info

Modified Sequence	AcD3-FQQGDM<Mox*>ESNGK<AcD3K*>YITK<AcD3K*>SGAR-COOH	Ion score	72.74
Protein	P06745	Confidence level	99.00%
Precursor m/z	1134.5509	Identity Threshold	46.97
Charge	2+	Mass error	0.003401



Info

Modified Sequence	AcD3-FTK<AcD3K*>NQK<AcD3K*>DPGVLR-COOH	Ion score	91.92
Protein	P50543	Confidence level	99.00%
Precursor m/z	826.95226	Identity Threshold	47.53
Charge	2+	Mass error	0.001983



Info

Modified Sequence AcD3-LIIK<AcD3K*>DQDR-COOH

Ion score 55.27

Protein P80316

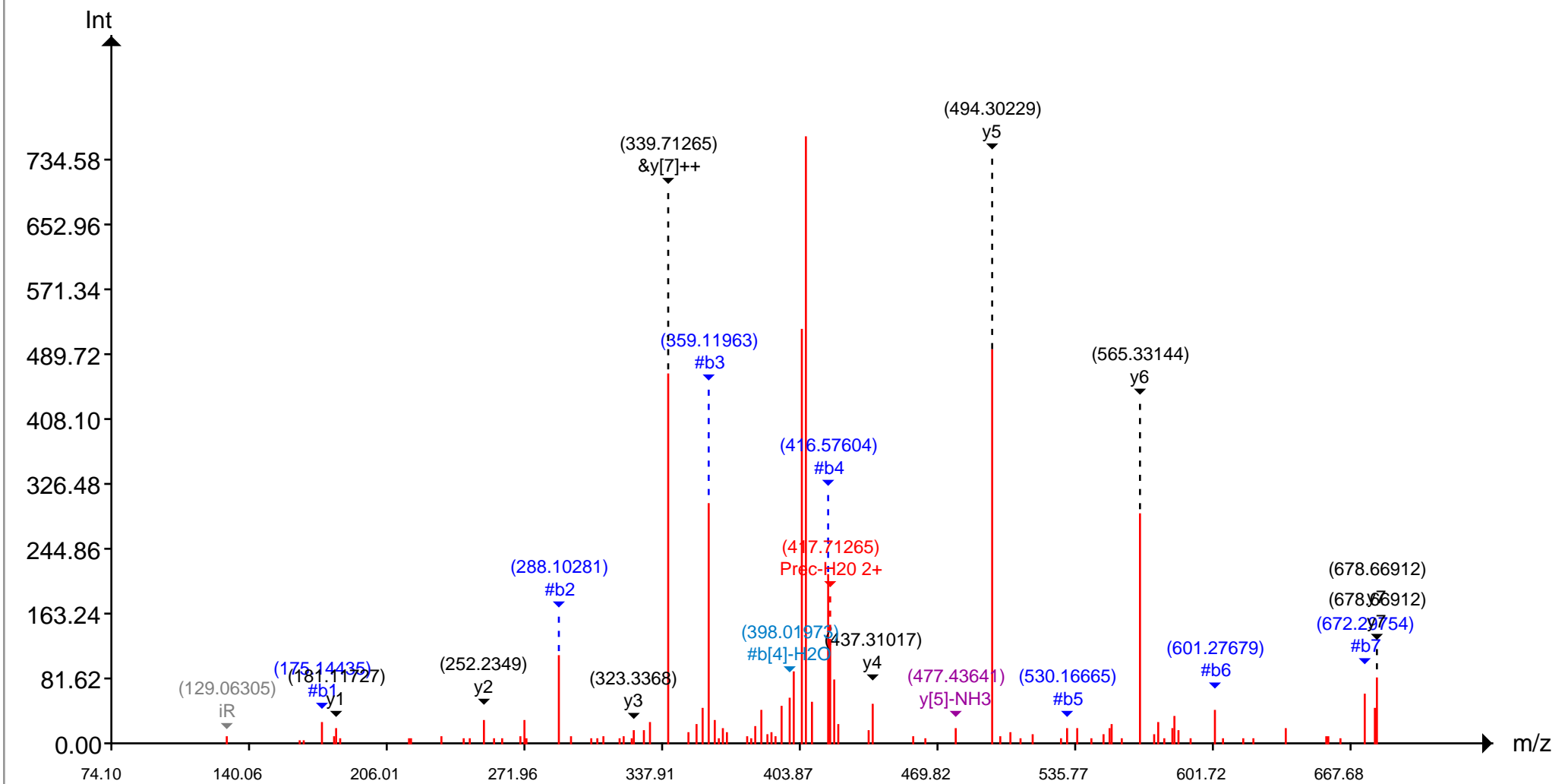
Confidence level 99.00%

Precursor m/z 545.82225

Identity Threshold 45.60

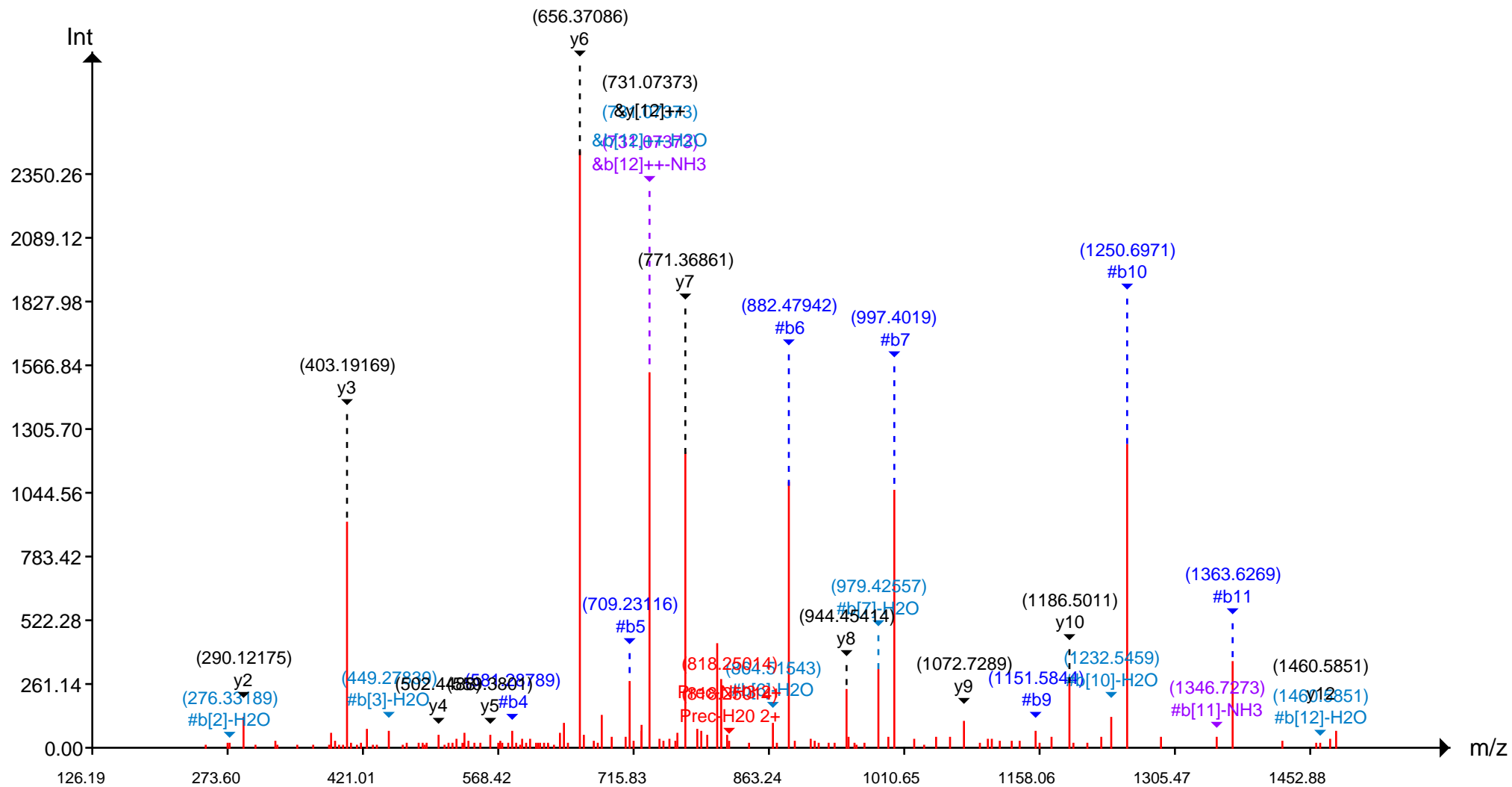
Charge 2+

Mass error -1.18E-4



Info

Modif...	AcD3-ELAGNAAR<C13>-COOH	Ion score
P...	Q8CGP5, Q8CGP6, Q8CGP7, P22752, Q6GSS7, Q64522, Q64523, Q8BFU2, Q8R1M2, Q8CCK0, P27661, Q9QZQ8	Confidence level
	426.73914	Identity Threshold
	2+	Mass error



Info

Modified Sequence AcD3-FTK<AcD3K*>NQK<AcD3K*>DPGVLR-COOH

Ion score 64.12

Protein P50543

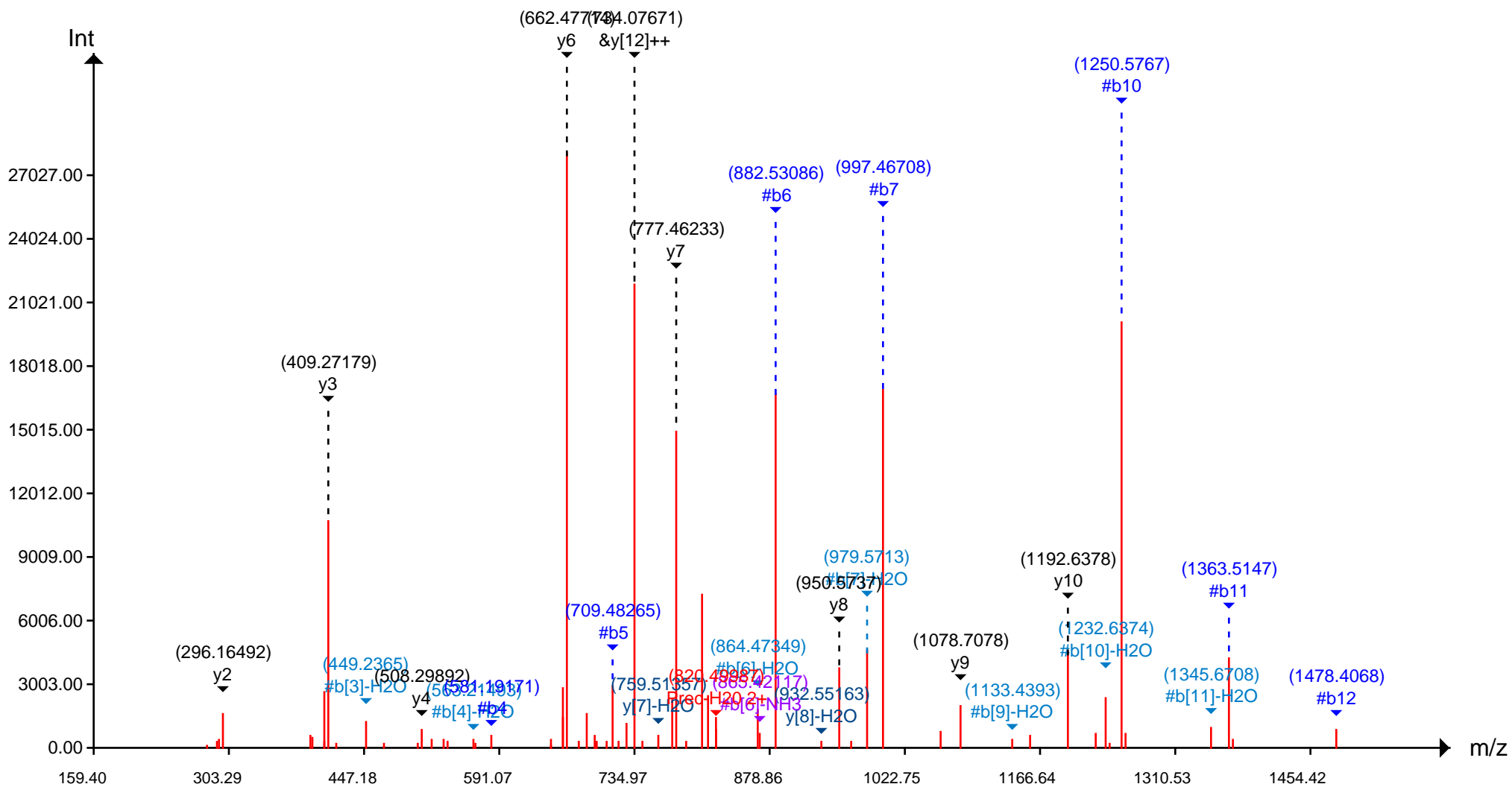
Confidence level 99.00%

Precursor m/z 826.95223

Identity Threshold 47.53

Charge 2+

Mass error 0.001923



Info

Modified Sequence AcD3-FTK<AcD3K*>NQK<AcD3K*>DPGVLDK<C13>-COOH

Ion score 58.36

Protein P50543

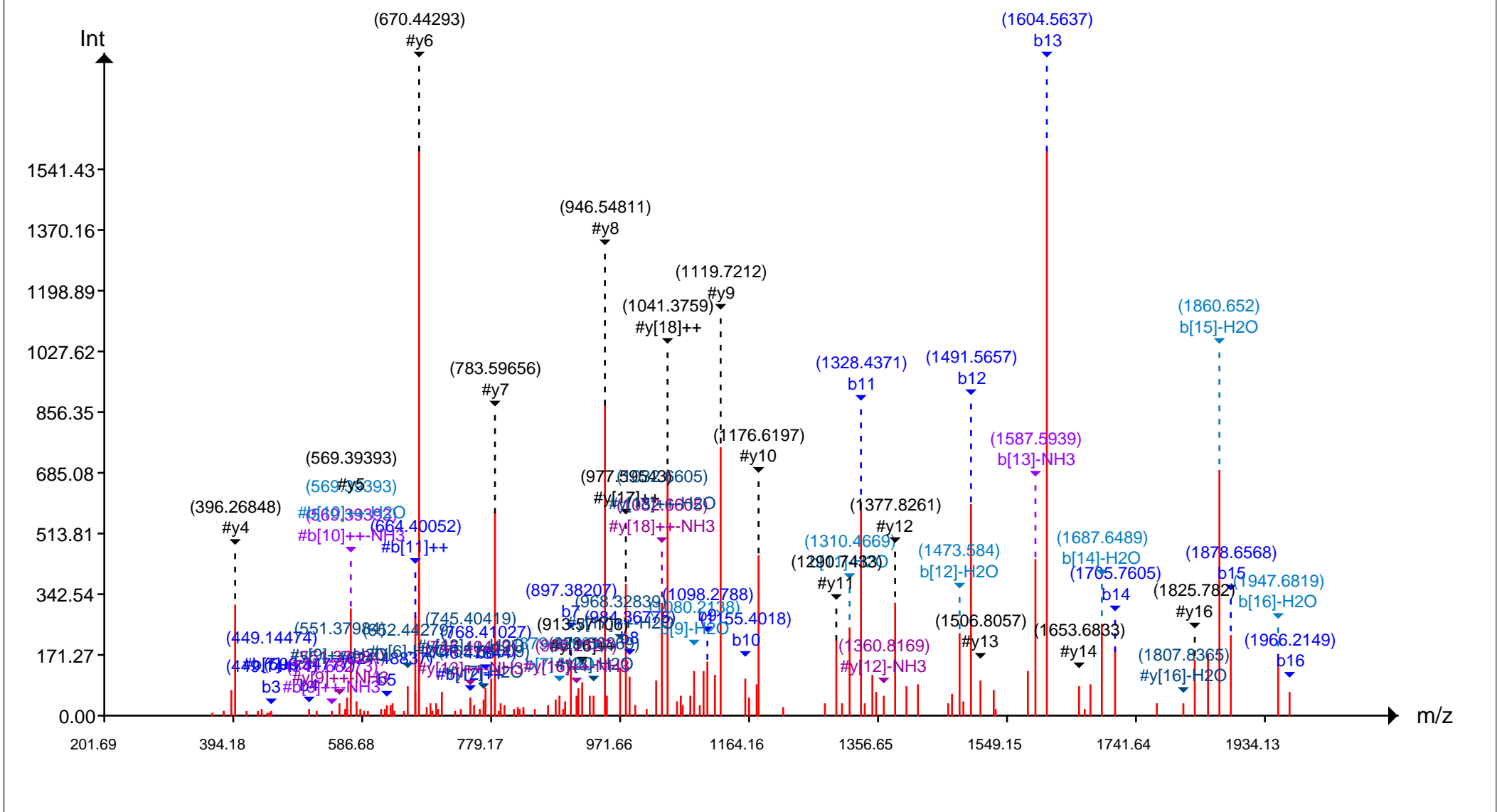
Confidence level 99.00%

Precursor m/z 829.96208

Identity Threshold 47.11

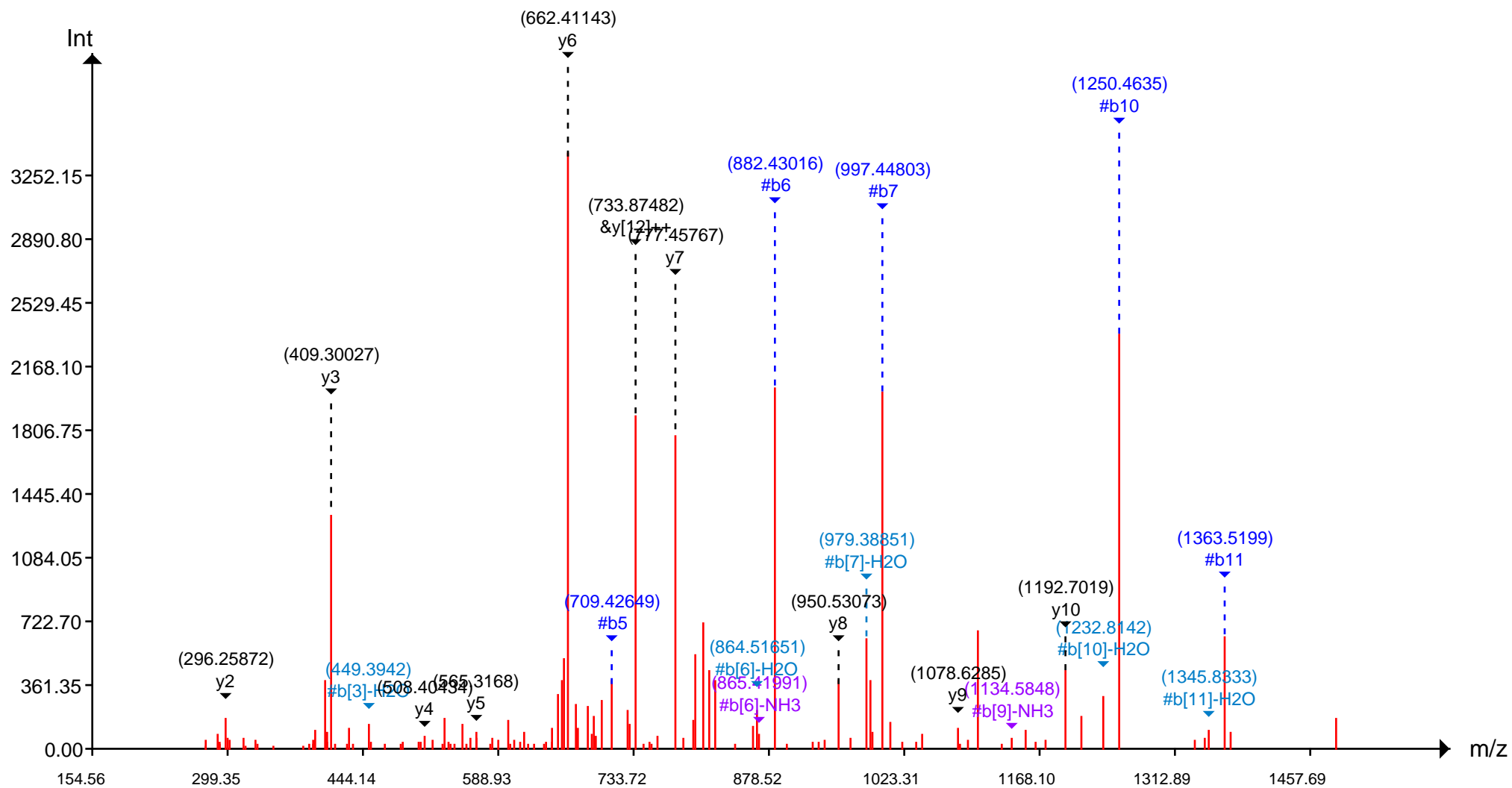
Charge 2+

Mass error 0.001481



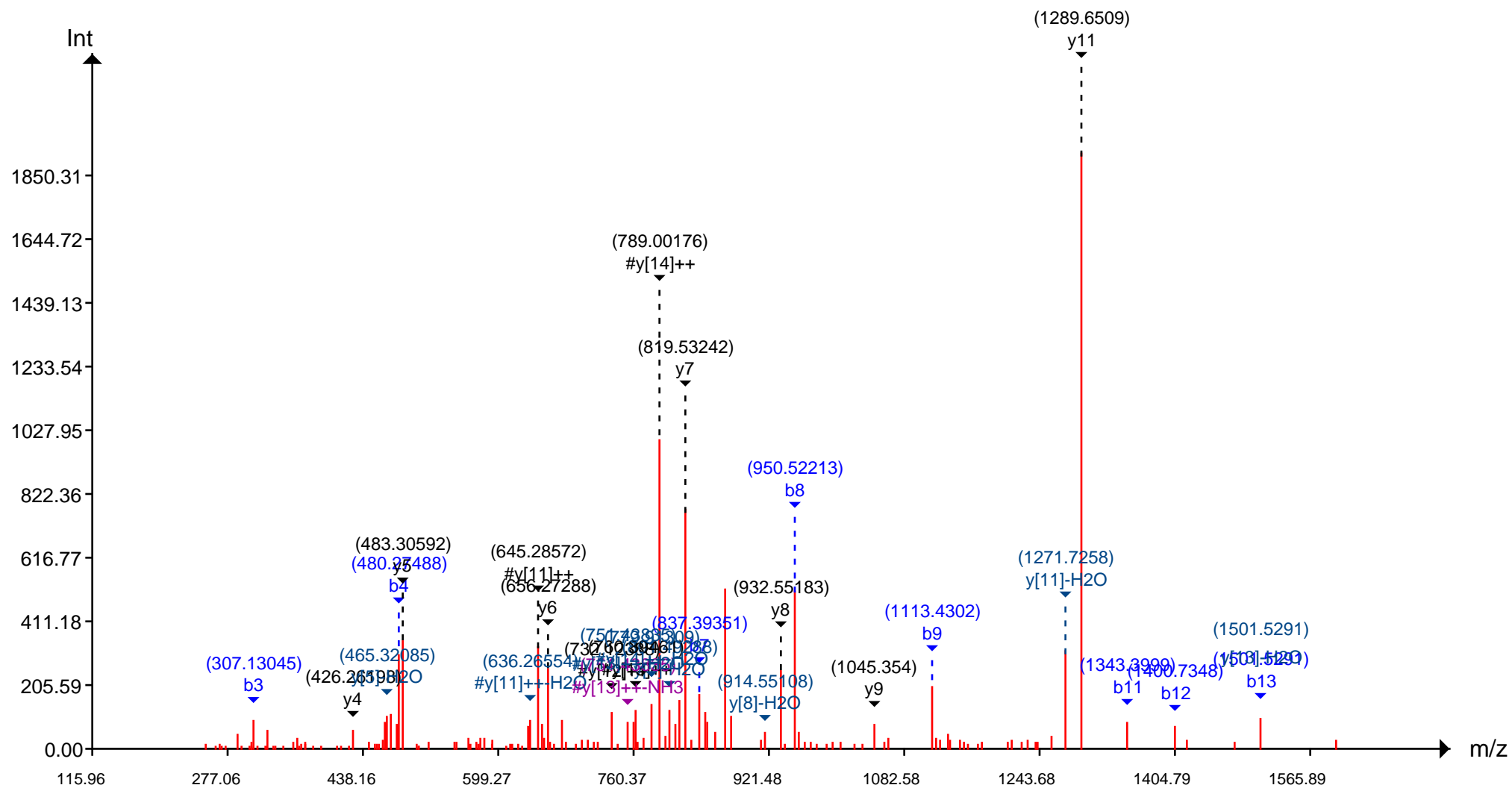
Info

Modified Sequence	AcD3-FQQGDM<Mox*>ESNGK<AcD3K*>YITK<AcD3K*>SGAR<C13>-COOH	Ion score	86.79
Protein	P06745	Confidence level	99.00%
Precursor m/z	1137.5606	Identity Threshold	47.34
Charge	2+	Mass error	0.002659



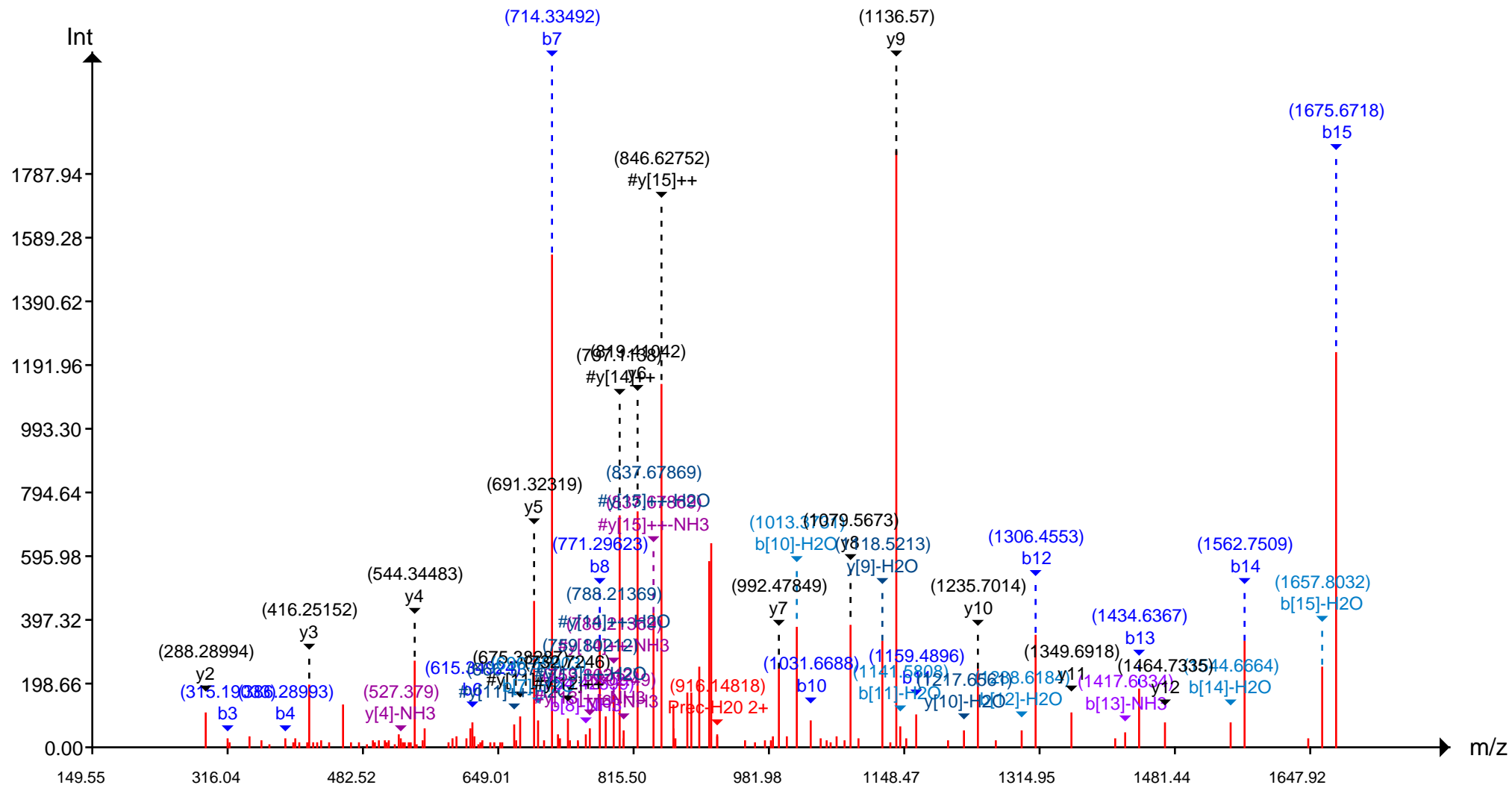
Info

Modified Sequence	AcD3-FTK<AcD3K*>NQK<AcD3K*>DPGVLDLDR<C13>-COOH	Ion score	49.18
Protein	P50543	Confidence level	99.00%
Precursor m/z	829.96235	Identity Threshold	47.14
Charge	2+	Mass error	0.002021



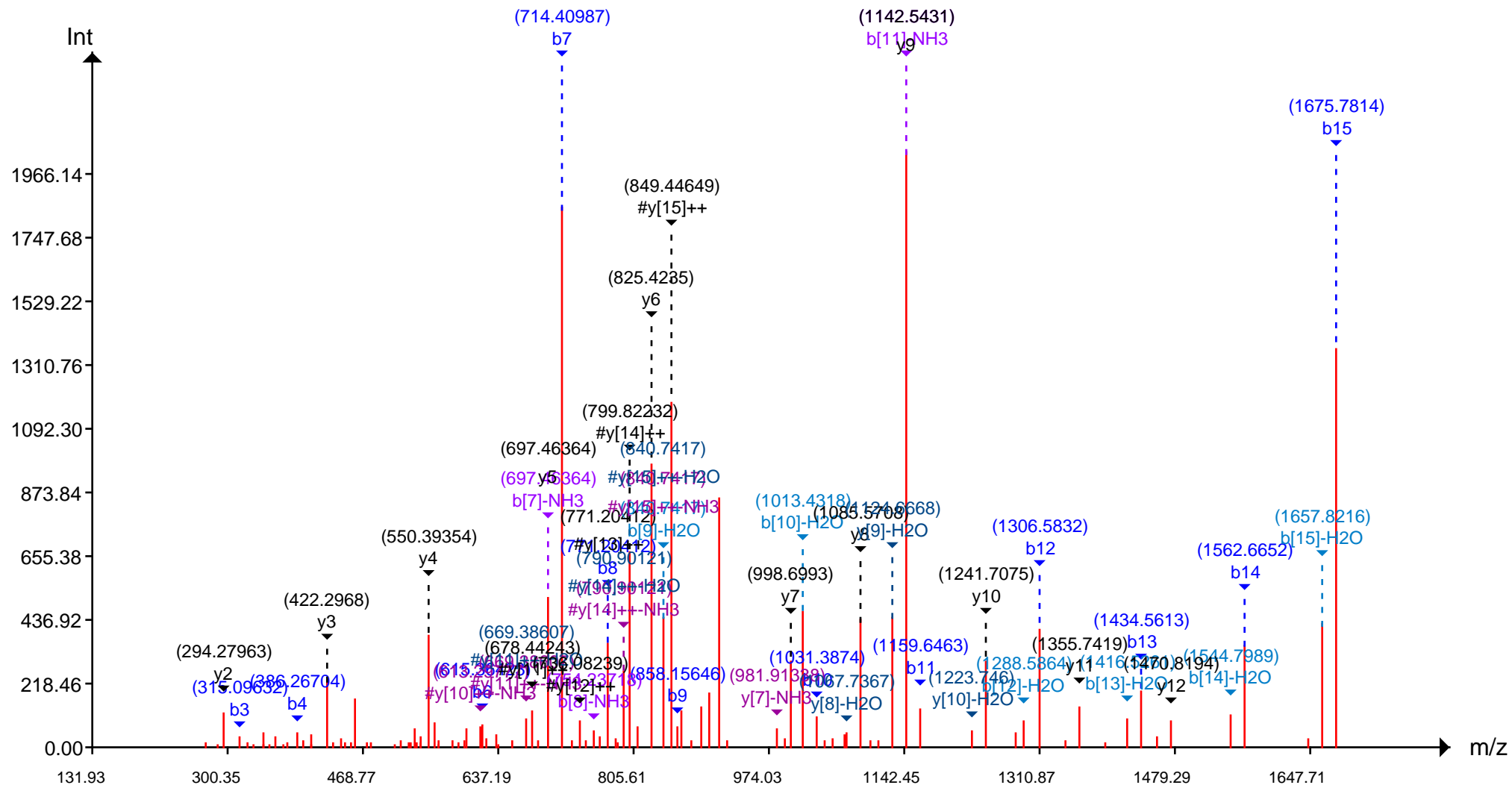
Info

Modified Sequence	AcD3-FGGK<AcD3K*>PM<Mox*>IIYK<AcD3K*>GGTSR<C13>-COOH	Ion score	57.05
Protein	P13020	Confidence level	99.00%
Precursor m/z	884.98959	Identity Threshold	47.13
Charge	2+	Mass error	0.001097



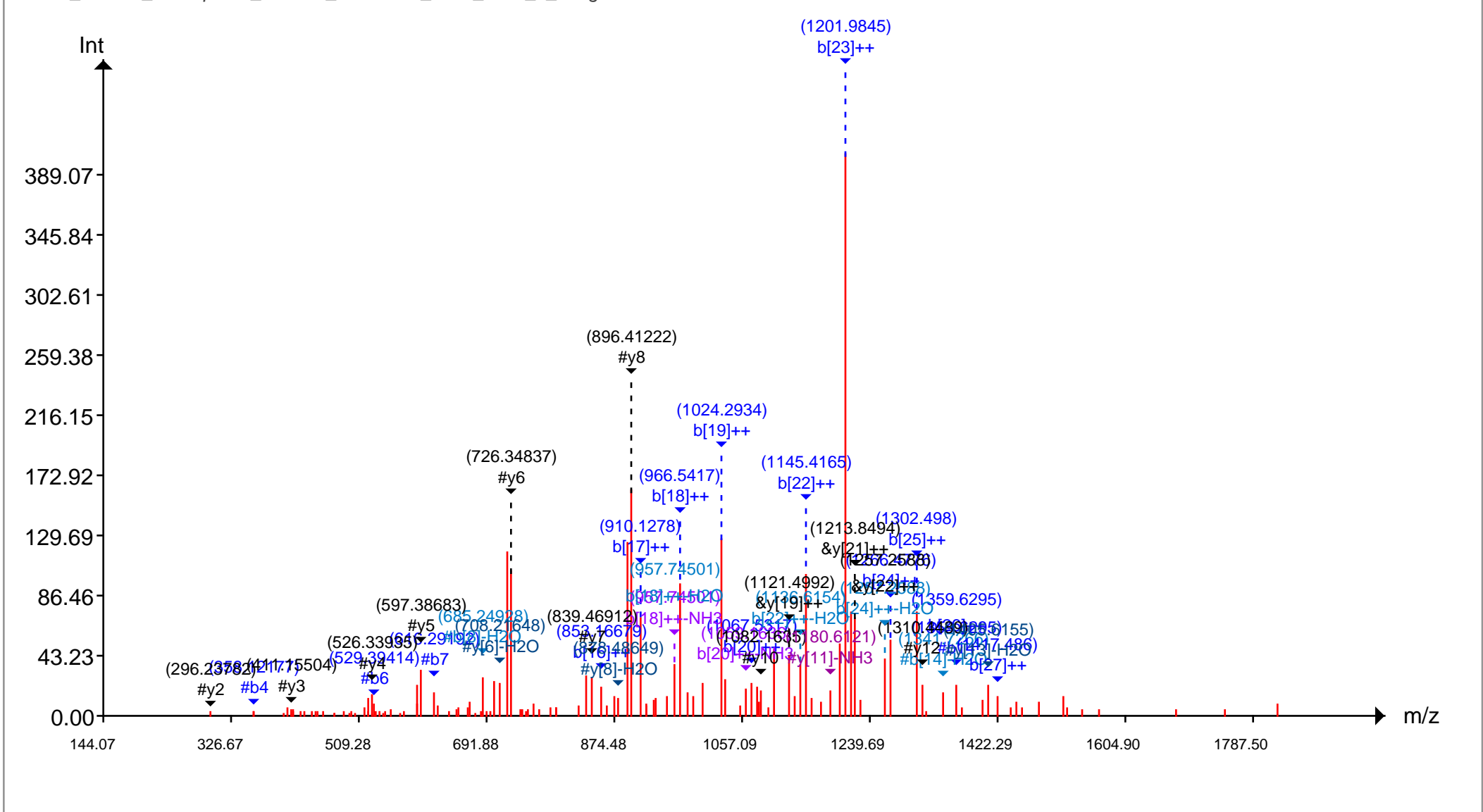
Info

Modified Sequence	AcD3-IVGADNVGSK<AcD3K*>QM<Mox*>QQIR-COOH	Ion score	93.24
Protein	P14869	Confidence level	99.00%
Precursor m/z	925.48981	Identity Threshold	47.99
Charge	2+	Mass error	0.001613



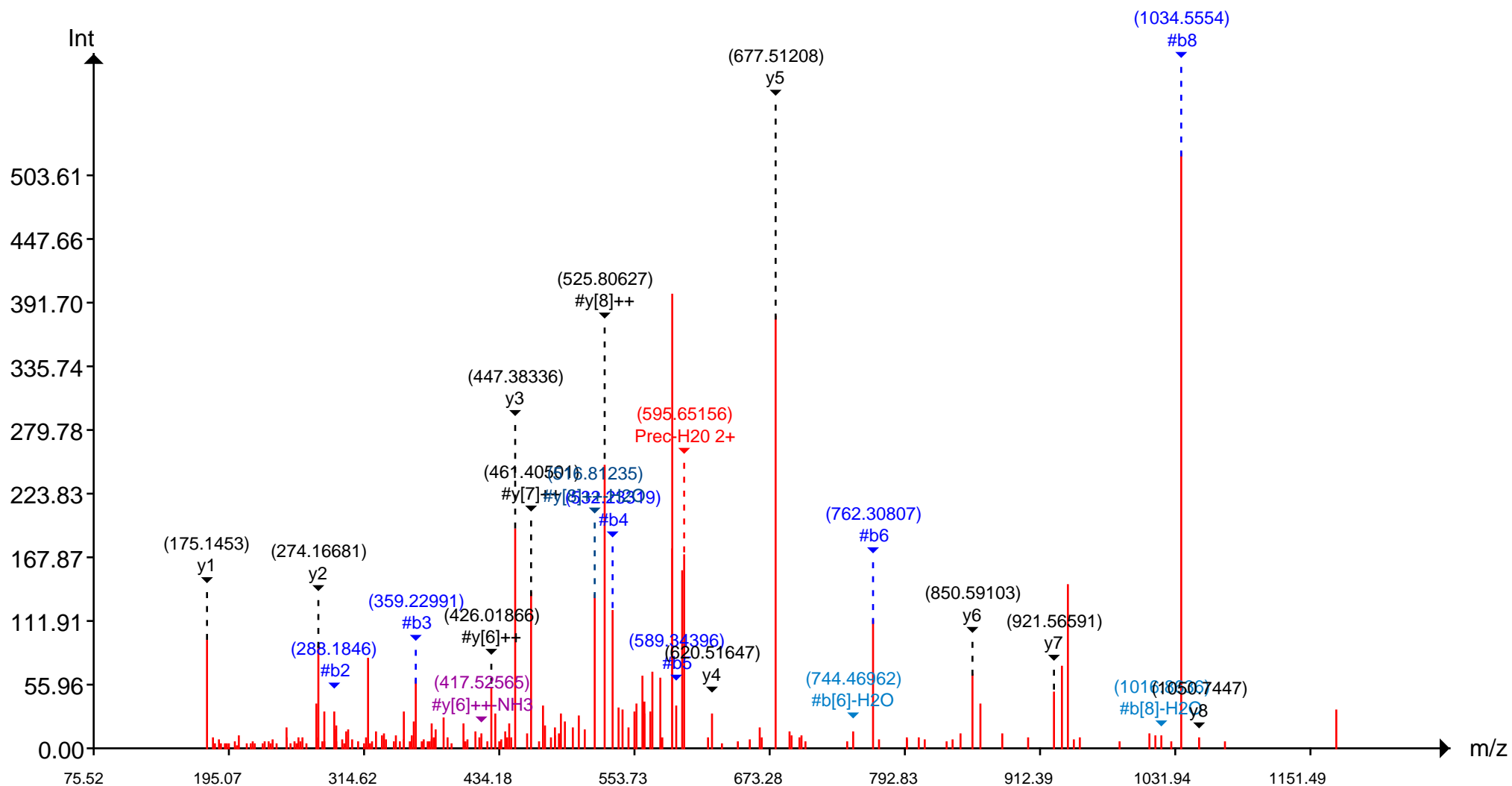
Info

Modified Sequence	AcD3-IVGADNVGSK<AcD3K*>QM<Mox*>QQIR<C13>-COOH	Ion score	103.07
Protein	P14869	Confidence level	99.00%
Precursor m/z	928.4997	Identity Threshold	47.76
Charge	2+	Mass error	0.001251



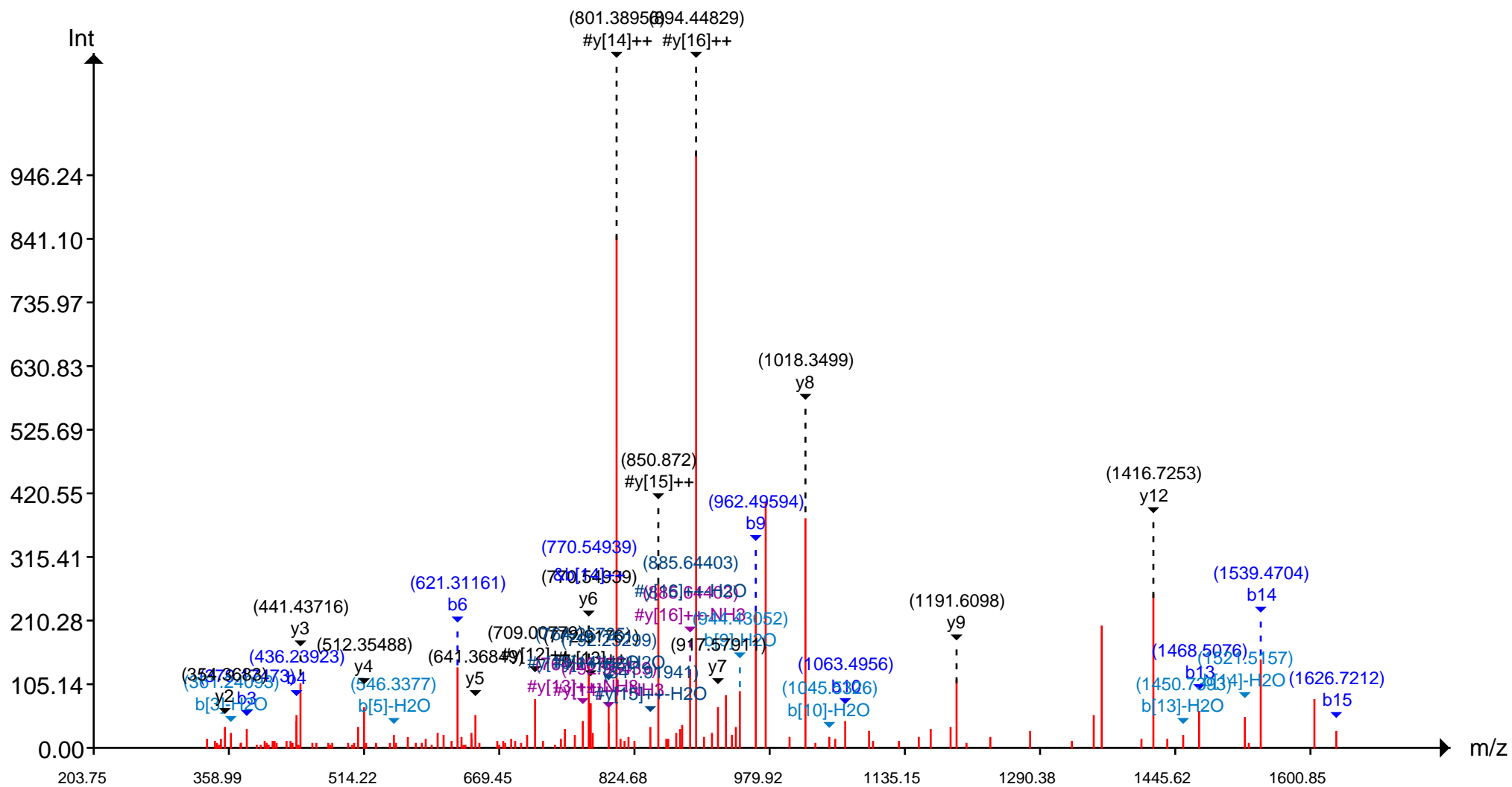
Info

Modified Sequence	AcD3-AALGGNSSPSAK<AcD3K*>DIK<AcD3K*>K<AcD3K*>ILDSVGIEADDDR<C13>-COOH	Ion score	49.07
Protein	P99027	Confidence level	99.00%
Precursor m/z	1043.5532	Identity Threshold	47.22
Charge	3+	Mass error	8.09E-4



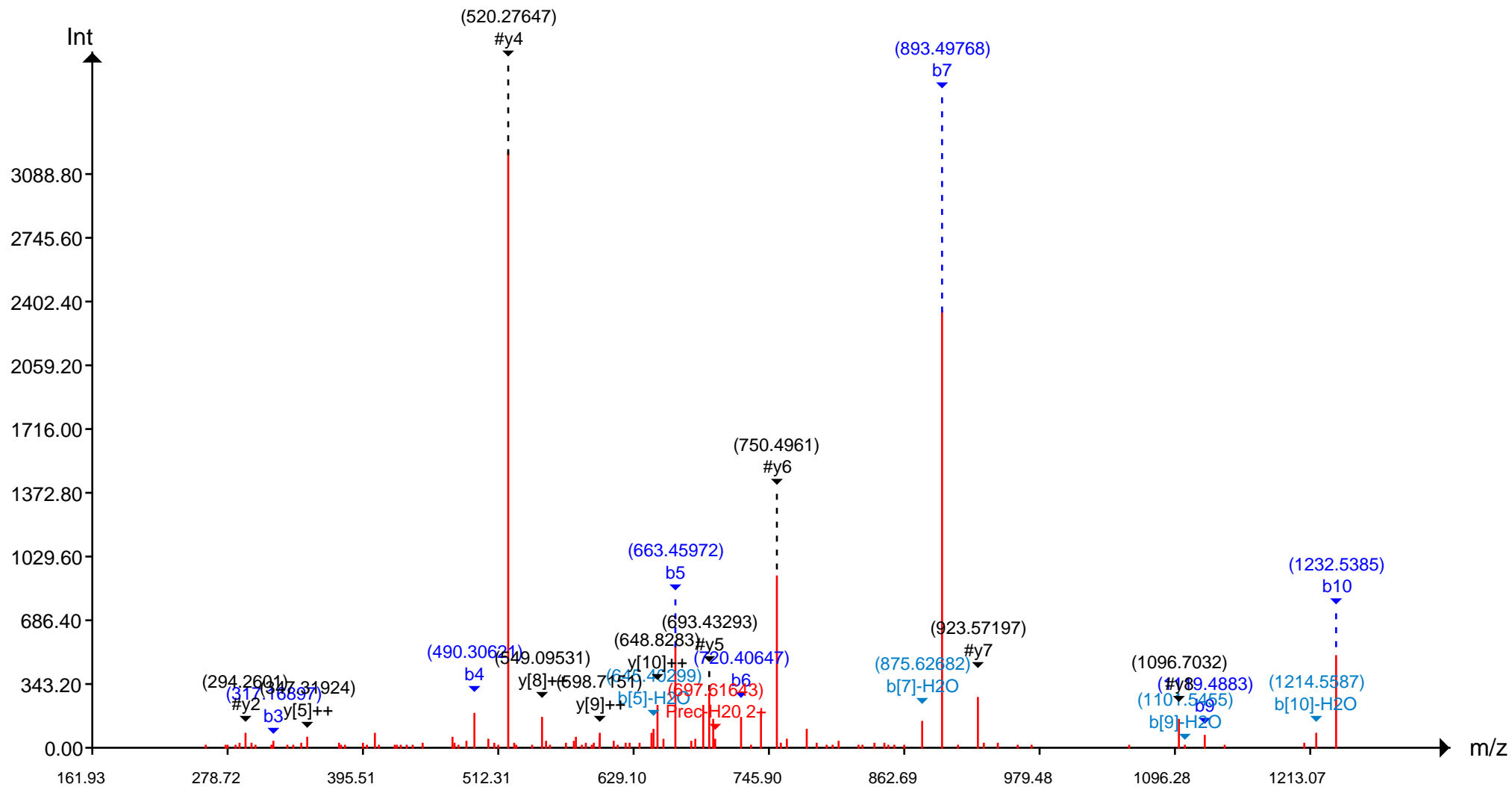
Info

Modified Sequence	AcD3-IEAK<AcD3K*>GK<AcD3K*>K<AcD3K*>VR-COOH	Ion score	39.35
Protein	P42208	Confidence level	99.00%
Precursor m/z	604.8913	Identity Threshold	38.50
Charge	2+	Mass error	2.72E-4



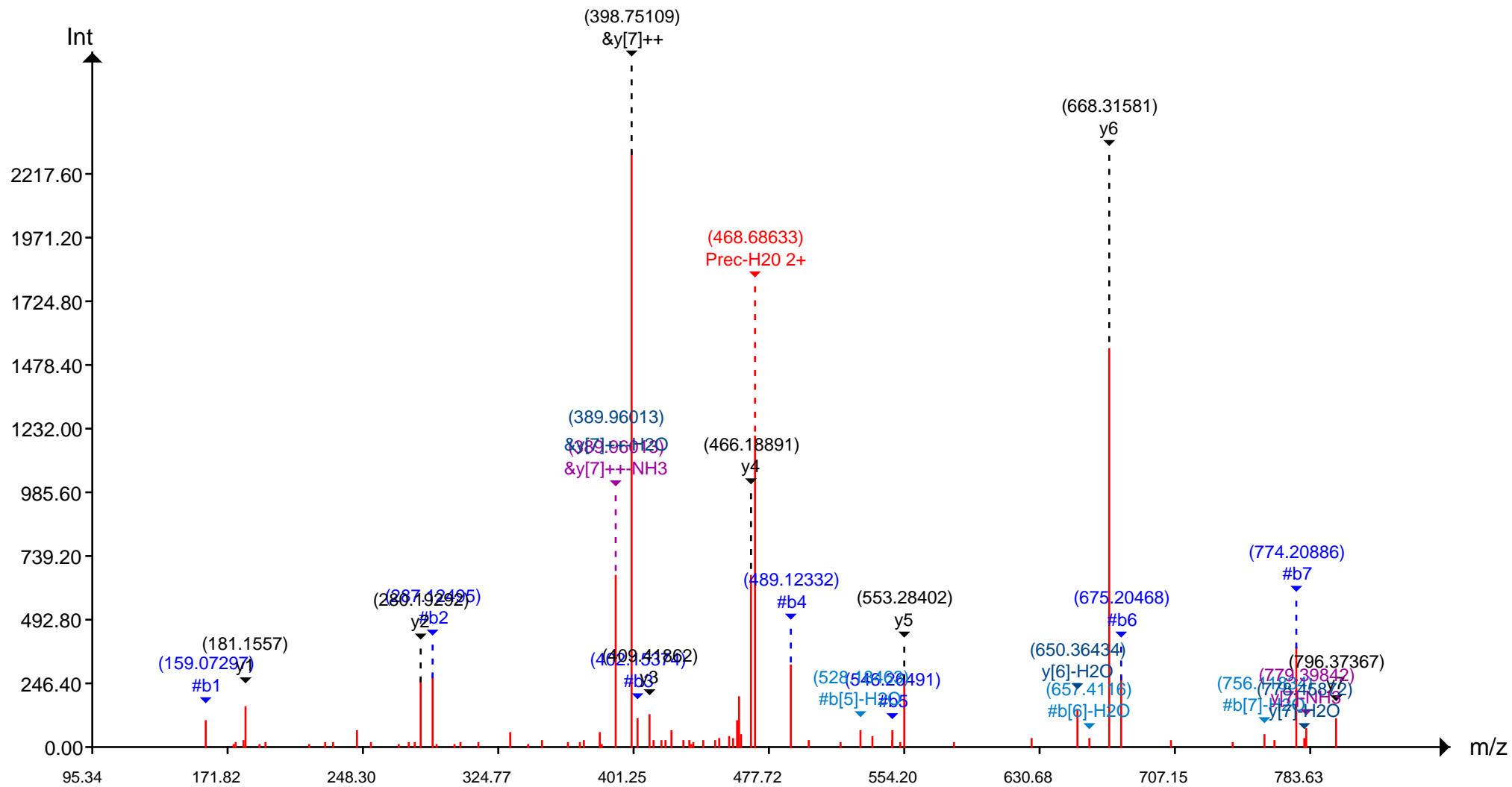
Info

Modified Sequence	AcD3-FSVGQGPAK<AcD3K*>TM<Mox*>EEASK<AcD3K*>R<C13>-COOH	Ion score	68.68
Protein	O70310	Confidence level	99.00%
Precursor m/z	990.51234	Identity Threshold	48.06
Charge	2+	Mass error	0.00256



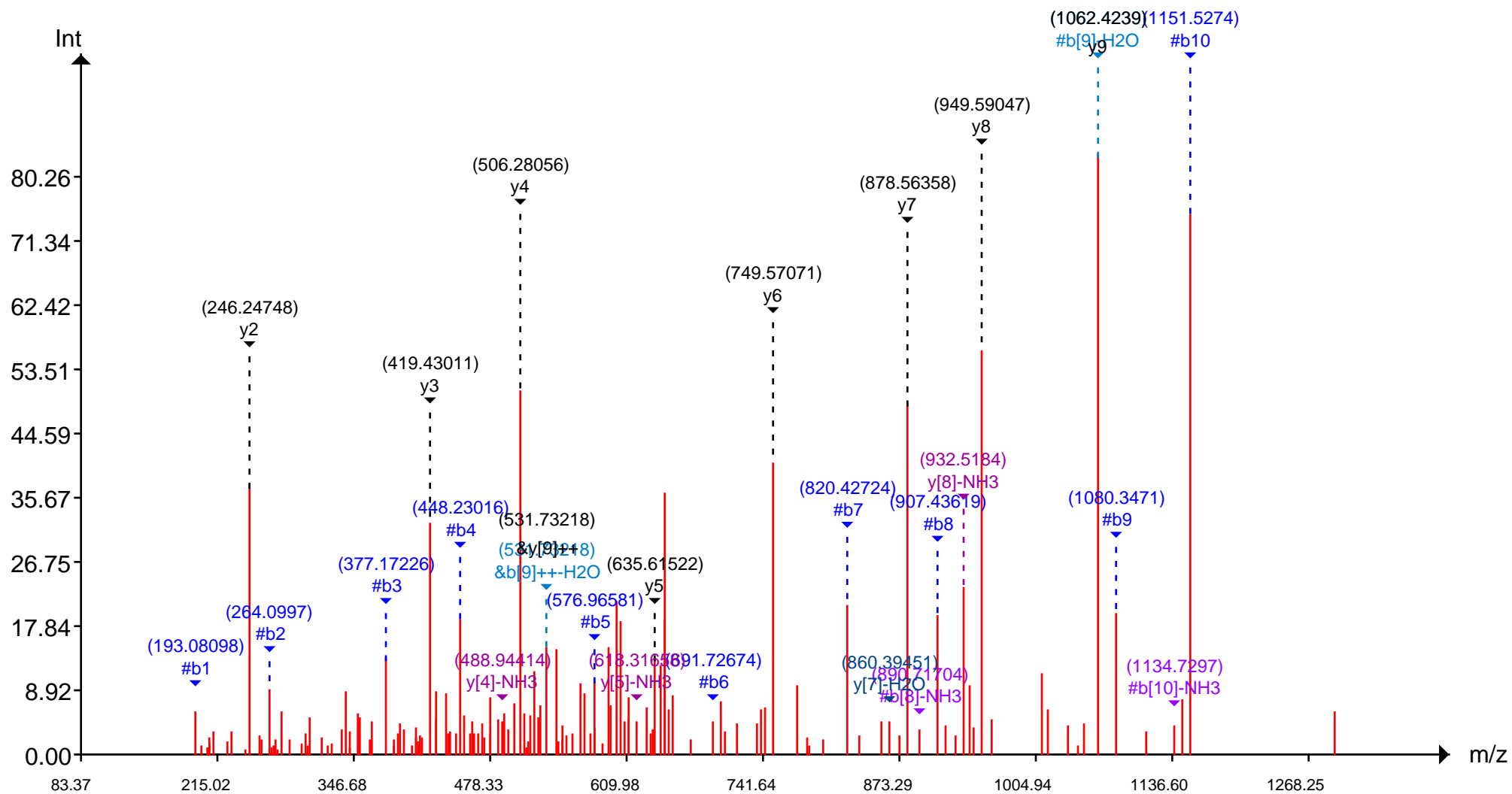
Info

Modified Sequence	AcD3-ATVK<AcD3K*>K<AcD3K*>GK<AcD3K*>PELR<C13>-COOH	Ion score	46.67
Protein	P62830	Confidence level	99.00%
Precursor m/z	706.9518	Identity Threshold	38.29
Charge	2+	Mass error	6.81E-4



Info

Modified Sequence	AcD3-LQDSGEVR<C13>-COOH	Ion score	57.44
Protein	P63242	Confidence level	99.00%
Precursor m/z	477.75459	Identity Threshold	47.48
Charge	2+	Mass error	-6.57E-4



Info

Modified Sequence AcD3-FAIAENESK<AcD3K*>AR-COOH

Ion score 73.26

Protein Q923D4

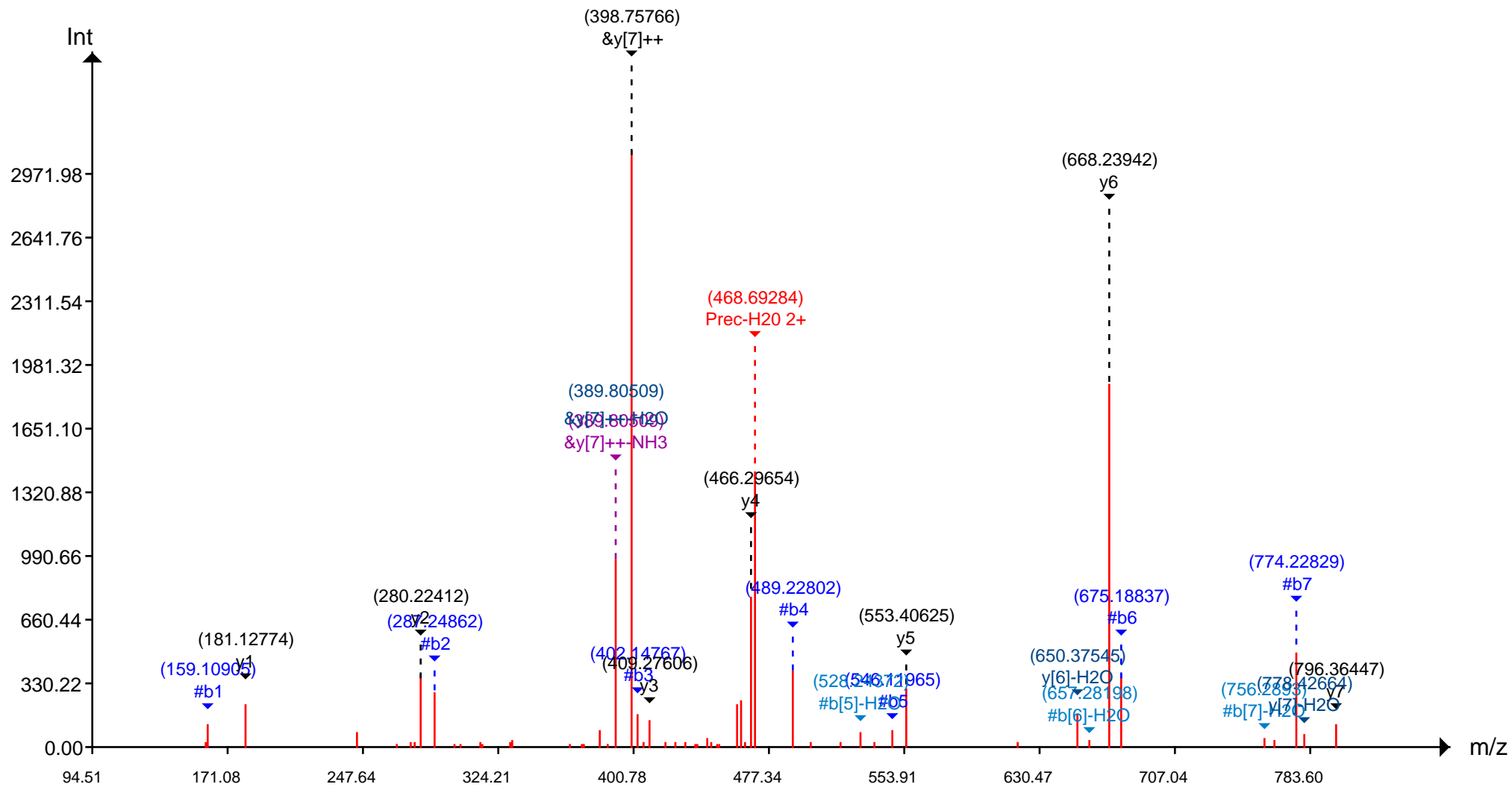
Confidence level 99.00%

Precursor m/z 663.35233

Identity Threshold 47.97

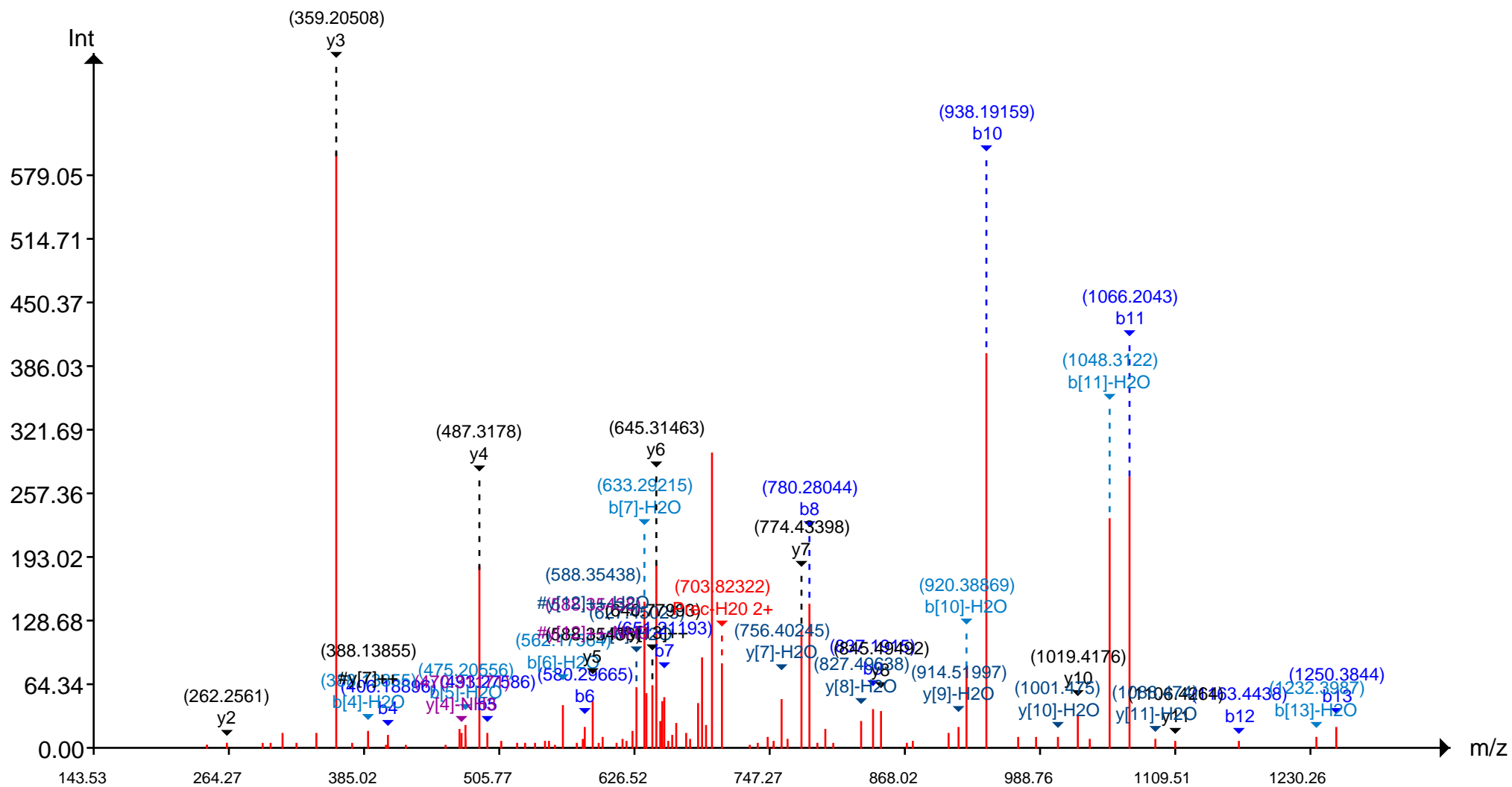
Charge 2+

Mass error 7.62E-4



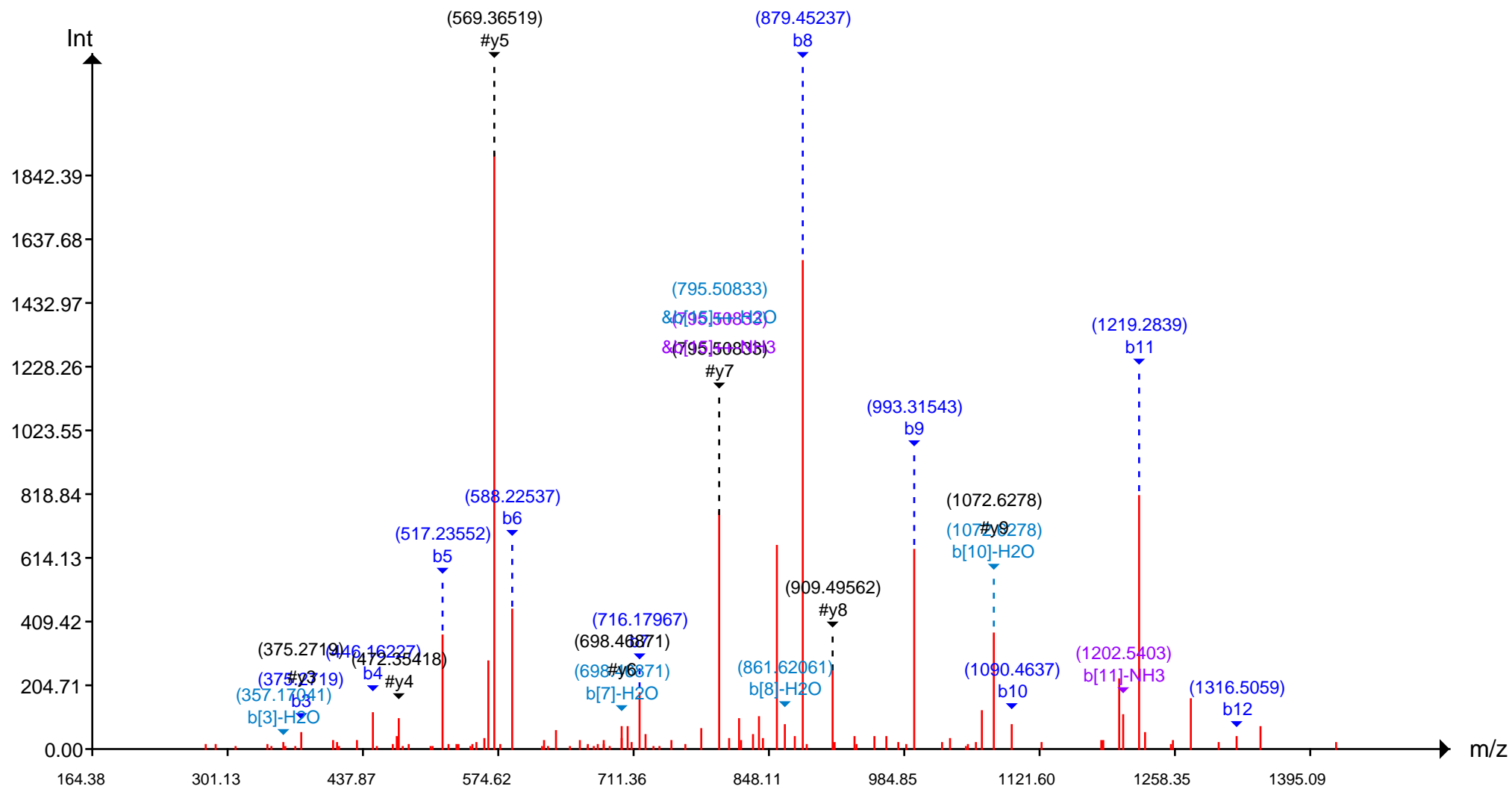
Info

Modified Sequence	AcD3-LQDSGEVR<C13>-COOH	Ion score	56.08
Protein	P63242	Confidence level	99.00%
Precursor m/z	477.75447	Identity Threshold	47.50
Charge	2+	Mass error	-8.97E-4



Info

Modified Sequence	AcD3-VSSSSSAEGTQPSR-COOH	Ion score	77.51
Protein	Q8R317	Confidence level	99.00%
Precursor m/z	712.8386	Identity Threshold	46.08
Charge	2+	Mass error	8.13E-4



Info

Modified Sequence AcD3-ISEAAQYNPEPPPPR<C13>-COOH

Ion score 78.99

Protein O88456

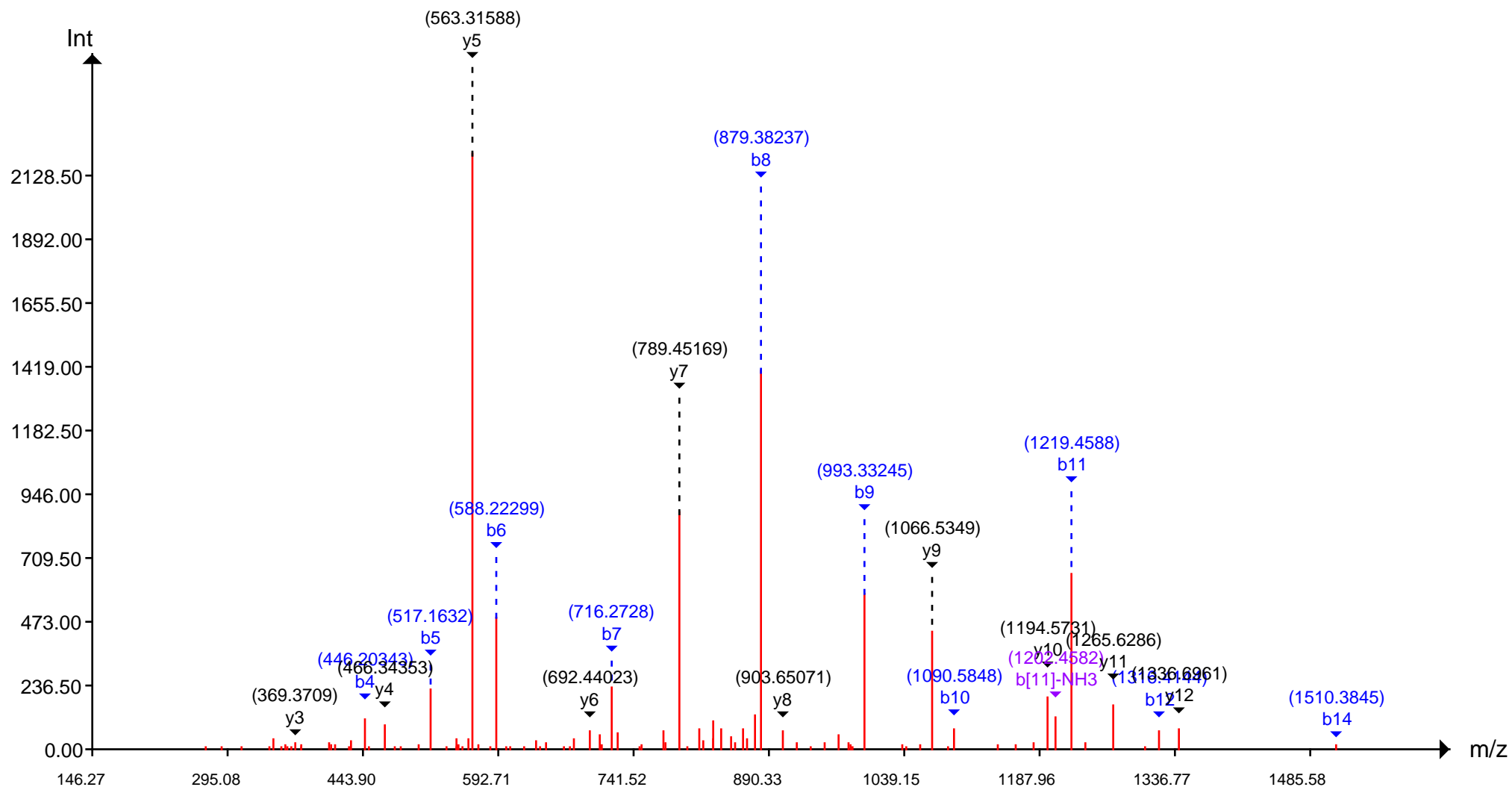
Confidence level 99.00%

Precursor m/z 894.45945

Identity Threshold 48.05

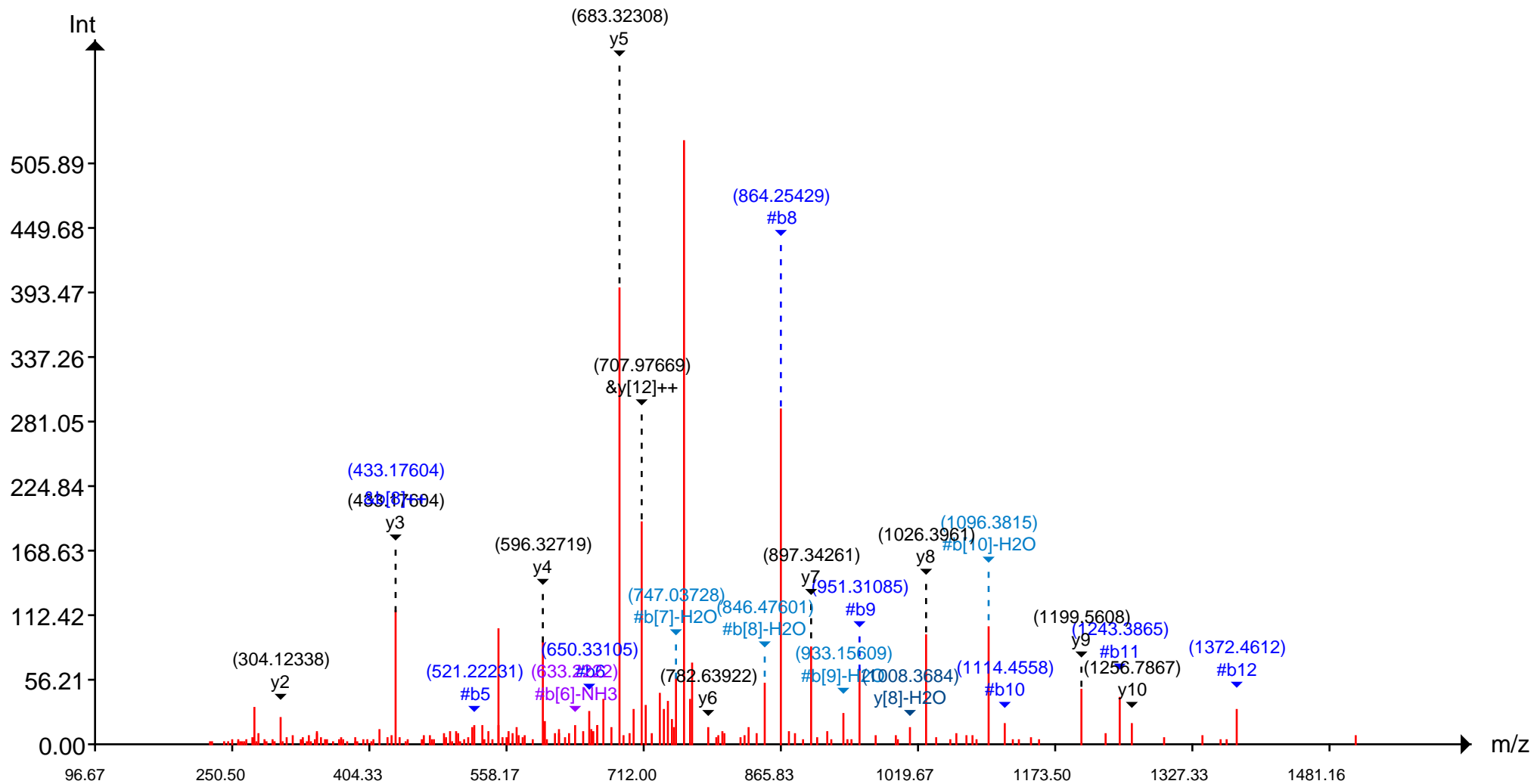
Charge 2+

Mass error 0.001913



Info

Modified Sequence	AcD3-ISEAAQYNPEPPPPR-COOH	Ion score	85.98
Protein	O88456	Confidence level	99.00%
Precursor m/z	891.44917	Identity Threshold	47.86
Charge	2+	Mass error	0.001494



Info

Modified Sequence AcD3-STGGK<AcD3K*>EDVSYEER-COOH

Ion score 58.92

Protein Q9R257

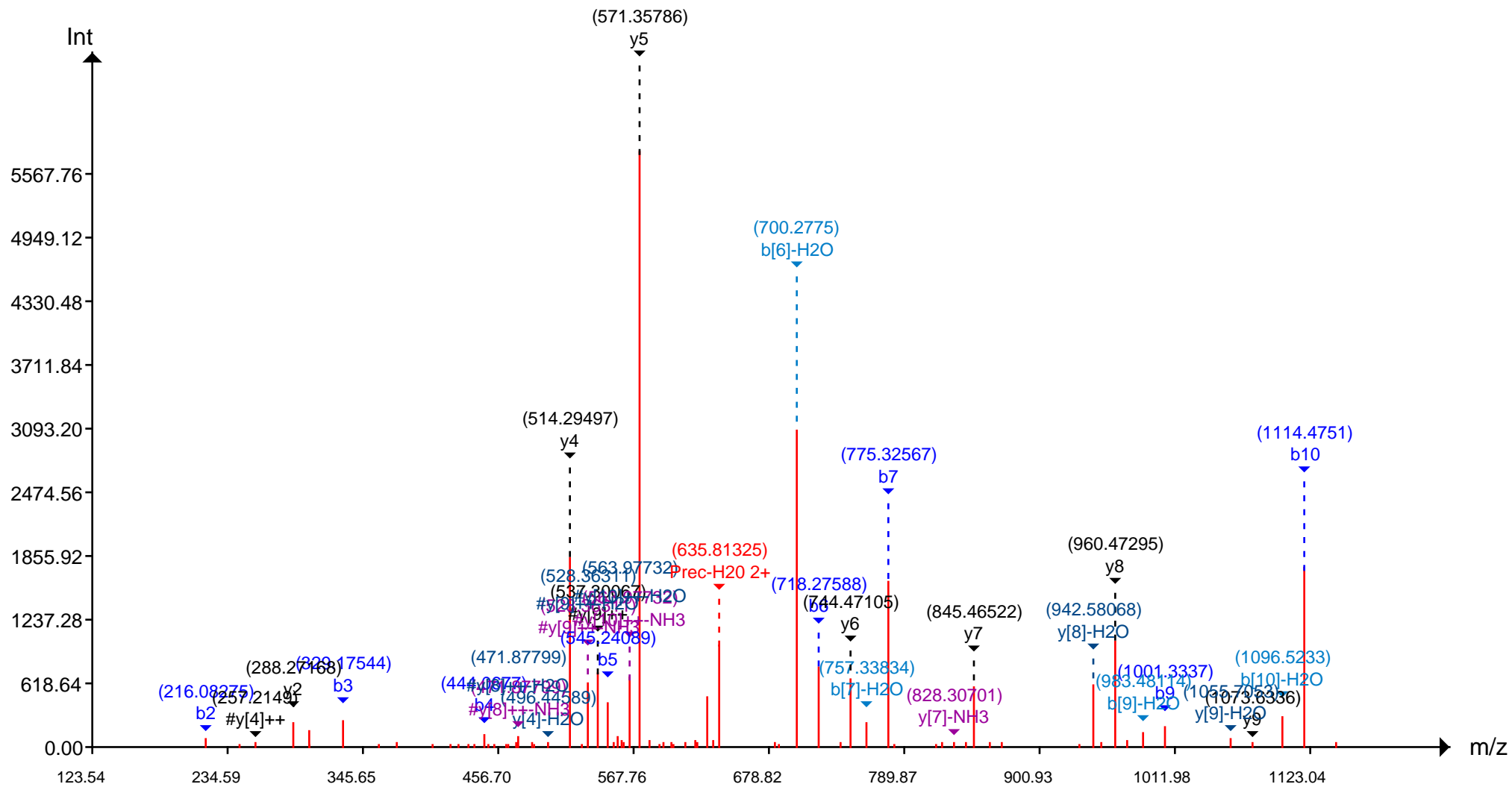
Confidence level 99.00%

Precursor m/z 773.86043

Identity Threshold 45.36

Charge 2+

Mass error -2.35E-4



Info

Modified Sequence AcD3-VALDTK<AcD3K*>GPEIR-COOH

Ion score 57.07

Protein P52480

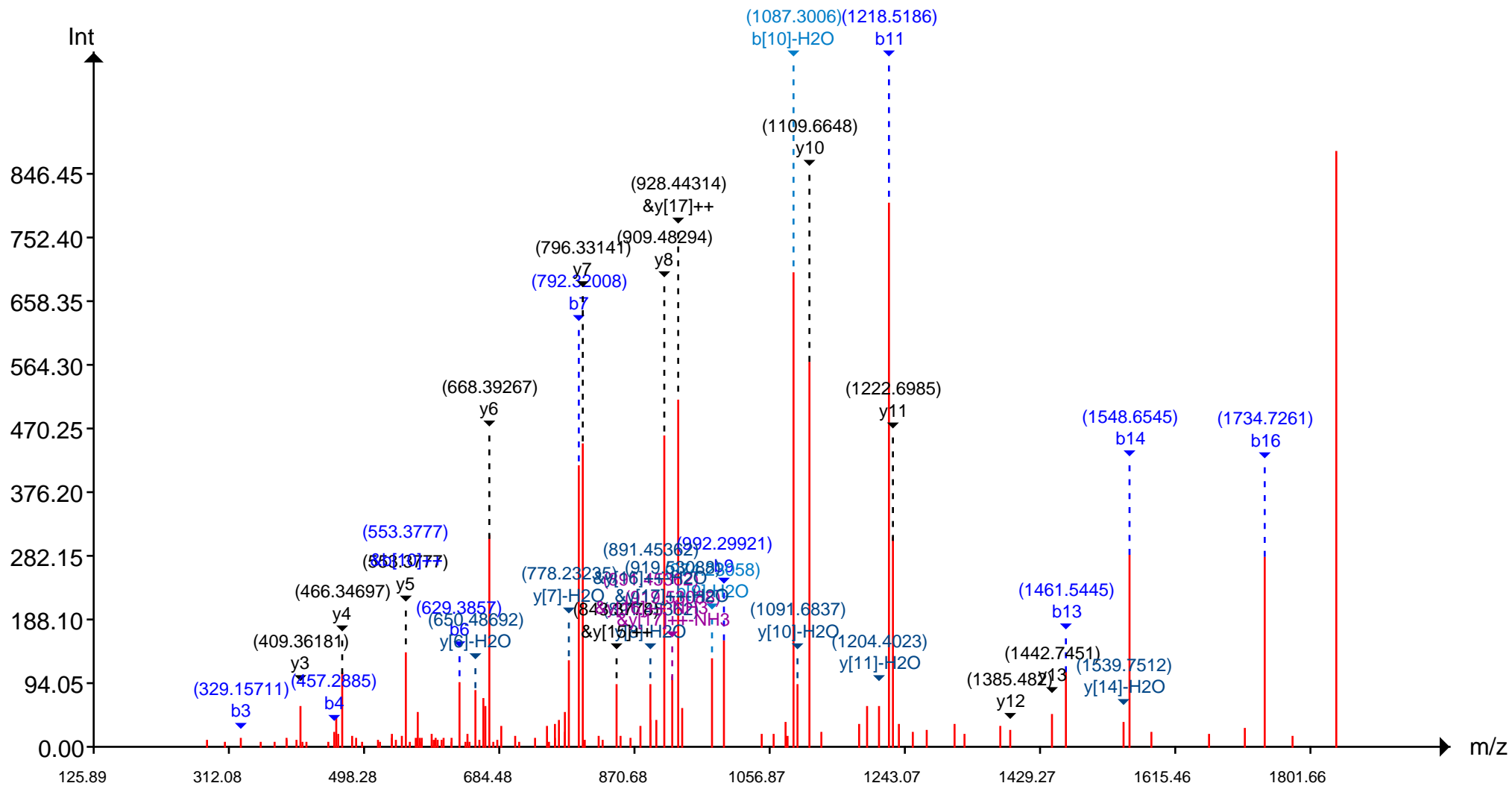
Confidence level 99.00%

Precursor m/z 644.8731

Identity Threshold 46.56

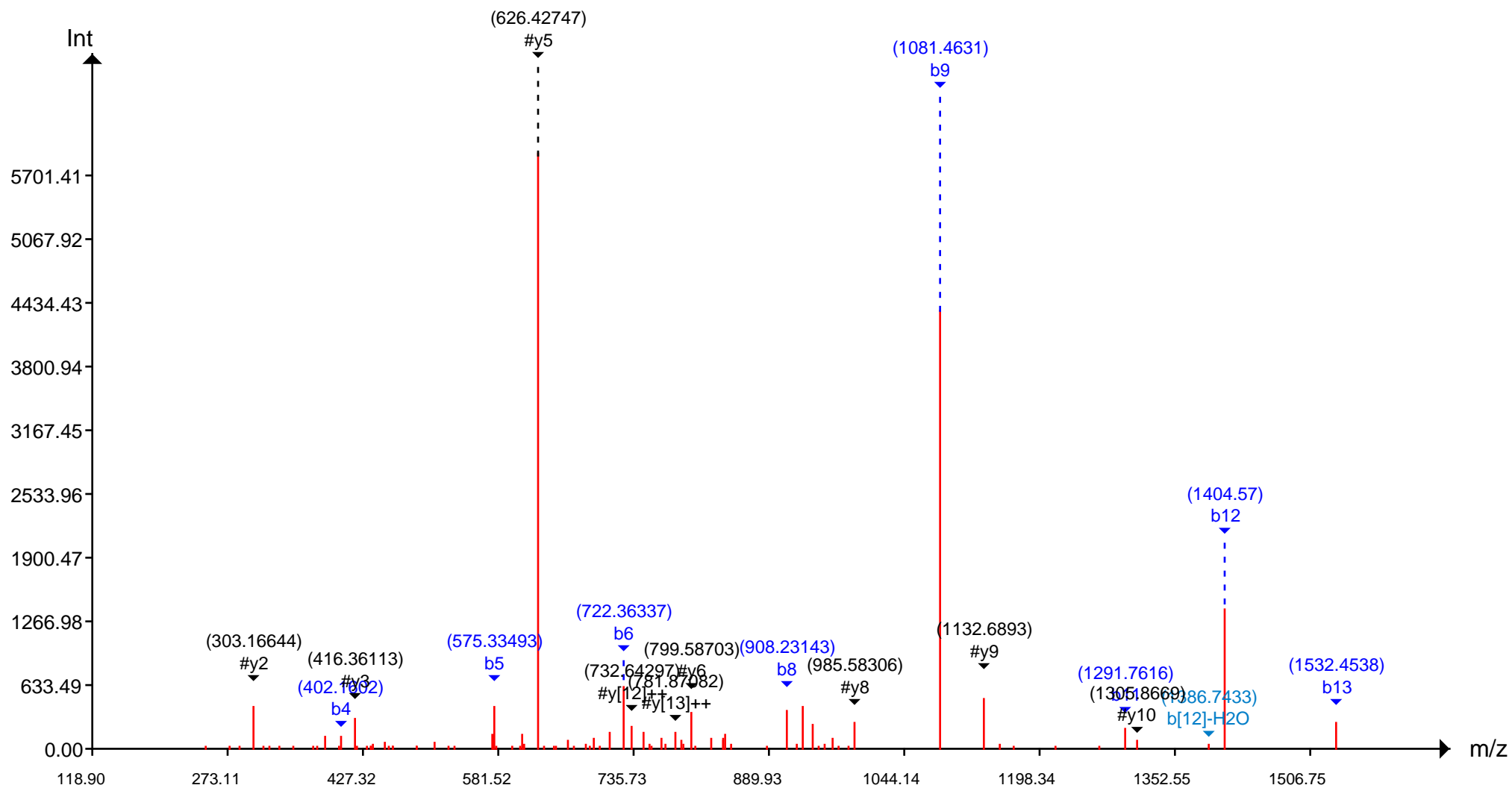
Charge 2+

Mass error 0.001133



Info

Modified Sequence	AcD3-IGIQDGYLSLLQDSGEVR<C13>-COOH	Ion score	75.43
Protein	P63242	Confidence level	99.00%
Precursor m/z	1007.5359	Identity Threshold	47.88
Charge	2+	Mass error	0.001904



Info

Modified Sequence AcD3-VVSAK<AcD3K*>FEGK<AcD3K*>PLLQR-COOH

Ion score 59.46

Protein Q8BGS2

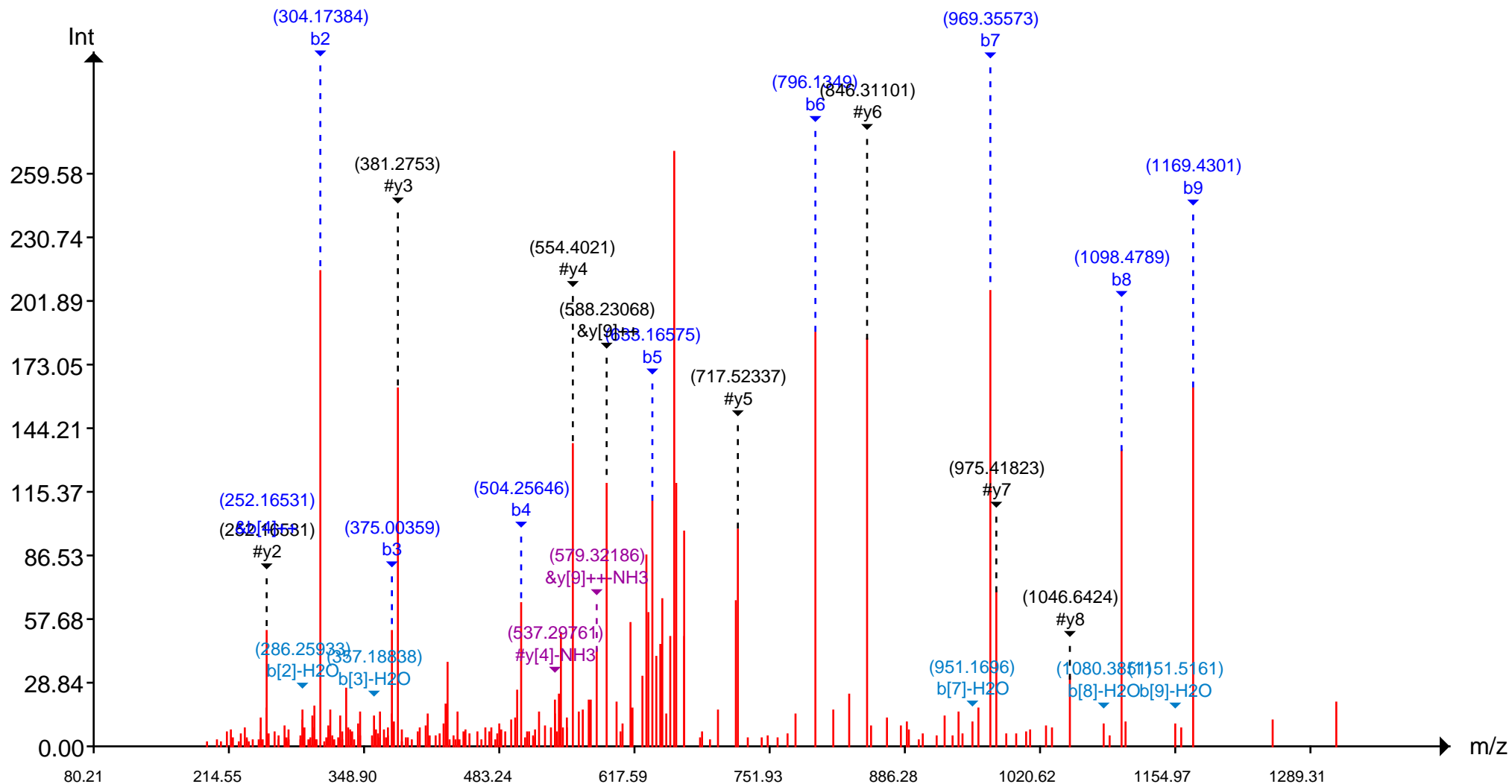
Confidence level 99.00%

Precursor m/z 854.01221

Identity Threshold 42.77

Charge 2+

Mass error 0.002178



Info

Modified Sequence AcD3-EEAEEYK<AcD3K*>EAR<C13>-COOH

Ion score 56.0

Protein P48428

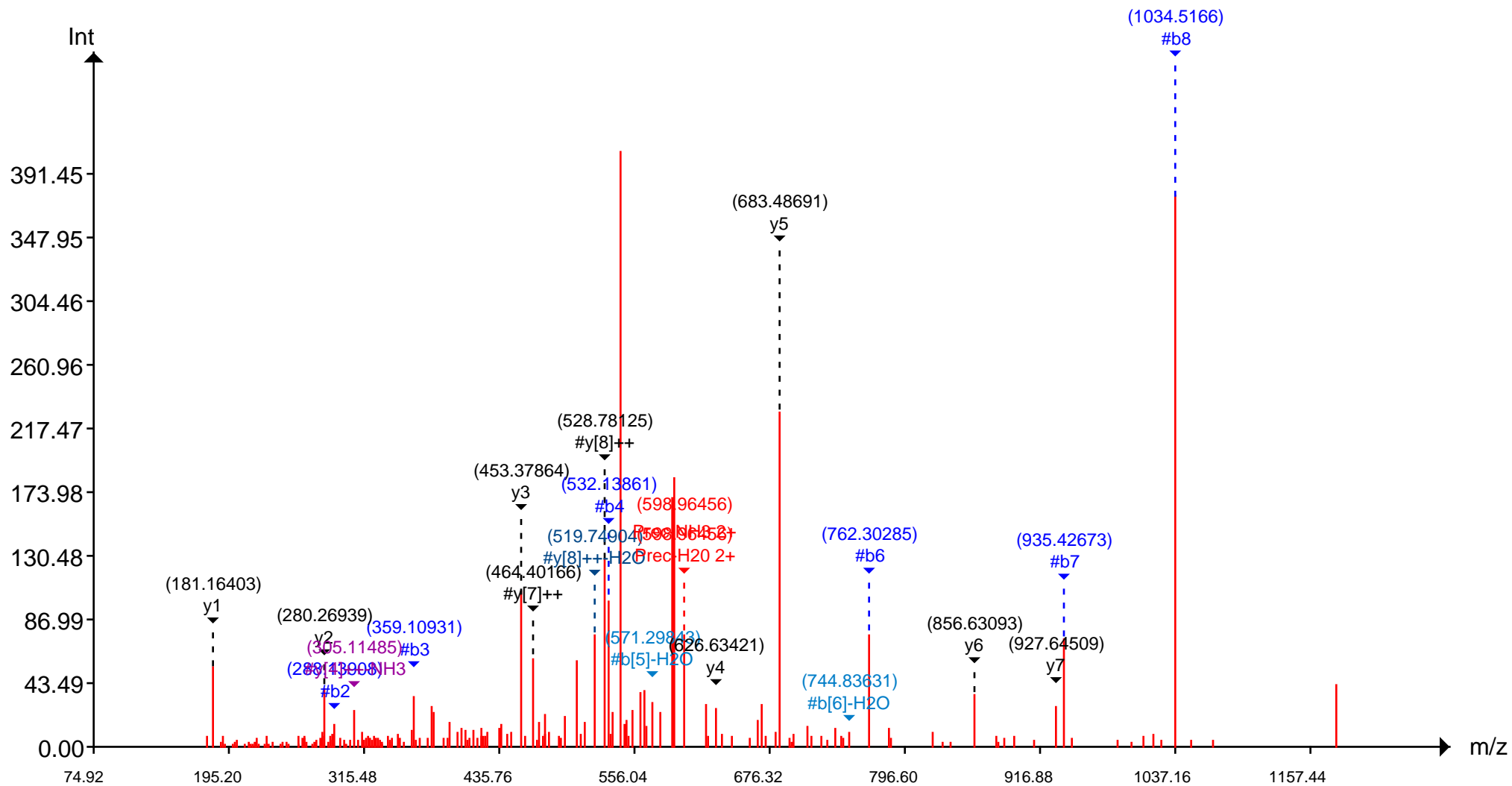
Confidence level 99.00%

Precursor m/z 675.32552

Identity Threshold 46.65

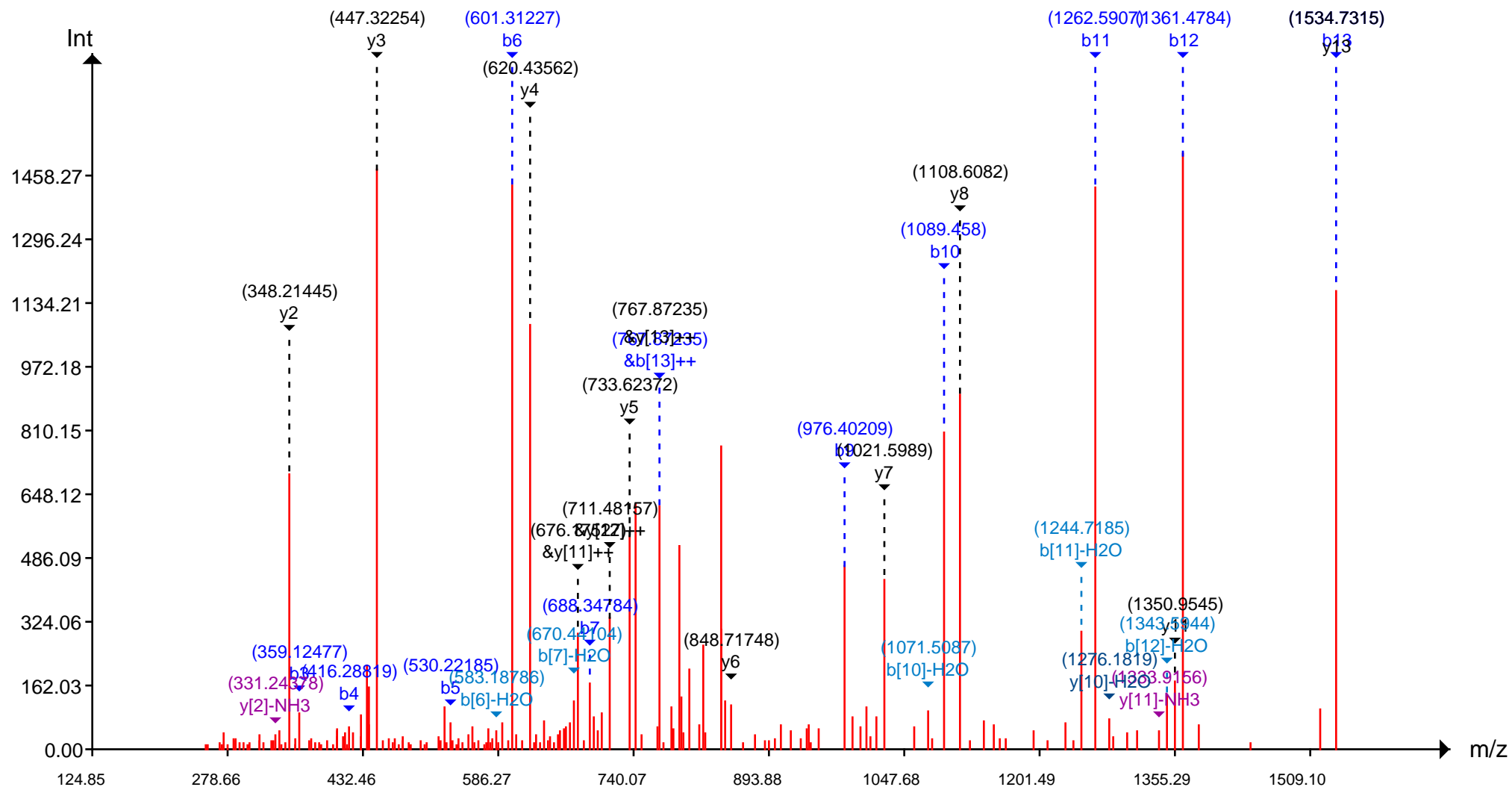
Charge 2+

Mass error 4.56E-4



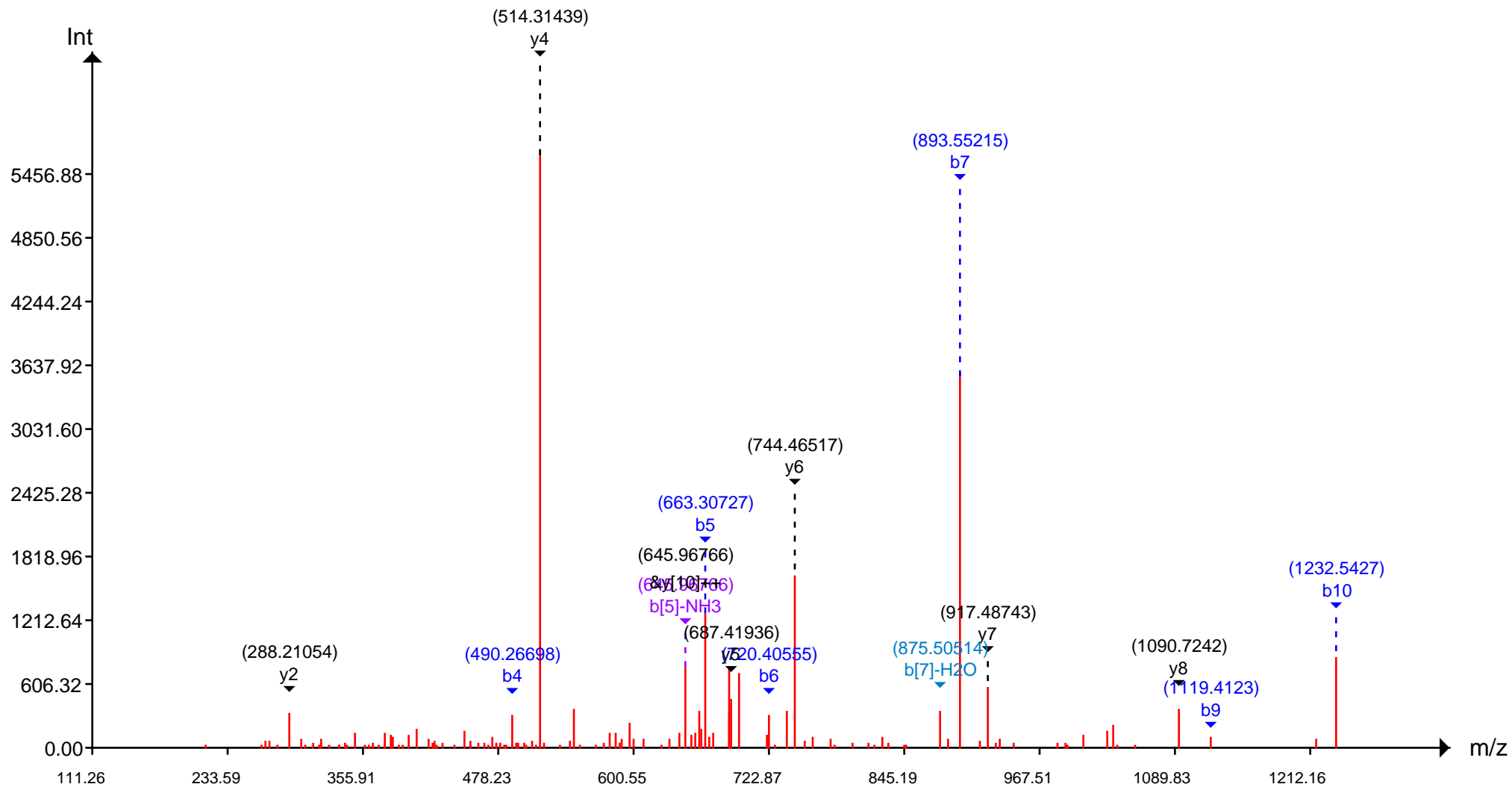
Info

Modified Sequence	AcD3-IEAK<AcD3K*>GK<AcD3K*>K<AcD3K*>VR<C13>-COOH	Ion score	42.69
Protein	P42208	Confidence level	99.00%
Precursor m/z	607.90136	Identity Threshold	37.45
Charge	2+	Mass error	2.5E-4



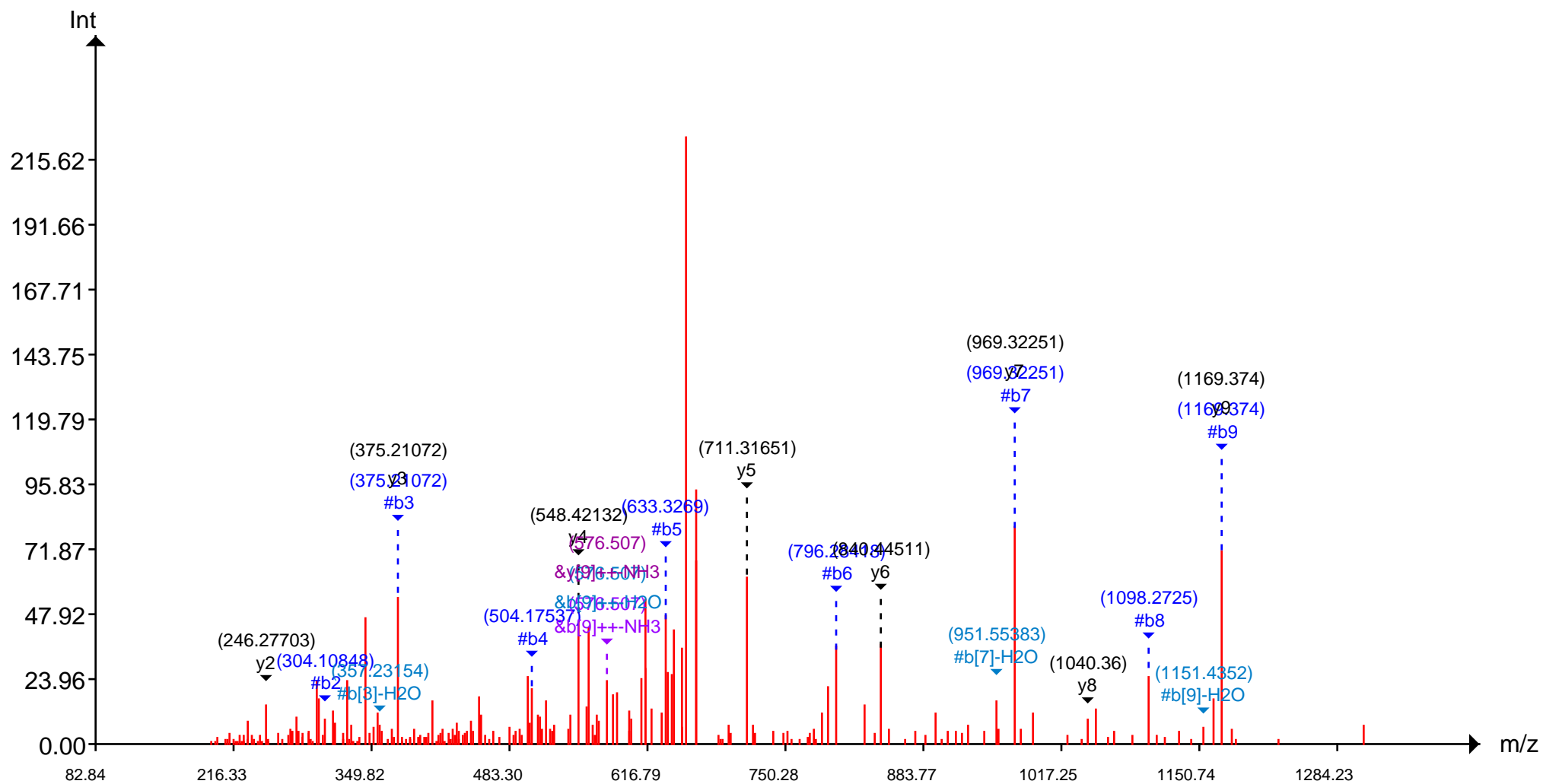
Info

Modified Sequence	AcD3-ELAGNASK<AcD3K*>DLK<AcD3K*>VK<AcD3K*>R-COOH	Ion score	54.47
Protein	Q3THW5, P0C0S6	Confidence level	99.00%
Precursor m/z	855.00551	Identity Threshold	43.83
Charge	2+	Mass error	0.005623



Info

Modified Sequence	AcD3-ATVK<AcD3K*>K<AcD3K*>GK<AcD3K*>PELR-COOH	Ion score	50.79
Protein	P62830	Confidence level	99.00%
Precursor m/z	703.94208	Identity Threshold	40.04
Charge	2+	Mass error	0.001383



Info

Modified Sequence AcD3-EEAEEYK<AcD3K*>EAR-COOH

Ion score 47.92

Protein P48428

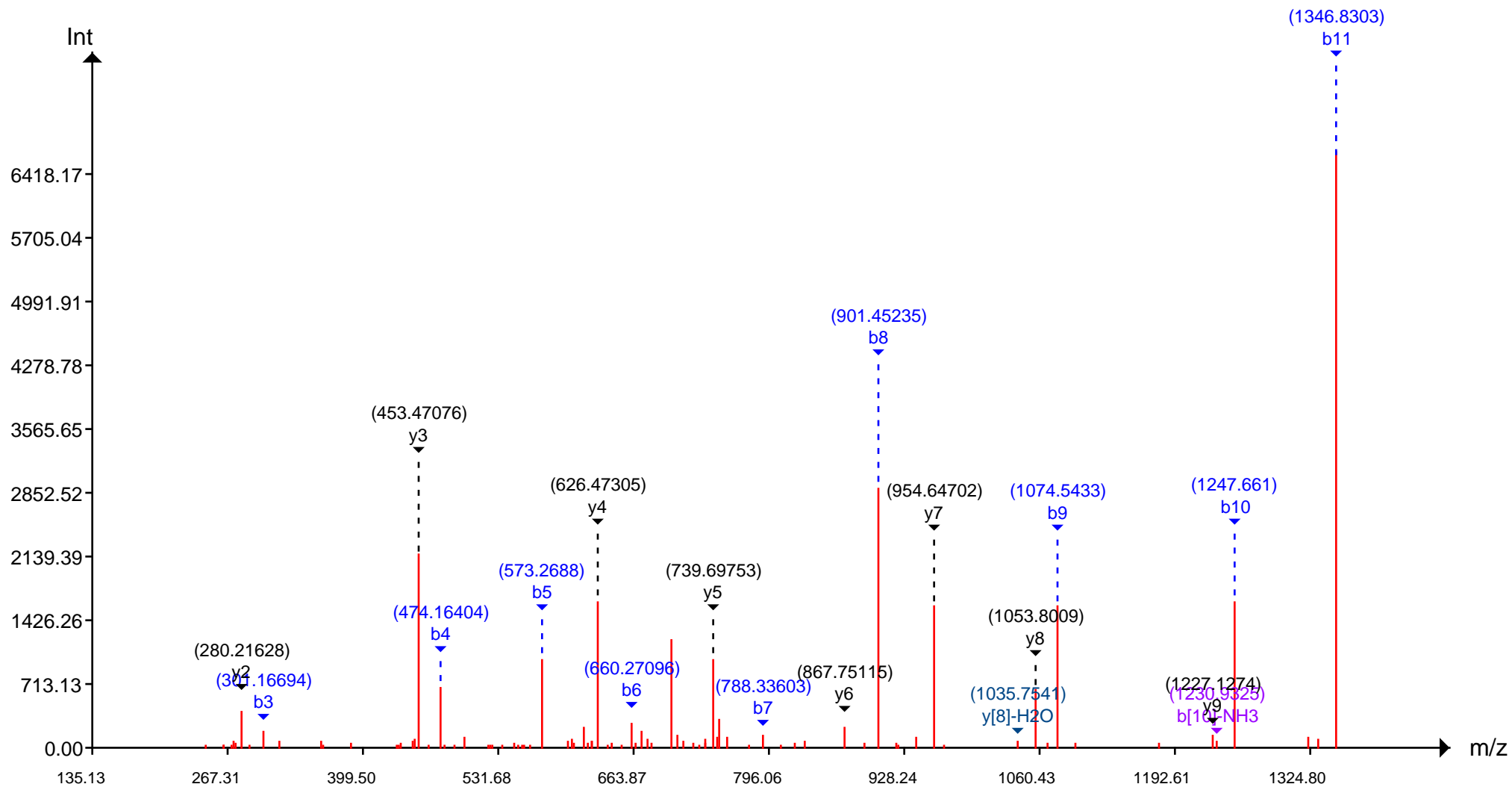
Confidence level 99.00%

Precursor m/z 672.31537

Identity Threshold 45.55

Charge 2+

Mass error 2.97E-4



Info

Modified Sequence AcD3-AALK<AcD3K*>VSQIK<AcD3K*>K<AcD3K*>VR<C13>-COOH

Ion score 76.67

Protein P26638

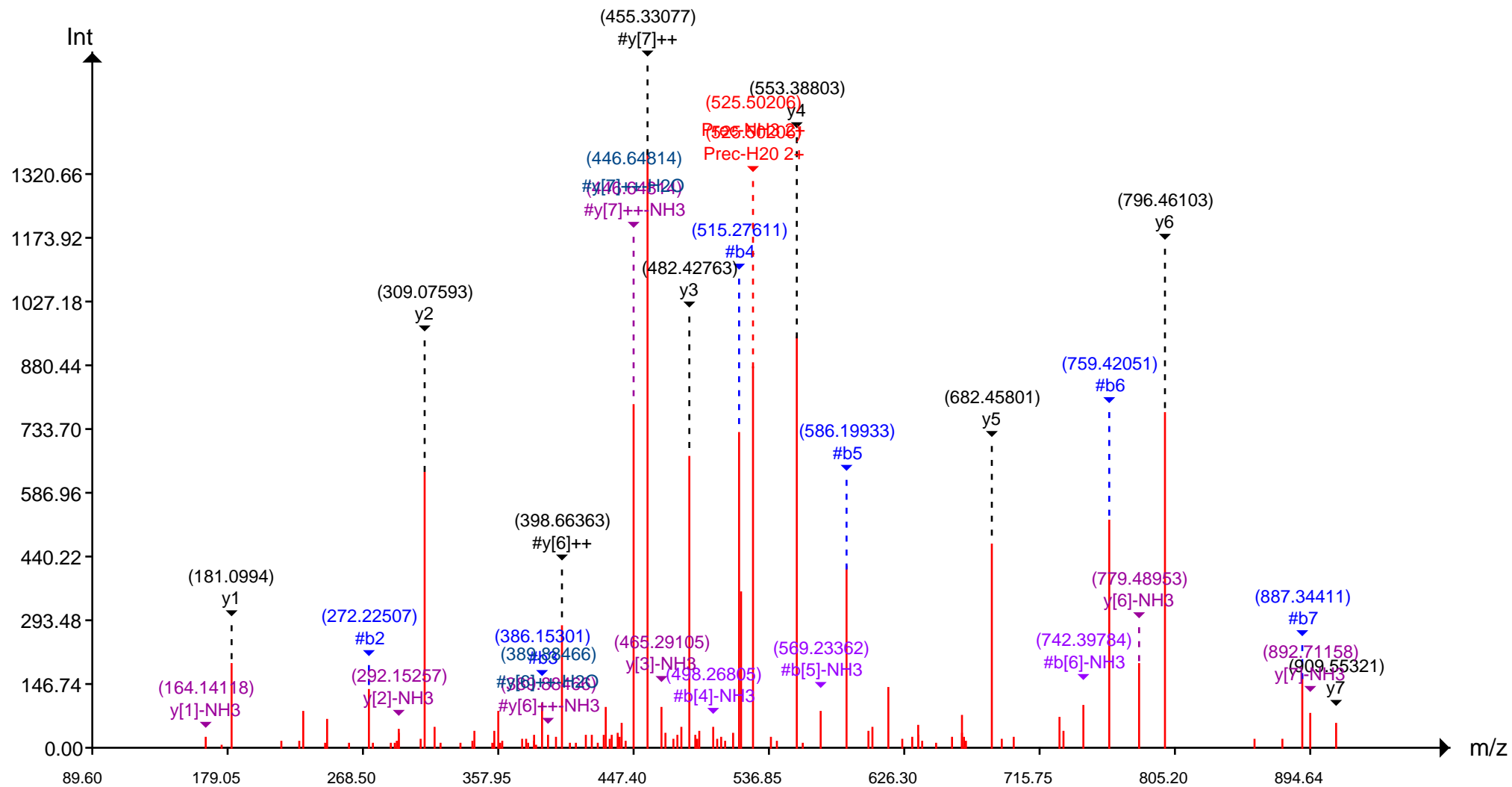
Confidence level 99.00%

Precursor m/z 764.01015

Identity Threshold 31.87

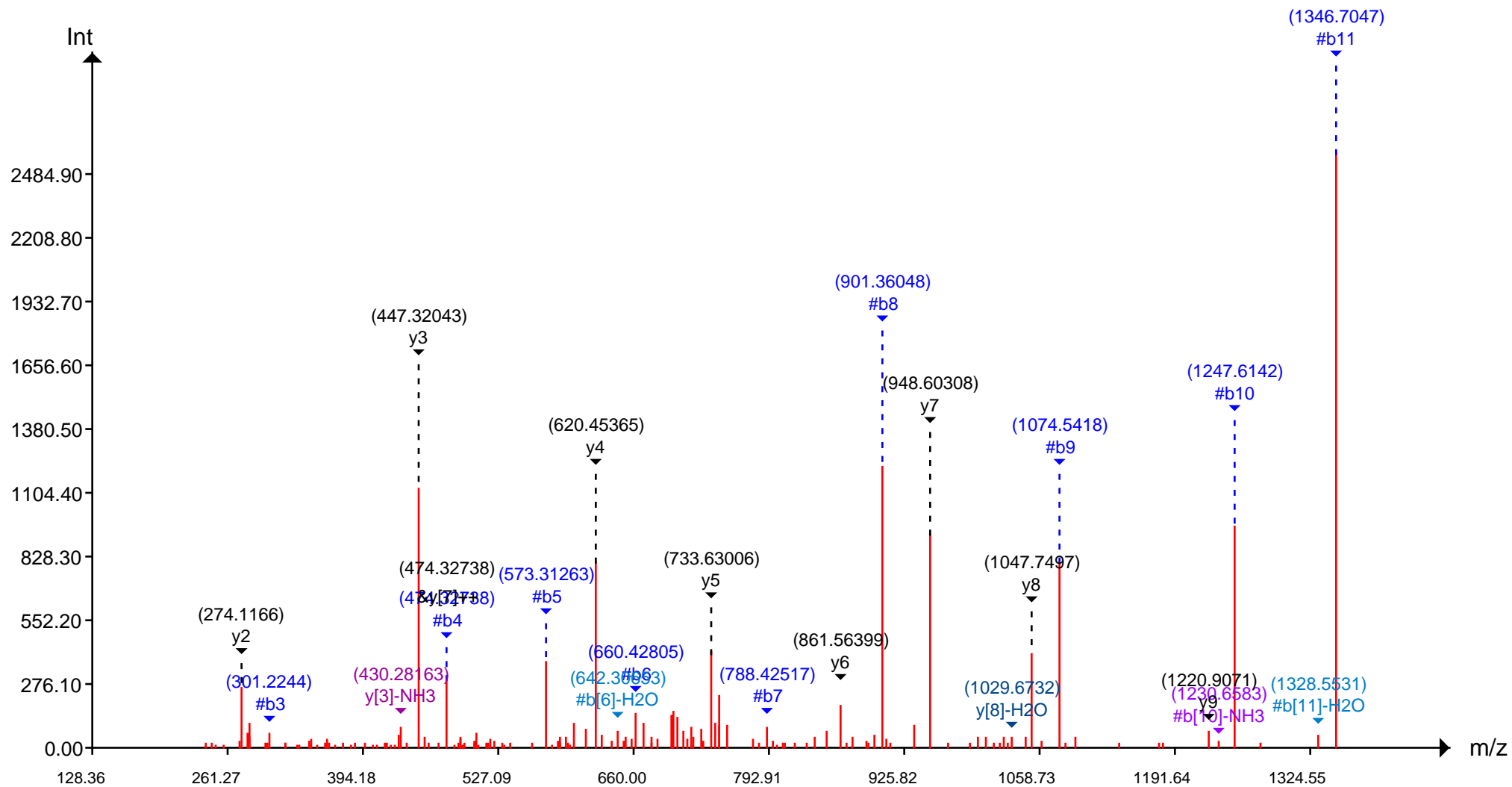
Charge 2+

Mass error 0.001689



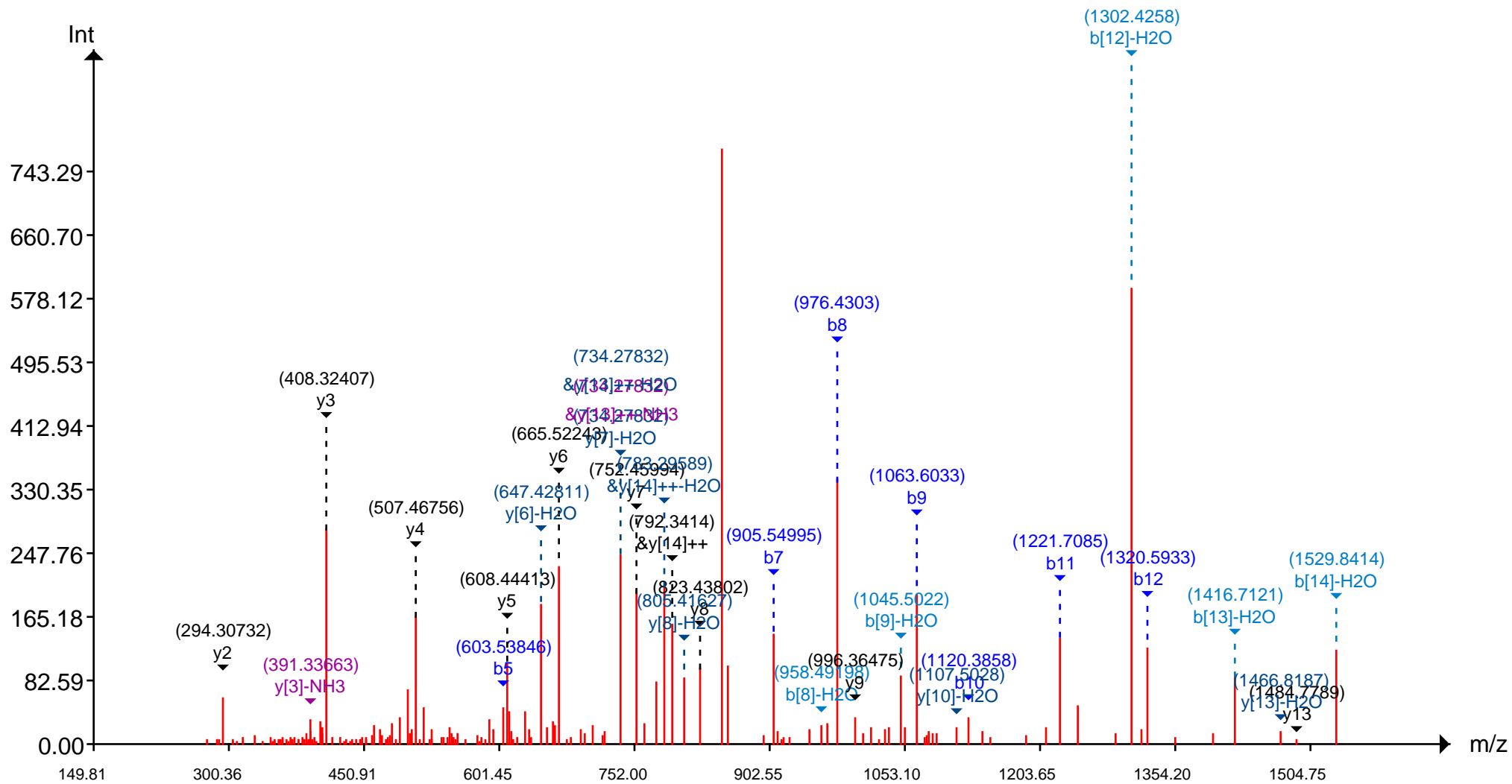
Info

Modified Sequence	AcD3-LLNEAK<AcD3K*>QR<C13>-COOH	Ion score	53.63
Protein	P50518	Confidence level	99.00%
Precursor m/z	534.32469	Identity Threshold	44.61
Charge	2+	Mass error	-4.4E-5



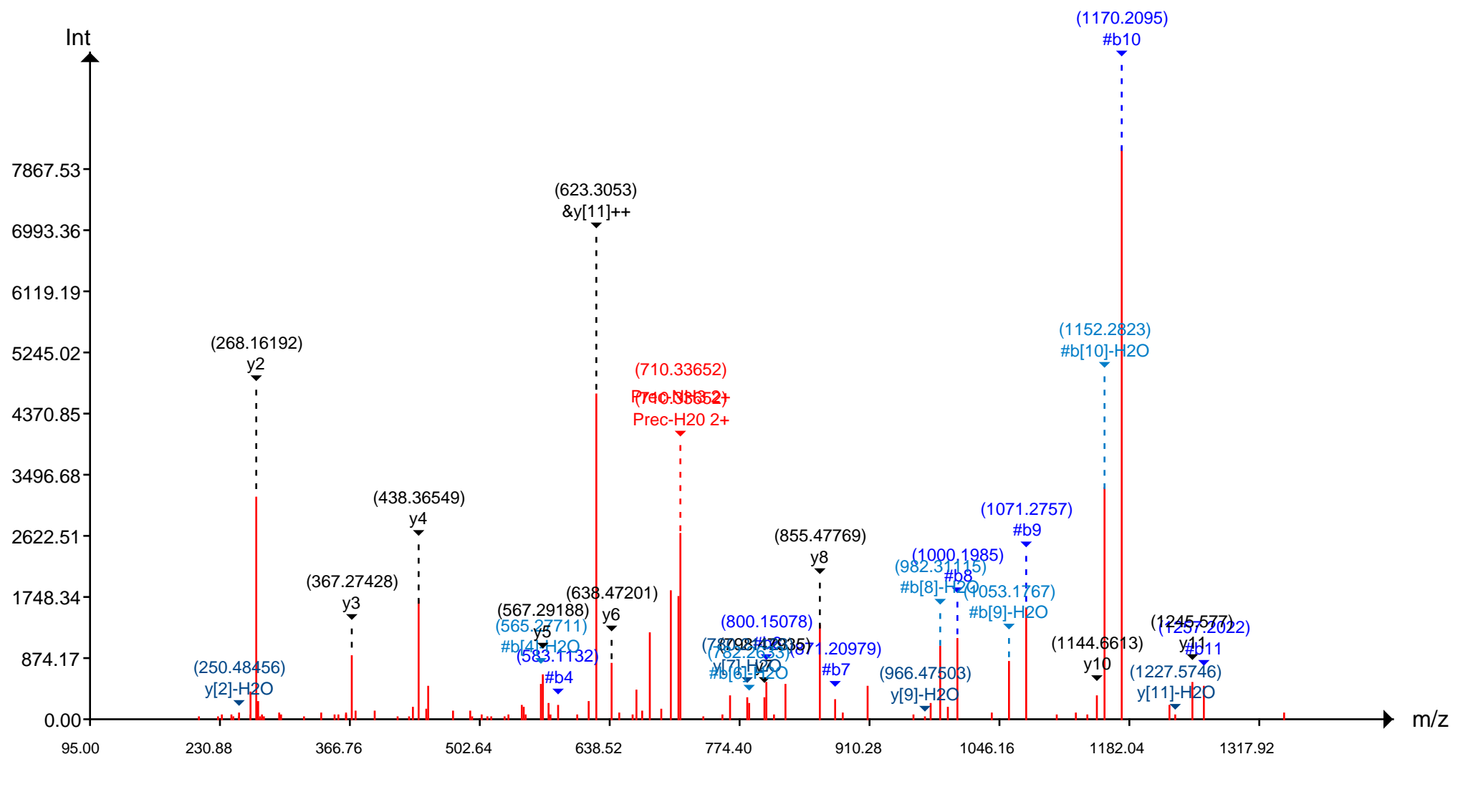
Info

Modified Sequence	AcD3-AALK<AcD3K*>VSQIK<AcD3K*>K<AcD3K*>VR-COOH	Ion score	66.47
Protein	P26638	Confidence level	99.00%
Precursor m/z	761.00017	Identity Threshold	33.87
Charge	2+	Mass error	0.001871

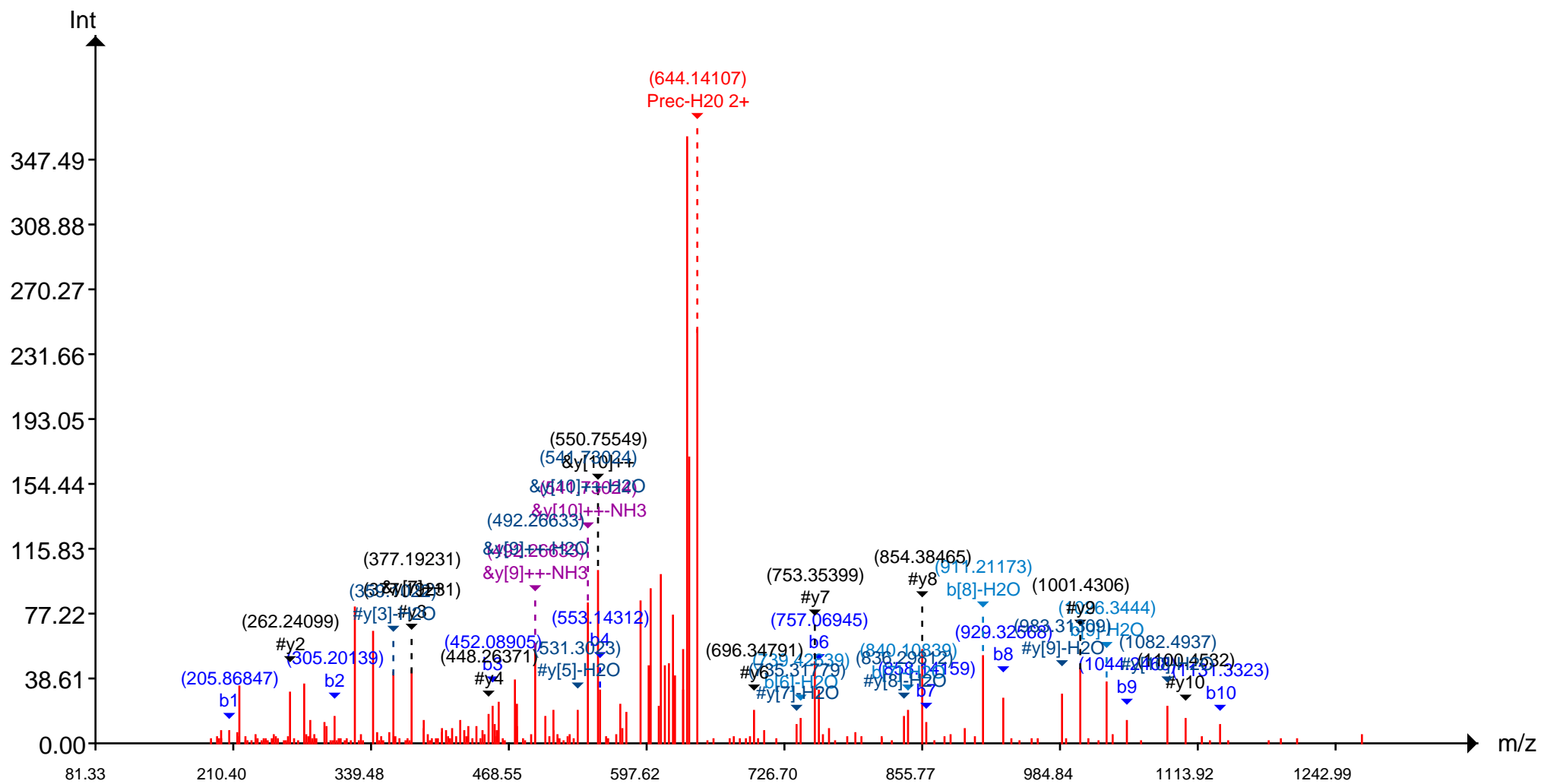


Info

Modified Sequence	AcD3-VVGEK<AcD3K*>EK<AcD3K*>ASGTVNIR<C13>-COOH	Ion score	54.88
Protein	Q9D0R2	Confidence level	99.00%
Precursor m/z	864.50233	Identity Threshold	45.02
Charge	2+	Mass error	0.003017



Info			
Modified Sequence	AcD3-M<Mox*>TC<Cmm*>EGC<Cmm*>AEAVSR<C13>-COOH	Ion score	47.52
Protein	O08997	Confidence level	99.00%
Precursor m/z	719.30043	Identity Threshold	40.00
Charge	2+	Mass error	-5.91E-4



Info

Modified Sequence AcD3-C<Cmm*>VM<Mox*>TGM<Mox*>TADSR-COOH

Ion score 57.52

Protein Q9QUM9

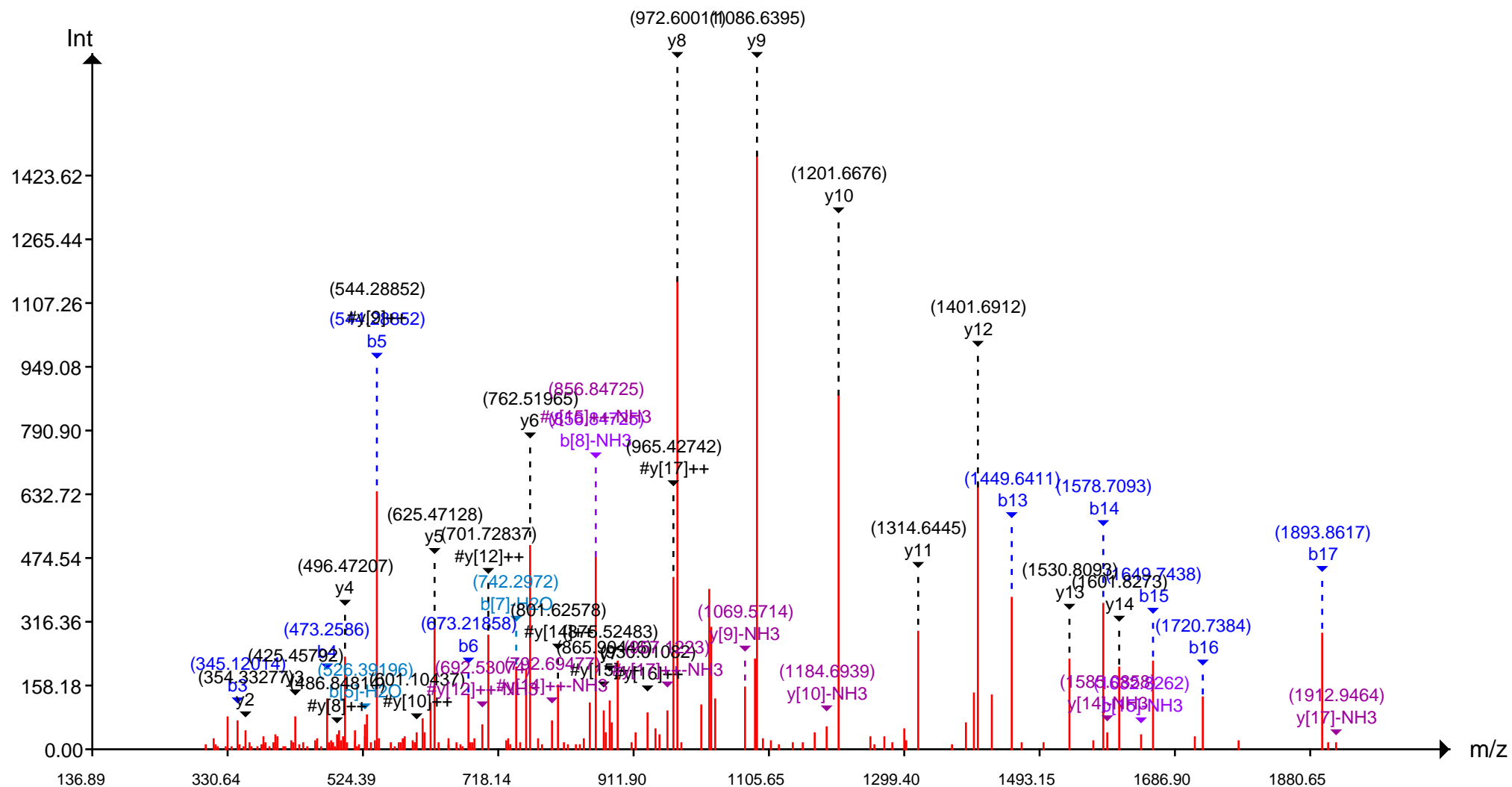
Confidence level 99.00%

Precursor m/z 653.26955

Identity Threshold 38.98

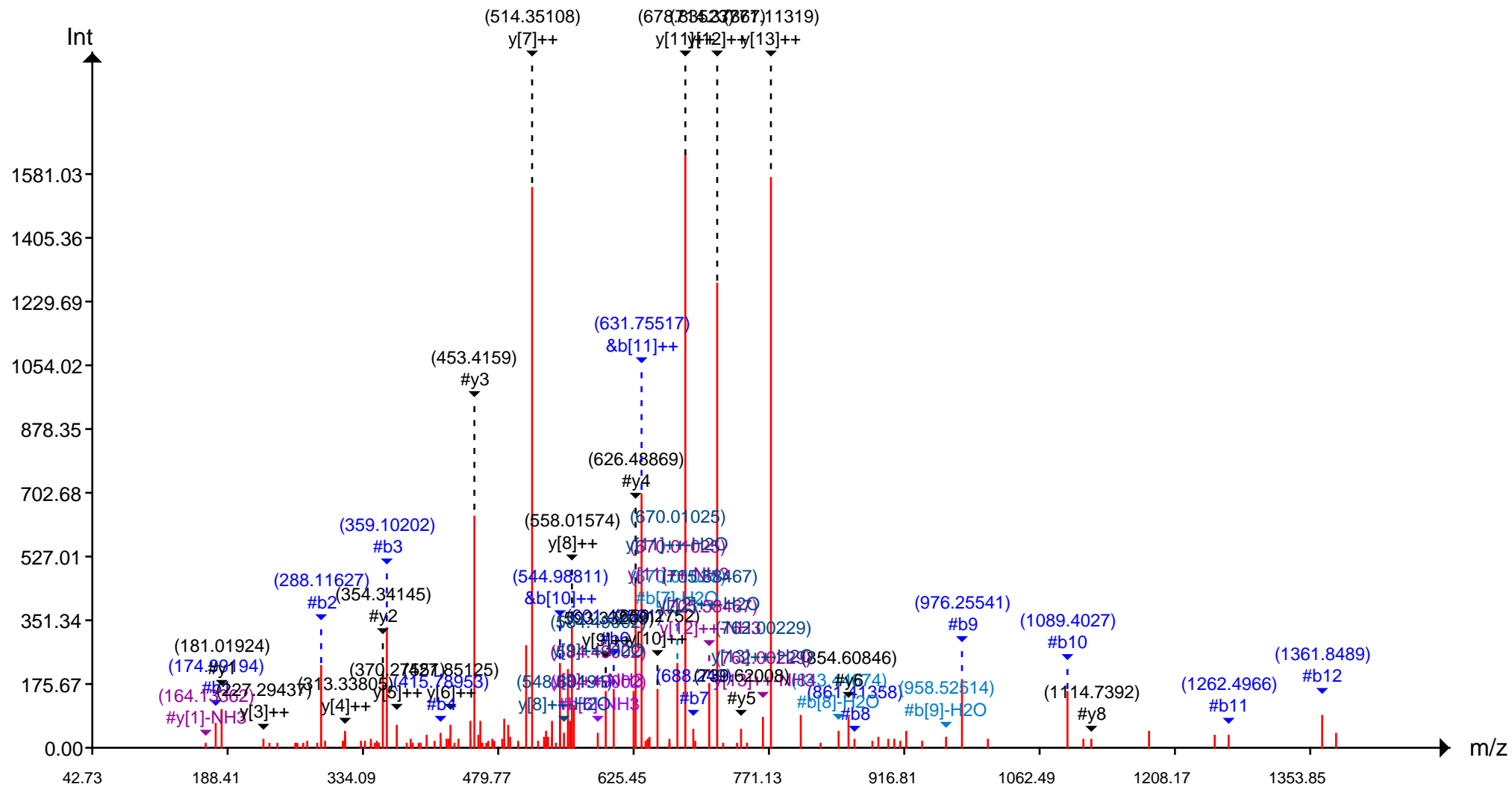
Charge 2+

Mass error 6.99E-4



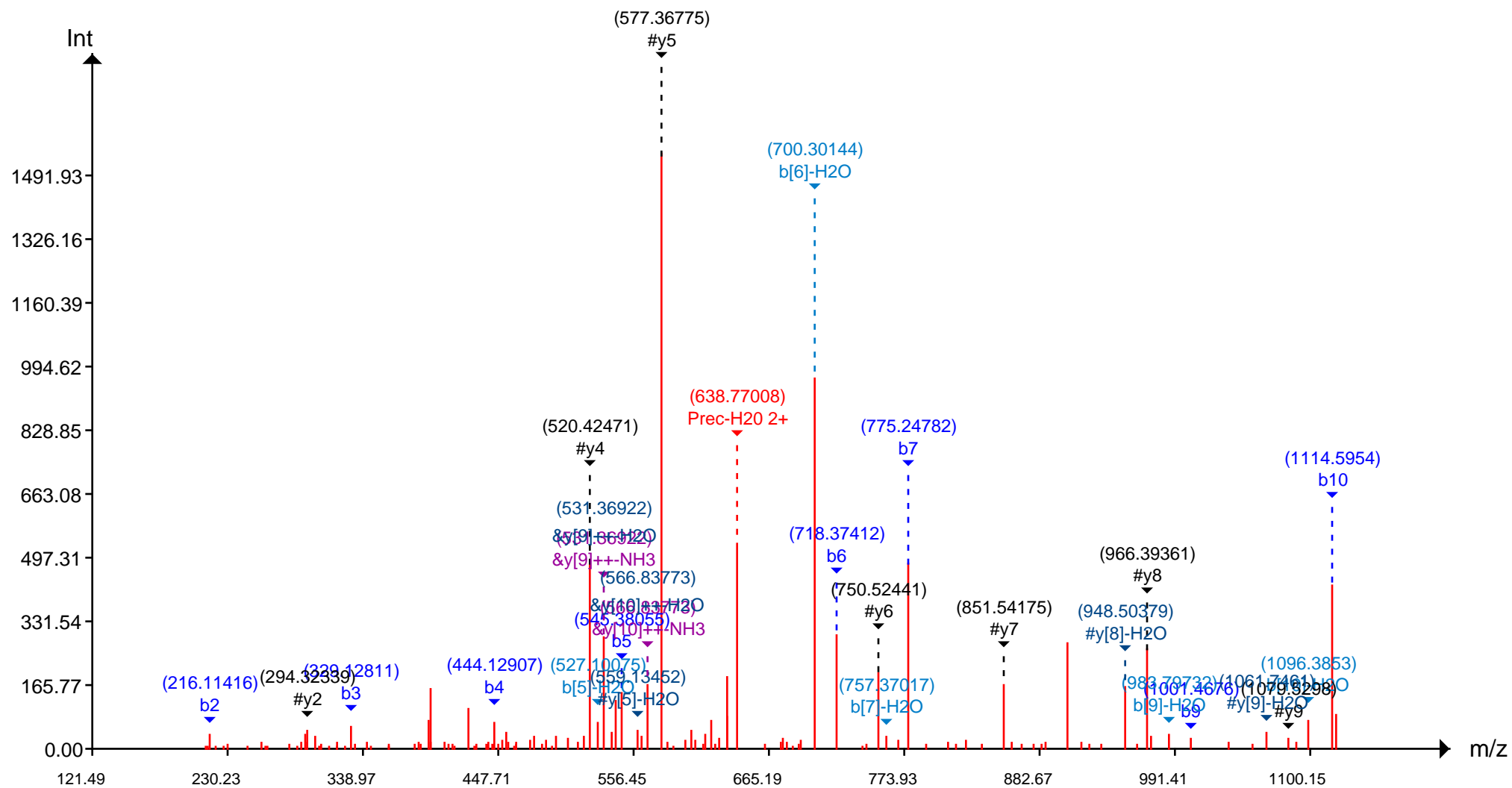
Info

Modified Sequence	AcD3-VAEQAESIDNPLHEAAK<AcD3K*>R<C13>-COOH	Ion score	95.84
Protein	Q62422	Confidence level	99.00%
Precursor m/z	1037.5438	Identity Threshold	48.05
Charge	2+	Mass error	0.002598



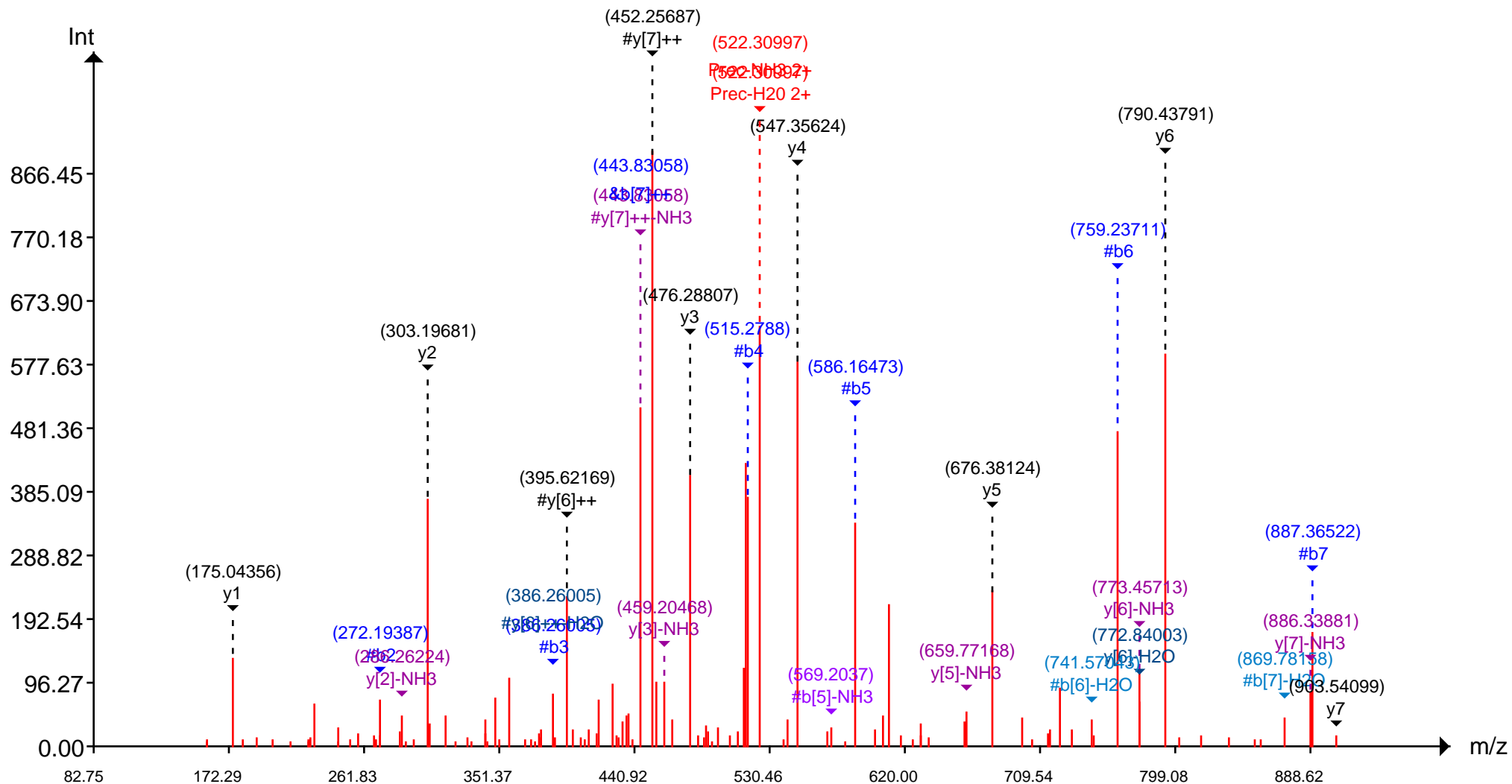
Info

Modified Sequence	AcD3-ELAGNASK<AcD3K*>DLK<AcD3K*>VK<AcD3K*>R<C13>-COOH	Ion score	77.94
Protein	Q3THW5, P0C0S6	Confidence level	99.00%
Precursor m/z	572.34421	Identity Threshold	42.96
Charge	3+	Mass error	-1.84E-4



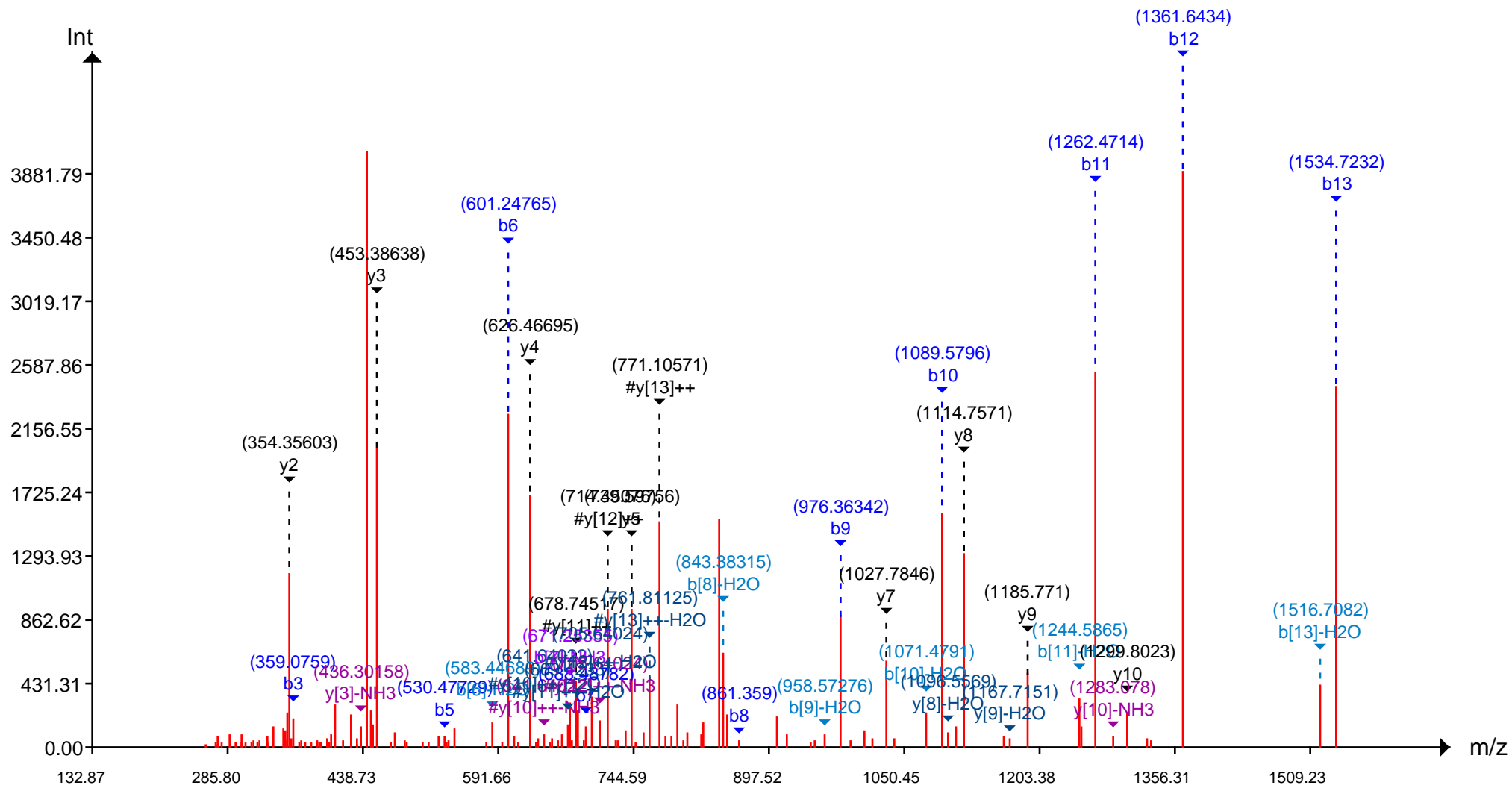
Info

Modified Sequence	AcD3-VALDTK<AcD3K*>GPEIR<C13>-COOH	Ion score	49.91
Protein	P52480	Confidence level	99.00%
Precursor m/z	647.88313	Identity Threshold	45.41
Charge	2+	Mass error	0.001052



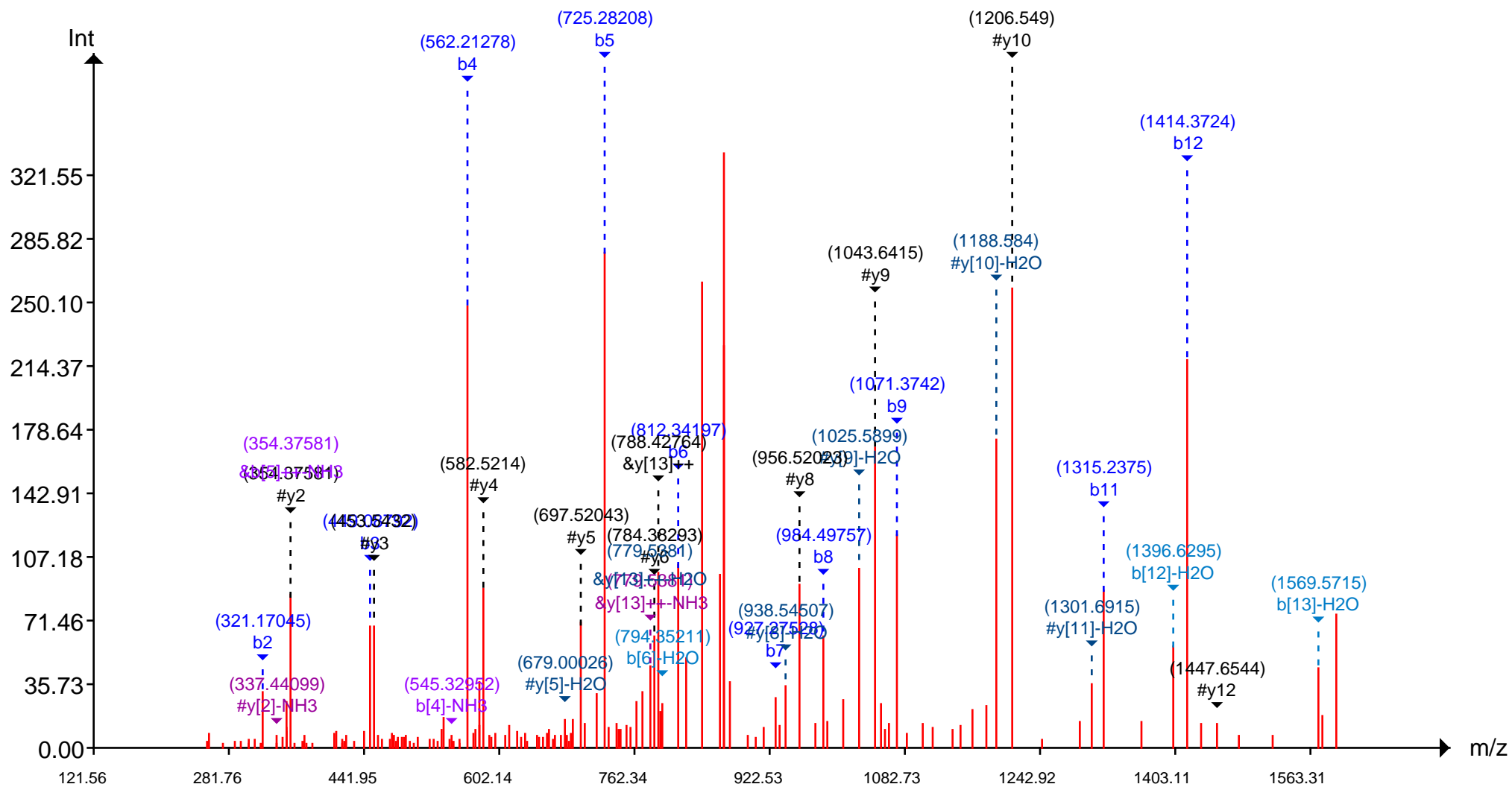
Info

Modified Sequence	AcD3-LLNEAK<AcD3K*>QR-COOH	Ion score	50.05
Protein	P50518	Confidence level	99.00%
Precursor m/z	531.31471	Identity Threshold	45.91
Charge	2+	Mass error	1.37E-4



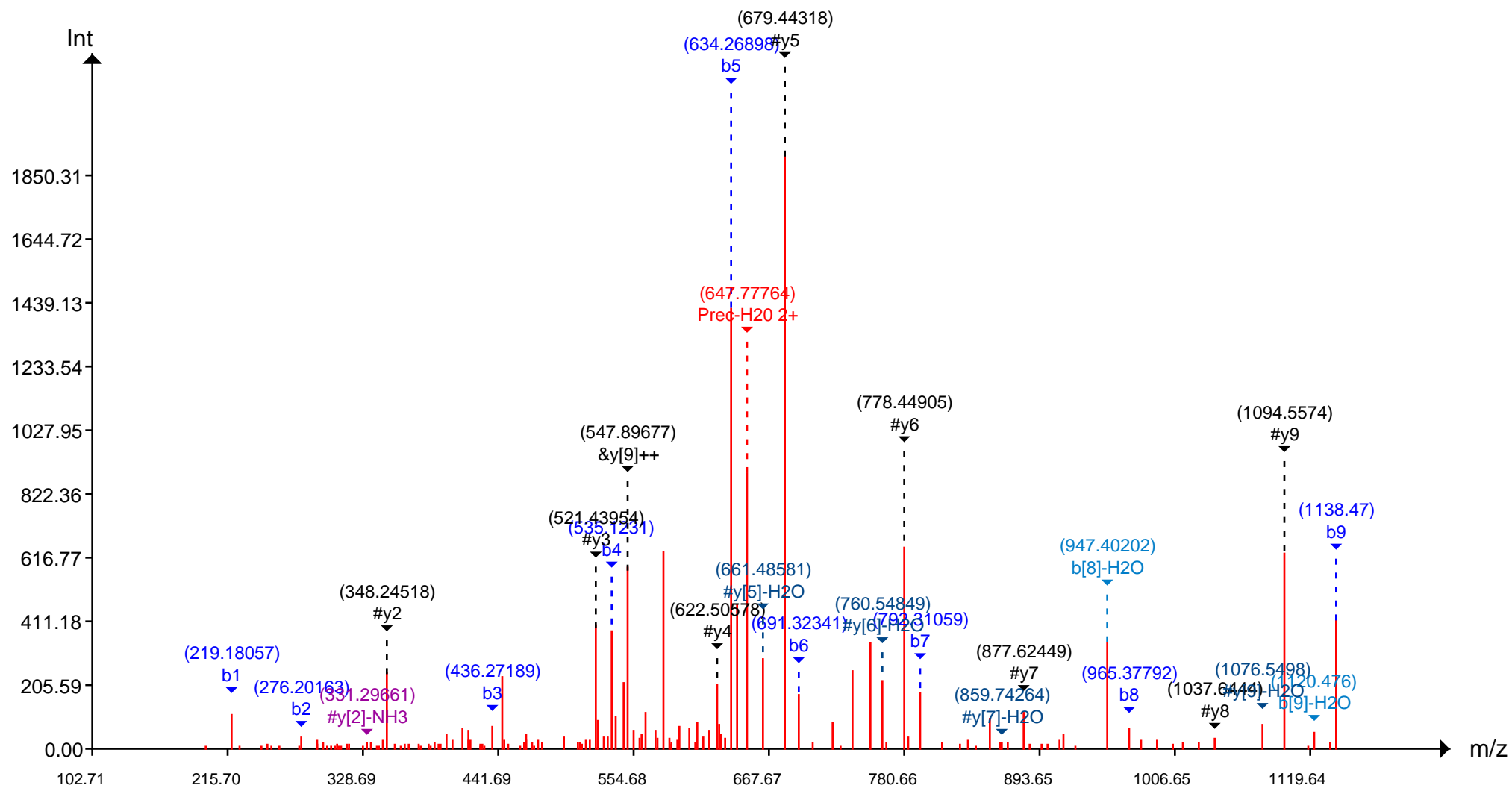
Info

Modified Sequence	AcD3-ELAGNASK<AcD3K*>DLK<AcD3K*>VK<AcD3K*>R<C13>-COOH	Ion score	64.7
Protein	Q3THW5, P0C0S6	Confidence level	99.00%
Precursor m/z	858.01518	Identity Threshold	42.63
Charge	2+	Mass error	0.004822



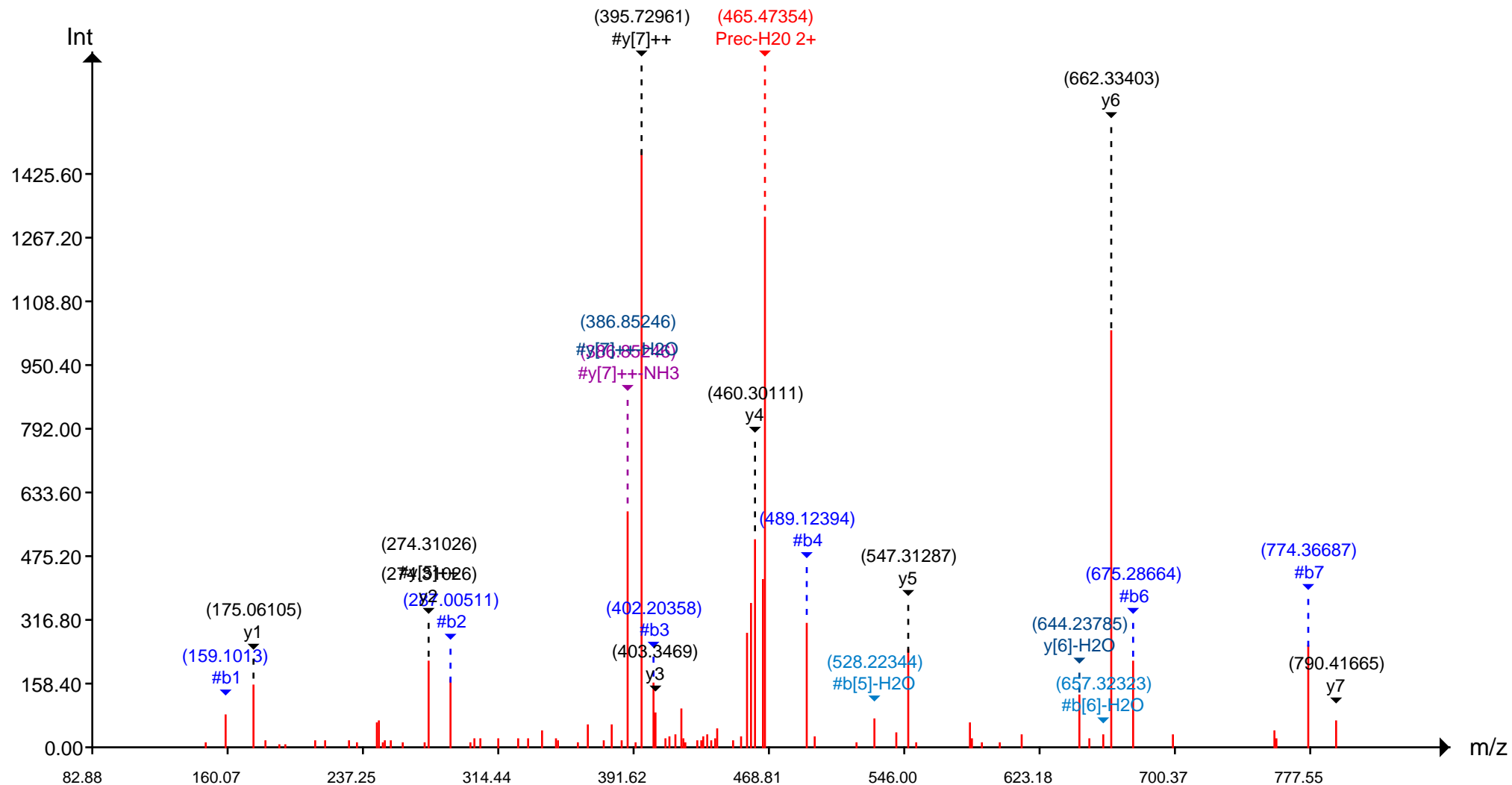
Info

Modified Sequence	AcD3-FQQIYSDGSDEVK<AcD3K*>R<C13>-COOH	Ion score	56.3
Protein	Q9CX34	Confidence level	99.00%
Precursor m/z	884.44298	Identity Threshold	47.70
Charge	2+	Mass error	0.002466



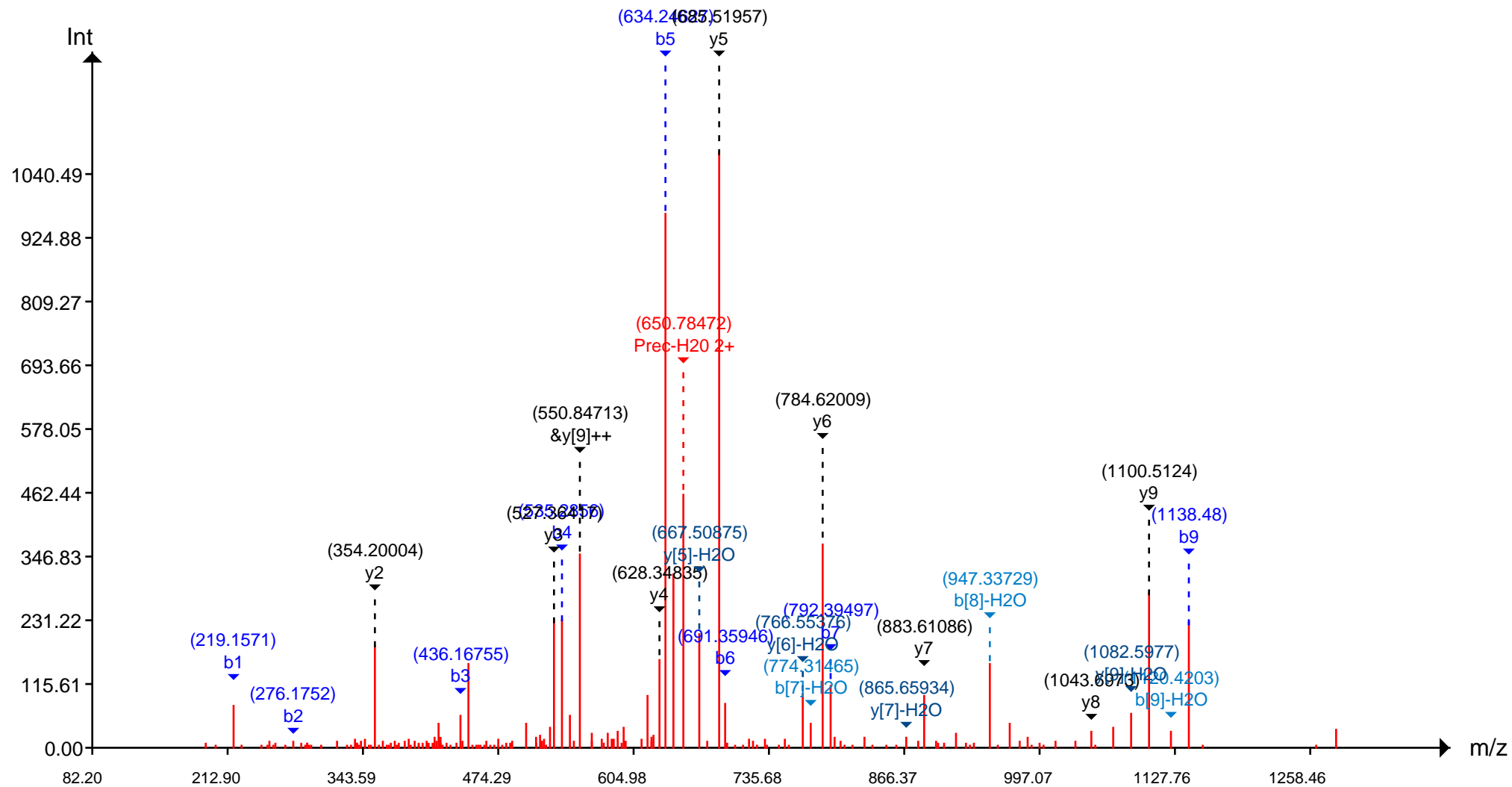
Info

Modified Sequence	AcD3-K<AcD3K*>GC<Cmm*>VVGTK<AcD3K*>K<AcD3K*>R-COOH	Ion score	45.21
Protein	P27659	Confidence level	99.00%
Precursor m/z	656.8936	Identity Threshold	44.49
Charge	2+	Mass error	4.01E-4



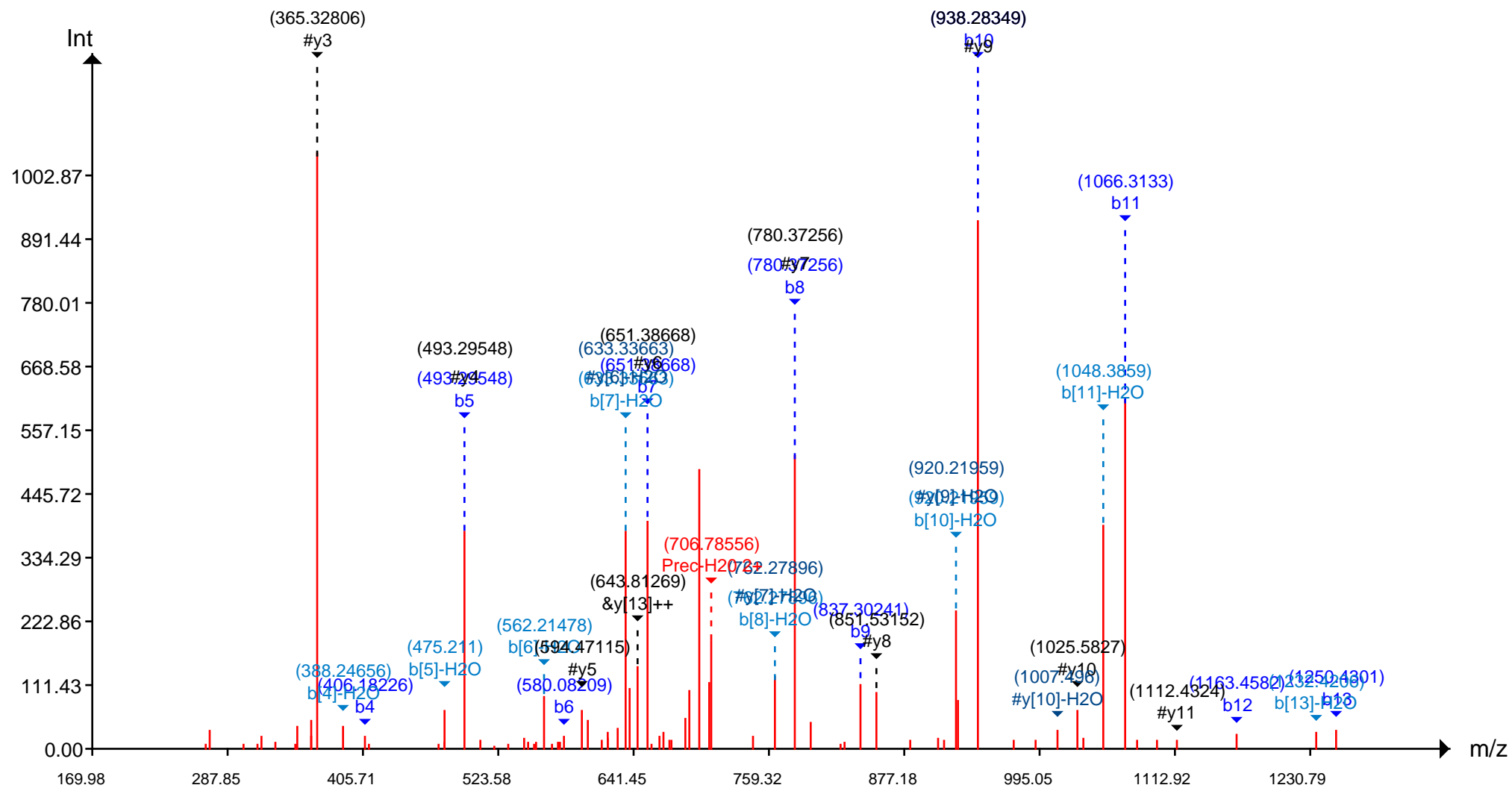
Info

Modified Sequence	AcD3-LQDSGEVR-COOH	Ion score	61.79
Protein	P63242	Confidence level	99.00%
Precursor m/z	474.74484	Identity Threshold	47.54
Charge	2+	Mass error	-1.5E-5



Info

Modified Sequence	AcD3-K<AcD3K*>GC<Cmm*>VVGTK<AcD3K*>K<AcD3K*>R<C13>-COOH	Ion score	57.71
Protein	P27659	Confidence level	99.00%
Precursor m/z	659.90388	Identity Threshold	42.81
Charge	2+	Mass error	8.19E-4



Info

Modified Sequence AcD3-VSSSSSAEGTQPSR<C13>-COOH

Ion score 88.95

Protein Q8R317

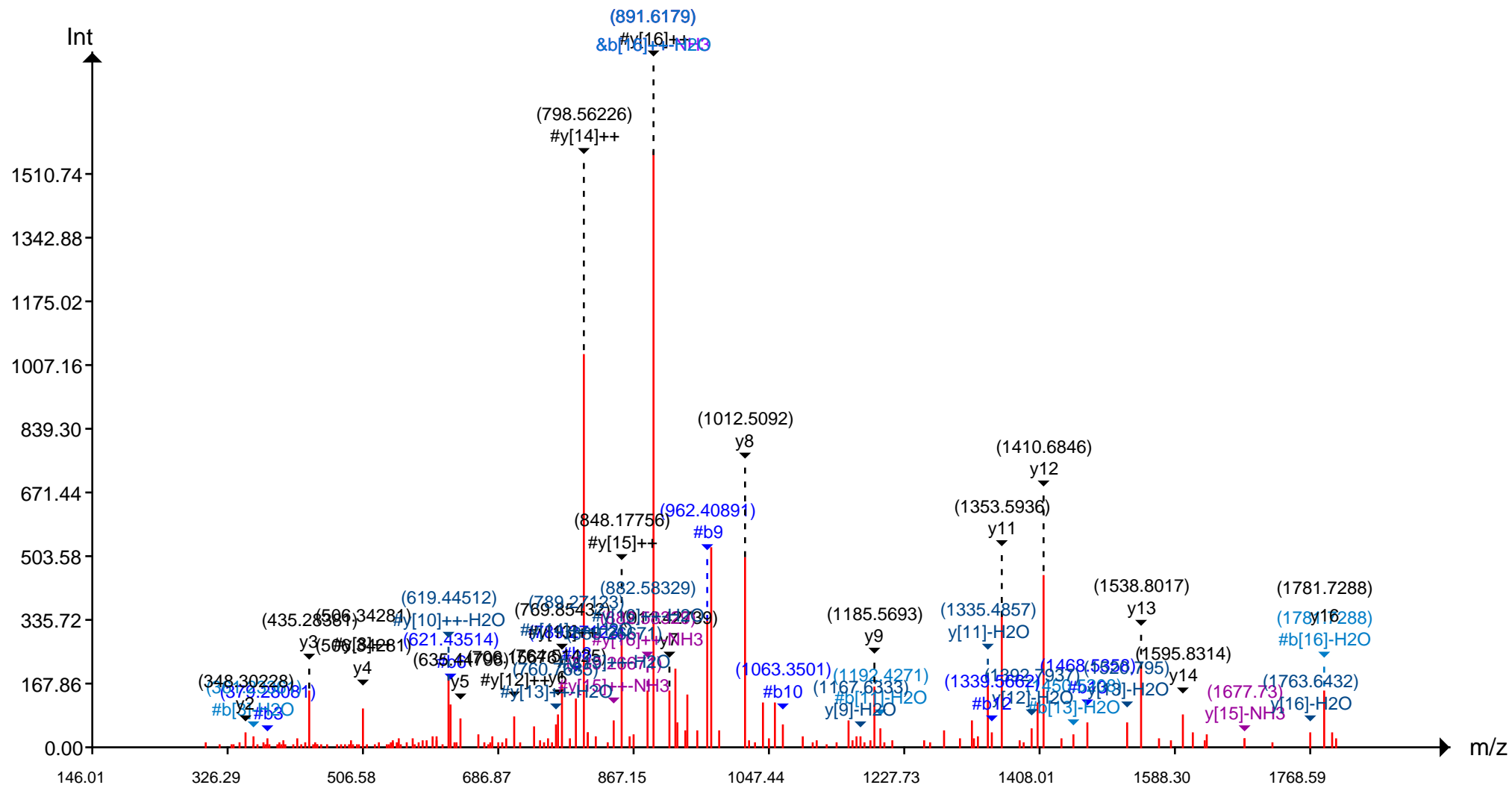
Confidence level 99.00%

Precursor m/z 715.84863

Identity Threshold 46.90

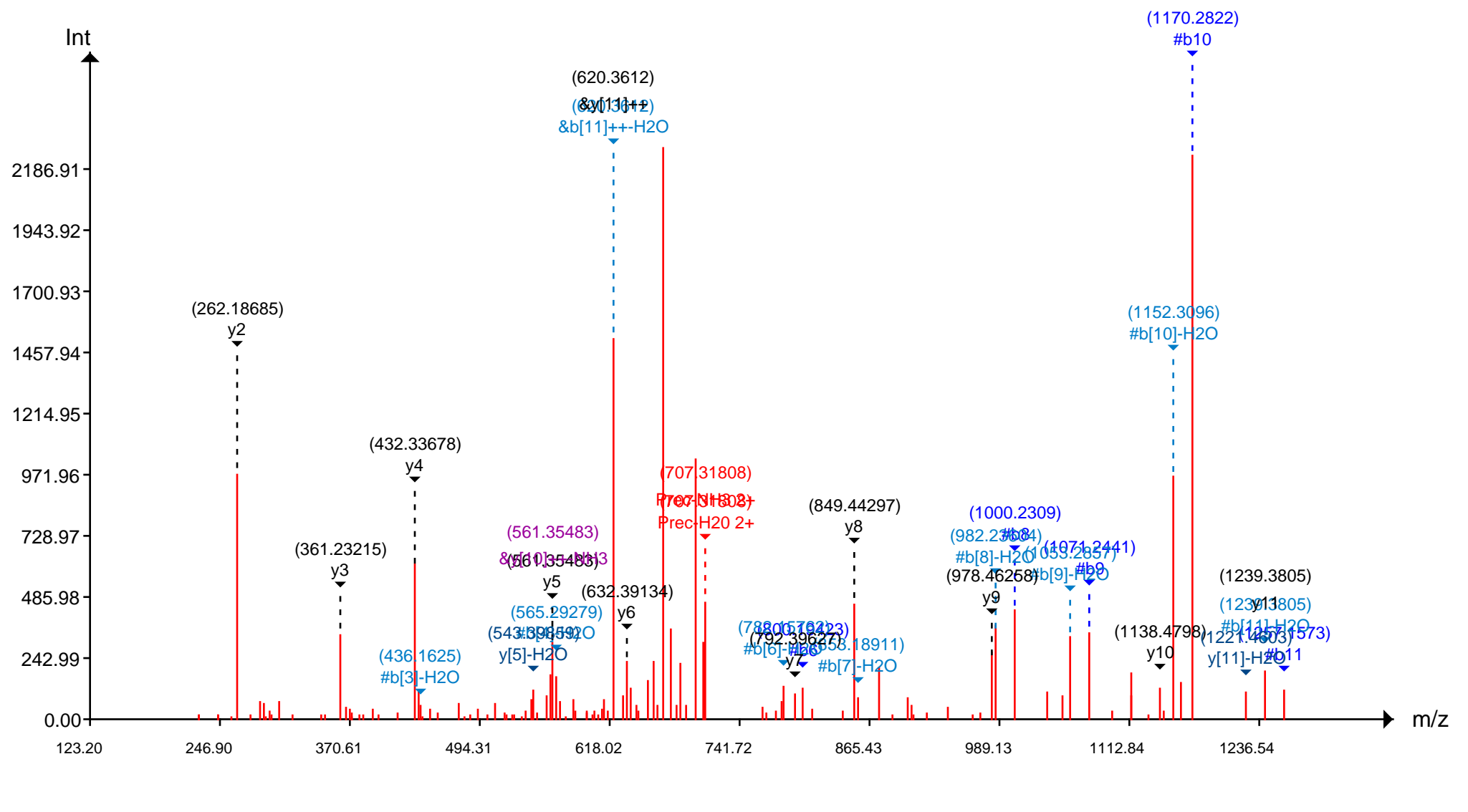
Charge 2+

Mass error 7.32E-4

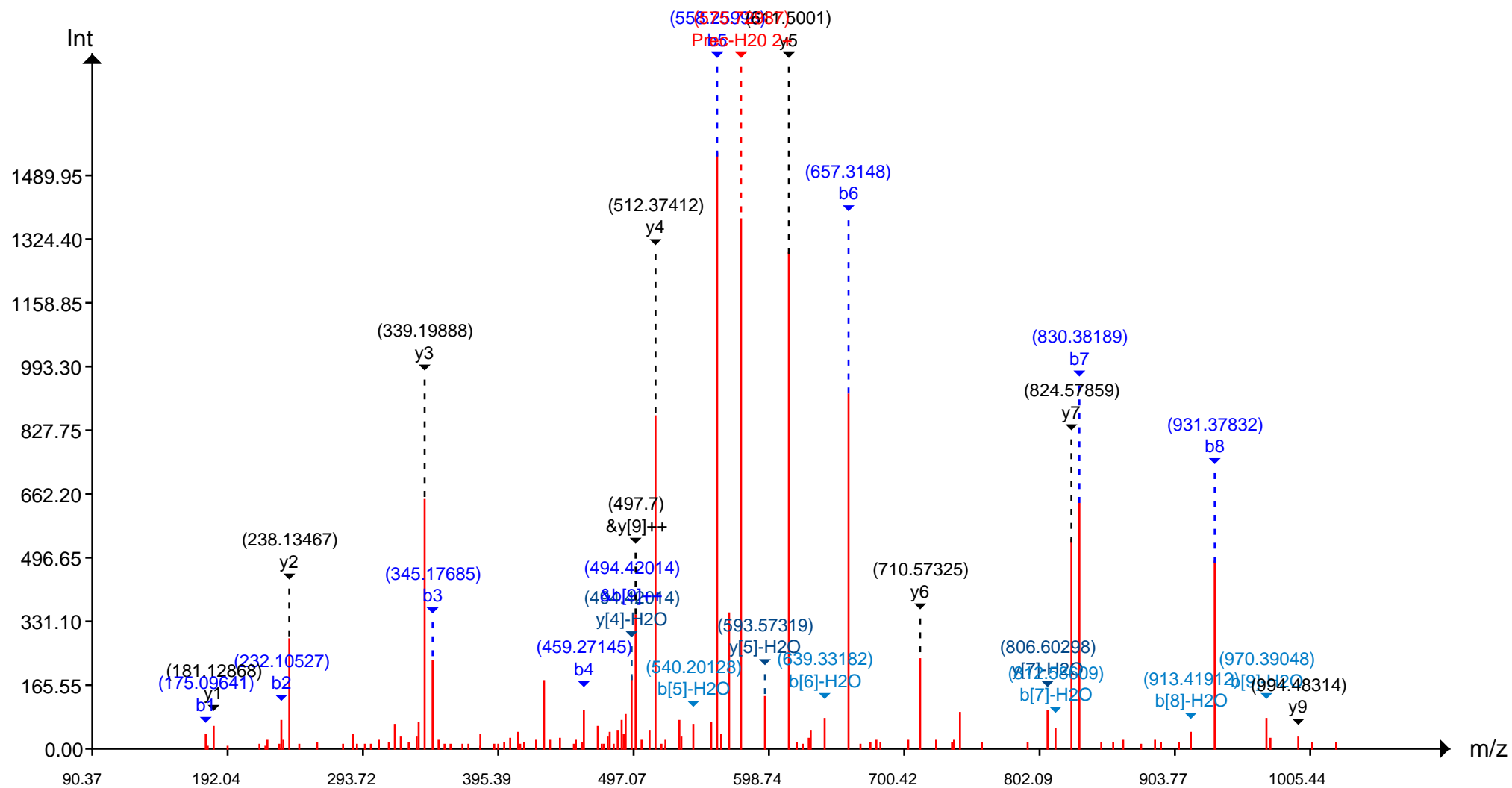


Info

Modified Sequence	AcD3-FSVGQGPAK<AcD3K*>TM<Mox*>EEASK<AcD3K*>R-COOH	Ion score	88.86
Protein	O70310	Confidence level	99.00%
Precursor m/z	987.50214	Identity Threshold	47.92
Charge	2+	Mass error	0.002301

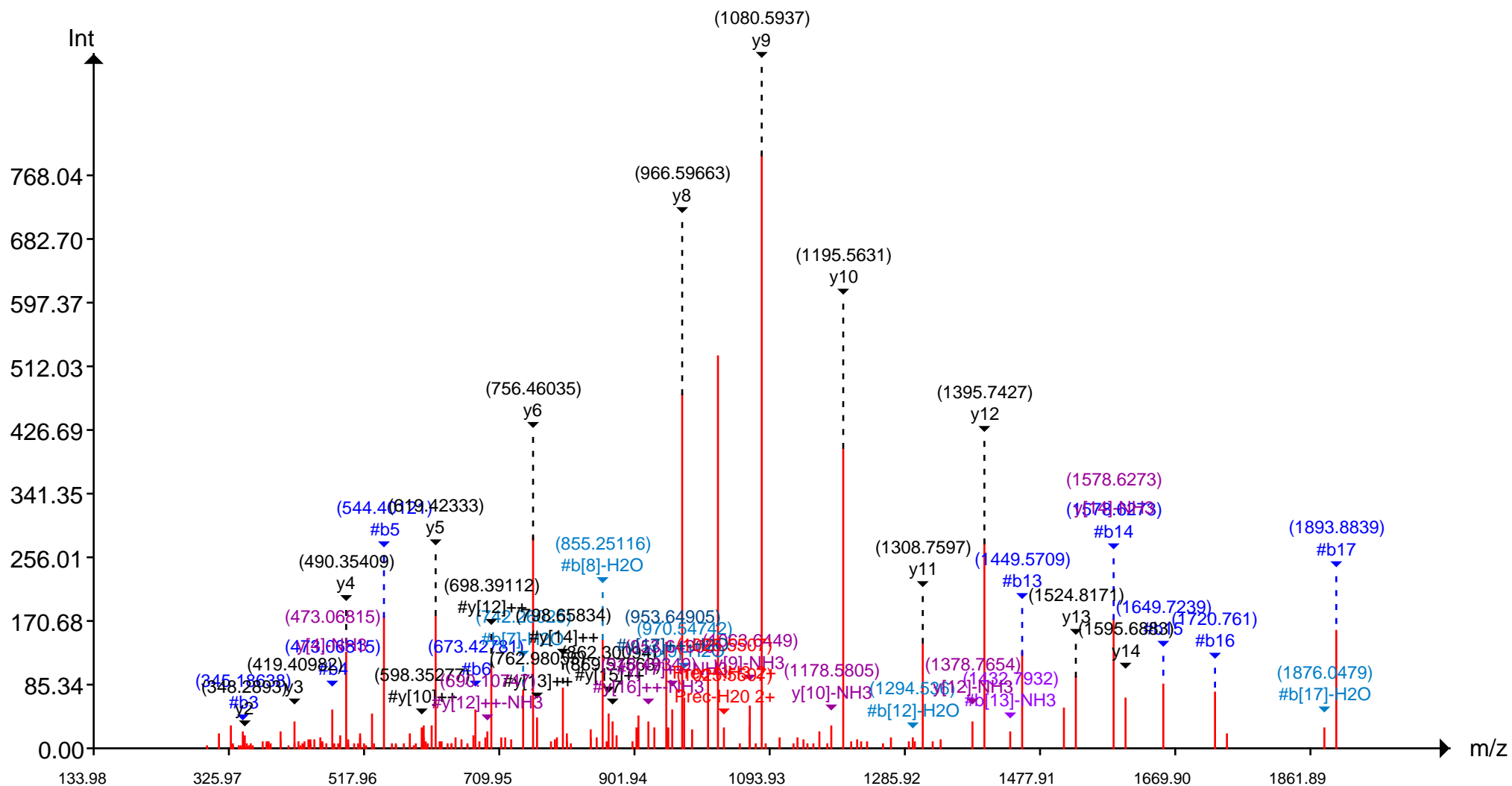


Info			
Modified Sequence	AcD3-M<Mox*>TC<Cmm*>EGC<Cmm*>AEAVSR-COOH	Ion score	57.24
Protein	O08997	Confidence level	99.00%
Precursor m/z	716.29096	Identity Threshold	38.08
Charge	2+	Mass error	6.11E-4



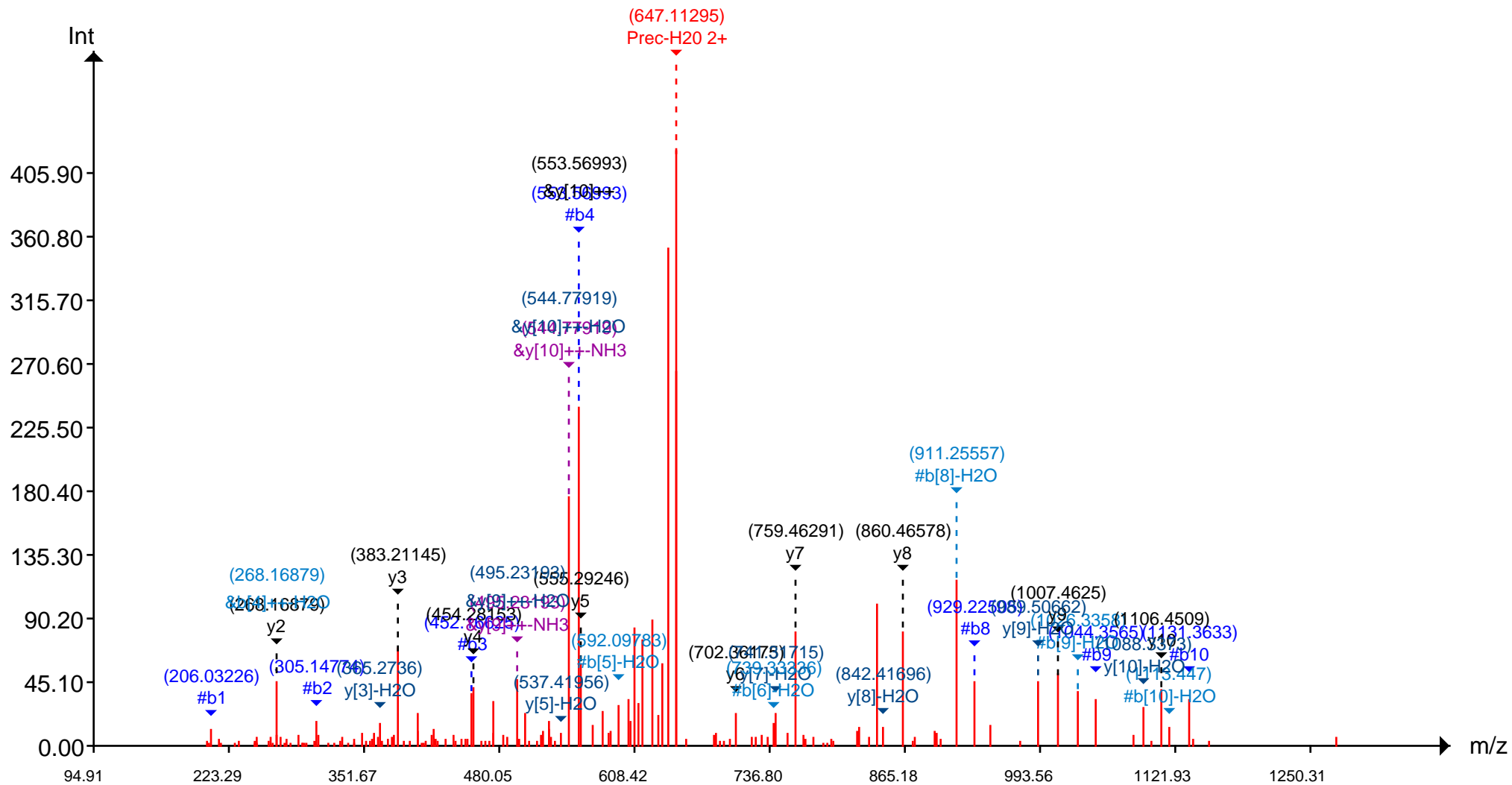
Info

Modified Sequence	AcD3-EGLNVVK<AcD3K*>TGR<C13>-COOH	Ion score	67.22
Protein	P15532, Q01768	Confidence level	99.00%
Precursor m/z	584.8492	Identity Threshold	45.23
Charge	2+	Mass error	0.001261



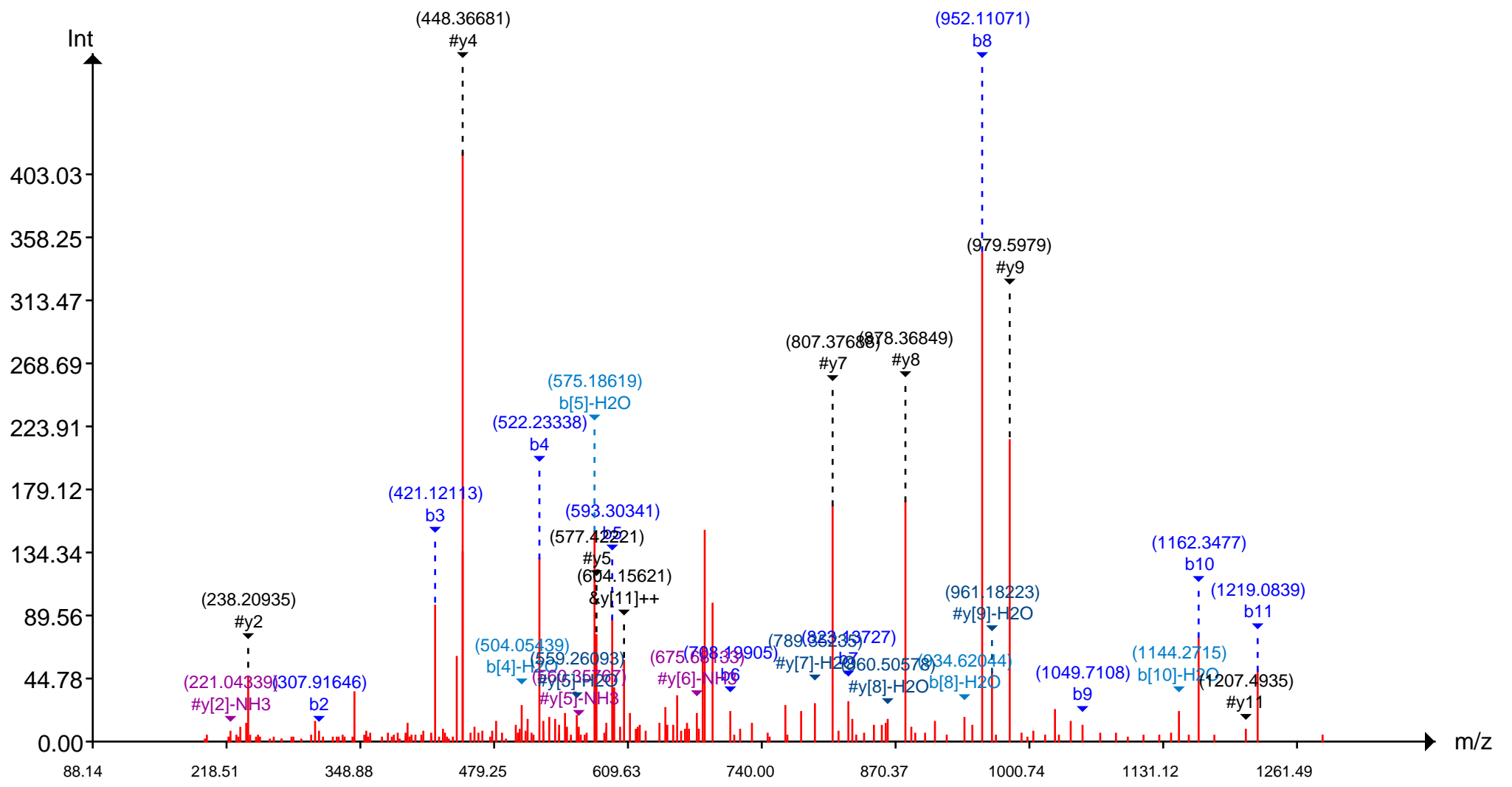
Info

Modified Sequence	AcD3-VAEQAESIDNPLHEAAK<AcD3K*>R-COOH	Ion score	99.32
Protein	Q62422	Confidence level	99.00%
Precursor m/z	1034.5337	Identity Threshold	47.96
Charge	2+	Mass error	0.00254



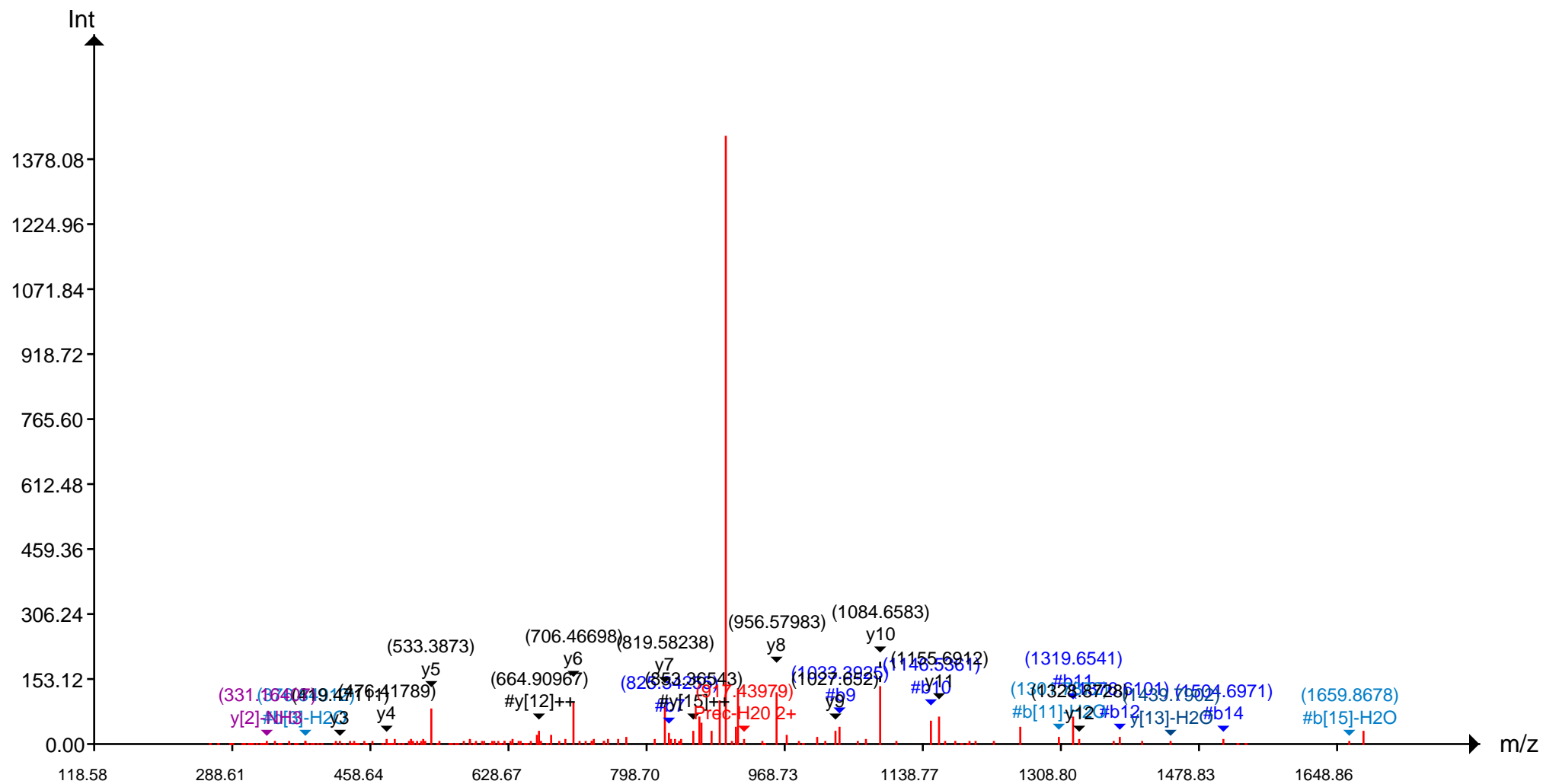
Info

Modified Sequence	AcD3-C<Cmm*>VM<Mox*>TGM<Mox*>TADSR<C13>-COOH	Ion score	81.14
Protein	Q9QUM9	Confidence level	99.00%
Precursor m/z	656.28003	Identity Threshold	41.20
Charge	2+	Mass error	0.001517



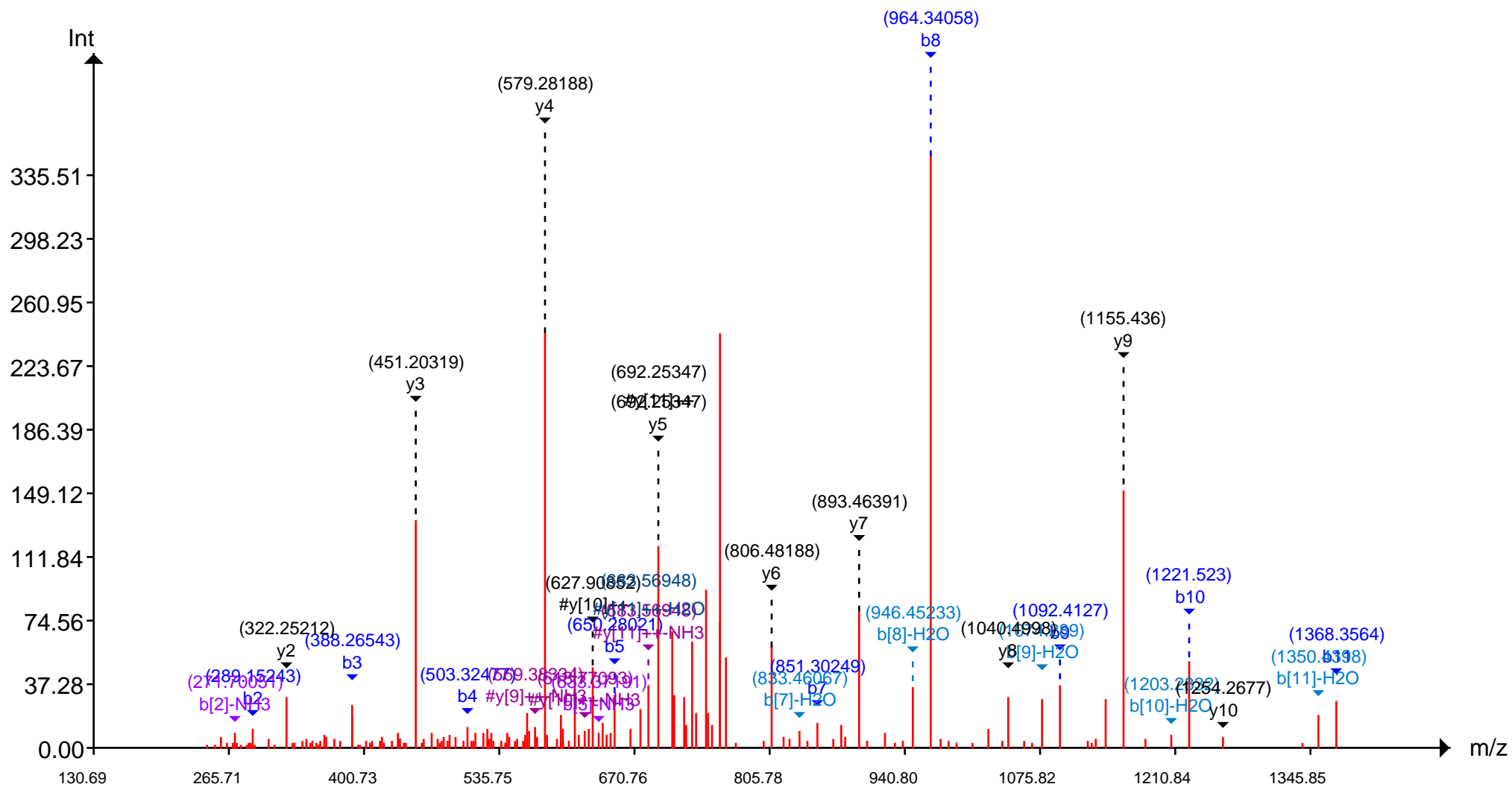
Info

Modified Sequence	AcD3-FDITADDEPLGR<C13>-COOH	Ion score	70.8
Protein	P17742	Confidence level	99.00%
Precursor m/z	700.34778	Identity Threshold	47.39
Charge	2+	Mass error	8.17E-4



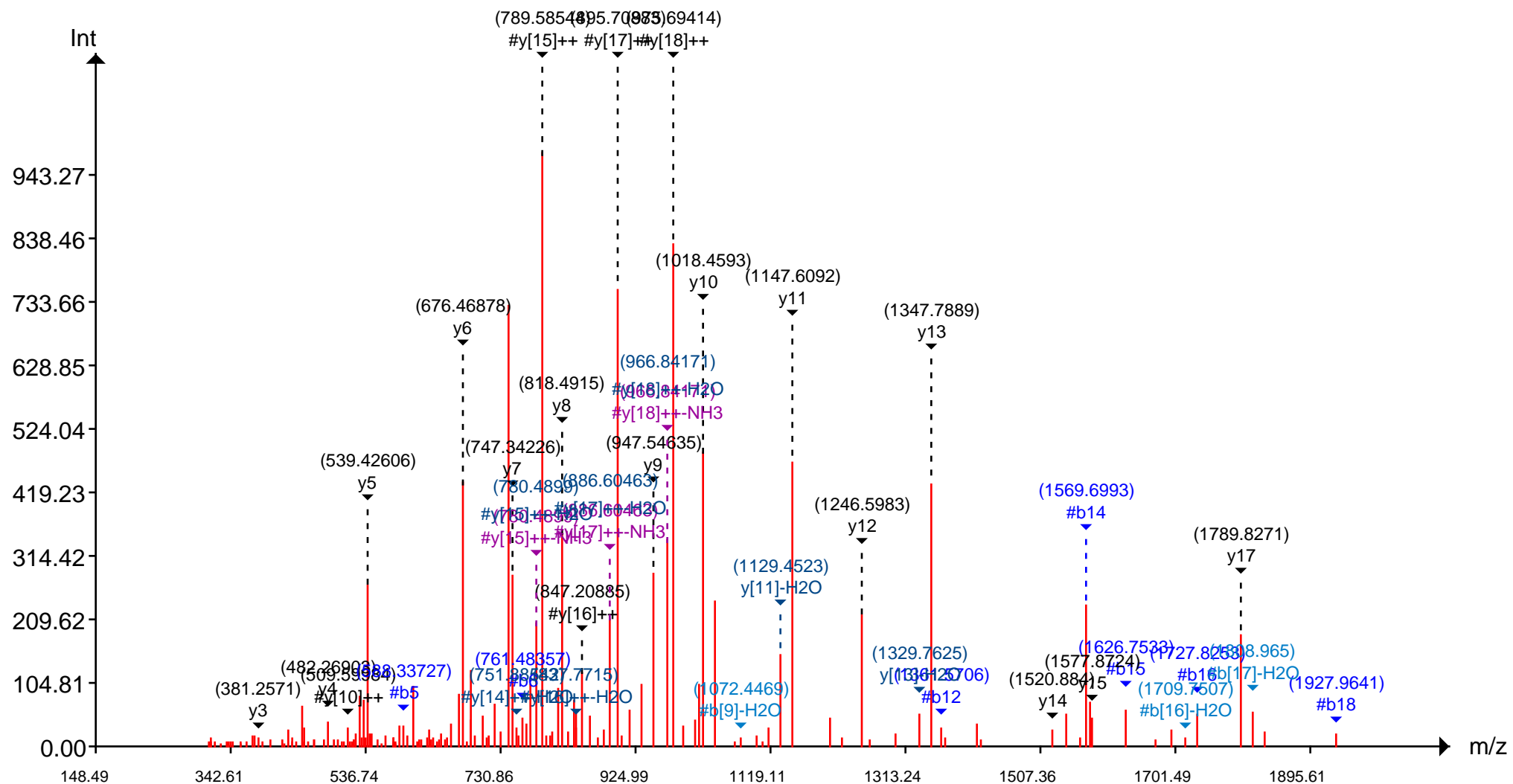
Info

Modified Sequence	AcD3-TTM<Mox*>EK<AcD3K*>AG AHLK<AcD3K*>GGAK<AcD3K*>R-COOH	Ion score	65.57
Protein	P16858	Confidence level	99.00%
Precursor m/z	926.51113	Identity Threshold	47.20
Charge	2+	Mass error	0.001513



Info

Modified Sequence	AcD3-NEVDM<Mox*>SNIQEM<Mox*>R-COOH	Ion score	79.36
Protein	Q9D2G2	Confidence level	99.00%
Precursor m/z	771.83454	Identity Threshold	41.85
Charge	2+	Mass error	0.00162



Info

Modified Sequence AcD3-IC<Cmm*>PDGK<AcD3K*>TVEAEAAHGTVTR<C13>-COOH

Ion score 83.44

Protein O88844

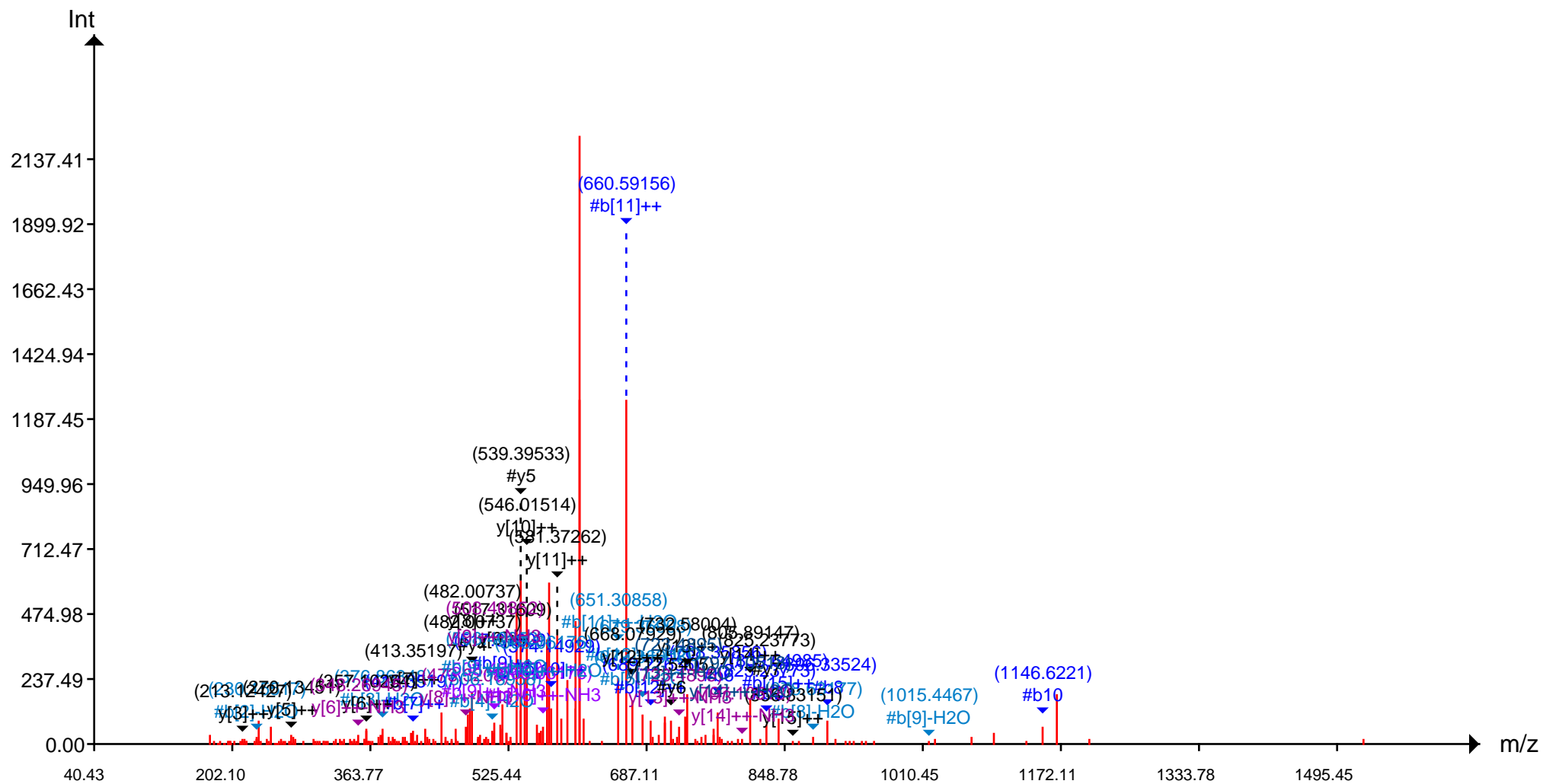
Confidence level 99.00%

Precursor m/z 1054.5374

Identity Threshold 47.91

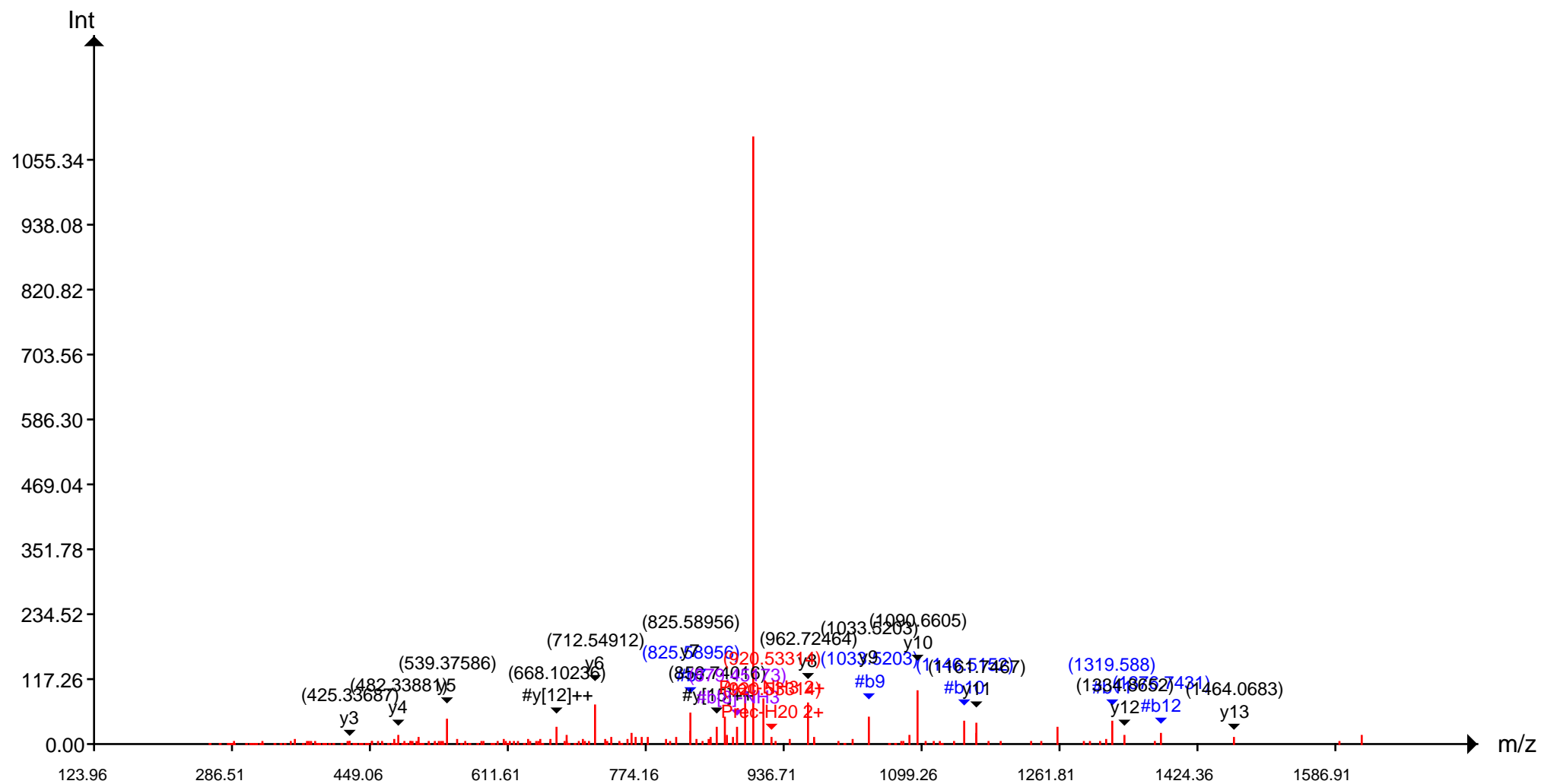
Charge 2+

Mass error 0.00202



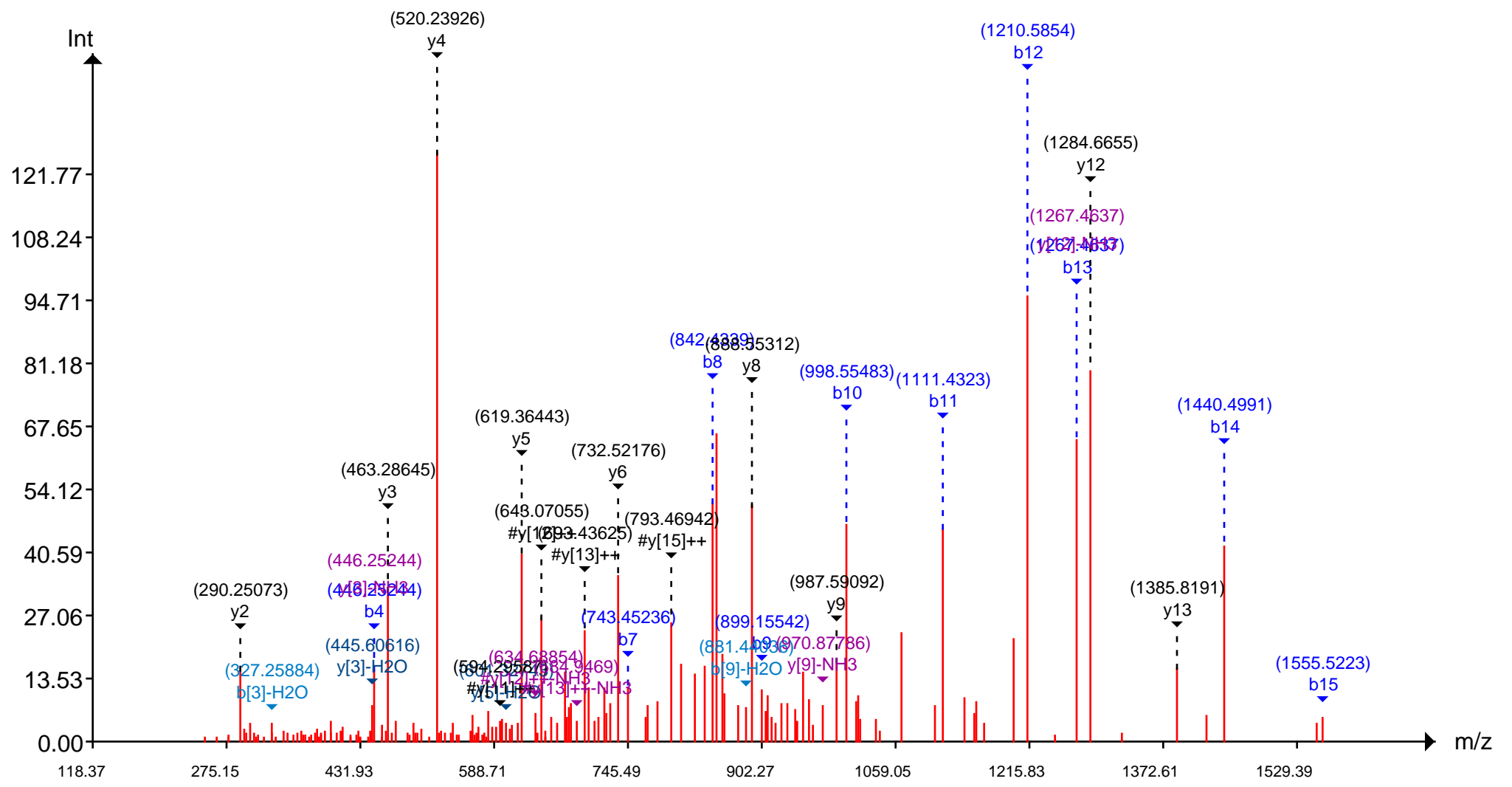
Info

Modified Sequence	AcD3-TTM<Mox*>EK<AcD3K*>AG AHLK<AcD3K*>GGAK<AcD3K*>R<C13>-COOH	Ion score	58.3
Protein	P16858	Confidence level	99.00%
Precursor m/z	620.01603	Identity Threshold	46.56
Charge	3+	Mass error	-7.5E-5

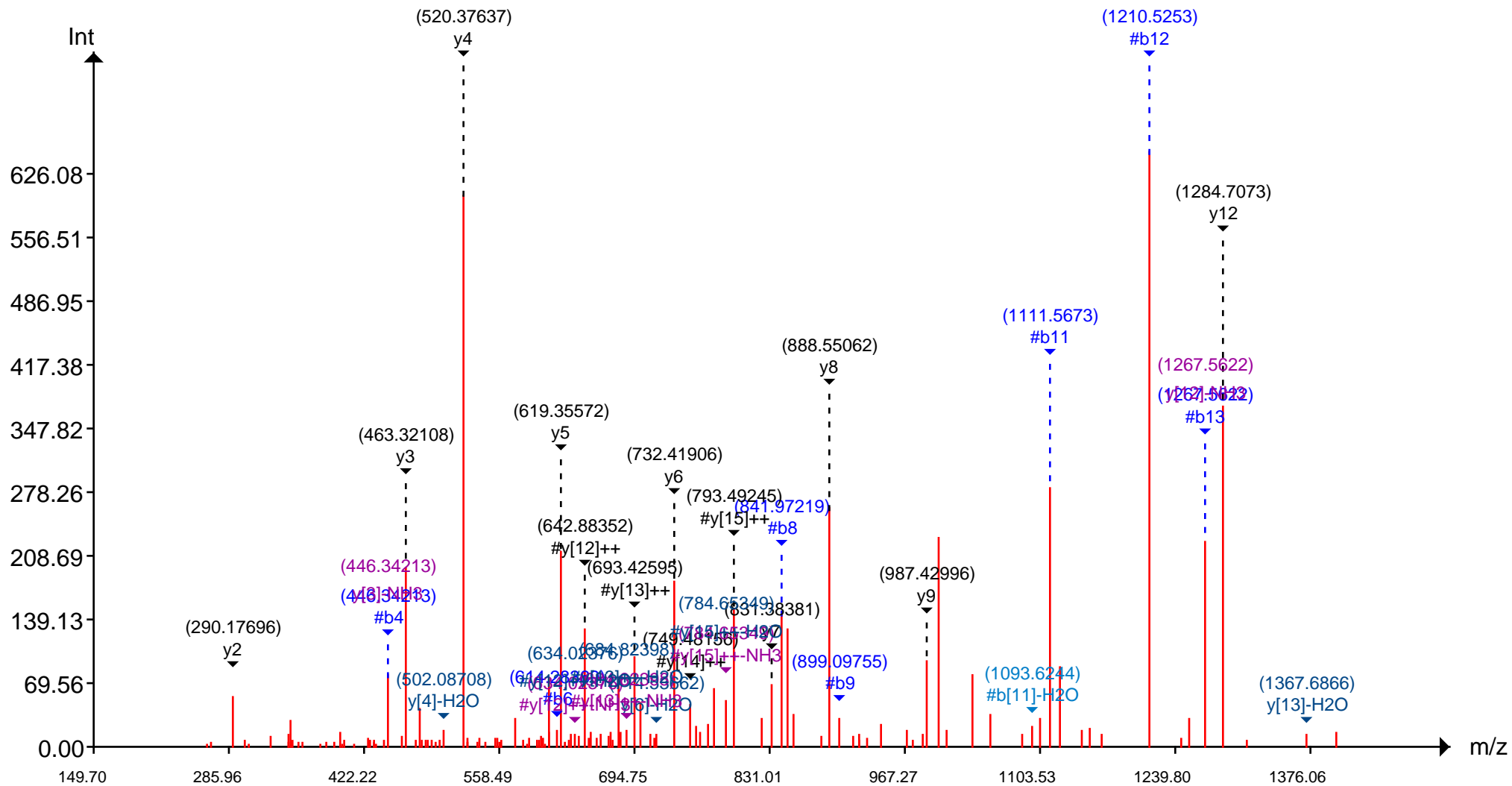


Info

Modified Sequence	AcD3-TTM<Mox*>EK<AcD3K*>AGHLK<AcD3K*>GGAK<AcD3K*>R<C13>-COOH	Ion score	83.71
Protein	P16858	Confidence level	99.00%
Precursor m/z	929.52181	Identity Threshold	46.47
Charge	2+	Mass error	0.002731



Info			
Modified Sequence	AcD3-VSITPAEVGVLVGK<AcD3K*>DR-COOH	Ion score	81.74
Protein	P62962	Confidence level	99.00%
Precursor m/z	865.50309	Identity Threshold	44.99
Charge	2+	Mass error	0.002355



Info

Modified Sequence AcD3-VSITPAEVLVVGK<AcD3K*>DR-COOH

Ion score 60.77

Protein P62962

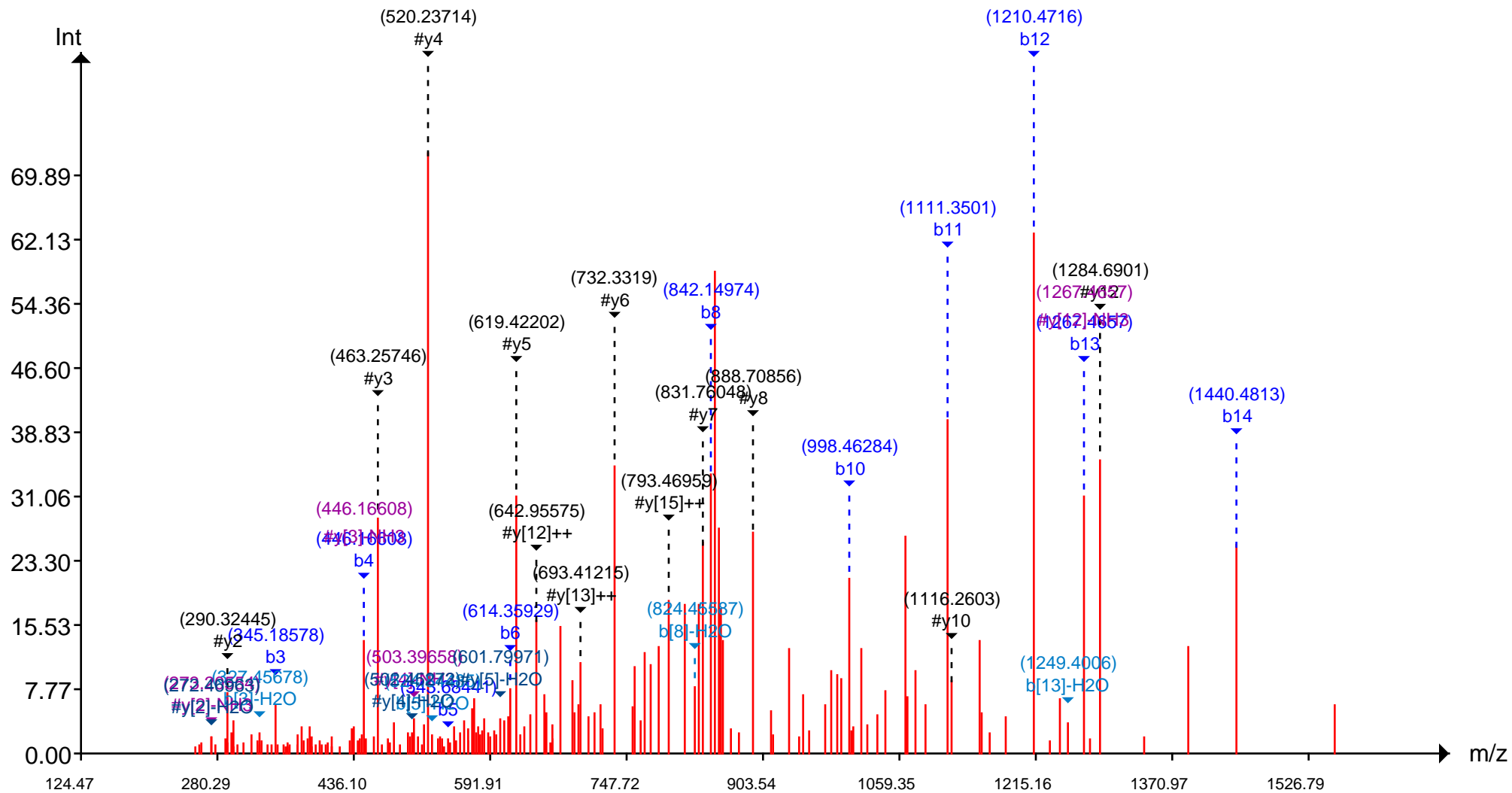
Confidence level 99.00%

Precursor m/z 865.50296

Identity Threshold 45.02

Charge 2+

Mass error 0.002095



Info

Modified Sequence AcD3-VSITPAEVGVLVGK<AcD3K*>DR-COOH

Ion score 51.32

Protein P62962

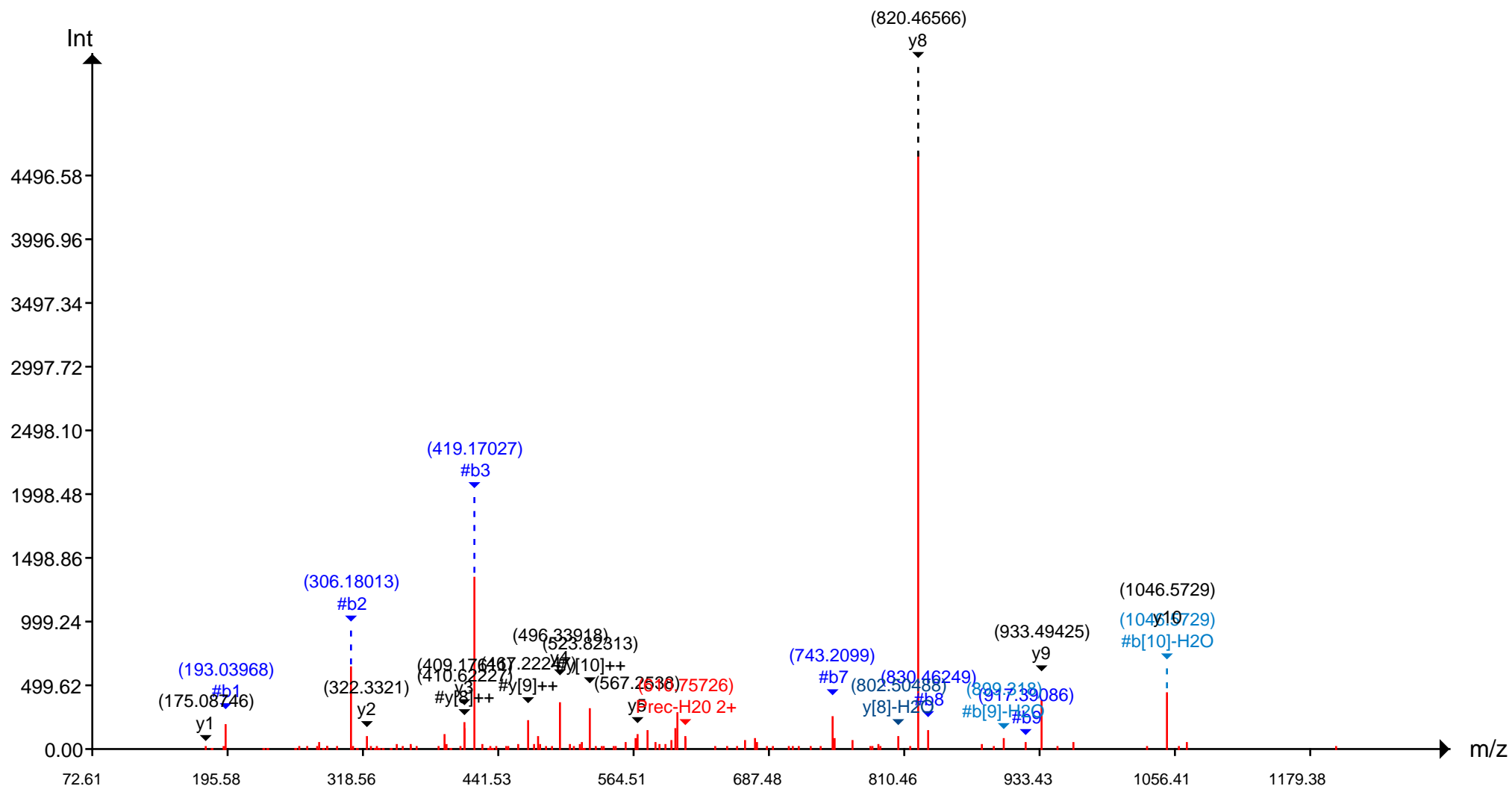
Confidence level 99.00%

Precursor m/z 865.50323

Identity Threshold 44.97

Charge 2+

Mass error 0.002635



Info

Modified Sequence AcD3-M<Mox*>ILPVGASSFR-COOH

Ion score 62.16

Protein P17182

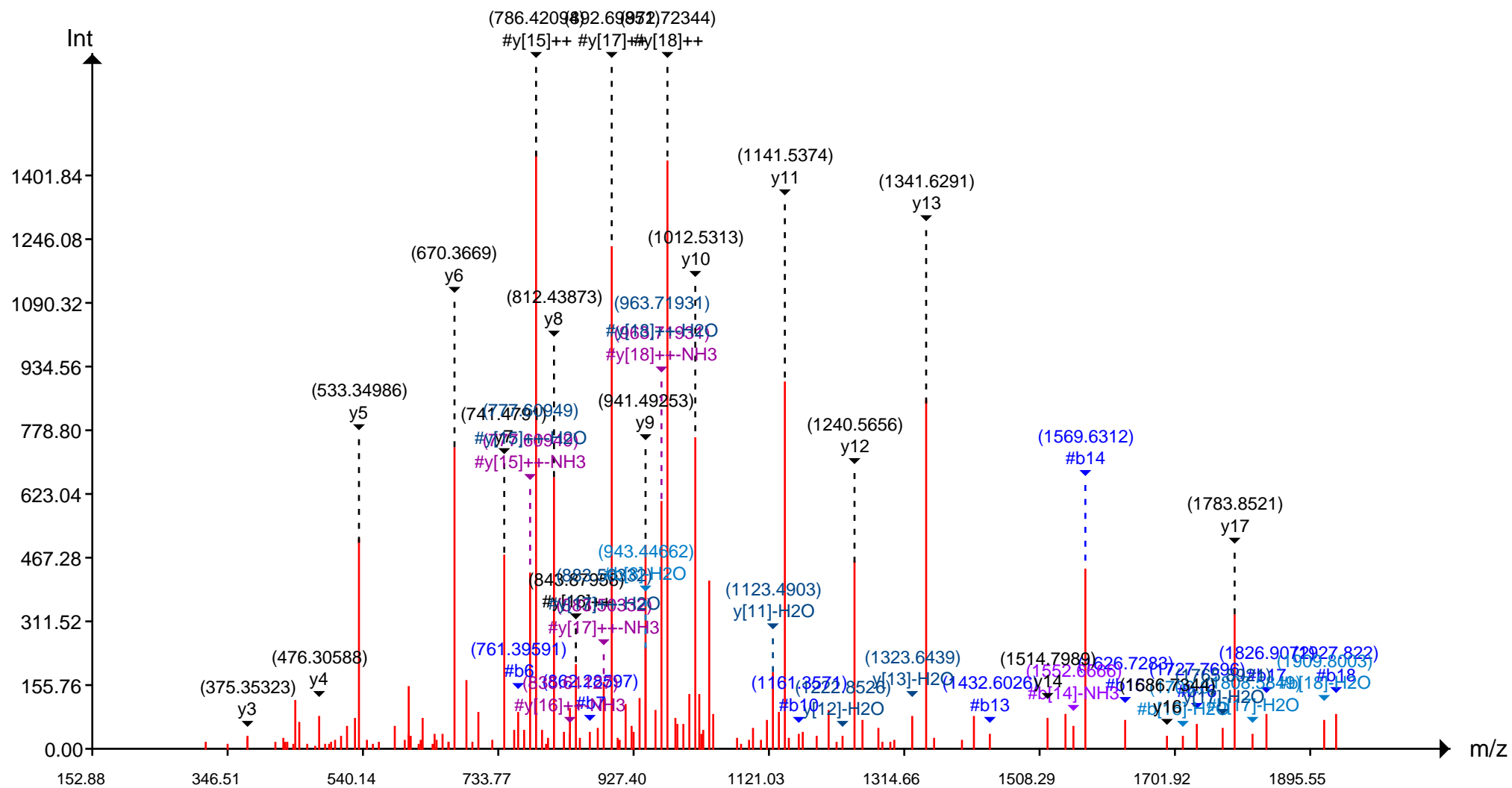
Confidence level 99.00%

Precursor m/z 619.83584

Identity Threshold 47.95

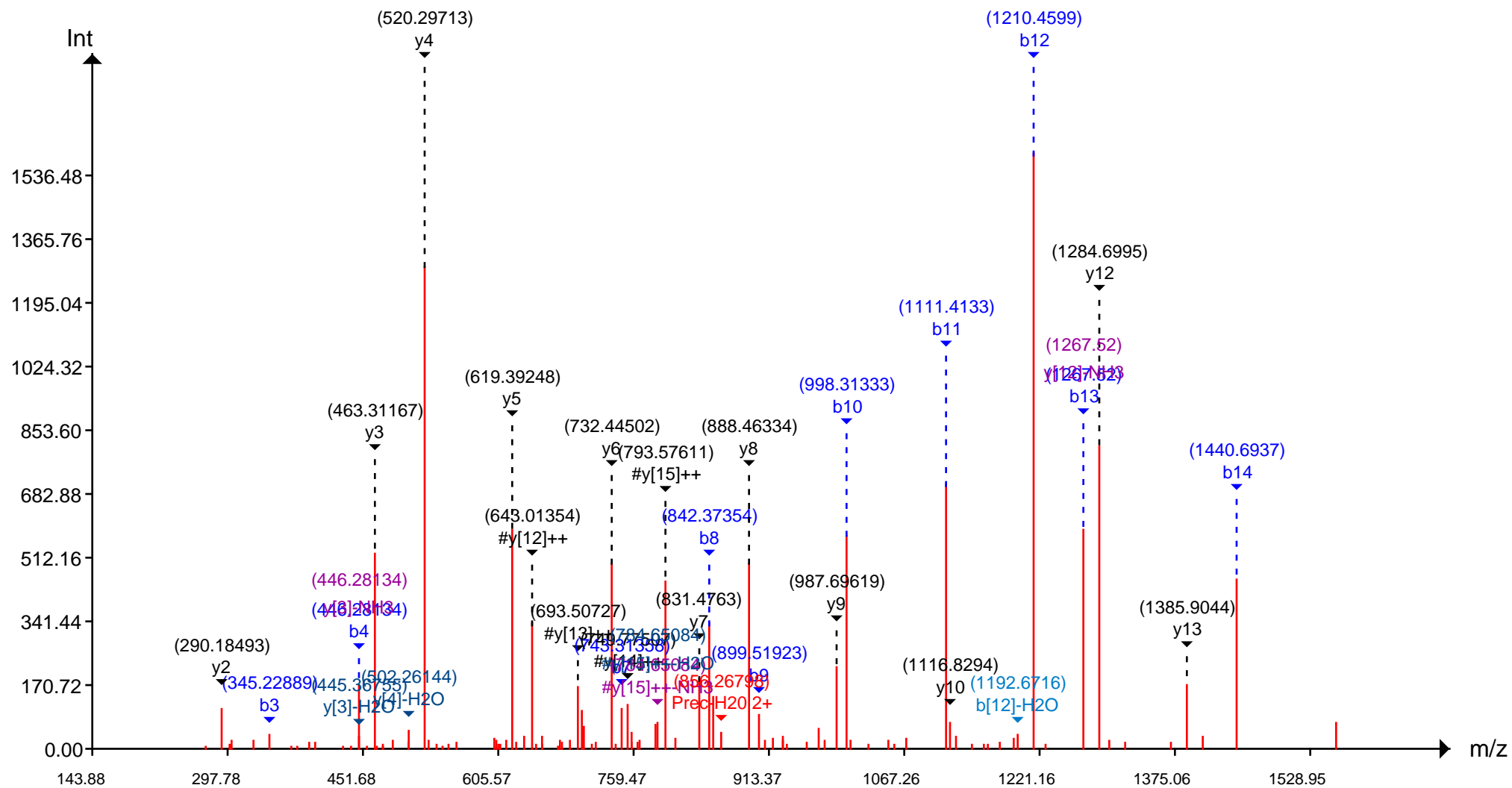
Charge 2+

Mass error 3.13E-4



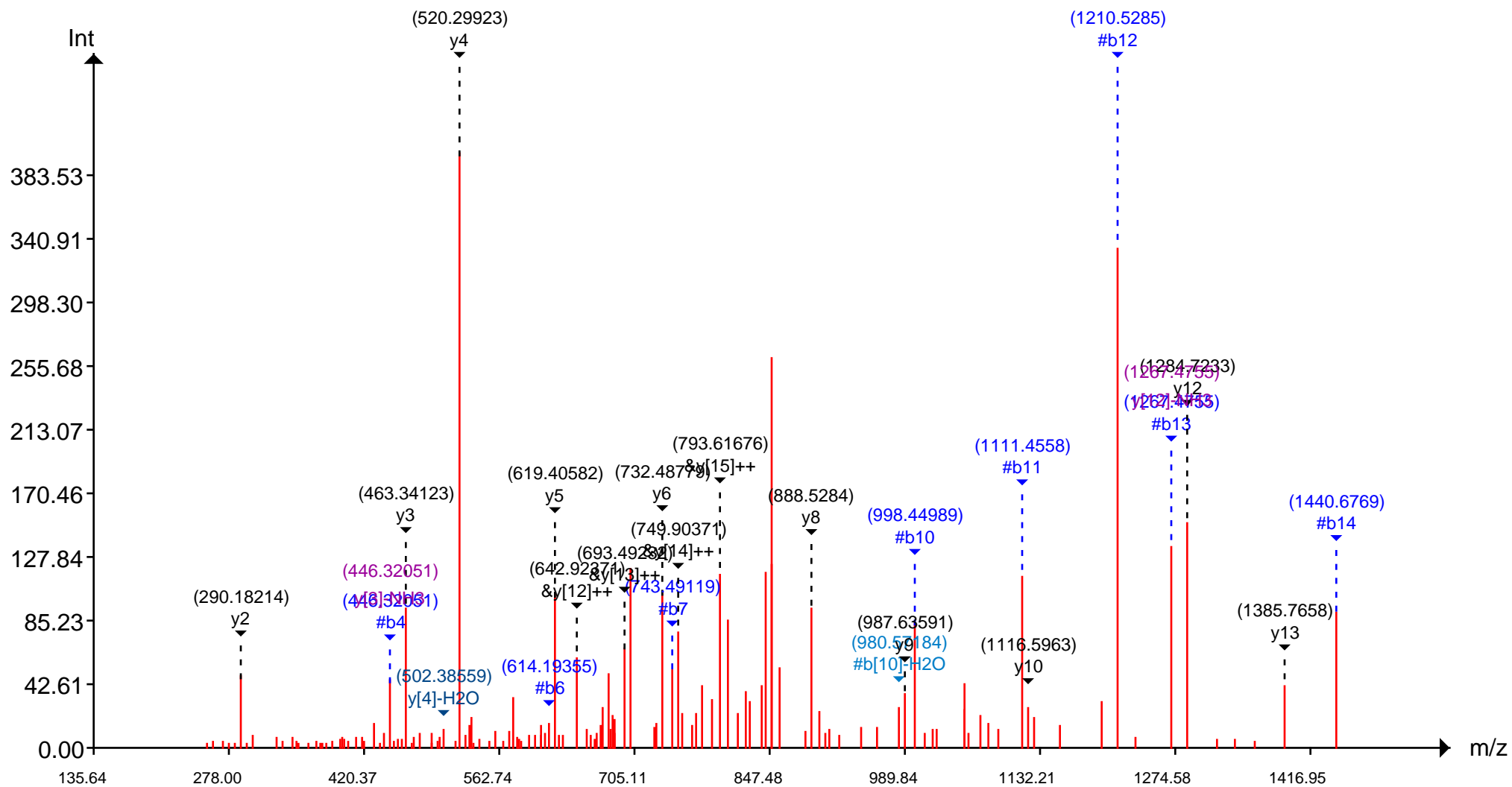
Info

Modified Sequence	AcD3-IC<Cmm*>PDGK<AcD3K*>TVEAEAAHGTVTR-COOH	Ion score	101.11
Protein	O88844	Confidence level	99.00%
Precursor m/z	1051.5267	Identity Threshold	47.69
Charge	2+	Mass error	7.62E-4



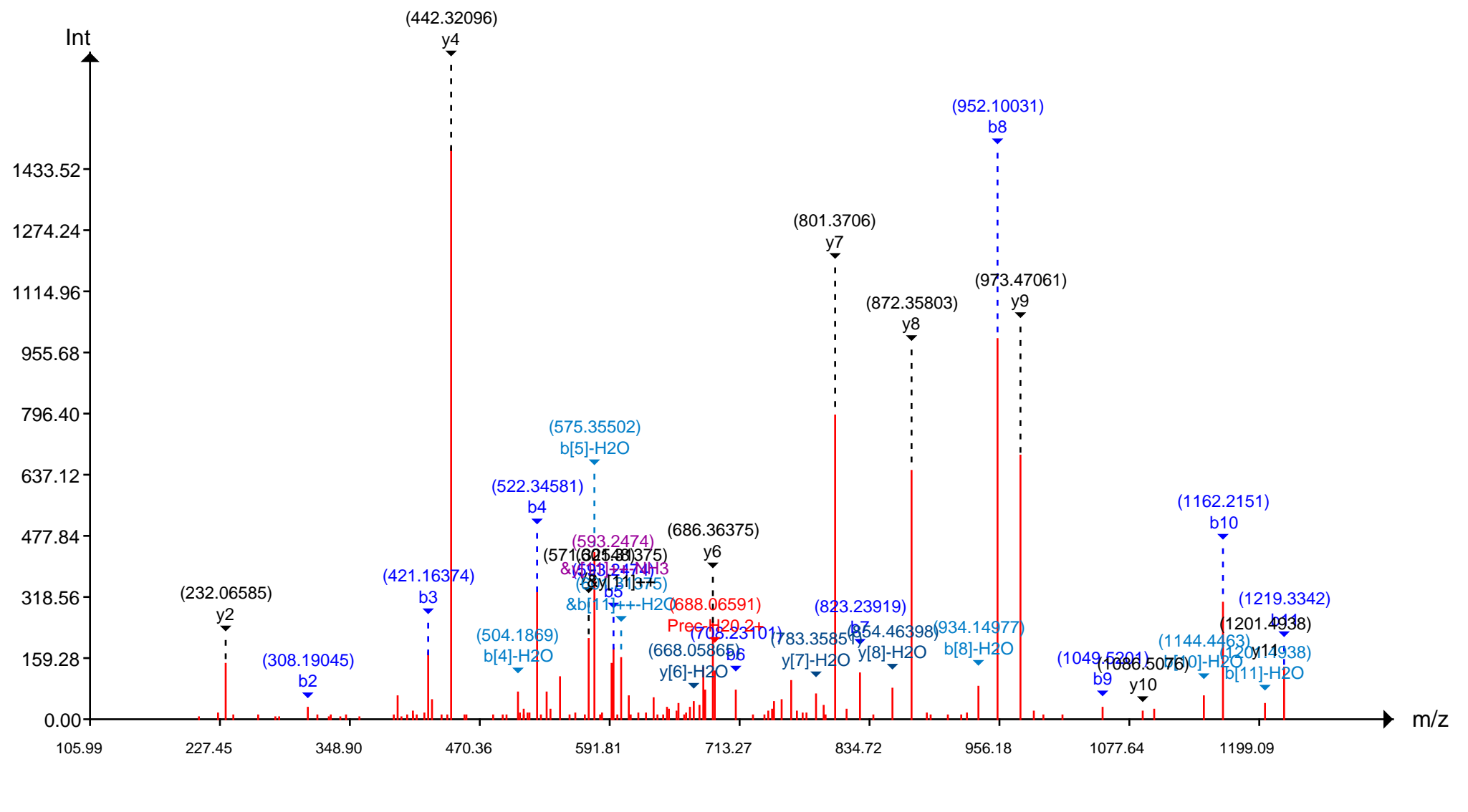
Info

Modified Sequence	AcD3-VSITPAEVGVLVGK<AcD3K*>DR-COOH	Ion score	104.01
Protein	P62962	Confidence level	99.00%
Precursor m/z	865.50316	Identity Threshold	44.97
Charge	2+	Mass error	0.002495

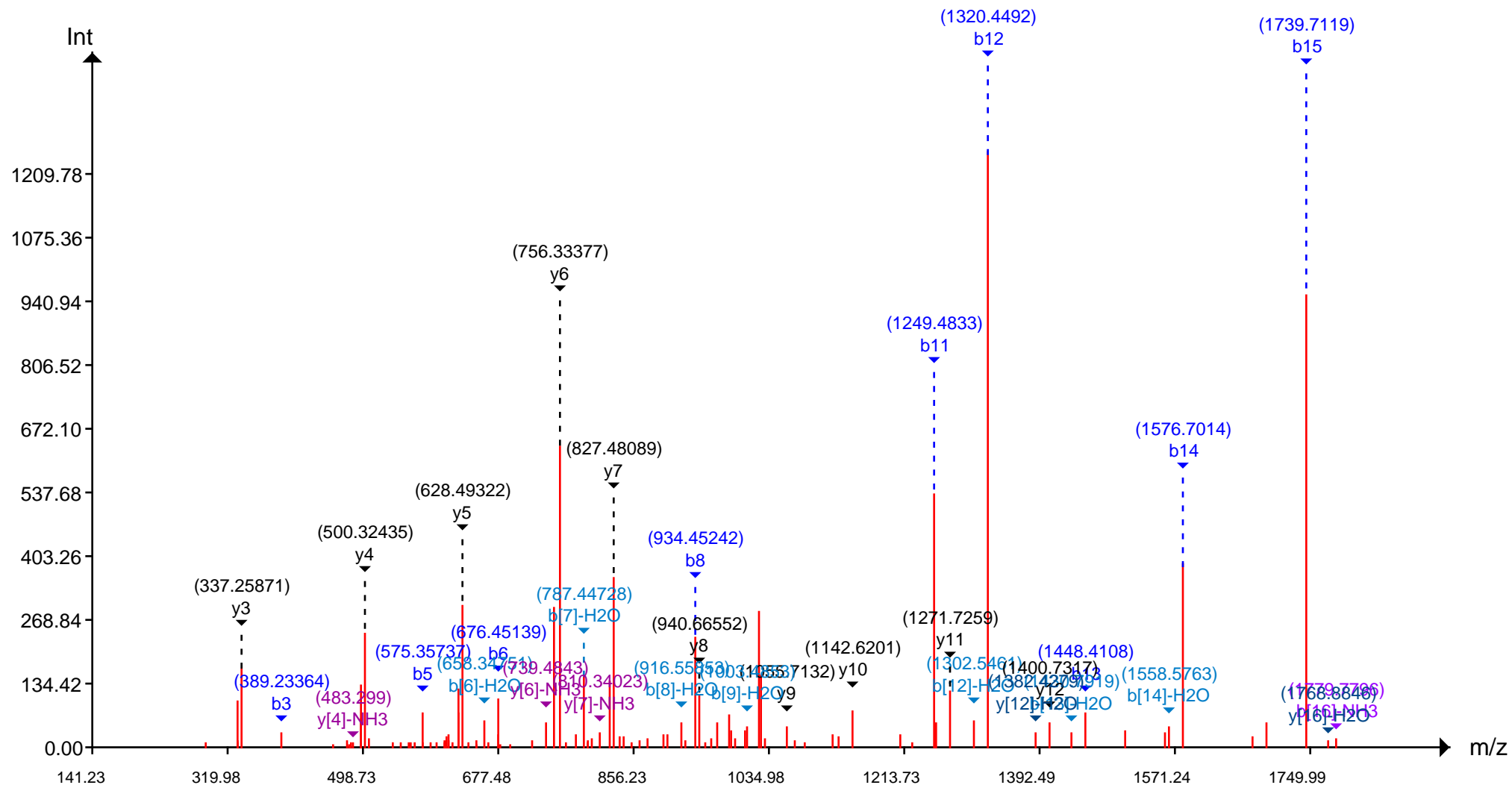


Info

Modified Sequence	AcD3-VSITPAEVLVVGK<AcD3K*>DR-COOH	Ion score	65.49
Protein	P62962	Confidence level	99.00%
Precursor m/z	865.50287	Identity Threshold	45.00
Charge	2+	Mass error	0.001915

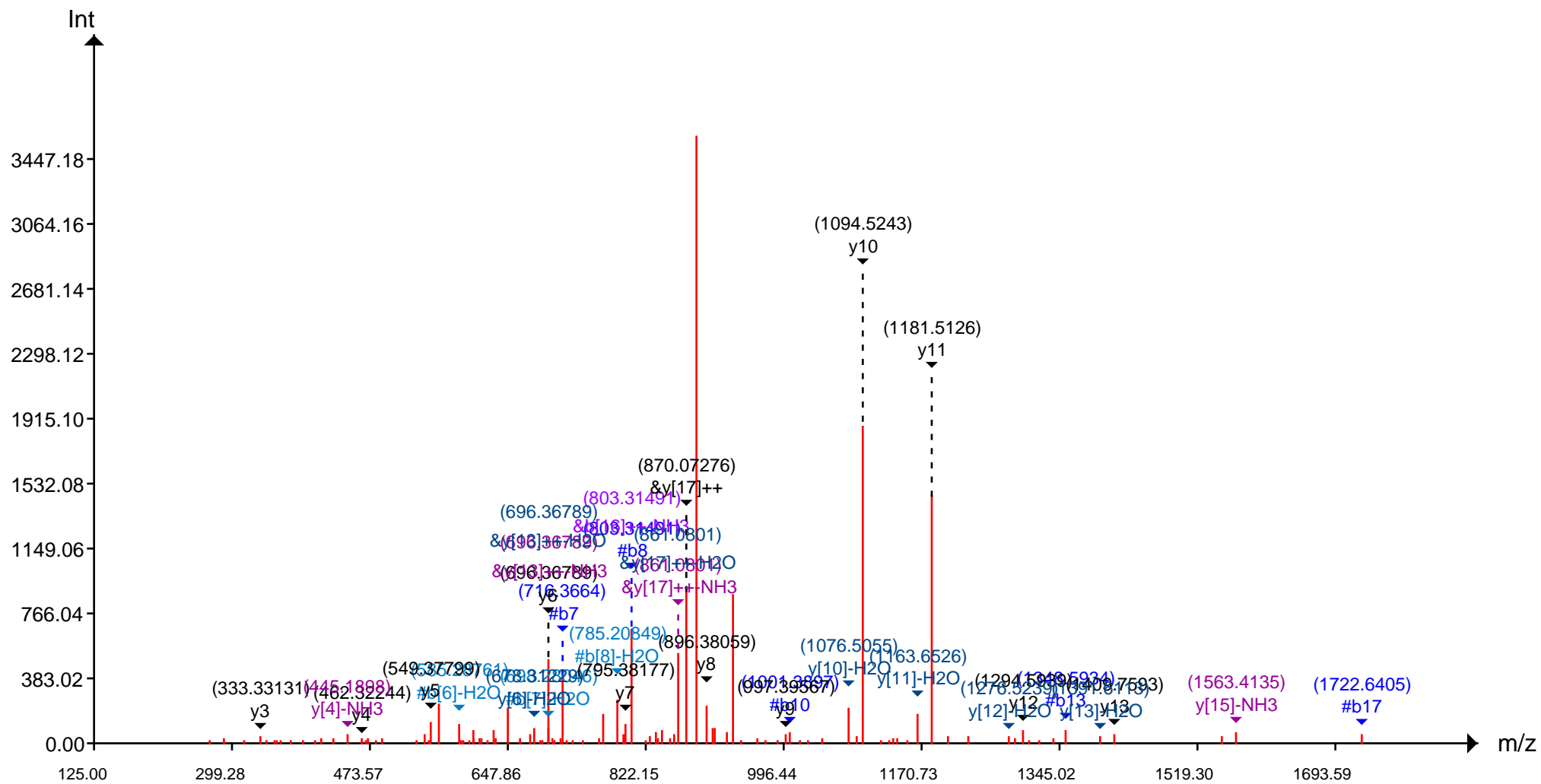


Info			
Modified Sequence	AcD3-FDITADDEPLGR-COOH	Ion score	87.89
Protein	P17742	Confidence level	99.00%
Precursor m/z	697.33843	Identity Threshold	46.90
Charge	2+	Mass error	0.002259



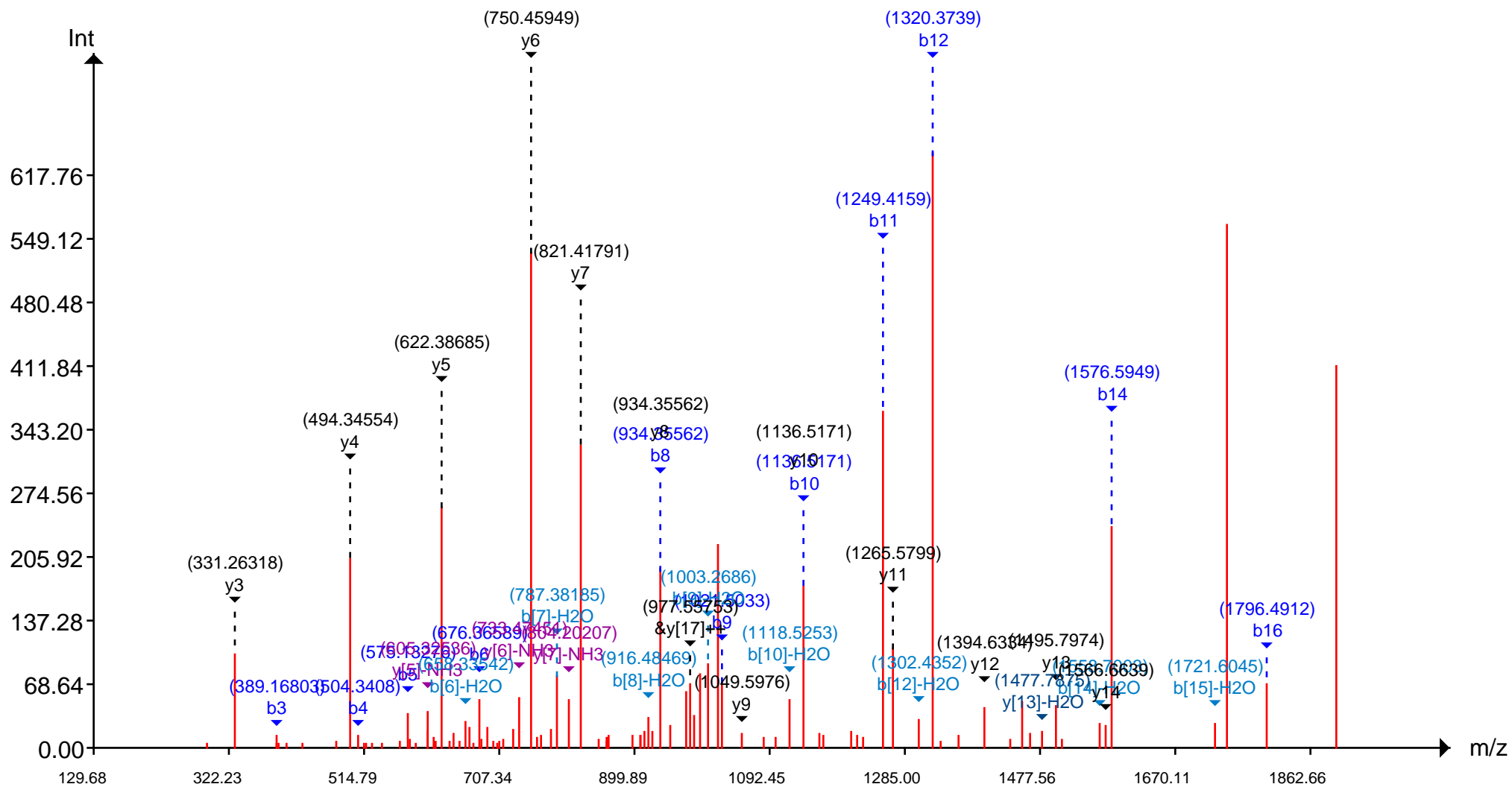
Info

Modified Sequence	AcD3-AK<AcD3K*>VDATEESDLAQQYGVR<C13>-COOH	Ion score	74.21
Protein	P09103	Confidence level	99.00%
Precursor m/z	1038.528	Identity Threshold	47.94
Charge	2+	Mass error	0.002935



Info

Modified Sequence	AcD3-LGTGNDLSPTTVM<Mox*>SEGTR-COOH	Ion score	65.42
Protein	Q923D2	Confidence level	99.00%
Precursor m/z	948.95874	Identity Threshold	46.88
Charge	2+	Mass error	0.005513



Info

Modified Sequence AcD3-AK<AcD3K*>VDATEESDLAQQYGVR-COOH

Ion score 94.12

Protein P09103

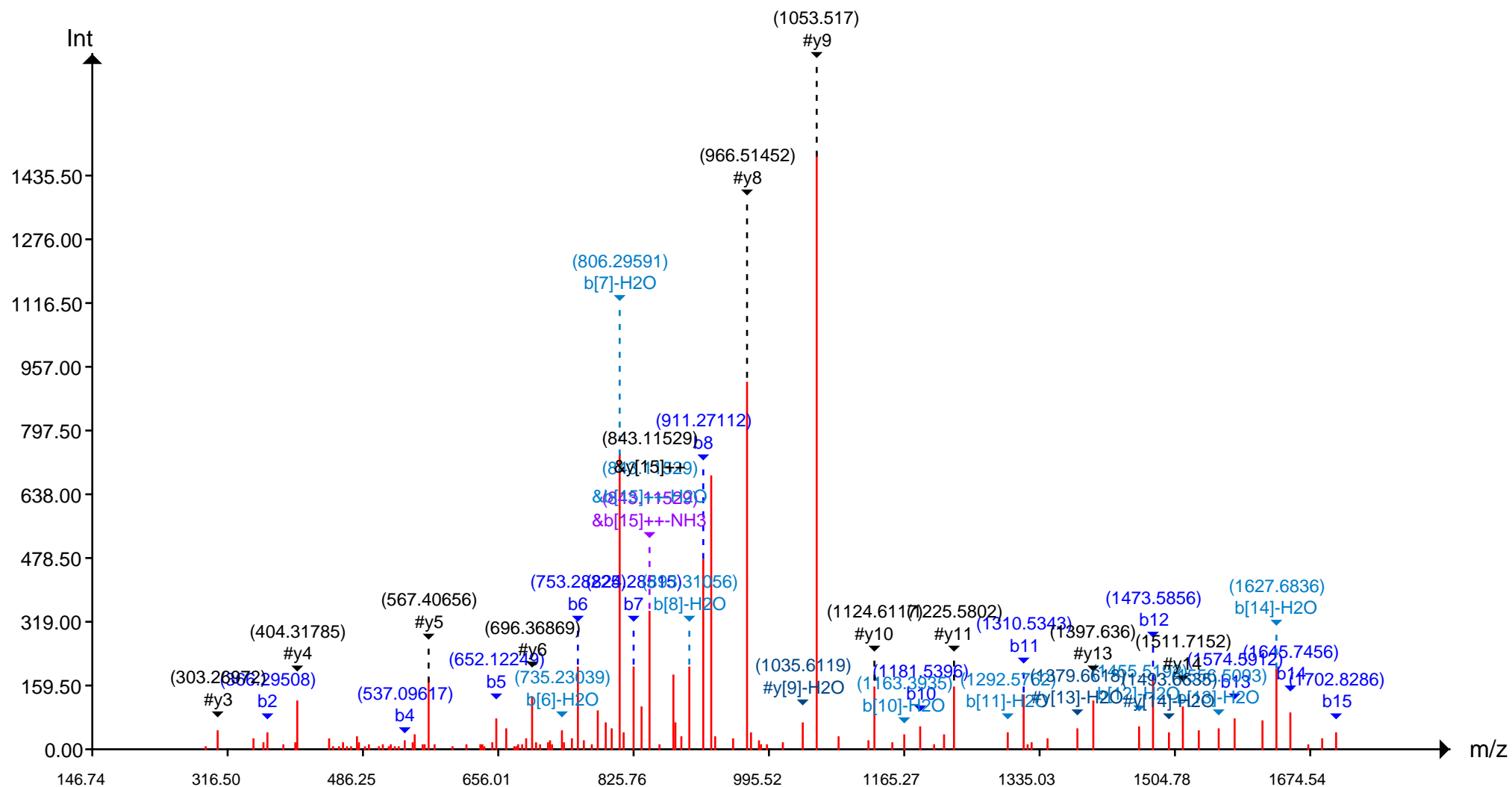
Confidence level 99.00%

Precursor m/z 1035.5179

Identity Threshold 47.68

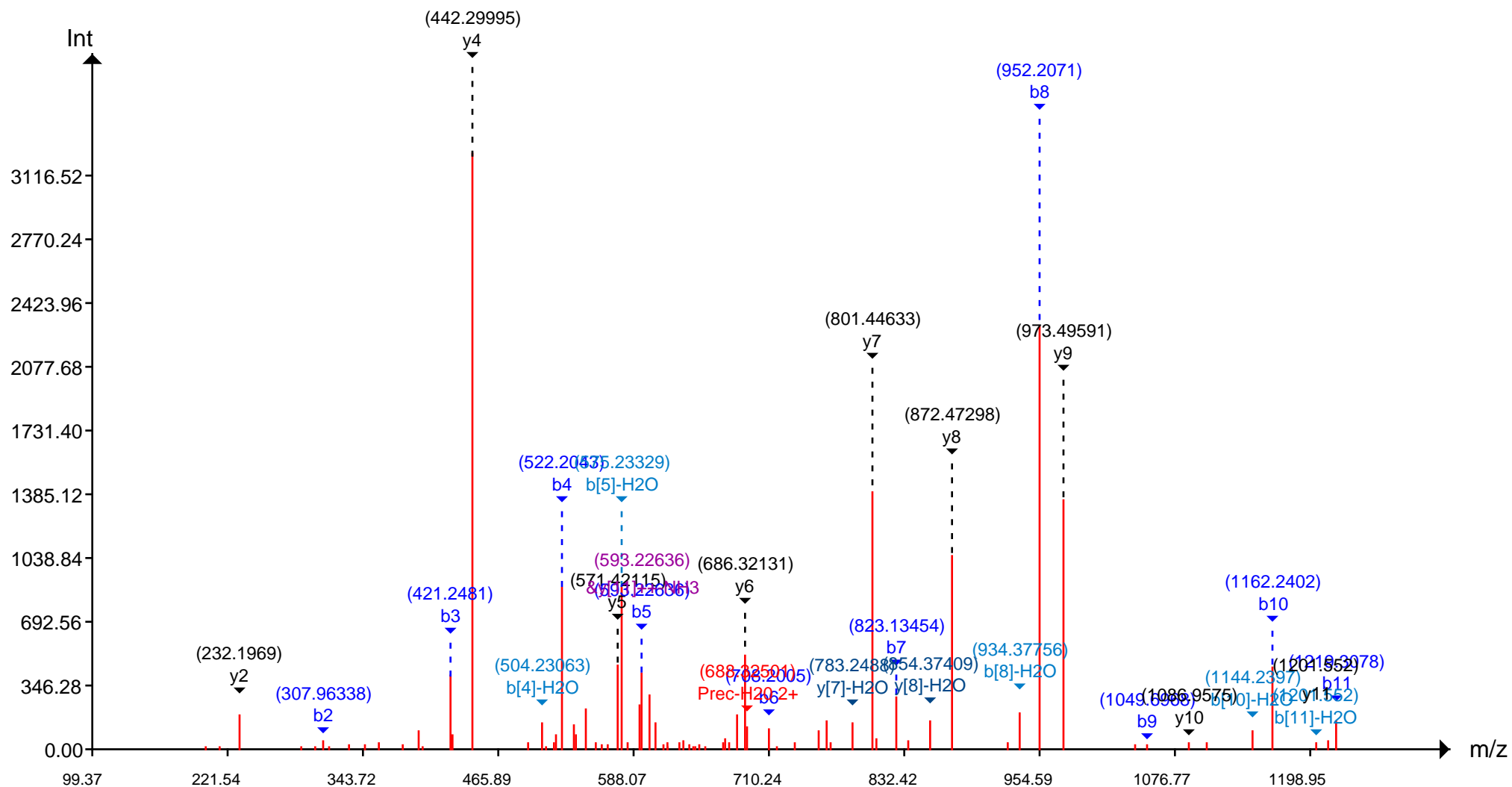
Charge 2+

Mass error 0.002876



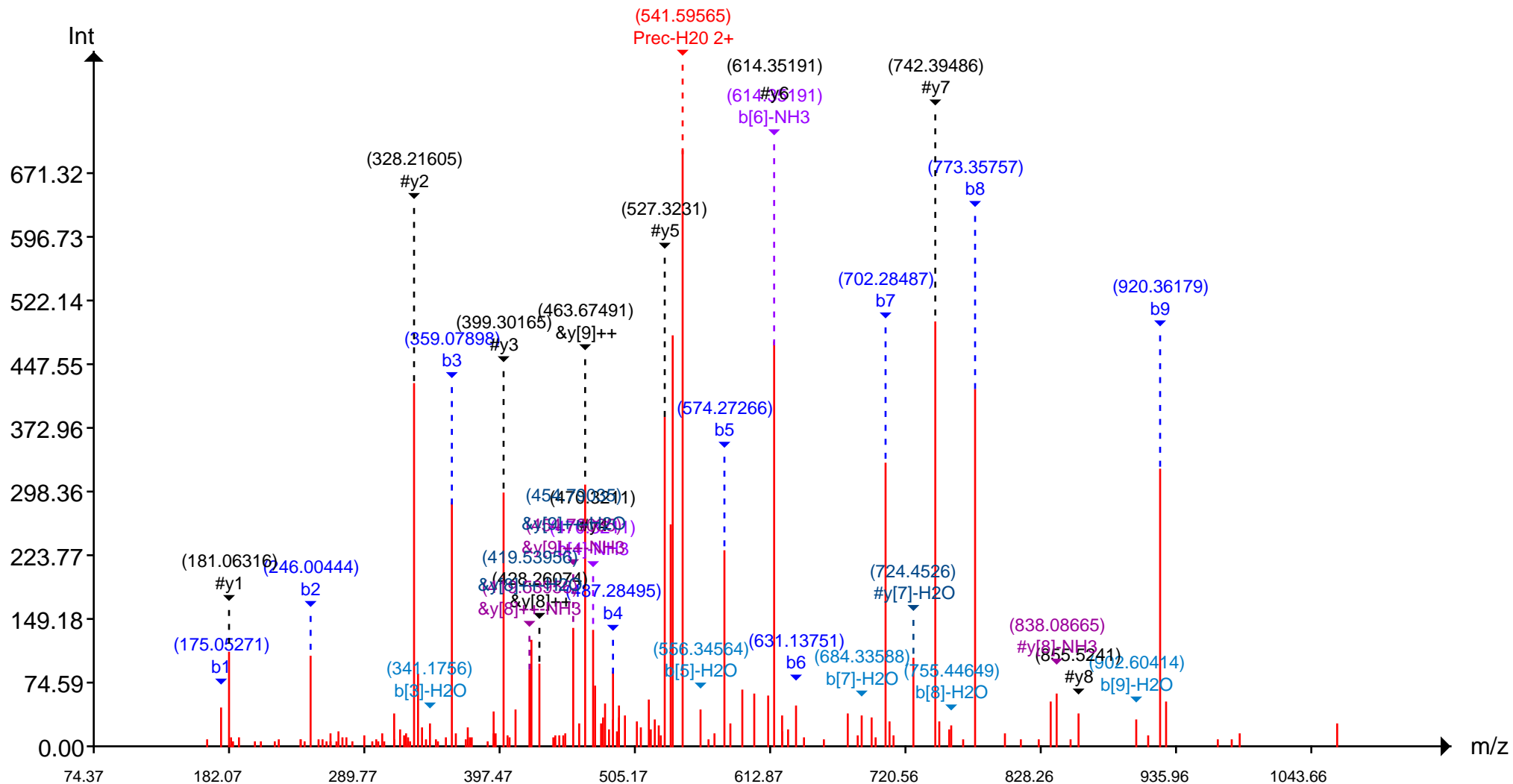
Info

Modified Sequence	AcD3-FK<AcD3K*>NGDTASPK<AcD3K*>EYTAGR-COOH	Ion score	106.15
Protein	P09103	Confidence level	99.00%
Precursor m/z	938.97439	Identity Threshold	47.93
Charge	2+	Mass error	0.002938



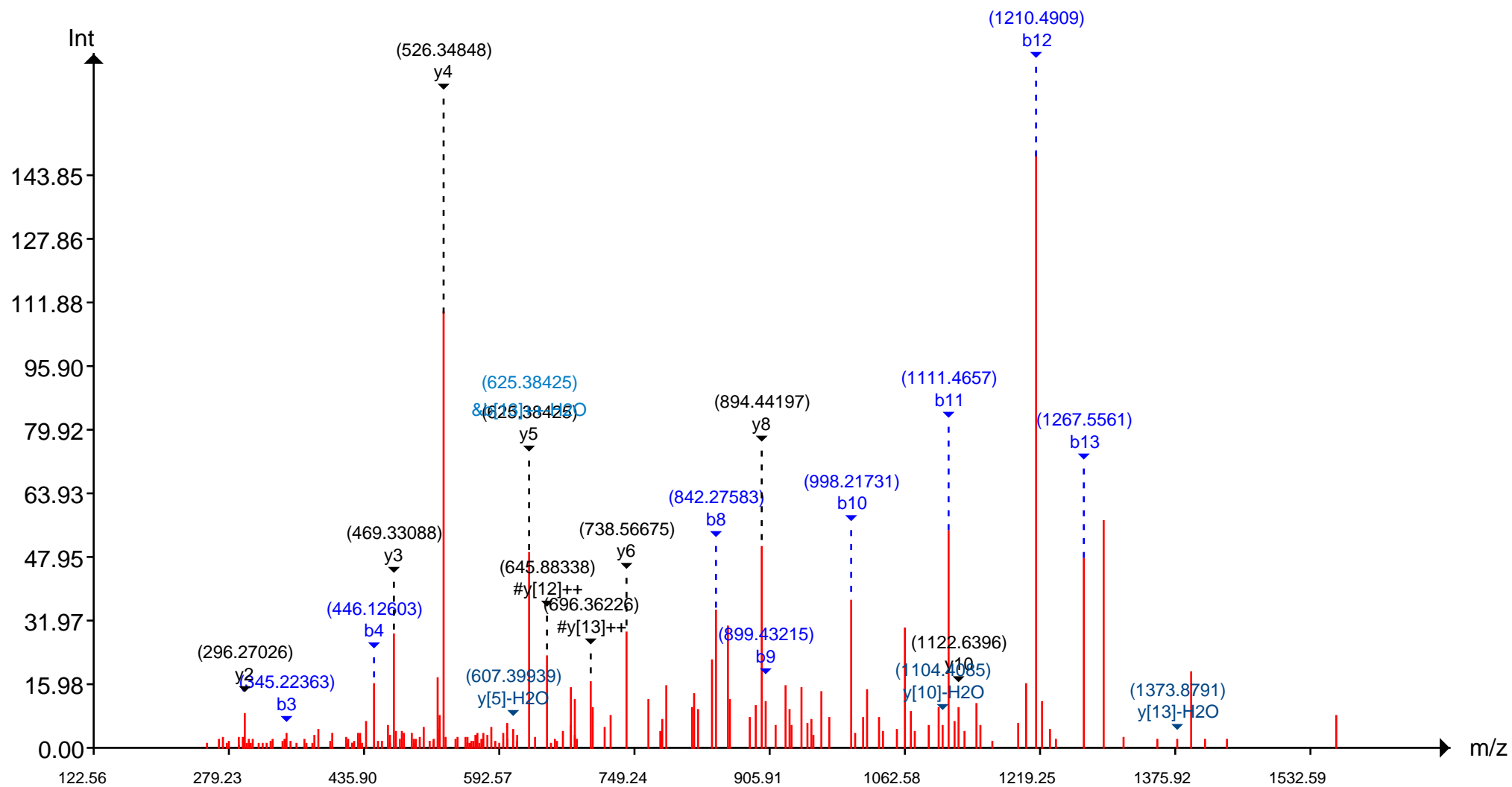
Info

Modified Sequence	AcD3-FDITADDEPLGR-COOH	Ion score	85.78
Protein	P17742	Confidence level	99.00%
Precursor m/z	697.33829	Identity Threshold	46.85
Charge	2+	Mass error	0.001979



Info

Modified Sequence	AcD3-EALQSGAAFR<C13>-COOH	Ion score	62.86
Protein	O08808, O70566, Q9Z207	Confidence level	99.00%
Precursor m/z	550.79699	Identity Threshold	48.00
Charge	2+	Mass error	-2.38E-4



Info

Modified Sequence AcD3-VSITPAEVLVVGK<AcD3K*>DR<C13>-COOH

Ion score 47.24

Protein P62962

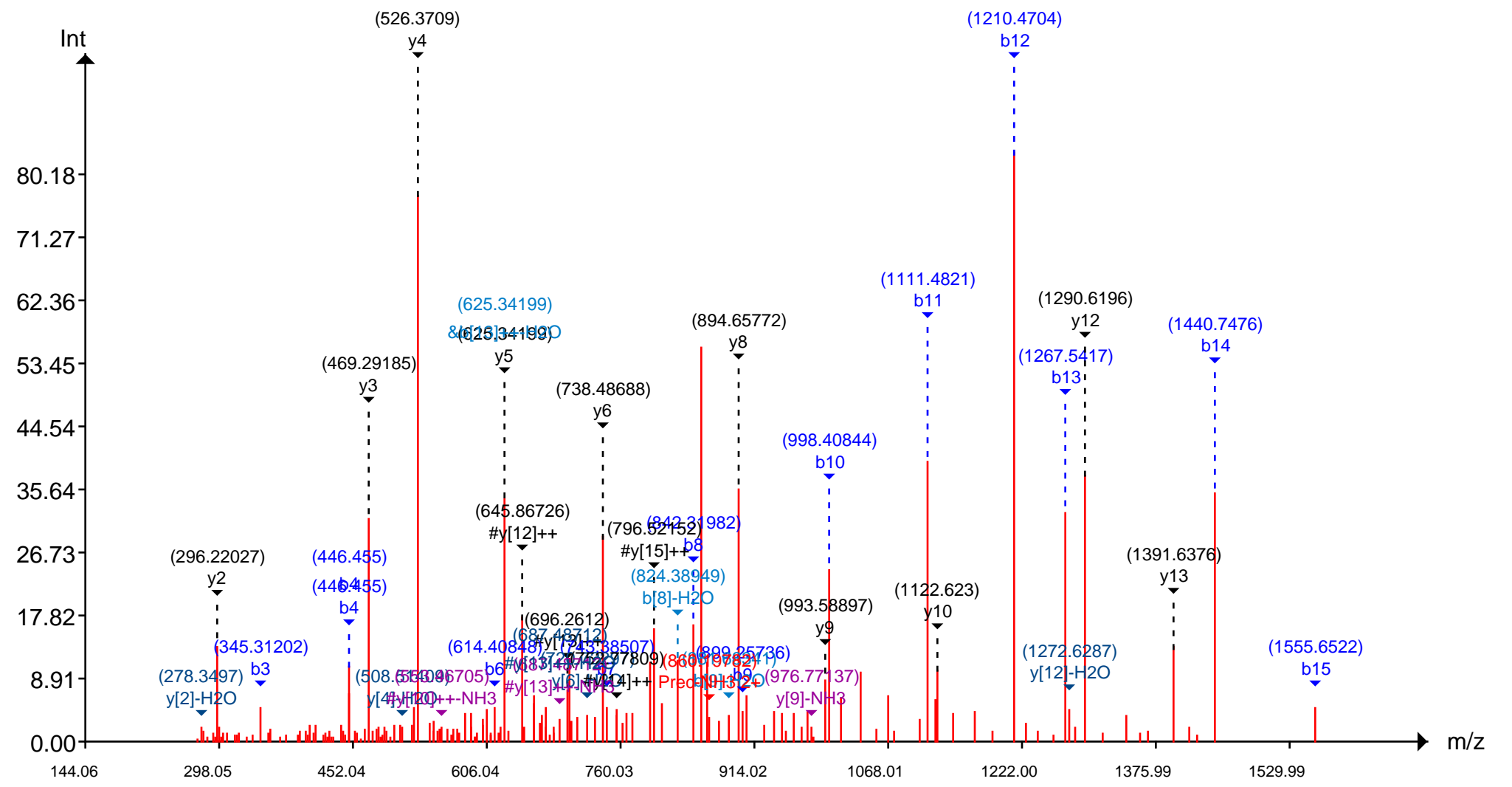
Confidence level 99.00%

Precursor m/z 868.51364

Identity Threshold 43.75

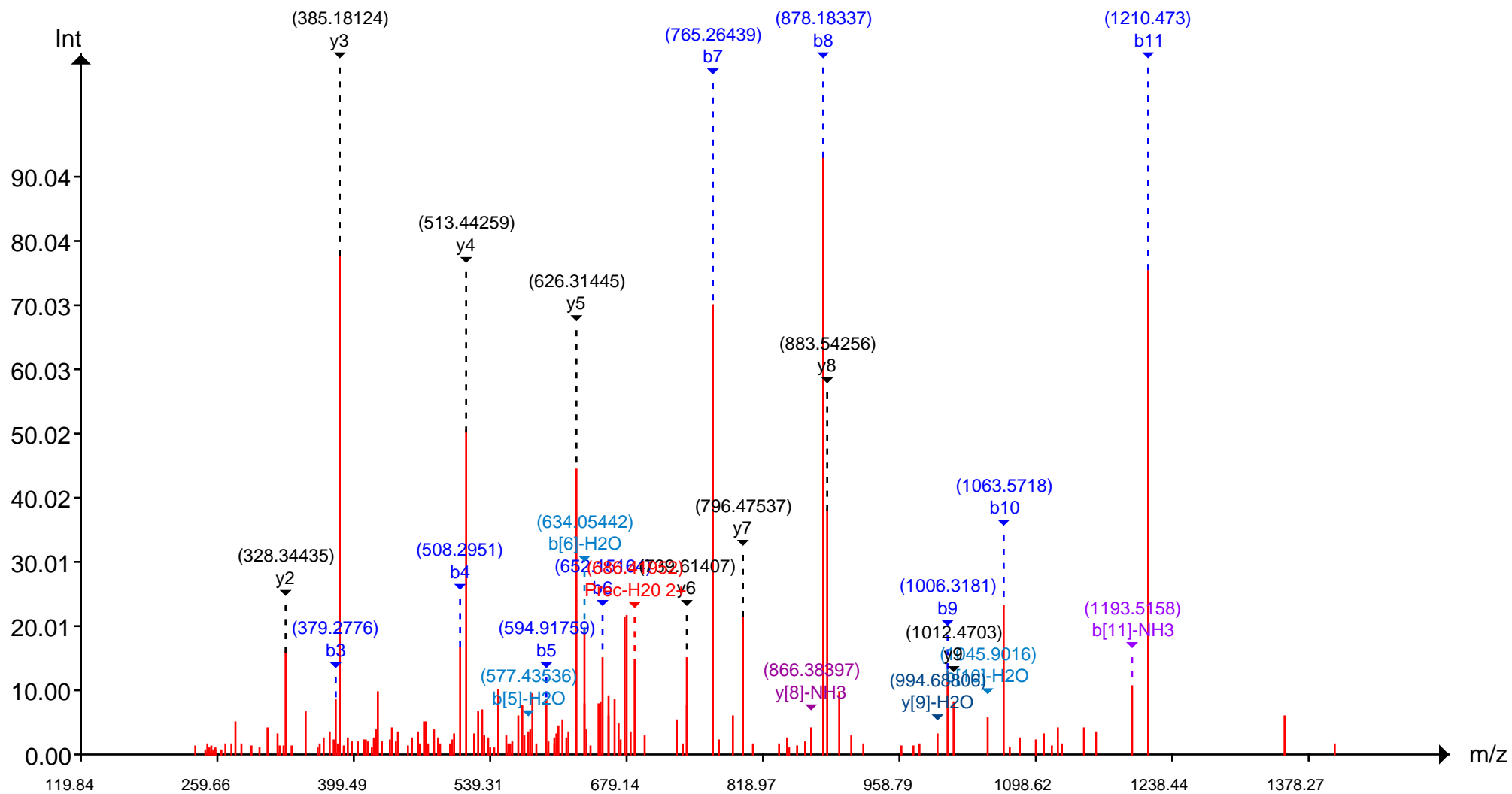
Charge 2+

Mass error 0.003313



Info

Modified Sequence	AcD3-VSITPAEVGVLVGK<AcD3K*>DR<C13>-COOH	Ion score	87.07
Protein	P62962	Confidence level	99.00%
Precursor m/z	868.51391	Identity Threshold	43.77
Charge	2+	Mass error	0.003853



Info

Modified Sequence AcD3-ADFESGLLQGFR<C13>-COOH

Ion score 62.58

Protein Q9JM14

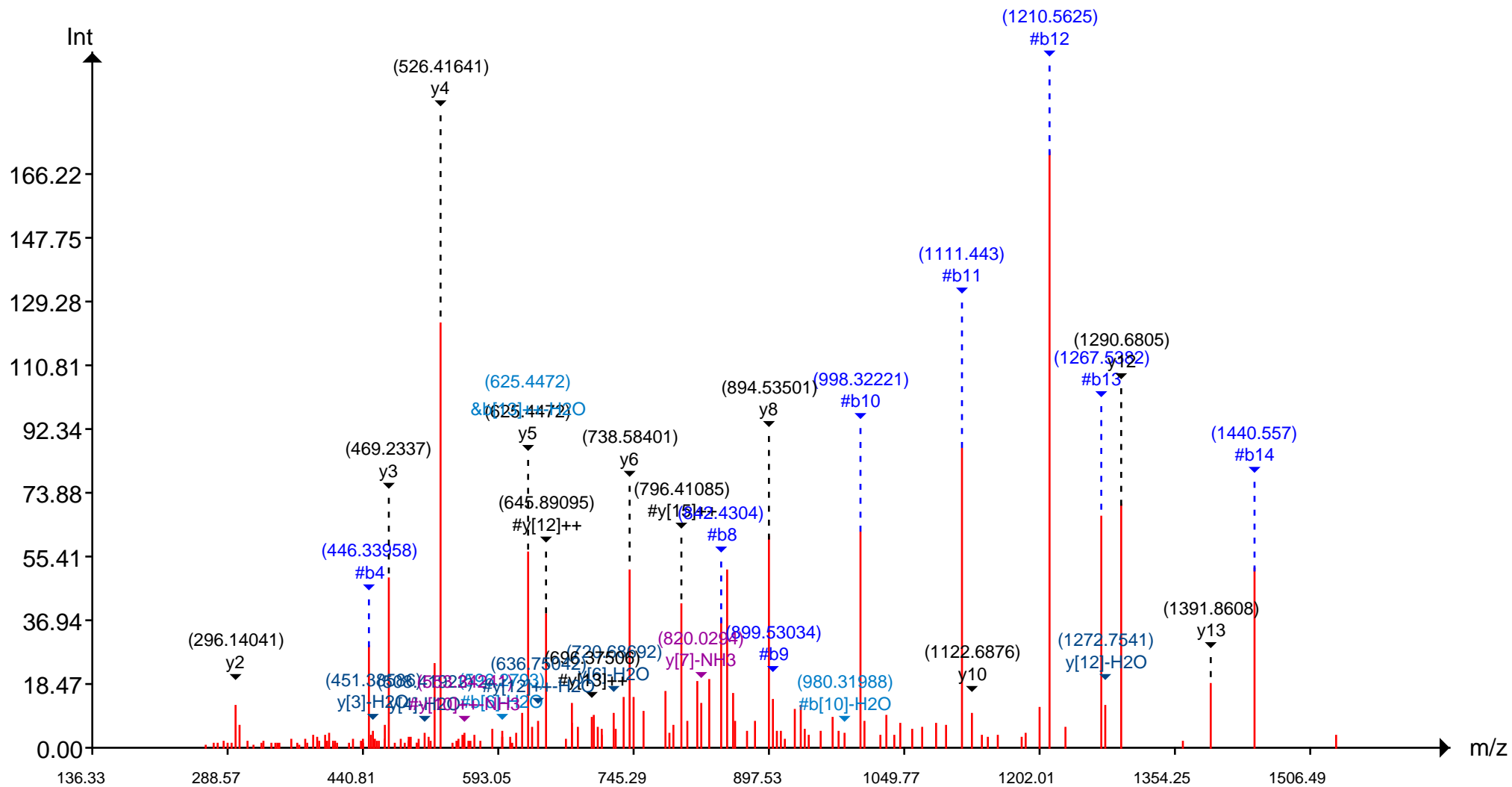
Confidence level 99.00%

Precursor m/z 695.86085

Identity Threshold 47.92

Charge 2+

Mass error 8.03E-4



Info

Modified Sequence AcD3-VSITPAEVGVLVGK<AcD3K*>DR<C13>-COOH

Ion score 72.05

Protein P62962

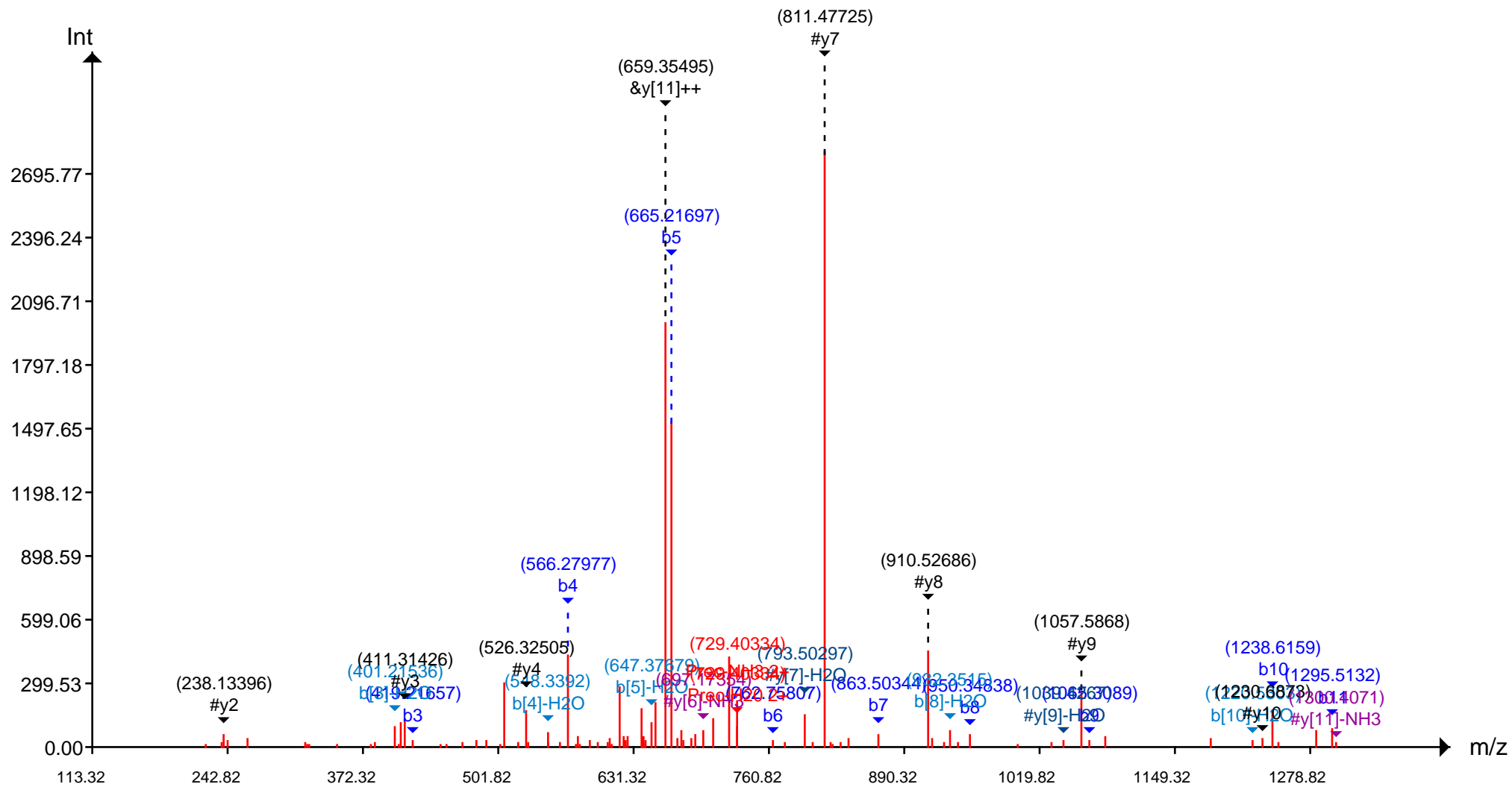
Confidence level 99.00%

Precursor m/z 868.51333

Identity Threshold 43.75

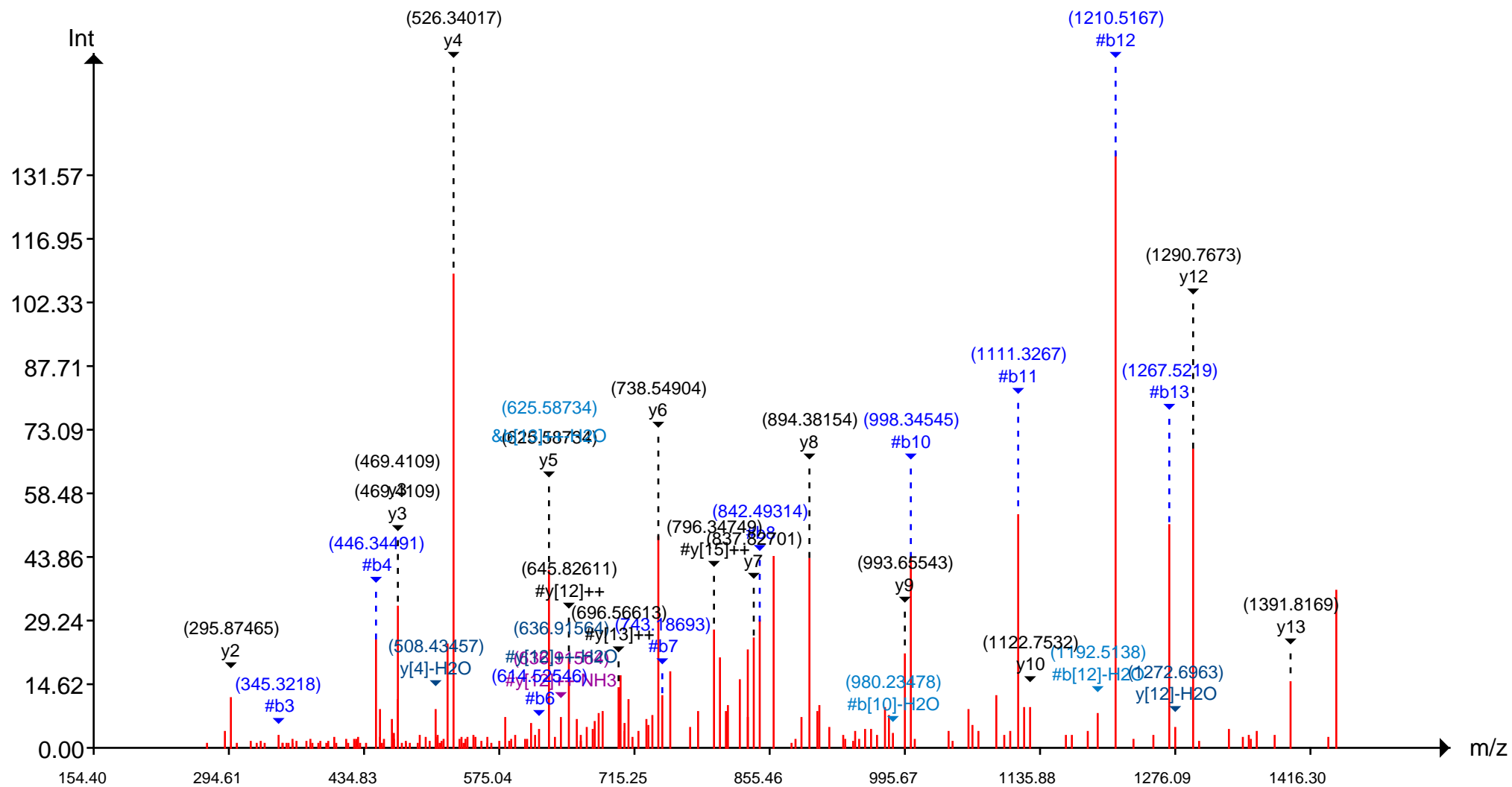
Charge 2+

Mass error 0.002693



Info

Modified Sequence	AcD3-ISK<AcD3K*>M<Mox*>VPTSDK<AcD3K*>GR<C13>-COOH	Ion score	51.67
Protein	P58252	Confidence level	99.00%
Precursor m/z	738.41379	Identity Threshold	47.30
Charge	2+	Mass error	0.002313



Info

Modified Sequence AcD3-VSITPAEVGVLVGK<AcD3K*>DR<C13>-COOH

Ion score 72.19

Protein P62962

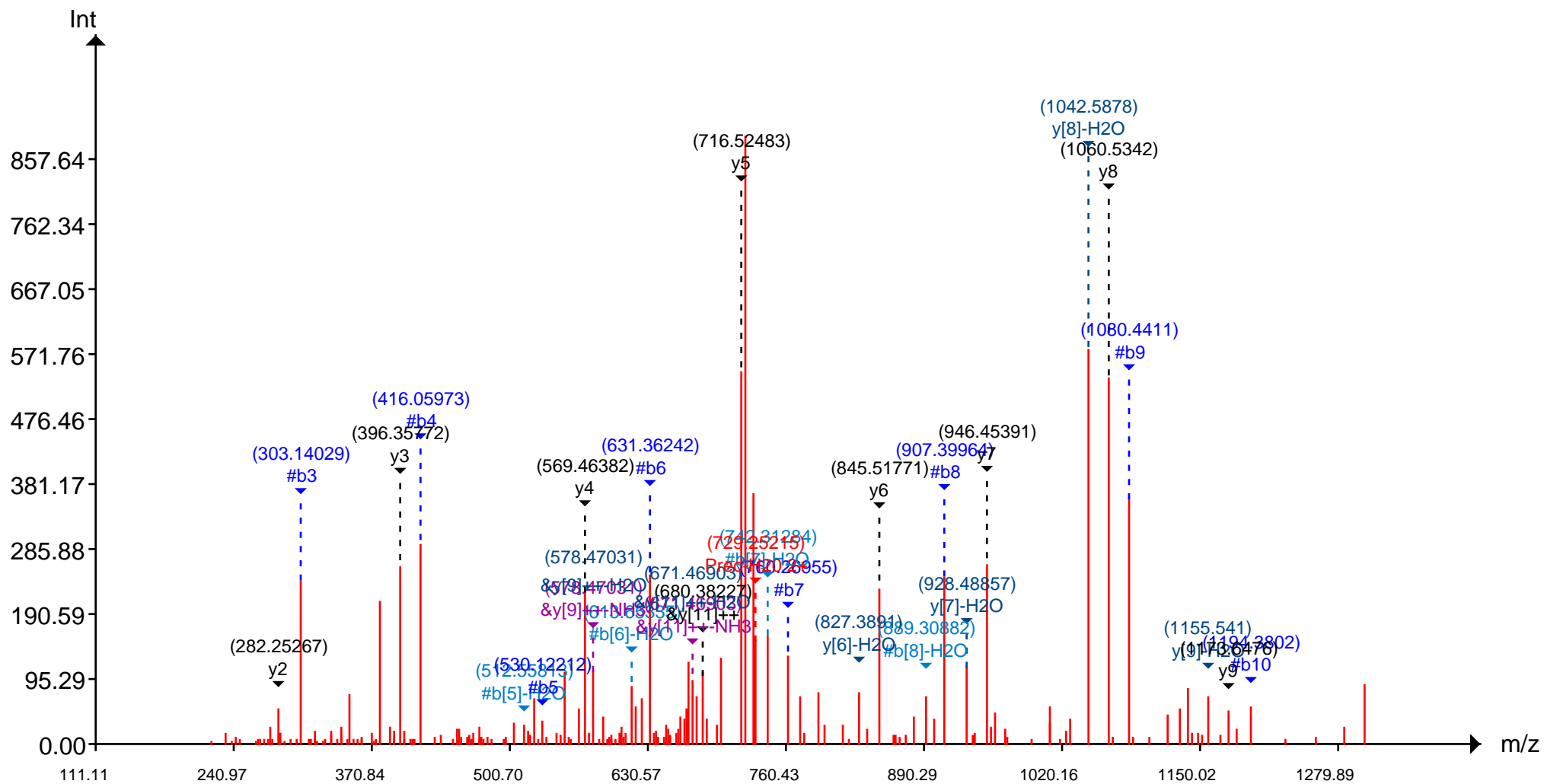
Confidence level 99.00%

Precursor m/z 868.51341

Identity Threshold 43.76

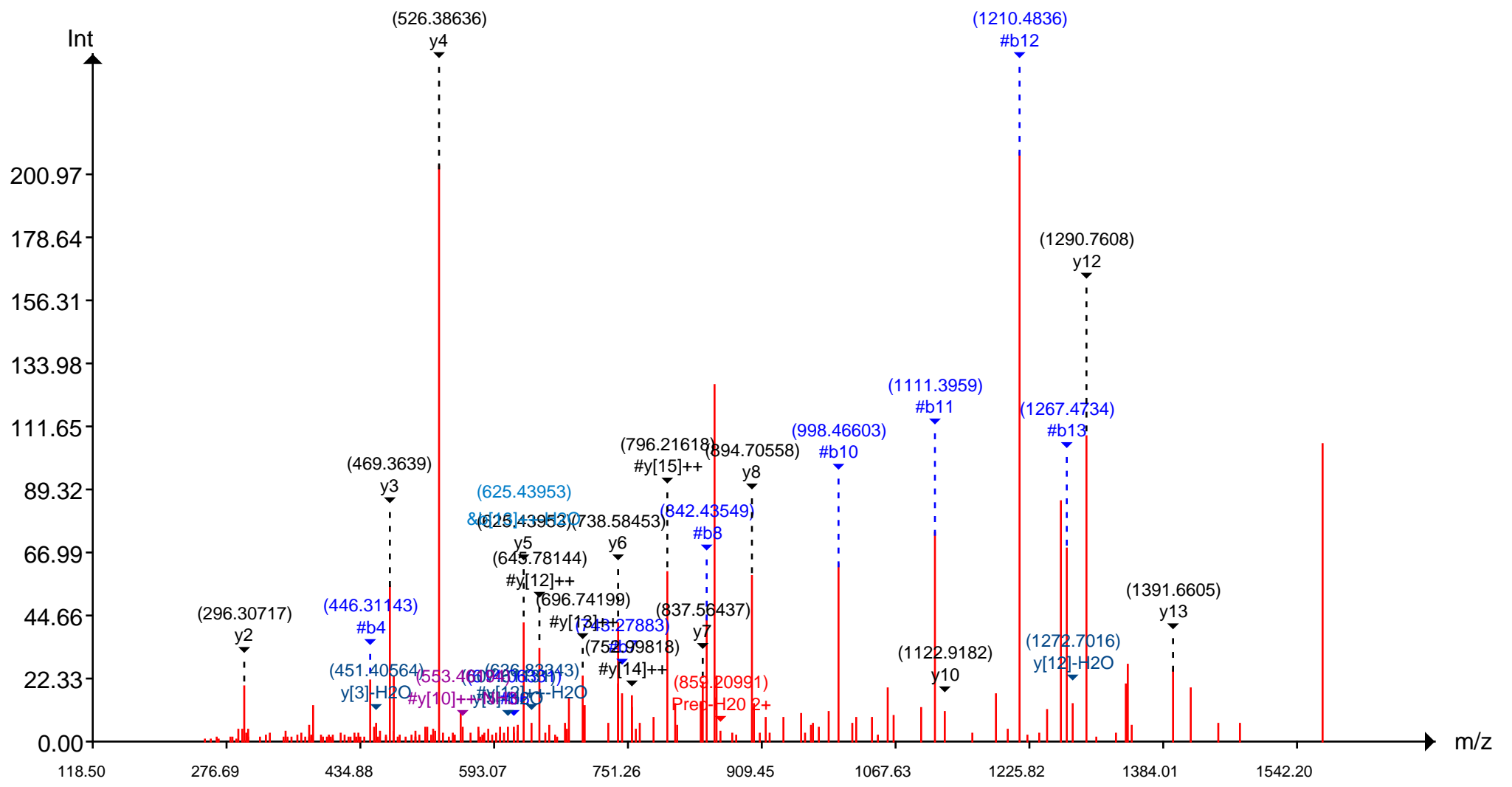
Charge 2+

Mass error 0.002853

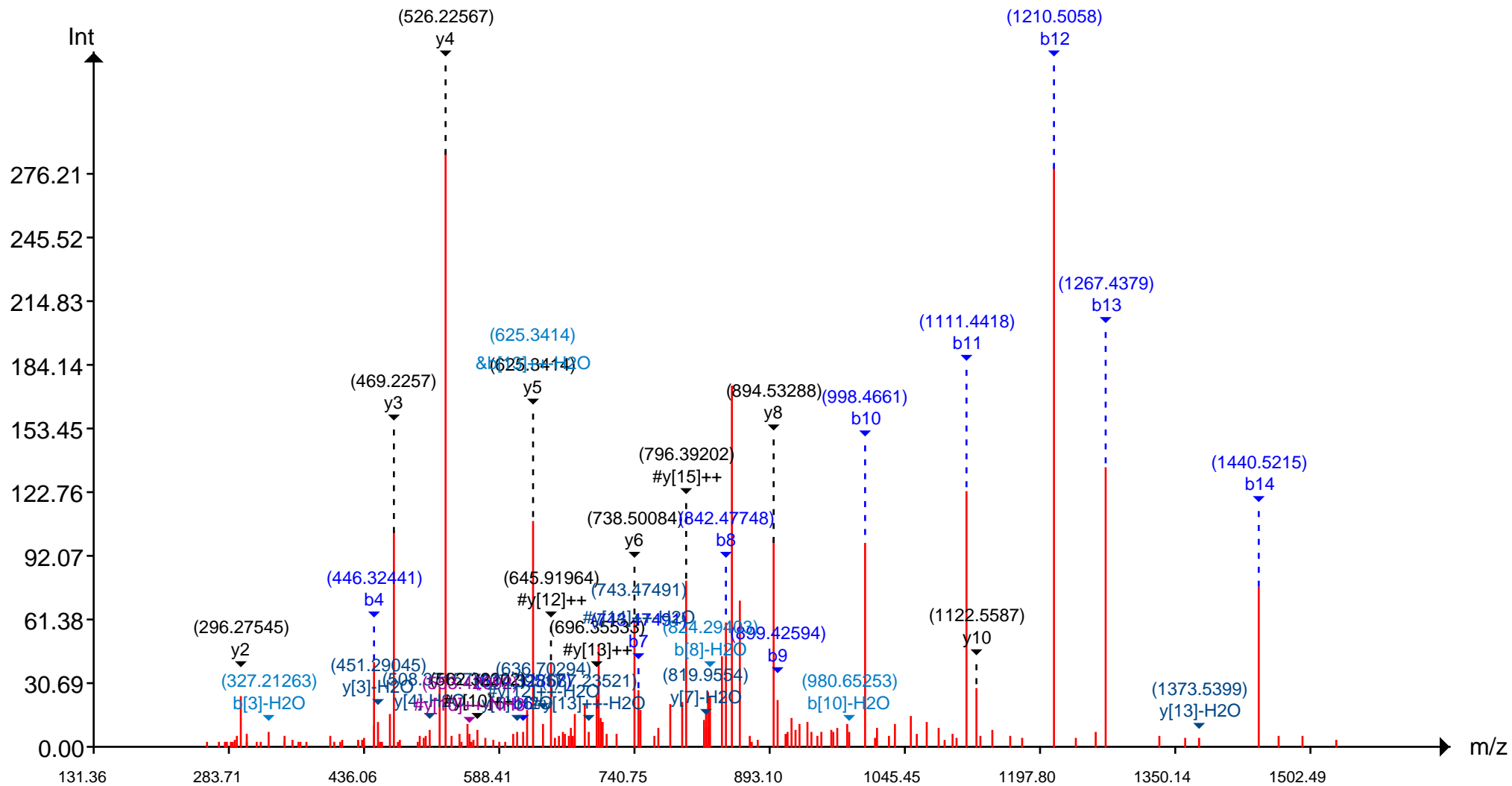


Info

Modified Sequence	AcD3-ADAINTEFK<AcD3K*>NTR<C13>-COOH	Ion score	53.41
Protein	P20152	Confidence level	99.00%
Precursor m/z	738.38832	Identity Threshold	48.15
Charge	2+	Mass error	-9.12E-4



Info			
Modified Sequence	AcD3-VSITPAEVLVVGK<AcD3K*>DR<C13>-COOH	Ion score	53.86
Protein	P62962	Confidence level	99.00%
Precursor m/z	868.51313	Identity Threshold	43.77
Charge	2+	Mass error	0.002293



Info

Modified Sequence AcD3-VSITPAEVGVLVGK<AcD3K*>DR<C13>-COOH

Ion score 57.6

Protein P62962

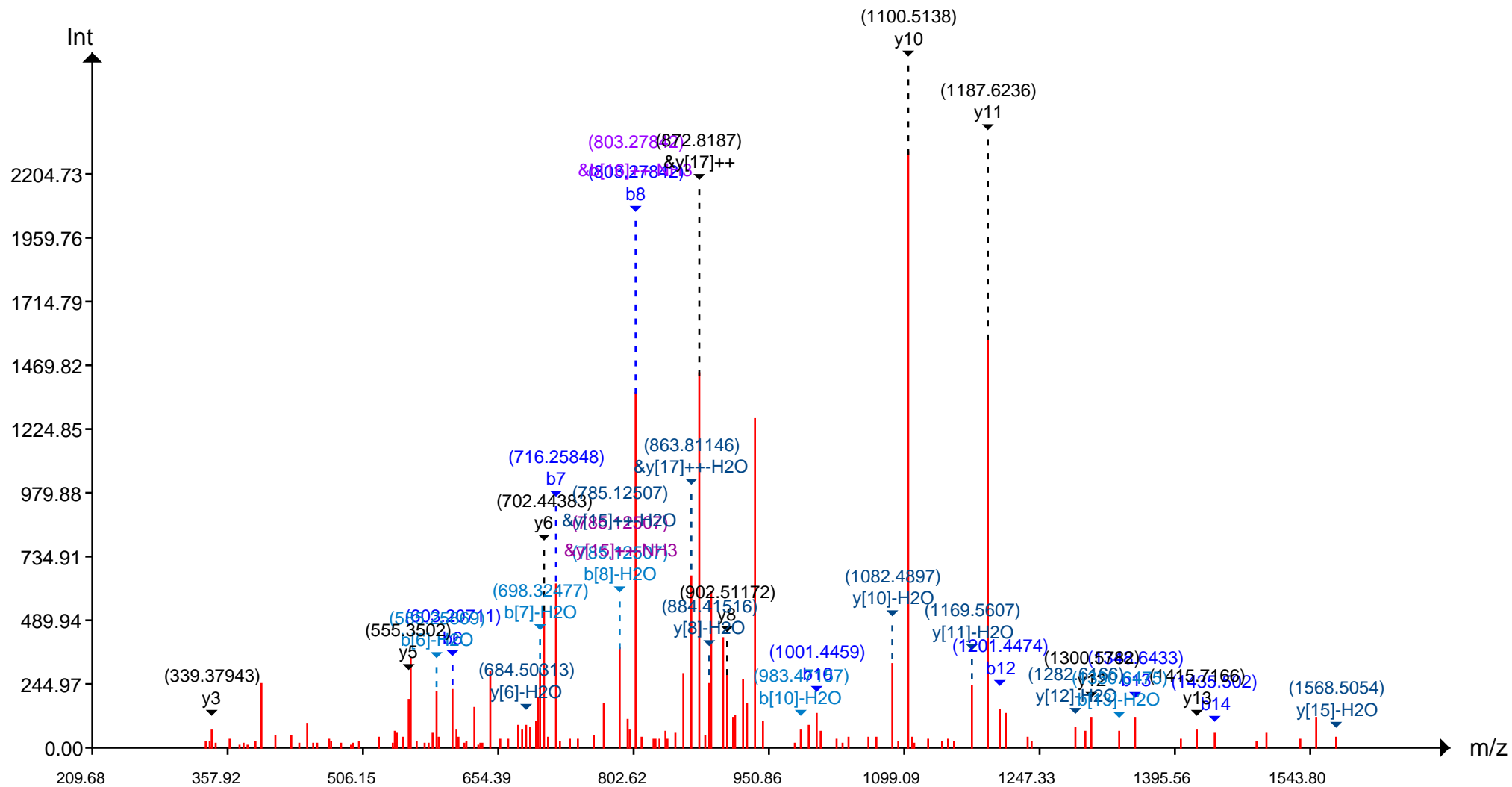
Confidence level 99.00%

Precursor m/z 868.51316

Identity Threshold 43.77

Charge 2+

Mass error 0.002353



Info

Modified Sequence AcD3-LGTGNDLSPTTVM<Mox*>SEGTR<C13>-COOH

Ion score 49.15

Protein Q923D2

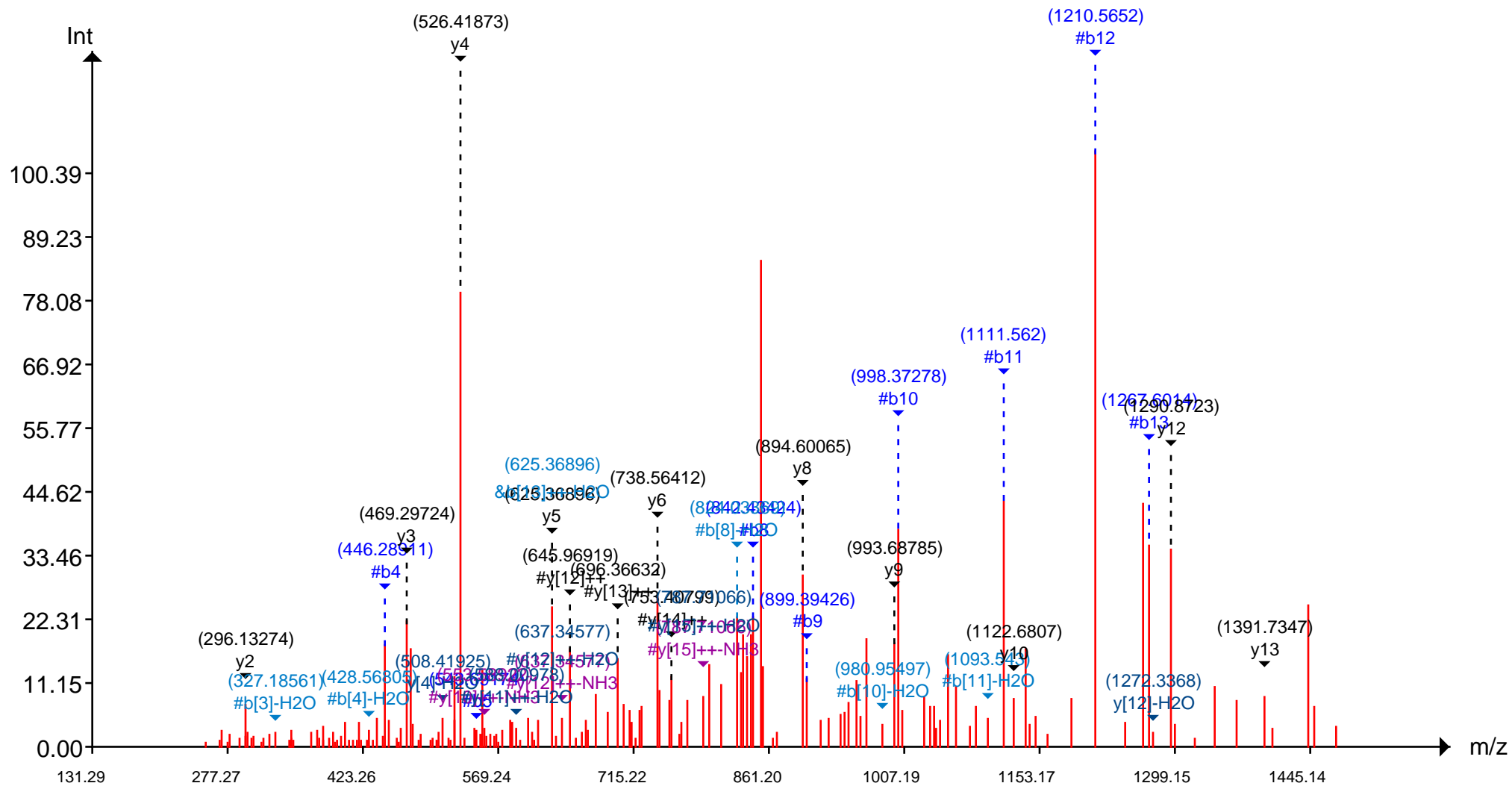
Confidence level 99.00%

Precursor m/z 951.96794

Identity Threshold 47.23

Charge 2+

Mass error 0.003771



Info

Modified Sequence AcD3-VSITPAEVLVVGK<AcD3K*>DR<C13>-COOH

Ion score 52.26

Protein P62962

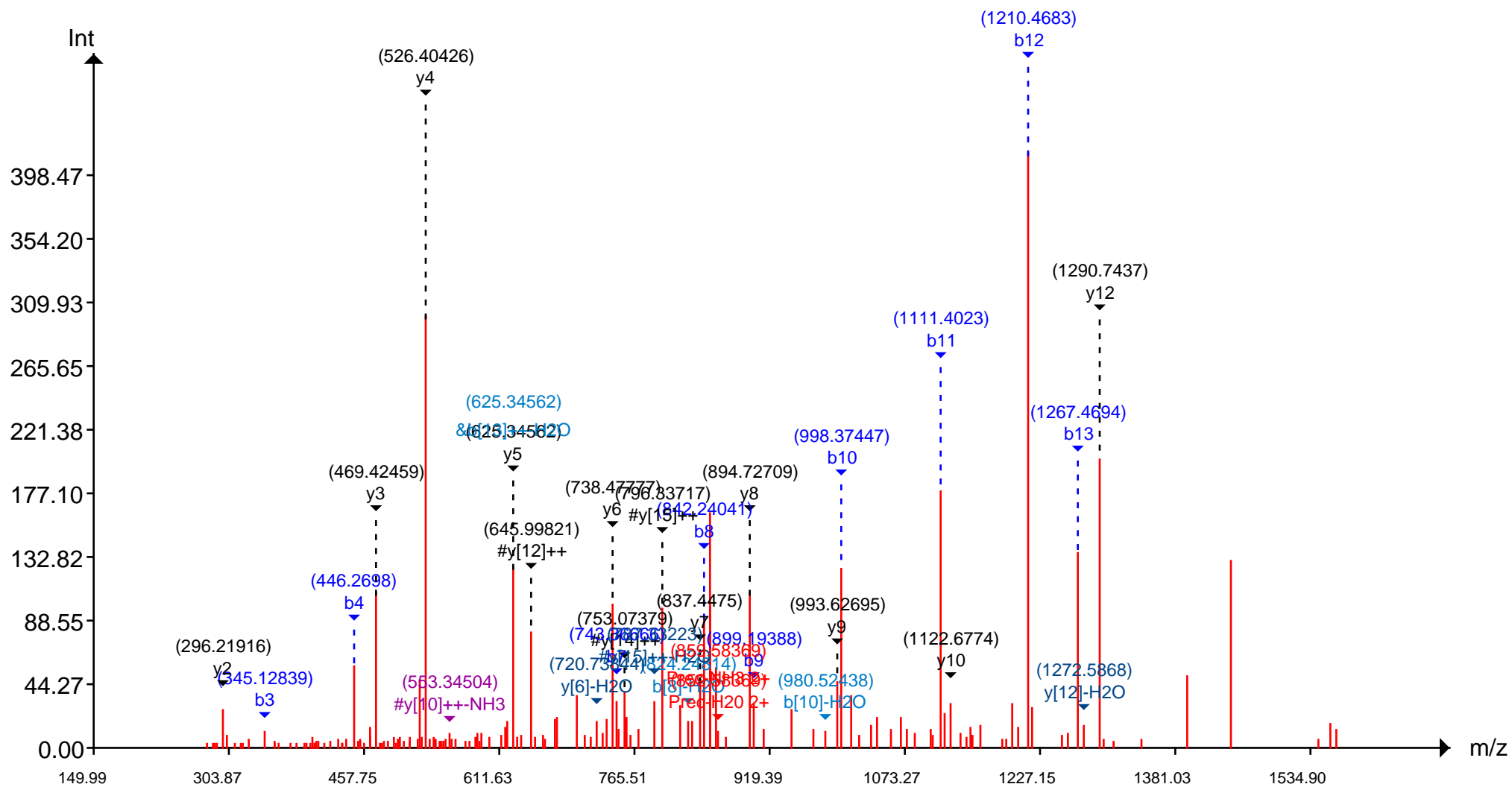
Confidence level 99.00%

Precursor m/z 868.51277

Identity Threshold 43.89

Charge 2+

Mass error 0.001573



Info

Modified Sequence AcD3-VSITPAEVLVVGK<AcD3K*>DR<C13>-COOH

Ion score 73.33

Protein P62962

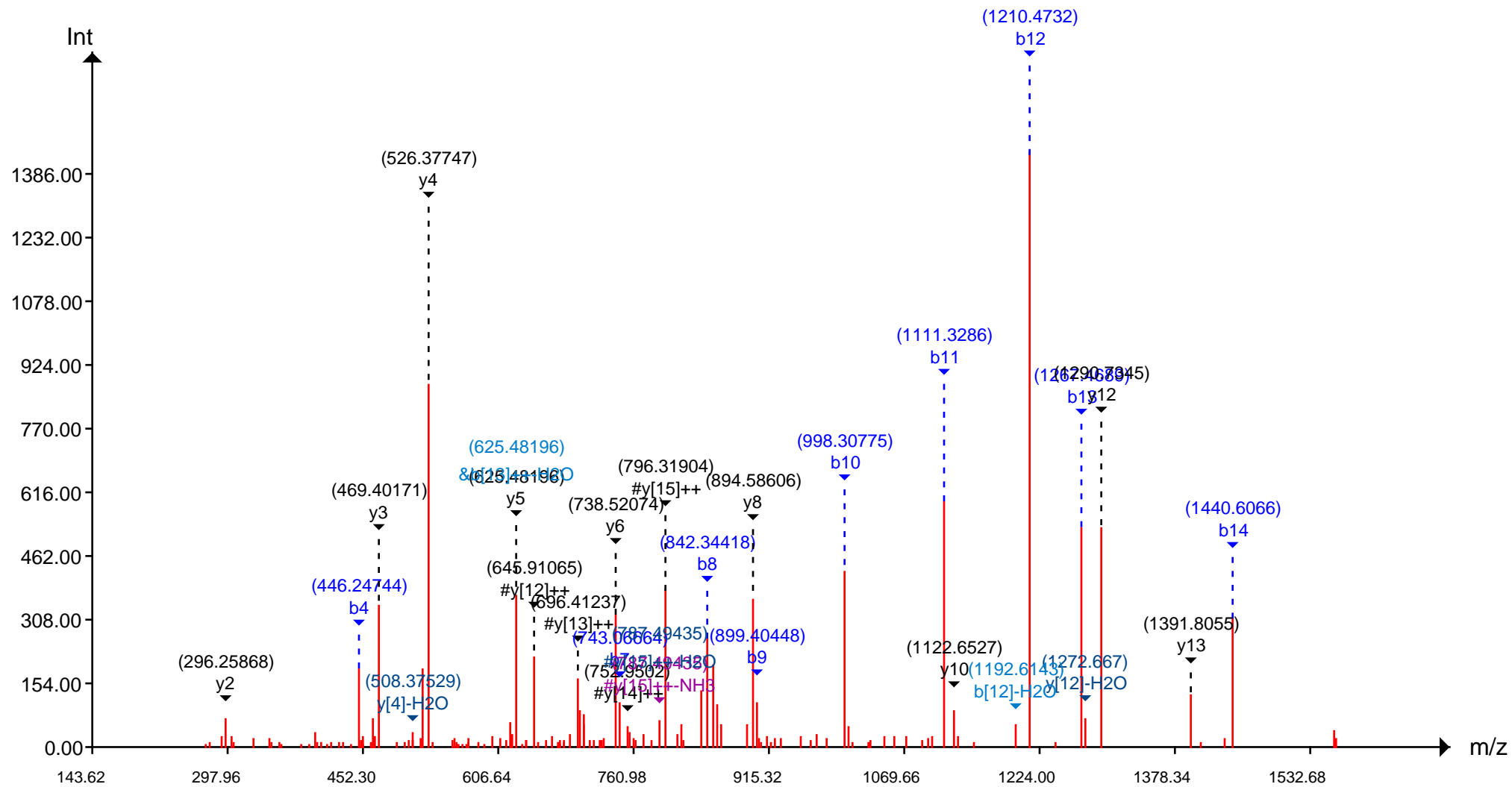
Confidence level 99.00%

Precursor m/z 868.51303

Identity Threshold 43.77

Charge 2+

Mass error 0.002093



Info

Modified Sequence AcD3-VSITPAEVGVLVGK<AcD3K*>DR<C13>-COOH

Ion score 79.07

Protein P62962

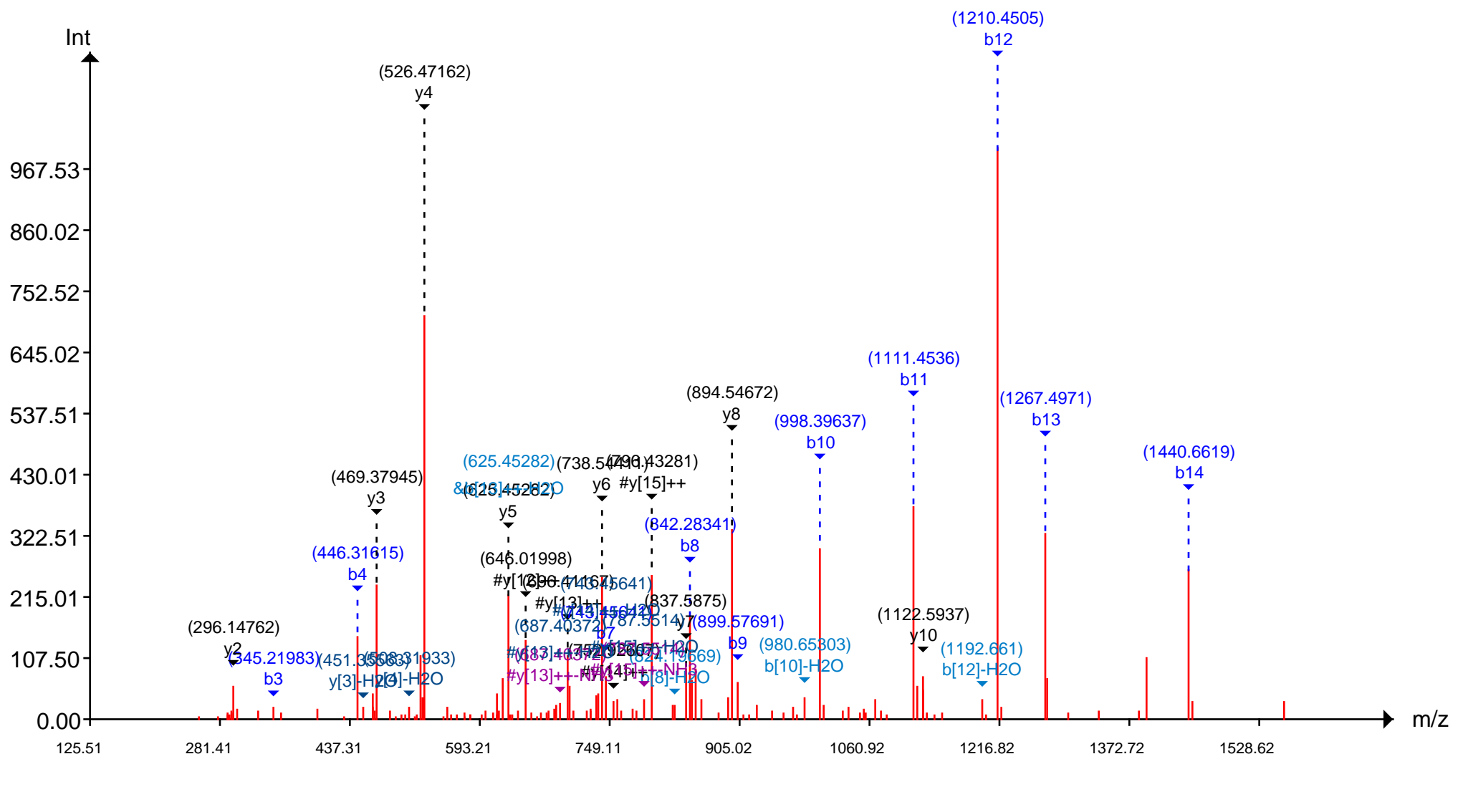
Confidence level 99.00%

Precursor m/z 868.51263

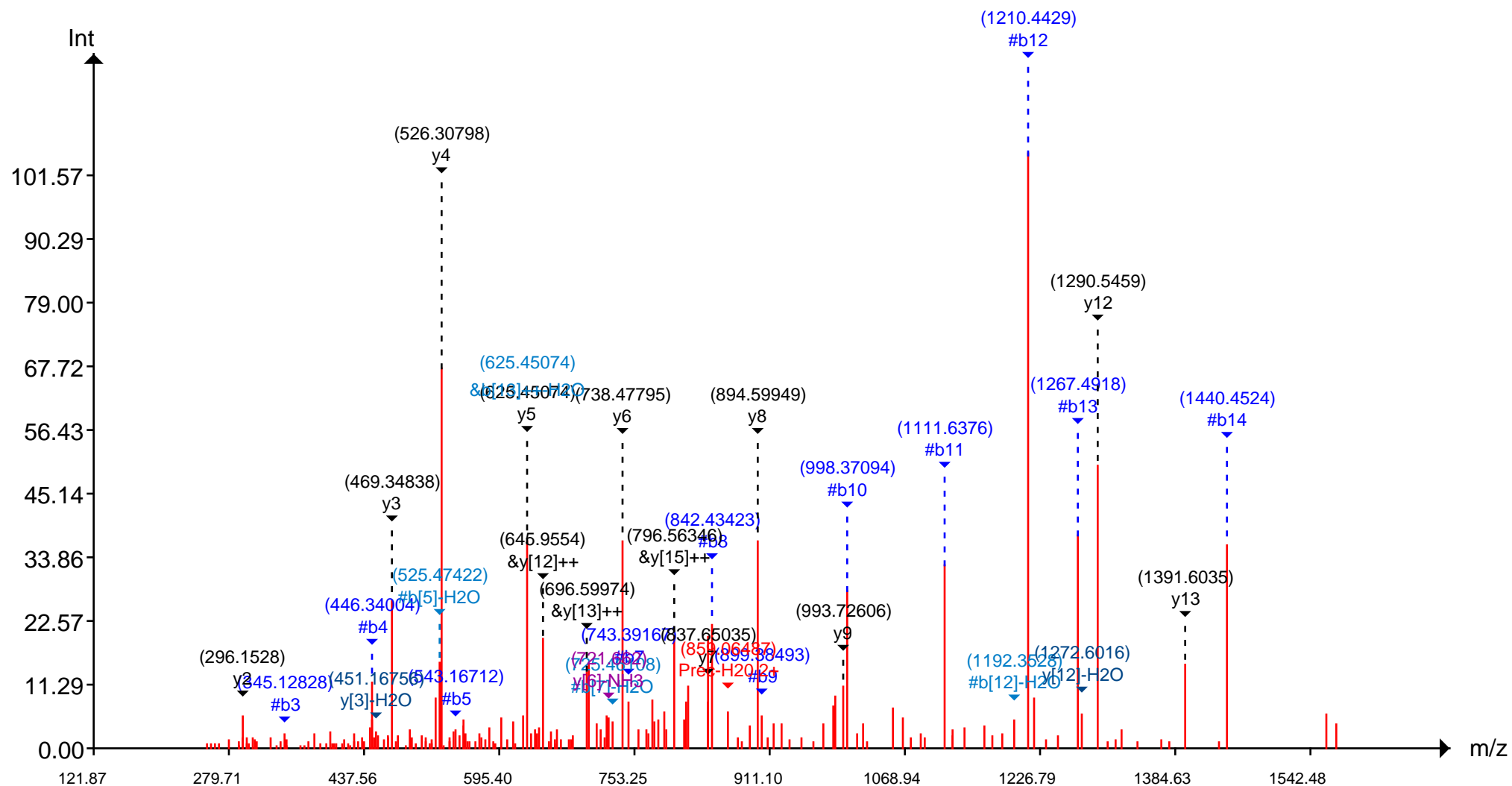
Identity Threshold 43.89

Charge 2+

Mass error 0.001293

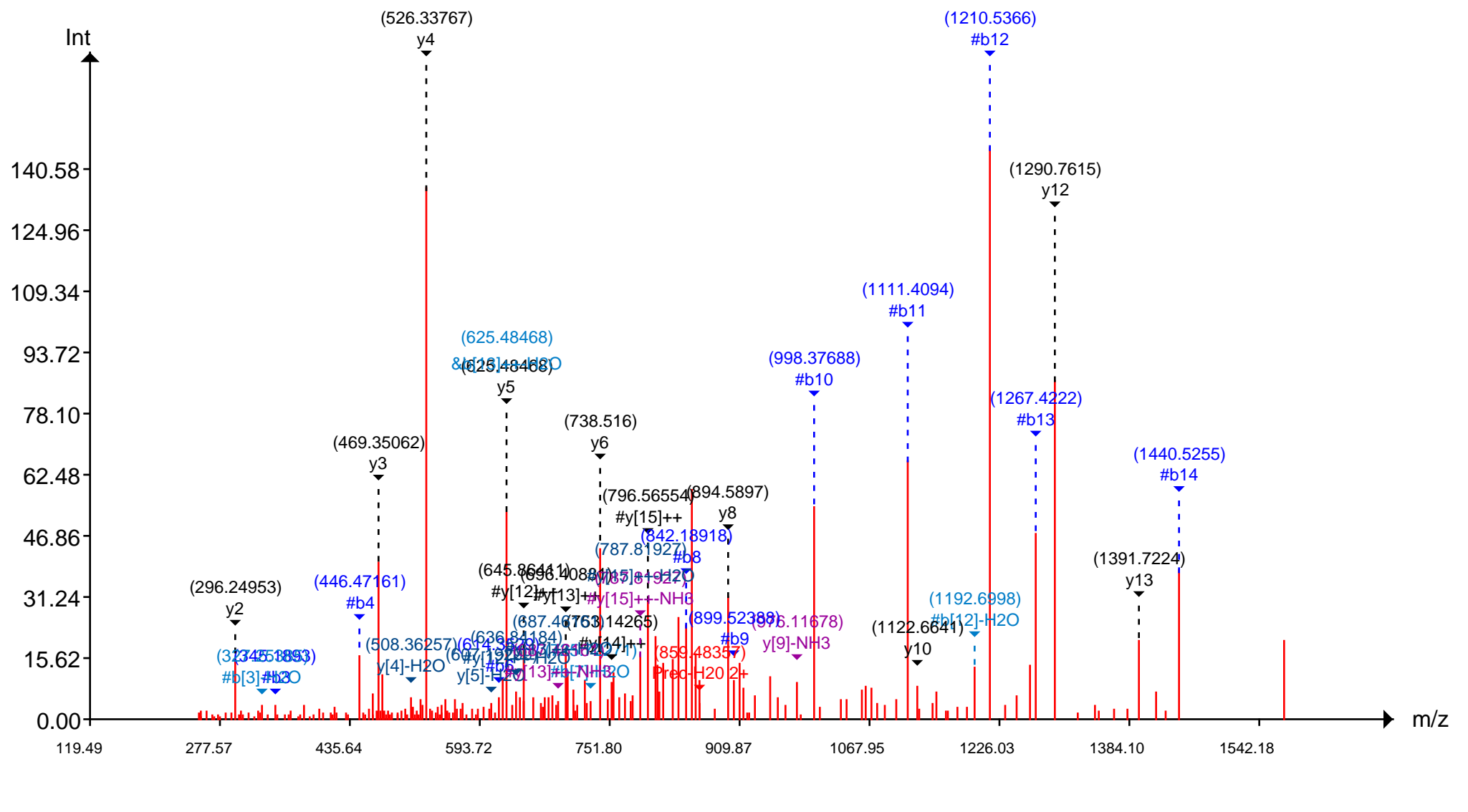


Info			
Modified Sequence	AcD3-VSITPAEVGVLVGK<AcD3K*>DR<C13>-COOH	Ion score	82.09
Protein	P62962	Confidence level	99.00%
Precursor m/z	868.5127	Identity Threshold	43.89
Charge	2+	Mass error	0.001433

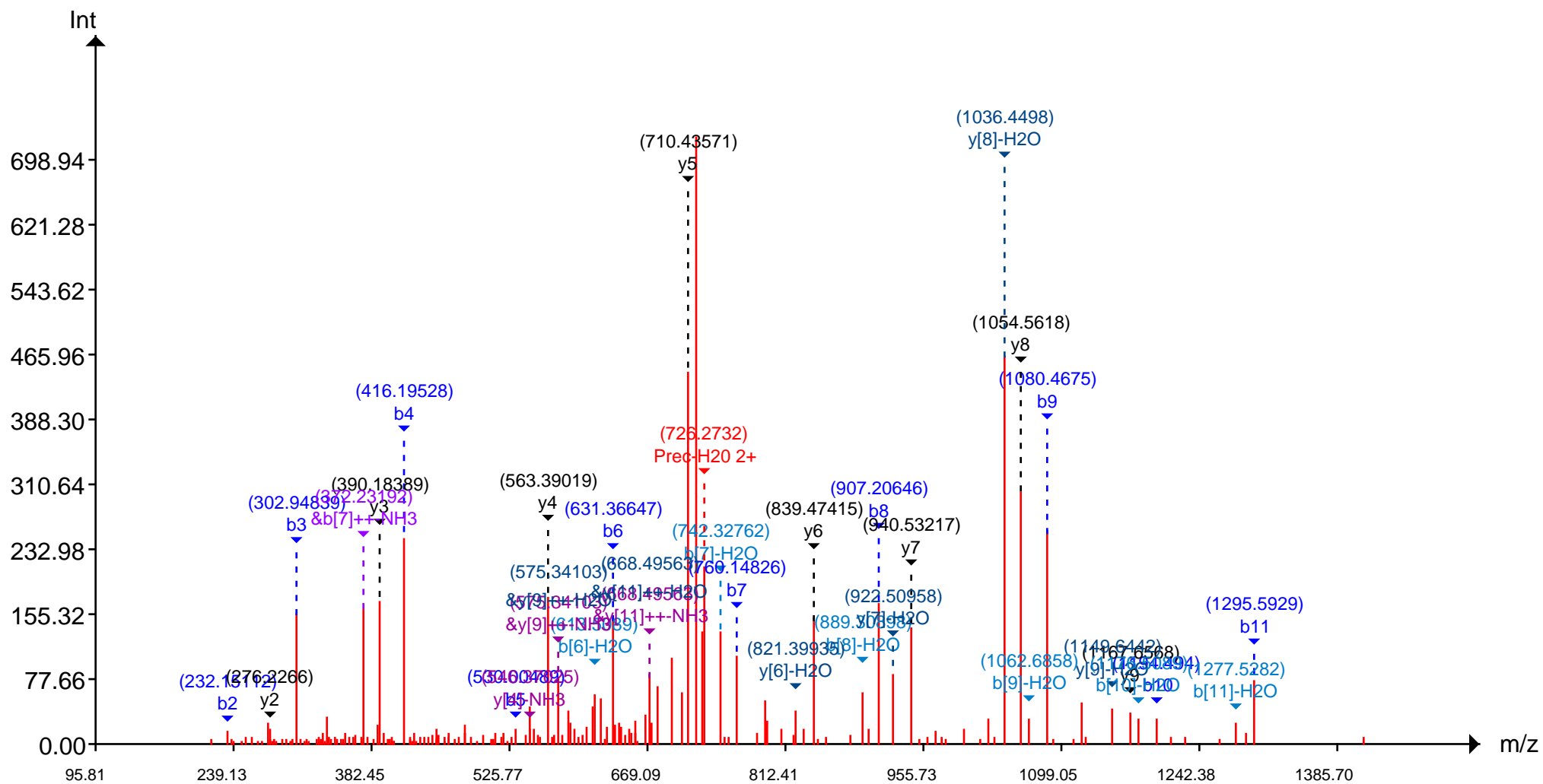


Info

Modified Sequence	AcD3-VSITPAEVGVLVGK<AcD3K*>DR<C13>-COOH	Ion score	64.43
Protein	P62962	Confidence level	99.00%
Precursor m/z	868.51245	Identity Threshold	43.93
Charge	2+	Mass error	9.33E-4

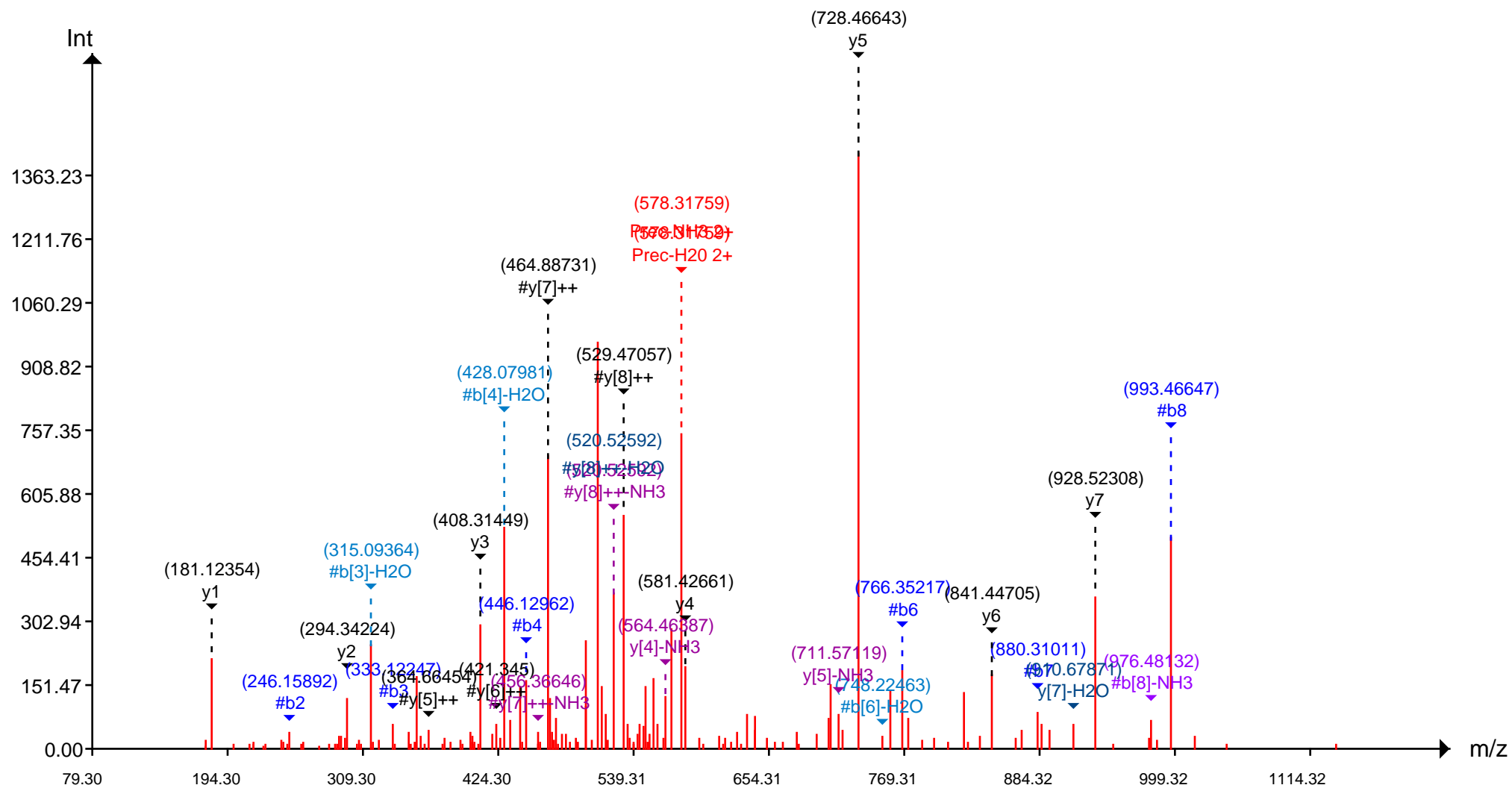


Info			
Modified Sequence	AcD3-VSITPAEVGVLVGK<AcD3K*>DR<C13>-COOH	Ion score	81.26
Protein	P62962	Confidence level	99.00%
Precursor m/z	868.51261	Identity Threshold	43.89
Charge	2+	Mass error	0.001253



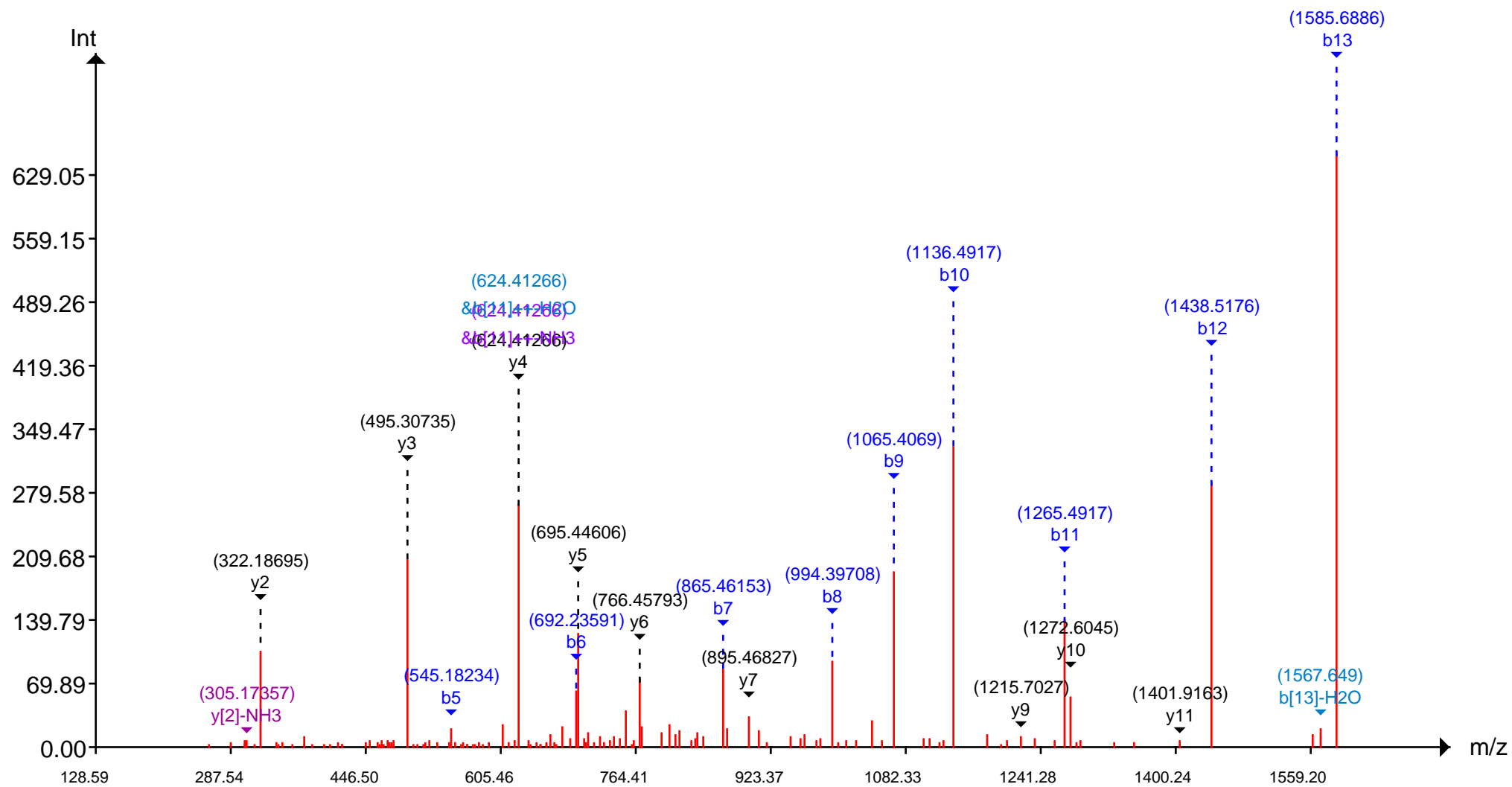
Info

Modified Sequence	AcD3-ADAINTEFK<AcD3K*>NTR-COOH	Ion score	59.97
Protein	P20152	Confidence level	99.00%
Precursor m/z	735.37949	Identity Threshold	47.83
Charge	2+	Mass error	0.001569



Info

Modified Sequence	AcD3-AESIM<Mox*>K<AcD3K*>NLR<C13>-COOH	Ion score	49.36
Protein	P06151	Confidence level	99.00%
Precursor m/z	587.32935	Identity Threshold	47.59
Charge	2+	Mass error	4.25E-4



Info

Modified Sequence AcD3-LVTEGFK<AcD3K*>EAAEK<AcD3K*>FR-COOH

Ion score 88.04

Protein Q9D7M1

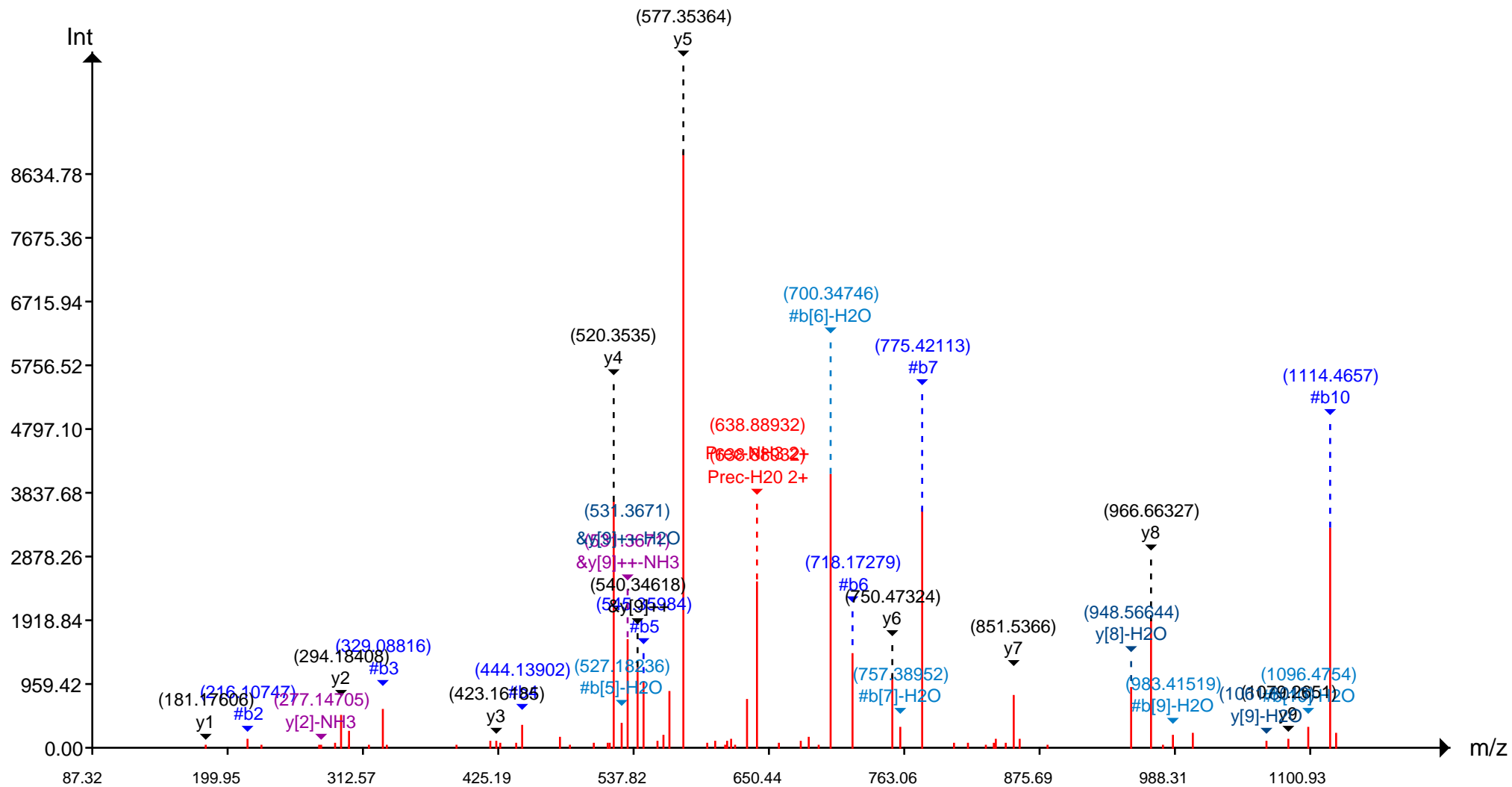
Confidence level 99.00%

Precursor m/z 880.48399

Identity Threshold 47.47

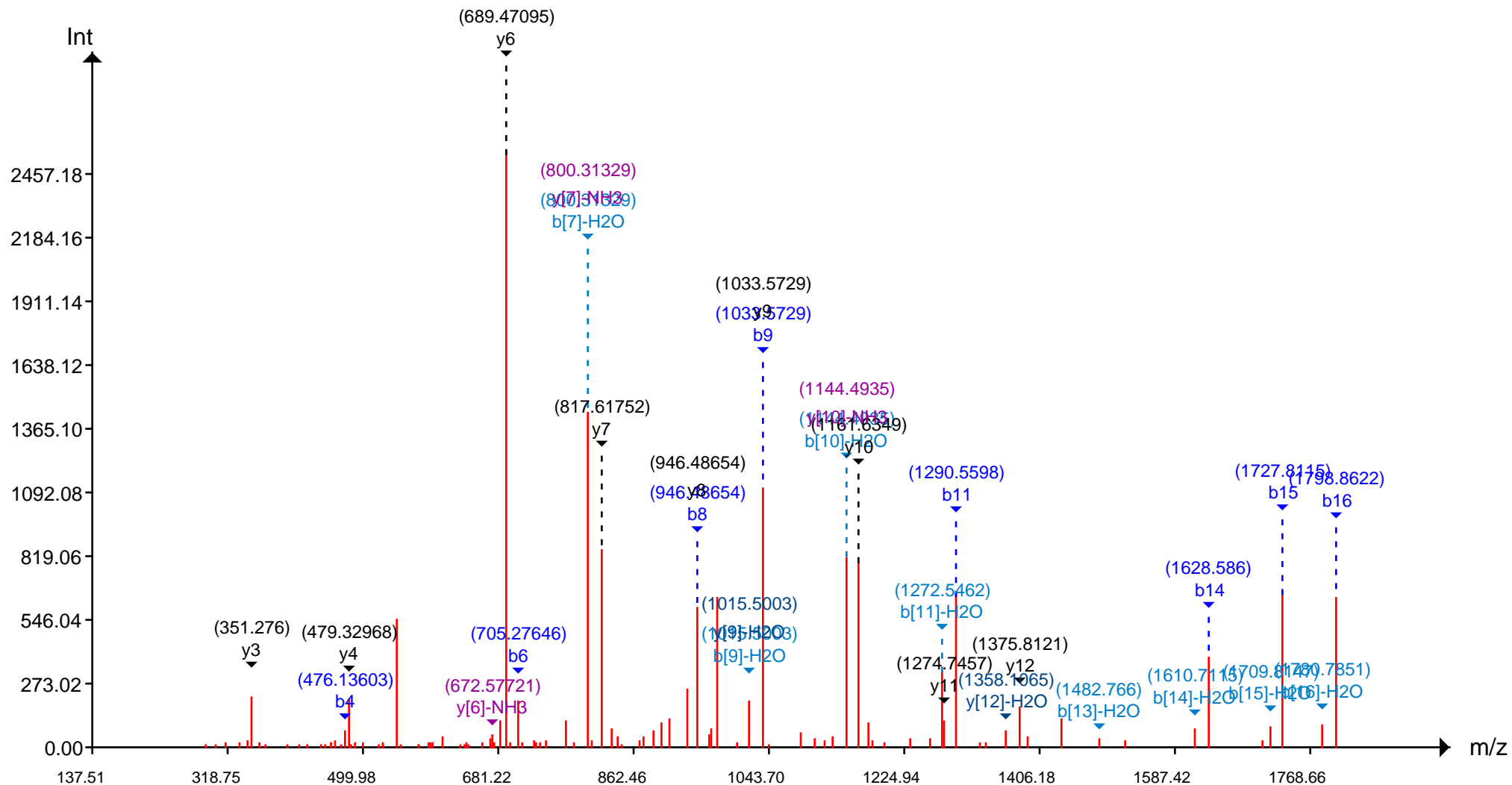
Charge 2+

Mass error 0.003202



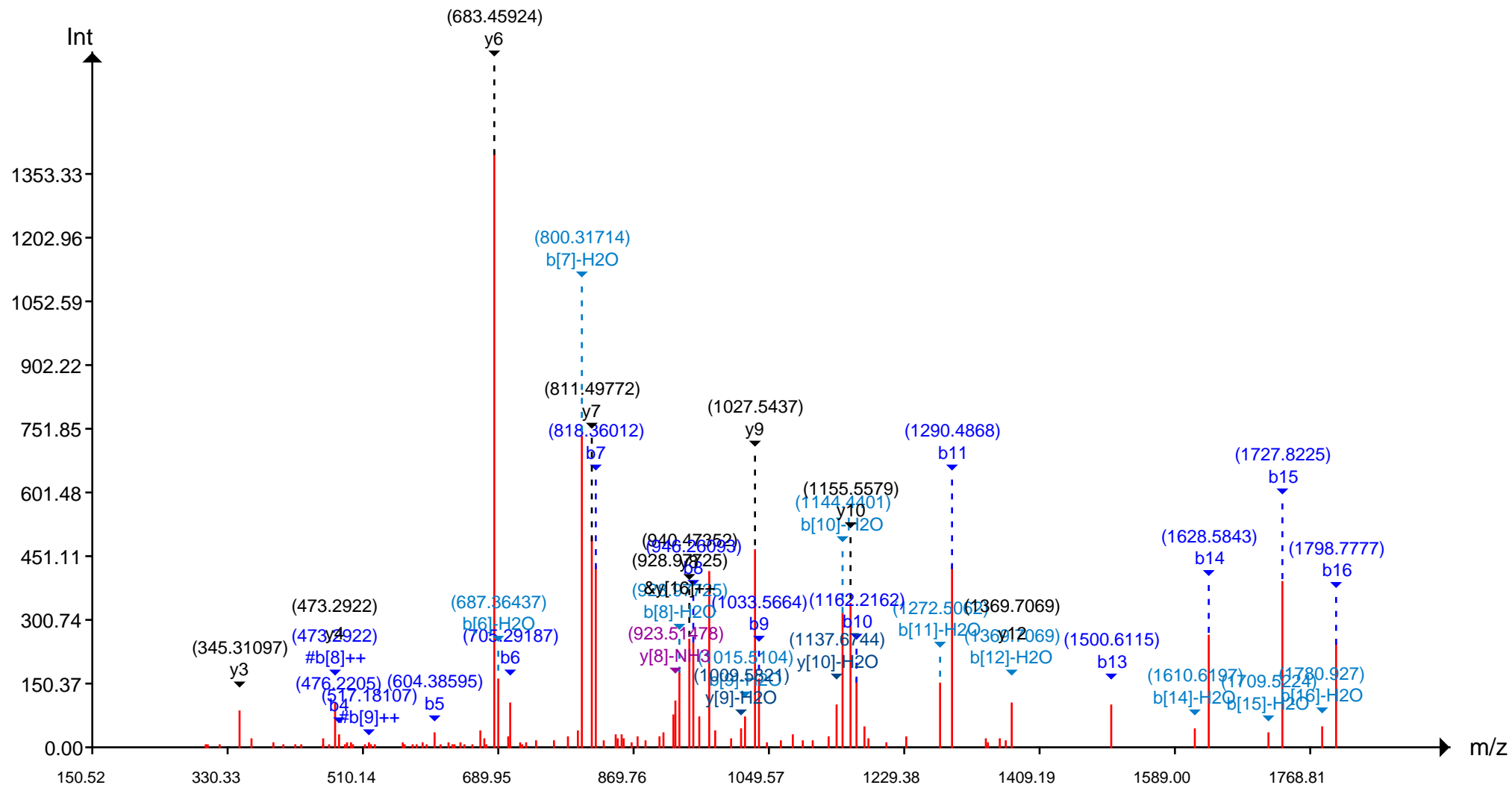
Info

Modified Sequence	AcD3-VALDTK<AcD3K*>GPEIR<C13>-COOH	Ion score	71.32
Protein	P52480	Confidence level	99.00%
Precursor m/z	647.88252	Identity Threshold	45.38
Charge	2+	Mass error	-1.68E-4



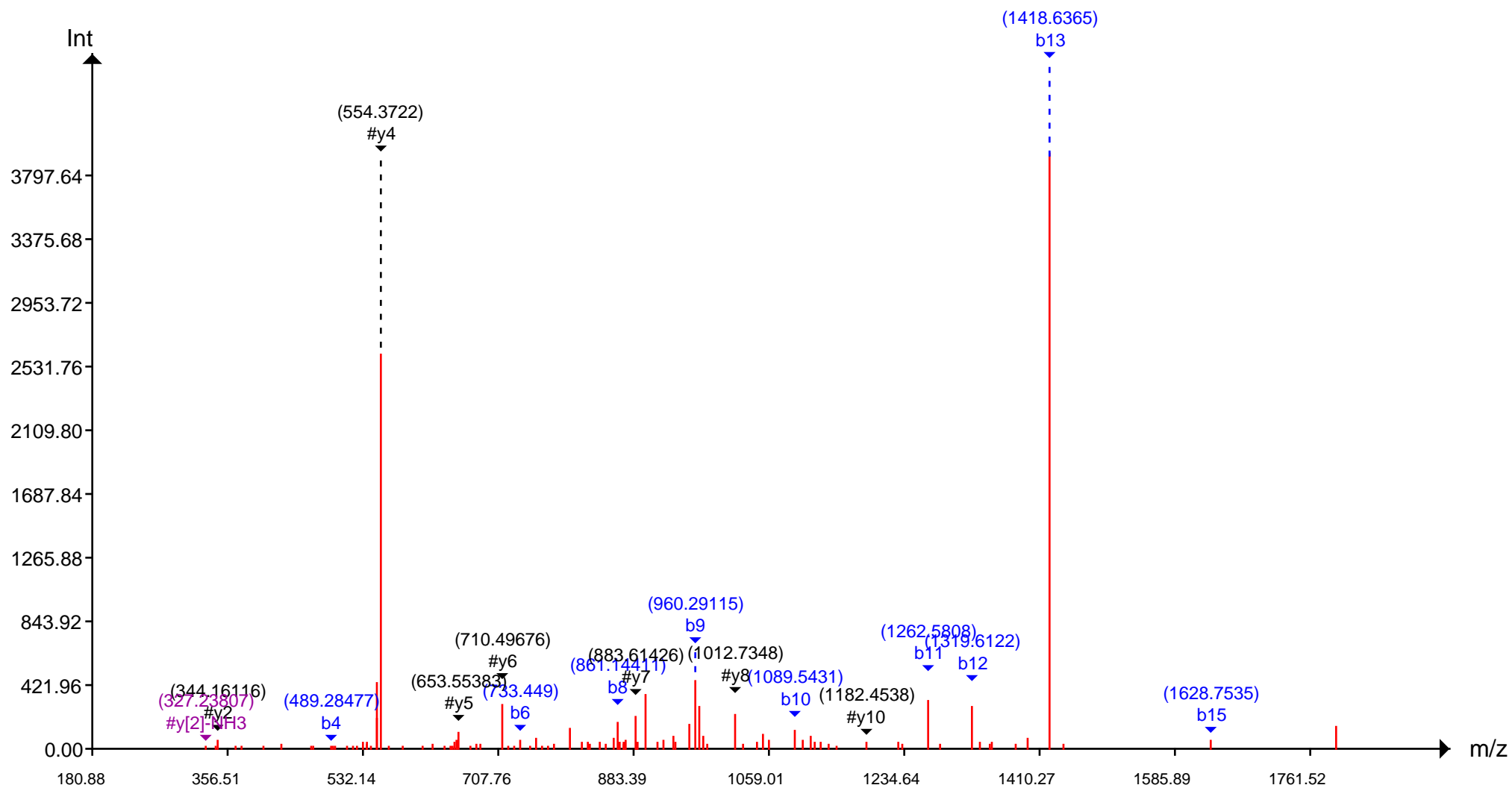
Info

Modified Sequence	AcD3-AADK<AcD3K*>QTLQSEQPLQVAR<C13>-COOH	Ion score	83.62
Protein	P46471	Confidence level	99.00%
Precursor m/z	990.04329	Identity Threshold	47.18
Charge	2+	Mass error	0.00228



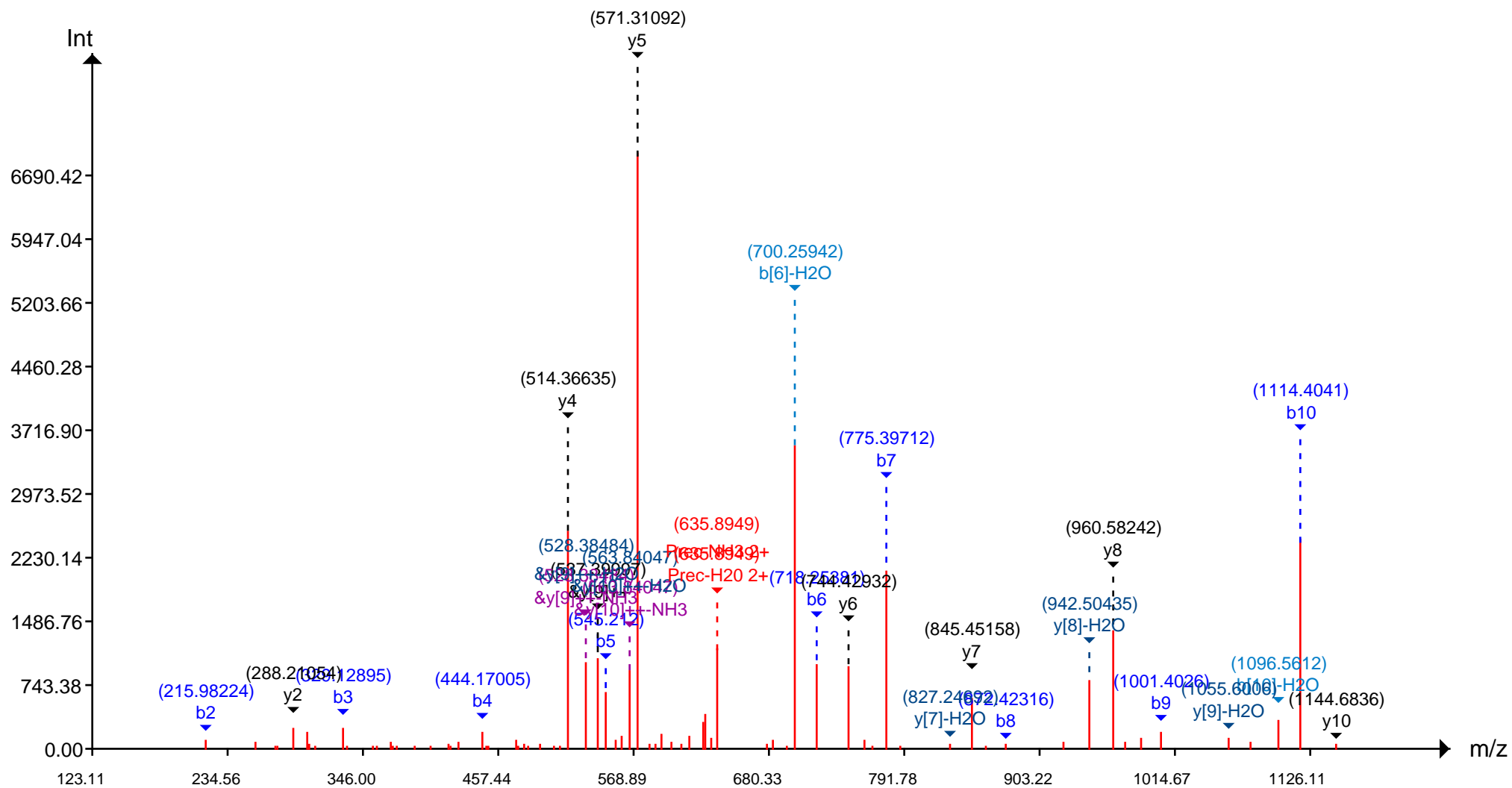
Info

Modified Sequence	AcD3-AADK<AcD3K*>QTLQSEQPLQVAR-COOH	Ion score	93.5
Protein	P46471	Confidence level	99.00%
Precursor m/z	987.03361	Identity Threshold	47.61
Charge	2+	Mass error	0.003062



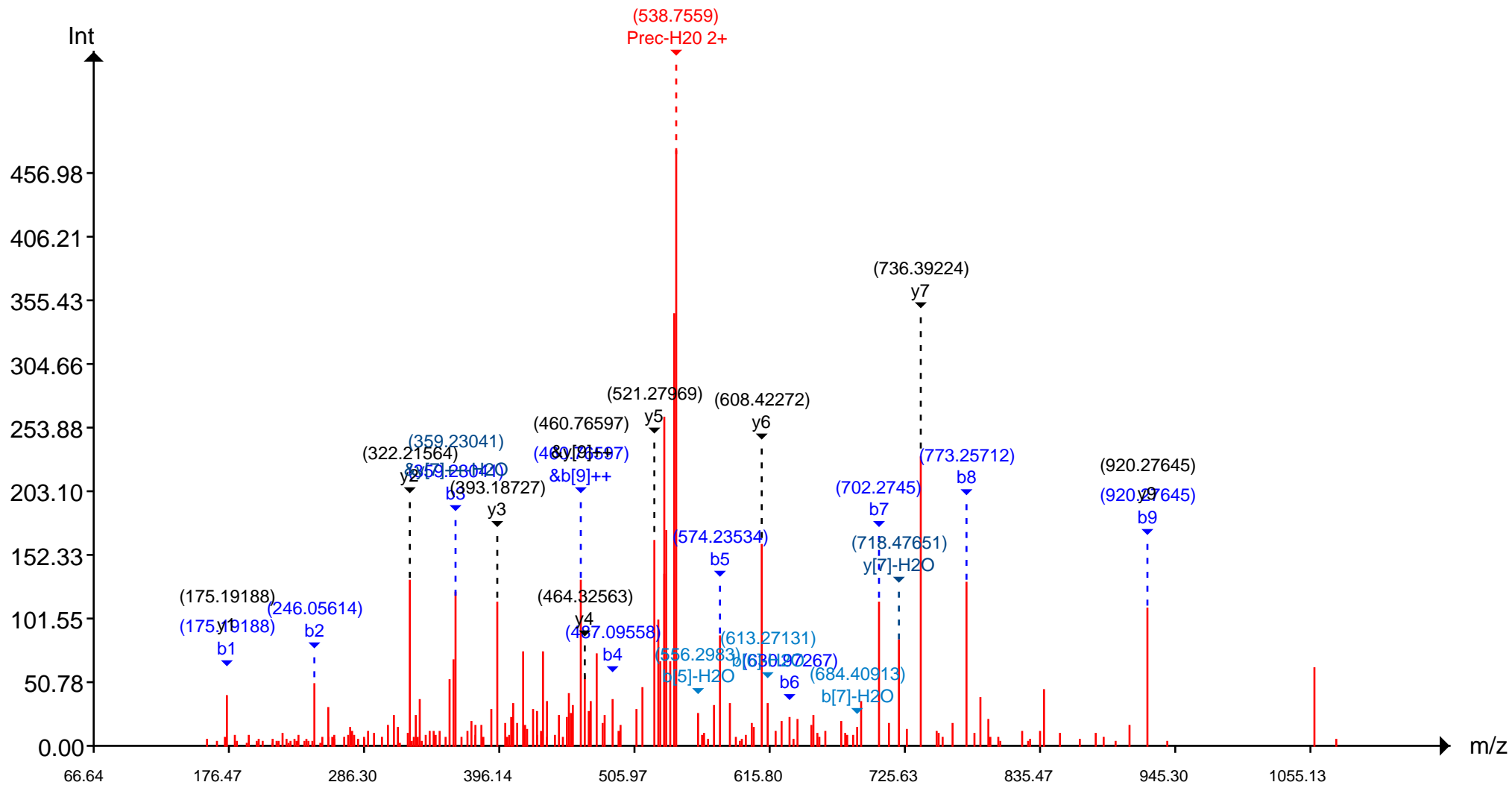
Info

Modified Sequence	AcD3-LAVC<Cmm*>K<AcD3K*>AGAVEK<AcD3K*>GVPLYR<C13>-COOH	Ion score	48.76
Protein	P17182	Confidence level	99.00%
Precursor m/z	986.57188	Identity Threshold	44.33
Charge	2+	Mass error	0.002316



Info

Modified Sequence	AcD3-VALDTK<AcD3K*>GPEIR-COOH	Ion score	79.22
Protein	P52480	Confidence level	99.00%
Precursor m/z	644.87292	Identity Threshold	46.67
Charge	2+	Mass error	7.73E-4



Info

Modified Sequence AcD3-EALQSGAAFR-COOH

Ion score 53.98

Protein O08808, O70566, Q9Z207

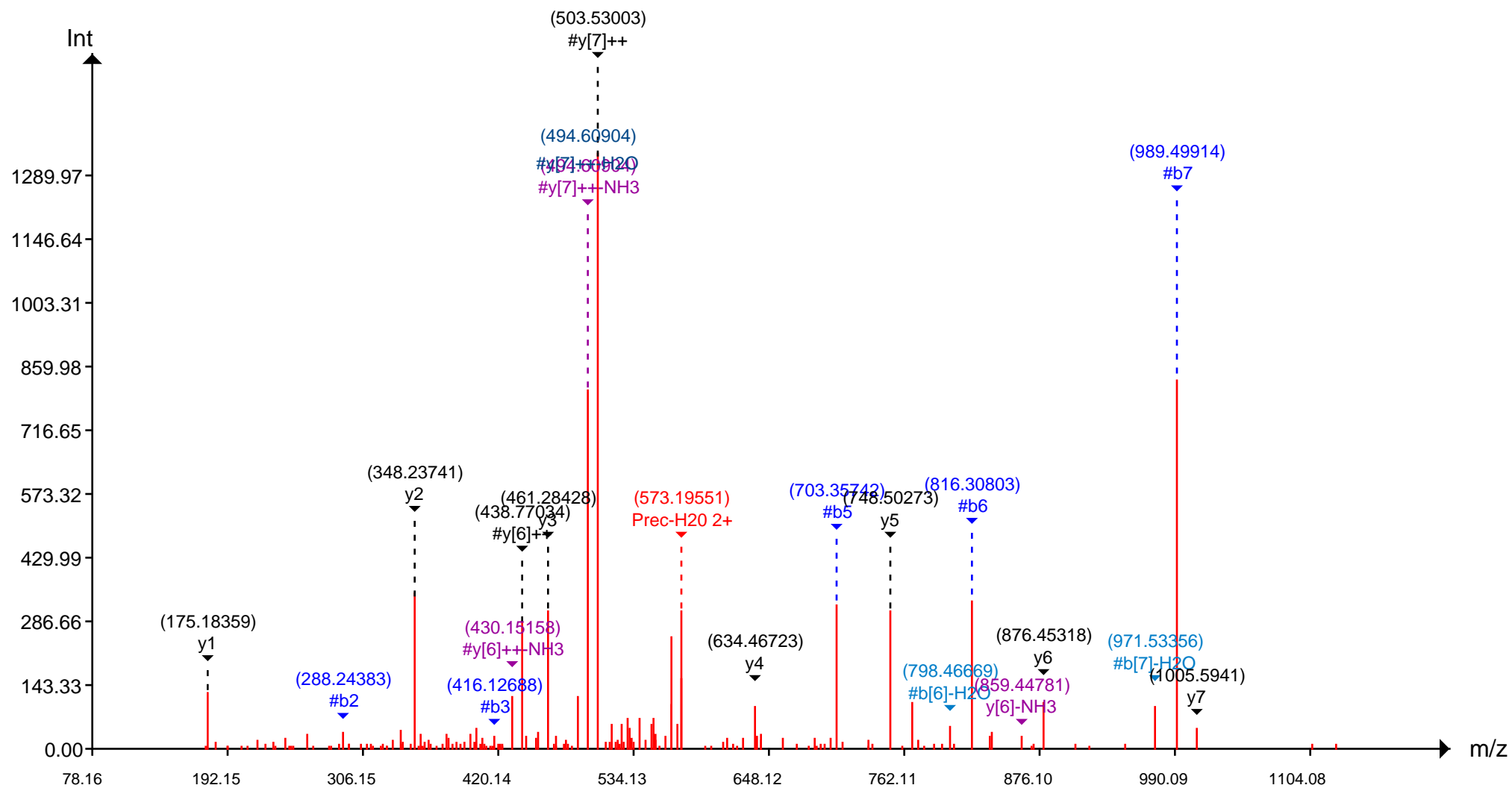
Confidence level 99.00%

Precursor m/z 547.78694

Identity Threshold 47.73

Charge 2+

Mass error -1.97E-4



Info

Modified Sequence AcD3-IEQNK<AcD3K*>LK<AcD3K*>R-COOH

Ion score 58.95

Protein Q8K2Q9

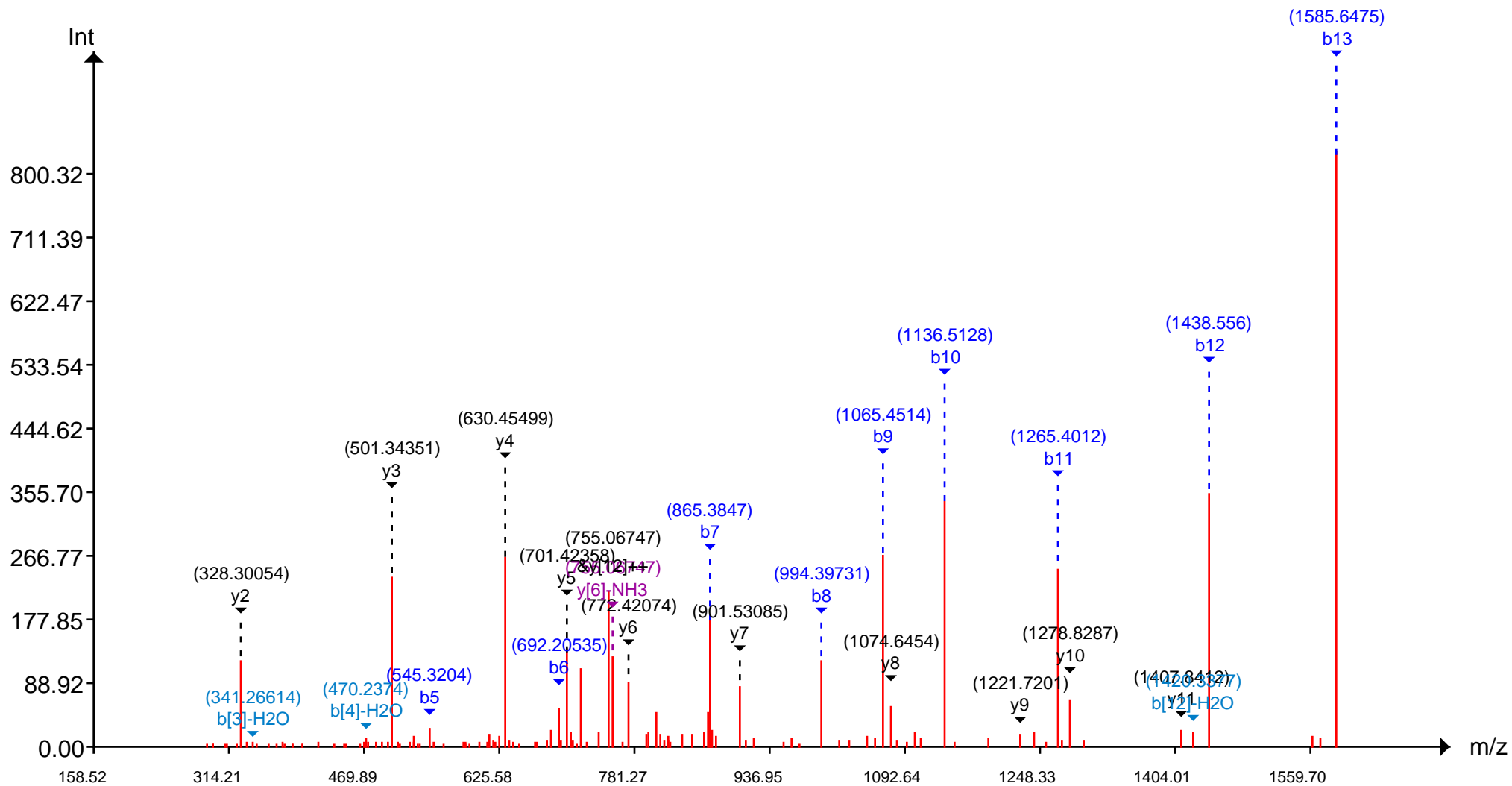
Confidence level 99.00%

Precursor m/z 582.35854

Identity Threshold 43.63

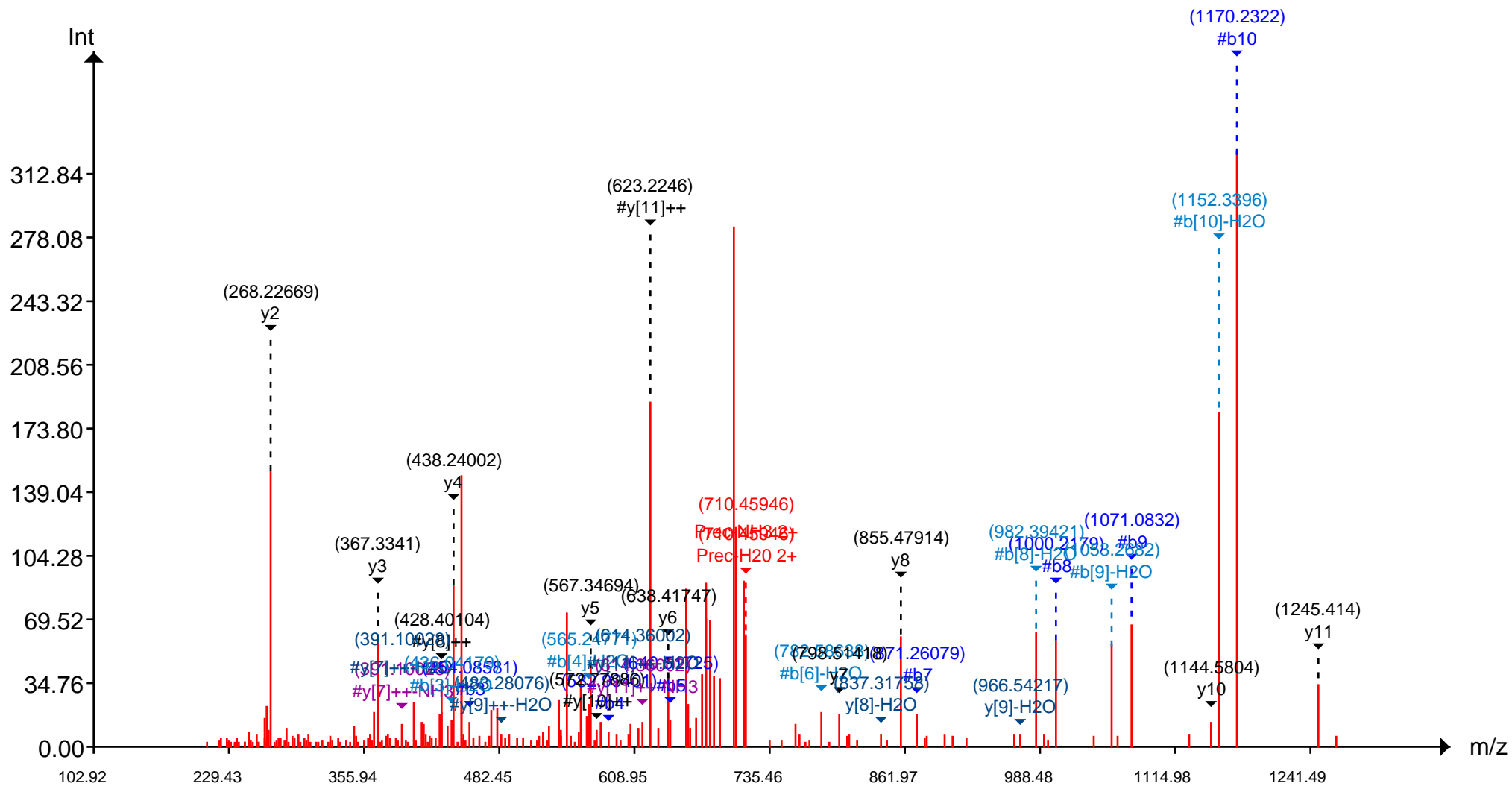
Charge 2+

Mass error 5.47E-4



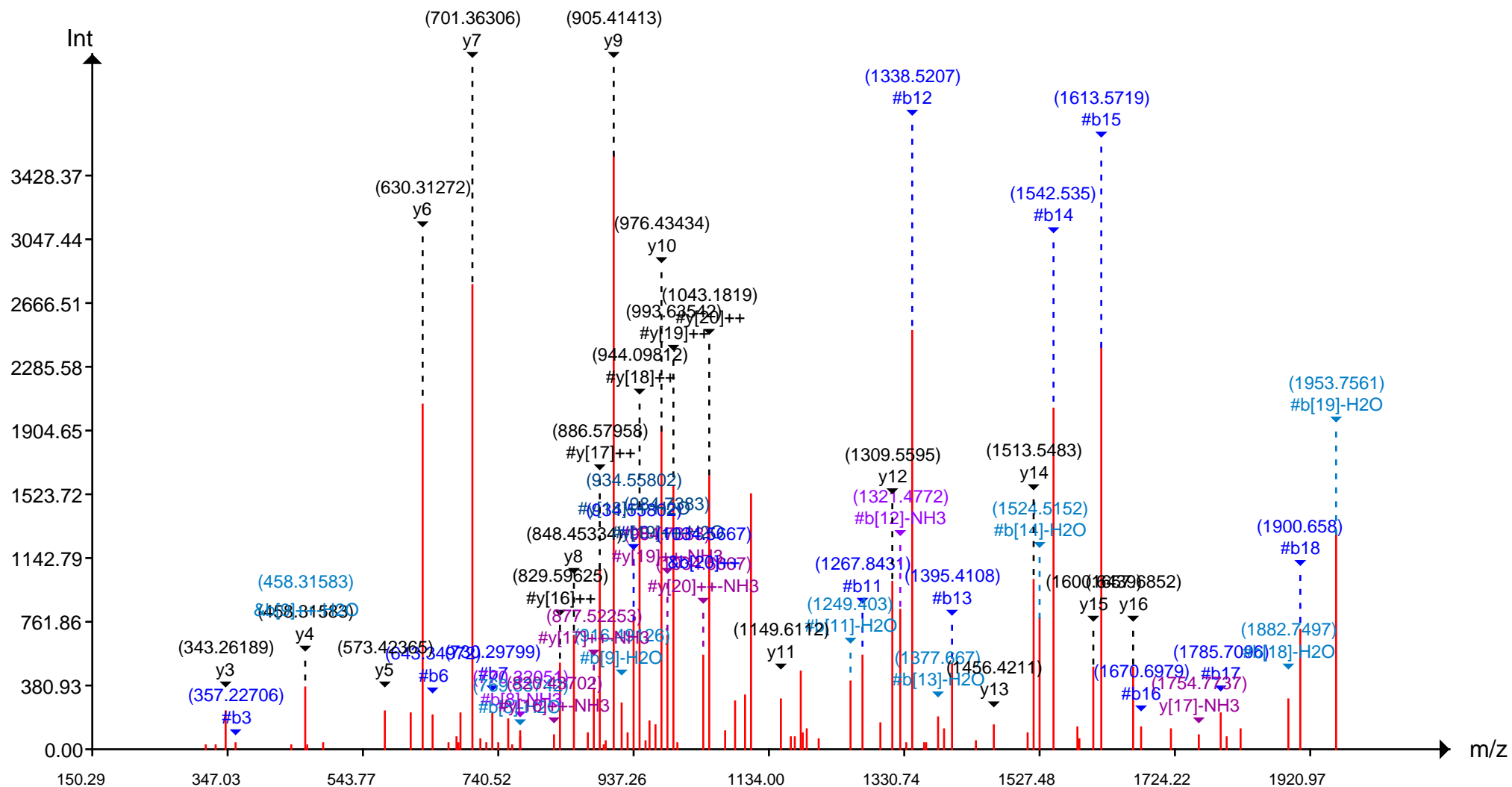
Info

Modified Sequence	AcD3-LVTEGFK<AcD3K*>EAAEK<AcD3K*>FR<C13>-COOH	Ion score	65.79
Protein	Q9D7M1	Confidence level	99.00%
Precursor m/z	883.49387	Identity Threshold	46.90
Charge	2+	Mass error	0.002821



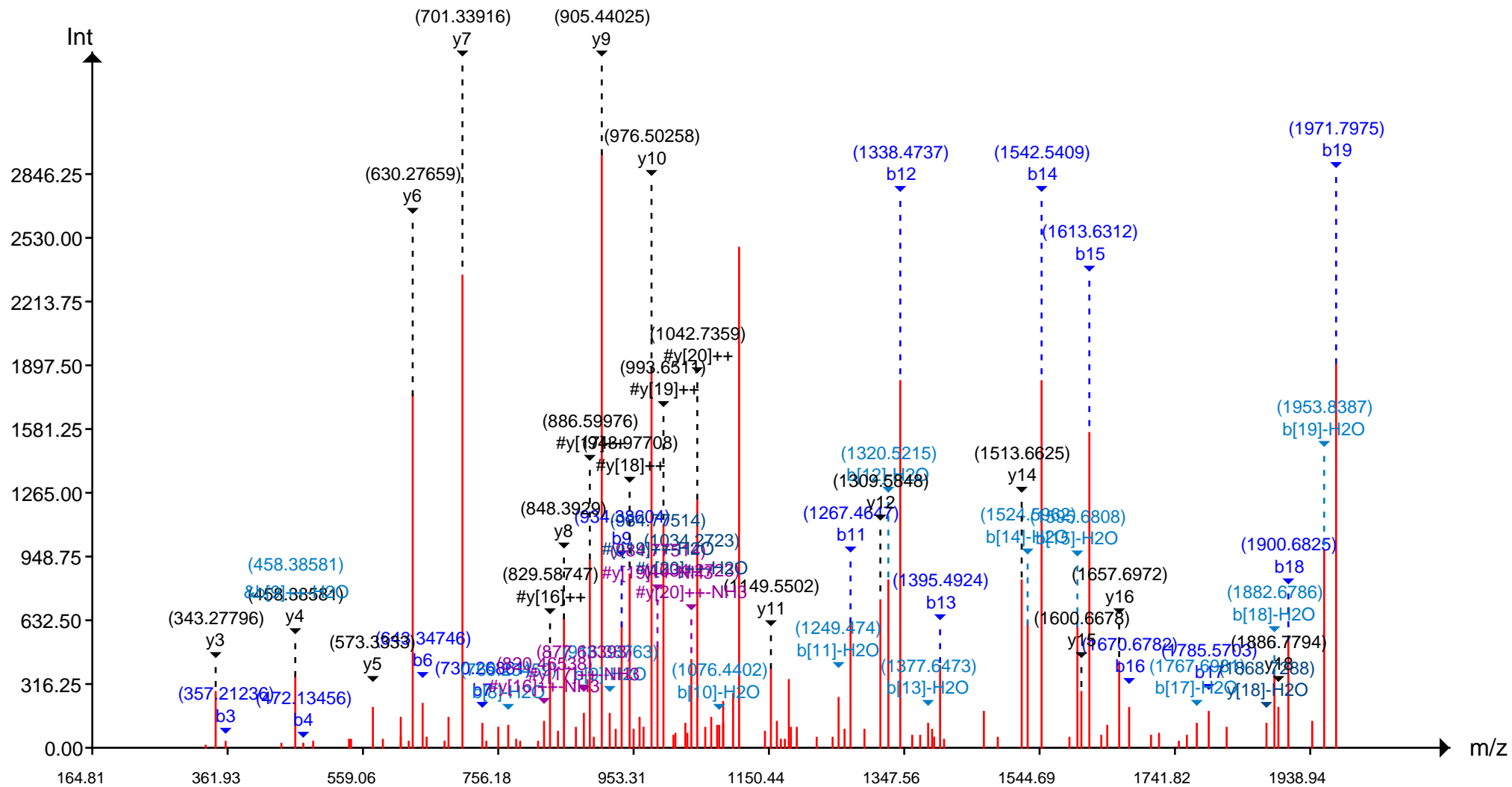
Info

Modified Sequence	AcD3-M<Mox*>TC<Cmm*>EGC<Cmm*>AEAVSR<C13>-COOH	Ion score	44.74
Protein	O08997	Confidence level	99.00%
Precursor m/z	719.30093	Identity Threshold	39.98
Charge	2+	Mass error	4.09E-4



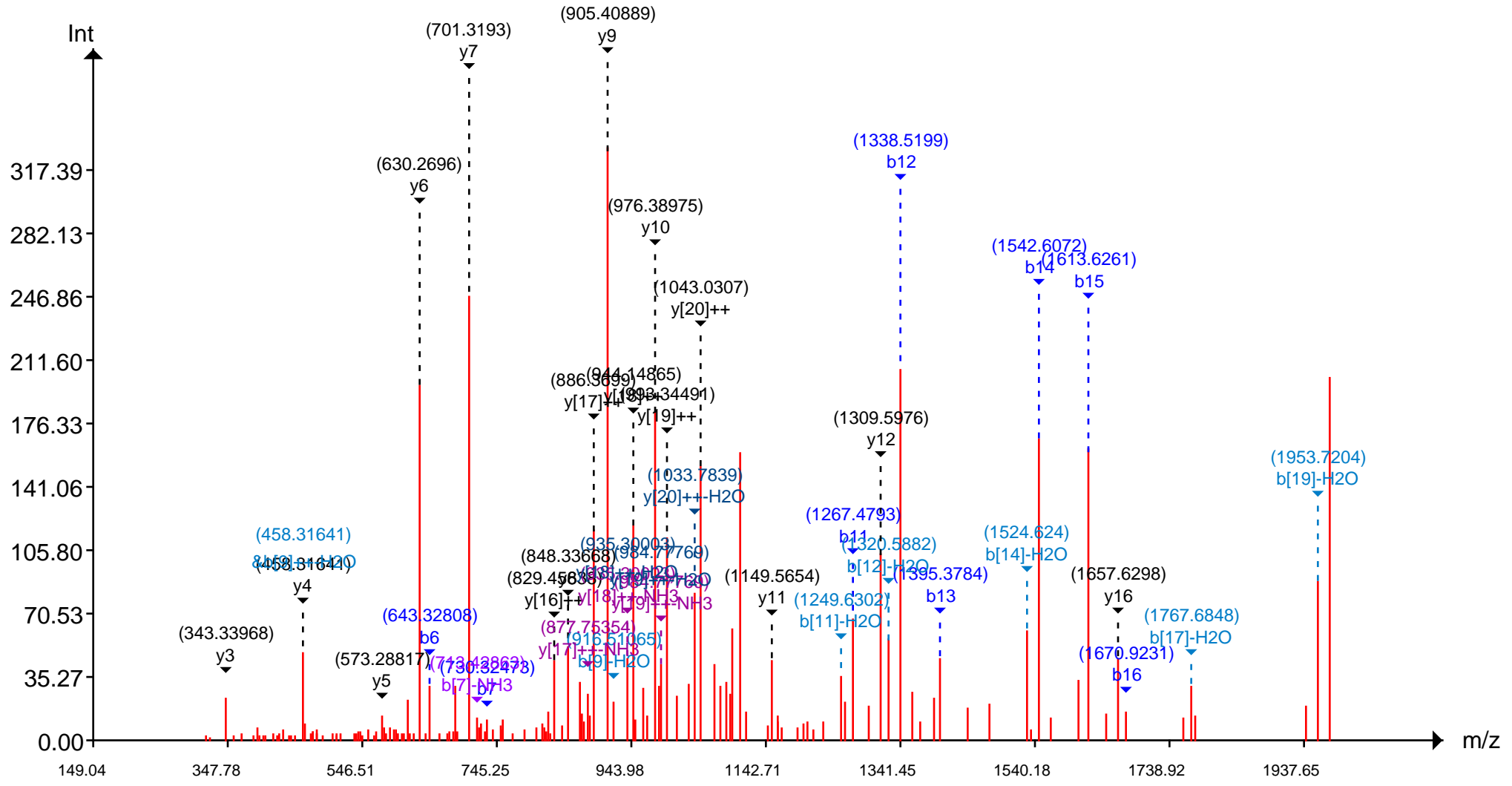
Info

Modified Sequence	AcD3-LVVDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR-COOH	Ion score	111.69
Protein	P60710	Confidence level	99.00%
Precursor m/z	1122.0215	Identity Threshold	45.09
Charge	2+	Mass error	0.001898



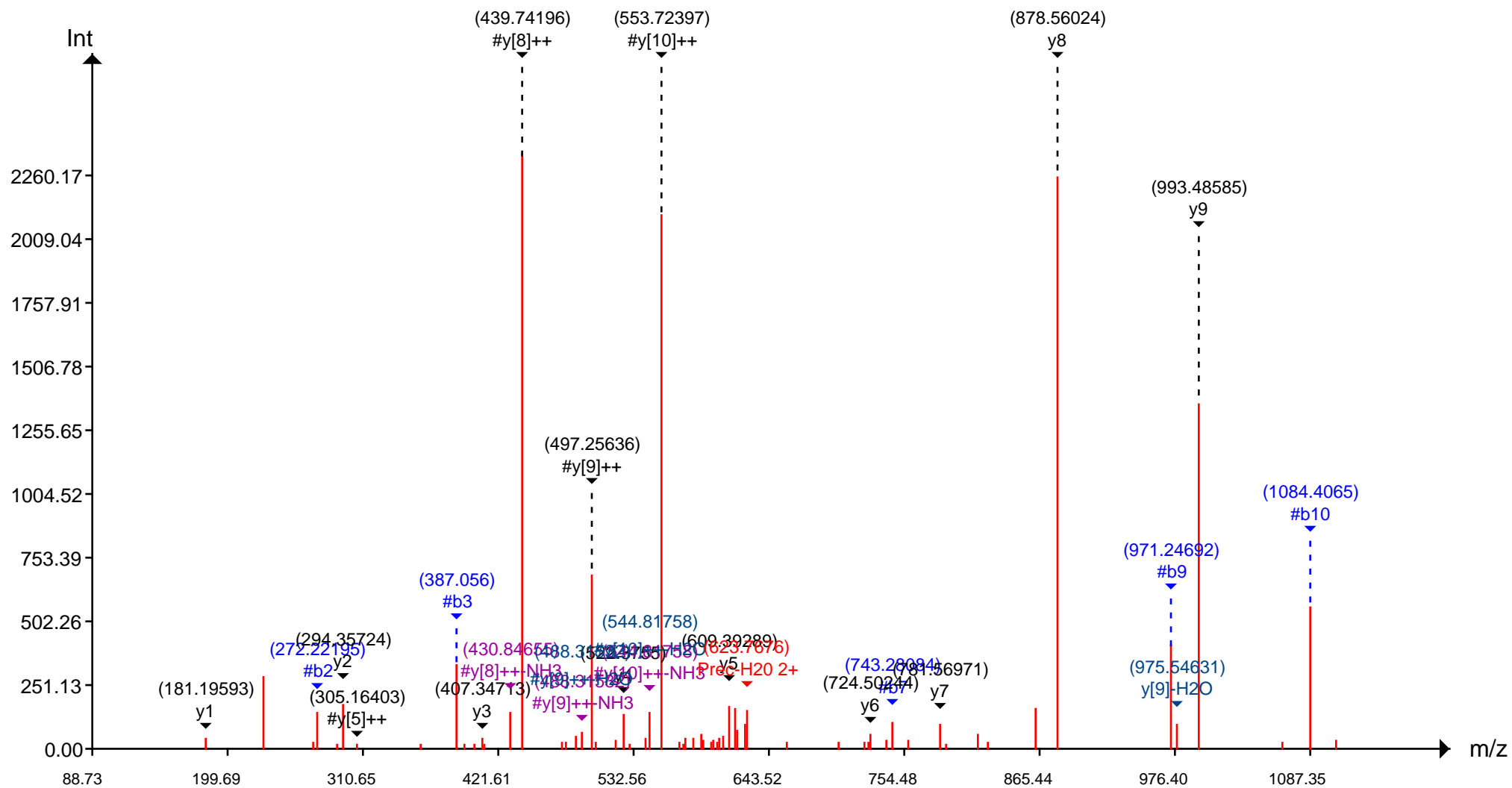
Info

Modified Sequence	AcD3-LVVDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR-COOH	Ion score	121.21
Protein	P60710	Confidence level	99.00%
Precursor m/z	1122.0219	Identity Threshold	45.15
Charge	2+	Mass error	0.002698



Info

Modified Sequence	AcD3-LVVDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR-COOH	Ion score	75.26
Protein	P60710	Confidence level	99.00%
Precursor m/z	1122.0221	Identity Threshold	45.20
Charge	2+	Mass error	0.003098



Info

Modified Sequence AcD3-IIDPGDSDIIR<C13>-COOH

Ion score 70.28

Protein P62889

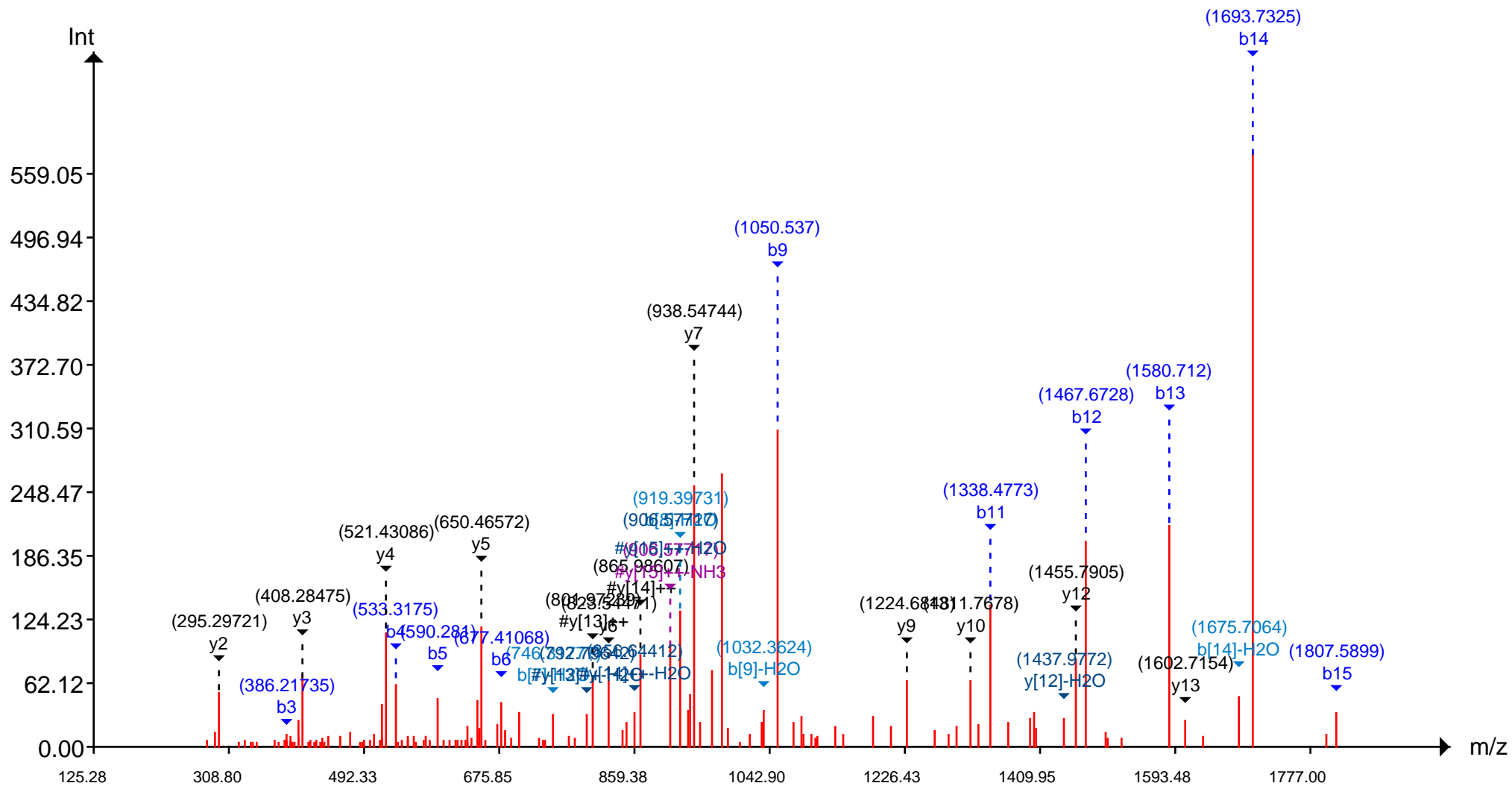
Confidence level 99.00%

Precursor m/z 632.84983

Identity Threshold 47.84

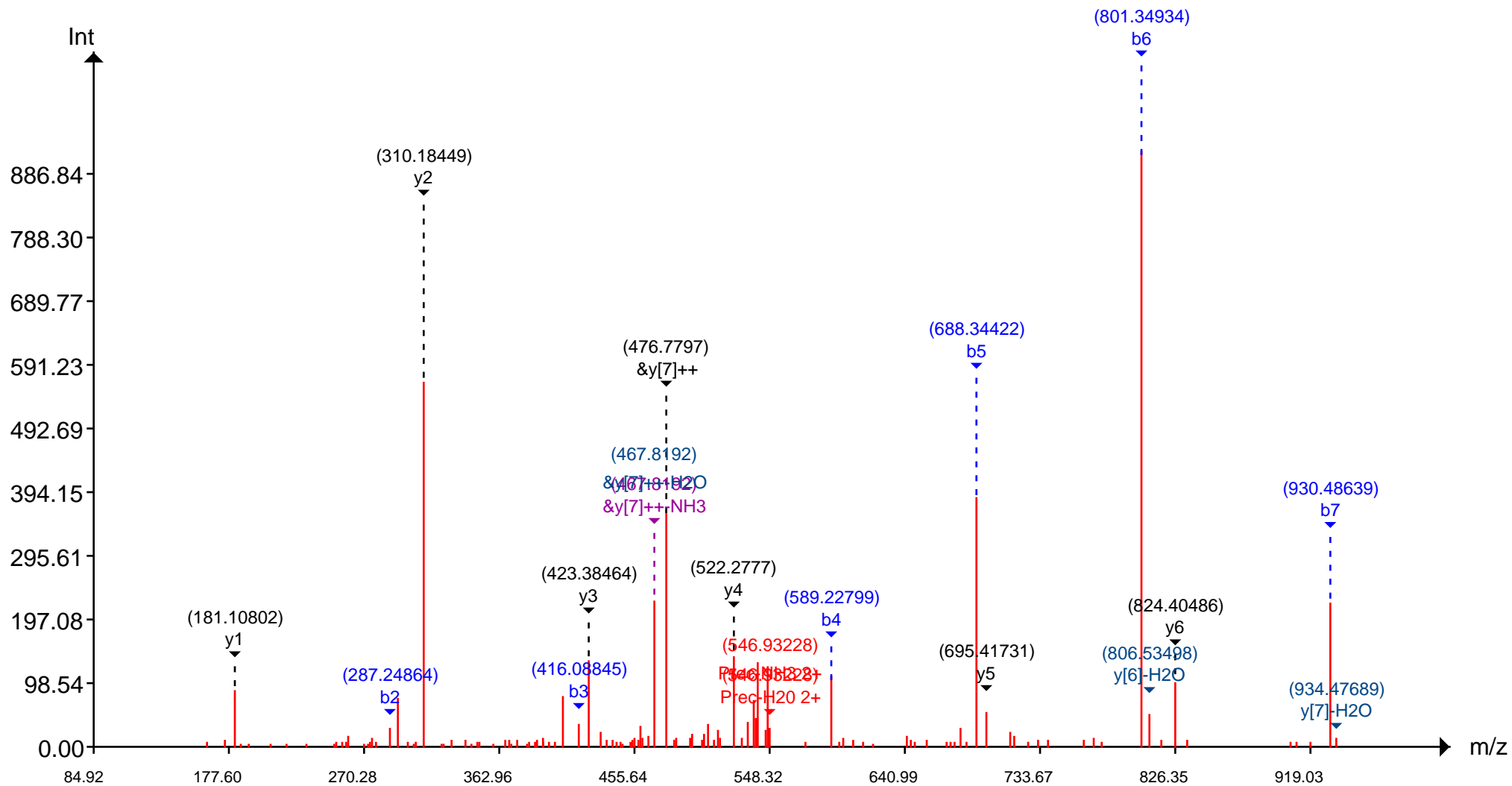
Charge 2+

Mass error 5.68E-4



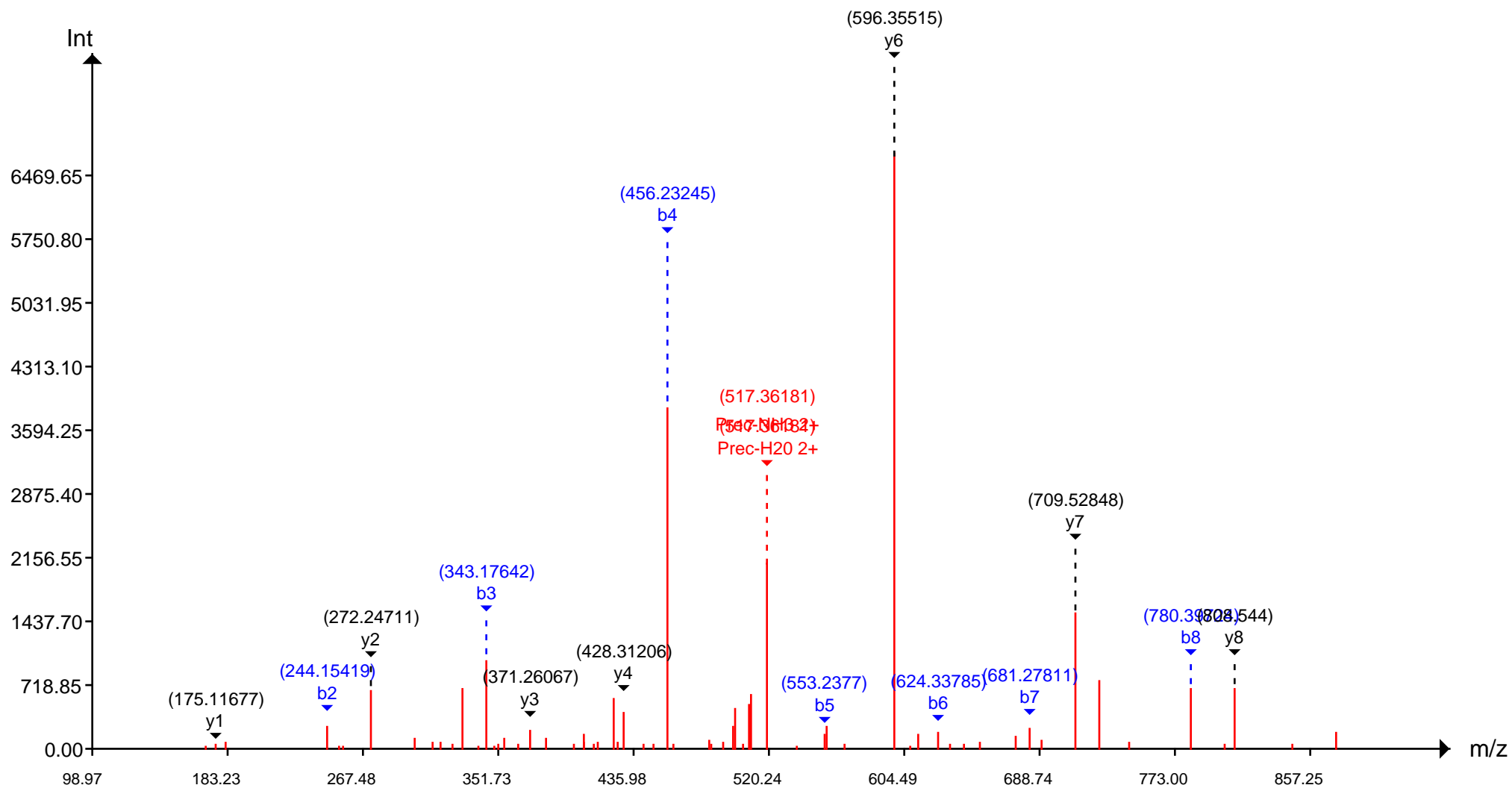
Info

Modified Sequence	AcD3-IVQFGSSK<AcD3K*>IDK<AcD3K*>ELINR<C13>-COOH	Ion score	60.2
Protein	P32921	Confidence level	99.00%
Precursor m/z	994.57835	Identity Threshold	44.13
Charge	2+	Mass error	0.002591



Info

Modified Sequence	AcD3-IQEK<AcD3K*>VLER<C13>-COOH	Ion score	47.96
Protein	P97372	Confidence level	99.00%
Precursor m/z	555.84016	Identity Threshold	43.94
Charge	2+	Mass error	-8.0E-5



Info

Modified Sequence AcD3-VVVIPAGVPR-COOH

Ion score 43.45

Protein P08249

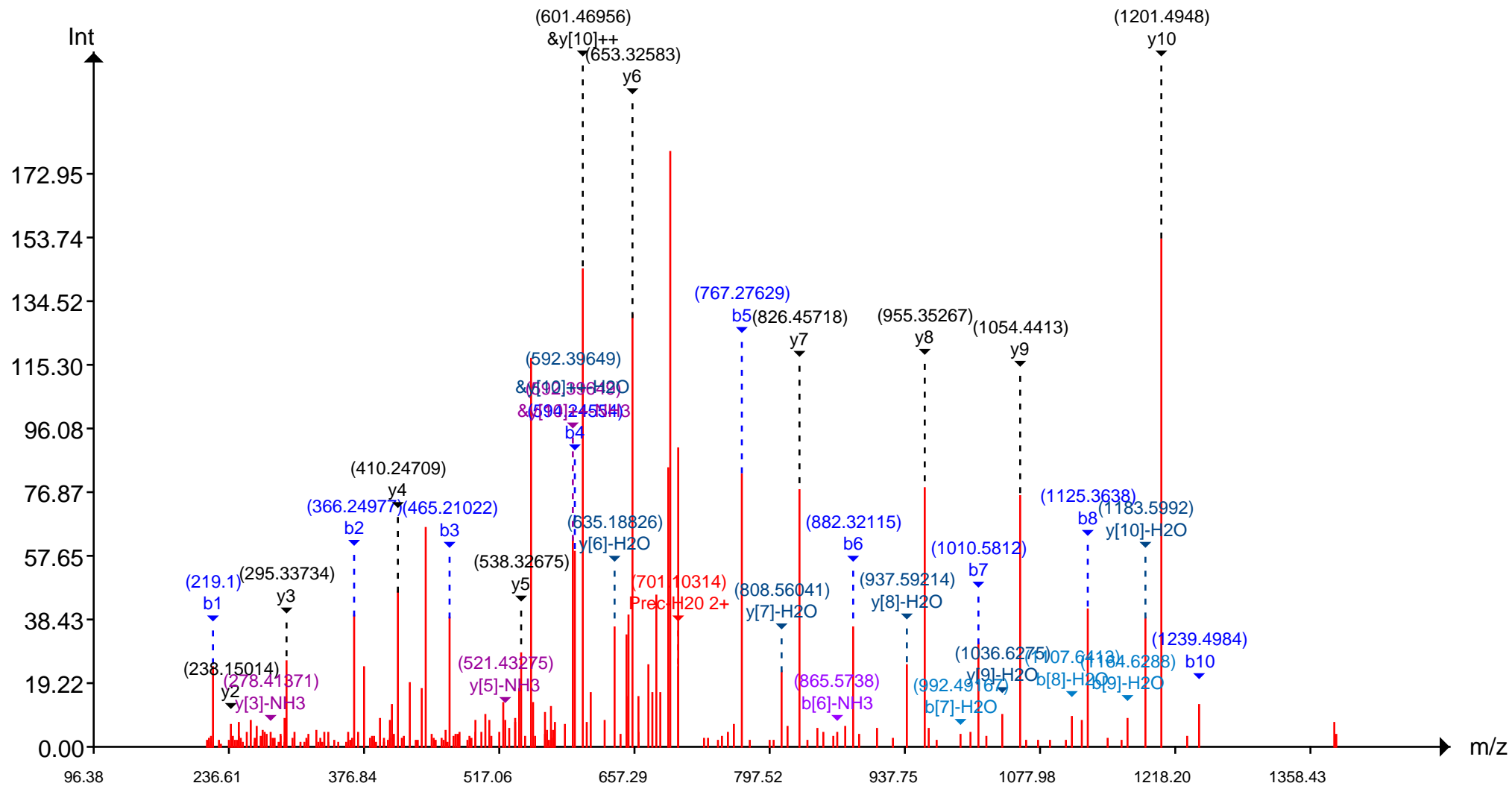
Confidence level 99.00%

Precursor m/z 526.33867

Identity Threshold 41.19

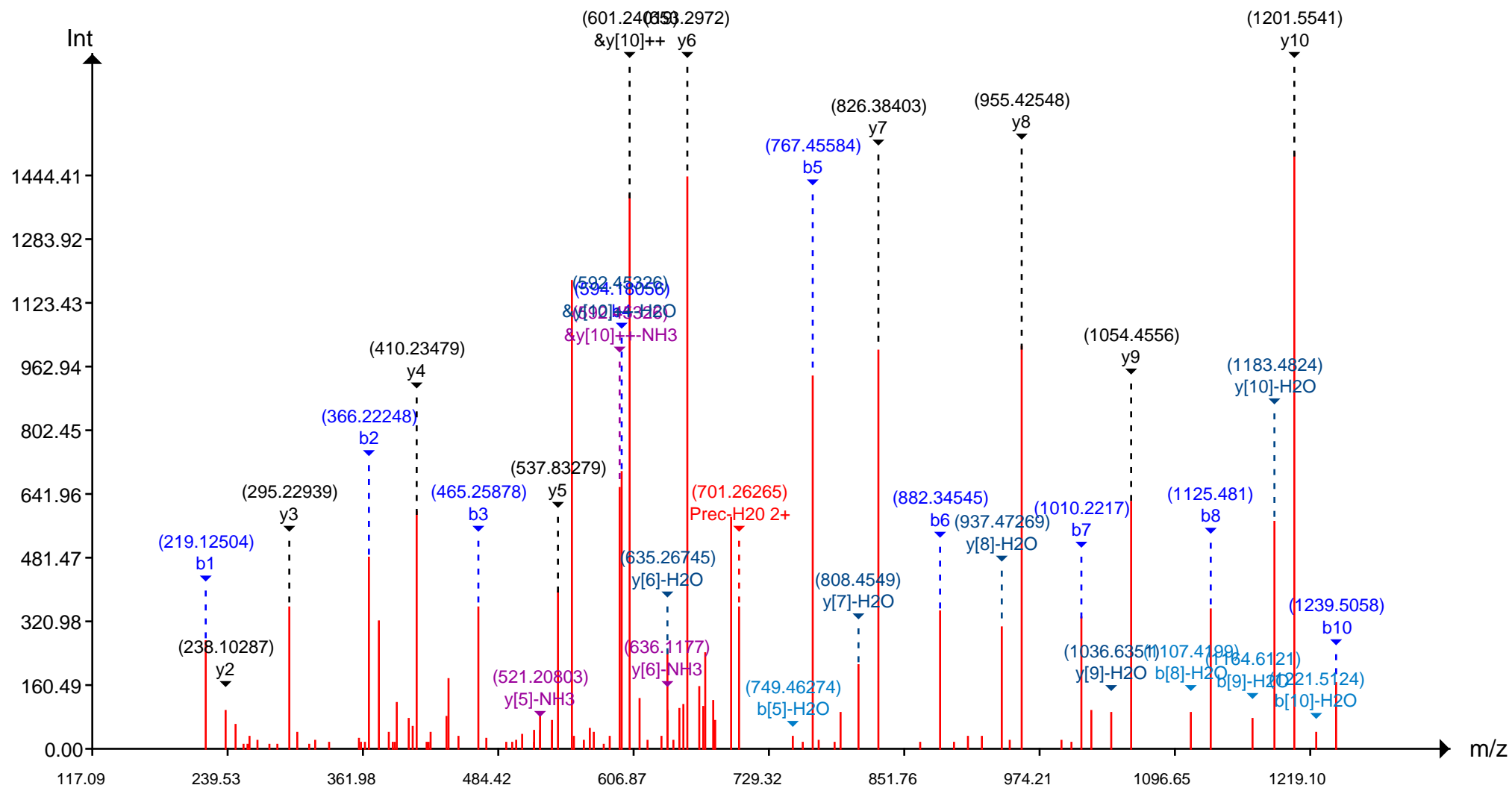
Charge 2+

Mass error -1.15E-4



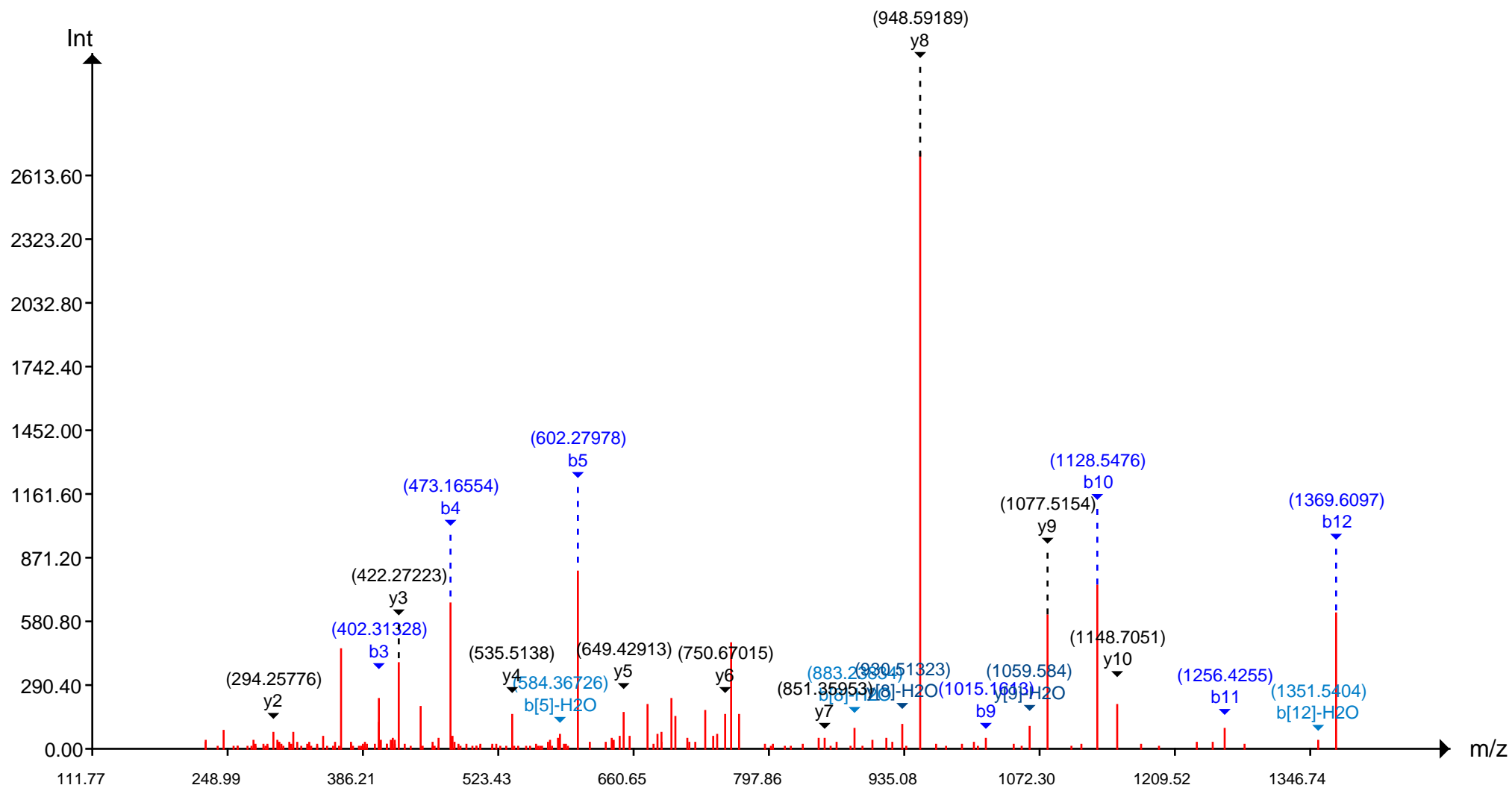
Info

Modified Sequence	AcD3-K<AcD3K*>M<Mox*>VEK<AcD3K*>DQDGGR<C13>-COOH	Ion score	66.33
Protein	Q9CZX8	Confidence level	99.00%
Precursor m/z	710.36367	Identity Threshold	47.84
Charge	2+	Mass error	0.001057



Info

Modified Sequence	AcD3-K<AcD3K*>M<Mox*>VEK<AcD3K*>DQDGGR<C13>-COOH	Ion score	75.89
Protein	Q9CZX8	Confidence level	99.00%
Precursor m/z	710.36365	Identity Threshold	47.84
Charge	2+	Mass error	0.001017



Info

Modified Sequence AcD3-INEAEPTTNIQIR<C13>-COOH

Ion score 51.88

Protein Q9CZ44

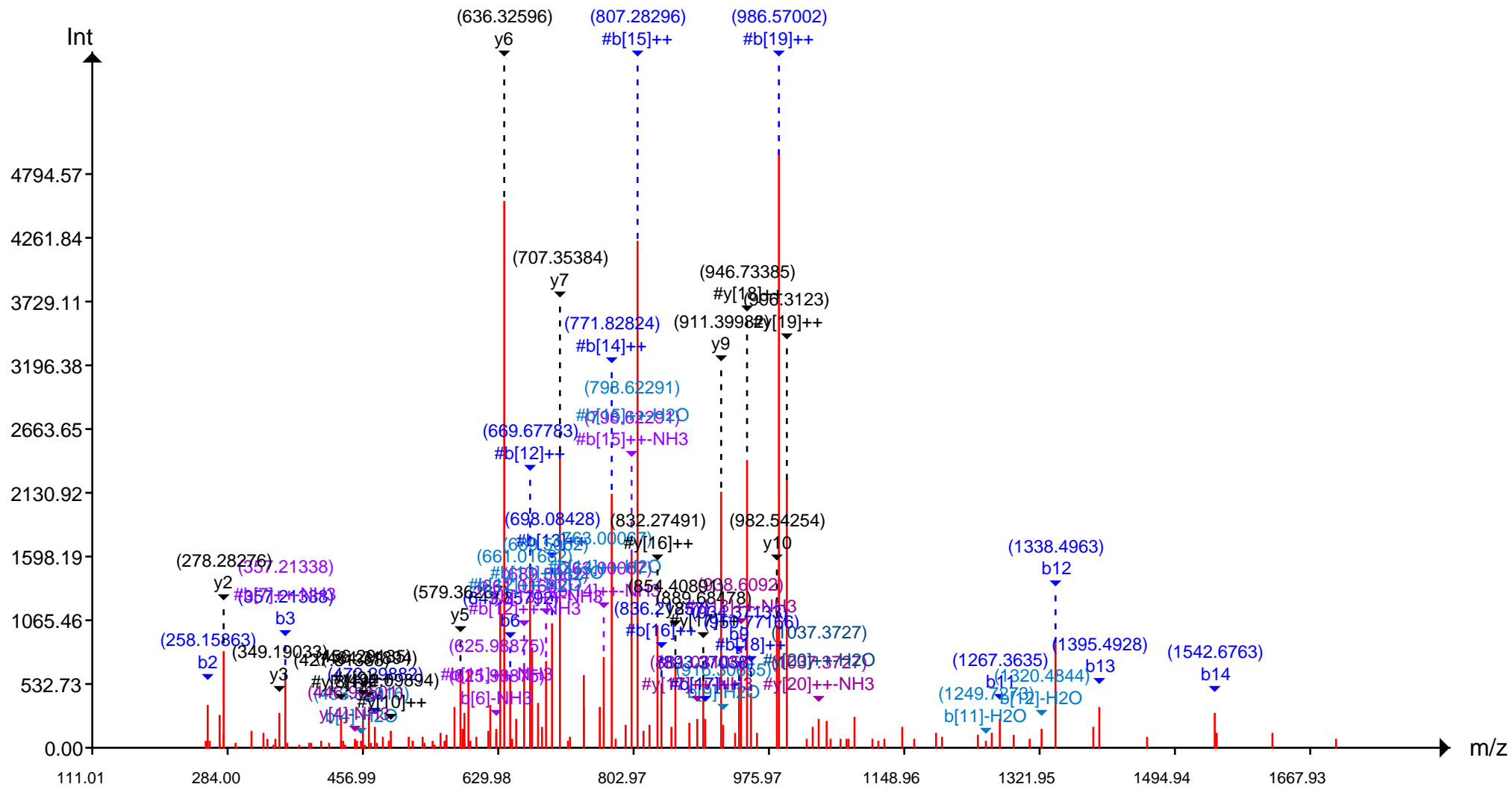
Confidence level 99.00%

Precursor m/z 775.42202

Identity Threshold 47.68

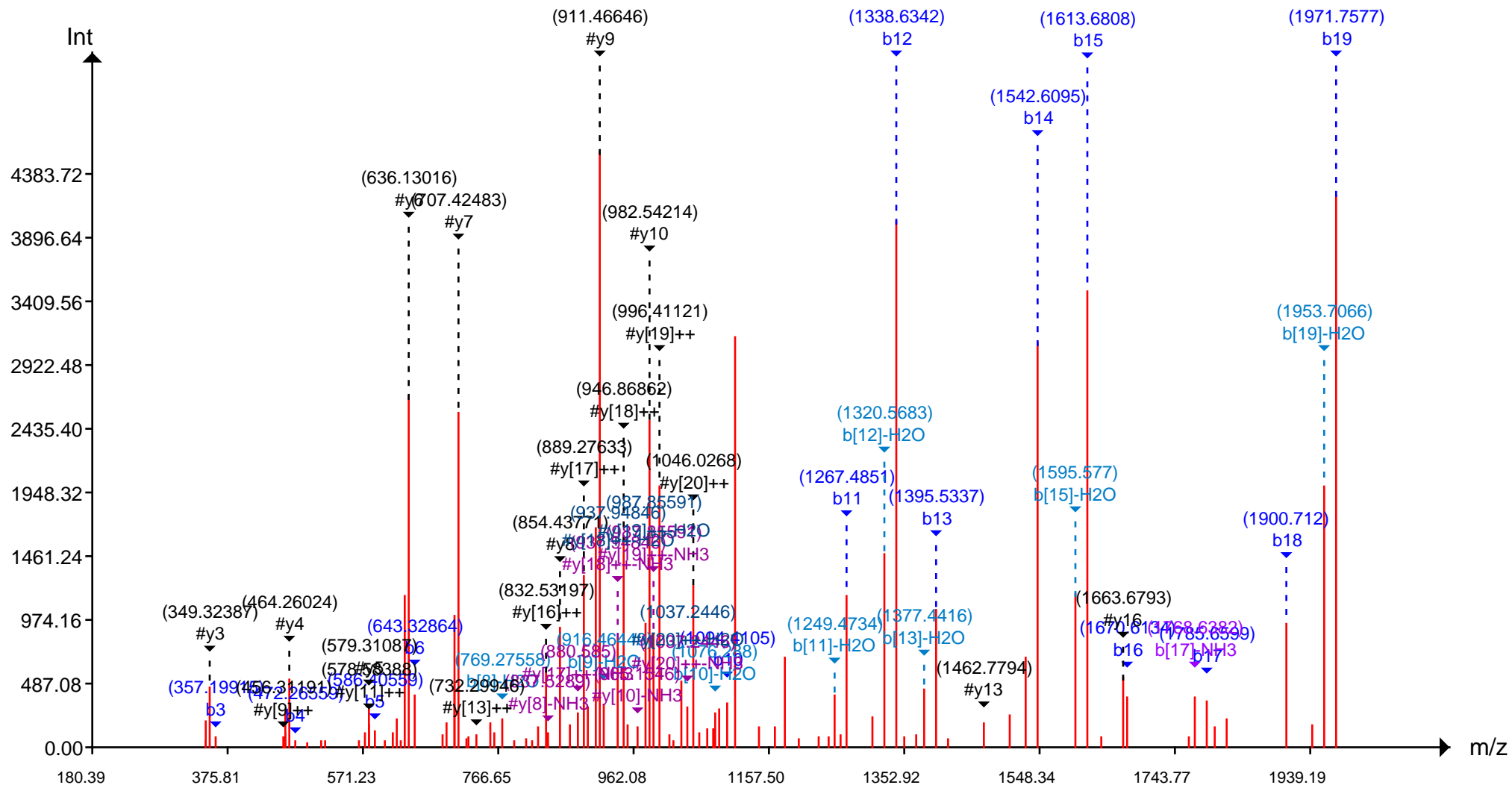
Charge 2+

Mass error 0.001271



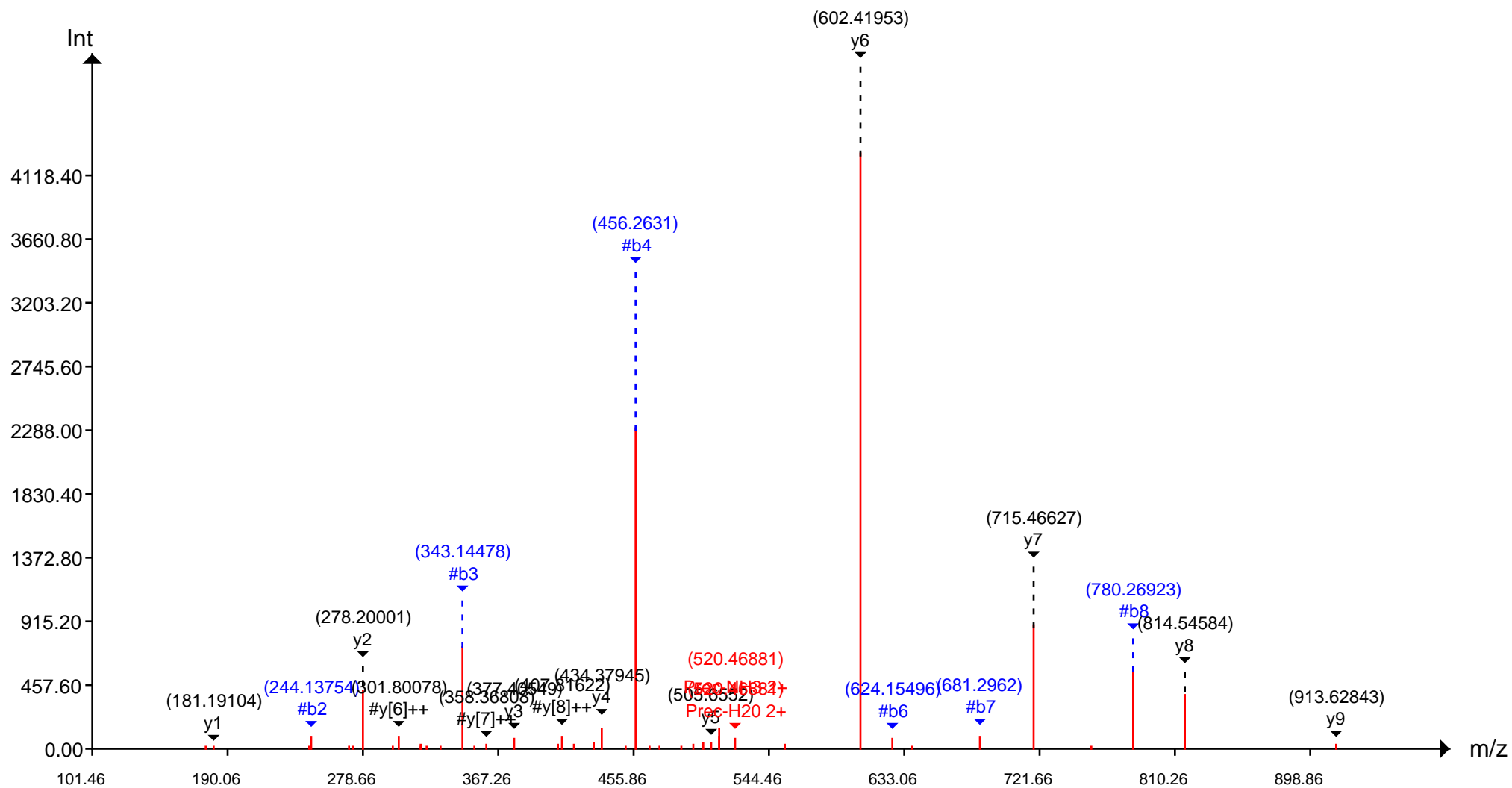
Info

Modified Sequence	AcD3-LVVDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH	Ion score	62.73
Protein	P60710	Confidence level	99.00%
Precursor m/z	750.35645	Identity Threshold	45.77
Charge	3+	Mass error	8.3E-4



Info

Modified Sequence	AcD3-LVVDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH	Ion score	91.34
Protein	P60710	Confidence level	99.00%
Precursor m/z	1125.0312	Identity Threshold	45.83
Charge	2+	Mass error	0.001156



Info

Modified Sequence AcD3-VVVIPAGVPR<C13>-COOH

Ion score 77.18

Protein P08249

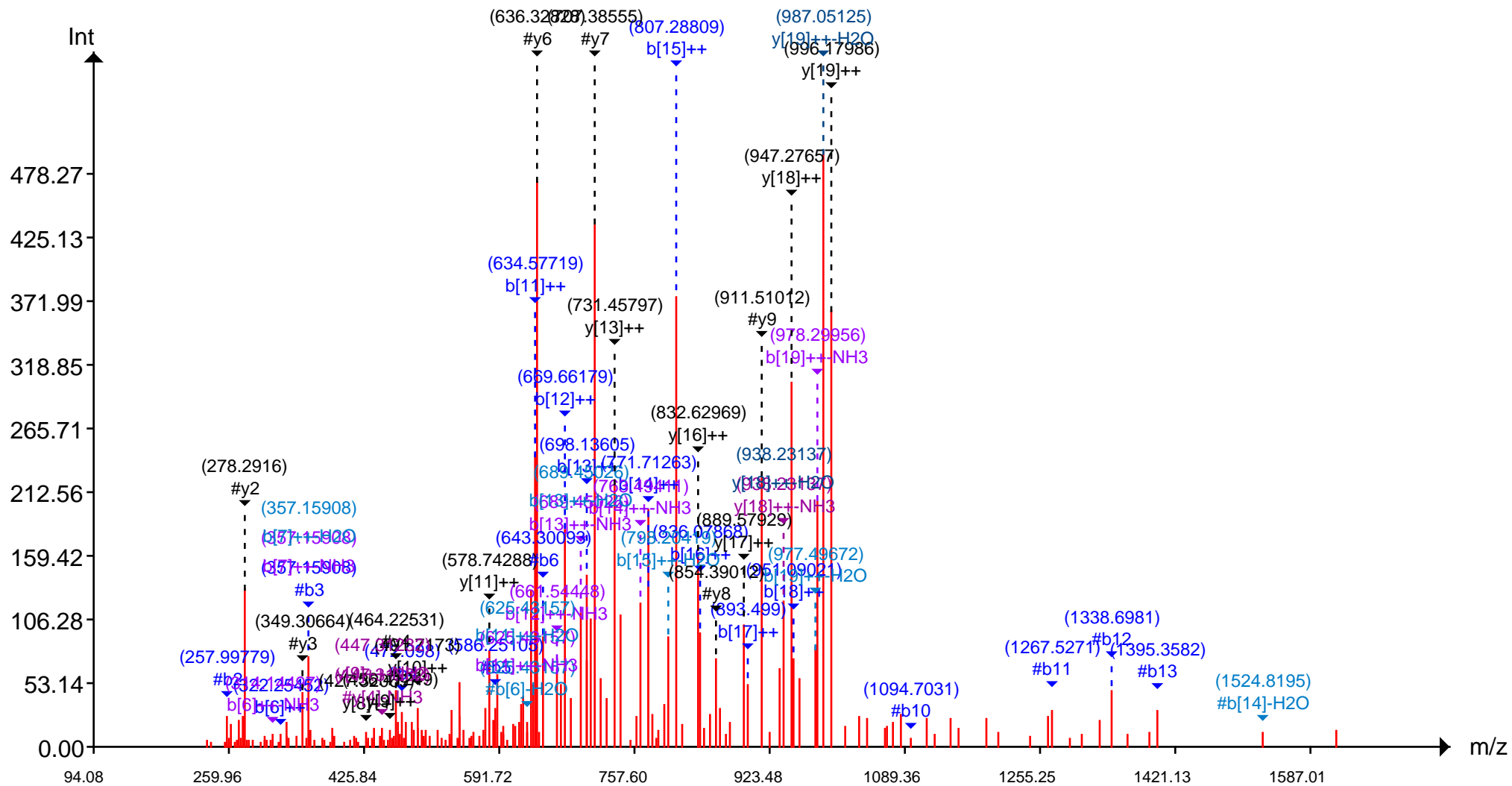
Confidence level 99.00%

Precursor m/z 529.34884

Identity Threshold 38.80

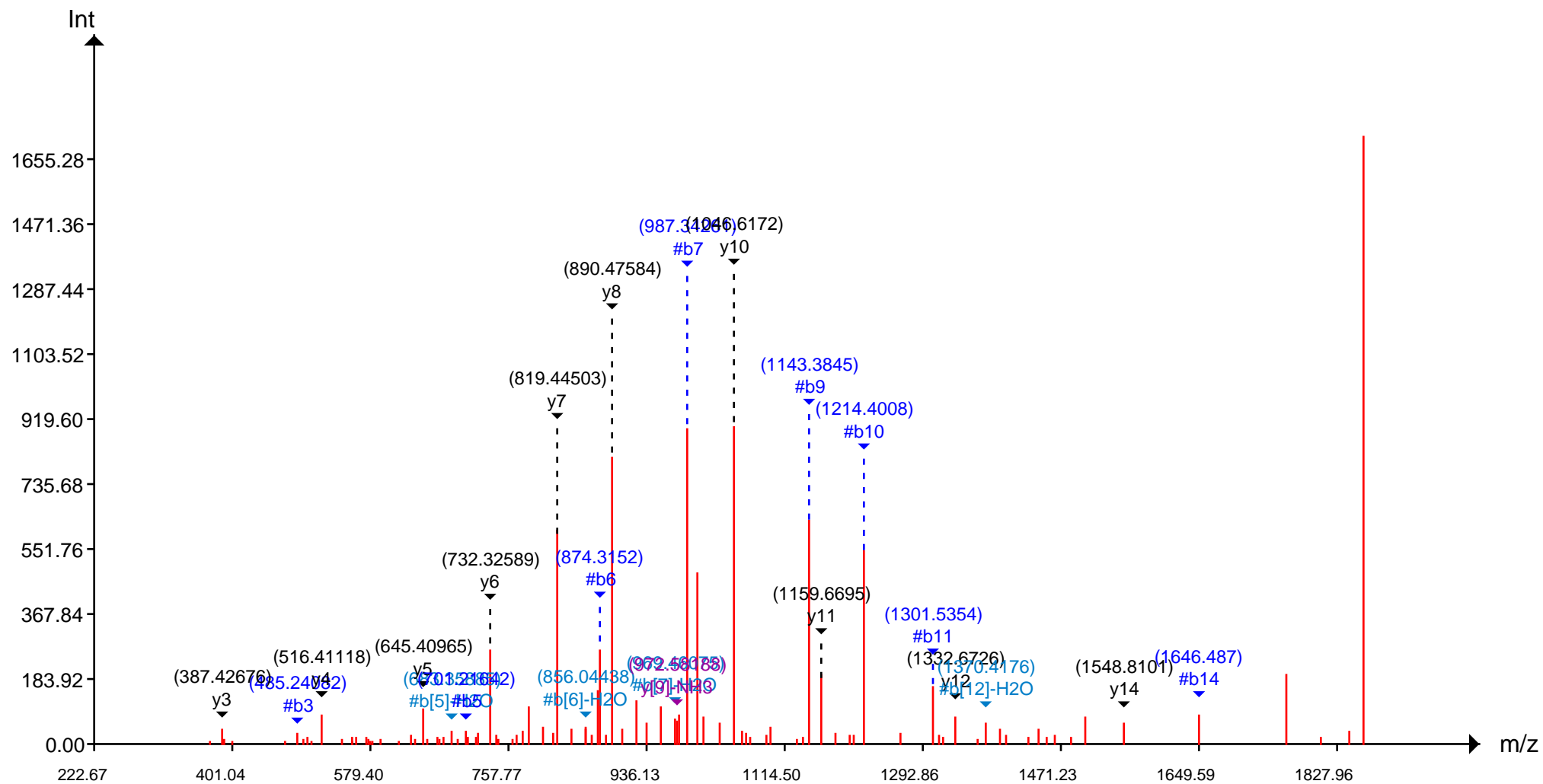
Charge 2+

Mass error 8.4E-5



Info

Modified Sequence	AcD3-LVVDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH	Ion score	50.09
Protein	P60710	Confidence level	99.00%
Precursor m/z	750.3567	Identity Threshold	45.83
Charge	3+	Mass error	0.00158



Info

Modified Sequence AcD3-M<Mox*>YESEK<AcD3K*>IGVASSEELVR-COOH

Ion score 68.8

Protein Q9ERB0

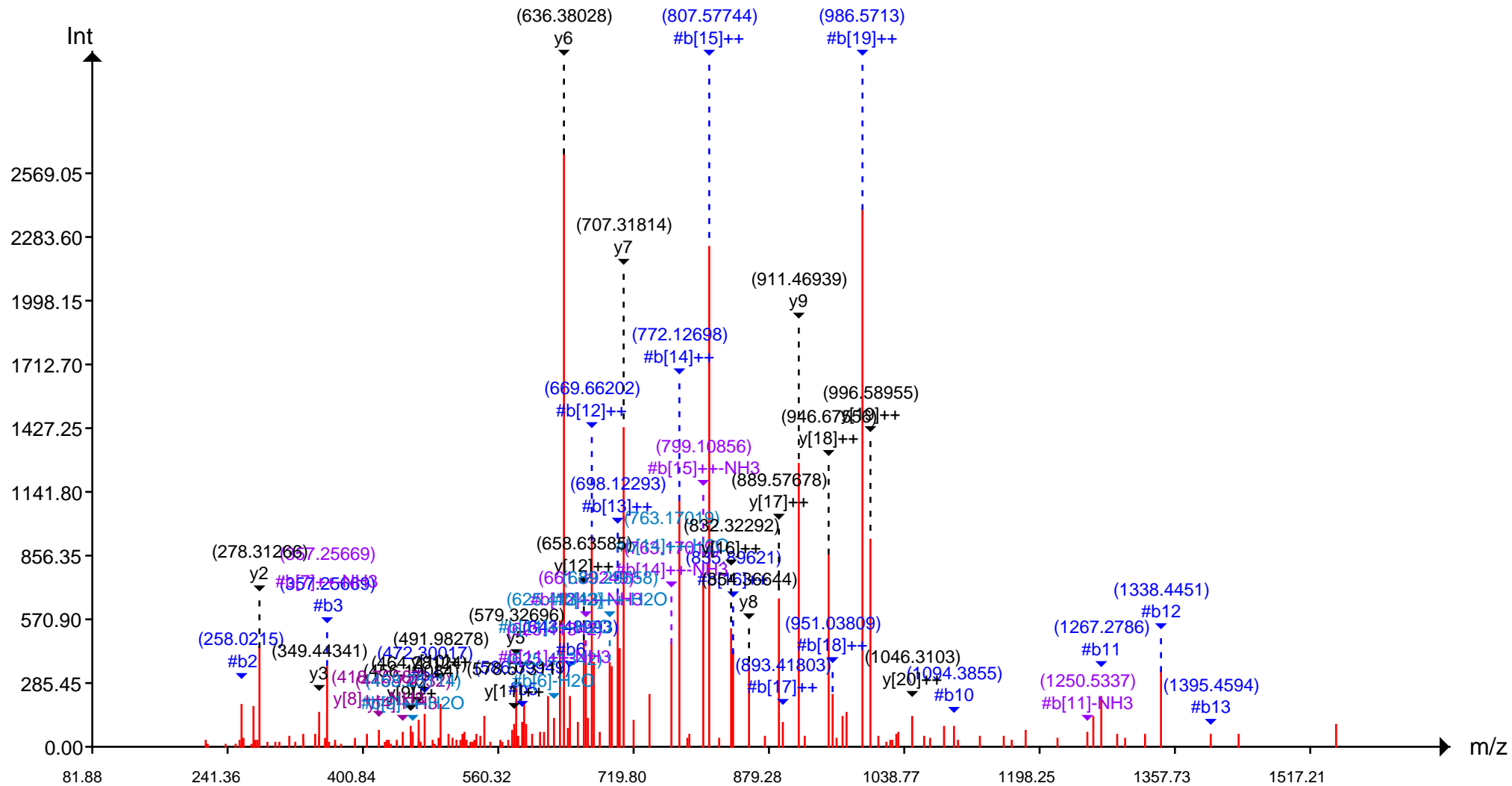
Confidence level 99.00%

Precursor m/z 1017.0055

Identity Threshold 47.50

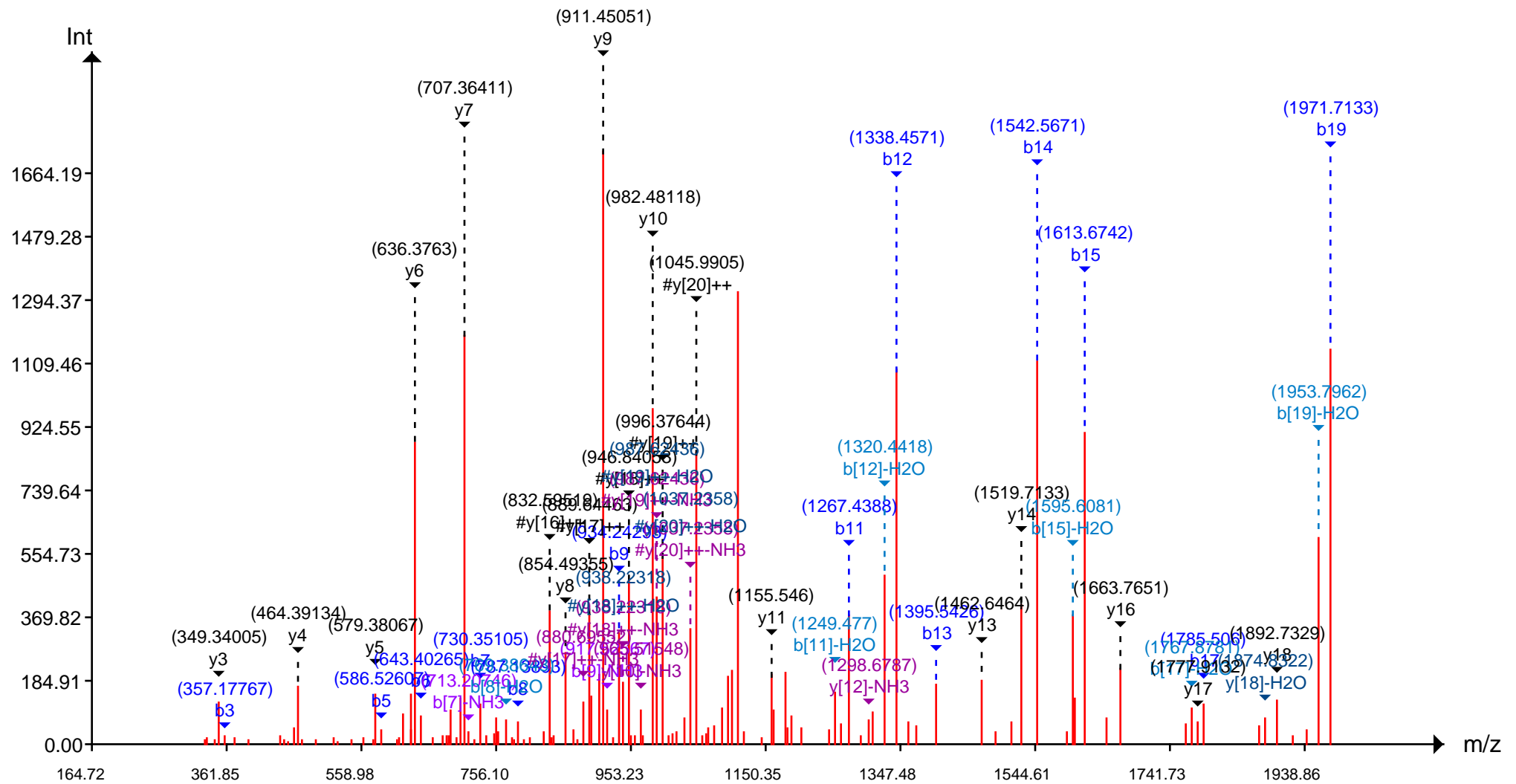
Charge 2+

Mass error 0.002338



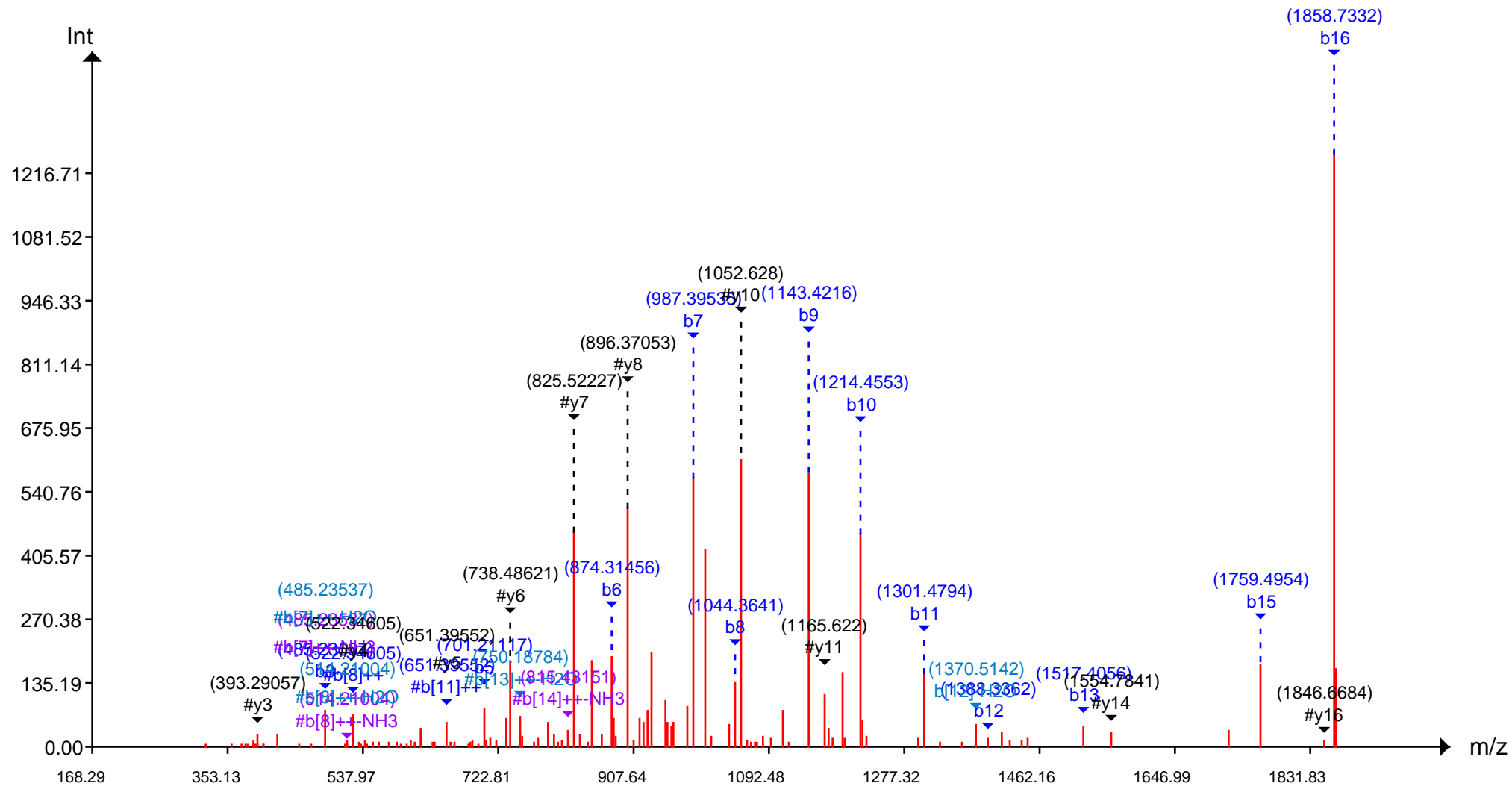
Info

Modified Sequence	AcD3-LVVDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH	Ion score	65.62
Protein	P60710	Confidence level	99.00%
Precursor m/z	750.35677	Identity Threshold	45.82
Charge	3+	Mass error	0.00179



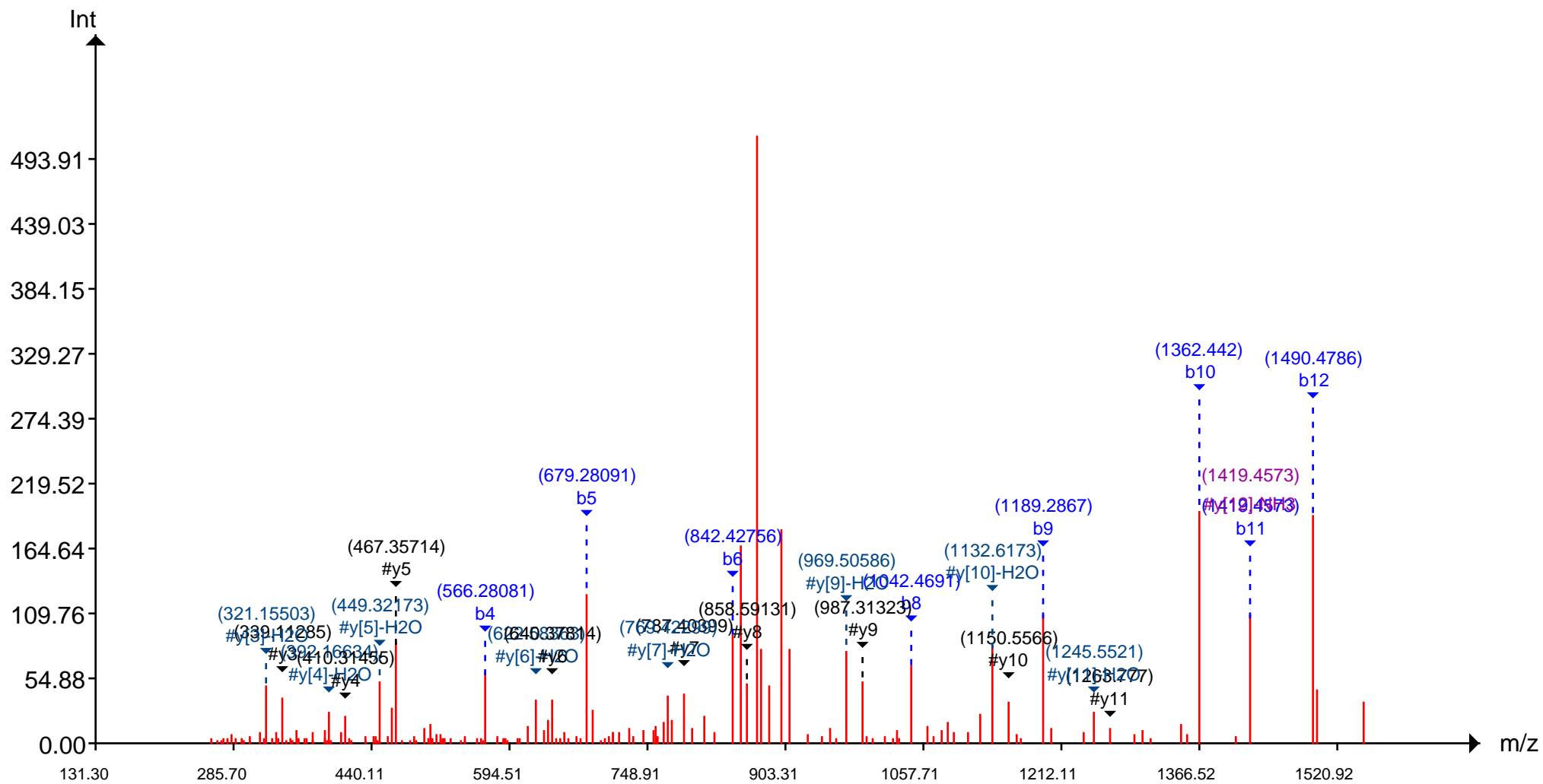
Info

Modified Sequence	AcD3-LVVDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH	Ion score	119.81
Protein	P60710	Confidence level	99.00%
Precursor m/z	1125.0315	Identity Threshold	45.83
Charge	2+	Mass error	0.001756



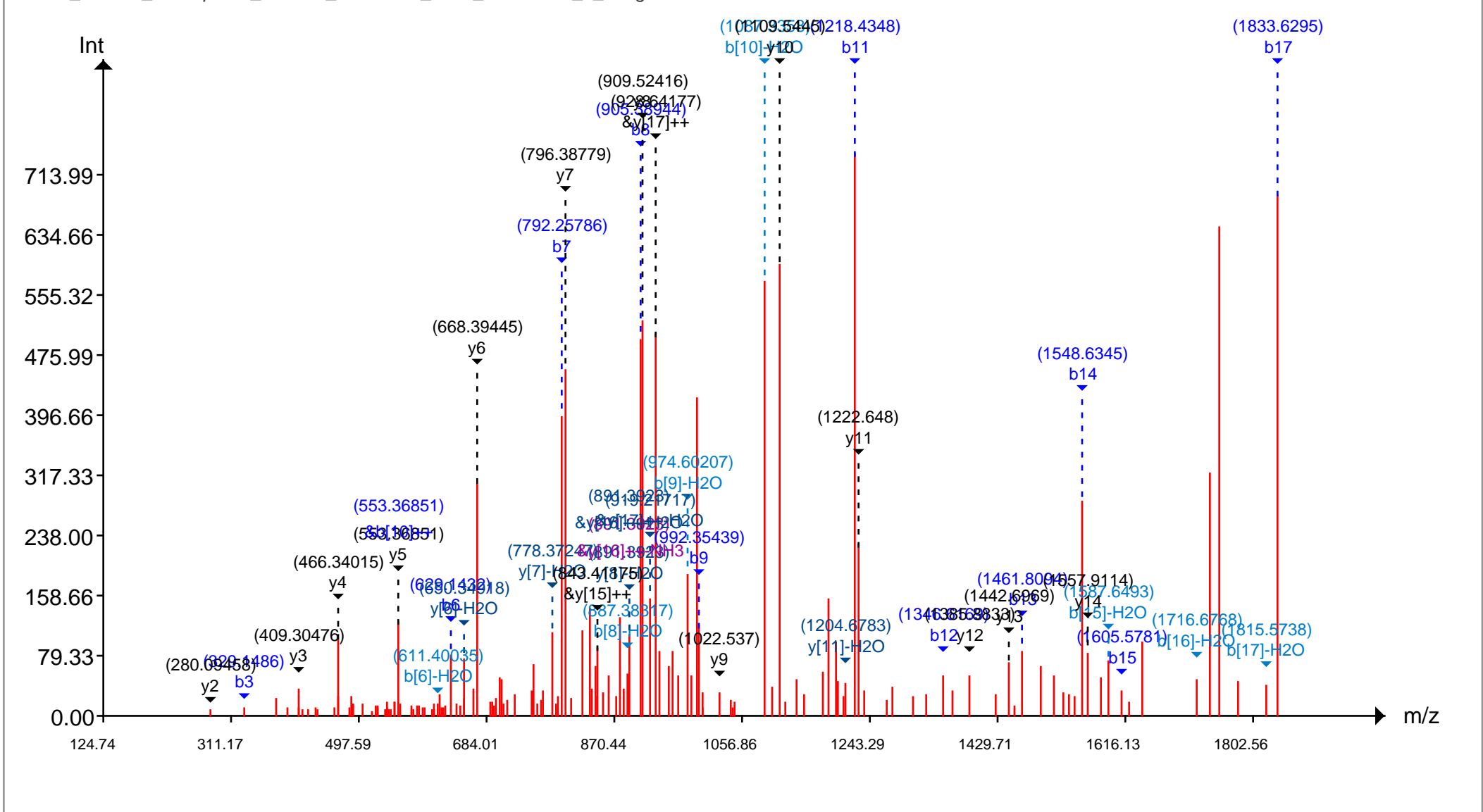
Info

Modified Sequence	AcD3-M<Mox*>YESEK<AcD3K*>IGVASSEELVR<C13>-COOH	Ion score	81.5
Protein	Q9ERB0	Confidence level	99.00%
Precursor m/z	1020.0157	Identity Threshold	47.95
Charge	2+	Mass error	0.002596



Info

Modified Sequence	AcD3-FAEK<AcD3K*>LYEAM<Mox*>K<AcD3K*>GAGTR<C13>-COOH	Ion score	58.38
Protein	P10107	Confidence level	99.00%
Precursor m/z	914.98286	Identity Threshold	48.12
Charge	2+	Mass error	0.002926



Info

Modified Sequence AcD3-IGIQDGYLSLLQDSGEVR<C13>-COOH

Ion score 95.96

Protein P63242

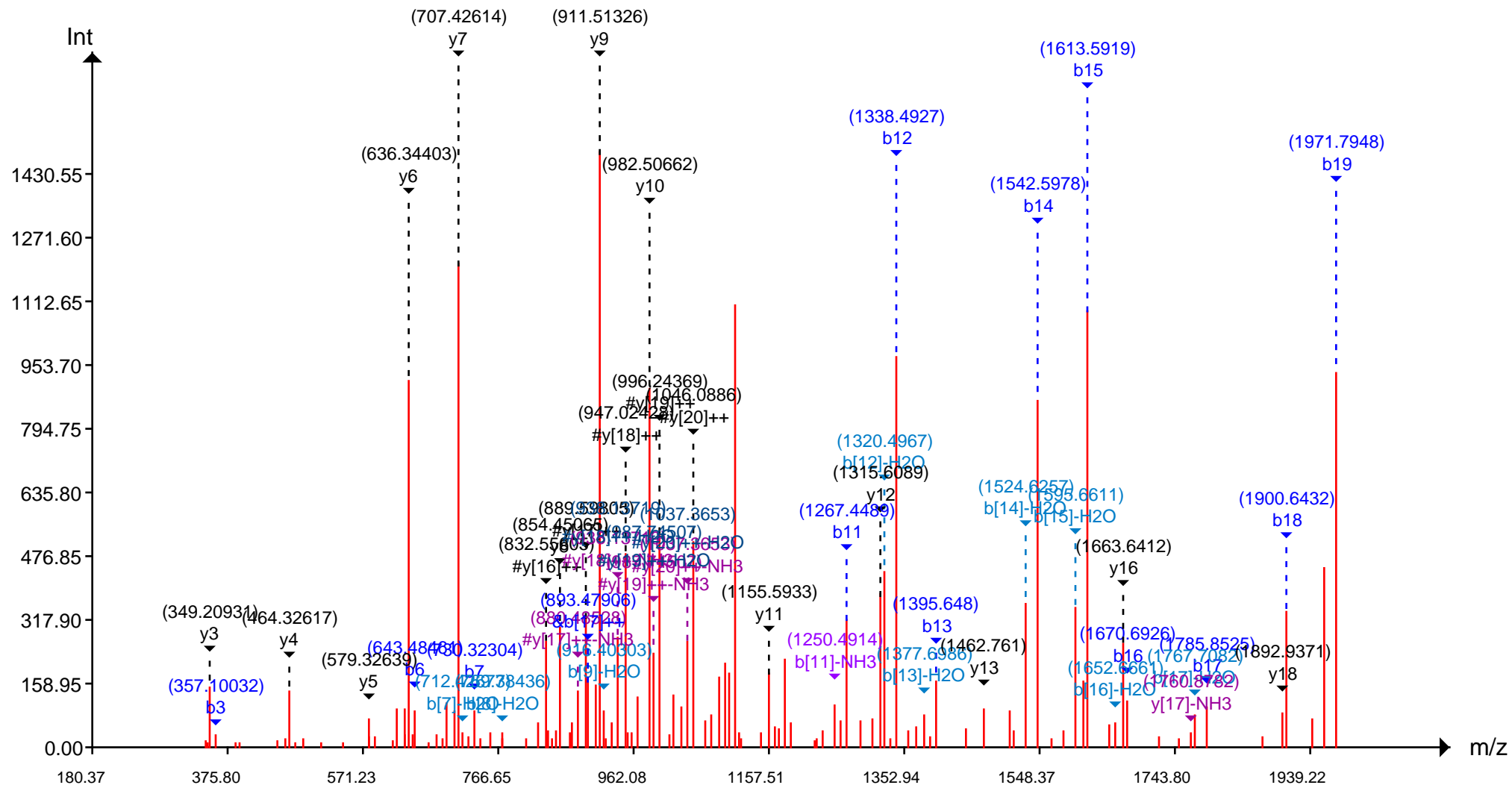
Confidence level 99.00%

Precursor m/z 1007.5356

Identity Threshold 47.90

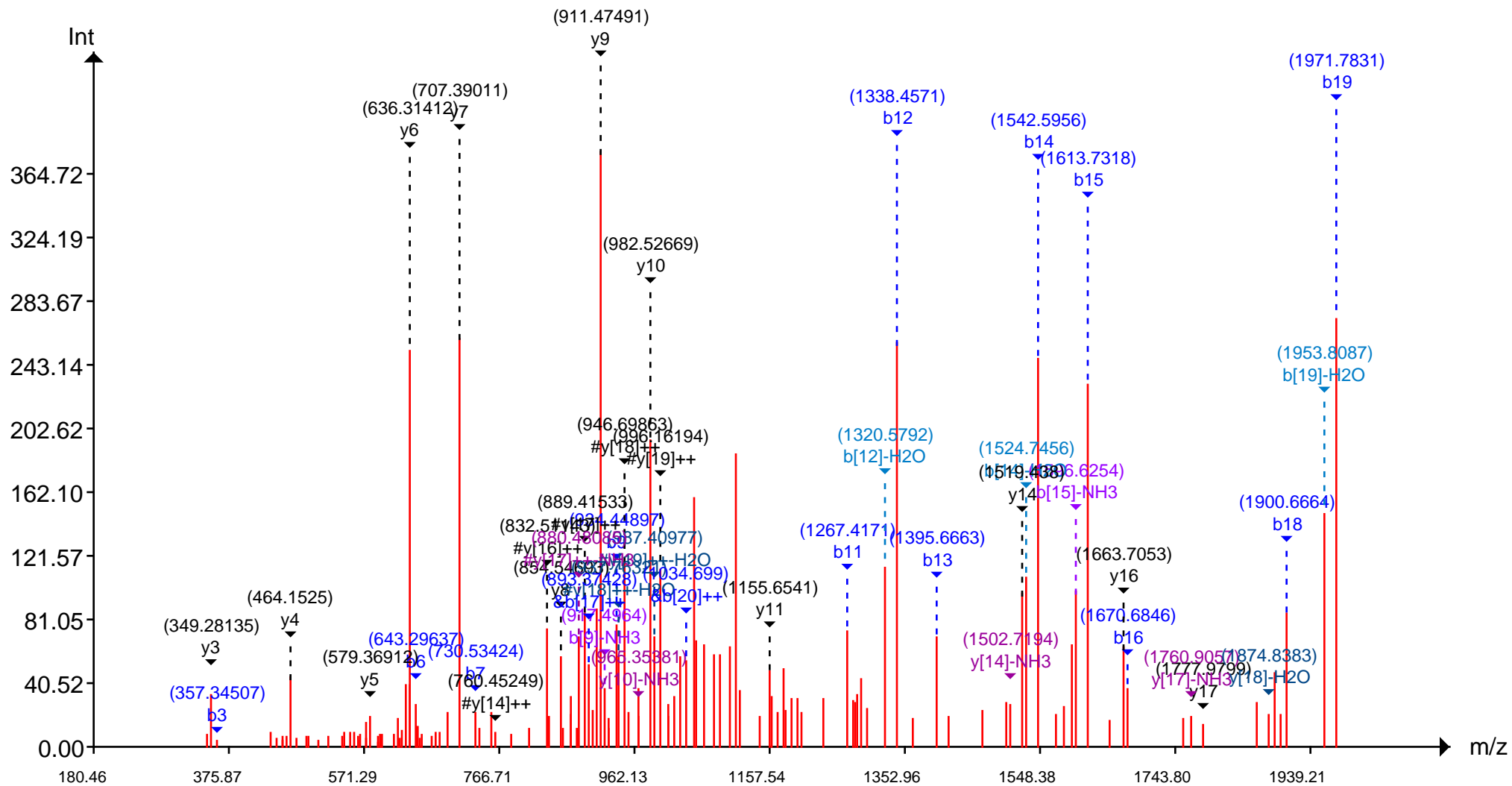
Charge 2+

Mass error 0.001304



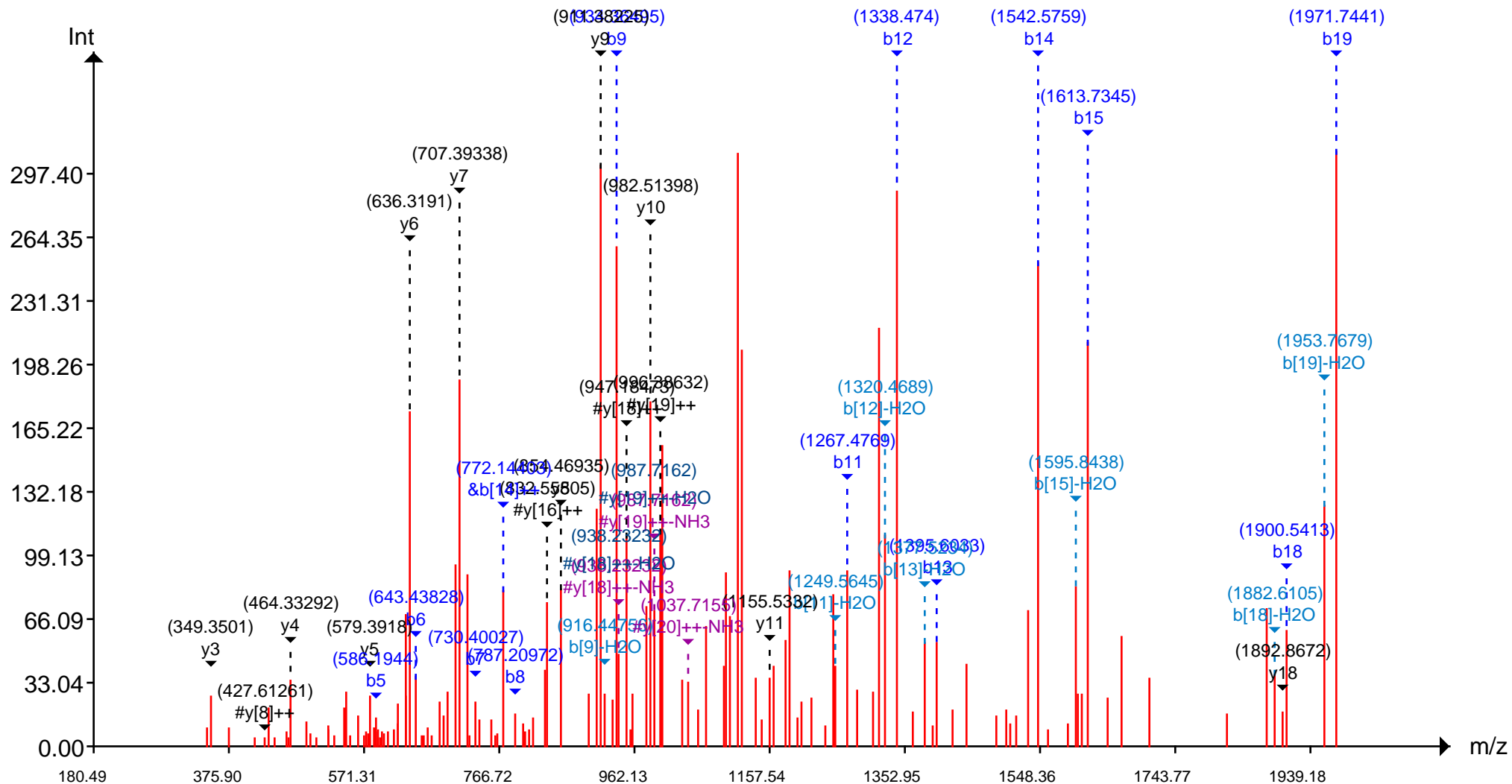
Info

Modified Sequence	AcD3-LVVDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH	Ion score	102.78
Protein	P60710	Confidence level	99.00%
Precursor m/z	1125.0317	Identity Threshold	45.83
Charge	2+	Mass error	0.002156



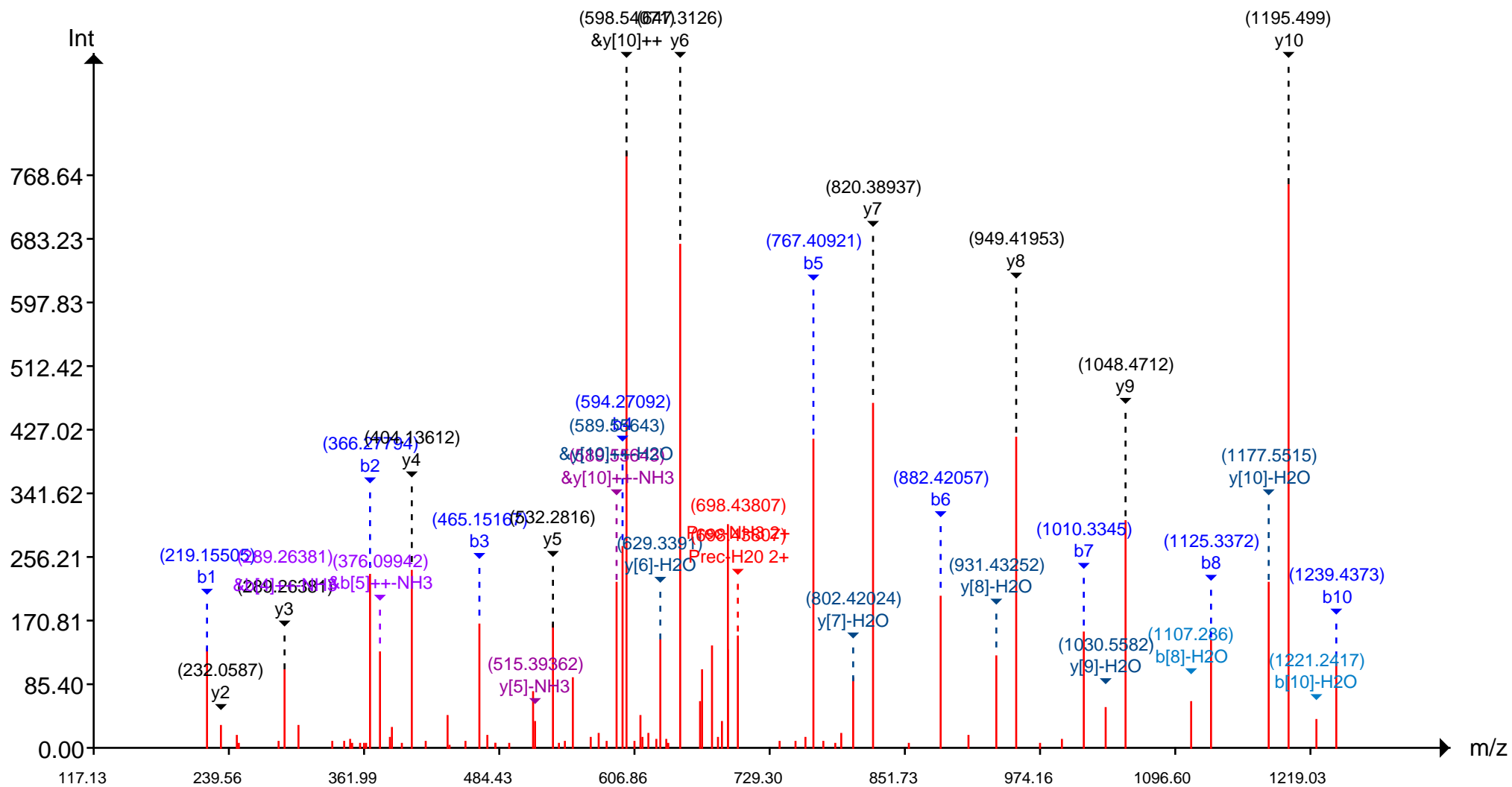
Info

Modified Sequence	AcD3-LVVDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH	Ion score	77.5
Protein	P60710	Confidence level	99.00%
Precursor m/z	1125.0321	Identity Threshold	45.87
Charge	2+	Mass error	0.002956



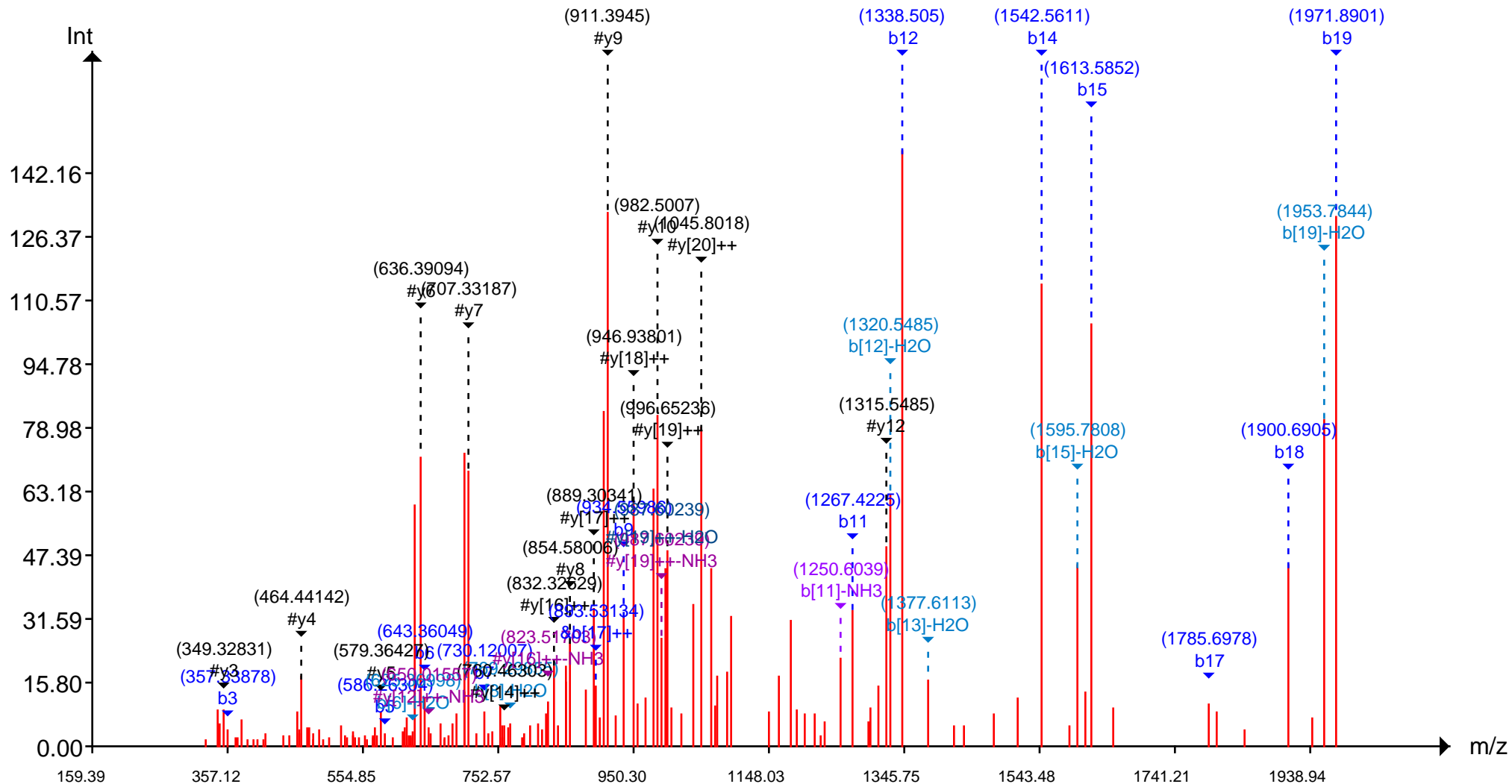
Info

Modified Sequence	AcD3-LVVDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH	Ion score	53.26
Protein	P60710	Confidence level	99.00%
Precursor m/z	1125.032	Identity Threshold	45.84
Charge	2+	Mass error	0.002756



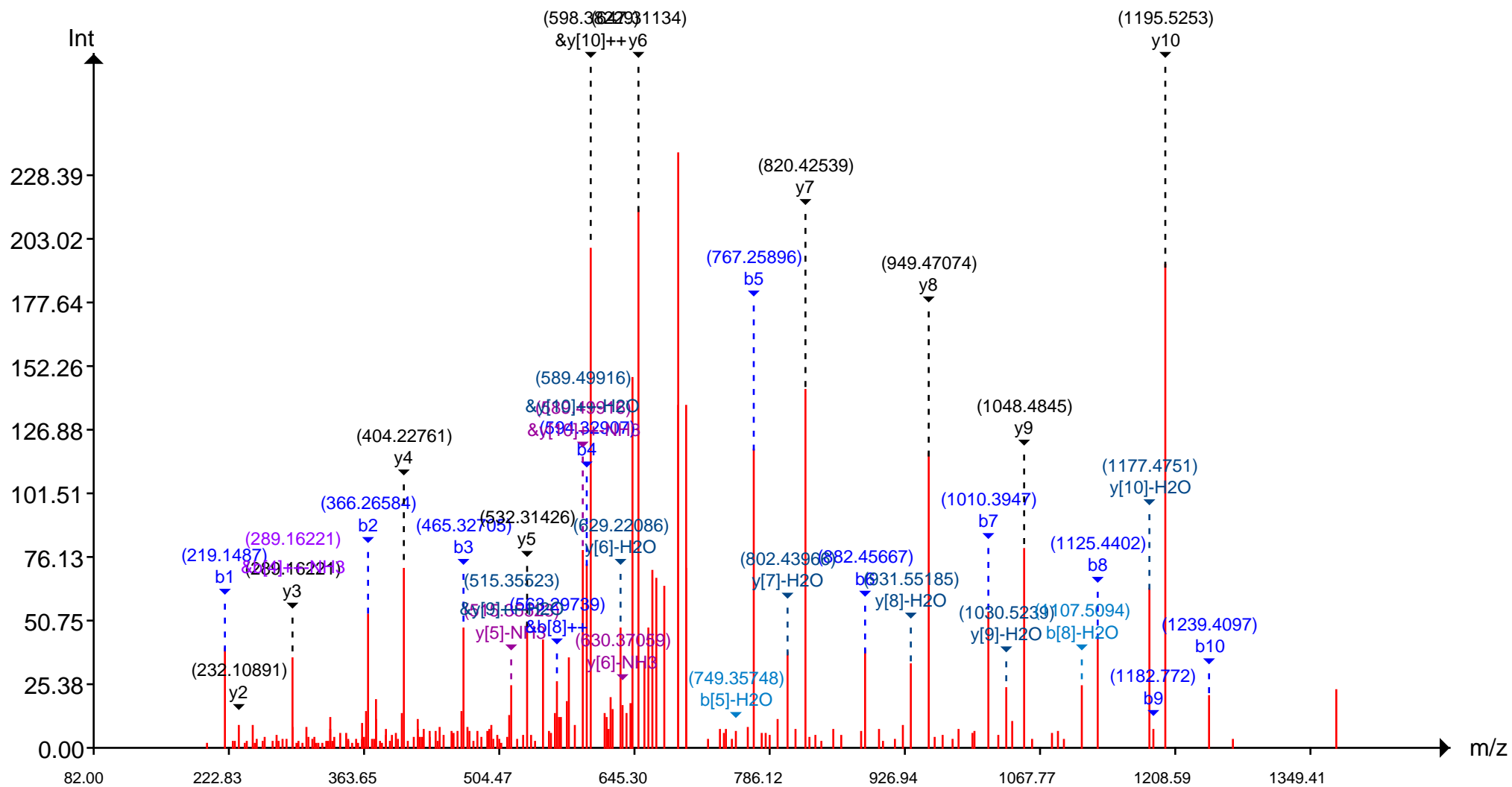
Info

Modified Sequence	AcD3-K<AcD3K*>M<Mox*>VEK<AcD3K*>DQDGGR-COOH	Ion score	86.37
Protein	Q9CZX8	Confidence level	99.00%
Precursor m/z	707.35404	Identity Threshold	47.65
Charge	2+	Mass error	0.001939



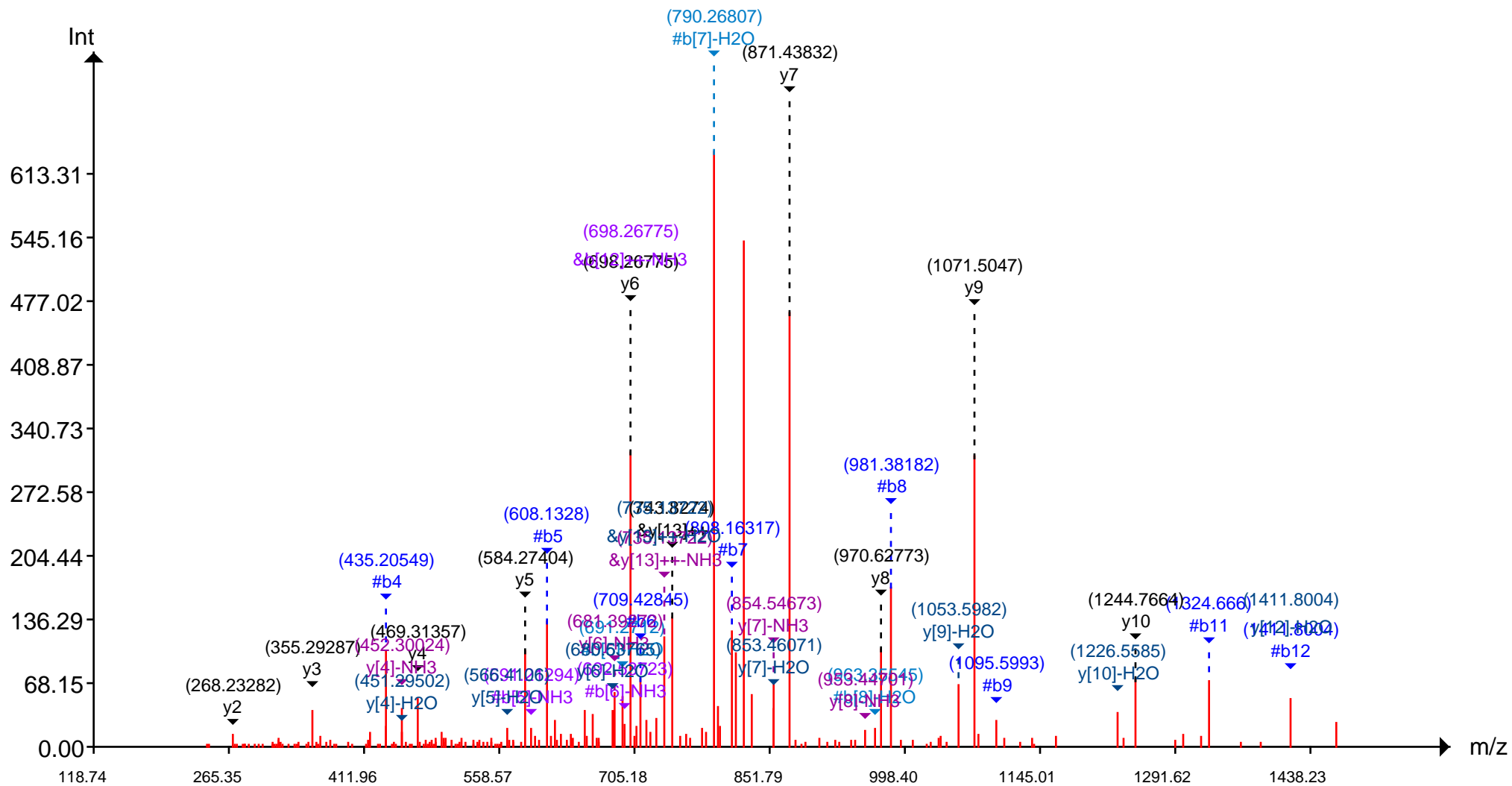
Info

Modified Sequence	AcD3-LVVDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH	Ion score	51.91
Protein	P60710	Confidence level	99.00%
Precursor m/z	1125.0323	Identity Threshold	45.87
Charge	2+	Mass error	0.003356



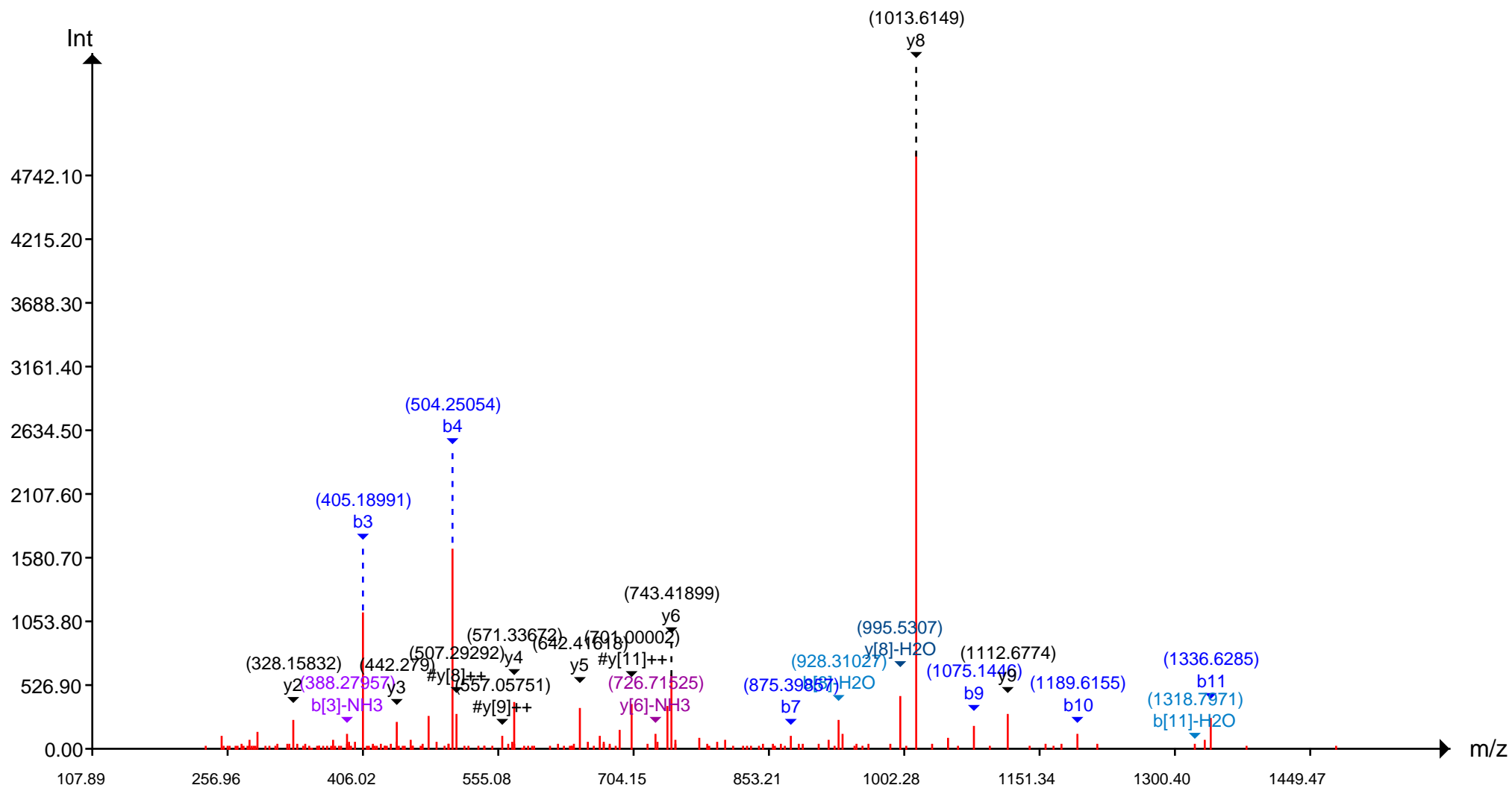
Info

Modified Sequence	AcD3-K<AcD3K*>M<Mox*>VEK<AcD3K*>DQDGGR-COOH	Ion score	70.1
Protein	Q9CZX8	Confidence level	99.00%
Precursor m/z	707.3534	Identity Threshold	47.51
Charge	2+	Mass error	6.59E-4



Info

Modified Sequence	AcD3-FGNAK<AcD3K*>TVK<AcD3K*>NDNSSR<C13>-COOH	Ion score	78.37
Protein	Q61879, O08638, Q6URW6, Q8VDD5	Confidence level	99.00%
Precursor m/z	839.94371	Identity Threshold	48.14
Charge	2+	Mass error	8.1E-5



Info

Modified Sequence AcD3-ADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH

Ion score 48.76

Protein P17742

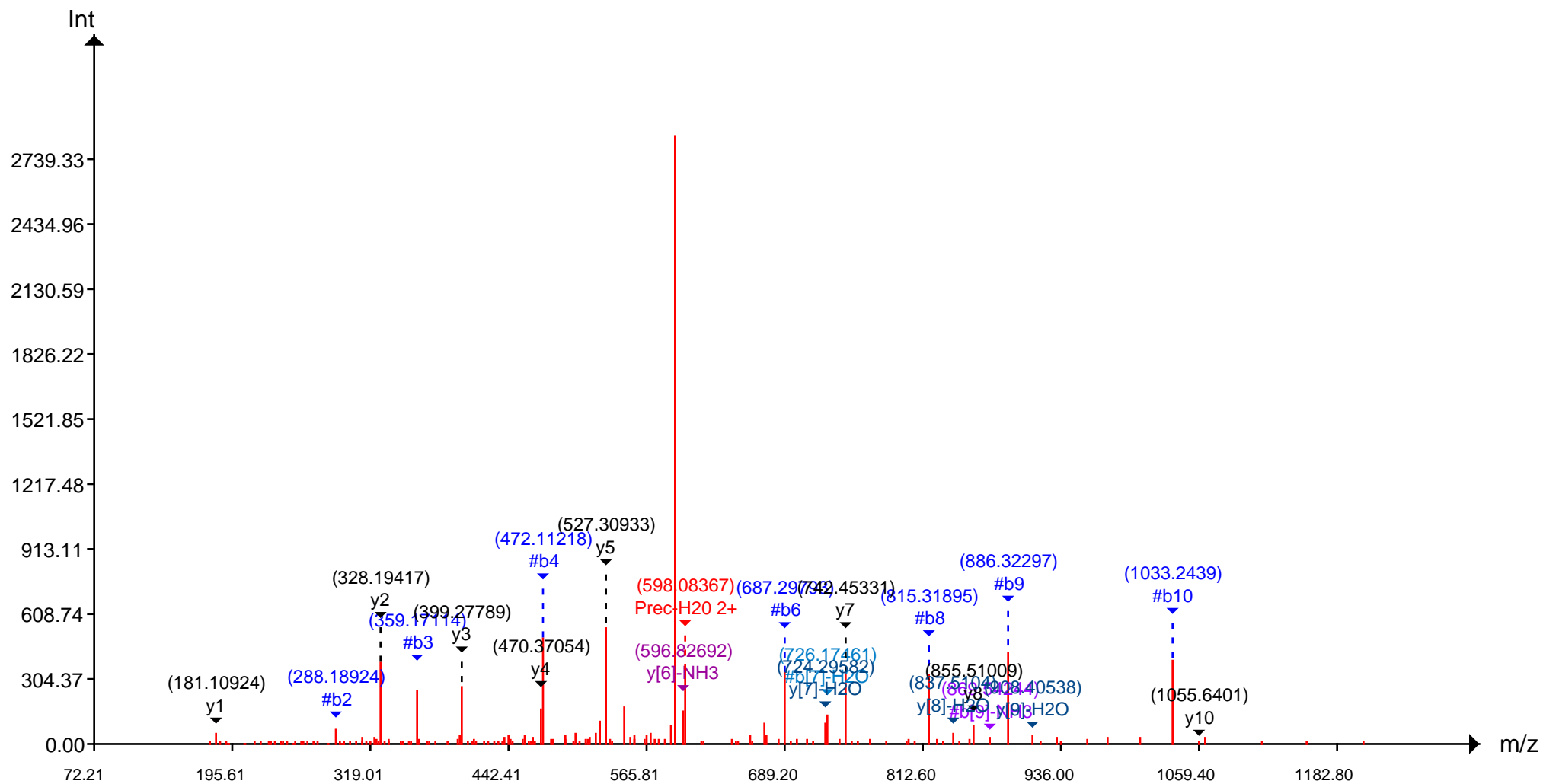
Confidence level 99.00%

Precursor m/z 758.9248

Identity Threshold 47.20

Charge 2+

Mass error 0.00117



Info

Modified Sequence AcD3-LEALQSGAAFR<C13>-COOH

Ion score 55.51

Protein O08808, O70566, Q9Z207

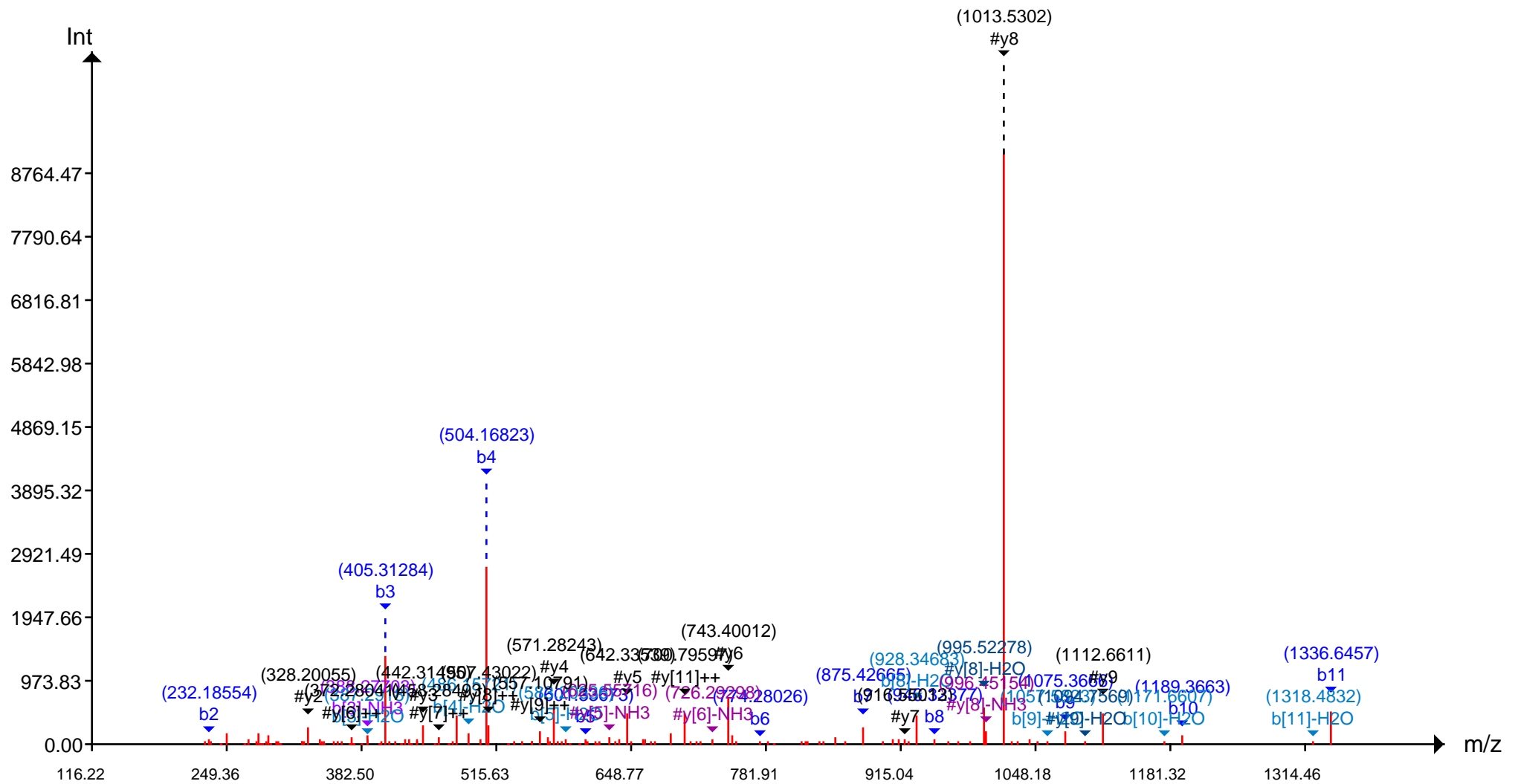
Confidence level 99.00%

Precursor m/z 607.33894

Identity Threshold 47.49

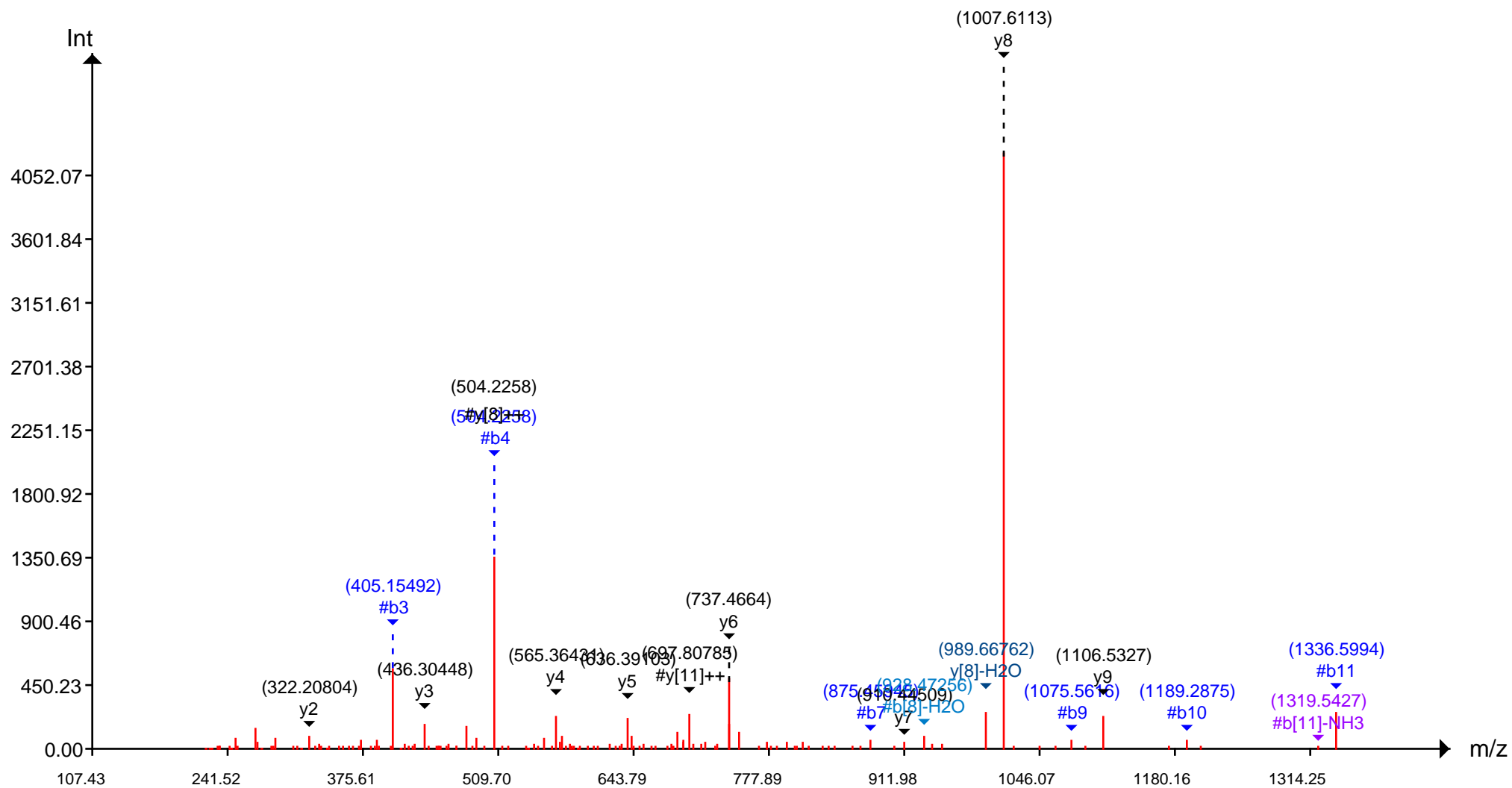
Charge 2+

Mass error -3.99E-4



Info

Modified Sequence	AcD3-ADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH	Ion score	67.63
Protein	P17742	Confidence level	99.00%
Precursor m/z	758.92525	Identity Threshold	47.18
Charge	2+	Mass error	0.00207



Info

Modified Sequence AcD3-ADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH

Ion score 66.44

Protein P17742

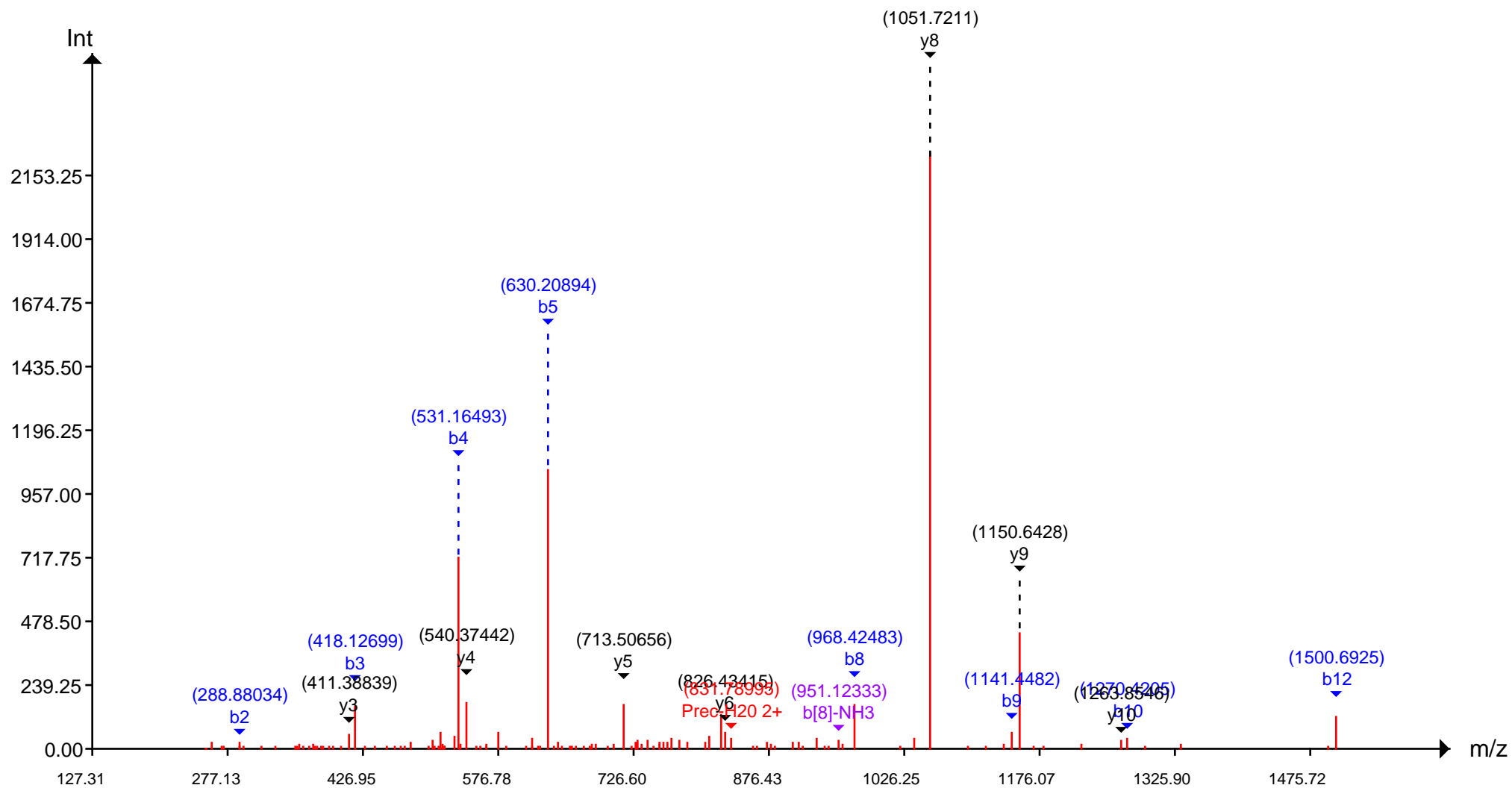
Confidence level 99.00%

Precursor m/z 755.91523

Identity Threshold 47.58

Charge 2+

Mass error 0.002172



Info

Modified Sequence AcD3-NEEIVPQIK<AcD3K*>EGK<AcD3K*>R<C13>-COOH

Ion score 62.01

Protein Q9DBJ1

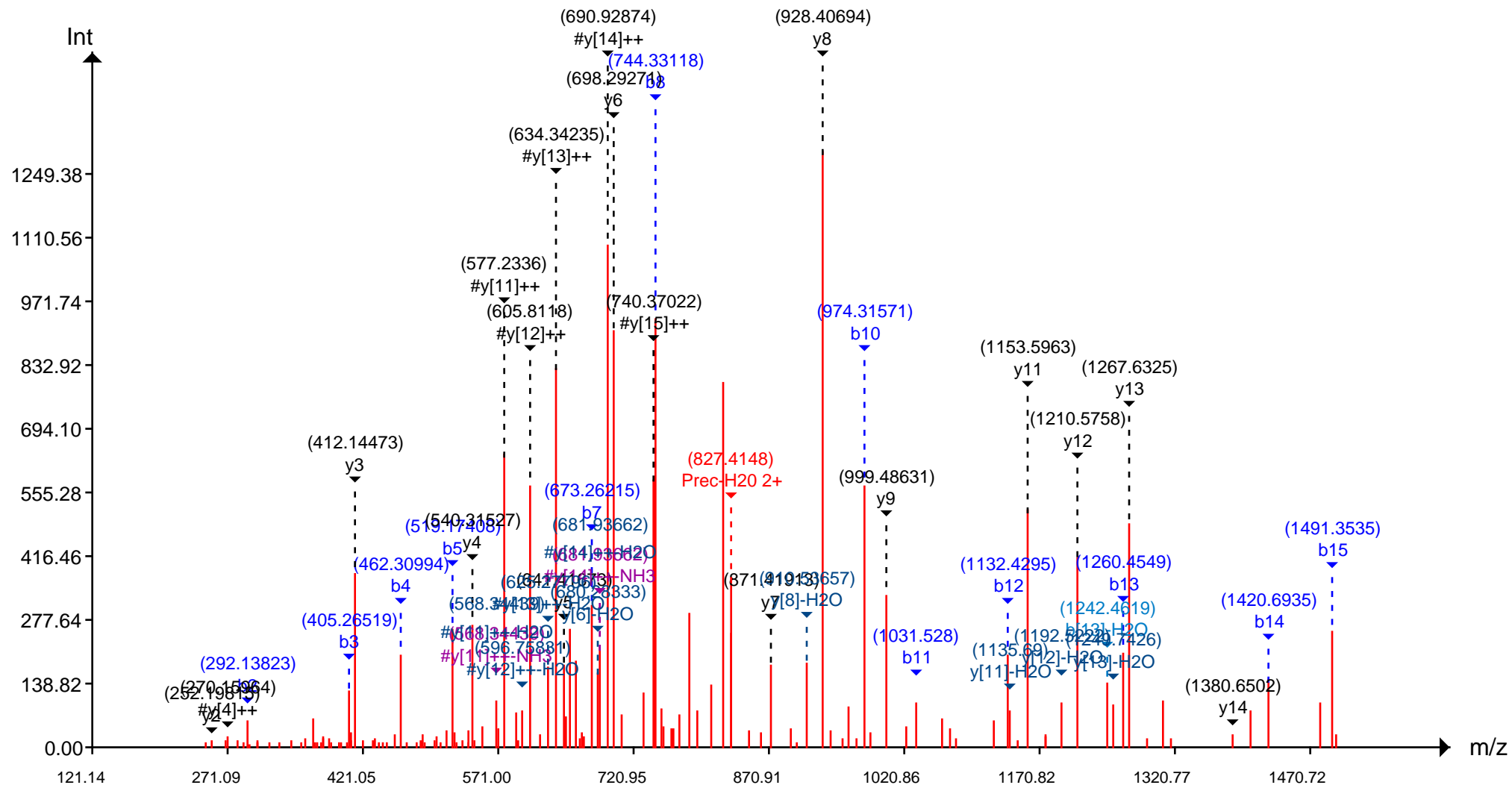
Confidence level 99.00%

Precursor m/z 840.98272

Identity Threshold 45.84

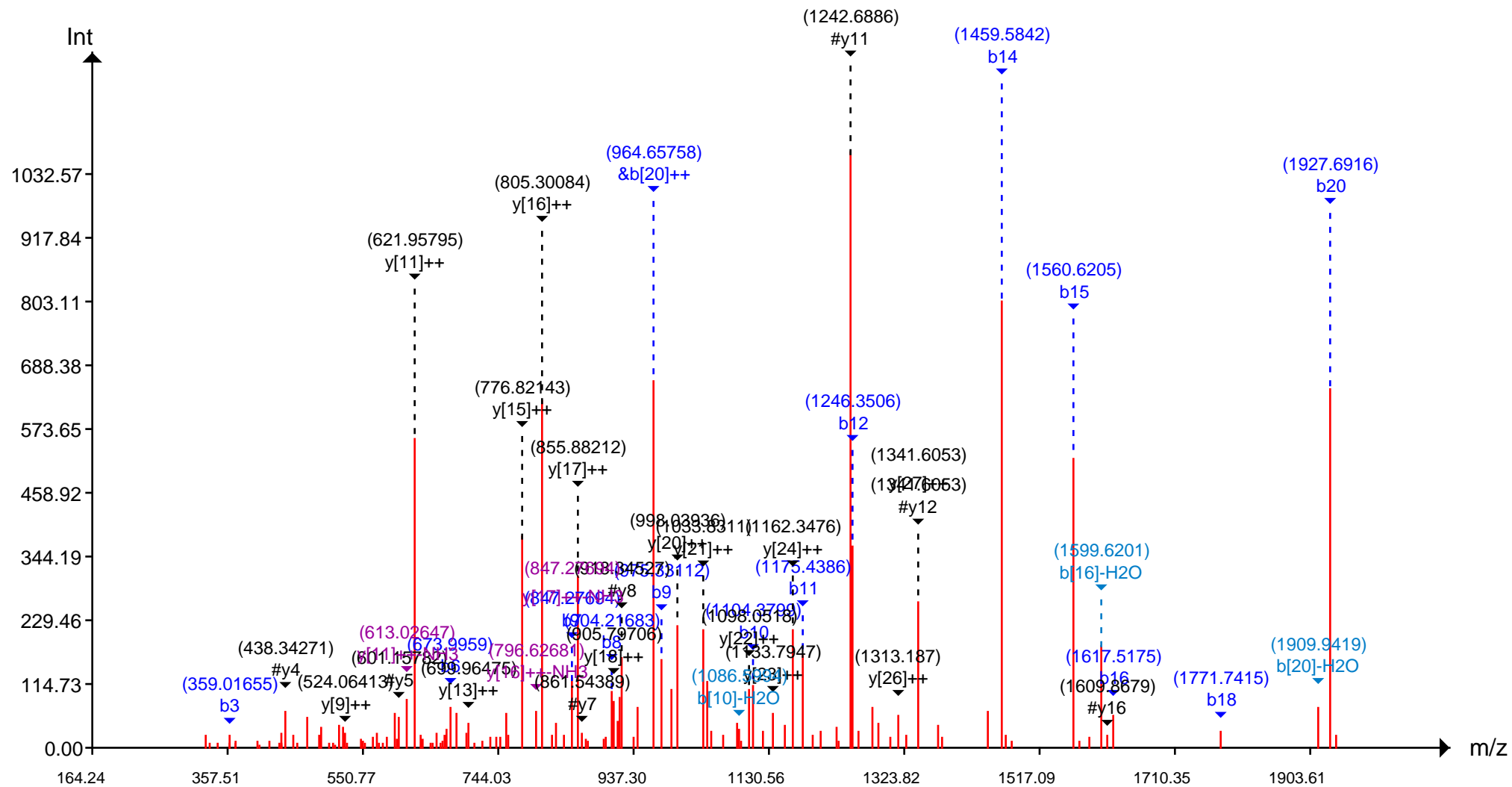
Charge 2+

Mass error 9.37E-4



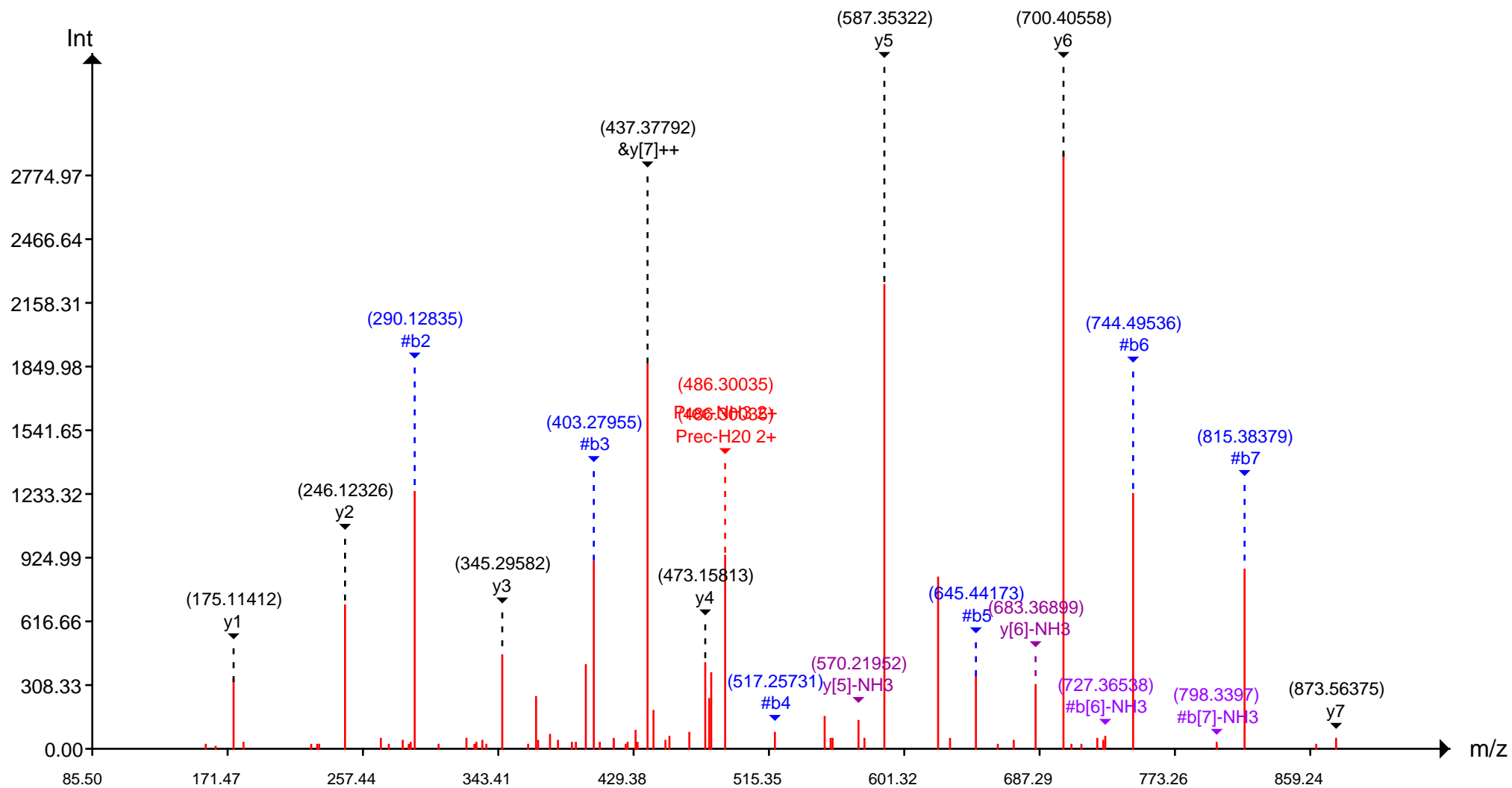
Info

Modified Sequence	AcD3-FVLGGPGAGK<AcD3K*>GTQC<Cmm*>AR<C13>-COOH	Ion score	104.99
Protein	Q9DBP5	Confidence level	99.00%
Precursor m/z	836.44703	Identity Threshold	48.01
Charge	2+	Mass error	0.001807



Info

Modified Sequence	AcD3-DVVEGEK<AcD3K*>GAEAANVTGPGGVPVQGSK<AcD3K*>YAADR<C13>-COOH	Ion score	83.26
Protein	P62960	Confidence level	99.00%
Precursor m/z	1057.2077	Identity Threshold	47.40
Charge	3+	Mass error	0.003173



Info

Modified Sequence AcD3-AK<AcD3K*>LNQVAR-COOH

Ion score 65.39

Protein P60335

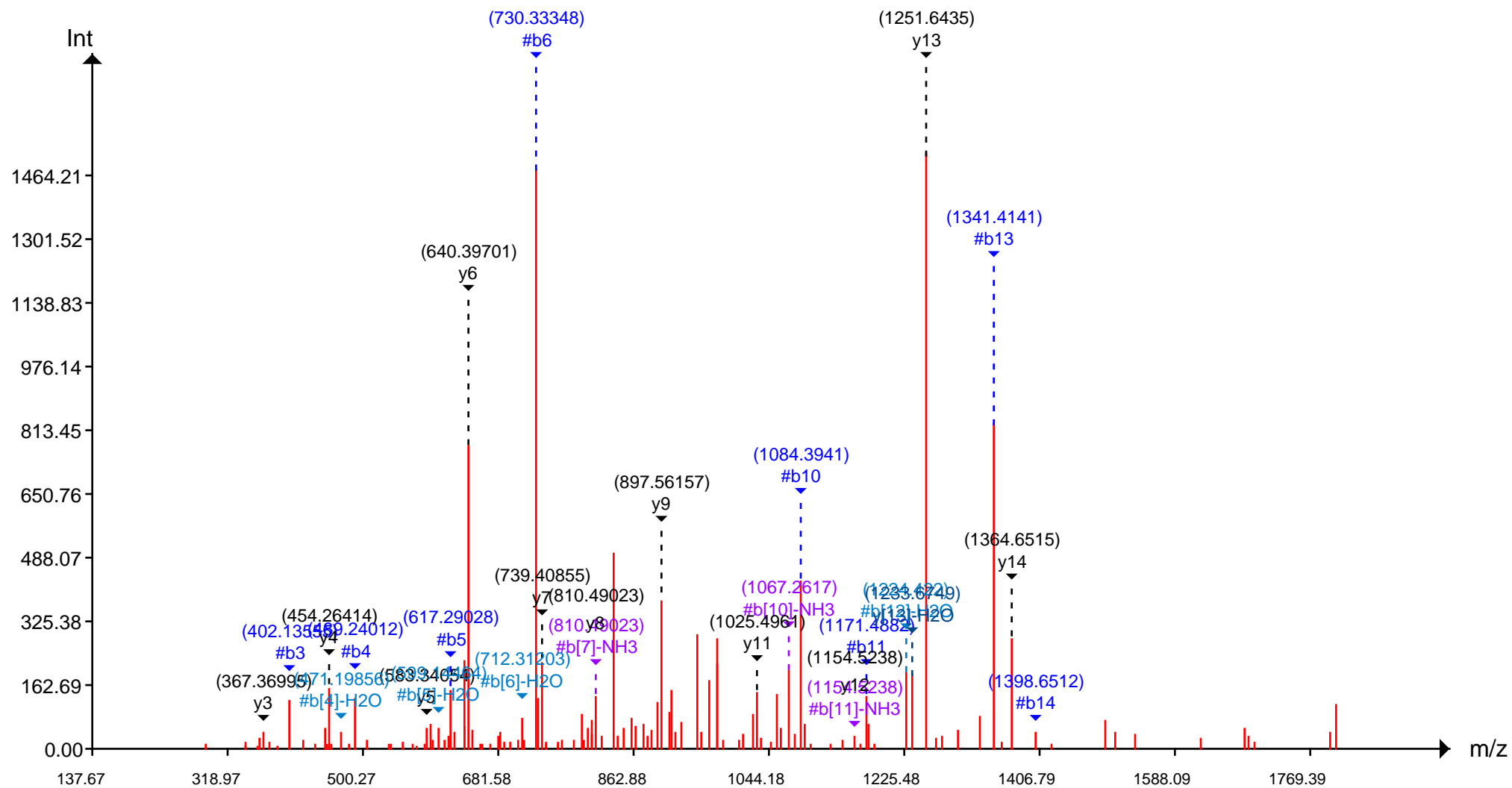
Confidence level 99.00%

Precursor m/z 495.30364

Identity Threshold 44.72

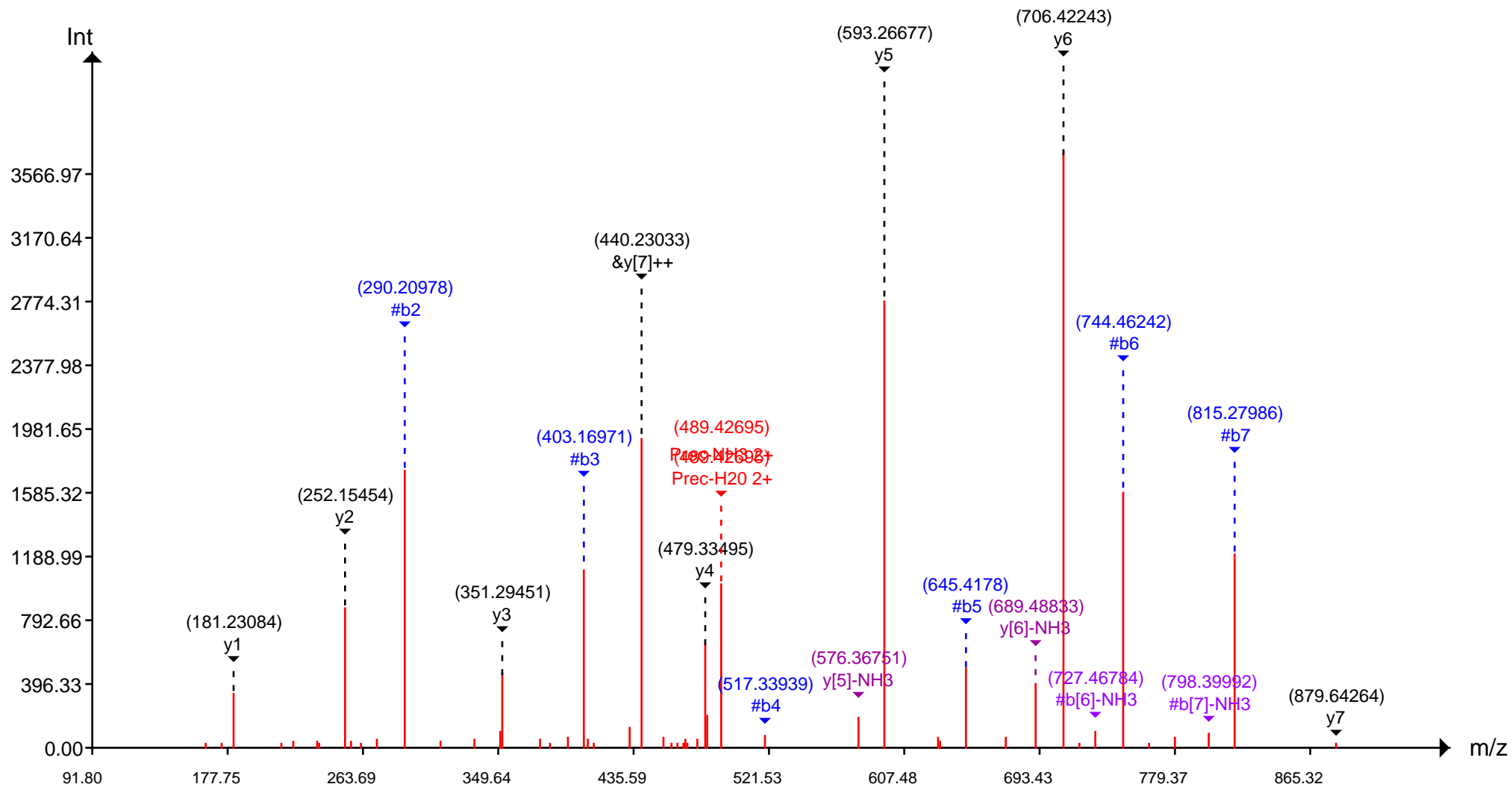
Charge 2+

Mass error -8.85E-4



Info

Modified Sequence	AcD3-DQLSQLPEGASAVGESSVR<C13>-COOH	Ion score	65.89
Protein	Q921W0	Confidence level	99.00%
Precursor m/z	991.0054	Identity Threshold	47.97
Charge	2+	Mass error	0.002779



Info

Modified Sequence AcD3-AK<AcD3K*>LNQVAR<C13>-COOH

Ion score 72.82

Protein P60335

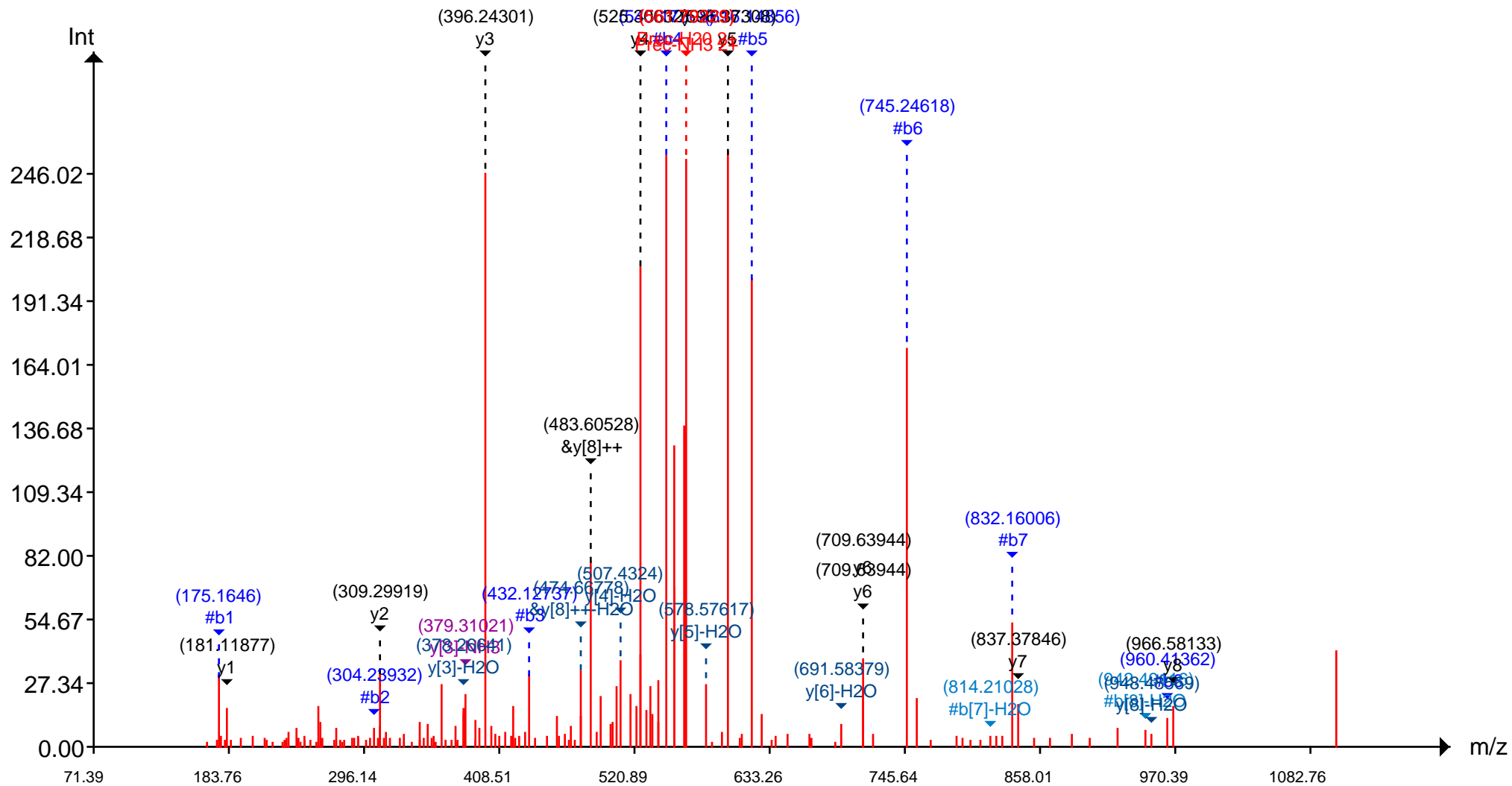
Confidence level 99.00%

Precursor m/z 498.3135

Identity Threshold 43.31

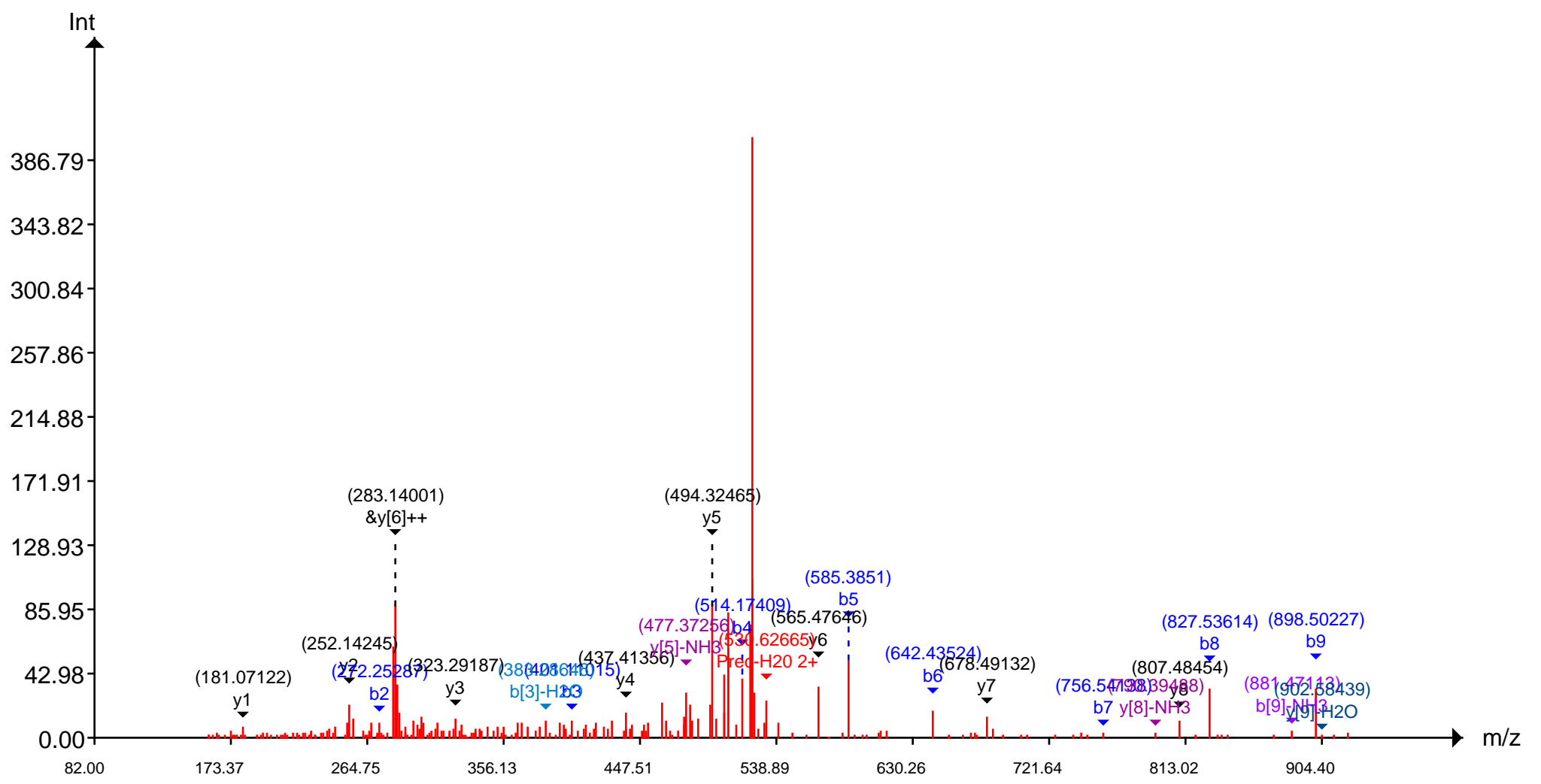
Charge 2+

Mass error -0.001306

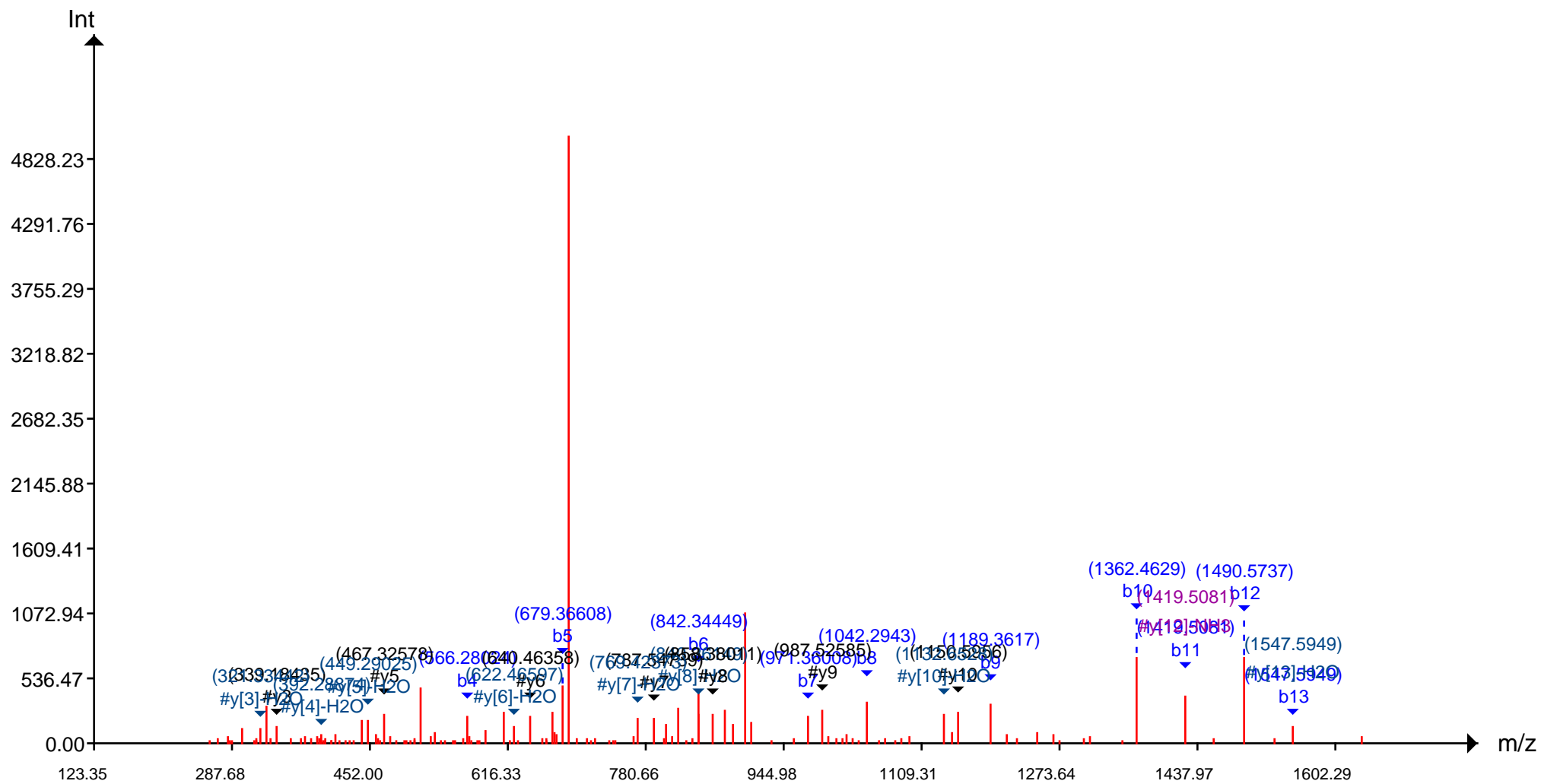


Info

Modified Sequence	AcD3-EEQLAESQR<C13>-COOH	Ion score	69.91
Protein	Q3UZ39	Confidence level	99.00%
Precursor m/z	570.78689	Identity Threshold	47.17
Charge	2+	Mass error	-8.3E-5

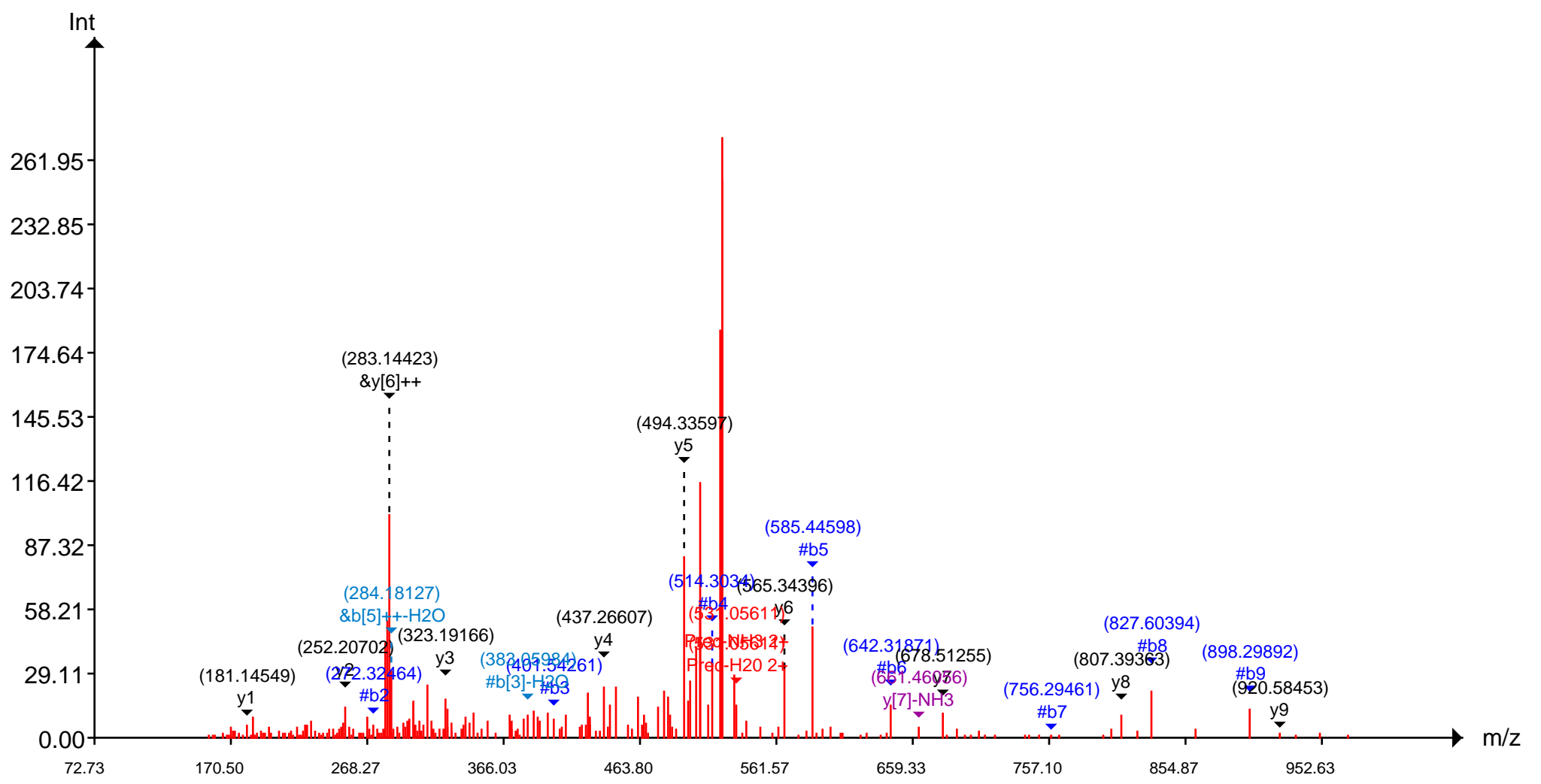


Info	
Modif...	AcD3-ILELAGNAAR<C13>-COOH
	Q8CGP5, Q8CGP6, Q8CGP7, P22752, Q6GSS7, Q64522, Q64523, Q8BFU2, Q8R1M2, Q8CCK0, P27661, Q9QZQ8
	539.82265
	2+
	Ion score
	Confidence level
	Identity Threshold
	Mass error

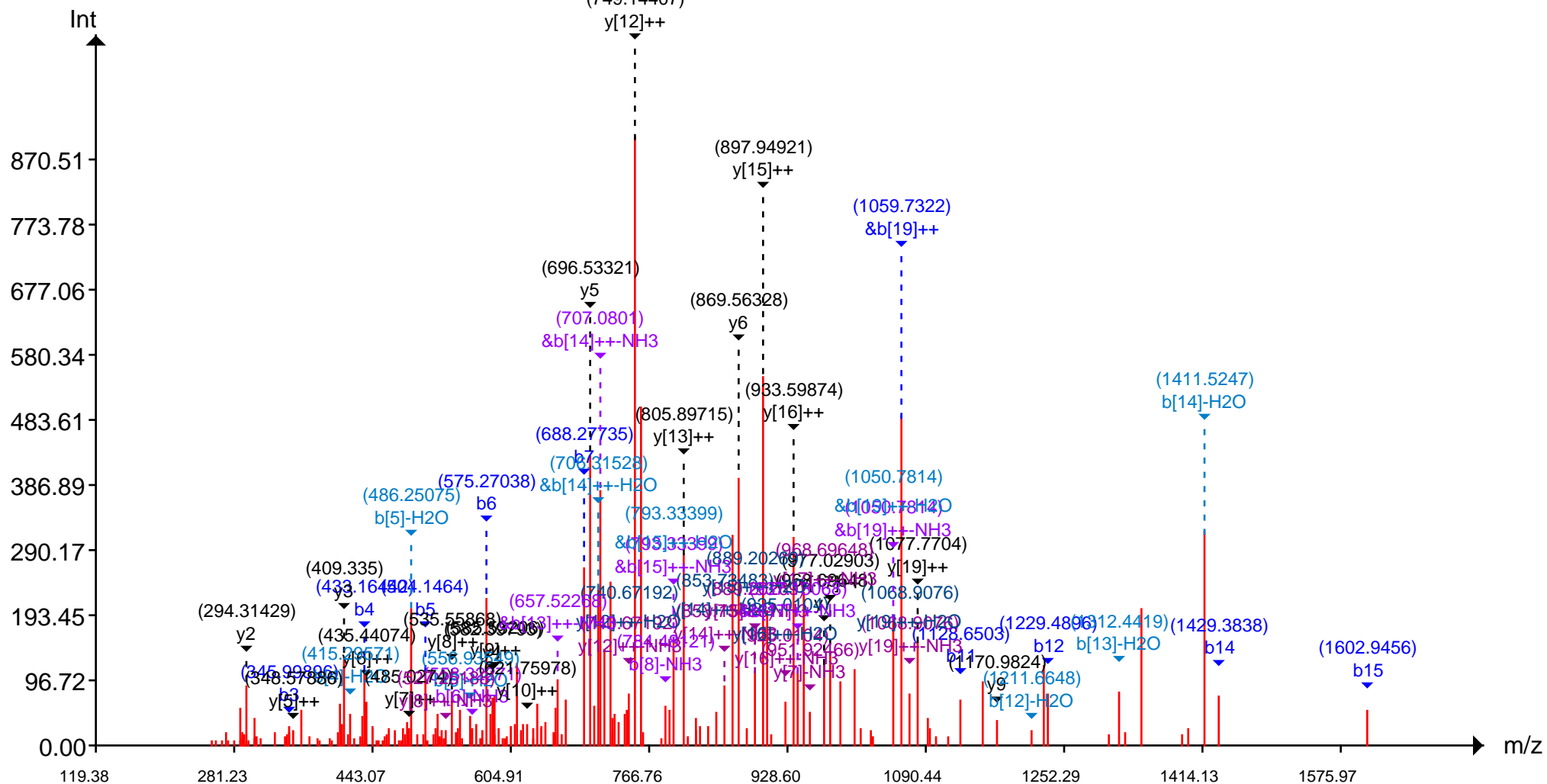


Info

Modified Sequence	AcD3-FAEK<AcD3K*>LYEAM<Mox*>K<AcD3K*>GAGTR<C13>-COOH	Ion score	57.14
Protein	P10107	Confidence level	99.00%
Precursor m/z	914.98302	Identity Threshold	48.12
Charge	2+	Mass error	0.003246

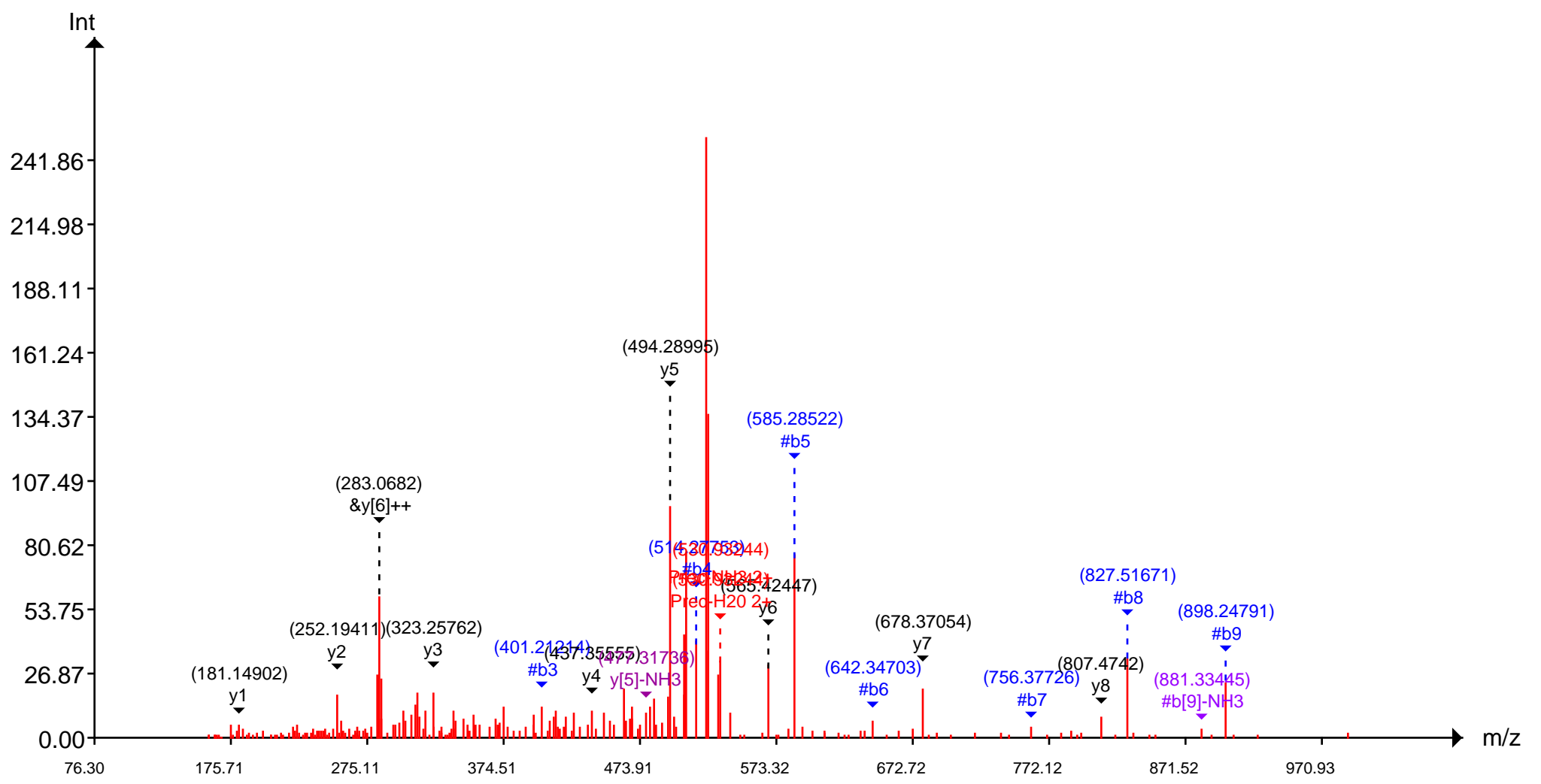


Info		Ion score
Modif...	AcD3-ILELAGNAAR<C13>-COOH	
	Q8CGP5, Q8CGP6, Q8CGP7, P22752, Q6GSS7, Q64522, Q64523, Q8BFU2, Q8R1M2, Q8CCK0, P27661, Q9QZQ8	Confidence level
	539.82274	Identity Threshold
	2+	Mass error

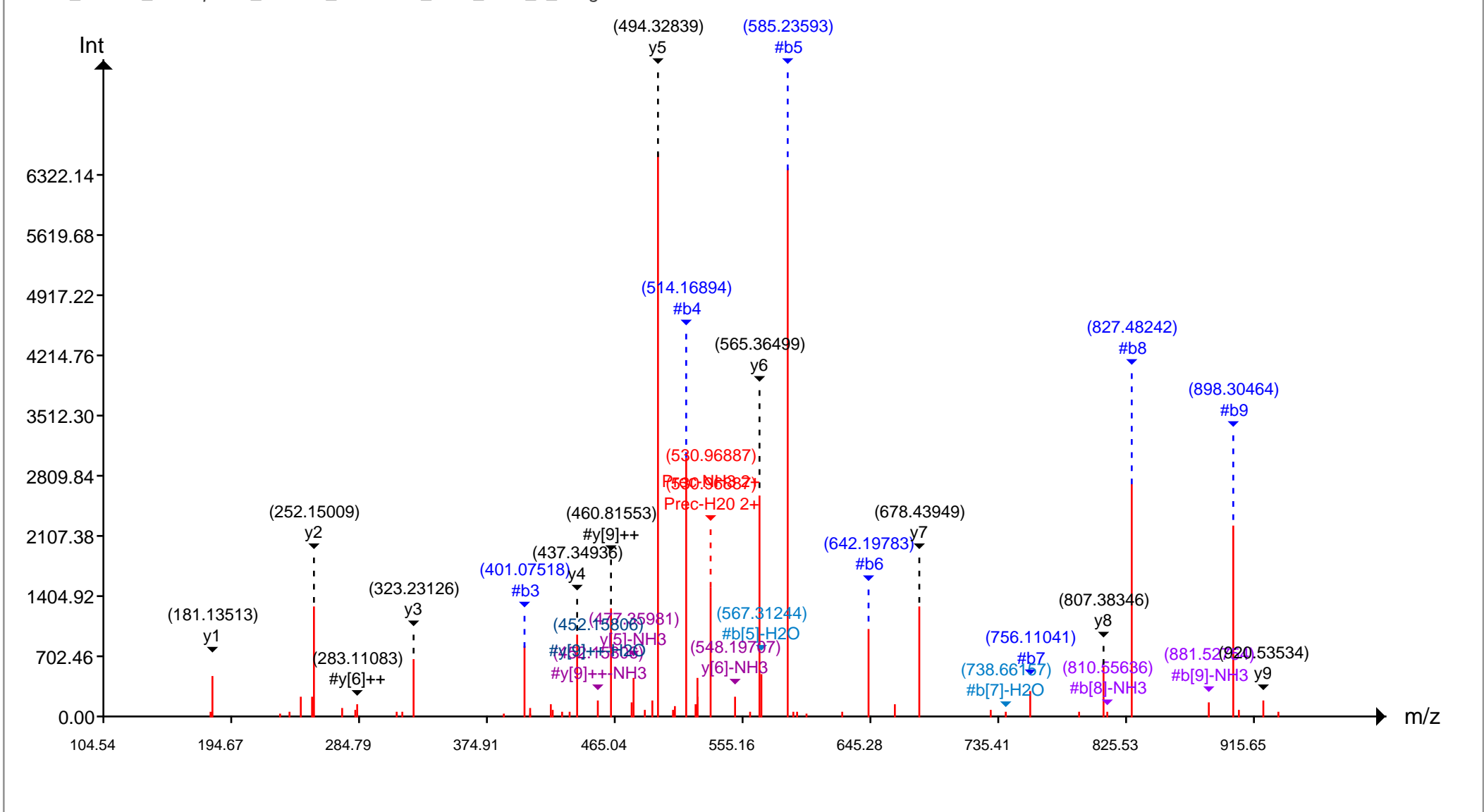


Info

Modified Sequence	AcD3-VSNSAALIQQATTVK<AcD3K*>NK<AcD3K*>DIR<C13>-COOH	Ion score	67.12
Protein	P51410	Confidence level	99.00%
Precursor m/z	766.77436	Identity Threshold	44.45
Charge	3+	Mass error	0.001615



Info		Ion score
Modif...	AcD3-ILELAGNAAR<C13>-COOH	
	Q8CGP5, Q8CGP6, Q8CGP7, P22752, Q6GSS7, Q64522, Q64523, Q8BFU2, Q8R1M2, Q8CCK0, P27661, Q9QZQ8	Confidence level
	539.82294	Identity Threshold
	2+	Mass error



Info

Modif... AcD3-ILELAGNAAR<C13>-COOH

Ion score

Q8CGP5, Q8CGP6, Q8CGP7, P22752, Q6GSS7, Q64522, Q64523, Q8BFU2, Q8R1M2, Q8CCK0, P27661, Q9QZQ8

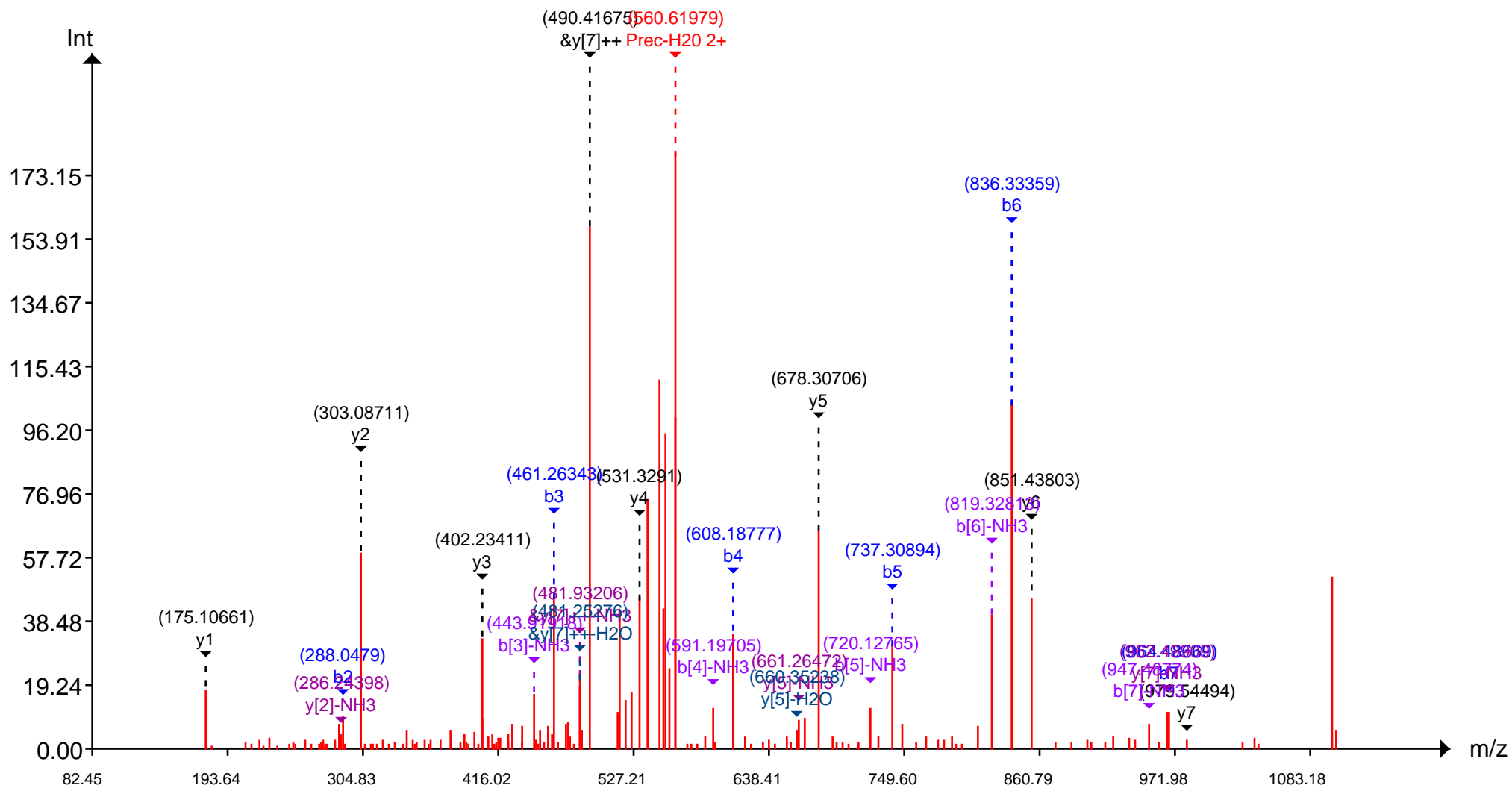
Confidence level

539.82304

Identity Threshold

2+

Mass error



Info

Modified Sequence AcD3-NQK<AcD3K*>M<Mox*>EVQR-COOH

Ion score 48.38

Protein P16675

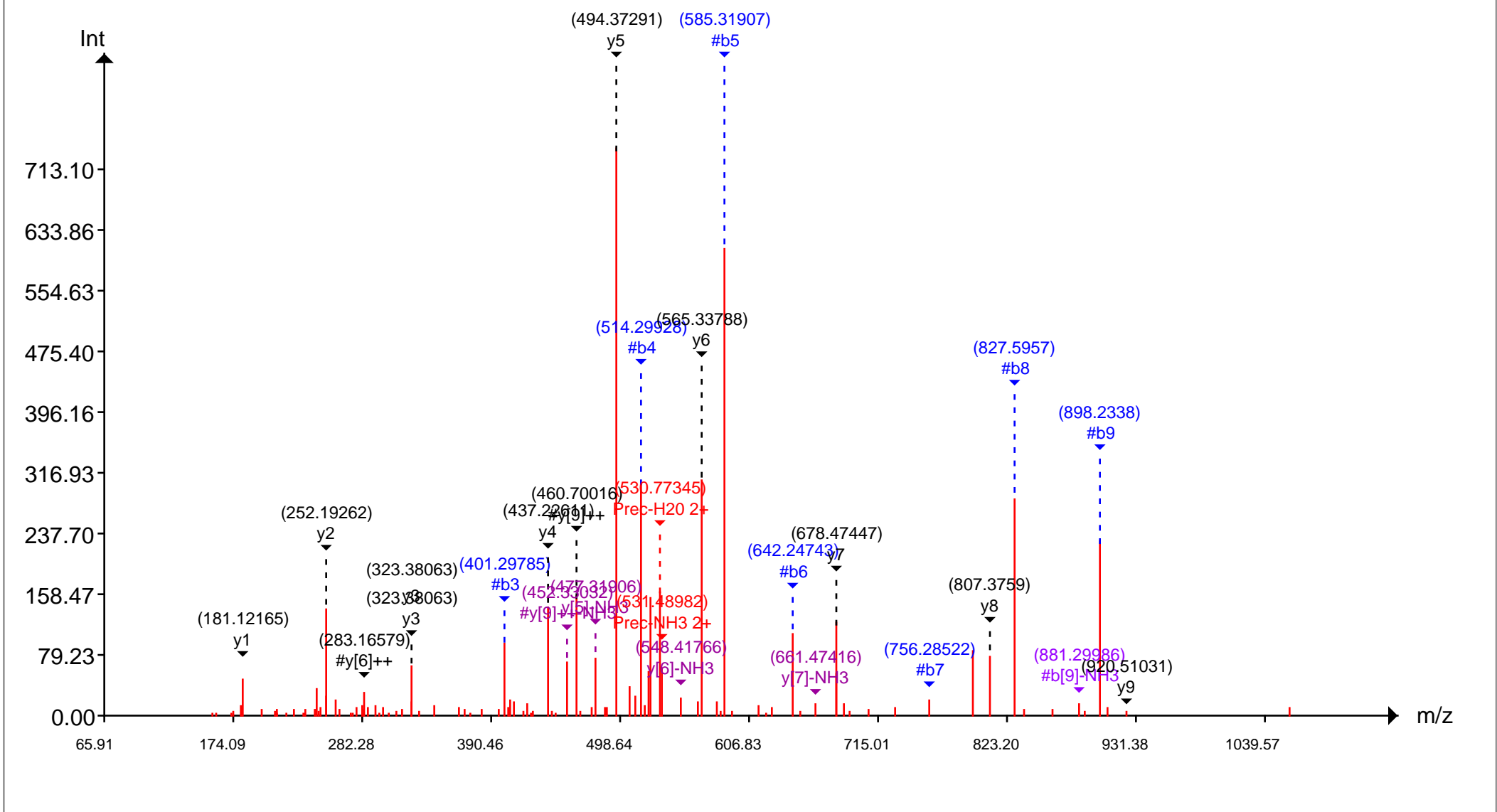
Confidence level 99.00%

Precursor m/z 569.79276

Identity Threshold 47.80

Charge 2+

Mass error -9.31E-4



Info

Modif... AcD3-ILELAGNAAR<C13>-COOH

Ion score

Q8CGP5, Q8CGP6, Q8CGP7, P22752, Q6GSS7, Q64522, Q64523, Q8BFU2, Q8R1M2, Q8CCK0, P27661, Q9QZQ8

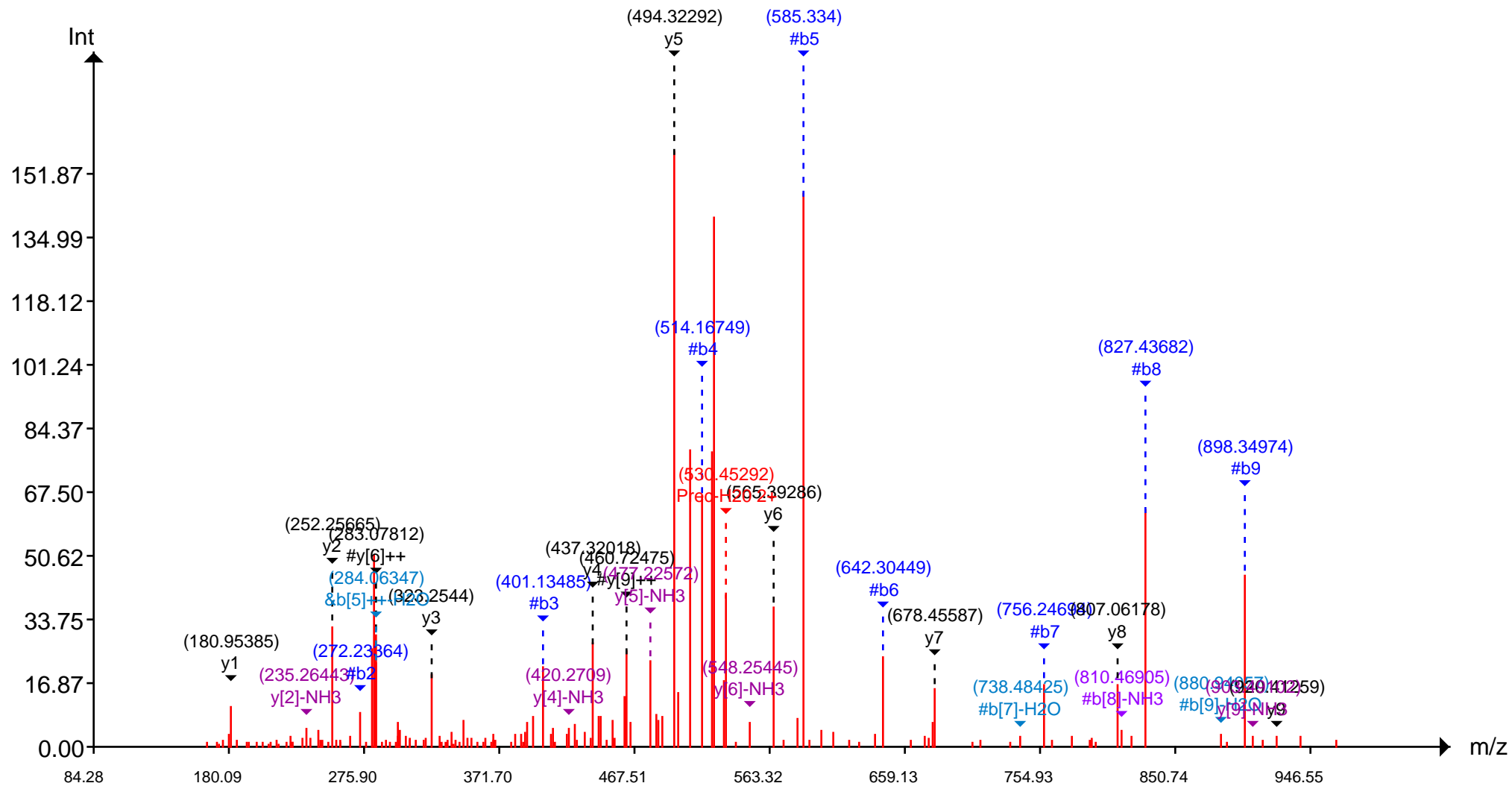
Confidence level

539.82303

Identity Threshold

2+

Mass error



Info

Modif... AcD3-ILELAGNAAR<C13>-COOH

Ion score

Q8CGP5, Q8CGP6, Q8CGP7, P22752, Q6GSS7, Q64522, Q64523, Q8BFU2, Q8R1M2, Q8CCK0, P27661, Q9QZQ8

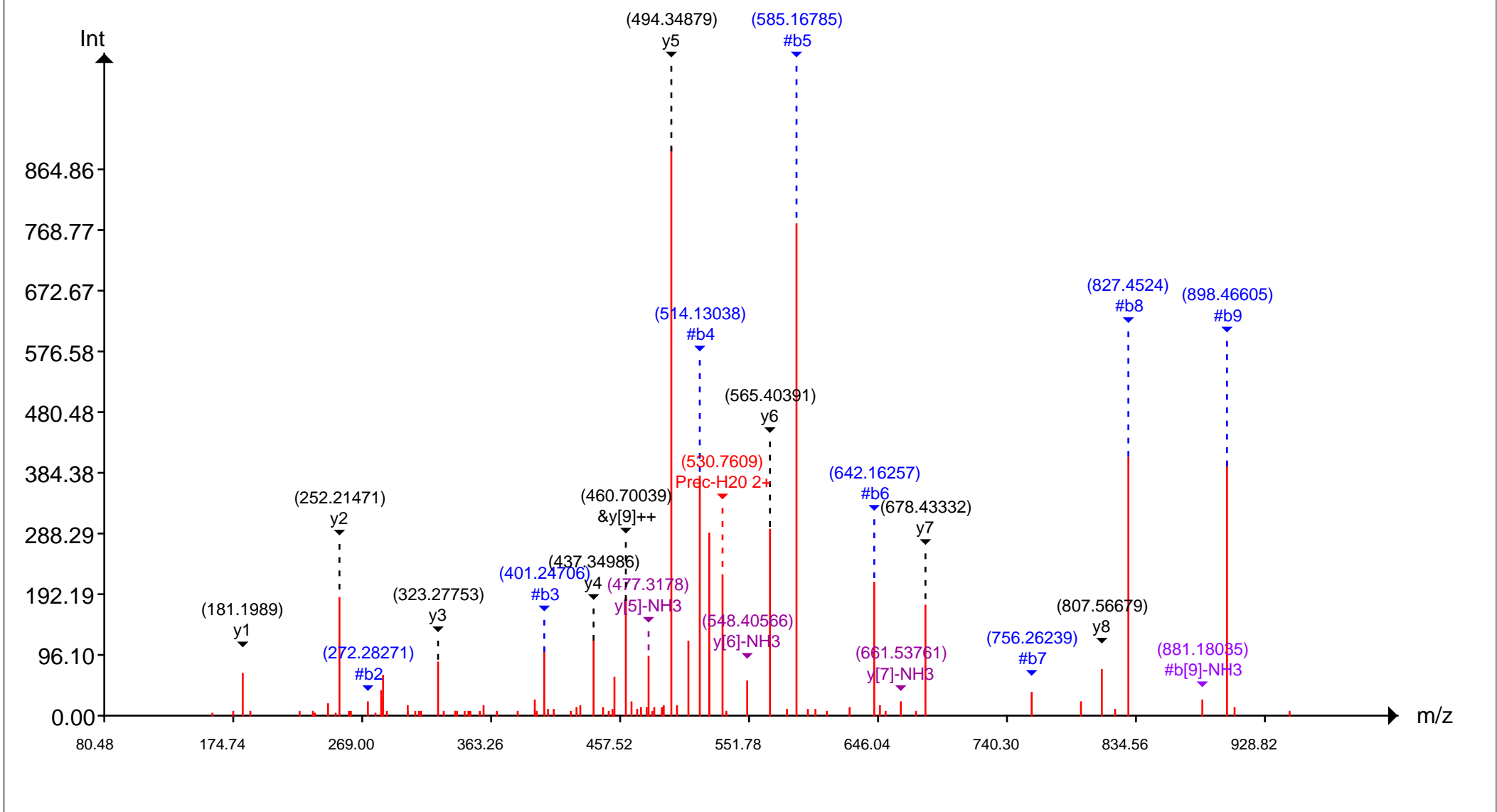
Confidence level

P... 539.82311

Identity Threshold

2+

Mass error



Info

Modif... AcD3-ILELAGNAAR<C13>-COOH

Ion score

Q... Q8CGP5, Q8CGP6, Q8CGP7, P22752, Q6GSS7, Q64522, Q64523, Q8BFU2, Q8R1M2, Q8CCK0, P27661, Q9QZQ8

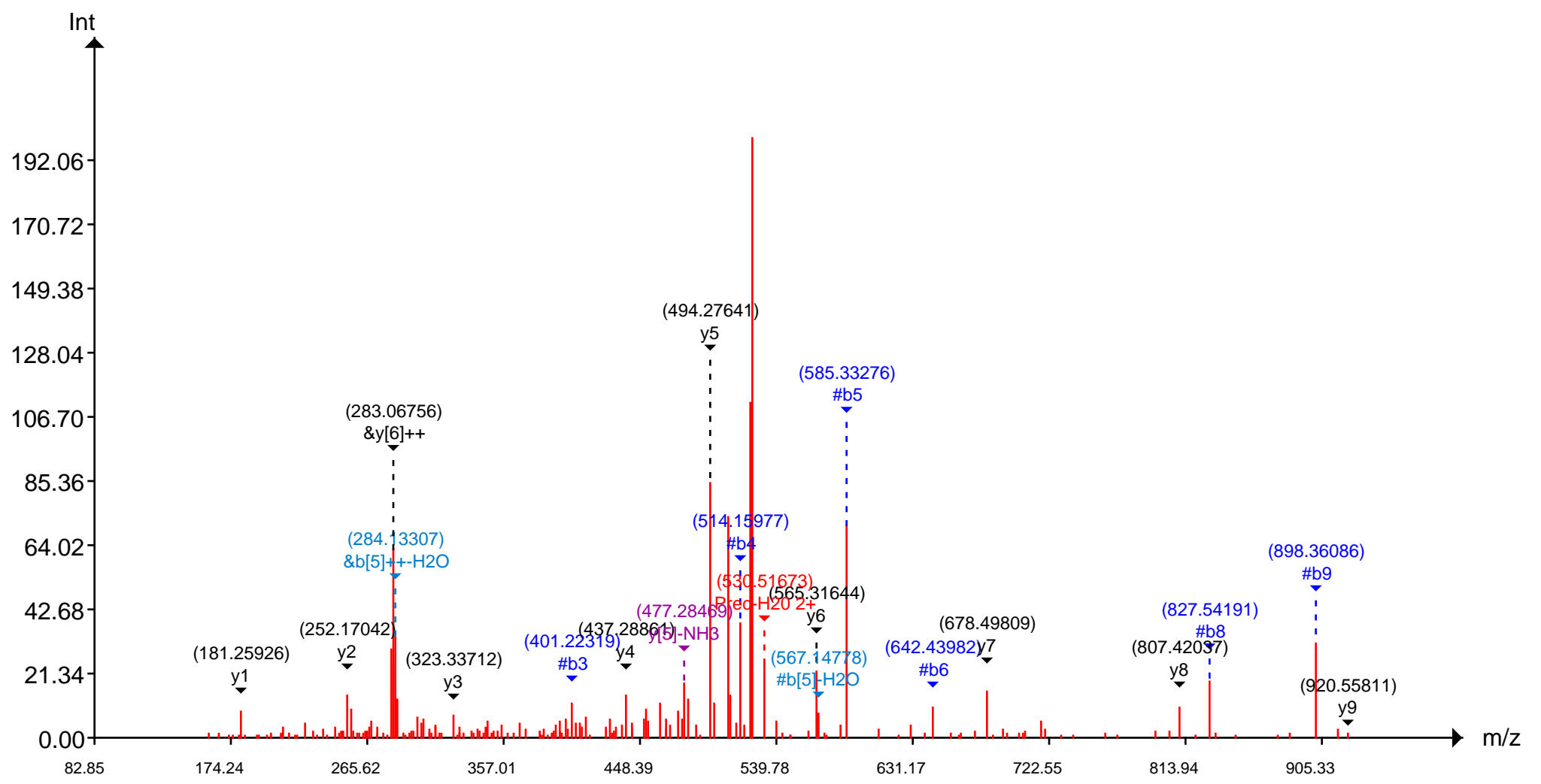
Confidence level

P... 539.82308

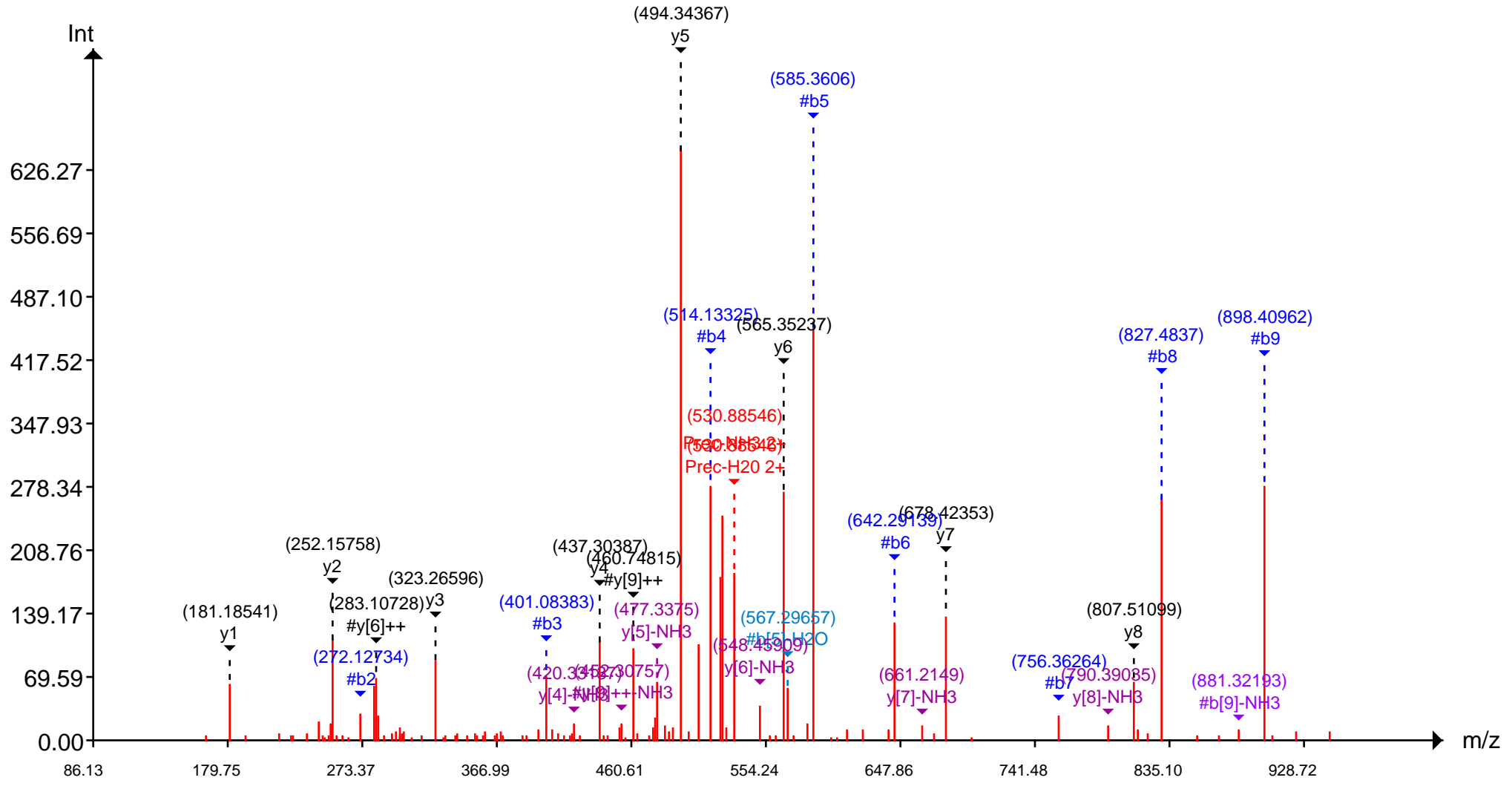
Identity Thresho

2+

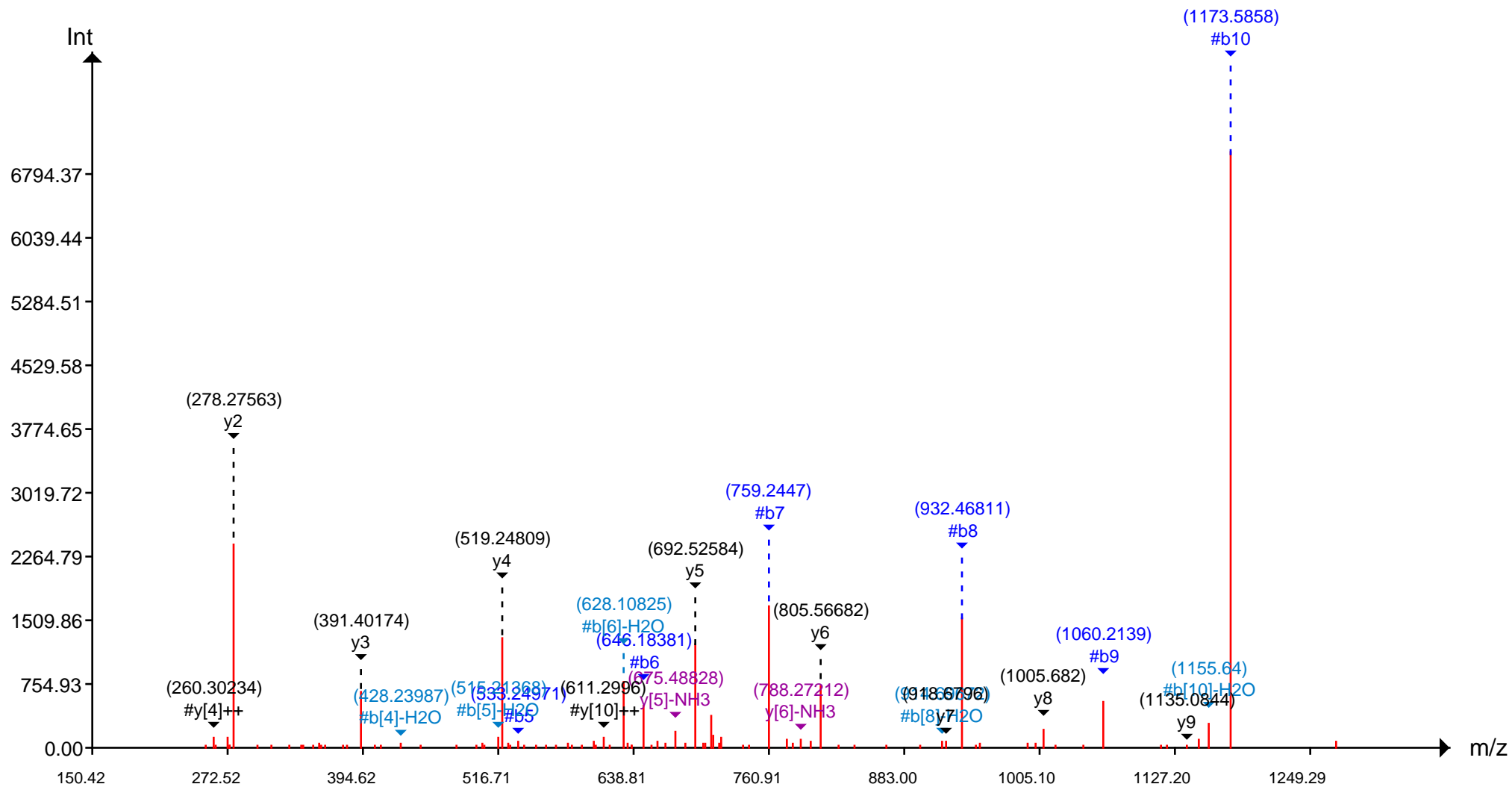
Mass error



Info	
Modif...	AcD3-ILELAGNAAR<C13>-COOH
P...	539.82314
	2+
Ion score	
Confidence level	Q8CGP5, Q8CGP6, Q8CGP7, P22752, Q6GSS7, Q64522, Q64523, Q8BFU2, Q8R1M2, Q8CCK0, P27661, Q9QZQ8
Identity Threshold	
Mass error	

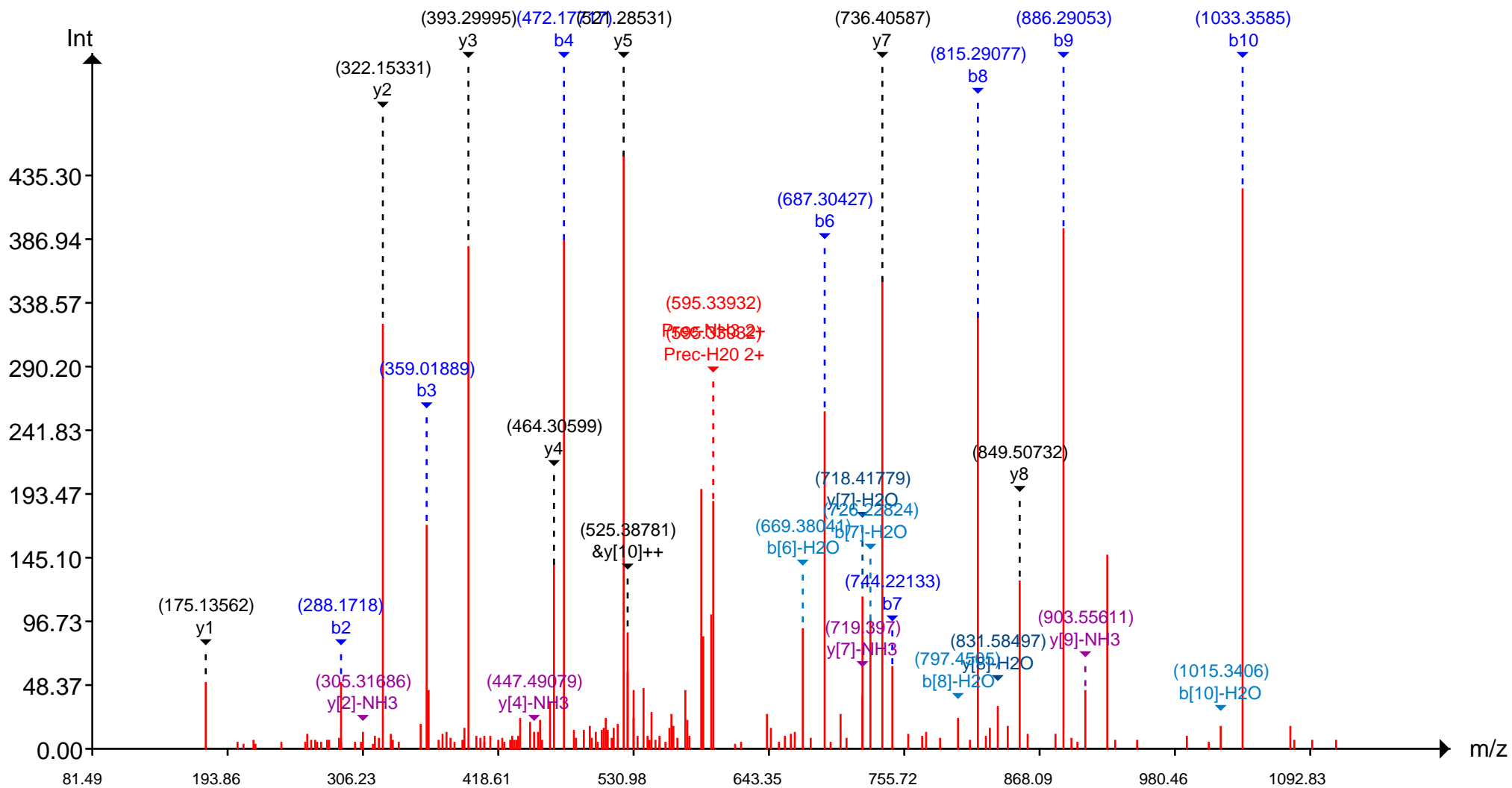


Info		
Modif...	AcD3-ILELAGNAAR<C13>-COOH	Ion score
P...	Q8CGP5, Q8CGP6, Q8CGP7, P22752, Q6GSS7, Q64522, Q64523, Q8BFU2, Q8R1M2, Q8CCK0, P27661, Q9QZQ8	Confidence level
	539.82311	Identity Thresho
	2+	Mass error



Info

Modified Sequence	AcD3-LASESLIK<AcD3K*>QIPR<C13>-COOH	Ion score	63.64
Protein	P53026	Confidence level	99.00%
Precursor m/z	725.94645	Identity Threshold	41.83
Charge	2+	Mass error	0.001456



Info

Modified Sequence AcD3-LEALQSGAAFR-COOH

Protein O08808, O70566, Q9Z207

Precursor m/z 604.32924

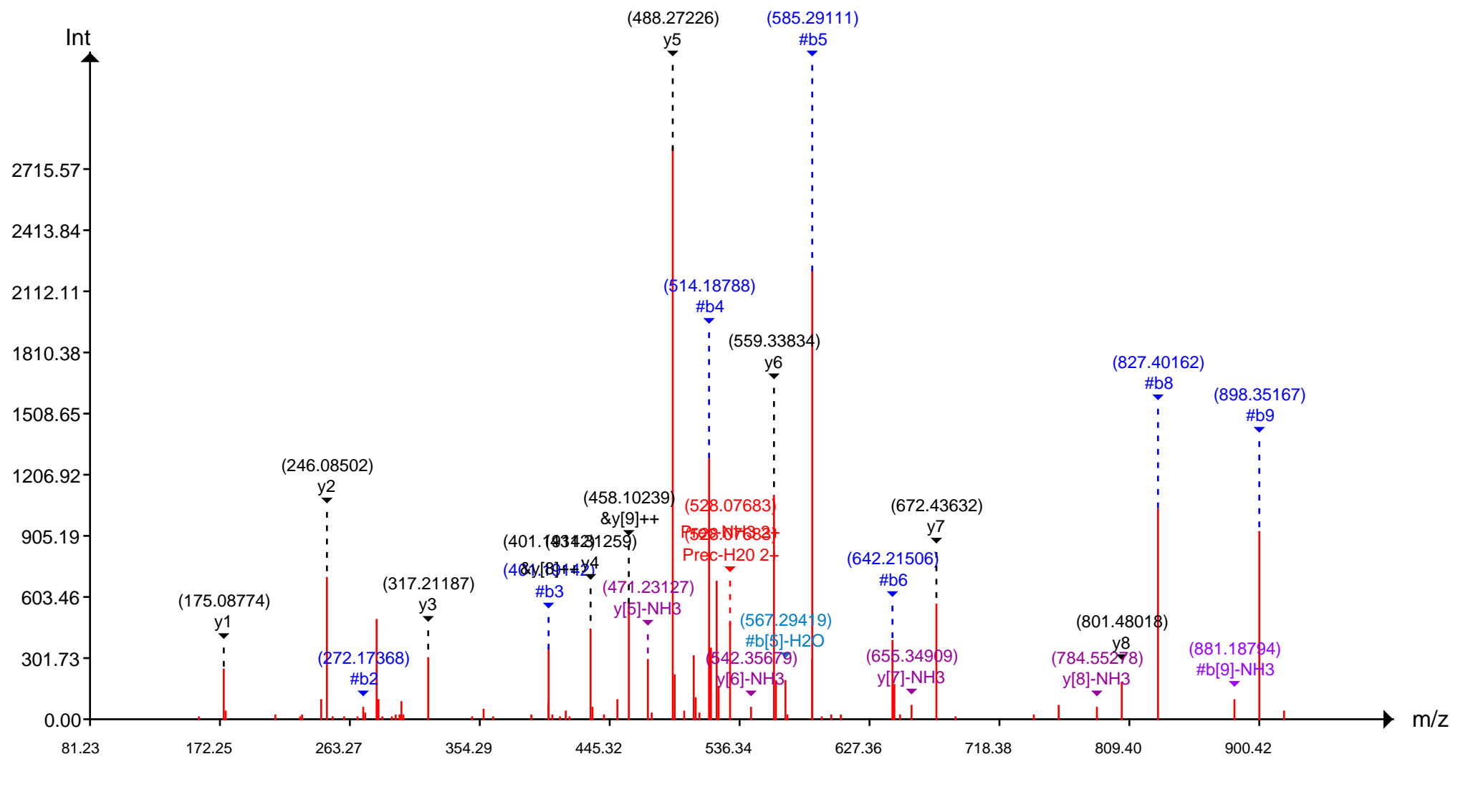
Charge 2+

Ion score 52.97

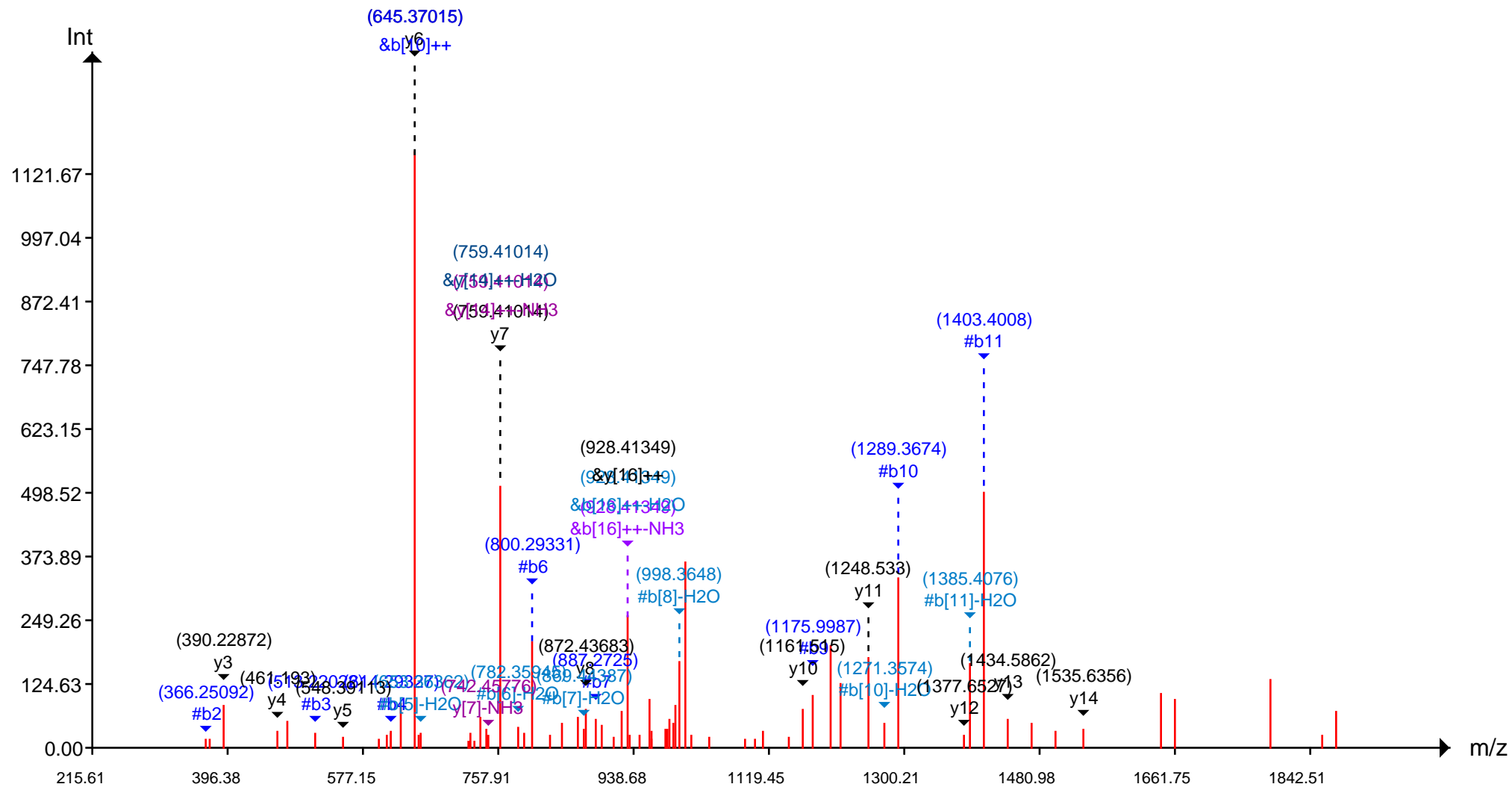
Confidence level 99.00%

Identity Threshold 47.98

Mass error 3.43E-4

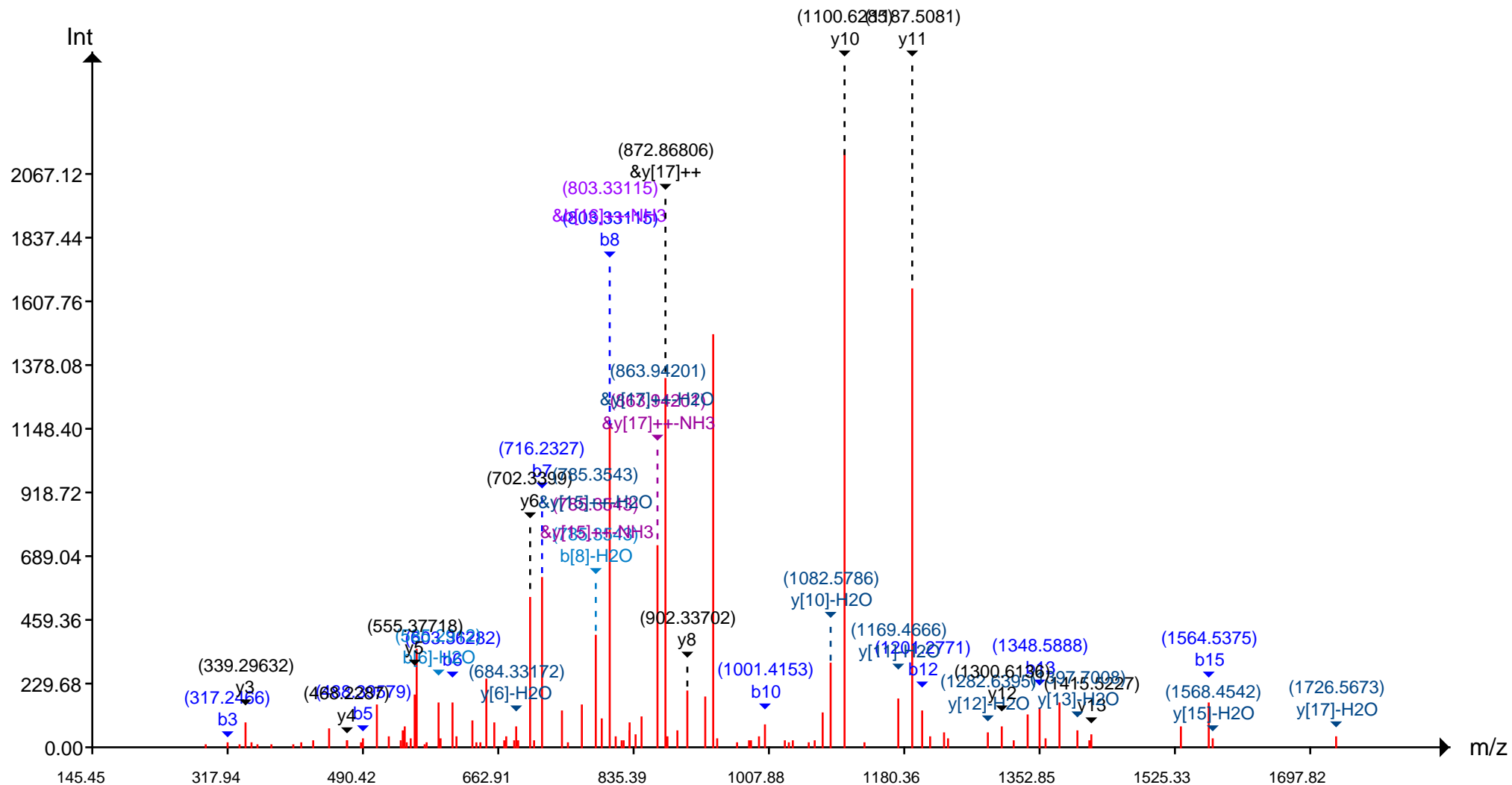


Info		Ion score
Modif...	AcD3-ILELAGNAAR-COOH	
P...	Q8CGP5, Q8CGP6, Q8CGP7, P22752, Q6GSS7, Q64522, Q64523, Q8BFU2, Q8R1M2, Q8CCK0, P27661, Q9QZQ8	Confidence level
	536.81328	Identity Thresho
	2+	Mass error



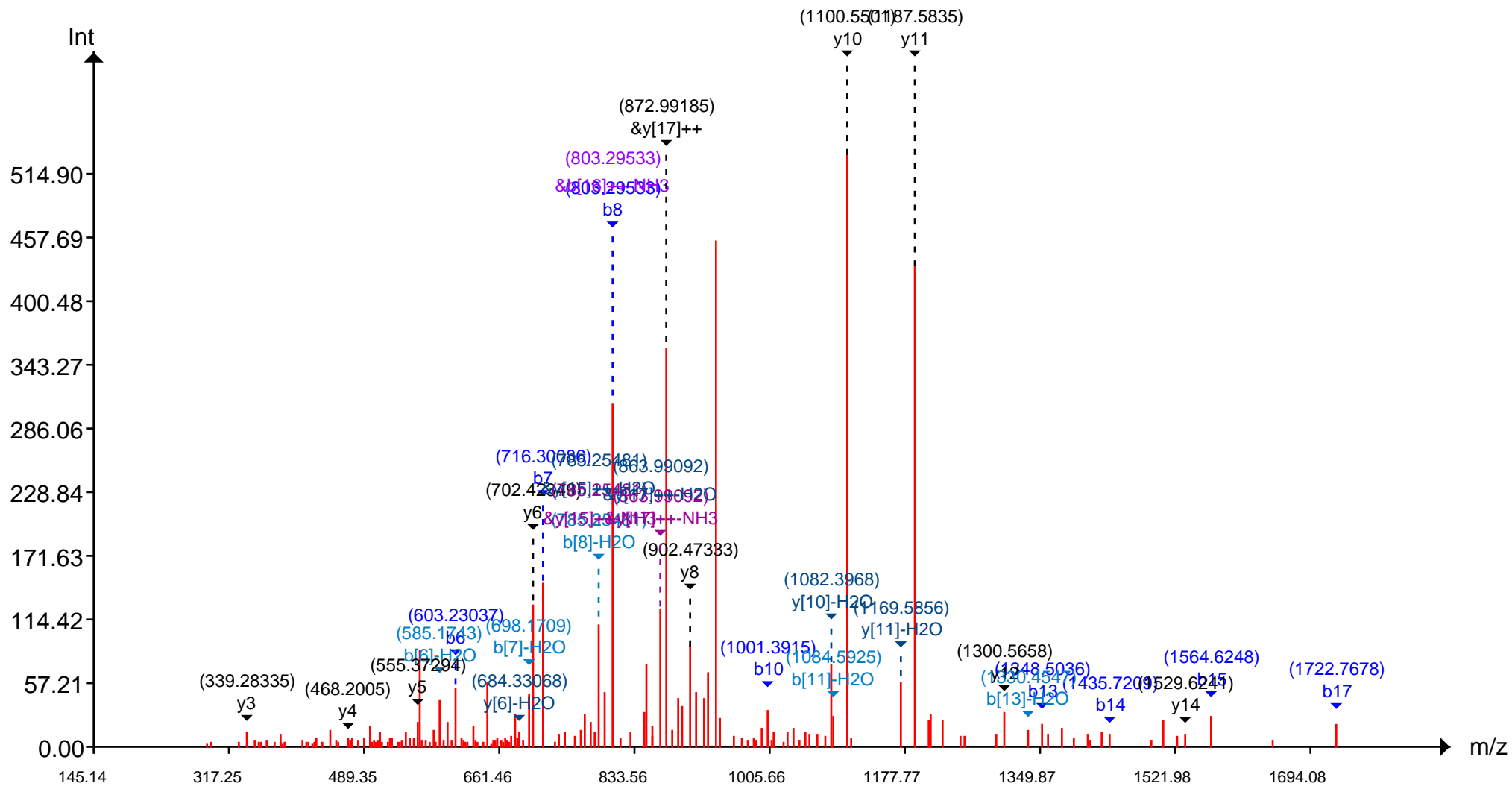
Info

Modified Sequence	AcD3-FK<AcD3K*>M<Mox*>TGESEC<Cmm*>LNPSAQSR-COOH	Ion score	63.88
Protein	O54988	Confidence level	99.00%
Precursor m/z	1024.4714	Identity Threshold	44.72
Charge	2+	Mass error	0.002513



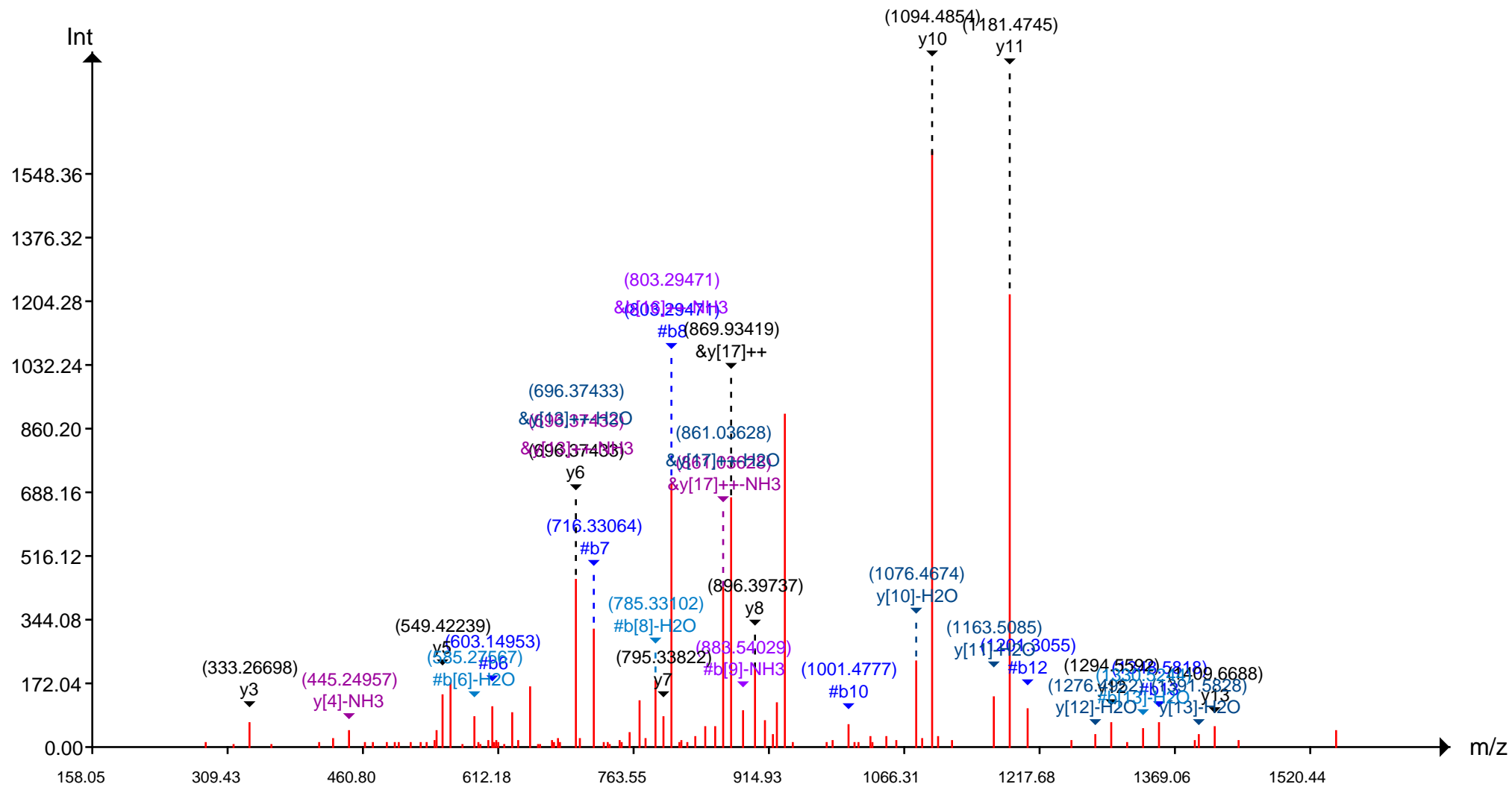
Info

Modified Sequence	AcD3-LGTGNDLSPTTVM<Mox*>SEGTR<C13>-COOH	Ion score	60.39
Protein	Q923D2	Confidence level	99.00%
Precursor m/z	951.96759	Identity Threshold	47.26
Charge	2+	Mass error	0.003071



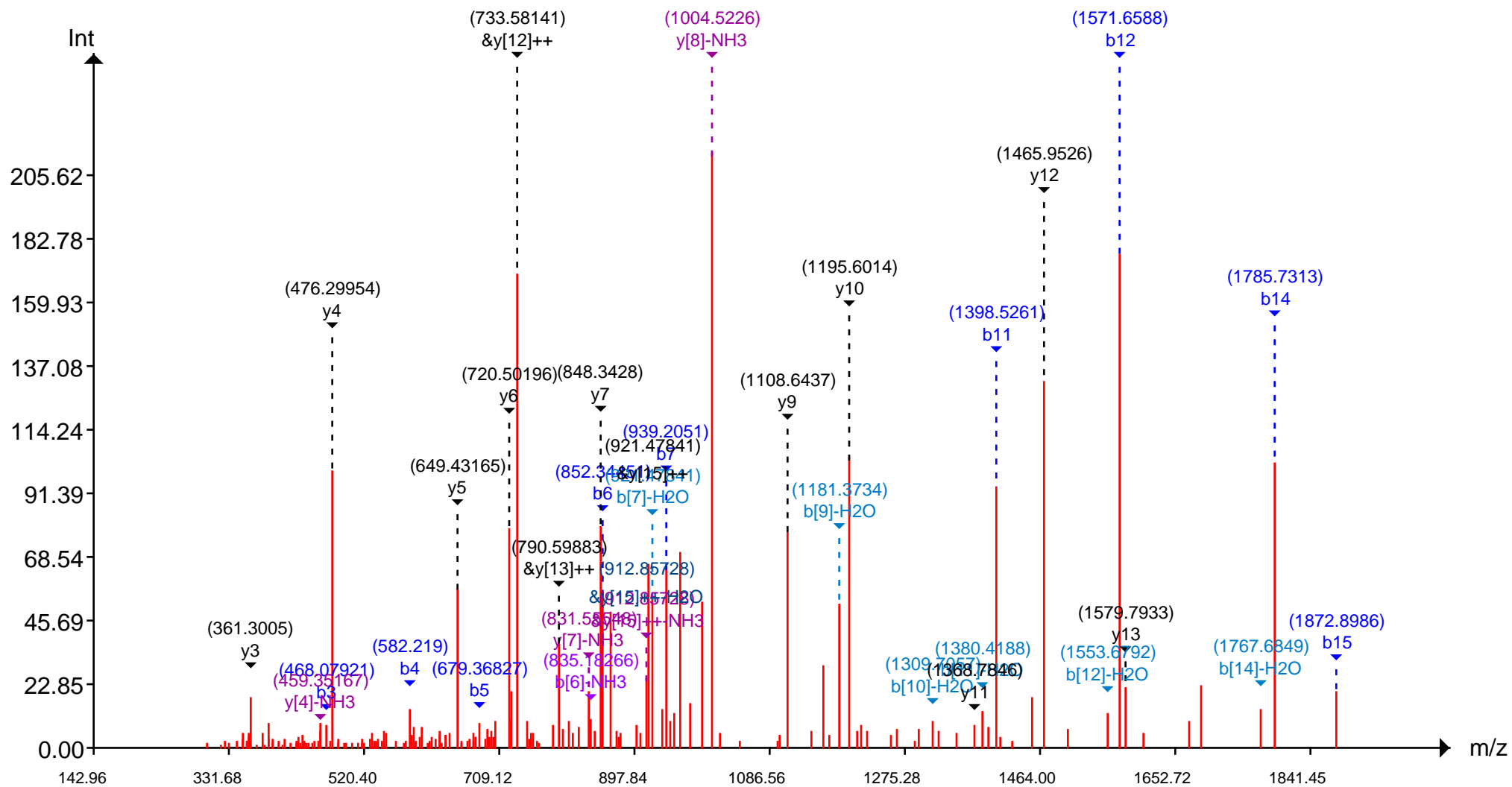
Info

Modified Sequence	AcD3-LGTGNDLSPTTVM<Mox*>SEGTR<C13>-COOH	Ion score	53.51
Protein	Q923D2	Confidence level	99.00%
Precursor m/z	951.96786	Identity Threshold	47.26
Charge	2+	Mass error	0.003611



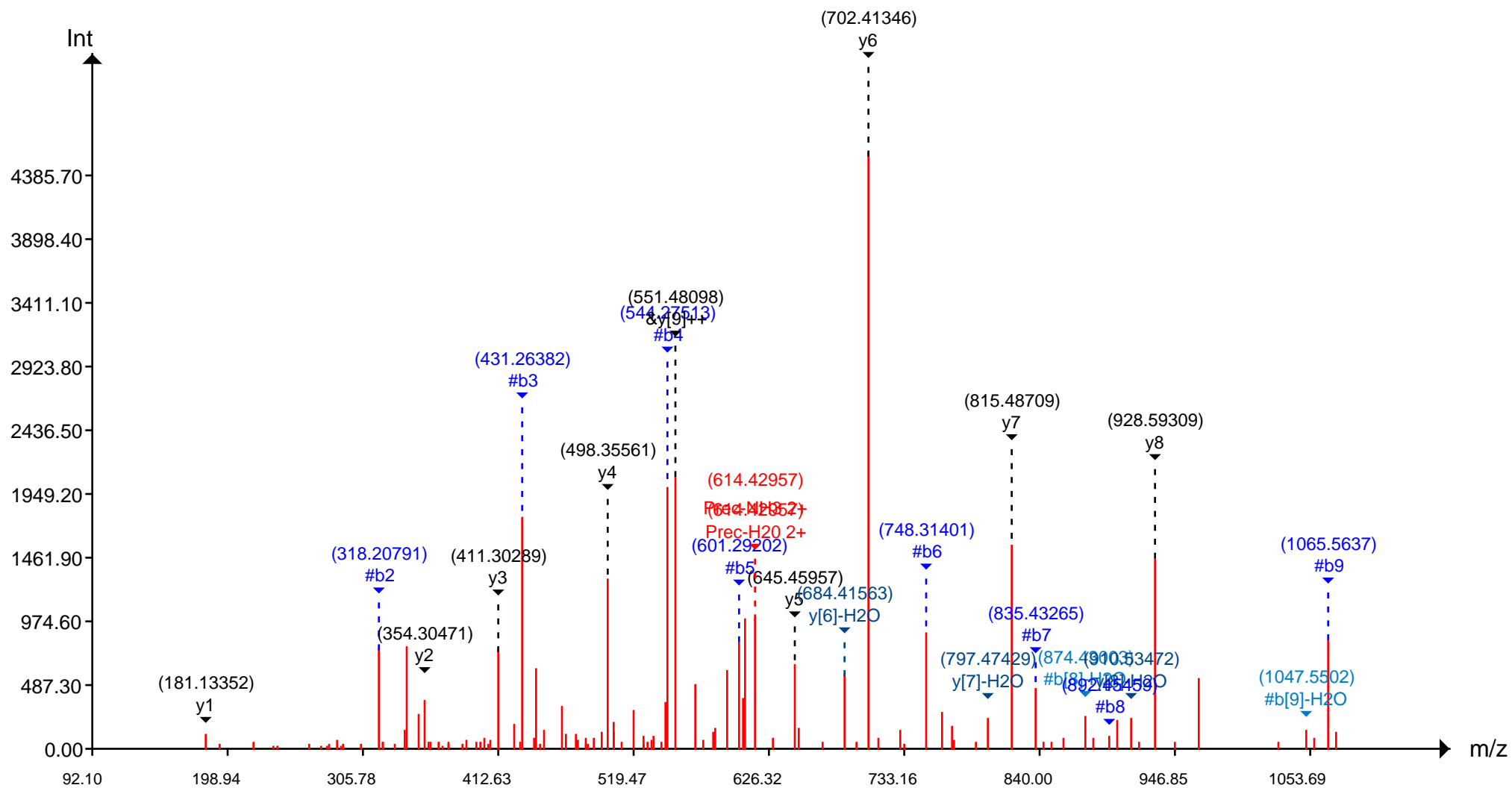
Info

Modified Sequence	AcD3-LGTGNDLSPTTVM<Mox*>SEGTR-COOH	Ion score	58.95
Protein	Q923D2	Confidence level	99.00%
Precursor m/z	948.95817	Identity Threshold	46.88
Charge	2+	Mass error	0.004373



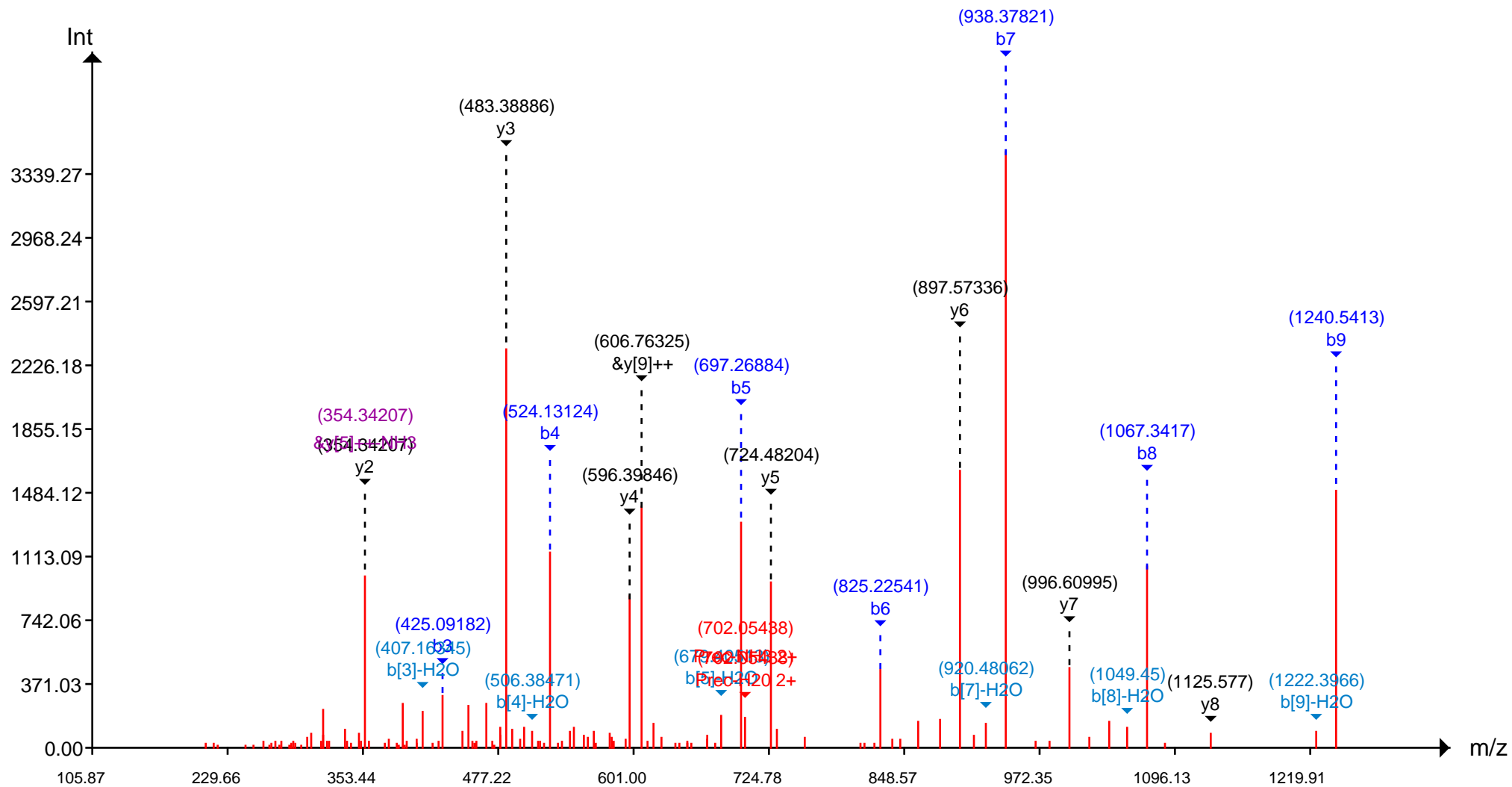
Info

Modified Sequence	AcD3-C<Cmm*>DFNPK<AcD3K*>SSK<AcD3K*>QAK<AcD3K*>DVSR-COOH	Ion score	67.67
Protein	O08585	Confidence level	99.00%
Precursor m/z	1024.0205	Identity Threshold	47.94
Charge	2+	Mass error	0.003468



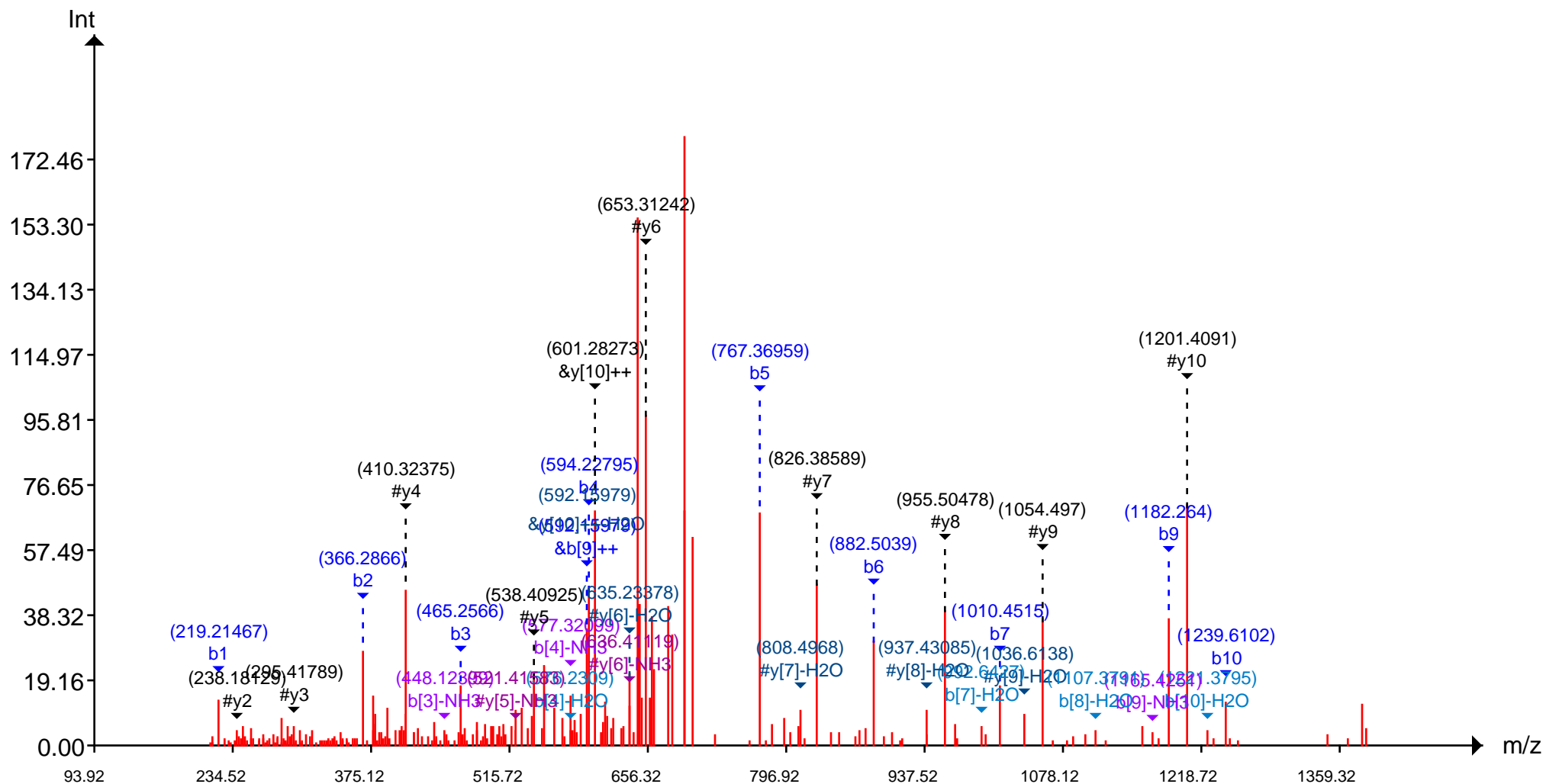
Info

Modified Sequence	AcD3-VK<AcD3K*>LLGM<Mox*>SGK<AcD3K*>R<C13>-COOH	Ion score	53.4
Protein	Q61937	Confidence level	99.00%
Precursor m/z	623.38587	Identity Threshold	43.11
Charge	2+	Mass error	3.67E-4



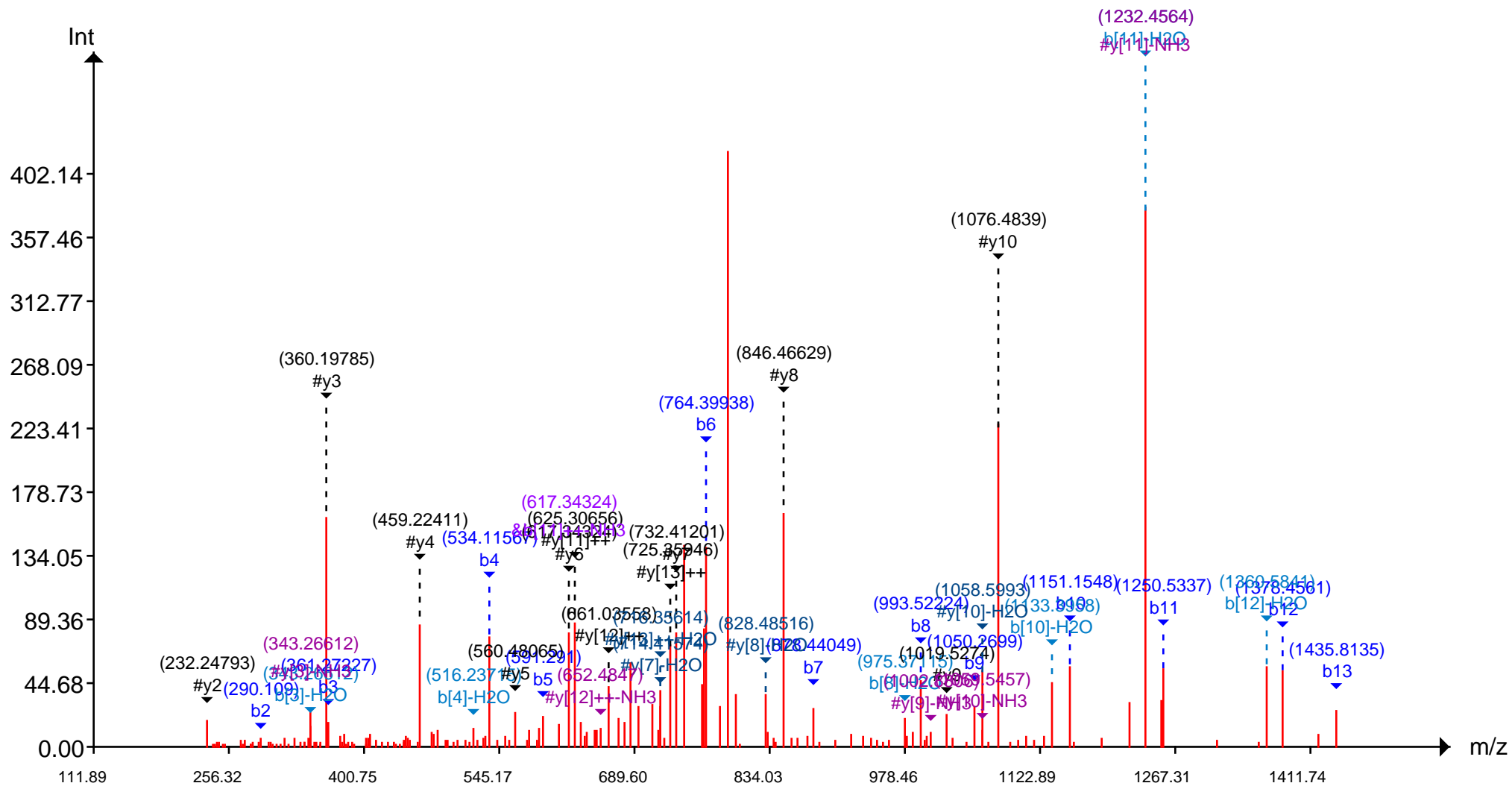
Info

Modified Sequence	AcD3-YSEVK<AcD3K*>QIEK<AcD3K*>R<C13>-COOH	Ion score	55.83
Protein	Q6NZB0	Confidence level	99.00%
Precursor m/z	710.90819	Identity Threshold	46.33
Charge	2+	Mass error	3.3E-4



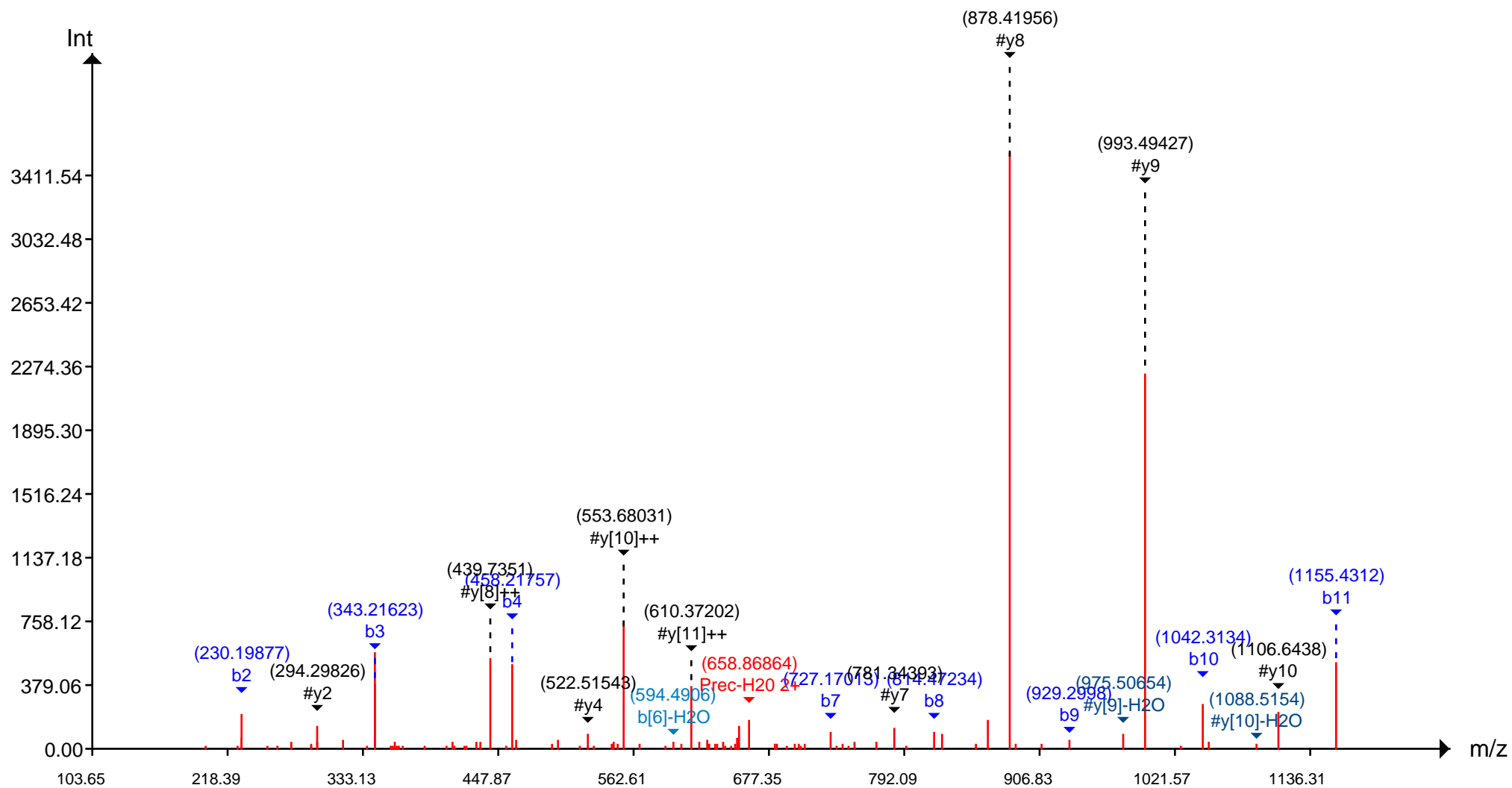
Info

Modified Sequence	AcD3-K<AcD3K*>M<Mox*>VEK<AcD3K*>DQDGGR<C13>-COOH	Ion score	71.94
Protein	Q9CZX8	Confidence level	99.00%
Precursor m/z	710.36325	Identity Threshold	47.91
Charge	2+	Mass error	2.17E-4



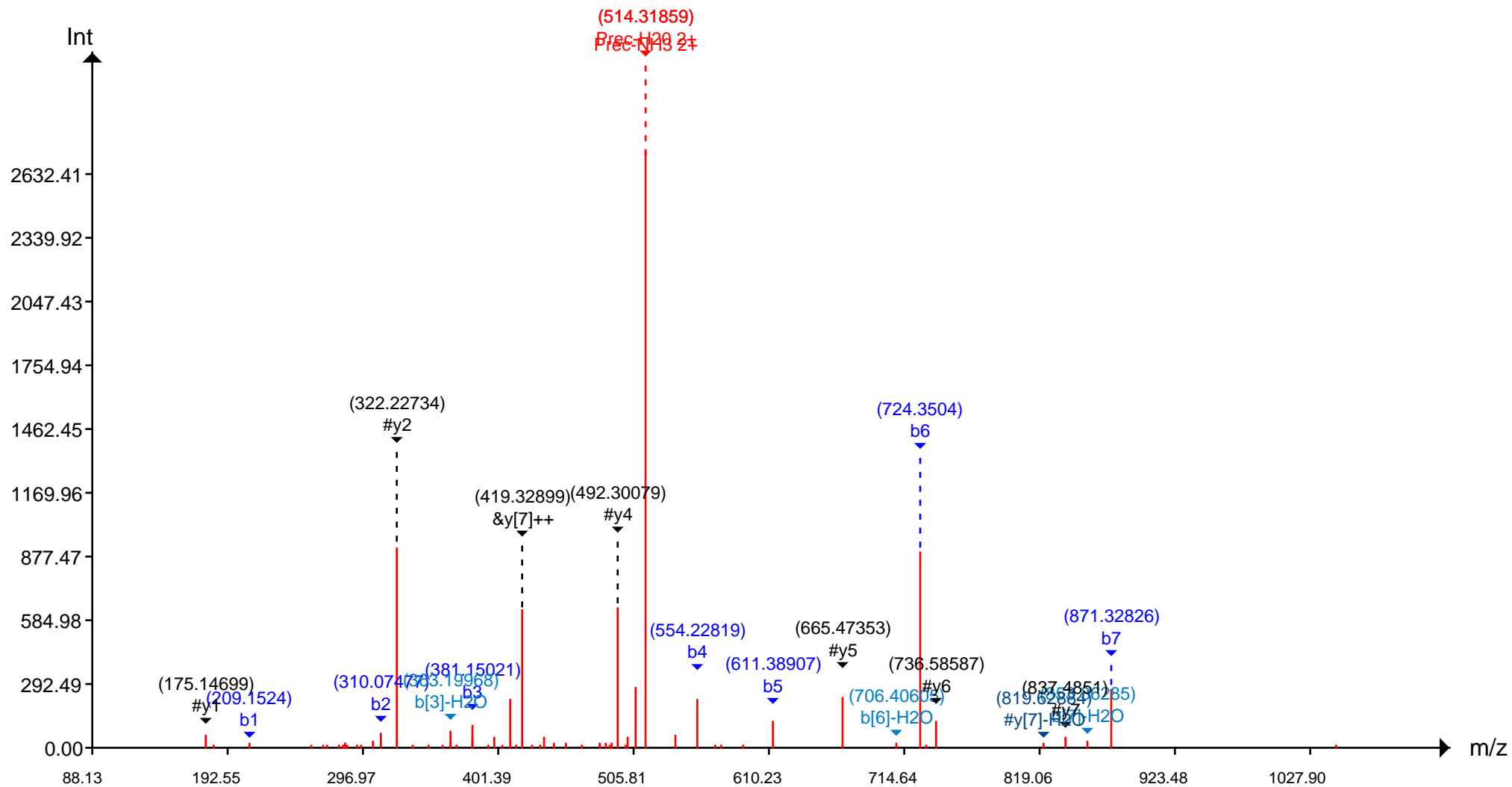
Info

Modified Sequence	AcD3-DEAK<AcD3K*>GK<AcD3K*>NDGTVQGR-COOH	Ion score	93.81
Protein	O08788	Confidence level	99.00%
Precursor m/z	805.4108	Identity Threshold	47.98
Charge	2+	Mass error	0.001674



Info

Modified Sequence	AcD3-AIIDPGDSDIIR<C13>-COOH	Ion score	64.97
Protein	P62889	Confidence level	99.00%
Precursor m/z	668.36849	Identity Threshold	47.86
Charge	2+	Mass error	7.79E-4



Info

Modified Sequence AcD3-YTAK<AcD3K*>GLFR-COOH

Ion score 58.93

Protein P17182, P17183

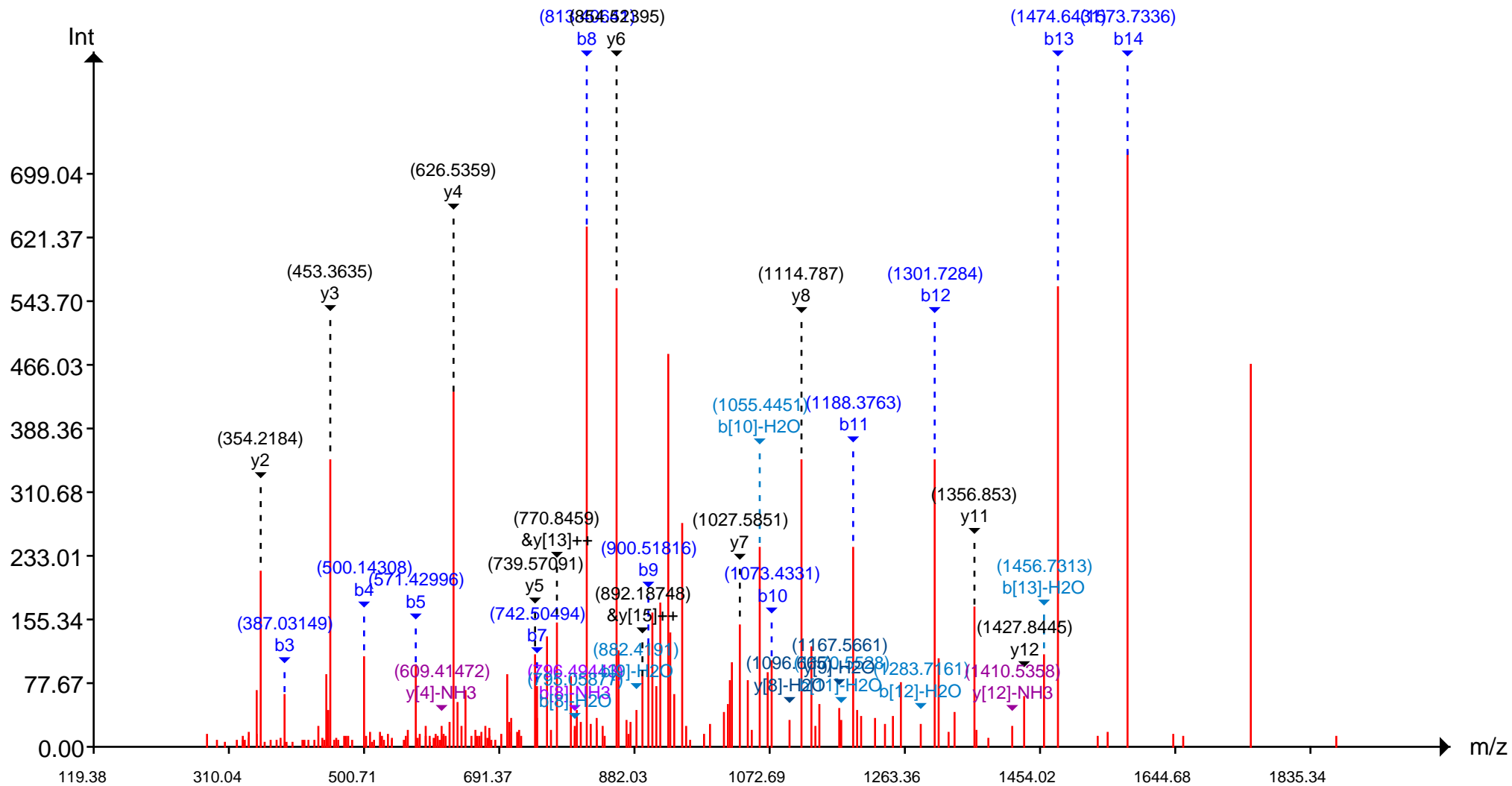
Confidence level 99.00%

Precursor m/z 523.30111

Identity Threshold 47.13

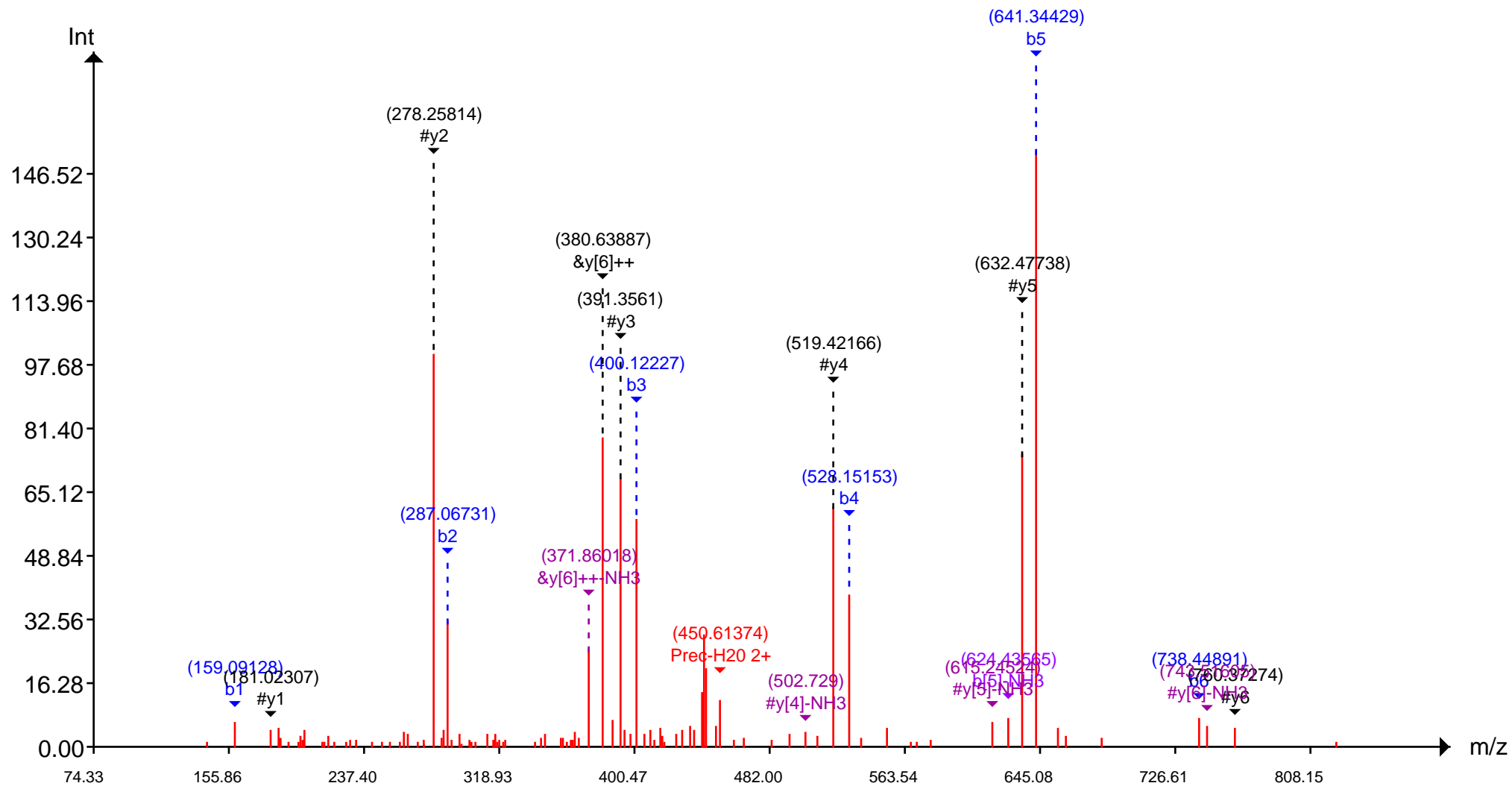
Charge 2+

Mass error 1.89E-4



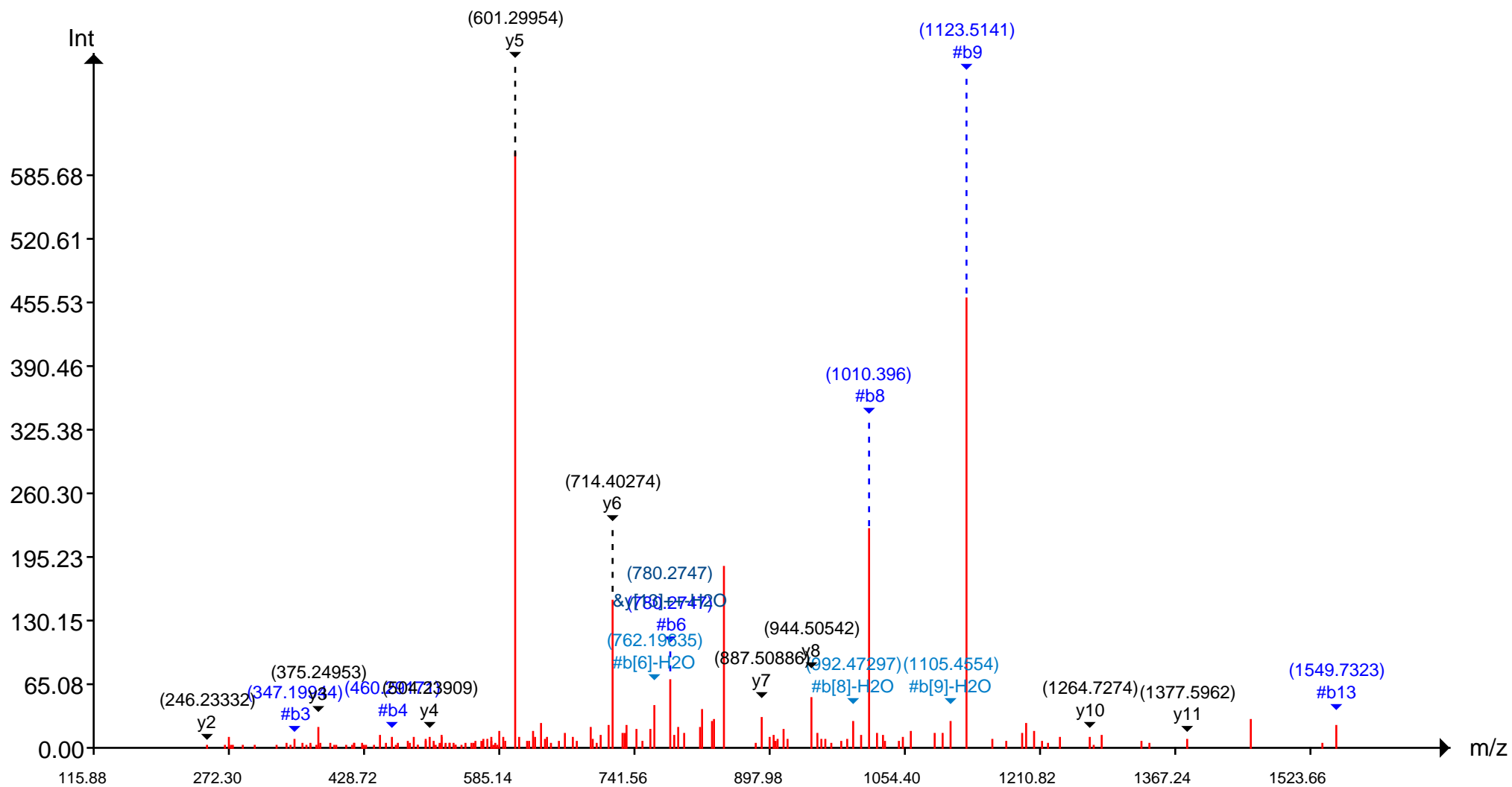
Info

Modified Sequence	AcD3-VLELAGNASK<AcD3K*>DLK<AcD3K*>VK<AcD3K*>R<C13>-COOH	Ion score	49.82
Protein	Q3THW5, P0C0S6	Confidence level	99.00%
Precursor m/z	964.08977	Identity Threshold	39.41
Charge	2+	Mass error	0.001521



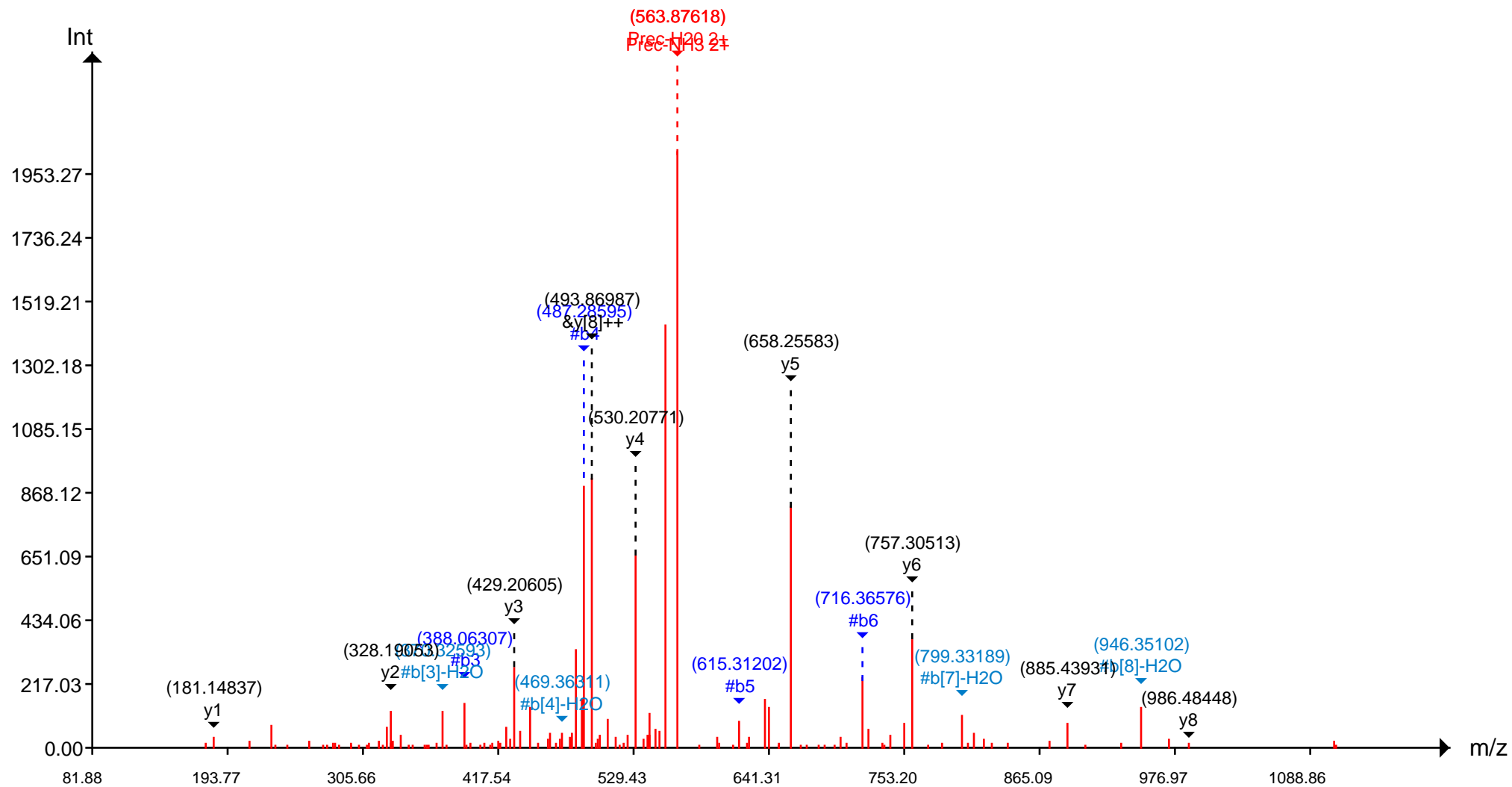
Info

Modified Sequence	AcD3-LQLQIPR<C13>-COOH	Ion score	48.97
Protein	P97371	Confidence level	99.00%
Precursor m/z	459.79873	Identity Threshold	41.63
Charge	2+	Mass error	-3.9E-4



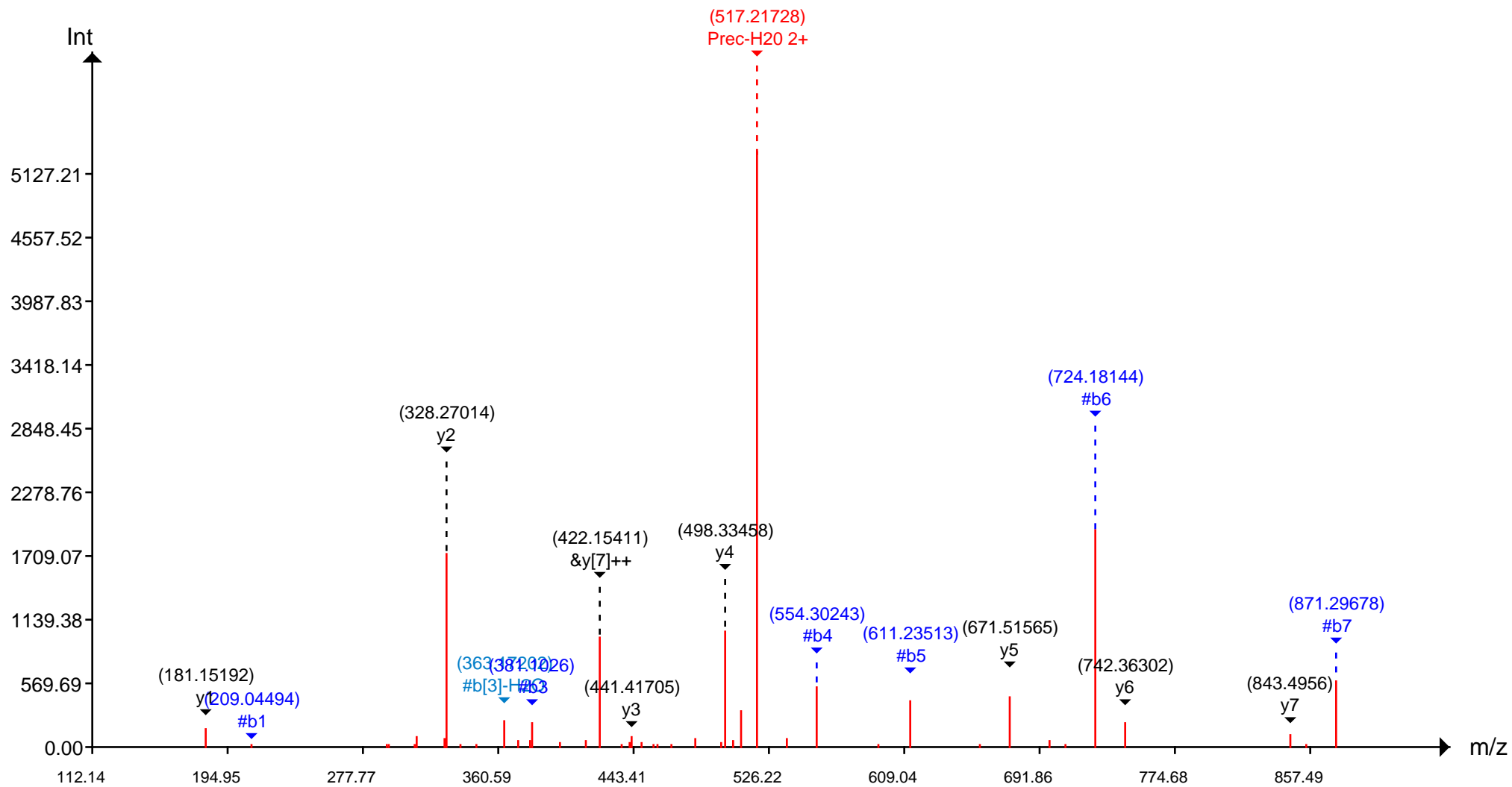
Info

Modified Sequence	AcD3-TEALM<Mox*>K<AcD3K*>GK<AcD3K*>LPPEAR-COOH	Ion score	51.45
Protein	P06745	Confidence level	99.00%
Precursor m/z	862.46699	Identity Threshold	47.80
Charge	2+	Mass error	0.002237



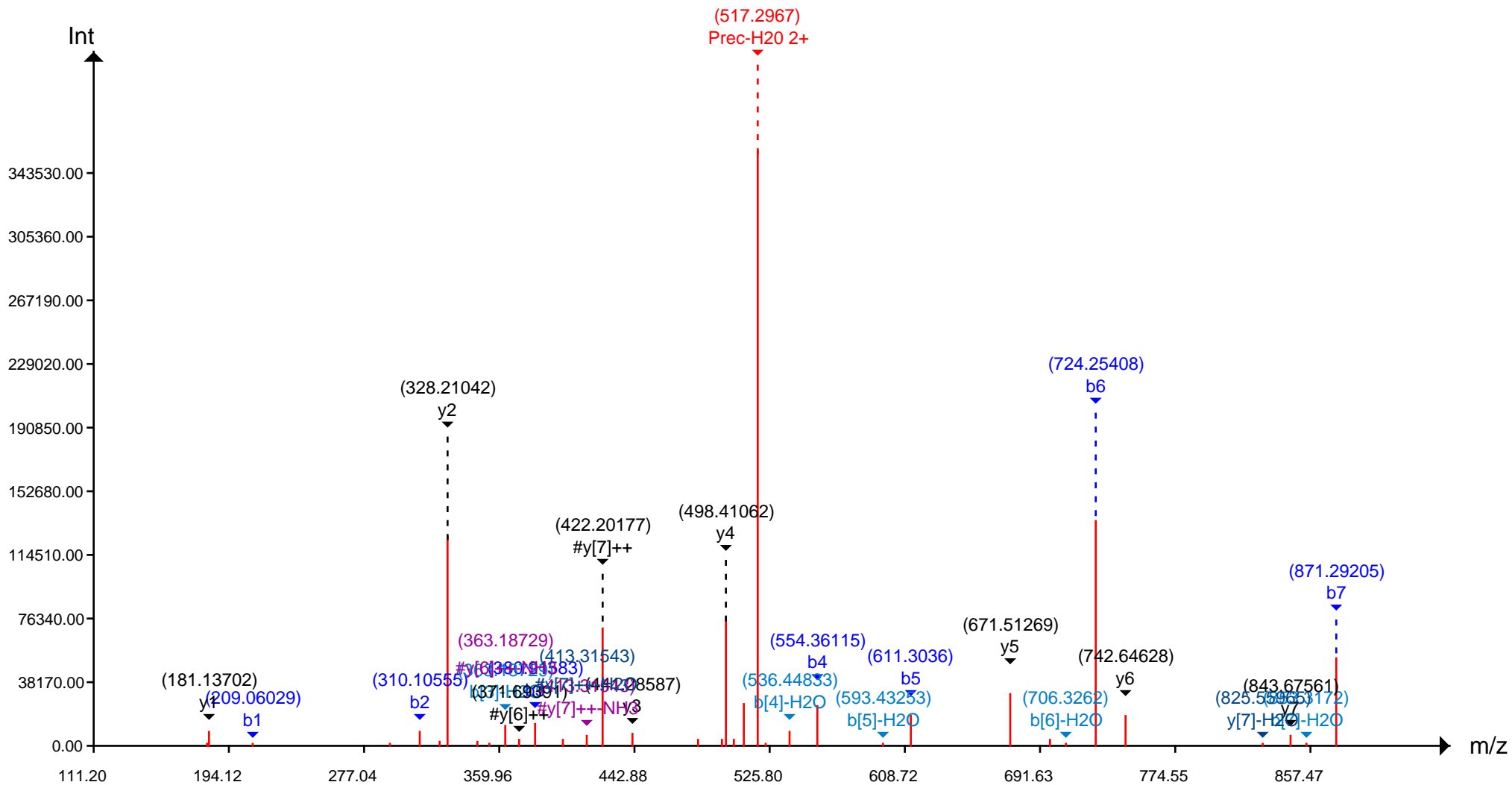
Info

Modified Sequence	AcD3-LTQVQTTM<Mox*>R<C13>-COOH	Ion score	54.74
Protein	Q99KJ8	Confidence level	99.00%
Precursor m/z	572.81168	Identity Threshold	47.87
Charge	2+	Mass error	-4.75E-4



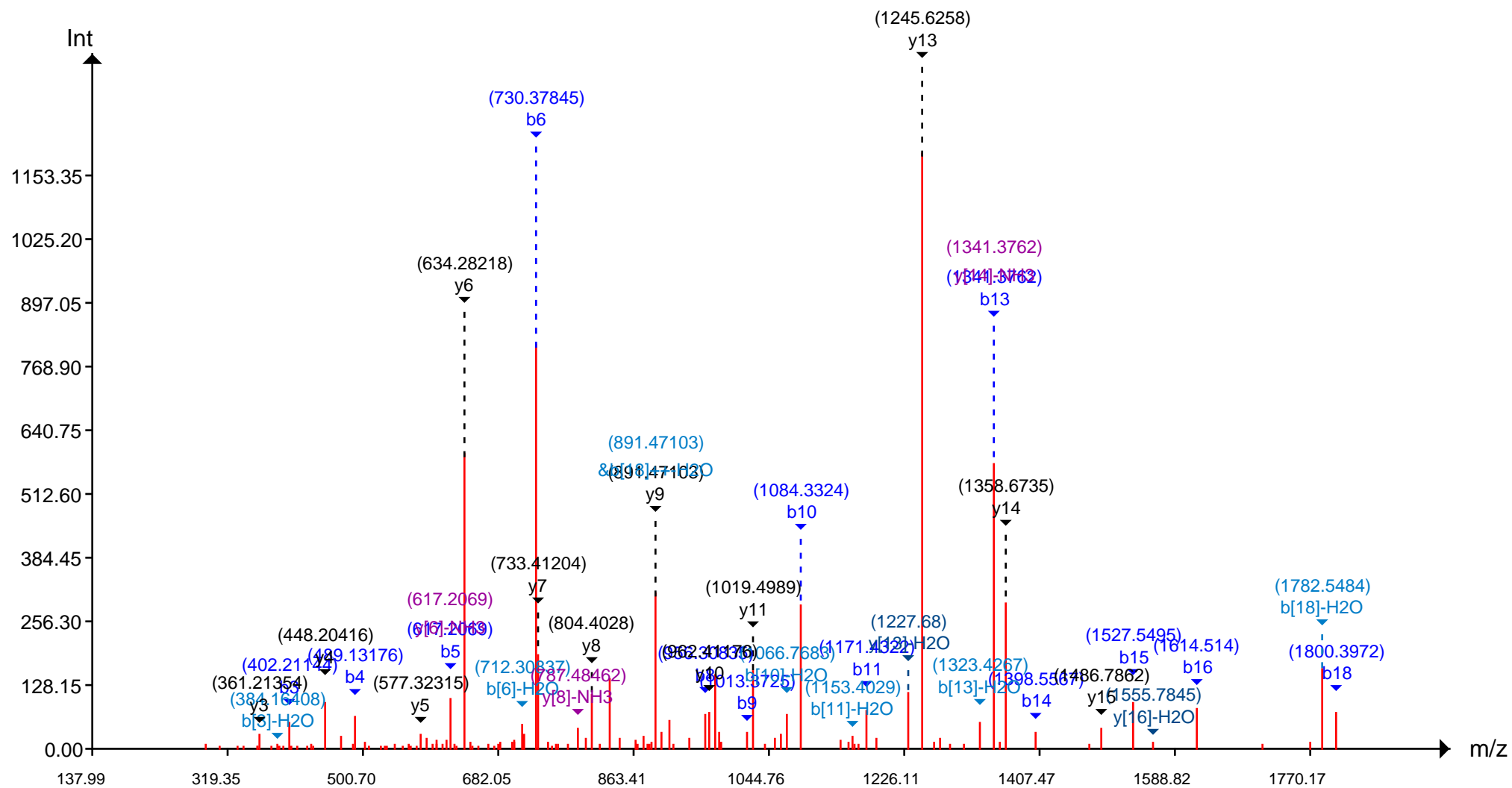
Info

Modified Sequence	AcD3-YTAK<AcD3K*>GLFR<C13>-COOH	Ion score	70.09
Protein	P17182, P17183	Confidence level	99.00%
Precursor m/z	526.31094	Identity Threshold	46.03
Charge	2+	Mass error	-2.92E-4



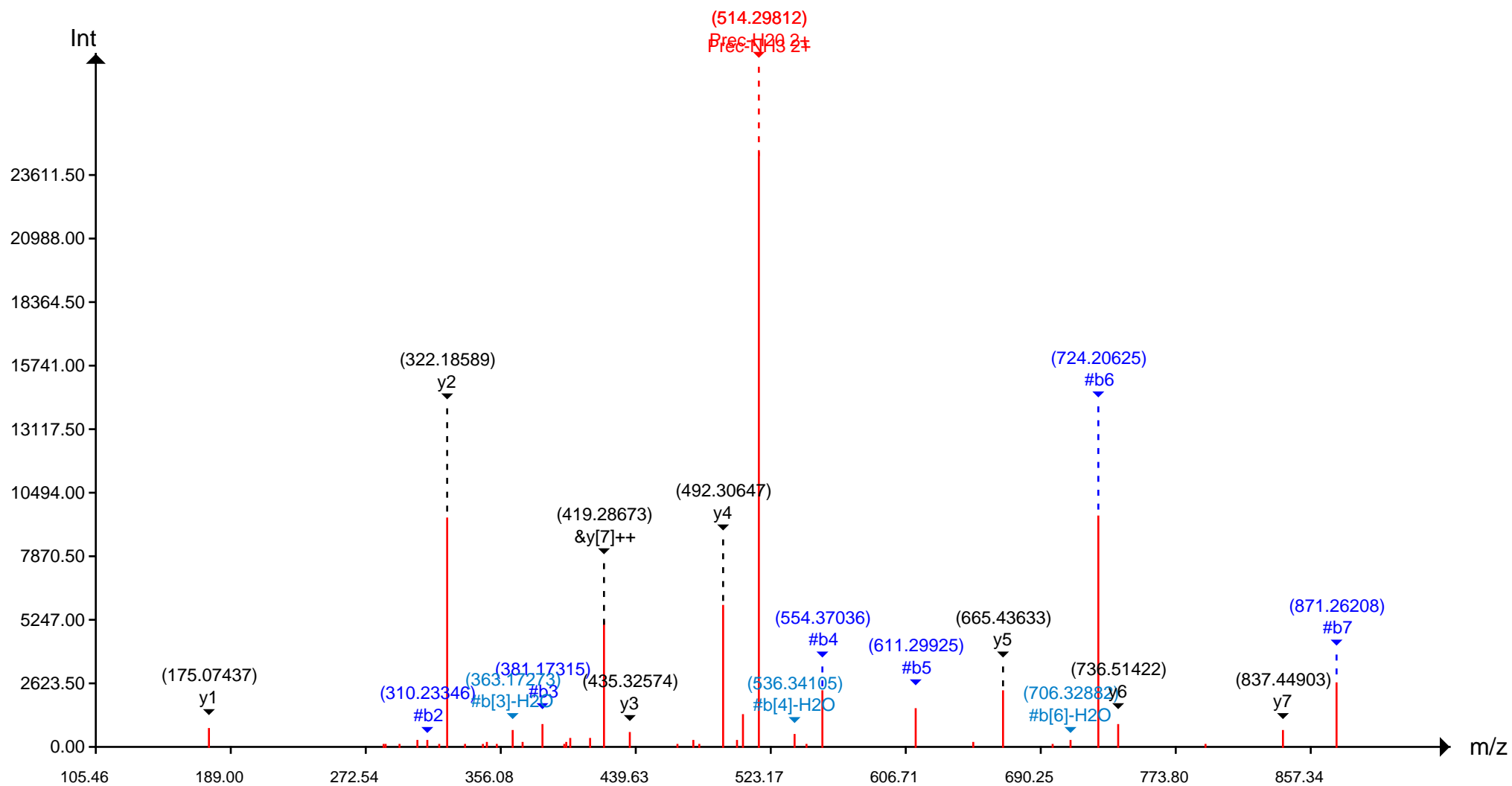
Info

Modified Sequence	AcD3-YTAK<AcD3K*>GLFR<C13>-COOH	Ion score	69.11
Protein	P17182, P17183	Confidence level	99.00%
Precursor m/z	526.3103	Identity Threshold	45.99
Charge	2+	Mass error	-0.001572



Info

Modified Sequence	AcD3-DQLSQLPEGASAVGESSVR-COOH	Ion score	122.16
Protein	Q921W0	Confidence level	99.00%
Precursor m/z	987.99529	Identity Threshold	47.72
Charge	2+	Mass error	0.0027



Info

Modified Sequence AcD3-YTAK<AcD3K*>GLFR-COOH

Ion score 64.61

Protein P17182, P17183

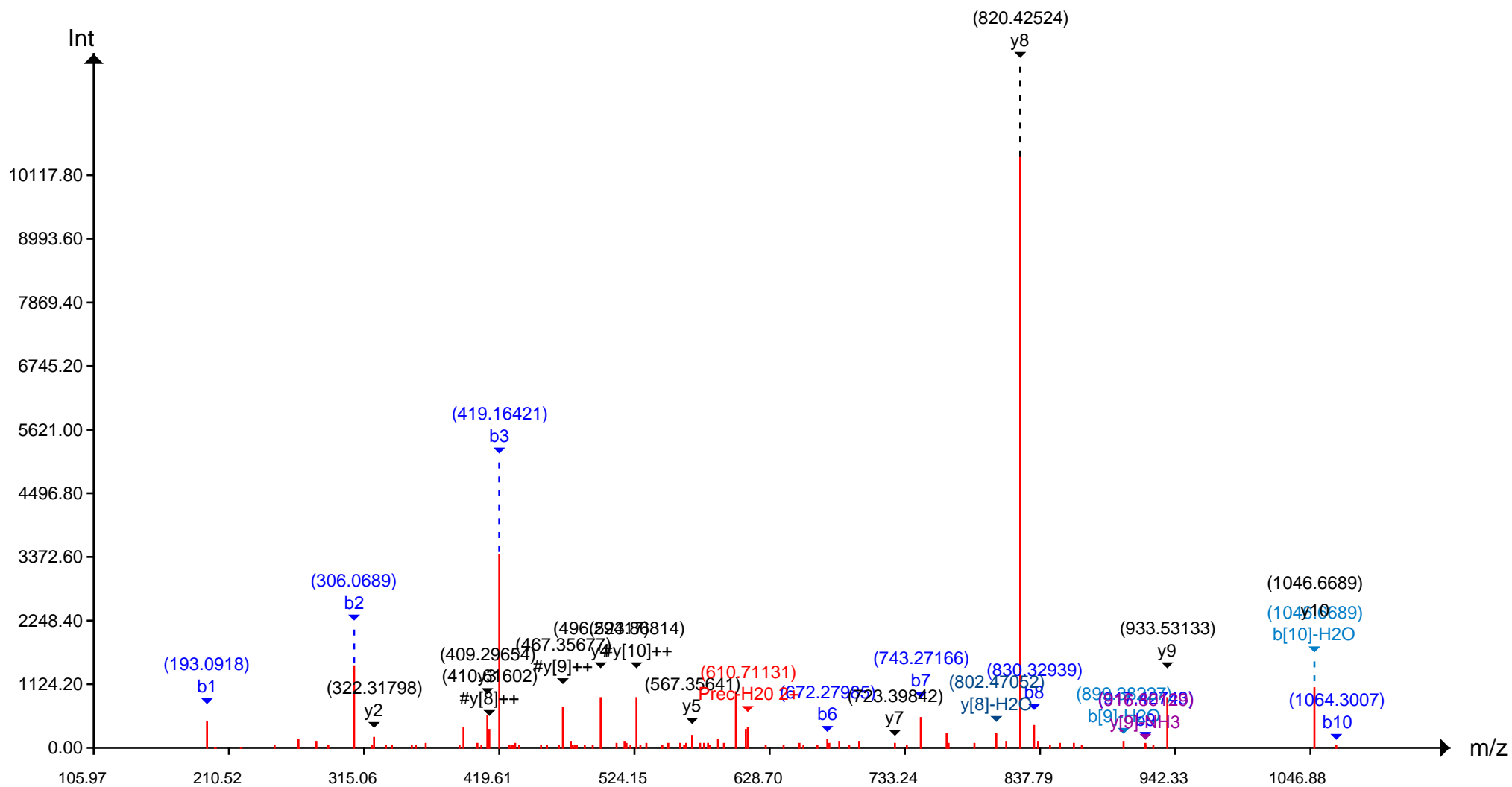
Confidence level 99.00%

Precursor m/z 523.30141

Identity Threshold 47.15

Charge 2+

Mass error 7.89E-4



Info

Modified Sequence AcD3-M<Mox*>ILPVGASSFR-COOH

Ion score 65.87

Protein P17182

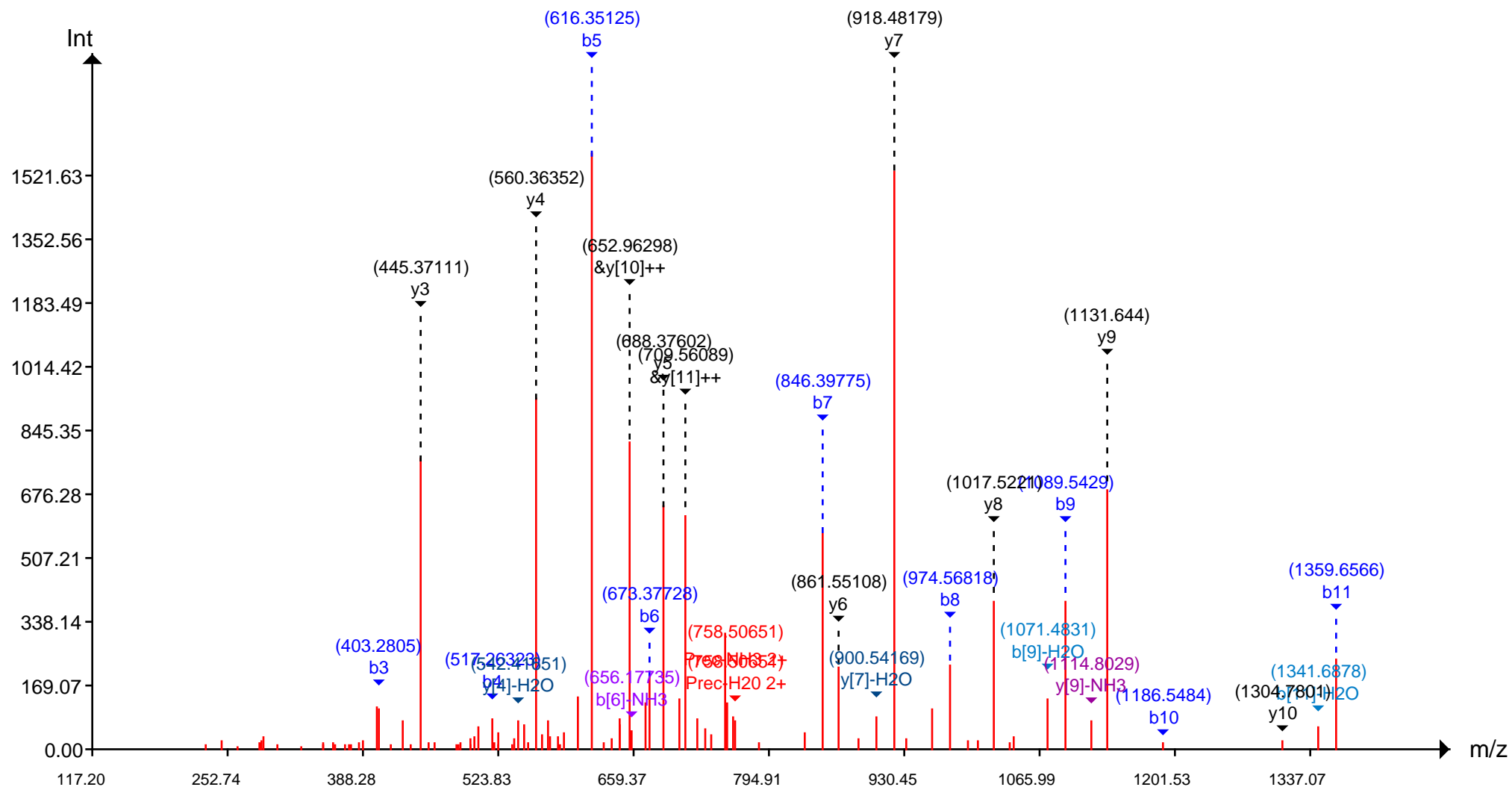
Confidence level 99.00%

Precursor m/z 619.836

Identity Threshold 47.94

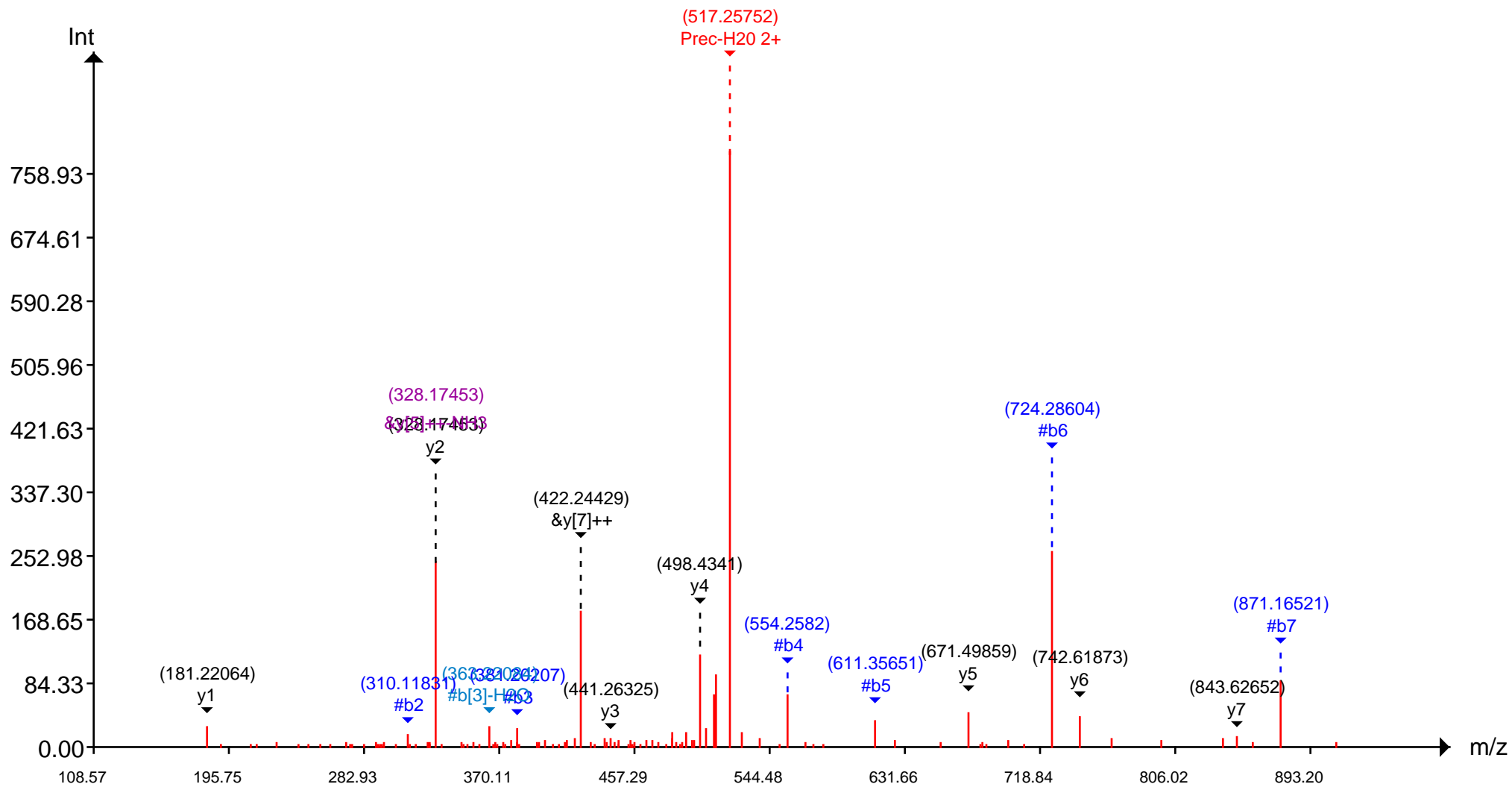
Charge 2+

Mass error 6.33E-4



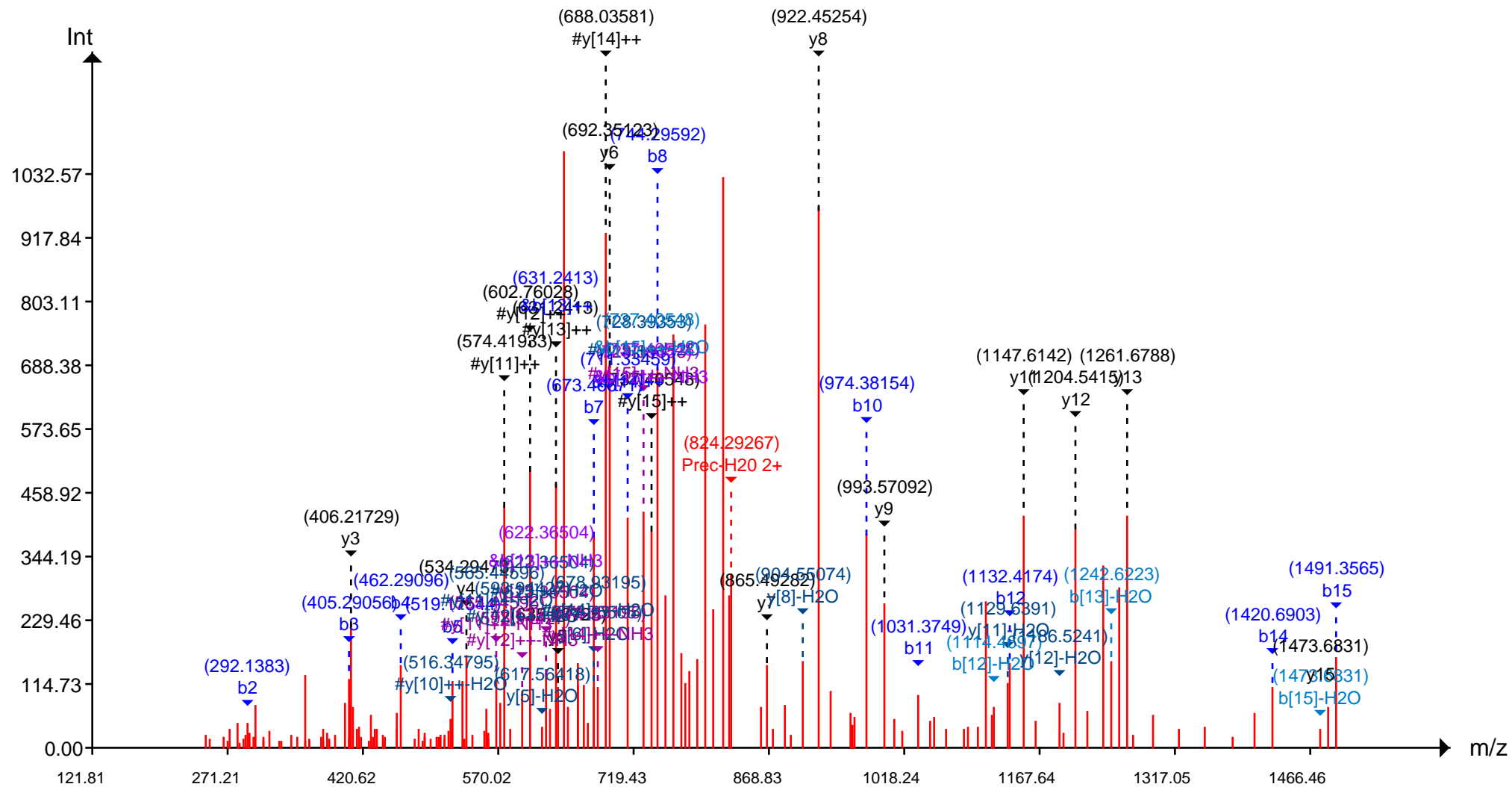
Info

Modified Sequence	AcD3-AIK<AcD3K*>NVGK<AcD3K*>QDPK<AcD3K*>R-COOH	Ion score	72.42
Protein	O35593	Confidence level	99.00%
Precursor m/z	767.4599	Identity Threshold	43.74
Charge	2+	Mass error	-0.001155



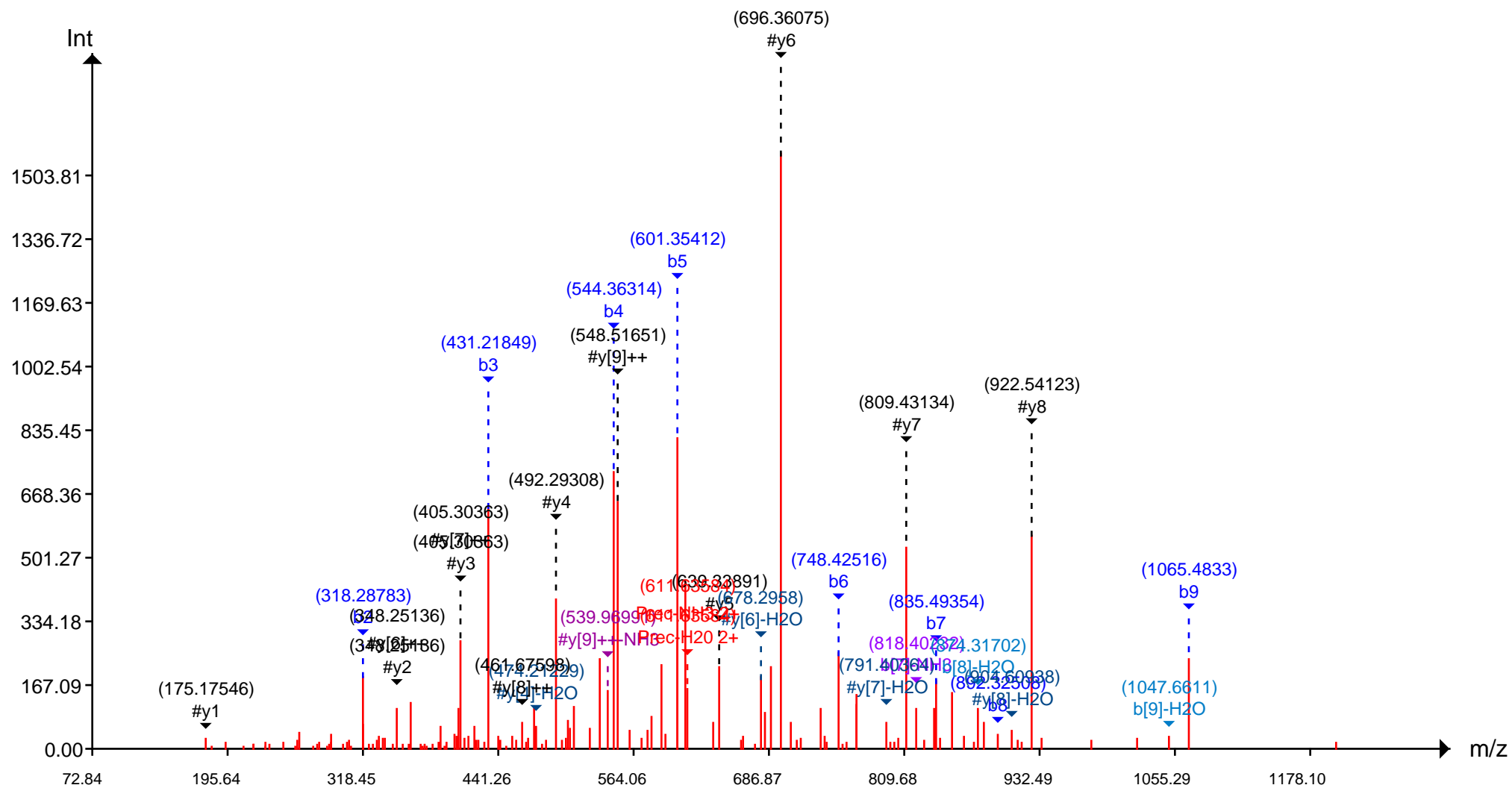
Info

Modified Sequence	AcD3-YTAK<AcD3K*>GLFR<C13>-COOH	Ion score	52.64
Protein	P17182, P17183	Confidence level	99.00%
Precursor m/z	526.3112	Identity Threshold	45.93
Charge	2+	Mass error	2.28E-4



Info

Modified Sequence	AcD3-FVLGGPGAGK<AcD3K*>GTQC<Cmm*>AR-COOH	Ion score	89.89
Protein	Q9DBP5	Confidence level	99.00%
Precursor m/z	833.43709	Identity Threshold	48.17
Charge	2+	Mass error	0.002069



Info

Modified Sequence AcD3-VK<AcD3K*>LLGM<Mox*>SGK<AcD3K*>R-COOH

Ion score 55.65

Protein Q61937

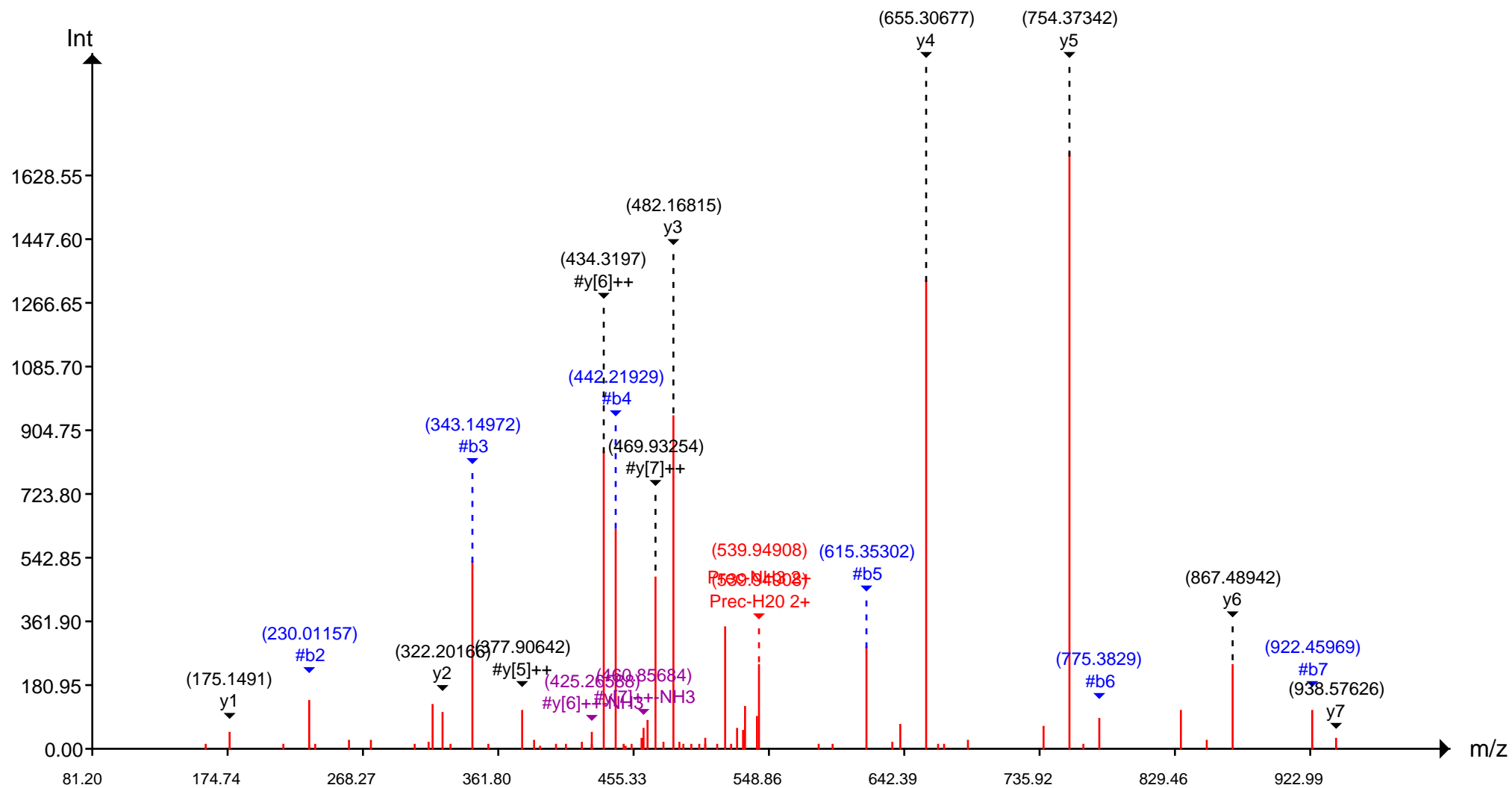
Confidence level 99.00%

Precursor m/z 620.37588

Identity Threshold 43.96

Charge 2+

Mass error 5.29E-4



Info

Modified Sequence AcD3-LAIVK<AcD3K*>C<Cmm*>M<Mox*>R-COOH

Ion score 55.5

Protein P97429

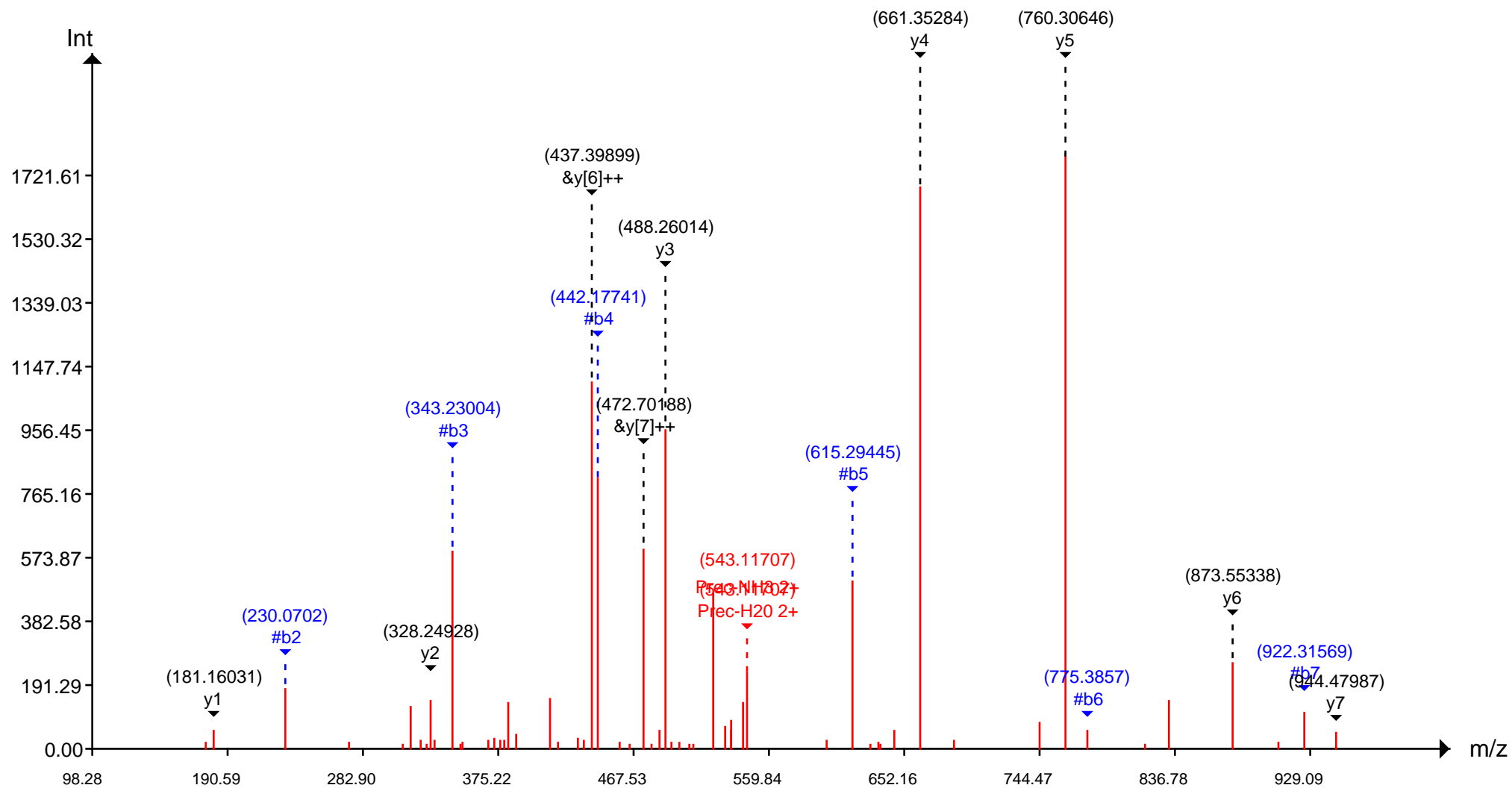
Confidence level 99.00%

Precursor m/z 548.80993

Identity Threshold 47.37

Charge 2+

Mass error 1.9E-4



Info

Modified Sequence AcD3-LAIVK<AcD3K*>C<Cmm*>M<Mox*>R<C13>-COOH

Ion score 62.82

Protein P97429

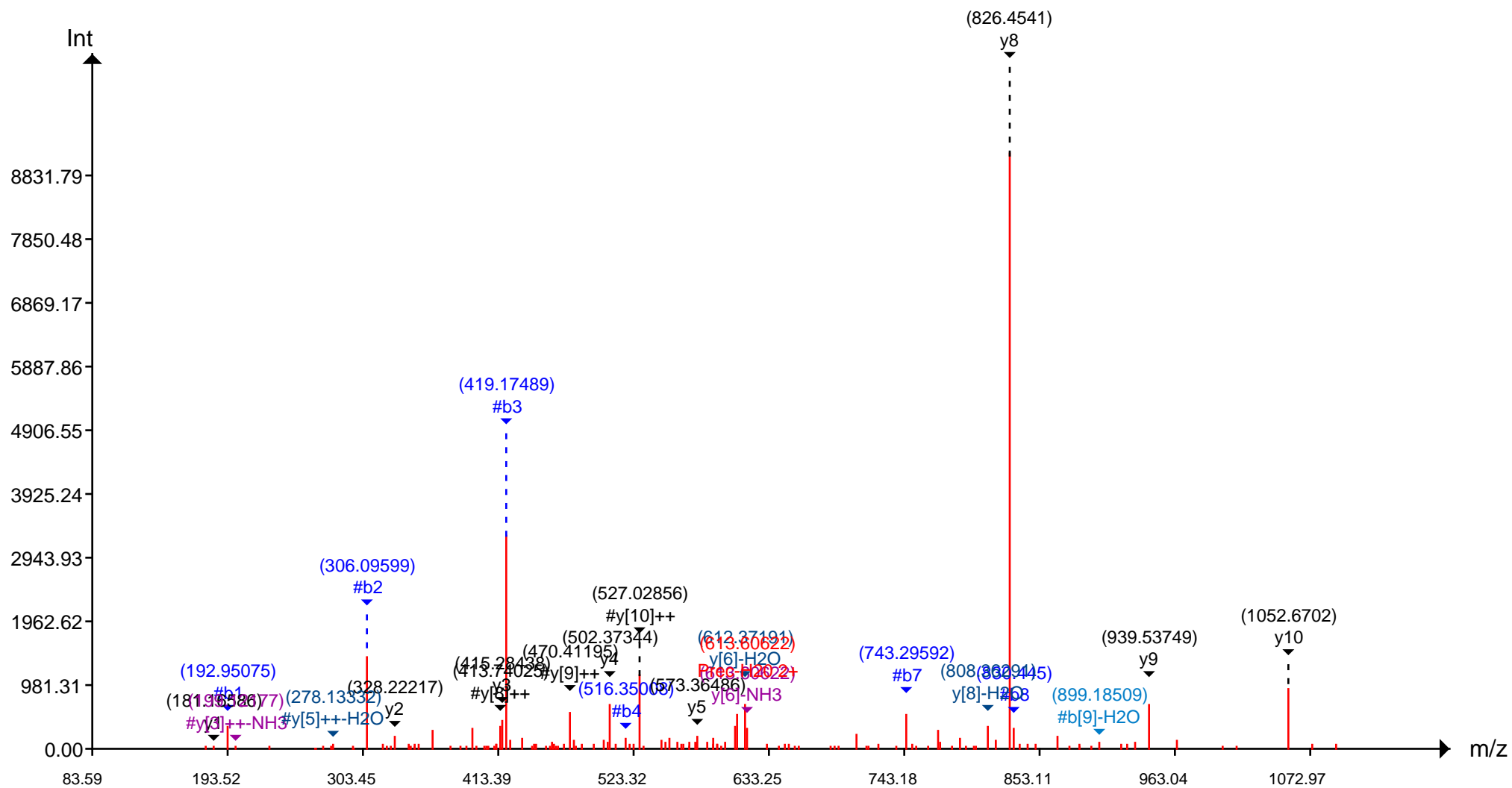
Confidence level 99.00%

Precursor m/z 551.82

Identity Threshold 46.72

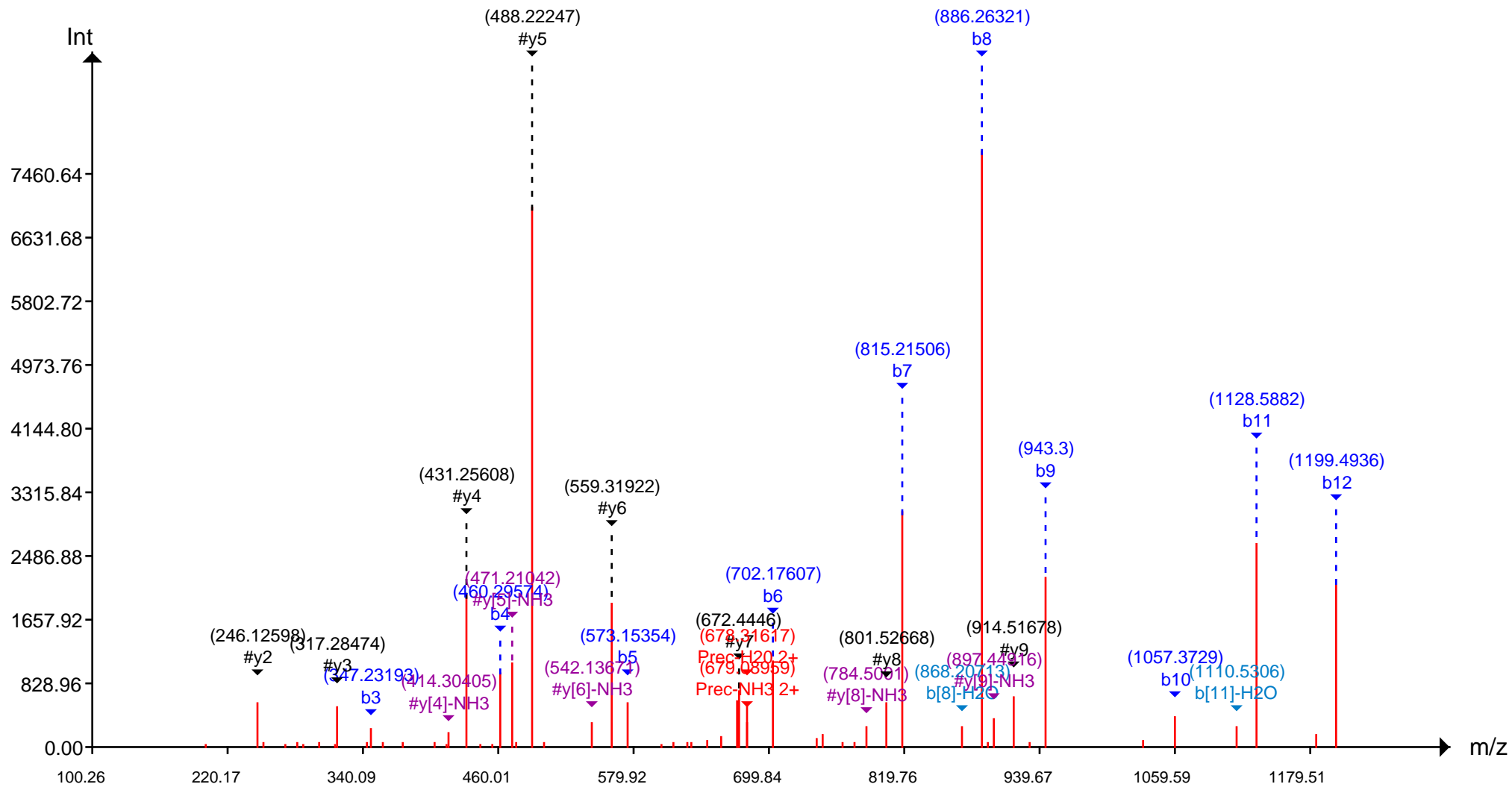
Charge 2+

Mass error 1.89E-4



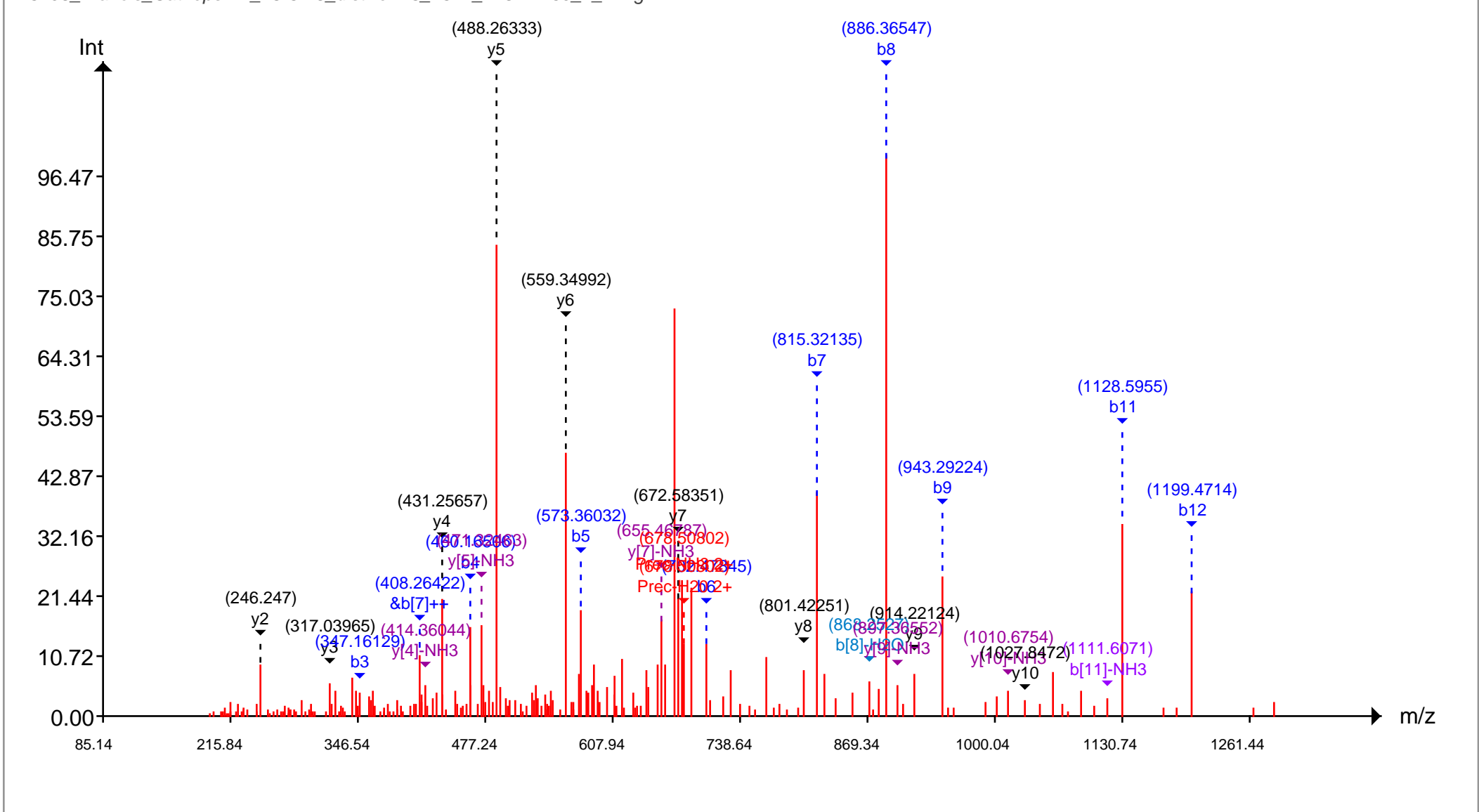
Info

Modified Sequence	AcD3-M<Mox*>ILPVGASSFR<C13>-COOH	Ion score	57.26
Protein	P17182	Confidence level	99.00%
Precursor m/z	622.84576	Identity Threshold	47.59
Charge	2+	Mass error	1.2E-5



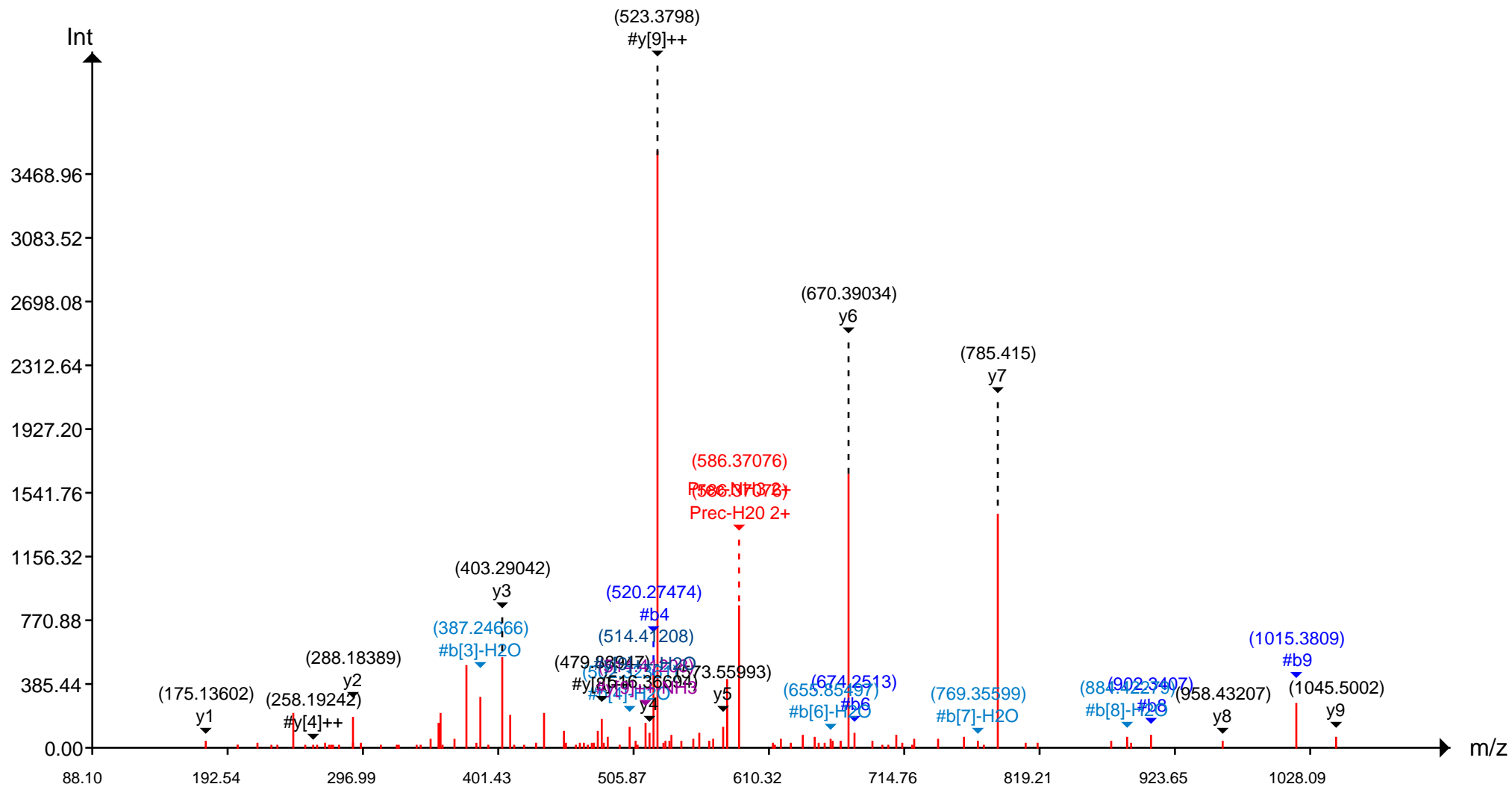
Info

Modified ...	AcD3-TAEILELAGNAAR-COOH	Ion score	91
	Q8CGP5, Q8CGP6, Q8CGP7, P22752, Q6GSS7, Q64522, Q64523, Q8BFU2, Q8R1M2, P27661, Q9QZQ8	Confidence level	99
Prec...	687.37849	Identity Threshold	47
	2+	Mass error	0.



Info

Modified ...	AcD3-TAEILELAGNAAR-COOH	Ion score	59
	Q8CGP5, Q8CGP6, Q8CGP7, P22752, Q6GSS7, Q64522, Q64523, Q8BFU2, Q8R1M2, P27661, Q9QZQ8	Confidence level	99
Prec...	687.37766	Identity Threshold	47
	2+	Mass error	0.



Info

Modified Sequence AcD3-VSK<AcD3K*>DPGLDLR-COOH

Ion score 48.9

Protein Q91V76

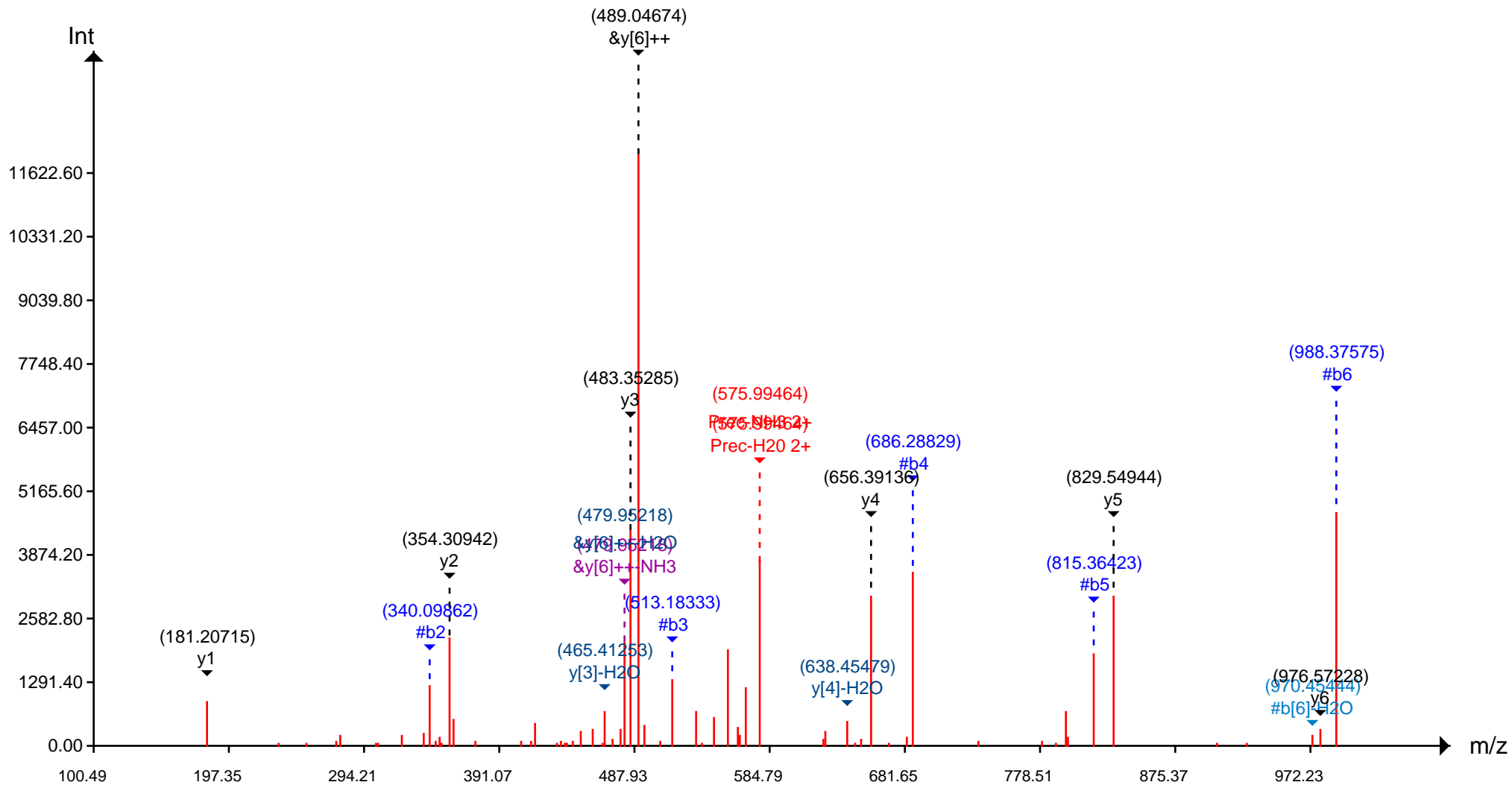
Confidence level 99.00%

Precursor m/z 595.33852

Identity Threshold 47.44

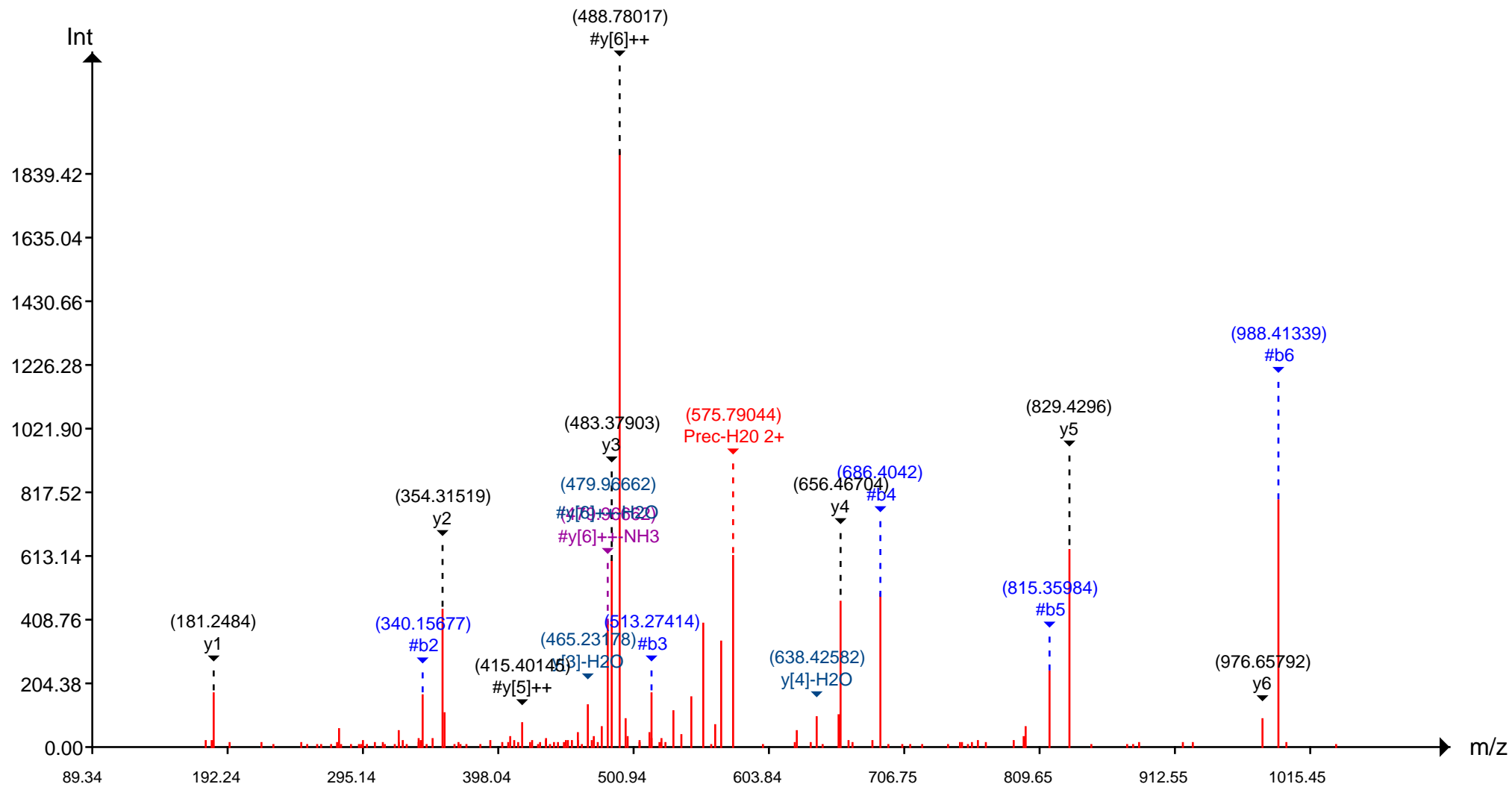
Charge 2+

Mass error 3.79E-4



Info

Modified Sequence	AcD3-FM<Mox*>K<AcD3K*>K<AcD3K*>EK<AcD3K*>R<C13>-COOH	Ion score	51.79
Protein	P31725	Confidence level	99.00%
Precursor m/z	584.84835	Identity Threshold	45.51
Charge	2+	Mass error	0.001469



Info

Modified Sequence AcD3-FM<Mox*>K<AcD3K*>K<AcD3K*>EK<AcD3K*>R<C13>-COOH

Ion score 46.53

Protein P31725

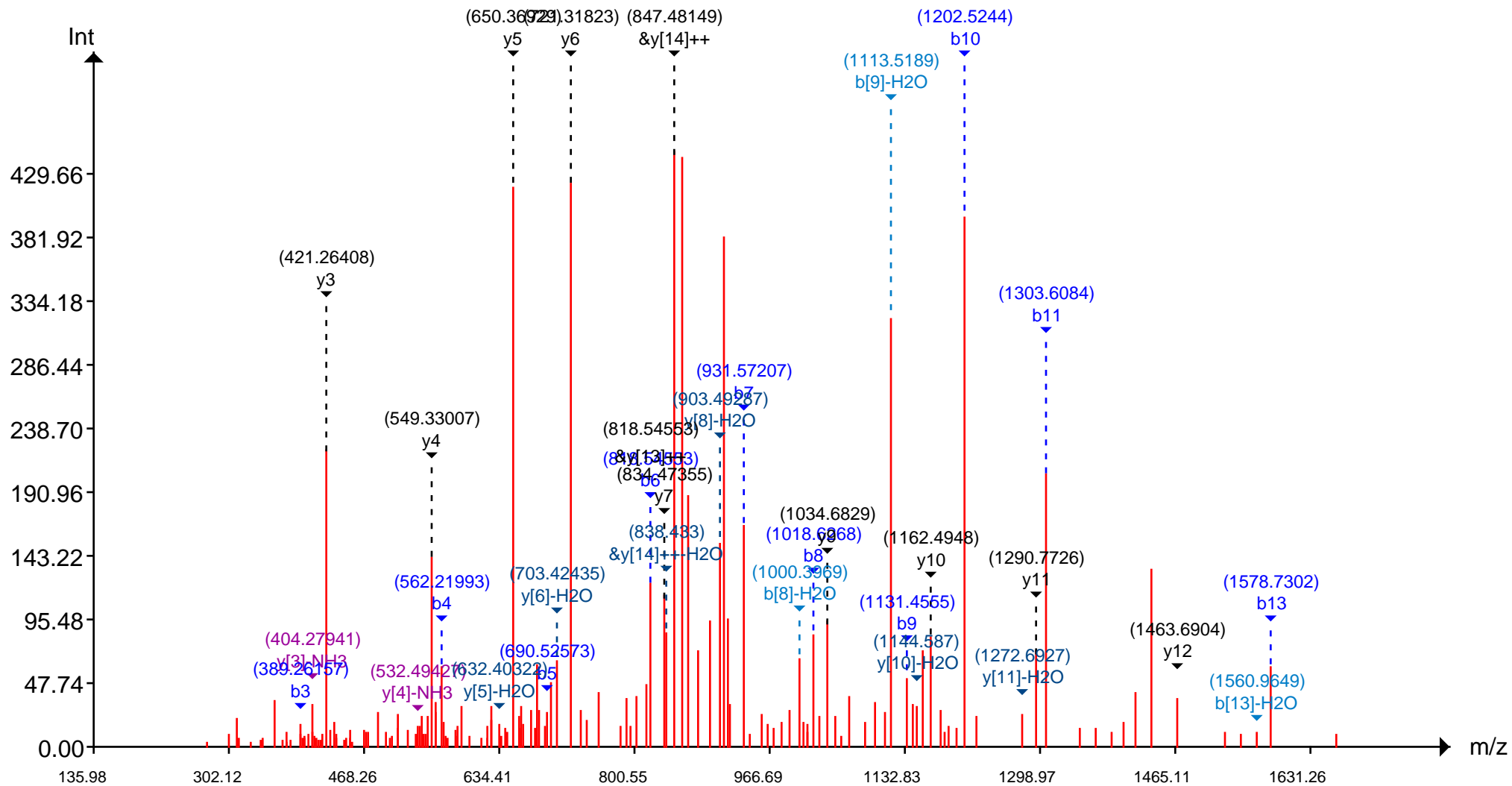
Confidence level 99.00%

Precursor m/z 584.84733

Identity Threshold 45.68

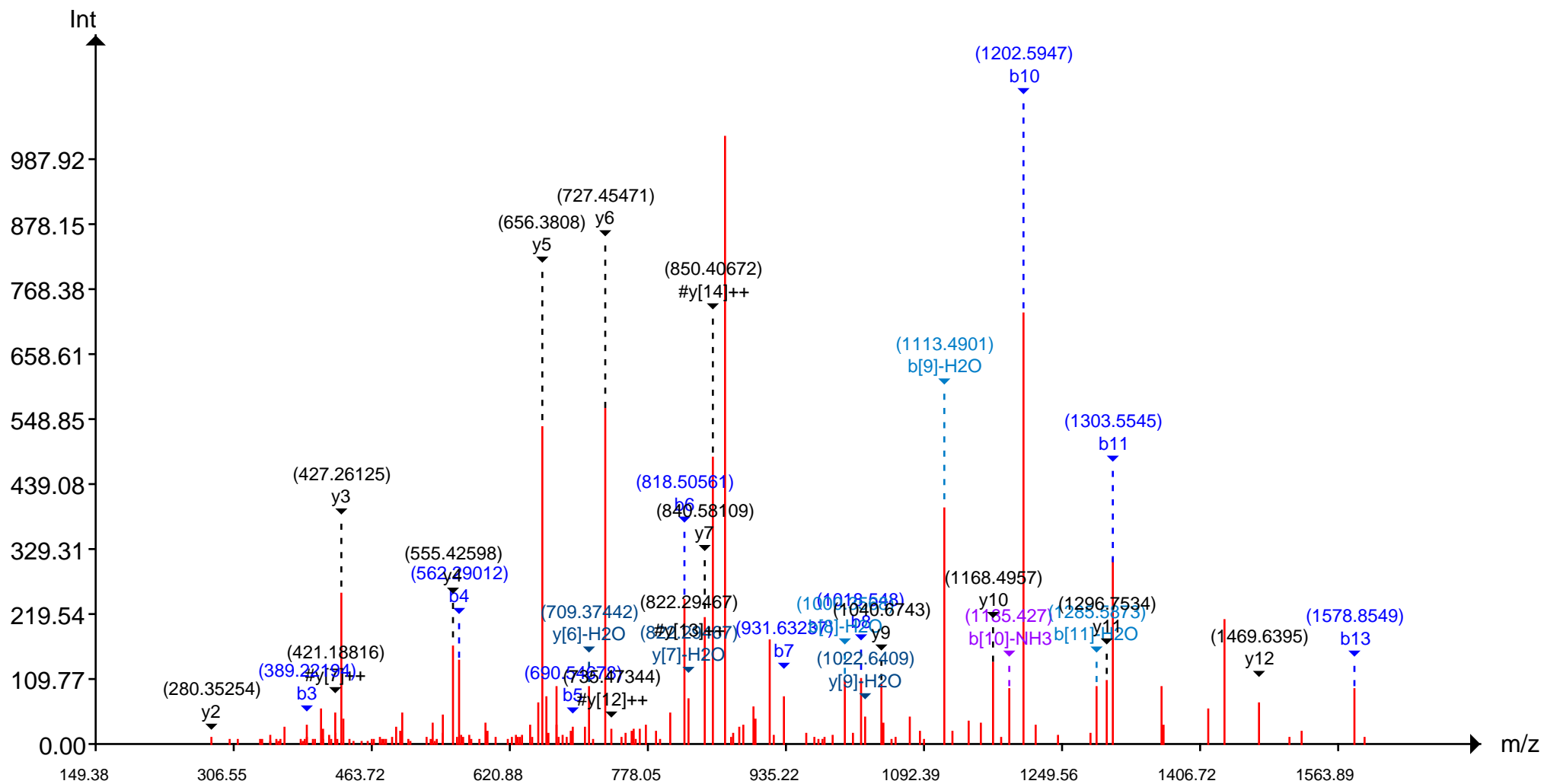
Charge 2+

Mass error -5.71E-4



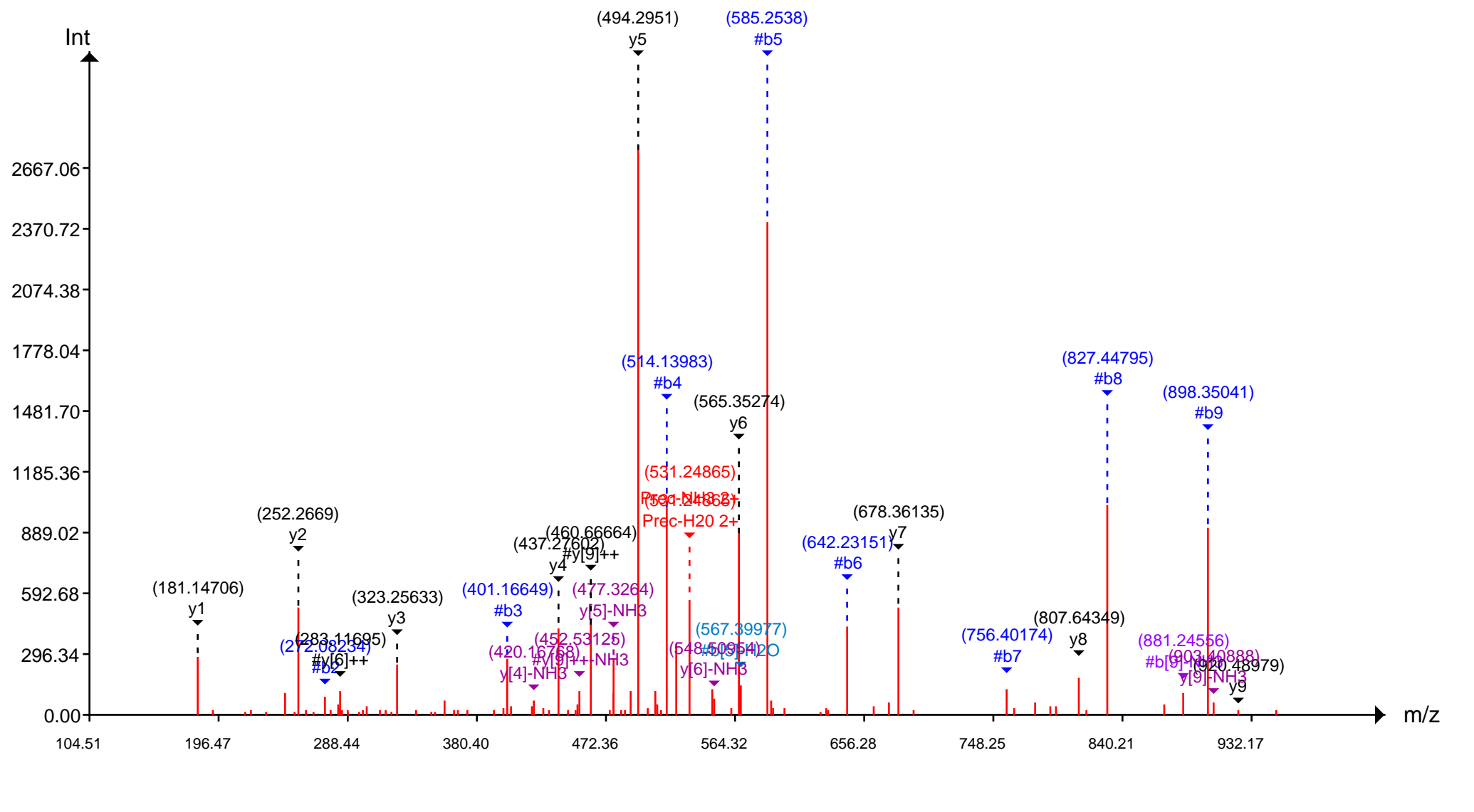
Info

Modified Sequence	AcD3-IGK<AcD3K*>K<AcD3K*>QQISLATQM<Mox*>VR-COOH	Ion score	50.31
Protein	P80316	Confidence level	99.00%
Precursor m/z	926.53798	Identity Threshold	44.56
Charge	2+	Mass error	0.001609

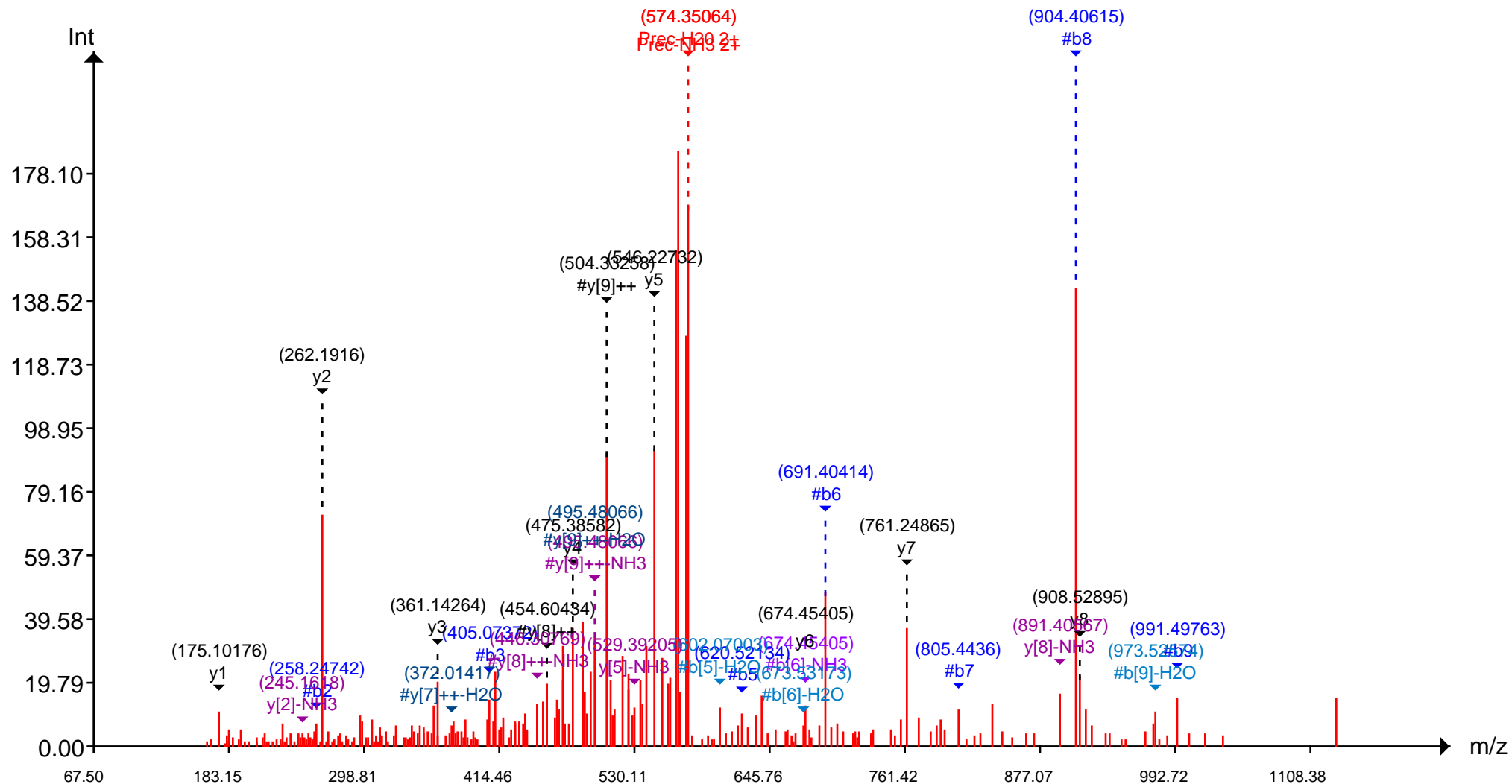


Info

Modified Sequence	AcD3-IGK<AcD3K*>K<AcD3K*>QQISLATQM<Mox*>VR<C13>-COOH	Ion score	54.84
Protein	P80316	Confidence level	99.00%
Precursor m/z	929.54808	Identity Threshold	43.34
Charge	2+	Mass error	0.001667

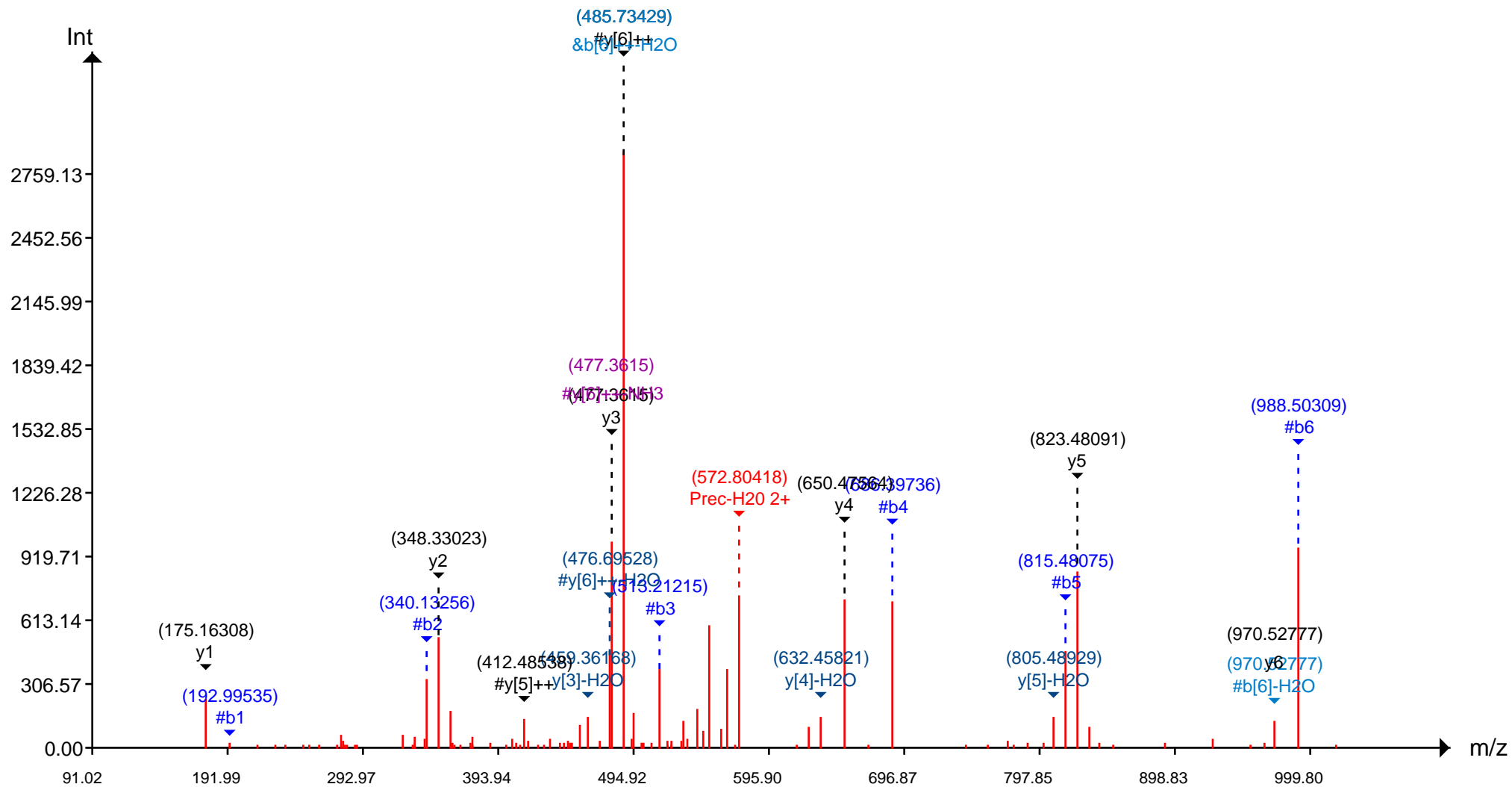


Info	
Modif...	AcD3-ILELAGNAAR<C13>-COOH
	Q8CGP5, Q8CGP6, Q8CGP7, P22752, Q6GSS7, Q64522, Q64523, Q8BFU2, Q8R1M2, Q8CCK0, P27661, Q9QZQ8
	539.82303
	2+
Ion score	
Confidence level	
Identity Threshold	
Mass error	



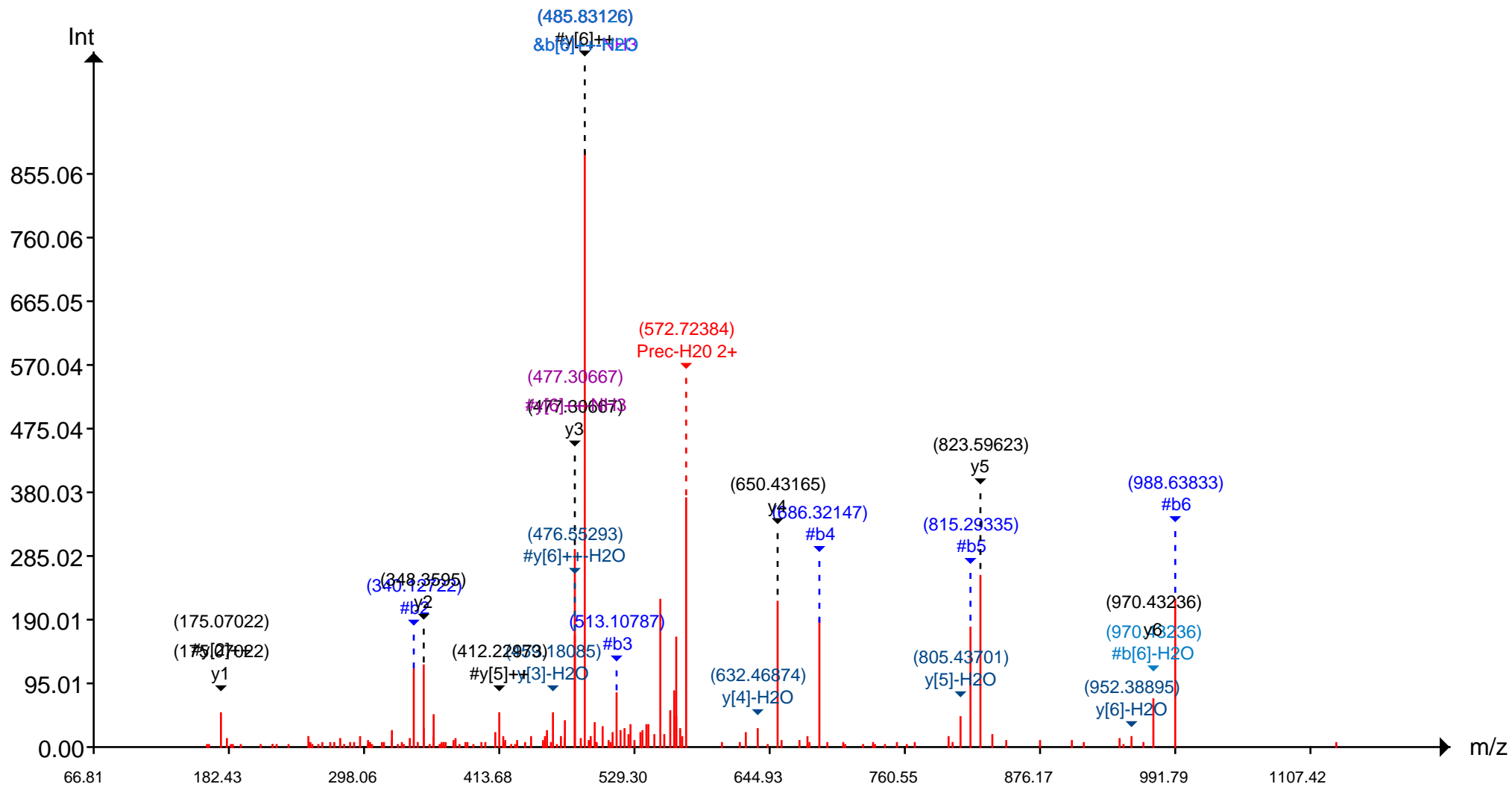
Info

Modified Sequence	AcD3-LVM<Mox*>SQANVSR-COOH	Ion score	51.37
Protein	P70670, Q60817	Confidence level	99.00%
Precursor m/z	583.30805	Identity Threshold	48.06
Charge	2+	Mass error	0.001527



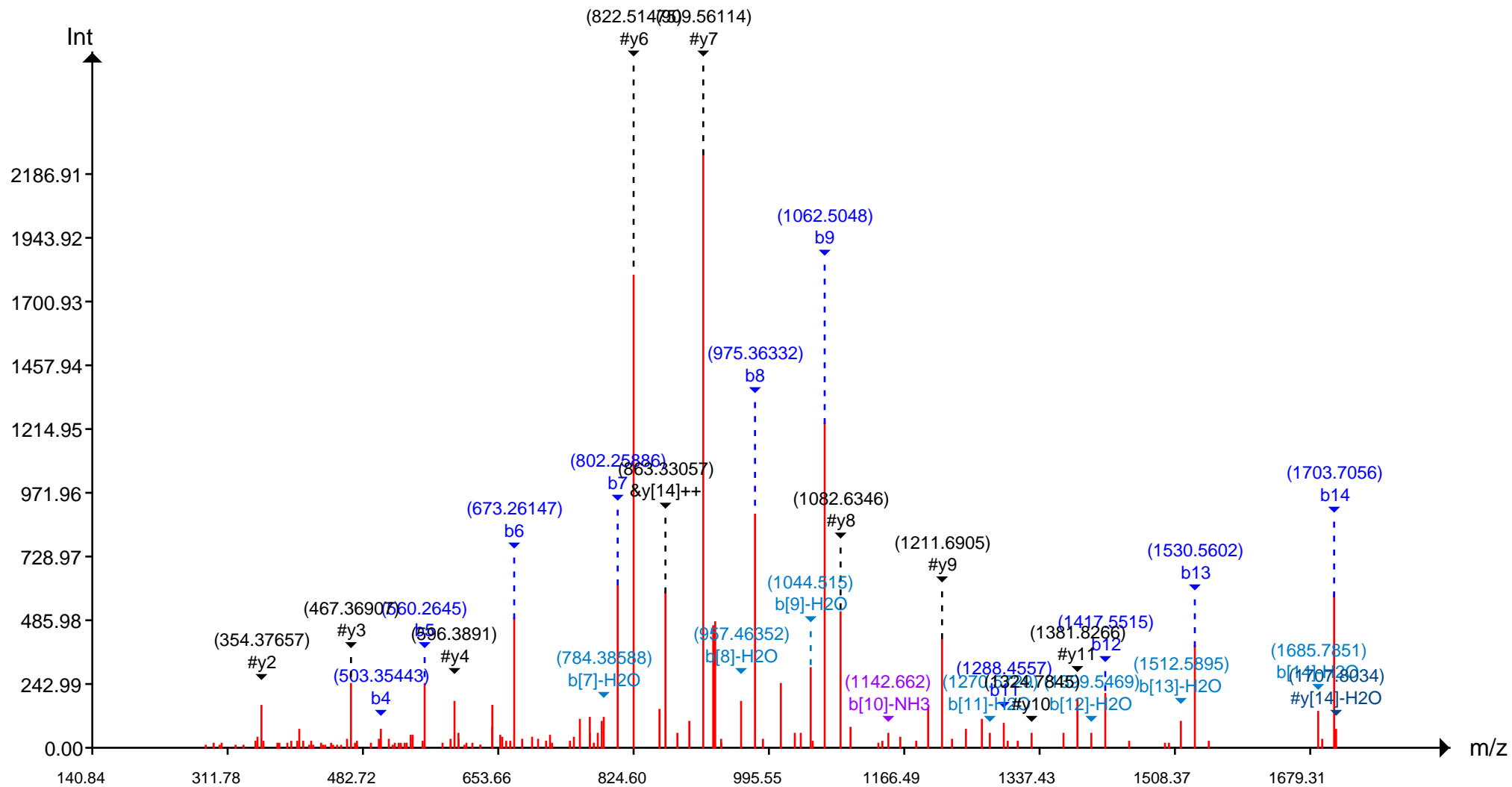
Info

Modified Sequence	AcD3-FM<Mox*>K<AcD3K*>K<AcD3K*>EK<AcD3K*>R-COOH	Ion score	51.64
Protein	P31725	Confidence level	99.00%
Precursor m/z	581.8372	Identity Threshold	46.60
Charge	2+	Mass error	-6.9E-4



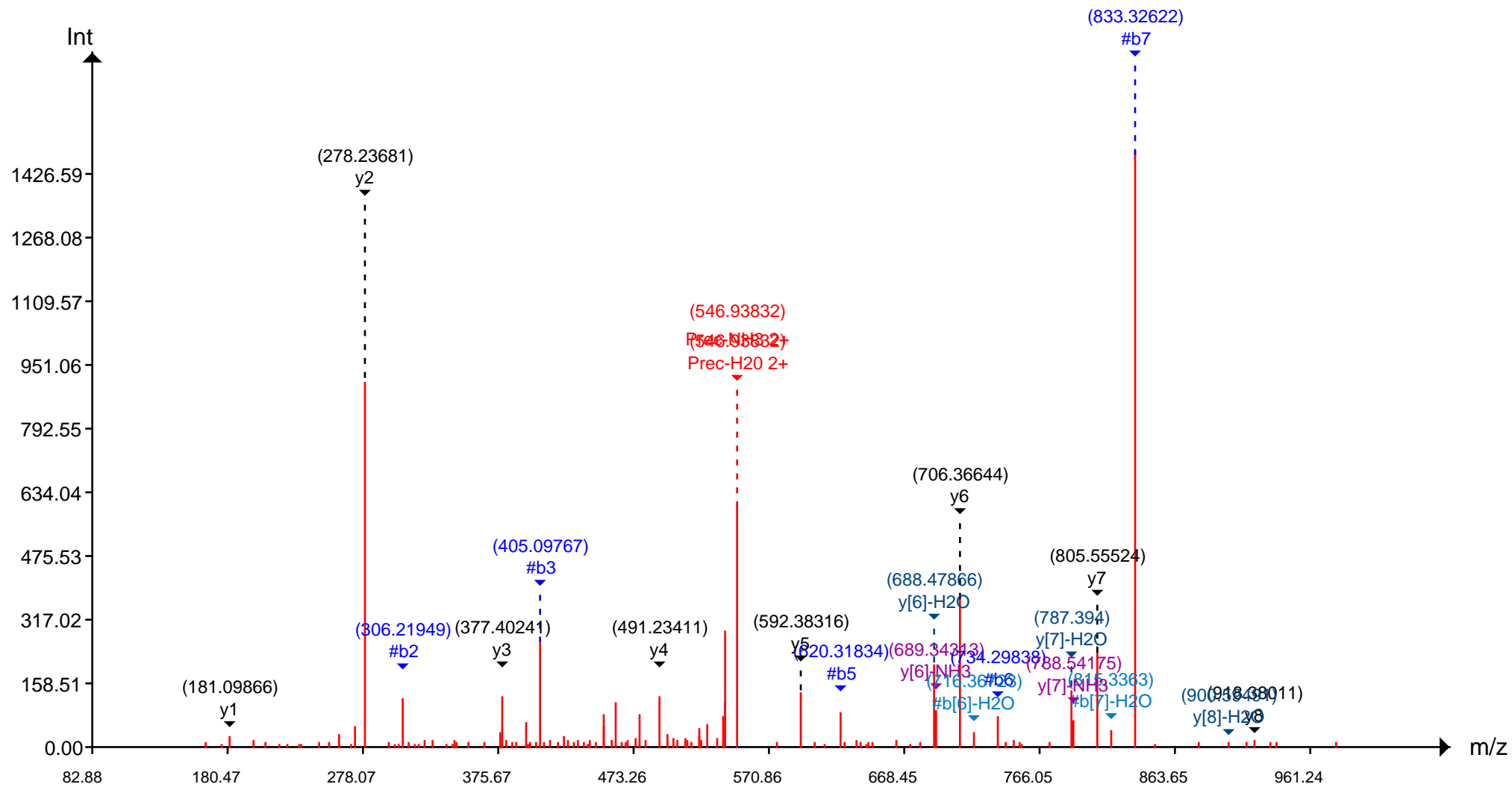
Info

Modified Sequence	AcD3-FM<Mox*>K<AcD3K*>K<AcD3K*>EK<AcD3K*>R-COOH	Ion score	49.03
Protein	P31725	Confidence level	99.00%
Precursor m/z	581.83814	Identity Threshold	46.65
Charge	2+	Mass error	0.00119



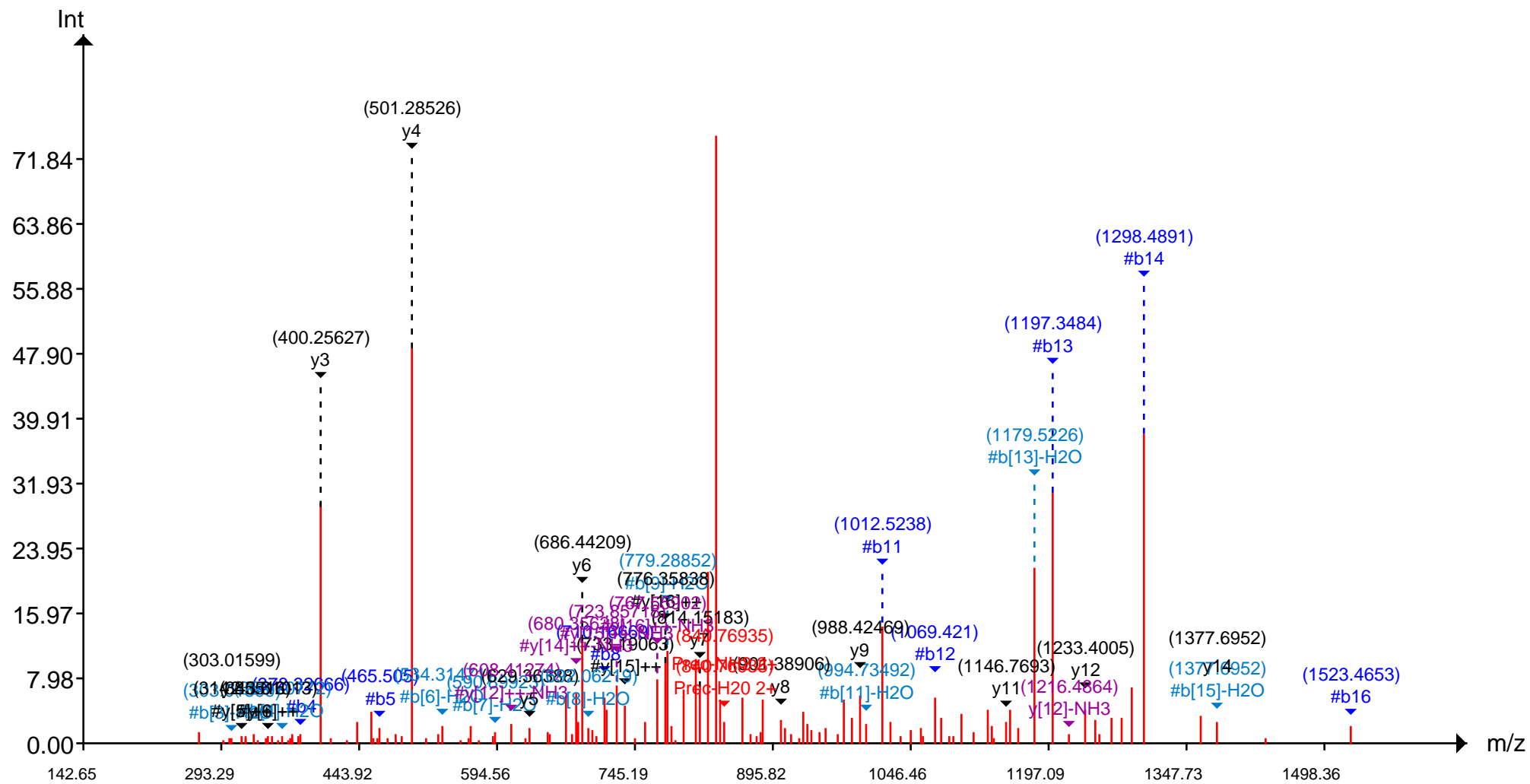
Info

Modified Sequence	AcD3-LTNEGLEK<AcD3K*>SPEEIK<AcD3K*>R<C13>-COOH	Ion score	86.2
Protein	Q99J99	Confidence level	99.00%
Precursor m/z	942.52322	Identity Threshold	46.94
Charge	2+	Mass error	0.002576

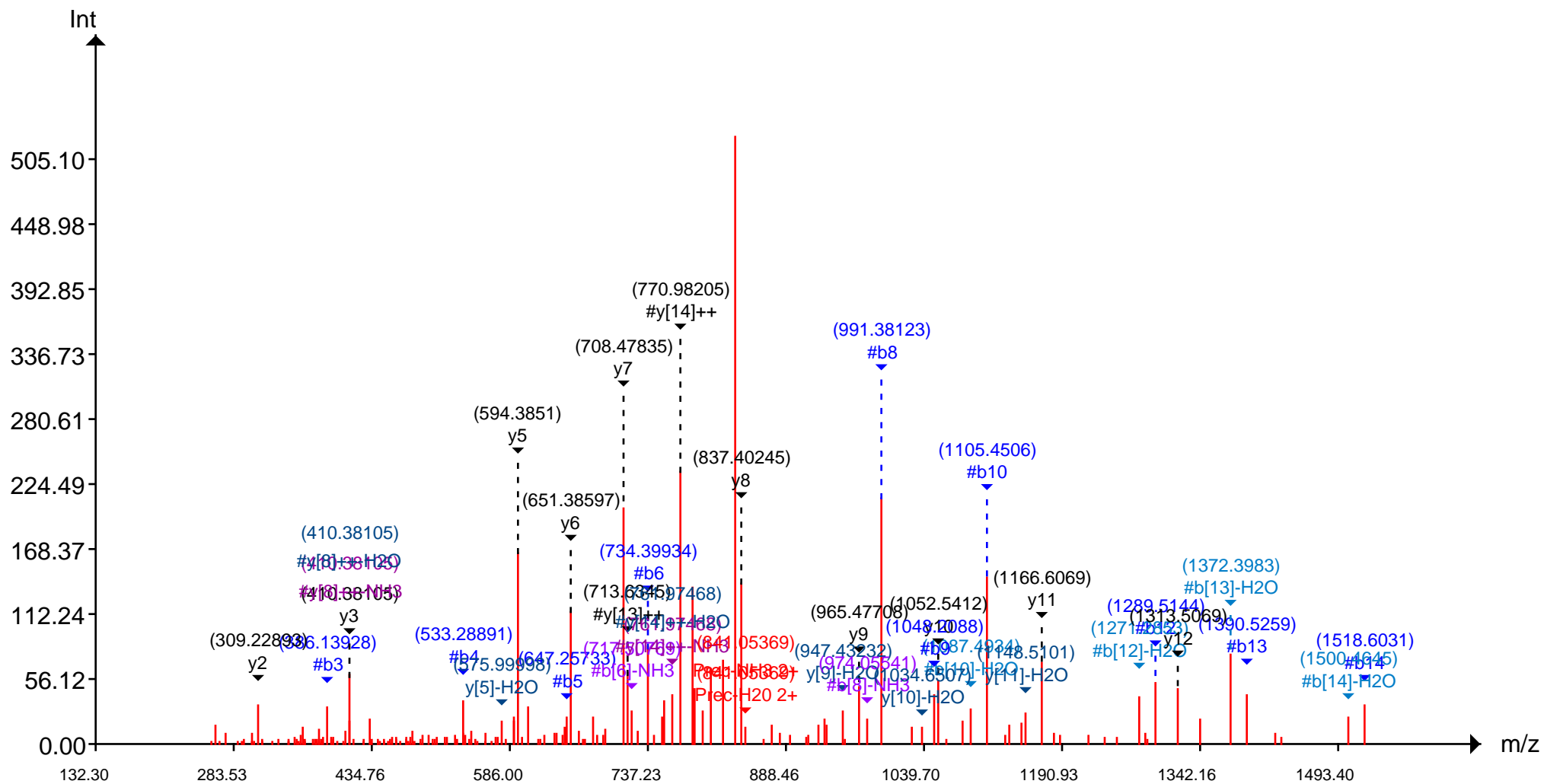


Info

Modified Sequence	AcD3-FIVNTNVPR<C13>-COOH	Ion score	68.3
Protein	P34884	Confidence level	99.00%
Precursor m/z	555.8258	Identity Threshold	46.18
Charge	2+	Mass error	2.38E-4

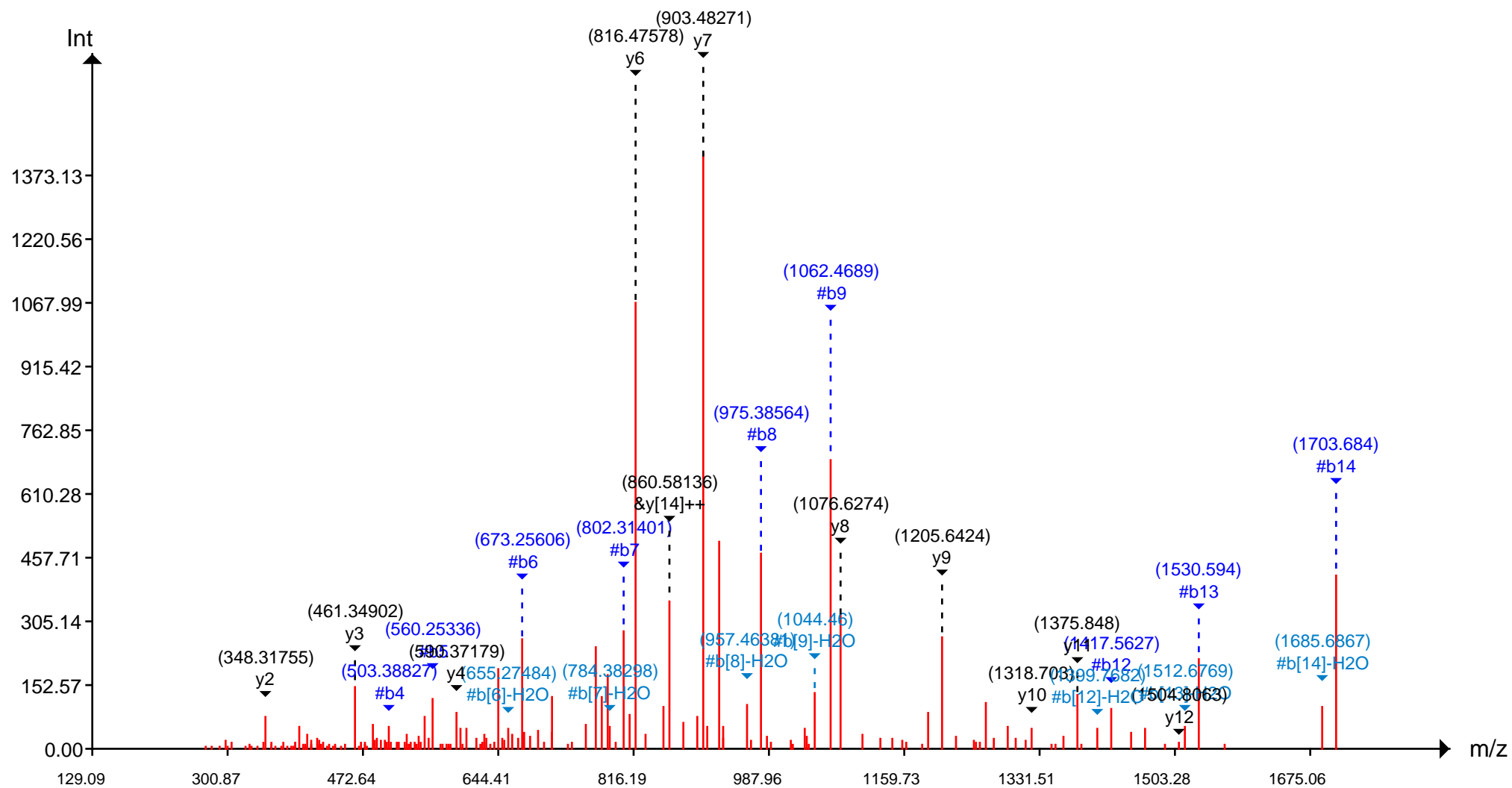


Info			
Modified Sequence	AcD3-TSSGSSGTSSQGQTPQR-COOH	Ion score	55.45
Protein	O89090	Confidence level	99.00%
Precursor m/z	849.3926	Identity Threshold	45.14
Charge	2+	Mass error	0.001468



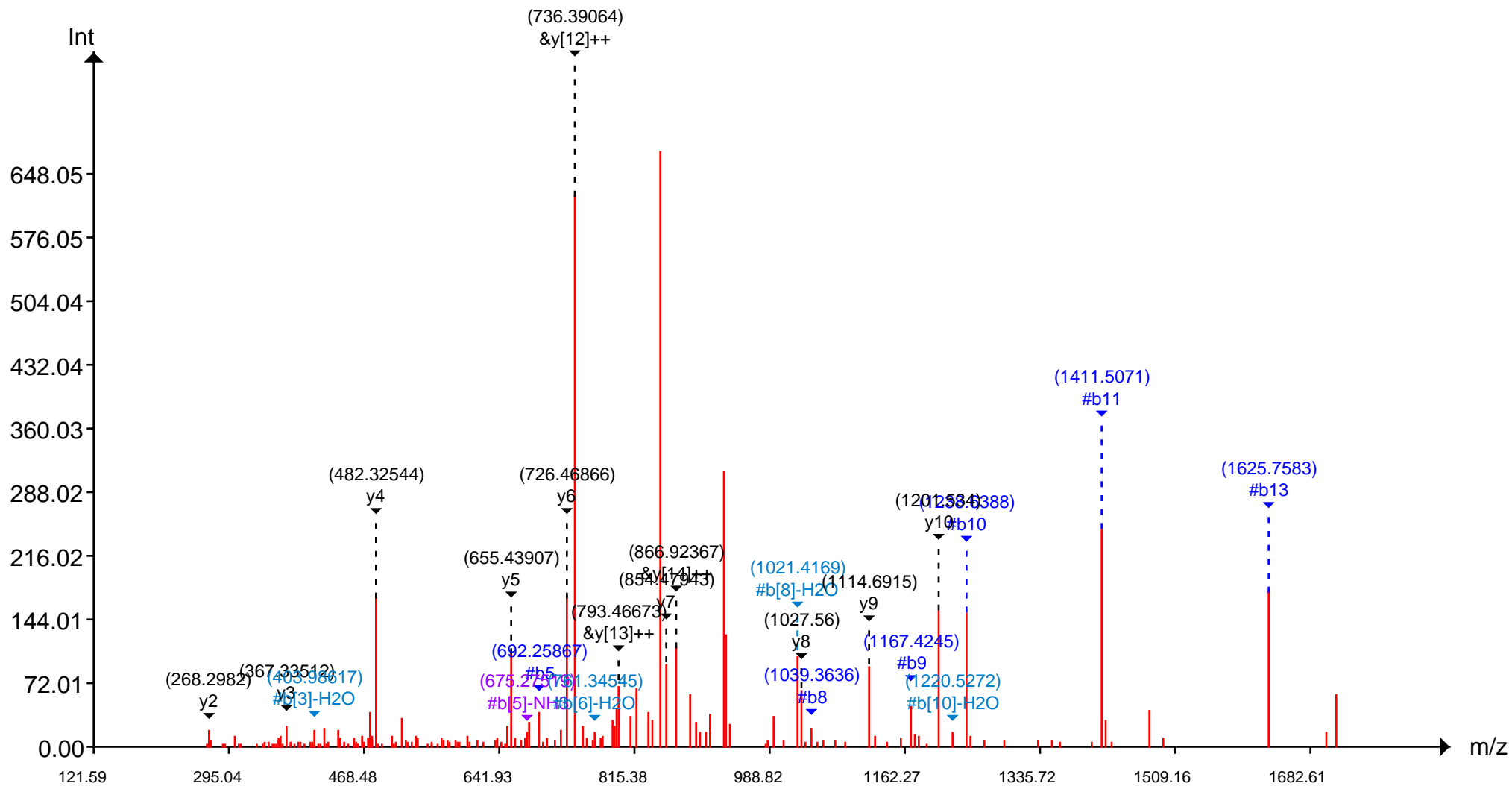
Info

Modified Sequence	AcD3-LNIM<Mox*>NSQEGGPSTQR<C13>-COOH	Ion score	75.24
Protein	P70429	Confidence level	99.00%
Precursor m/z	849.91548	Identity Threshold	47.09
Charge	2+	Mass error	-0.001341



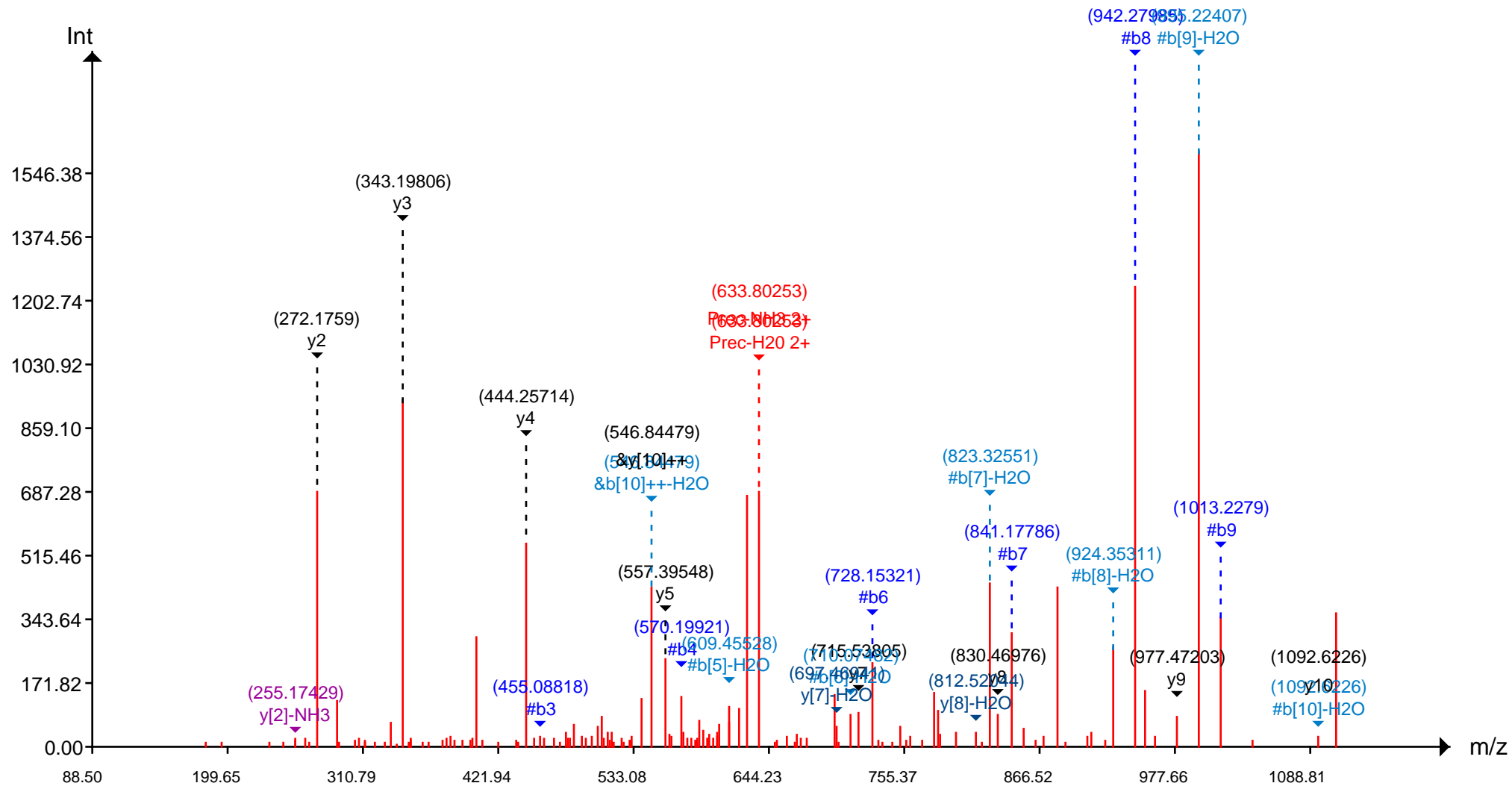
Info

Modified Sequence	AcD3-LTNEGLEK<AcD3K*>SPEEIK<AcD3K*>R-COOH	Ion score	66.89
Protein	Q99J99	Confidence level	99.00%
Precursor m/z	939.51307	Identity Threshold	47.51
Charge	2+	Mass error	0.002417



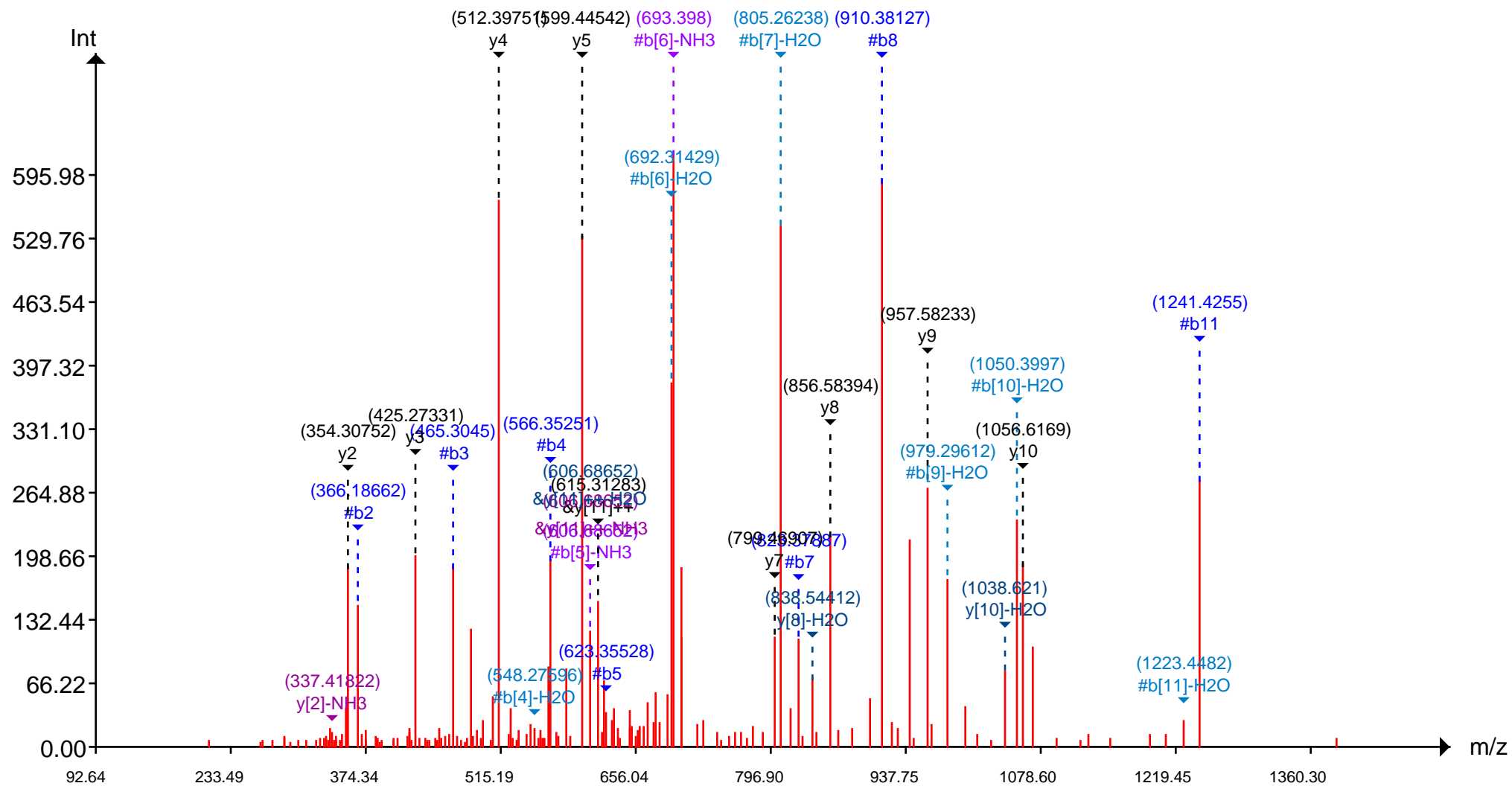
Info

Modified Sequence	AcD3-DFNPK<AcD3K*>SSK<AcD3K*>QAK<AcD3K*>DVSR<C13>-COOH	Ion score	48.85
Protein	O08585	Confidence level	99.00%
Precursor m/z	947.01434	Identity Threshold	47.65
Charge	2+	Mass error	0.001662



Info

Modified Sequence	AcD3-FDM<Mox*>DGTLTAPR-COOH	Ion score	46.6
Protein	Q9Z2M7	Confidence level	99.00%
Precursor m/z	642.80288	Identity Threshold	45.76
Charge	2+	Mass error	0.001669



Info

Modified Sequence AcD3-FK<AcD3K*>VTGSLSSAK<AcD3K*>R<C13>-COOH

Ion score 67.6

Protein P39749

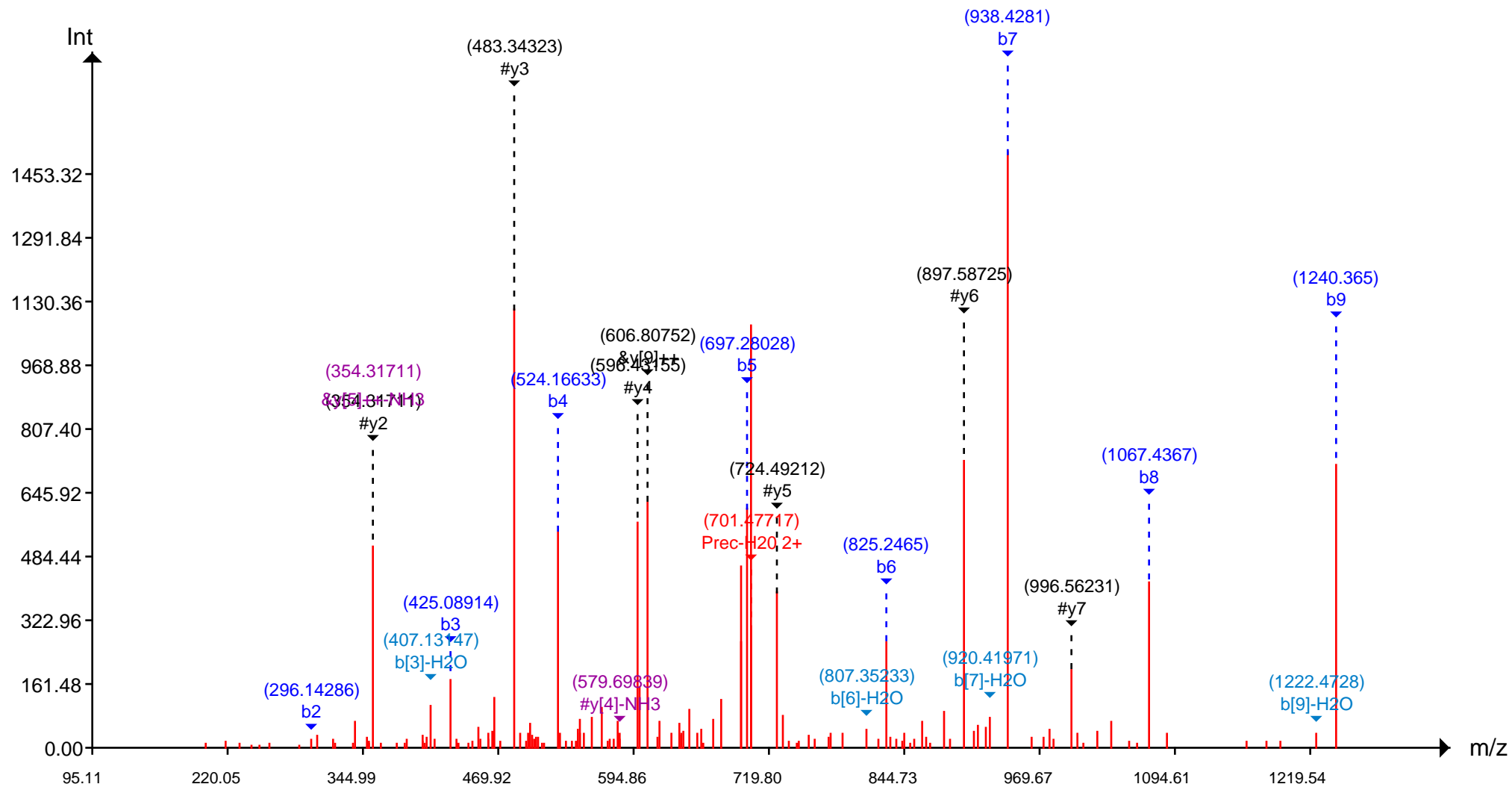
Confidence level 99.00%

Precursor m/z 711.42423

Identity Threshold 44.53

Charge 2+

Mass error 7.48E-4



Info

Modified Sequence AcD3-YSEVK<AcD3K*>QIEK<AcD3K*>R<C13>-COOH

Ion score 61.86

Protein Q6NZB0

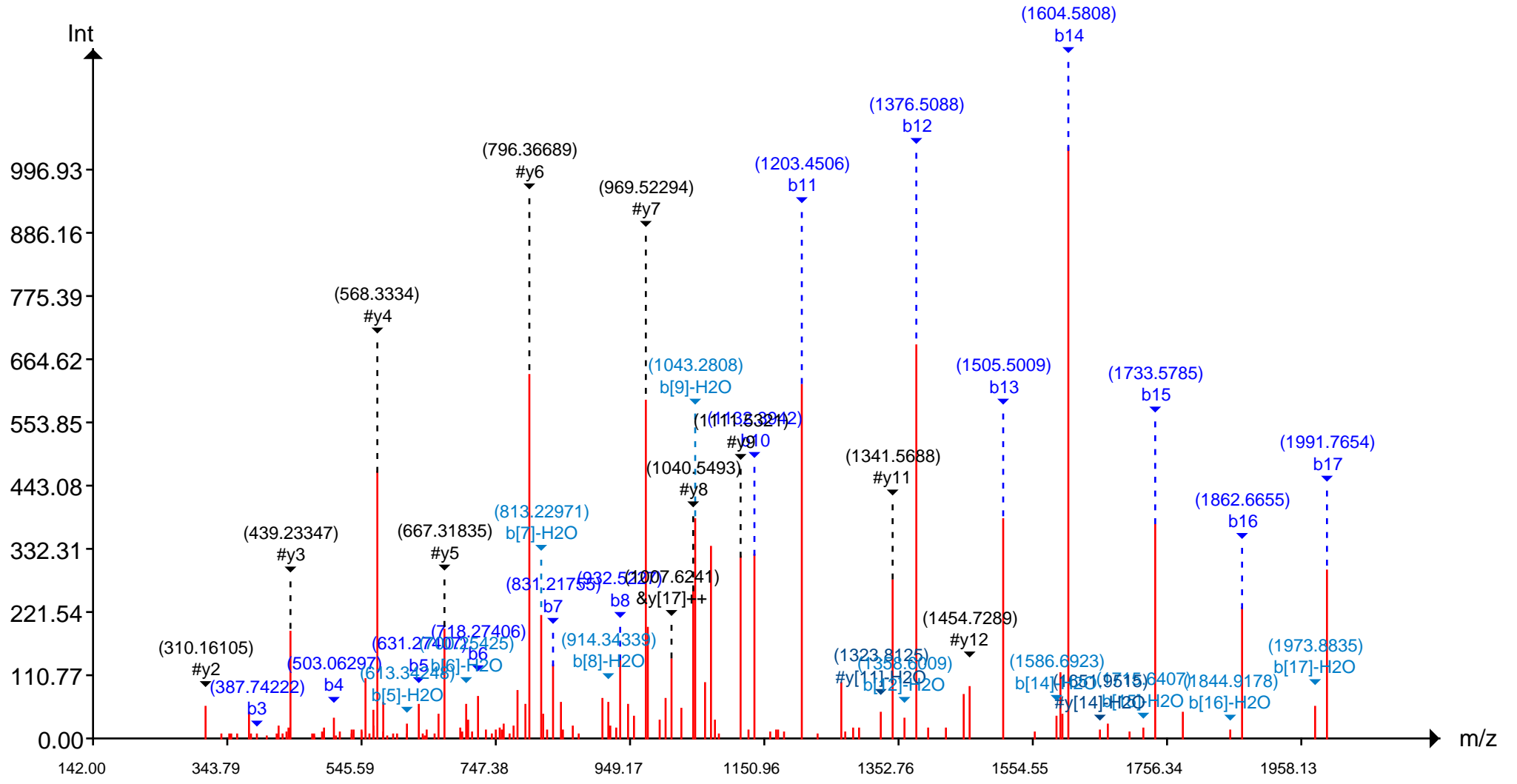
Confidence level 99.00%

Precursor m/z 710.90837

Identity Threshold 46.28

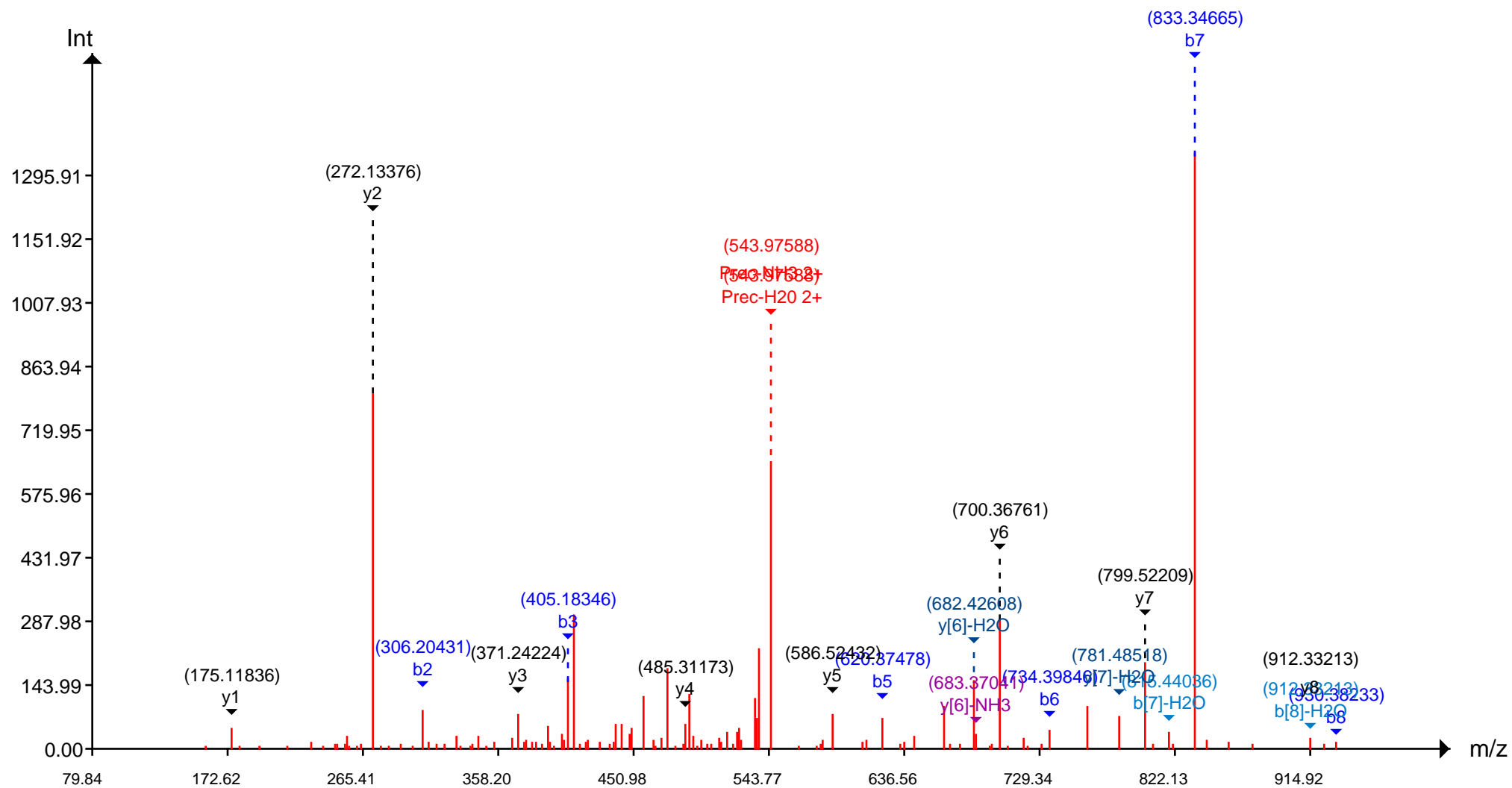
Charge 2+

Mass error 6.9E-4



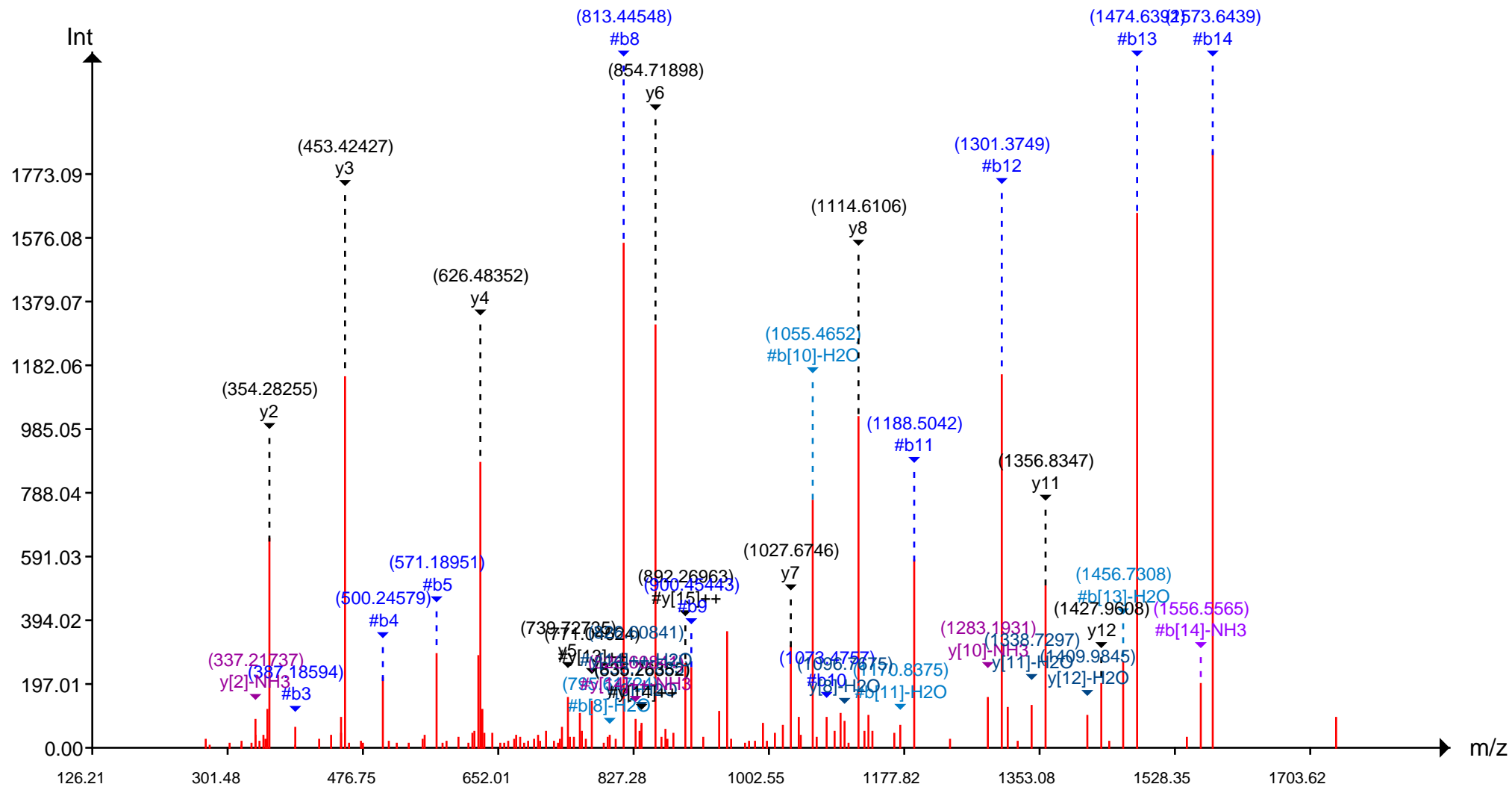
Info

Modified Sequence	AcD3-LQTDQSLTEAAK<AcD3K*>EVEEER<C13>-COOH	Ion score	99.47
Protein	Q9Z0E6	Confidence level	99.00%
Precursor m/z	1086.5489	Identity Threshold	47.77
Charge	2+	Mass error	0.002498



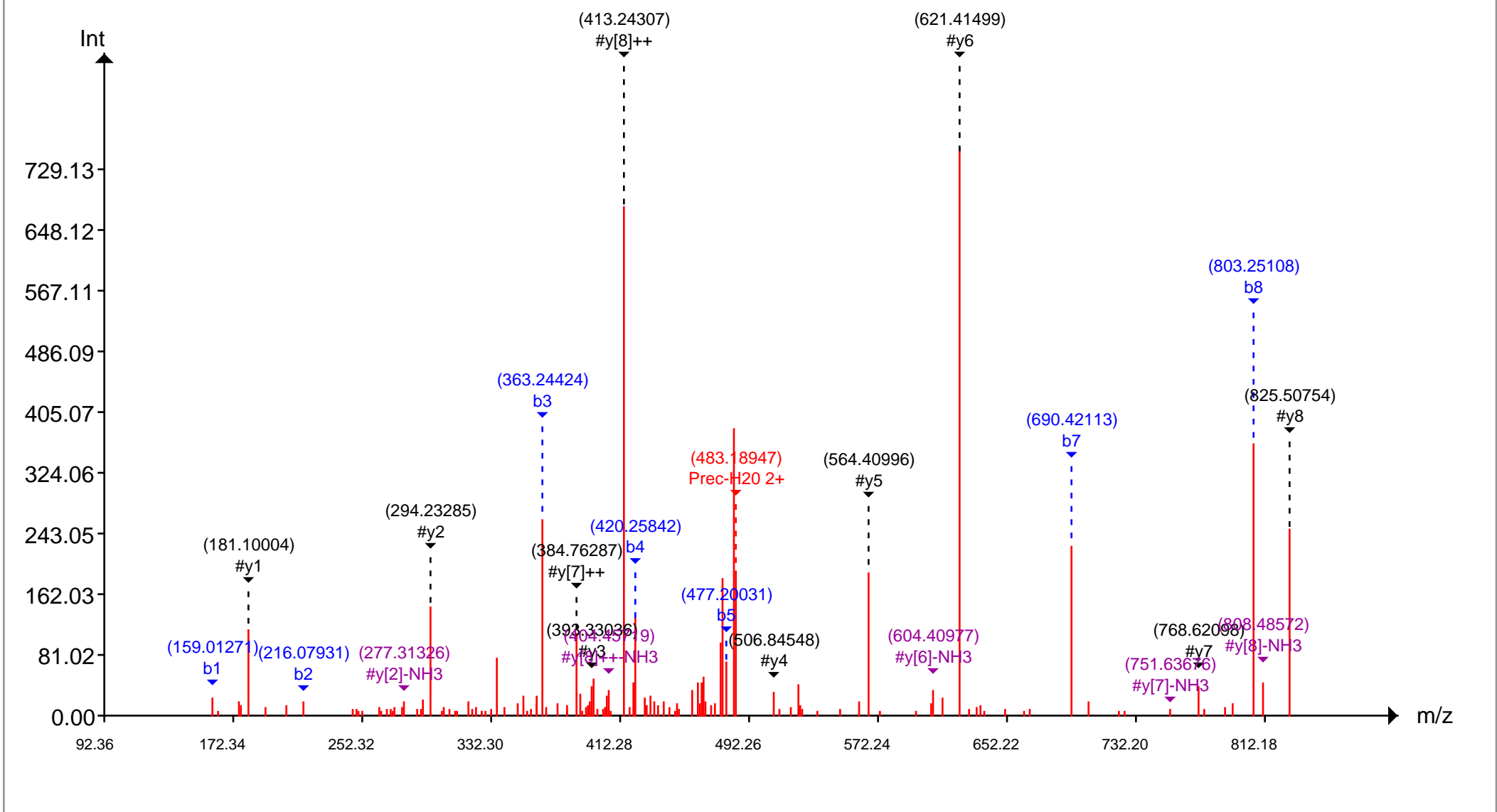
Info

Modified Sequence	AcD3-FIVNTNVPR-COOH	Ion score	55.23
Protein	P34884	Confidence level	99.00%
Precursor m/z	552.81589	Identity Threshold	47.33
Charge	2+	Mass error	5.59E-4



Info

Modified Sequence	AcD3-VLELAGNASK<AcD3K*>DLK<AcD3K*>VK<AcD3K*>R<C13>-COOH	Ion score	70.71
Protein	Q3THW5, P0C0S6	Confidence level	99.00%
Precursor m/z	964.09064	Identity Threshold	39.21
Charge	2+	Mass error	0.003261



Info

Modified Sequence AcD3-IGFGGNVIR<C13>-COOH

Protein Q99LS3

Precursor m/z 492.29382

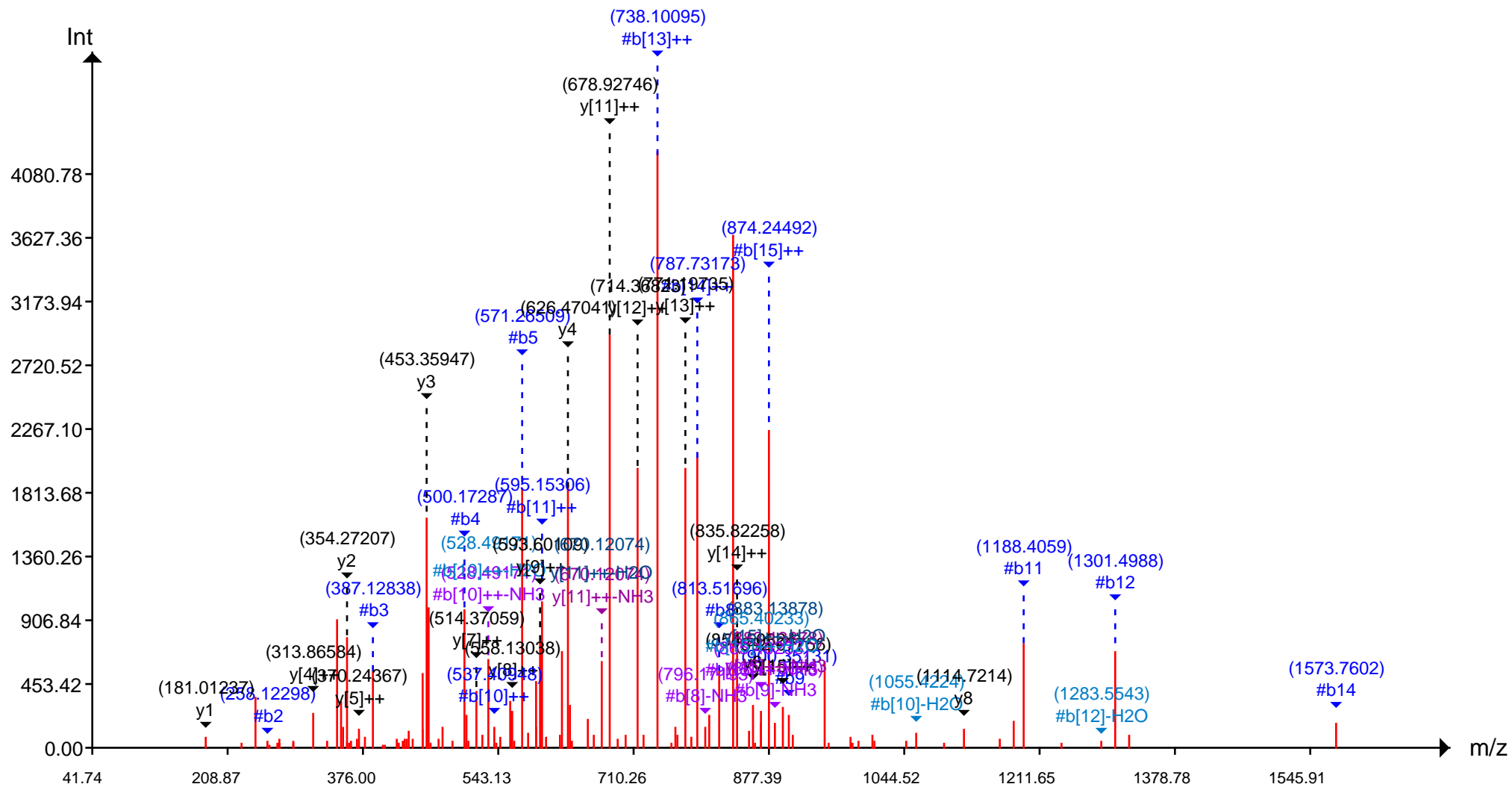
Charge 2+

Ion score 47.5

Confidence level 99.00%

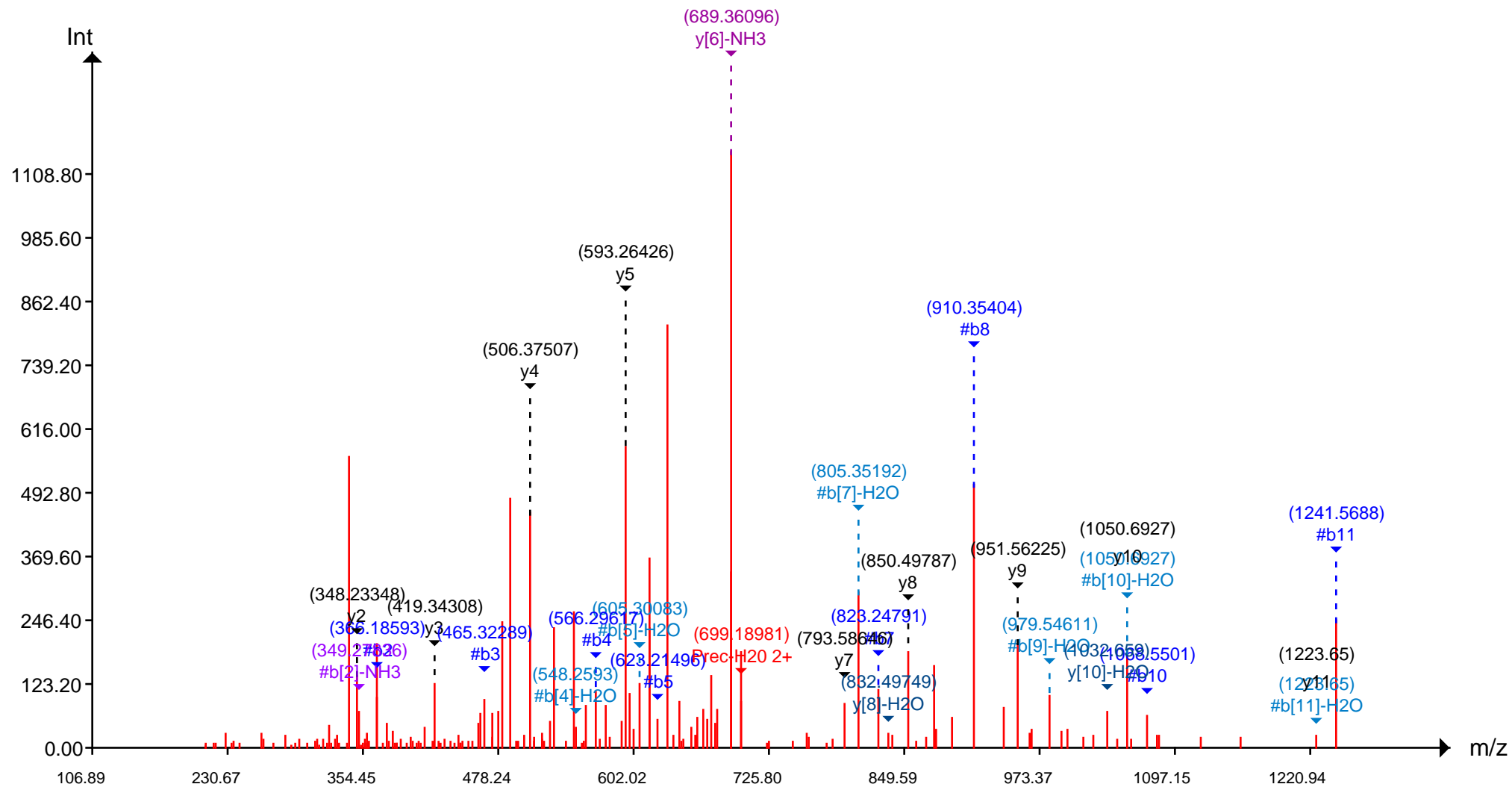
Identity Threshold 46.01

Mass error -3.98E-4



Info

Modified Sequence	AcD3-VLELAGNASK<AcD3K*>DLK<AcD3K*>VK<AcD3K*>R<C13>-COOH	Ion score	58.67
Protein	Q3THW5, P0C0S6	Confidence level	99.00%
Precursor m/z	643.06234	Identity Threshold	39.39
Charge	3+	Mass error	0.001725



Info

Modified Sequence AcD3-FK<AcD3K*>VTGSLSSAK<AcD3K*>R-COOH

Ion score 55.45

Protein P39749

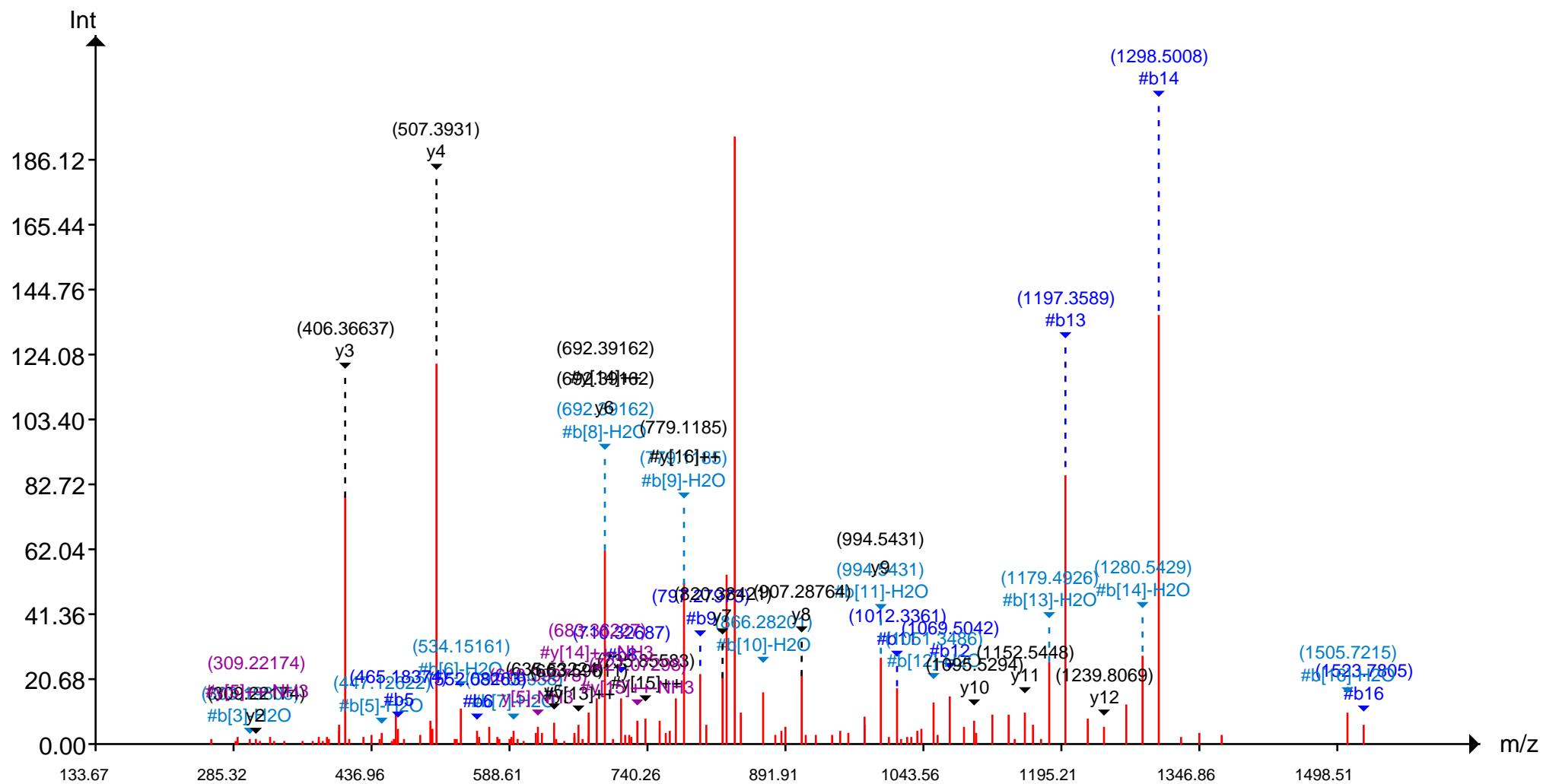
Confidence level 99.00%

Precursor m/z 708.41412

Identity Threshold 45.67

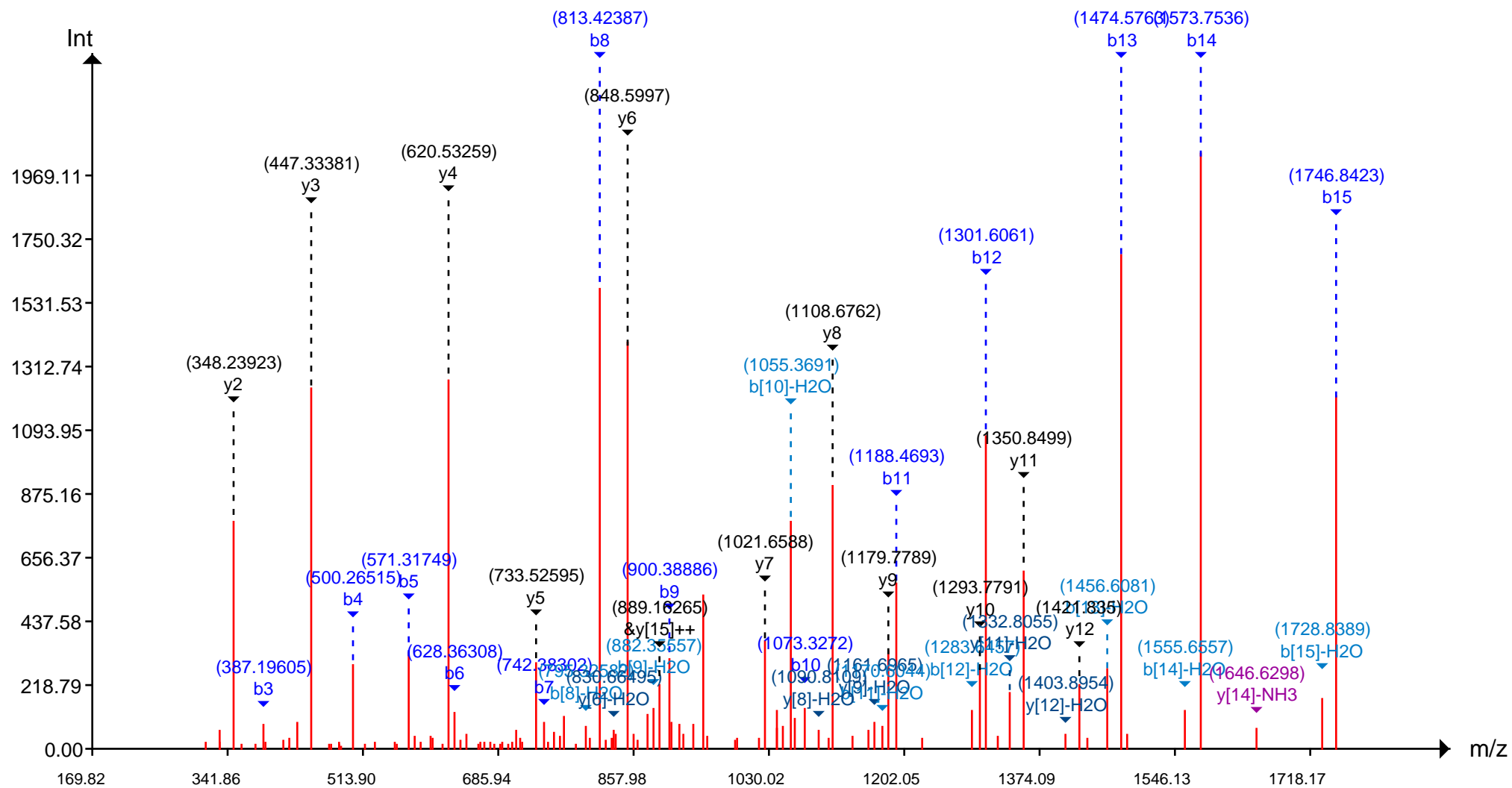
Charge 2+

Mass error 6.69E-4



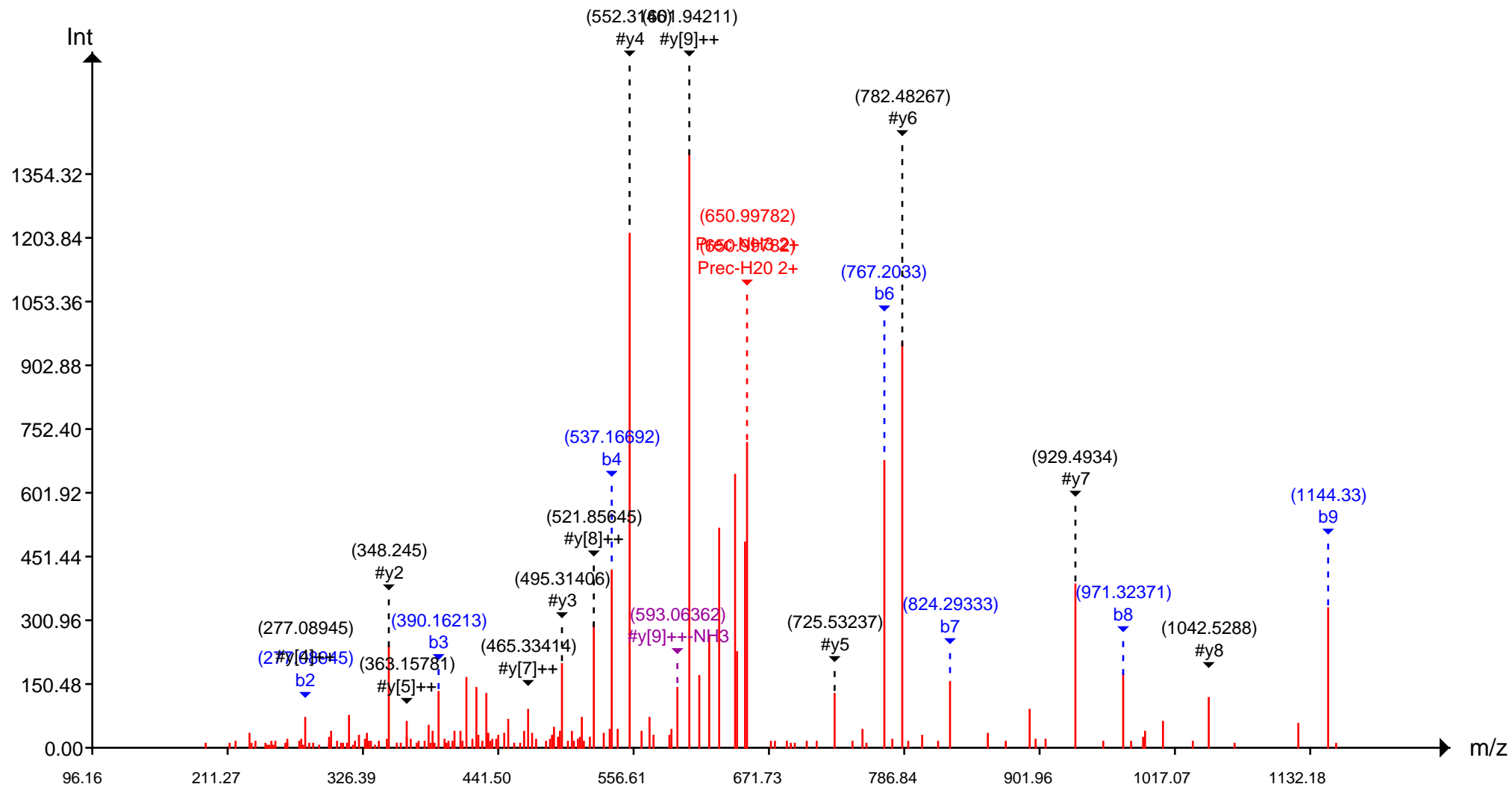
Info

Modified Sequence	AcD3-TSSGSSGTSSQGQTPQR<C13>-COOH	Ion score	74.45
Protein	O89090	Confidence level	99.00%
Precursor m/z	852.40281	Identity Threshold	46.01
Charge	2+	Mass error	0.001746



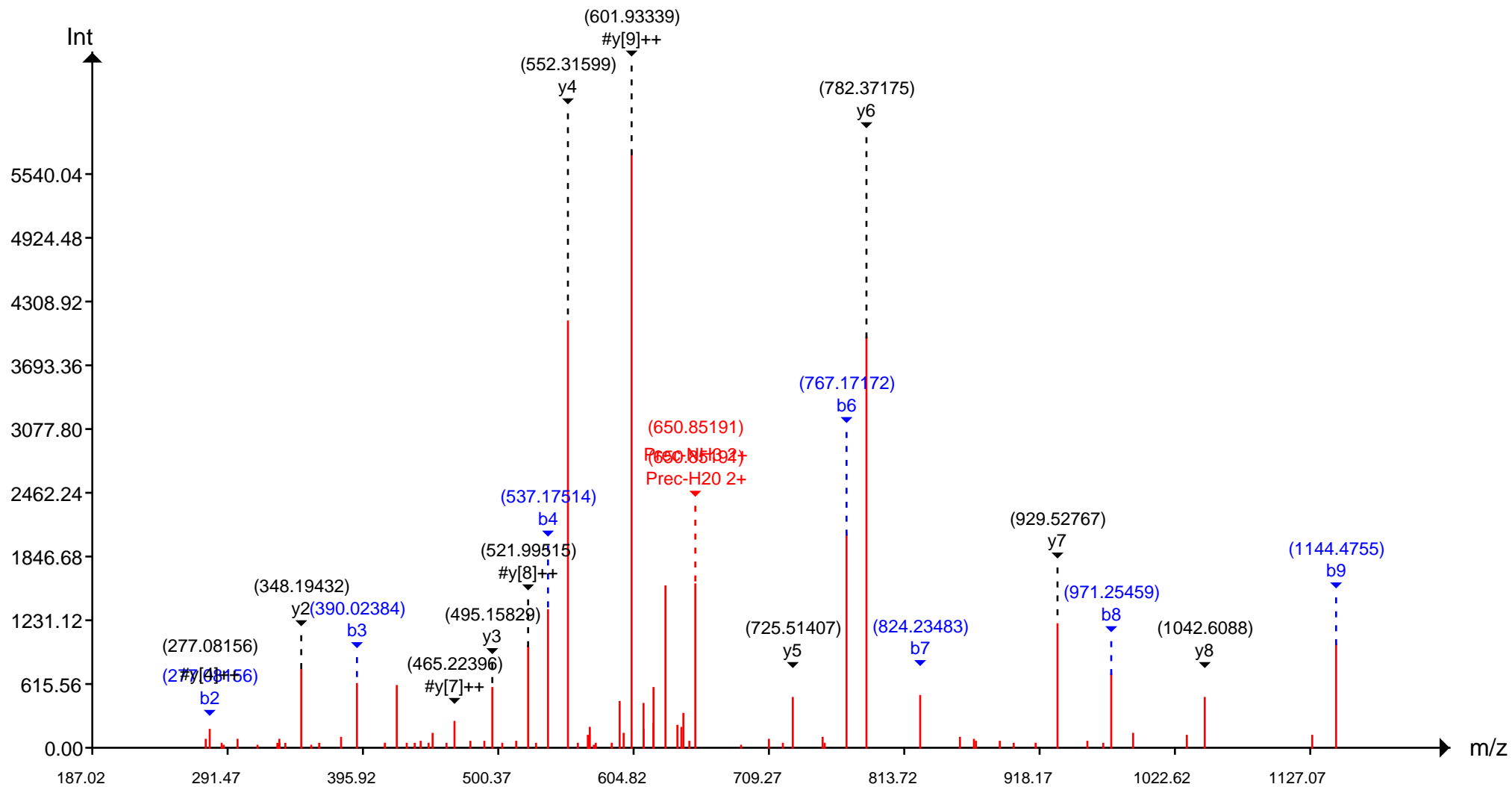
Info

Modified Sequence	AcD3-VLELAGNASK<AcD3K*>DLK<AcD3K*>VK<AcD3K*>R-COOH	Ion score	105.89
Protein	Q3THW5, P0C0S6	Confidence level	99.00%
Precursor m/z	961.08039	Identity Threshold	41.03
Charge	2+	Mass error	0.002902



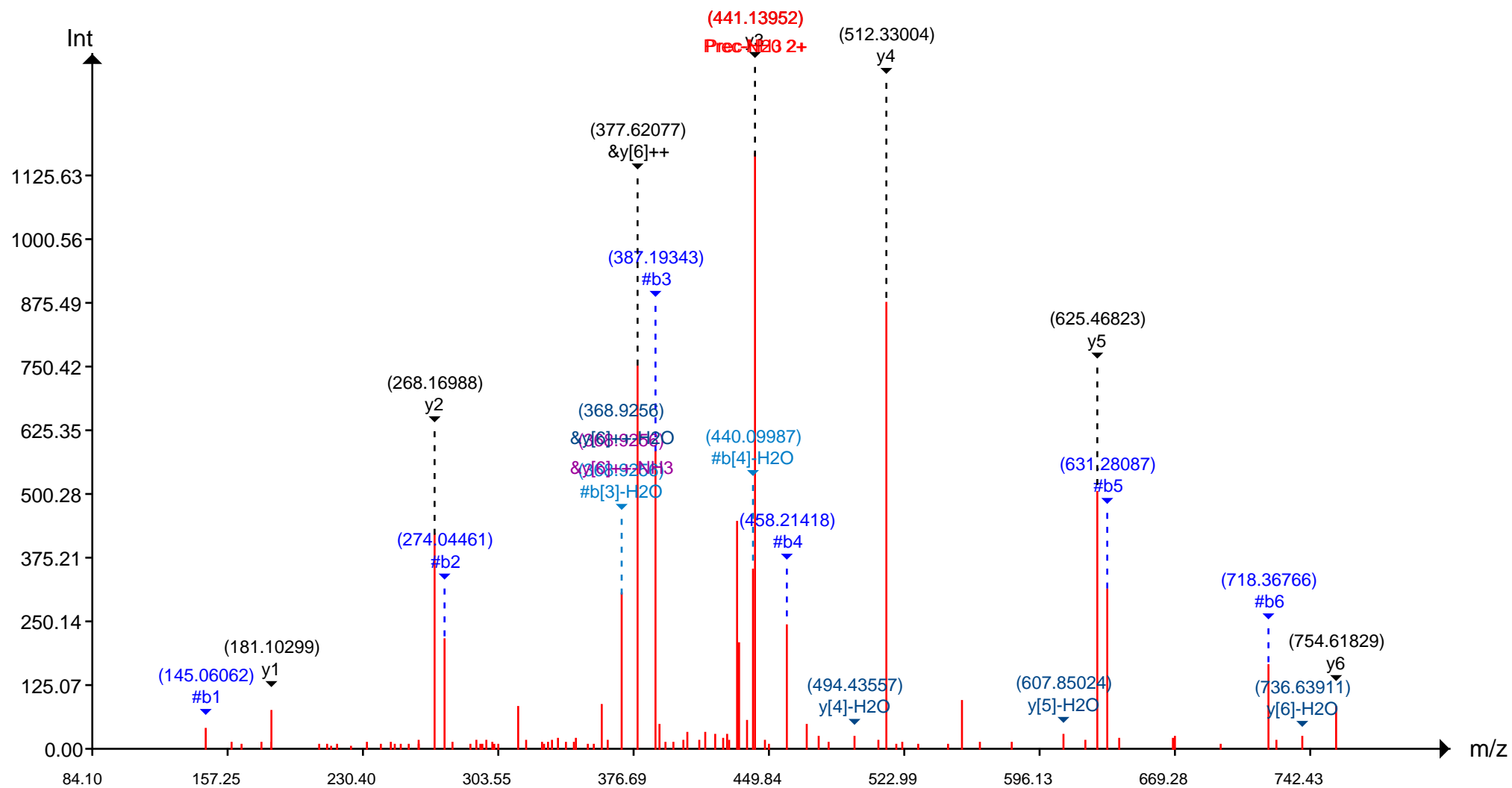
Info

Modified Sequence	AcD3-AC<Cmm*>LM<Mox*>GK<AcD3K*>GM<Mox*>K<AcD3K*>R-COOH	Ion score	51.11
Protein	Q61233	Confidence level	99.00%
Precursor m/z	659.83487	Identity Threshold	47.79
Charge	2+	Mass error	-1.46E-4



Info

Modified Sequence	AcD3-AC<Cmm*>LM<Mox*>GK<AcD3K*>GM<Mox*>K<AcD3K*>R-COOH	Ion score	62.23
Protein	Q61233	Confidence level	99.00%
Precursor m/z	659.83454	Identity Threshold	47.68
Charge	2+	Mass error	-8.06E-4



Info

Modified Sequence AcD3-VELAK<AcD3K*>SR<C13>-COOH

Ion score 61.28

Protein P06745, Q6GQX2

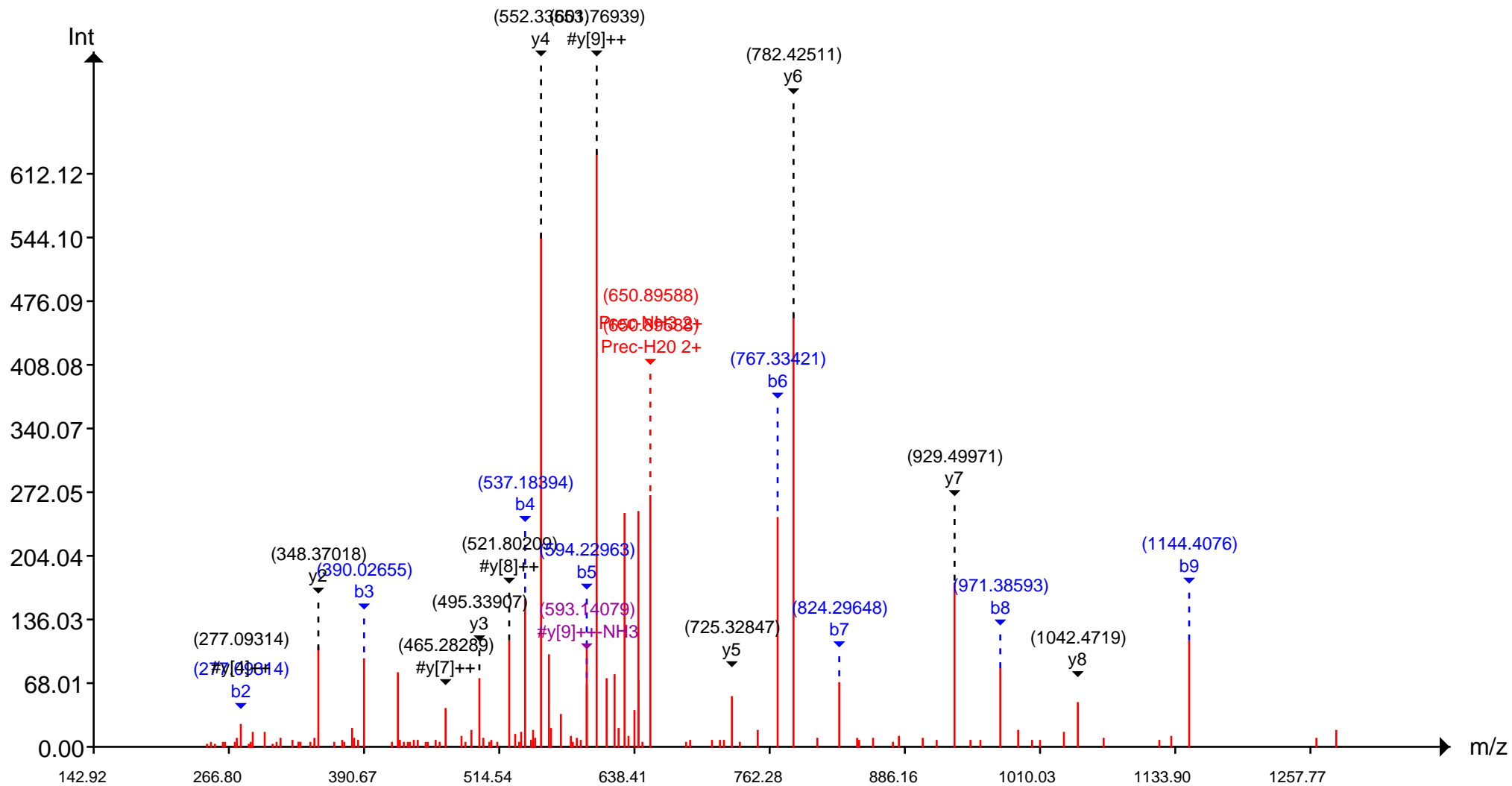
Confidence level 99.00%

Precursor m/z 449.78209

Identity Threshold 44.03

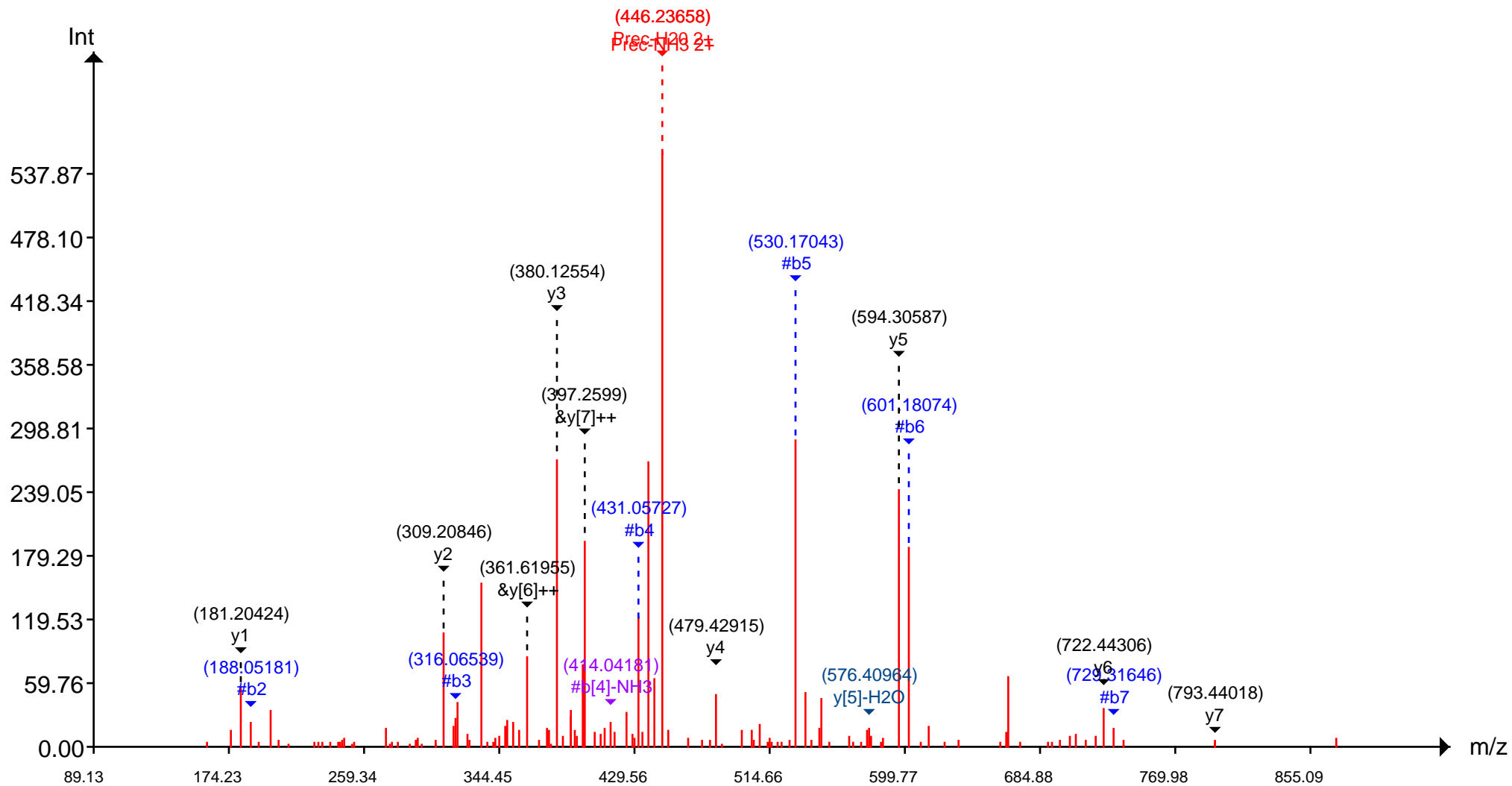
Charge 2+

Mass error -1.31E-4



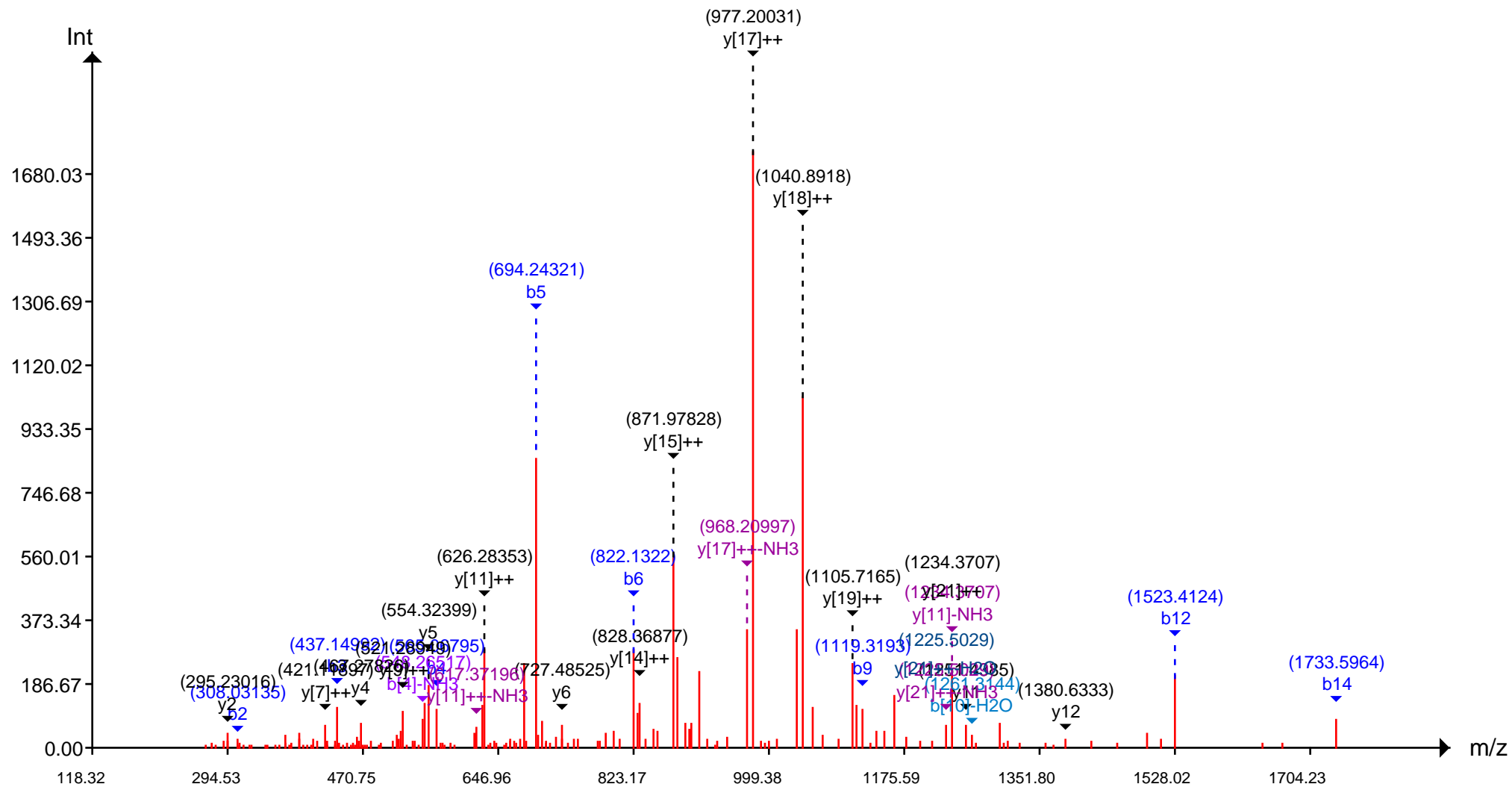
Info

Modified Sequence	AcD3-AC<Cmm*>LM<Mox*>GK<AcD3K*>GM<Mox*>K<AcD3K*>R-COOH	Ion score	48.49
Protein	Q61233	Confidence level	99.00%
Precursor m/z	659.83562	Identity Threshold	47.76
Charge	2+	Mass error	0.001354



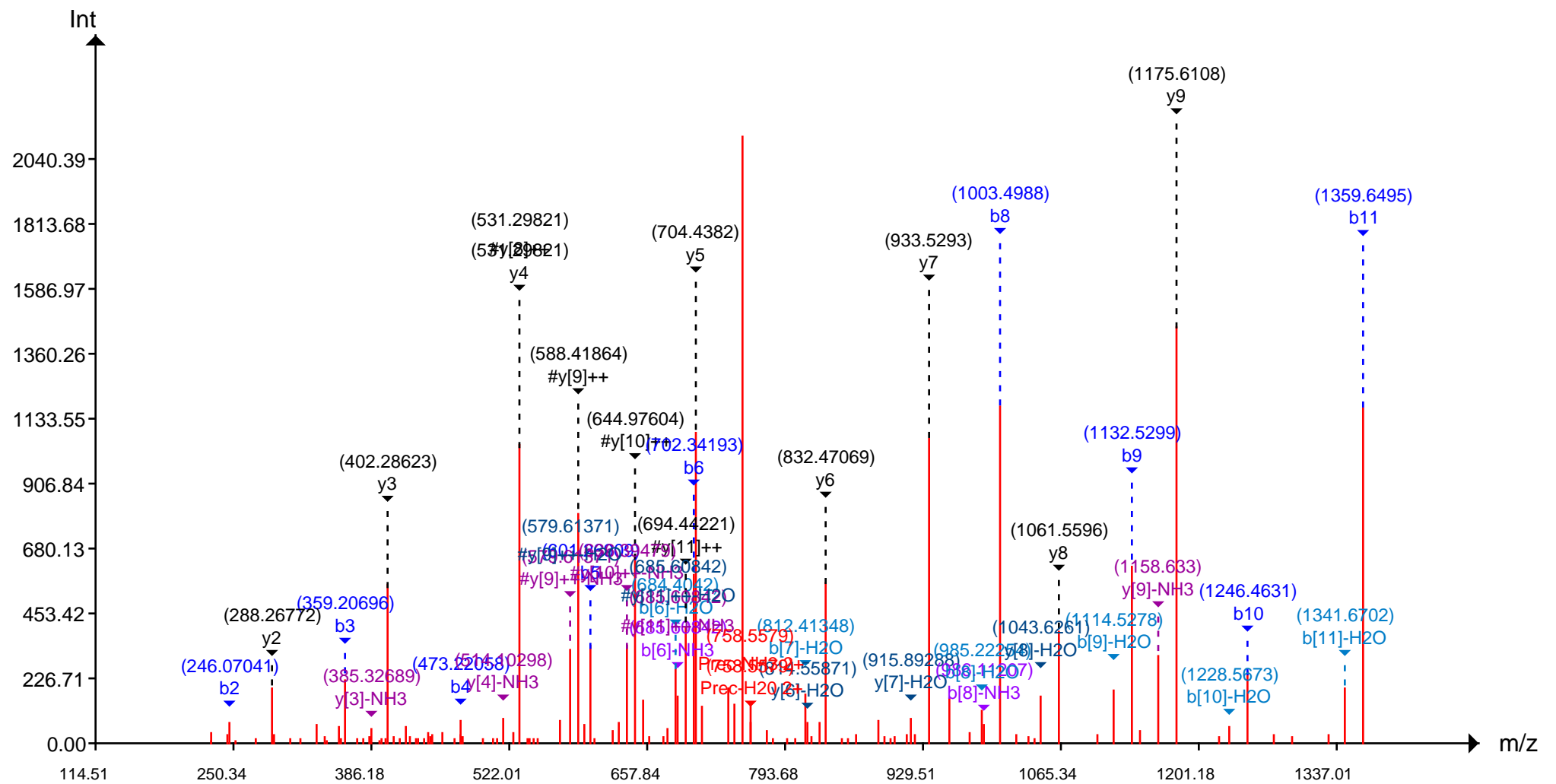
Info

Modified Sequence	AcD3-AAQDVAQR<C13>-COOH	Ion score	53.19
Protein	P62264	Confidence level	99.00%
Precursor m/z	455.24969	Identity Threshold	47.83
Charge	2+	Mass error	-2.18E-4

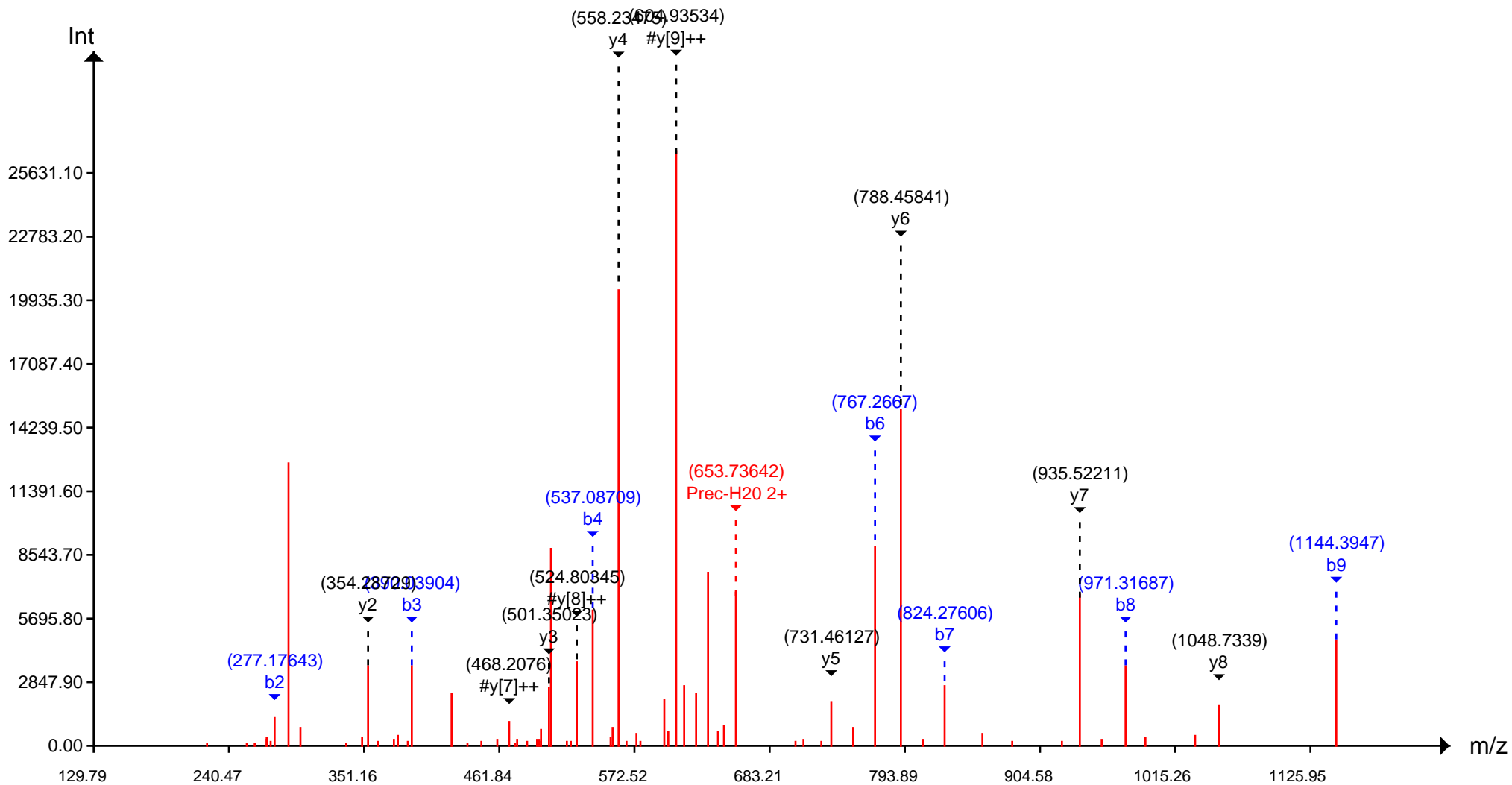


Info

Modified Sequence	AcD3-VYEQQLPSC<Cmm*>DEPILSNK<AcD3K*>SGDNR<C13>-COOH	Ion score	49.0
Protein	P70296	Confidence level	99.00%
Precursor m/z	925.44356	Identity Threshold	46.00
Charge	3+	Mass error	9.75E-4

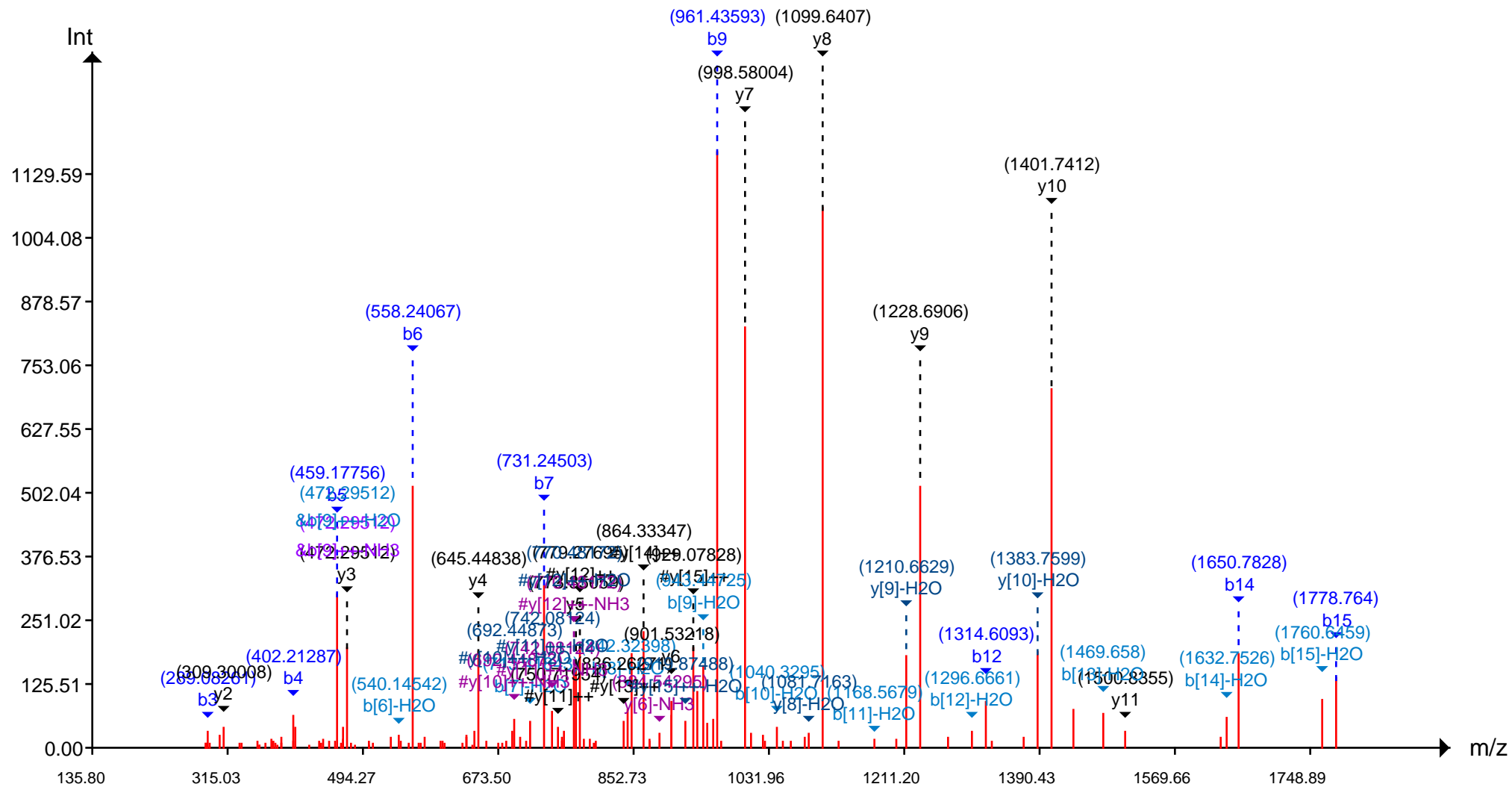


Info		Ion score	96.67
Modified Sequence	AcD3-TVINQTQK<AcD3K*>ENLR-COOH	Confidence level	99.00%
Protein	Q6ZWW7	Identity Threshold	47.36
Precursor m/z	767.4287	Mass error	-6.3E-5
Charge	2+		



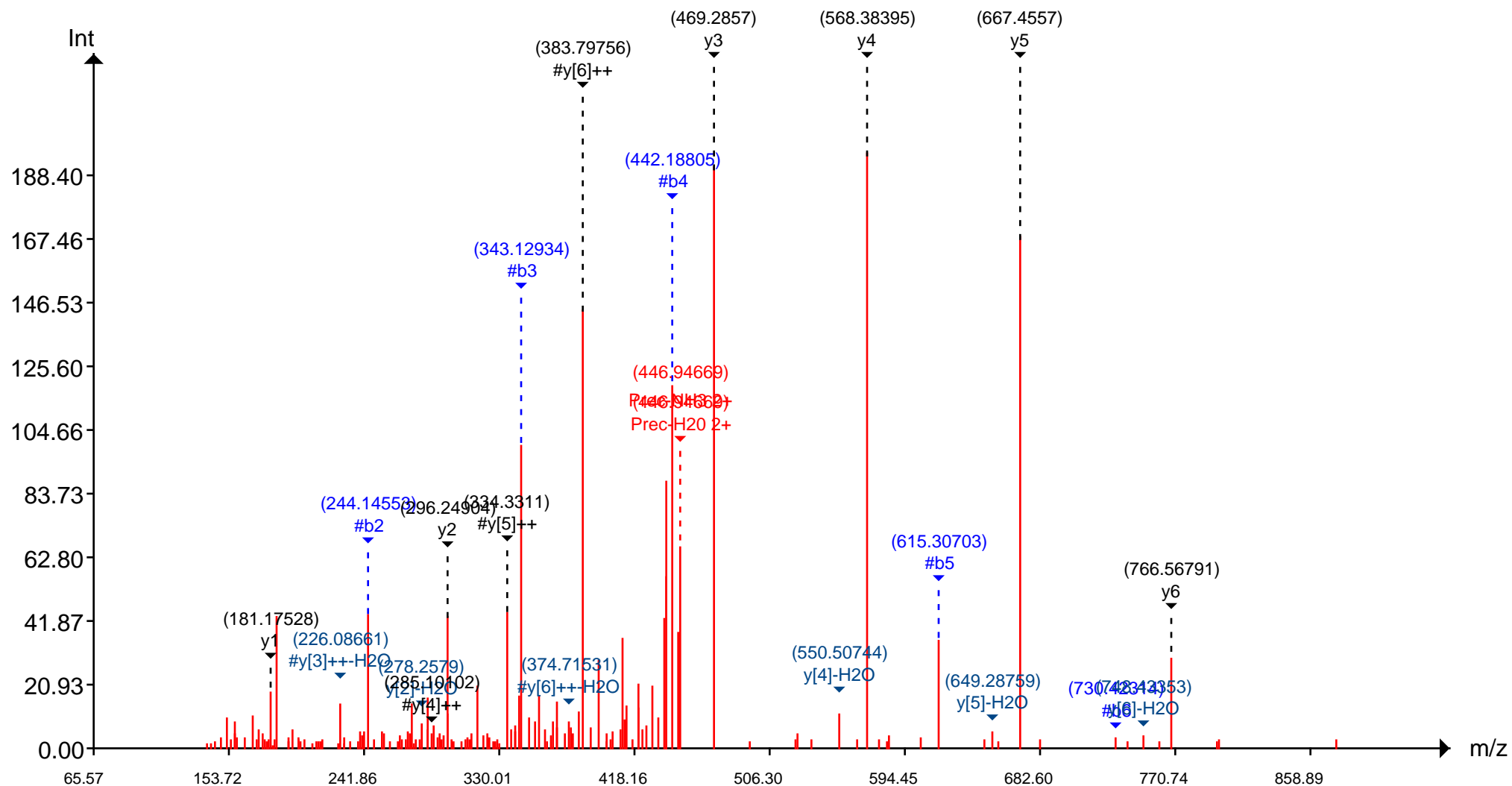
Info

Modified Sequence	AcD3-AC<Cmm*>LM<Mox*>GK<AcD3K*>GM<Mox*>K<AcD3K*>R<C13>-COOH	Ion score	50.35
Protein	Q61233	Confidence level	99.00%
Precursor m/z	662.84532	Identity Threshold	47.88
Charge	2+	Mass error	6.12E-4



Info

Modified Sequence	AcD3-GEGLGVK<AcD3K*>ETPQQK<AcD3K*>YQR<C13>-COOH	Ion score	80.19
Protein	Q99KJ8	Confidence level	99.00%
Precursor m/z	980.03369	Identity Threshold	47.45
Charge	2+	Mass error	0.001345



Info

Modified Sequence AcD3-VVVVK<AcD3K*>DR<C13>-COOH

Ion score 53.32

Protein P60824, O89086

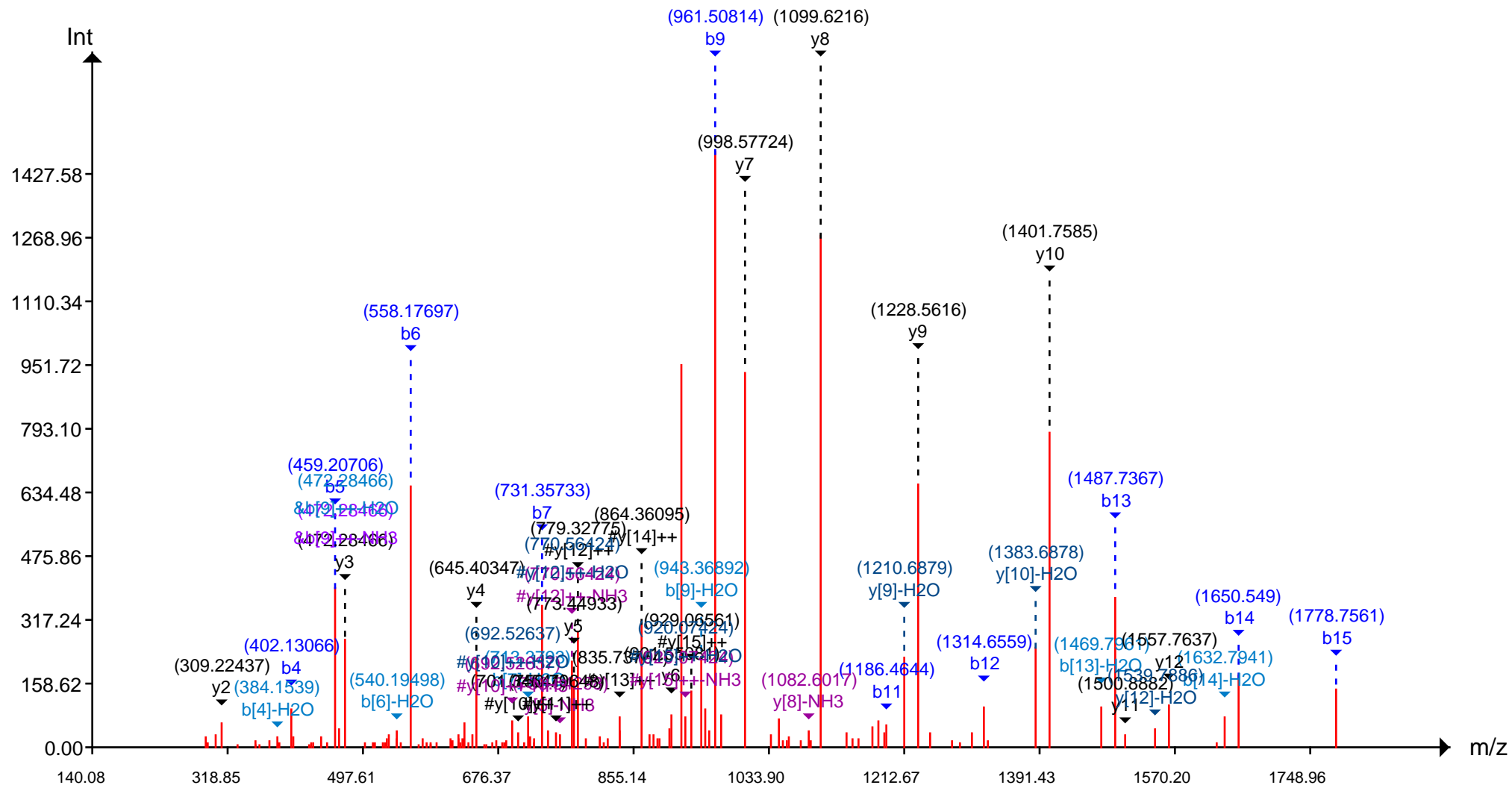
Confidence level 99.00%

Precursor m/z 455.79988

Identity Threshold 40.79

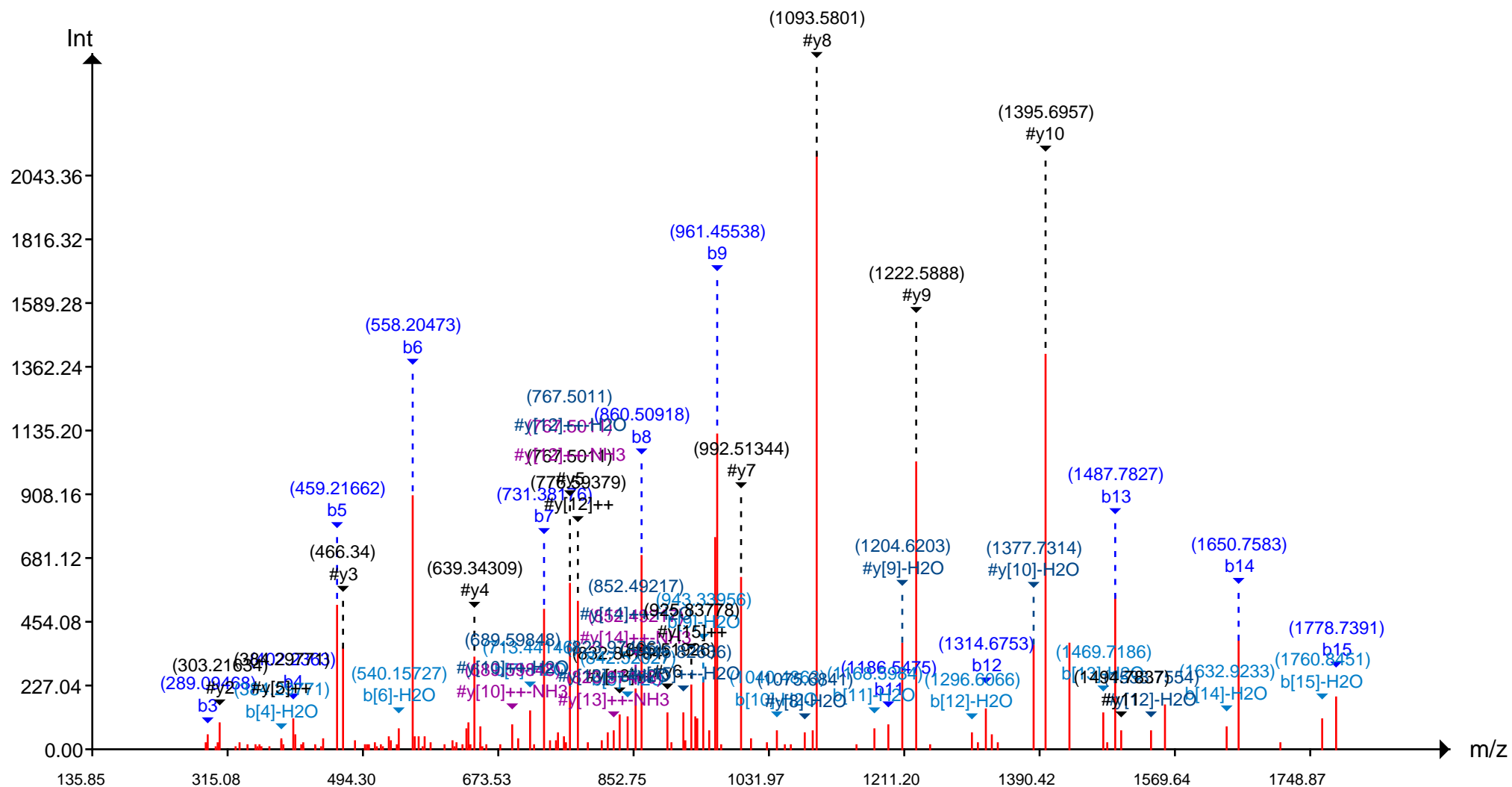
Charge 2+

Mass error -9.74E-4



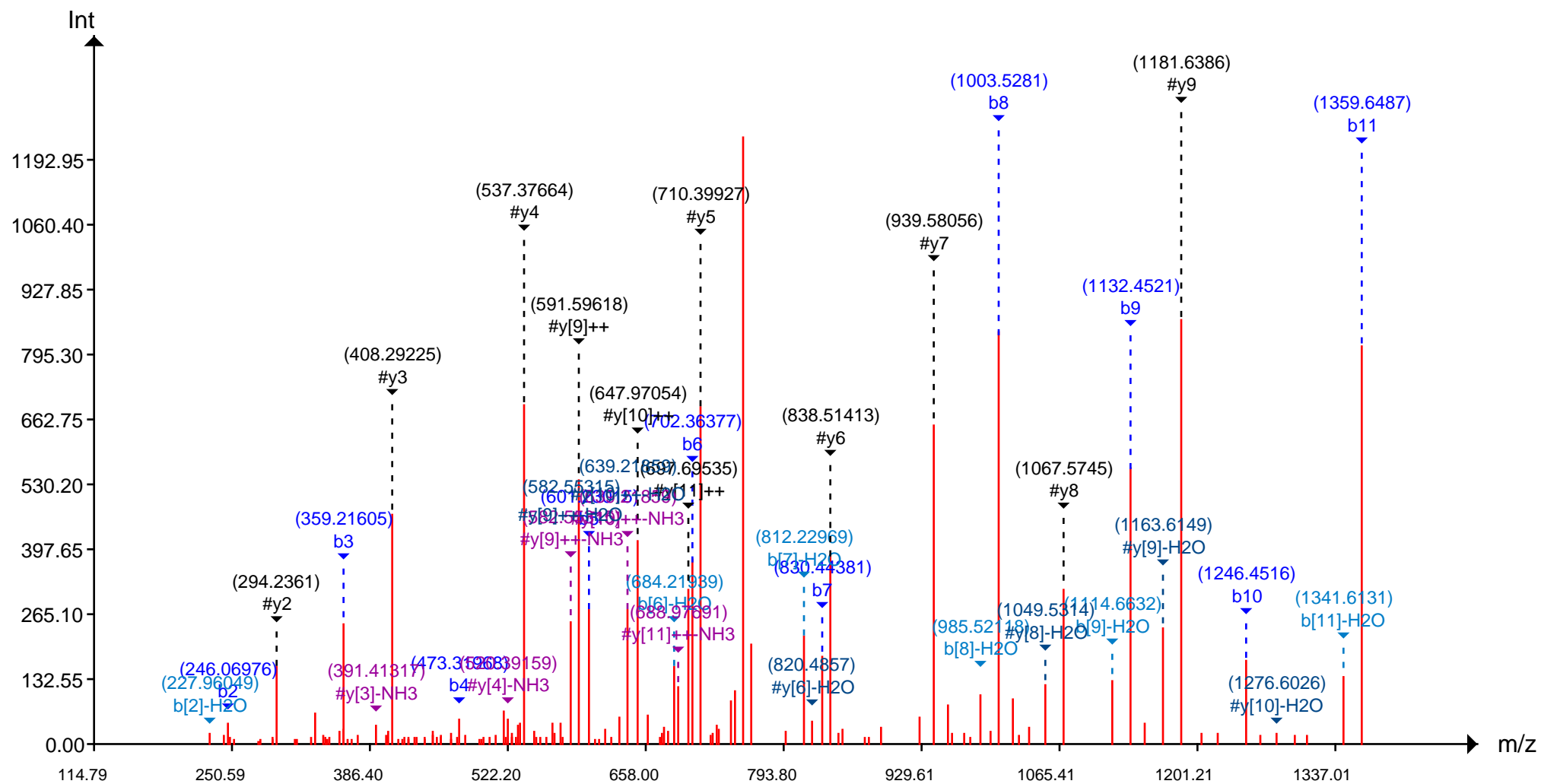
Info

Modified Sequence	AcD3-GEGLGVK<AcD3K*>ETPQQK<AcD3K*>YQR<C13>-COOH	Ion score	74.95
Protein	Q99KJ8	Confidence level	99.00%
Precursor m/z	980.03377	Identity Threshold	47.46
Charge	2+	Mass error	0.001505



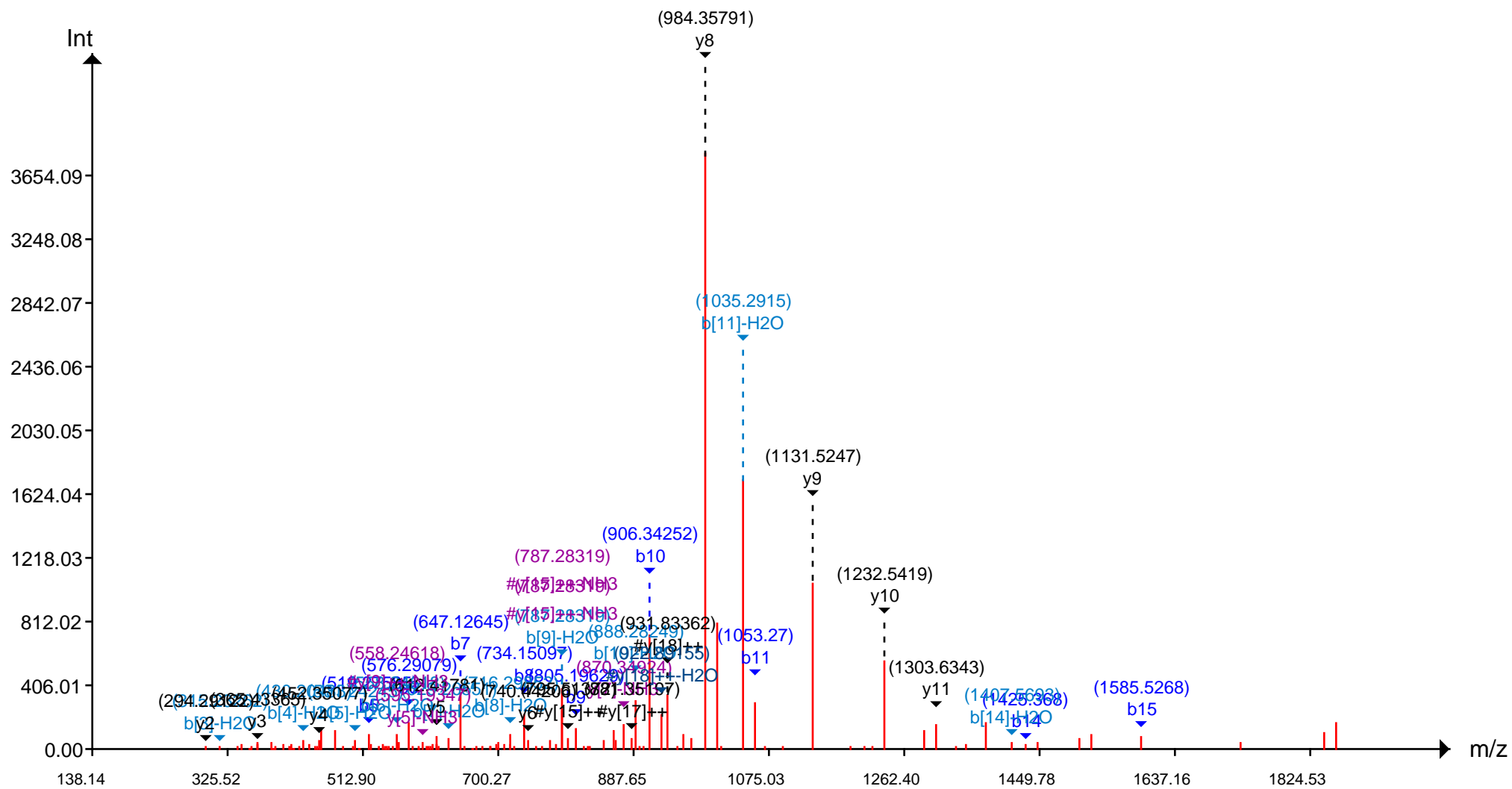
Info

Modified Sequence	AcD3-GEGLGVK<AcD3K*>ETPQQK<AcD3K*>YQR-COOH	Ion score	106.66
Protein	Q99KJ8	Confidence level	99.00%
Precursor m/z	977.0237	Identity Threshold	47.75
Charge	2+	Mass error	0.001506



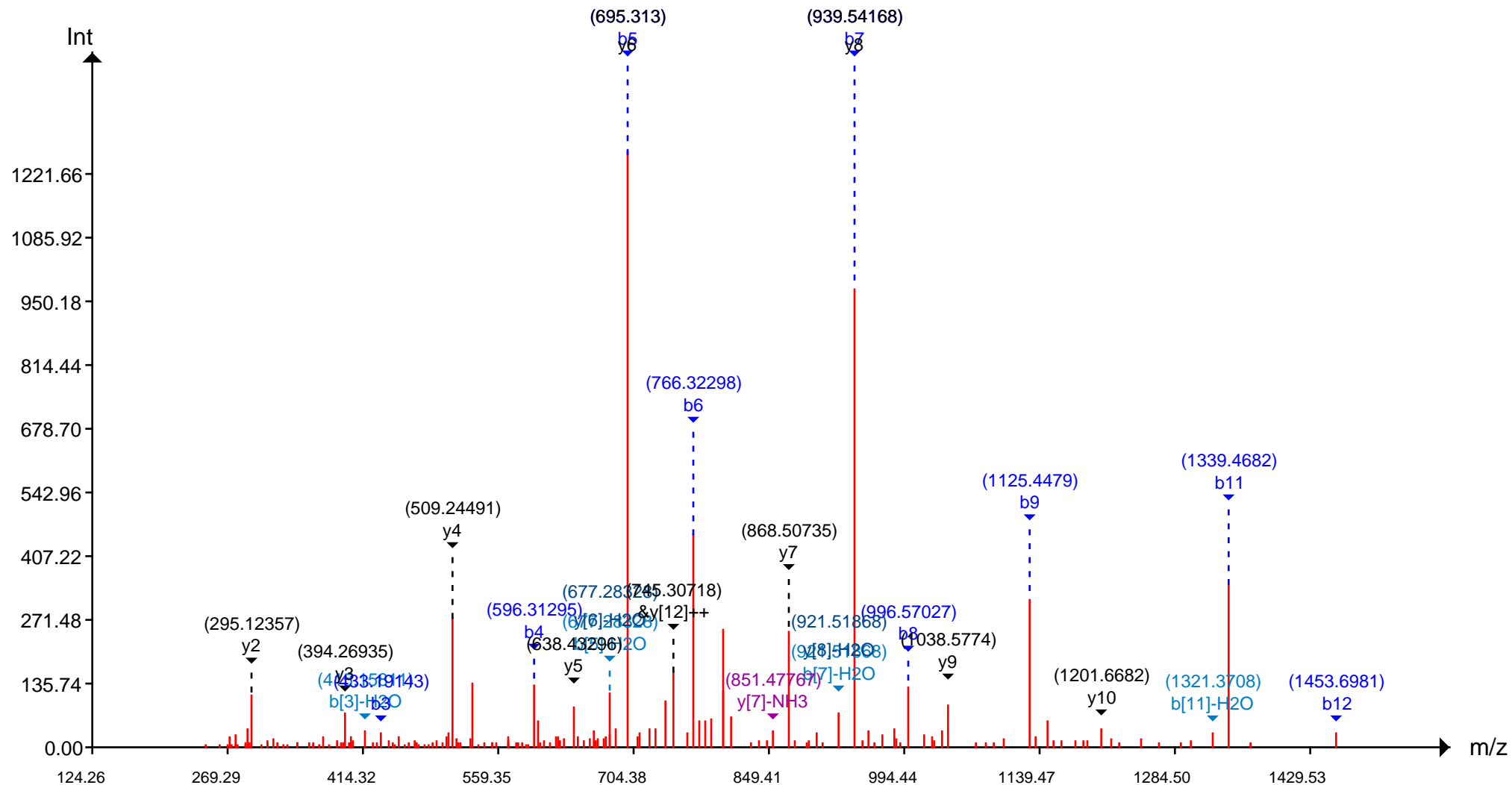
Info

Modified Sequence	AcD3-TVINQTQK<AcD3K*>ENLR<C13>-COOH	Ion score	93.7
Protein	Q6ZWW7	Confidence level	99.00%
Precursor m/z	770.43861	Identity Threshold	46.62
Charge	2+	Mass error	-3.84E-4



Info

Modified Sequence	AcD3-ETGDAGASATFPM<Mox*>QC<Cmm*>SALR<C13>-COOH	Ion score	57.11
Protein	P63242	Confidence level	99.00%
Precursor m/z	1018.9635	Identity Threshold	44.06
Charge	2+	Mass error	0.001041



Info

Modified Sequence AcD3-VK<AcD3K*>DYVAK<AcD3K*>GEDVNR<C13>-COOH

Ion score 83.21

Protein P62774

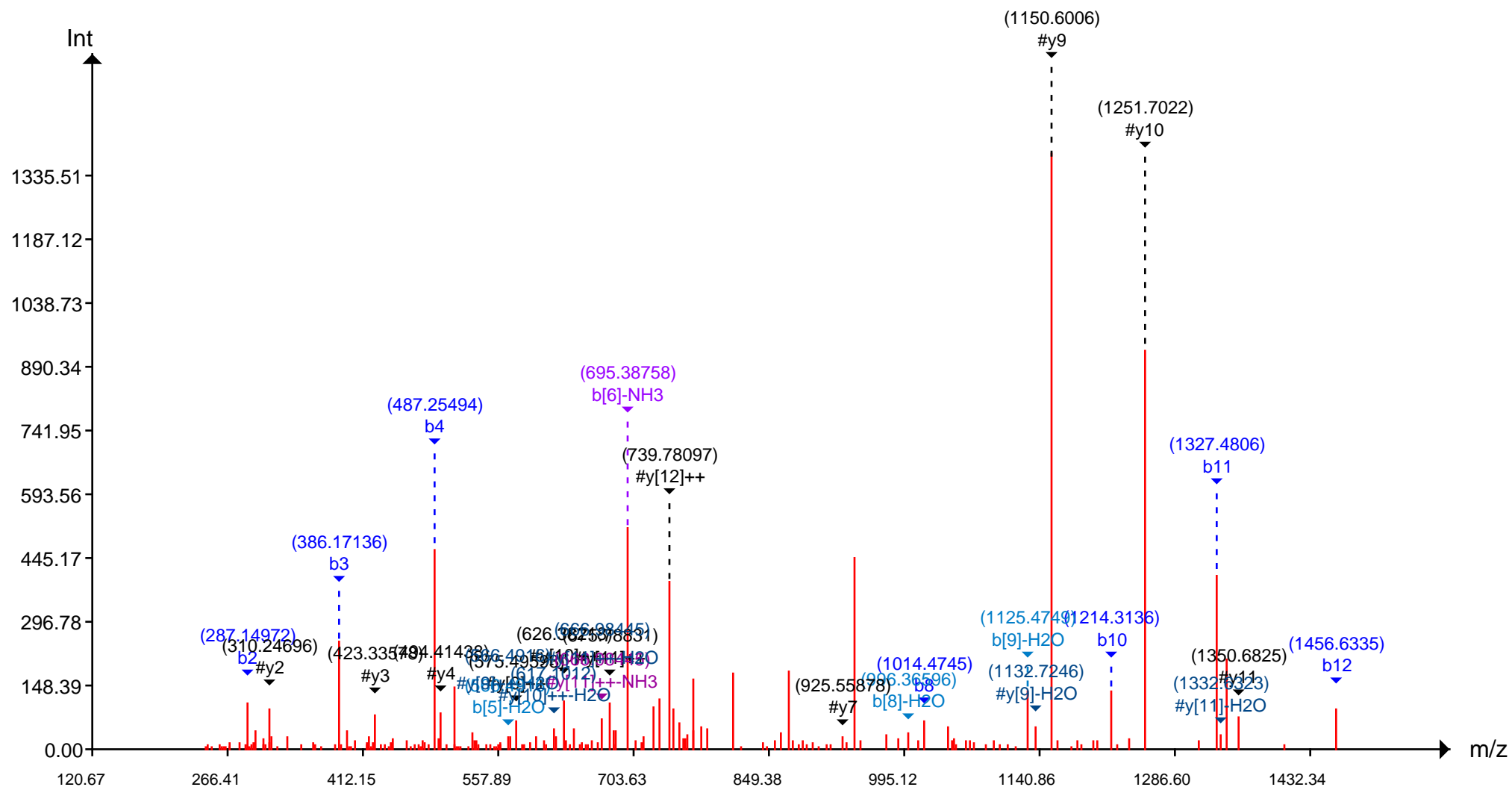
Confidence level 99.00%

Precursor m/z 817.44659

Identity Threshold 47.71

Charge 2+

Mass error 0.002148



Info

Modified Sequence AcD3-IQVTPQEK<AcD3K*>EAIER<C13>-COOH

Ion score 48.67

Protein P54726, P54728

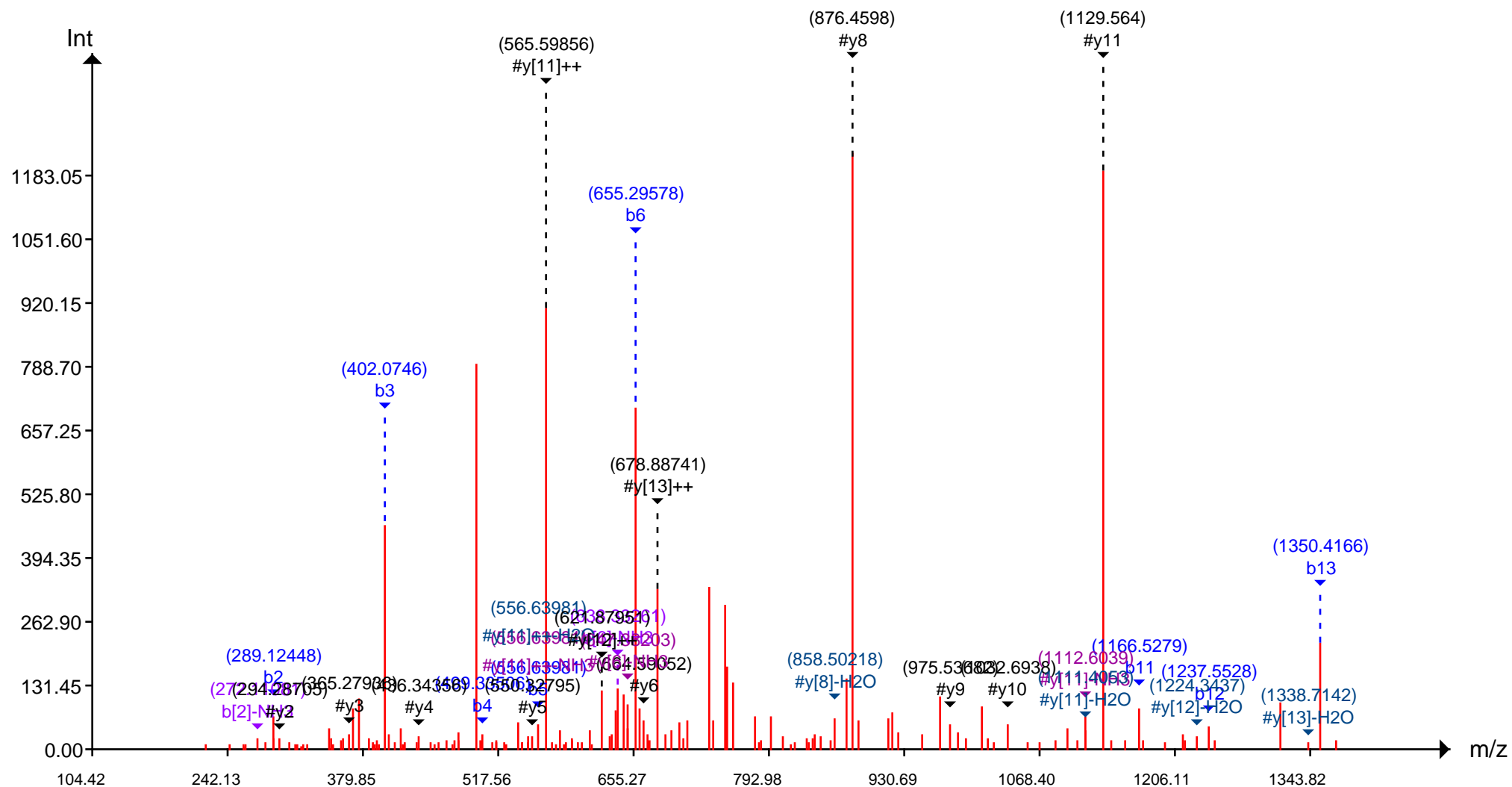
Confidence level 99.00%

Precursor m/z 818.9607

Identity Threshold 46.90

Charge 2+

Mass error 0.002276



Info

Modified Sequence AcD3-ENLPGVDPNNAAIR<C13>-COOH

Ion score 48.76

Protein O35226

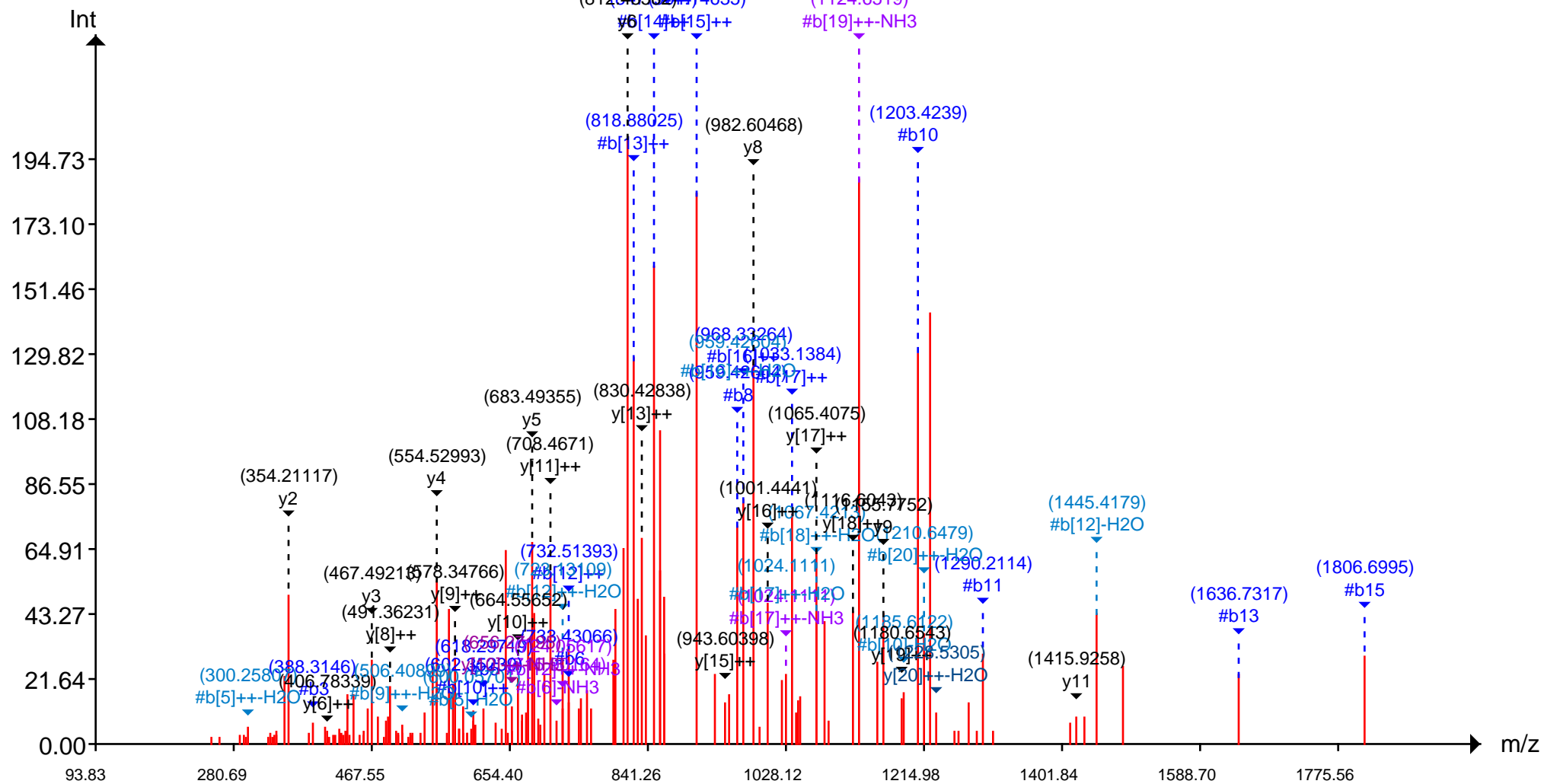
Confidence level 99.00%

Precursor m/z 765.90677

Identity Threshold 48.15

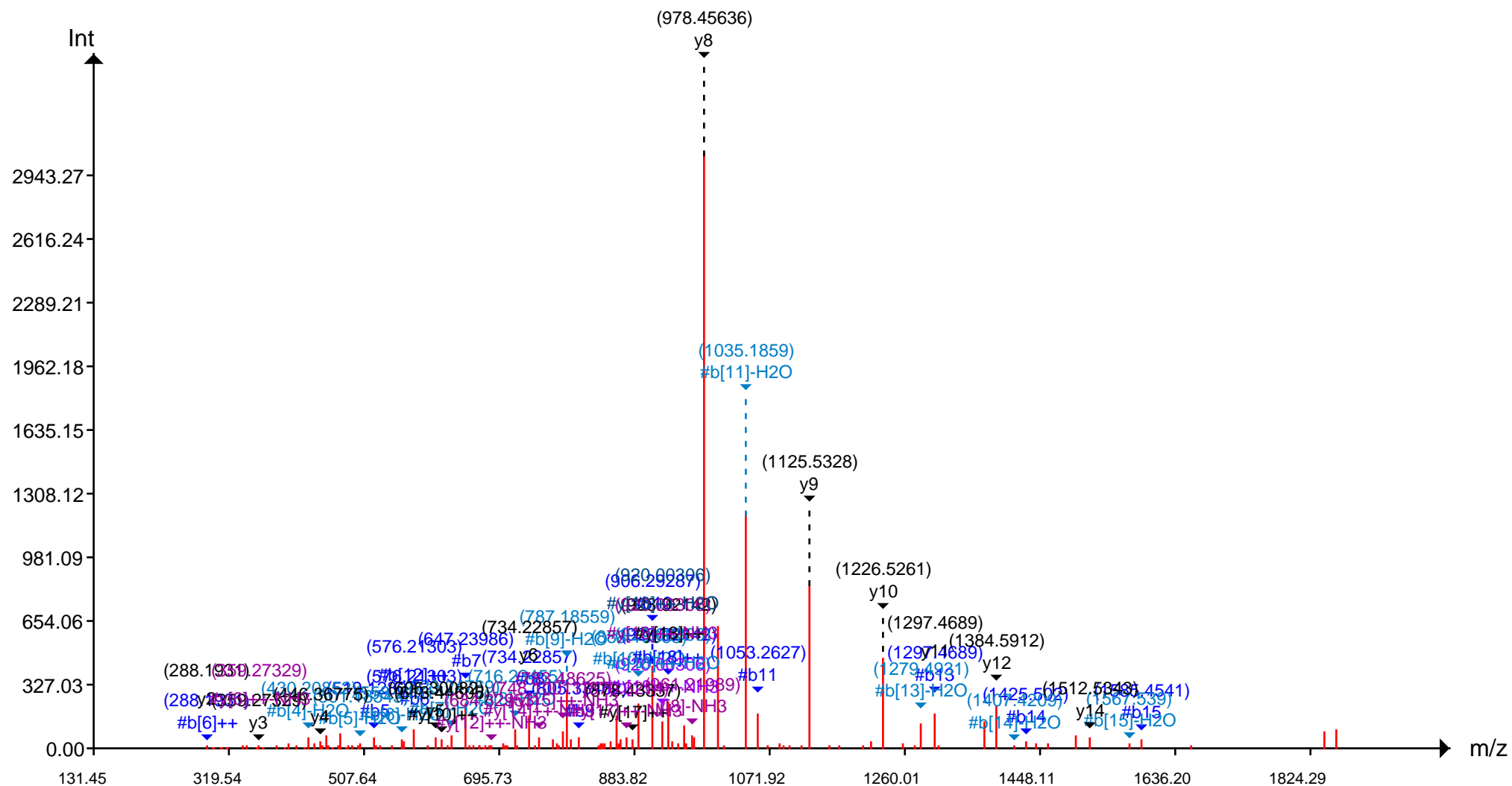
Charge 2+

Mass error 0.001732



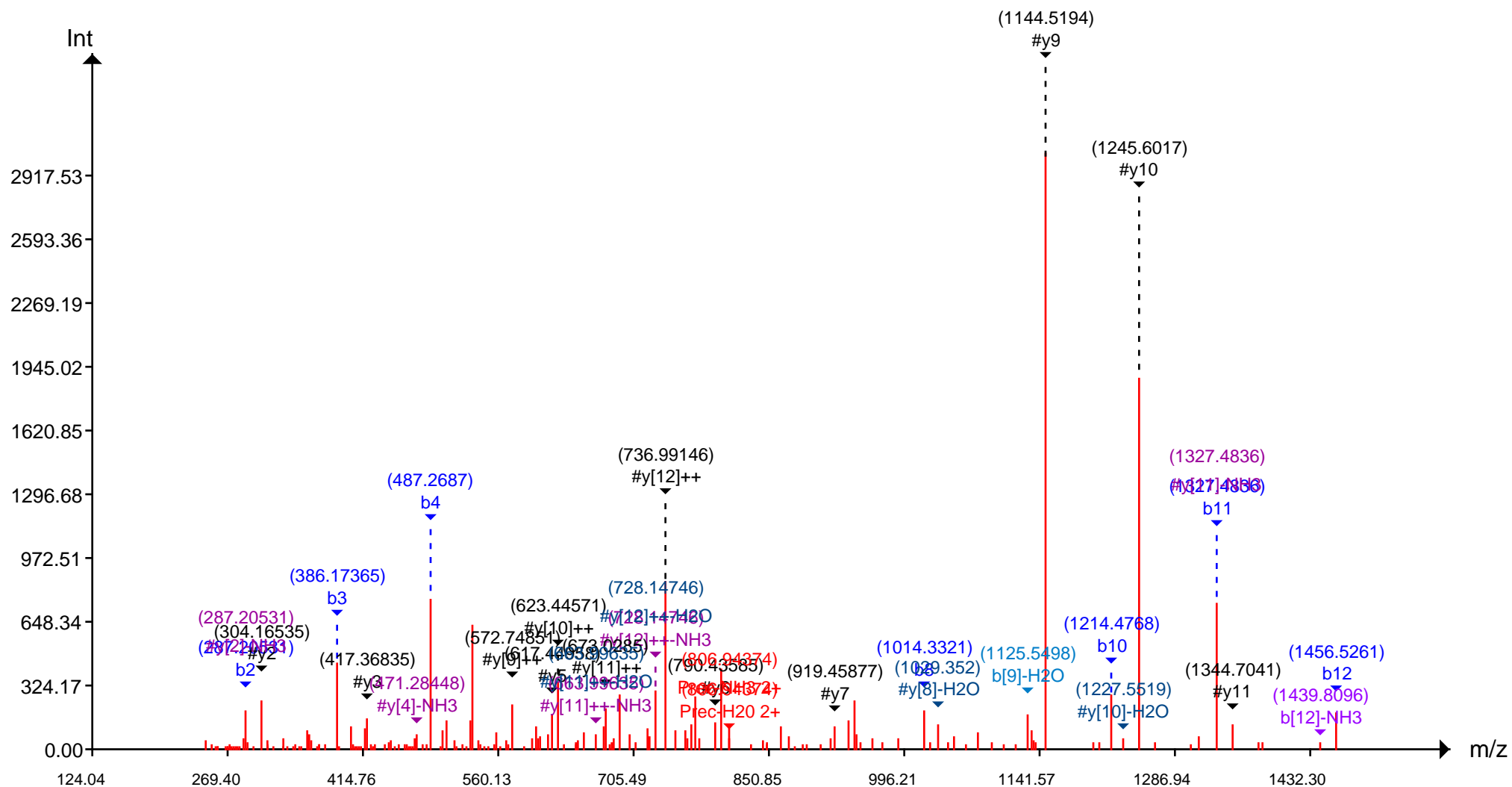
Info

Modified Sequen...	AcD3-VDQTEDILAK<AcD3K*>SK<AcD3K*>K<AcD3K*>GIEESLK<AcD3K*>R<C13>-COOH	Ion score	49.03
Prote...	Q61425	Confidence level	99.00%
Precursor m...	873.49906	Identity Threshold	43.99
Char...	3+	Mass error	0.001506



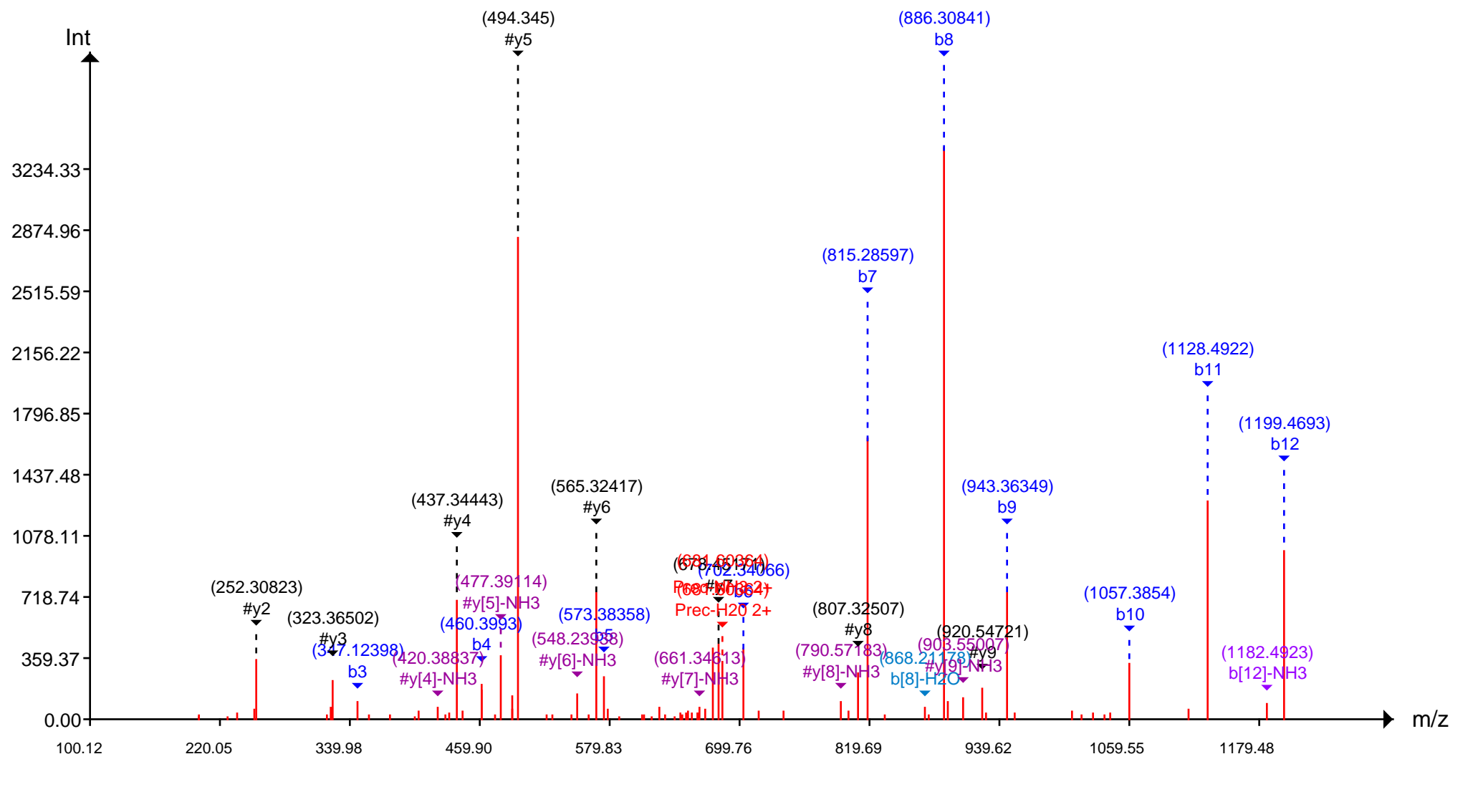
Info

Modified Sequence	AcD3-ETGDAGASATFPM<Mox*>QC<Cmm*>SALR-COOH	Ion score	69.95
Protein	P63242	Confidence level	99.00%
Precursor m/z	1015.9533	Identity Threshold	43.08
Charge	2+	Mass error	7.82E-4

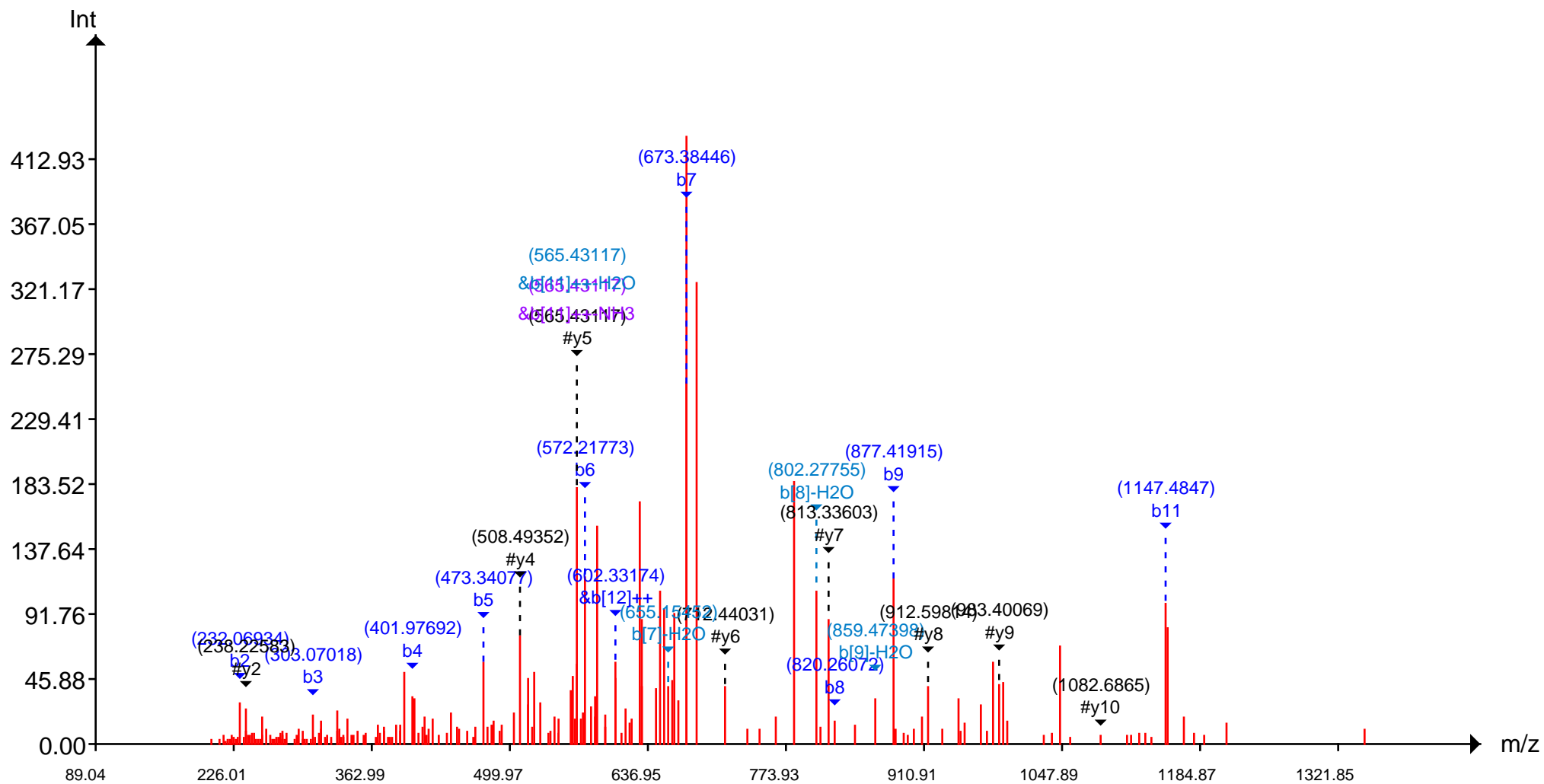


Info

Modified Sequence	AcD3-IQVTPQEK<AcD3K*>EAIER-COOH	Ion score	53.31
Protein	P54726, P54728	Confidence level	99.00%
Precursor m/z	815.95045	Identity Threshold	47.43
Charge	2+	Mass error	0.001918

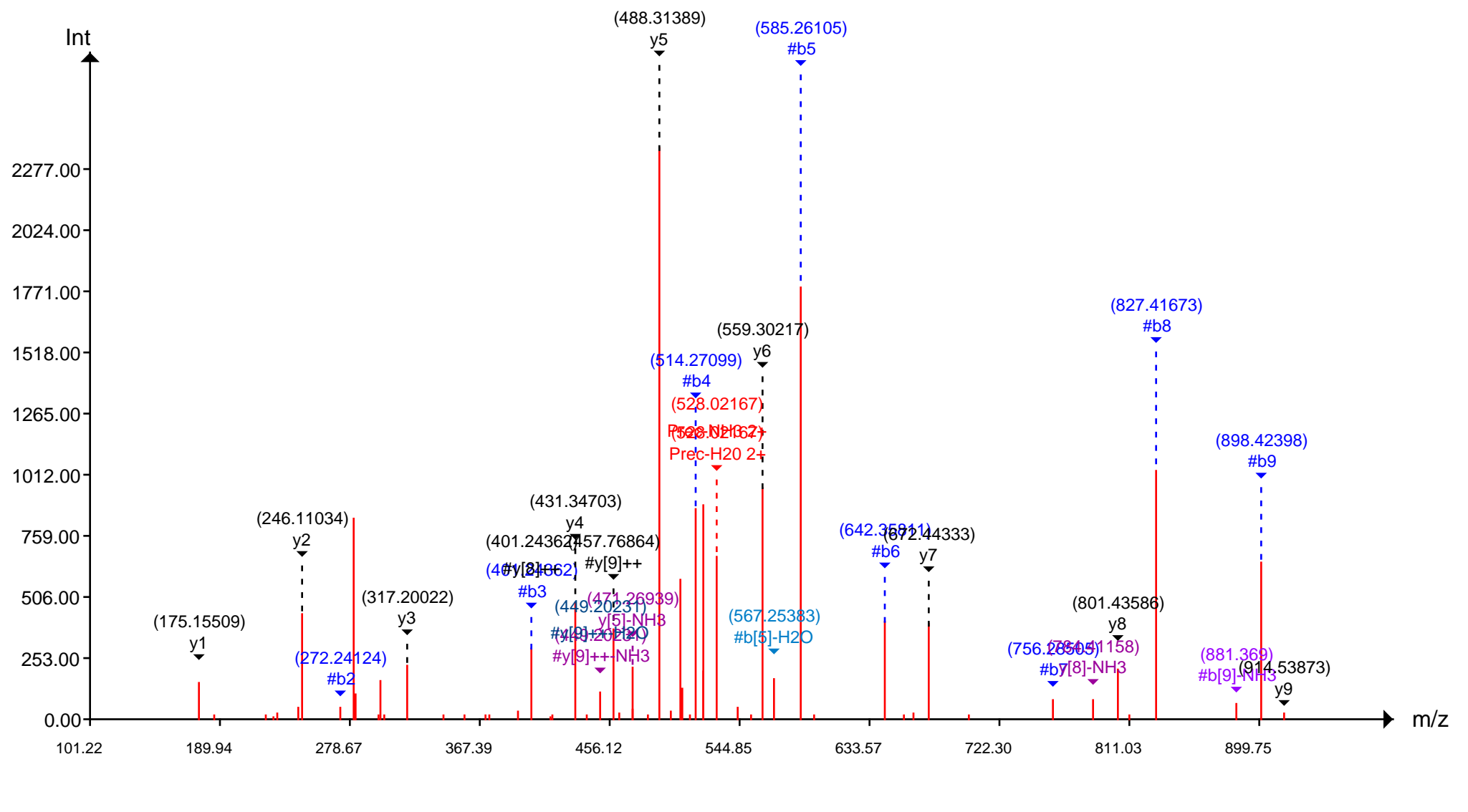


Info			
Modified ...	AcD3-TAEILELAGNAAR<C13>-COOH	Ion score	91
	Q8CGP5, Q8CGP6, Q8CGP7, P22752, Q6GSS7, Q64522, Q64523, Q8BFU2, Q8R1M2, P27661, Q9QZQ8	Confidence level	99
Prec...	690.38869	Identity Threshold	47
	2+	Mass error	0.

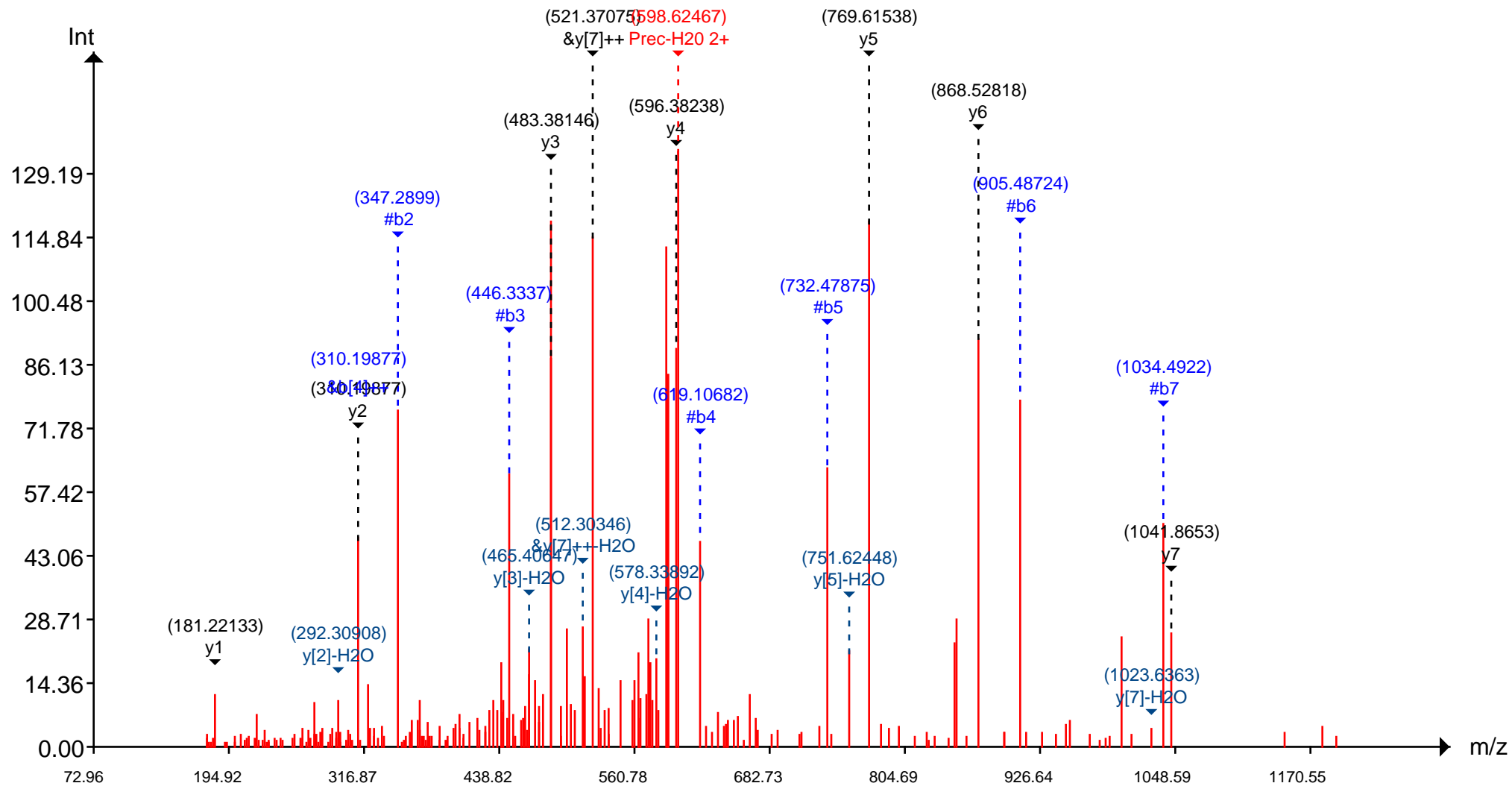


Info

Modified Sequence	AcD3-ADAVAVTM<Mox*>GPK<AcD3K*>GR<C13>-COOH	Ion score	49.69
Protein	P63038	Confidence level	99.00%
Precursor m/z	692.87654	Identity Threshold	47.66
Charge	2+	Mass error	-9.28E-4

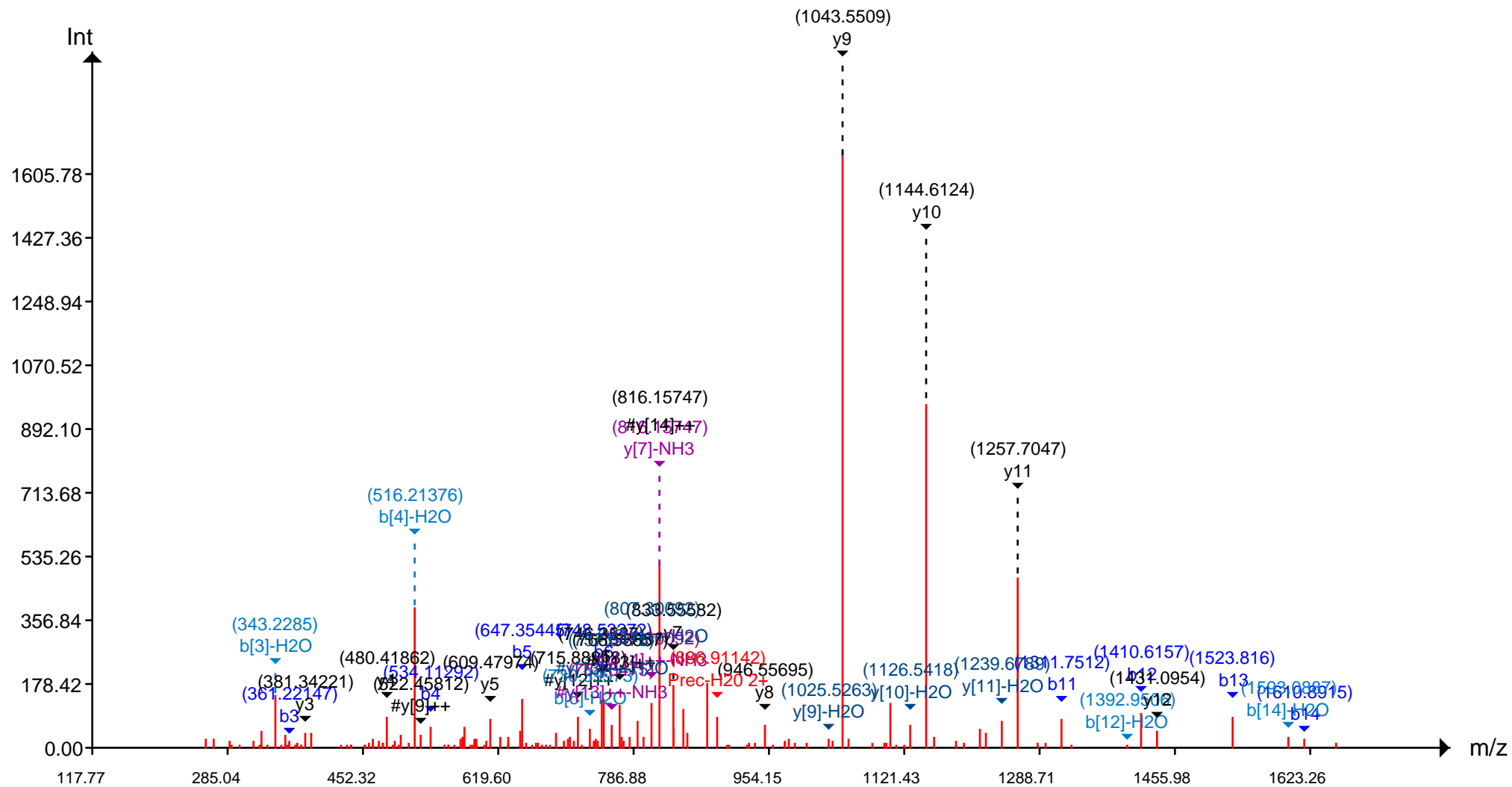


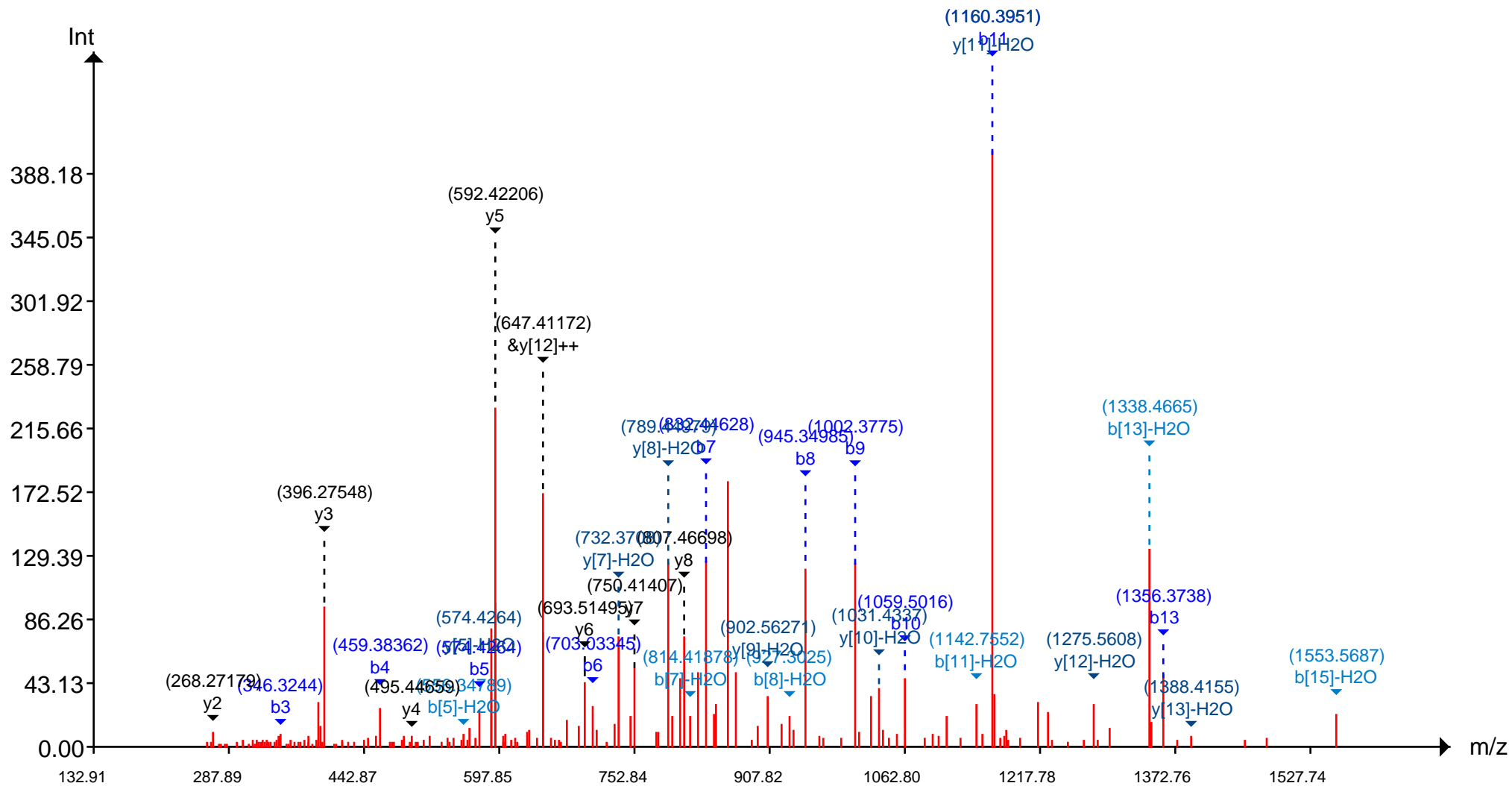
Info	
Modif...	AcD3-ILELAGNAAR-COOH
P...	536.81322
	2+
Ion score	
Confidence level	Q8CGP5, Q8CGP6, Q8CGP7, P22752, Q6GSS7, Q64522, Q64523, Q8BFU2, Q8R1M2, Q8CCK0, P27661, Q9QZQ8
Identity Threshold	
Mass error	



Info

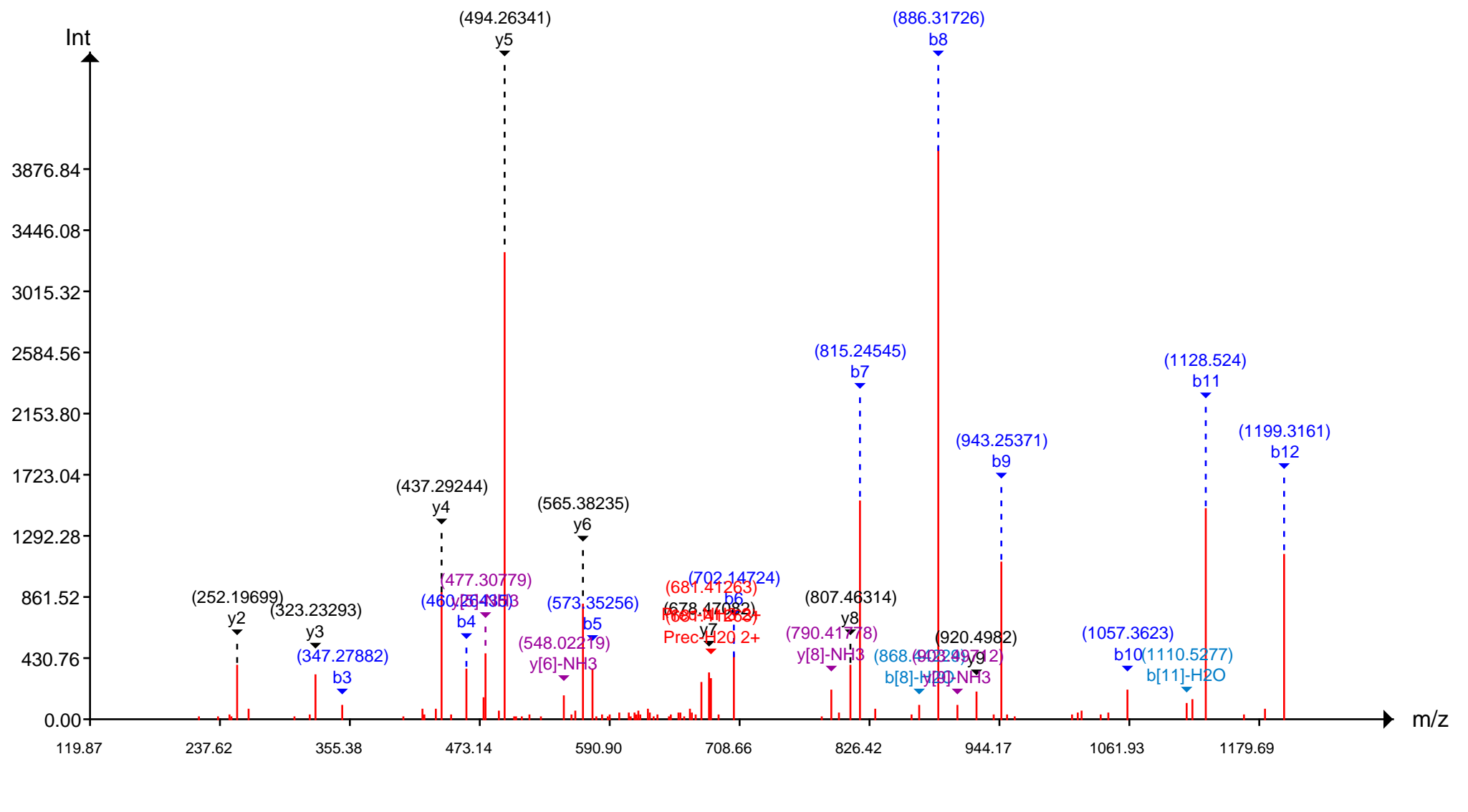
Modified Sequence	AcD3-QK<AcD3K*>VK<AcD3K*>LK<AcD3K*>ER<C13>-COOH	Ion score	41.19
Protein	P57784	Confidence level	99.00%
Precursor m/z	607.90128	Identity Threshold	37.45
Charge	2+	Mass error	9.0E-5



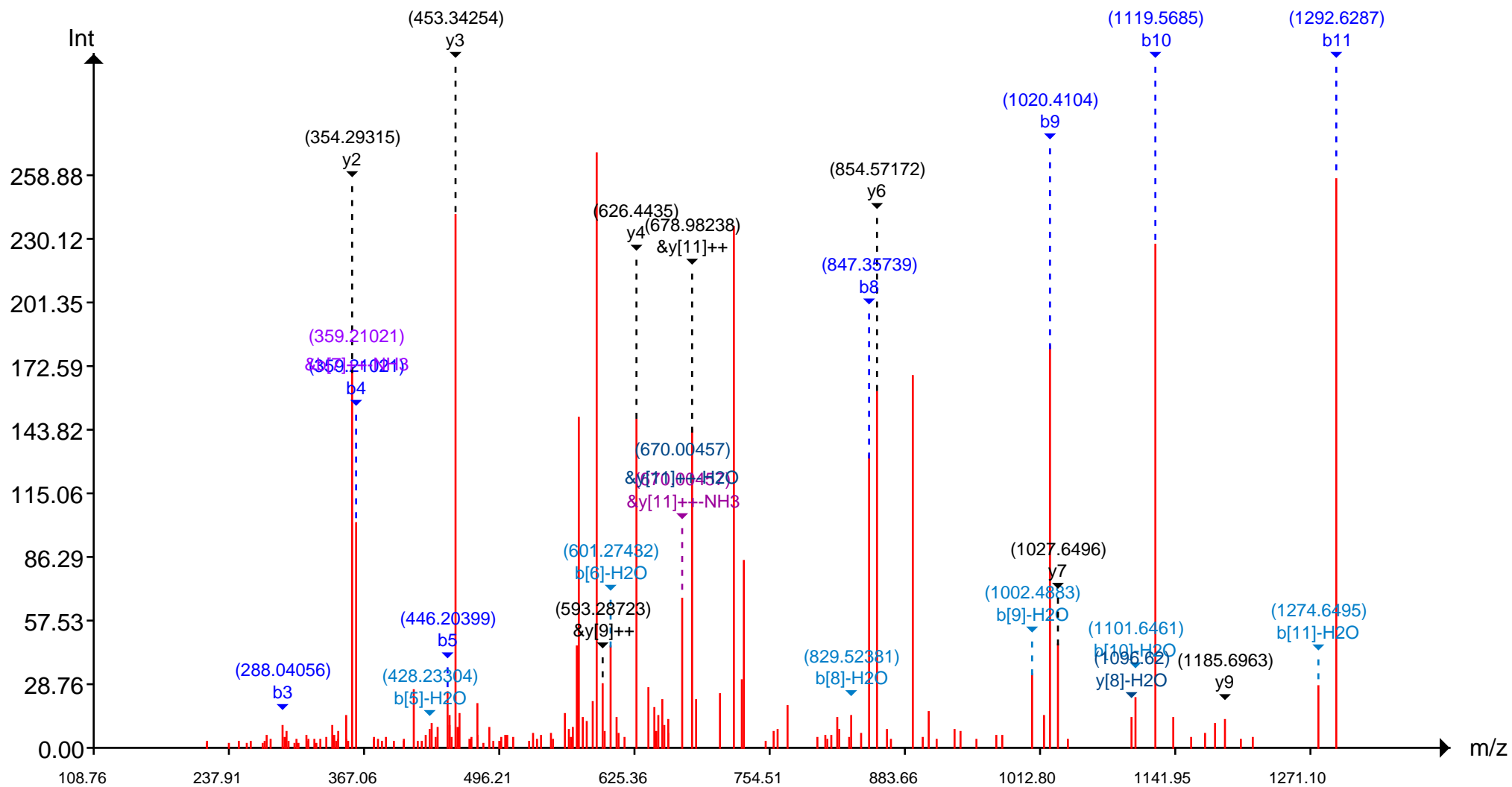


Info

Modified Sequence	AcD3-TAQLDEELGGTPVQSR<C13>-COOH	Ion score	54.72
Protein	P13020	Confidence level	99.00%
Precursor m/z	876.45184	Identity Threshold	48.08
Charge	2+	Mass error	0.001906

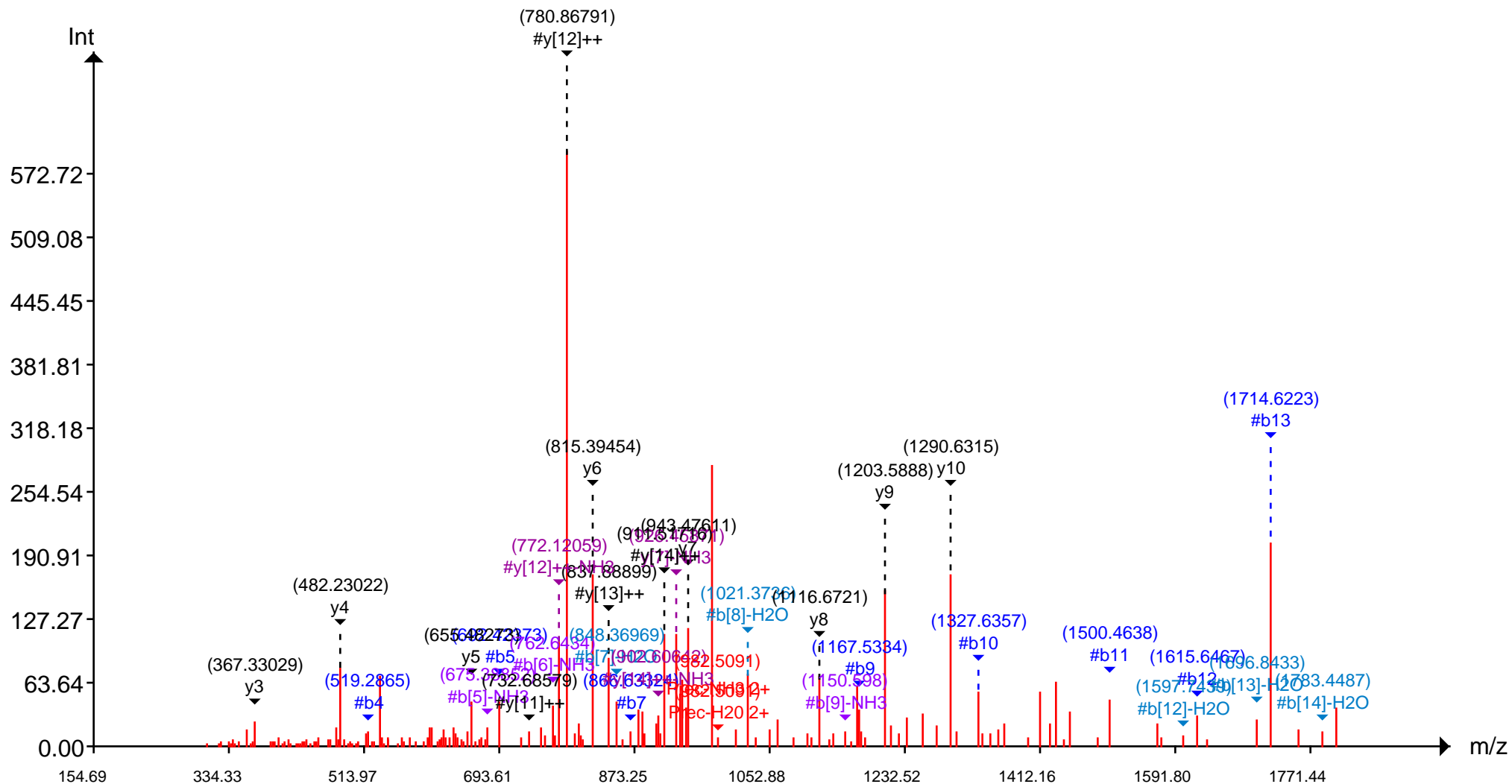


Info			
Modified ...	AcD3-TAEILELAGNAAR<C13>-COOH	Ion score	95
	Q8CGP5, Q8CGP6, Q8CGP7, P22752, Q6GSS7, Q64522, Q64523, Q8BFU2, Q8R1M2, P27661, Q9QZQ8	Confidence level	99
Prec...	690.38732	Identity Threshold	47
	2+	Mass error	0.



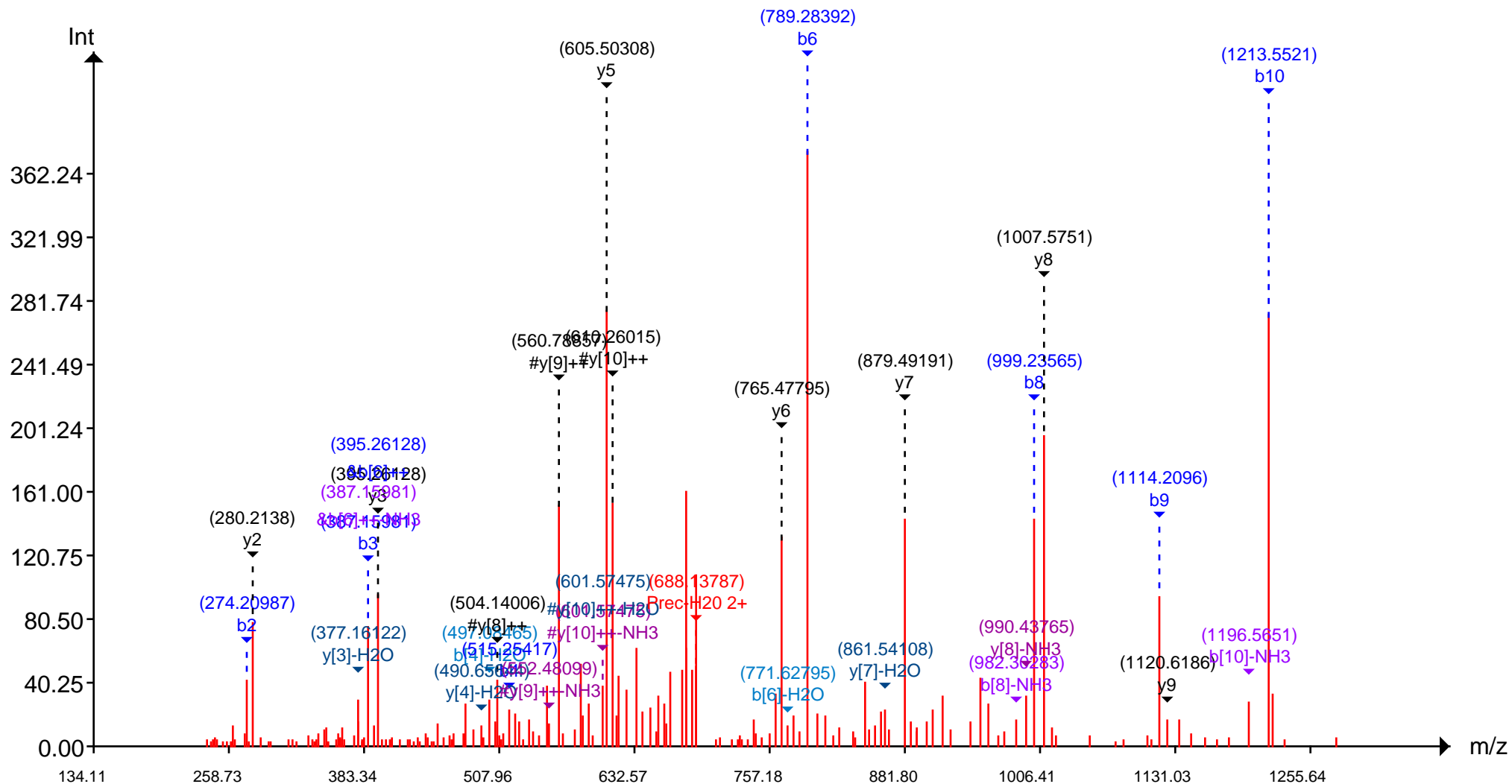
Info

Modified Sequence	AcD3-AGNASK<AcD3K*>DLK<AcD3K*>VK<AcD3K*>R<C13>-COOH	Ion score	42.43
Protein	Q3THW5, P0C0S6	Confidence level	99.00%
Precursor m/z	736.95032	Identity Threshold	42.37
Charge	2+	Mass error	0.00175



Info

Modified Sequence	AcD3-DFNPK<AcD3K*>SSK<AcD3K*>QC<Cmm*>K<AcD3K*>DVSR<C13>-COOH	Ion score	48.47
Protein	Q6IRU5	Confidence level	99.00%
Precursor m/z	991.51284	Identity Threshold	48.07
Charge	2+	Mass error	0.005116



Info

Modified Sequence AcD3-EVIQNC<Cmm*>PIDVR<C13>-COOH

Ion score 56.98

Protein Q99JY9

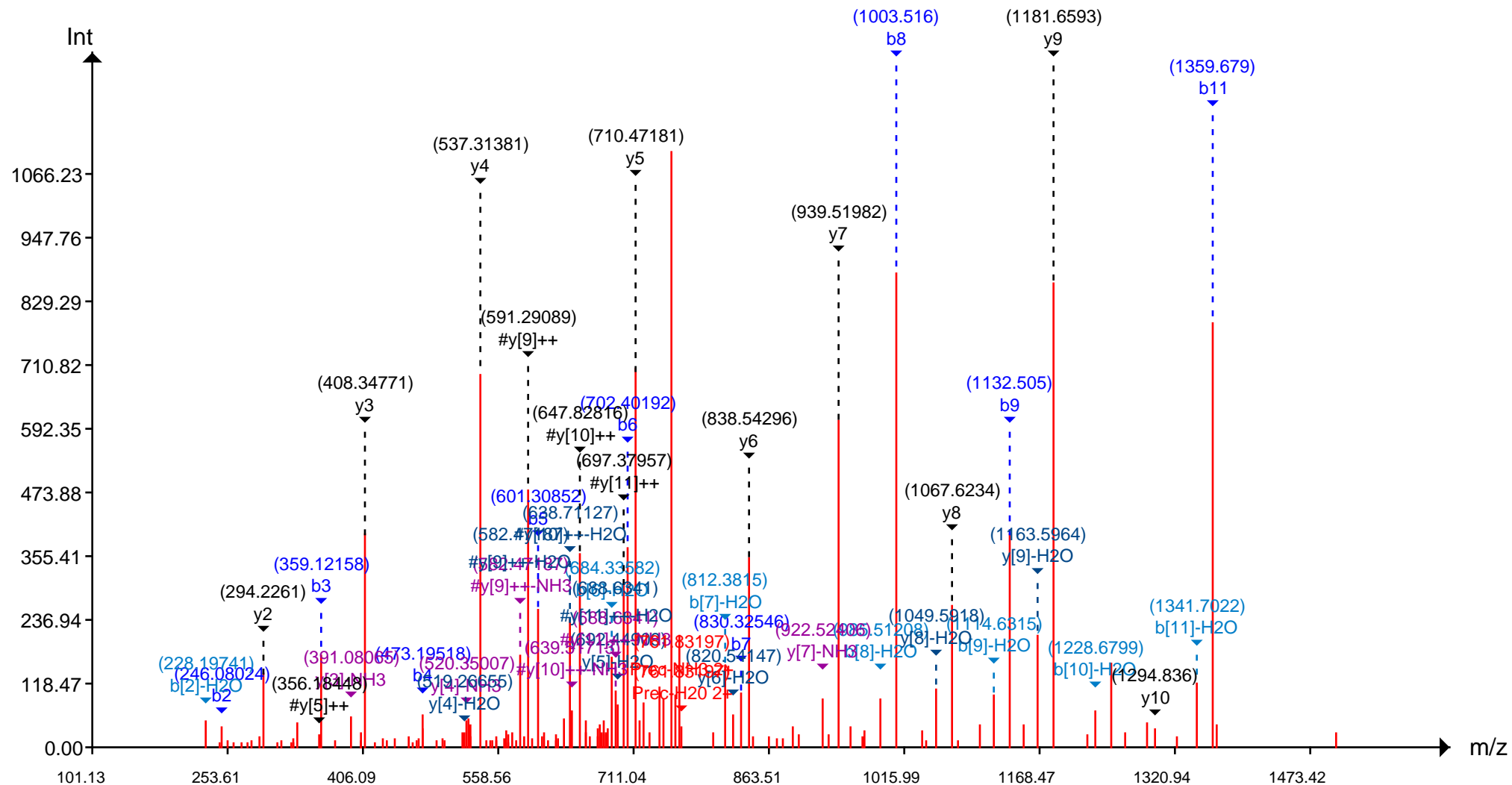
Confidence level 99.00%

Precursor m/z 697.36812

Identity Threshold 48.13

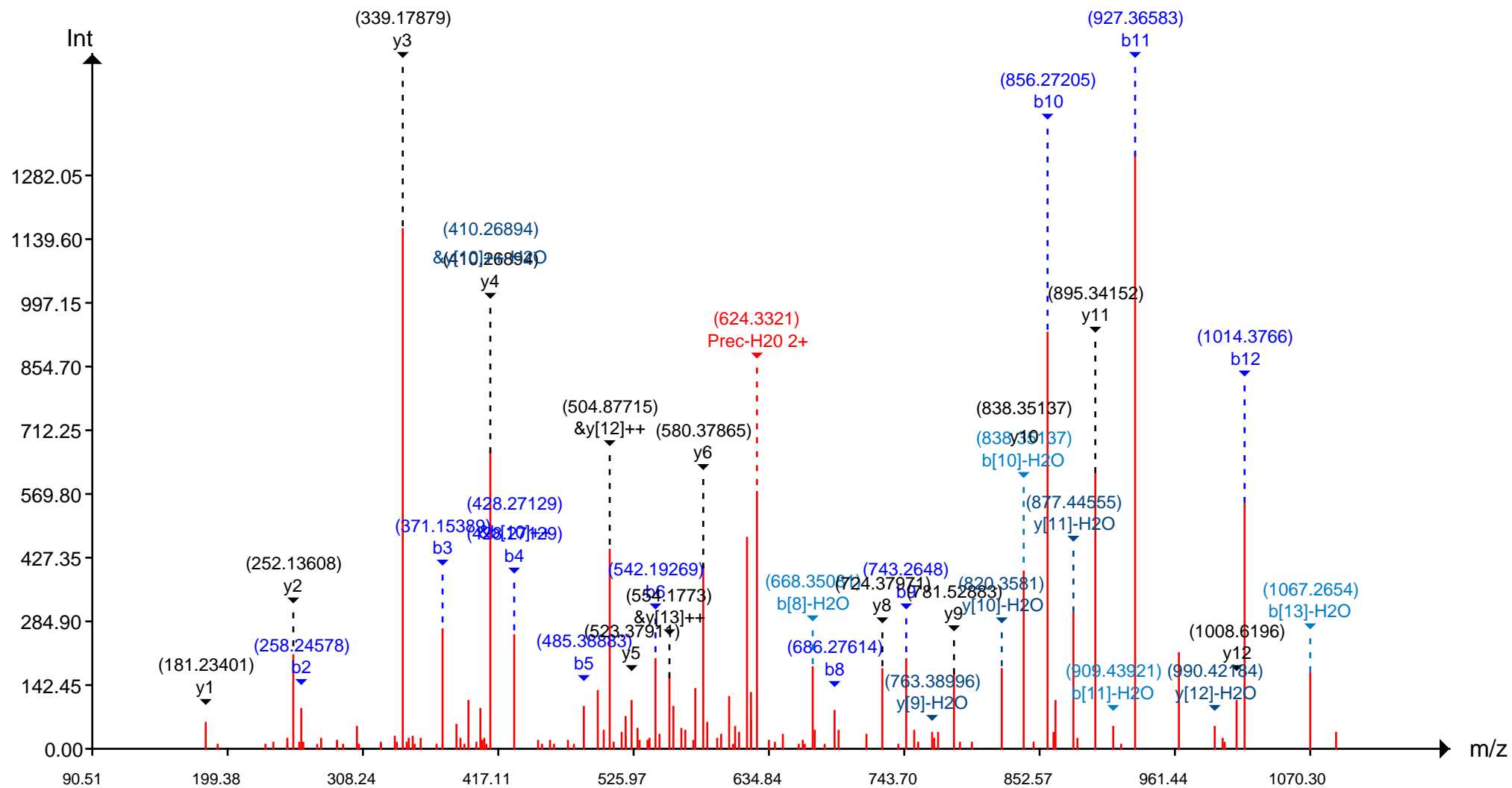
Charge 2+

Mass error 0.001076



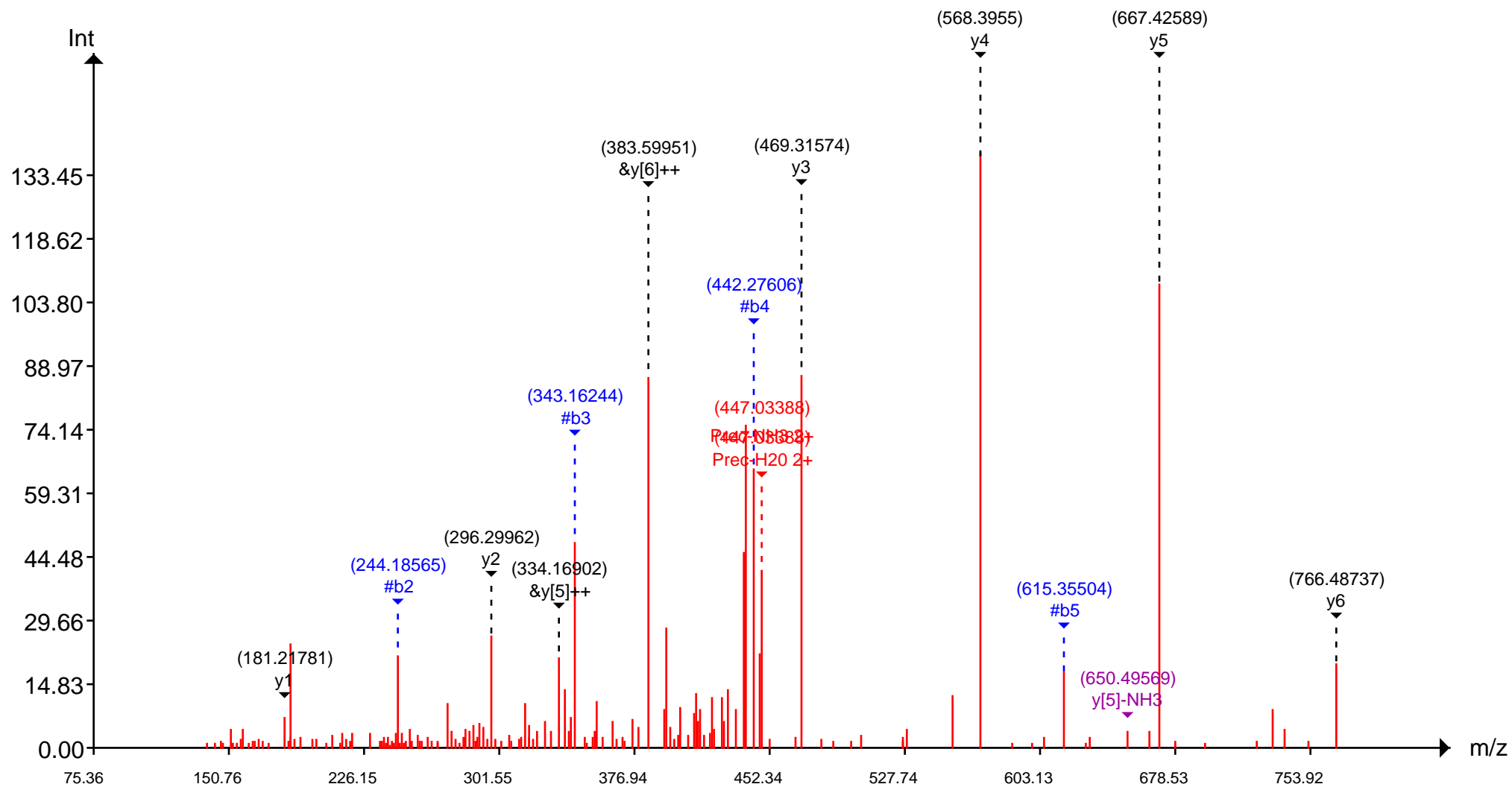
Info

Modified Sequence	AcD3-TVINQTQK<AcD3K*>ENLR<C13>-COOH	Ion score	80.07
Protein	Q6ZWW7	Confidence level	99.00%
Precursor m/z	770.43914	Identity Threshold	46.47
Charge	2+	Mass error	6.76E-4



Info

Modified Sequence	AcD3-LVIGGGSGGLASAR<C13>-COOH	Ion score	89.35
Protein	P47791	Confidence level	99.00%
Precursor m/z	633.37101	Identity Threshold	45.97
Charge	2+	Mass error	5.1E-5



Info

Modified Sequence AcD3-VVVVK<AcD3K*>DR<C13>-COOH

Ion score 50.61

Protein P60824, O89086

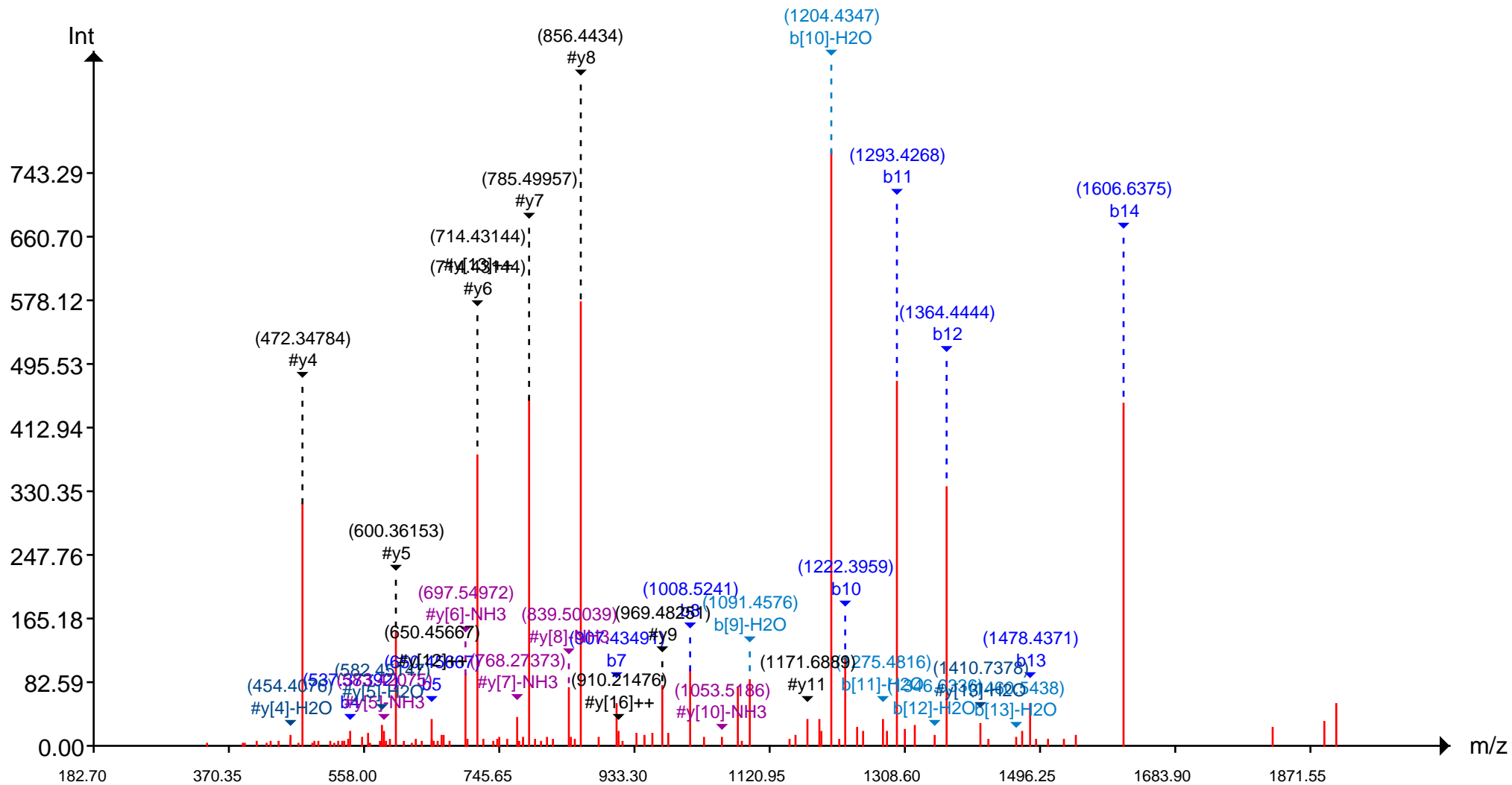
Confidence level 99.00%

Precursor m/z 455.80034

Identity Threshold 40.89

Charge 2+

Mass error -5.4E-5



Info

Modified Sequence AcD3-TINYLQETTLAANQPLSR-COOH

Ion score 59.83

Protein O09159

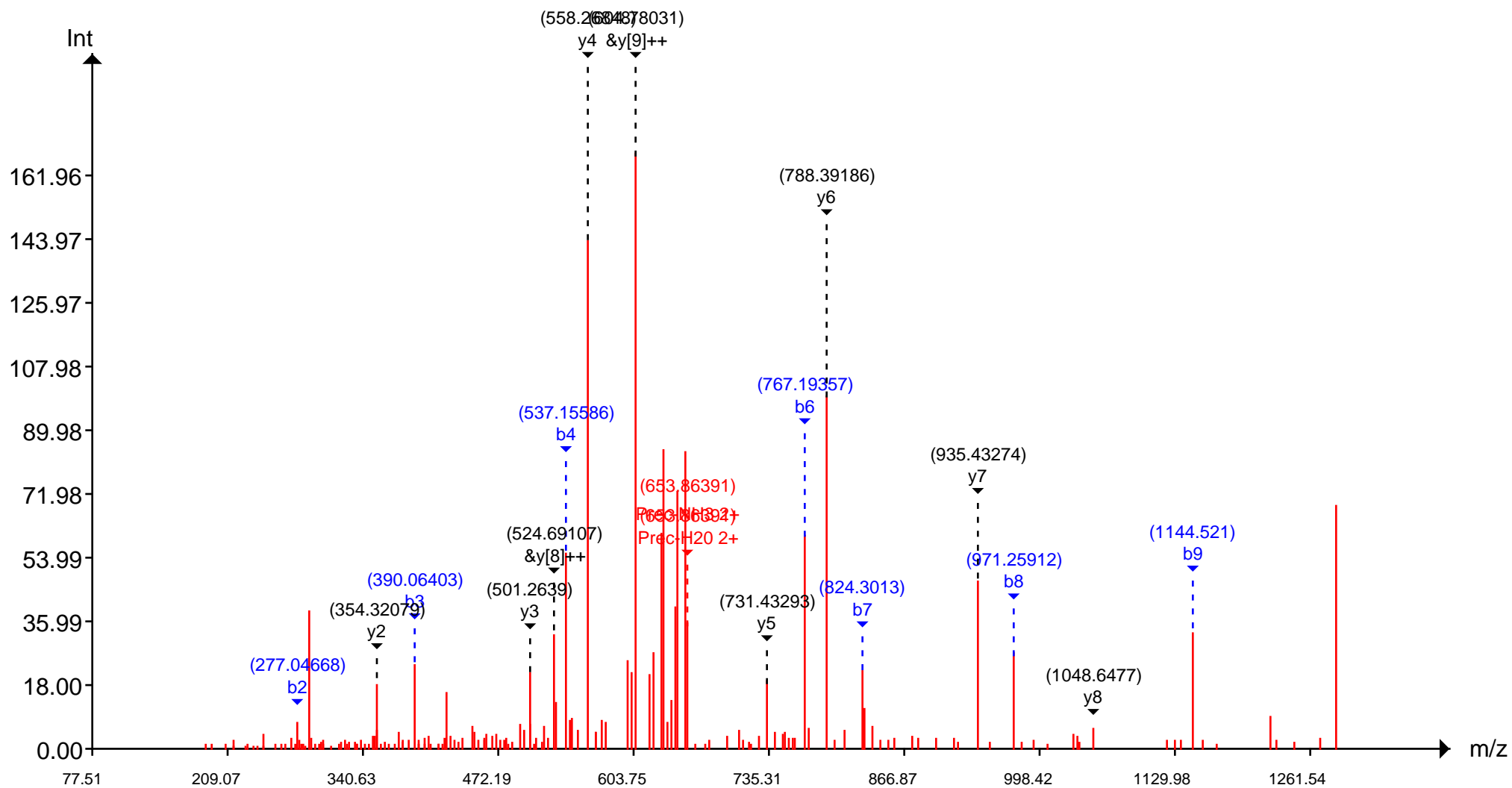
Confidence level 99.00%

Precursor m/z 1039.5525

Identity Threshold 47.89

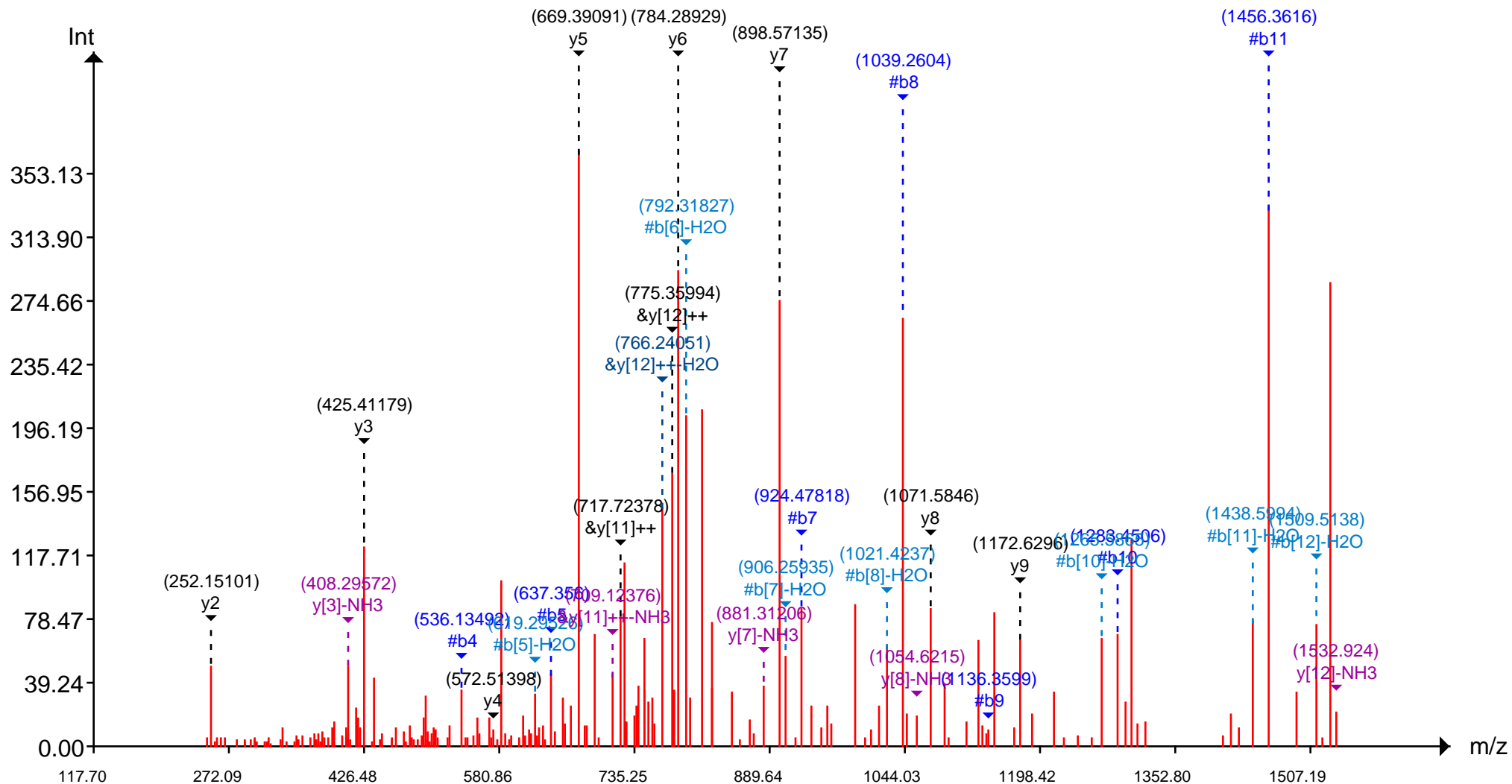
Charge 2+

Mass error 0.002176



Info

Modified Sequence	AcD3-AC<Cmm*>LM<Mox*>GK<AcD3K*>GM<Mox*>K<AcD3K*>R<C13>-COOH	Ion score	48.11
Protein	Q61233	Confidence level	99.00%
Precursor m/z	662.84511	Identity Threshold	47.96
Charge	2+	Mass error	1.92E-4



Info

Modified Sequence AcD3-LDDM<Mox*>TK<AcD3K*>NDPFK<AcD3K*>AR<C13>-COOH

Ion score 56.84

Protein Q62523

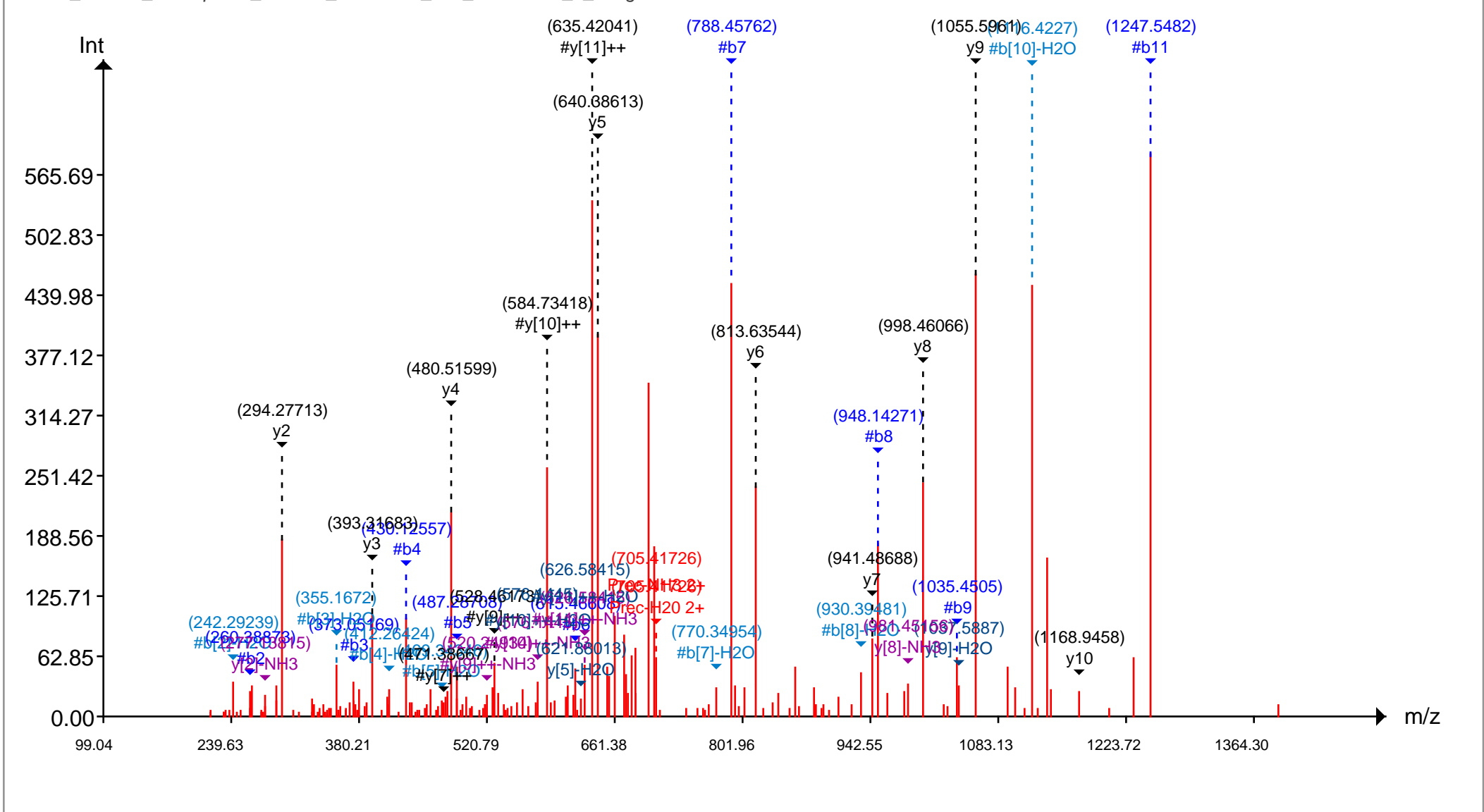
Confidence level 99.00%

Precursor m/z 854.43764

Identity Threshold 47.93

Charge 2+

Mass error 0.001612



Info

Modified Sequence AcD3-LTLGGQK<AcD3K*>C<Cmm*>SVIR<C13>-COOH

Ion score 64.27

Protein P62962

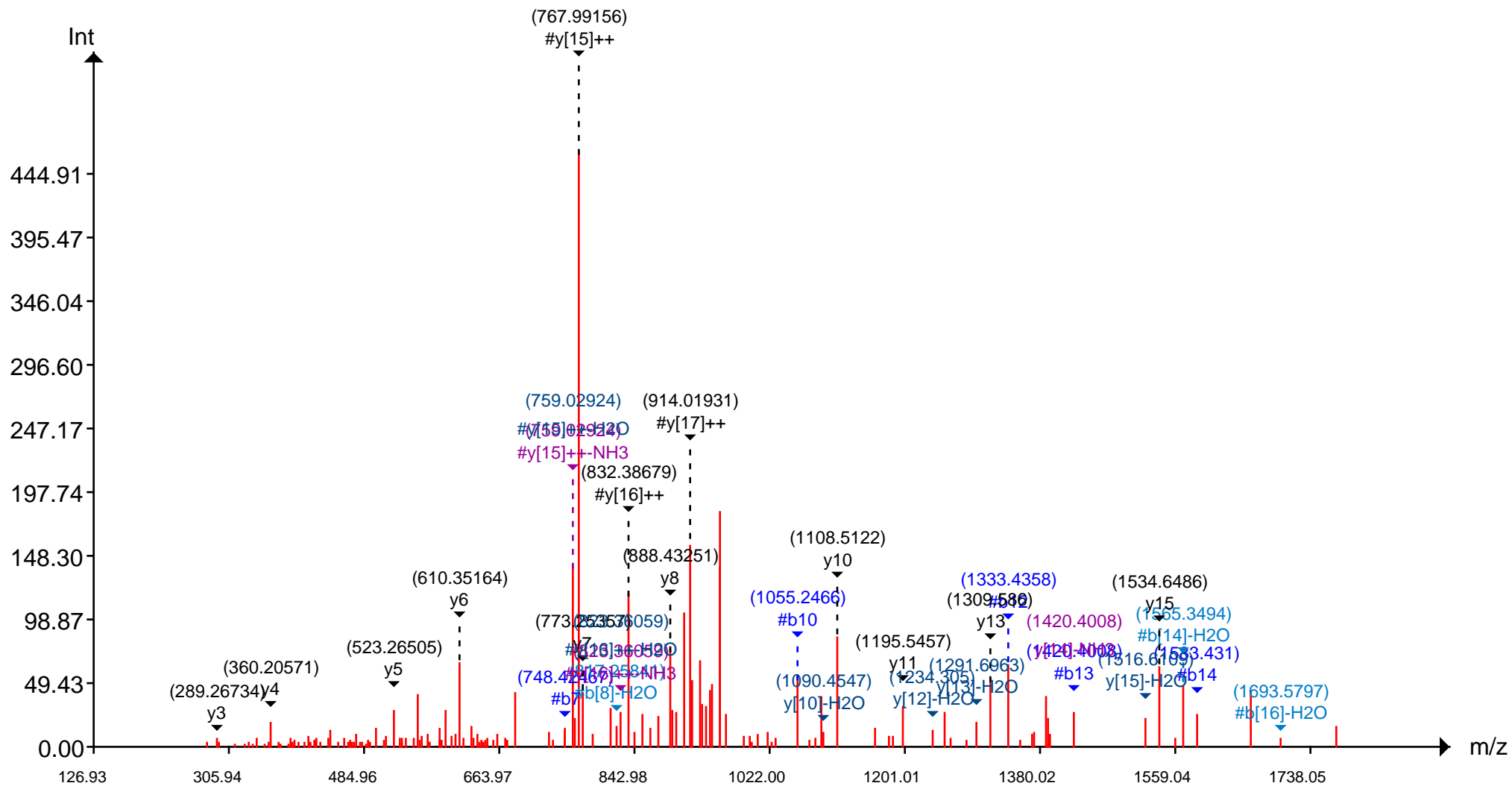
Confidence level 99.00%

Precursor m/z 714.41715

Identity Threshold 45.44

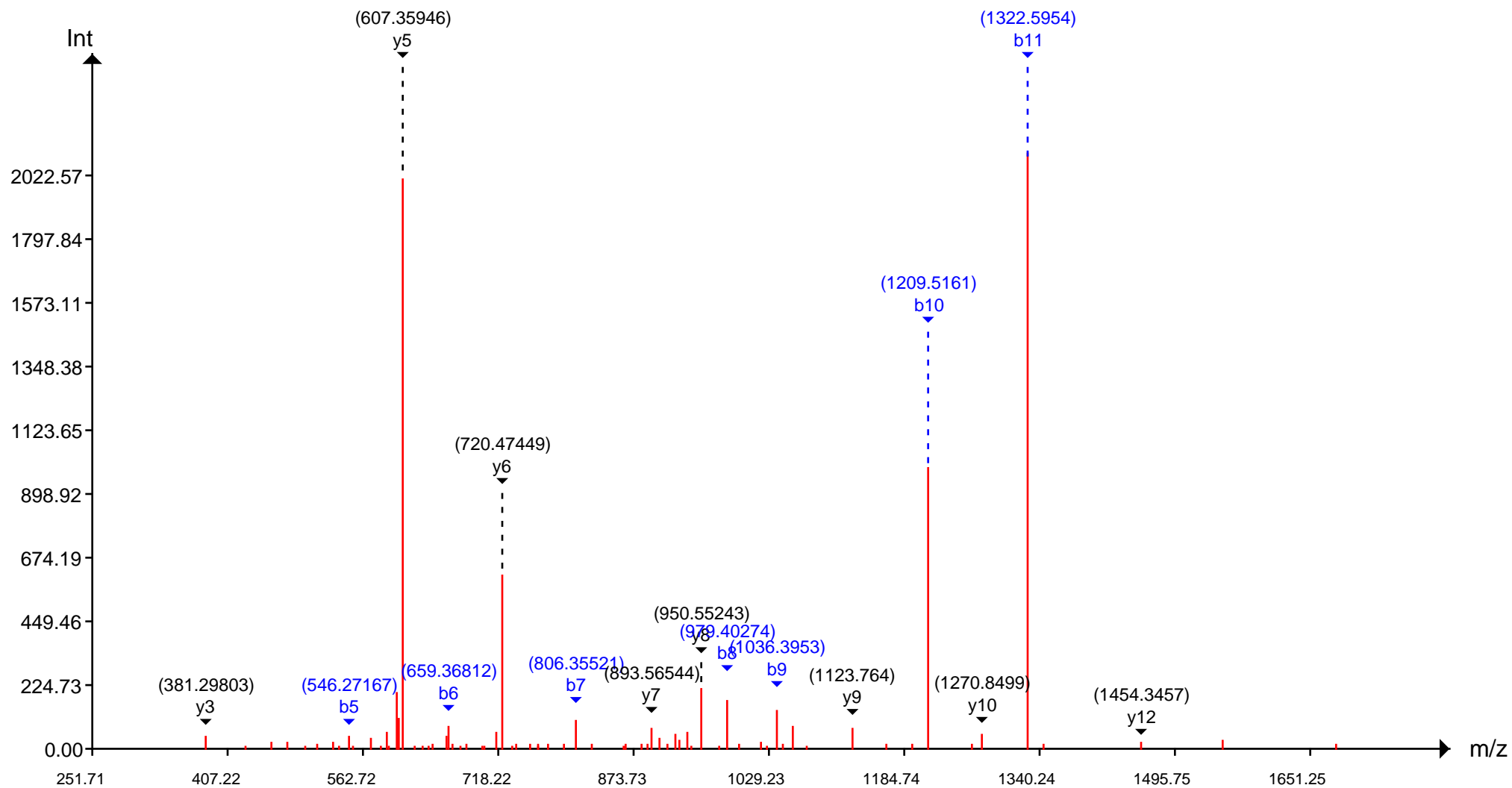
Charge 2+

Mass error 0.001709



Info

Modified Sequence	AcD3-AYEPQGGSGYDYSYAGGR-COOH	Ion score	47.67
Protein	P61979	Confidence level	99.00%
Precursor m/z	971.91874	Identity Threshold	40.93
Charge	2+	Mass error	0.002081



Info

Modified Sequence AcD3-AQTEALM<Mox*>K<AcD3K*>GK<AcD3K*>LPPEAR<C13>-COOH

Ion score 55.55

Protein P06745

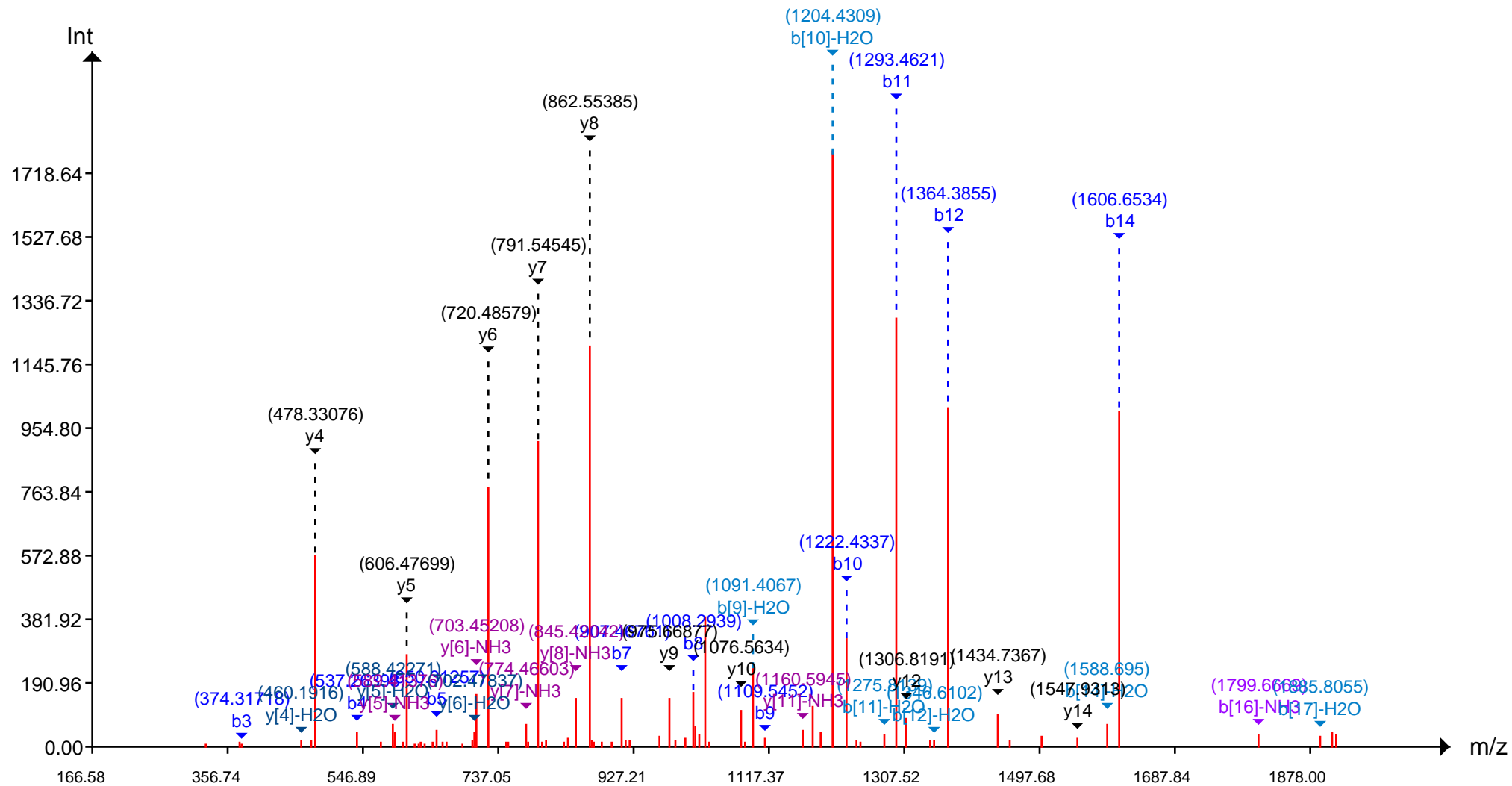
Confidence level 99.00%

Precursor m/z 965.02521

Identity Threshold 47.53

Charge 2+

Mass error 0.002848



Info

Modified Sequence AcD3-TINYLQETTLAANQPLSR<C13>-COOH

Ion score 84.5

Protein O09159

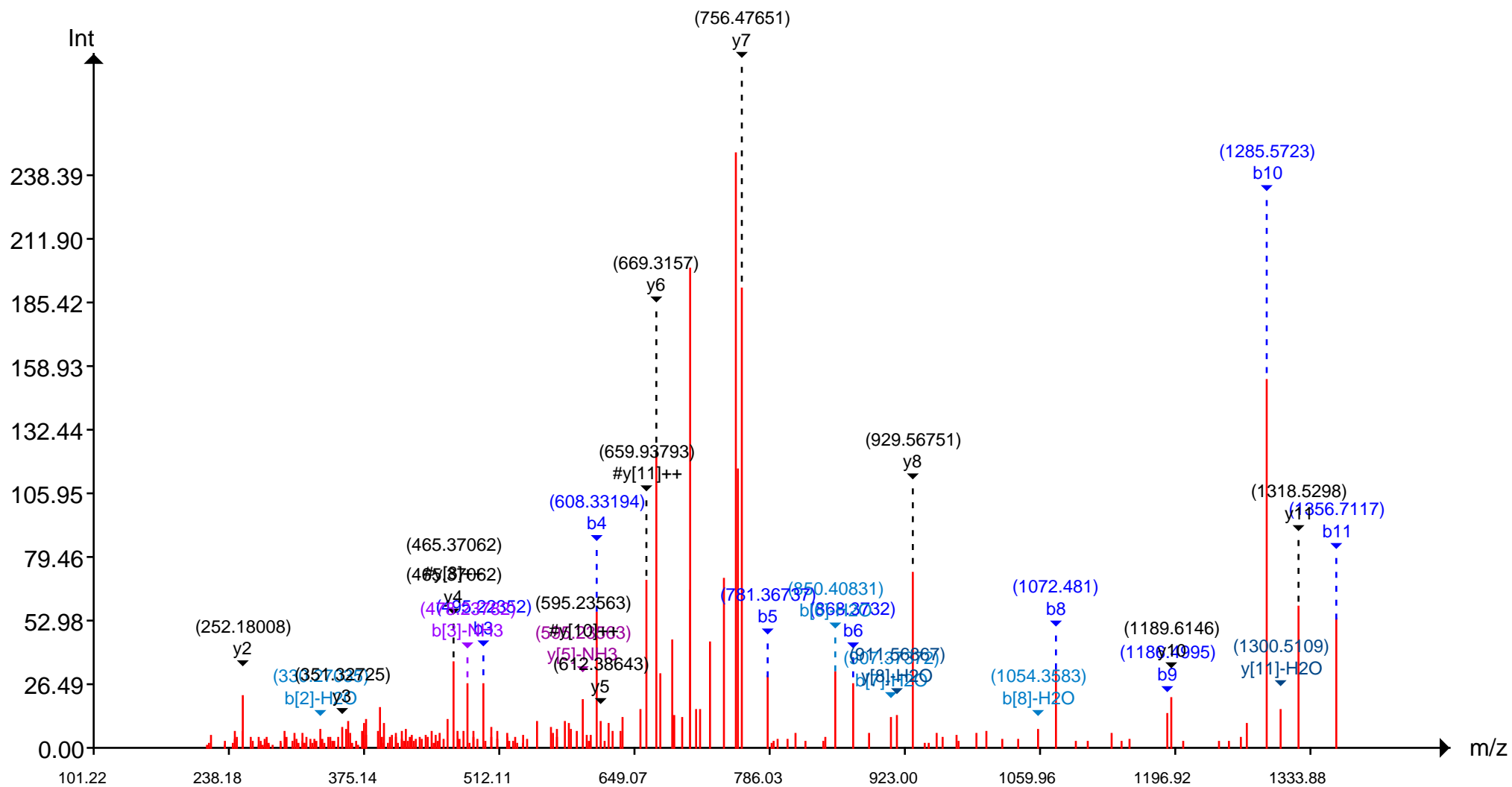
Confidence level 99.00%

Precursor m/z 1042.5627

Identity Threshold 47.60

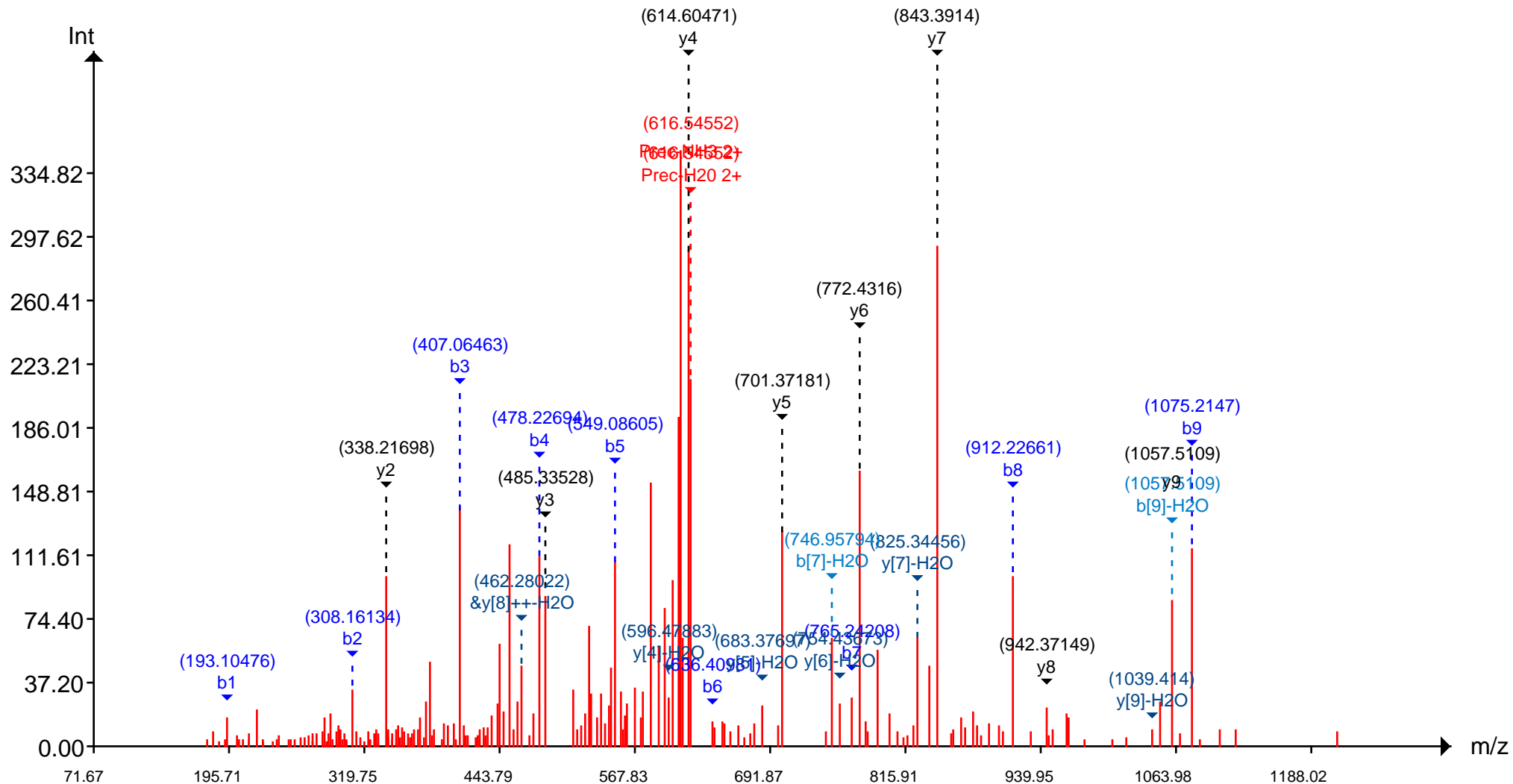
Charge 2+

Mass error 0.002434



Info

Modified Sequence	AcD3-K<AcD3K*>EM<Mox*>IK<AcD3K*>SGM<Mox*>NVAR<C13>-COOH	Ion score	65.19
Protein	P52480	Confidence level	99.00%
Precursor m/z	768.91241	Identity Threshold	48.05
Charge	2+	Mass error	9.57E-4



Info

Modified Sequence AcD3-M<Mox*>DVAASEFYR-COOH

Ion score 61.34

Protein P17182, P21550, P17183

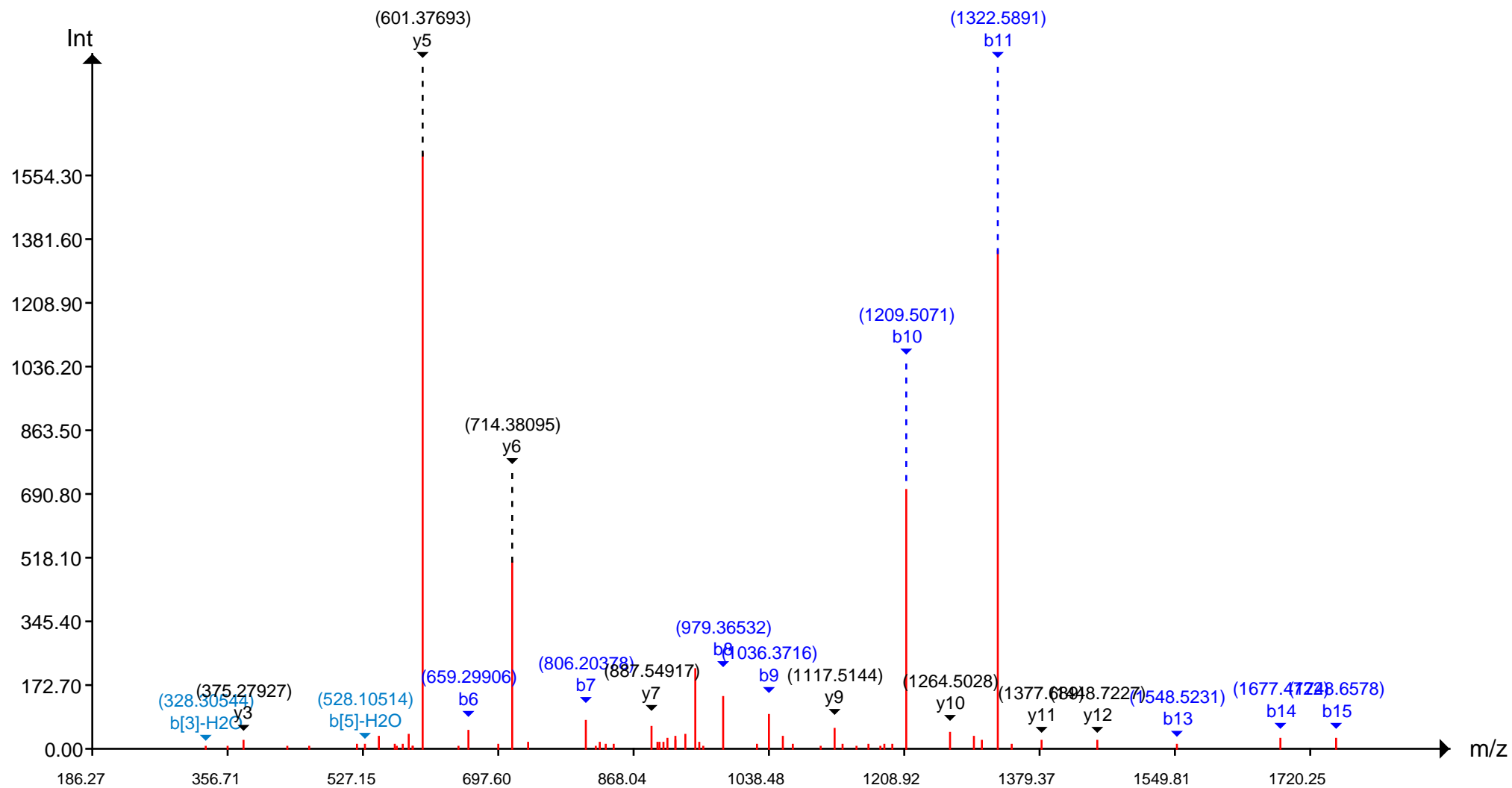
Confidence level 99.00%

Precursor m/z 625.28374

Identity Threshold 44.23

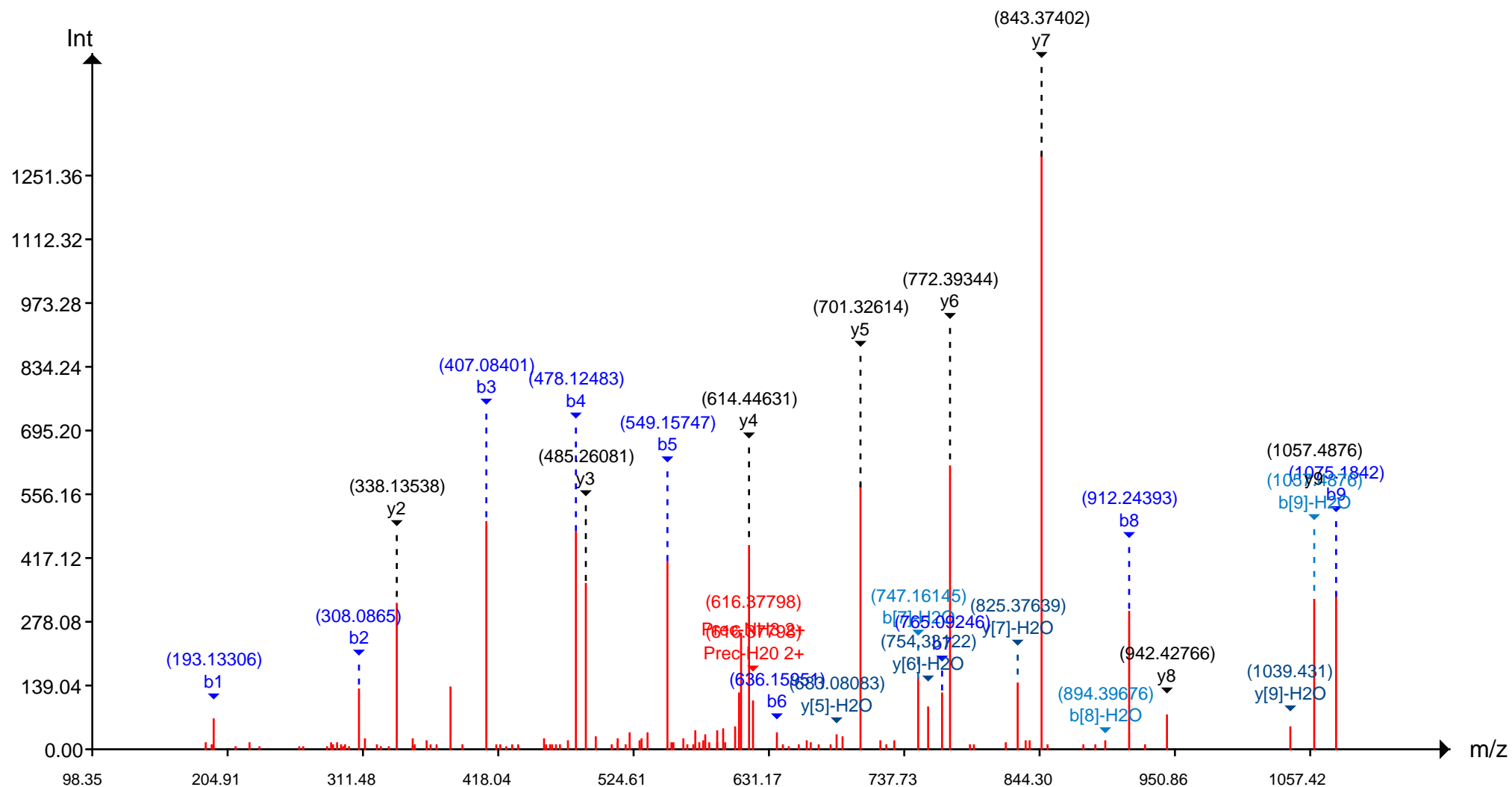
Charge 2+

Mass error 5.29E-4



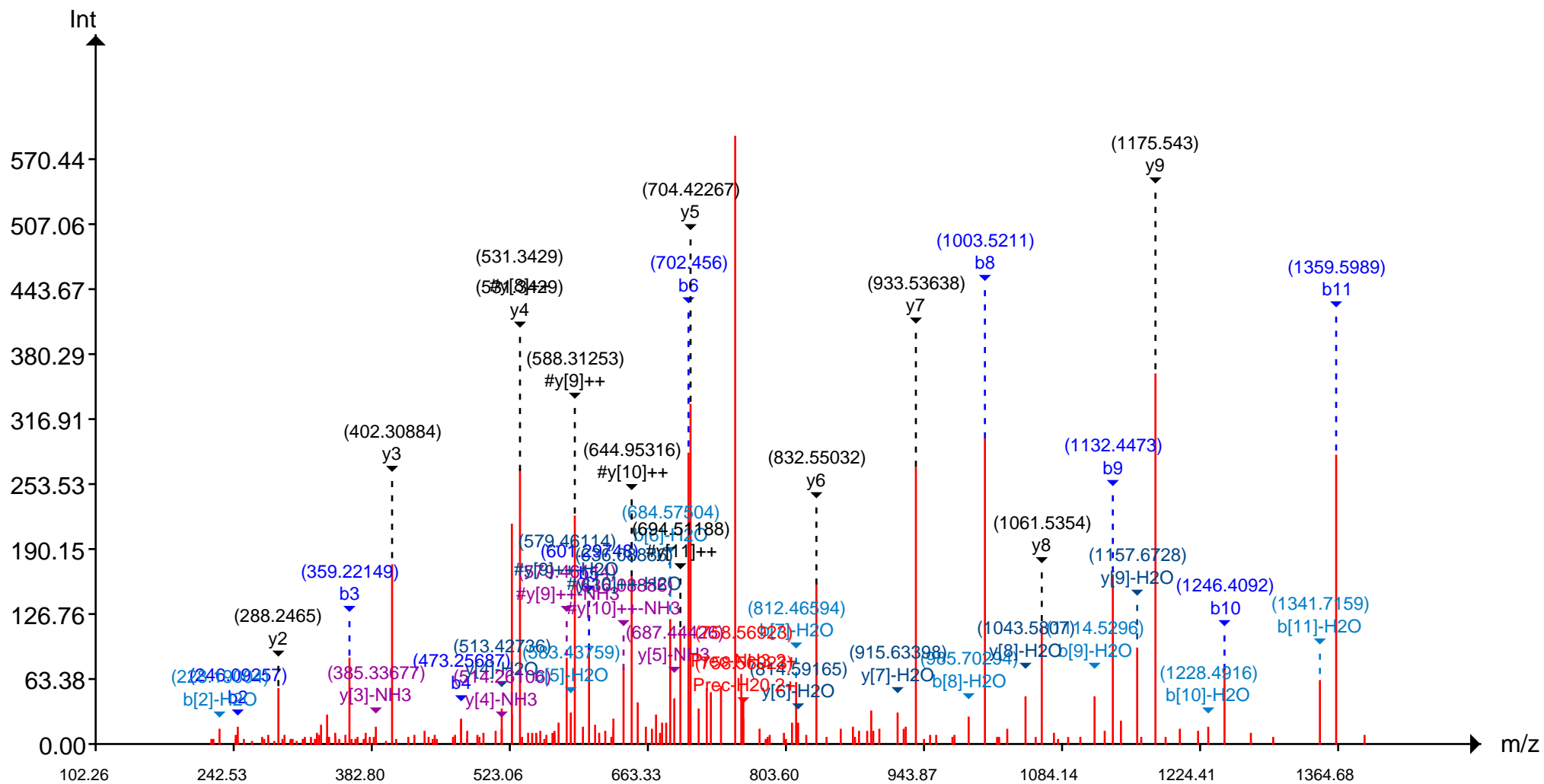
Info

Modified Sequence	AcD3-AQTEALM<Mox*>K<AcD3K*>GK<AcD3K*>LPPEAR-COOH	Ion score	81.98
Protein	P06745	Confidence level	99.00%
Precursor m/z	962.01535	Identity Threshold	47.91
Charge	2+	Mass error	0.00327



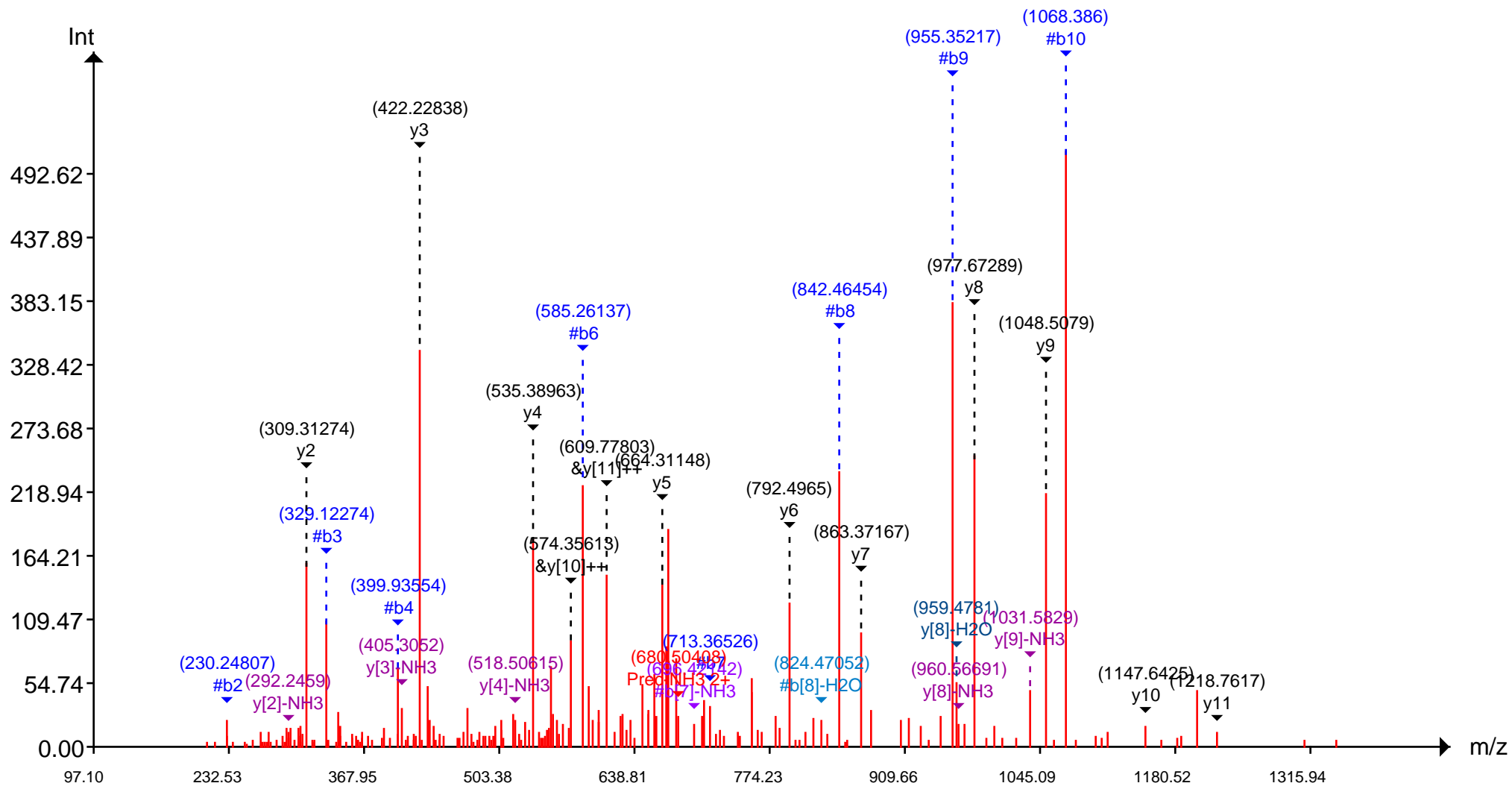
Info

Modified Sequence	AcD3-M<Mox*>DVAASEFYR-COOH	Ion score	67.85
Protein	P17182, P21550, P17183	Confidence level	99.00%
Precursor m/z	625.28412	Identity Threshold	44.31
Charge	2+	Mass error	0.001289



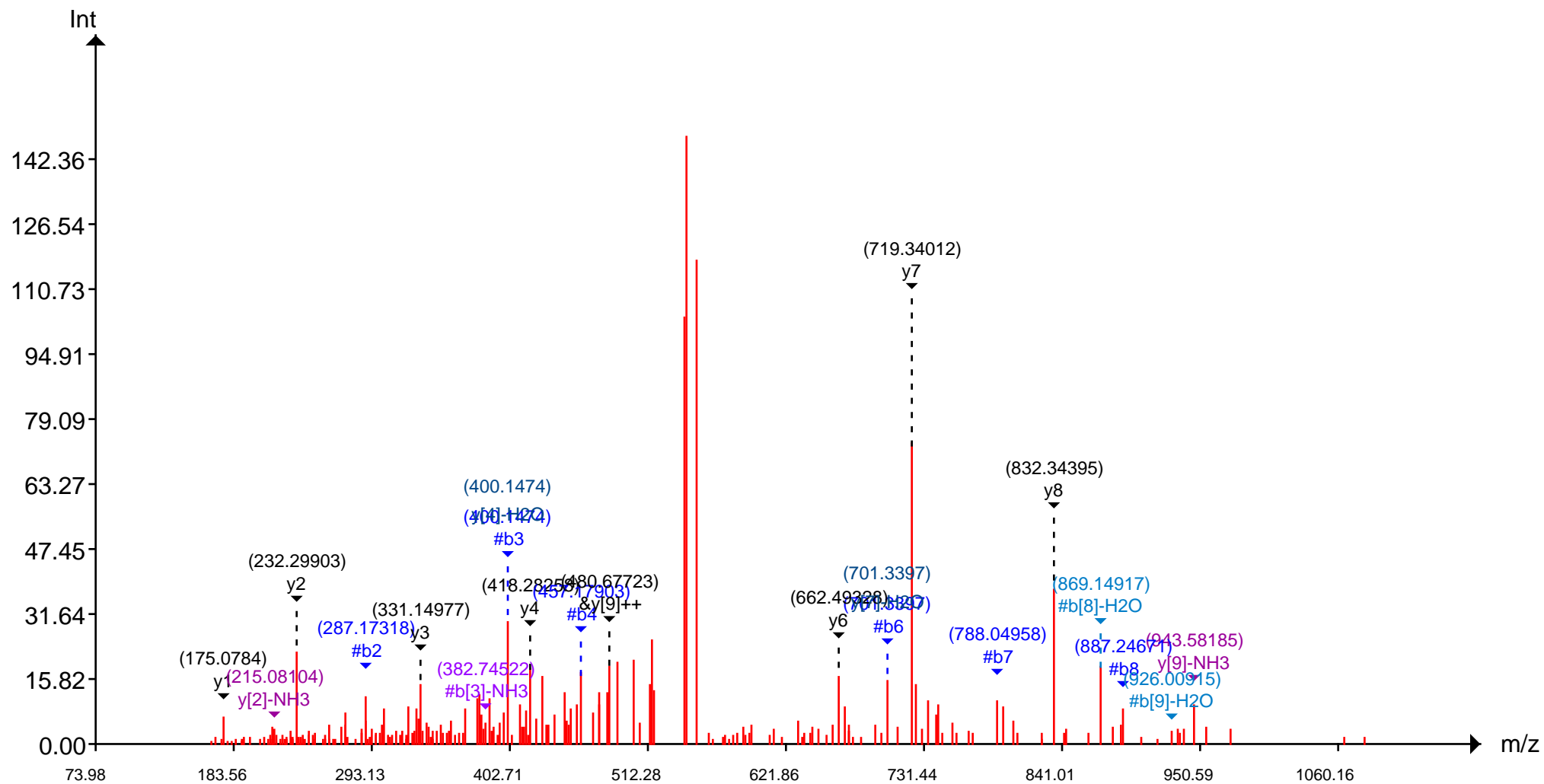
Info

Modified Sequence	AcD3-TVINQTQK<AcD3K*>ENLR-COOH	Ion score	88.51
Protein	Q6ZWW7	Confidence level	99.00%
Precursor m/z	767.42918	Identity Threshold	47.35
Charge	2+	Mass error	8.97E-4



Info

Modified Sequence	AcD3-IAVANAQELLQR<C13>-COOH	Ion score	89.84
Protein	P62075	Confidence level	99.00%
Precursor m/z	688.90551	Identity Threshold	45.42
Charge	2+	Mass error	6.76E-4



Info

Modified Sequence AcD3-LQIGDESVGR-COOH

Protein P24369

Precursor m/z 559.79789

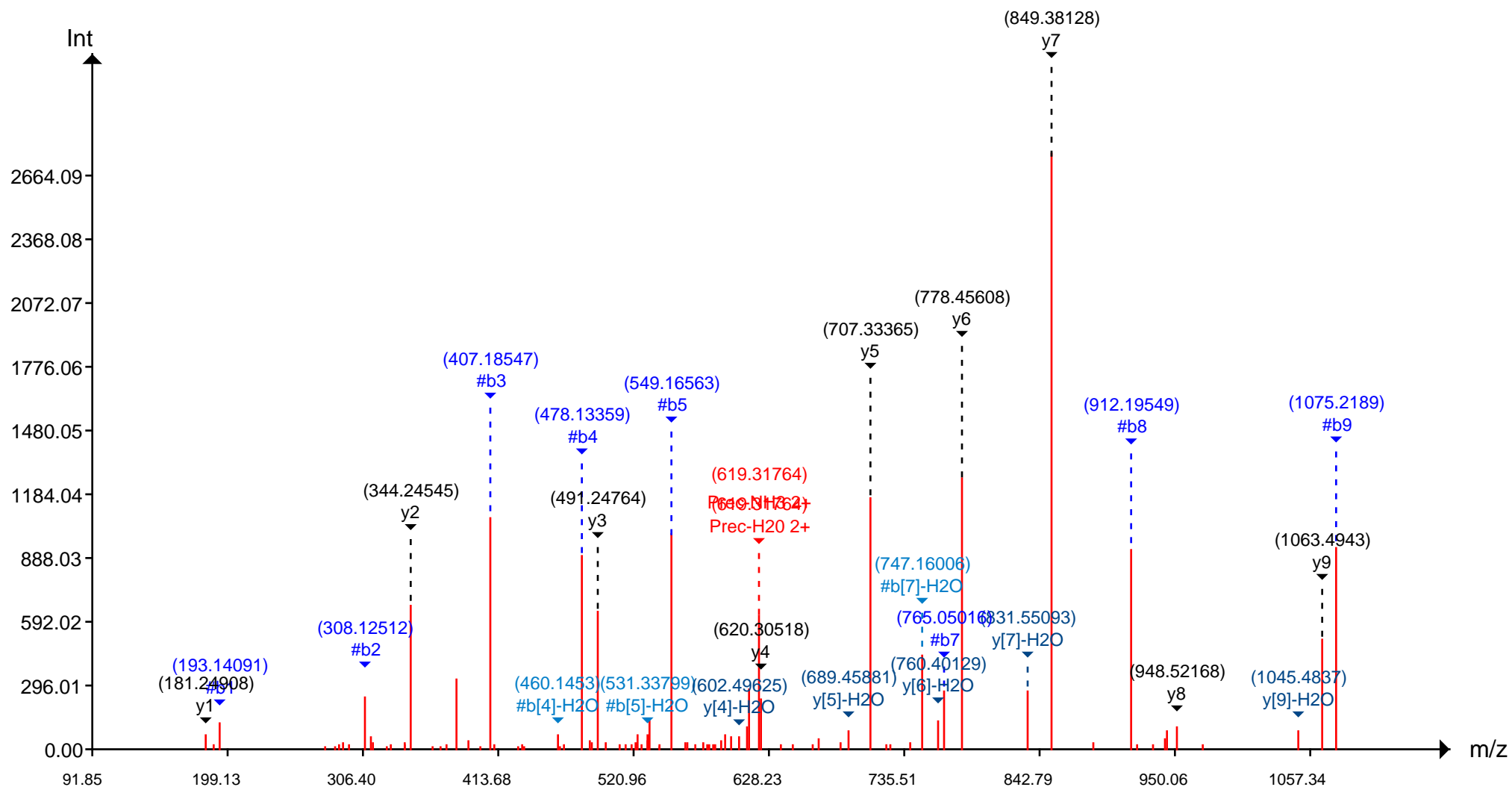
Charge 2+

Ion score 51.42

Confidence level 99.00%

Identity Threshold 48.09

Mass error 5.55E-4



Info

Modified Sequence AcD3-M<Mox*>DVAASEFYR<C13>-COOH

Ion score 84.47

Protein P17182, P21550, P17183

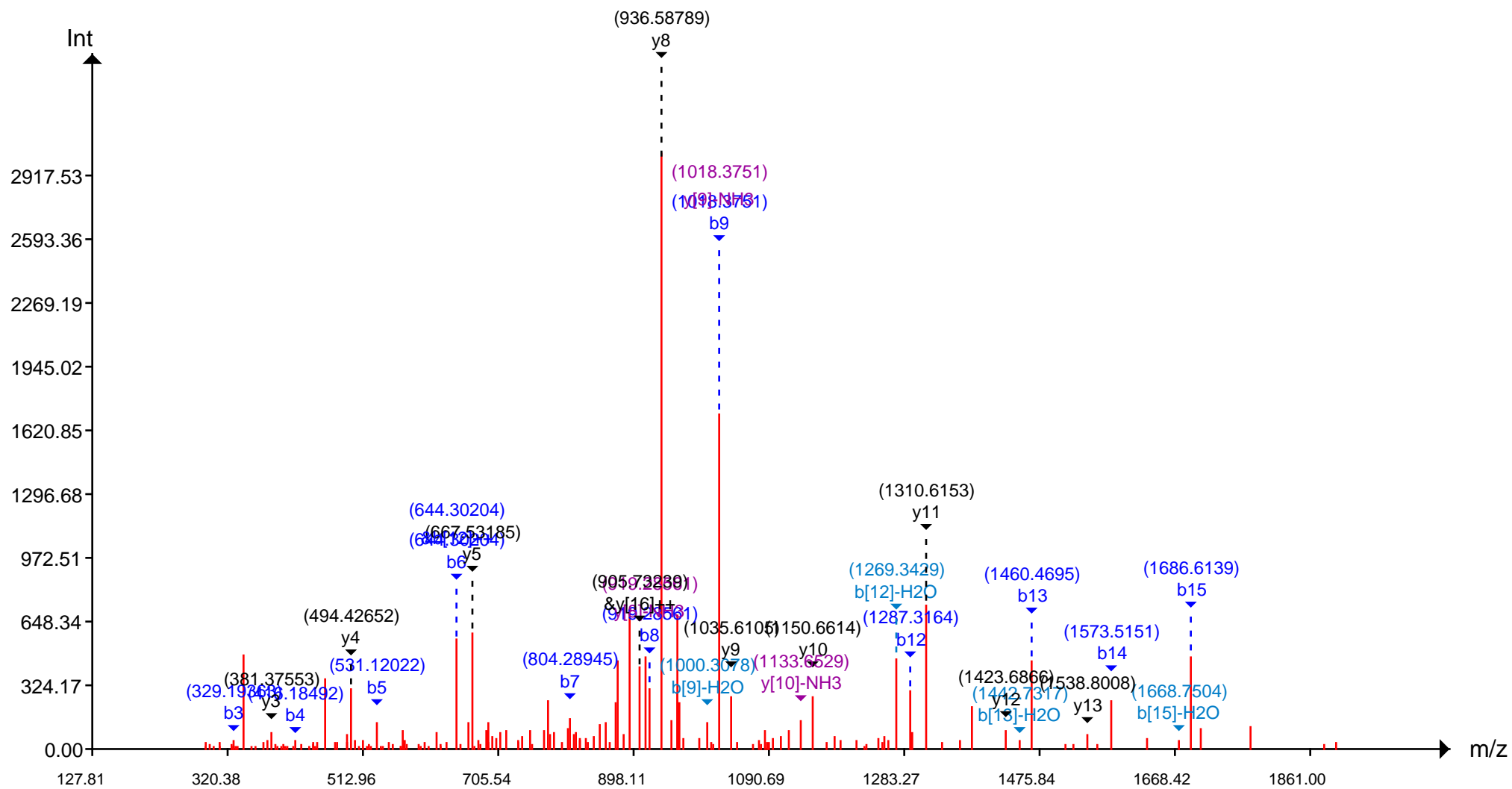
Confidence level 99.00%

Precursor m/z 628.29363

Identity Threshold 45.39

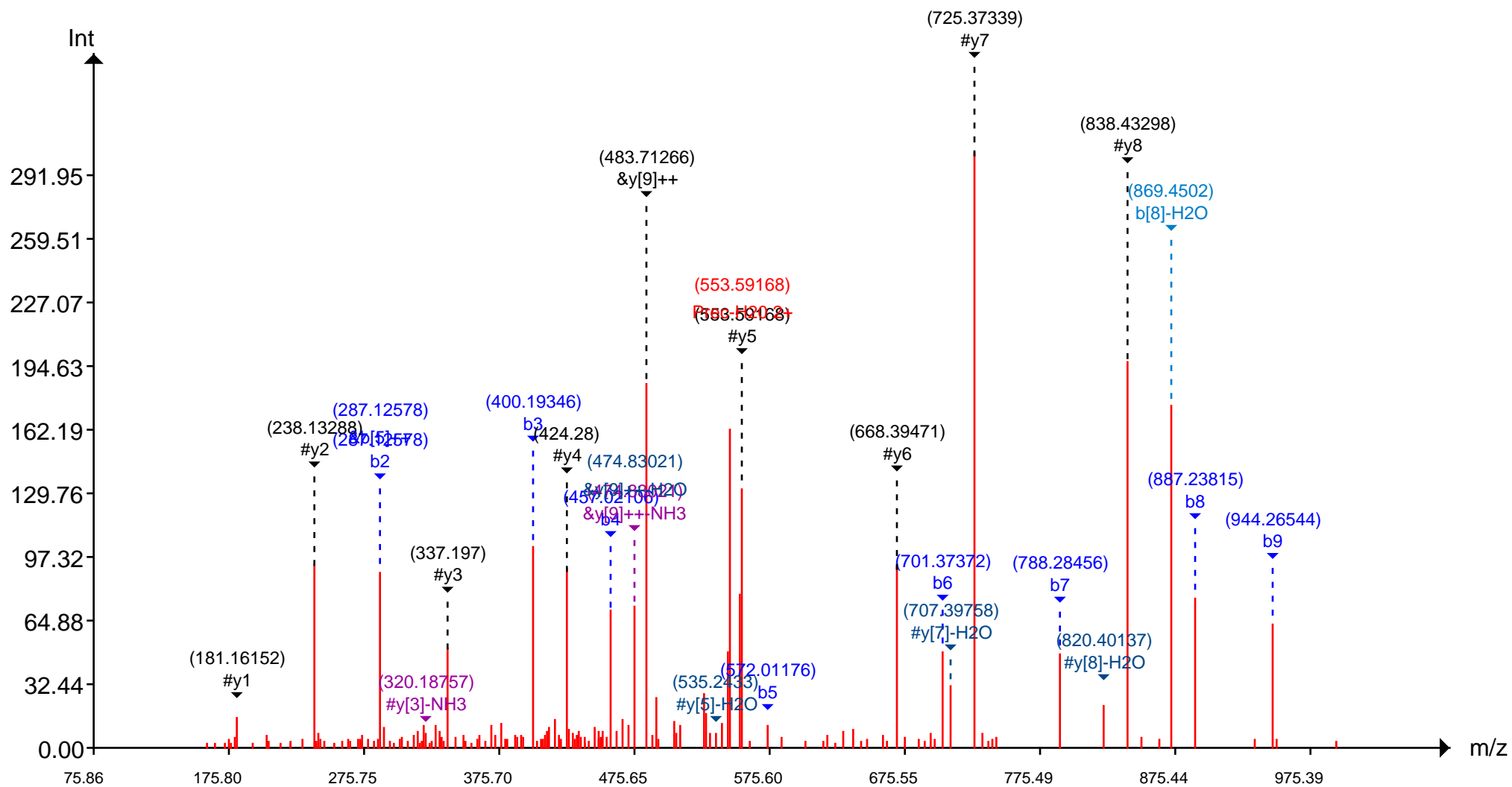
Charge 2+

Mass error 1.68E-4



Info

Modified Sequence	AcD3-VALSDIC<Cmm*>DVPTAK<AcD3K*>IISR<C13>-COOH	Ion score	56.86
Protein	Q9CWJ9	Confidence level	99.00%
Precursor m/z	977.54815	Identity Threshold	46.44
Charge	2+	Mass error	-1.61E-4



Info

Modified Sequence AcD3-LQIGDESVGR<C13>-COOH

Ion score 70.3

Protein P24369

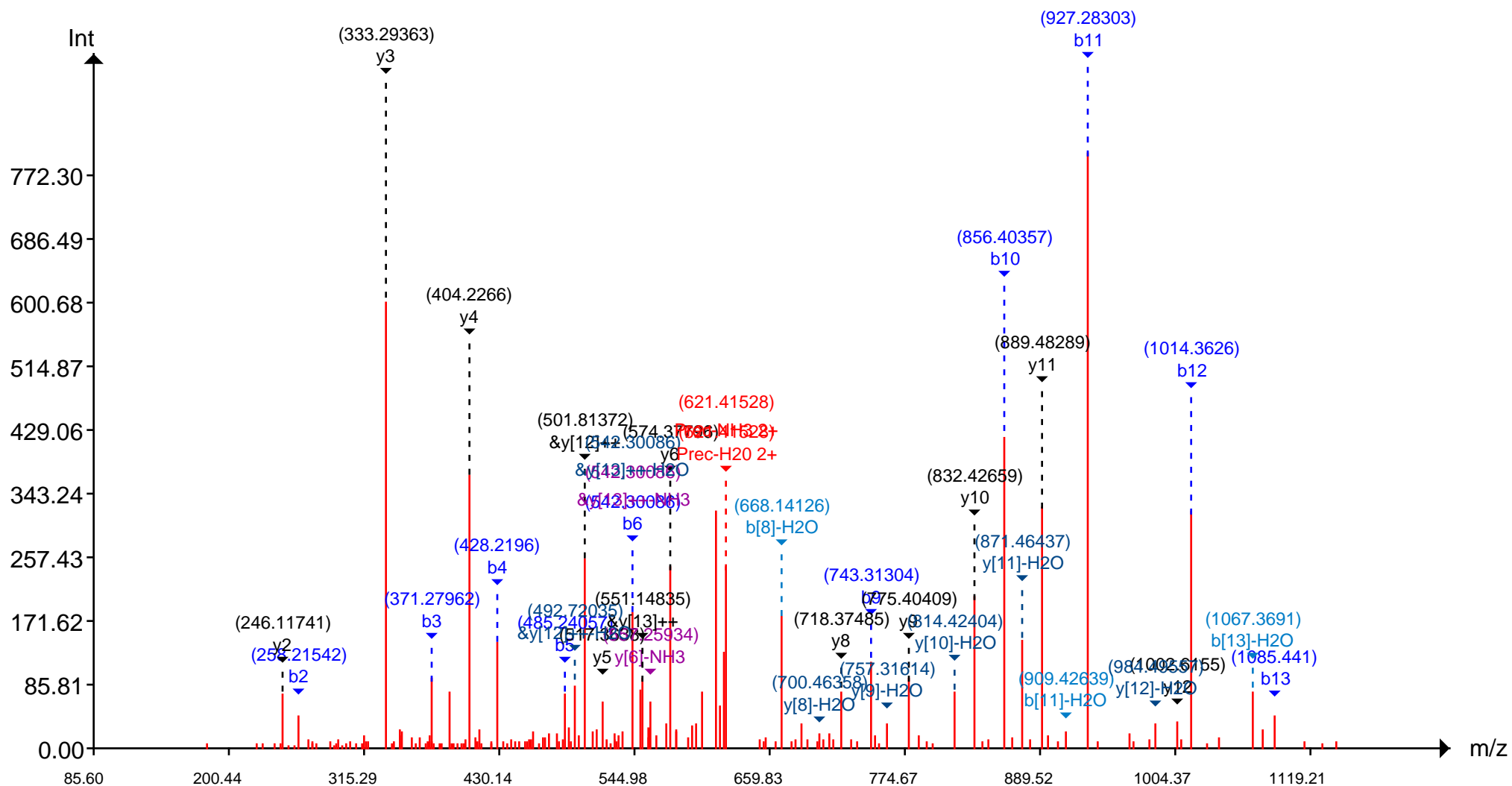
Confidence level 99.00%

Precursor m/z 562.80806

Identity Threshold 47.57

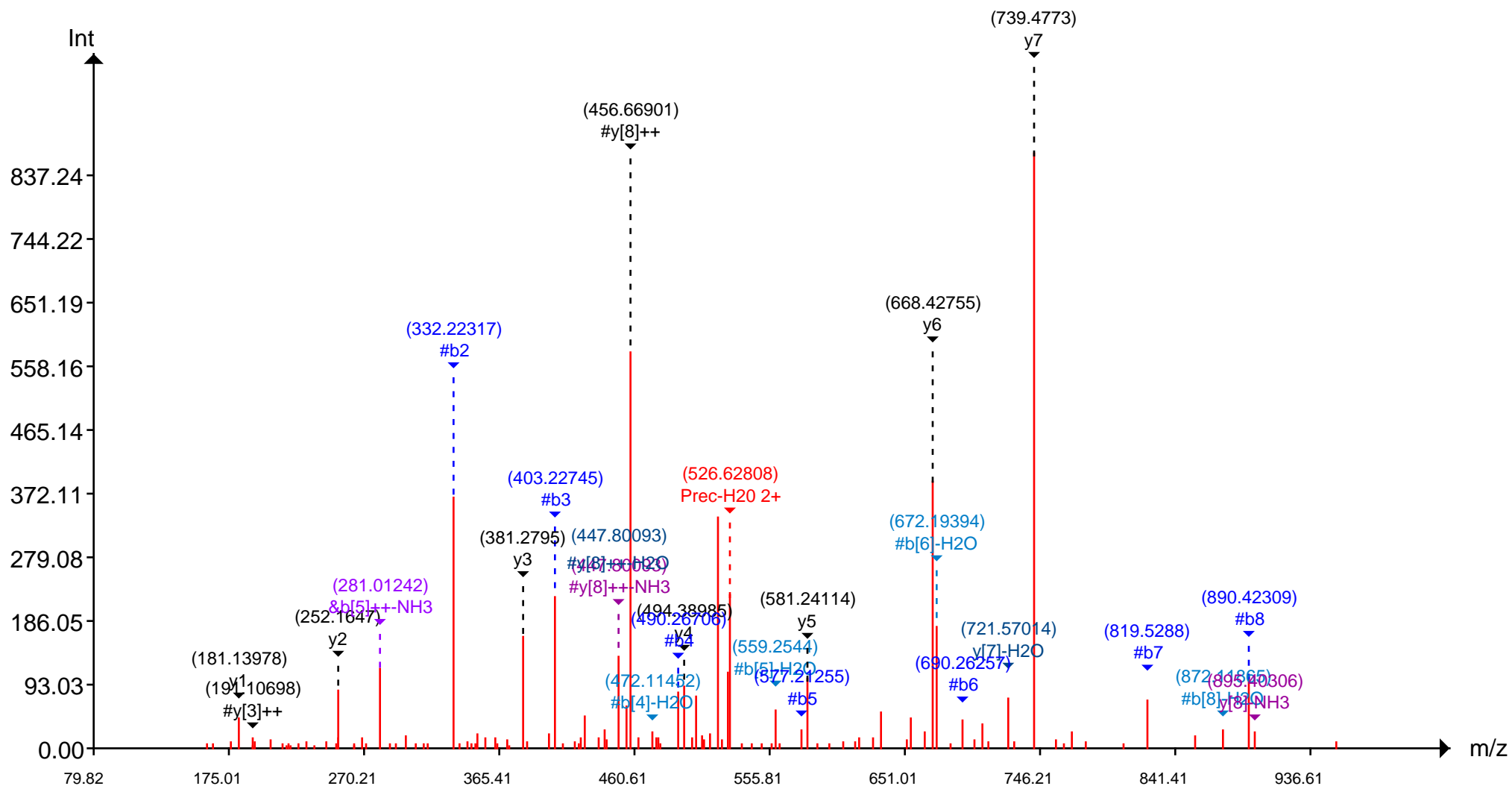
Charge 2+

Mass error 7.53E-4



Info

Modified Sequence	AcD3-LVIGGGSGGLASAR-COOH	Ion score	88.45
Protein	P47791	Confidence level	99.00%
Precursor m/z	630.36108	Identity Threshold	46.98
Charge	2+	Mass error	3.33E-4



Info

Modified Sequence AcD3-IK<AcD3K*>ASSIEAR<C13>-COOH

Ion score 52.08

Protein P57759

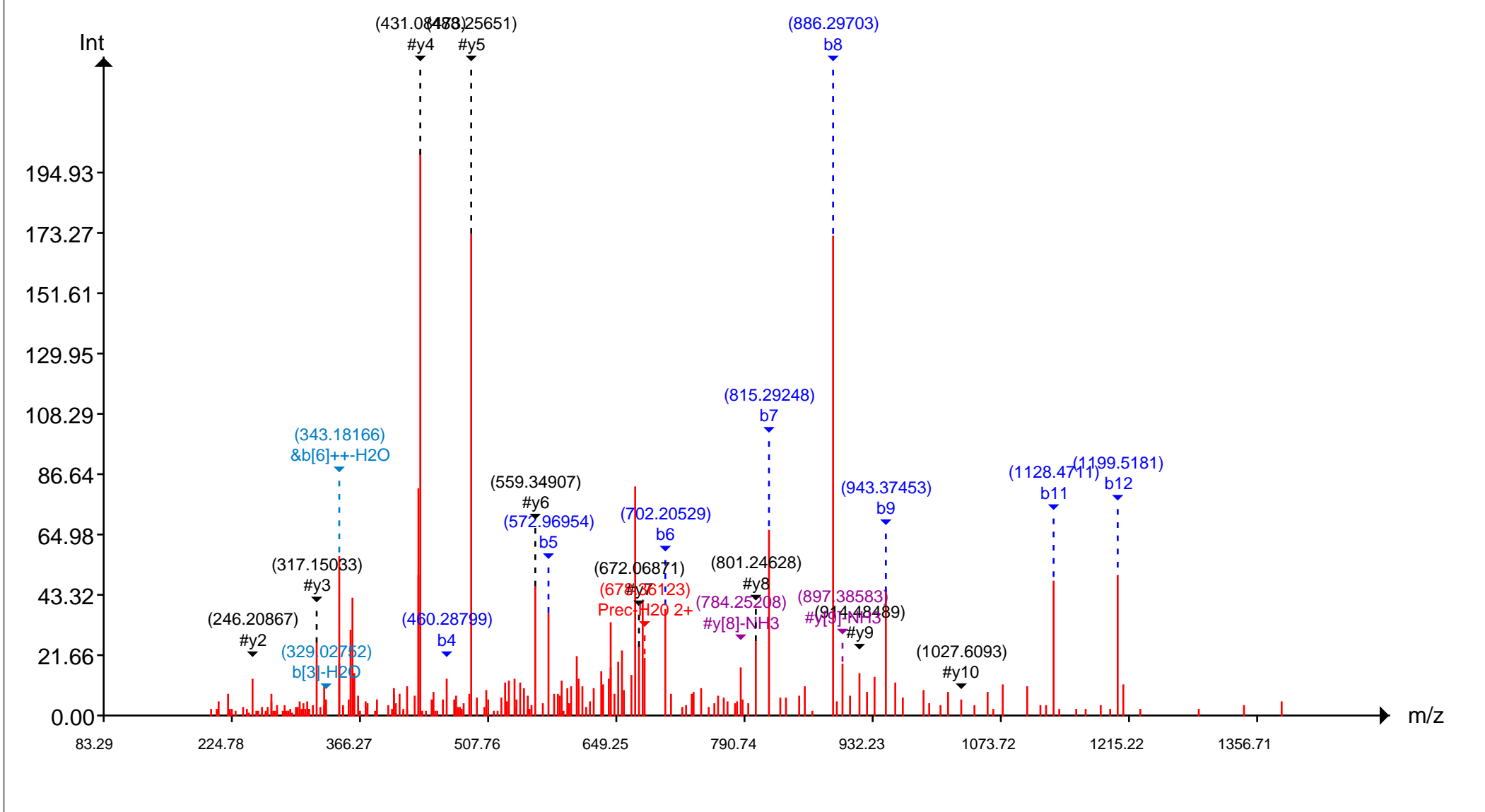
Confidence level 99.00%

Precursor m/z 535.82466

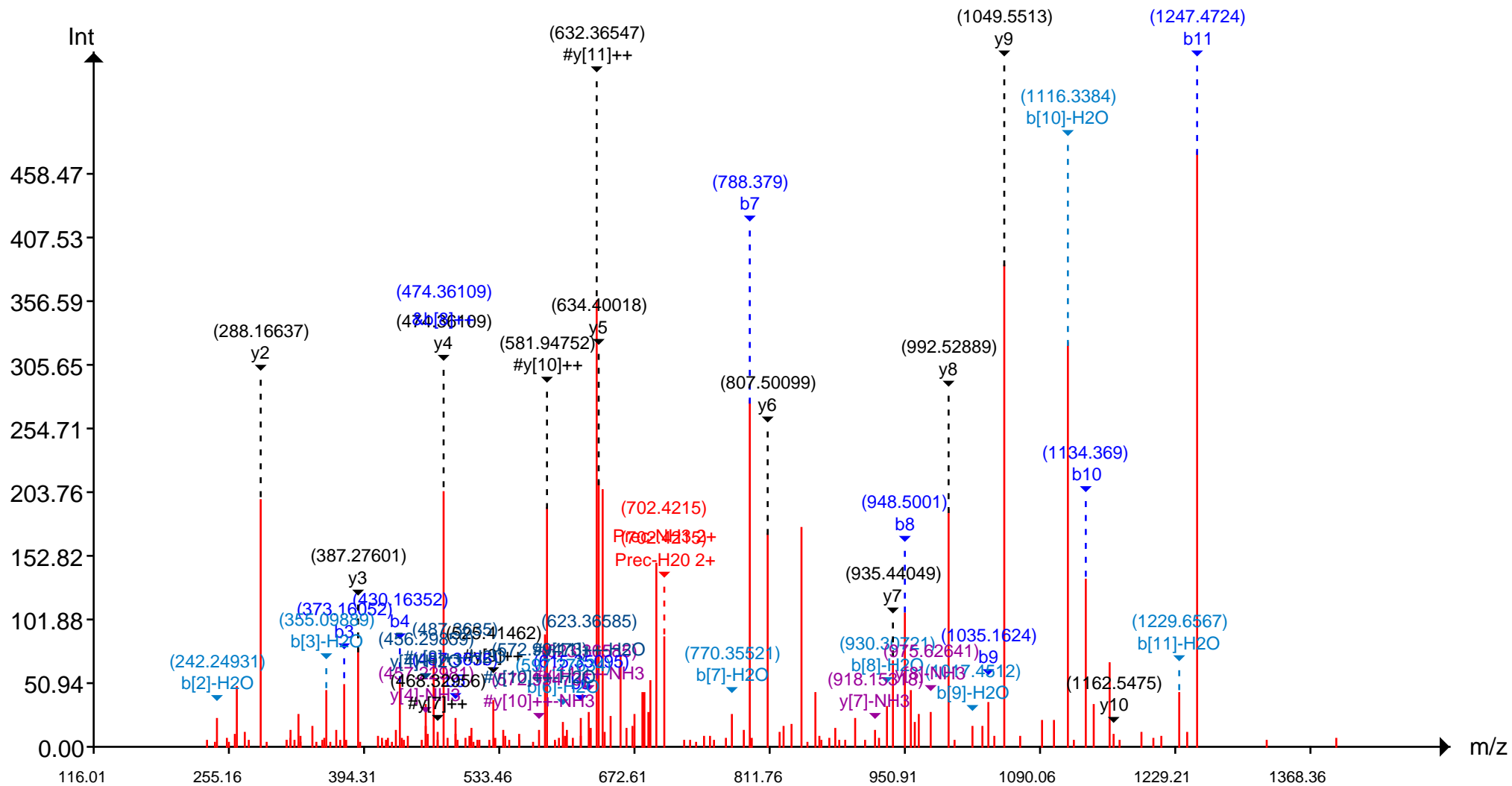
Identity Threshold 44.68

Charge 2+

Mass error 2.31E-4

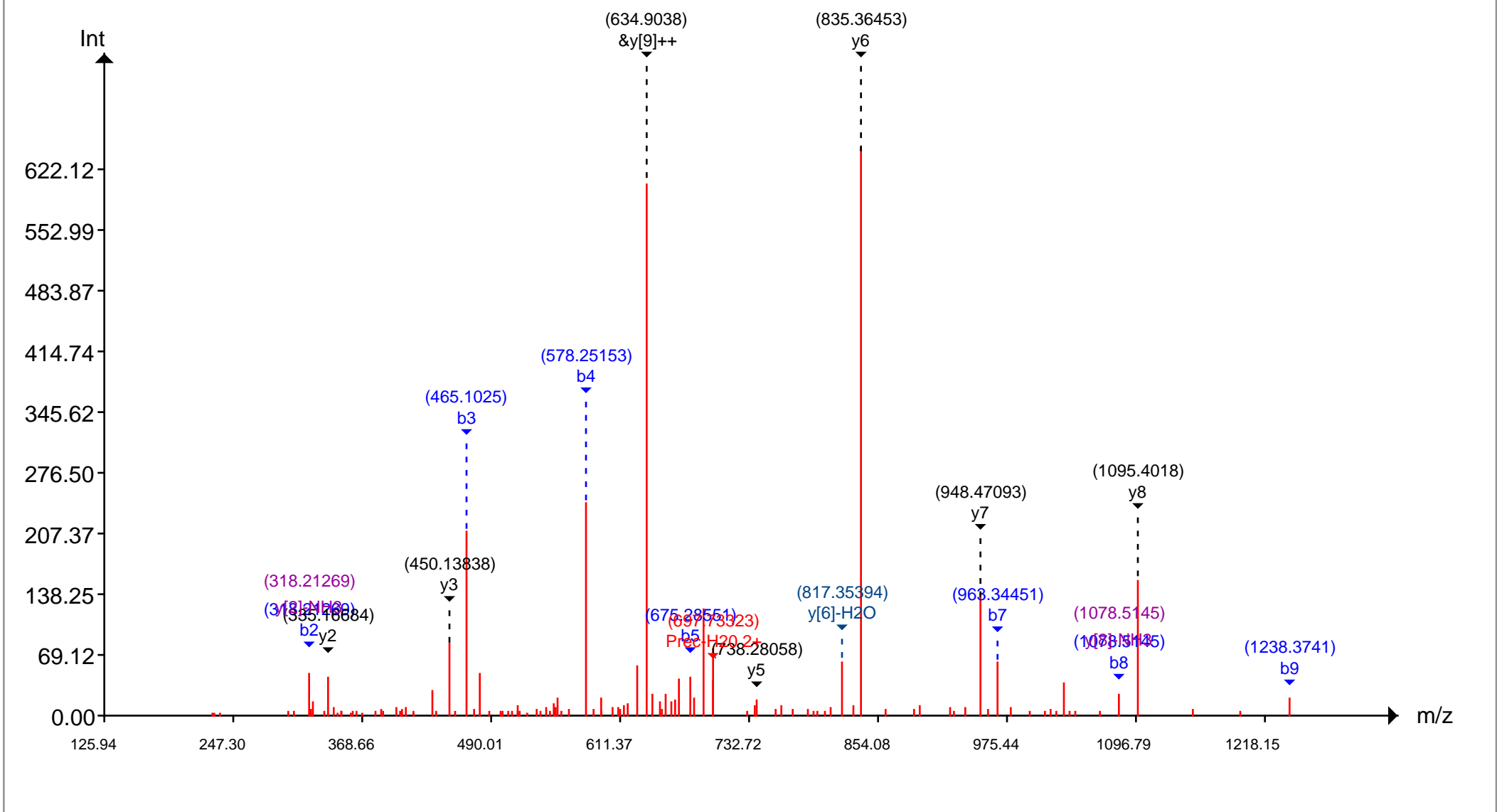


Info			
Modified ...	AcD3-TAEILELAGNAAR-COOH	Ion score	47
	Q8CGP5, Q8CGP6, Q8CGP7, P22752, Q6GSS7, Q64522, Q64523, Q8BFU2, Q8R1M2, P27661, Q9QZQ8	Confidence level	99
Prec...	687.37735	Identity Threshold	47
	2+	Mass error	0.



Info

Modified Sequence	AcD3-LTLGGQK<AcD3K*>C<Cmm*>SVIR-COOH	Ion score	57.82
Protein	P62962	Confidence level	99.00%
Precursor m/z	711.40726	Identity Threshold	46.70
Charge	2+	Mass error	0.002071



Info

Modified Sequence AcD3-VK<AcD3K*>M<Mox*>LPDK<AcD3K*>DC<Cmm*>R-COOH

Ion score 50.83

Protein P18760

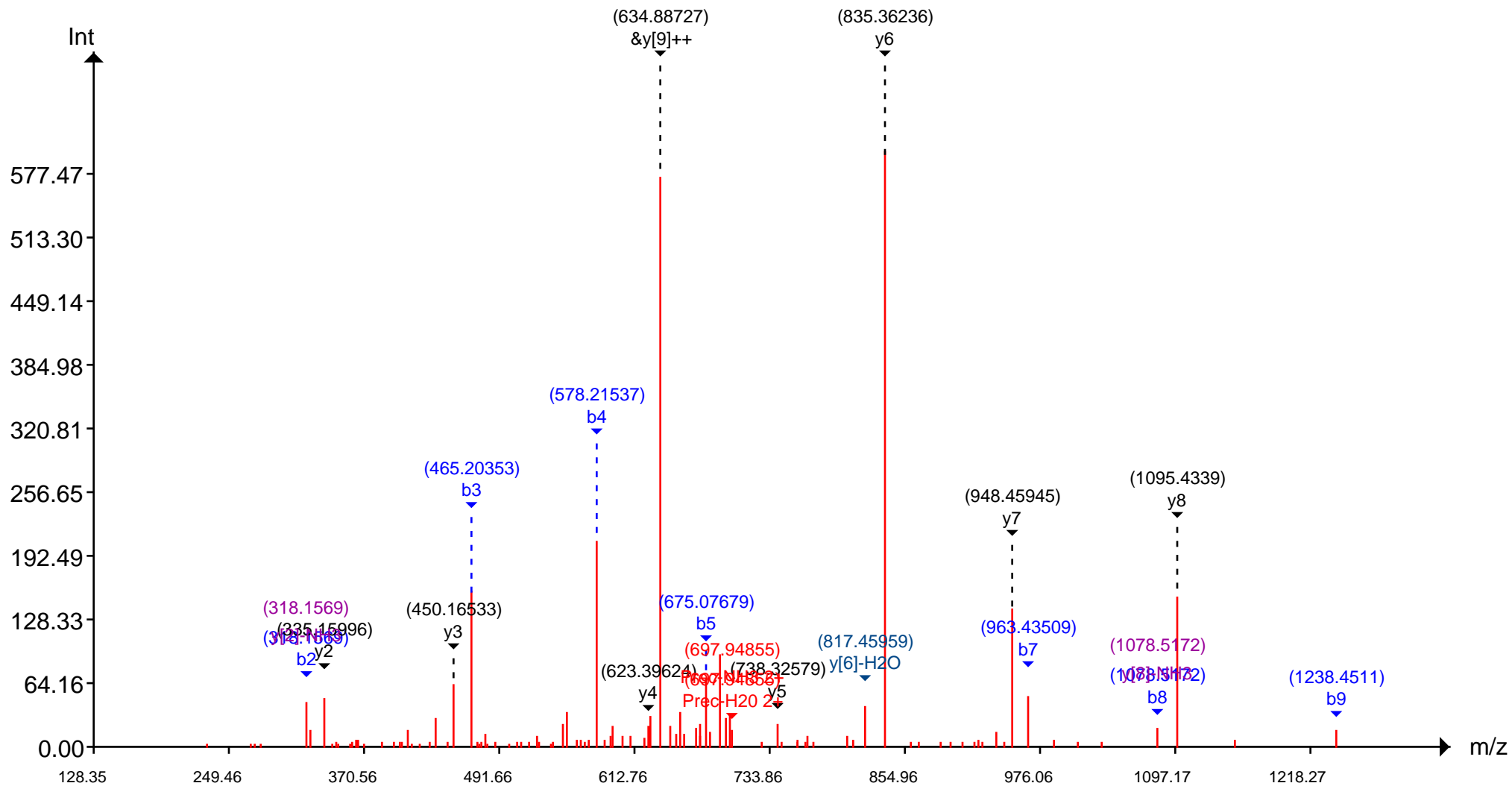
Confidence level 99.00%

Precursor m/z 706.8651

Identity Threshold 48.08

Charge 2+

Mass error 6.82E-4



Info

Modified Sequence AcD3-VK<AcD3K*>M<Mox*>LPDK<AcD3K*>DC<Cmm*>R-COOH

Ion score 48.06

Protein P18760

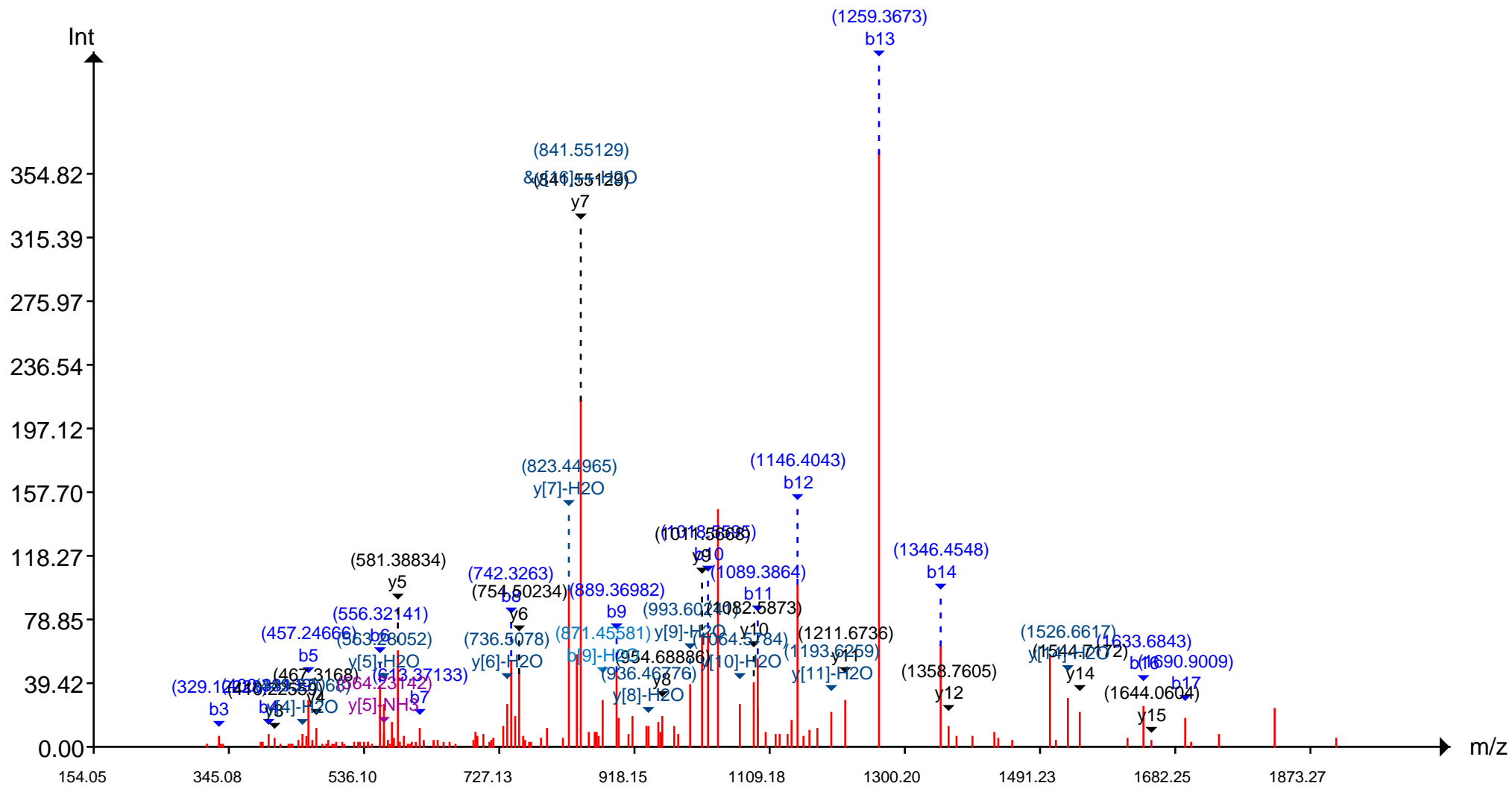
Confidence level 99.00%

Precursor m/z 706.86606

Identity Threshold 48.03

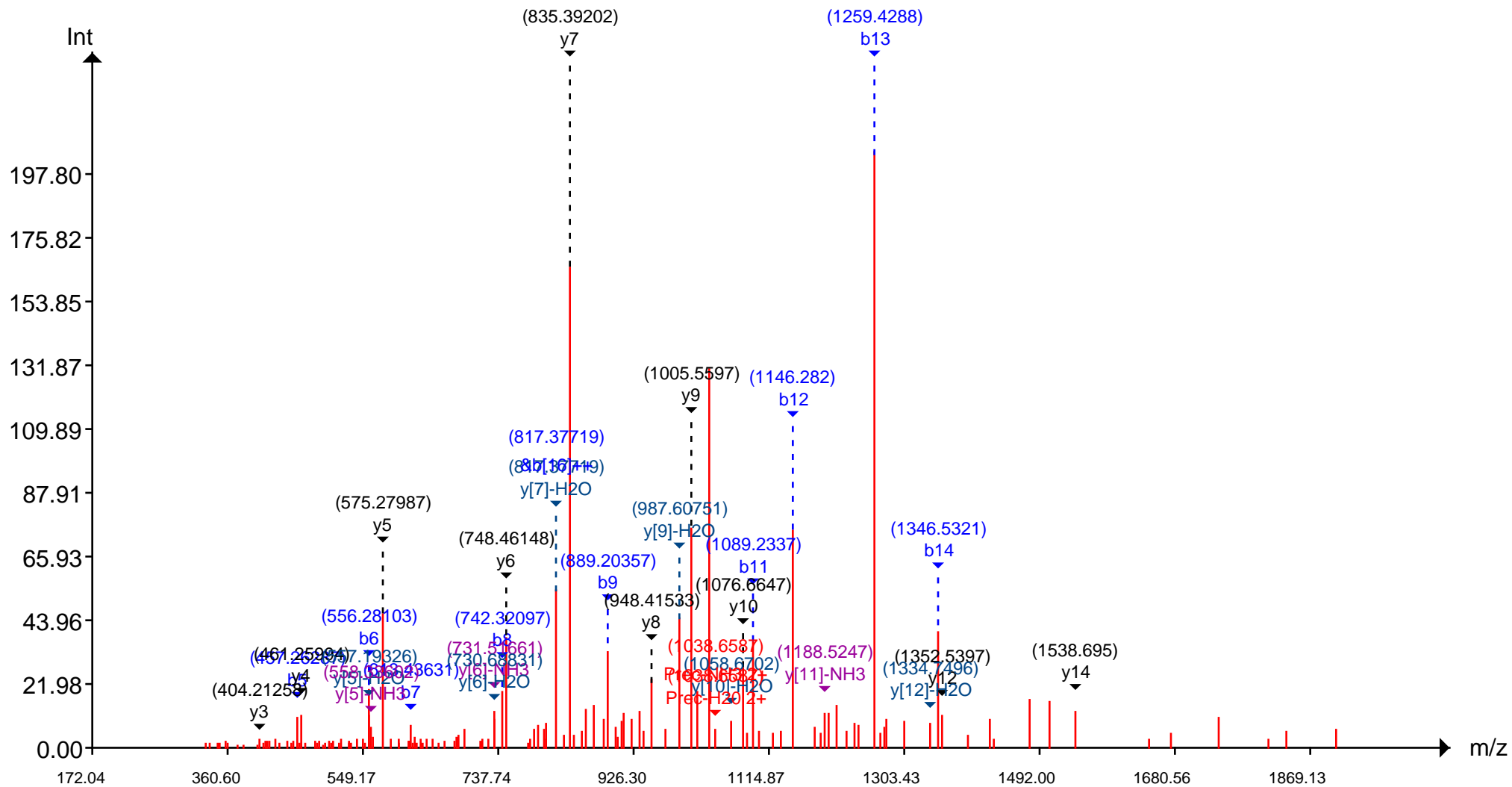
Charge 2+

Mass error 0.002602



Info

Modified Sequence	AcD3-IVAAGVGEFEAGISK<AcD3K*>NGQTR<C13>-COOH	Ion score	107.65
Protein	P10126, P62631	Confidence level	99.00%
Precursor m/z	1050.5697	Identity Threshold	47.33
Charge	2+	Mass error	0.00232



Info

Modified Sequence AcD3-IVAAGVGEFEAGISK<AcD3K*>NGQTR-COOH

Ion score 58.42

Protein P10126, P62631

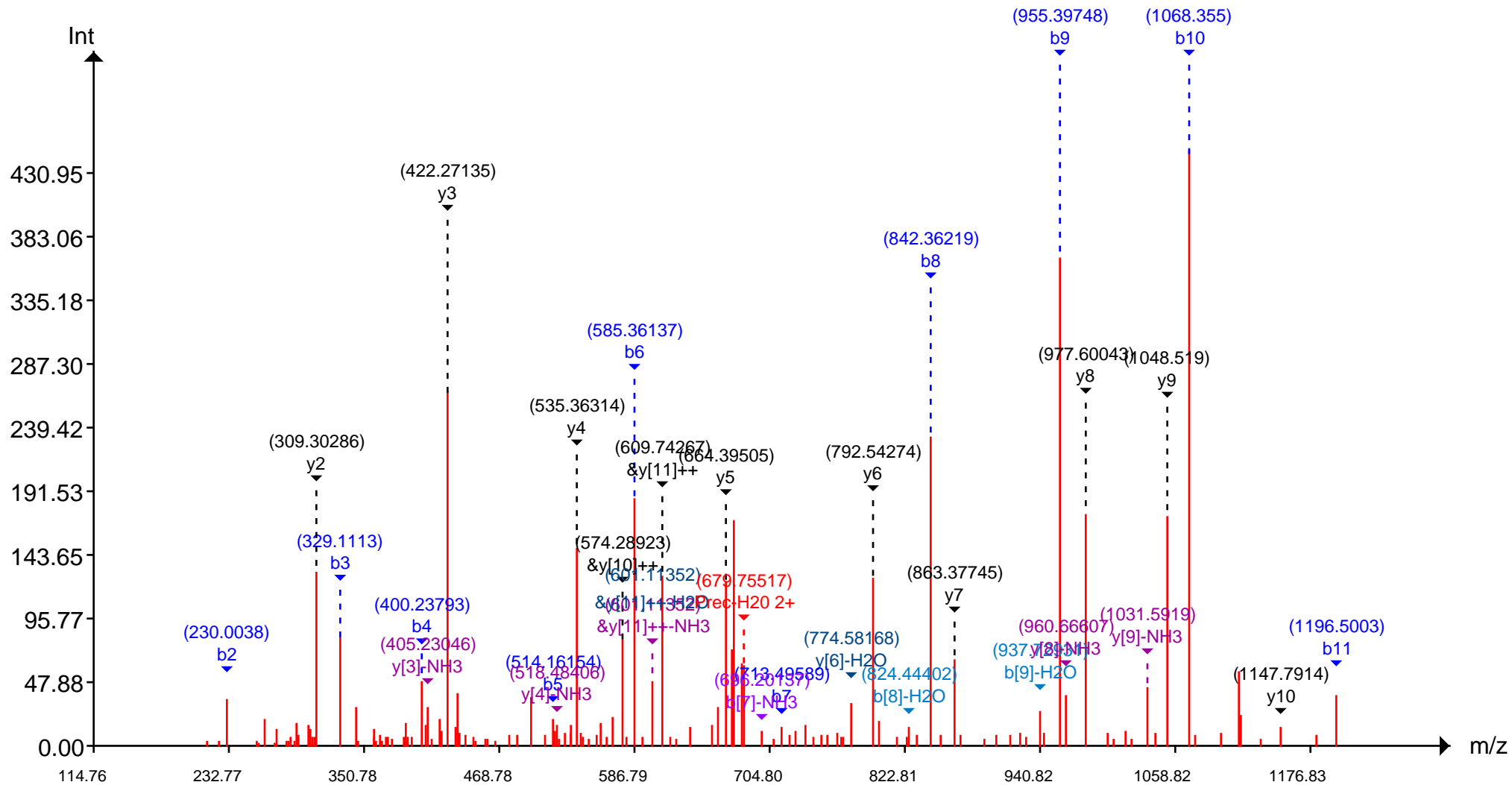
Confidence level 99.00%

Precursor m/z 1047.5593

Identity Threshold 47.79

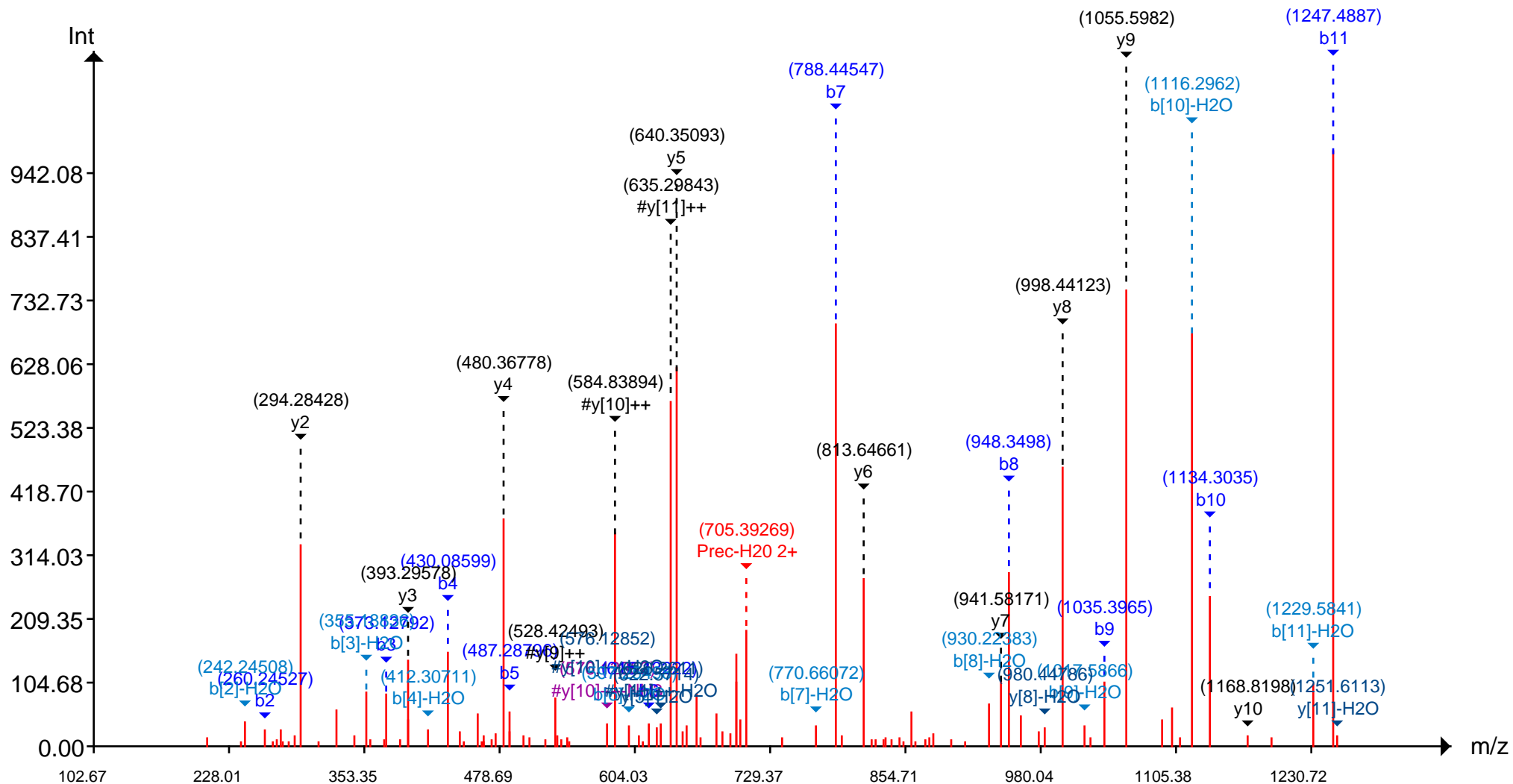
Charge 2+

Mass error 0.001662



Info

Modified Sequence	AcD3-IAVANAQELLQR<C13>-COOH	Ion score	90.28
Protein	P62075	Confidence level	99.00%
Precursor m/z	688.90555	Identity Threshold	45.42
Charge	2+	Mass error	7.56E-4



Info

Modified Sequence AcD3-LTLGGQK<AcD3K*>C<Cmm*>SVIR<C13>-COOH

Ion score 88.44

Protein P62962

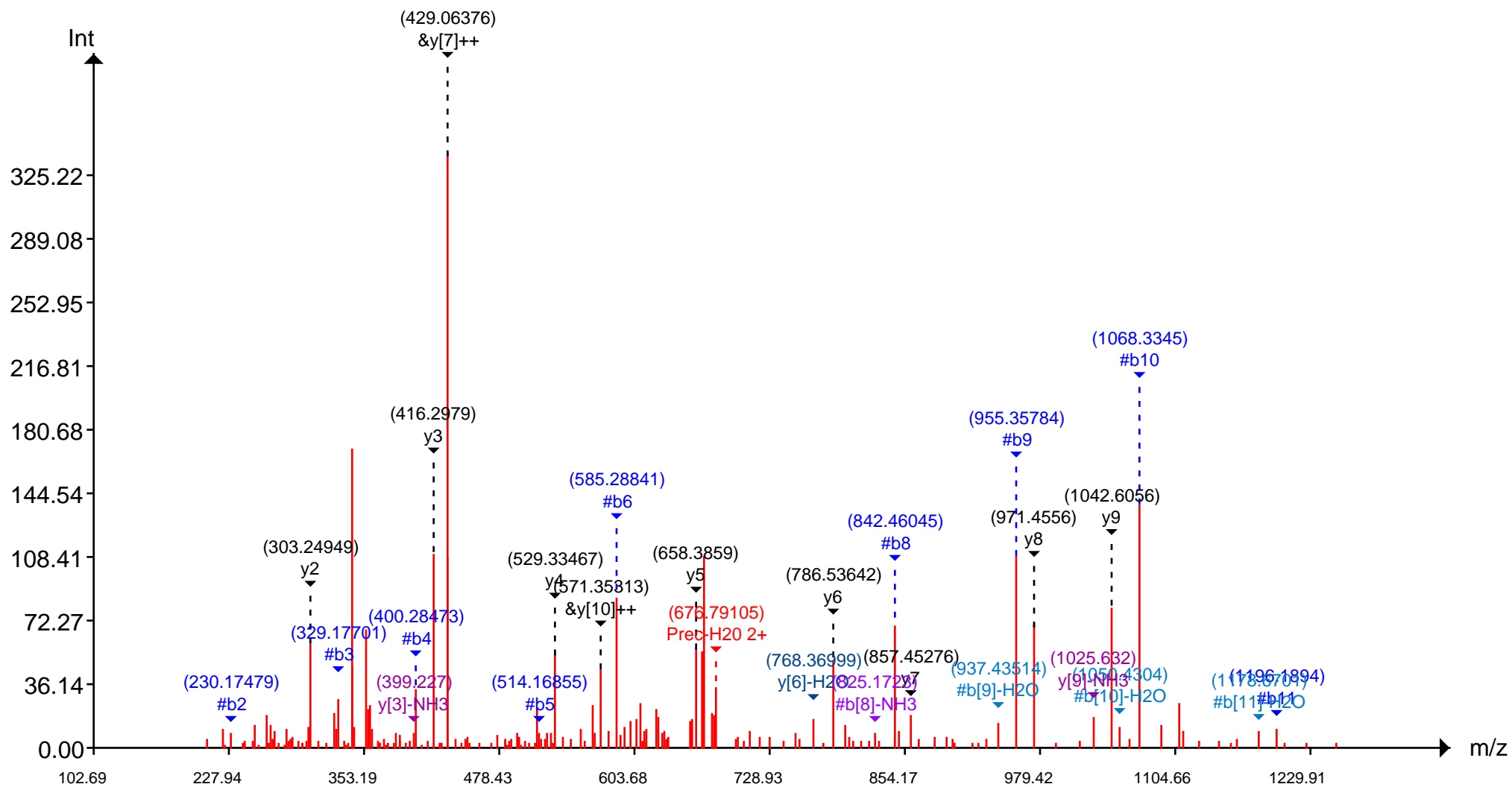
Confidence level 99.00%

Precursor m/z 714.4173

Identity Threshold 45.44

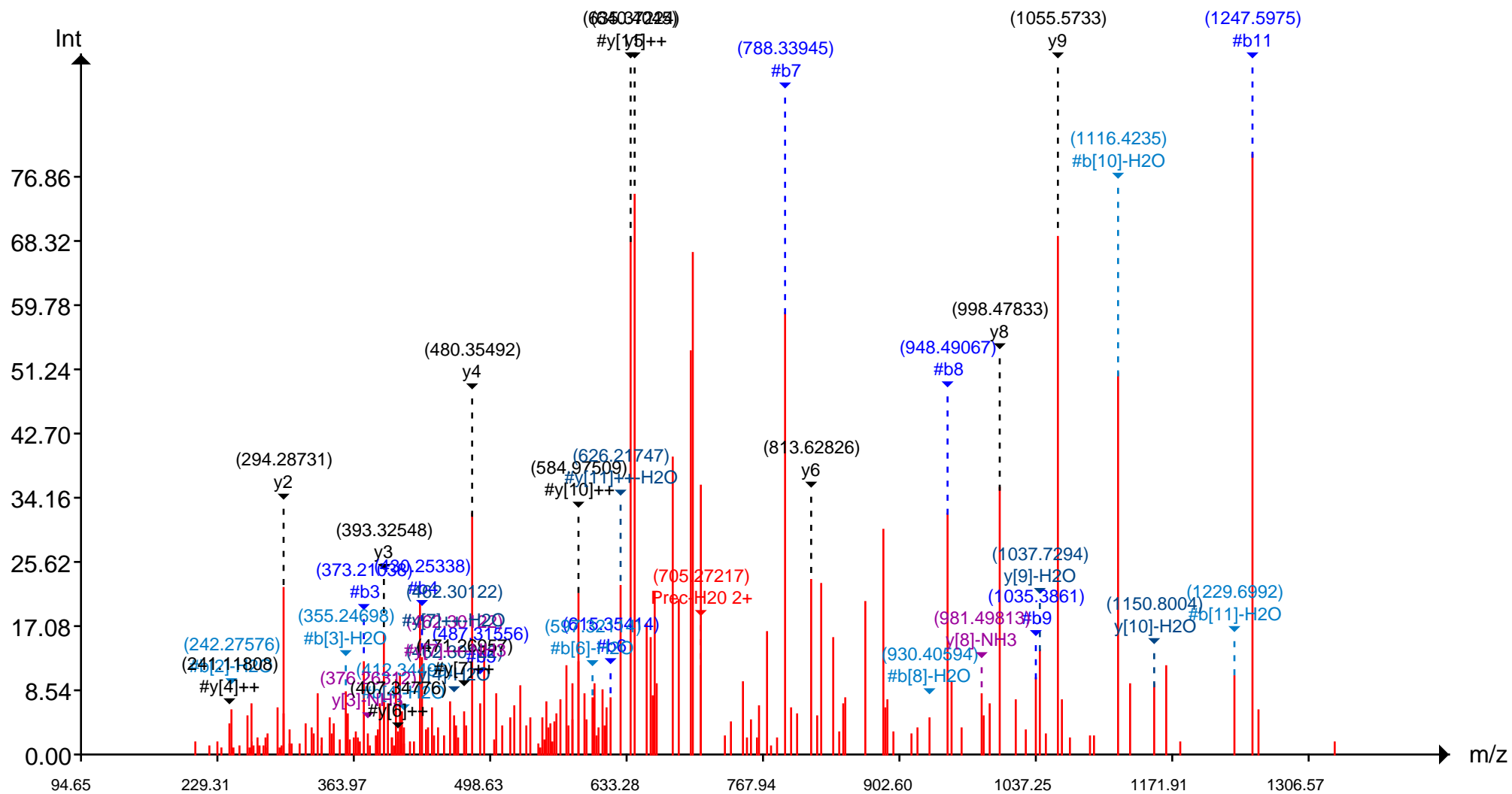
Charge 2+

Mass error 0.002009



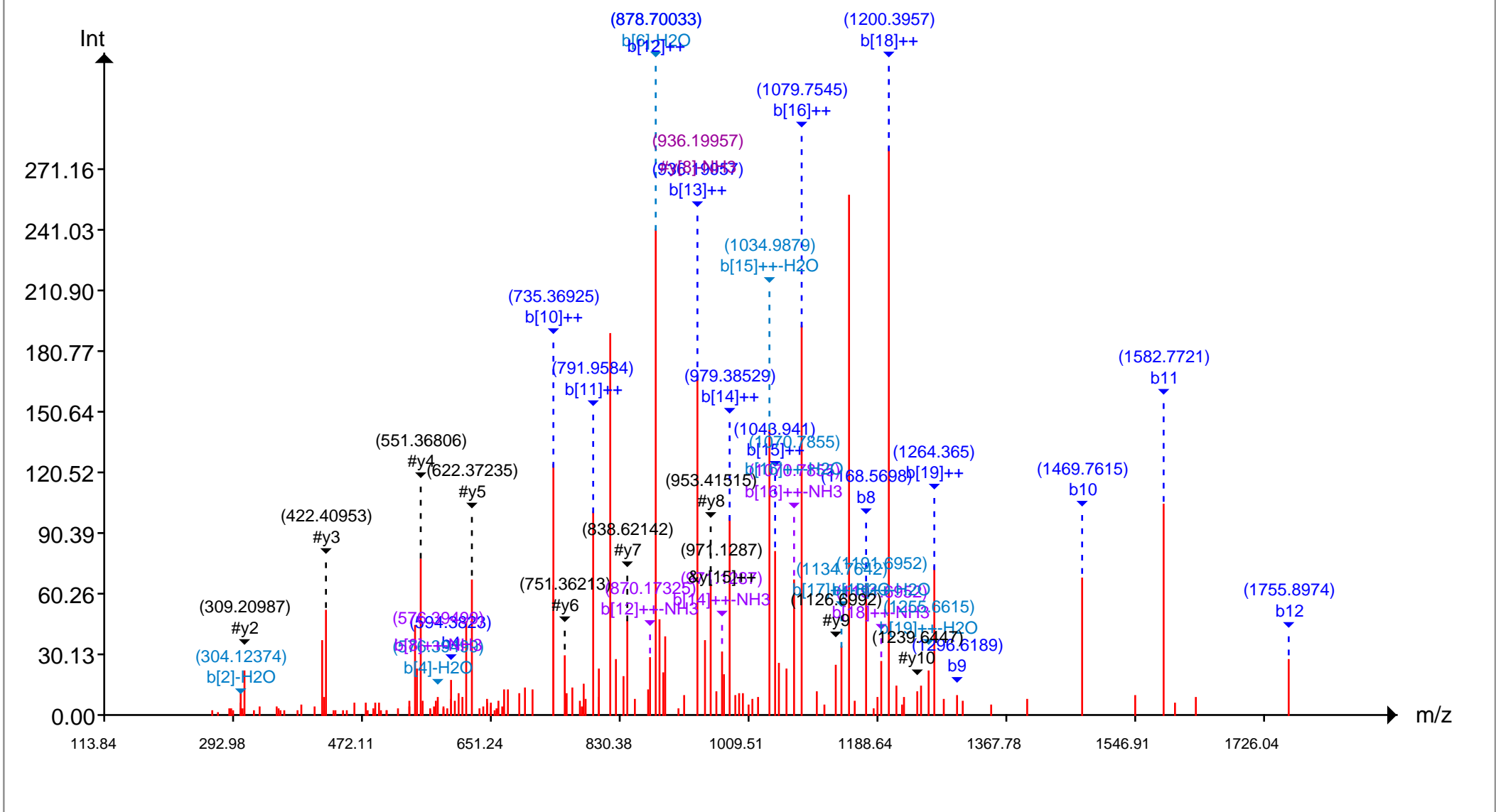
Info

Modified Sequence	AcD3-IAVANAQELLQR-COOH	Ion score	58.18
Protein	P62075	Confidence level	99.00%
Precursor m/z	685.89543	Identity Threshold	46.39
Charge	2+	Mass error	6.58E-4



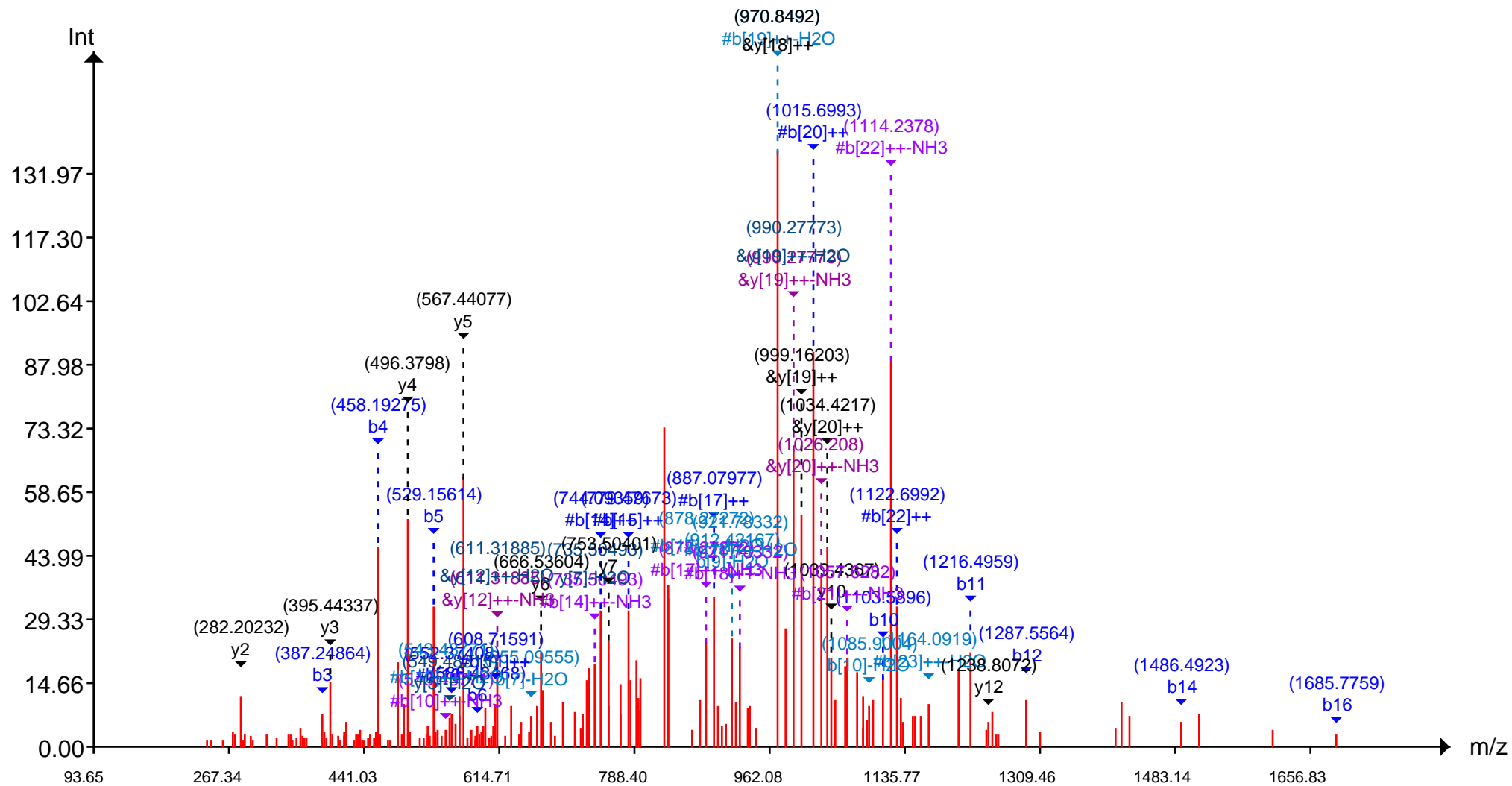
Info

Modified Sequence	AcD3-LTLGGQK<AcD3K*>C<Cmm*>SVIR<C13>-COOH	Ion score	49.26
Protein	P62962	Confidence level	99.00%
Precursor m/z	714.41731	Identity Threshold	45.44
Charge	2+	Mass error	0.002029



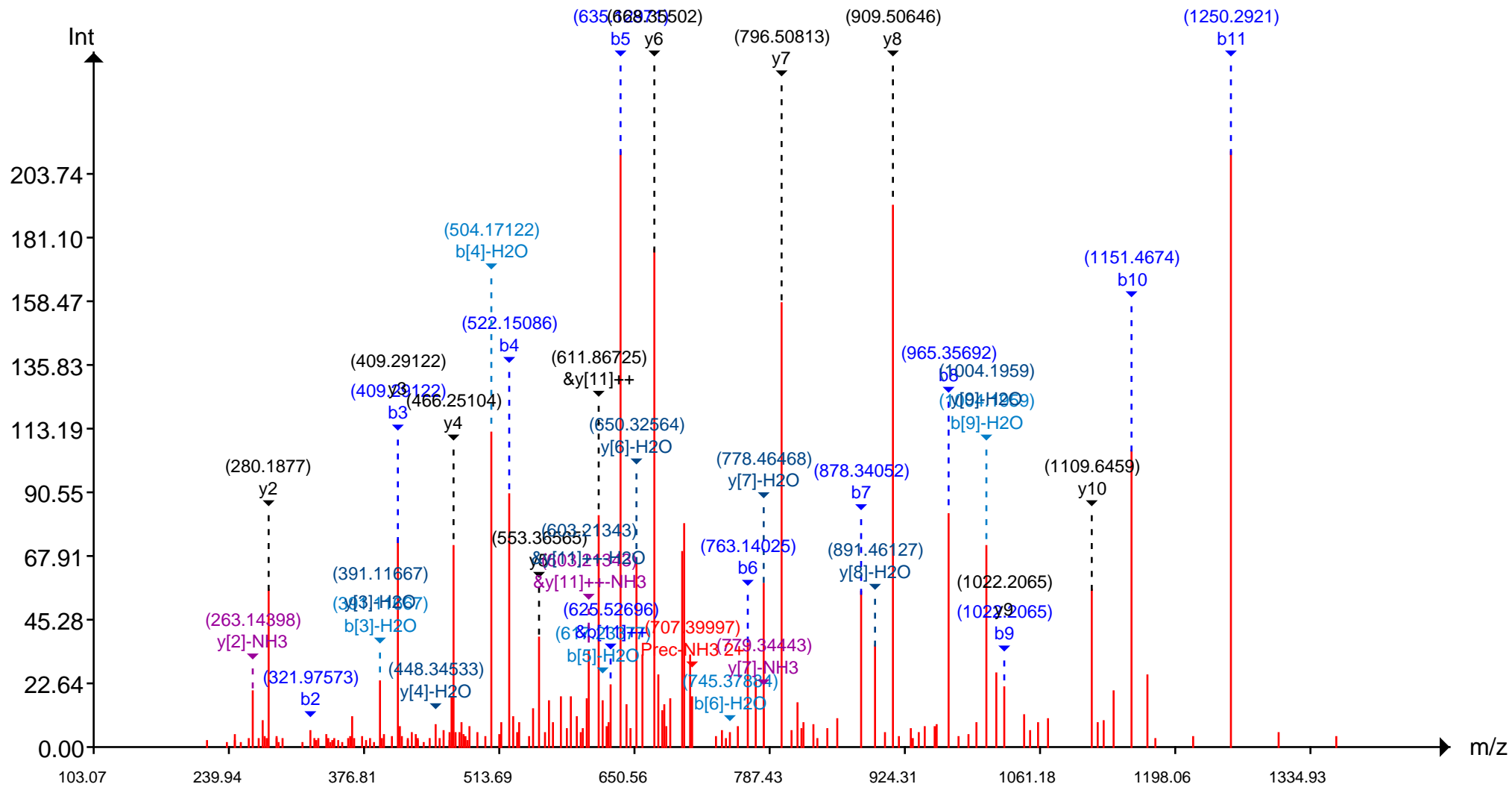
Info

Modifie...	AcD3-EM<Mox*>K<AcD3K*>VK<AcD3K*>EK<AcD3K*>VQK<AcD3K*>LK<AcD3K*>DSEAELQR<C13>-COOH	Ion score
	O55131	Confidence level
Pre...	903.51078	Identity Threshold
	3+	Mass error



Info

Modified Sequence	AcD3-VQNAAGSQEK<AcD3K*>LAQAAQSSVATITR<C13>-COOH	Ion score	48.1
Protein	P26039	Confidence level	99.00%
Precursor m/z	842.45699	Identity Threshold	47.11
Charge	3+	Mass error	0.003225



Info

Modified Sequence AcD3-YLSLLQDSGEVR<C13>-COOH

Ion score 82.26

Protein P63242

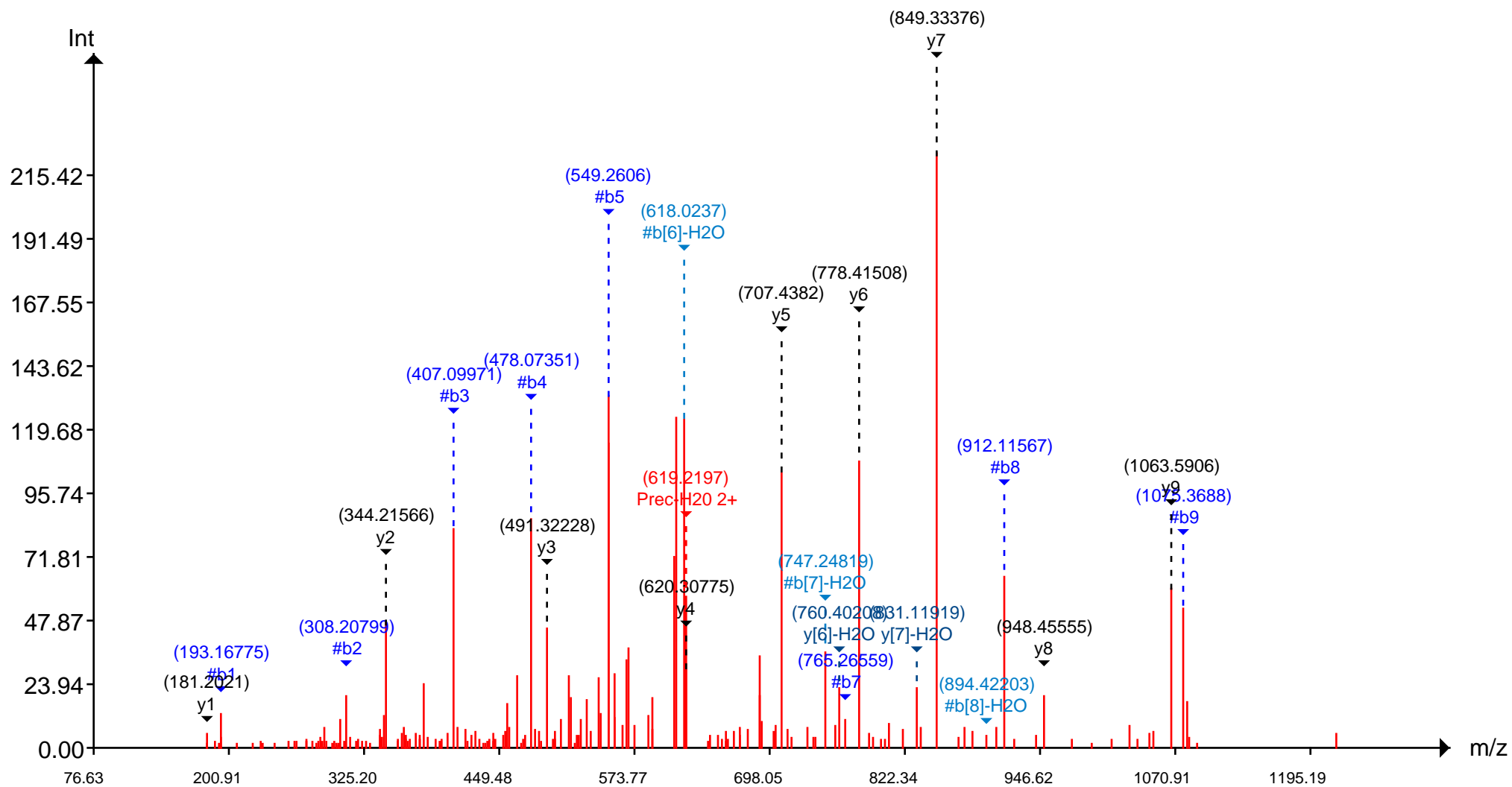
Confidence level 99.00%

Precursor m/z 715.88726

Identity Threshold 47.84

Charge 2+

Mass error 0.001209



Info

Modified Sequence AcD3-M<Mox*>DVAASEFYR<C13>-COOH

Ion score 57.43

Protein P17182, P21550, P17183

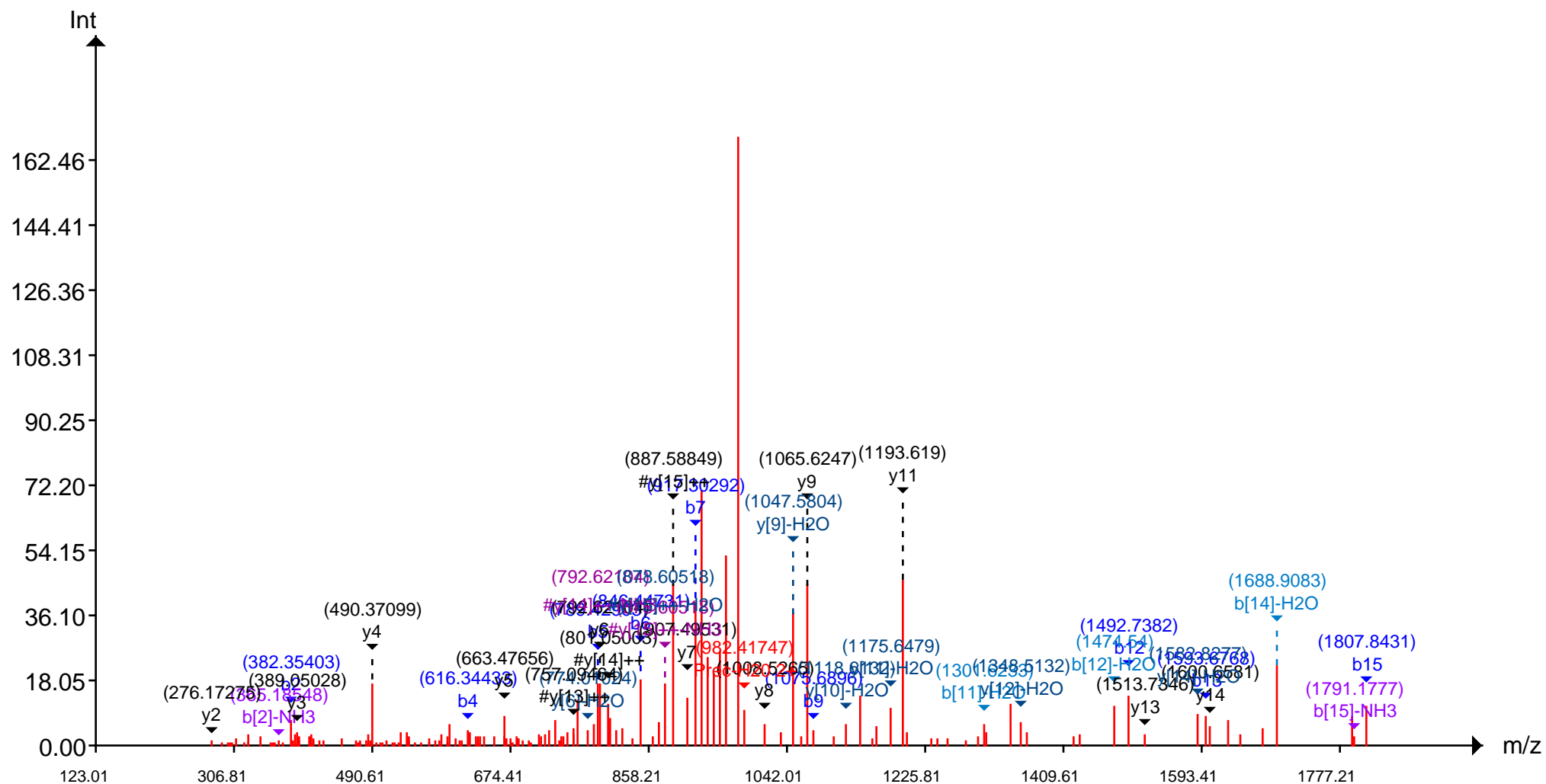
Confidence level 99.00%

Precursor m/z 628.29389

Identity Threshold 45.52

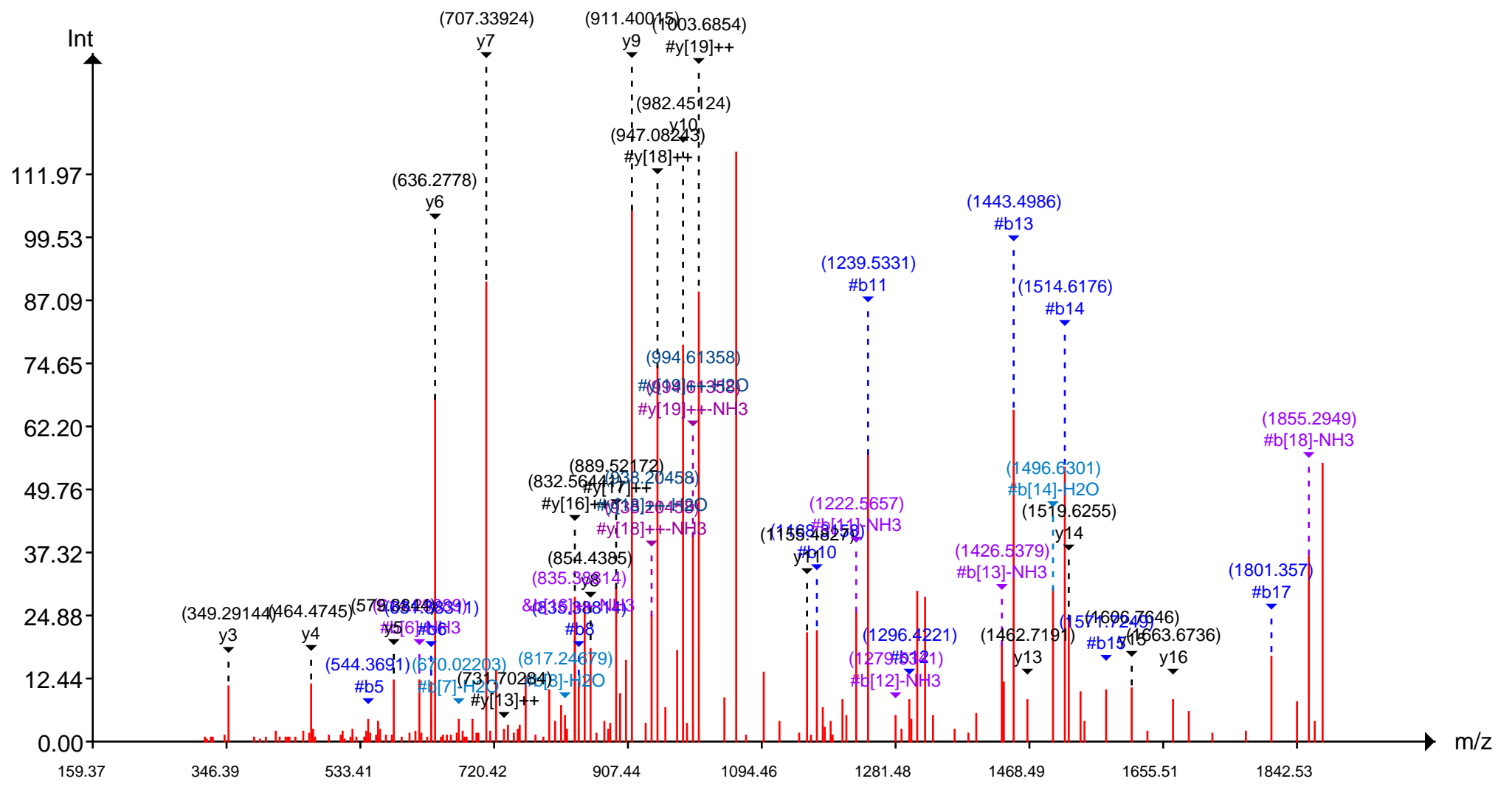
Charge 2+

Mass error 6.88E-4

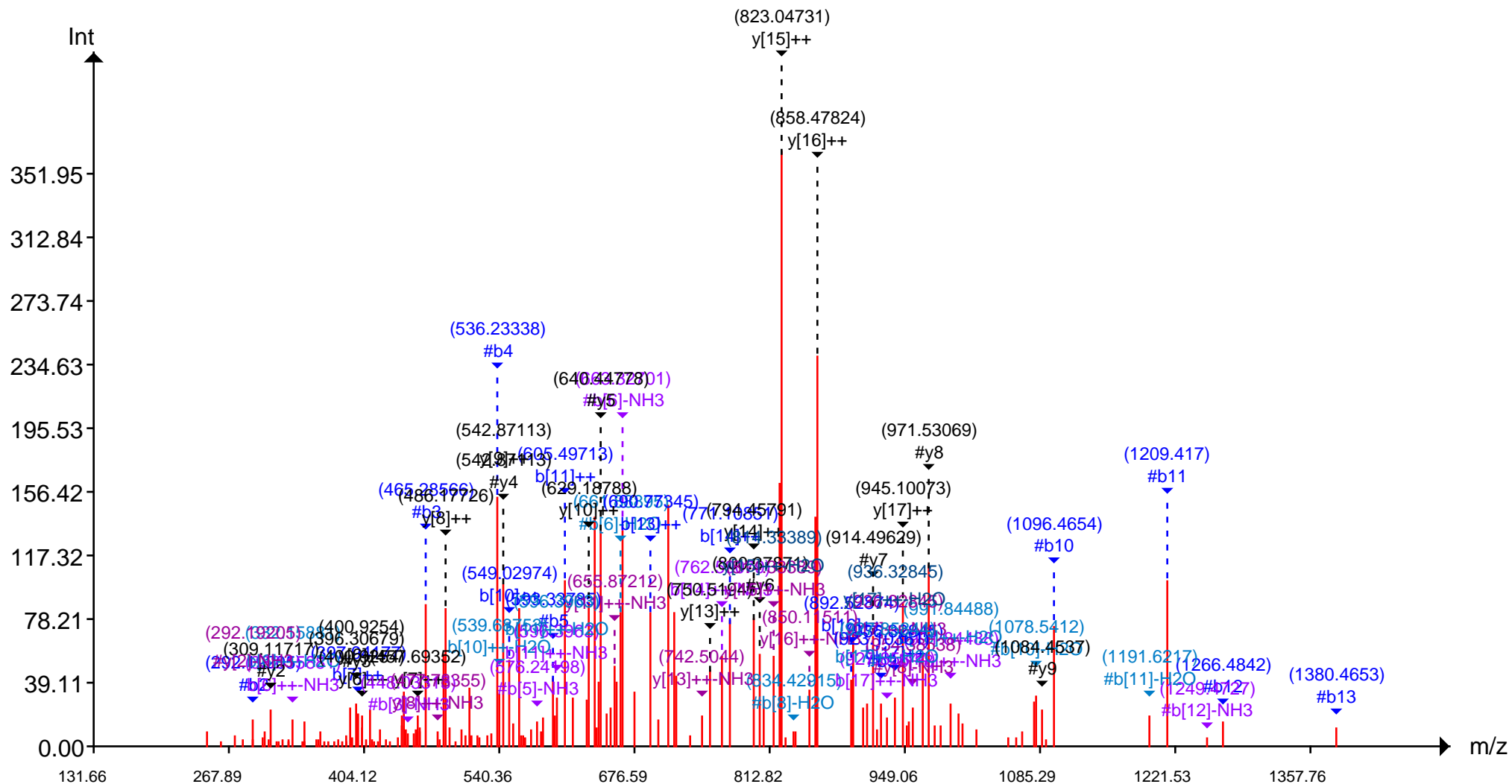


Info

Modified Sequence	AcD3-YK<AcD3K*>SM<Mox*>K<AcD3K*>GAGTDEK<AcD3K*>TLTR-COOH	Ion score	59.93
Protein	P14824	Confidence level	99.00%
Precursor m/z	991.51968	Identity Threshold	48.06
Charge	2+	Mass error	0.003232

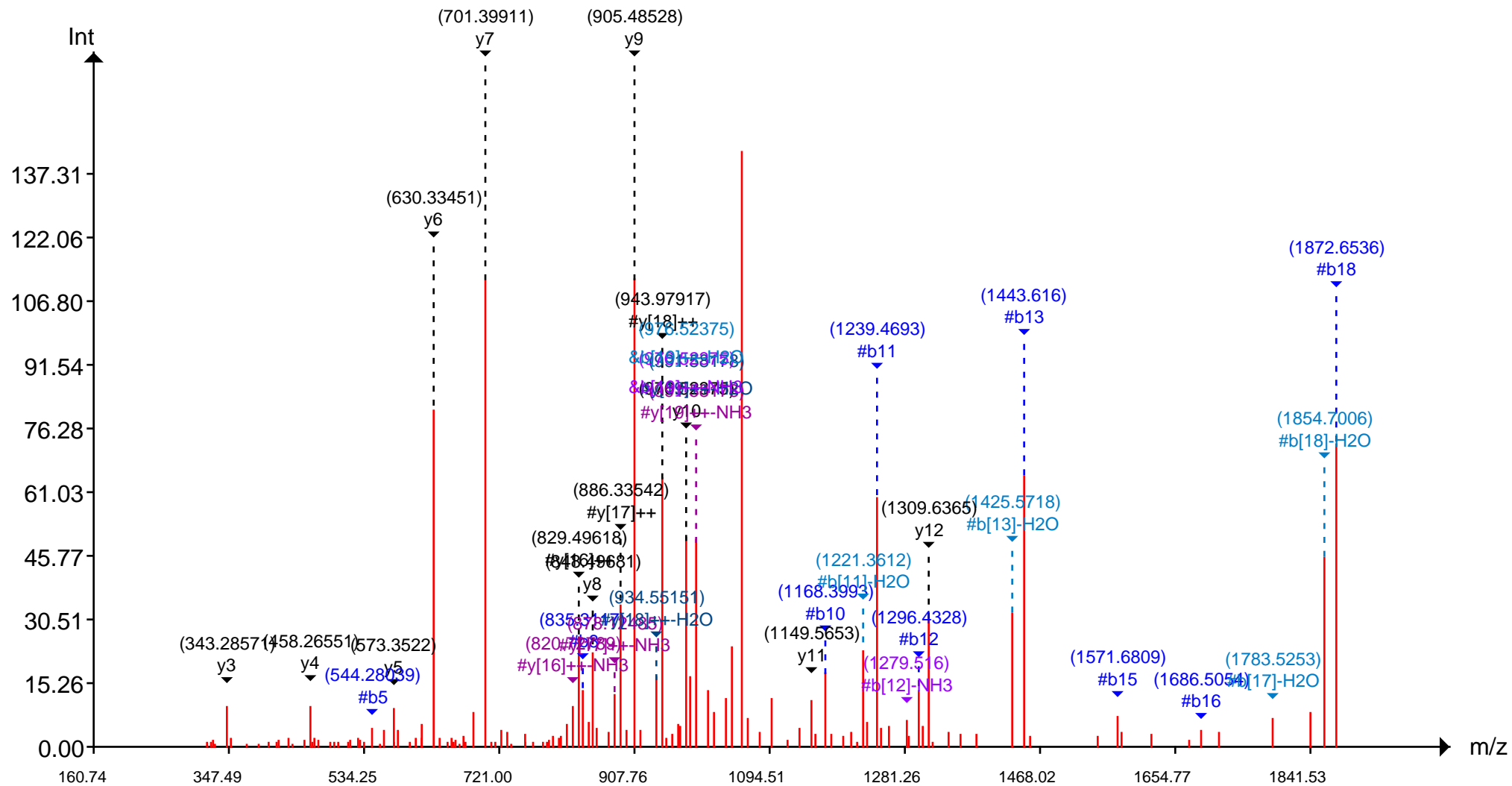


Info			
Modified Sequence	AcD3-VIDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH	Ion score	77.1
Protein	P63260	Confidence level	99.00%
Precursor m/z	1075.4984	Identity Threshold	45.05
Charge	2+	Mass error	0.003977



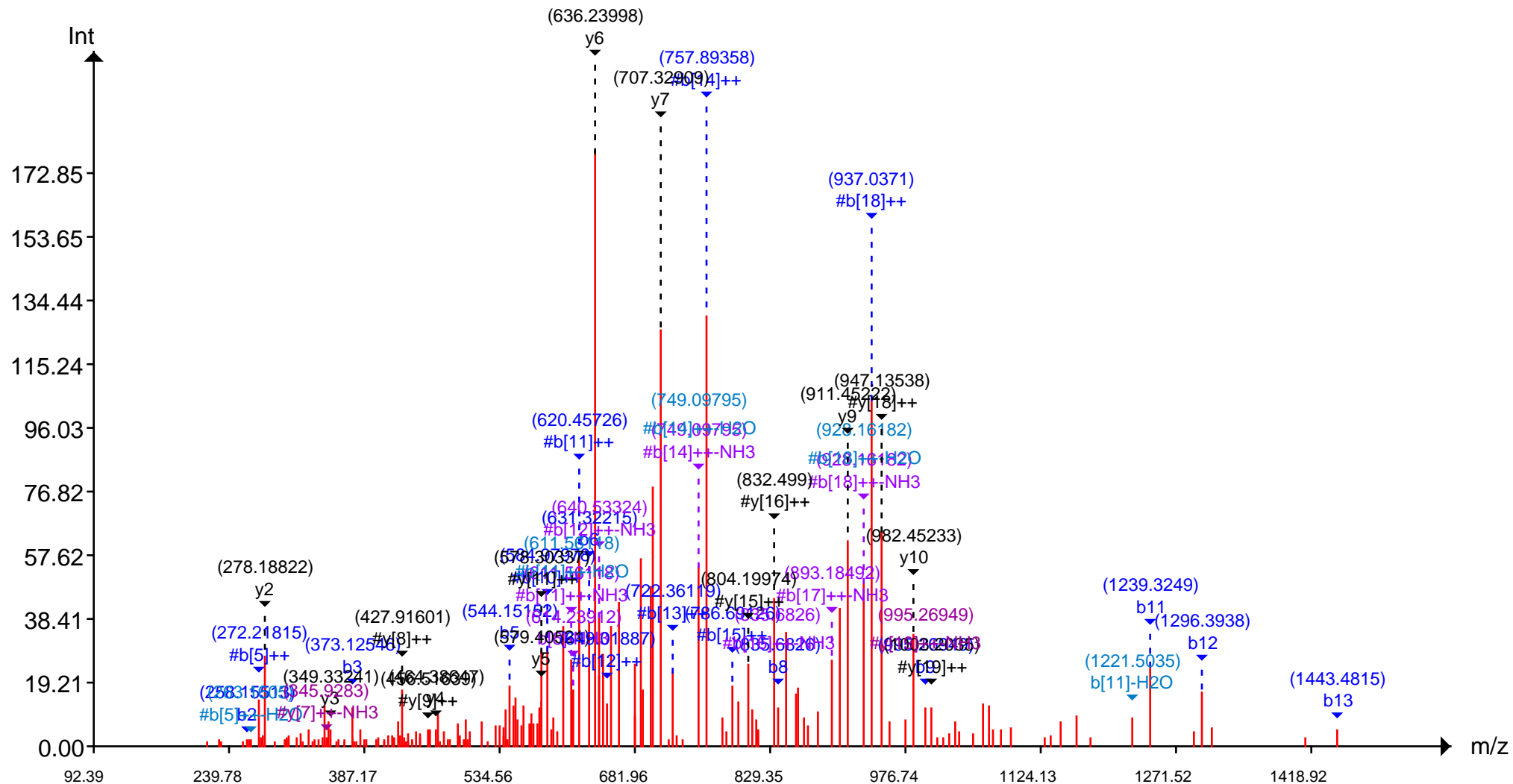
Info

Modified Sequence	AcD3-FVK<AcD3K*>AGSDGAK<AcD3K*>IGNC<Cmm*>PFSQR<C13>-COOH	Ion score	56.77
Protein	Q9Z1Q5	Confidence level	99.00%
Precursor m/z	727.37866	Identity Threshold	48.07
Charge	3+	Mass error	3.67E-4



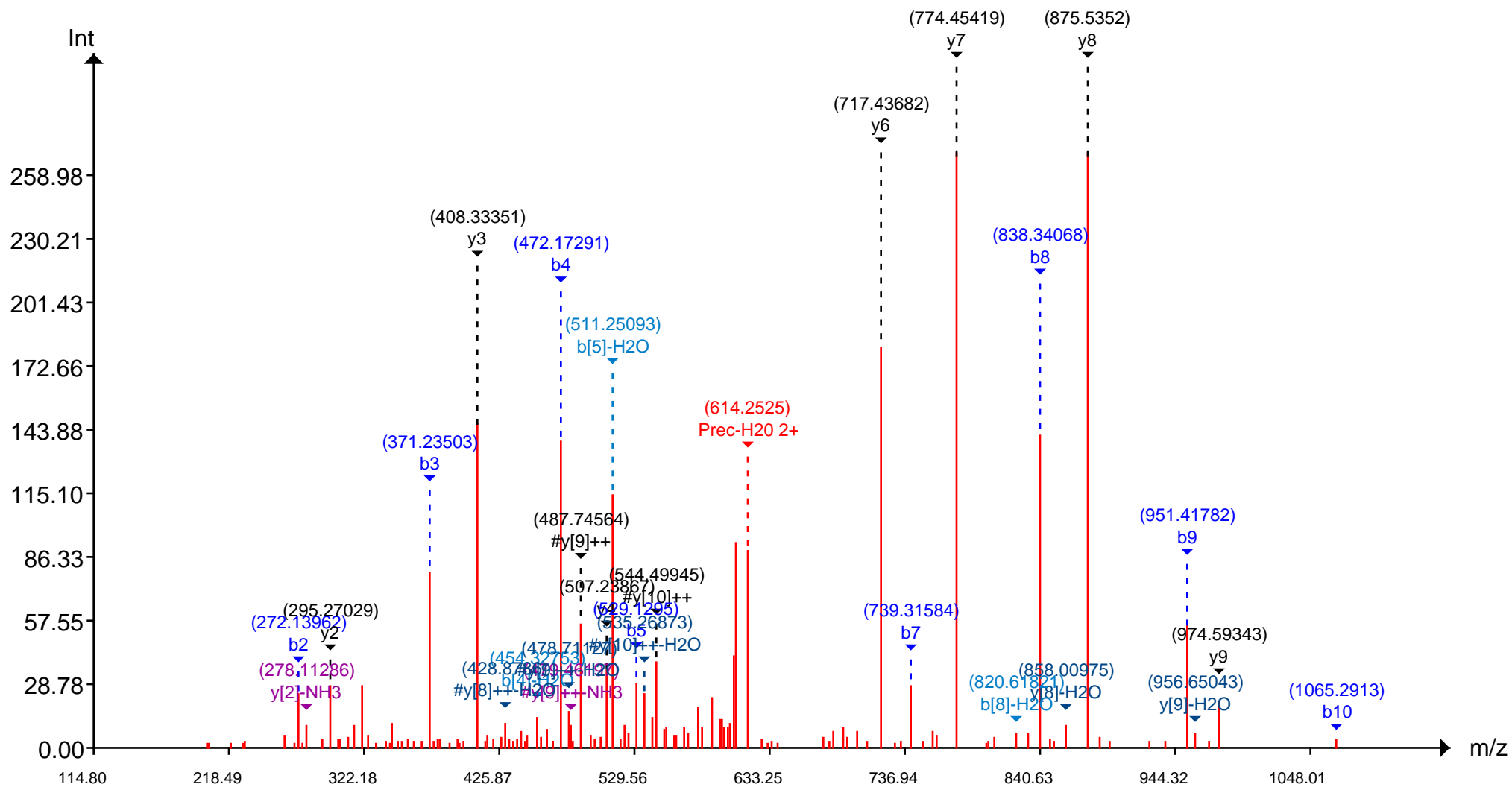
Info

Modified Sequence	AcD3-VIDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR-COOH	Ion score	63.58
Protein	P63260	Confidence level	99.00%
Precursor m/z	1072.4883	Identity Threshold	44.11
Charge	2+	Mass error	0.003918



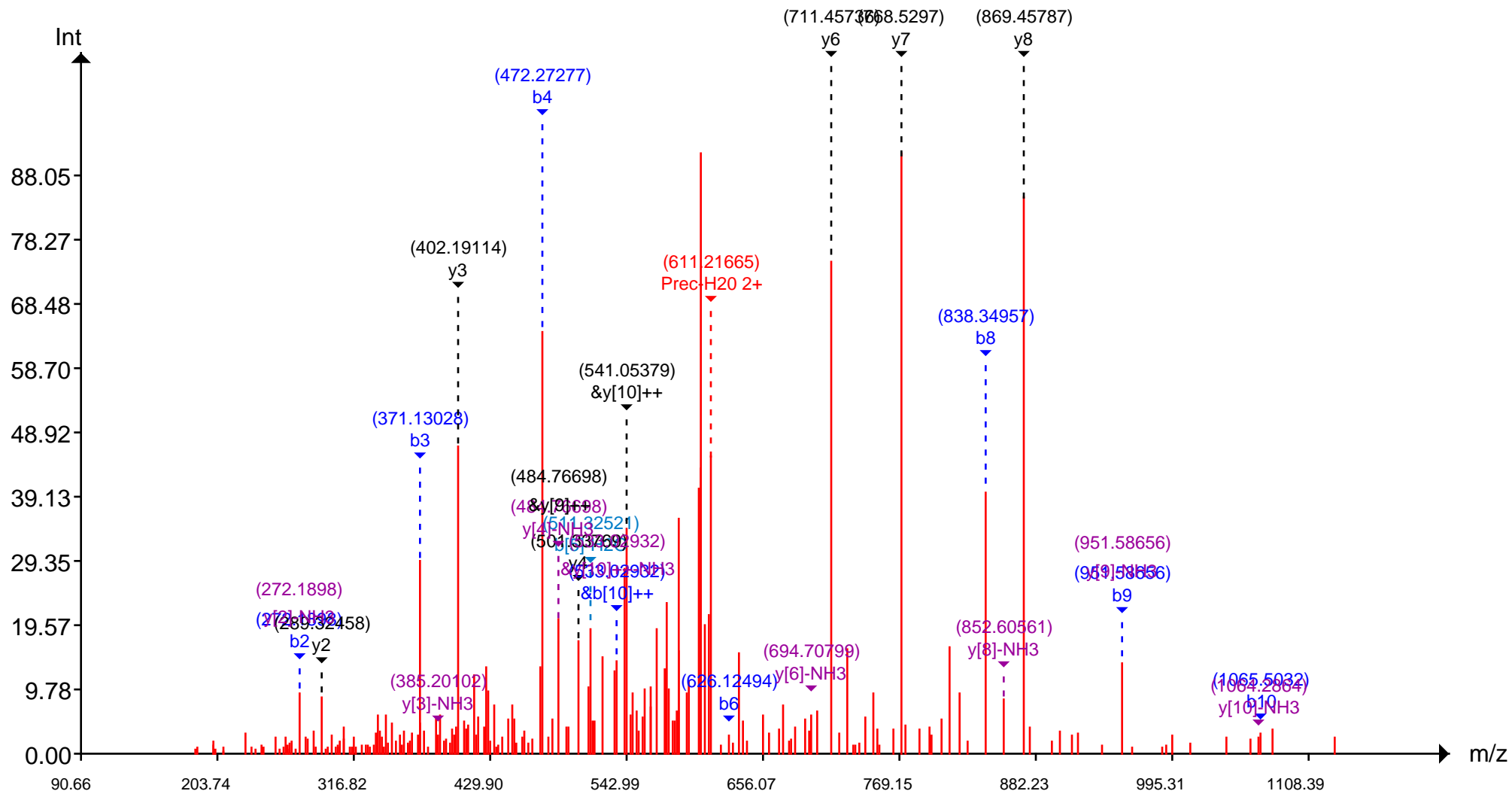
Info

Modified Sequence	AcD3-VIDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH	Ion score	46.91
Protein	P63260	Confidence level	99.00%
Precursor m/z	717.3339	Identity Threshold	44.94
Charge	3+	Mass error	0.001601



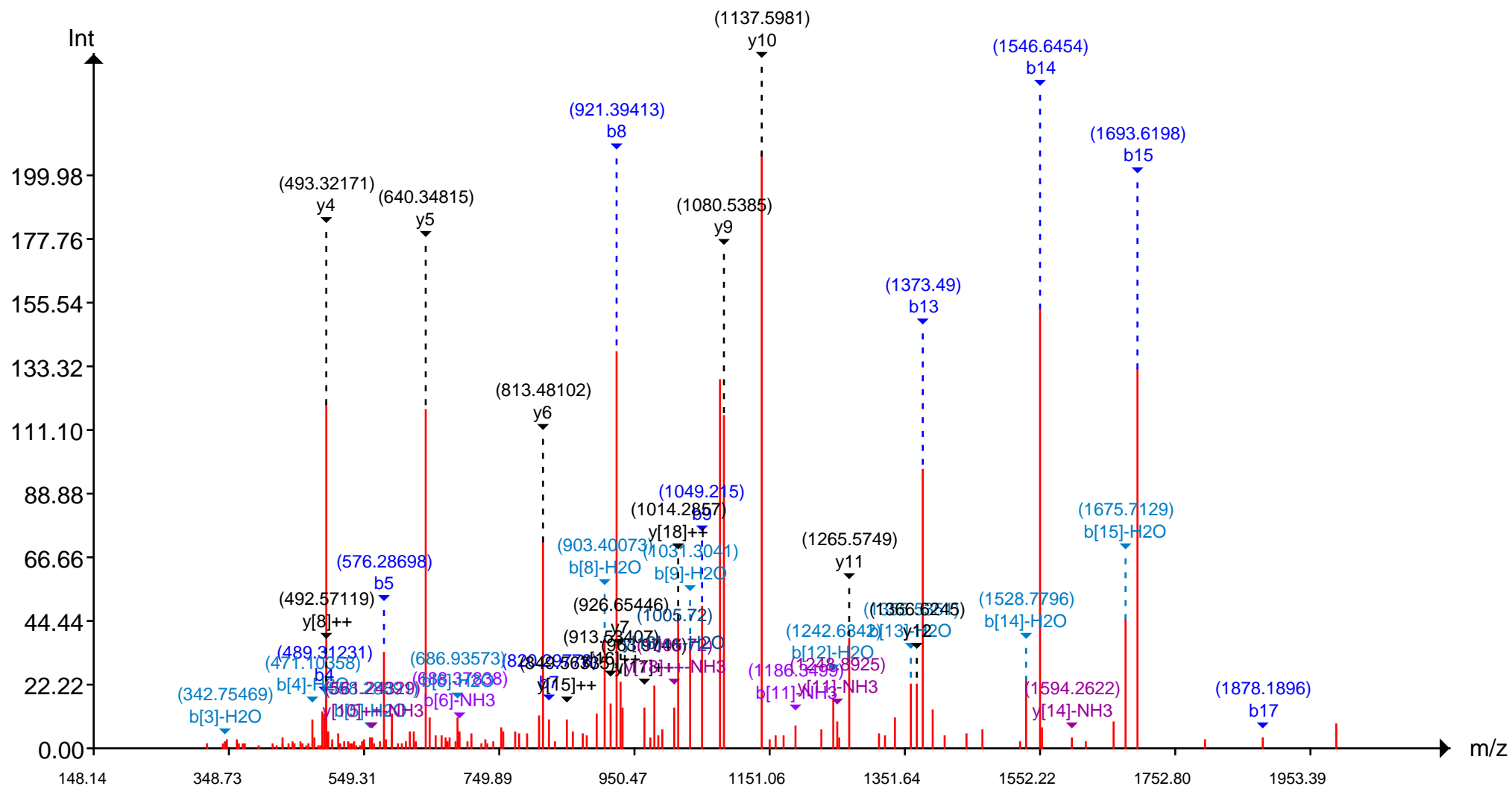
Info

Modified Sequence	AcD3-LLVTGPLVINR<C13>-COOH	Ion score	67.14
Protein	P47911	Confidence level	99.00%
Precursor m/z	623.40721	Identity Threshold	37.90
Charge	2+	Mass error	7.5E-4



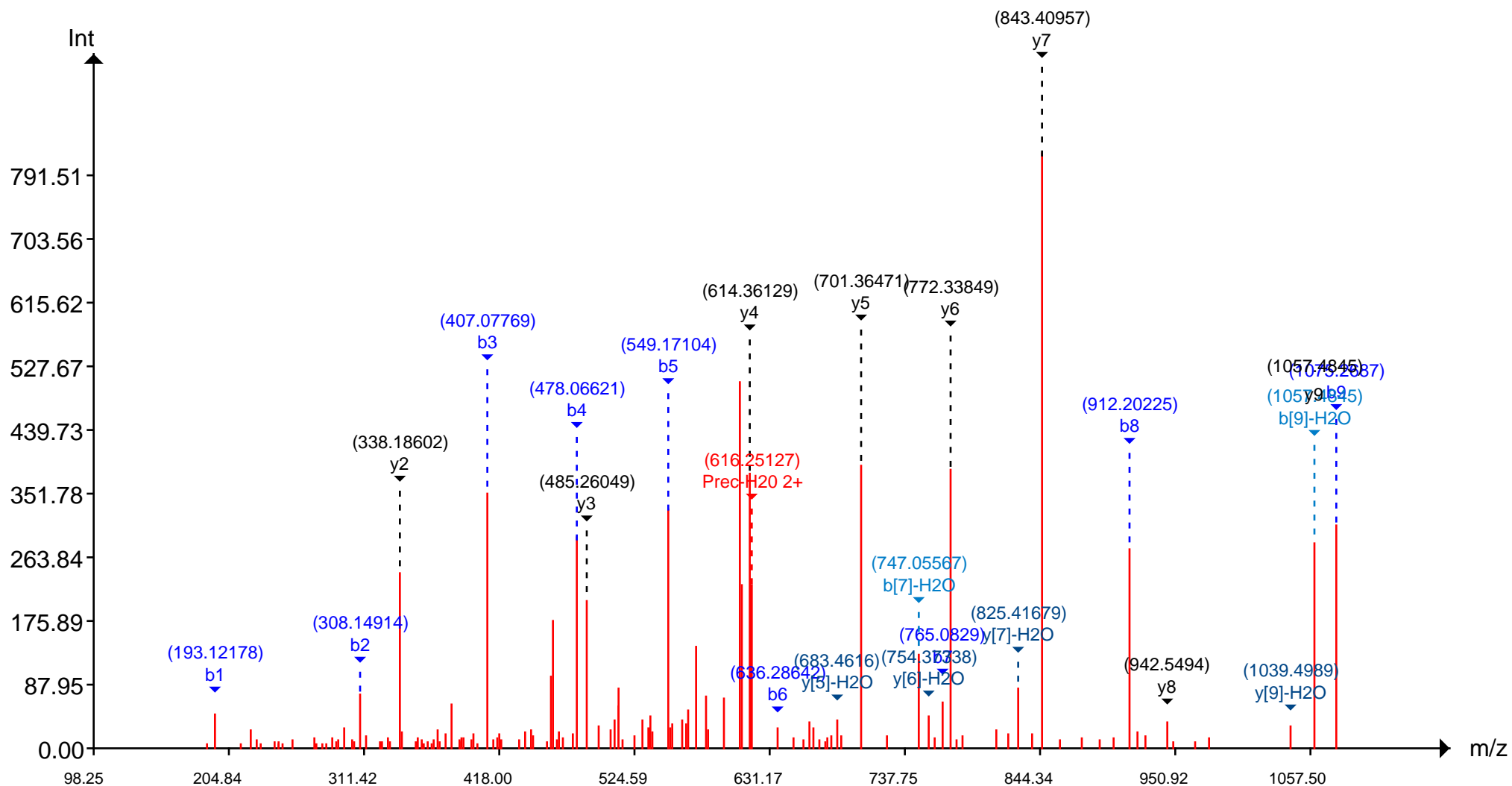
Info

Modified Sequence	AcD3-LLVTGPLVNR-COOH	Ion score	41.51
Protein	P47911	Confidence level	99.00%
Precursor m/z	620.39705	Identity Threshold	39.10
Charge	2+	Mass error	5.72E-4



Info

Modified Sequence	AcD3-LTTQSEDTQGPKLK<AcD3K*>M<Mox*>PSQR<C13>-COOH	Ion score	52.71
Protein	O54824	Confidence level	99.00%
Precursor m/z	1093.5537	Identity Threshold	47.82
Charge	2+	Mass error	0.002928



Info

Modified Sequence AcD3-M<Mox*>DVAASEFYR-COOH

Ion score 68.39

Protein P17182, P21550, P17183

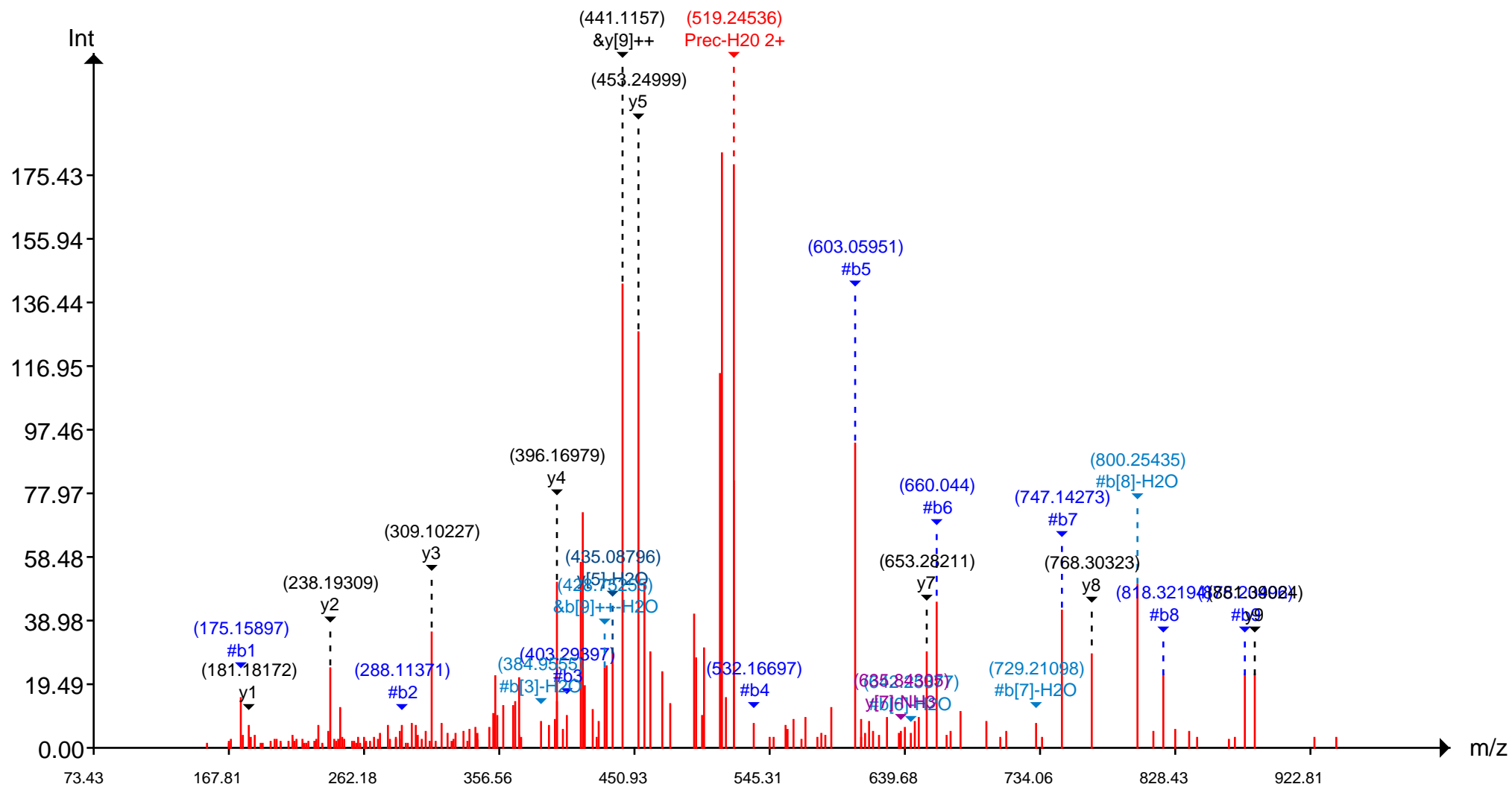
Confidence level 99.00%

Precursor m/z 625.28381

Identity Threshold 44.16

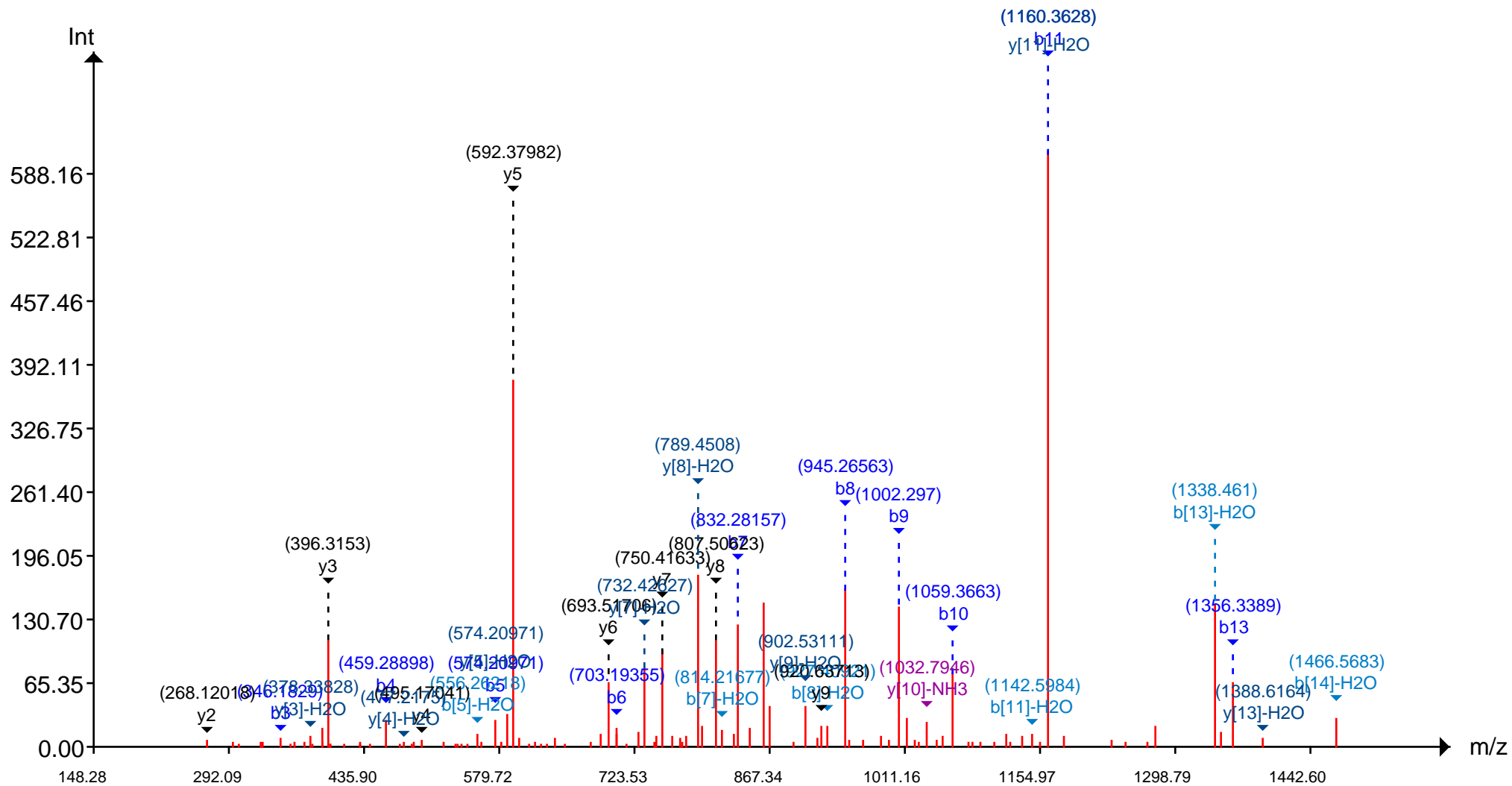
Charge 2+

Mass error 6.69E-4



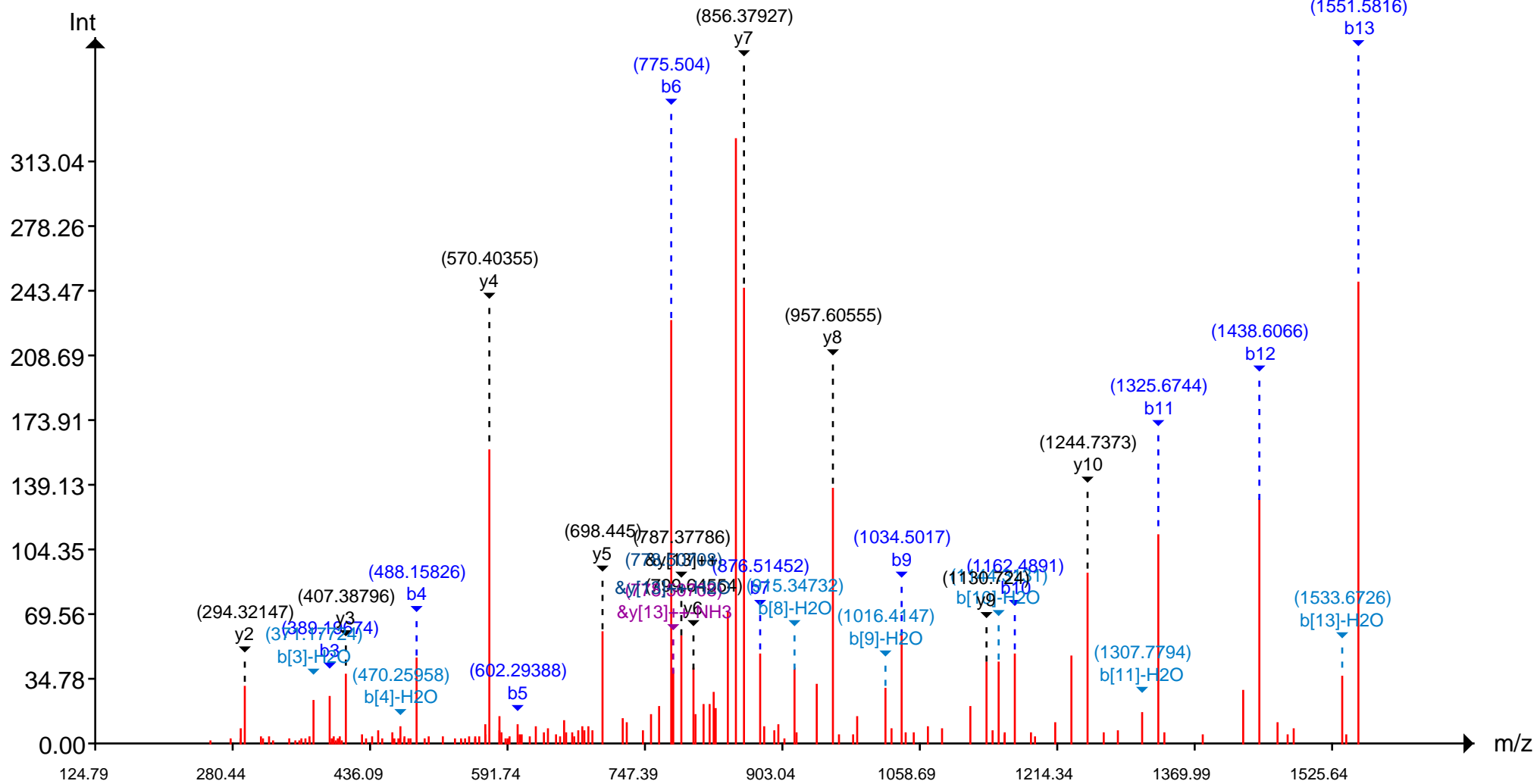
Info

Modified Sequence	AcD3-ELDEAGSAGR<C13>-COOH	Ion score	53.61
Protein	Q9QZK7	Confidence level	99.00%
Precursor m/z	528.26059	Identity Threshold	46.69
Charge	2+	Mass error	6.7E-5



Info

Modified Sequence	AcD3-TAQLDEELGGTPVQSR<C13>-COOH	Ion score	74.72
Protein	P13020	Confidence level	99.00%
Precursor m/z	876.45205	Identity Threshold	47.99
Charge	2+	Mass error	0.002326



Info

Modified Sequence AcD3-LTEVNK<AcD3K*>TGTQYLLR<C13>-COOH

Ion score 82.17

Protein Q60854

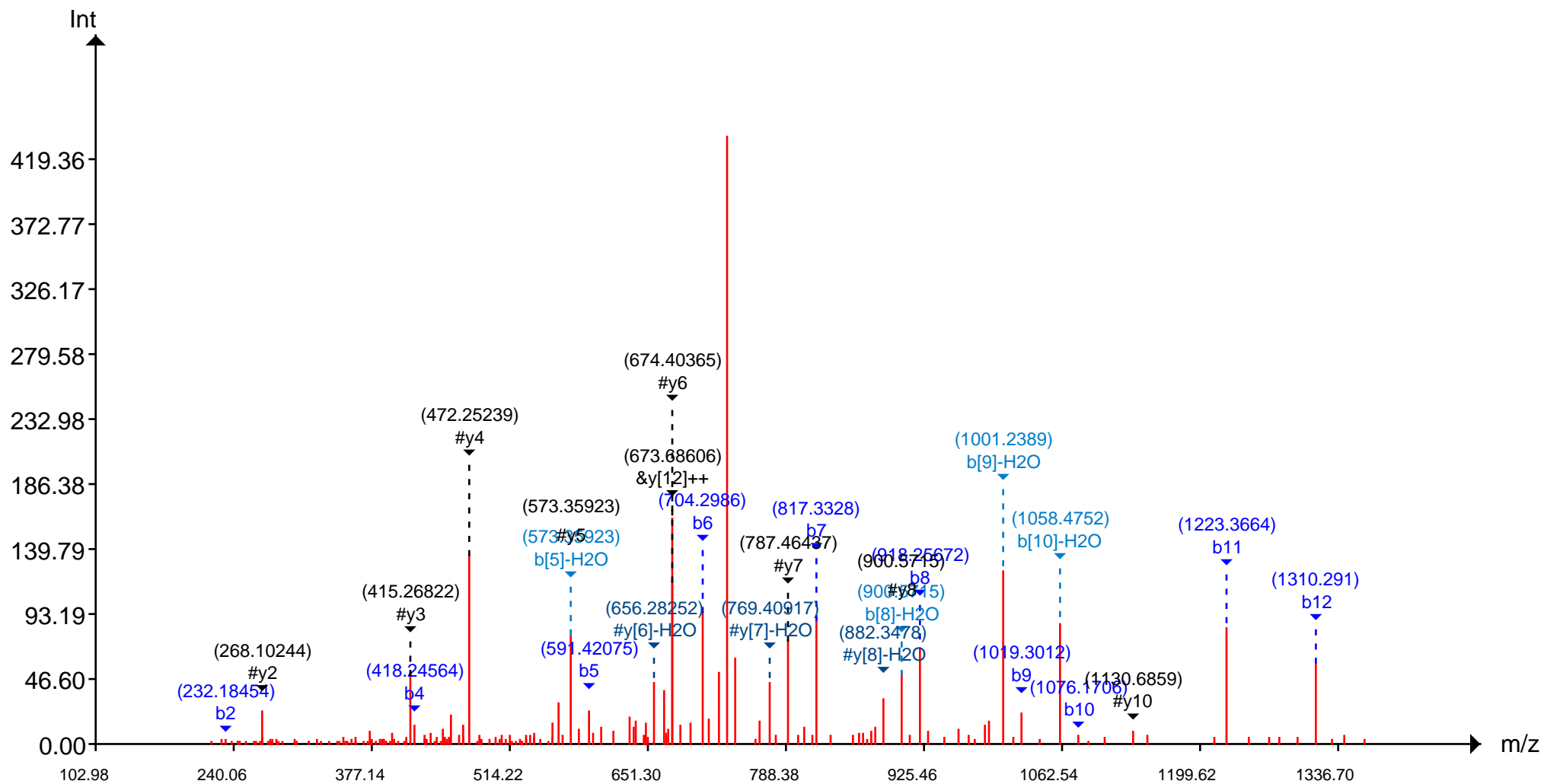
Confidence level 99.00%

Precursor m/z 866.49739

Identity Threshold 45.65

Charge 2+

Mass error 0.00214



Info

Modified Sequence AcD3-VSEGK<AcD3K*>ILTTGFSR<C13>-COOH

Ion score 66.42

Protein O89053

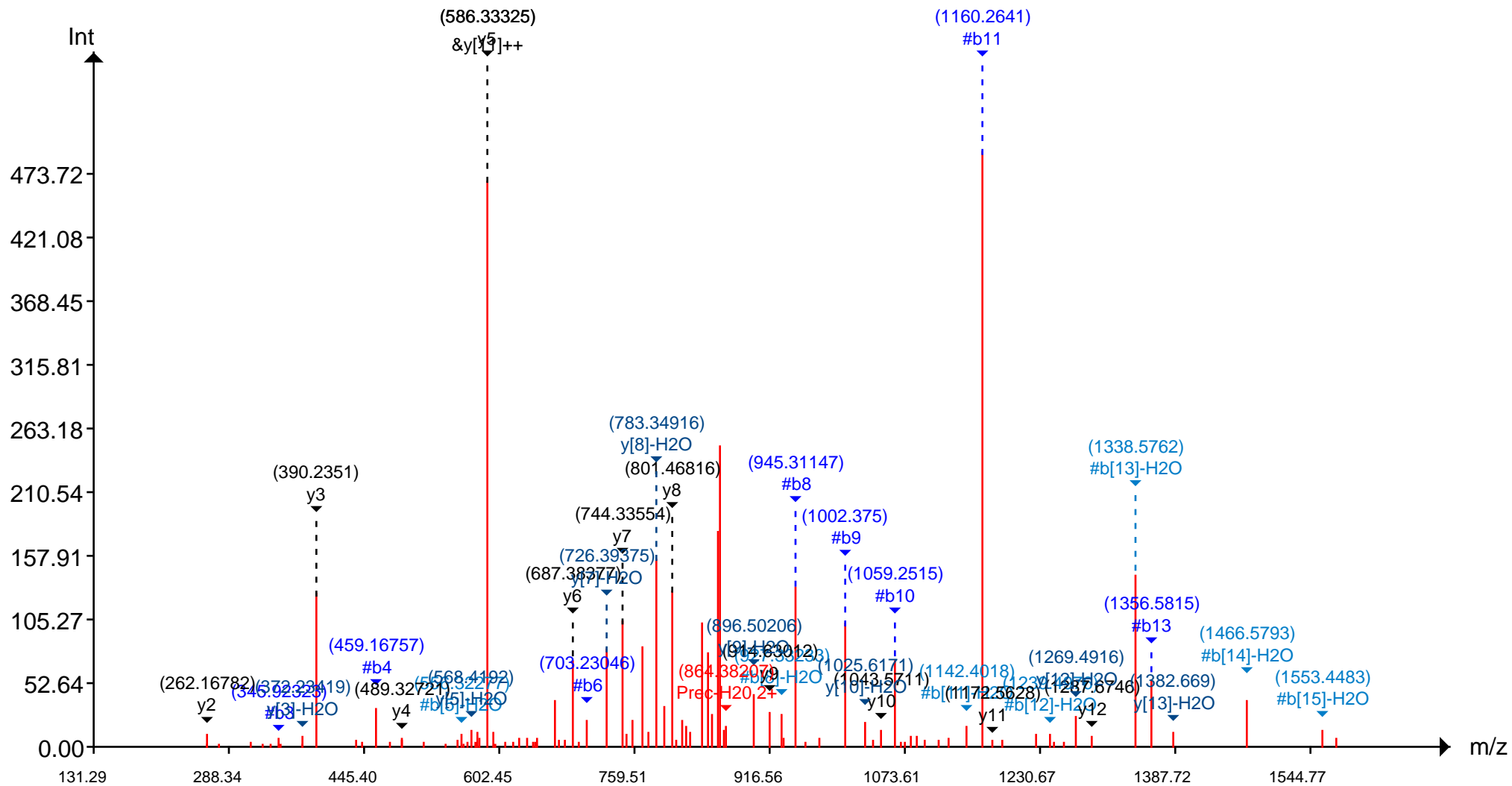
Confidence level 99.00%

Precursor m/z 745.92491

Identity Threshold 46.51

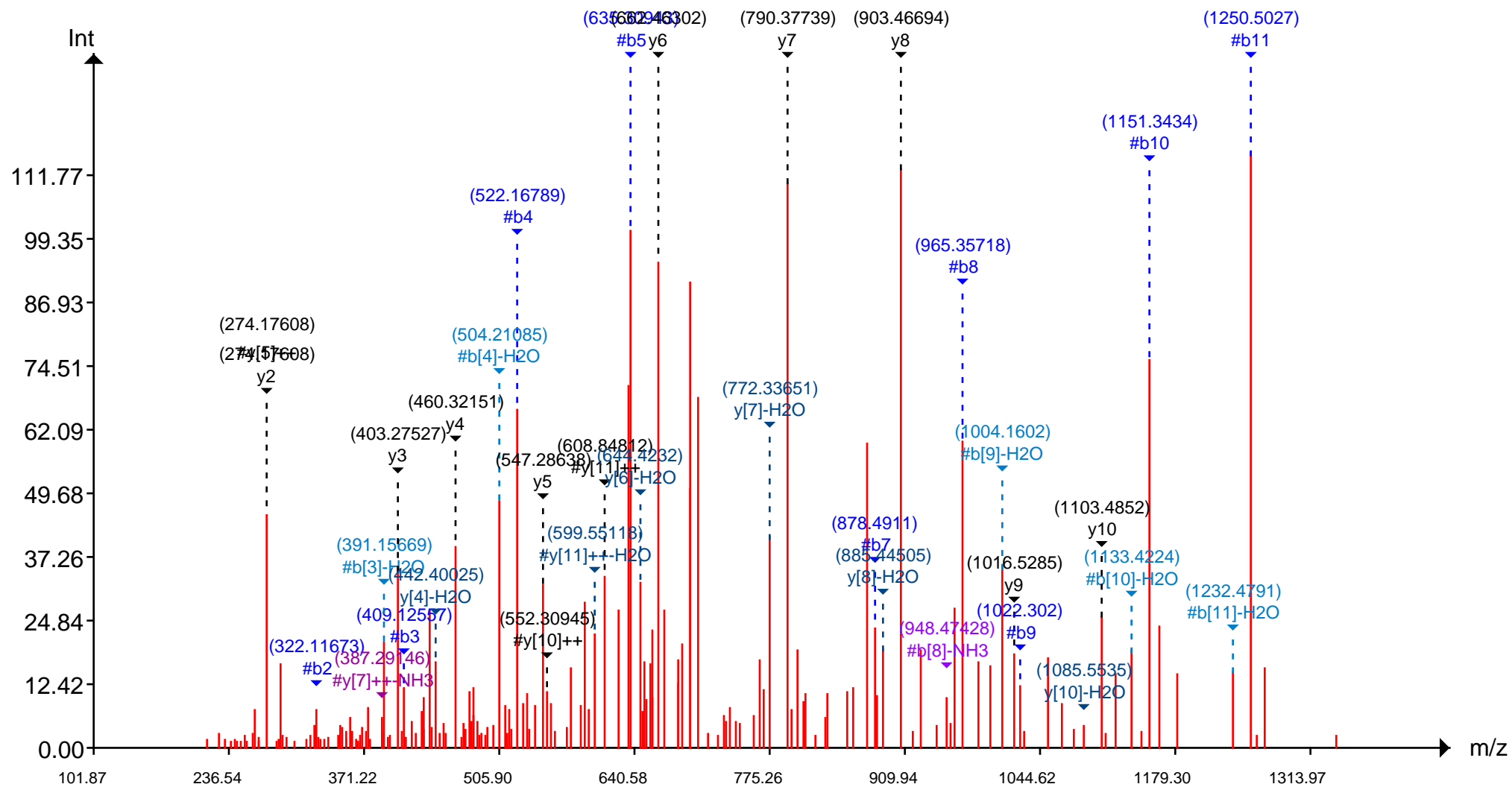
Charge 2+

Mass error -1.96E-4



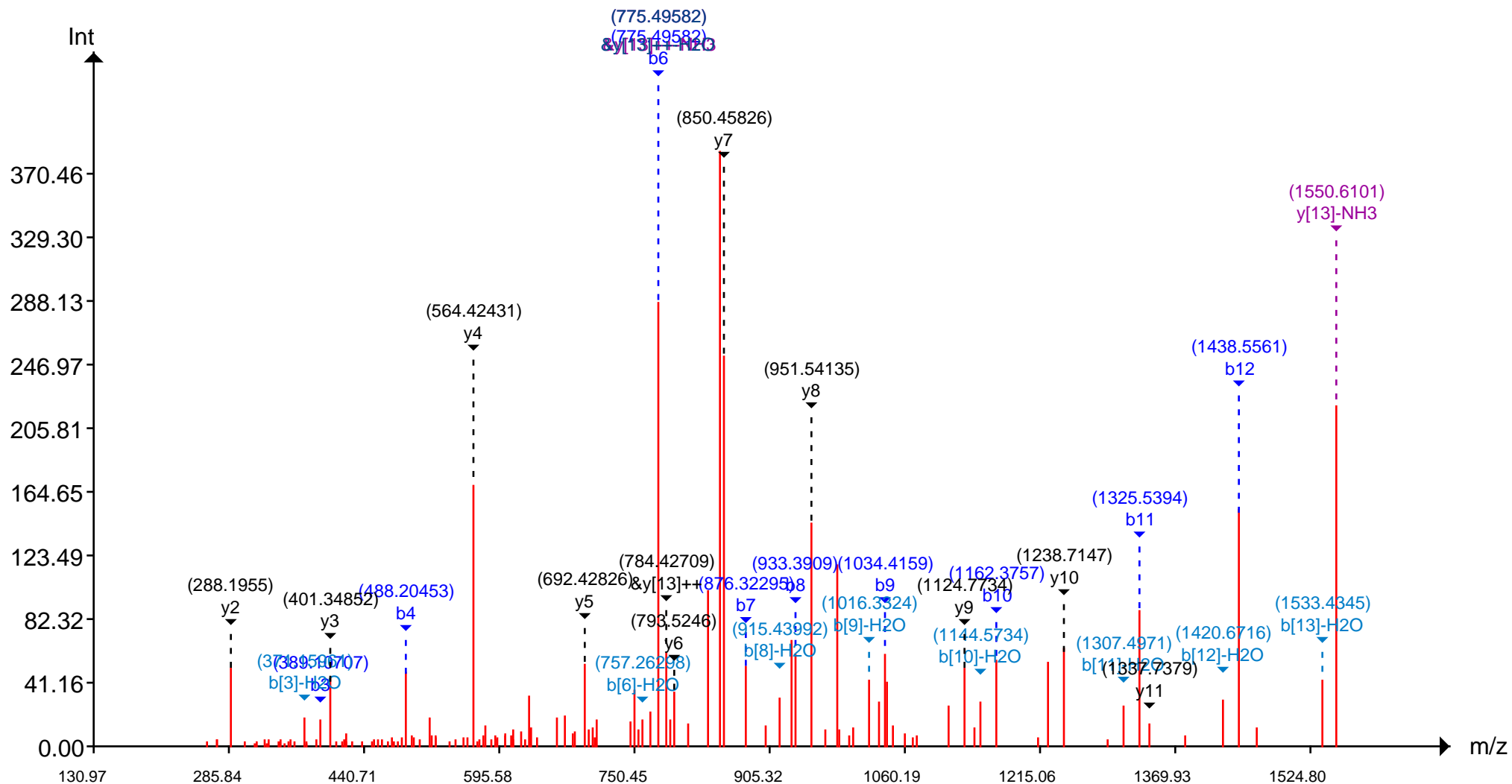
Info

Modified Sequence	AcD3-TAQLDEELGGTPVQSR-COOH	Ion score	83.91
Protein	P13020	Confidence level	99.00%
Precursor m/z	873.44195	Identity Threshold	47.93
Charge	2+	Mass error	0.002267



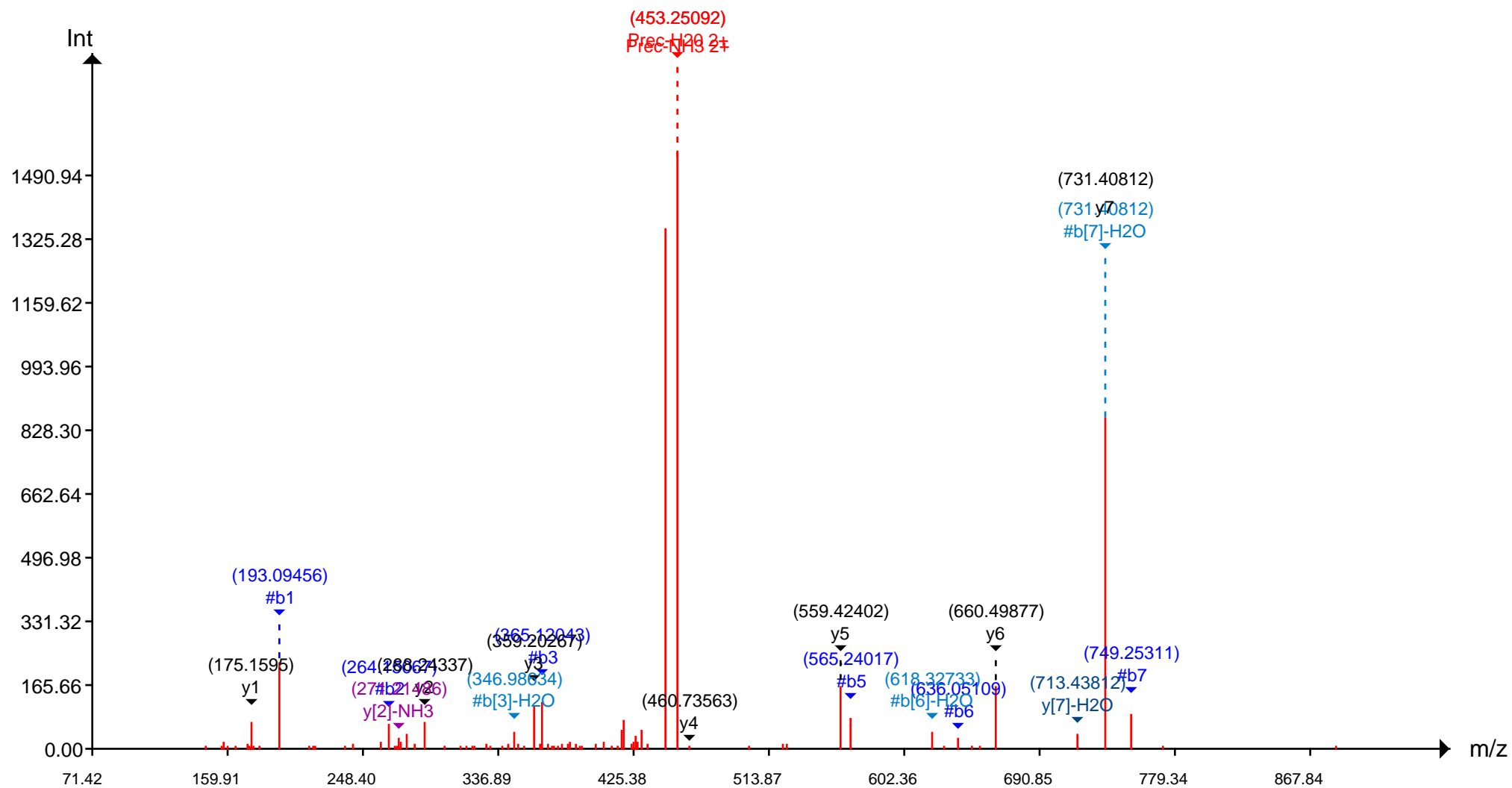
Info

Modified Sequence	AcD3-YLSLLQDSGEVR-COOH	Ion score	69.8
Protein	P63242	Confidence level	99.00%
Precursor m/z	712.87725	Identity Threshold	48.03
Charge	2+	Mass error	0.001331



Info

Modified Sequence	AcD3-LTEVNK<AcD3K*>TGTQYLLR-COOH	Ion score	65.59
Protein	Q60854	Confidence level	99.00%
Precursor m/z	863.48714	Identity Threshold	46.55
Charge	2+	Mass error	0.001781



Info

Modified Sequence AcD3-M<Mox*>ATVTALR-COOH

Ion score 55.43

Protein P05064, P05063

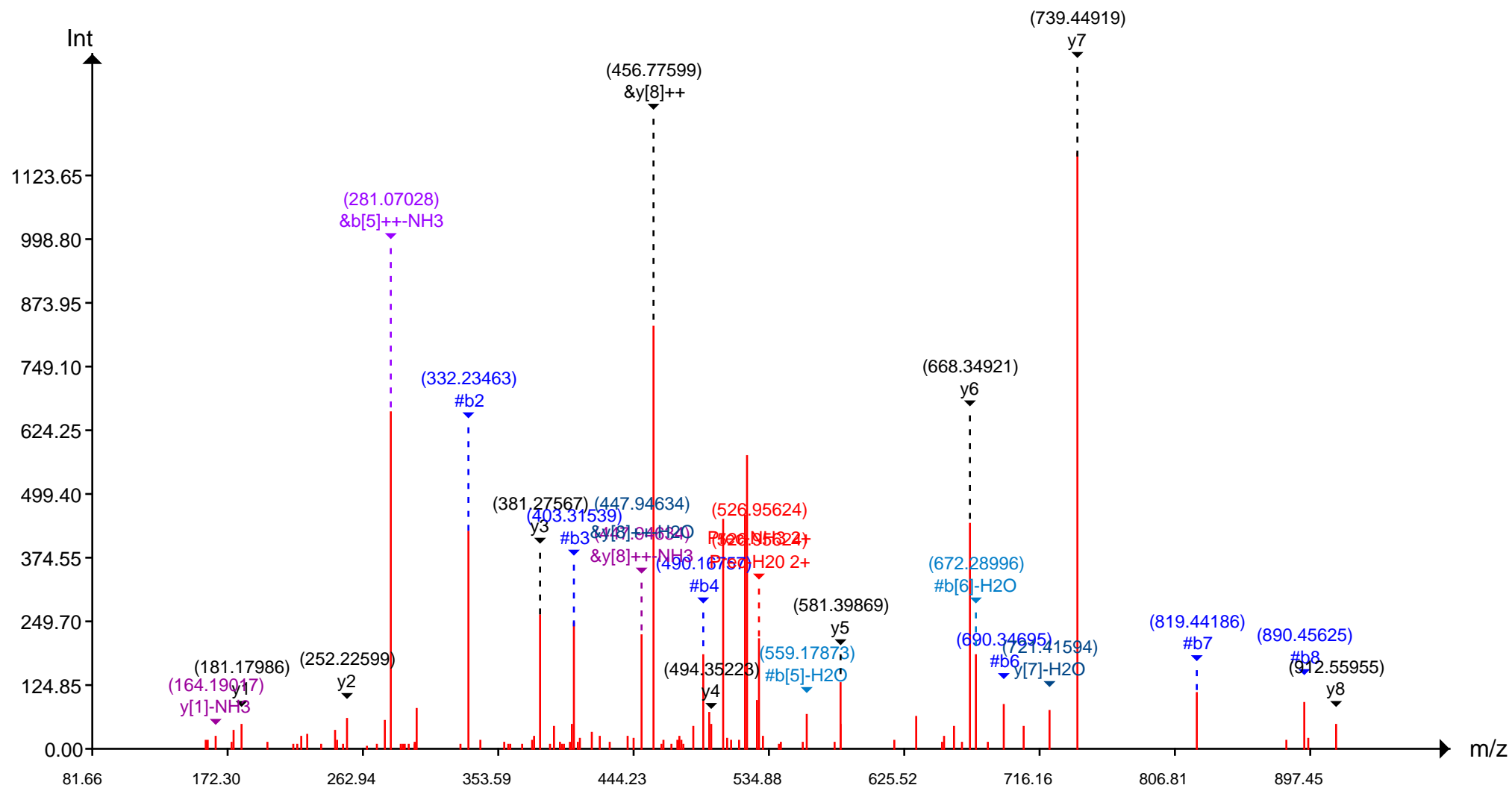
Confidence level 99.00%

Precursor m/z 462.25618

Identity Threshold 47.82

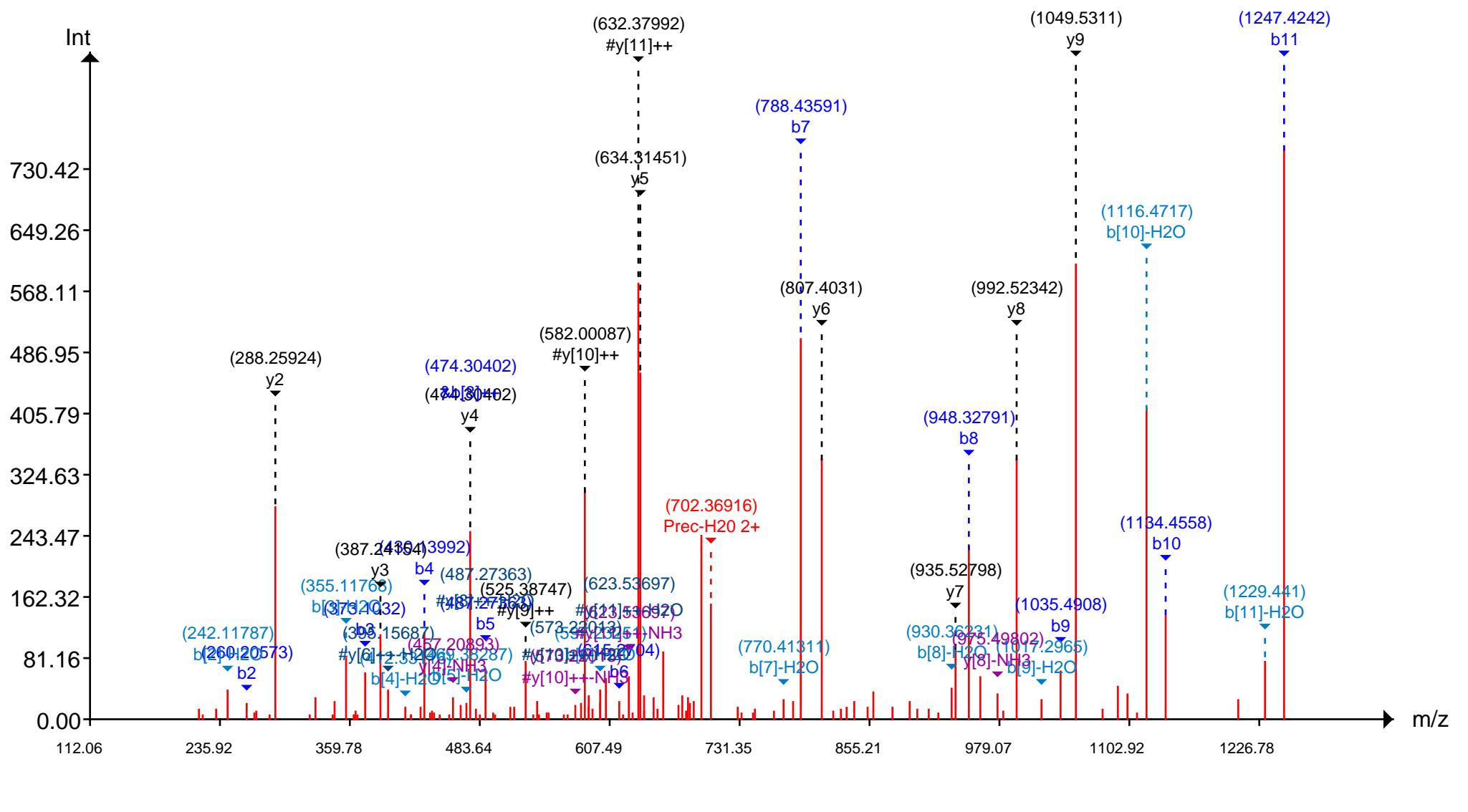
Charge 2+

Mass error -7.12E-4

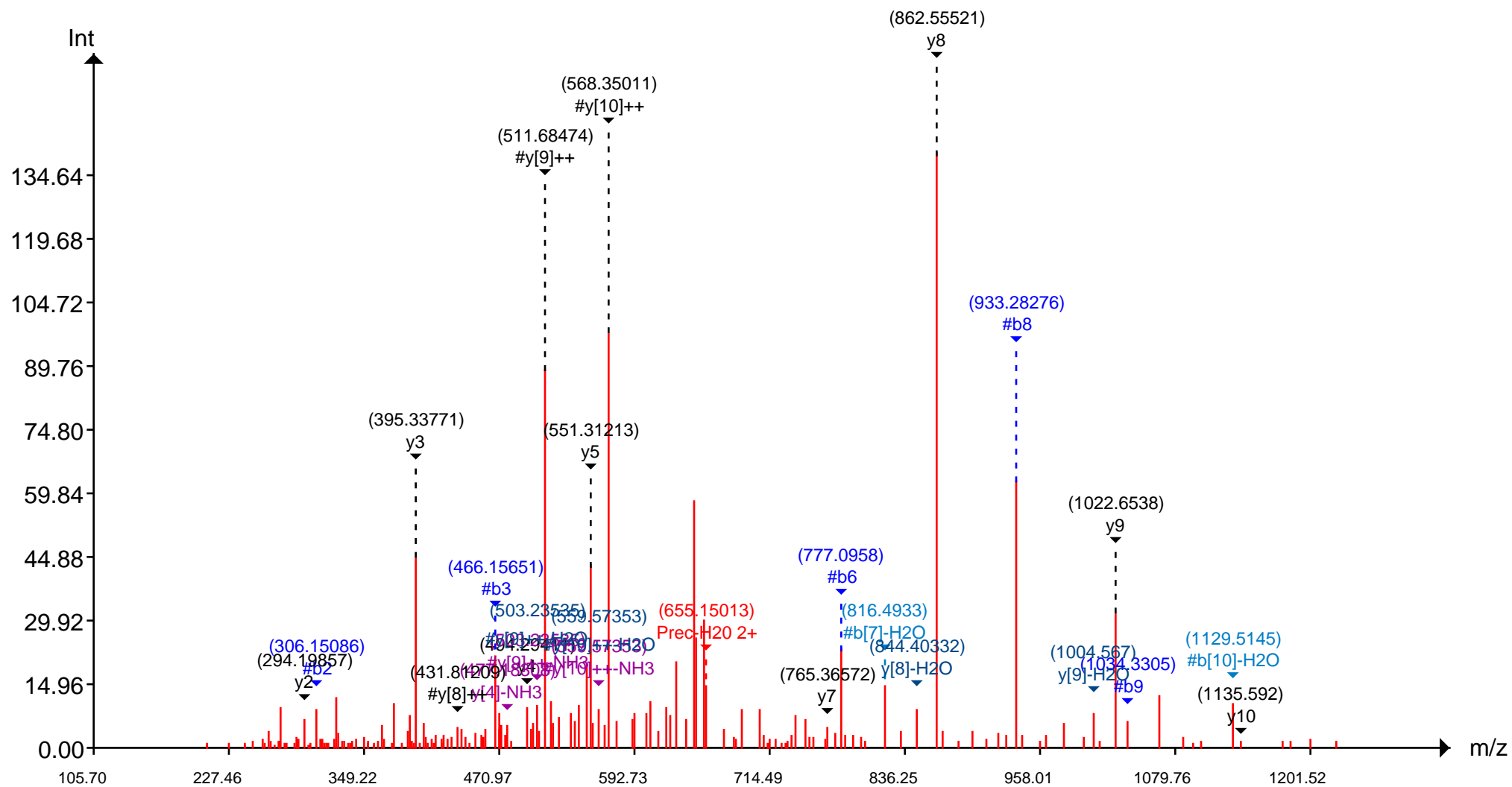


Info

Modified Sequence	AcD3-IK<AcD3K*>ASSIEAR<C13>-COOH	Ion score	68.53
Protein	P57759	Confidence level	99.00%
Precursor m/z	535.82438	Identity Threshold	44.70
Charge	2+	Mass error	-3.29E-4

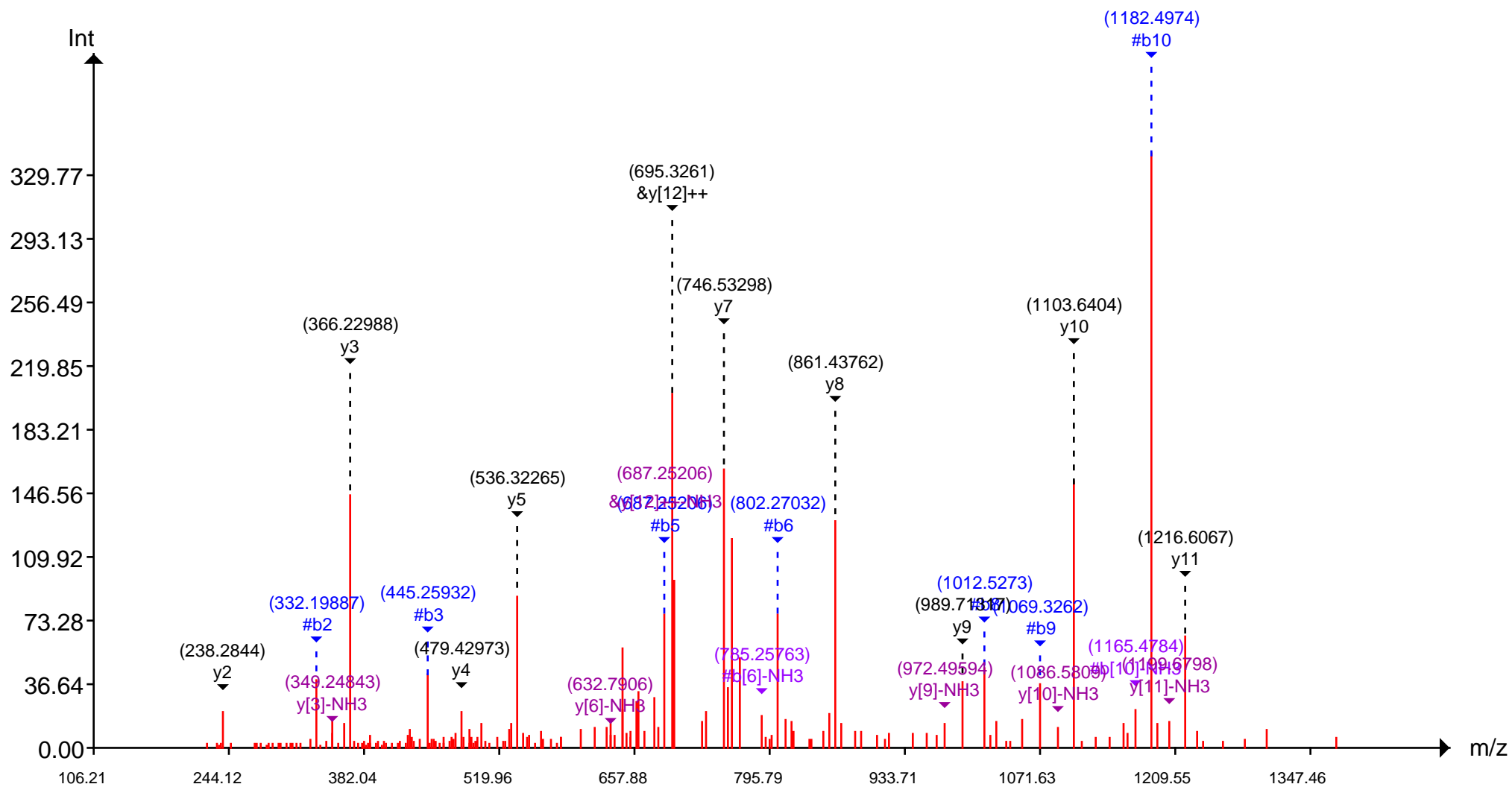


Info			
Modified Sequence	AcD3-LTLGGQK<AcD3K*>C<Cmm*>SVIR-COOH	Ion score	81.62
Protein	P62962	Confidence level	99.00%
Precursor m/z	711.4068	Identity Threshold	46.68
Charge	2+	Mass error	0.001151



Info

Modified Sequence	AcD3-FIC<Cmm*>PLTGVTLR<C13>-COOH	Ion score	46.97
Protein	Q99PL6	Confidence level	99.00%
Precursor m/z	664.38302	Identity Threshold	46.28
Charge	2+	Mass error	0.001015



Info

Modified Sequence AcD3-IK<AcD3K*>INQDPLGIQGR<C13>-COOH

Ion score 66.17

Protein Q9QWR8

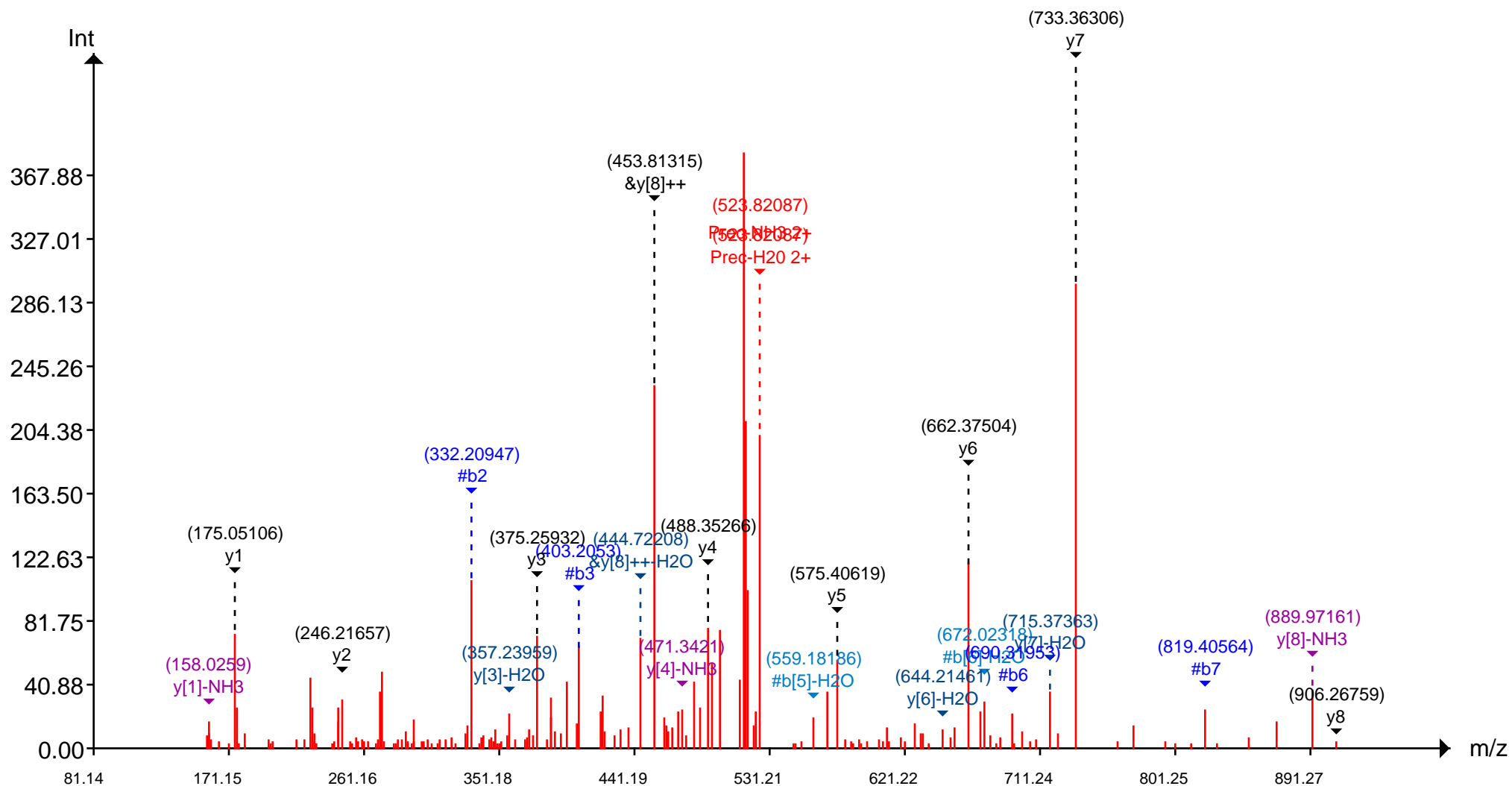
Confidence level 99.00%

Precursor m/z 774.45994

Identity Threshold 44.18

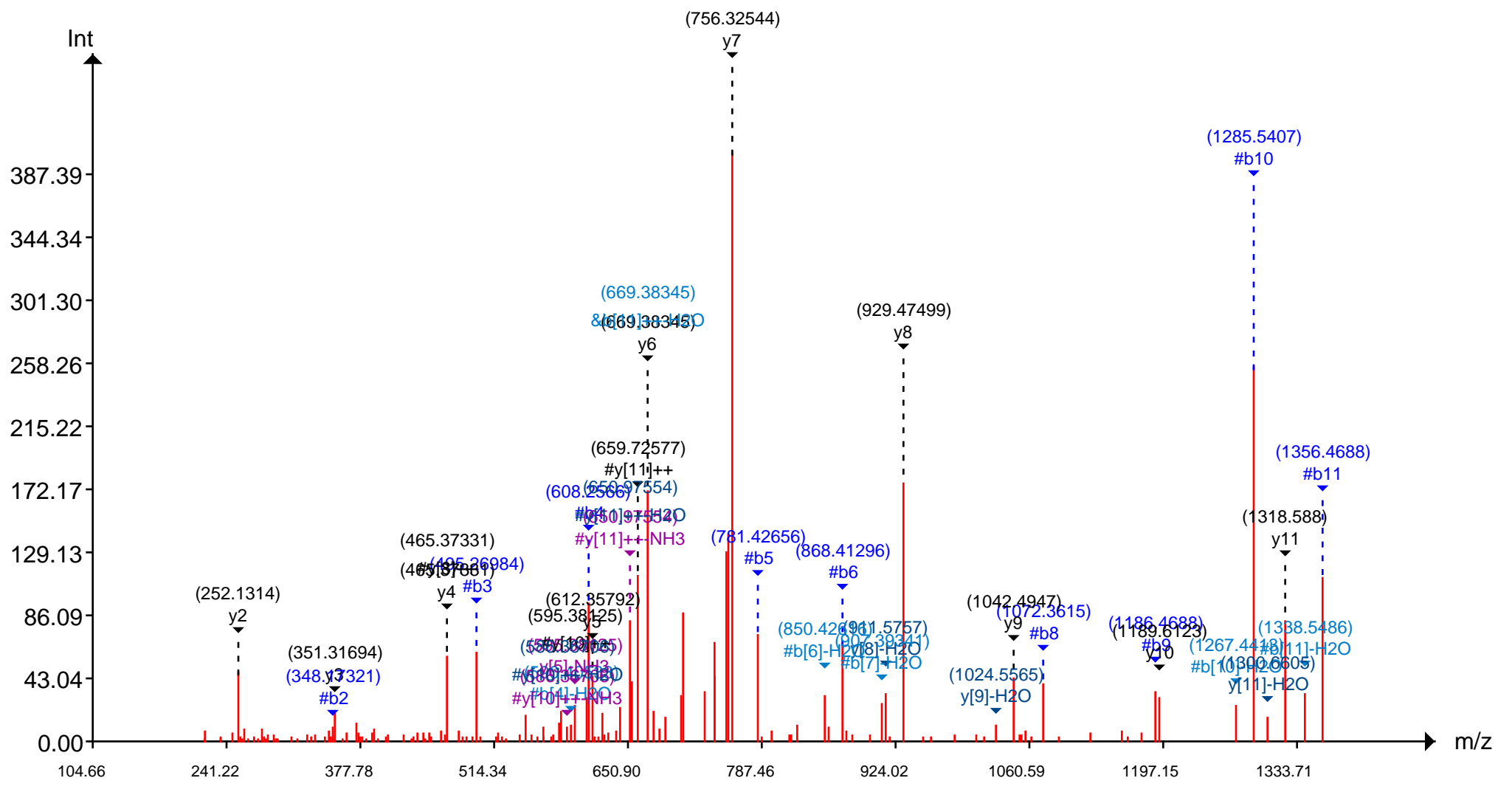
Charge 2+

Mass error 8.02E-4

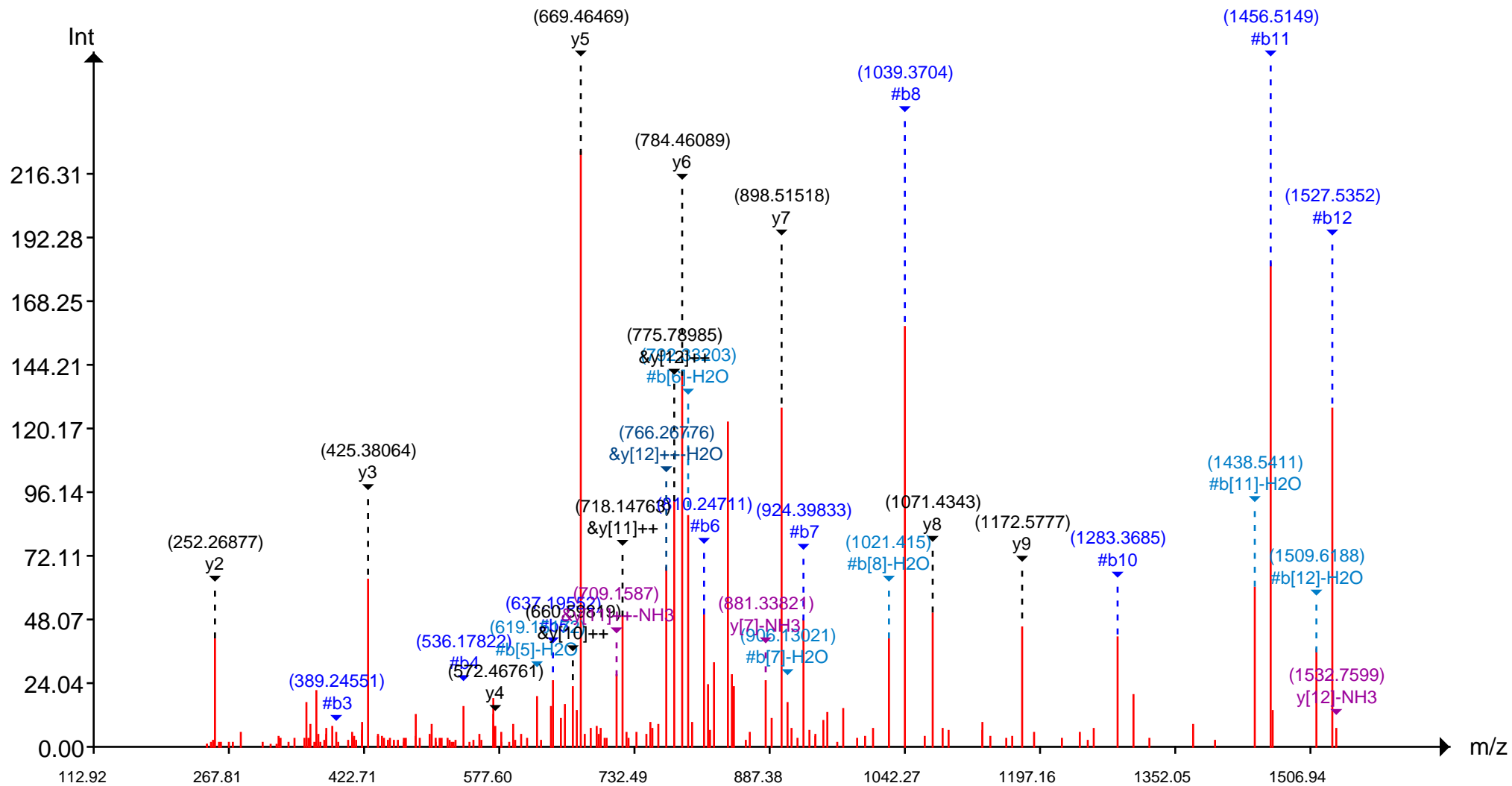


Info

Modified Sequence	AcD3-IK<AcD3K*>ASSIEAR-COOH	Ion score	50.18
Protein	P57759	Confidence level	99.00%
Precursor m/z	532.81598	Identity Threshold	45.99
Charge	2+	Mass error	0.003013

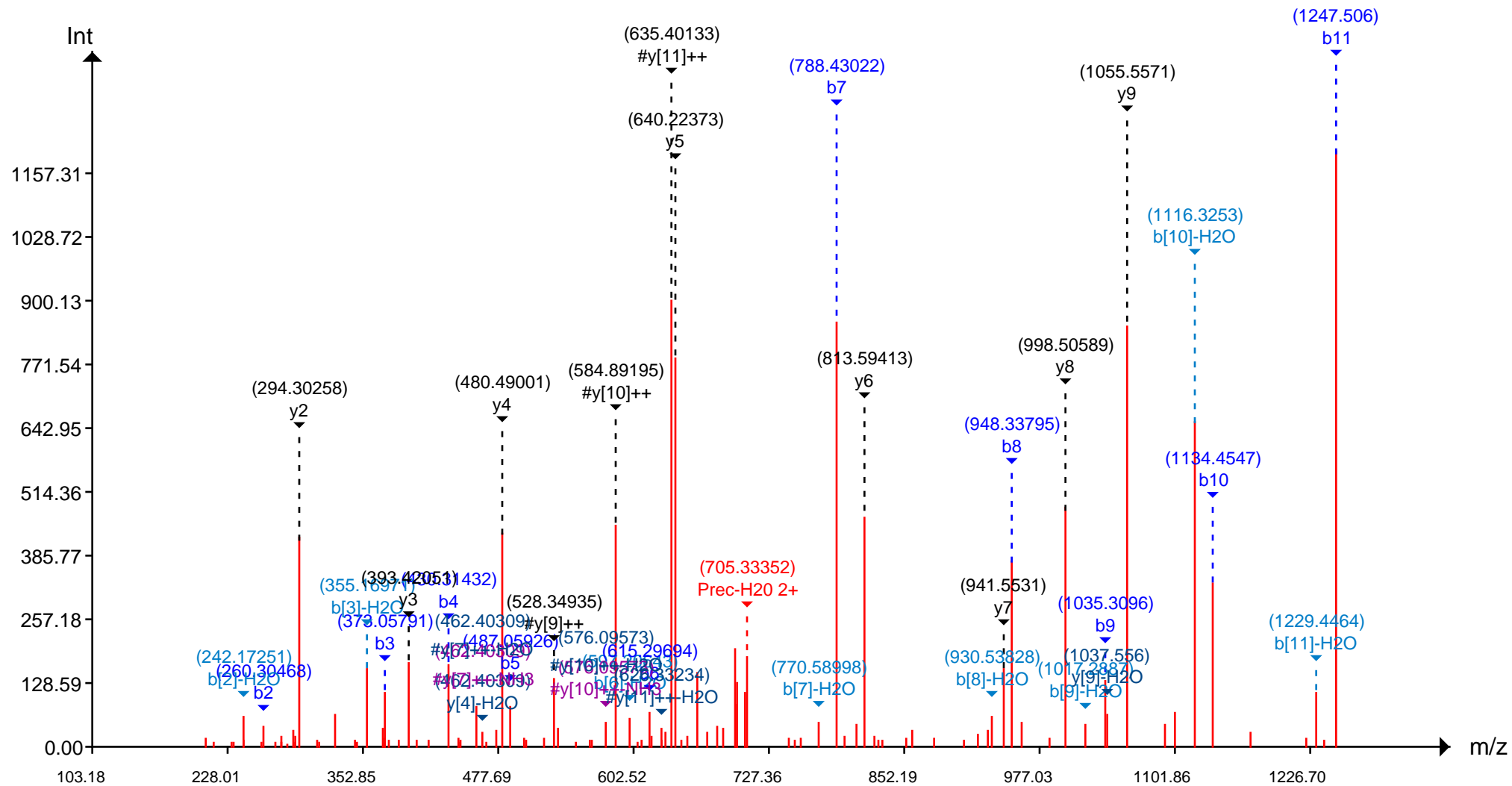


Info			
Modified Sequence	AcD3-K<AcD3K*>EM<Mox*>IK<AcD3K*>SGM<Mox*>NVAR<C13>-COOH	Ion score	98.12
Protein	P52480	Confidence level	99.00%
Precursor m/z	768.91209	Identity Threshold	48.07
Charge	2+	Mass error	3.17E-4



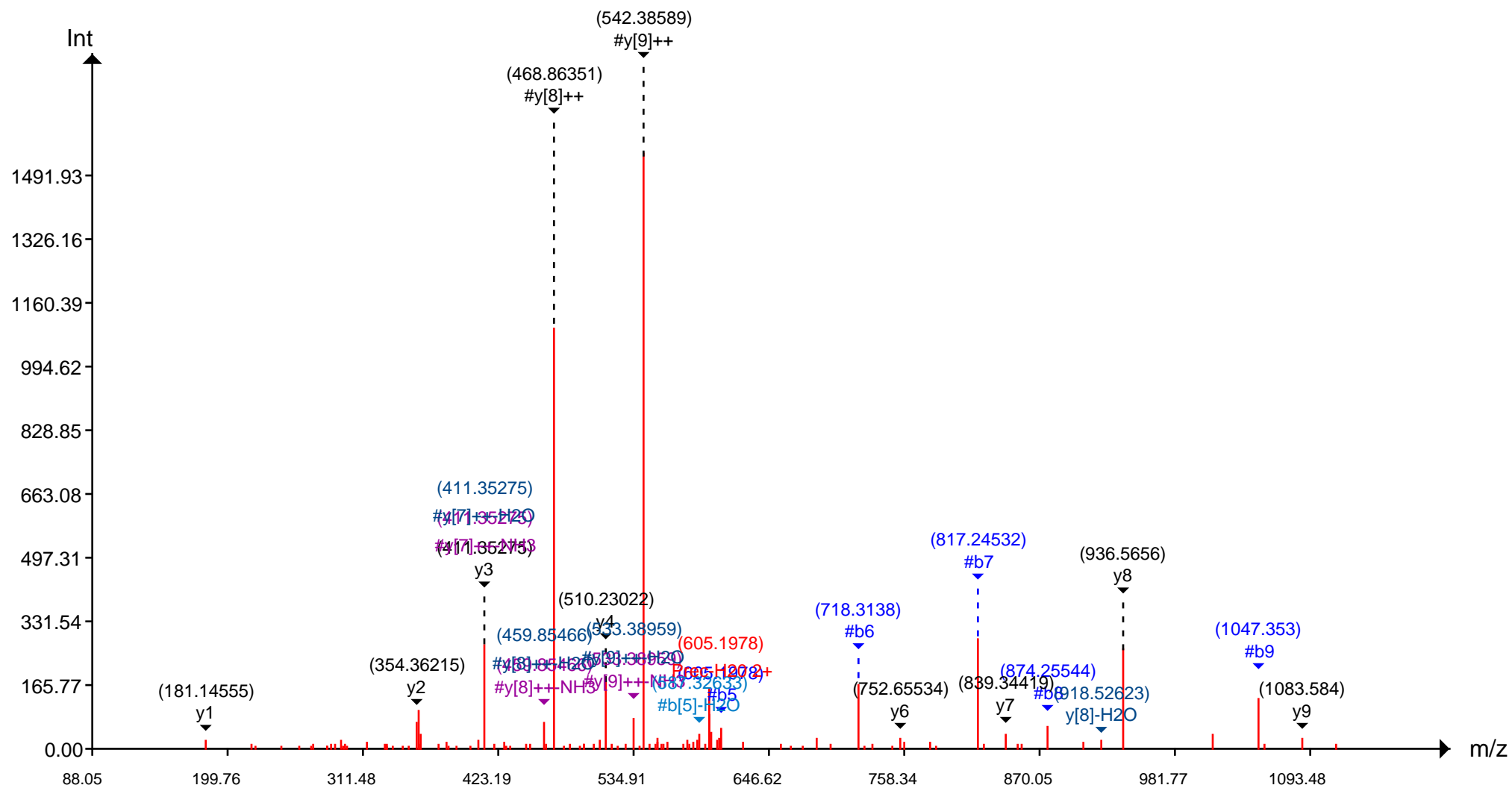
Info

Modified Sequence	AcD3-LDDM<Mox*>TK<AcD3K*>NDPFK<AcD3K*>AR<C13>-COOH	Ion score	67.85
Protein	Q62523	Confidence level	99.00%
Precursor m/z	854.43757	Identity Threshold	47.93
Charge	2+	Mass error	0.001472



Info

Modified Sequence	AcD3-LTLGGQK<AcD3K*>C<Cmm*>SVIR<C13>-COOH	Ion score	79.04
Protein	P62962	Confidence level	99.00%
Precursor m/z	714.41674	Identity Threshold	45.51
Charge	2+	Mass error	8.89E-4



Info

Modified Sequence AcD3-VFPSEIVGK<AcD3K*>R<C13>-COOH

Ion score 60.46

Protein P62082

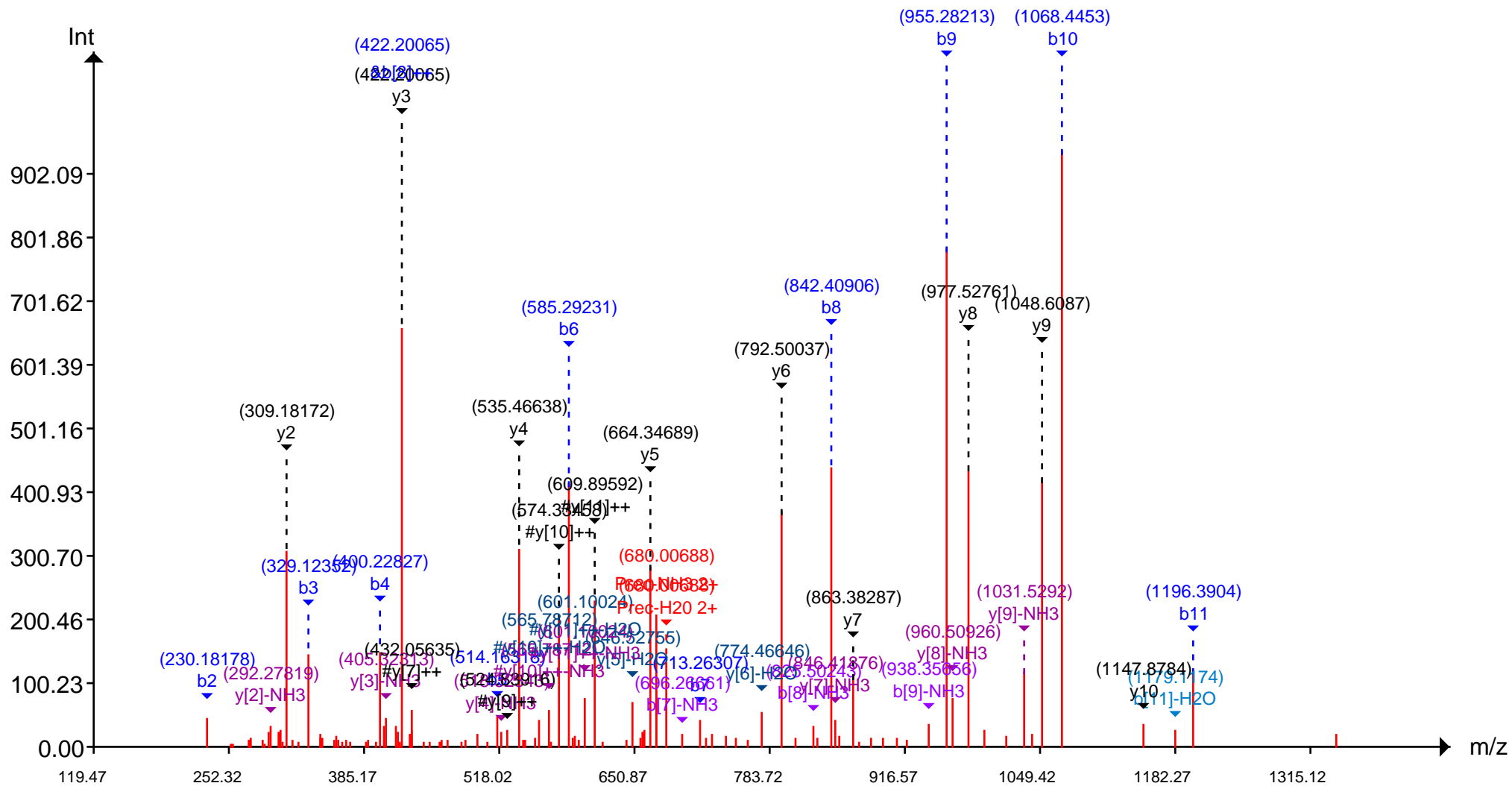
Confidence level 99.00%

Precursor m/z 614.36936

Identity Threshold 44.61

Charge 2+

Mass error 4.44E-4



Info

Modified Sequence AcD3-IAVANAQELLQR<C13>-COOH

Ion score 95.45

Protein P62075

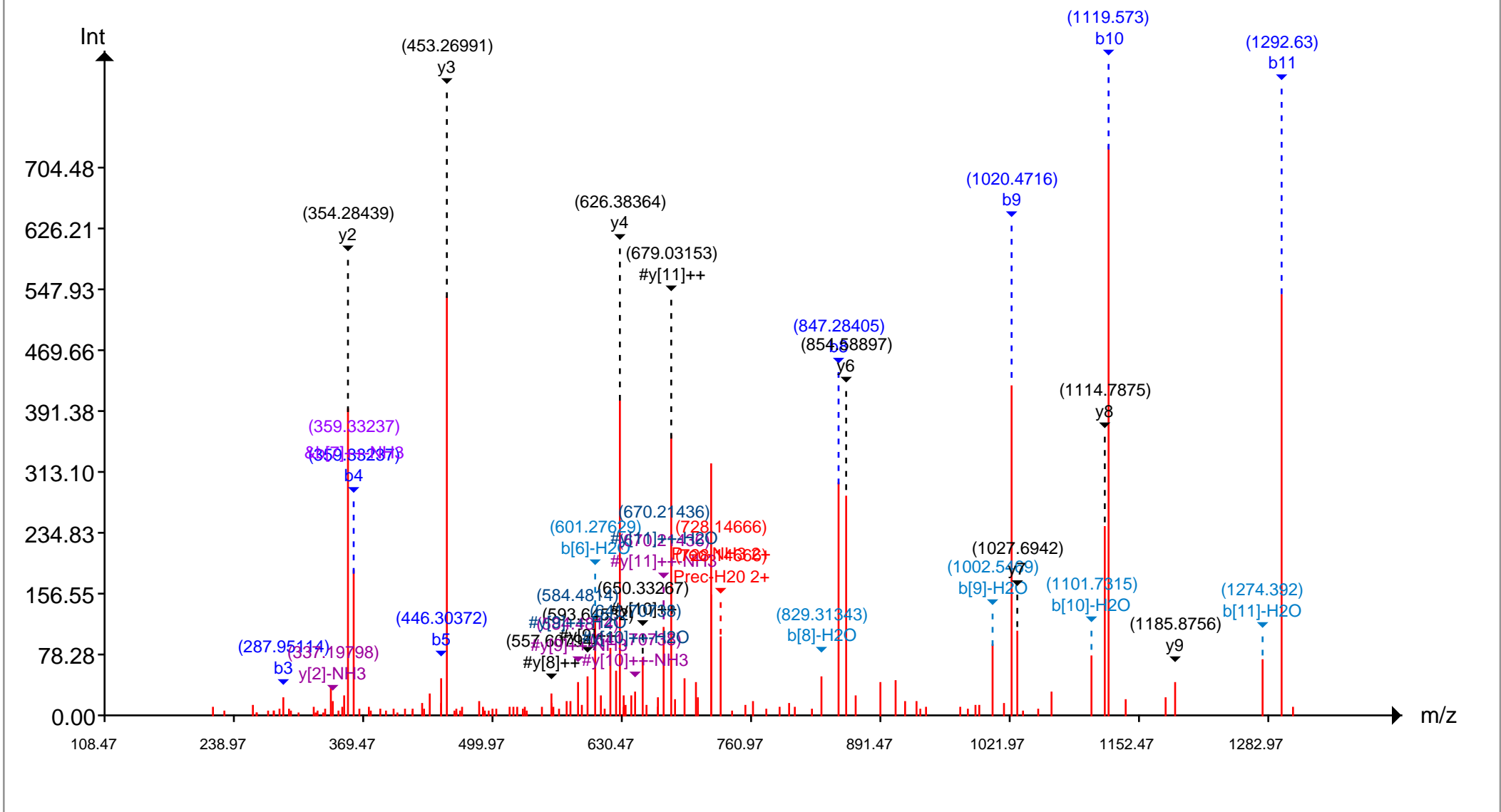
Confidence level 99.00%

Precursor m/z 688.90549

Identity Threshold 45.39

Charge 2+

Mass error 6.36E-4



Info

Modified Sequence AcD3-AGNASK<AcD3K*>DLK<AcD3K*>VK<AcD3K*>R<C13>-COOH

Ion score 67.51

Protein Q3THW5, P0C0S6

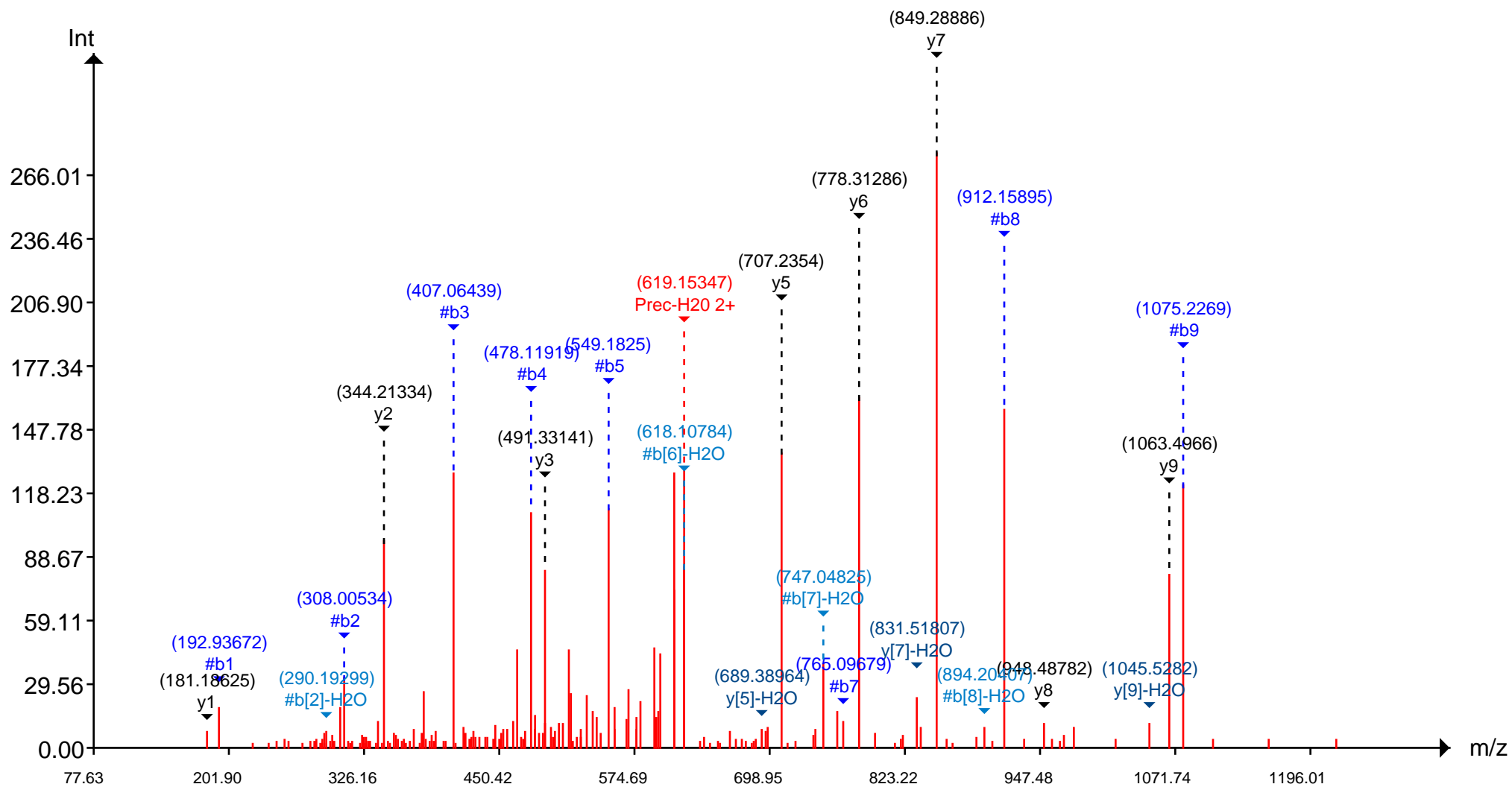
Confidence level 99.00%

Precursor m/z 736.95017

Identity Threshold 42.40

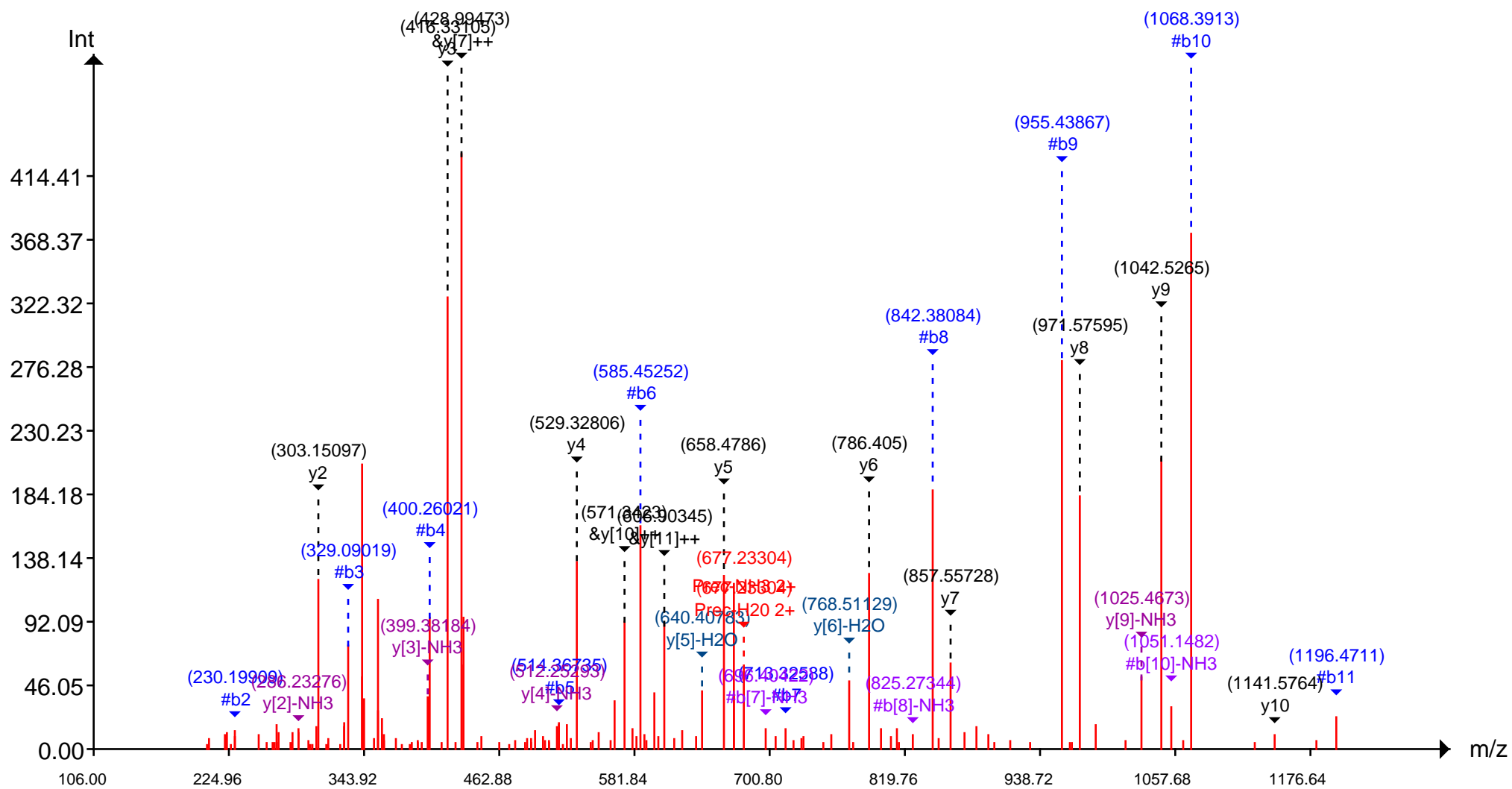
Charge 2+

Mass error 0.00145



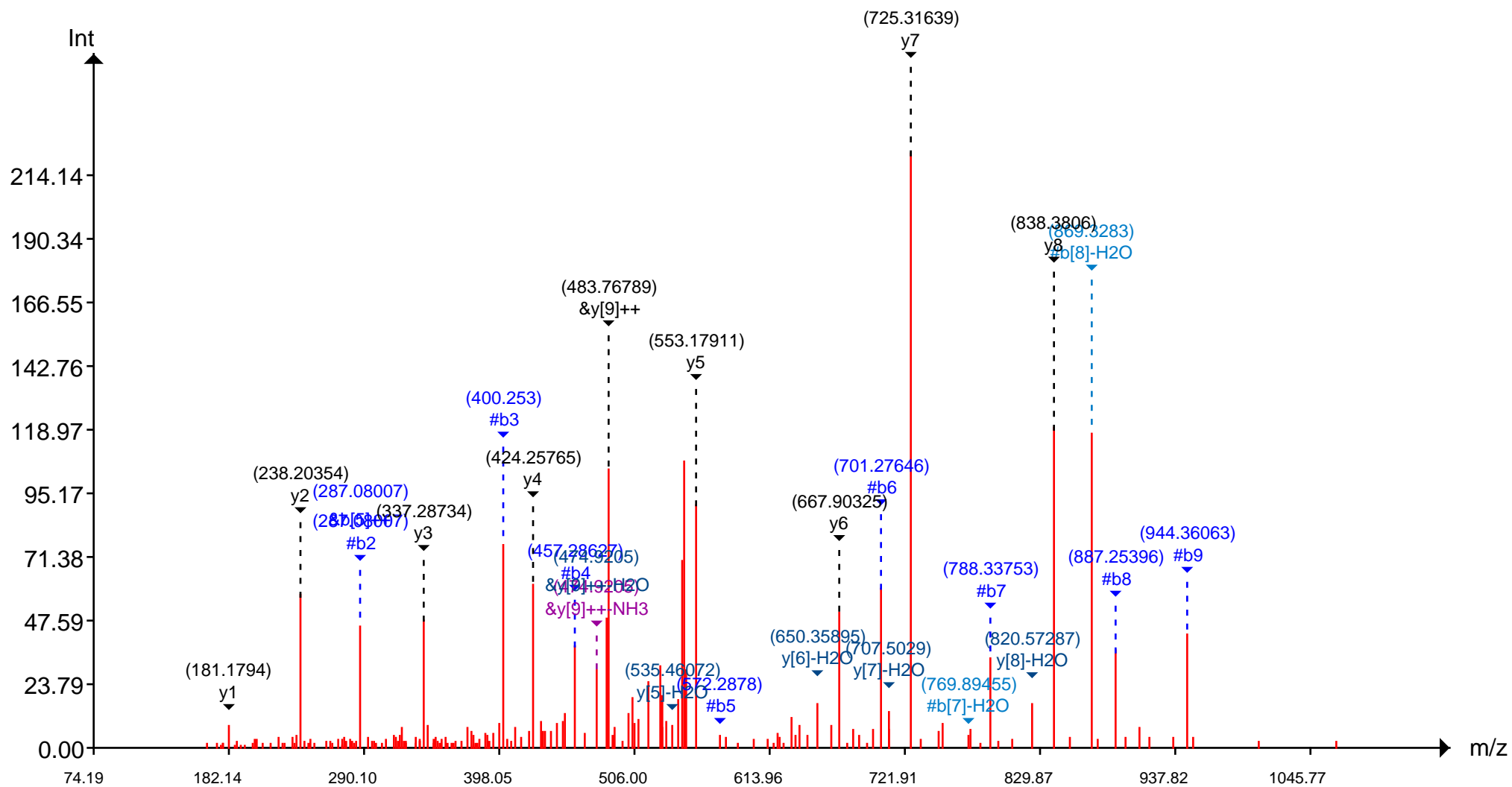
Info

Modified Sequence	AcD3-M<Mox*>DVAASEFYR<C13>-COOH	Ion score	68.87
Protein	P17182, P21550, P17183	Confidence level	99.00%
Precursor m/z	628.29374	Identity Threshold	45.46
Charge	2+	Mass error	3.88E-4



Info

Modified Sequence	AcD3-IAVANAQELLQR-COOH	Ion score	78.97
Protein	P62075	Confidence level	99.00%
Precursor m/z	685.89545	Identity Threshold	46.38
Charge	2+	Mass error	6.98E-4



Info

Modified Sequence AcD3-LQIGDESVGR<C13>-COOH

Ion score 65.09

Protein P24369

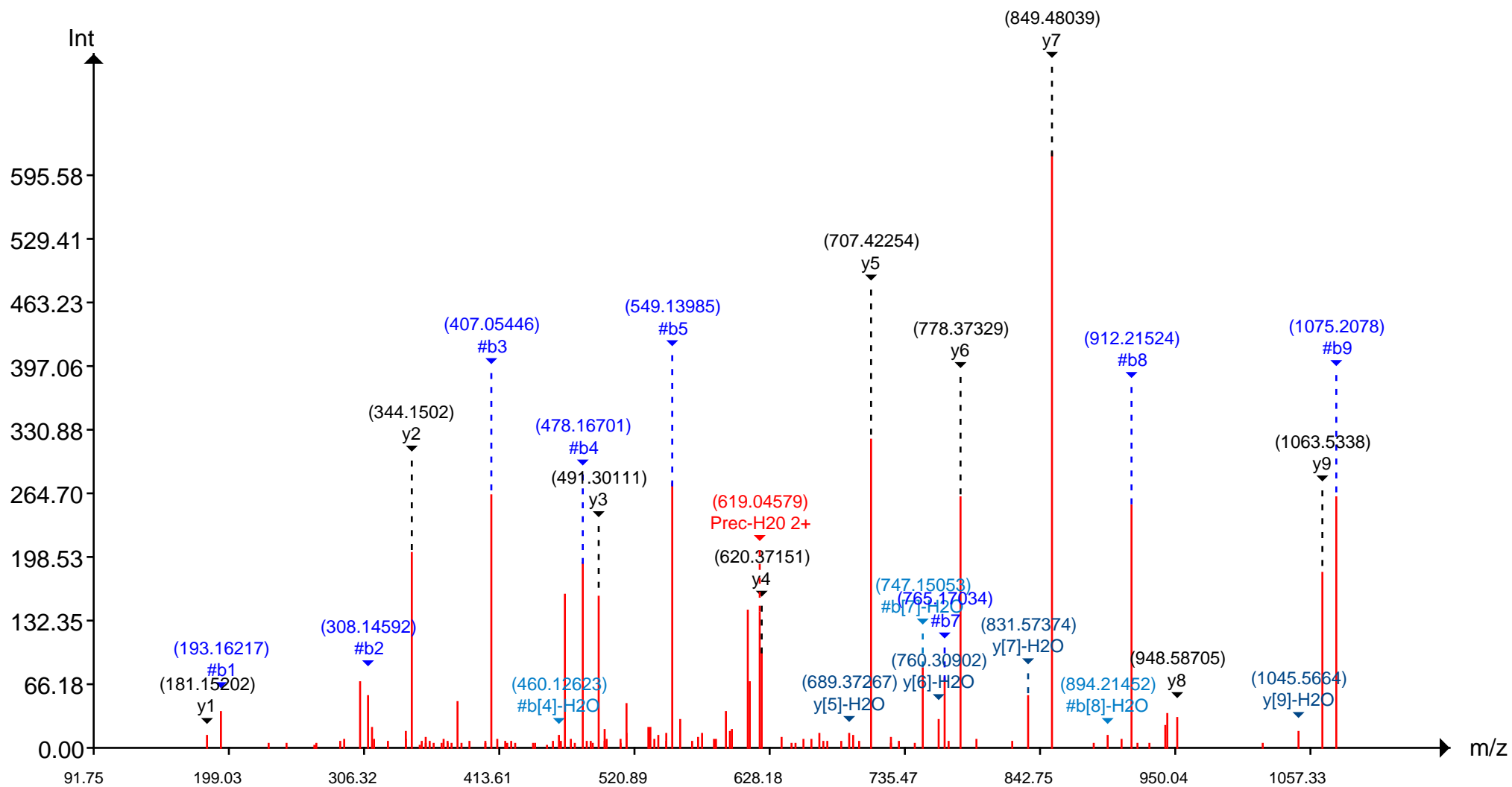
Confidence level 99.00%

Precursor m/z 562.80787

Identity Threshold 47.56

Charge 2+

Mass error 3.73E-4



Info

Modified Sequence AcD3-M<Mox*>DVAASEFYR<C13>-COOH

Ion score 76.1

Protein P17182, P21550, P17183

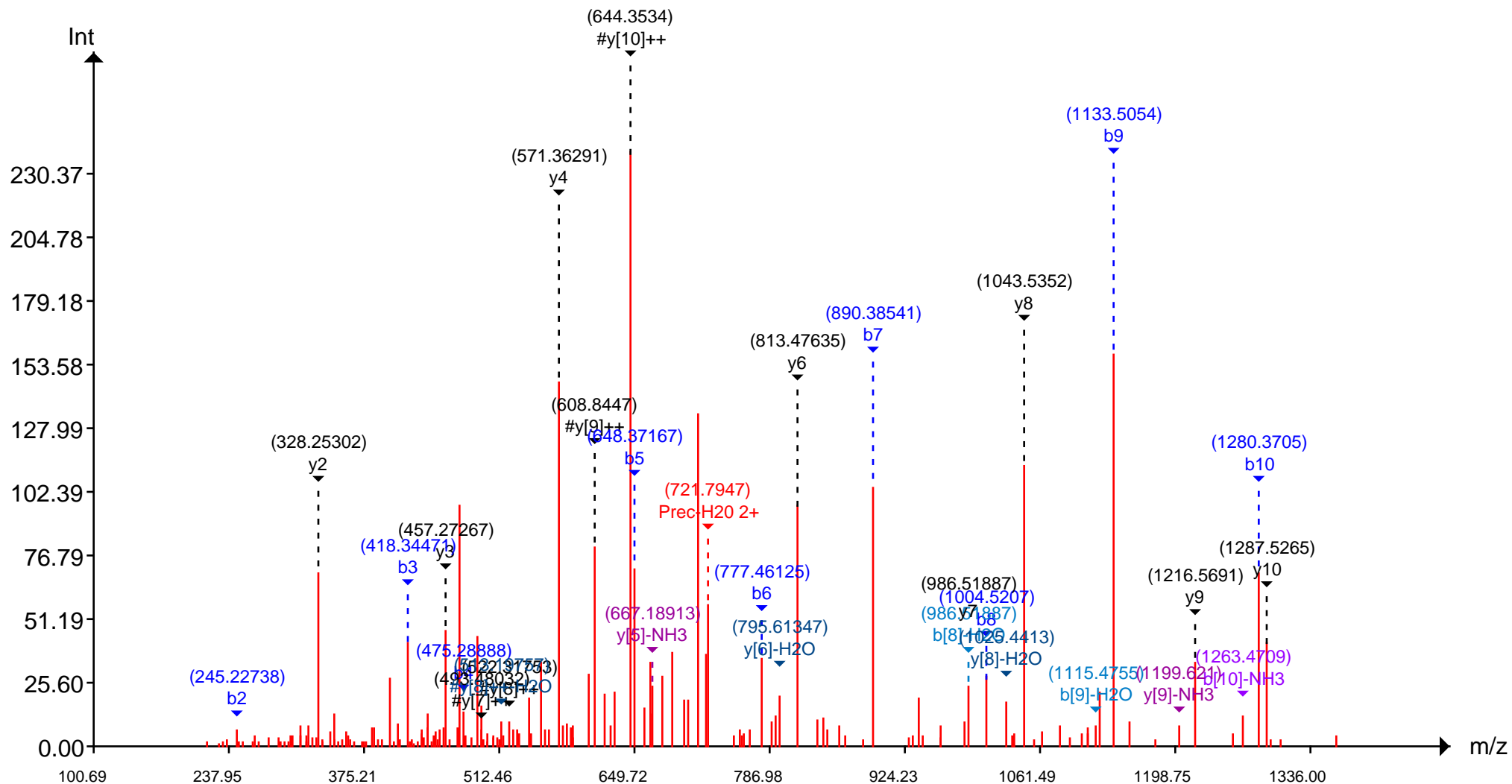
Confidence level 99.00%

Precursor m/z 628.29379

Identity Threshold 45.46

Charge 2+

Mass error 4.88E-4



Info

Modified Sequence AcD3-QAK<AcD3K*>GK<AcD3K*>ELNEFR<C13>-COOH

Ion score 52.5

Protein O70591

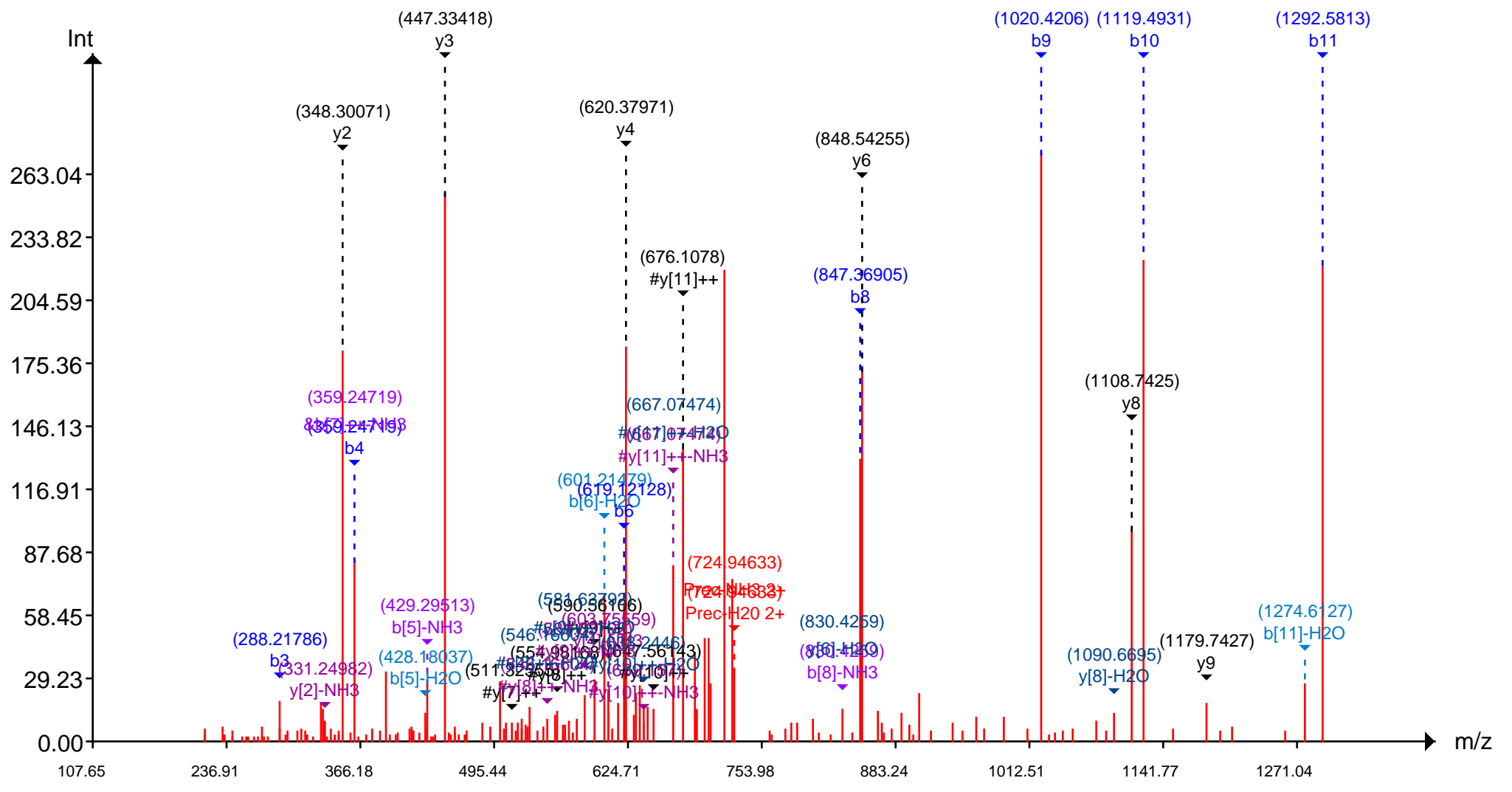
Confidence level 99.00%

Precursor m/z 730.91206

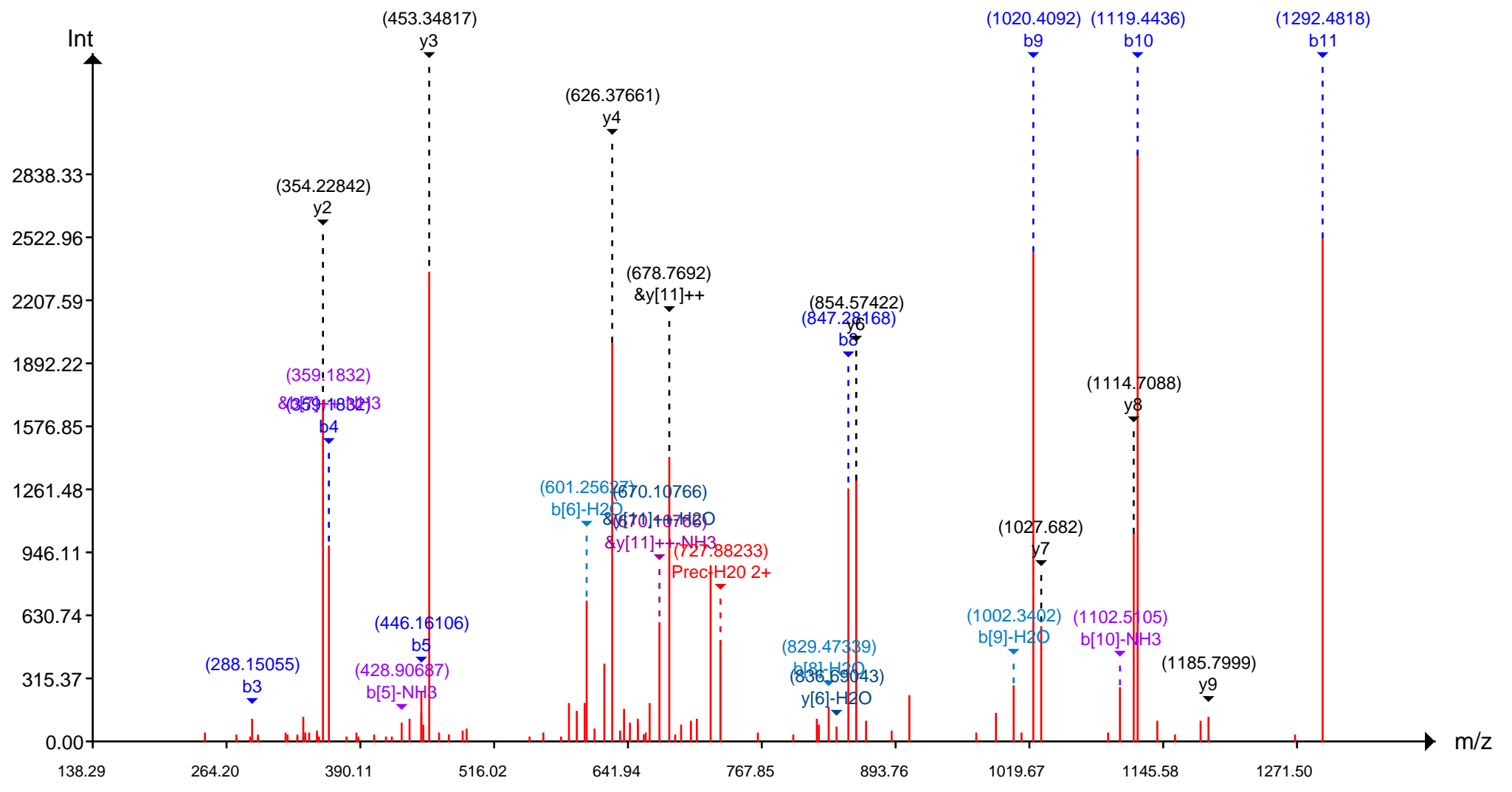
Identity Threshold 47.02

Charge 2+

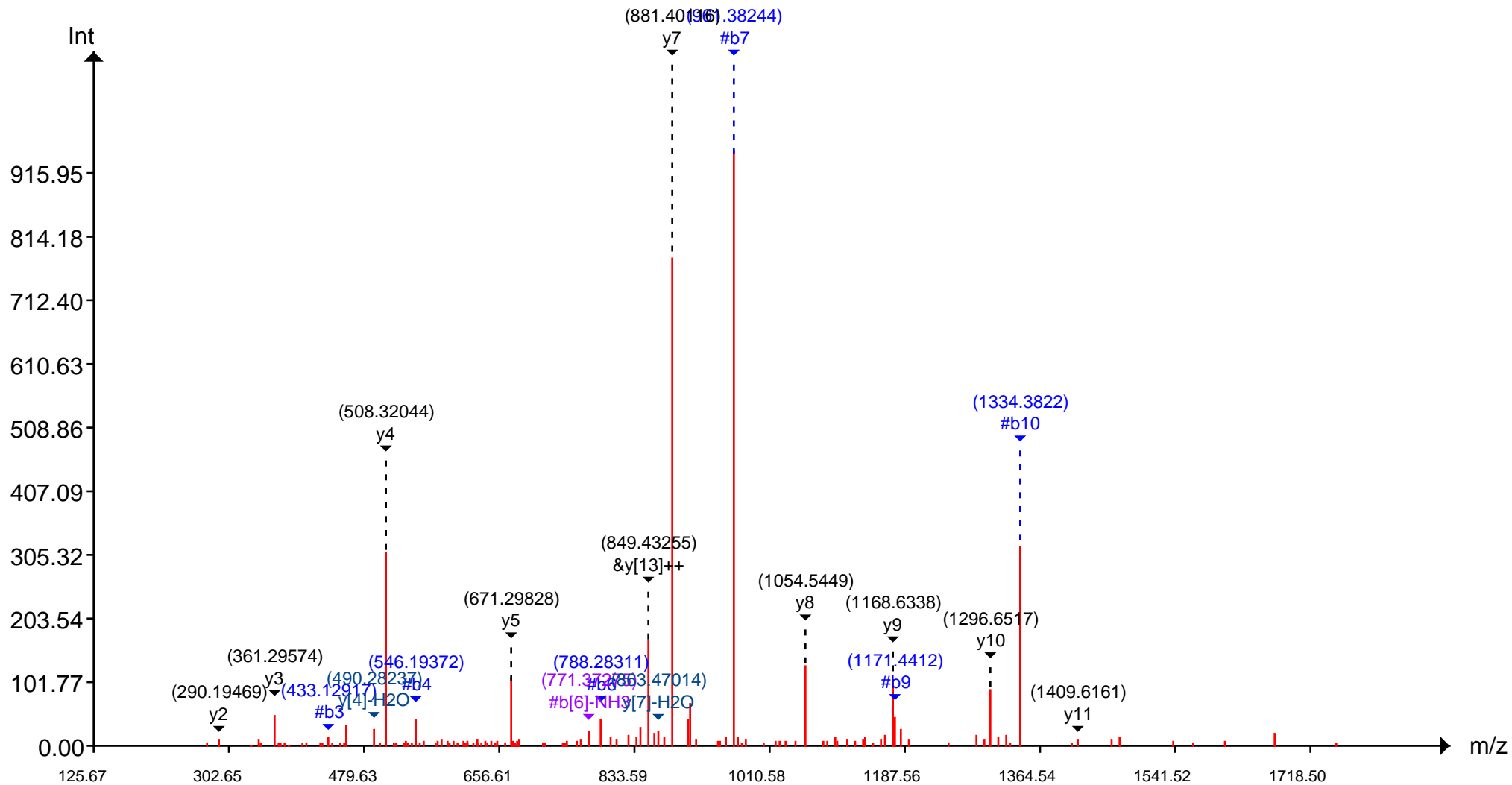
Mass error 0.00192



Info			
Modified Sequence	AcD3-AGNASK<AcD3K*>DLK<AcD3K*>VK<AcD3K*>R-COOH	Ion score	52.3
Protein	Q3THW5, P0C0S6	Confidence level	99.00%
Precursor m/z	733.9403	Identity Threshold	43.79
Charge	2+	Mass error	0.001851

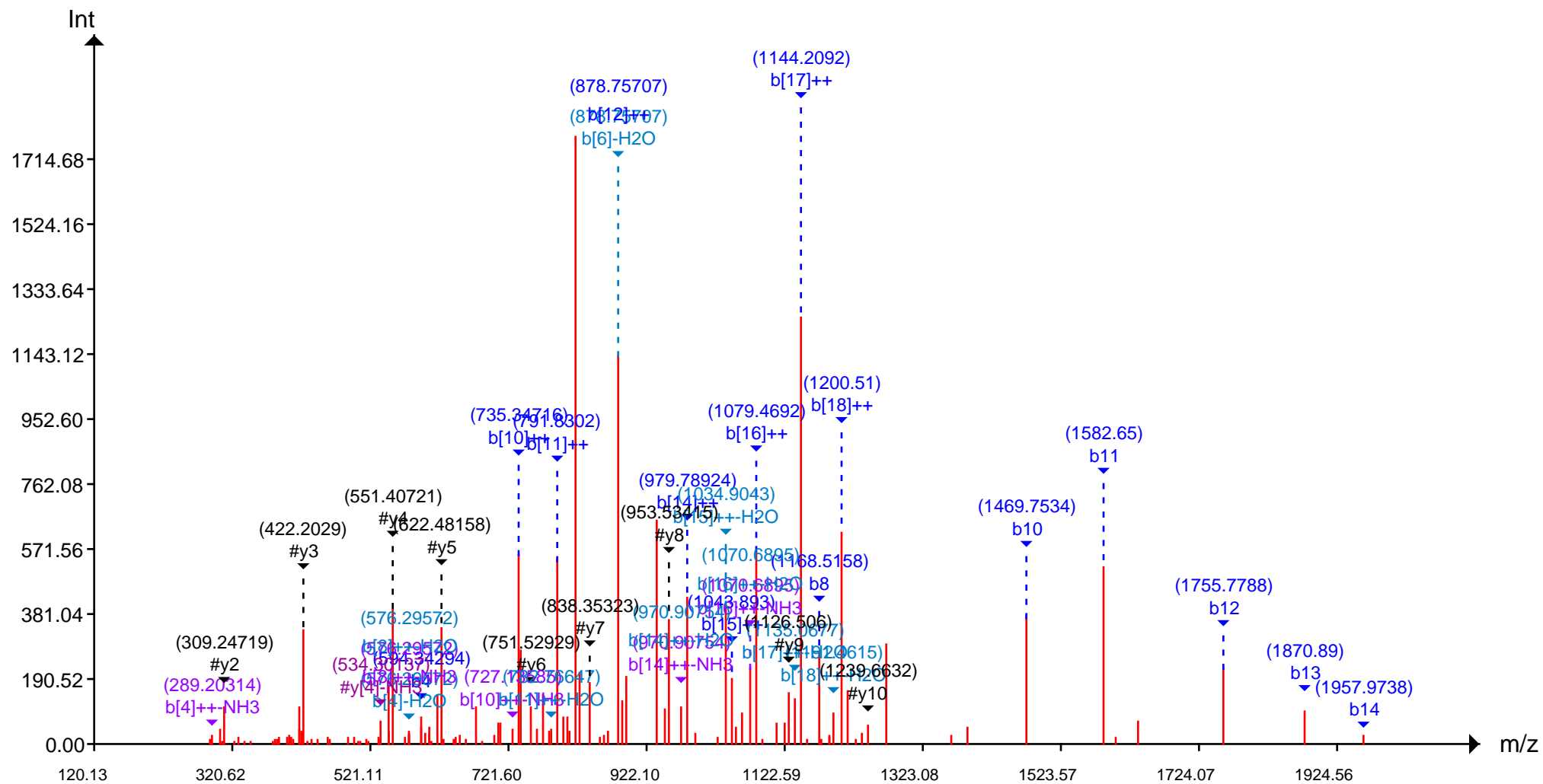


Info			
Modified Sequence	AcD3-AGNASK<AcD3K*>DLK<AcD3K*>VK<AcD3K*>R<C13>-COOH	Ion score	61.34
Protein	Q3THW5, P0C0S6	Confidence level	99.00%
Precursor m/z	736.94906	Identity Threshold	42.68
Charge	2+	Mass error	-7.7E-4



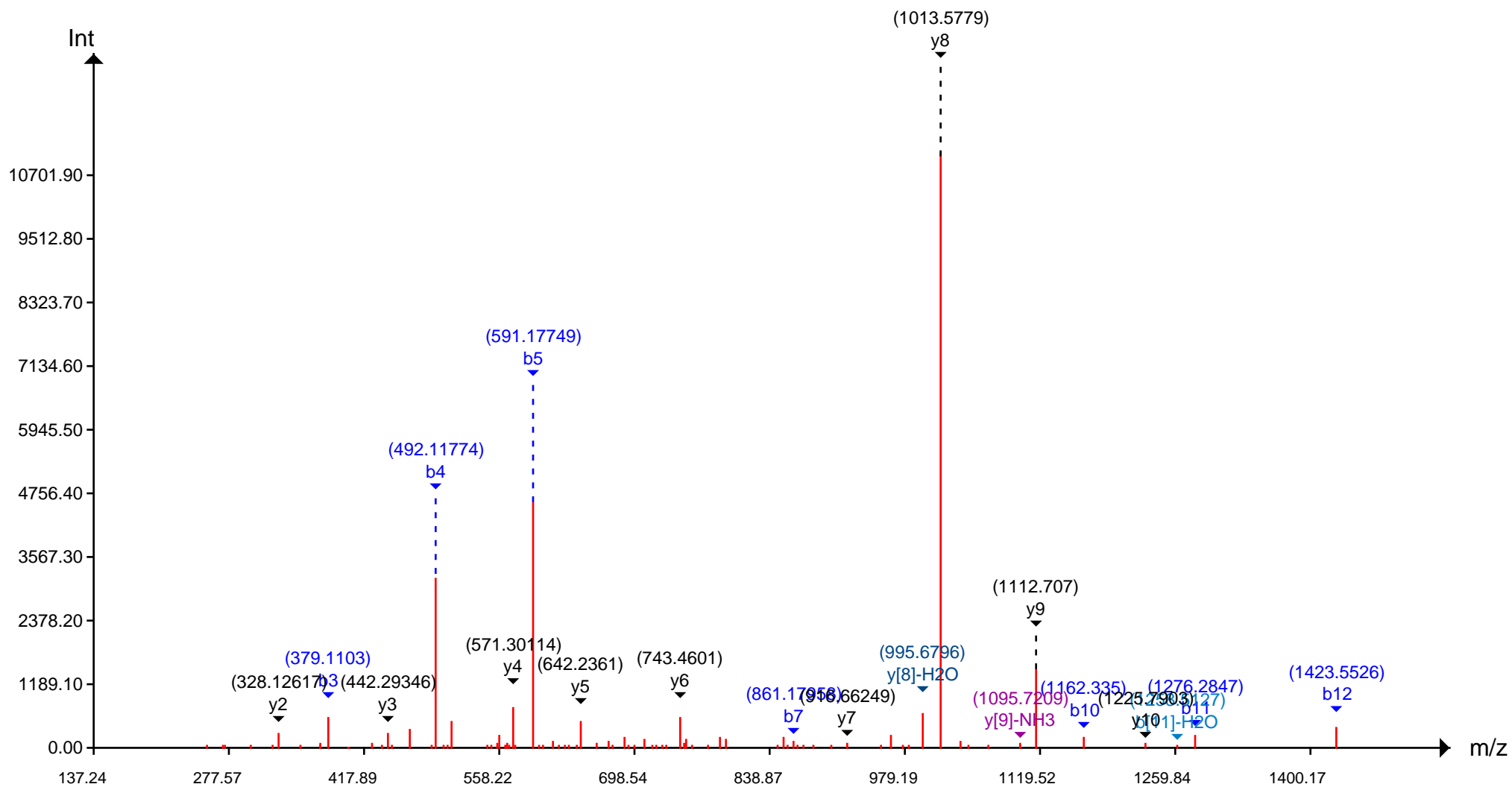
Info

Modified Sequence	AcD3-VQC<Cmm*>IQNK<AcD3K*>PLYFADR-COOH	Ion score	56.58
Protein	P07356	Confidence level	99.00%
Precursor m/z	921.47898	Identity Threshold	48.02
Charge	2+	Mass error	0.002154



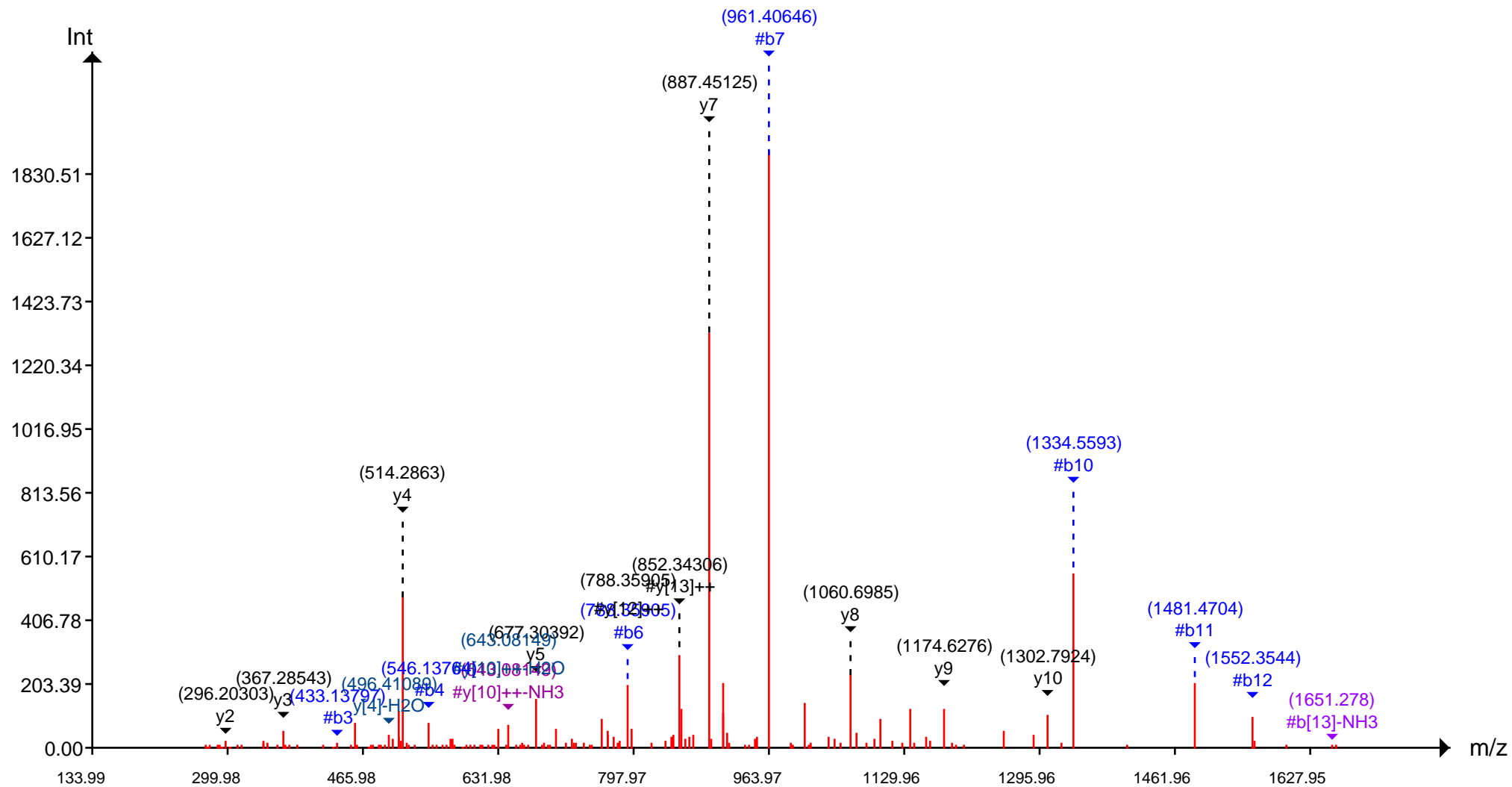
Info

Modifie...	AcD3-EM<Mox*>K<AcD3K*>VK<AcD3K*>EK<AcD3K*>VQK<AcD3K*>LK<AcD3K*>DSEAELQR<C13>-COOH	Ion score
	O55131	Confidence level
Pre...	903.51047	Identity Threshold
	3+	Mass error



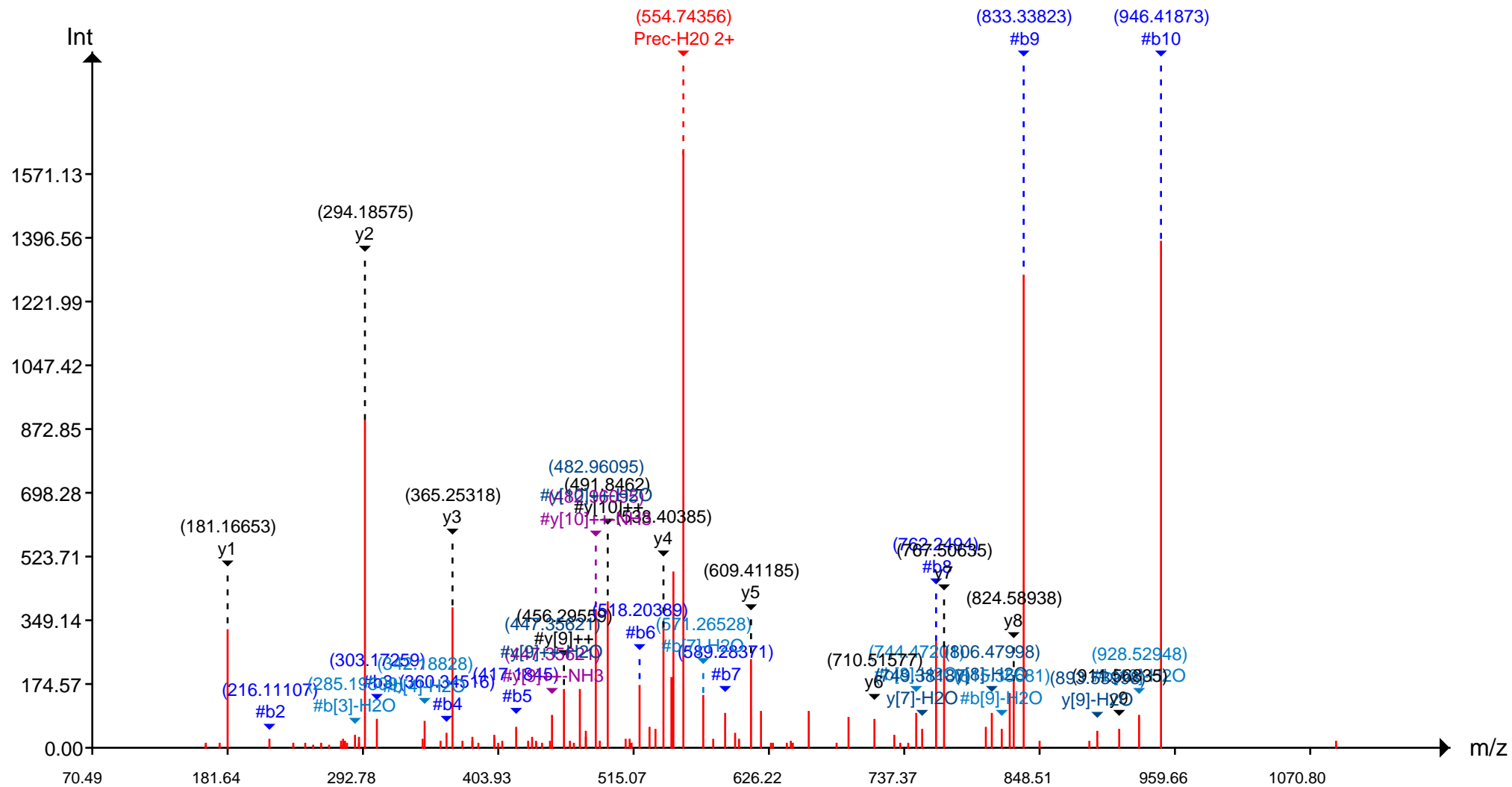
Info

Modified Sequence	AcD3-FADIVPK<AcD3K*>TAENFR<C13>-COOH	Ion score	71.83
Protein	Q9CR16	Confidence level	99.00%
Precursor m/z	802.43935	Identity Threshold	47.69
Charge	2+	Mass error	0.002148



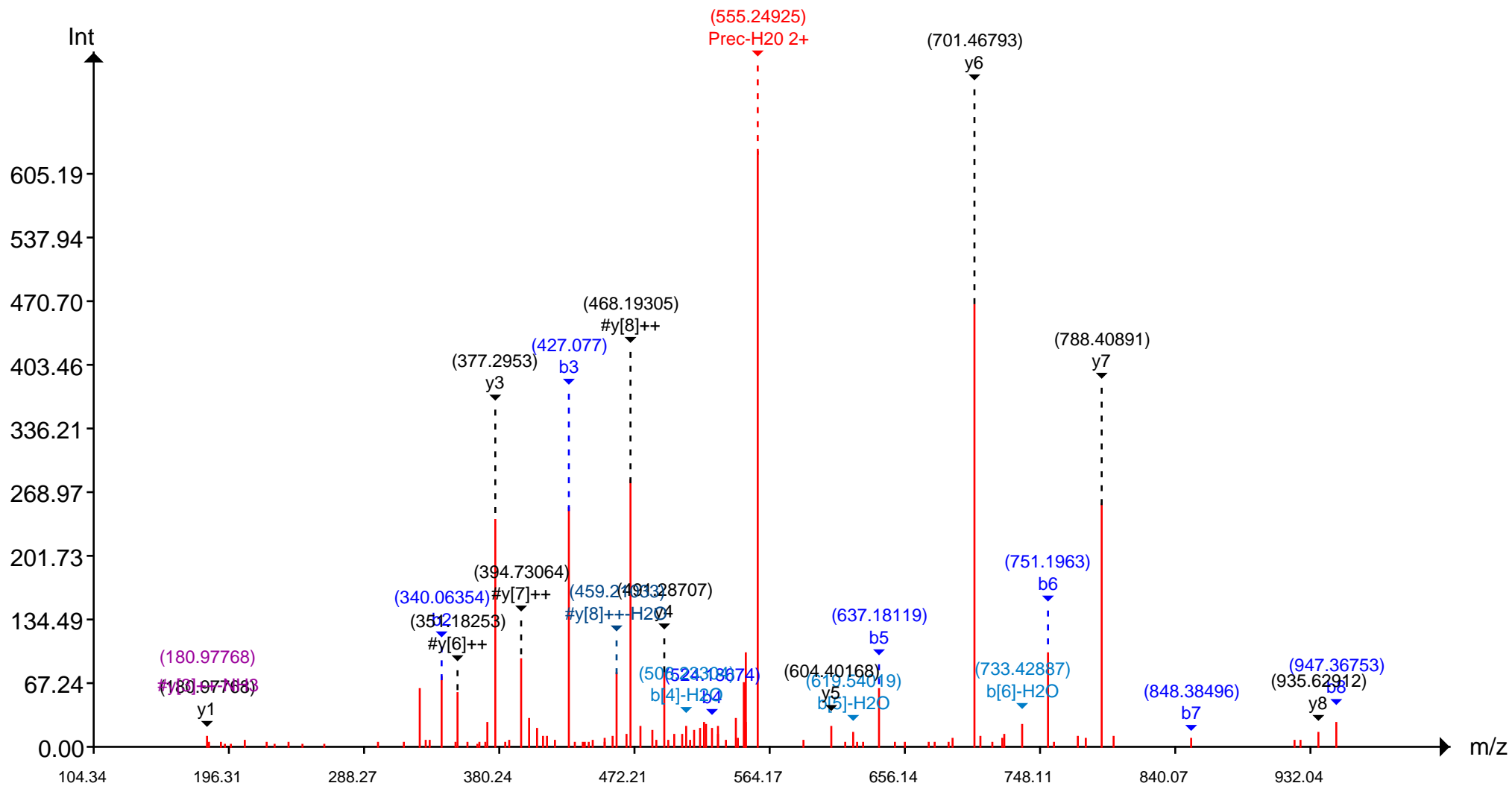
Info

Modified Sequence	AcD3-VQC<Cmm*>IQNK<AcD3K*>PLYFADR<C13>-COOH	Ion score	51.51
Protein	P07356	Confidence level	99.00%
Precursor m/z	924.48891	Identity Threshold	48.04
Charge	2+	Mass error	0.001873



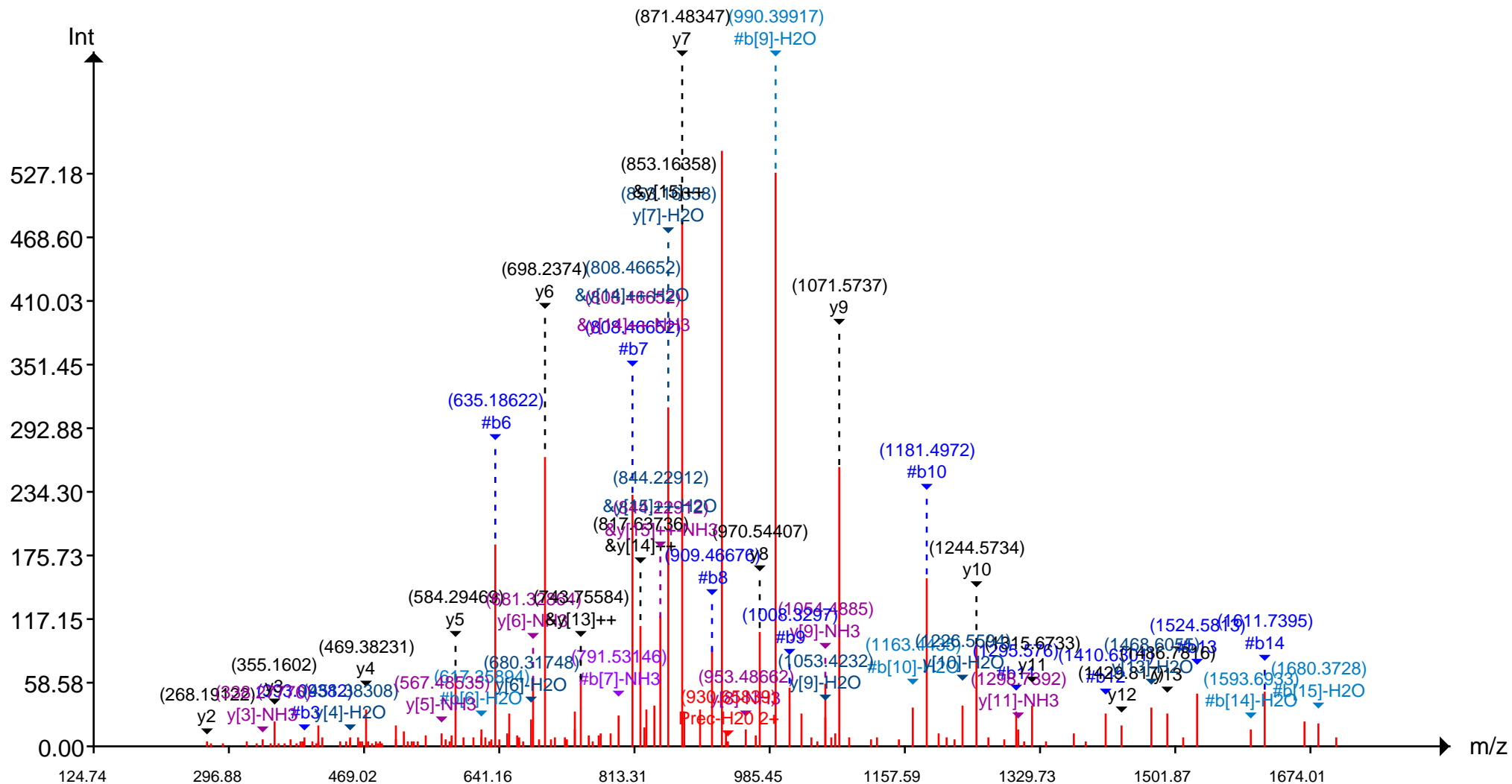
Info

Modified Sequence	AcD3-VASGGTAK<AcD3K*>AIR<C13>-COOH	Ion score	76.4
Protein	Q9CWJ9	Confidence level	99.00%
Precursor m/z	563.84288	Identity Threshold	44.89
Charge	2+	Mass error	-8.04E-4



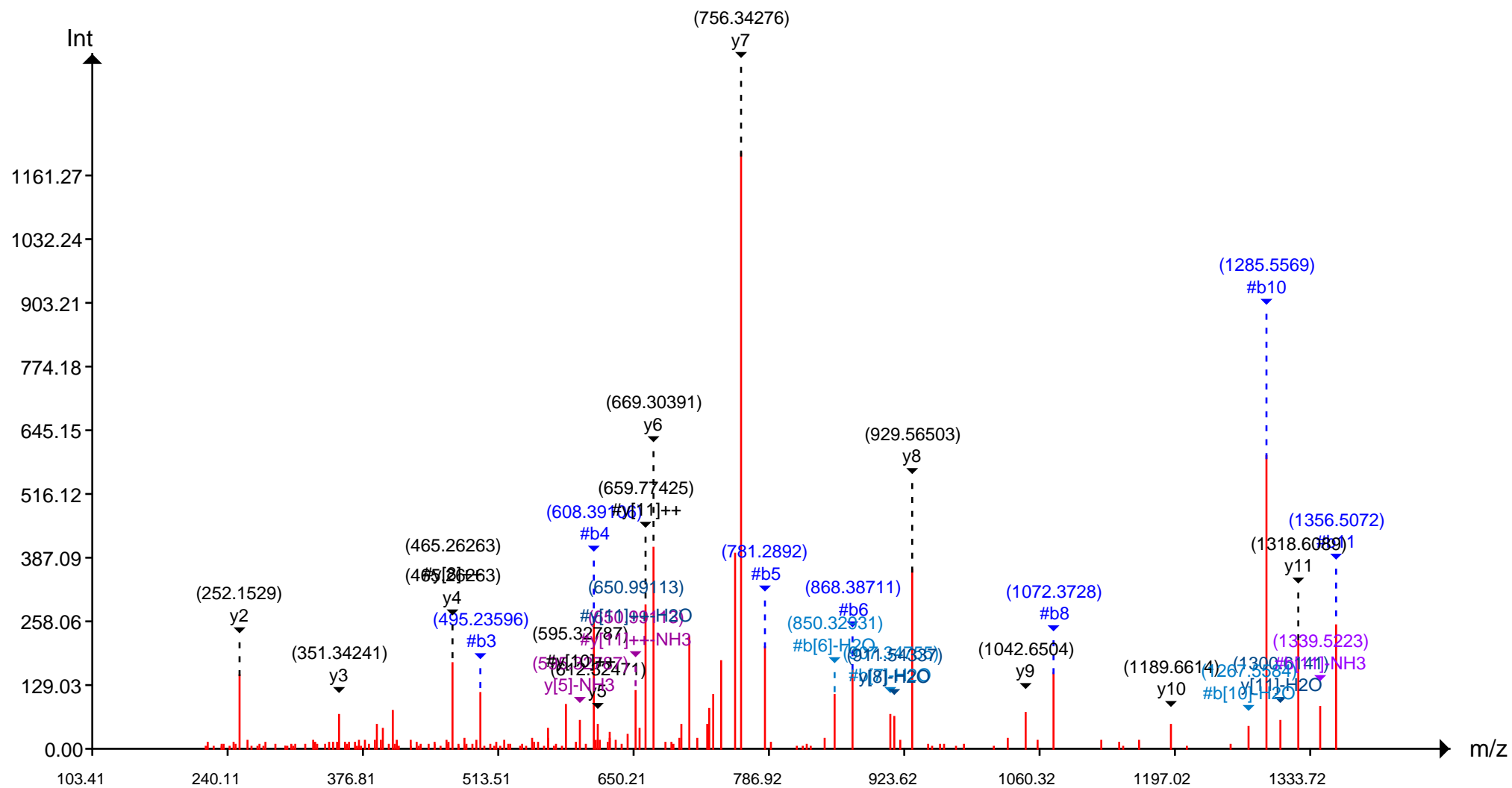
Info

Modified Sequence	AcD3-FFSPLNPVR<C13>-COOH	Ion score	59.41
Protein	O35737, Q9Z2X1	Confidence level	99.00%
Precursor m/z	564.32273	Identity Threshold	46.87
Charge	2+	Mass error	-8.9E-5



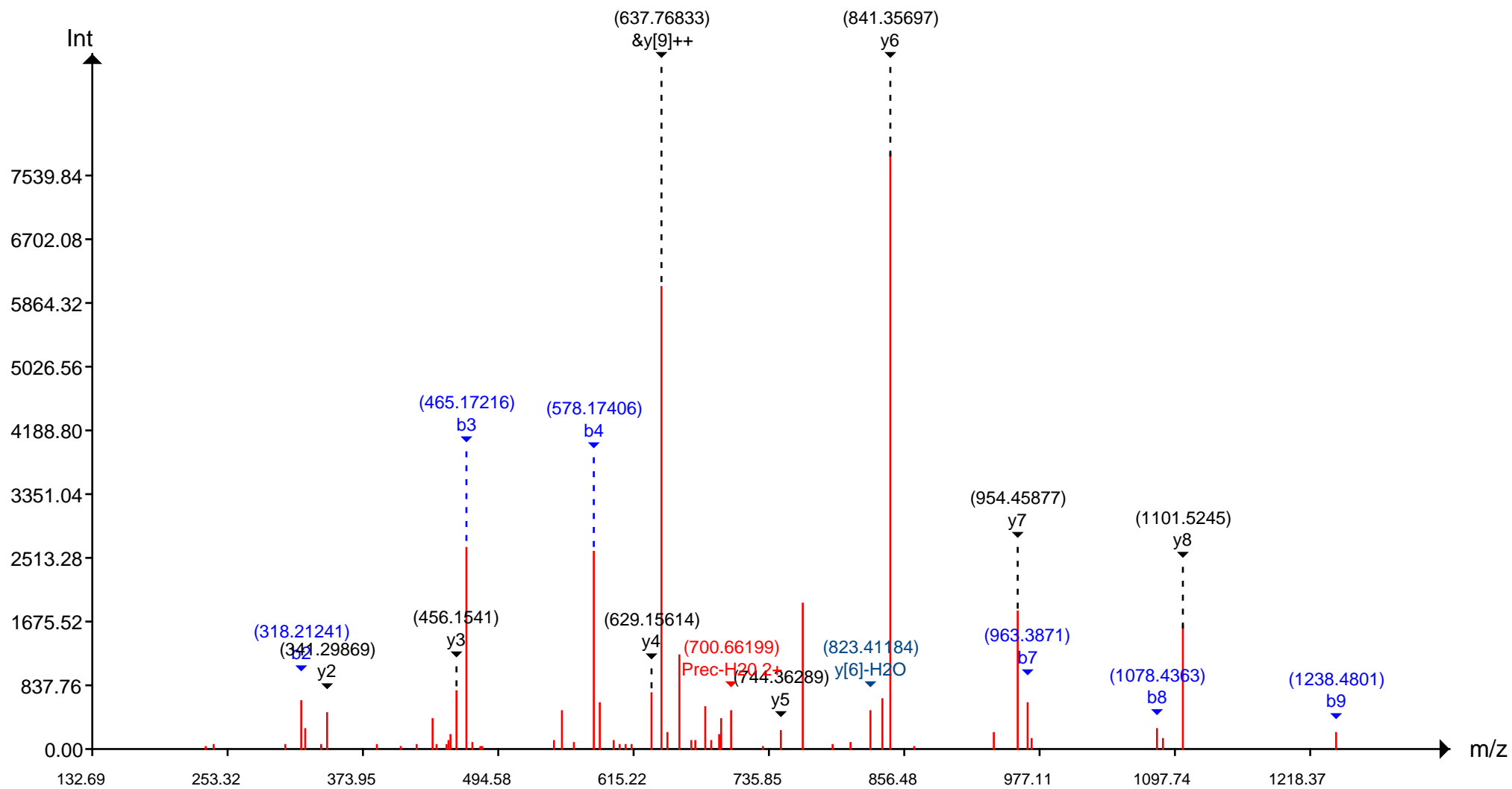
Info

Modified Sequence	AcD3-EAFGNAK<AcD3K*>TVK<AcD3K*>NDNSSR<C13>-COOH	Ion score	103.86
Protein	O08638, Q6URW6, Q8VDD5	Confidence level	99.00%
Precursor m/z	939.9842	Identity Threshold	48.09
Charge	2+	Mass error	0.001364



Info

Modified Sequence	AcD3-K<AcD3K*>EM<Mox*>IK<AcD3K*>SGM<Mox*>NVAR<C13>-COOH	Ion score	75.24
Protein	P52480	Confidence level	99.00%
Precursor m/z	768.91148	Identity Threshold	48.00
Charge	2+	Mass error	-9.03E-4



Info

Modified Sequence AcD3-VK<AcD3K*>M<Mox*>LPDK<AcD3K*>DC<Cmm*>R<C13>-COOH

Ion score 50.34

Protein P18760

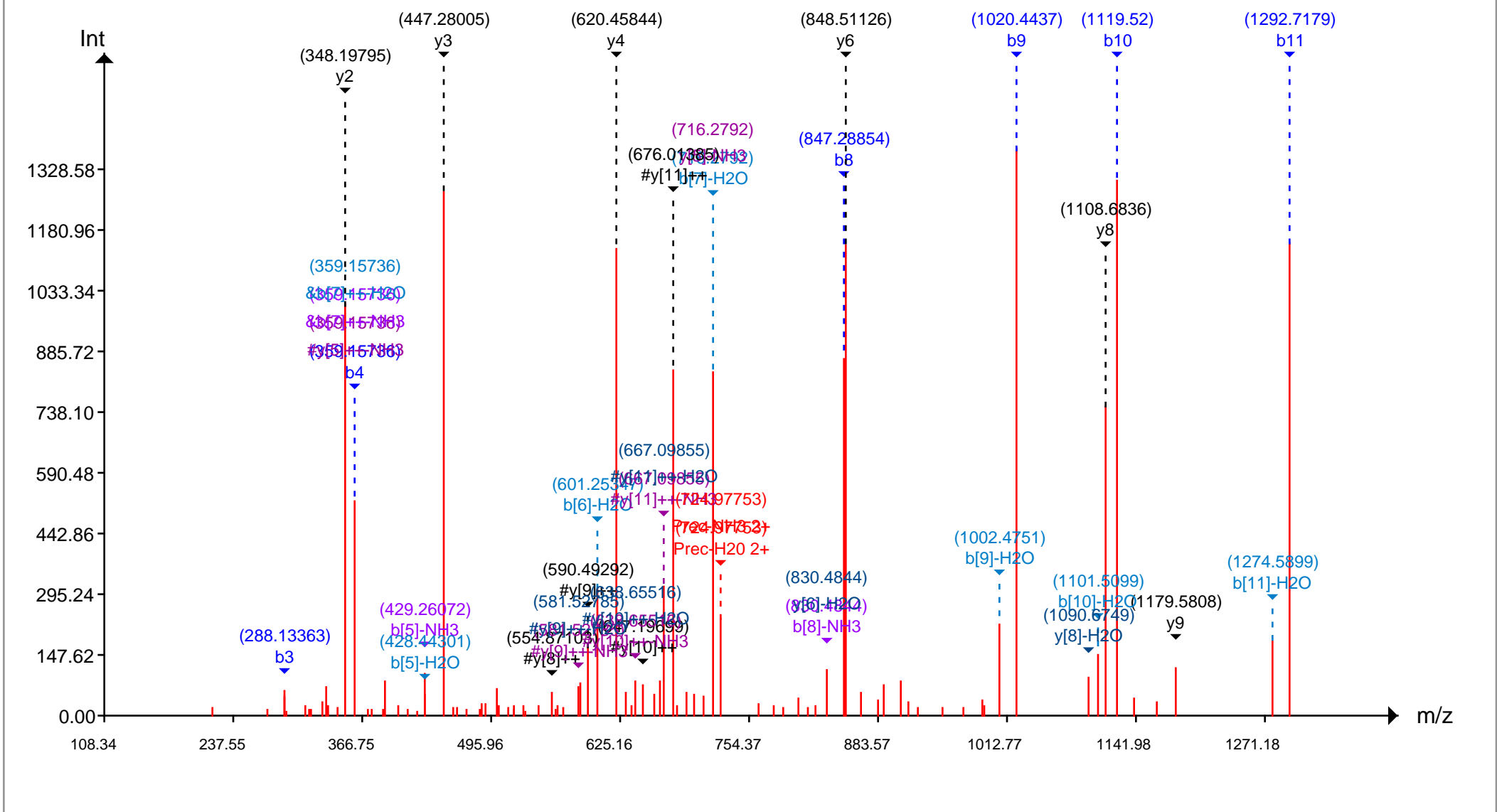
Confidence level 99.00%

Precursor m/z 709.87428

Identity Threshold 48.22

Charge 2+

Mass error -0.001099



Info

Modified Sequence AcD3-AGNASK<AcD3K*>DLK<AcD3K*>VK<AcD3K*>R-COOH

Ion score 56.14

Protein Q3THW5, P0C0S6

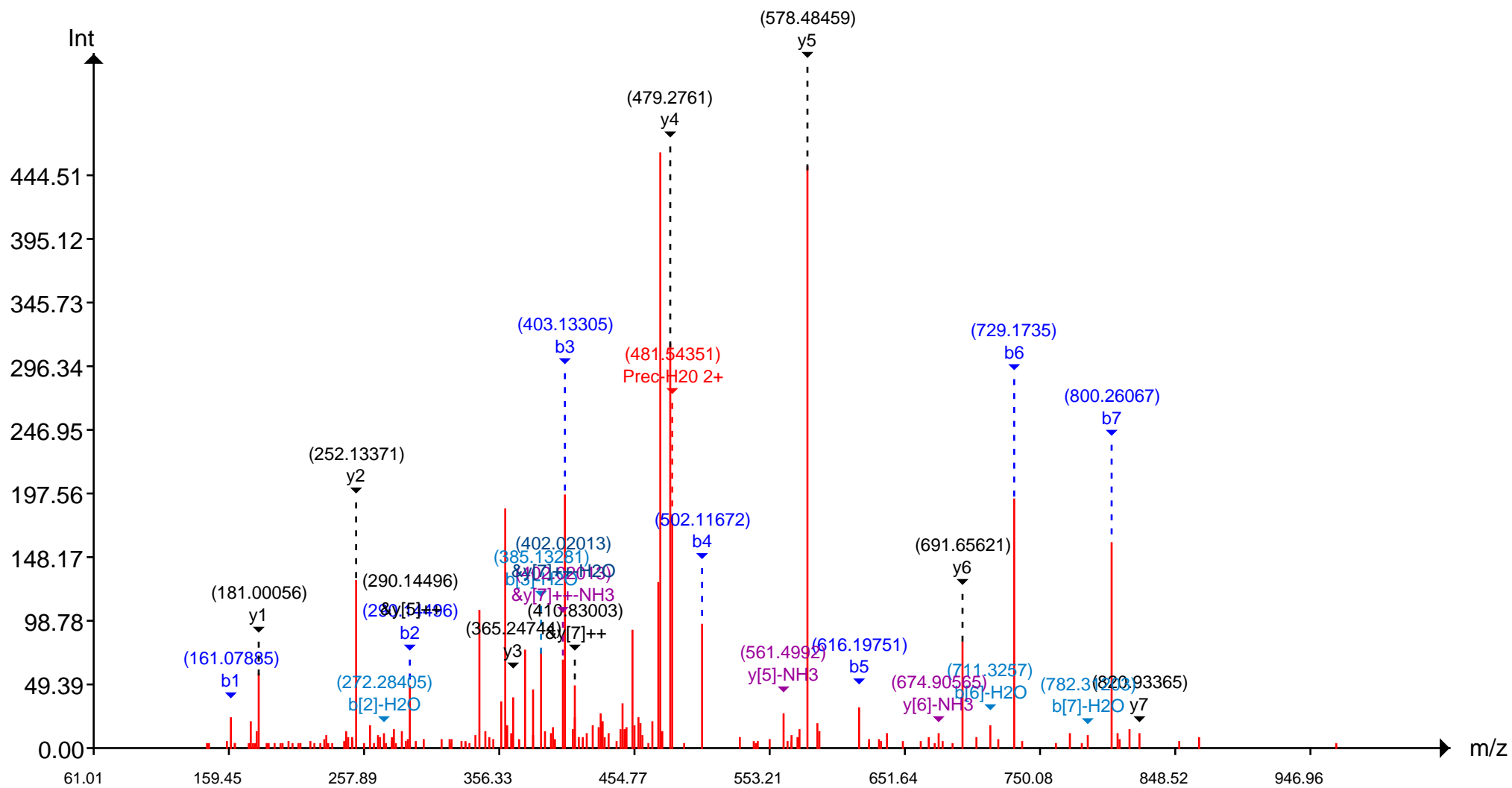
Confidence level 99.00%

Precursor m/z 733.93919

Identity Threshold 43.99

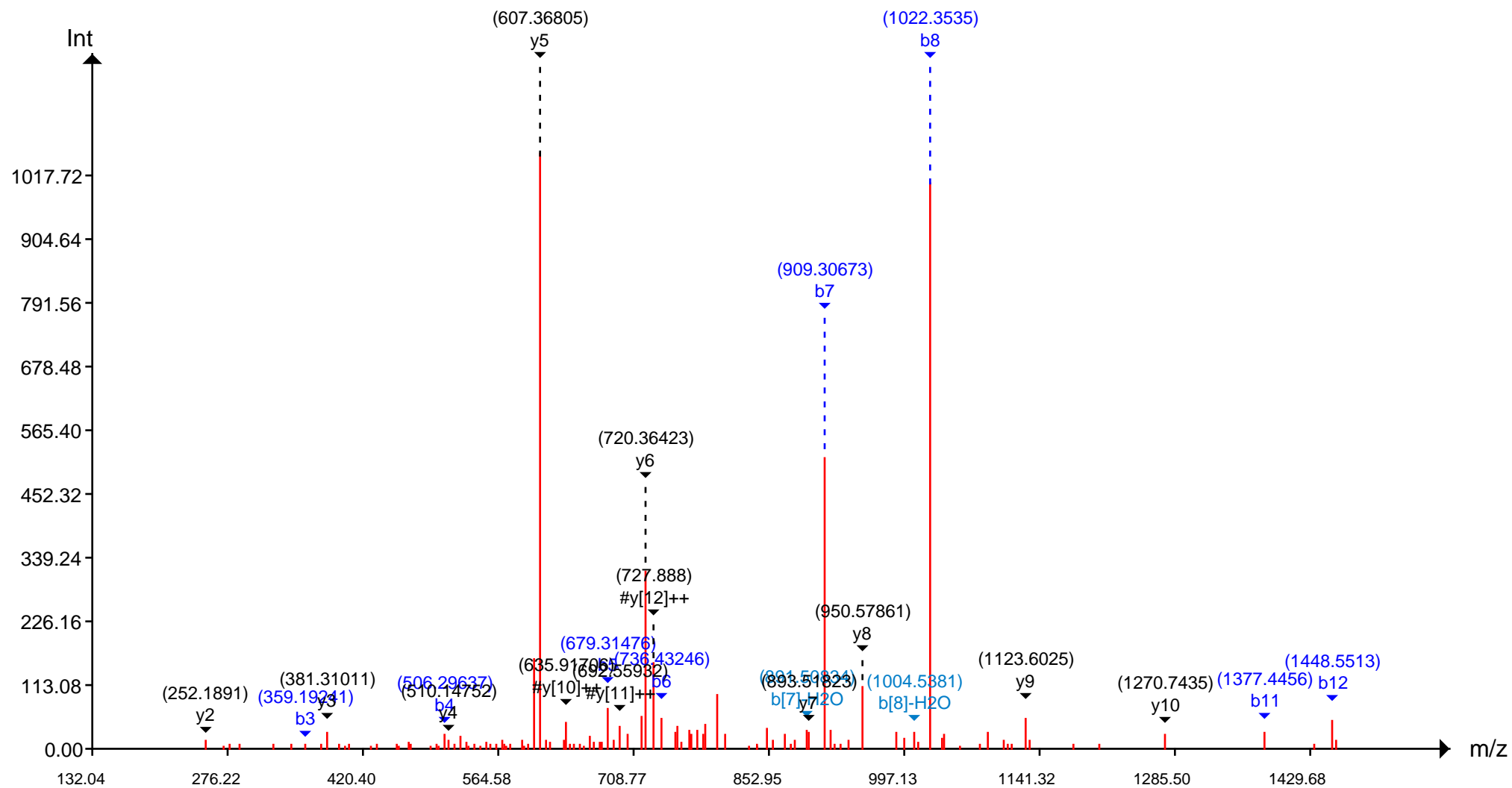
Charge 2+

Mass error -3.69E-4



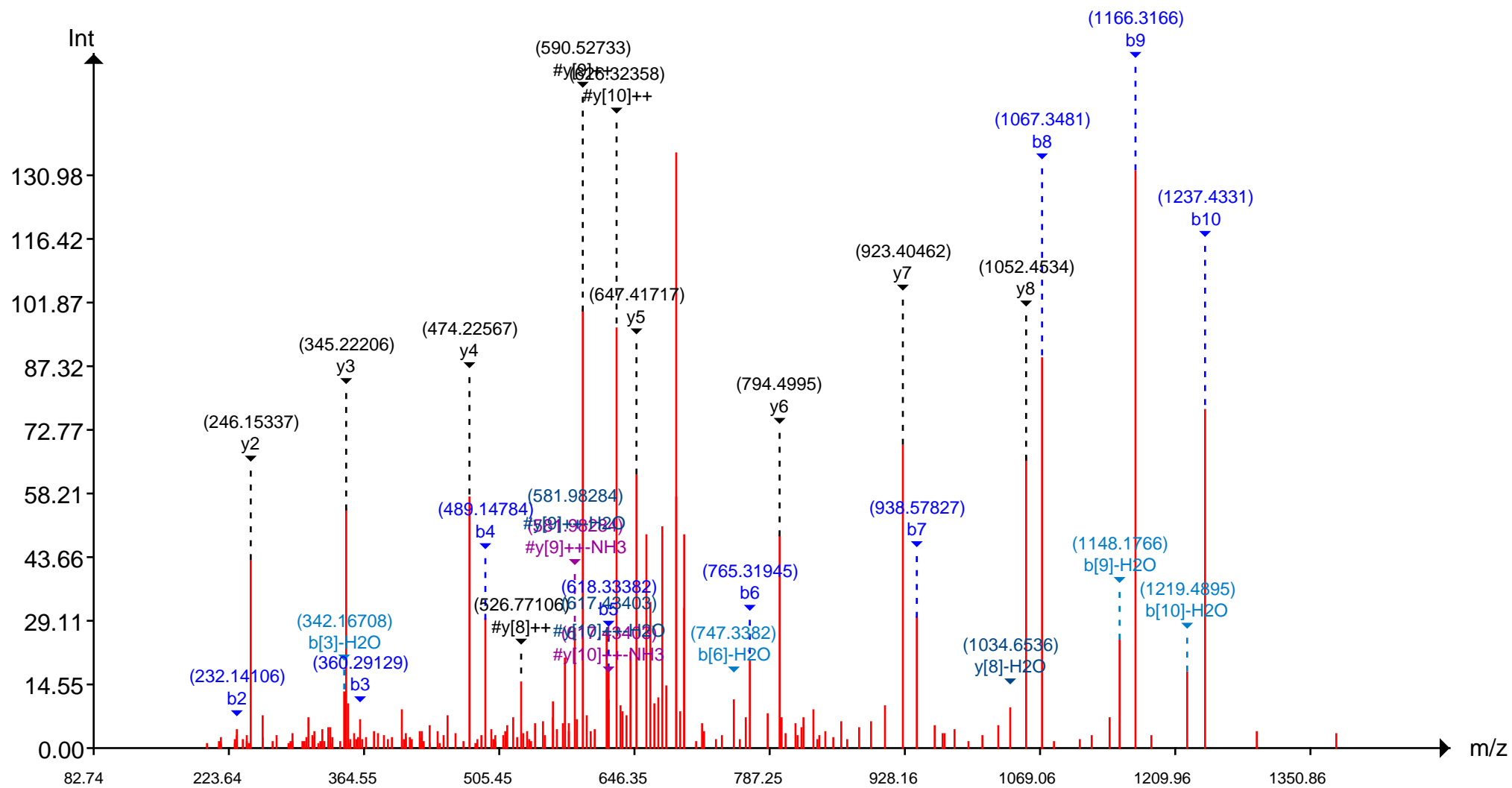
Info

Modified Sequence	AcD3-DEIVNIAR<C13>-COOH	Ion score	48.11
Protein	P35979	Confidence level	99.00%
Precursor m/z	490.78086	Identity Threshold	47.26
Charge	2+	Mass error	-1.34E-4



Info

Modified Sequence	AcD3-EALM<Mox*>K<AcD3K*>GK<AcD3K*>LPEEAR<C13>-COOH	Ion score	48.94
Protein	P06745	Confidence level	99.00%
Precursor m/z	814.95306	Identity Threshold	47.25
Charge	2+	Mass error	0.00192



Info

Modified Sequence AcD3-DAQEEM<Mox*>K<AcD3K*>EVAR-COOH

Ion score 54.42

Protein Q8CGC7

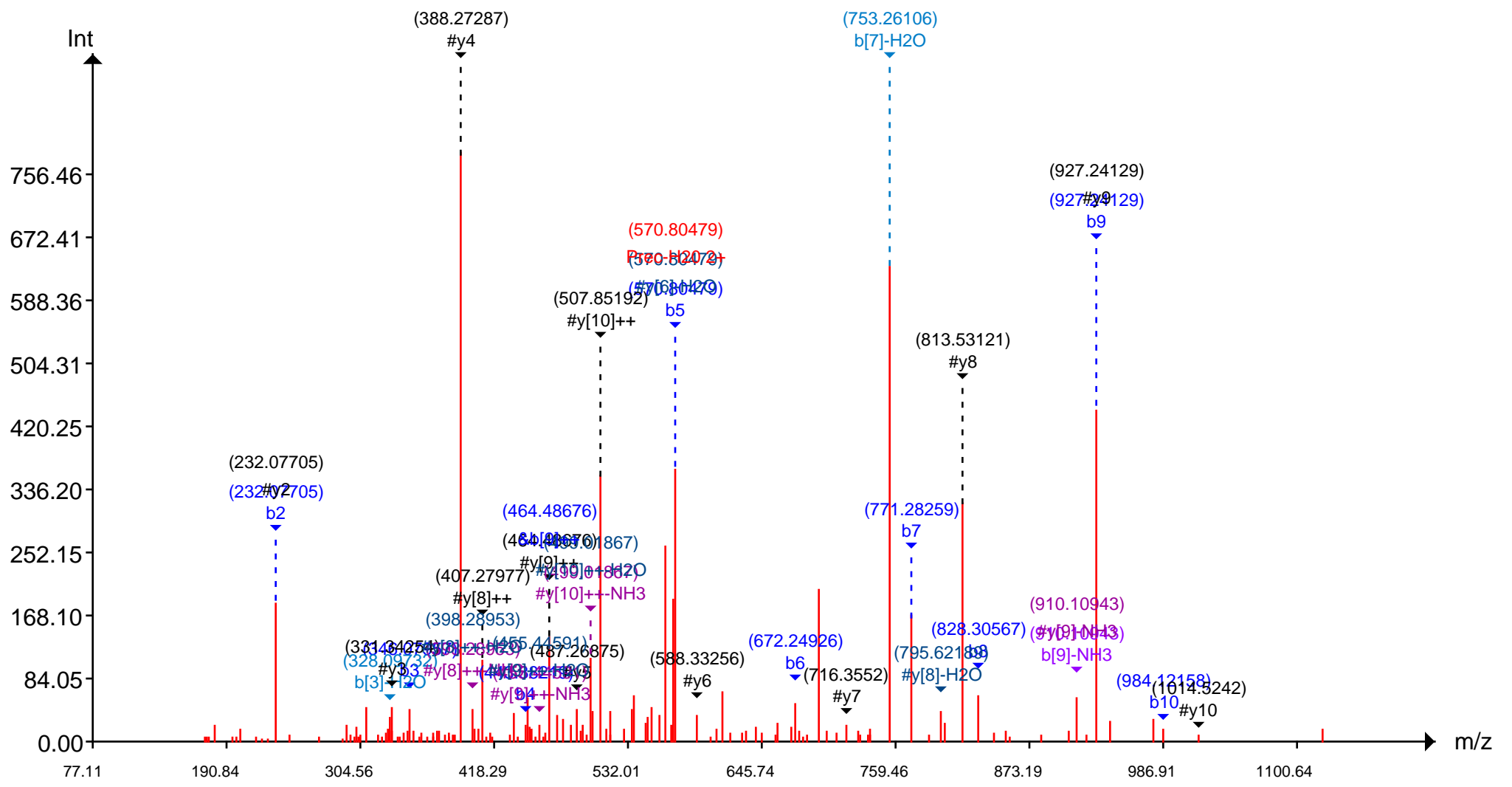
Confidence level 99.00%

Precursor m/z 706.33612

Identity Threshold 46.17

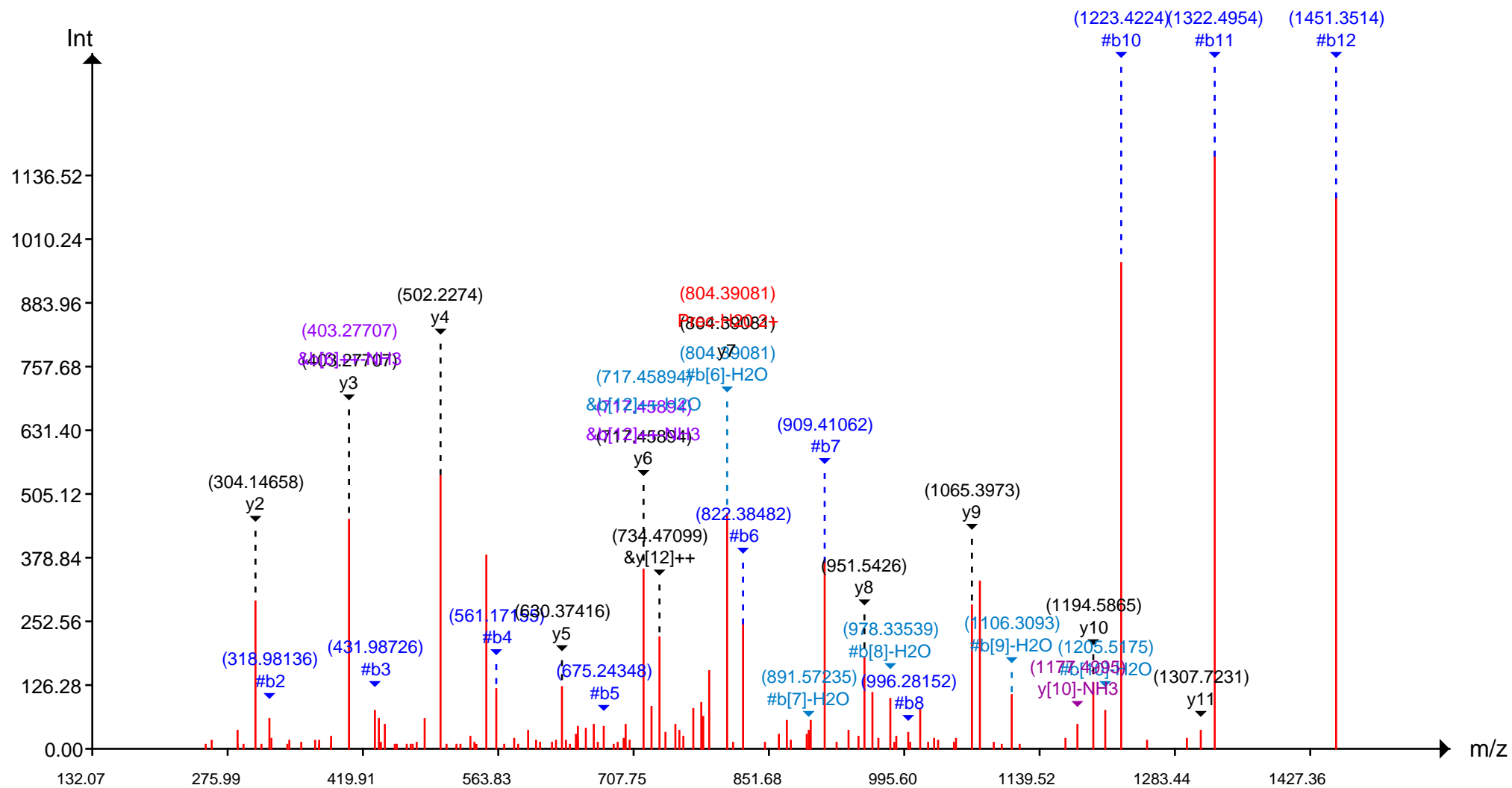
Charge 2+

Mass error 9.5E-4



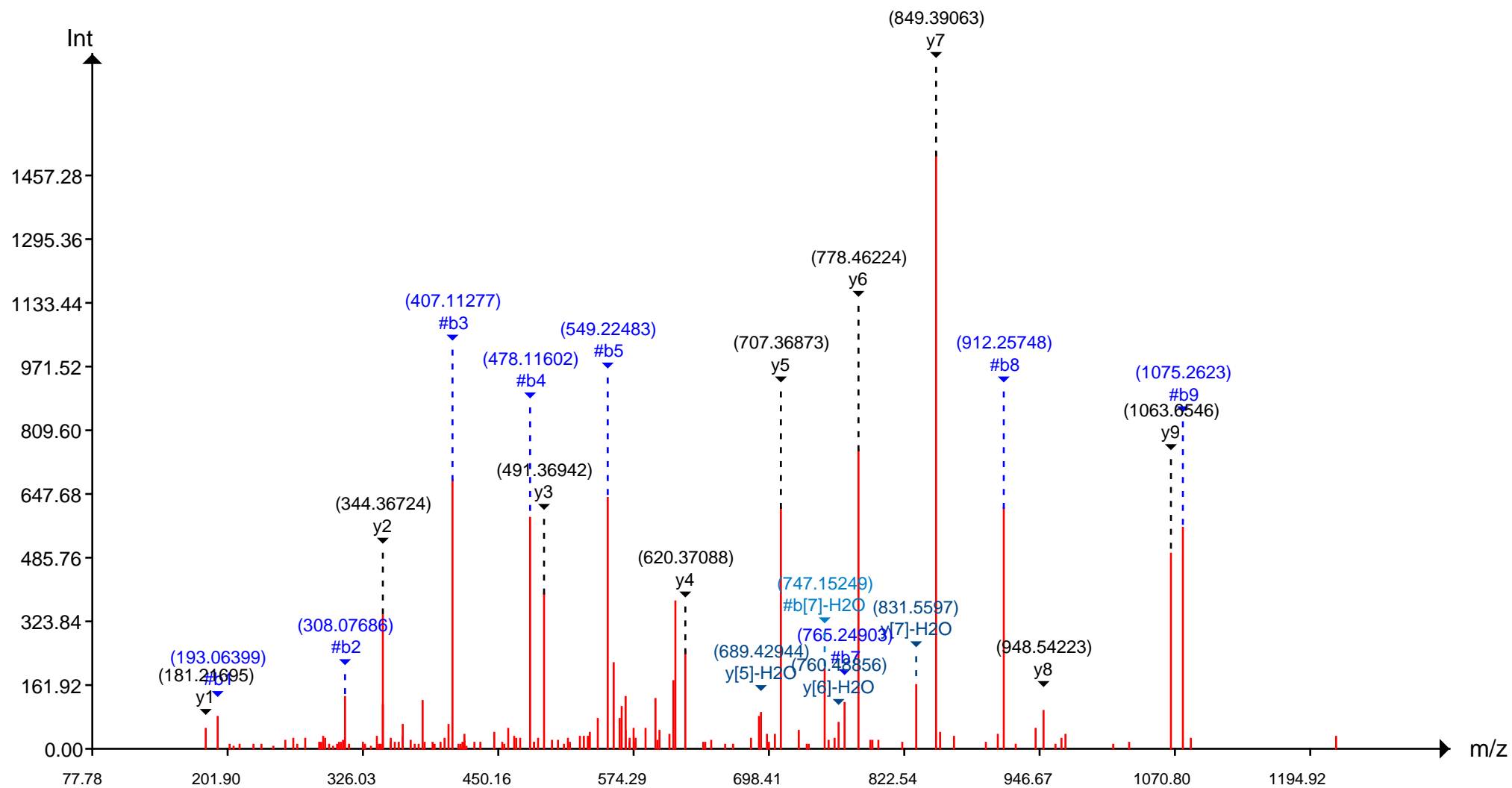
Info

Modified Sequence	AcD3-VSNPQTVGVGR-COOH	Ion score	55.49
Protein	O70492	Confidence level	99.00%
Precursor m/z	579.81838	Identity Threshold	47.82
Charge	2+	Mass error	-0.001022



Info

Modified Sequence	AcD3-LC<Cmm*>LENFSSQVVER-COOH	Ion score	92.47
Protein	P59999	Confidence level	99.00%
Precursor m/z	813.40693	Identity Threshold	47.57
Charge	2+	Mass error	0.003486



Info

Modified Sequence AcD3-M<Mox*>DVAASEFYR<C13>-COOH

Ion score 93.2

Protein P17182, P21550, P17183

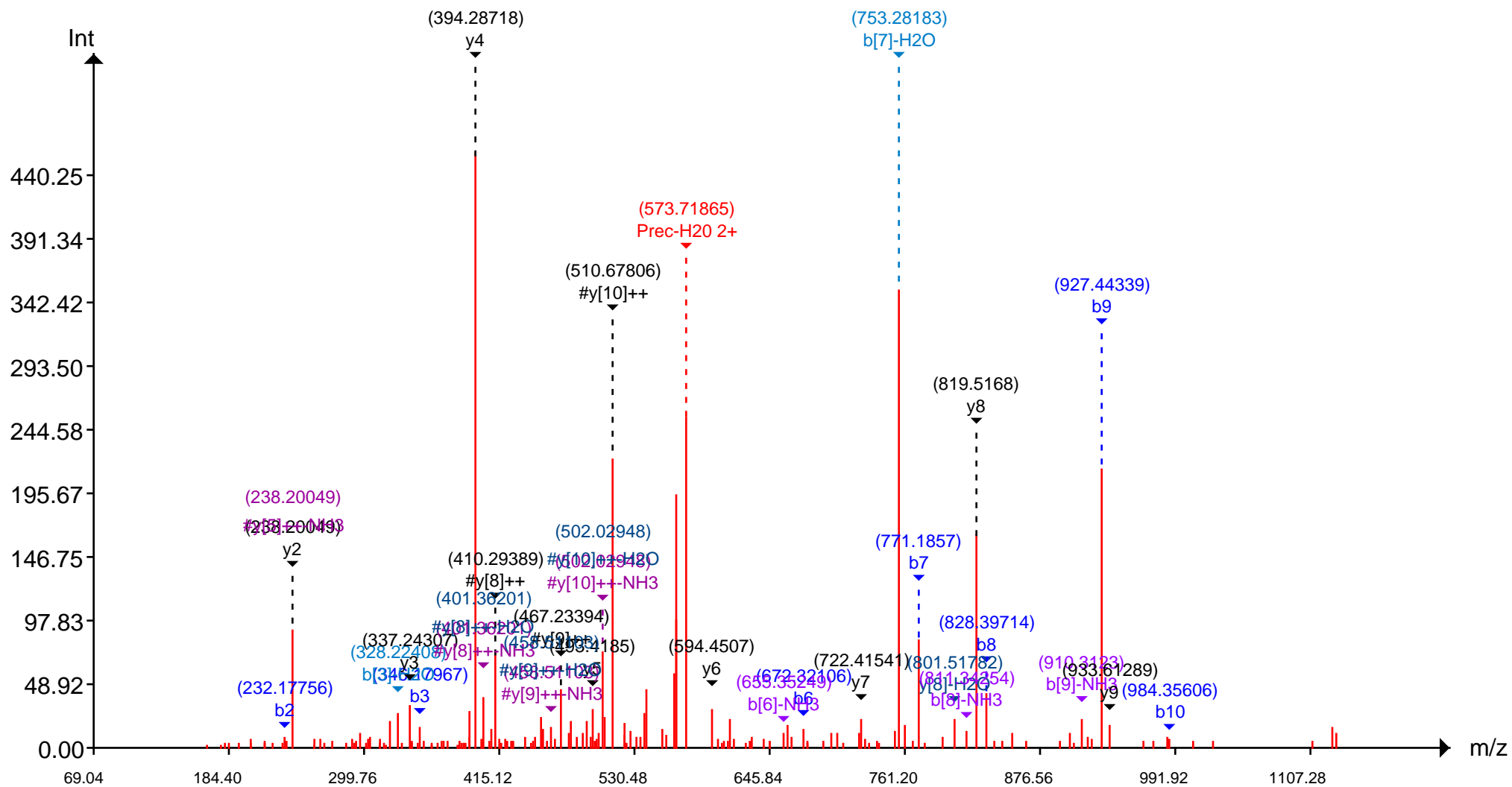
Confidence level 99.00%

Precursor m/z 628.29377

Identity Threshold 45.46

Charge 2+

Mass error 4.48E-4



Info

Modified Sequence AcD3-VSNPQTVGVGR<C13>-COOH

Ion score 53.63

Protein O70492

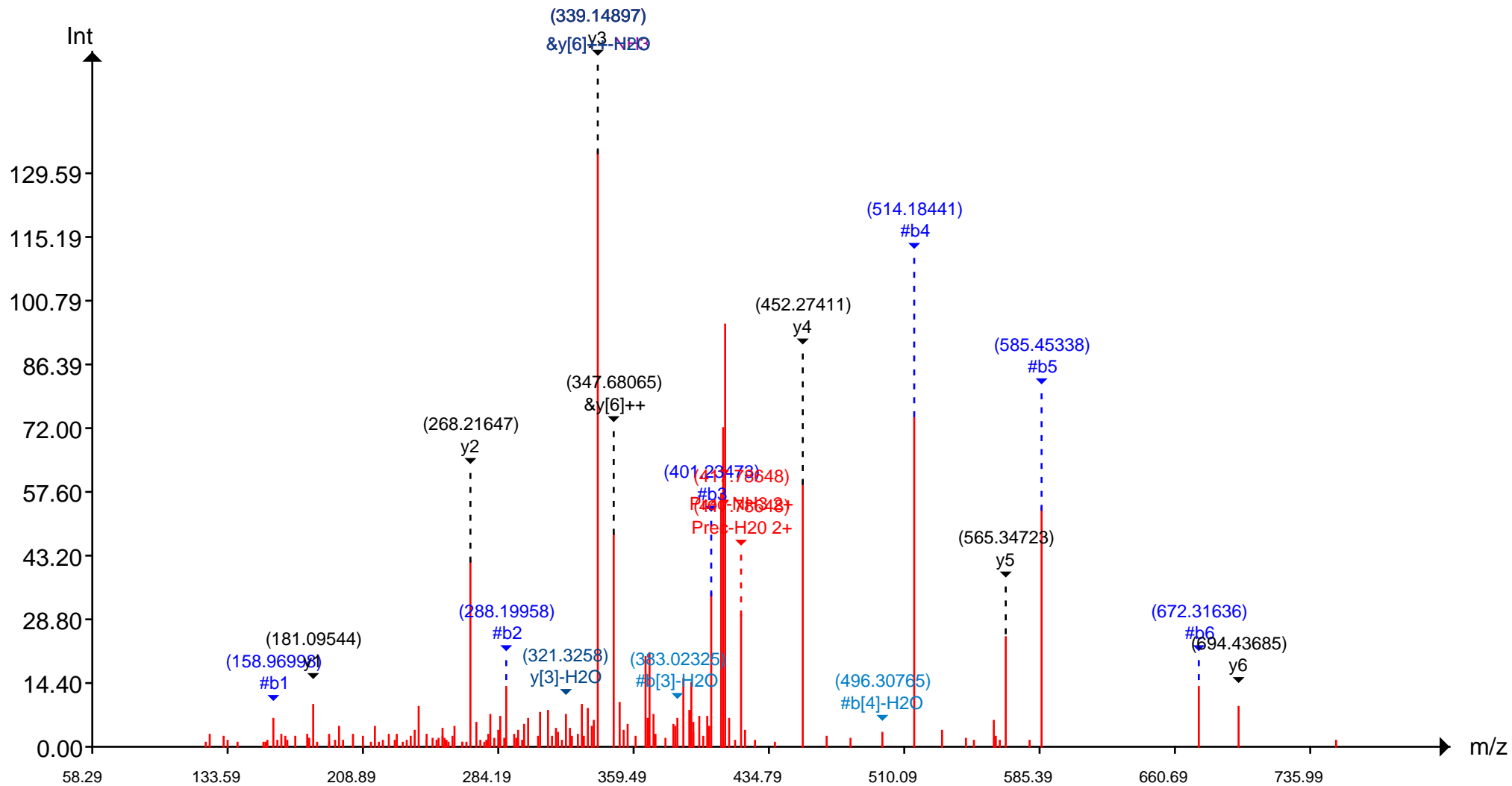
Confidence level 99.00%

Precursor m/z 582.82842

Identity Threshold 47.56

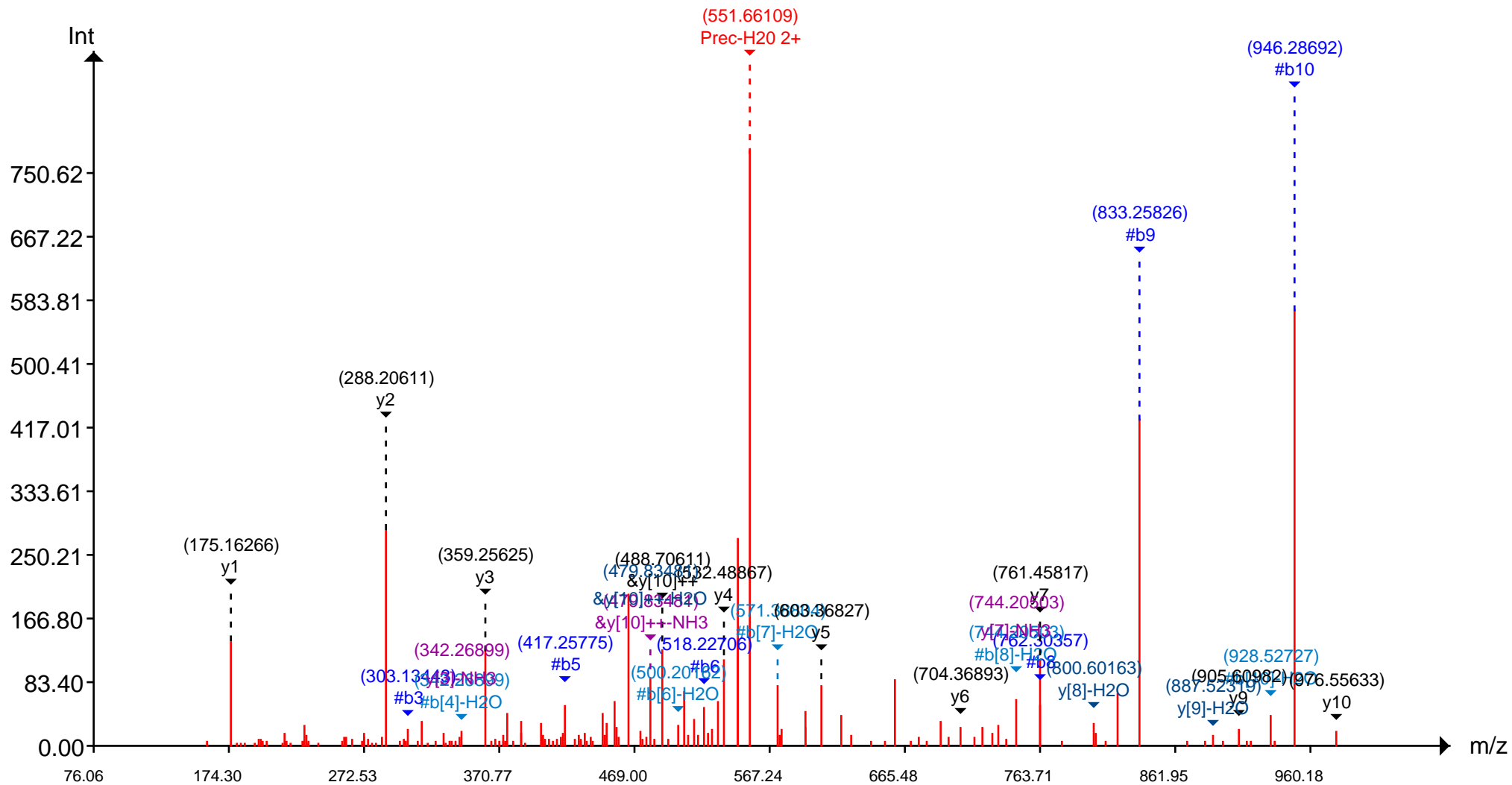
Charge 2+

Mass error -0.001084



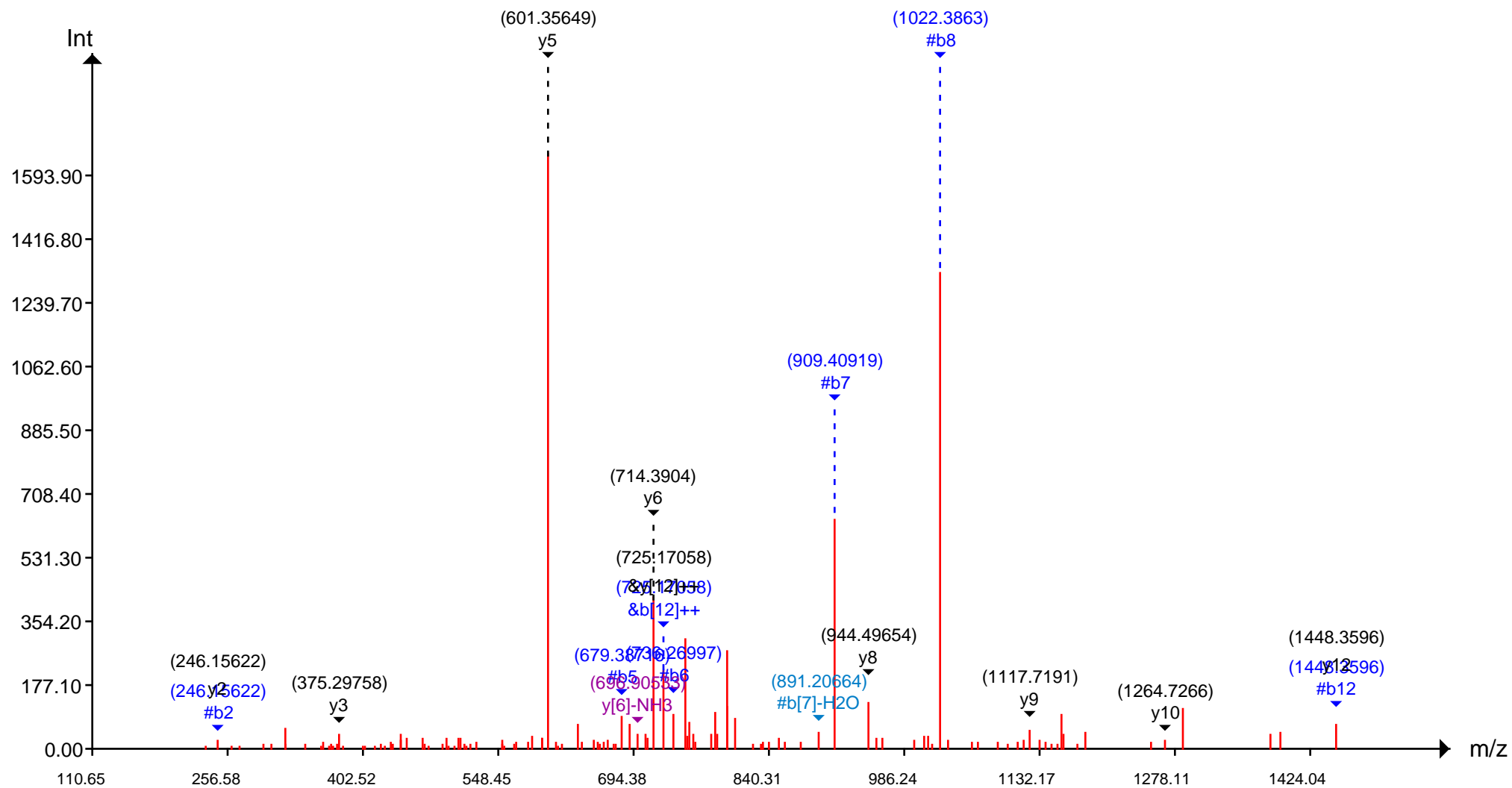
Info

Modified Sequence	AcD3-IEILASR<C13>-COOH	Ion score	63.06
Protein	Q9QZ10, P97429, P14824, O35640	Confidence level	99.00%
Precursor m/z	426.76982	Identity Threshold	42.85
Charge	2+	Mass error	-1.3E-5



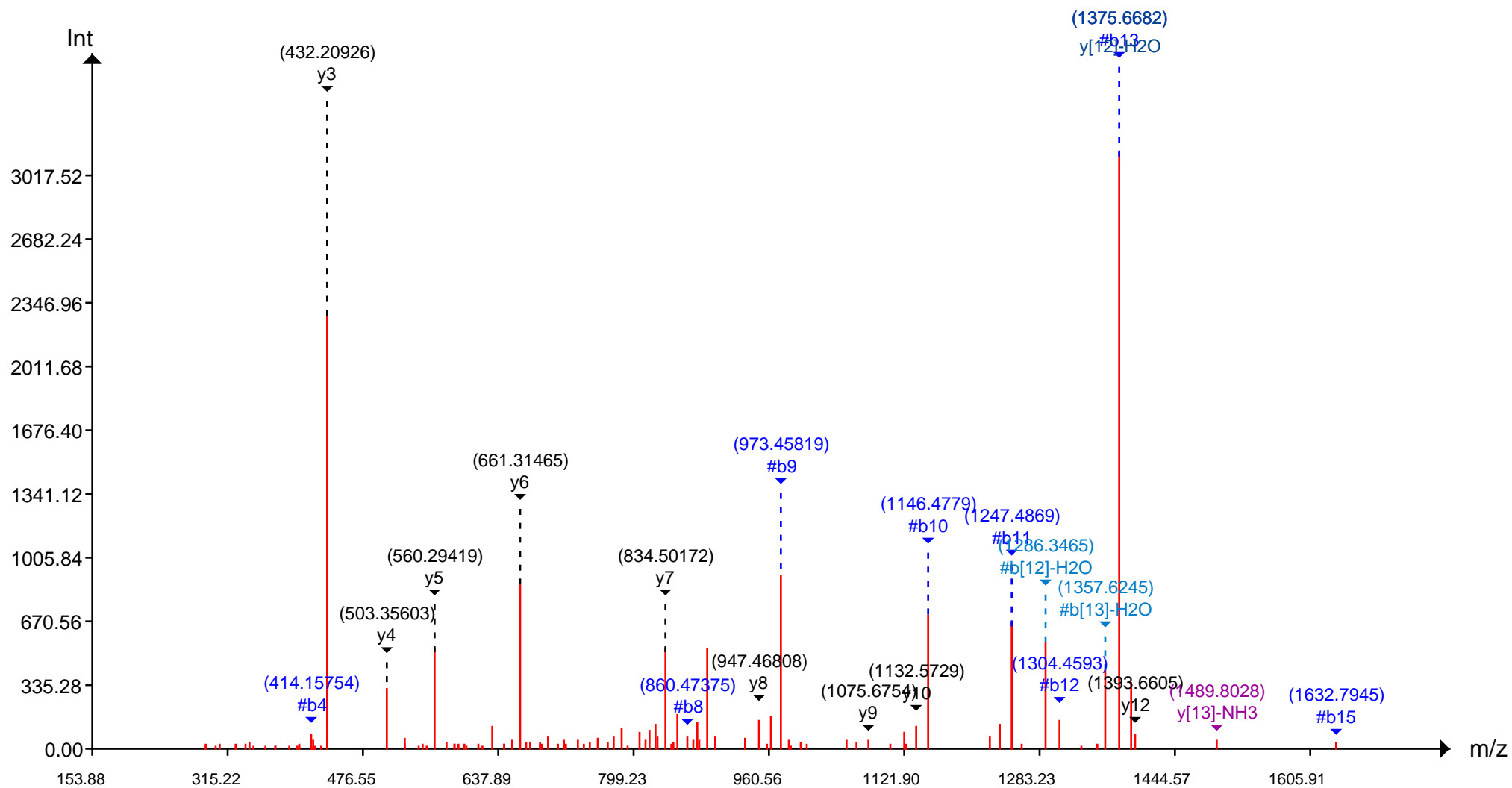
Info

Modified Sequence	AcD3-VASGGTAK<AcD3K*>AIR-COOH	Ion score	54.32
Protein	Q9CWJ9	Confidence level	99.00%
Precursor m/z	560.83308	Identity Threshold	46.04
Charge	2+	Mass error	-2.63E-4



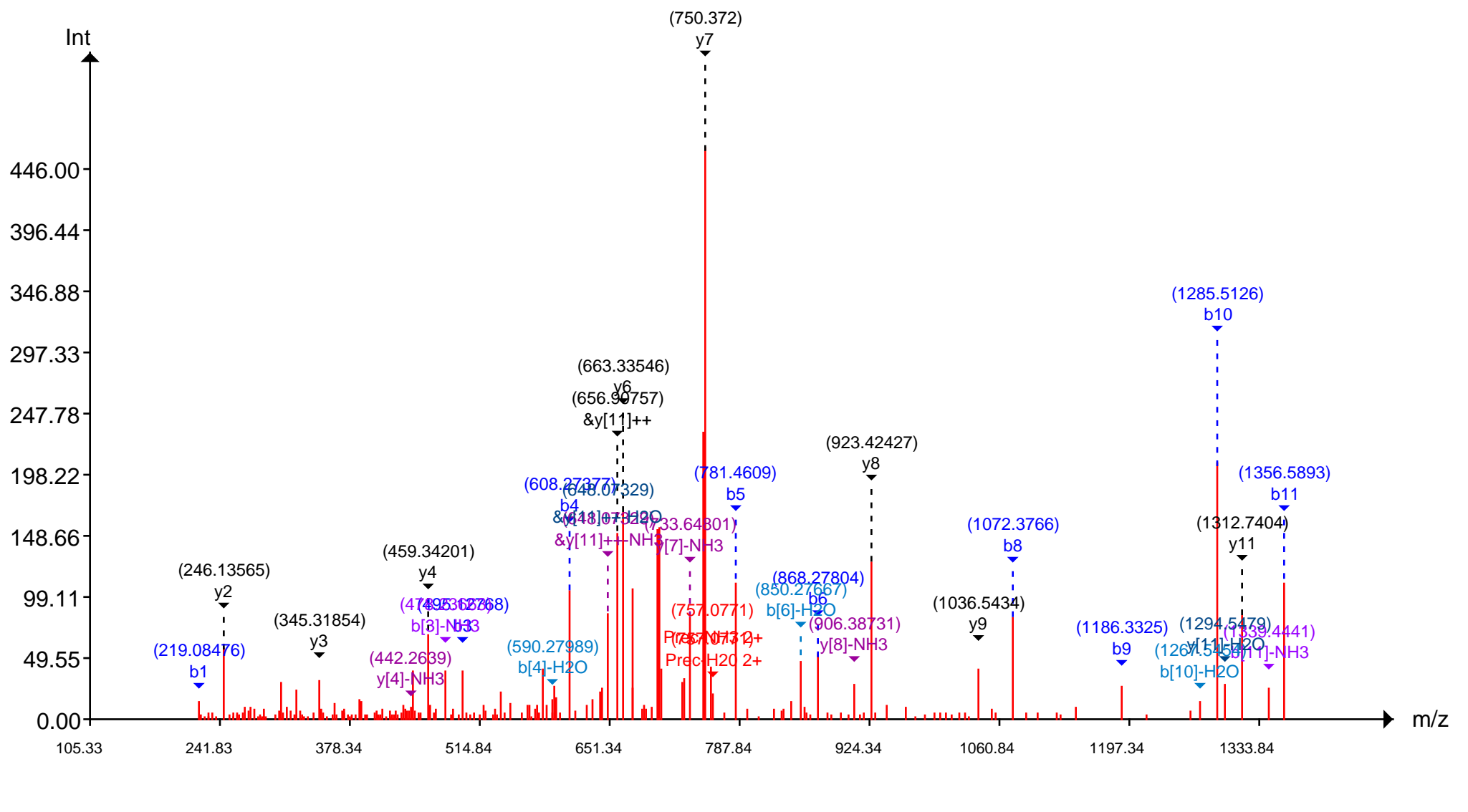
Info

Modified Sequence	AcD3-EALM<Mox*>K<AcD3K*>GK<AcD3K*>LPEEAR-COOH	Ion score	48.67
Protein	P06745	Confidence level	99.00%
Precursor m/z	811.94298	Identity Threshold	47.73
Charge	2+	Mass error	0.001901

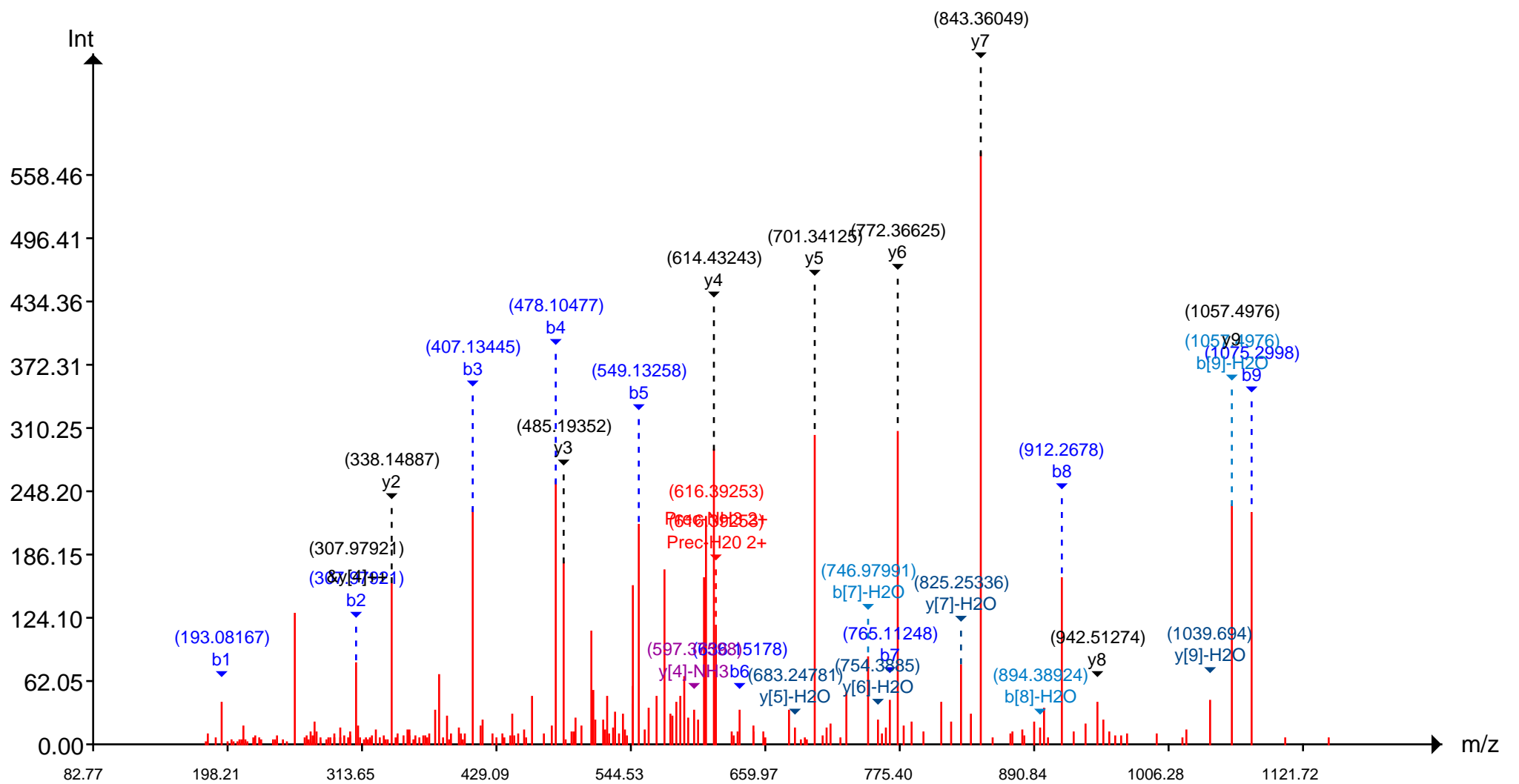


Info

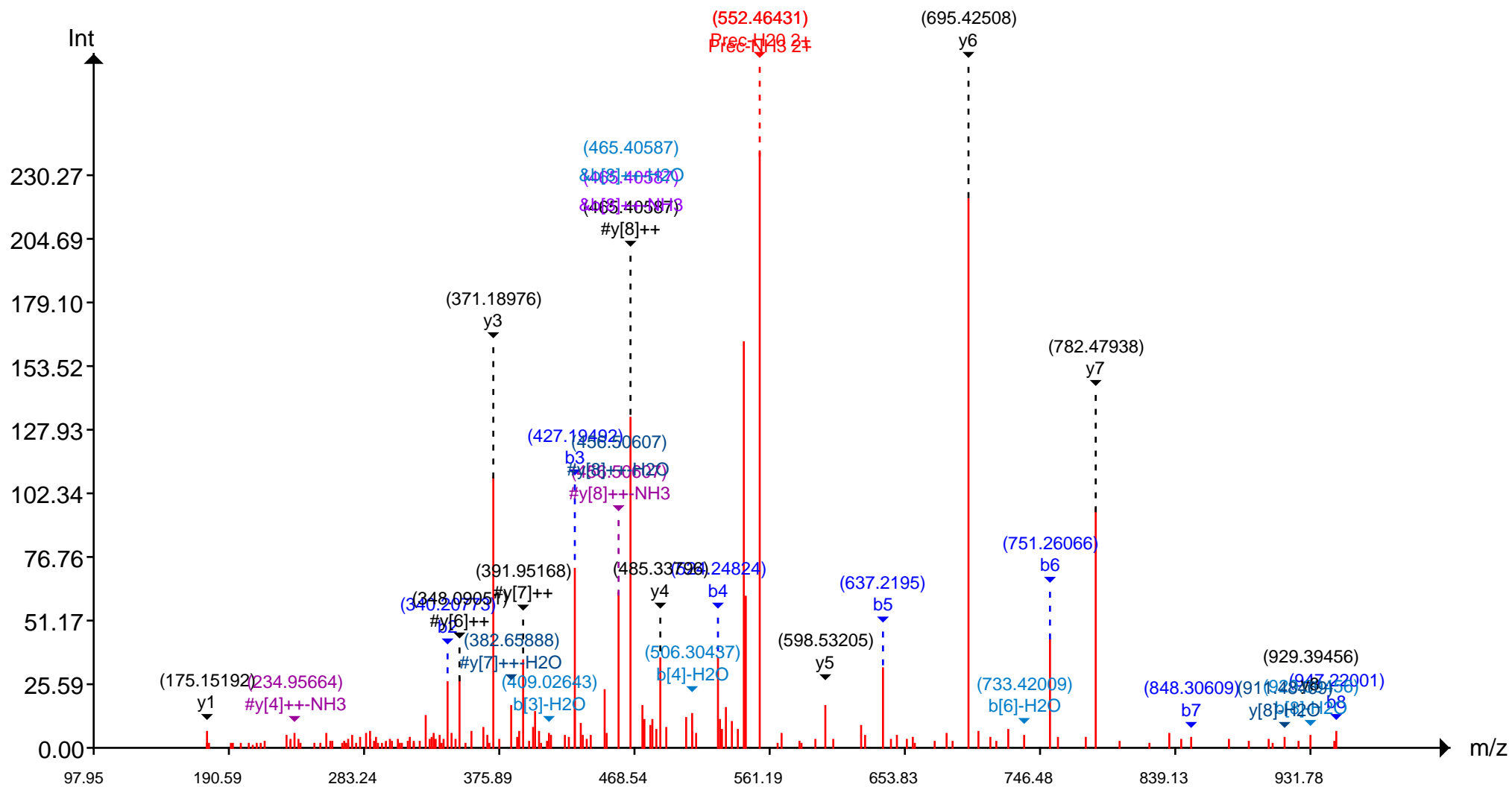
Modified Sequence	AcD3-VVGLC<Cmm*>TGQIK<AcD3K*>TGAPC<Cmm*>R-COOH	Ion score	58.21
Protein	P17182, P21550, P17183	Confidence level	99.00%
Precursor m/z	903.9781	Identity Threshold	48.03
Charge	2+	Mass error	0.001737



Info			
Modified Sequence	AcD3-K<AcD3K*>EM<Mox*>IK<AcD3K*>SGM<Mox*>NVAR-COOH	Ion score	66.67
Protein	P52480	Confidence level	99.00%
Precursor m/z	765.902	Identity Threshold	48.12
Charge	2+	Mass error	2.79E-4

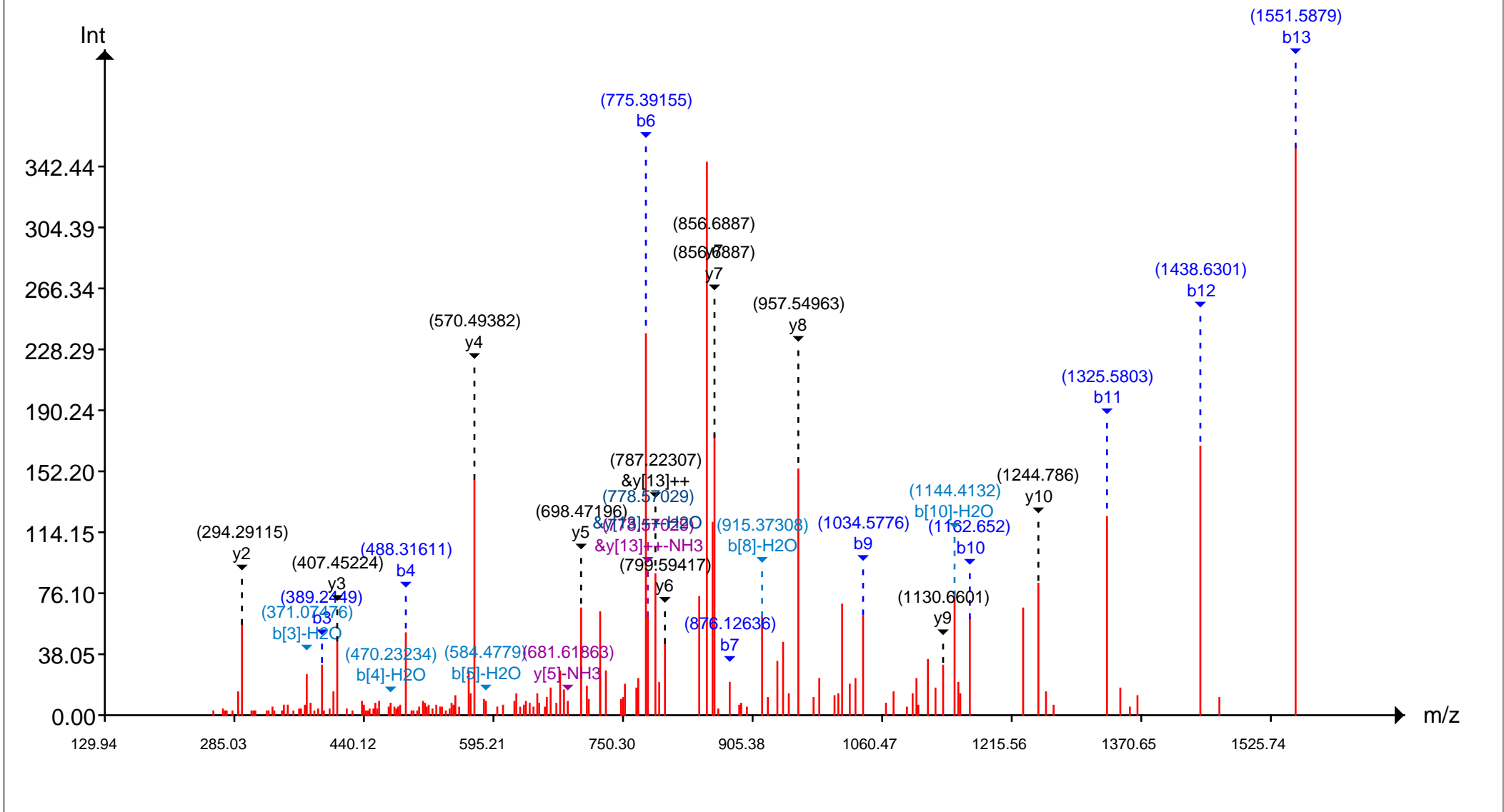


Info		Ion score	77.7
Modified Sequence	AcD3-M<Mox*>DVAASEFYR-COOH	Confidence level	99.00%
Protein	P17182, P21550, P17183	Identity Threshold	44.28
Precursor m/z	625.28393	Mass error	9.09E-4
Charge	2+		



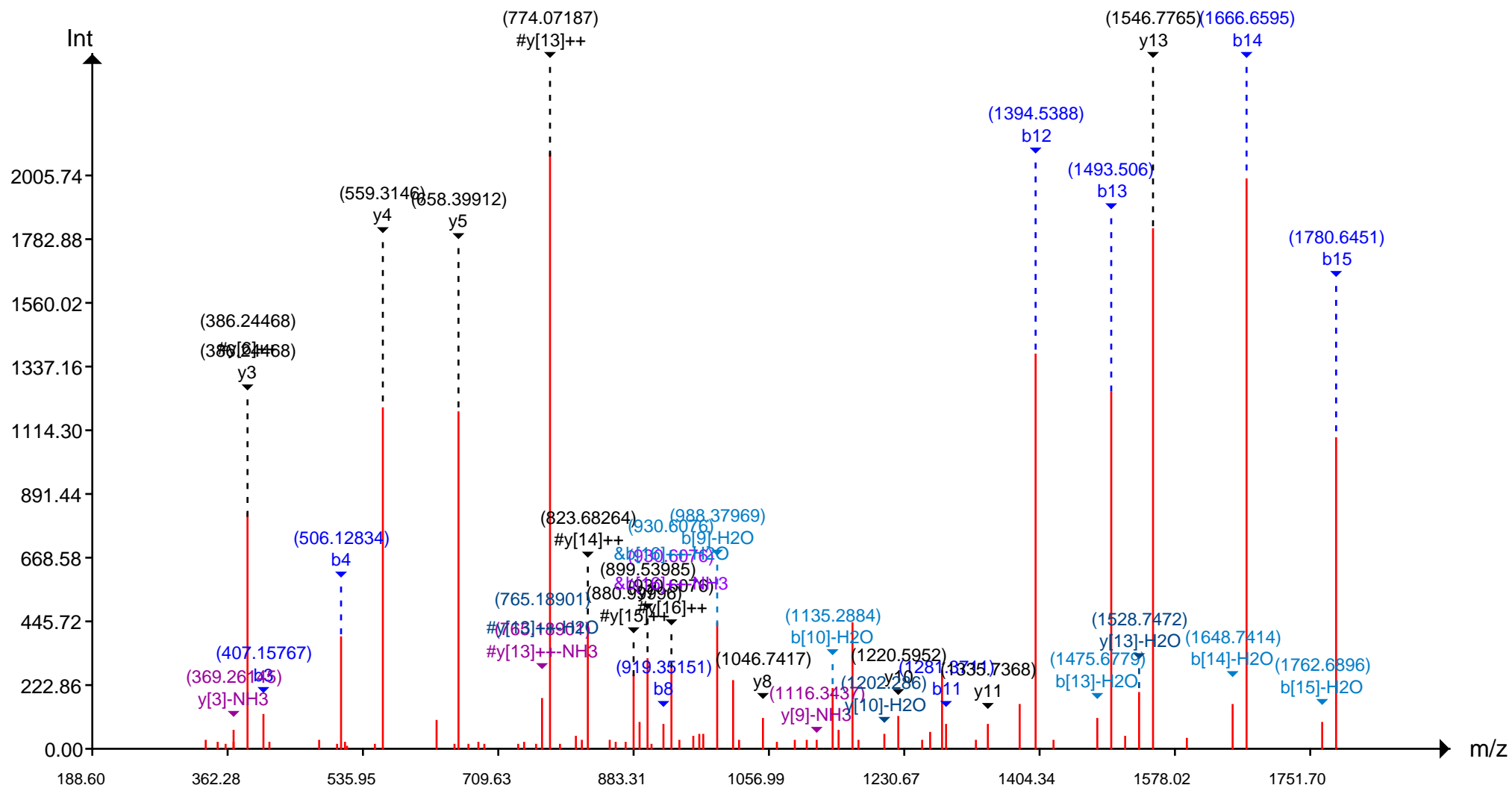
Info

Modified Sequence	AcD3-FFSPLNPVR-COOH	Ion score	55.14
Protein	O35737, Q9Z2X1	Confidence level	99.00%
Precursor m/z	561.31281	Identity Threshold	47.60
Charge	2+	Mass error	2.13E-4



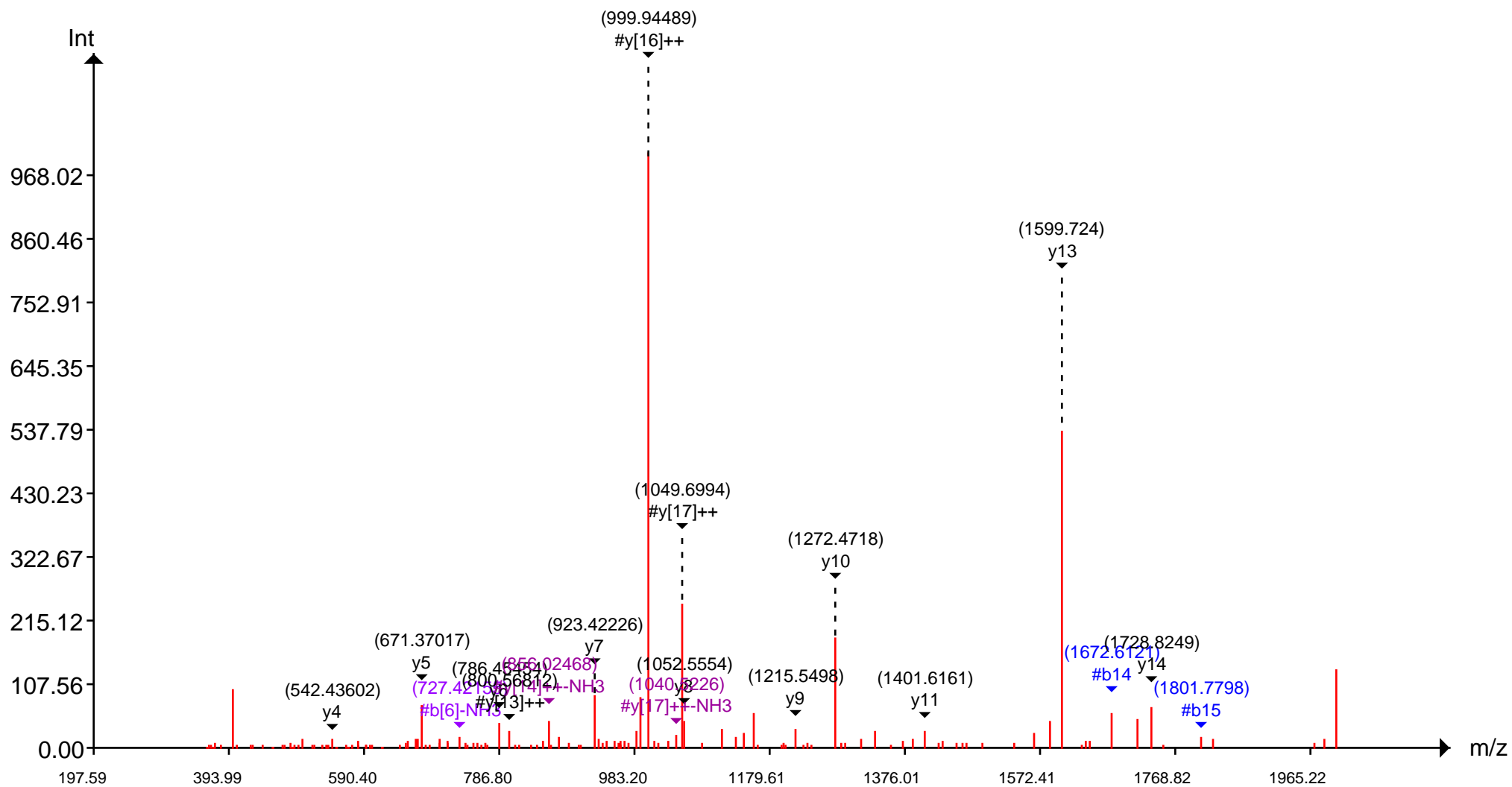
Info

Modified Sequence	AcD3-LTEVNK<AcD3K*>TGTQYLLR<C13>-COOH	Ion score	60.13
Protein	Q60854	Confidence level	99.00%
Precursor m/z	866.49658	Identity Threshold	45.82
Charge	2+	Mass error	5.2E-4



Info

Modified Sequence	AcD3-FVDVPNDSSFQIVK<AcD3K*>NPR-COOH	Ion score	58.3
Protein	Q61176	Confidence level	99.00%
Precursor m/z	1026.5386	Identity Threshold	48.00
Charge	2+	Mass error	0.003169



Info

Modified Sequence AcD3-AEVPK<AcD3K*>EPTEGYEHDETFLR<C13>-COOH

Ion score 51.67

Protein Q9DCG9

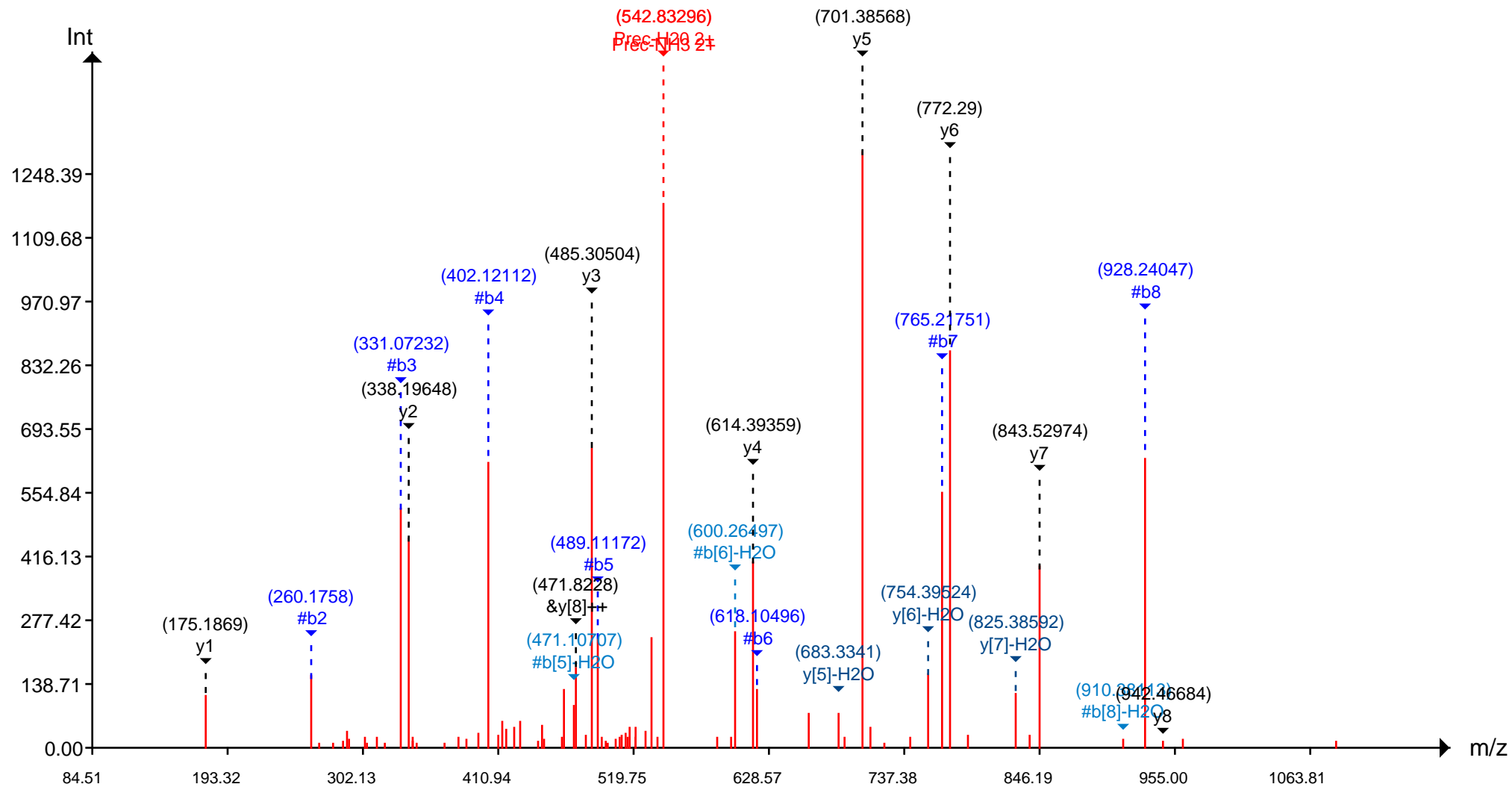
Confidence level 99.00%

Precursor m/z 1172.0738

Identity Threshold 47.08

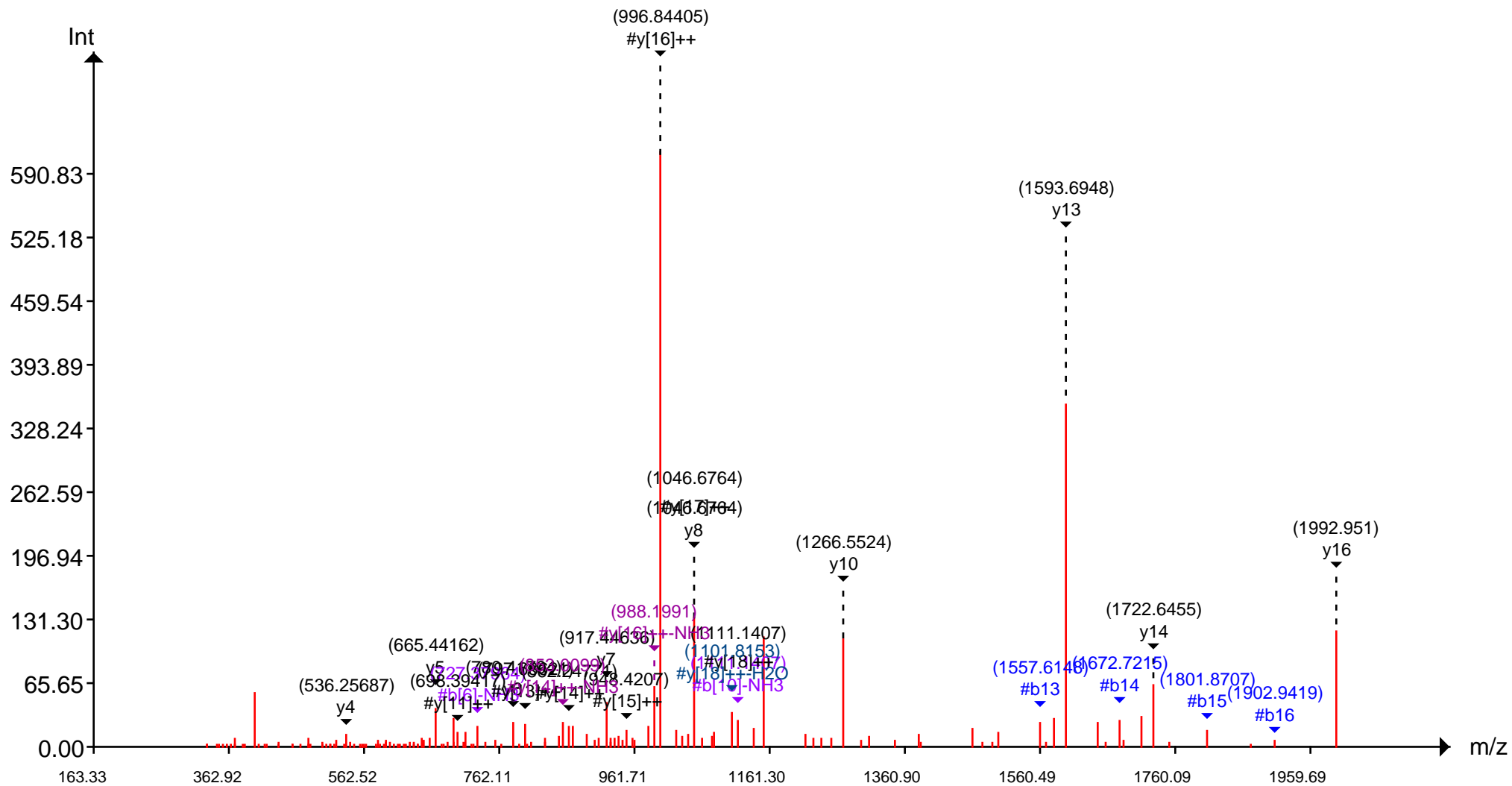
Charge 2+

Mass error 0.004996



Info

Modified Sequence	AcD3-DVAASEFYR-COOH	Ion score	66.57
Protein	P17182, P21550, P17183	Confidence level	99.00%
Precursor m/z	551.76574	Identity Threshold	46.19
Charge	2+	Mass error	-7.0E-5



Info

Modified Sequence AcD3-AEVPK<AcD3K*>EPTEGYEHDETFLR-COOH

Ion score 68.55

Protein Q9DCG9

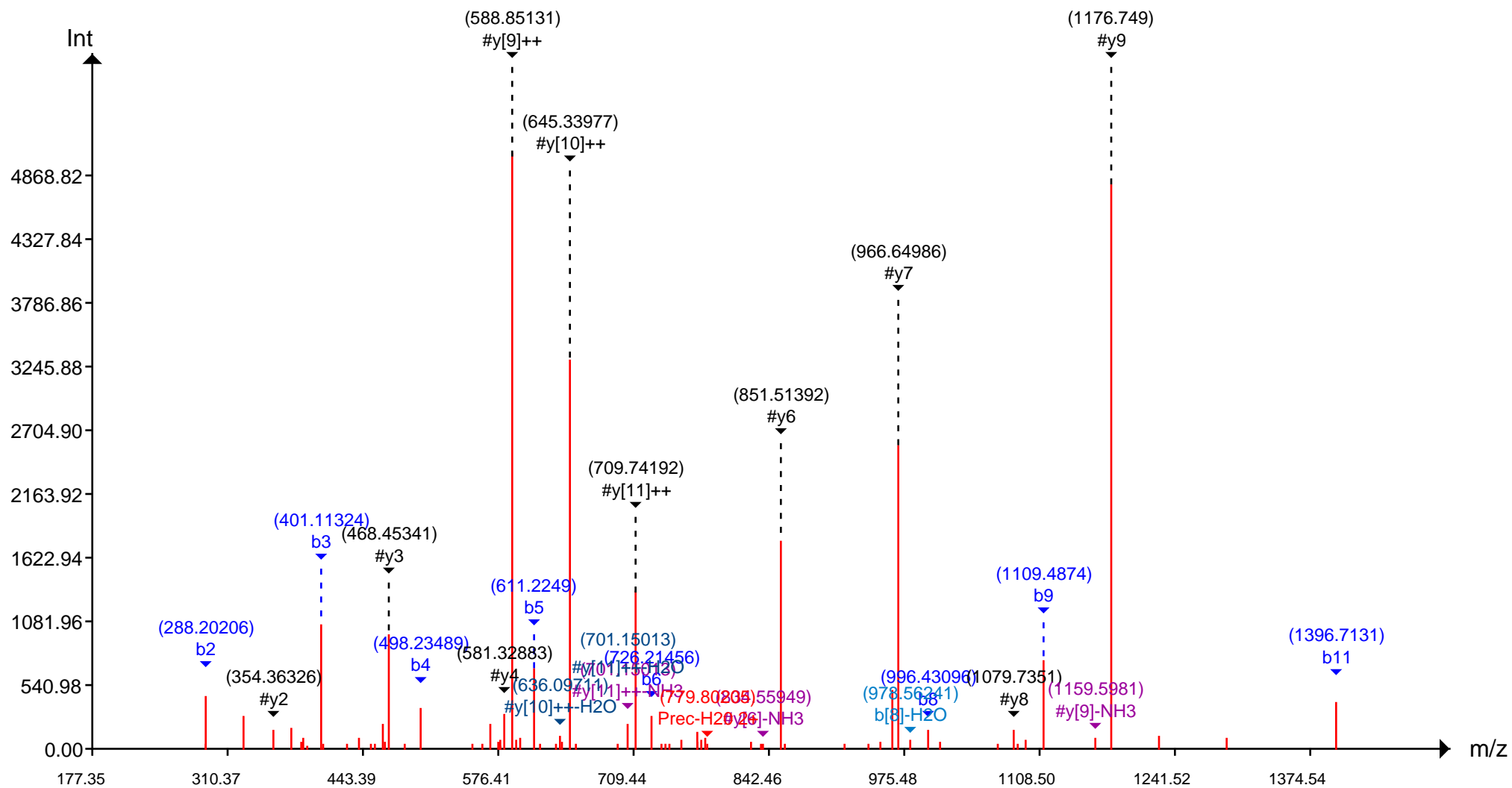
Confidence level 99.00%

Precursor m/z 1169.063

Identity Threshold 46.66

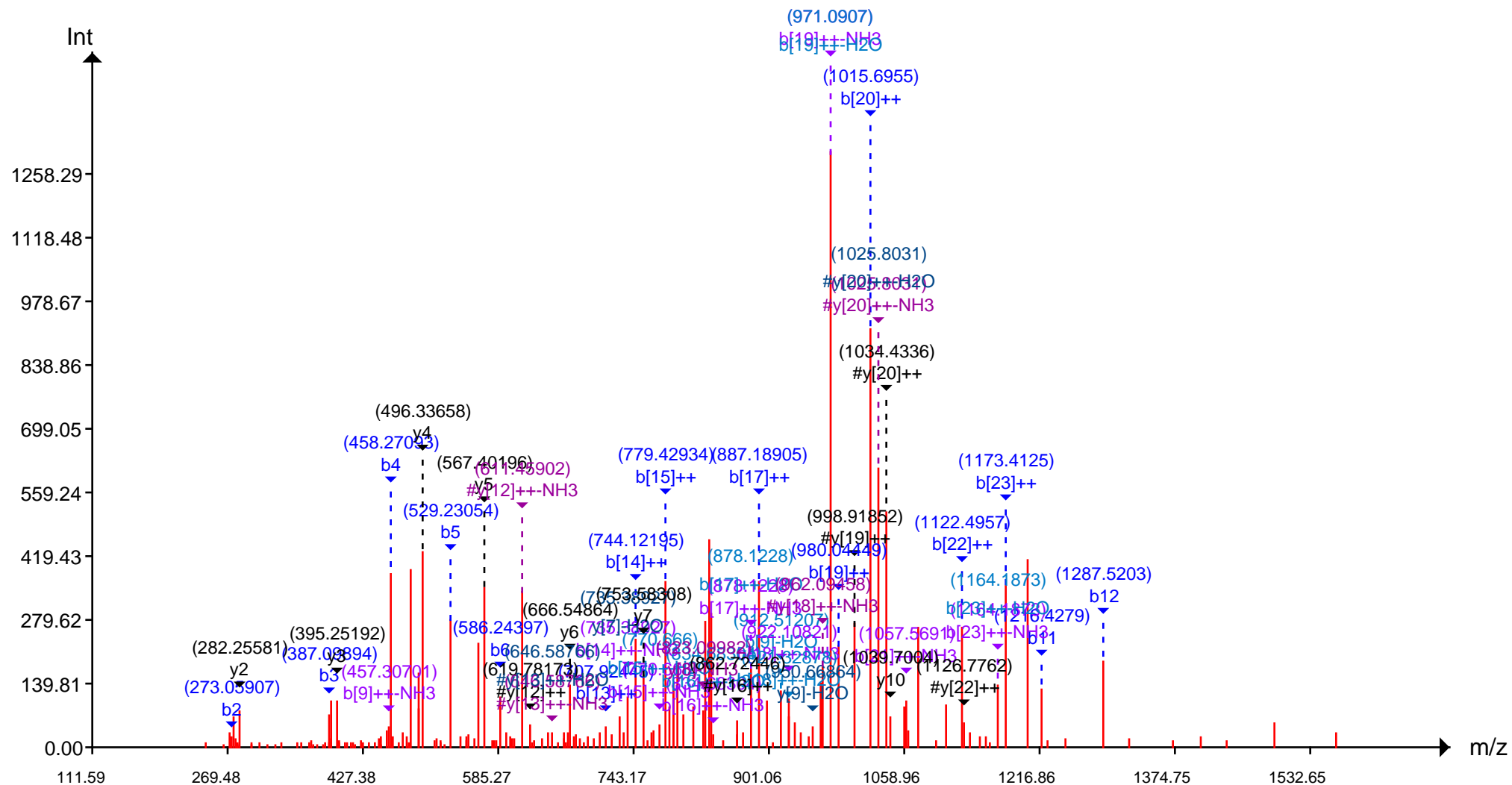
Charge 2+

Mass error 0.003538



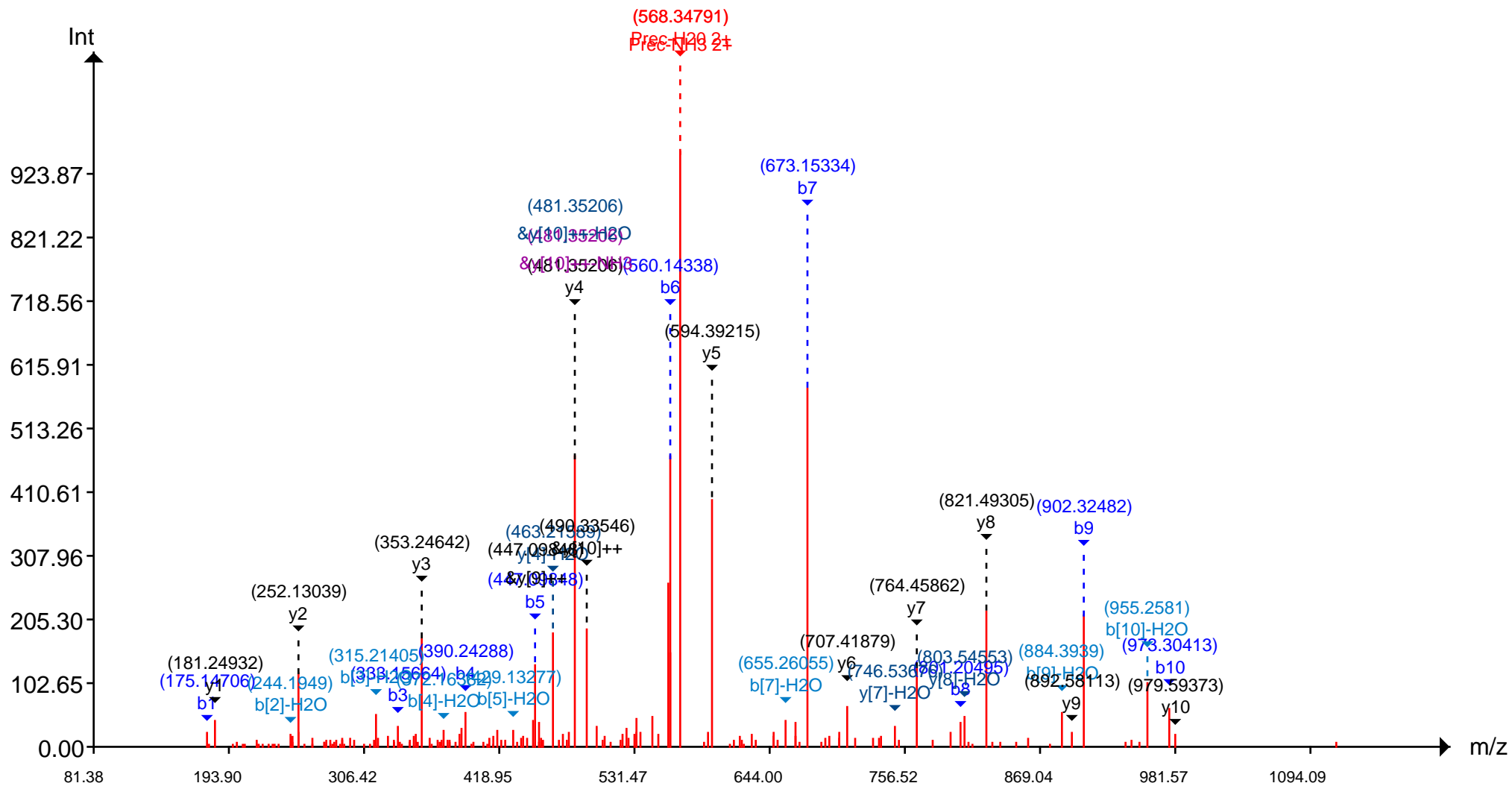
Info

Modified Sequence	AcD3-IELPIDPK<AcD3K*>LNK<AcD3K*>R<C13>-COOH	Ion score	57.92
Protein	Q99020	Confidence level	99.00%
Precursor m/z	788.9907	Identity Threshold	39.78
Charge	2+	Mass error	0.002706



Info

Modified Sequence	AcD3-VQNAAGSQEK<AcD3K*>LAQAAQSSVATITR<C13>-COOH	Ion score	48.41
Protein	P26039	Confidence level	99.00%
Precursor m/z	842.45666	Identity Threshold	47.14
Charge	3+	Mass error	0.002235



Info

Modified Sequence AcD3-ESAGGLIQTAR<C13>-COOH

Ion score 97.19

Protein P26039

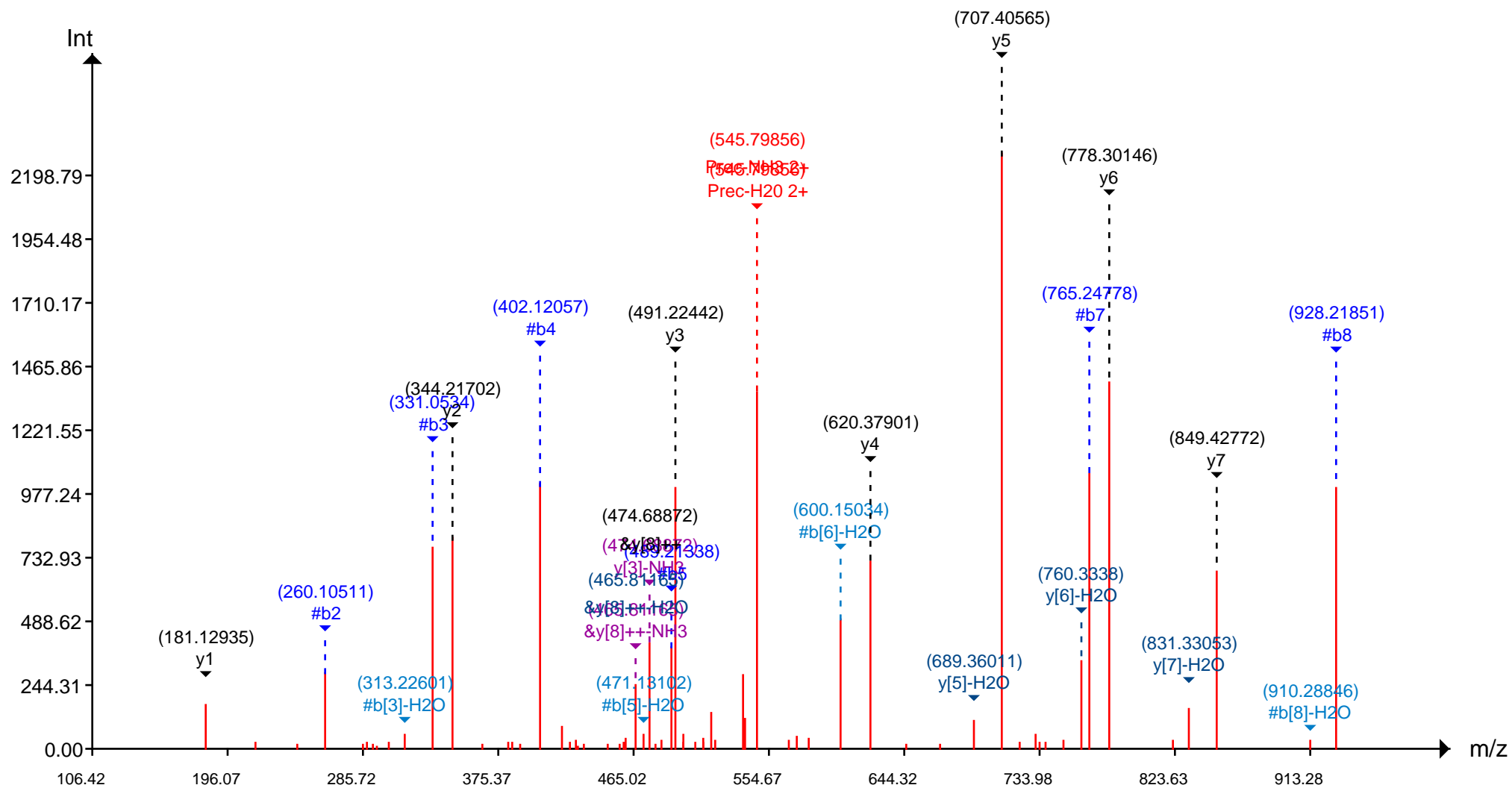
Confidence level 99.00%

Precursor m/z 577.32044

Identity Threshold 47.64

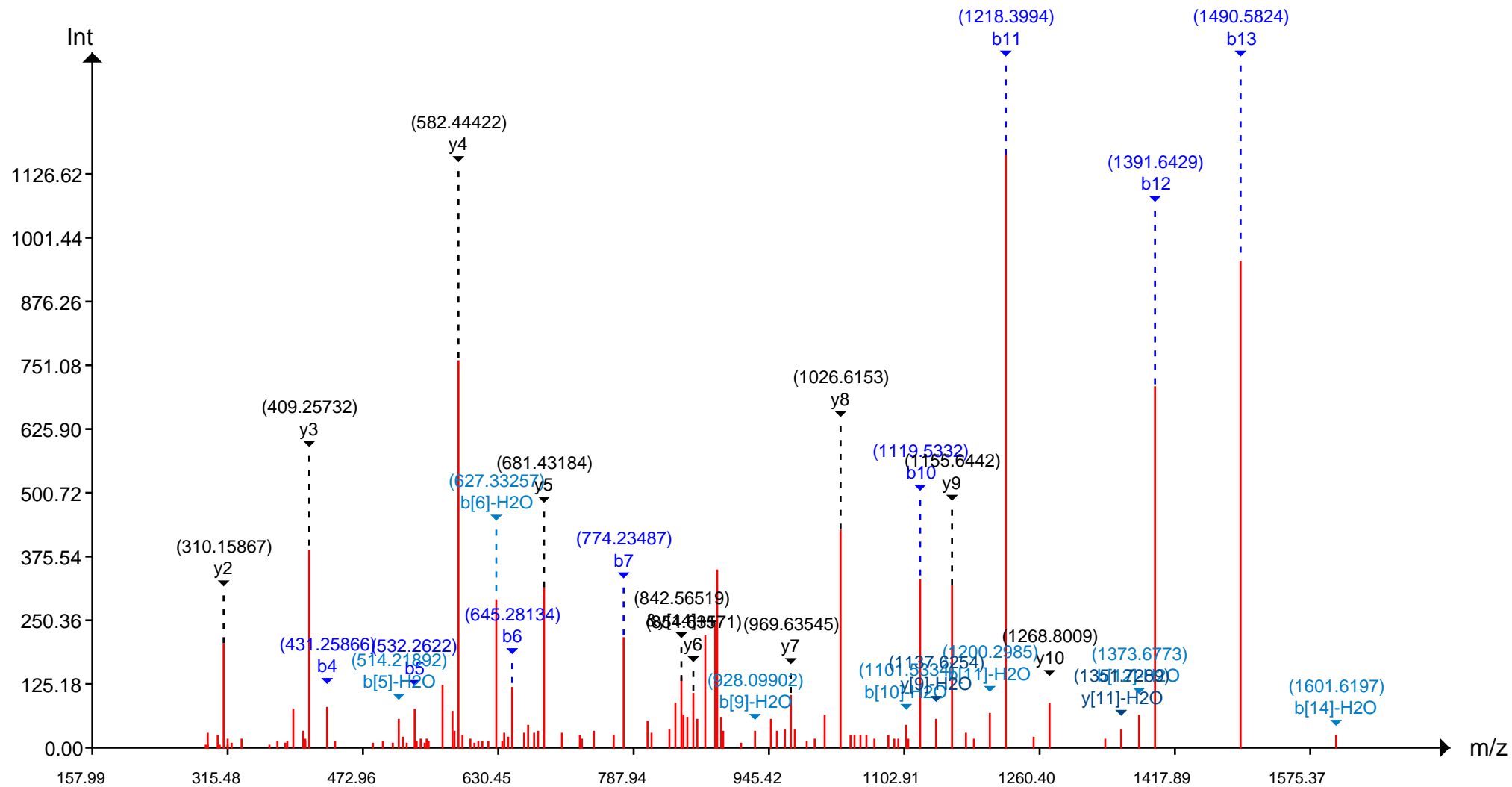
Charge 2+

Mass error -0.001022



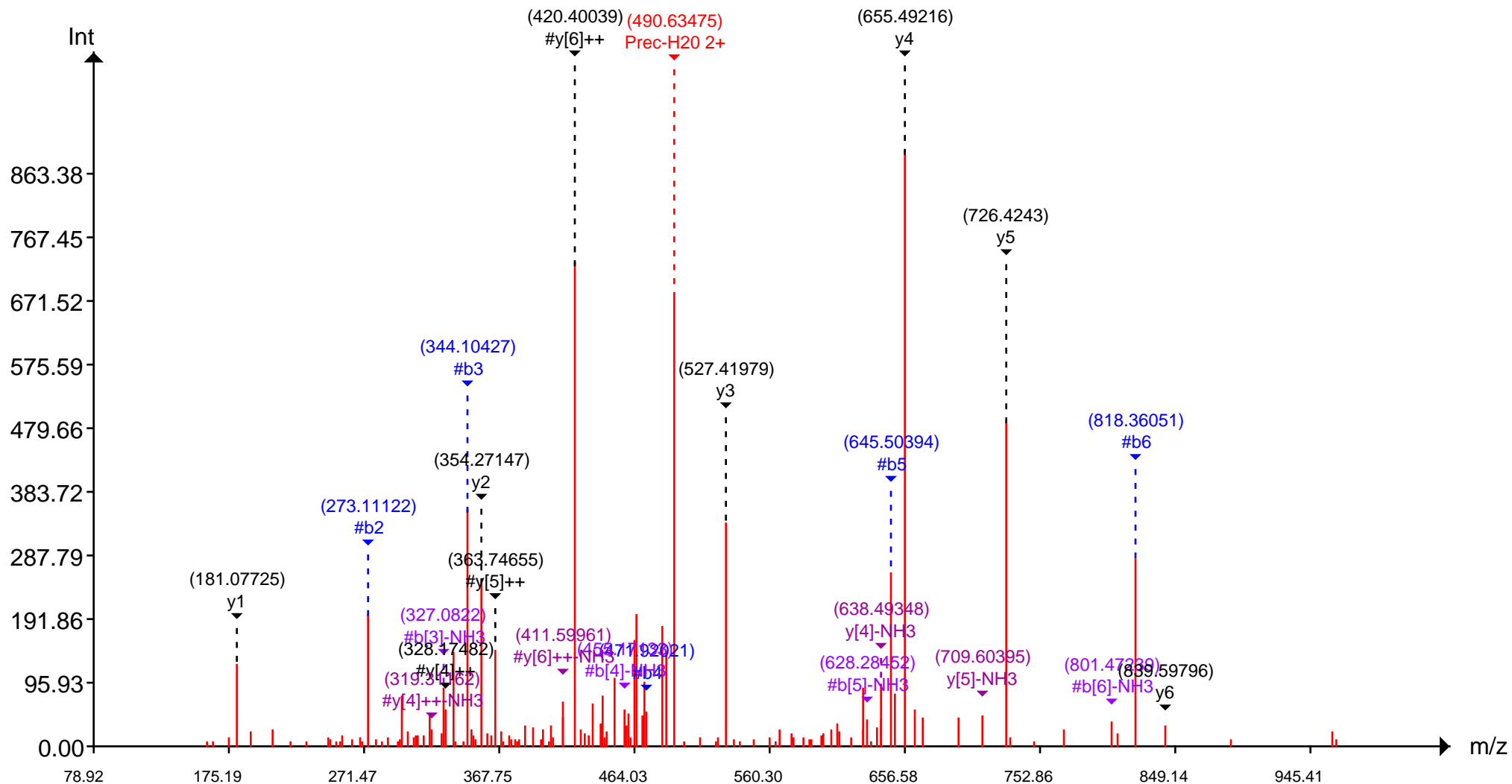
Info

Modified Sequence	AcD3-DVAASEFYR<C13>-COOH	Ion score	74.1
Protein	P17182, P21550, P17183	Confidence level	99.00%
Precursor m/z	554.77588	Identity Threshold	47.07
Charge	2+	Mass error	6.8E-5



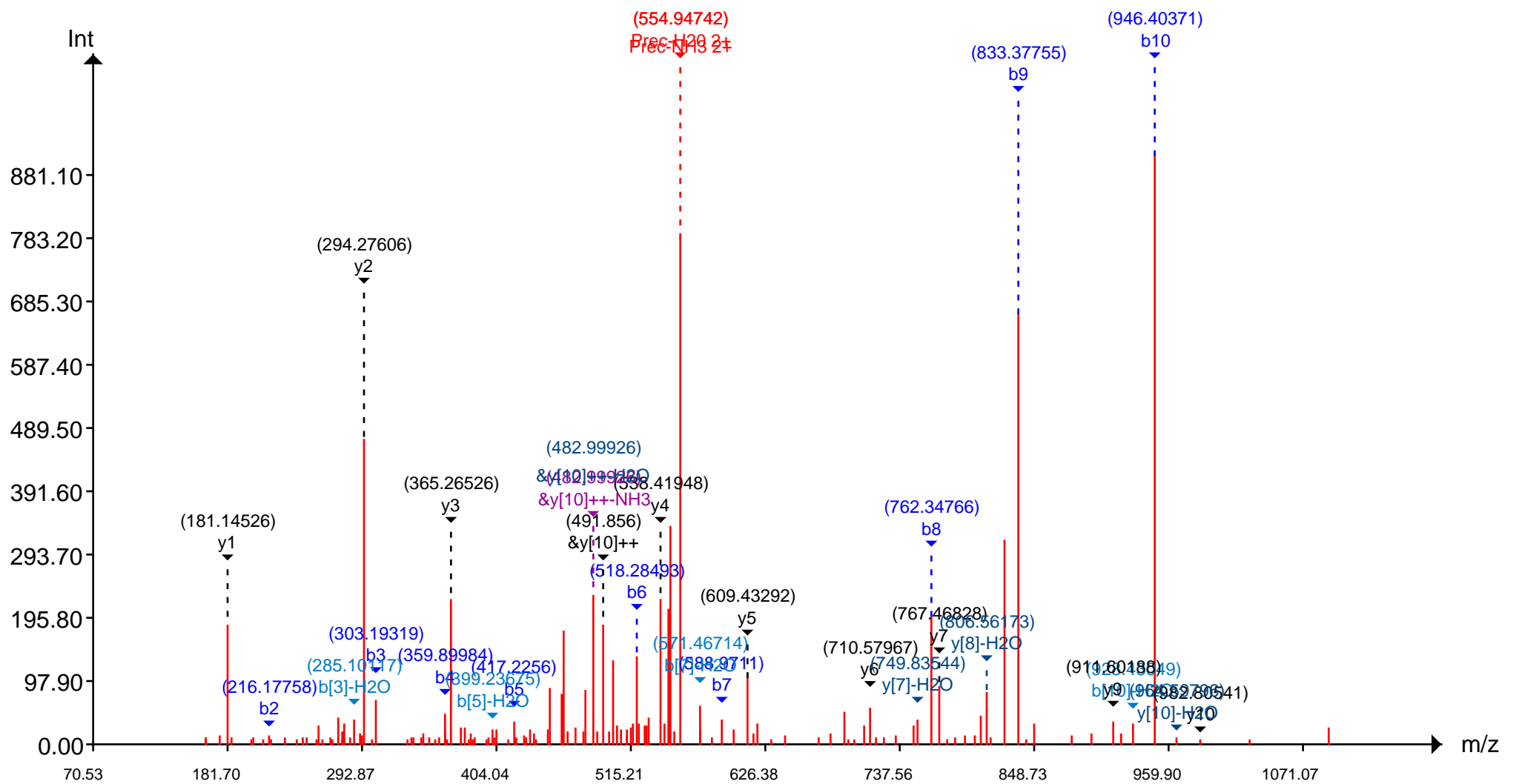
Info

Modified Sequence	AcD3-ASQVTLEGDK<AcD3K*>VK<AcD3K*>VER<C13>-COOH	Ion score	68.85
Protein	Q9DCW4	Confidence level	99.00%
Precursor m/z	900.51212	Identity Threshold	46.10
Charge	2+	Mass error	0.001464

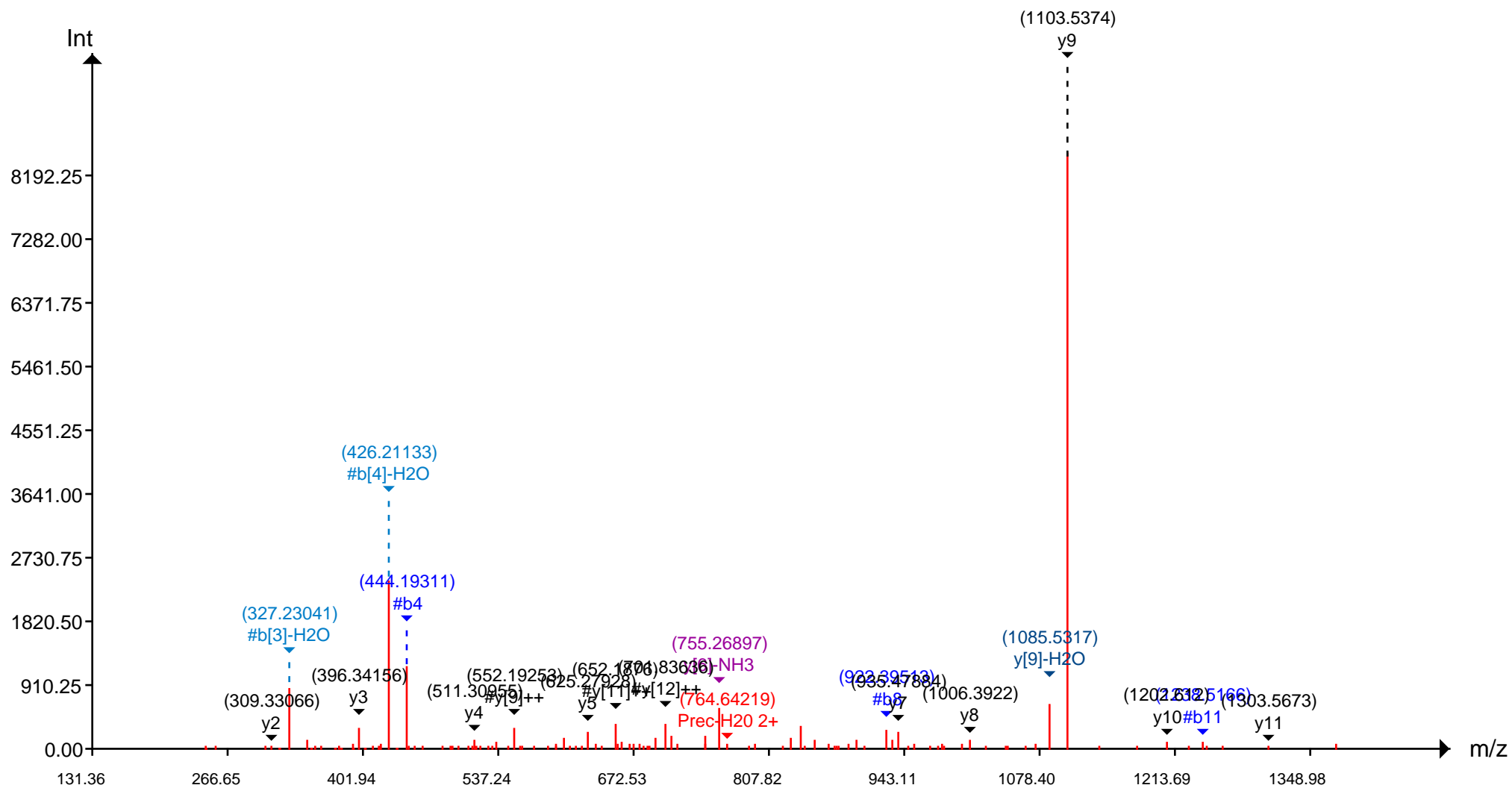


Info

Modified Sequence	AcD3-NLAQK<AcD3K*>K<AcD3K*>R<C13>-COOH	Ion score	43.2
Protein	P61759	Confidence level	99.00%
Precursor m/z	499.82338	Identity Threshold	40.67
Charge	2+	Mass error	-3.76E-4



Info			
Modified Sequence	AcD3-VASGGTAK<AcD3K*>AIR<C13>-COOH	Ion score	74.98
Protein	Q9CWJ9	Confidence level	99.00%
Precursor m/z	563.84333	Identity Threshold	44.51
Charge	2+	Mass error	9.6E-5



Info

Modified Sequence AcD3-VVTVPAYFNDSQR<C13>-COOH

Ion score 82.23

Protein P63017

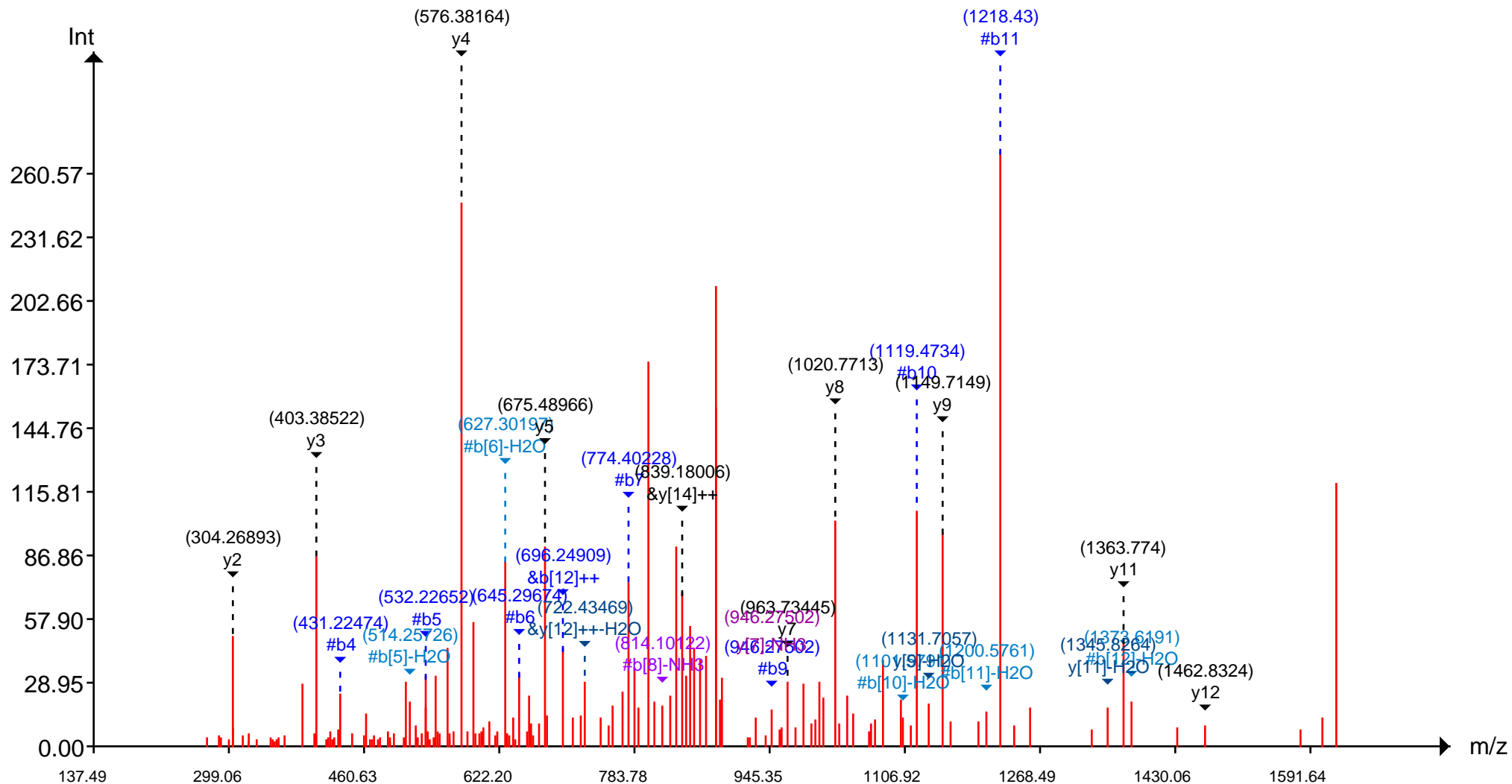
Confidence level 99.00%

Precursor m/z 773.90562

Identity Threshold 48.09

Charge 2+

Mass error 4.54E-4



Info

Modified Sequence AcD3-ASQVTLEGDK<AcD3K*>VK<AcD3K*>VER-COOH

Ion score 51.4

Protein Q9DCW4

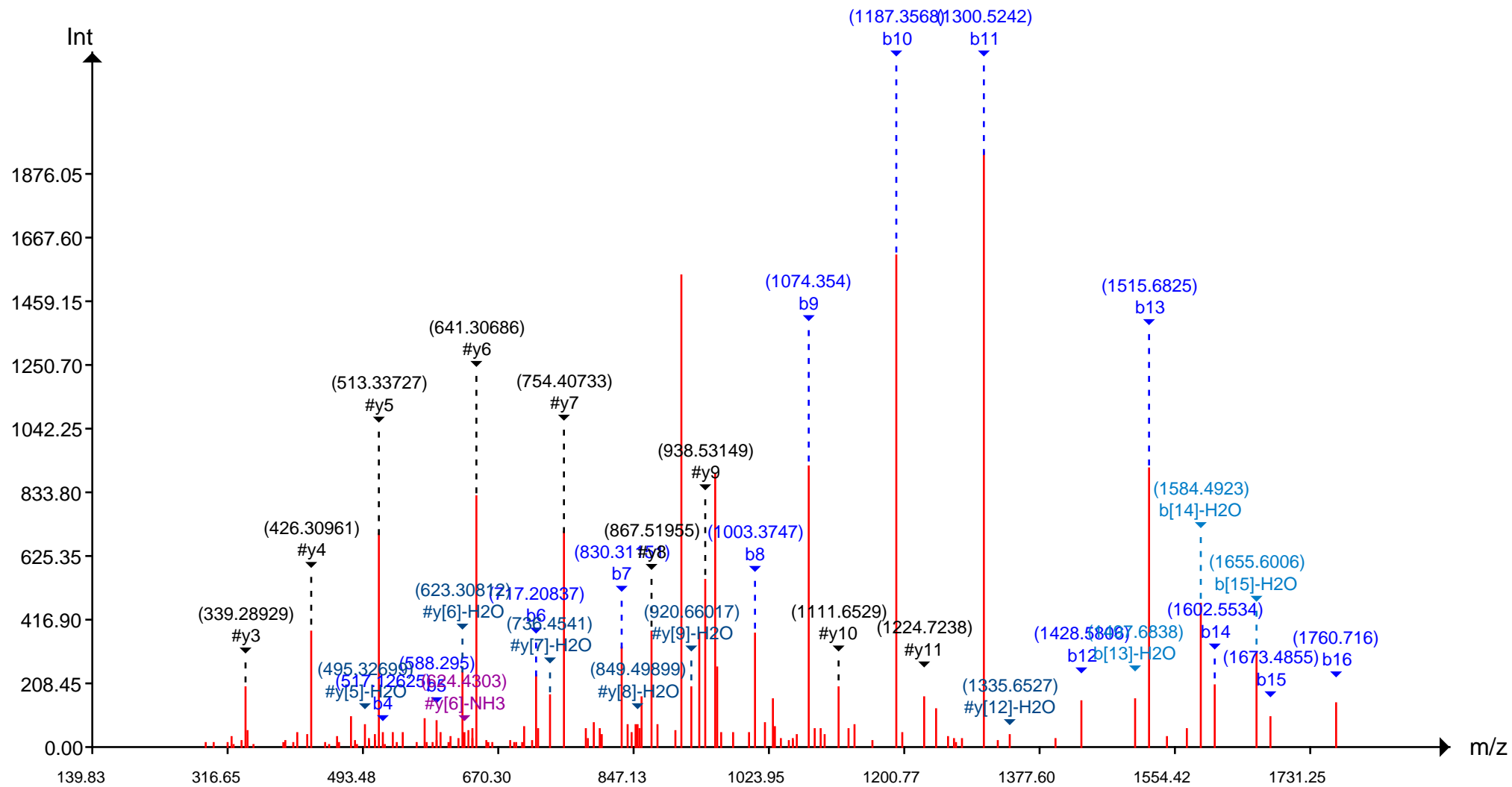
Confidence level 99.00%

Precursor m/z 897.50201

Identity Threshold 46.83

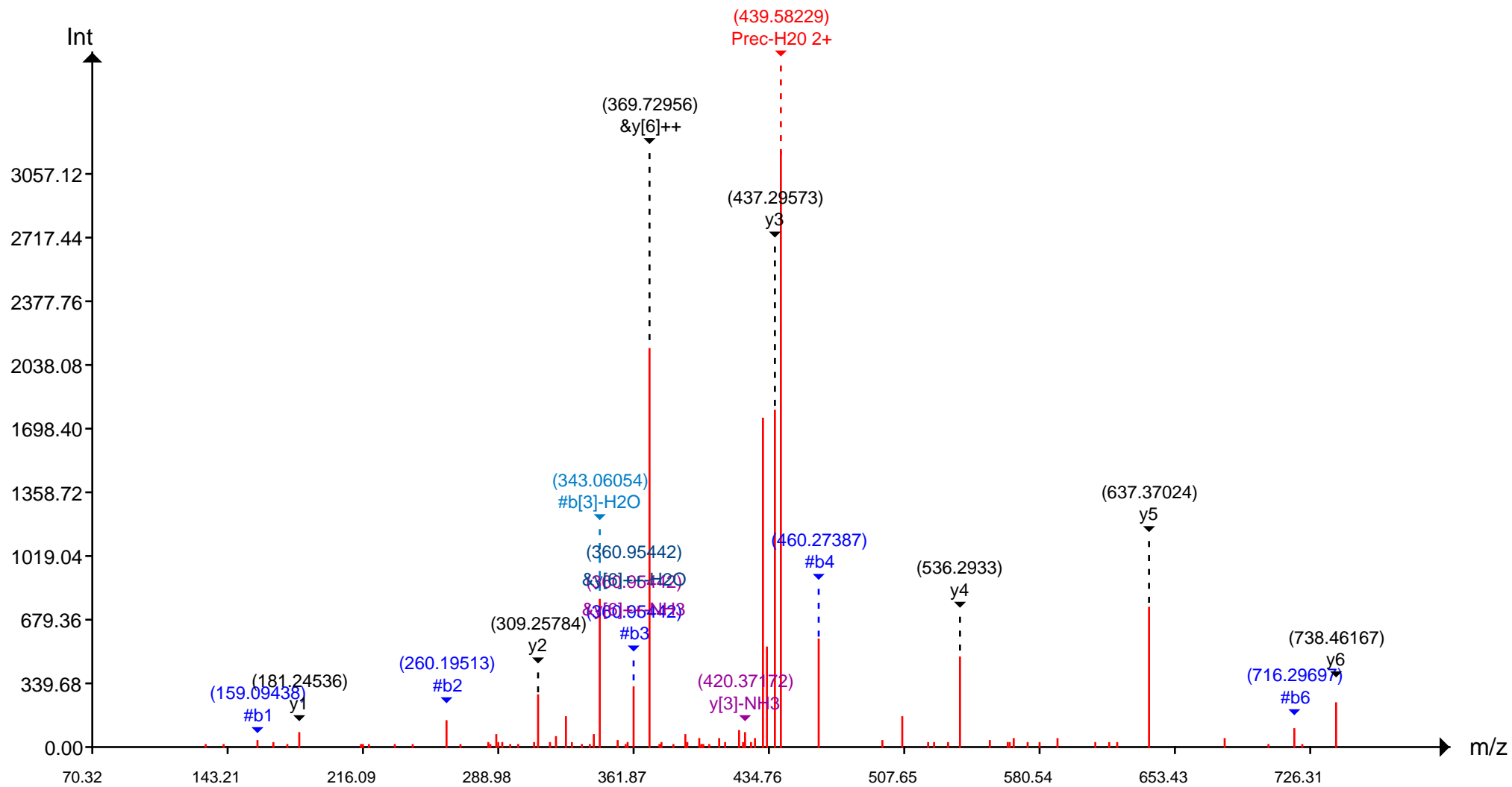
Charge 2+

Mass error 0.001385



Info

Modified Sequence	AcD3-EVQDAELK<AcD3K*>ALLQSSASR<C13>-COOH	Ion score	96.06
Protein	P50580	Confidence level	99.00%
Precursor m/z	971.02987	Identity Threshold	47.44
Charge	2+	Mass error	0.002341



Info

Modified Sequence AcD3-ITTVQQR<C13>-COOH

Ion score 51.27

Protein P14152

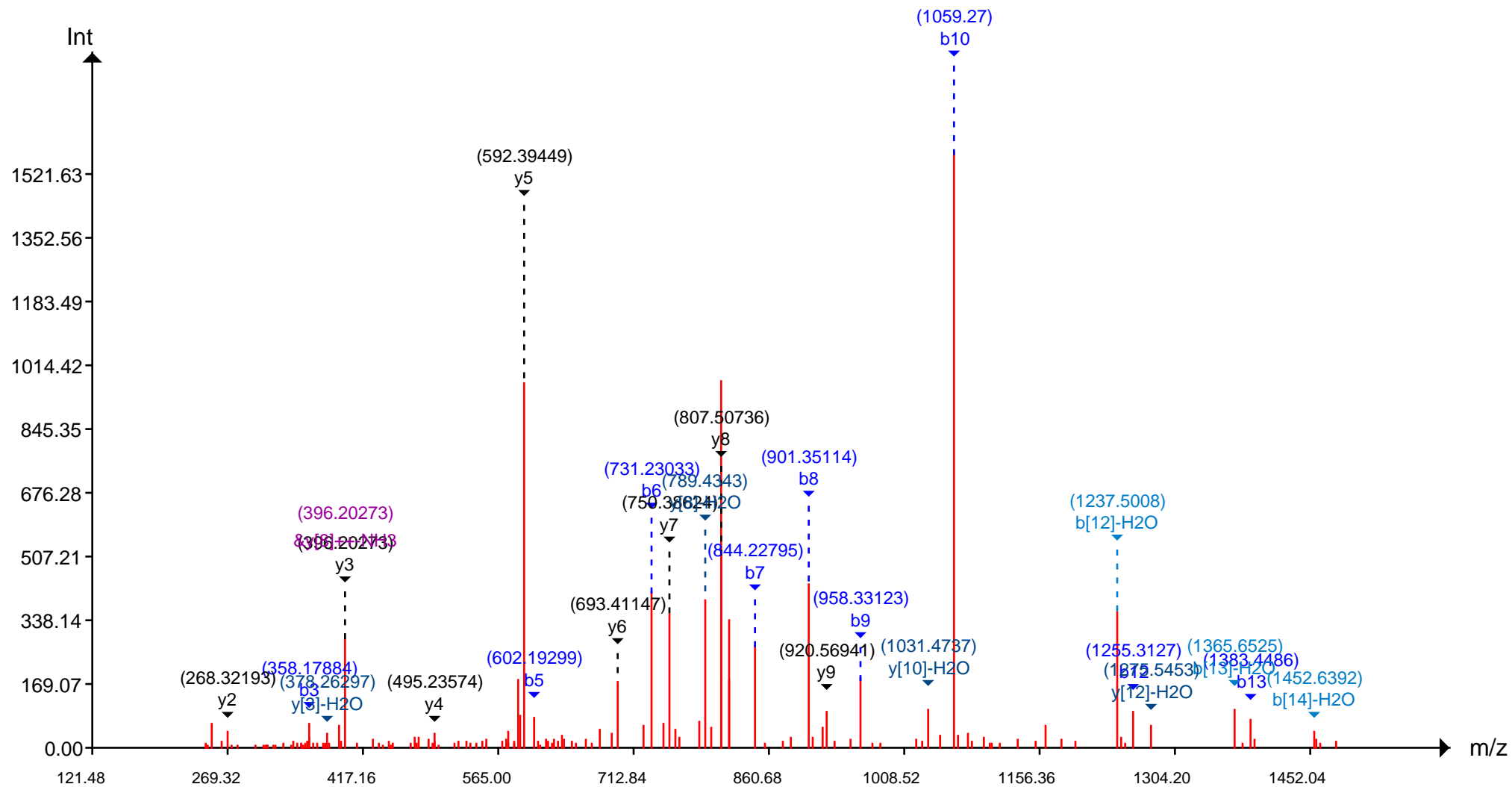
Confidence level 99.00%

Precursor m/z 448.77021

Identity Threshold 45.71

Charge 2+

Mass error -3.31E-4



Info

Modified Sequence AcD3-AQLDEELGGTPVQSR<C13>-COOH

Ion score 58.32

Protein P13020

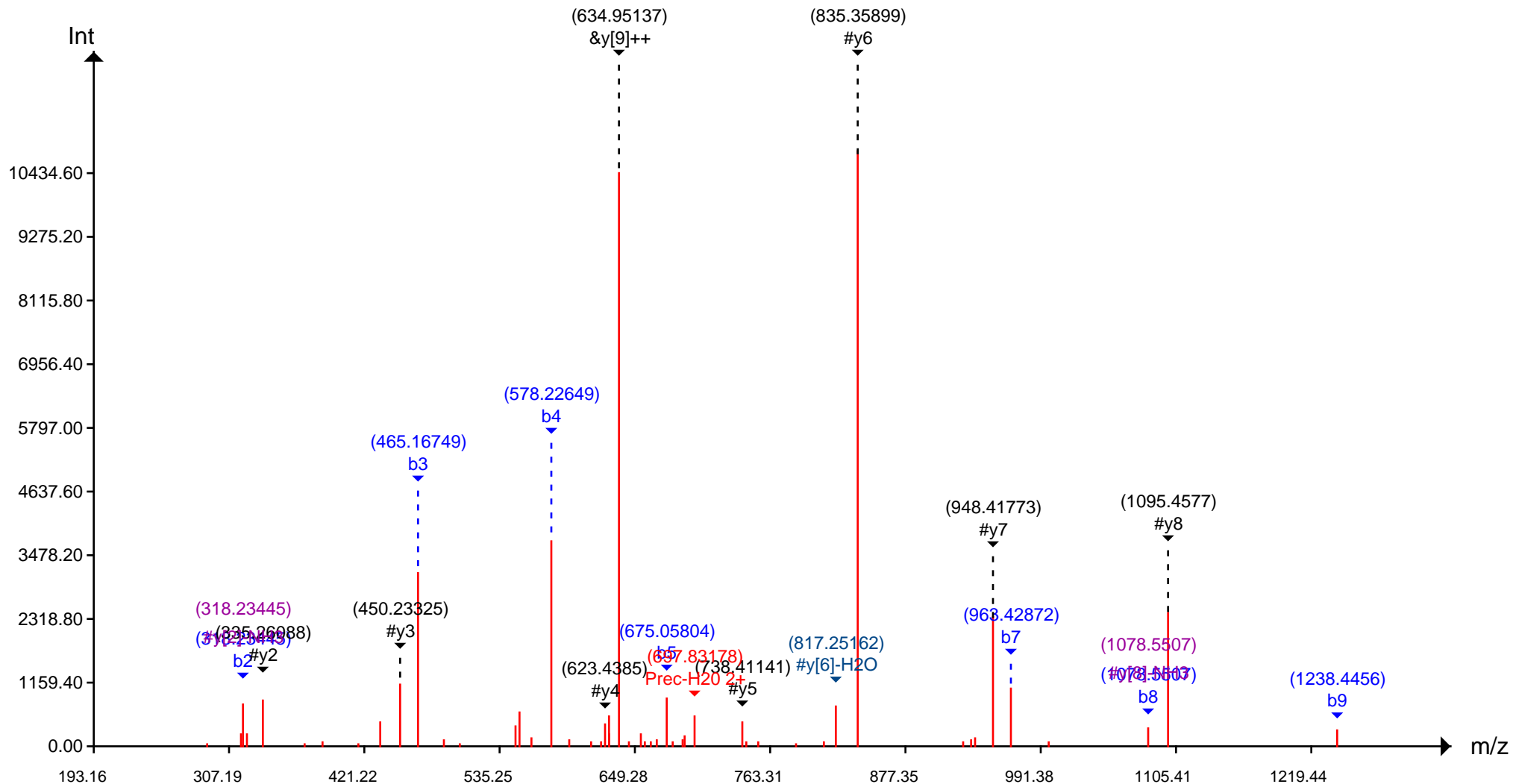
Confidence level 99.00%

Precursor m/z 825.92768

Identity Threshold 48.13

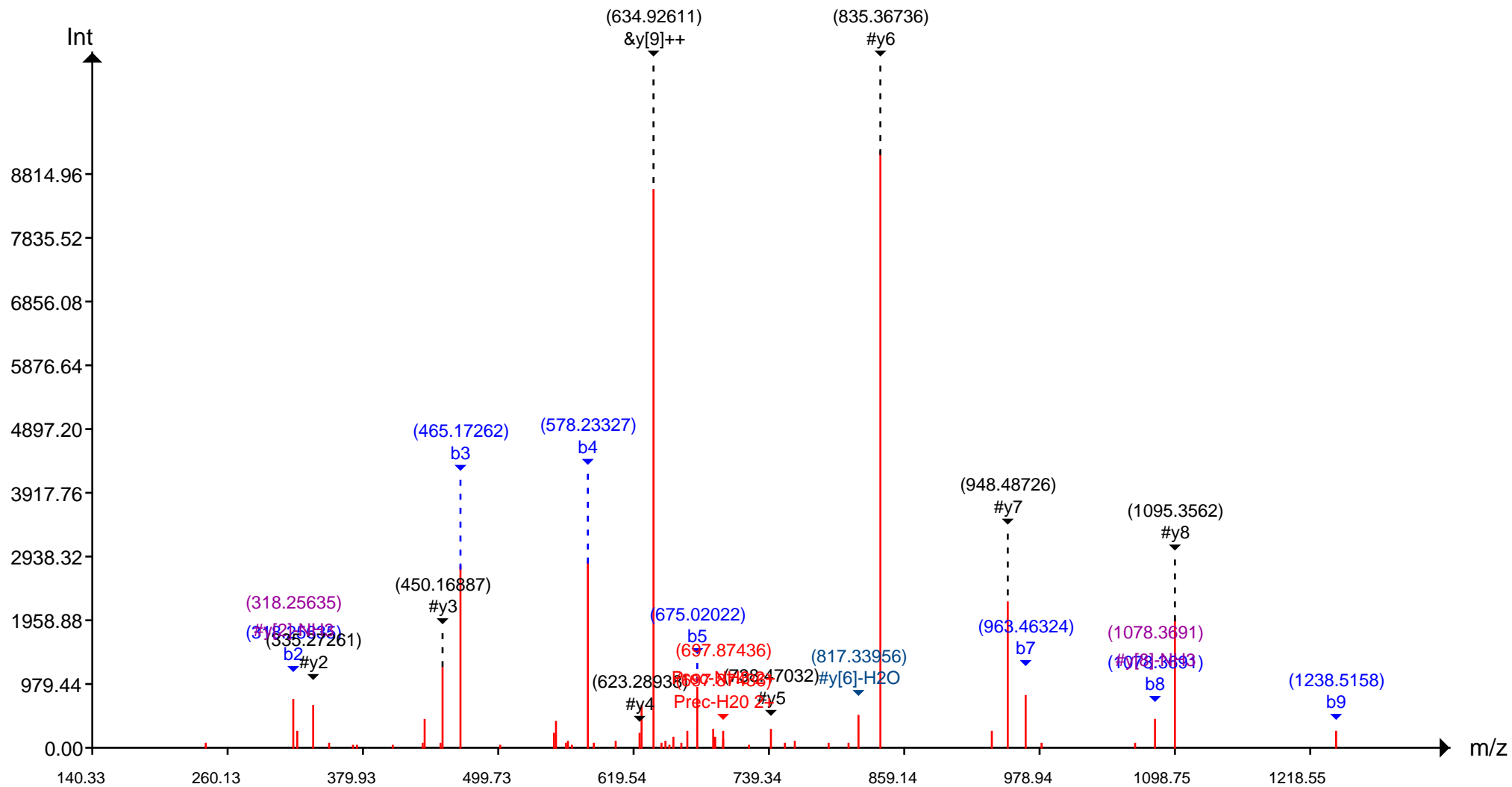
Charge 2+

Mass error 0.001269



Info

Modified Sequence	AcD3-VK<AcD3K*>M<Mox*>LPDK<AcD3K*>DC<Cmm*>R-COOH	Ion score	58.81
Protein	P18760	Confidence level	99.00%
Precursor m/z	706.86526	Identity Threshold	48.05
Charge	2+	Mass error	0.001002



Info

Modified Sequence AcD3-VK<AcD3K*>M<Mox*>LPDK<AcD3K*>DC<Cmm*>R-COOH

Ion score 61.83

Protein P18760

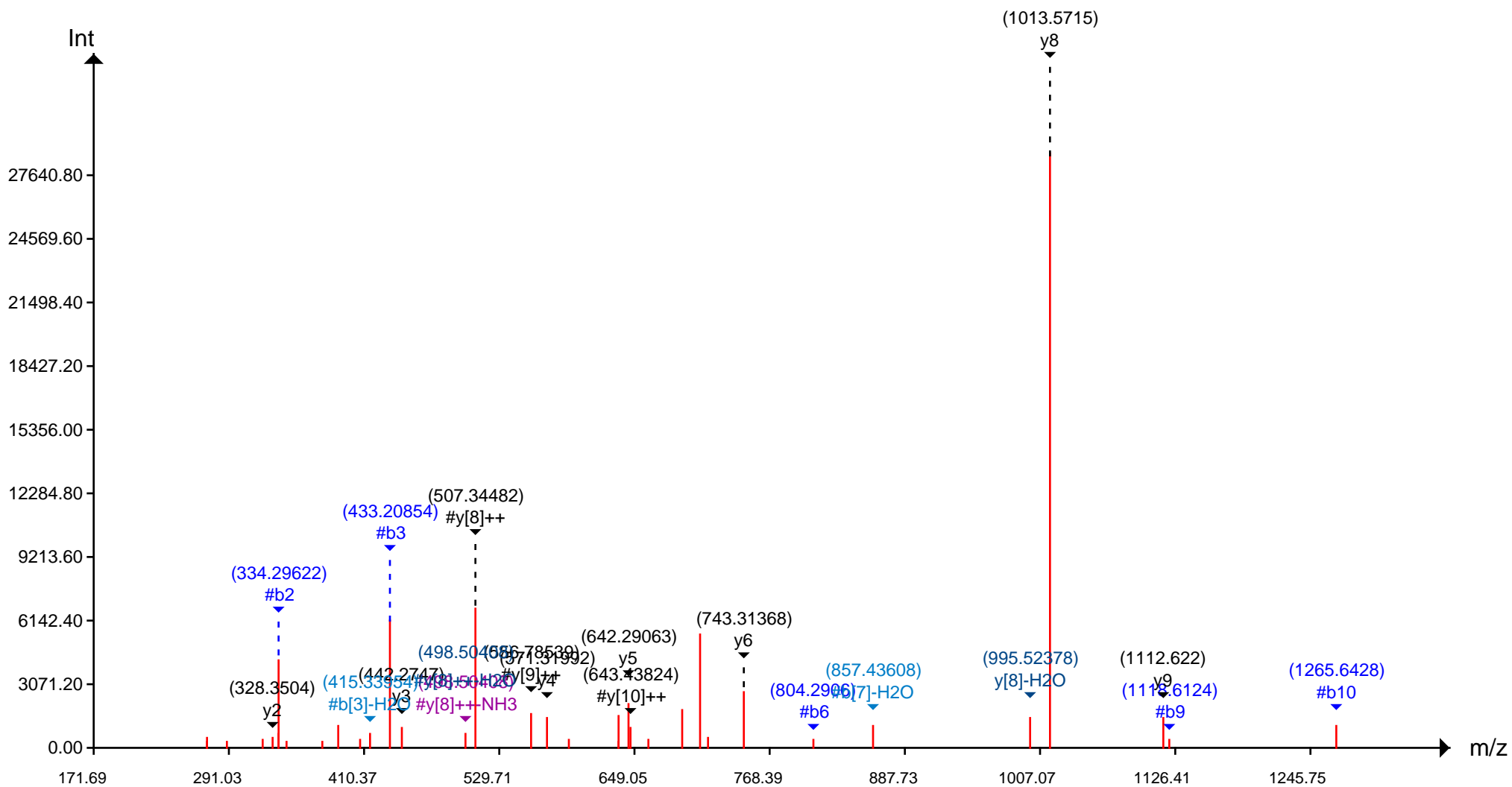
Confidence level 99.00%

Precursor m/z 706.86556

Identity Threshold 47.97

Charge 2+

Mass error 0.001602



Info

Modified Sequence AcD3-DK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH

Ion score 58.15

Protein P17742

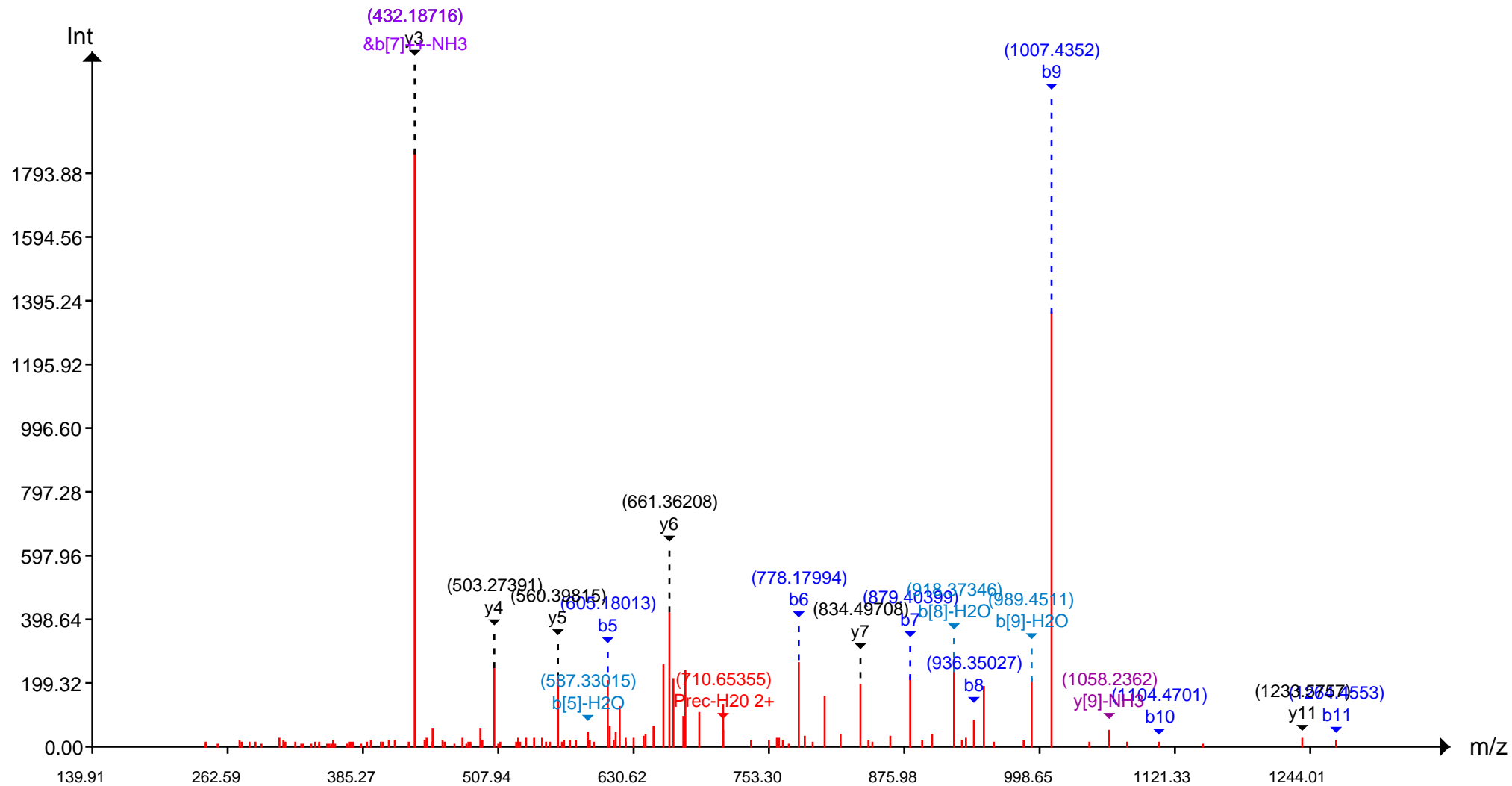
Confidence level 99.00%

Precursor m/z 723.40464

Identity Threshold 47.28

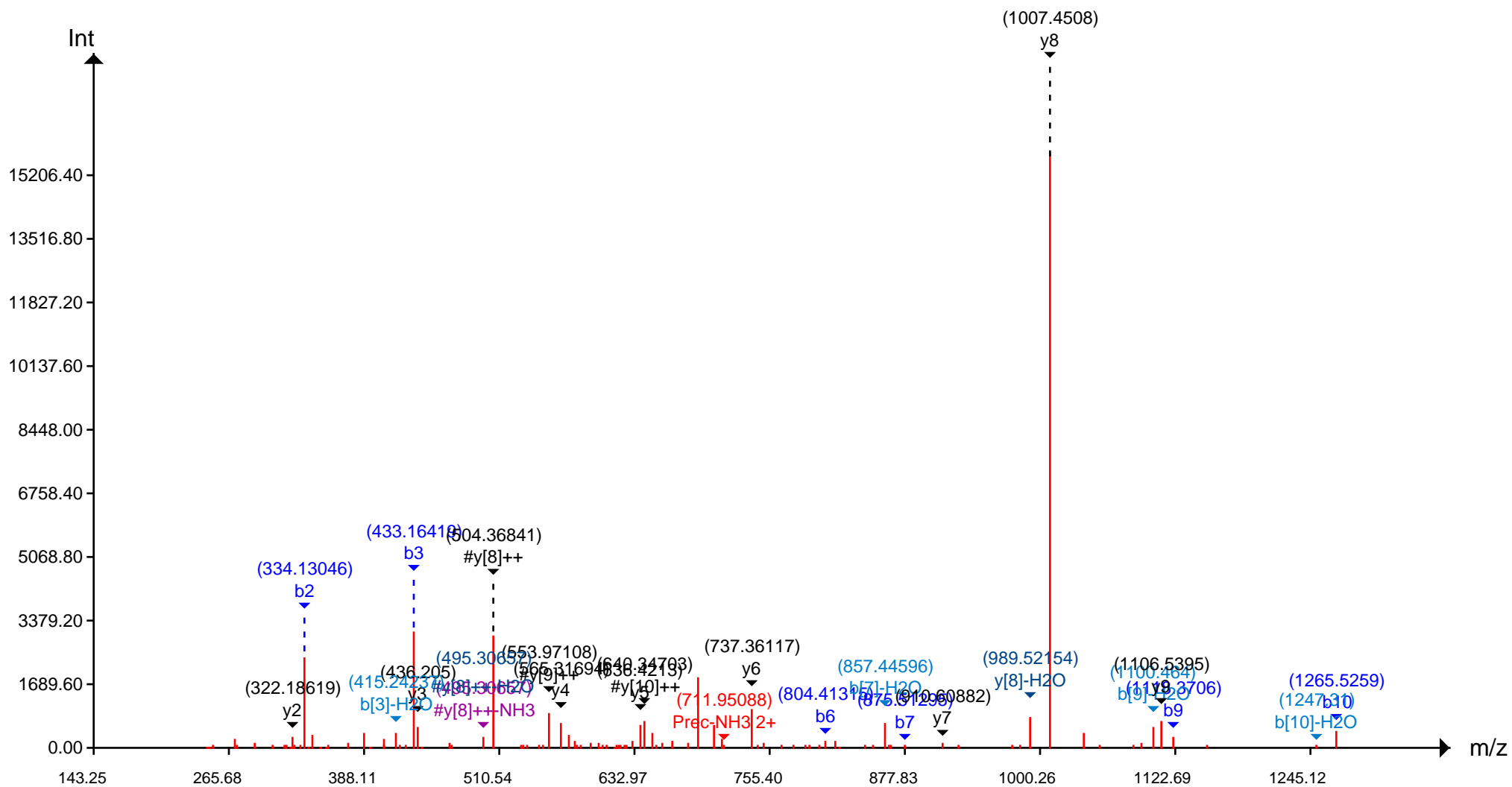
Charge 2+

Mass error -0.00204



Info

Modified Sequence	AcD3-C<Cmm*>TGQIK<AcD3K*>TGAPC<Cmm*>R-COOH	Ion score	54.44
Protein	P17182, P21550, P17183	Confidence level	99.00%
Precursor m/z	719.85721	Identity Threshold	47.44
Charge	2+	Mass error	0.002328



Info

Modified Sequence AcD3-DK<AcD3K*>VPK<AcD3K*>TAENFR-COOH

Ion score 60.08

Protein P17742

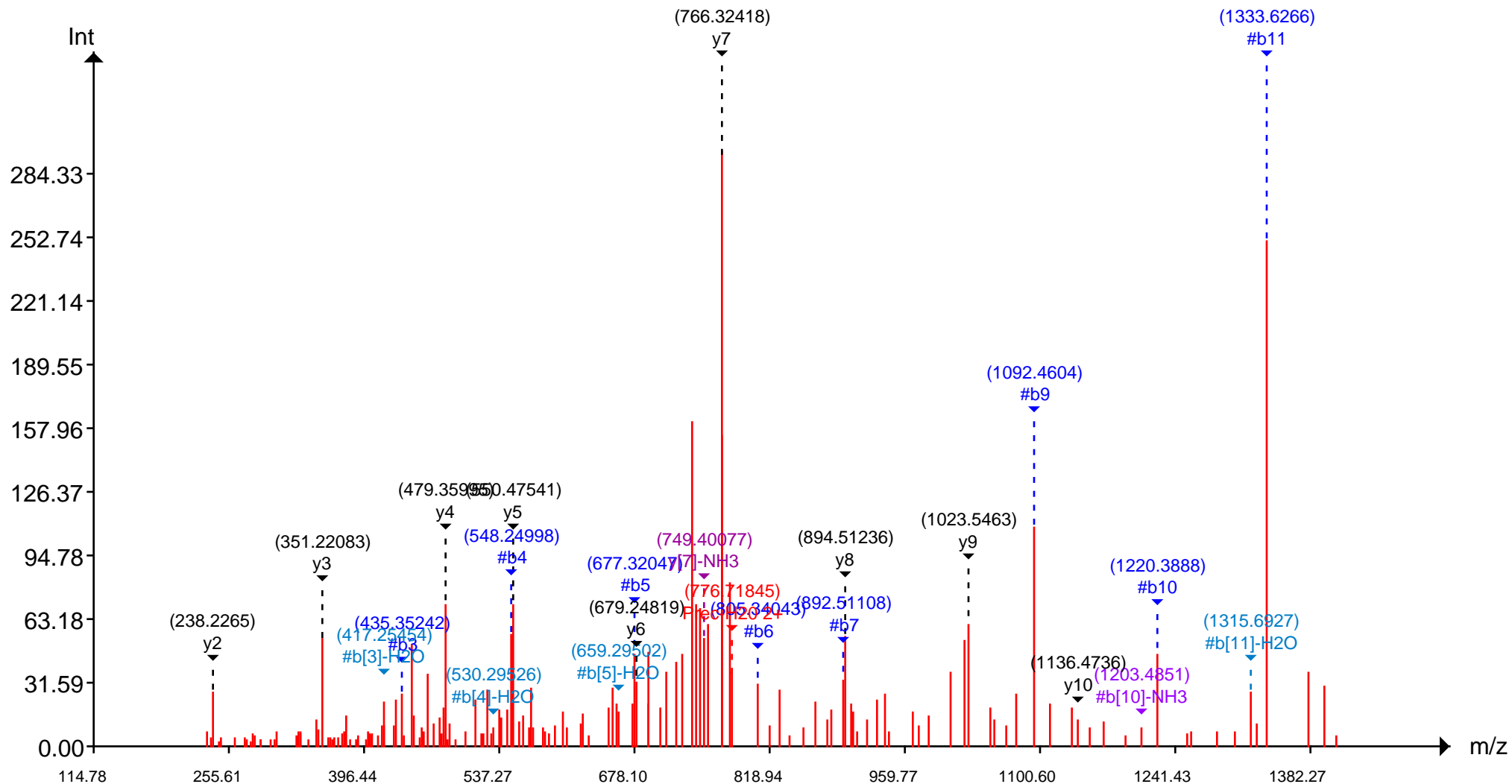
Confidence level 99.00%

Precursor m/z 720.39488

Identity Threshold 47.71

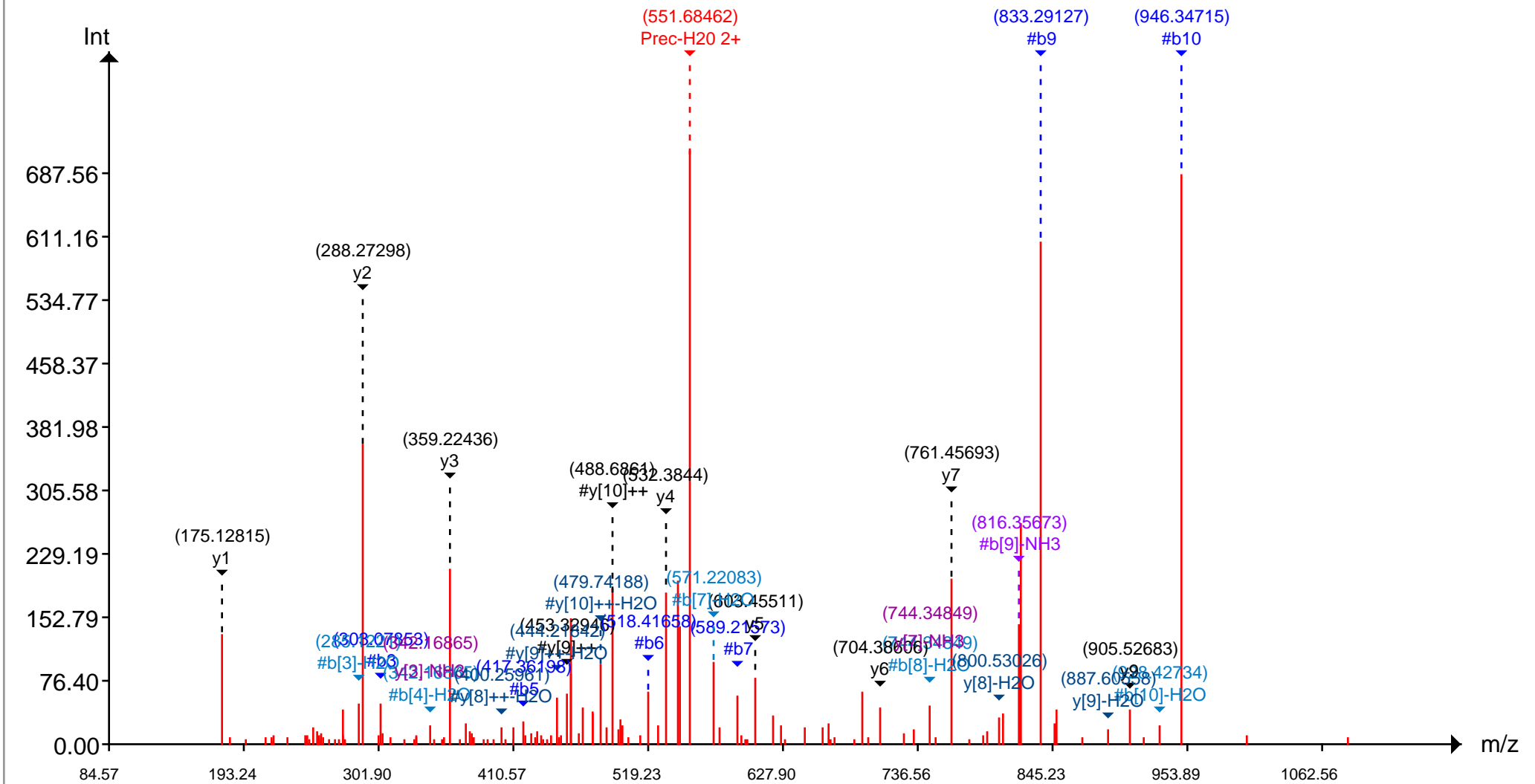
Charge 2+

Mass error -0.001419



Info

Modified Sequence	AcD3-SEK<AcD3K*>LEQSEAQLGR<C13>-COOH	Ion score	61.89
Protein	O35864	Confidence level	99.00%
Precursor m/z	785.91879	Identity Threshold	48.09
Charge	2+	Mass error	0.0018



Info

Modified Sequence AcD3-VASGGTAK<AcD3K*>AIR-COOH

Ion score 56.09

Protein Q9CWJ9

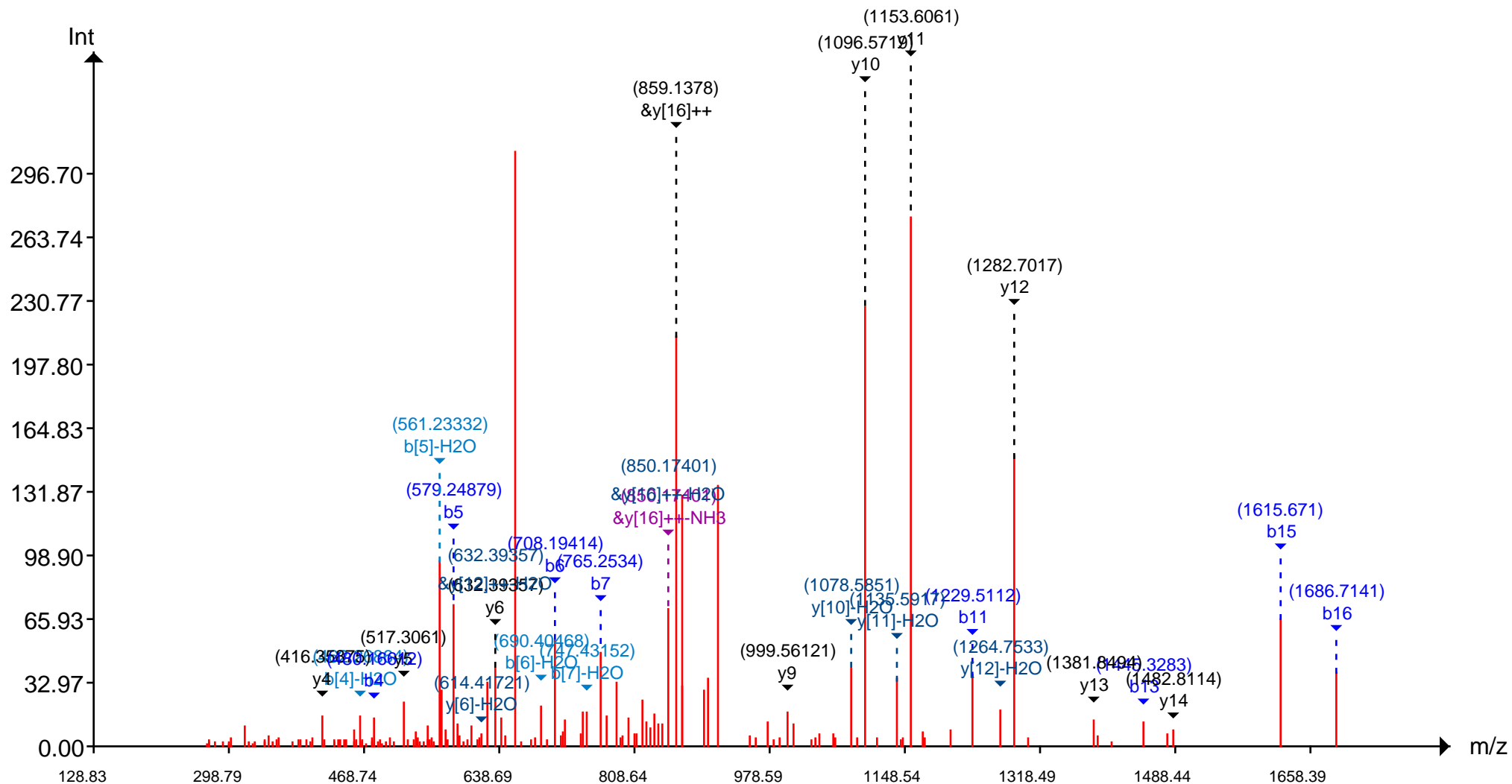
Confidence level 99.00%

Precursor m/z 560.8335

Identity Threshold 45.90

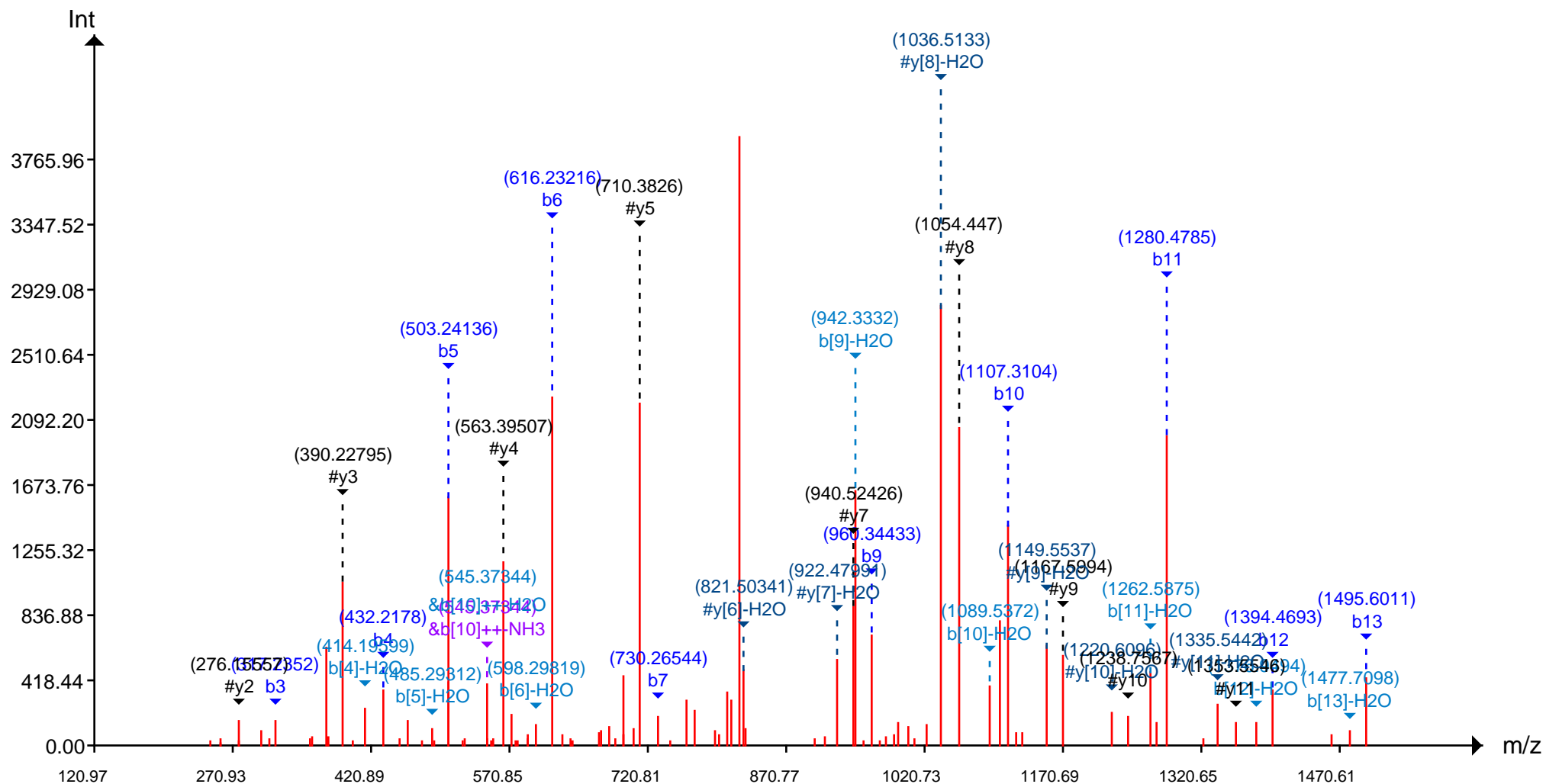
Charge 2+

Mass error 5.77E-4



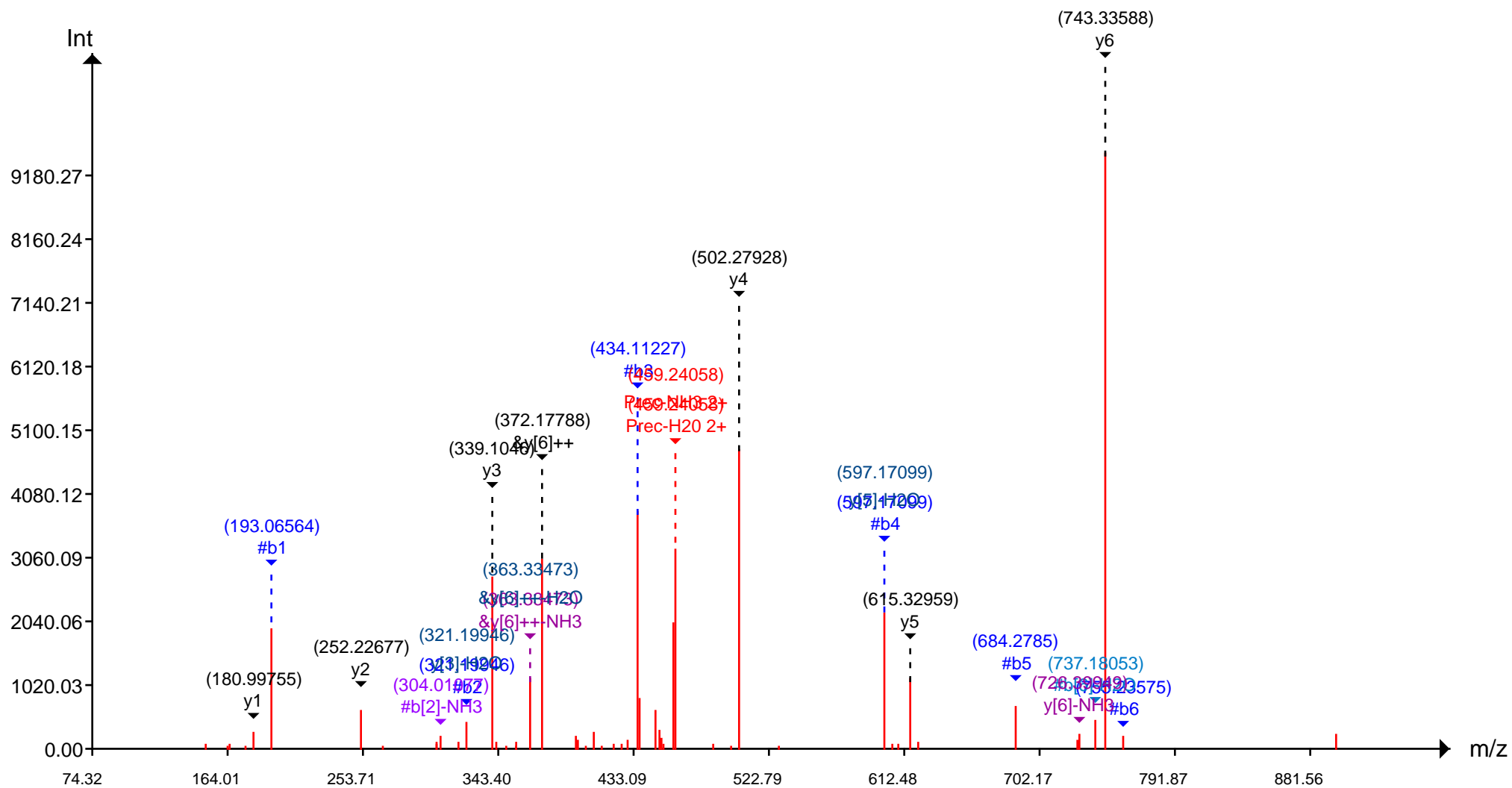
Info

Modified Sequence	AcD3-VSM<Mox*>TVEGPPPK<AcD3K*>DTGIAR-COOH	Ion score	54.45
Protein	P27048, P63163	Confidence level	99.00%
Precursor m/z	930.98666	Identity Threshold	48.09
Charge	2+	Mass error	0.001783



Info

Modified Sequence	AcD3-SLADAINTEFK<AcD3K*>NTR-COOH	Ion score	87.45
Protein	P20152	Confidence level	99.00%
Precursor m/z	835.4381	Identity Threshold	48.17
Charge	2+	Mass error	0.0027



Info

Modified Sequence AcD3-M<Mox*>QLYSAR<C13>-COOH

Ion score 55.08

Protein P62843

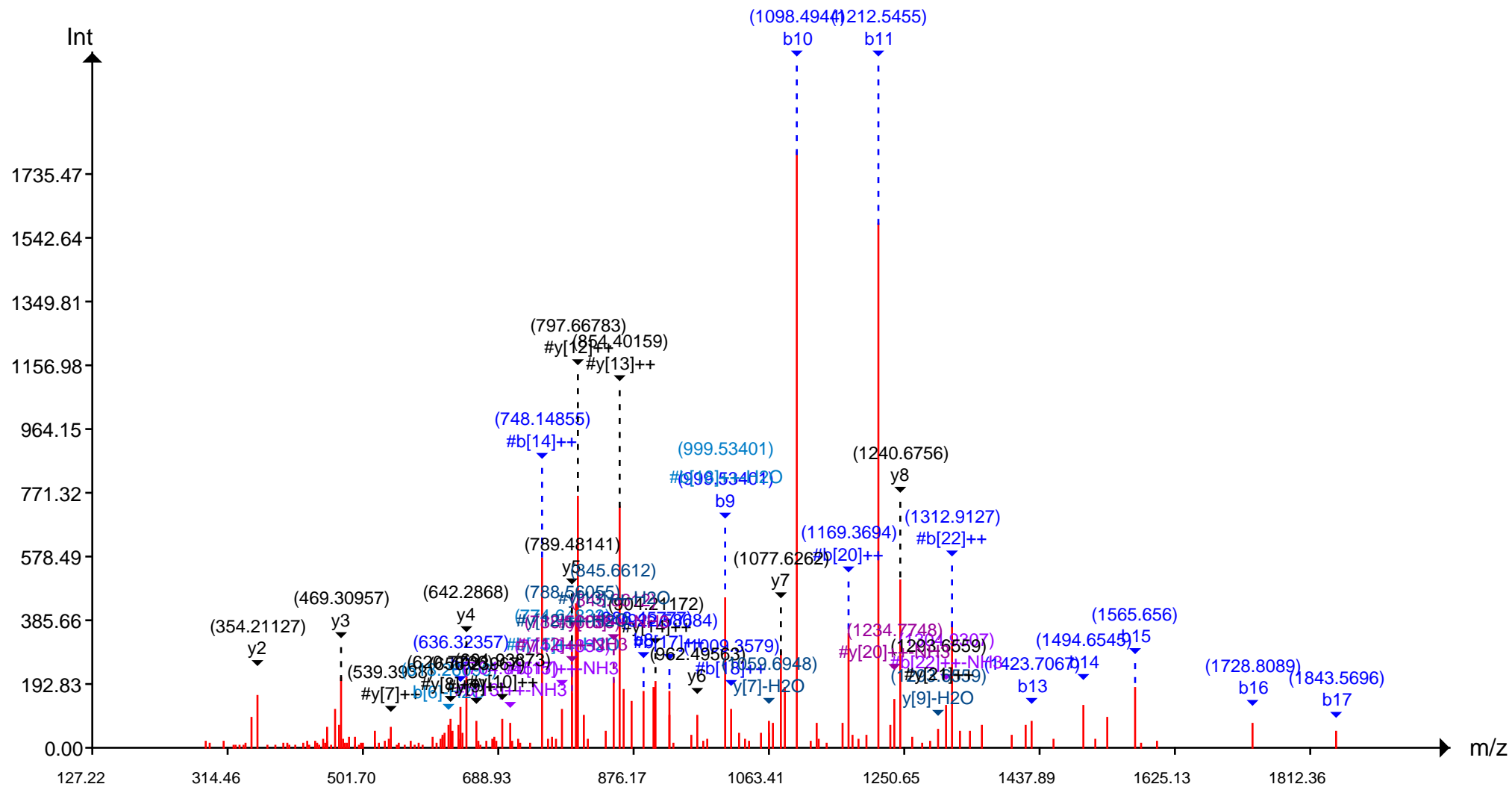
Confidence level 99.00%

Precursor m/z 468.24238

Identity Threshold 47.37

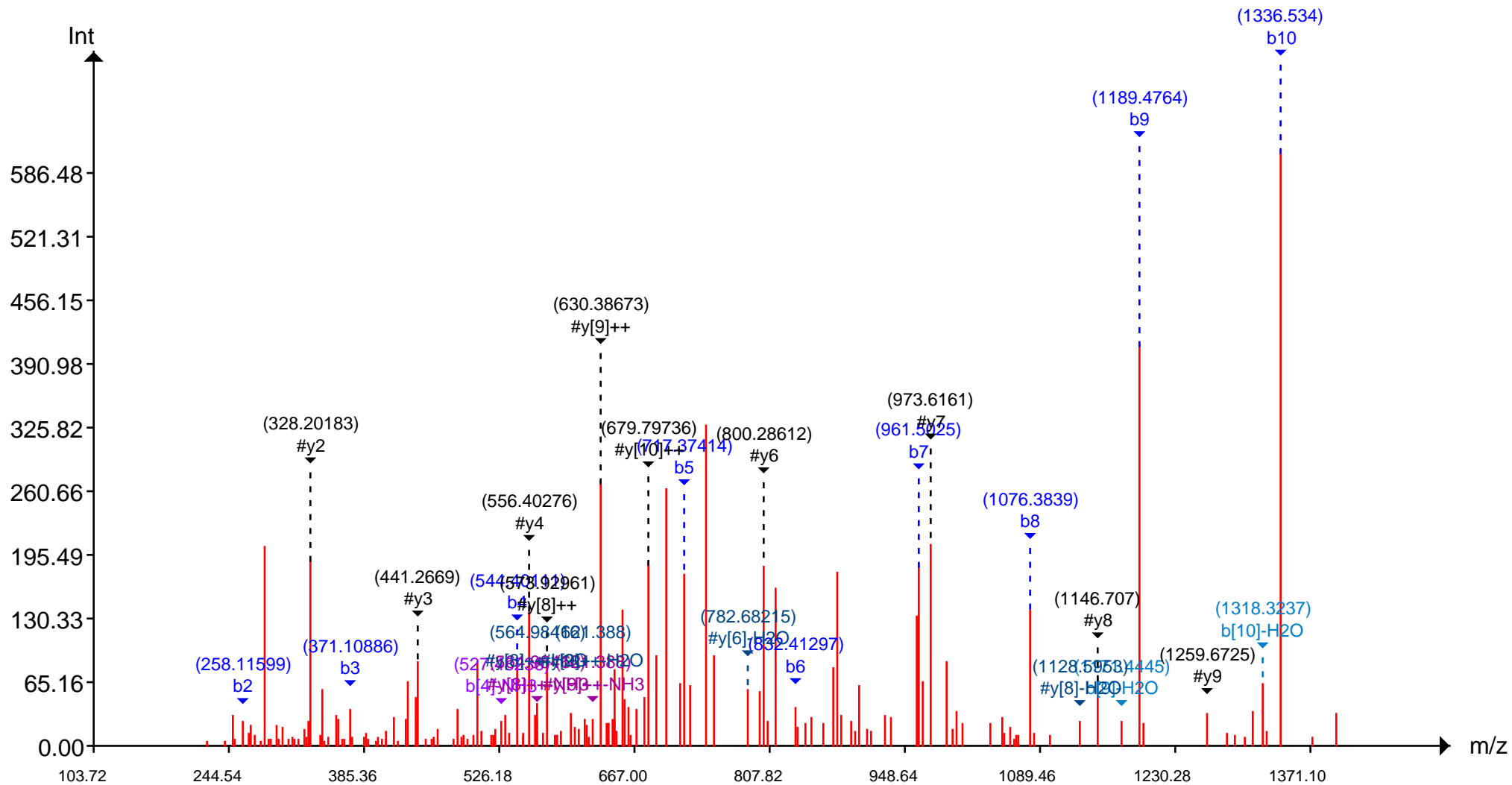
Charge 2+

Mass error -0.001487



Info

Modified Sequence	AcD3-SSTSVEHIIIVNPAAAYDK<AcD3K*>FK<AcD3K*>DK<AcD3K*>R<C13>-COOH	Ion score	63.83
Protein	Q99KJ8	Confidence level	99.00%
Precursor m/z	935.83627	Identity Threshold	47.17
Charge	3+	Mass error	0.004011



Info

Modified Sequence AcD3-LVLK<AcD3K*>K<AcD3K*>DEDLFR<C13>-COOH

Ion score 48.13

Protein O54962

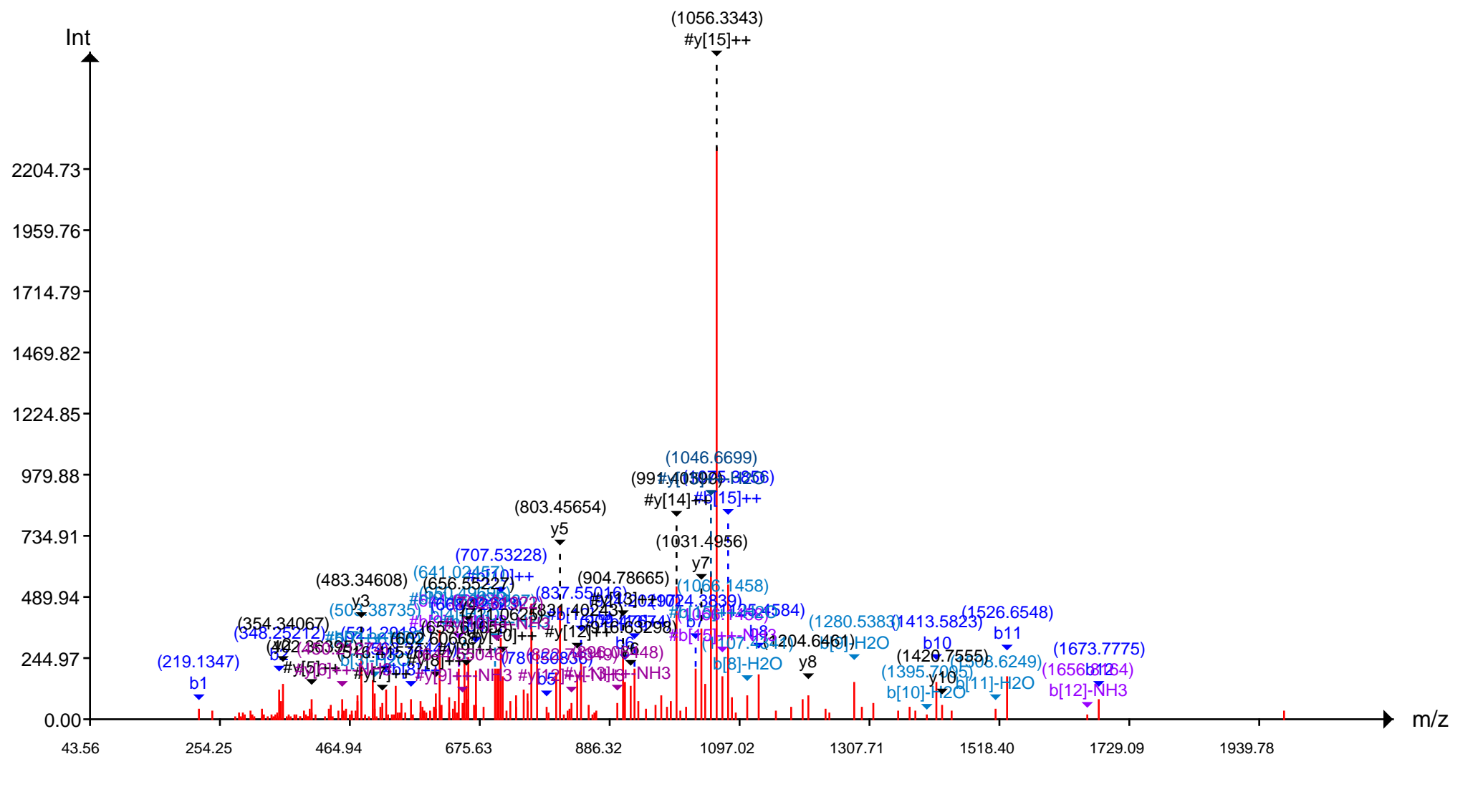
Confidence level 99.00%

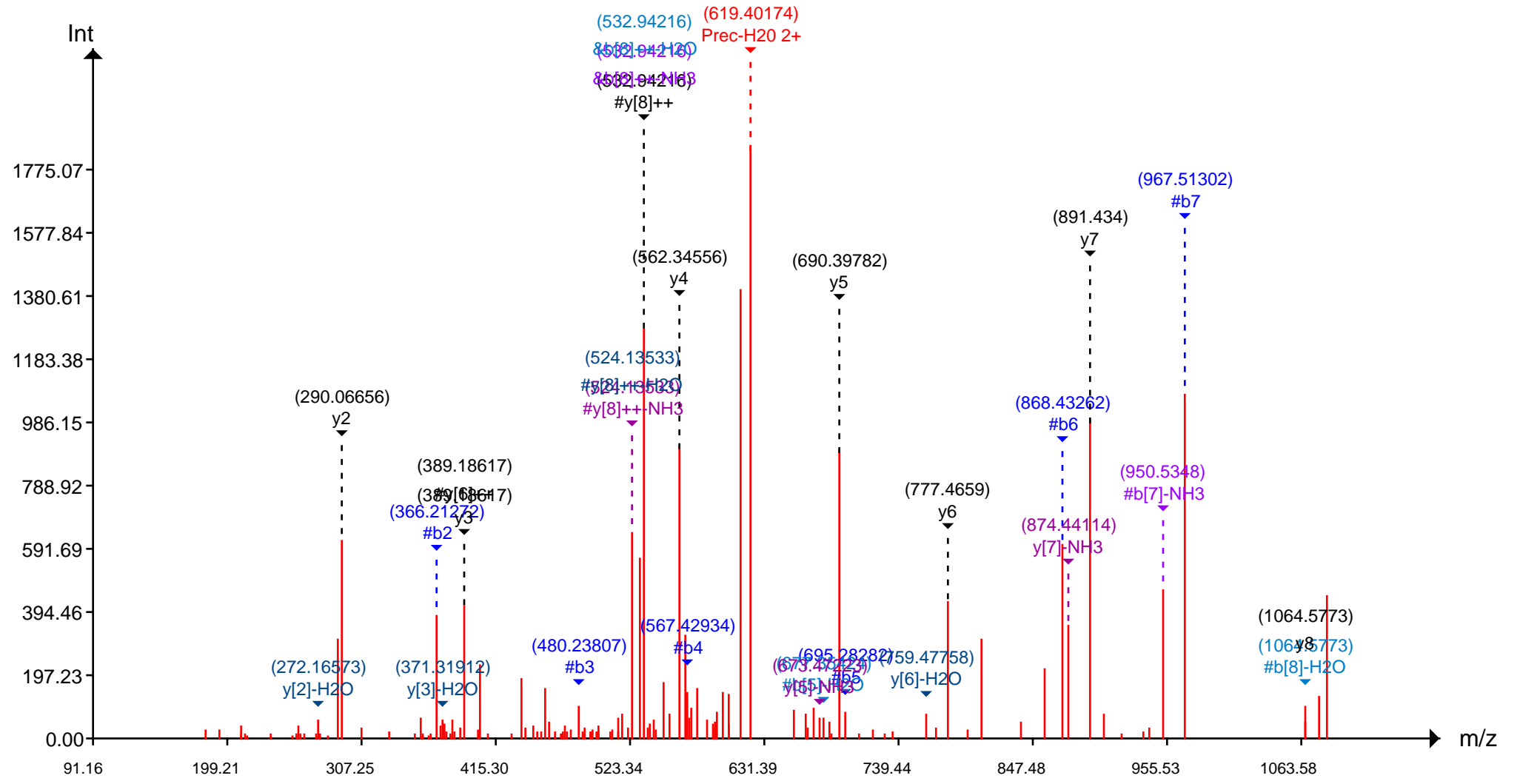
Precursor m/z 758.95592

Identity Threshold 43.50

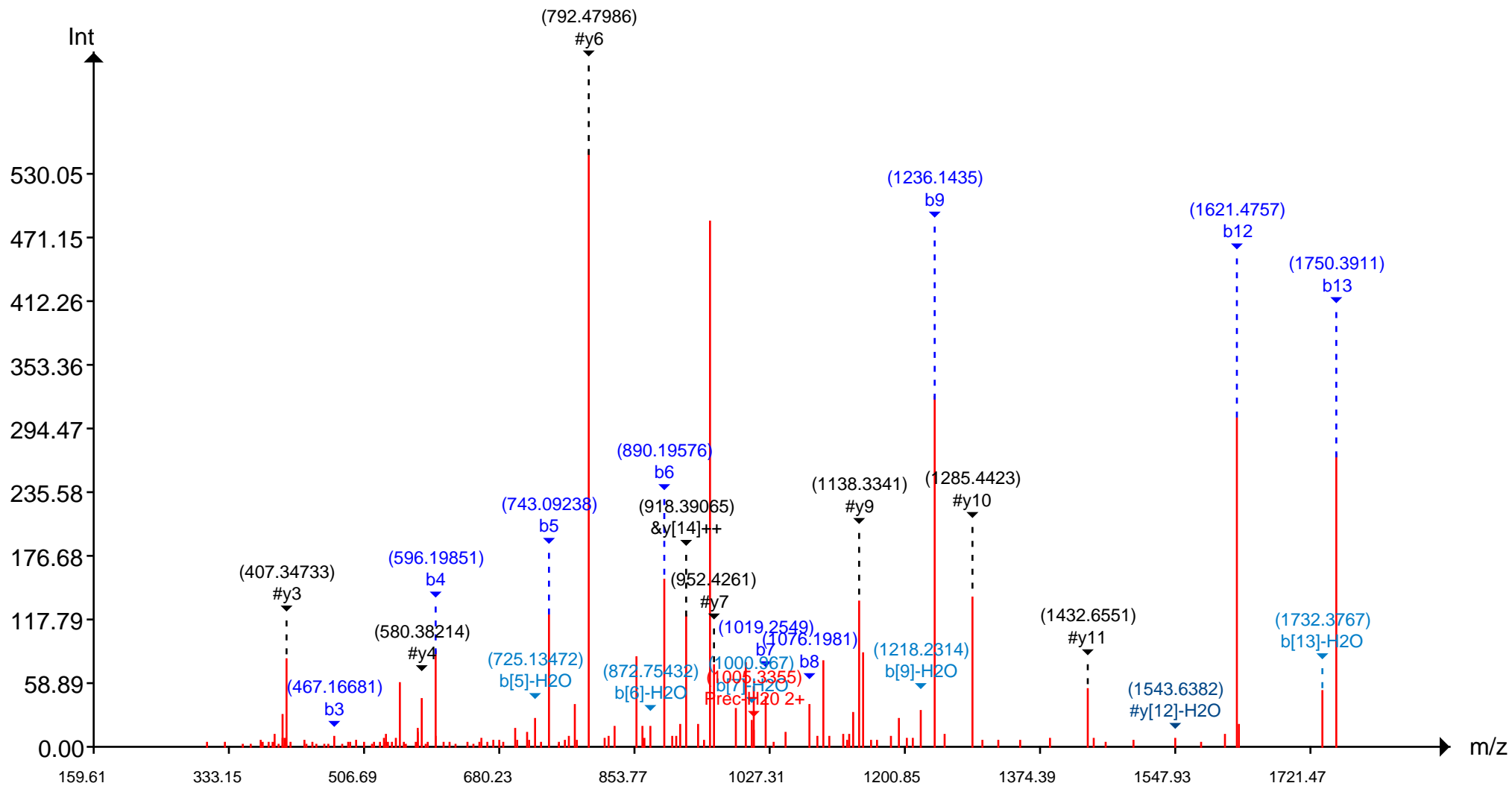
Charge 2+

Mass error 0.001872



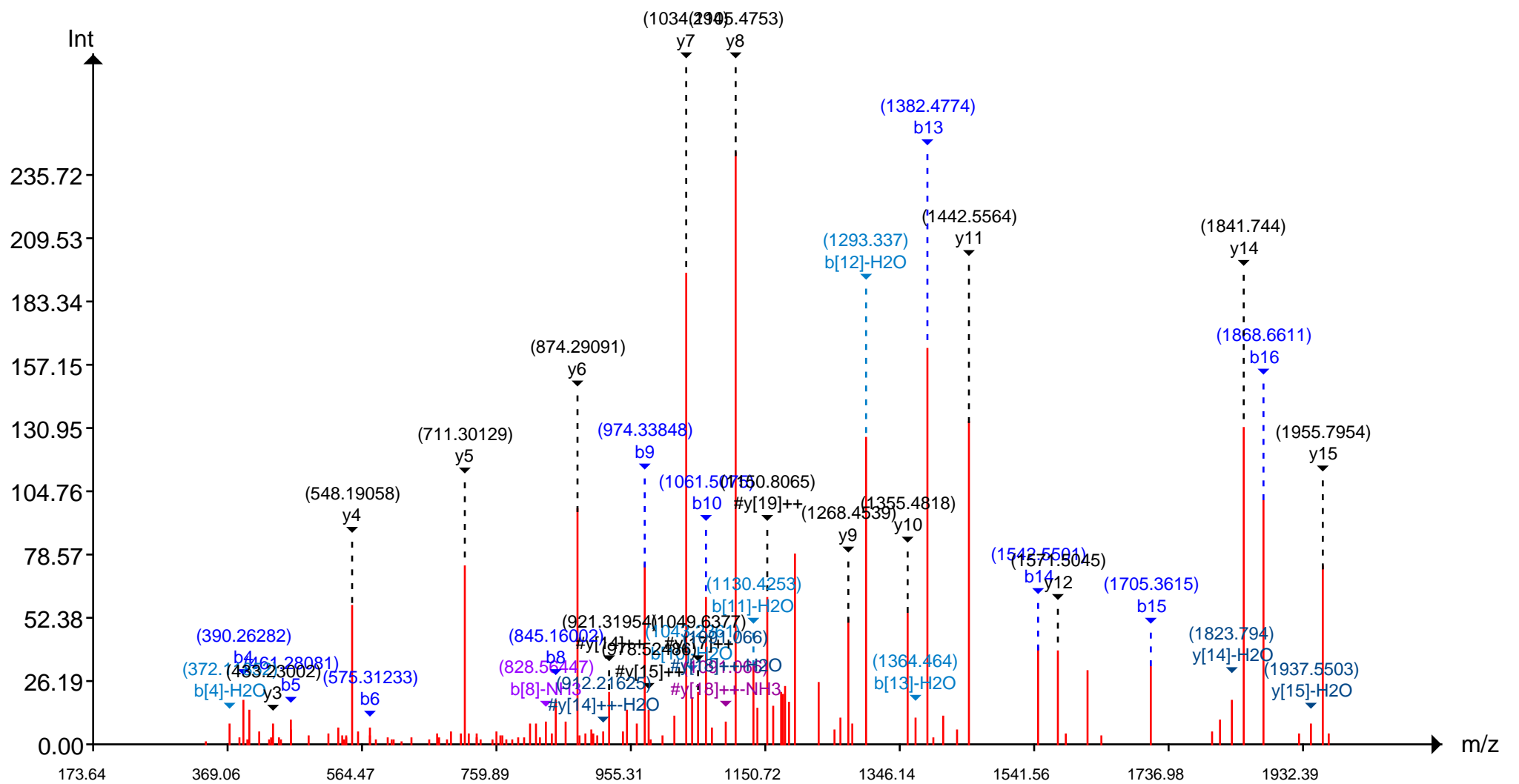


Info			
Modified Sequence	AcD3-FK<AcD3K*>NSQK<AcD3K*>VDR-COOH	Ion score	56.29
Protein	Q9CQM9	Confidence level	99.00%
Precursor m/z	628.84966	Identity Threshold	47.43
Charge	2+	Mass error	-0.00232

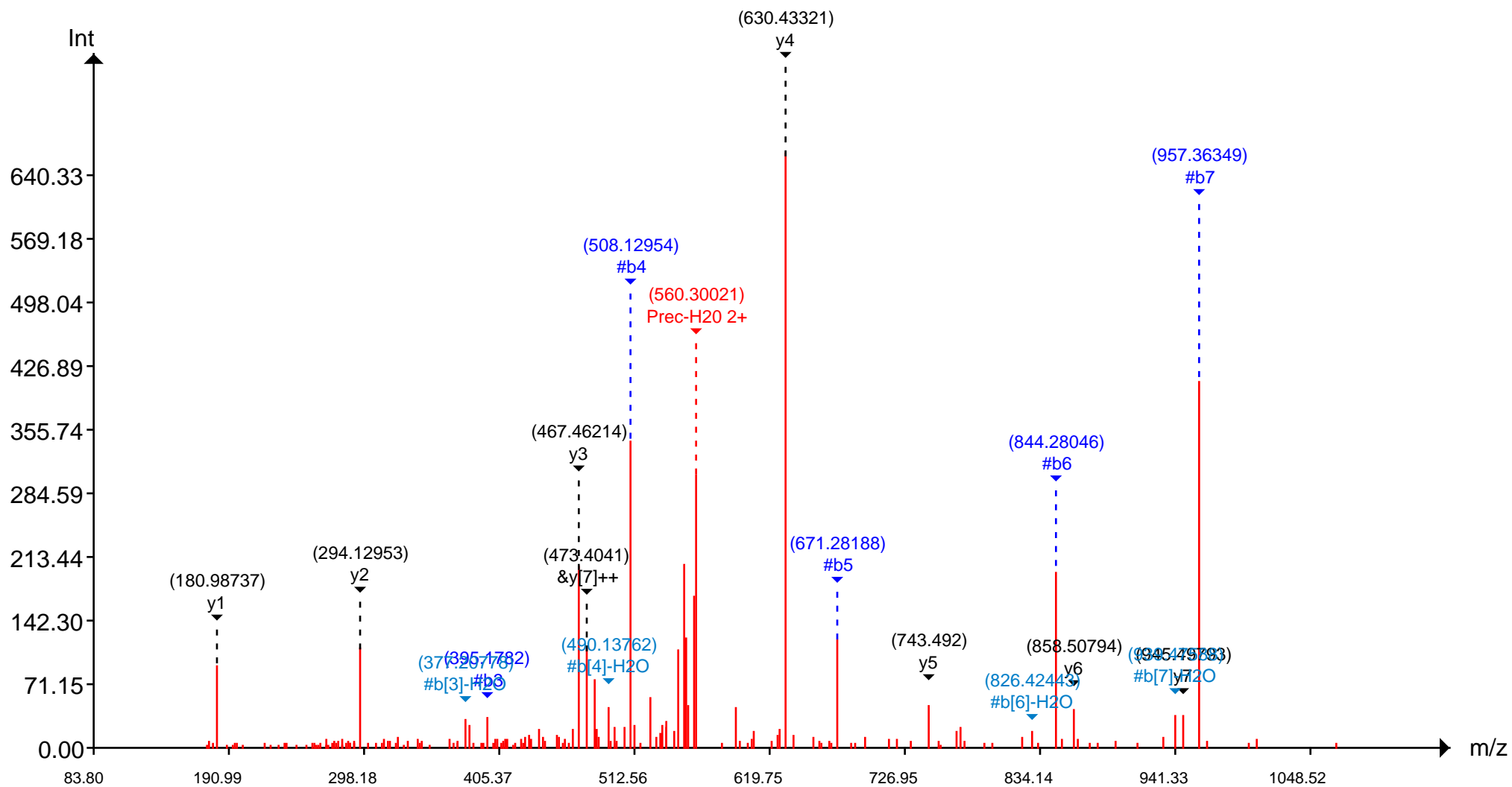


Info

Modified Sequence	AcD3-M<Mox*>C<Cmm*>NEFFEGC<Cmm*>PDK<AcD3K*>EPR<C13>-COOH	Ion score	40.88
Protein	P07091	Confidence level	99.00%
Precursor m/z	1014.4304	Identity Threshold	39.75
Charge	2+	Mass error	0.002513



Info			
Modified Sequence	AcD3-ATTAANPK<AcD3K*>ESSYAC<Cmm*>YYDEER-COOH	Ion score	96.54
Protein	O89017	Confidence level	99.00%
Precursor m/z	1208.5297	Identity Threshold	41.32
Charge	2+	Mass error	5.52E-4



Info

Modified Sequence AcD3-FSDIYK<AcD3K*>IR<C13>-COOH

Ion score 55.02

Protein P63028

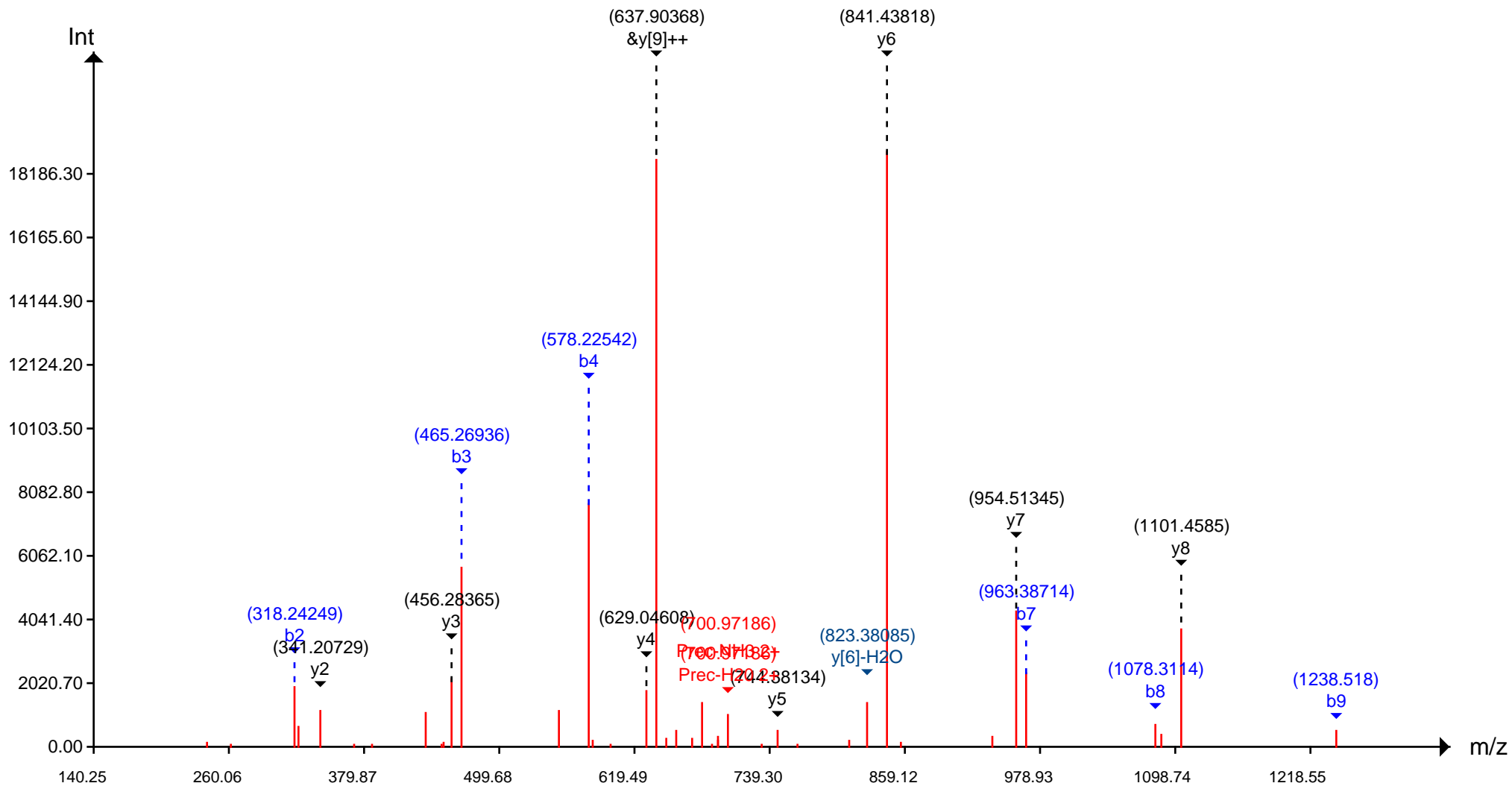
Confidence level 99.00%

Precursor m/z 569.32933

Identity Threshold 46.39

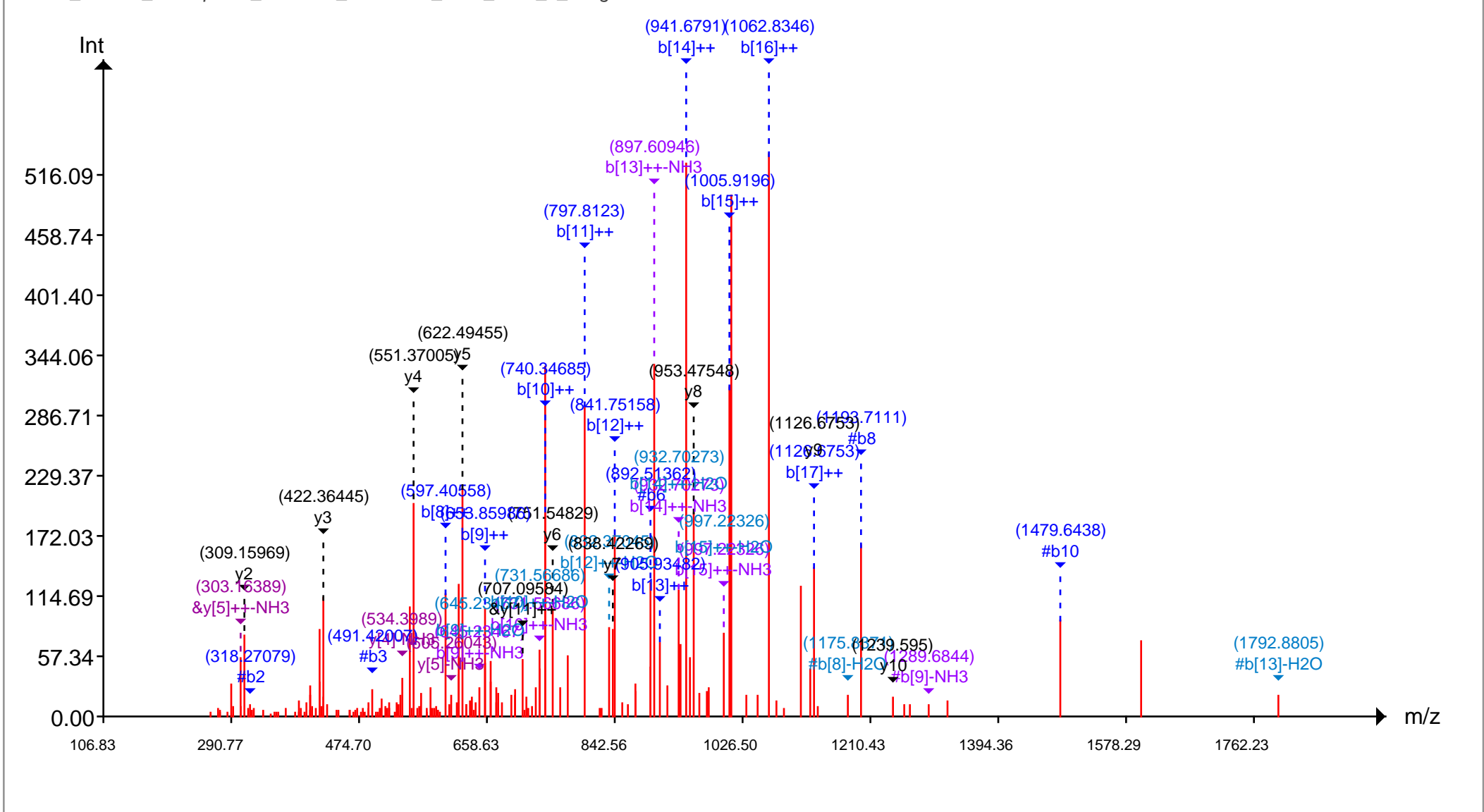
Charge 2+

Mass error -2.86E-4



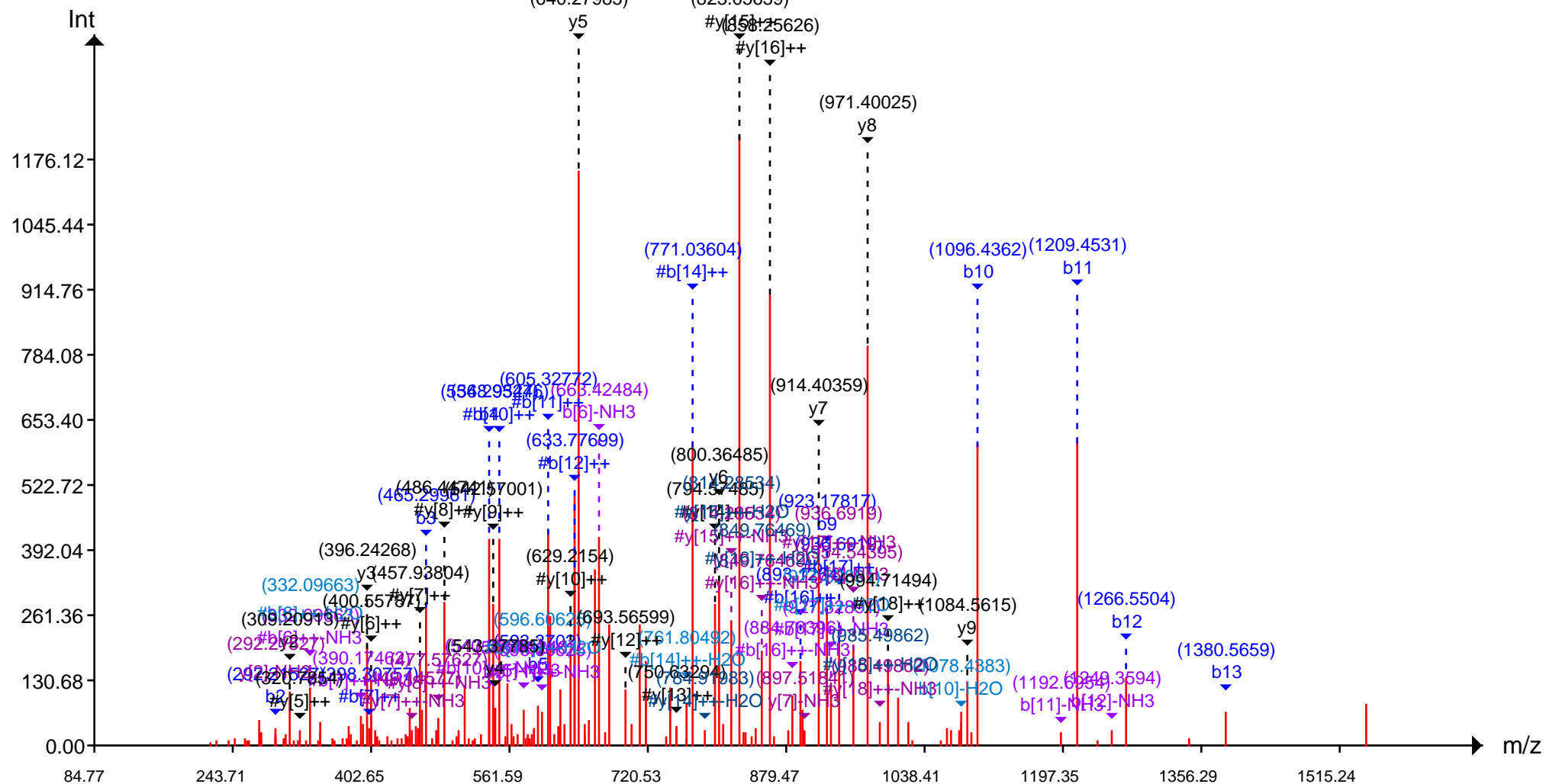
Info

Modified Sequence	AcD3-VK<AcD3K*>M<Mox*>LPDK<AcD3K*>DC<Cmm*>R<C13>-COOH	Ion score	48.59
Protein	P18760	Confidence level	99.00%
Precursor m/z	709.875	Identity Threshold	48.16
Charge	2+	Mass error	3.41E-4



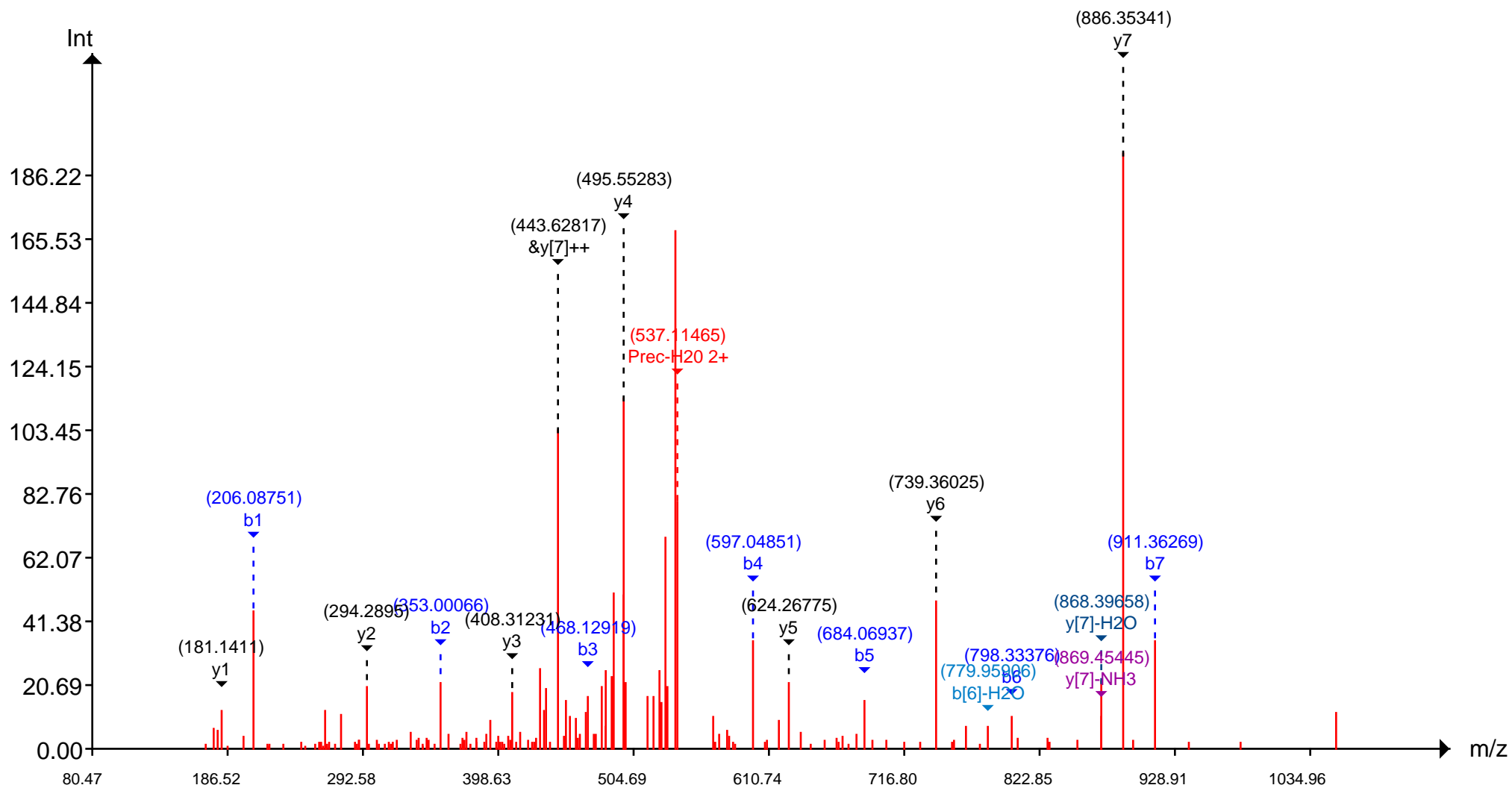
Info

Modified Seq...	AcD3-K<AcD3K*>VK<AcD3K*>EK<AcD3K*>VQK<AcD3K*>LK<AcD3K*>DSEAELQR<C13>-COOH	Ion score	54.18
P...	O55131	Confidence level	99.00
Precurso...	811.4845	Identity Threshold	40.27
C...	3+	Mass error	0.002



Info

Modified Sequence	AcD3-FVK<AcD3K*>AGSDGAK<AcD3K*>IGNC<Cmm*>PFSQR<C13>-COOH	Ion score	69.11
Protein	Q9Z1Q5	Confidence level	99.00%
Precursor m/z	727.37904	Identity Threshold	48.04
Charge	3+	Mass error	0.001507



Info

Modified Sequence AcD3-C<Cmm*>M<Mox*>DESNLR<C13>-COOH

Ion score 51.62

Protein Q9D358

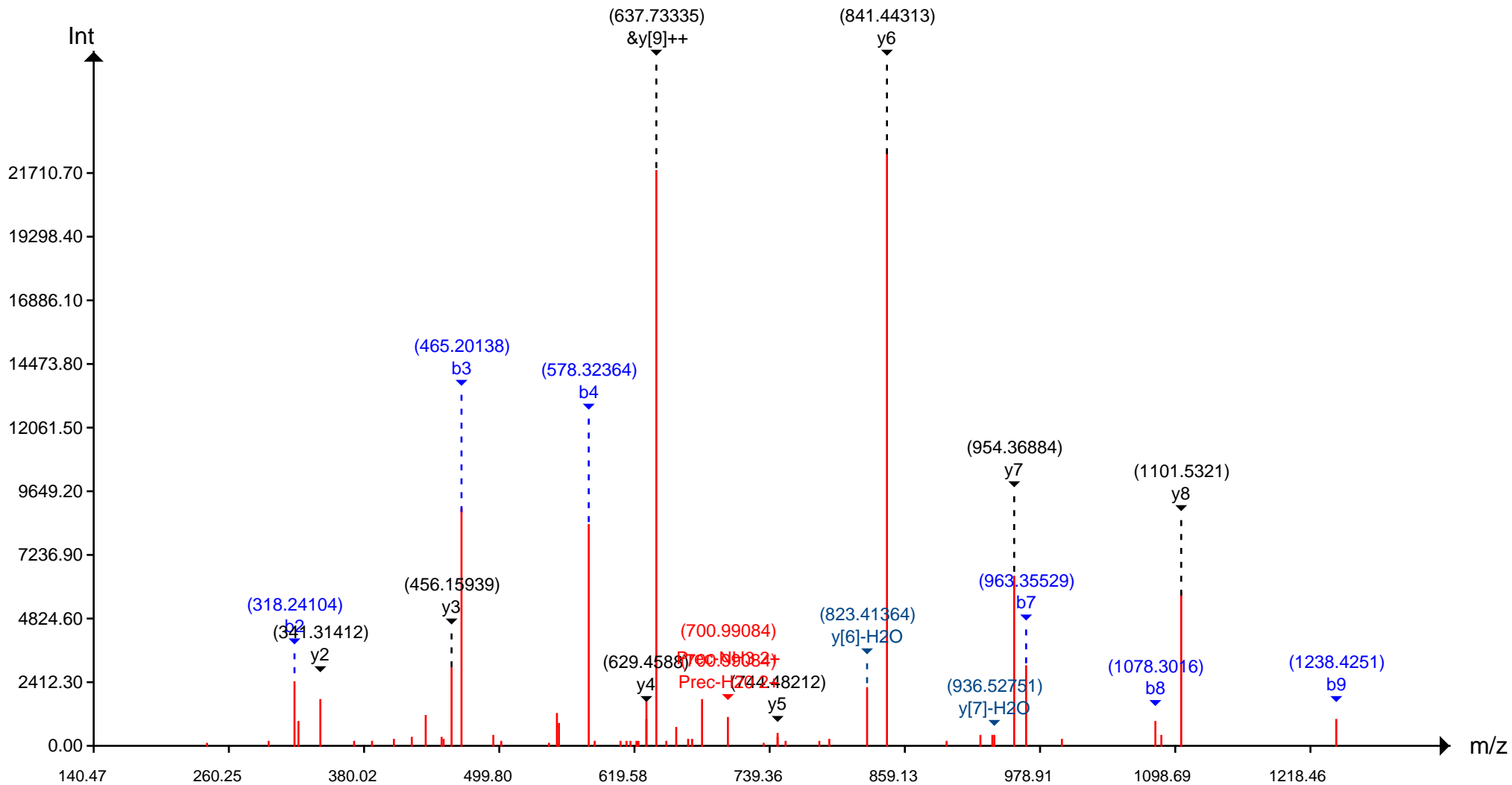
Confidence level 99.00%

Precursor m/z 546.2346

Identity Threshold 41.51

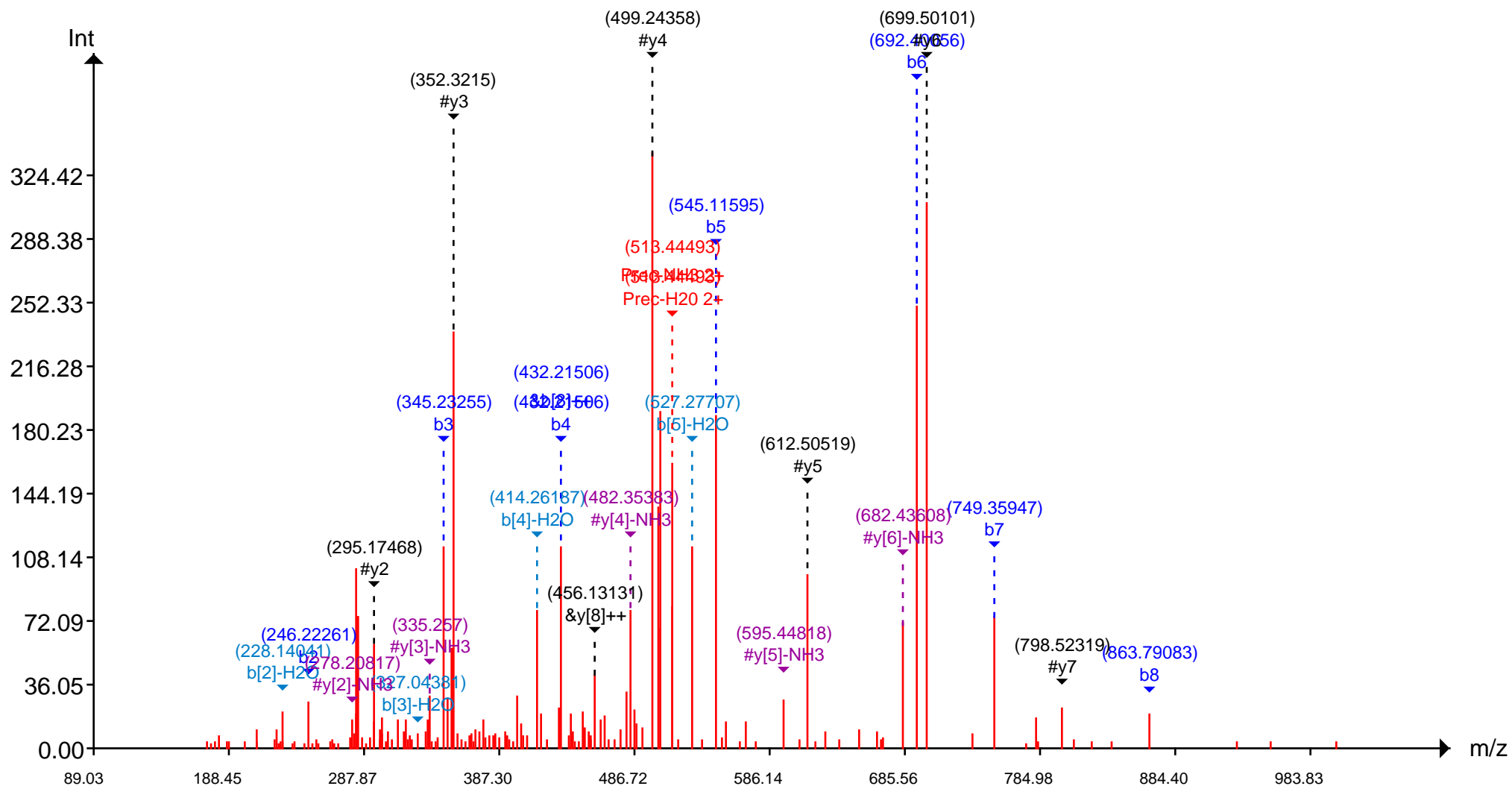
Charge 2+

Mass error -0.001147



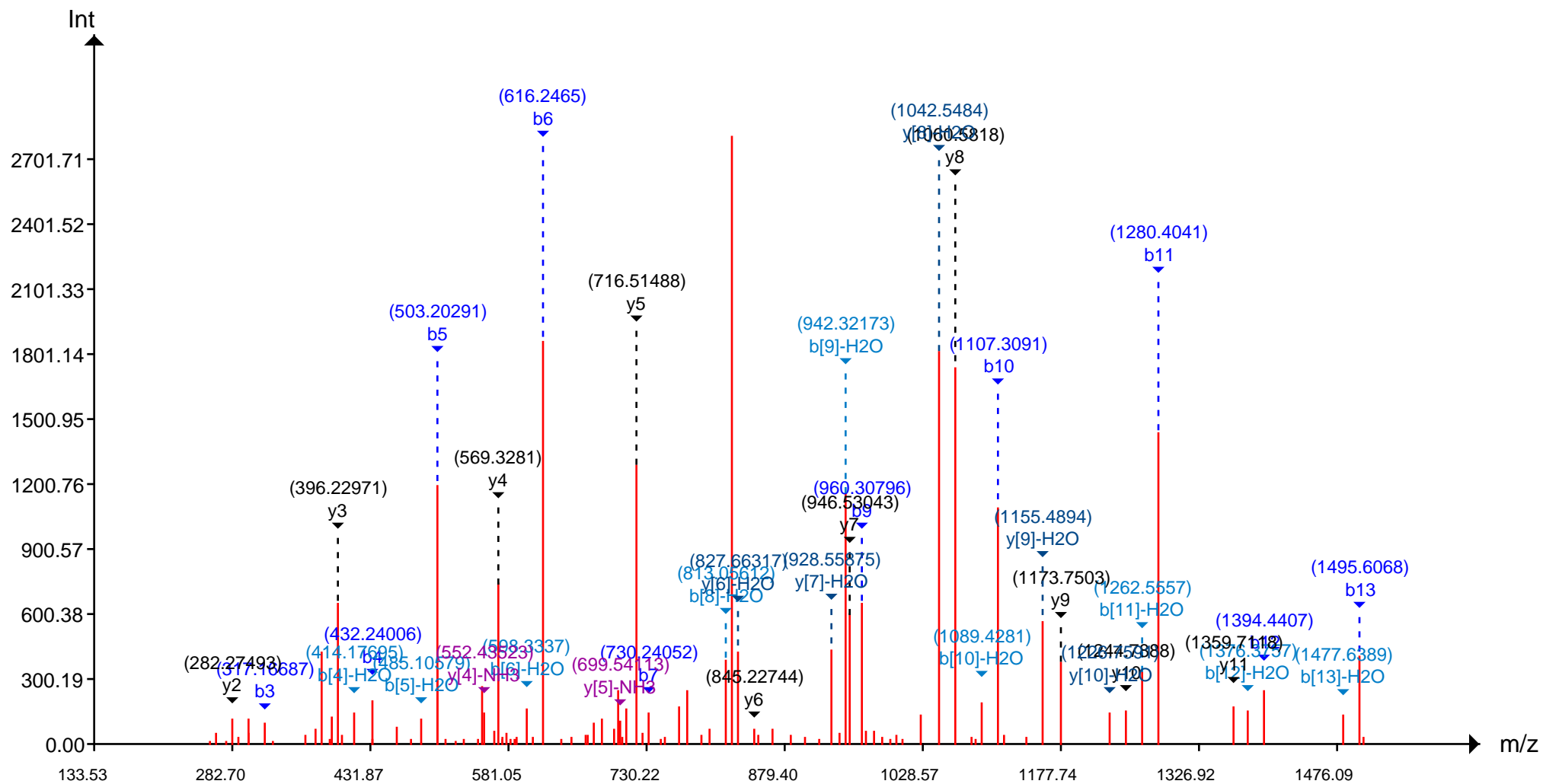
Info

Modified Sequence	AcD3-VK<AcD3K*>M<Mox*>LPDK<AcD3K*>DC<Cmm*>R<C13>-COOH	Ion score	53.9
Protein	P18760	Confidence level	99.00%
Precursor m/z	709.87486	Identity Threshold	48.15
Charge	2+	Mass error	6.1E-5



Info

Modified Sequence	AcD3-SLVSLFGNR<C13>-COOH	Ion score	56.38
Protein	P99029	Confidence level	99.00%
Precursor m/z	522.30434	Identity Threshold	46.87
Charge	2+	Mass error	-4.76E-4



Info

Modified Sequence AcD3-SLADAINTEFK<AcD3K*>NTR<C13>-COOH

Ion score 87.57

Protein P20152

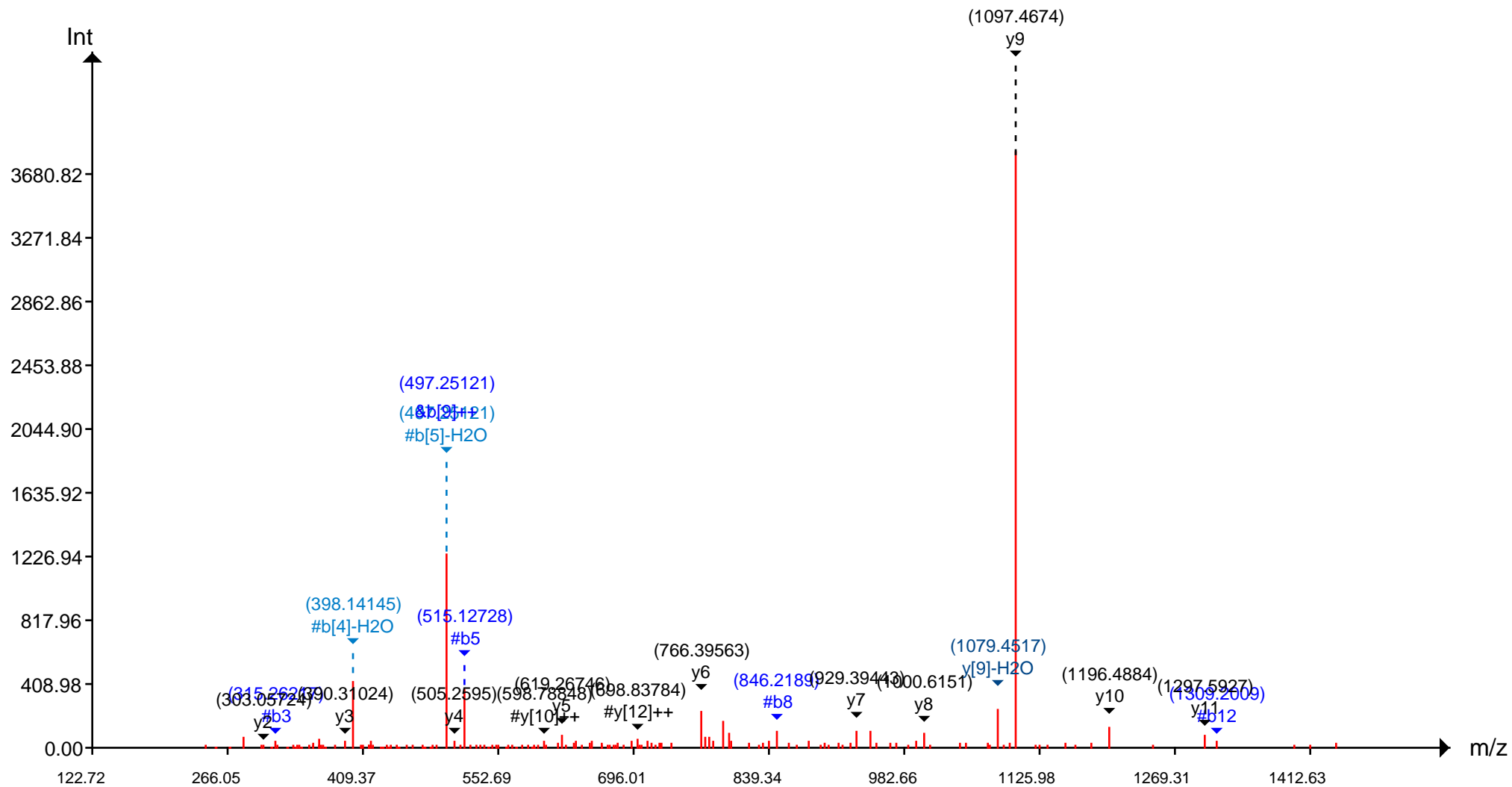
Confidence level 99.00%

Precursor m/z 838.44821

Identity Threshold 48.01

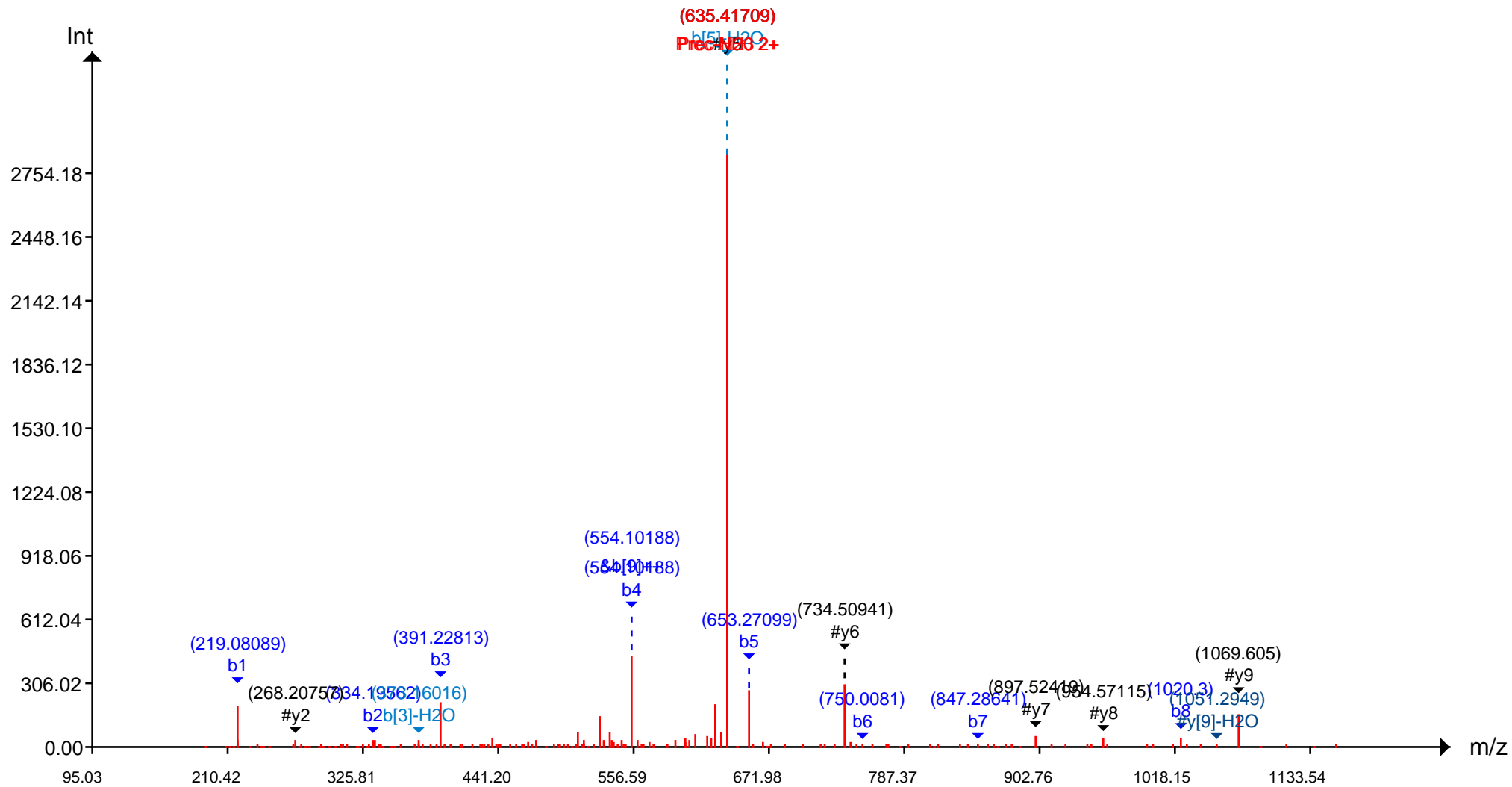
Charge 2+

Mass error 0.002779



Info

Modified Sequence	AcD3-AVVTVPAYFNDSQR-COOH	Ion score	67.51
Protein	P63017	Confidence level	99.00%
Precursor m/z	806.41473	Identity Threshold	48.09
Charge	2+	Mass error	0.001706



Info

Modified Sequence AcD3-K<AcD3K*>DGYVPPK<AcD3K*>SR<C13>-COOH

Ion score 54.92

Protein O89053

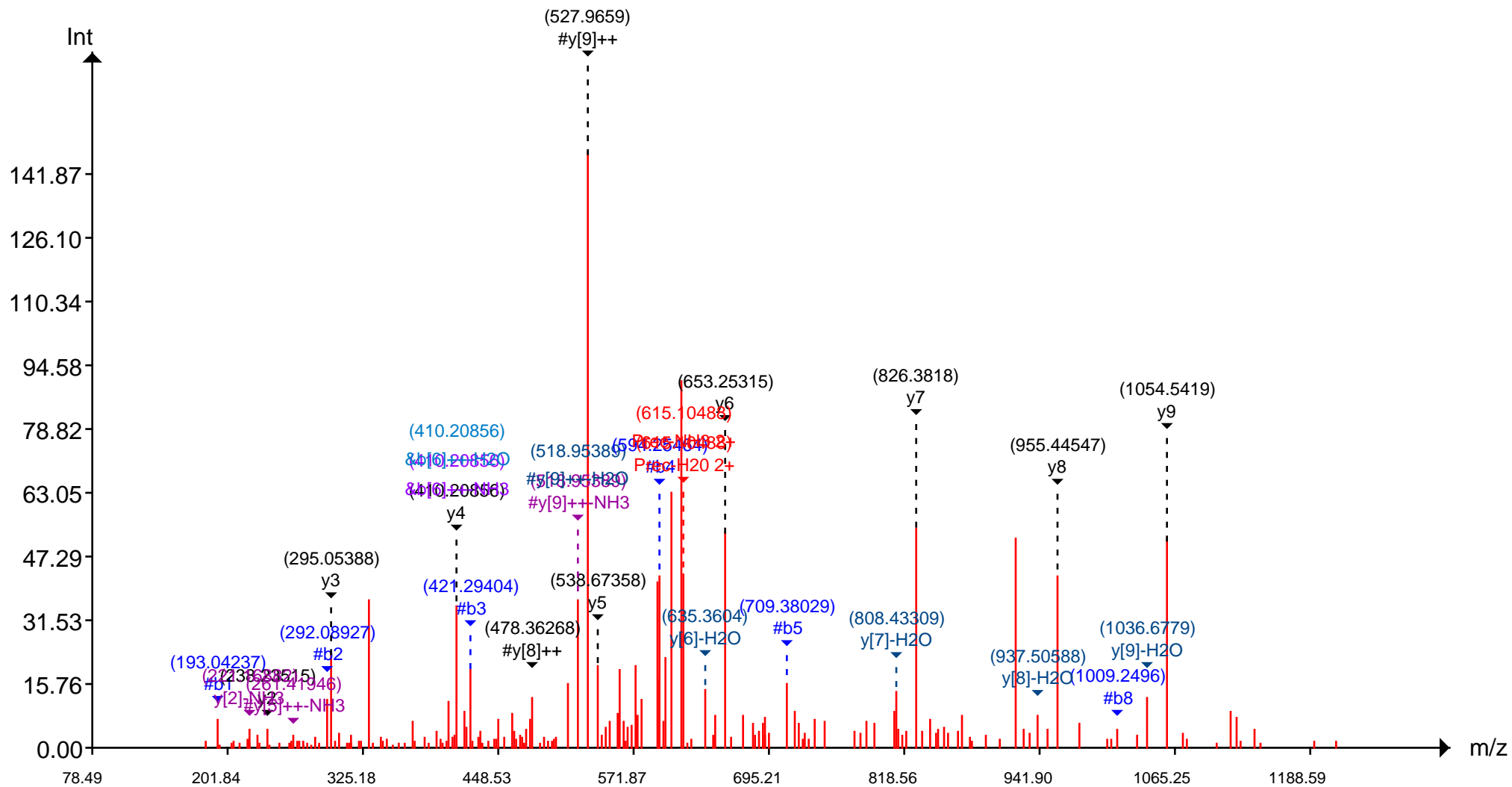
Confidence level 99.00%

Precursor m/z 644.37167

Identity Threshold 46.64

Charge 2+

Mass error 0.001157



Info

Modified Sequence AcD3-M<Mox*>VEK<AcD3K*>DQDGGR<C13>-COOH

Ion score 54.8

Protein Q9CZX8

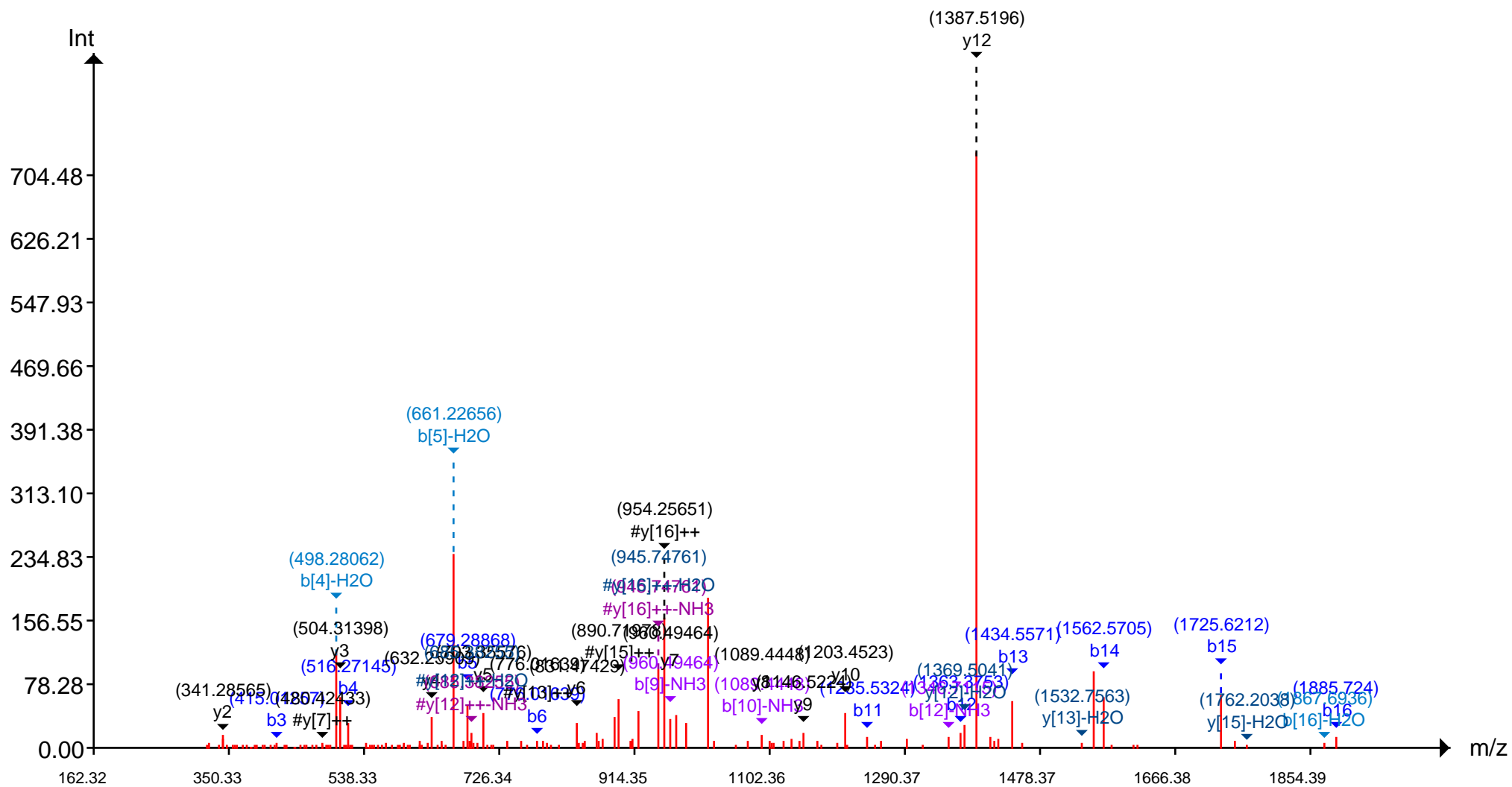
Confidence level 99.00%

Precursor m/z 623.80087

Identity Threshold 46.40

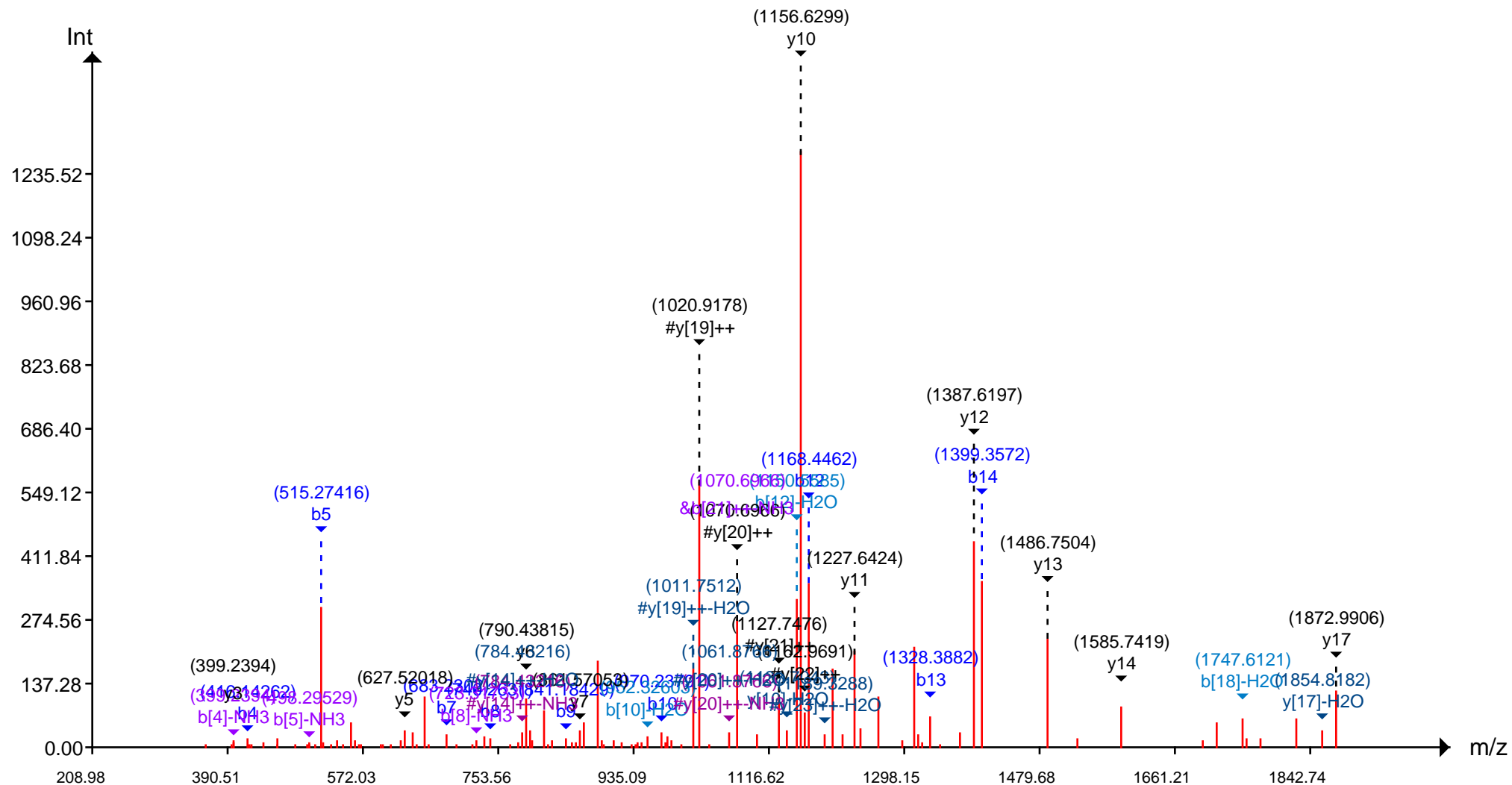
Charge 2+

Mass error -1.84E-4



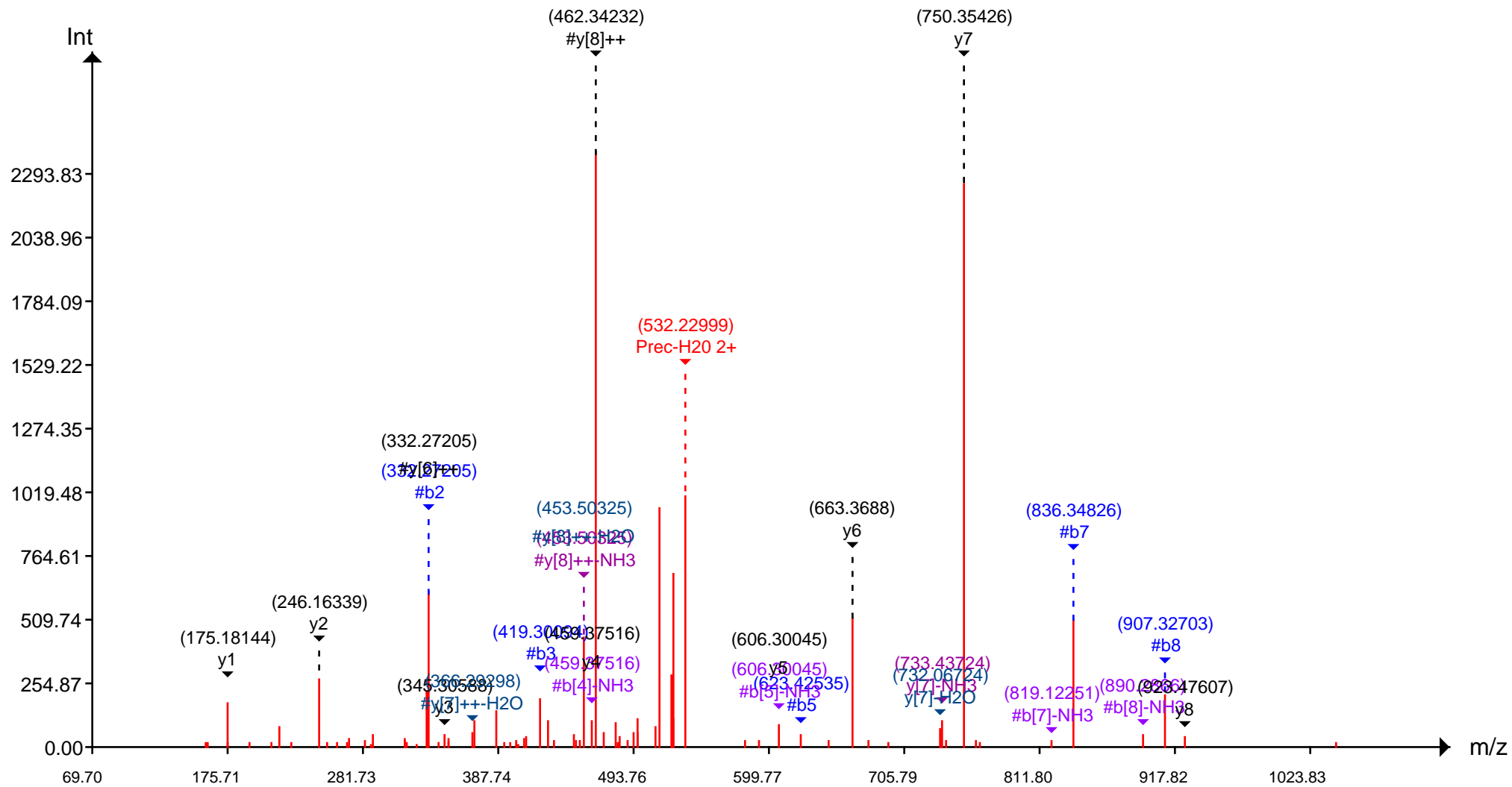
Info

Modified Sequence	AcD3-IQQTYPSSGEEQAQYC<Cmm*>R<C13>-COOH	Ion score	59.22
Protein	Q9WU78	Confidence level	99.00%
Precursor m/z	1033.4757	Identity Threshold	44.61
Charge	2+	Mass error	0.002278



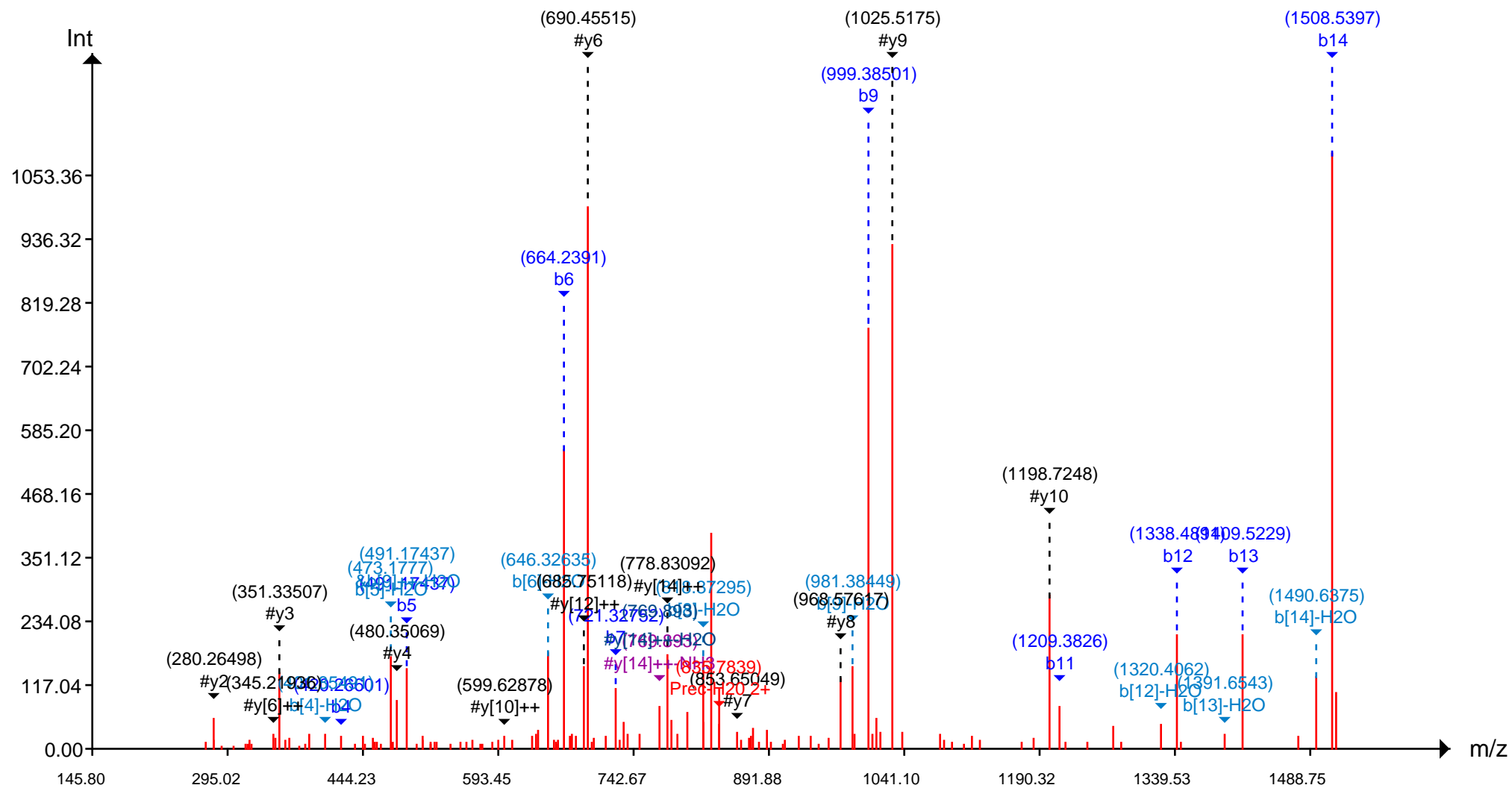
Info

Modified Sequence	AcD3-NAANVPAGTEVVC<Cmm*>APPTAYIDFAR<C13>-COOH	Ion score	54.71
Protein	P17751	Confidence level	99.00%
Precursor m/z	1278.1417	Identity Threshold	47.70
Charge	2+	Mass error	0.002765



Info

Modified Sequence	AcD3-IK<AcD3K*>SGM<Mox*>NVAR-COOH	Ion score	48.67
Protein	P52480	Confidence level	99.00%
Precursor m/z	541.30071	Identity Threshold	47.67
Charge	2+	Mass error	4.5E-5



Info

Modified Sequence AcD3-SGETAK<AcD3K*>GDYPLEAVR<C13>-COOH

Ion score 64.58

Protein P52480

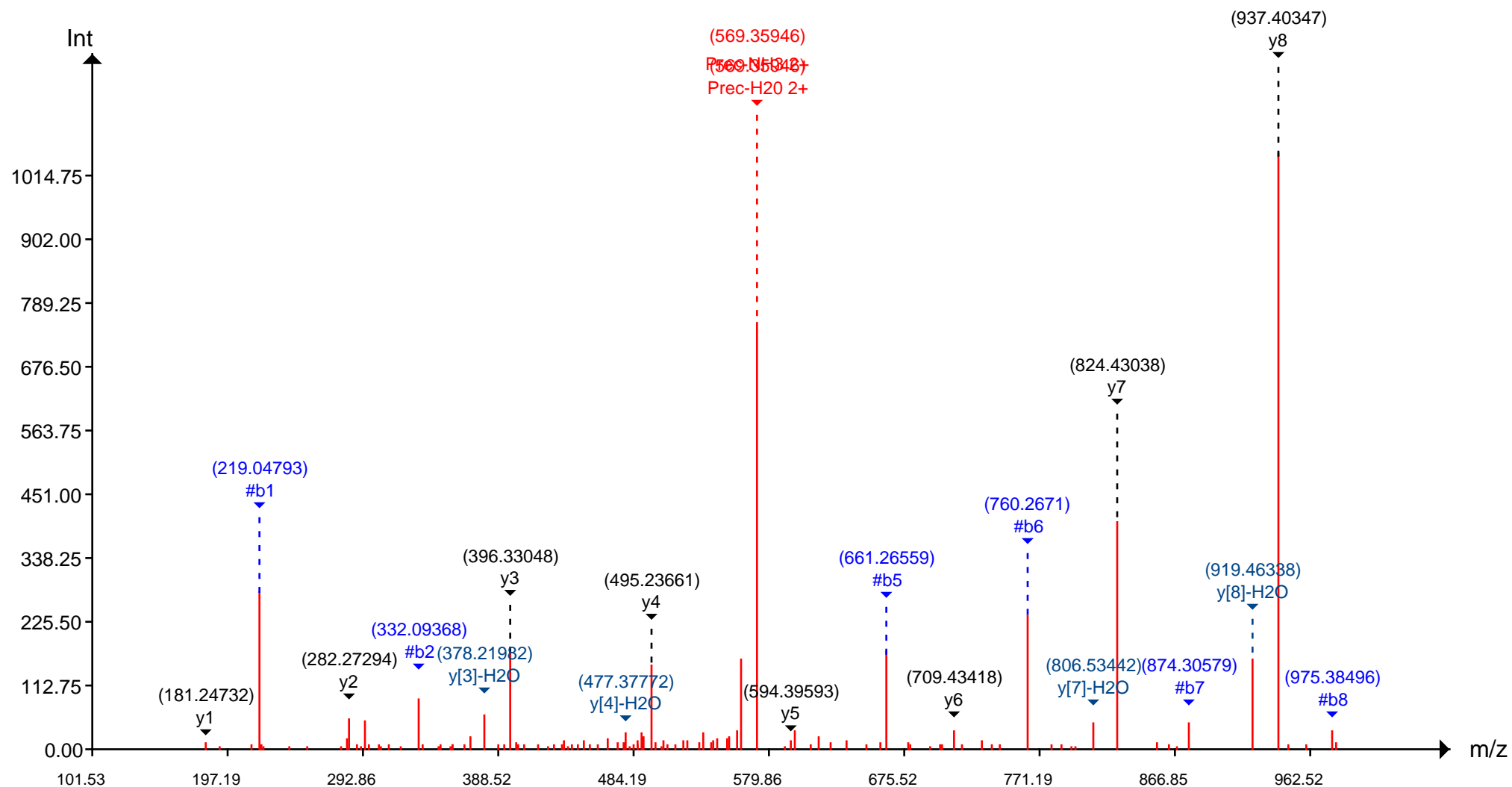
Confidence level 99.00%

Precursor m/z 844.93978

Identity Threshold 48.05

Charge 2+

Mass error 0.001895



Info

Modified Sequence AcD3-K<AcD3K*>IDDVVNTR<C13>-COOH

Ion score 56.32

Protein P80315

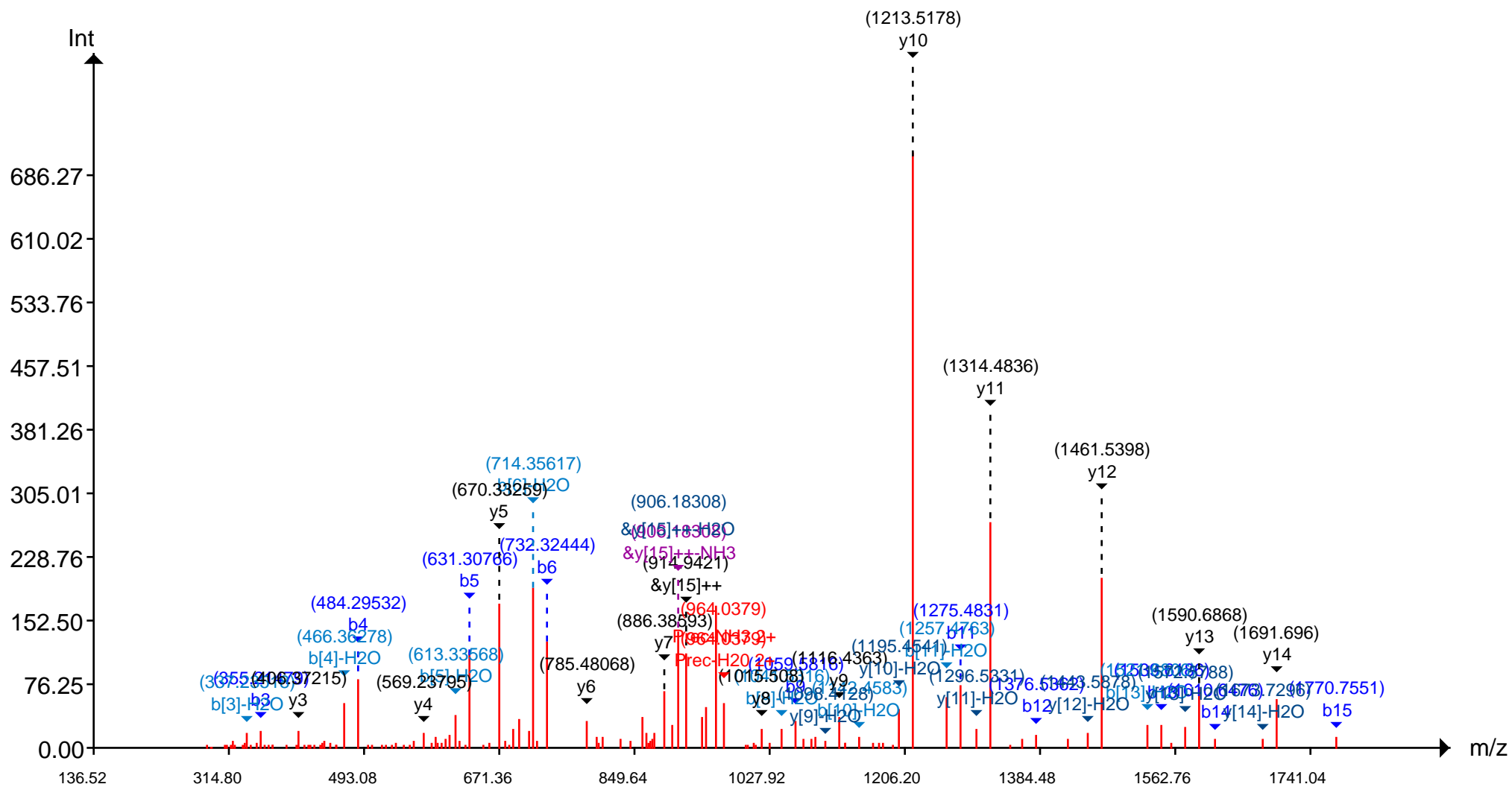
Confidence level 99.00%

Precursor m/z 578.33293

Identity Threshold 46.51

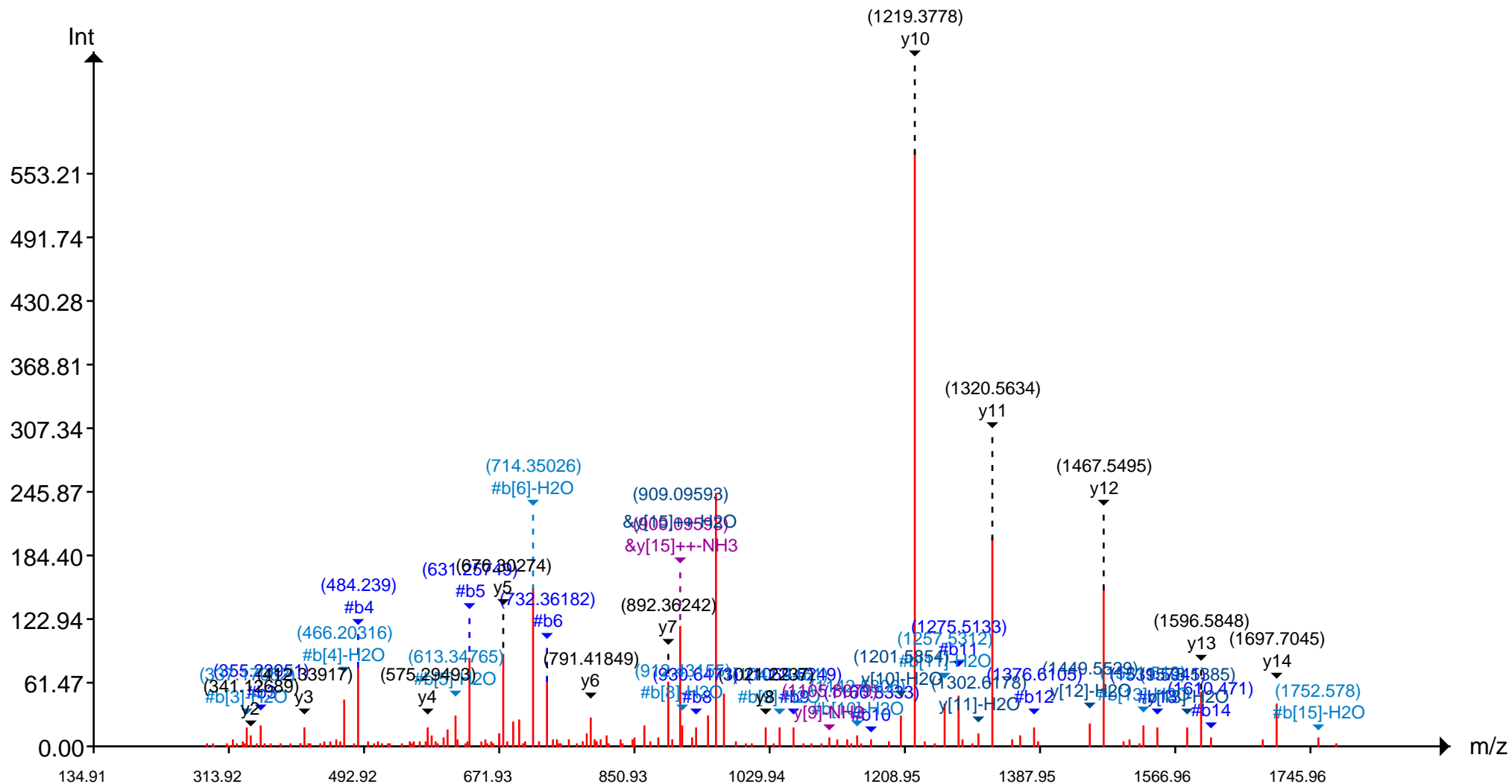
Charge 2+

Mass error 3.53E-4



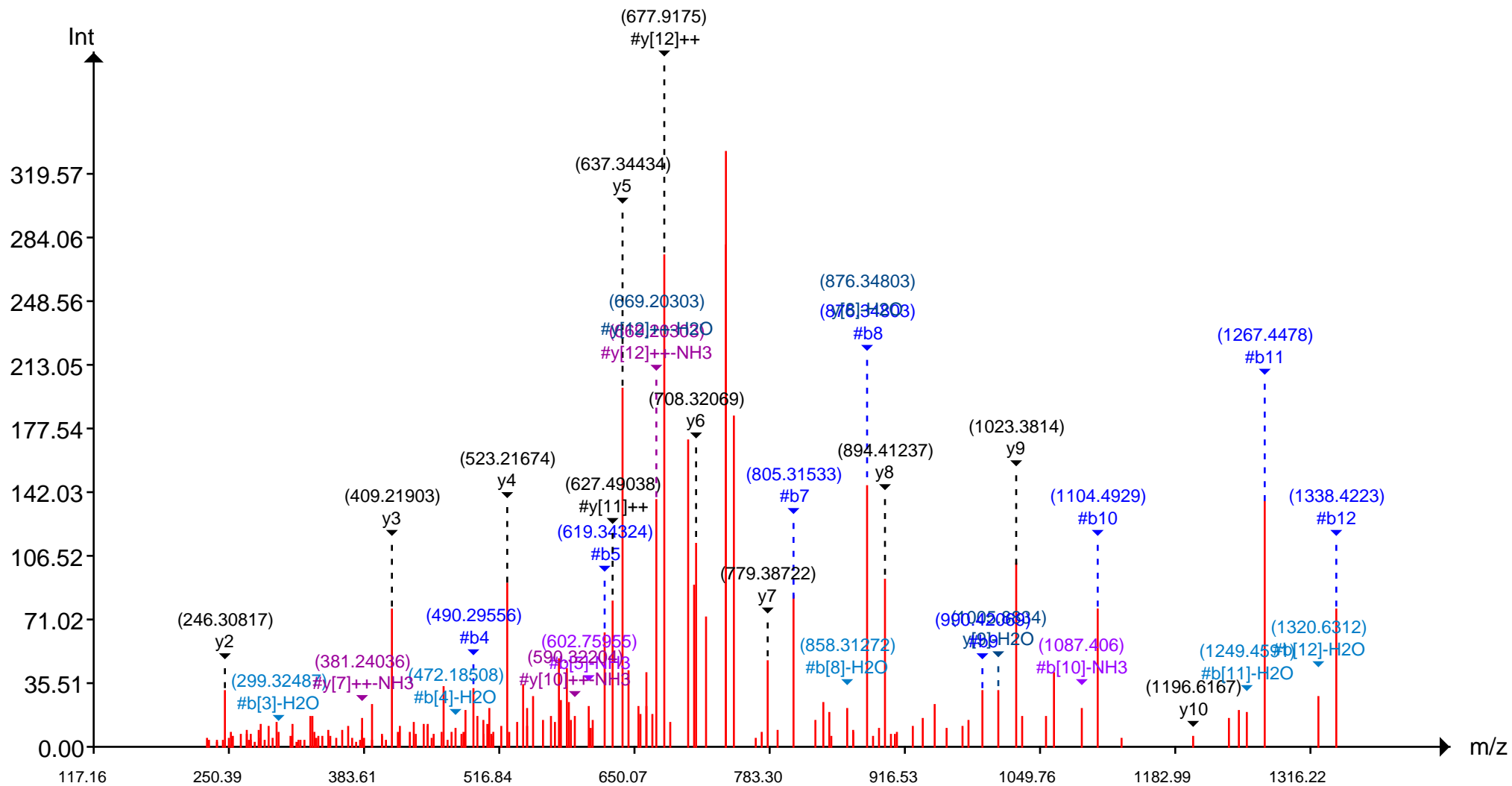
Info

Modified Sequence	AcD3-AHTEFTPTEPTYAC<Cmm*>R-COOH	Ion score	104.22
Protein	P01887	Confidence level	99.00%
Precursor m/z	972.92818	Identity Threshold	42.34
Charge	2+	Mass error	0.001918



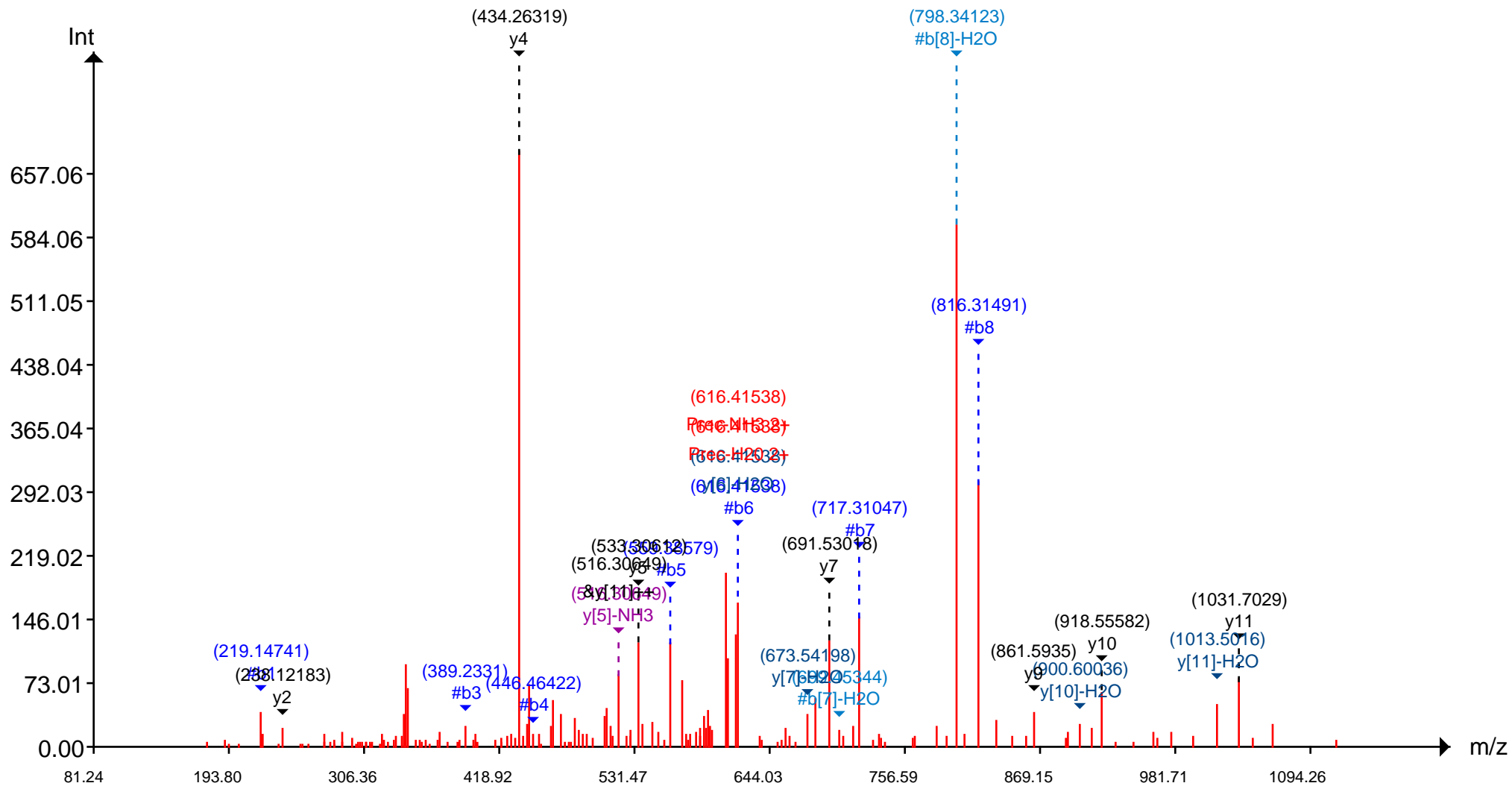
Info

Modified Sequence	AcD3-AHTEFTPTETDTYAC<Cmm*>R<C13>-COOH	Ion score	109.37
Protein	P01887	Confidence level	99.00%
Precursor m/z	975.9389	Identity Threshold	43.42
Charge	2+	Mass error	0.003217



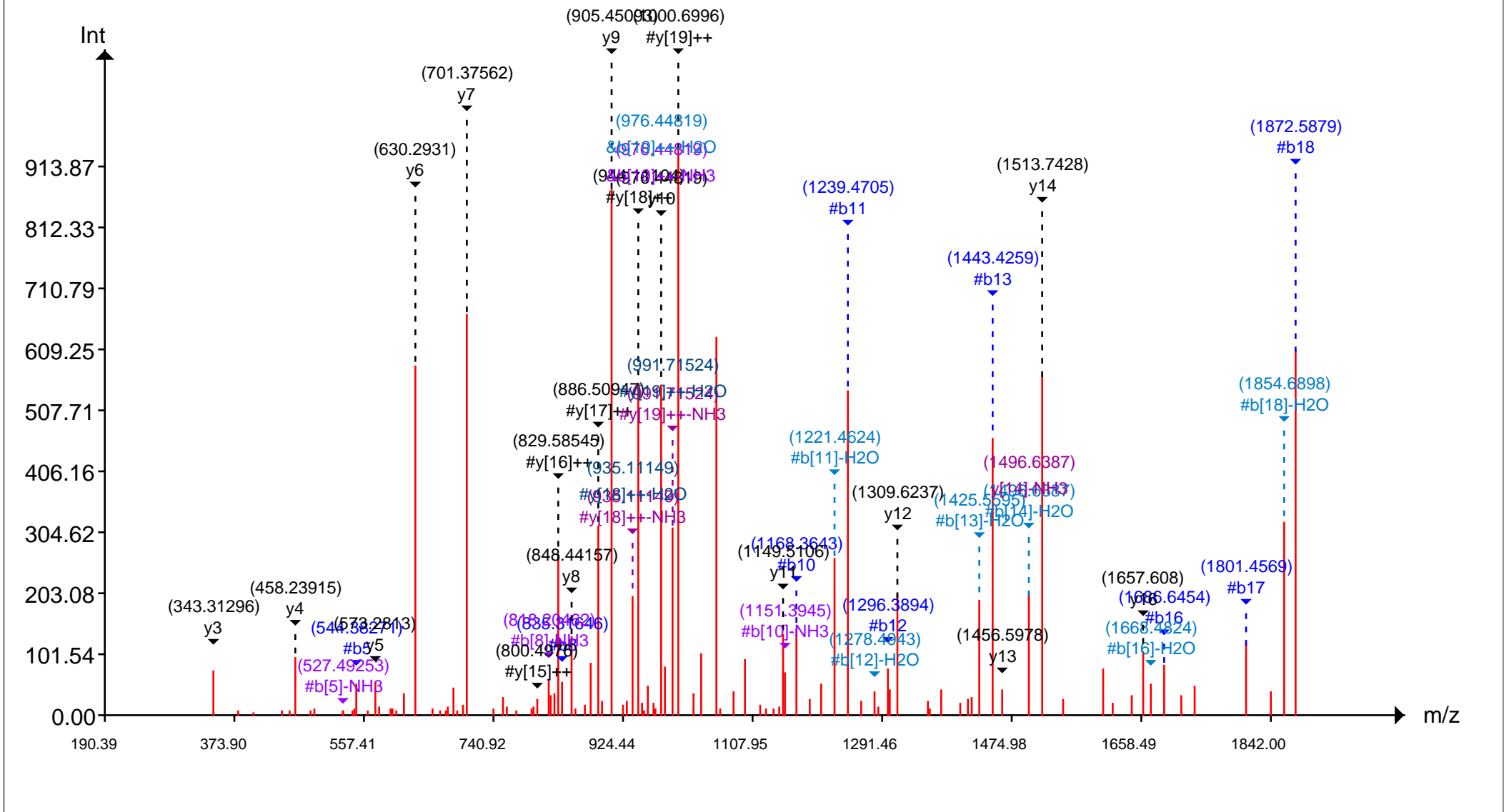
Info

Modified Sequence	AcD3-ITGK<AcD3K*>EDAANNYAR-COOH	Ion score	52.43
Protein	P68369, P05213, P68373, P05214, P68368, Q9JJZ2	Confidence level	99.00%
Precursor m/z	756.88279	Identity Threshold	47.85
Charge	2+	Mass error	0.002371



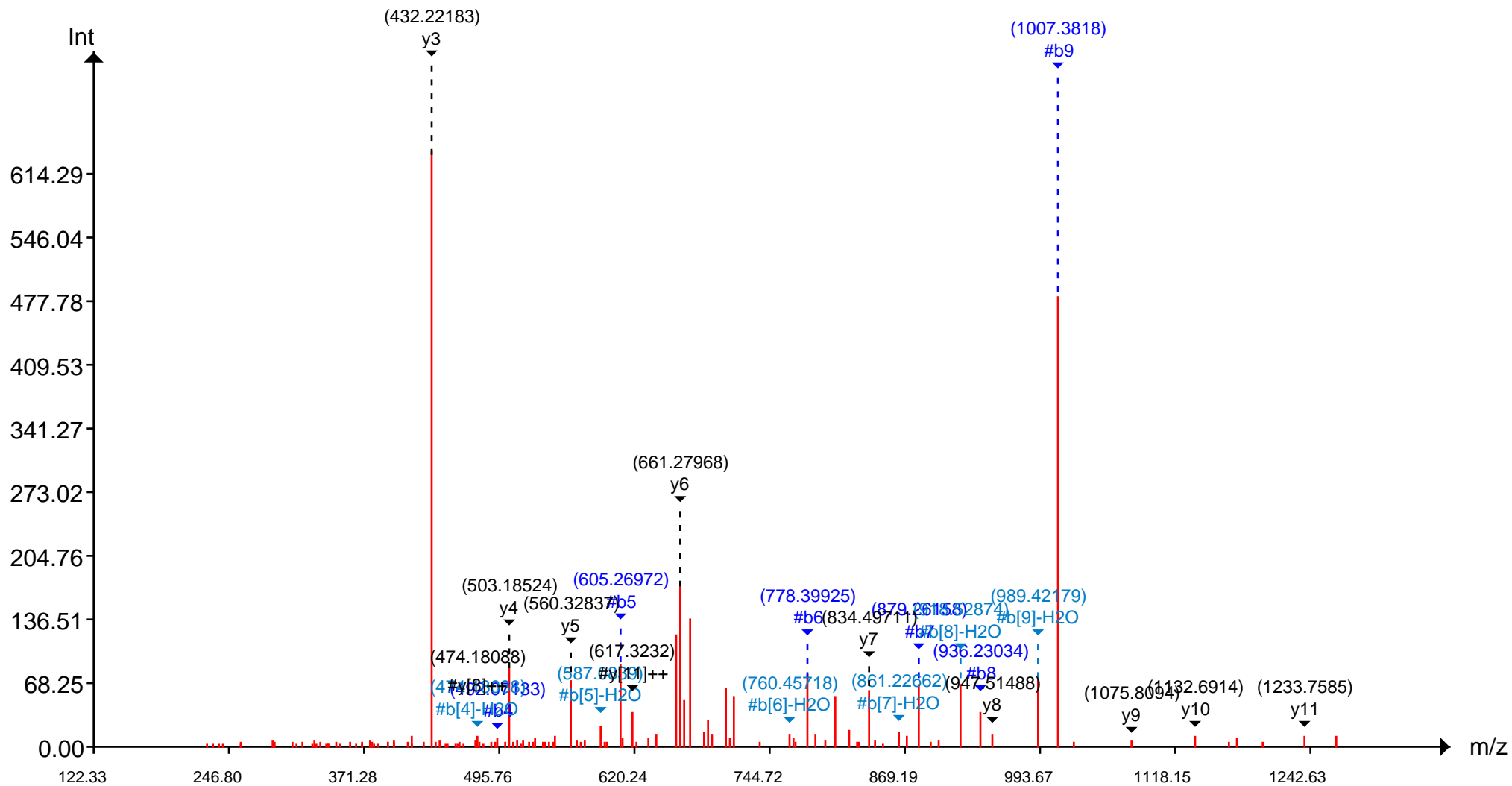
Info

Modified Sequence	AcD3-K<AcD3K*>IGGIGTVPVGR<C13>-COOH	Ion score	44.48
Protein	P10126, P62631	Confidence level	99.00%
Precursor m/z	625.39615	Identity Threshold	40.67
Charge	2+	Mass error	9.08E-4



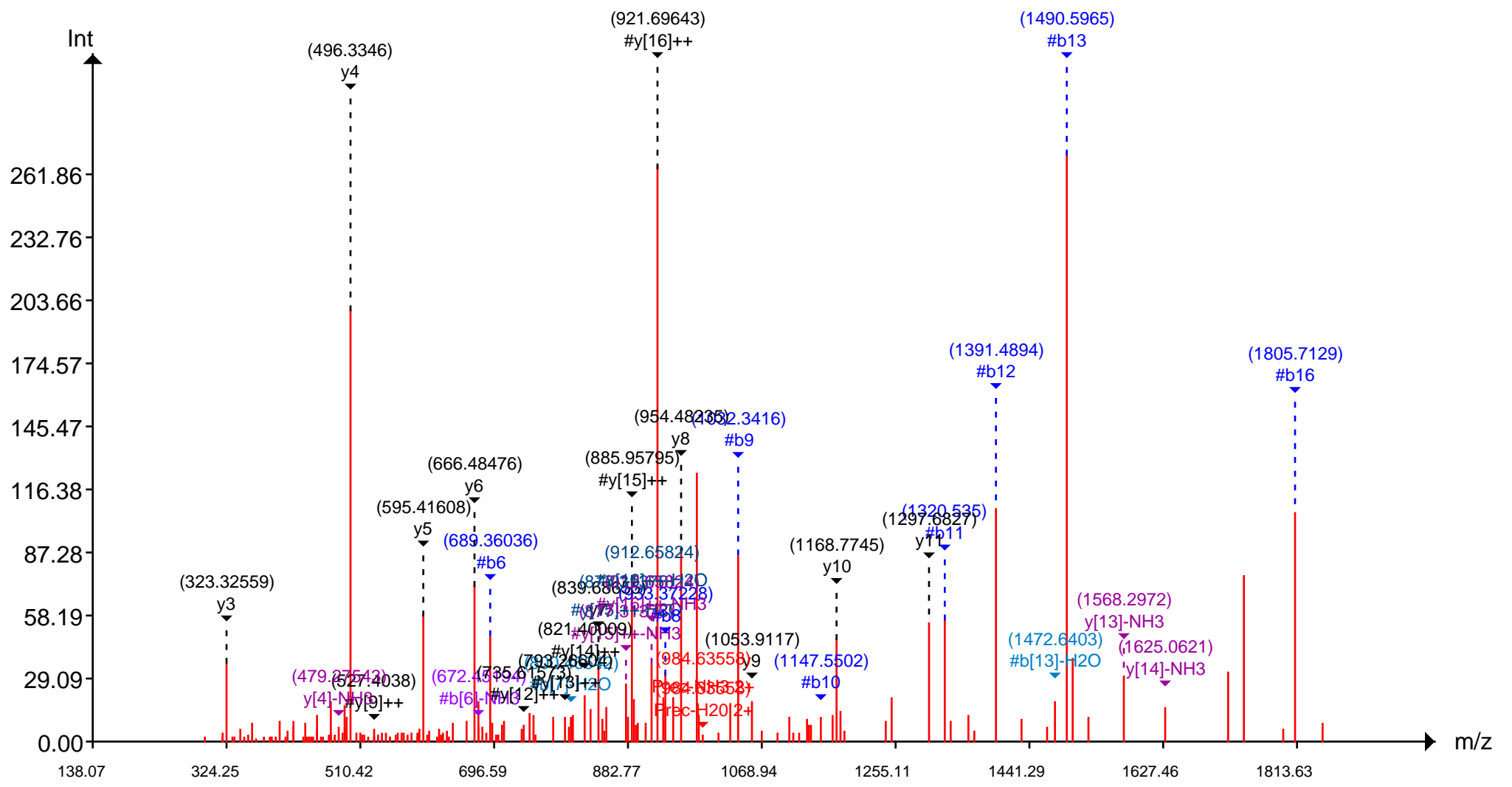
Info

Modified Sequence	AcD3-VIDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR-COOH	Ion score	93.05
Protein	P63260	Confidence level	99.00%
Precursor m/z	1072.4876	Identity Threshold	44.00
Charge	2+	Mass error	0.002518



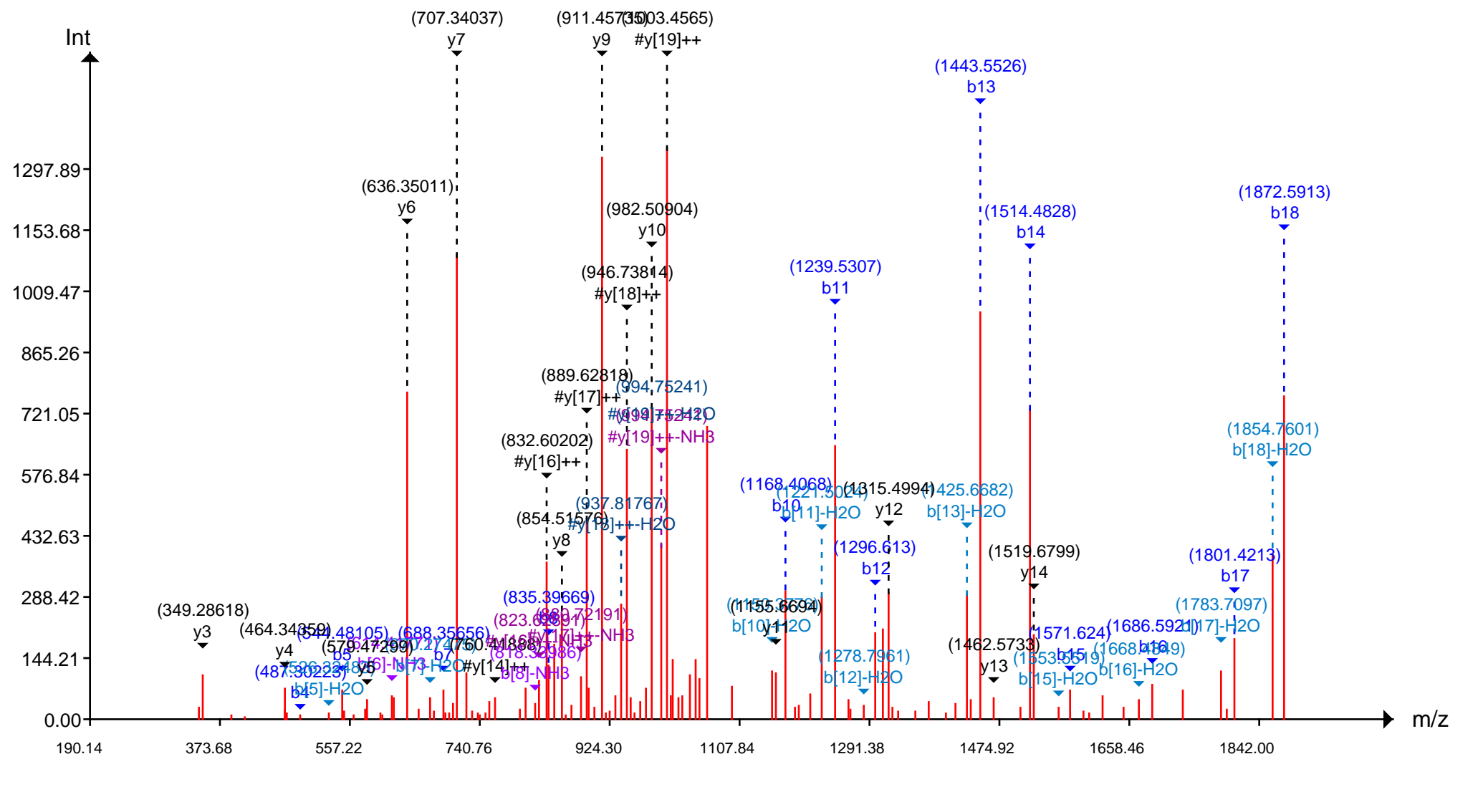
Info

Modified Sequence	AcD3-C<Cmm*>TGQIK<AcD3K*>TGAPC<Cmm*>R-COOH	Ion score	61.58
Protein	P17182, P21550, P17183	Confidence level	99.00%
Precursor m/z	719.8565	Identity Threshold	47.45
Charge	2+	Mass error	9.08E-4

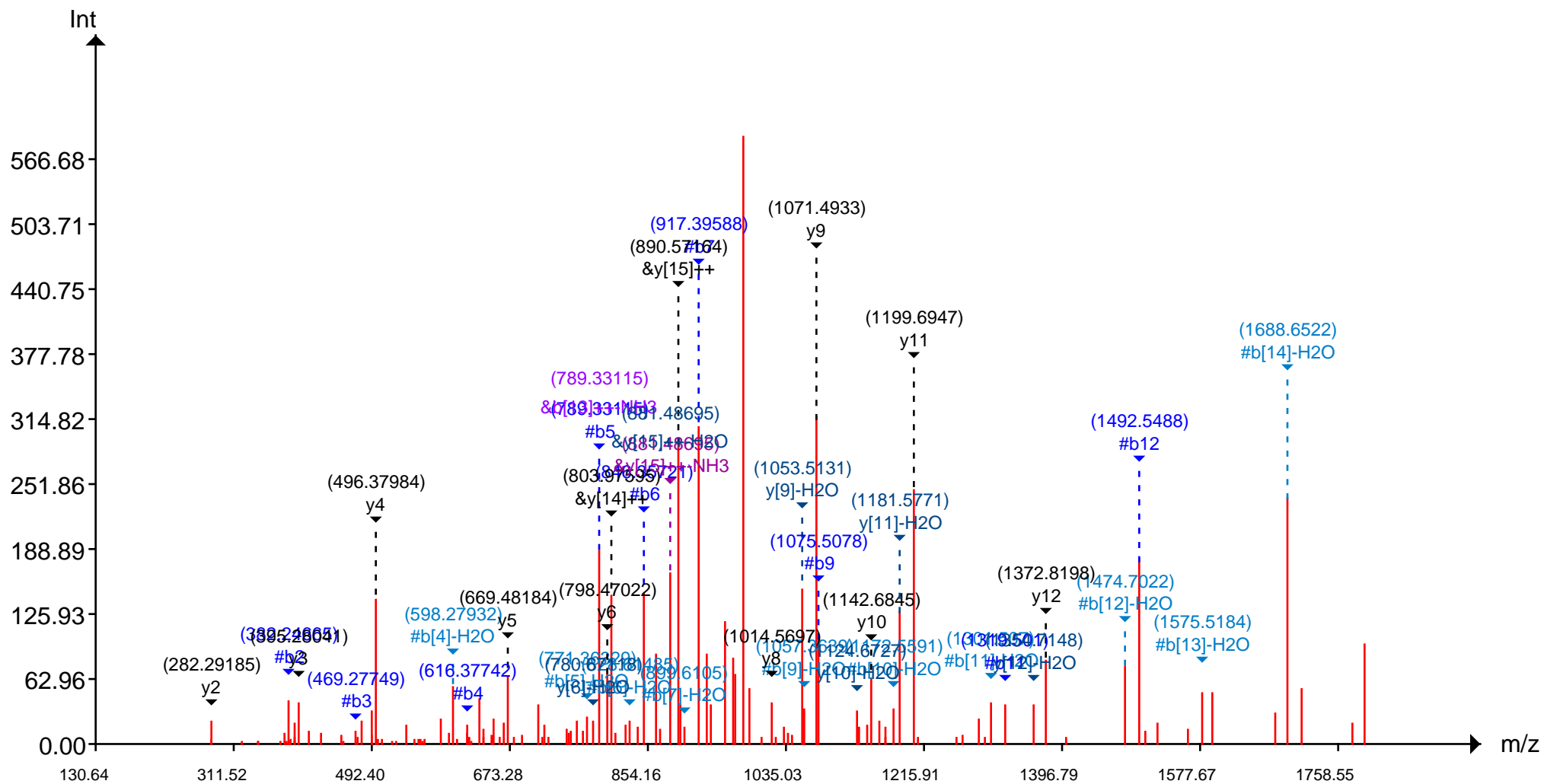


Info

Modified Sequence	AcD3-VAEGNK<AcD3K*>EDVDK<AcD3K*>AVK<AcD3K*>AAR<C13>-COOH	Ion score	47.28
Protein	P47738	Confidence level	99.00%
Precursor m/z	993.55433	Identity Threshold	46.51
Charge	2+	Mass error	0.002677

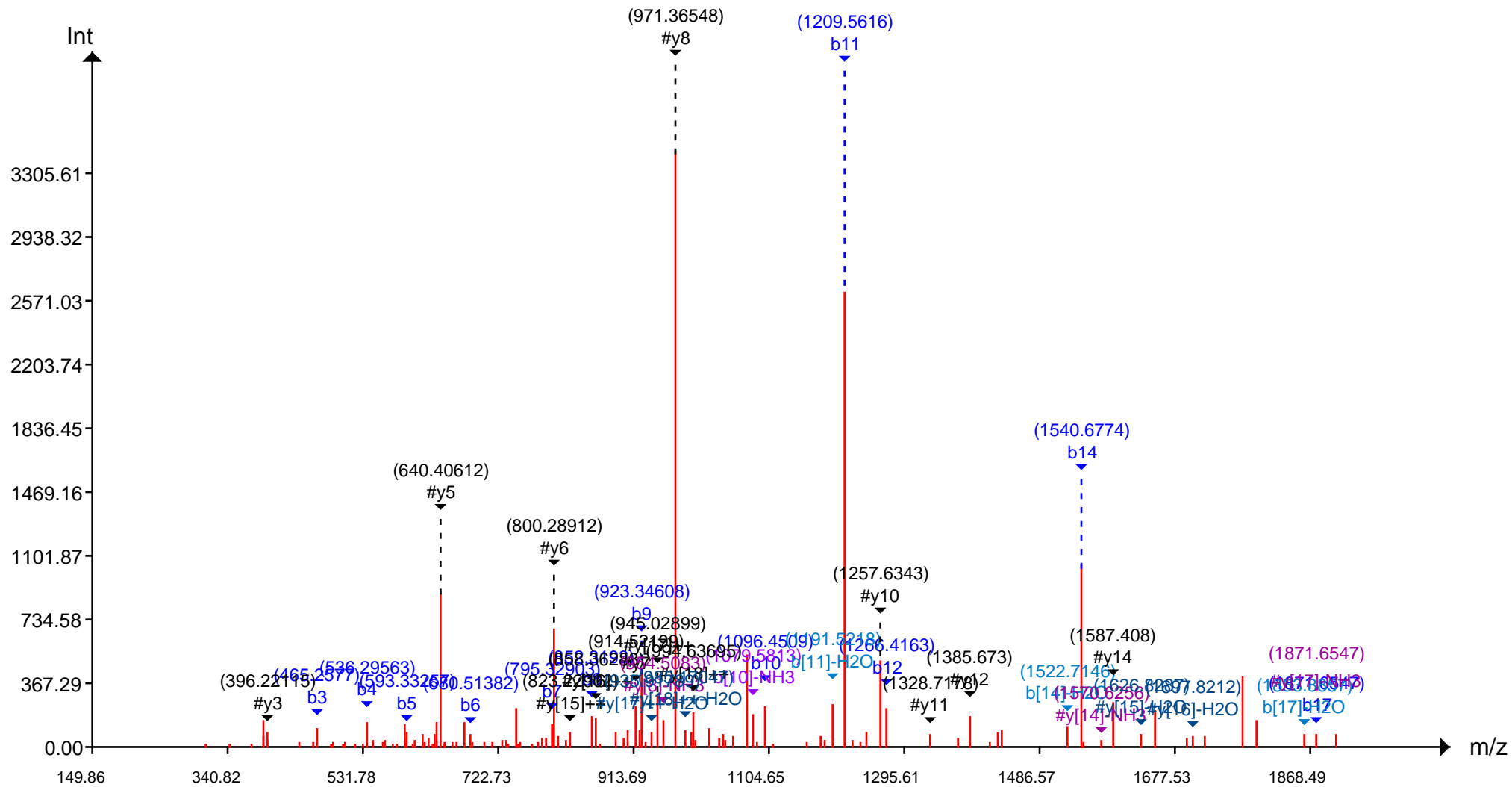


Info			
Modified Sequence	AcD3-VIDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH	Ion score	92.63
Protein	P63260	Confidence level	99.00%
Precursor m/z	1075.4977	Identity Threshold	45.01
Charge	2+	Mass error	0.002577



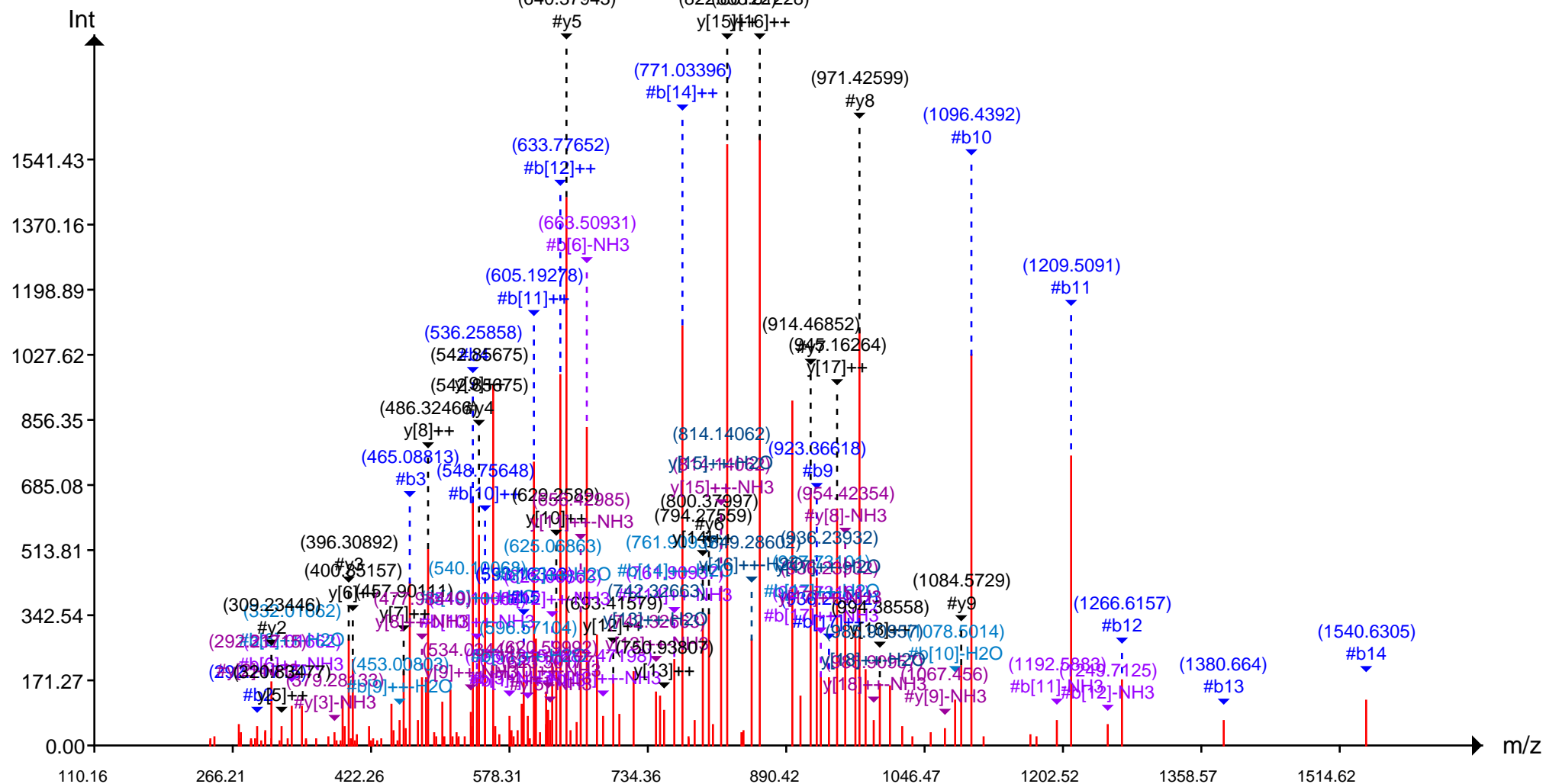
Info

Modified Sequence	AcD3-YK<AcD3K*>SM<Mox*>K<AcD3K*>GAGTDEK<AcD3K*>TLTR<C13>-COOH	Ion score	56.56
Protein	P14824	Confidence level	99.00%
Precursor m/z	994.52863	Identity Threshold	47.93
Charge	2+	Mass error	9.9E-4



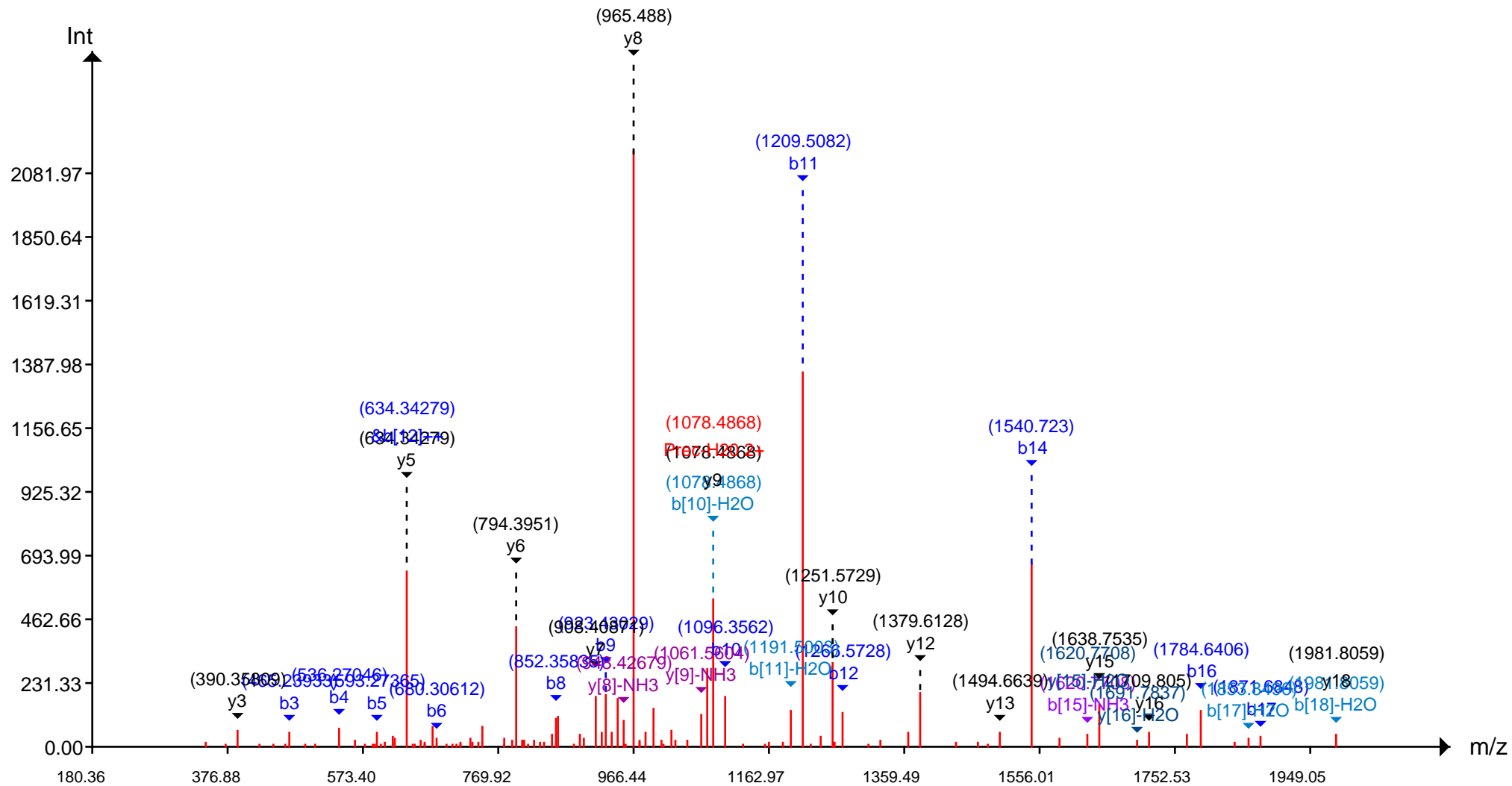
Info

Modified Sequence	AcD3-FVK<AcD3K*>AGSDGAK<AcD3K*>IGNC<Cmm*>PFSQR<C13>-COOH	Ion score	72.65
Protein	Q9Z1Q5	Confidence level	99.00%
Precursor m/z	1090.5653	Identity Threshold	48.07
Charge	2+	Mass error	0.002263



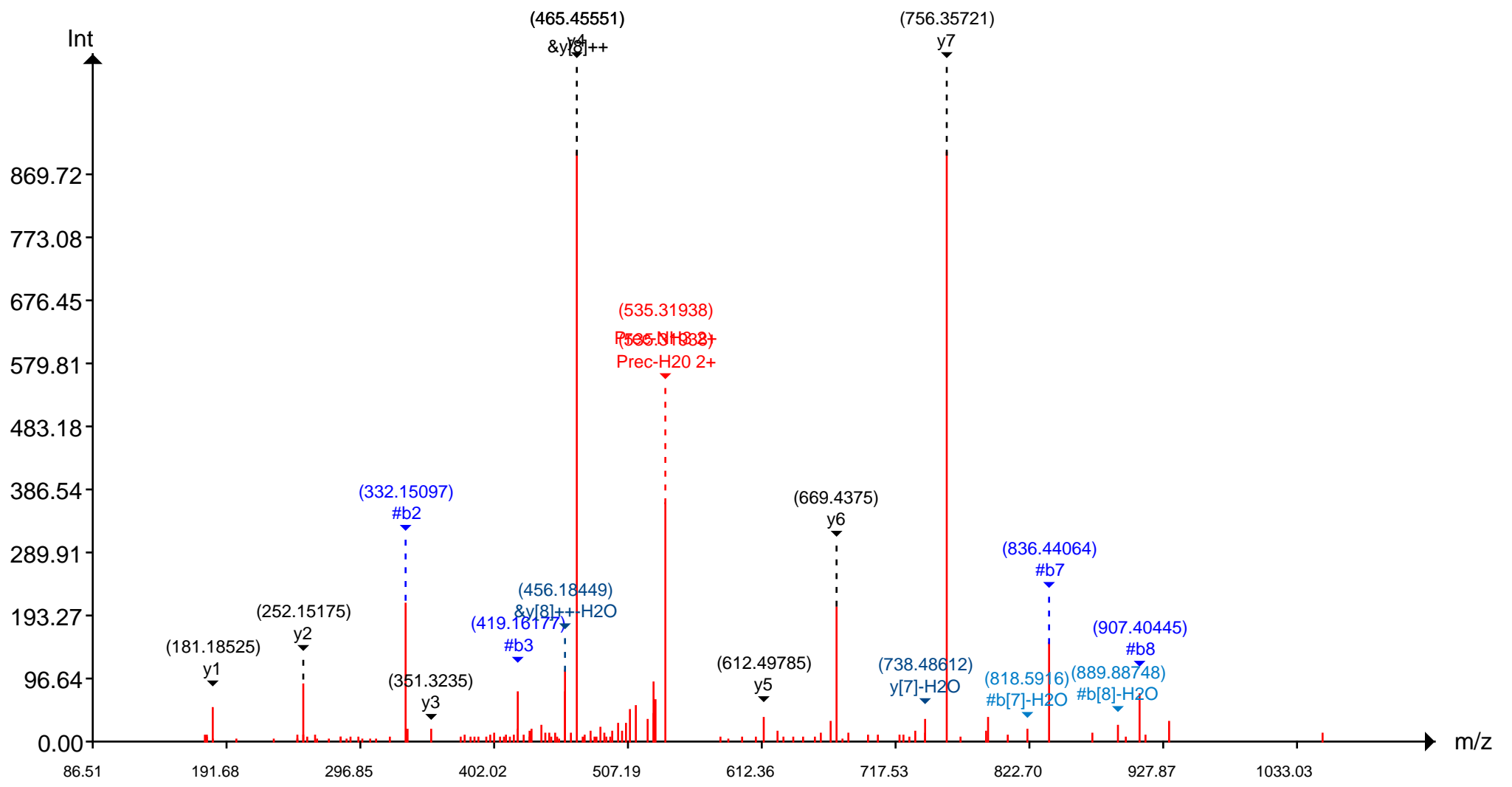
Info

Modified Sequence	AcD3-FVK<AcD3K*>AGSDGAK<AcD3K*>IGNC<Cmm*>PFSQR<C13>-COOH	Ion score	81.68
Protein	Q9Z1Q5	Confidence level	99.00%
Precursor m/z	727.37893	Identity Threshold	48.03
Charge	3+	Mass error	0.001177



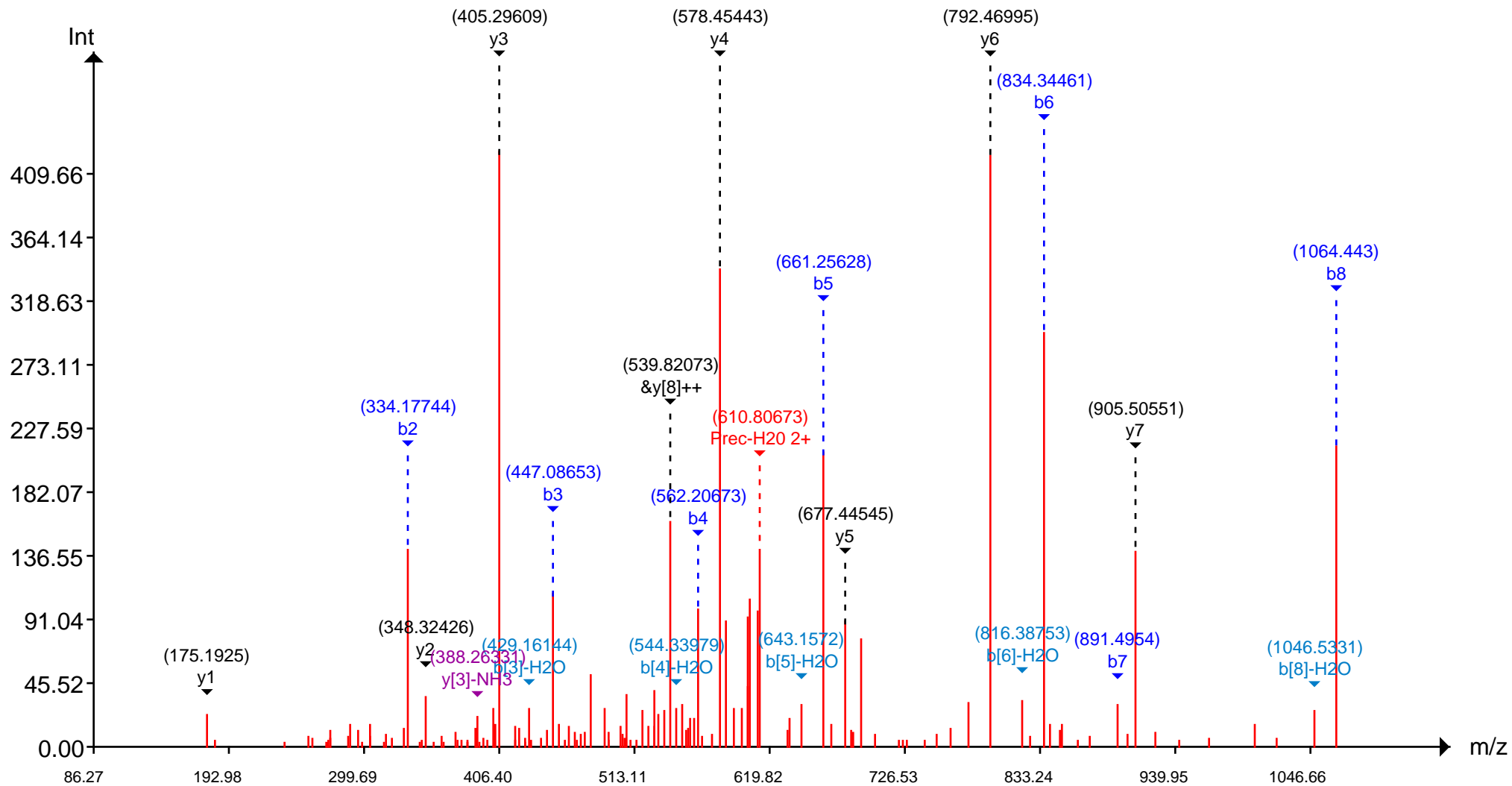
Info

Modified Sequence	AcD3-FVK<AcD3K*>AGSDGAK<AcD3K*>IGNC<Cmm*>PFSQR-COOH	Ion score	116.66
Protein	Q9Z1Q5	Confidence level	99.00%
Precursor m/z	1087.5556	Identity Threshold	47.90
Charge	2+	Mass error	0.003005



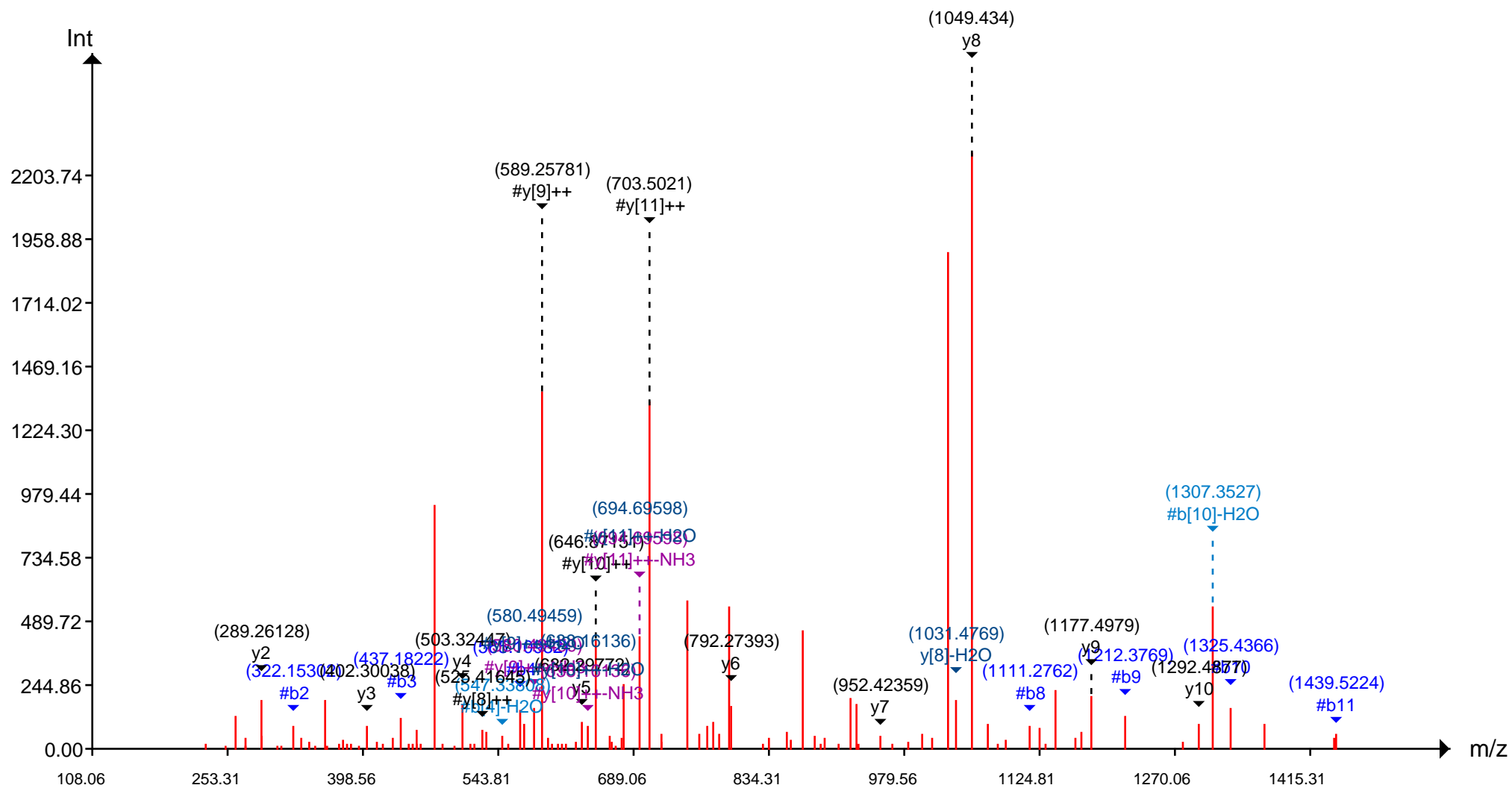
Info

Modified Sequence	AcD3-IK<AcD3K*>SGM<Mox*>NVAR<C13>-COOH	Ion score	60.13
Protein	P52480	Confidence level	99.00%
Precursor m/z	544.31121	Identity Threshold	47.07
Charge	2+	Mass error	9.04E-4



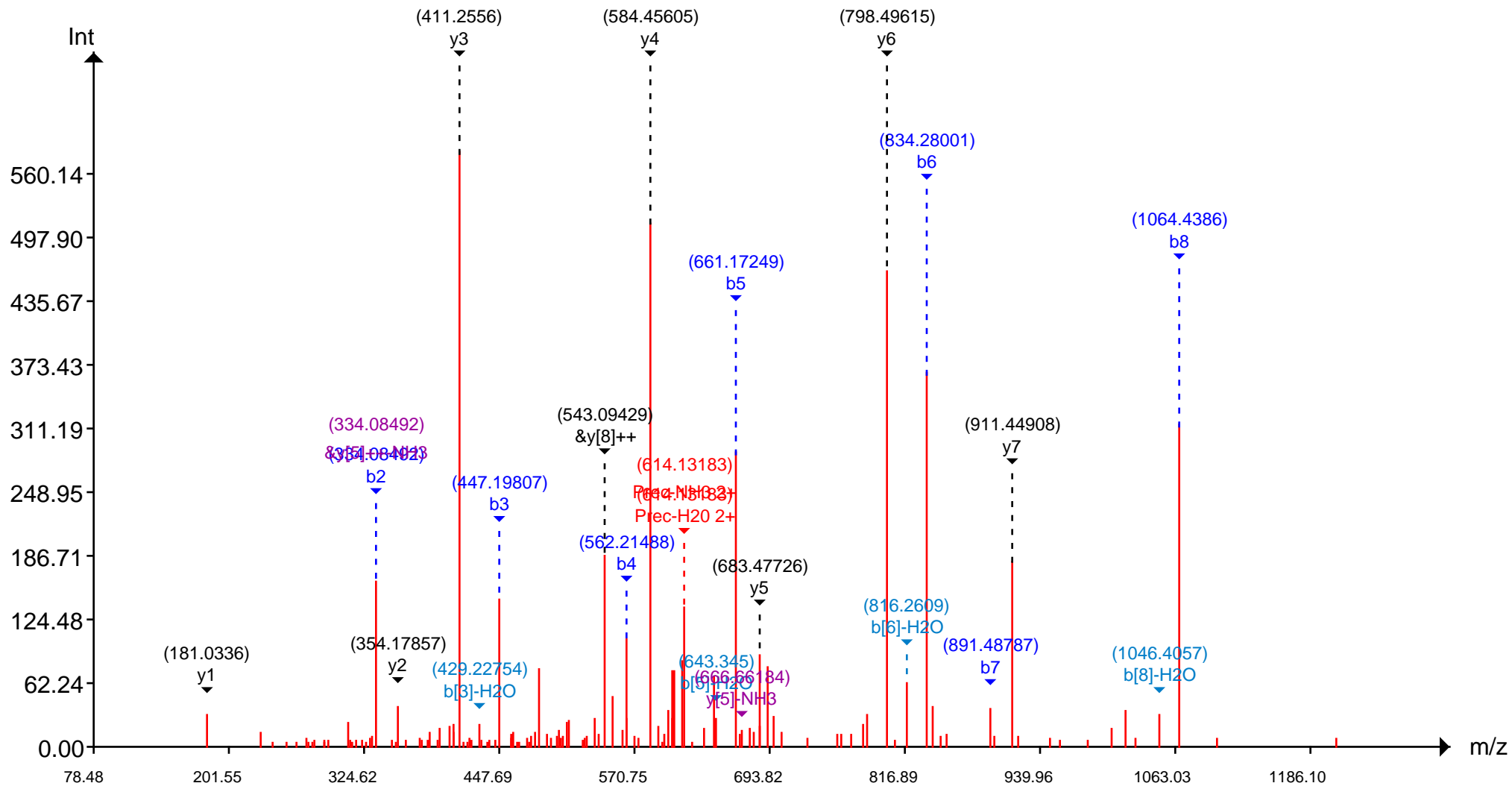
Info

Modified Sequence	AcD3-DK<AcD3K*>LDVK<AcD3K*>GK<AcD3K*>R-COOH	Ion score	68.24
Protein	P09411	Confidence level	99.00%
Precursor m/z	619.87894	Identity Threshold	43.15
Charge	2+	Mass error	0.001354



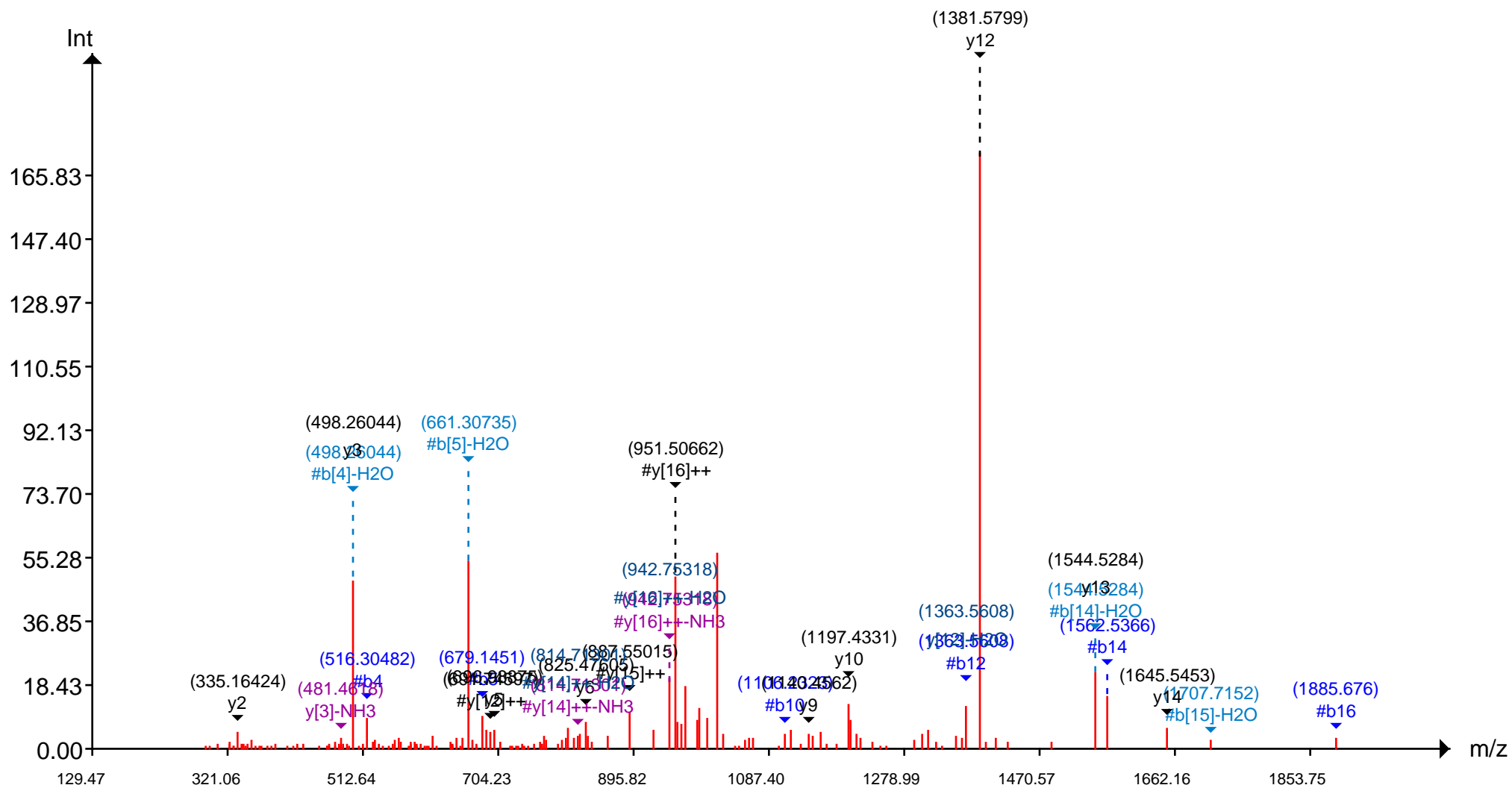
Info

Modified Sequence	AcD3-YLDQPC<Cmm*>C<Cmm*>ETINR-COOH	Ion score	48.69
Protein	Q61598	Confidence level	99.00%
Precursor m/z	807.36085	Identity Threshold	43.52
Charge	2+	Mass error	0.001856



Info

Modified Sequence	AcD3-DK<AcD3K*>LDVK<AcD3K*>GK<AcD3K*>R<C13>-COOH	Ion score	55.62
Protein	P09411	Confidence level	99.00%
Precursor m/z	622.8887	Identity Threshold	42.35
Charge	2+	Mass error	7.33E-4



Info

Modified Sequence AcD3-IQQTYPGGEEQAQYC<Cmm*>R-COOH

Ion score 53.93

Protein Q9WU78

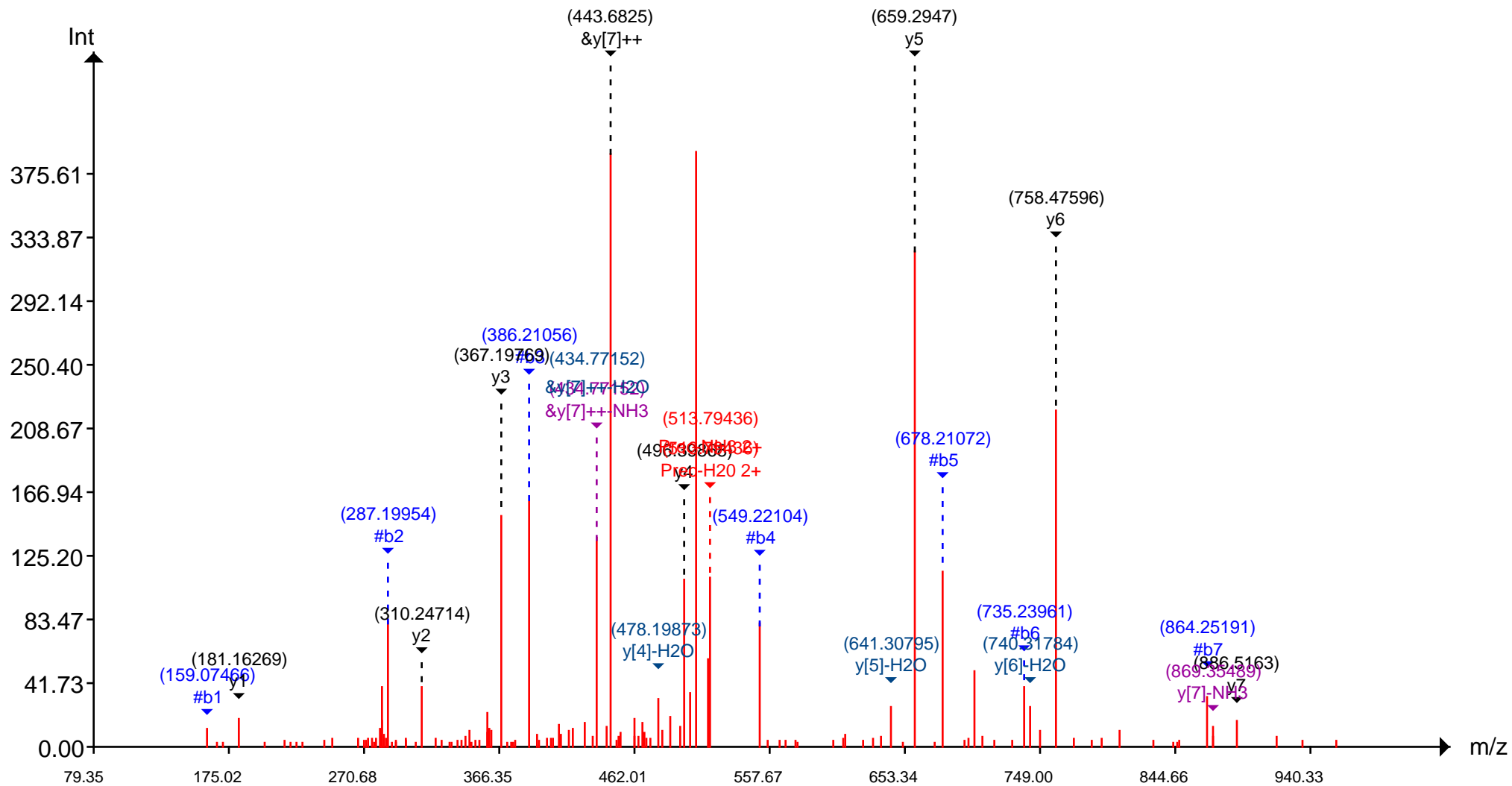
Confidence level 99.00%

Precursor m/z 1030.4642

Identity Threshold 43.56

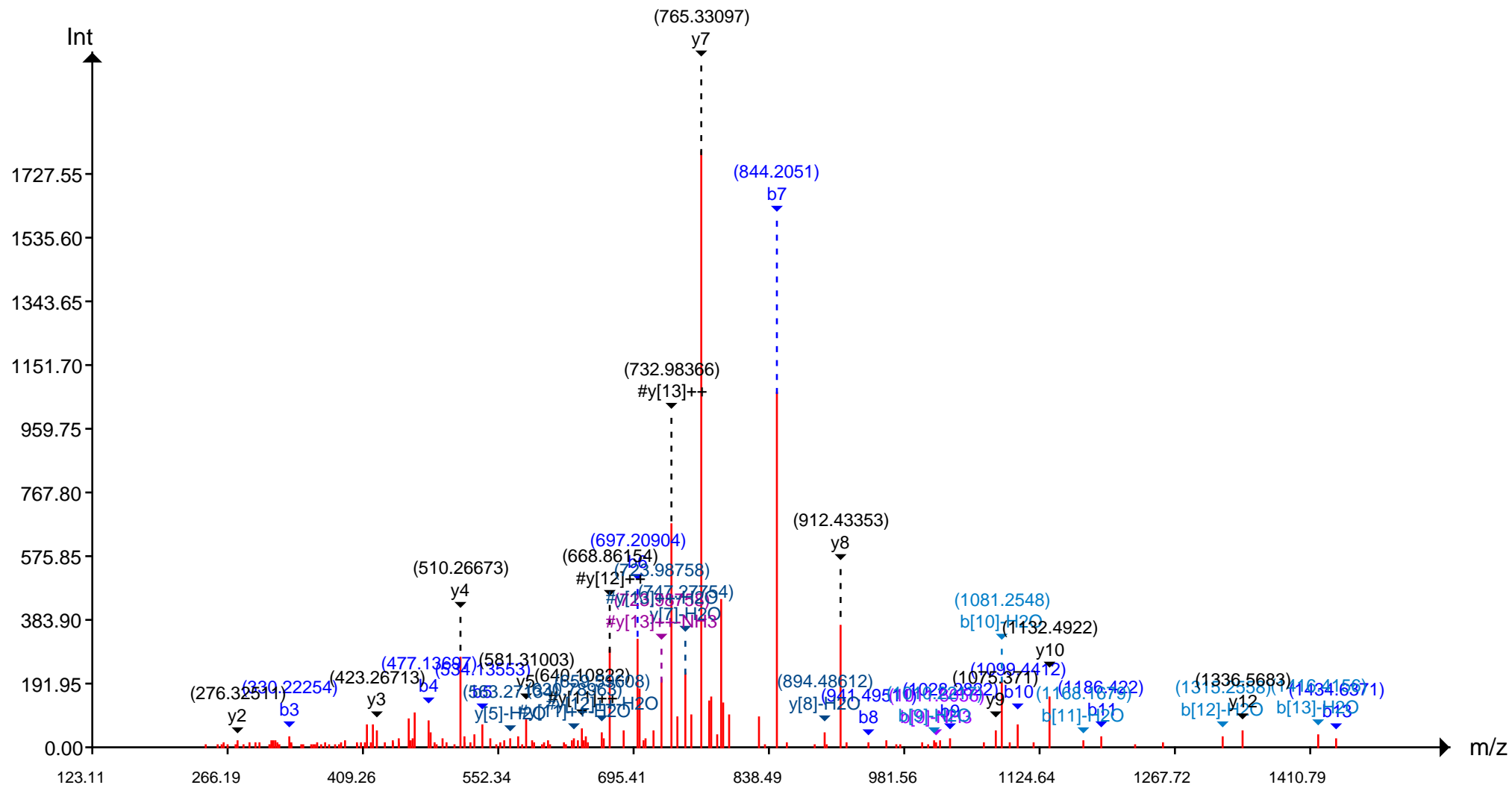
Charge 2+

Mass error -5.81E-4



Info

Modified Sequence	AcD3-IQVYEGER<C13>-COOH	Ion score	52.88
Protein	Q61696, P17879, P16627, P63017	Confidence level	99.00%
Precursor m/z	522.77847	Identity Threshold	47.89
Charge	2+	Mass error	1.67E-4



Info

Modified Sequence AcD3-VQGM<Mox*>GYM<Mox*>PSASM<Mox*>TR-COOH

Ion score 75.29

Protein Q62433

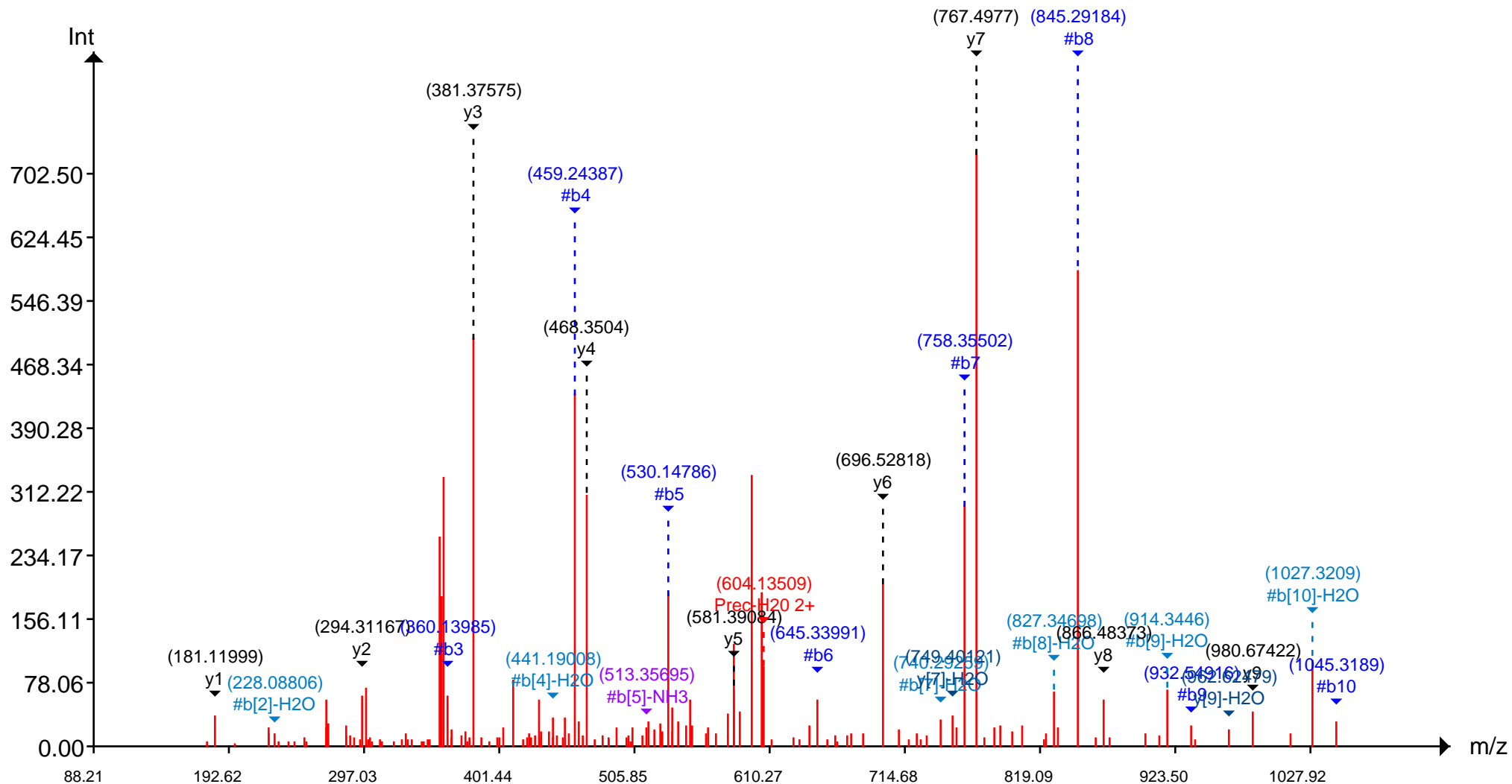
Confidence level 99.00%

Precursor m/z 804.84957

Identity Threshold 41.84

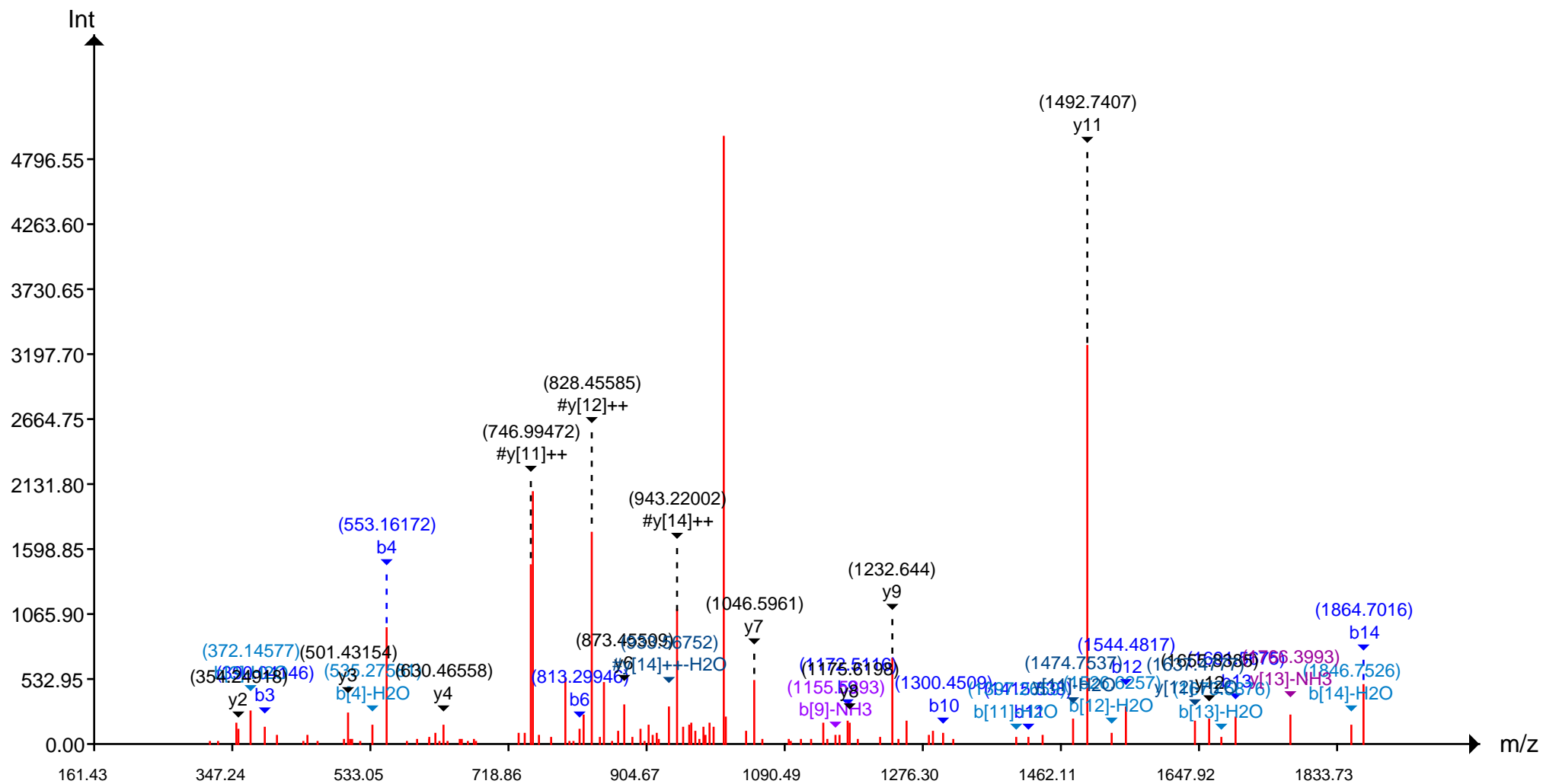
Charge 2+

Mass error 0.002459



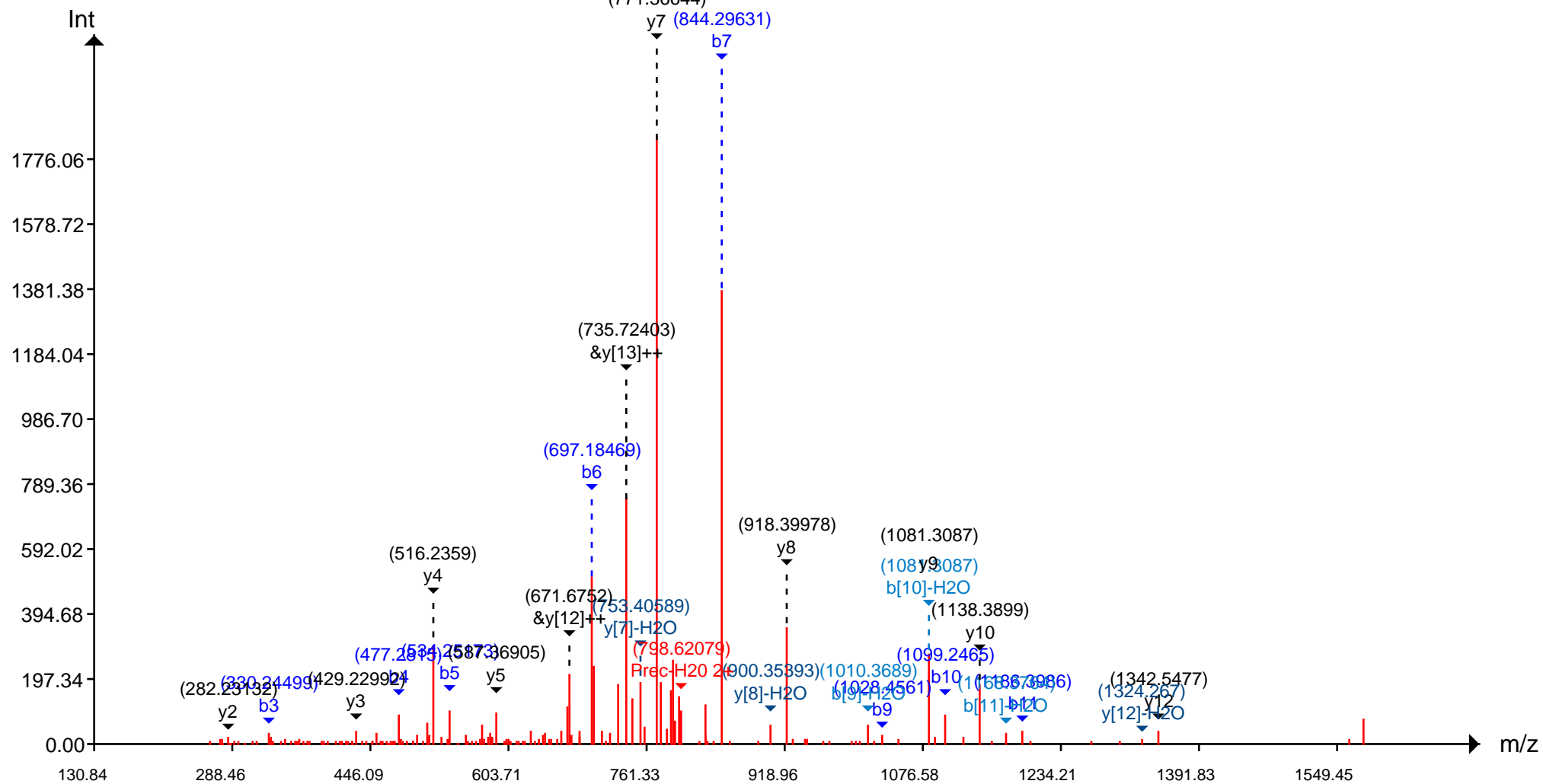
Info

Modified Sequence	AcD3-SLNVADLSSLR<C13>-COOH	Ion score	57.47
Protein	P97372	Confidence level	99.00%
Precursor m/z	613.35003	Identity Threshold	46.75
Charge	2+	Mass error	6.48E-4



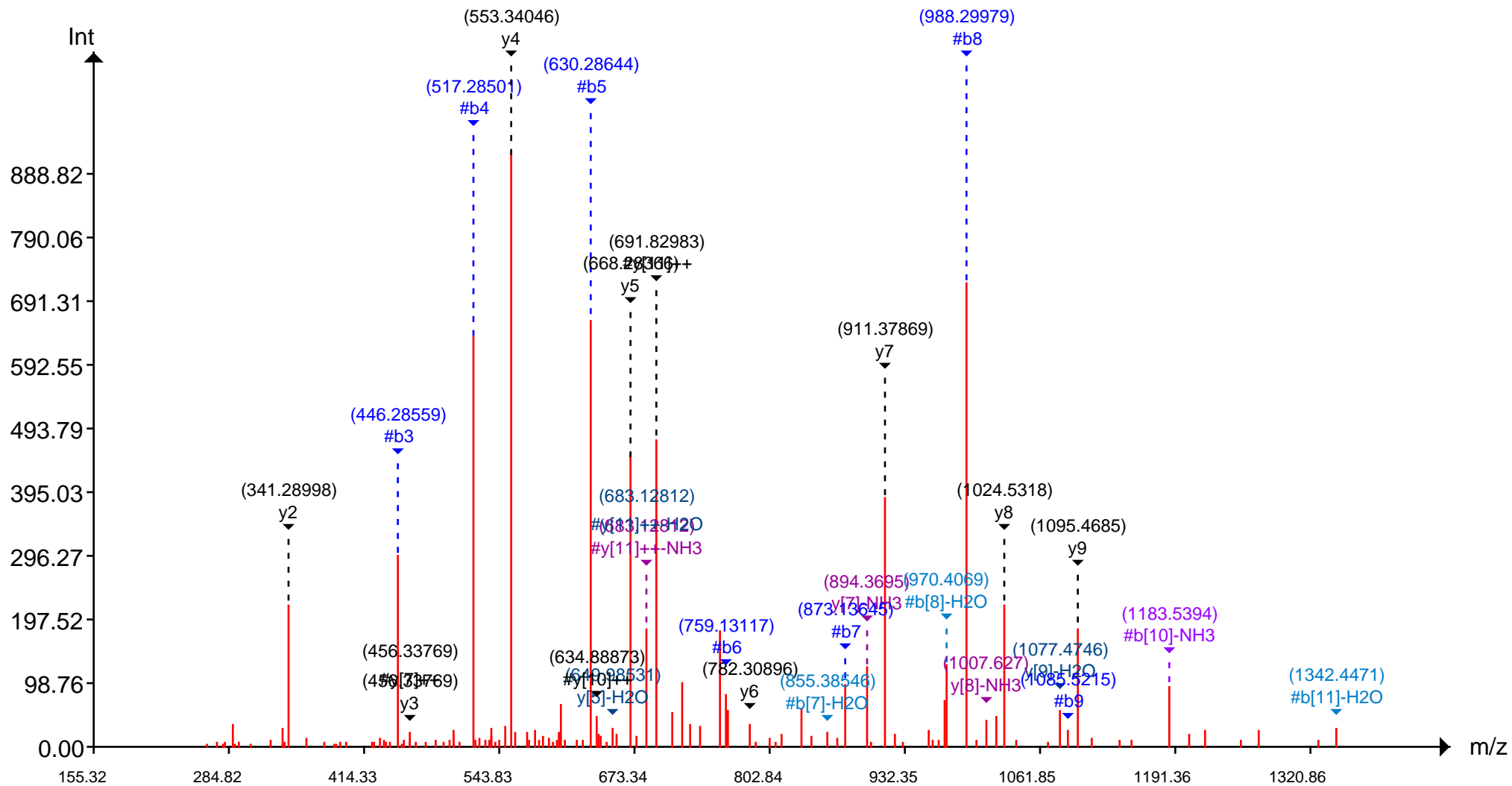
Info

Modified Sequence	AcD3-DTQYPYGEK<AcD3K*>QDEFK<AcD3K*>R<C13>-COOH	Ion score	57.52
Protein	P57759	Confidence level	99.00%
Precursor m/z	1023.0015	Identity Threshold	47.34
Charge	2+	Mass error	0.005324



Info

Modified Sequence	AcD3-VQGM<Mox*>GYM<Mox*>PSASM<Mox*>TR<C13>-COOH	Ion score	62.49
Protein	Q62433	Confidence level	99.00%
Precursor m/z	807.85917	Identity Threshold	43.16
Charge	2+	Mass error	0.001517



Info

Modified Sequence AcD3-INK<AcD3K*>ALENDPDC<Cmm*>R<C13>-COOH

Ion score 66.18

Protein Q61233, Q99K51

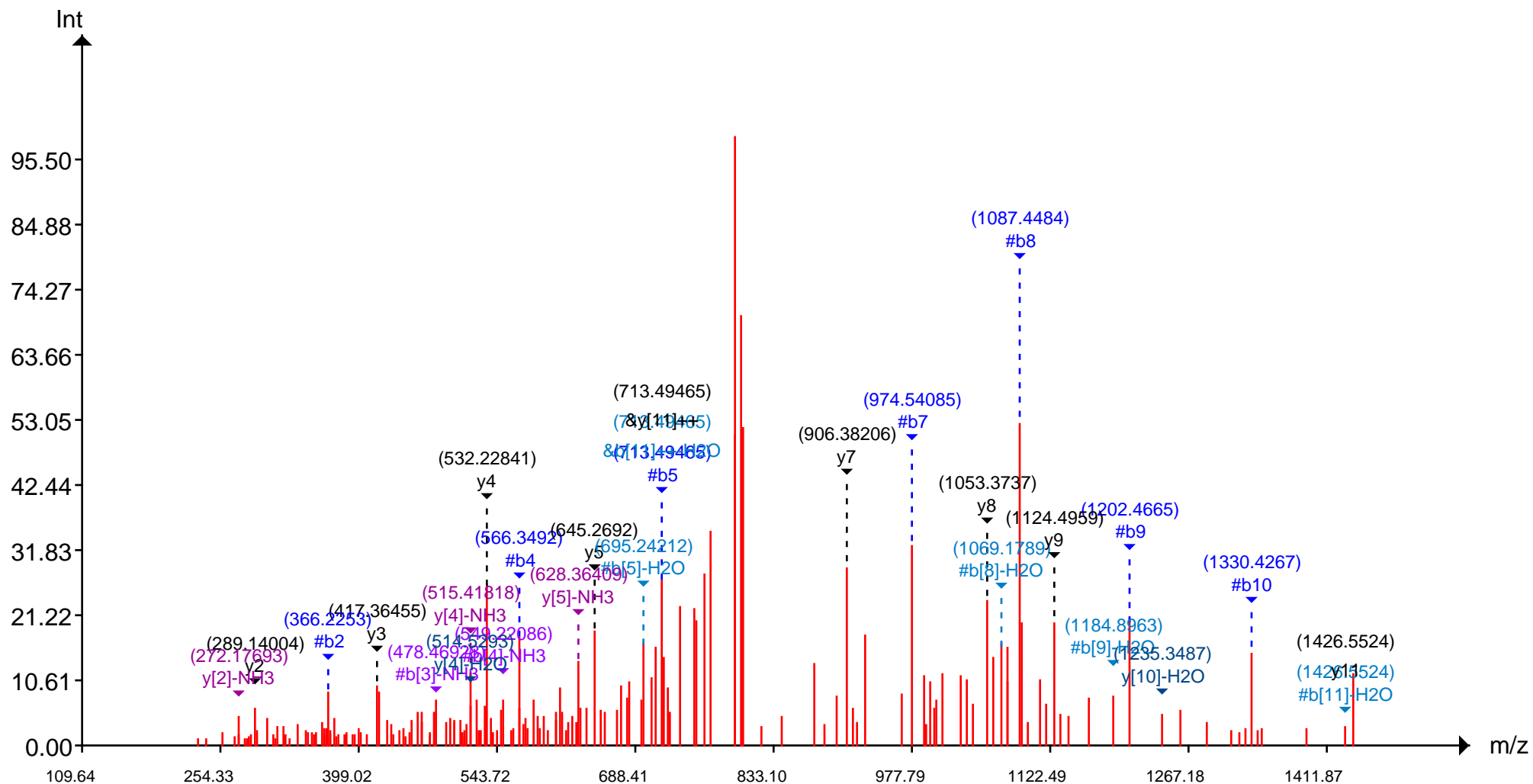
Confidence level 99.00%

Precursor m/z 770.88617

Identity Threshold 47.68

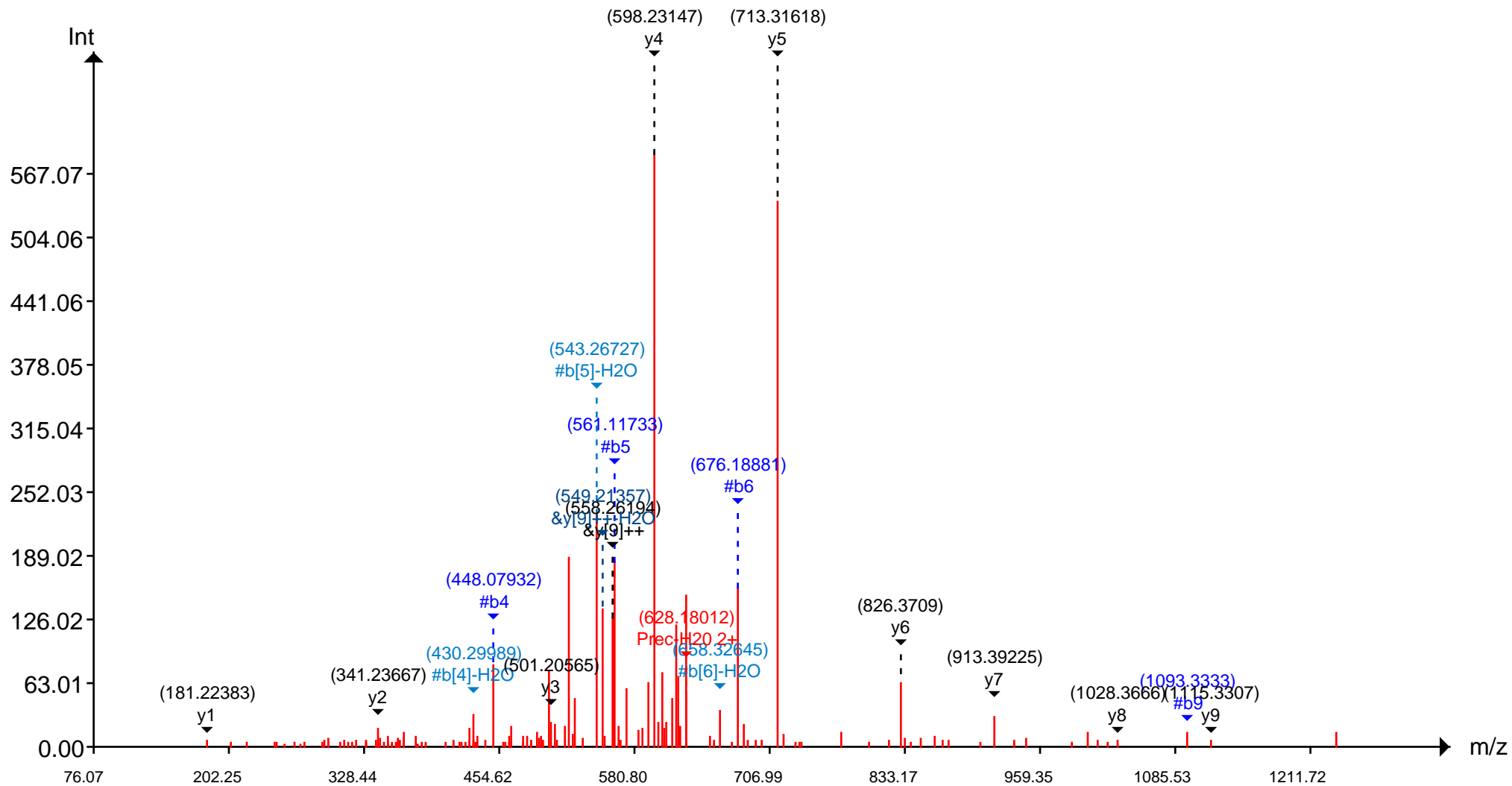
Charge 2+

Mass error 0.001257



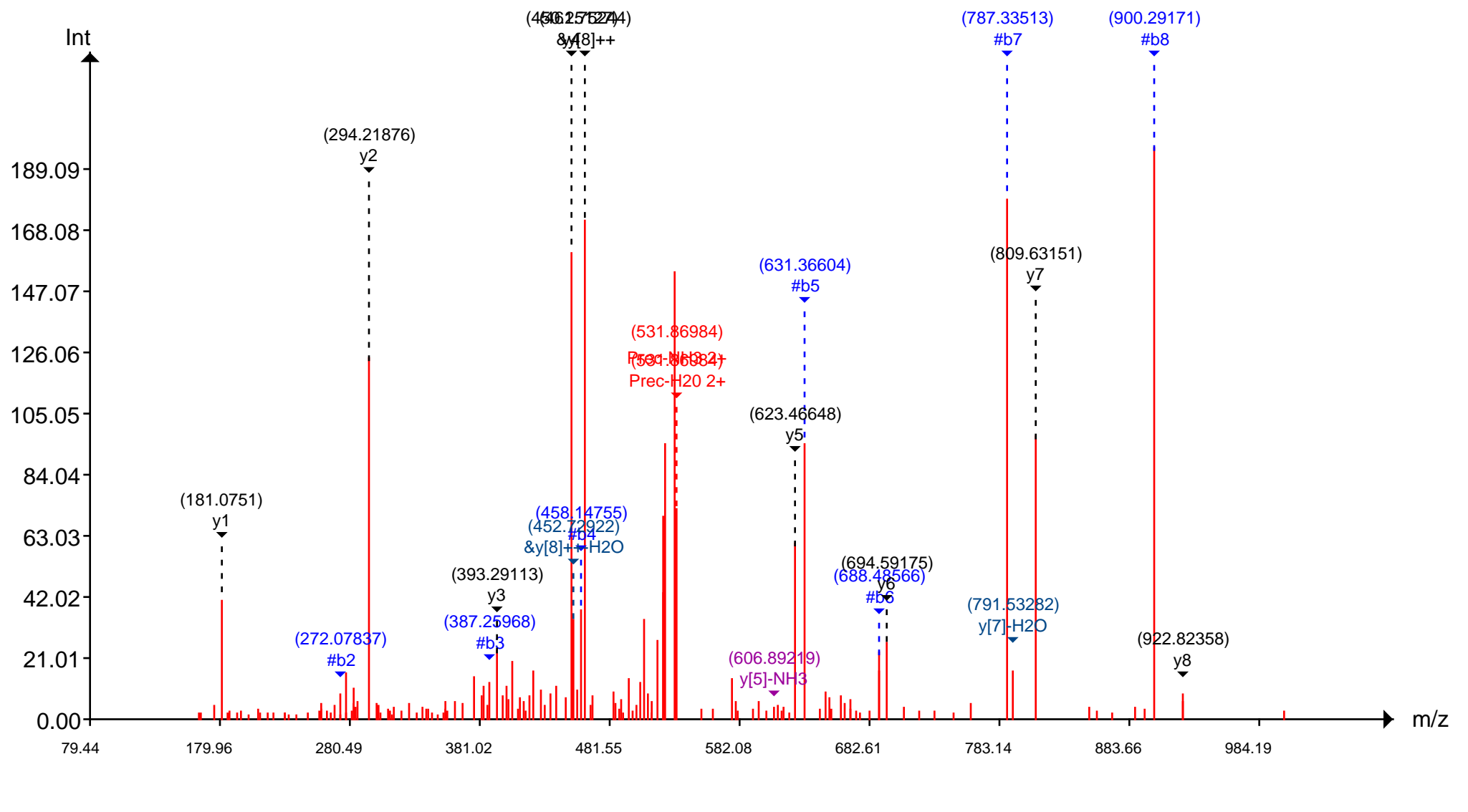
Info

Modified Sequence	AcD3-FK<AcD3K*>EAFNM<Mox*>IDQNR-COOH	Ion score	48.22
Protein	Q3THE2, Q9CQ19	Confidence level	99.00%
Precursor m/z	809.89459	Identity Threshold	46.99
Charge	2+	Mass error	0.001847

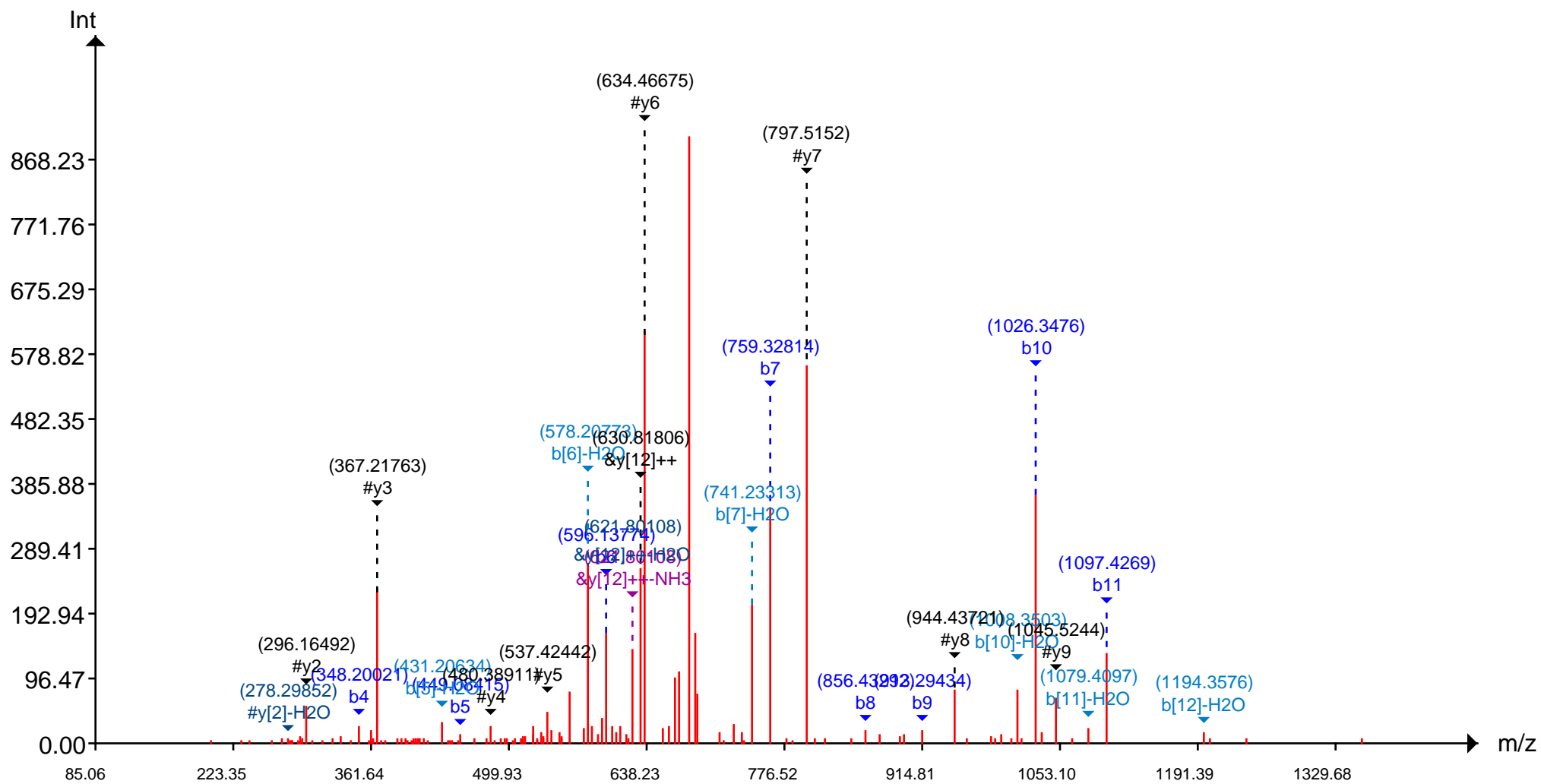


Info

Modified Sequence	AcD3-ISDSLDPC<Cmm*>C<Cmm*>R<C13>-COOH	Ion score	59.73
Protein	Q61753	Confidence level	99.00%
Precursor m/z	637.28832	Identity Threshold	43.98
Charge	2+	Mass error	7.48E-4

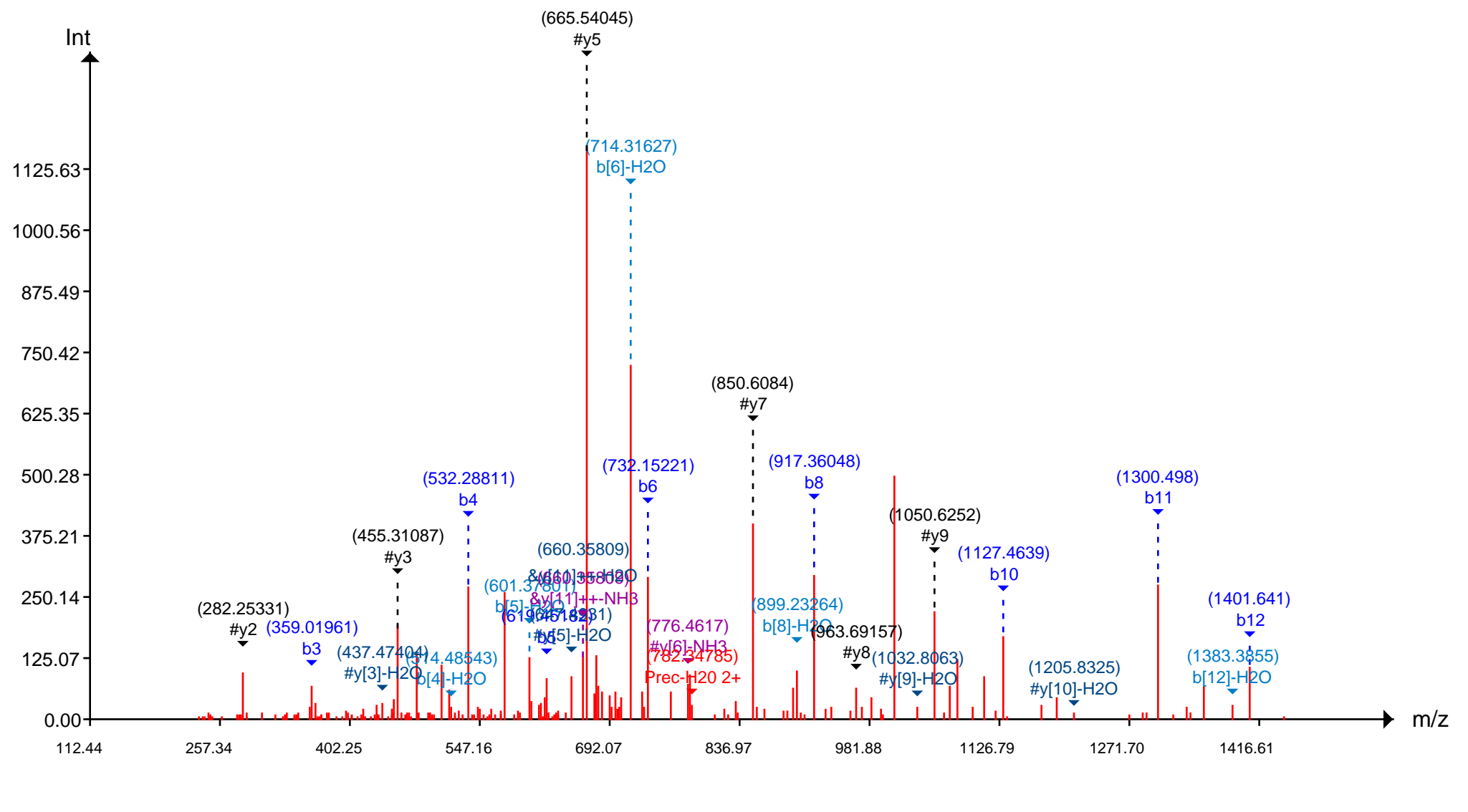


Info			
Modified Sequence	AcD3-IIDAK<AcD3K*>GVLR<C13>-COOH	Ion score	59.24
Protein	Q61171	Confidence level	99.00%
Precursor m/z	540.85249	Identity Threshold	39.19
Charge	2+	Mass error	-0.001253

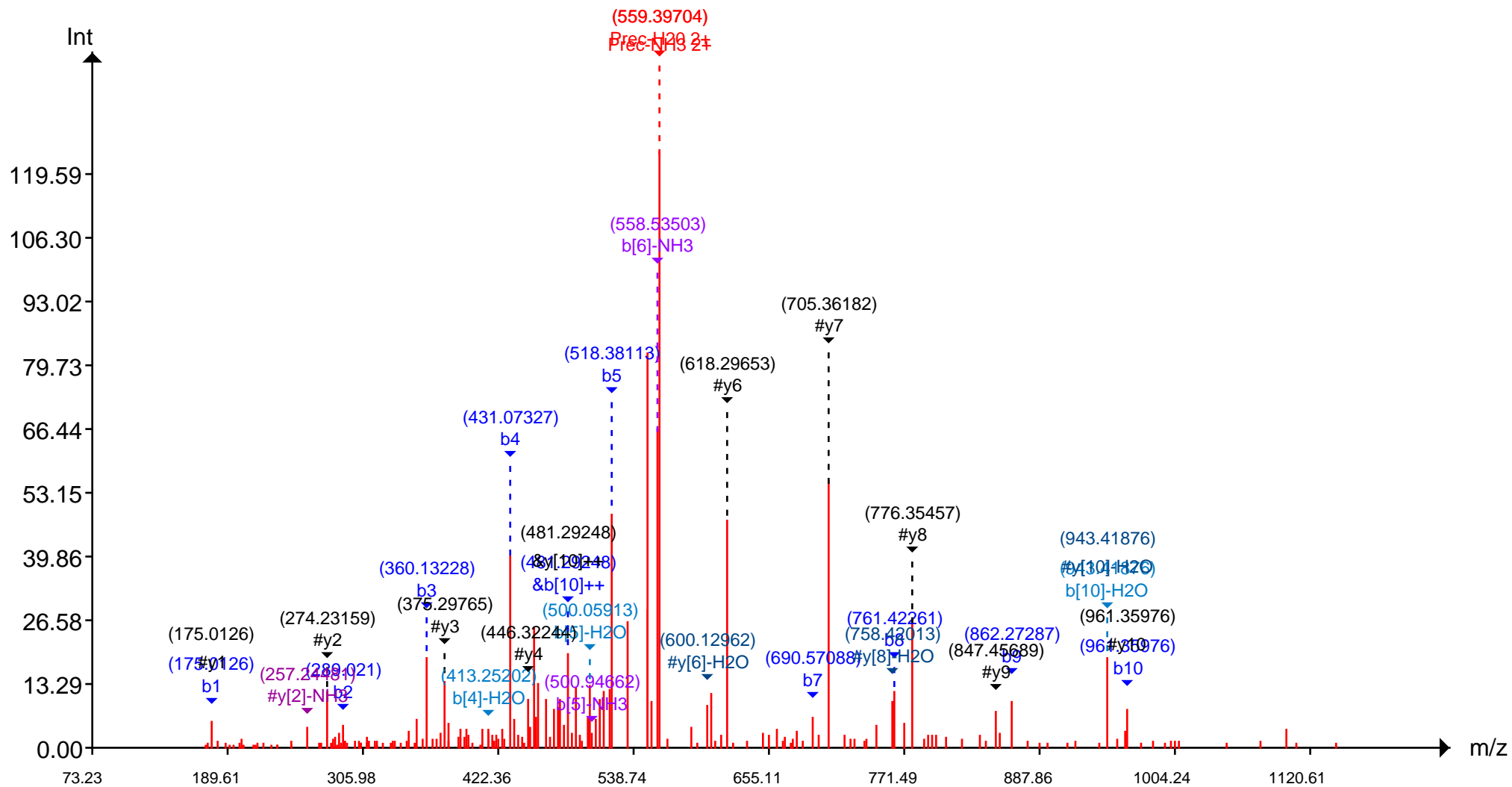


Info

Modified Sequence	AcD3-SGGTTM<Mox*>YPGIADR<C13>-COOH	Ion score	58.84
Protein	P62737, P60710, P68033, P63260, P63268, P68134	Confidence level	99.00%
Precursor m/z	696.83363	Identity Threshold	46.52
Charge	2+	Mass error	1.05E-4

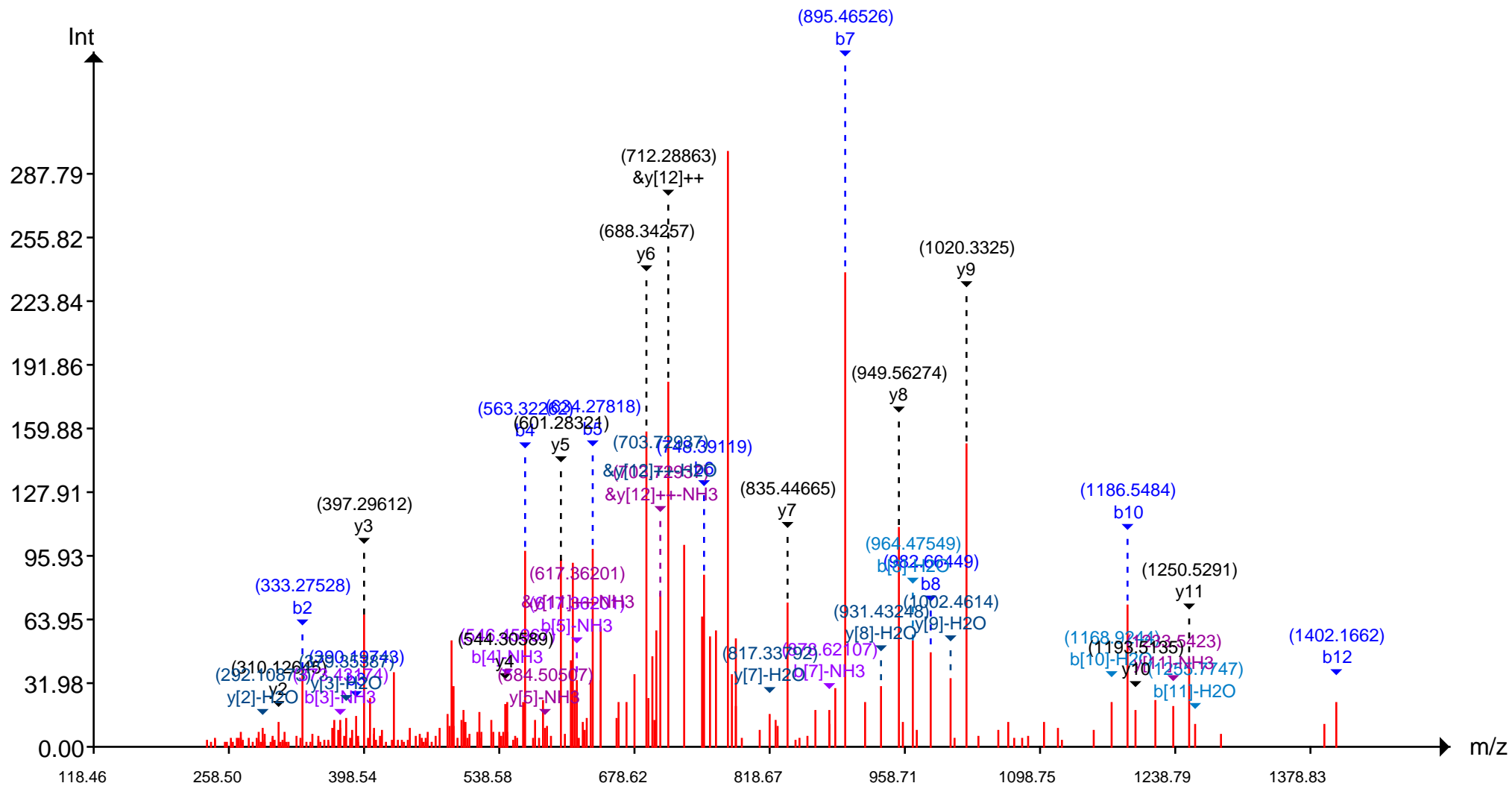


Info			
Modified Sequence	AcD3-AELK<AcD3K*>SLGQPIK<AcD3K*>TR<C13>-COOH	Ion score	44.74
Protein	Q61316	Confidence level	99.00%
Precursor m/z	791.48527	Identity Threshold	41.45
Charge	2+	Mass error	0.001673



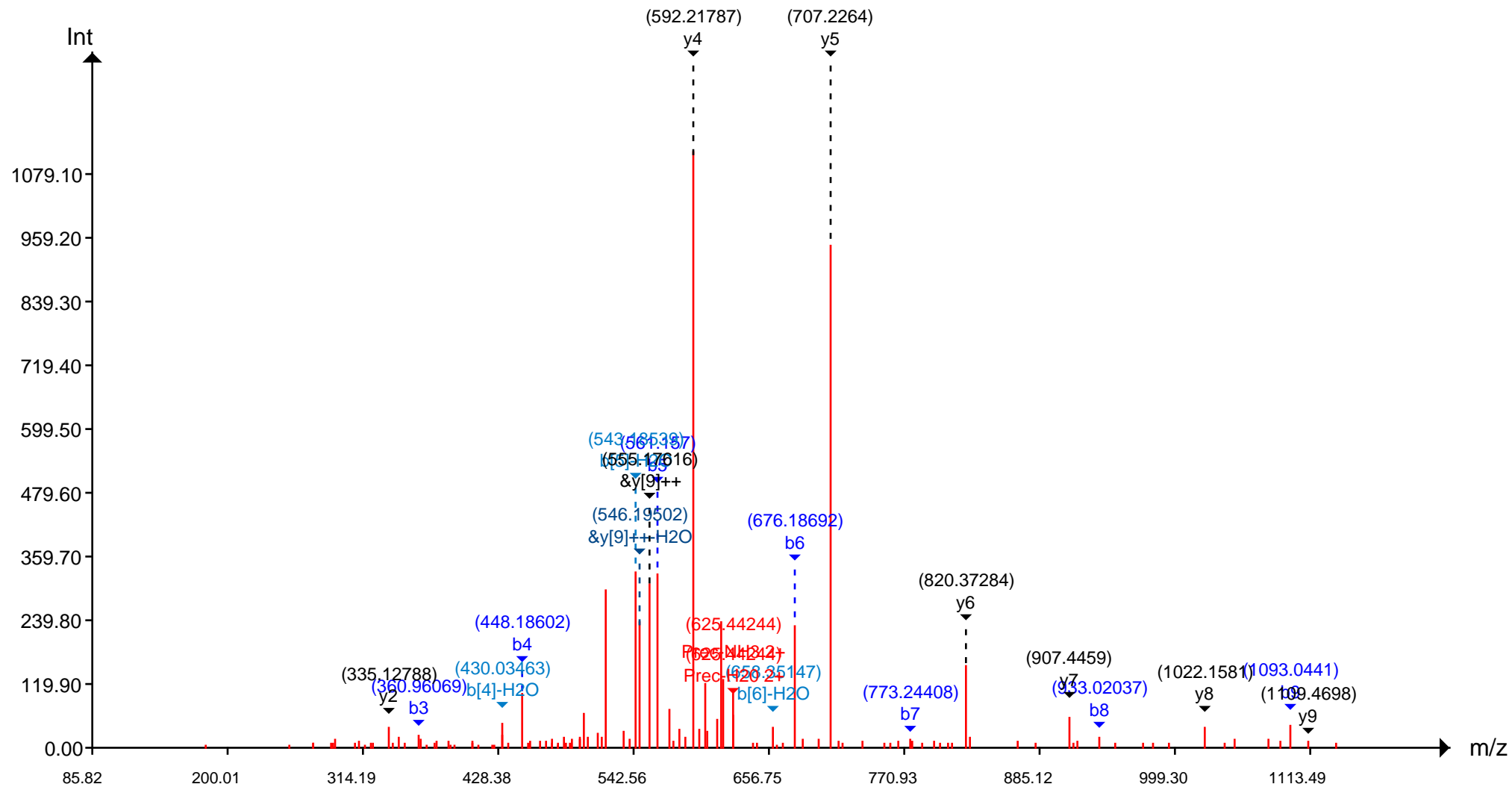
Info

Modified Sequence	AcD3-ENAASGDATVR-COOH	Ion score	63.03
Protein	Q9CSU0	Confidence level	99.00%
Precursor m/z	568.2748	Identity Threshold	46.56
Charge	2+	Mass error	6.09E-4



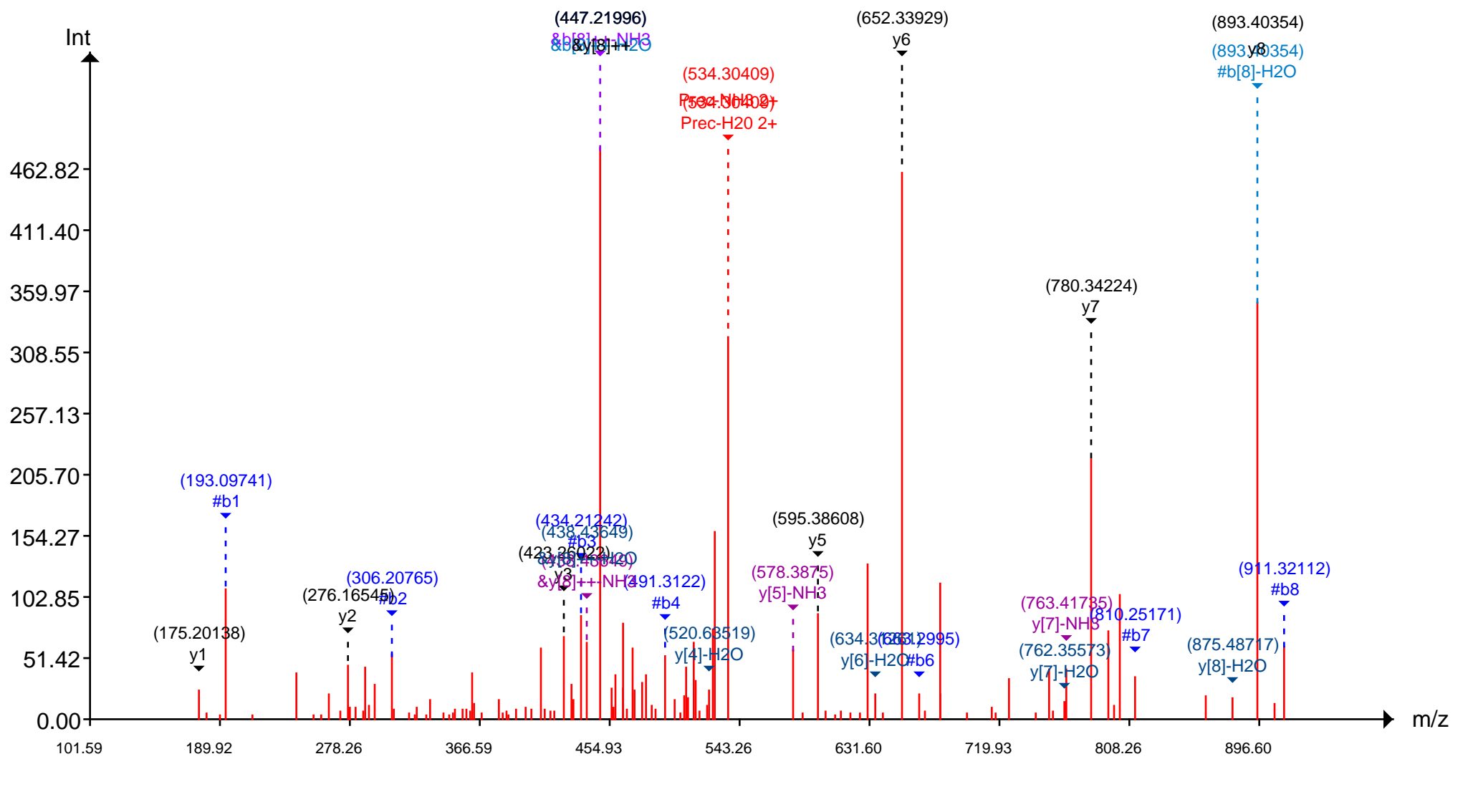
Info

Modified Sequence	AcD3-NK<AcD3K*>GK<AcD3K*>ANFSGM<Mox*>SER<C13>-COOH	Ion score	54.2
Protein	P12388	Confidence level	99.00%
Precursor m/z	791.90095	Identity Threshold	47.80
Charge	2+	Mass error	0.001078

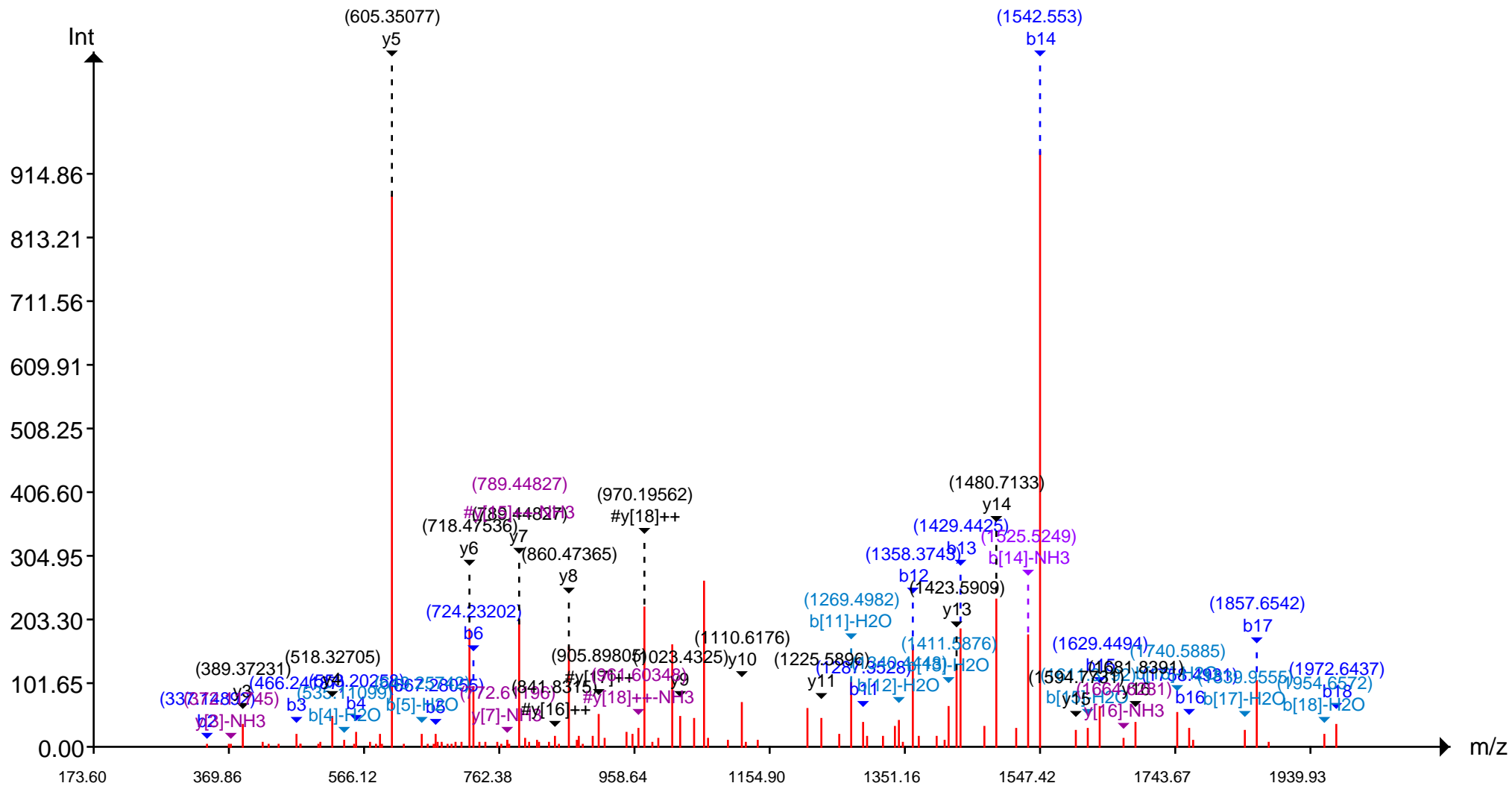


Info

Modified Sequence	AcD3-ISDSLDPCC<Cmm*>C<Cmm*>R-COOH	Ion score	66.45
Protein	Q61753	Confidence level	99.00%
Precursor m/z	634.27807	Identity Threshold	42.83
Charge	2+	Mass error	3.89E-4

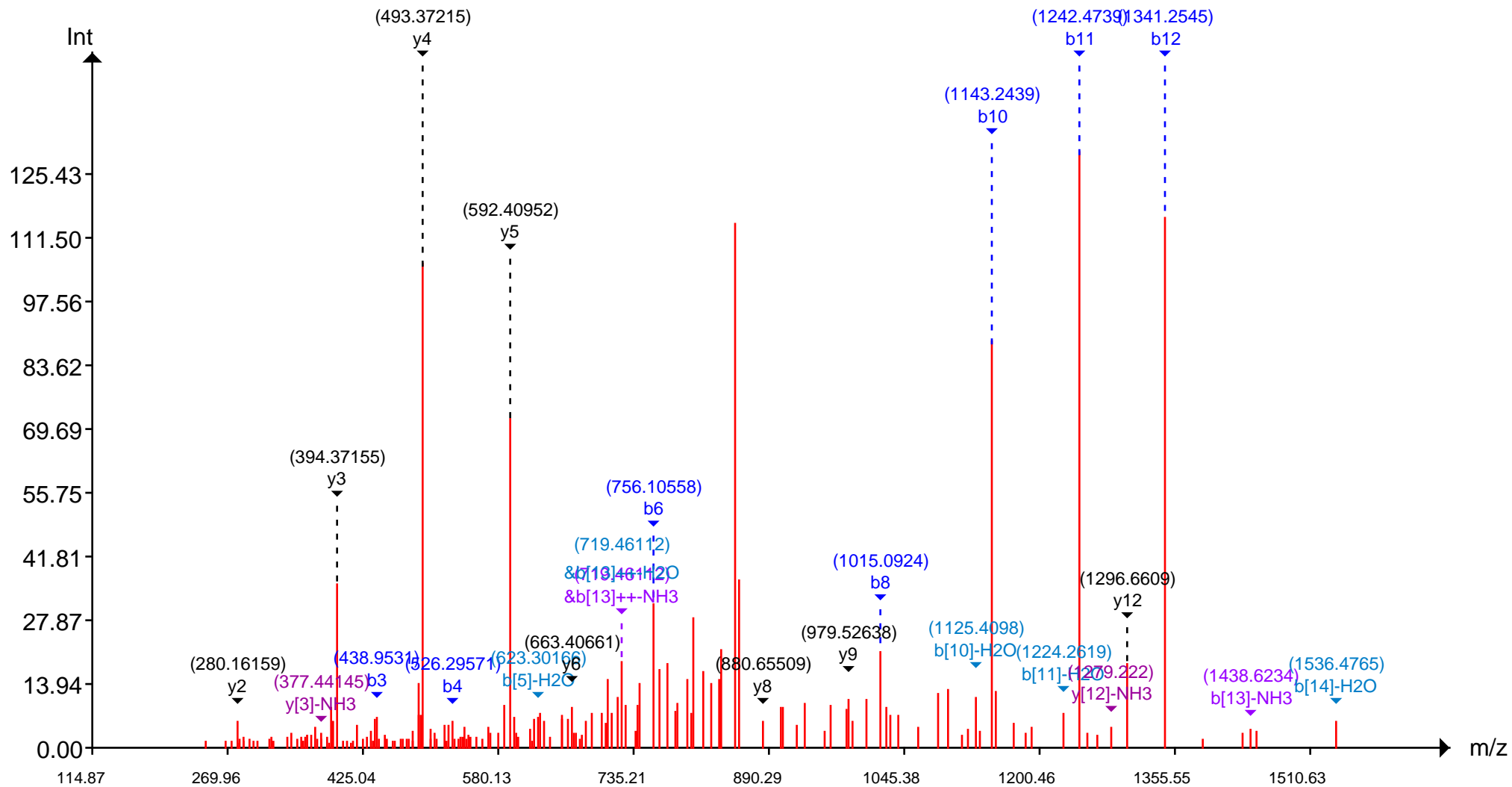


Info		Ion score	46.35
Modified Sequence	AcD3-M<Mox*>IQGGDFTR-COOH	Confidence level	99.00%
Protein	P24369	Identity Threshold	45.97
Precursor m/z	543.25941	Mass error	-8.13E-4
Charge	2+		



Info

Modified Sequence	AcD3-YQESNGPTDSYAAISEVDR-COOH	Ion score	120.61
Protein	O08585	Confidence level	99.00%
Precursor m/z	1073.9857	Identity Threshold	43.82
Charge	2+	Mass error	0.003876



Info

Modified Sequence AcD3-DDYSDDVC<Cmm*>GAVVNVR<C13>-COOH

Ion score 49.65

Protein P63073

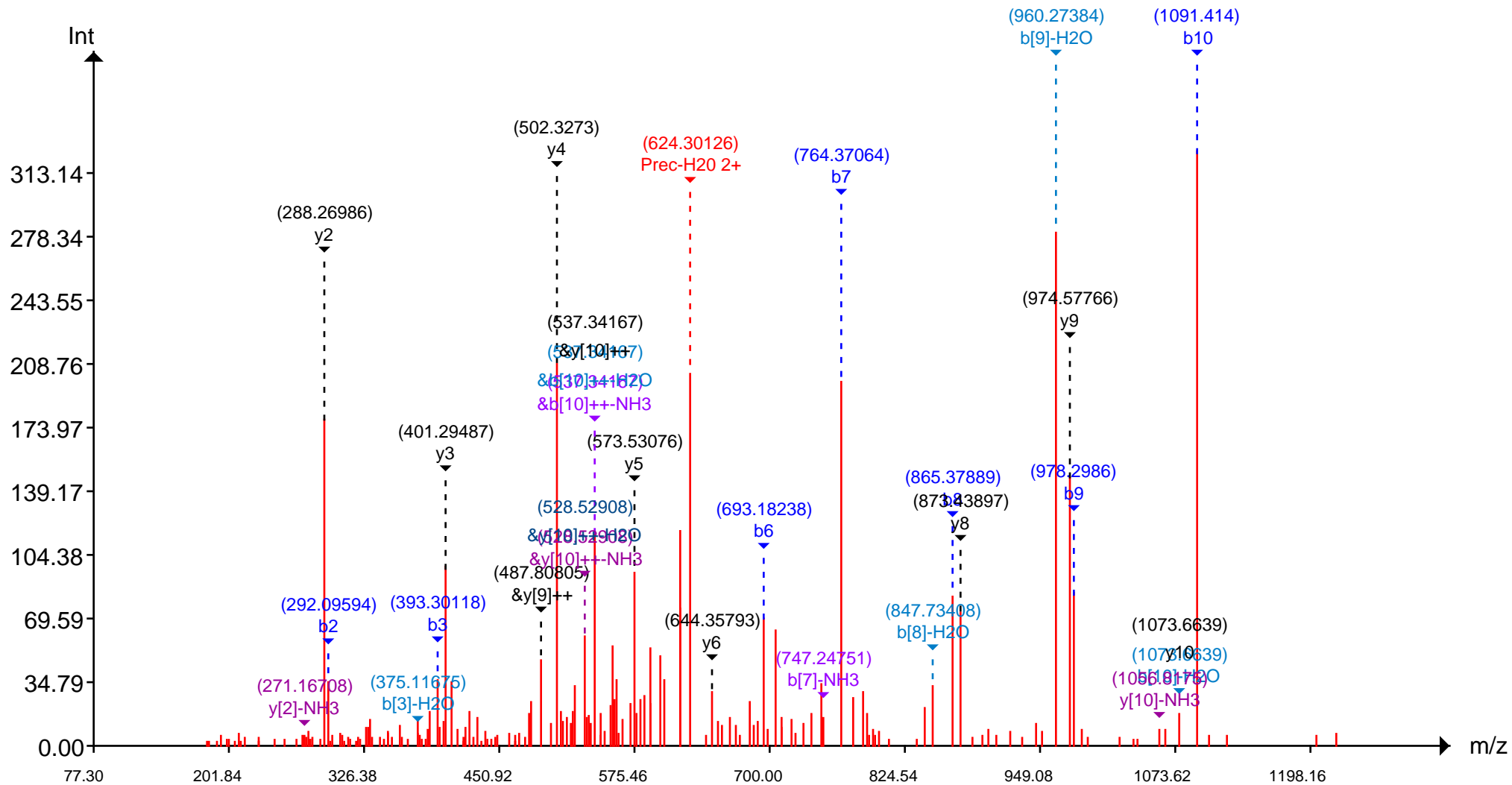
Confidence level 99.00%

Precursor m/z 867.89339

Identity Threshold 44.09

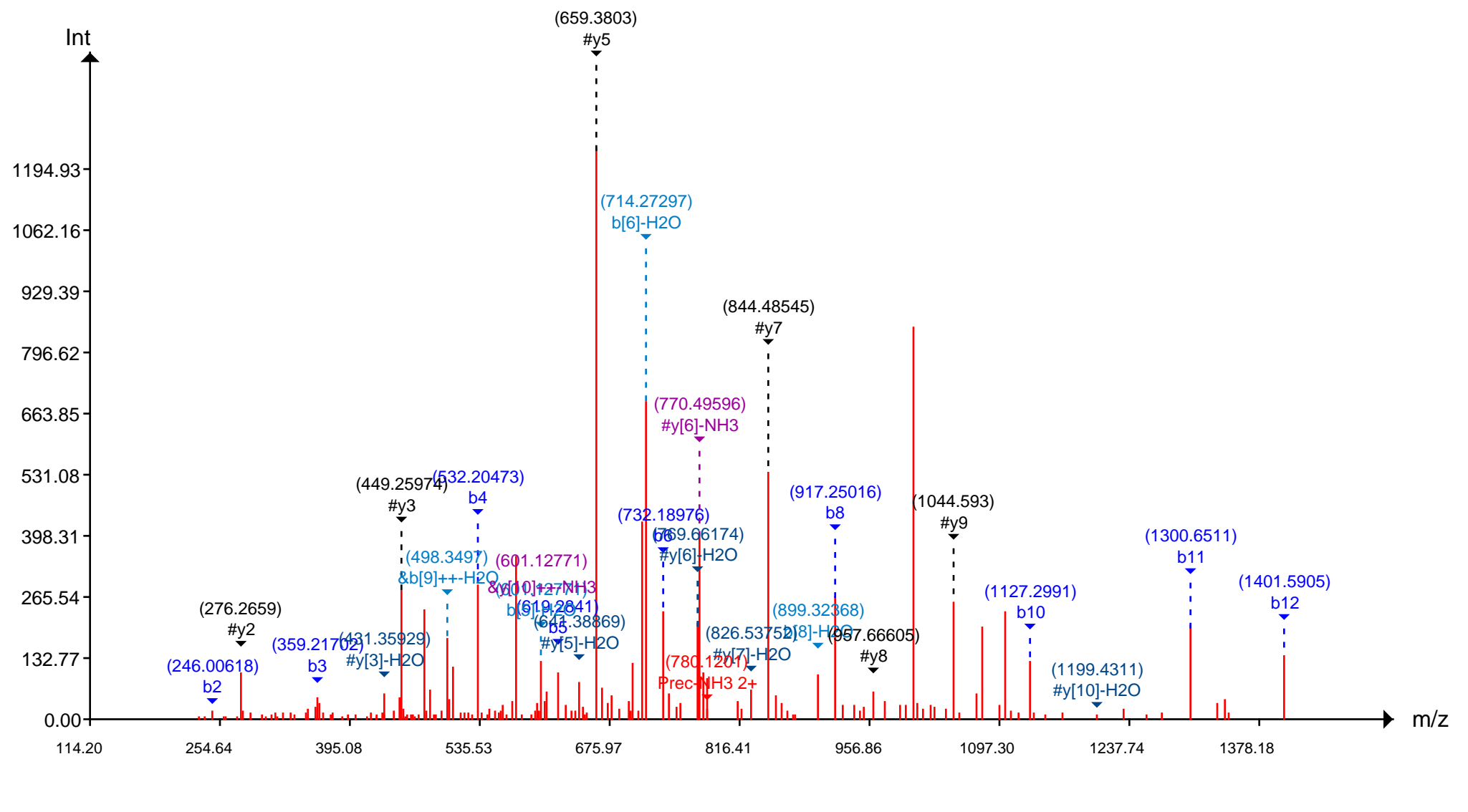
Charge 2+

Mass error 0.002056

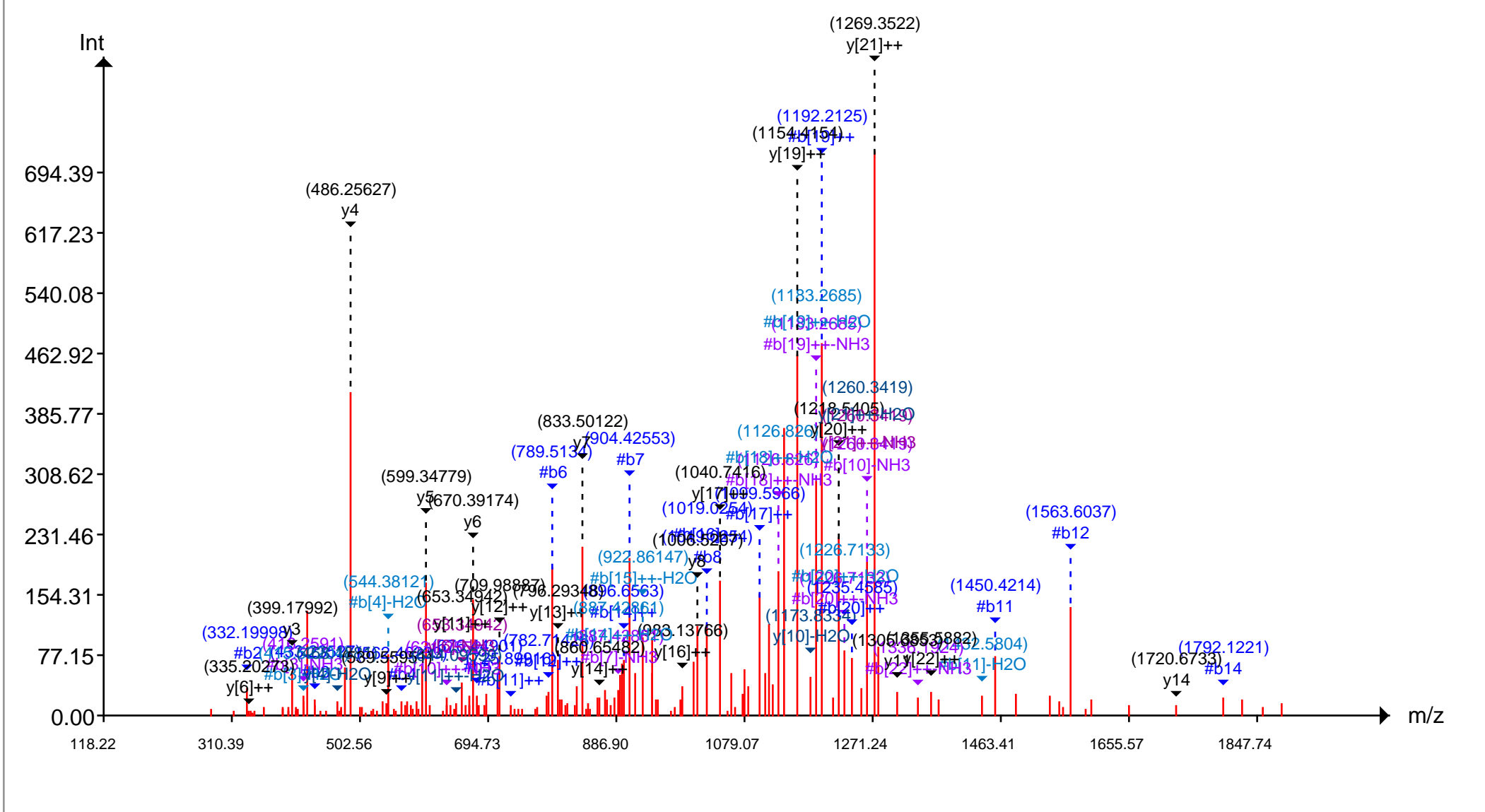


Info

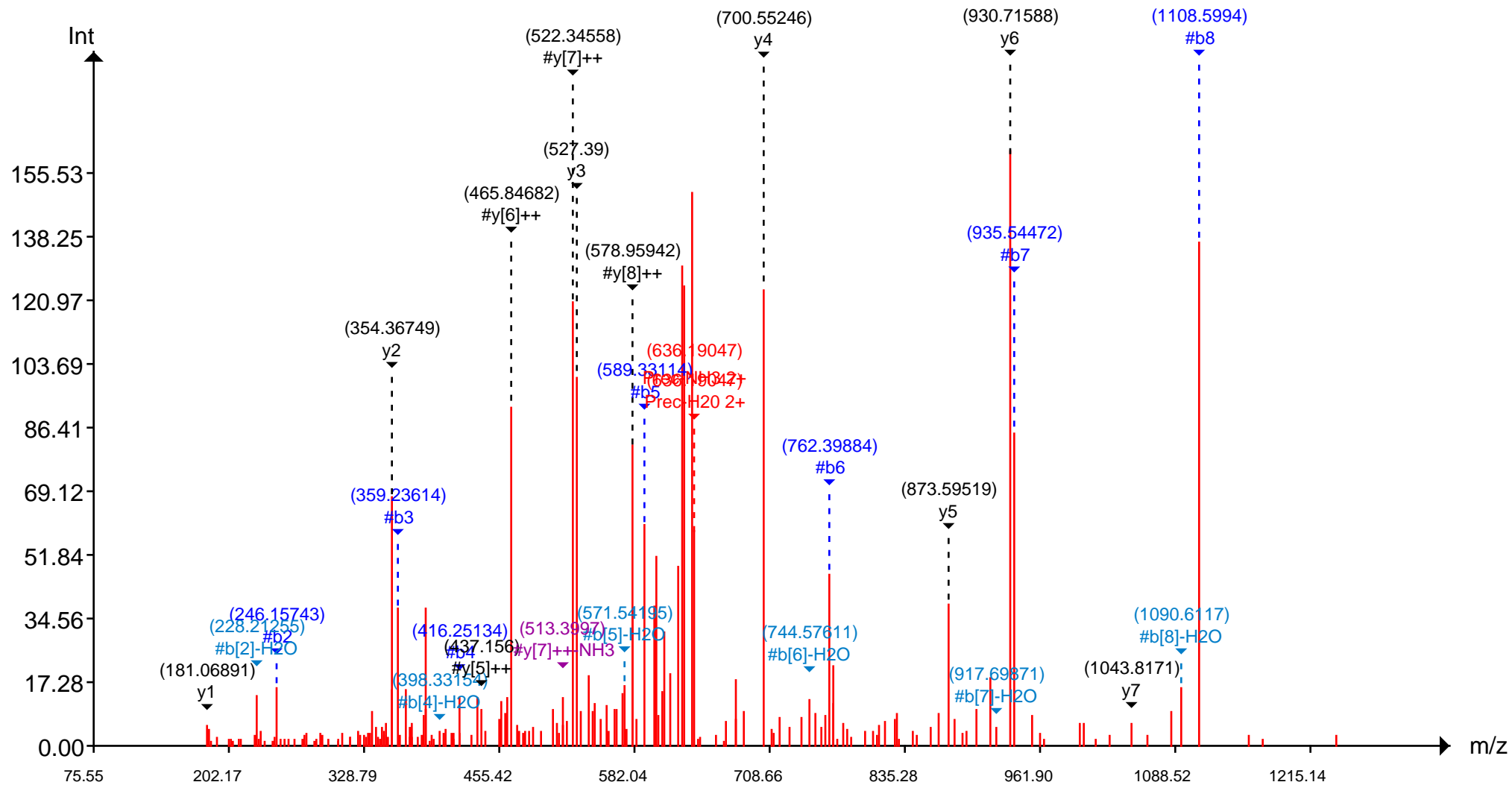
Modified Sequence	AcD3-FVTNDAATILR-COOH	Ion score	53.26
Protein	P42932	Confidence level	99.00%
Precursor m/z	633.34995	Identity Threshold	47.63
Charge	2+	Mass error	-1.23E-4



Info			
Modified Sequence	AcD3-AELK<AcD3K*>SLGQPIK<AcD3K*>TR-COOH	Ion score	46.26
Protein	Q61316	Confidence level	99.00%
Precursor m/z	788.47511	Identity Threshold	43.03
Charge	2+	Mass error	0.001494

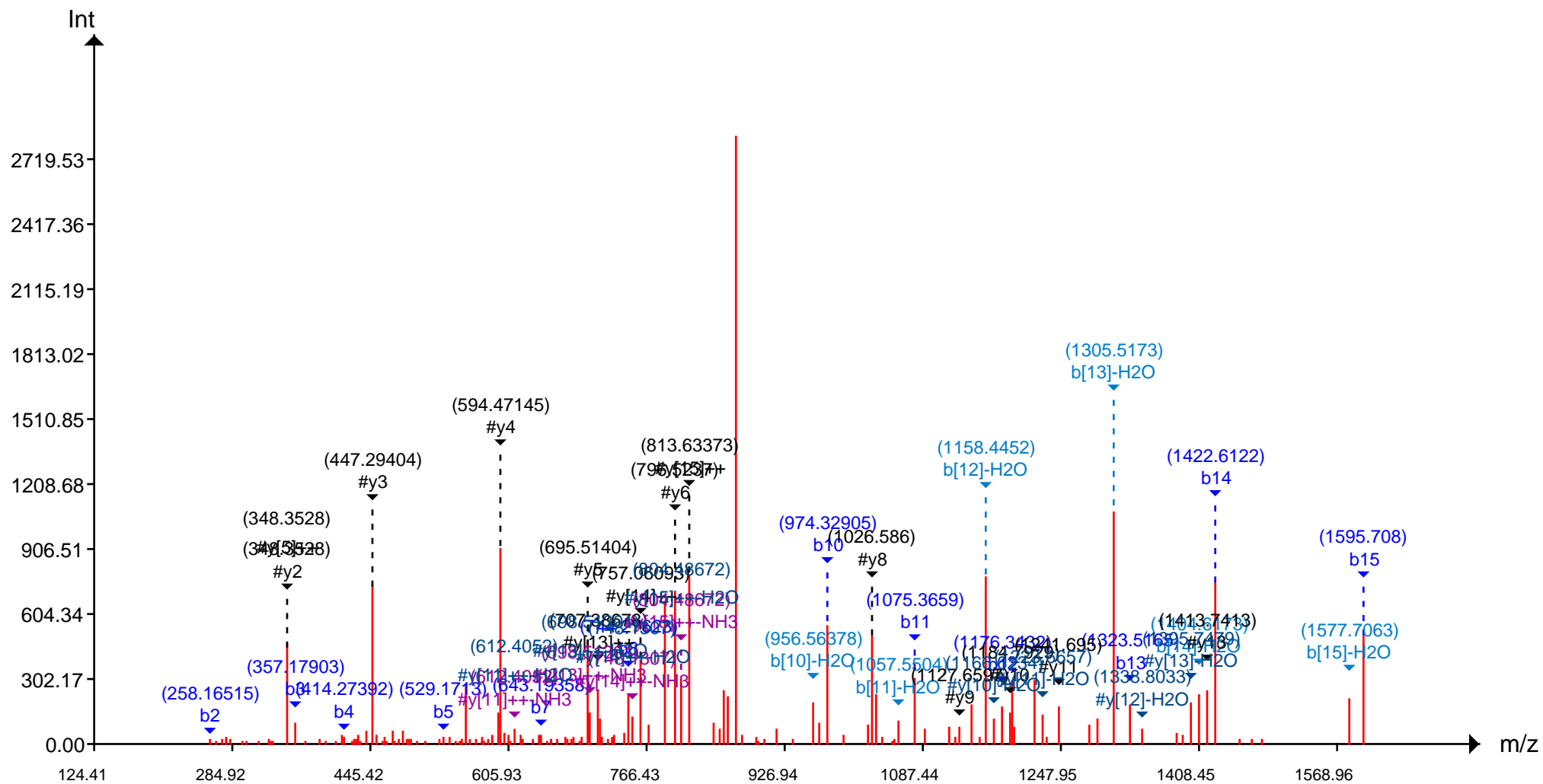


Info			
Modified Sequen...	AcD3-LK<AcD3K*>TENLDDEEK<AcD3K*>LNNAK<AcD3K*>YAISM<Mox*>AR<C13>-COOH	Ion score	58.71
Prot...	Q61233	Confidence level	99.00%
Precursor m...	956.83175	Identity Threshold	47.59
Char...	3+	Mass error	0.002719



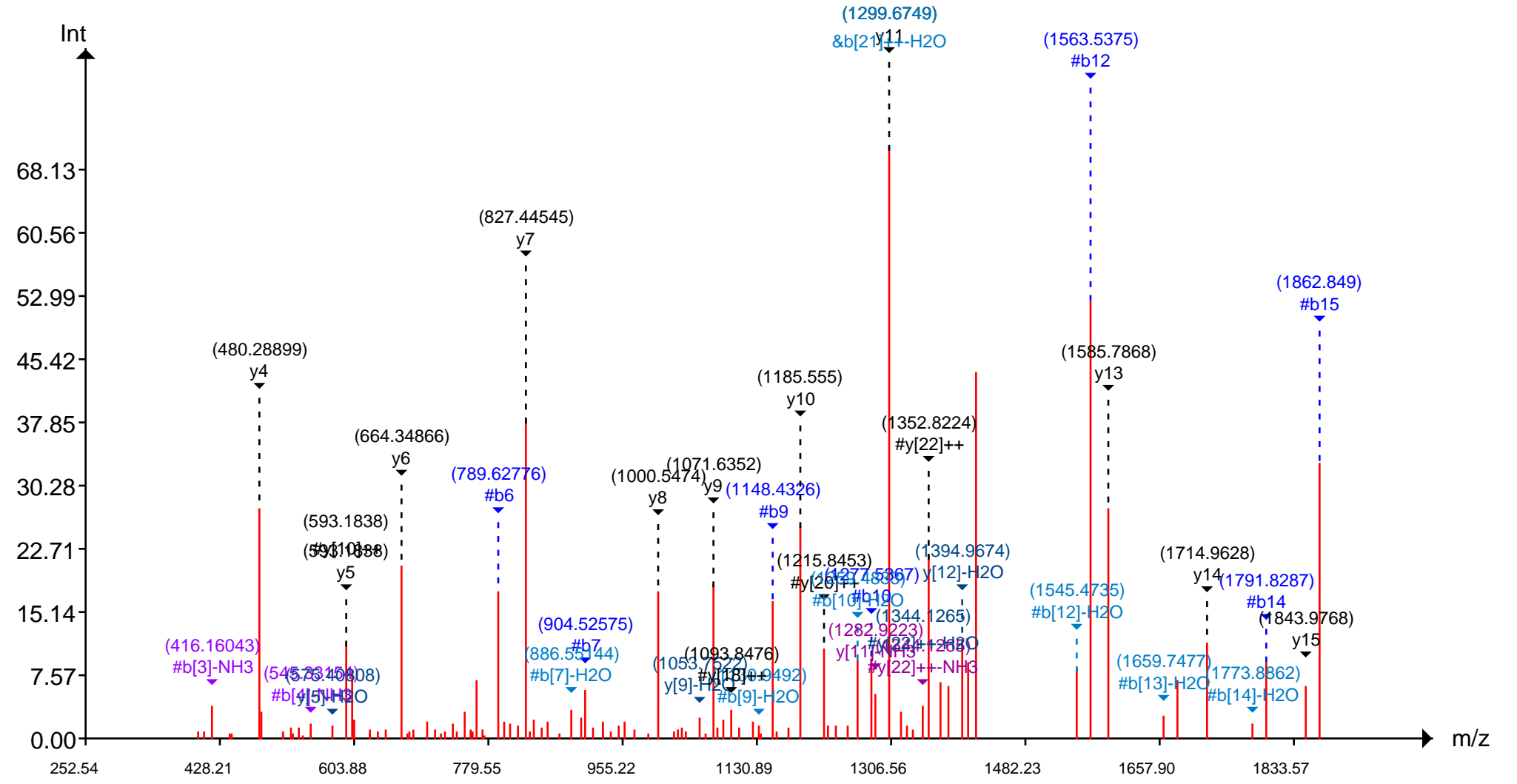
Info

Modified Sequence	AcD3-SLLGK<AcD3K*>K<AcD3K*>K<AcD3K*>K<AcD3K*>R<C13>-COOH	Ion score	43.05
Protein	P51410	Confidence level	99.00%
Precursor m/z	644.94776	Identity Threshold	29.30
Charge	2+	Mass error	7.19E-4



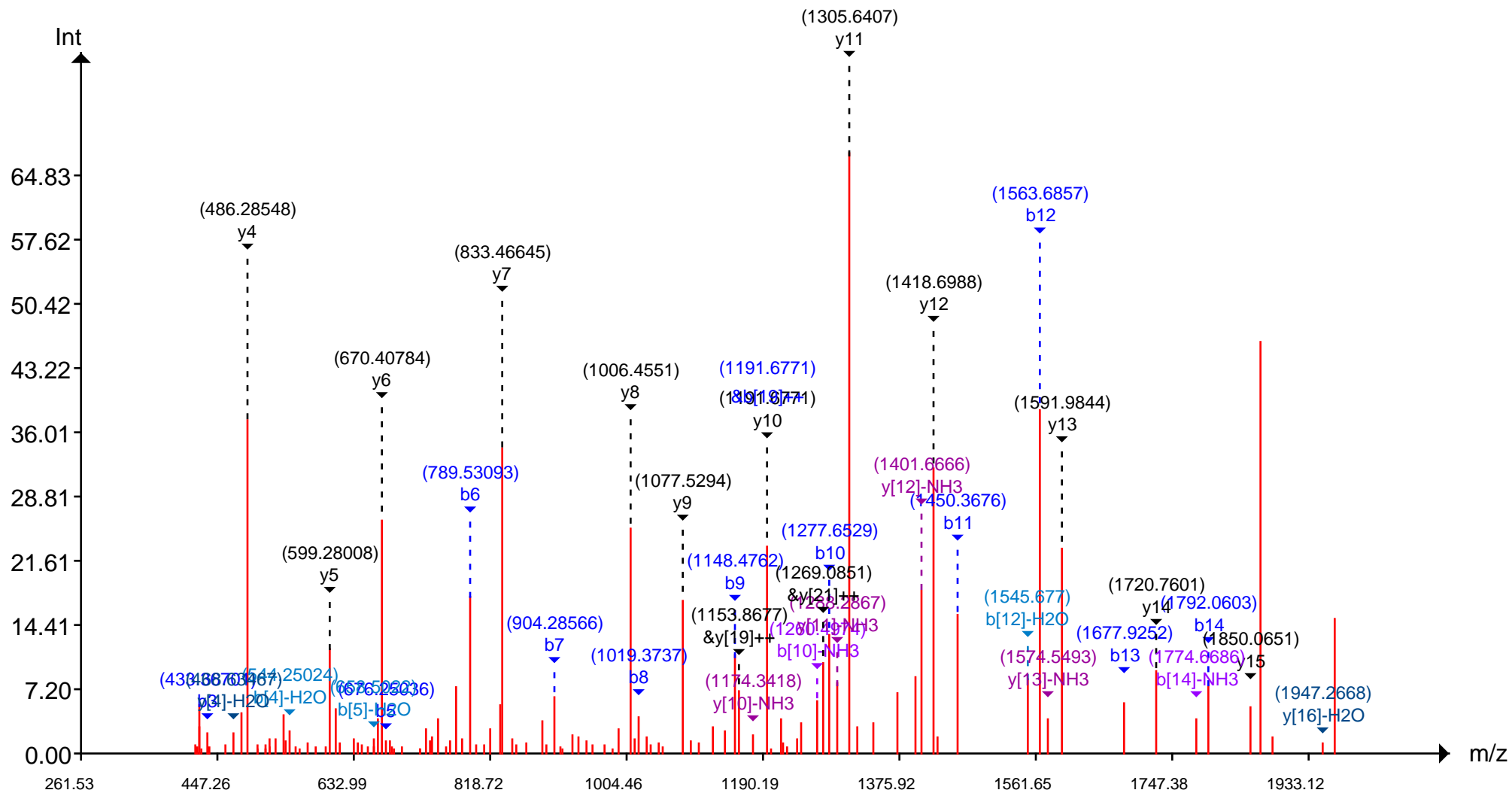
Info

Modified Sequence	AcD3-VLVGDGGTgk<AcD3K*>TTFVK<AcD3K*>R-COOH	Ion score	71.49
Protein	P62827	Confidence level	99.00%
Precursor m/z	885.51063	Identity Threshold	45.34
Charge	2+	Mass error	0.003321



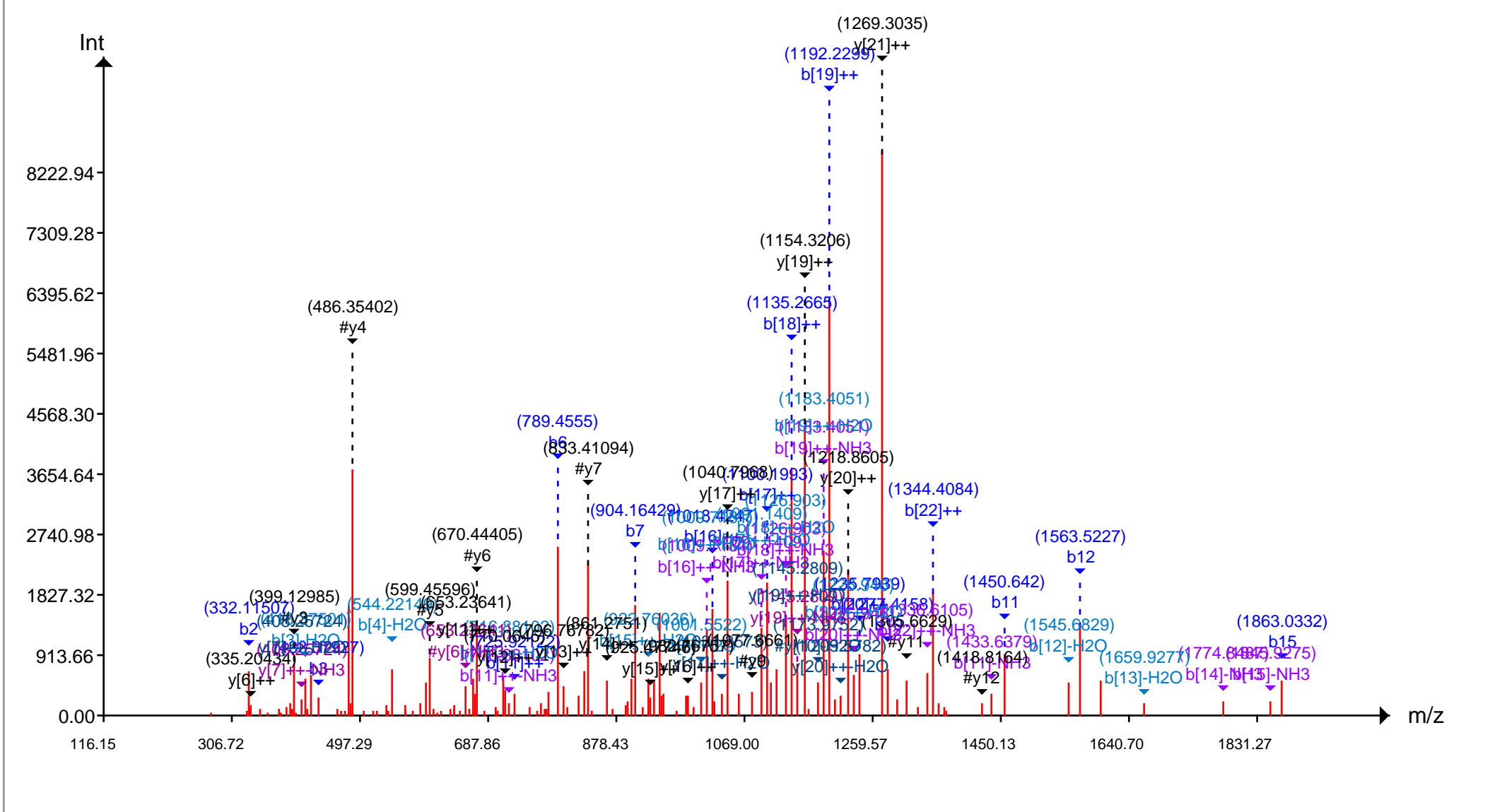
Info

Modified Sequence	AcD3-LK<AcD3K*>TENLDDEEK<AcD3K*>LNNAK<AcD3K*>YAISM<Mox*>AR-COOH	Ion score	86.75
Protein	Q61233	Confidence level	99.00%
Precursor m/z	1431.7338	Identity Threshold	47.74
Charge	2+	Mass error	0.002486

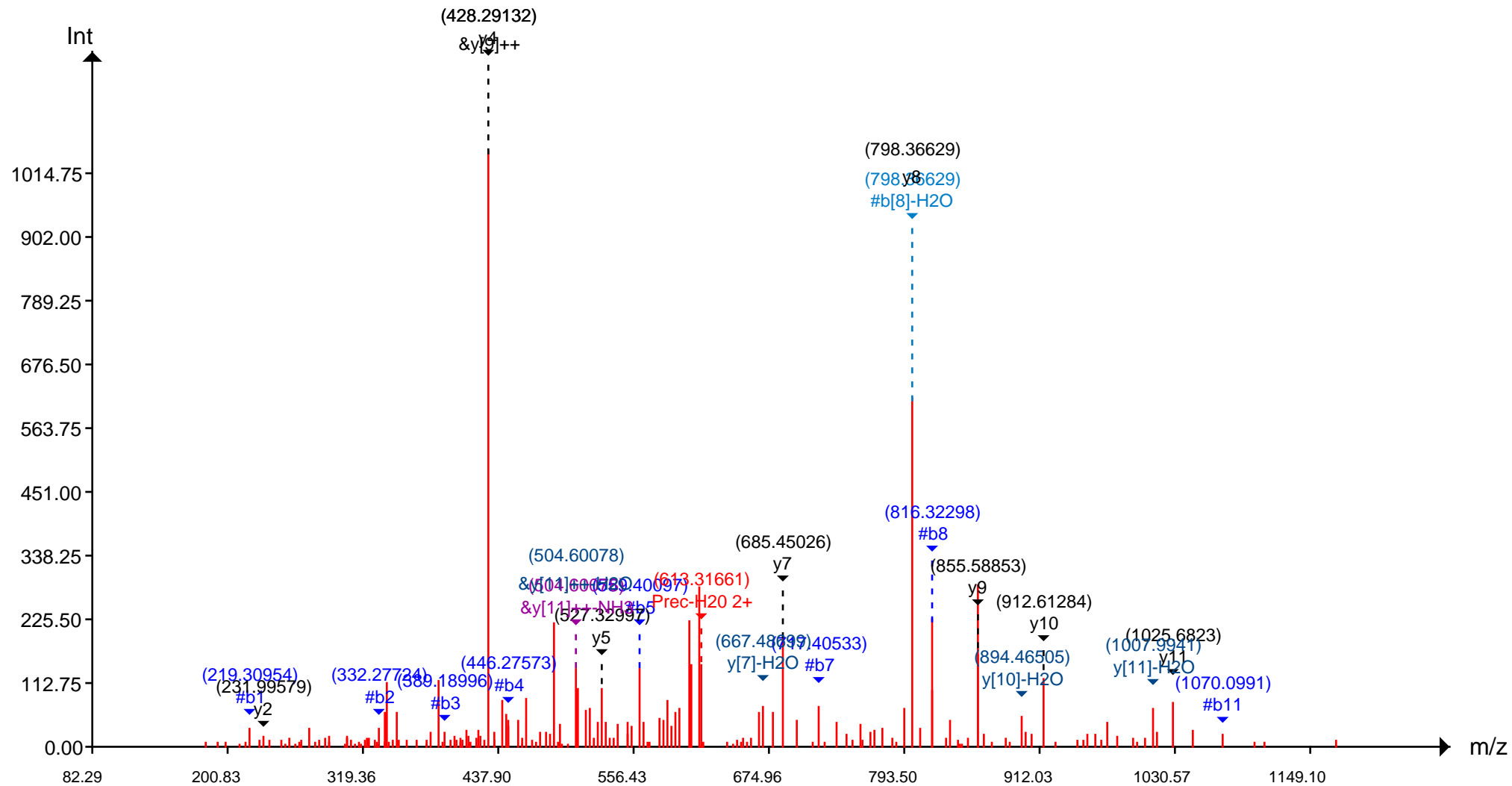


Info

Modified Sequen...	AcD3-LK<AcD3K*>TENLDDEEK<AcD3K*>LNNAK<AcD3K*>YAISM<Mox*>AR<C13>-COOH	Ion score	90.47
Prot...	Q61233	Confidence level	99.00%
Precursor m...	1434.7437	Identity Threshold	47.59
Char...	2+	Mass error	0.002145



Info			
Modified Sequen...	AcD3-LK<AcD3K*>TENLDDEEK<AcD3K*>LNNAK<AcD3K*>YAISM<Mox*>AR<C13>-COOH	Ion score	64.75
Prot...	Q61233	Confidence level	99.00%
Precursor m...	956.83167	Identity Threshold	47.60
Char...	3+	Mass error	0.002479



Info

Modified Sequence AcD3-K<AcD3K*>IGGIGTVPVGR-COOH

Ion score 45.5

Protein P10126, P62631

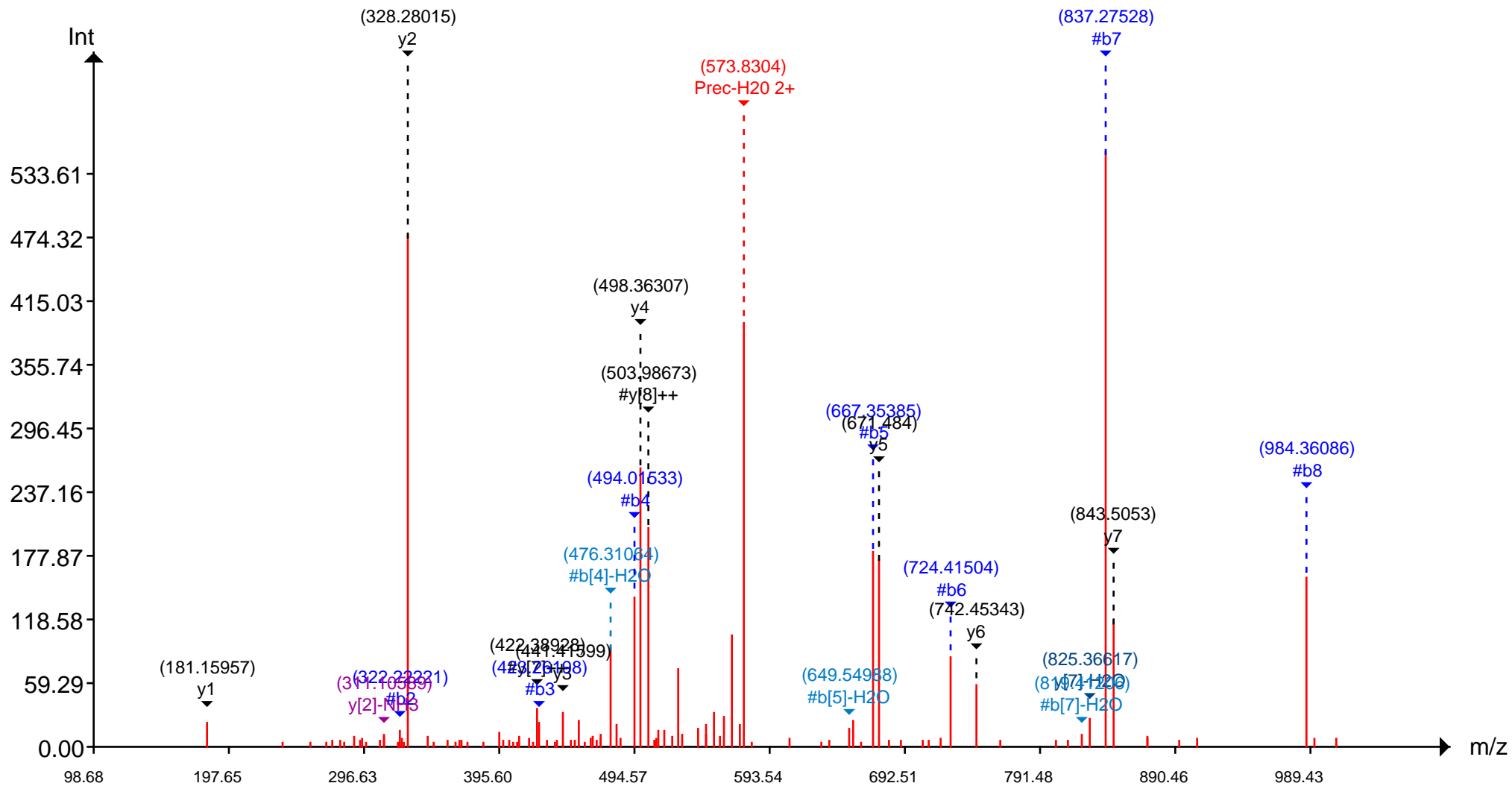
Confidence level 99.00%

Precursor m/z 622.38613

Identity Threshold 42.64

Charge 2+

Mass error 0.001009



Info

Modified Sequence AcD3-LYTAK<AcD3K*>GLFR<C13>-COOH

Ion score 51.94

Protein P17182, P17183

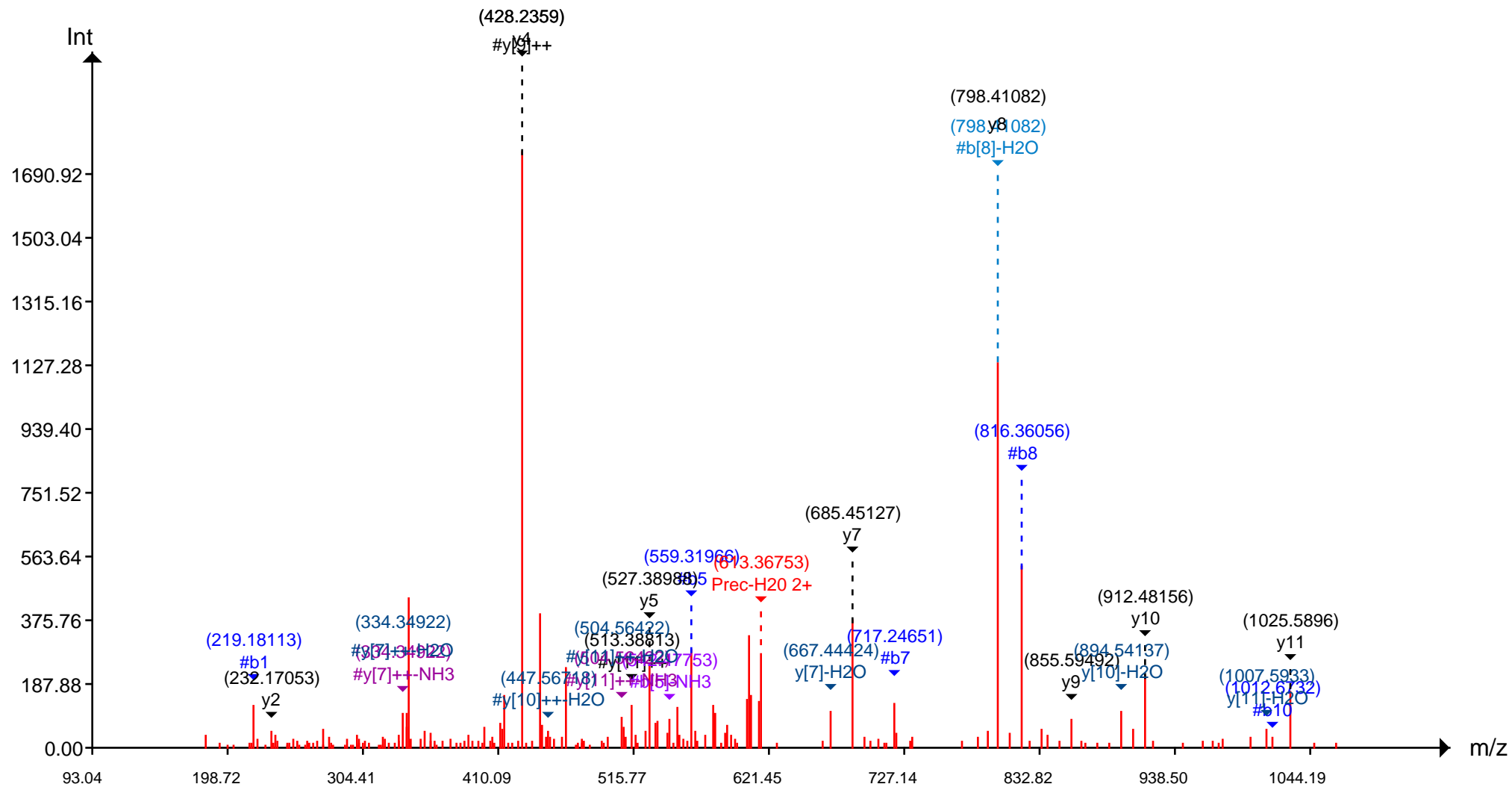
Confidence level 99.00%

Precursor m/z 582.85345

Identity Threshold 44.24

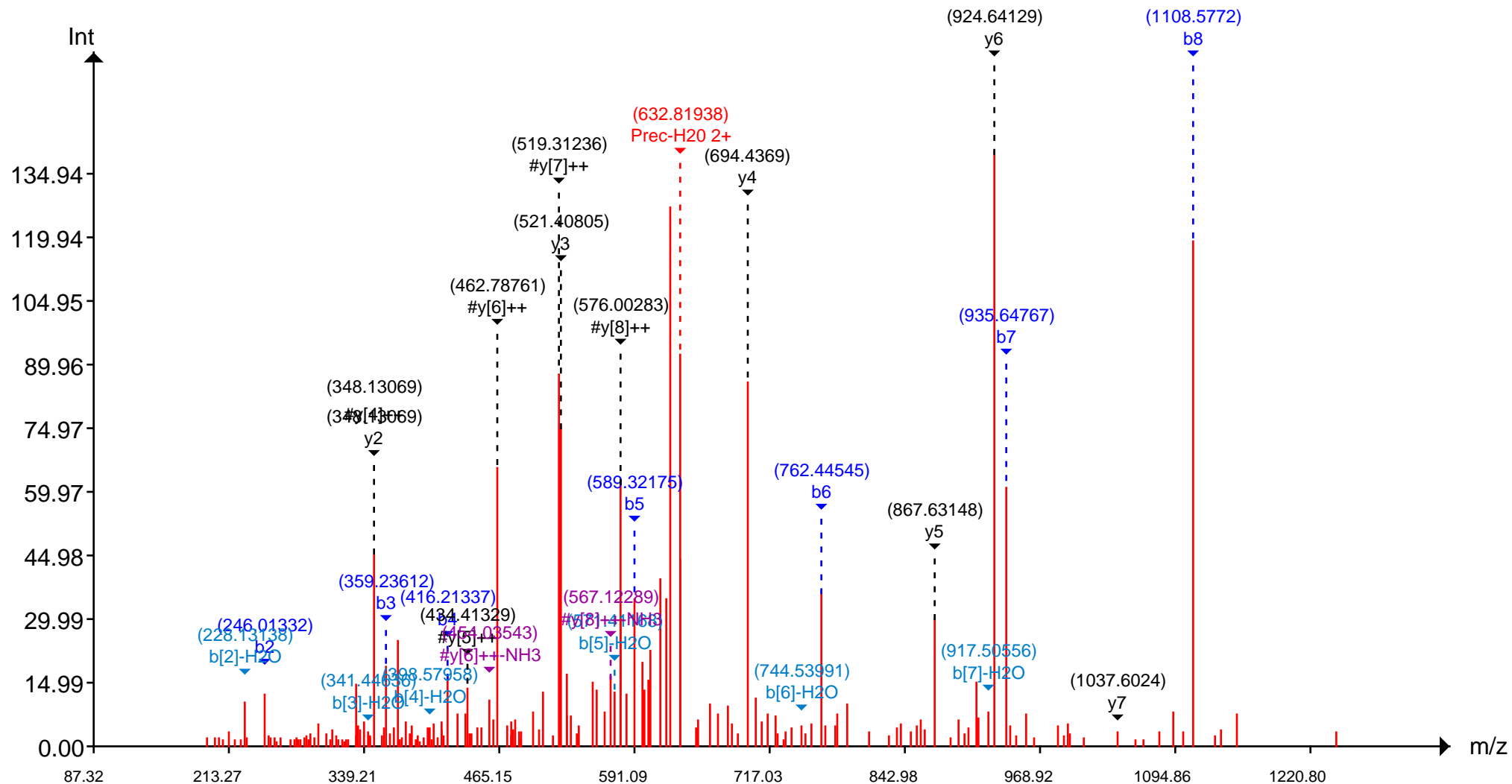
Charge 2+

Mass error 6.67E-4



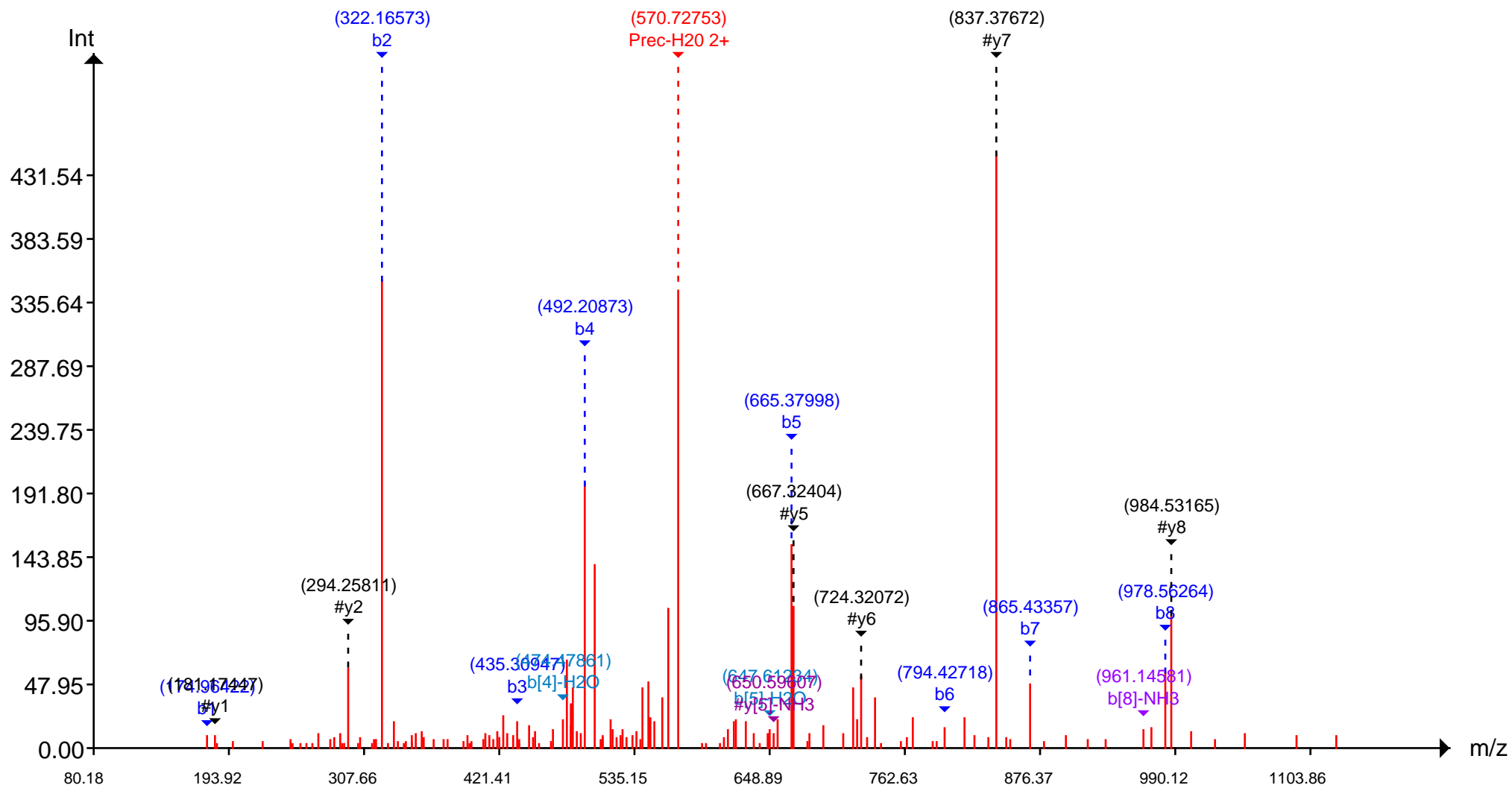
Info

Modified Sequence	AcD3-K<AcD3K*>IGGIGTVPVGR-COOH	Ion score	65.1
Protein	P10126, P62631	Confidence level	99.00%
Precursor m/z	622.38511	Identity Threshold	42.78
Charge	2+	Mass error	-0.001031



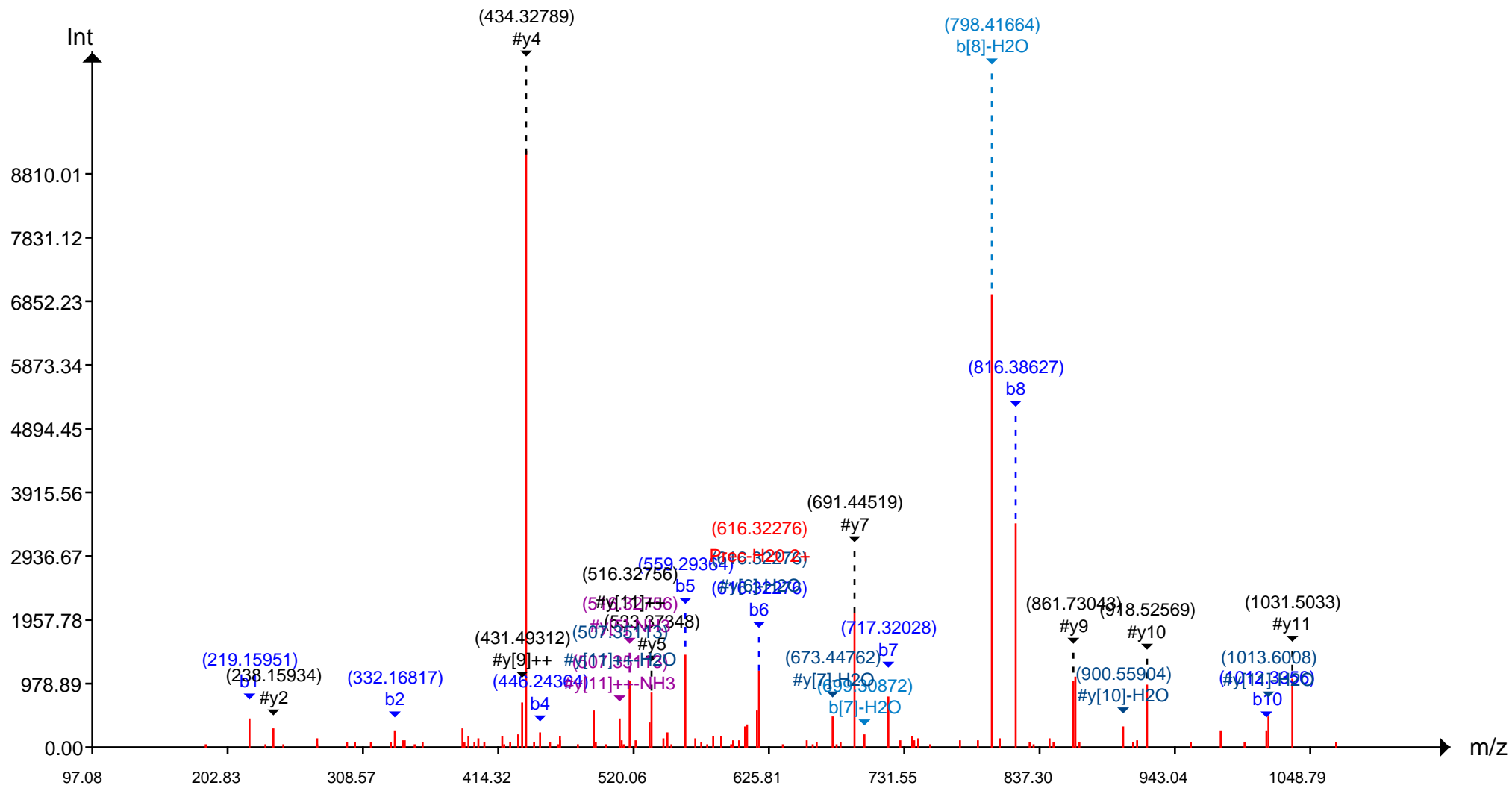
Info

Modified Sequence	AcD3-SLLGK<AcD3K*>K<AcD3K*>K<AcD3K*>K<AcD3K*>R-COOH	Ion score	44.81
Protein	P51410	Confidence level	99.00%
Precursor m/z	641.93754	Identity Threshold	31.01
Charge	2+	Mass error	4.21E-4



Info

Modified Sequence	AcD3-EFIGK<AcD3K*>EAIR<C13>-COOH	Ion score	49.75
Protein	Q8K448	Confidence level	99.00%
Precursor m/z	579.84274	Identity Threshold	46.05
Charge	2+	Mass error	0.00508



Info

Modified Sequence AcD3-K<AcD3K*>IGGIGTVPVGR<C13>-COOH

Ion score 58.64

Protein P10126, P62631

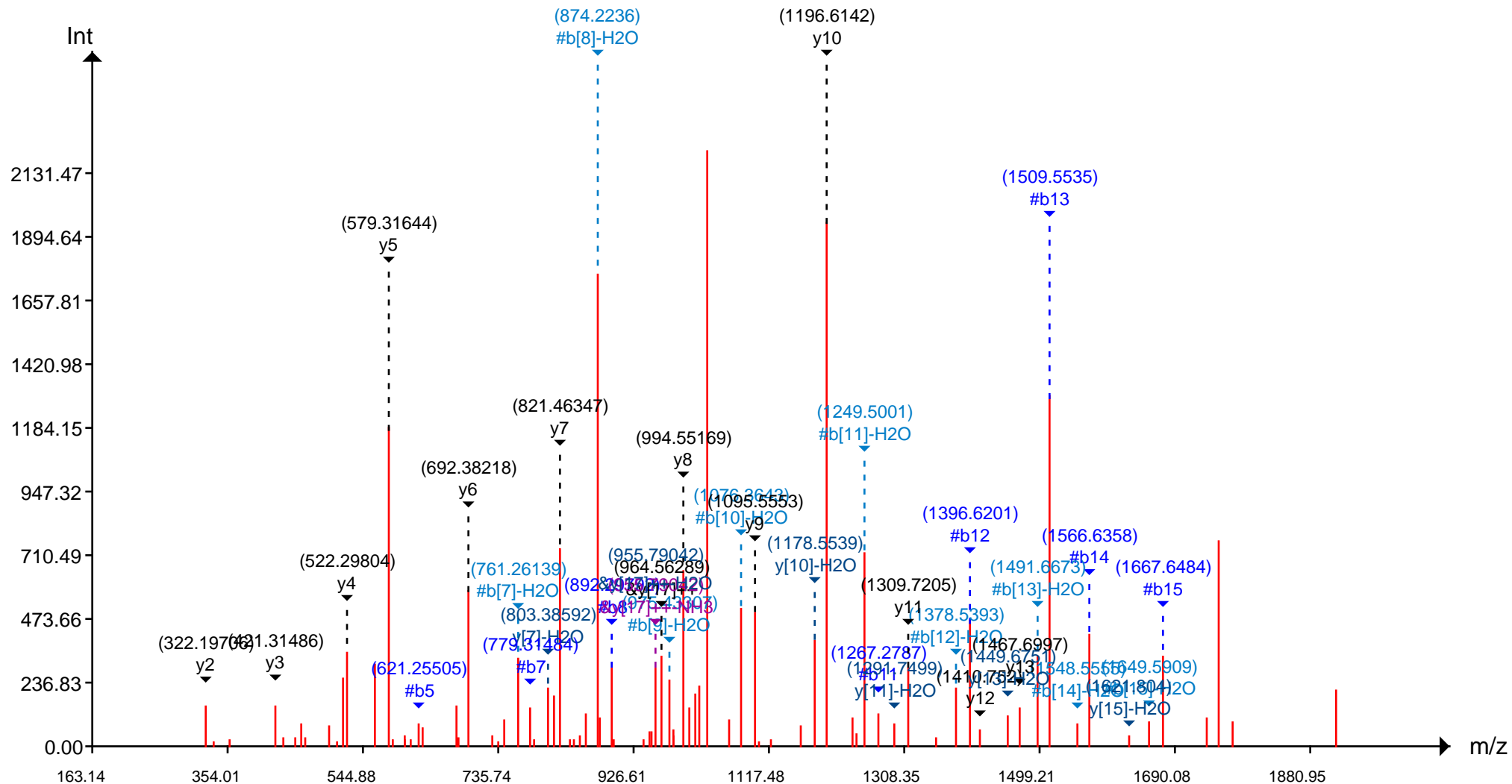
Confidence level 99.00%

Precursor m/z 625.39557

Identity Threshold 40.69

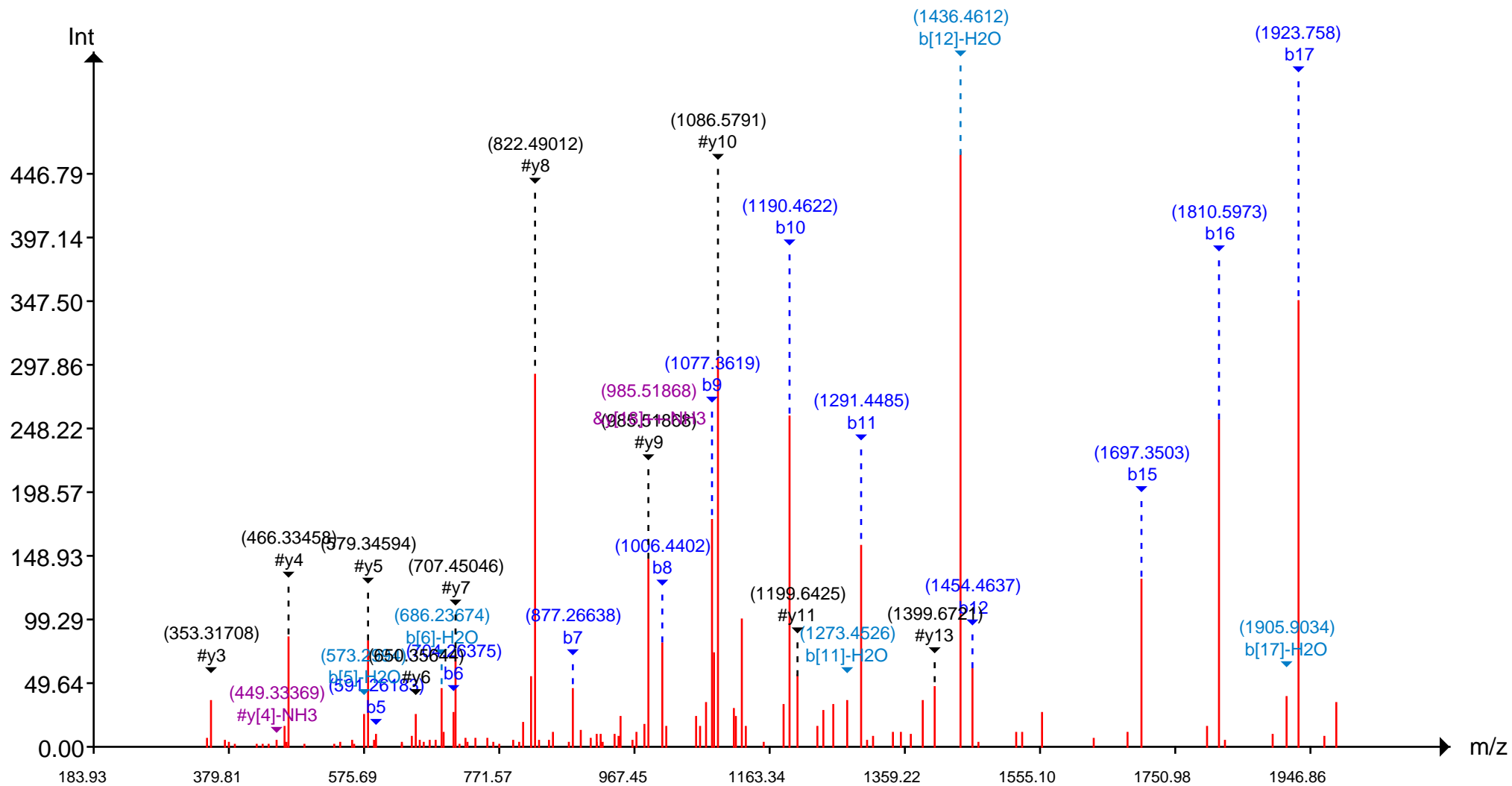
Charge 2+

Mass error -2.52E-4



Info

Modified Sequence	AcD3-DK<AcD3K*>DGDGTITTK<AcD3K*>ELGTVM<Mox*>R-COOH	Ion score	93.05
Protein	P62204	Confidence level	99.00%
Precursor m/z	1044.529	Identity Threshold	47.74
Charge	2+	Mass error	0.003165



Info

Modified Sequence AcD3-DEVDSLK<AcD3K*>EALTYDGALLGDR<C13>-COOH

Ion score 70.26

Protein Q9WUK2

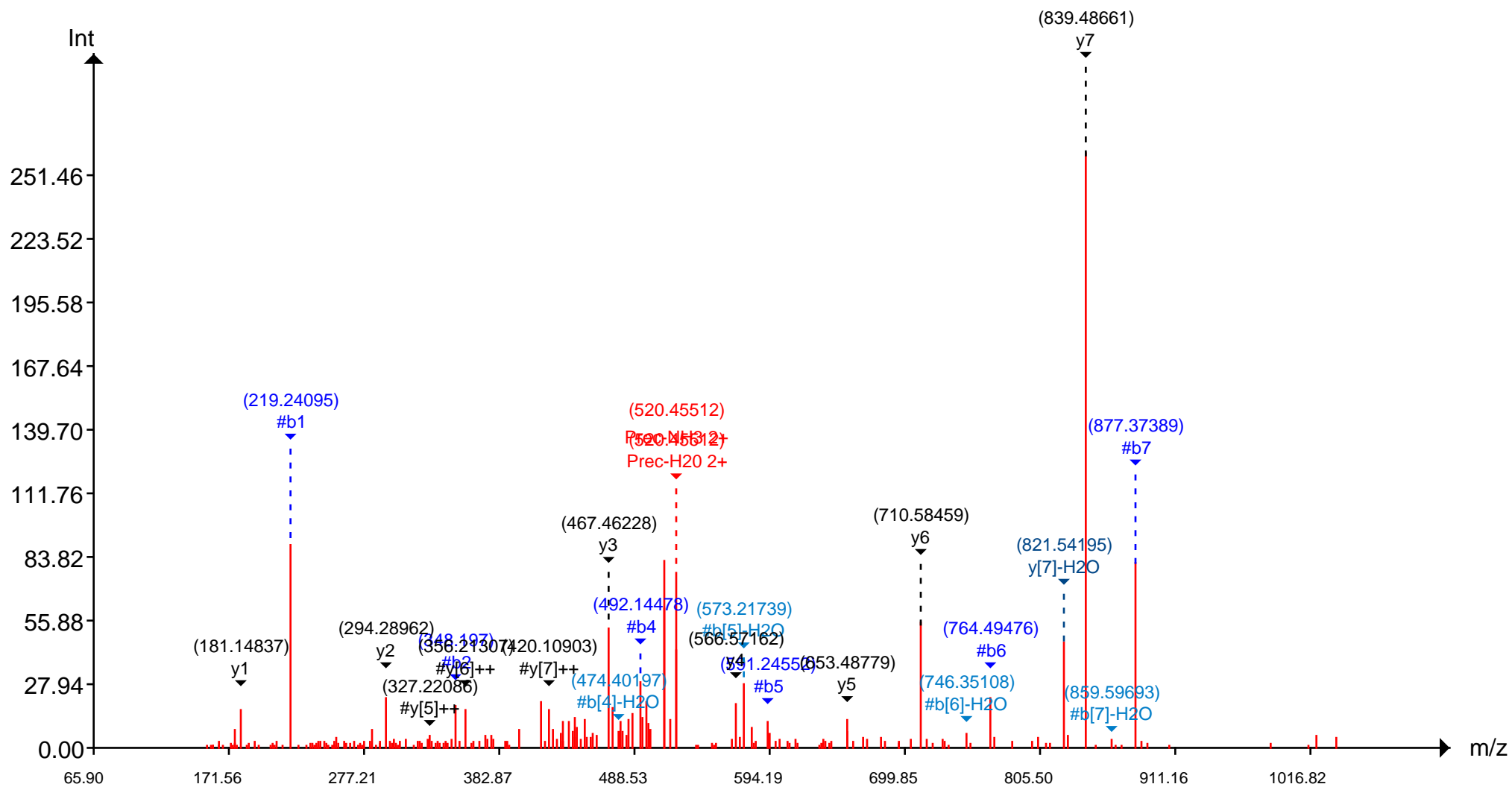
Confidence level 99.00%

Precursor m/z 1138.5808

Identity Threshold 47.91

Charge 2+

Mass error 0.003676



Info

Modified Sequence AcD3-K<AcD3K*>EGSVK<AcD3K*>LR<C13>-COOH

Ion score 46.43

Protein Q9JLQ0

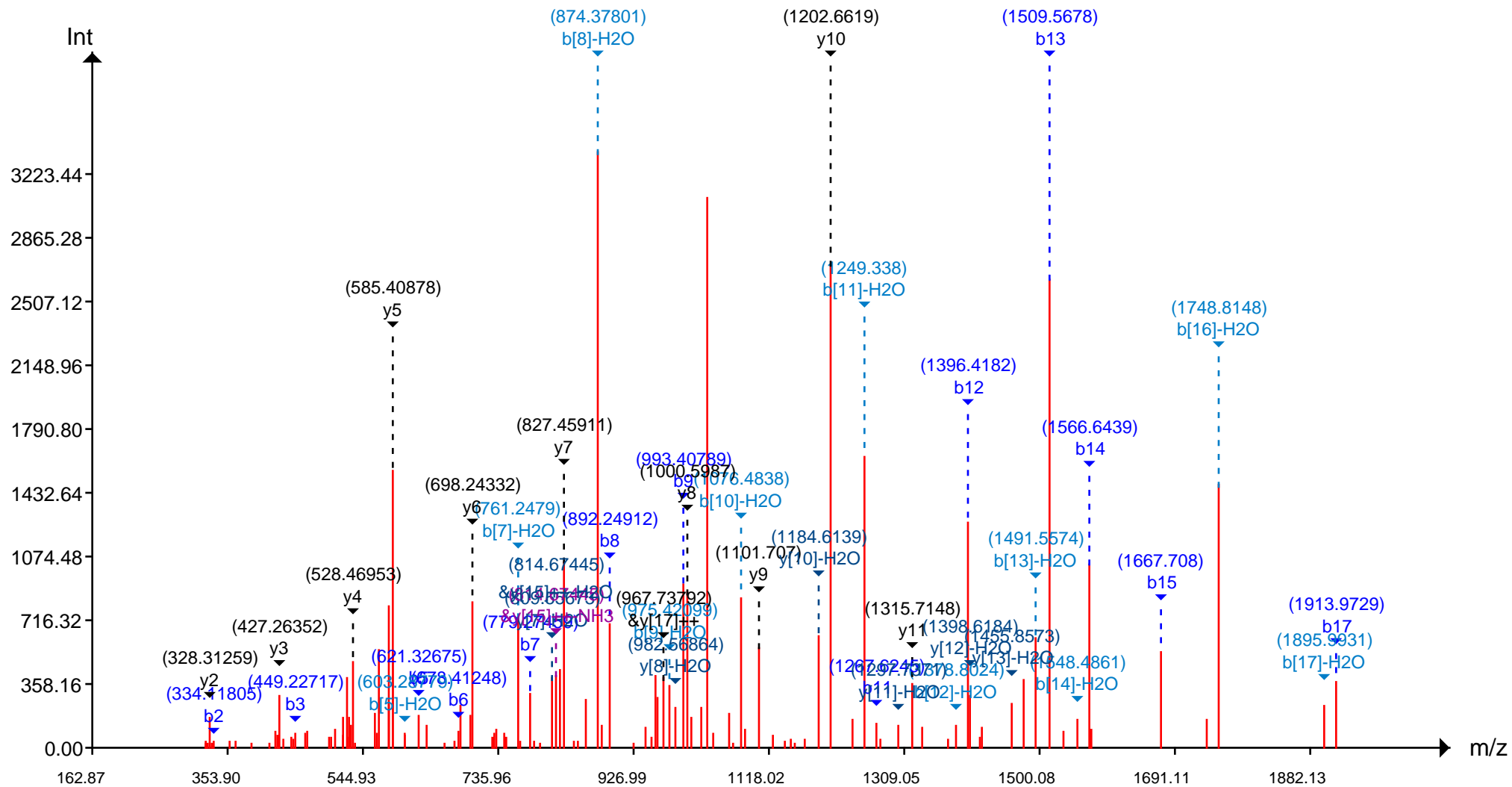
Confidence level 99.00%

Precursor m/z 529.33626

Identity Threshold 42.67

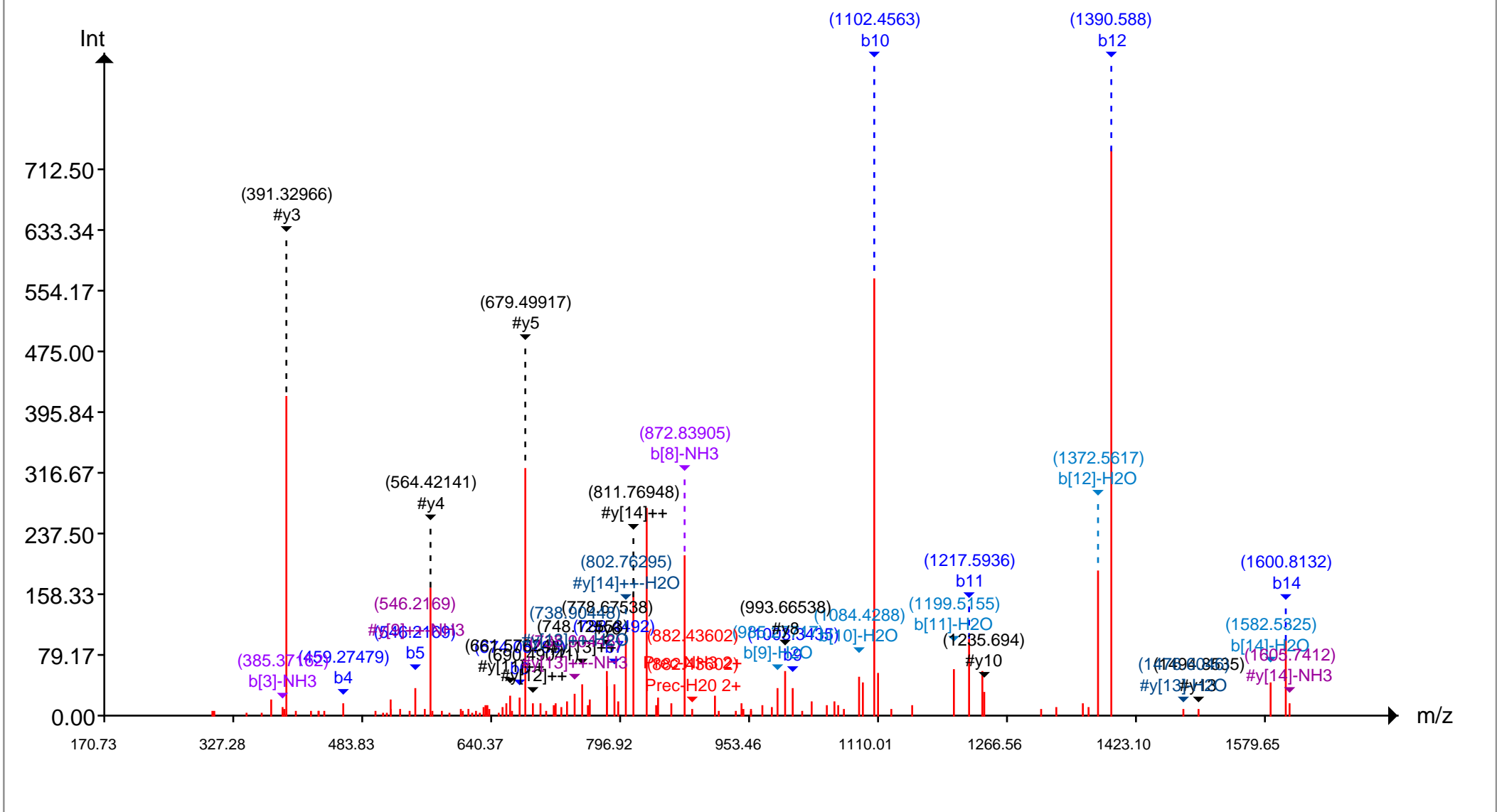
Charge 2+

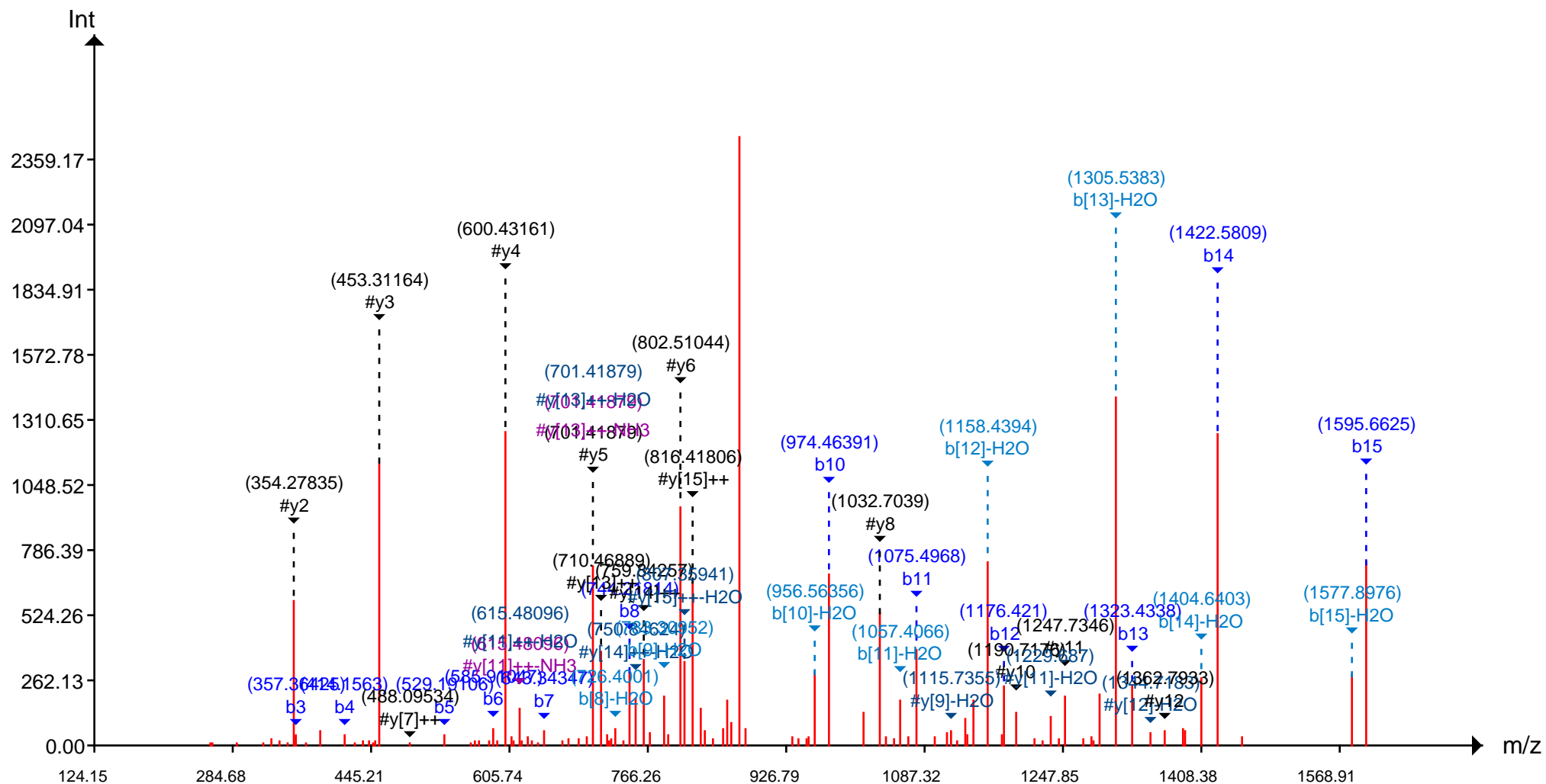
Mass error -5.1E-4



Info

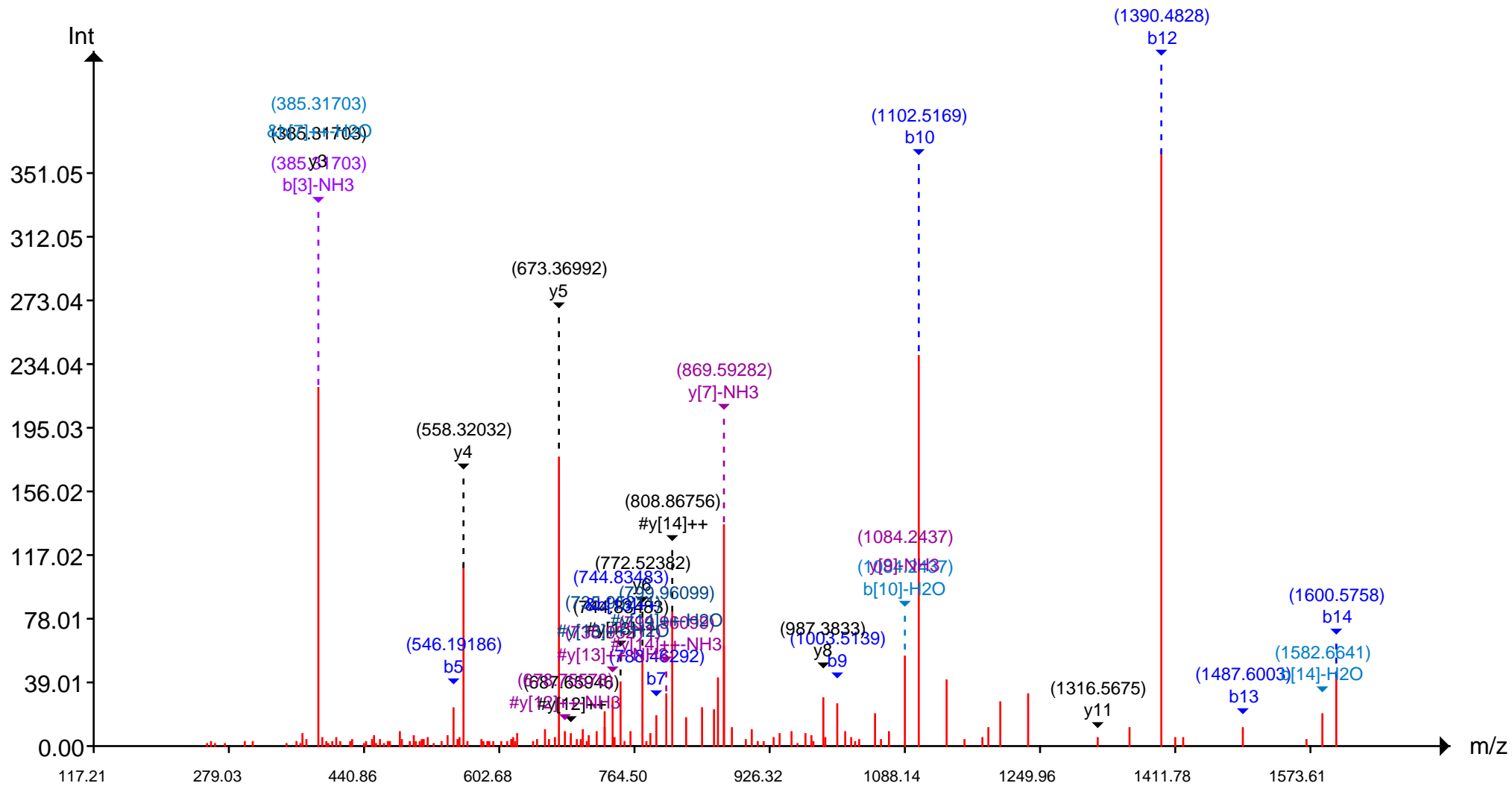
Modified Sequence	AcD3-DK<AcD3K*>DGDGTITTK<AcD3K*>ELGTVM<Mox*>R<C13>-COOH	Ion score	69.86
Protein	P62204	Confidence level	99.00%
Precursor m/z	1047.5382	Identity Threshold	47.99
Charge	2+	Mass error	0.001423





Info

Modified Sequence	AcD3-VLVGDGGTgk<AcD3K*>TTFVK<AcD3K*>R<C13>-COOH	Ion score	102.74
Protein	P62827	Confidence level	99.00%
Precursor m/z	888.52044	Identity Threshold	44.36
Charge	2+	Mass error	0.002799



Info

Modified Sequence AcD3-IQDGSQNTNVDK<AcD3K*>PLR-COOH

Ion score 49.63

Protein Q3U0V1

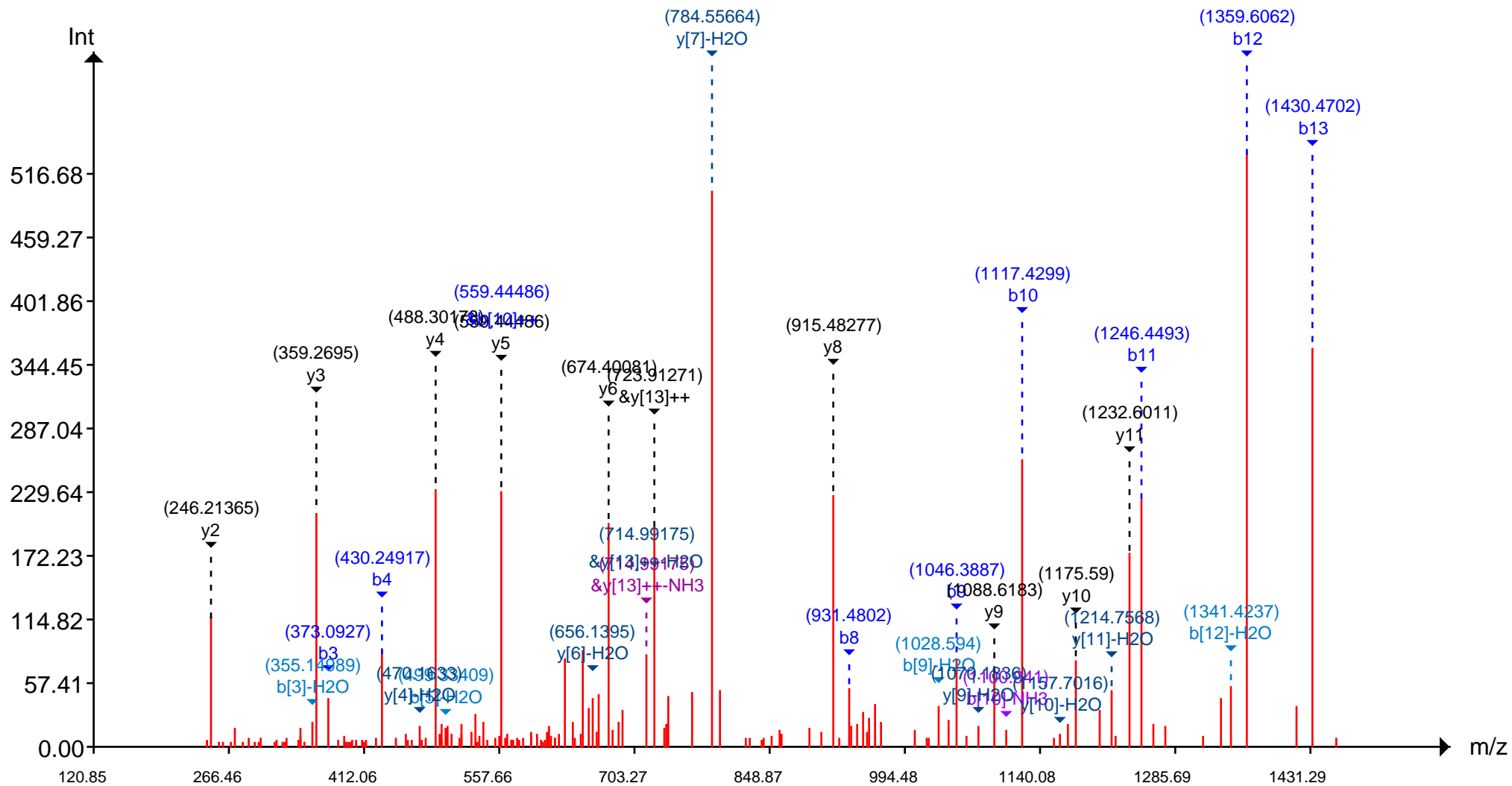
Confidence level 99.00%

Precursor m/z 887.96501

Identity Threshold 48.13

Charge 2+

Mass error 0.002672



Info

Modified Sequence AcD3-LTIGSK<AcD3K*>LQDAEIAR-COOH

Ion score 62.54

Protein P14069

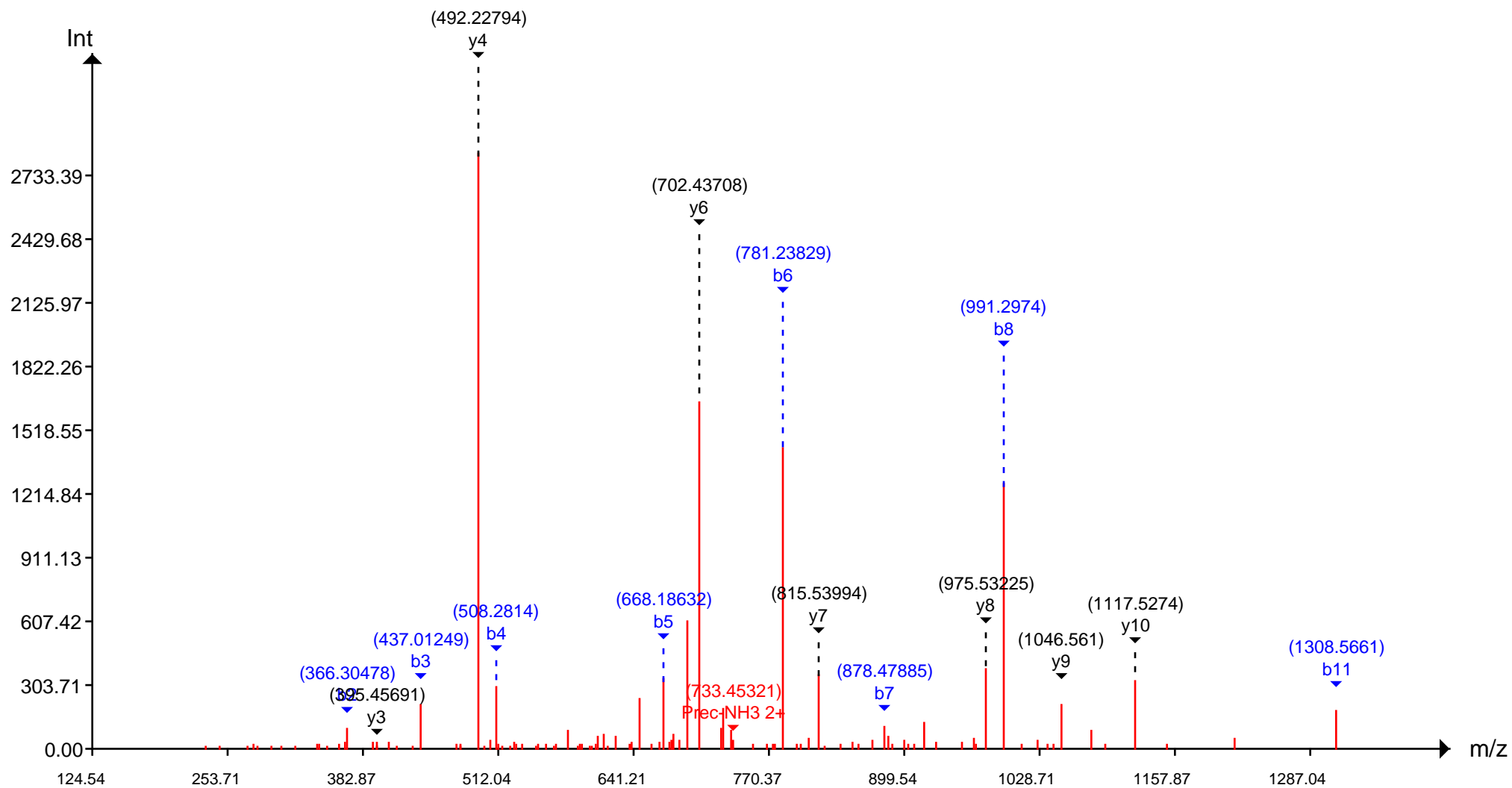
Confidence level 99.00%

Precursor m/z 802.96074

Identity Threshold 46.03

Charge 2+

Mass error 0.001761



Info

Modified Sequence AcD3-FK<AcD3K*>AAC<Cmm*>LPLPGYR-COOH

Ion score 53.75

Protein Q61233

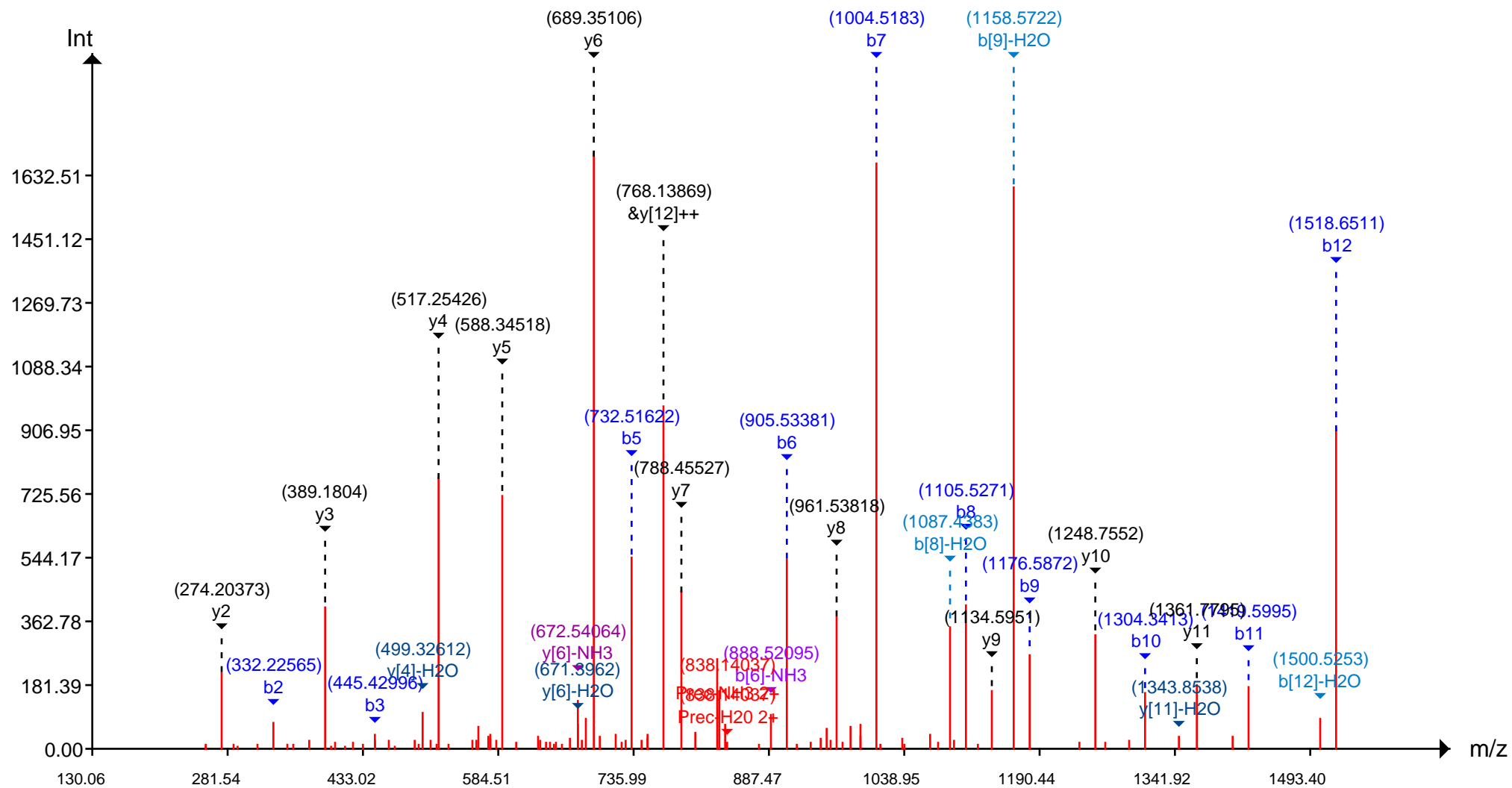
Confidence level 99.00%

Precursor m/z 741.90653

Identity Threshold 47.78

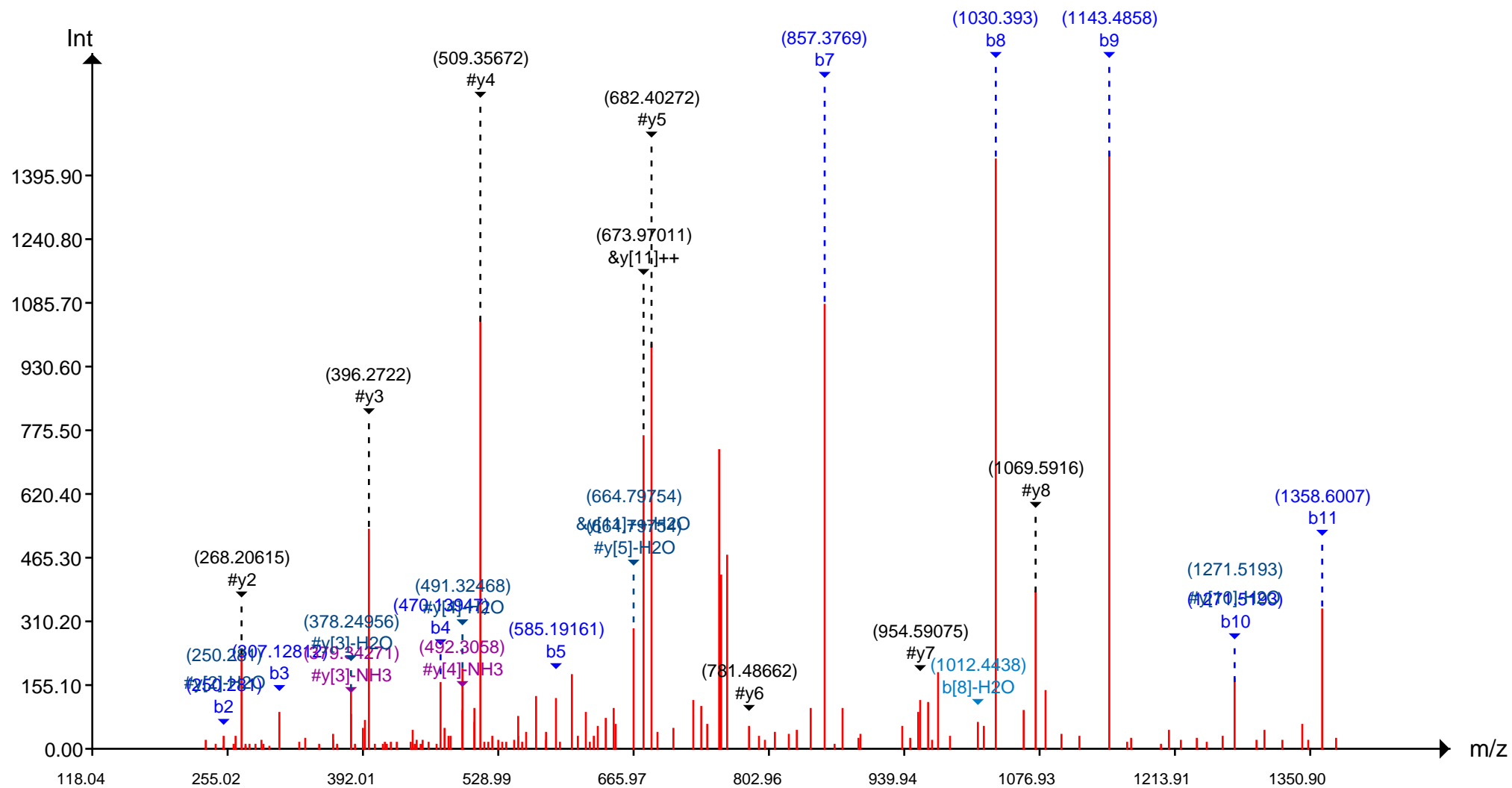
Charge 2+

Mass error 0.001358



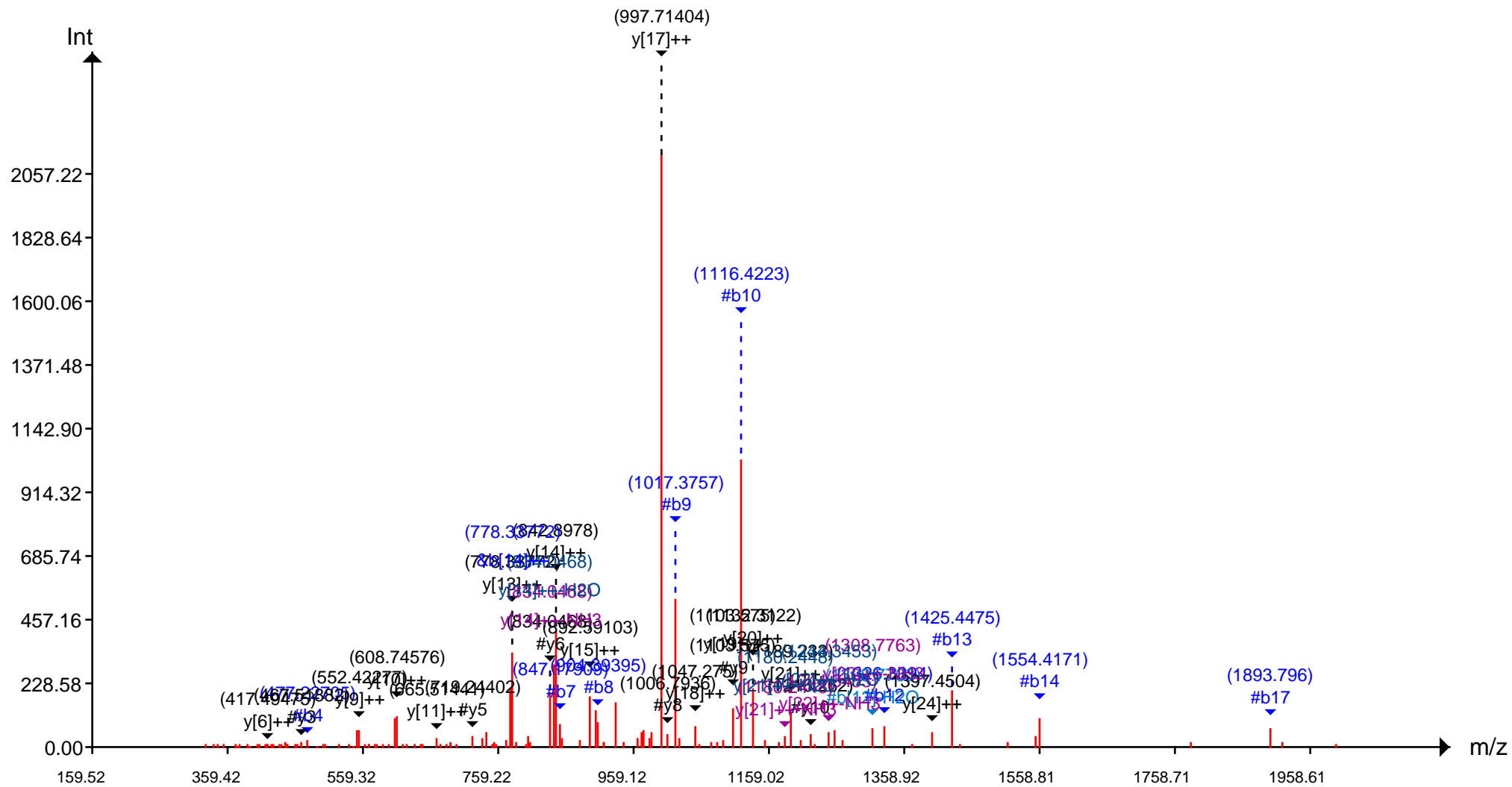
Info

Modified Sequence	AcD3-LK<AcD3K*>LNK<AcD3K*>K<AcD3K*>VTAQDVR-COOH	Ion score	95.0
Protein	P26041	Confidence level	99.00%
Precursor m/z	847.02432	Identity Threshold	39.48
Charge	2+	Mass error	0.001755



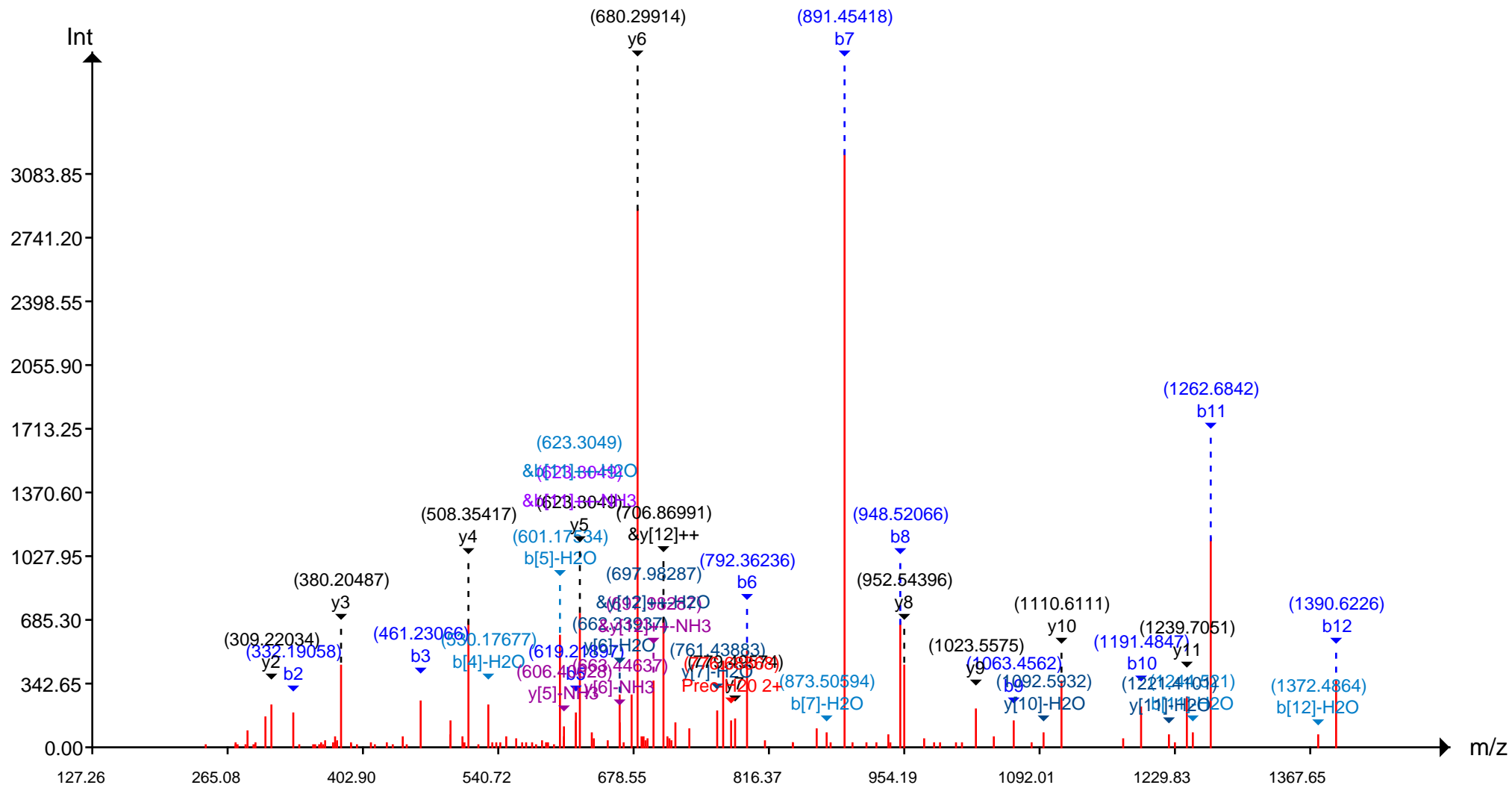
Info

Modified Sequence	AcD3-FGGYDK<AcD3K*>VK<AcD3K*>LQSR<C13>-COOH	Ion score	58.05
Protein	Q62465	Confidence level	99.00%
Precursor m/z	769.9351	Identity Threshold	46.90
Charge	2+	Mass error	0.001018



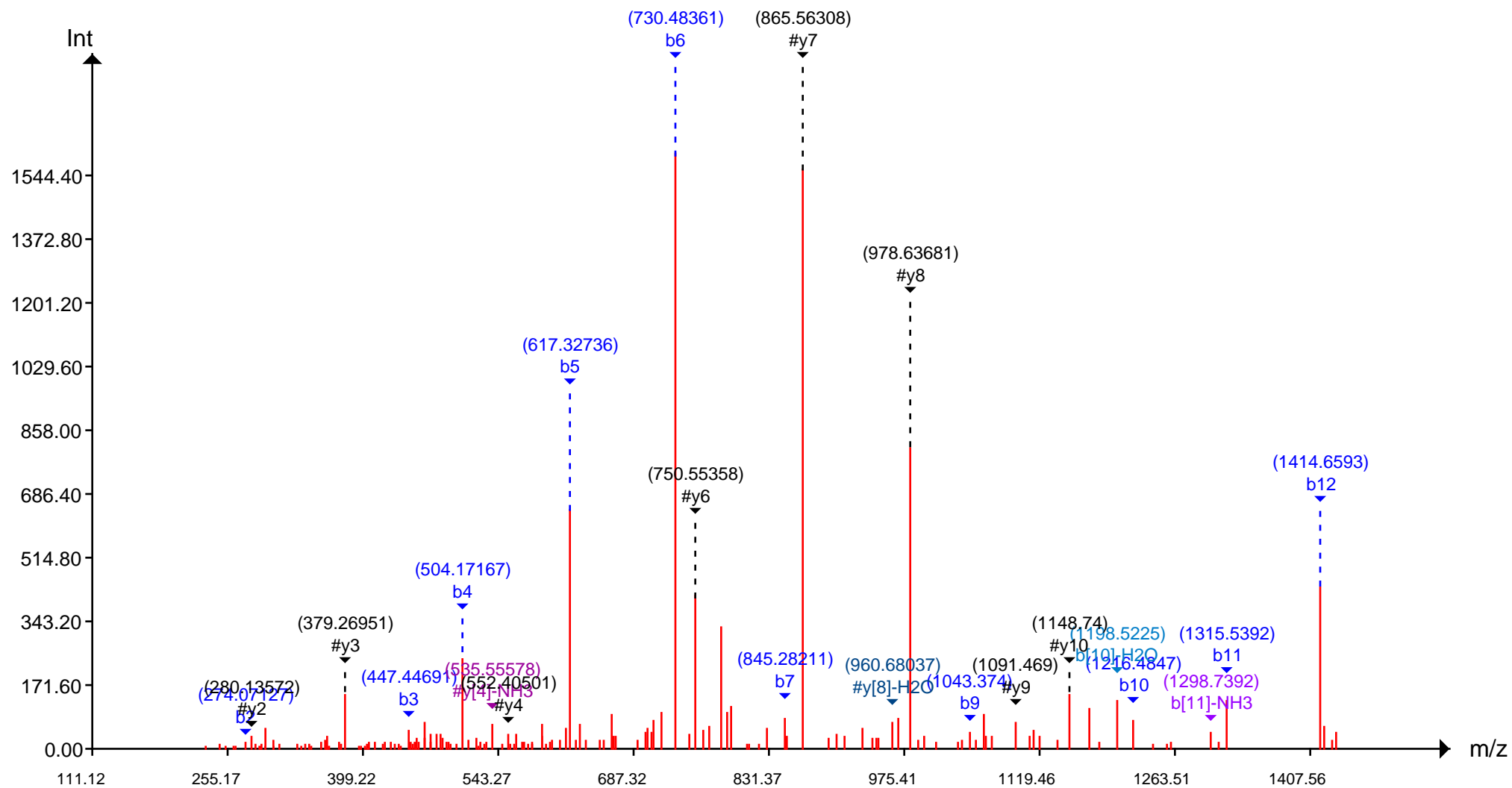
Info

Modified Sequence	AcD3-ASIC<Cmm*>QQNGIVPIVEPEILPDGDHDLK<AcD3K*>R<C13>-COOH	Ion score	70.48
Protein	P05064, P05063	Confidence level	99.00%
Precursor m/z	1037.2123	Identity Threshold	47.17
Charge	3+	Mass error	0.002279



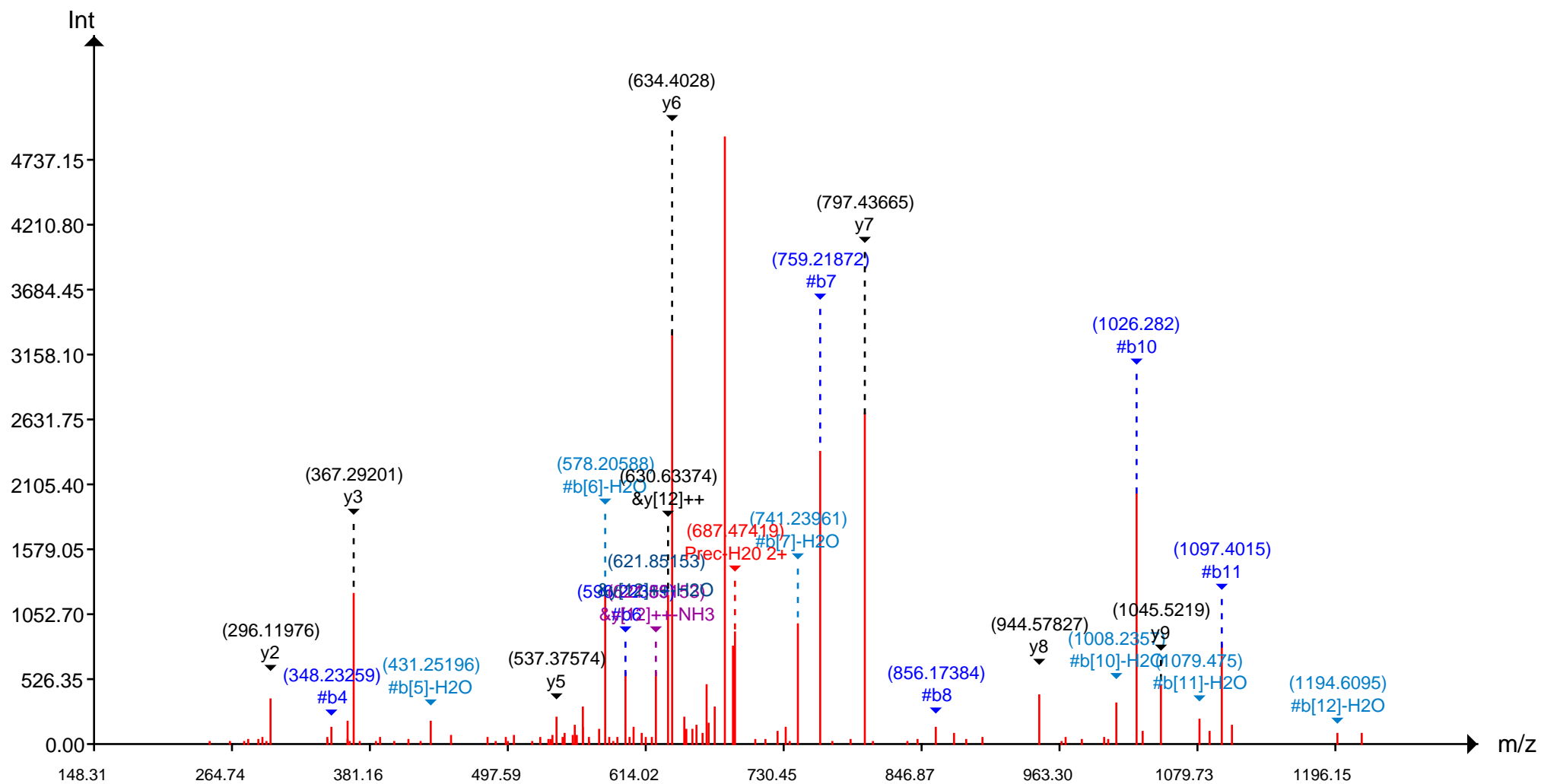
Info

Modified Sequence	AcD3-IK<AcD3K*>ESAK<AcD3K*>VGDAQQR<C13>-COOH	Ion score	94.02
Protein	Q99JY9	Confidence level	99.00%
Precursor m/z	785.94629	Identity Threshold	46.55
Charge	2+	Mass error	0.001227



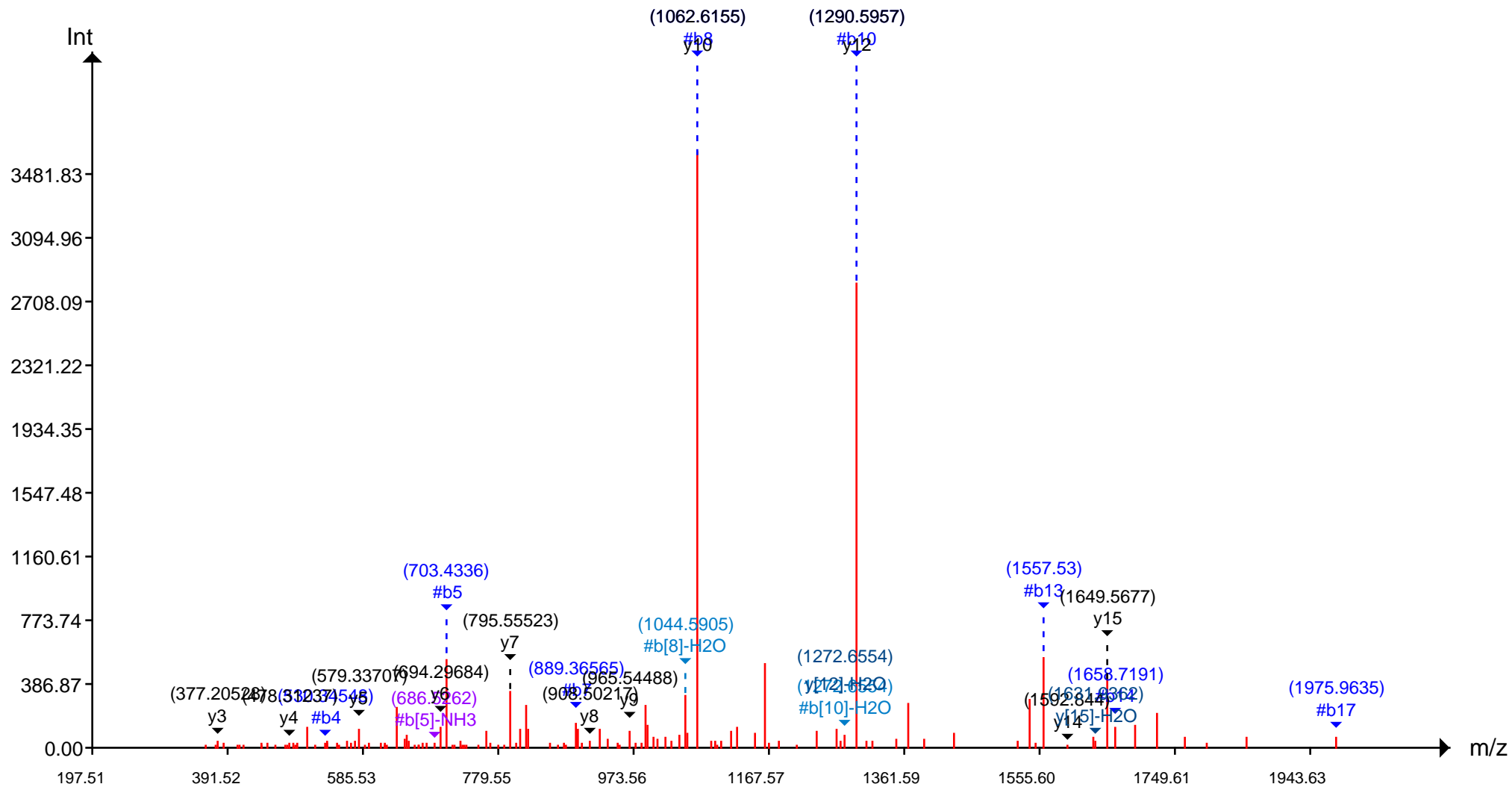
Info

Modified Sequence	AcD3-VEK<AcD3K*>GIIDPTK<AcD3K*>VVR<C13>-COOH	Ion score	55.74
Protein	P63038	Confidence level	99.00%
Precursor m/z	797.99519	Identity Threshold	40.04
Charge	2+	Mass error	0.001081



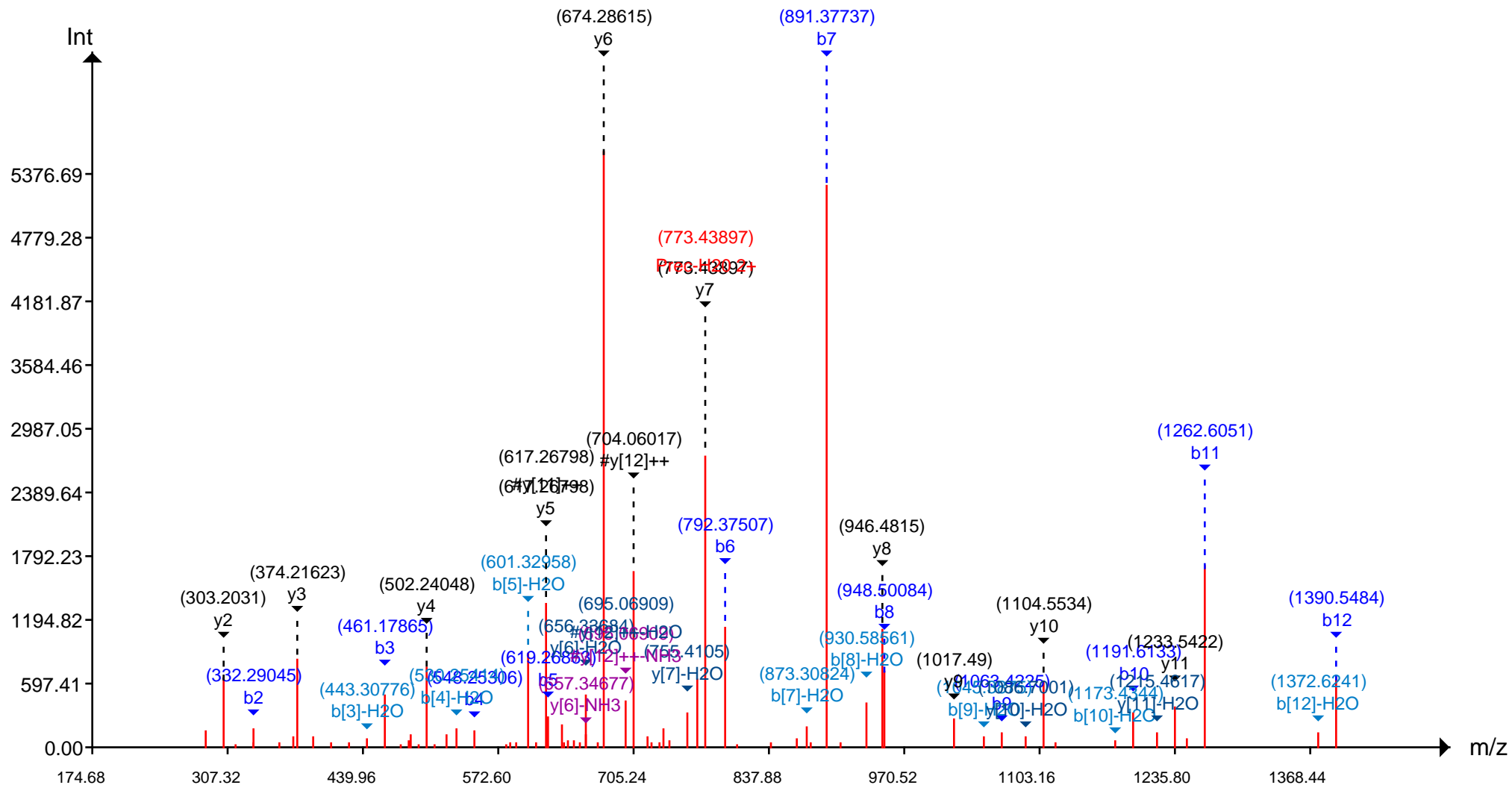
Info

Modified Sequence	AcD3-SGGTTM<Mox*>YPGIADR<C13>-COOH	Ion score	58.25
Protein	P62737, P60710, P68033, P63260, P63268, P68134	Confidence level	99.00%
Precursor m/z	696.83345	Identity Threshold	46.40
Charge	2+	Mass error	-2.55E-4



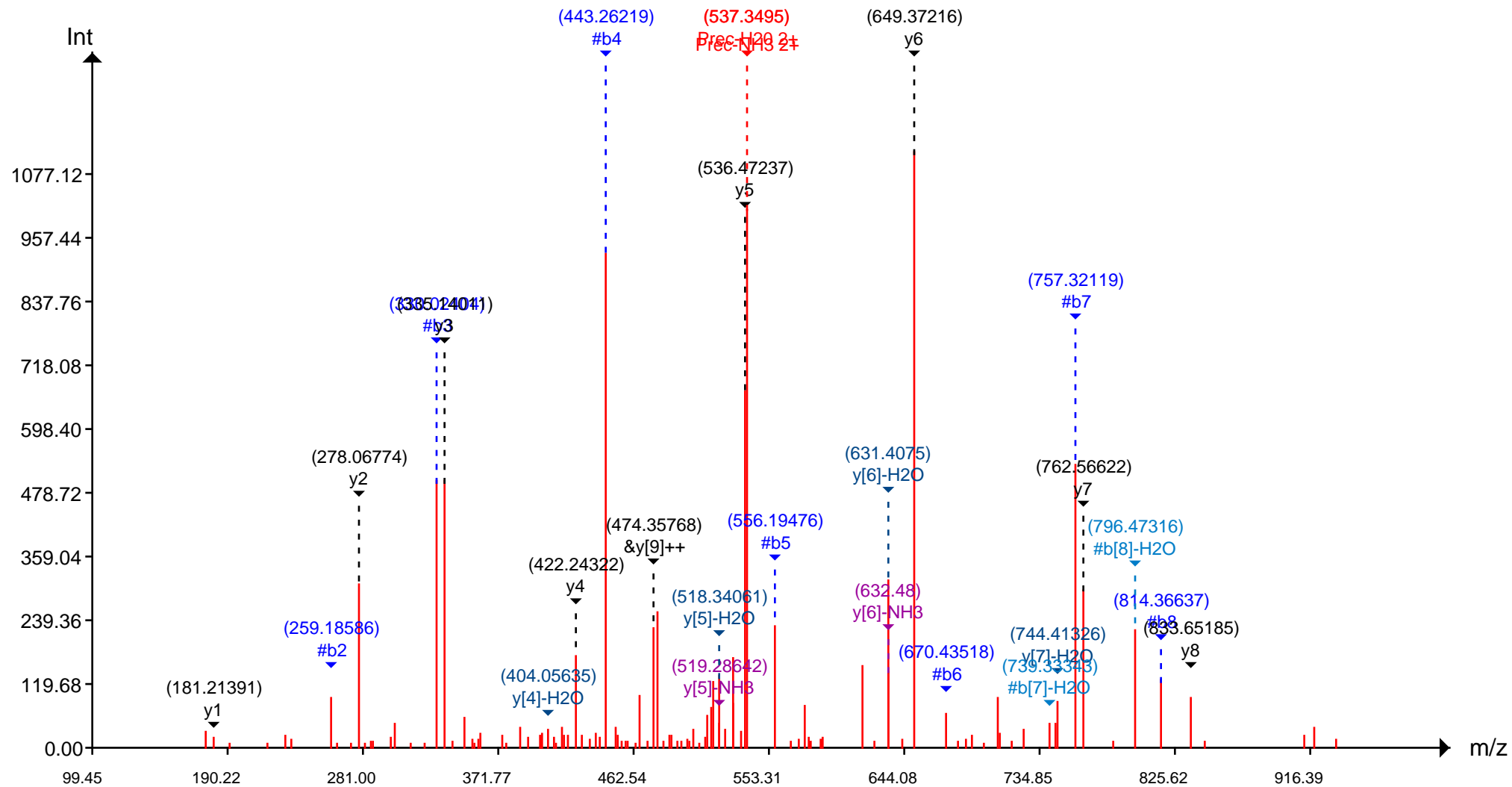
Info

Modified Sequence	AcD3-VIVK<AcD3K*>K<AcD3K*>GEK<AcD3K*>DIPGLTDTTVPR<C13>-COOH	Ion score	44.35
Protein	P62754	Confidence level	99.00%
Precursor m/z	1176.6987	Identity Threshold	40.72
Charge	2+	Mass error	0.003133



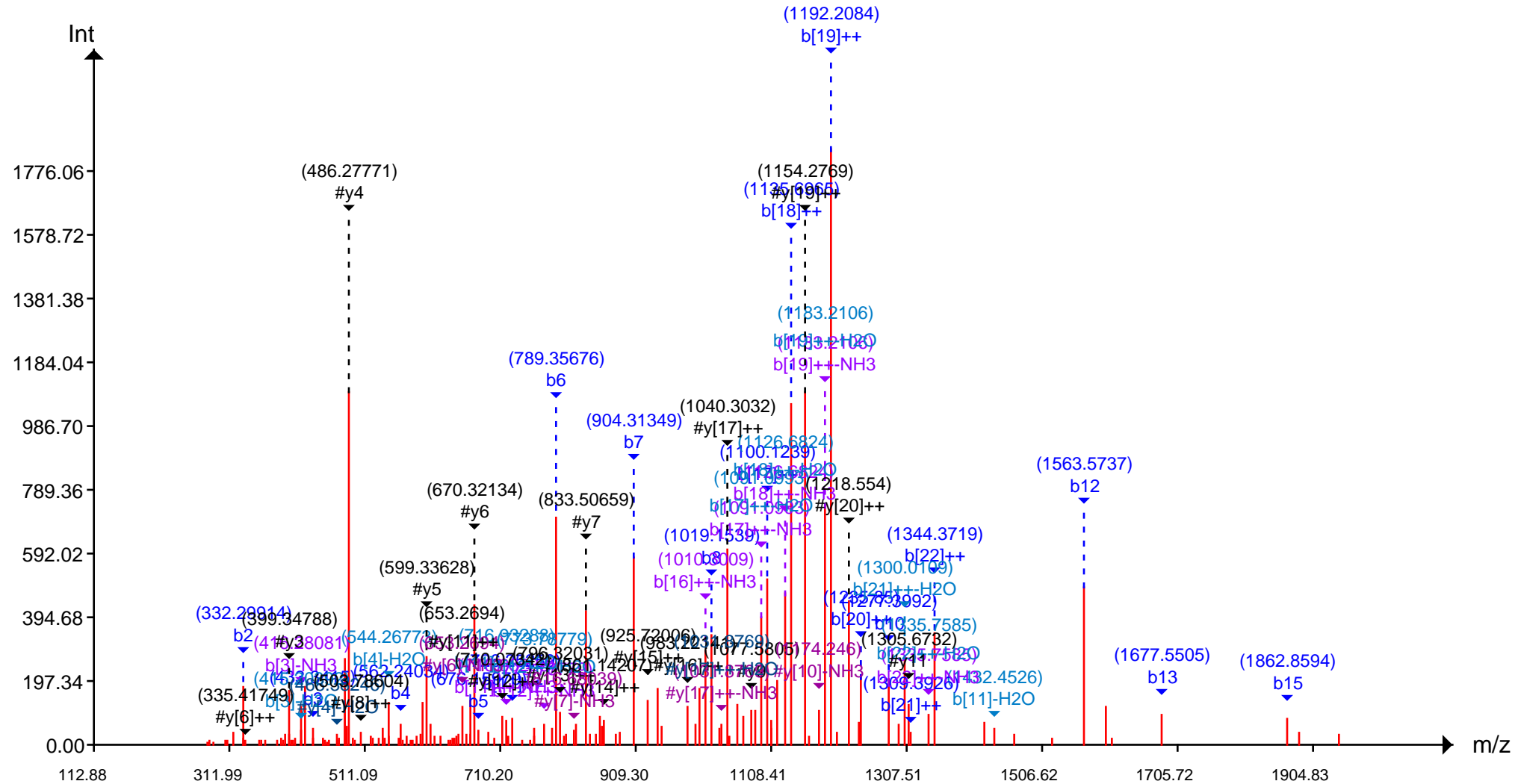
Info

Modified Sequence	AcD3-IK<AcD3K*>ESAK<AcD3K*>VGDQAQR-COOH	Ion score	110.15
Protein	Q99JY9	Confidence level	99.00%
Precursor m/z	782.93627	Identity Threshold	47.37
Charge	2+	Mass error	0.001329



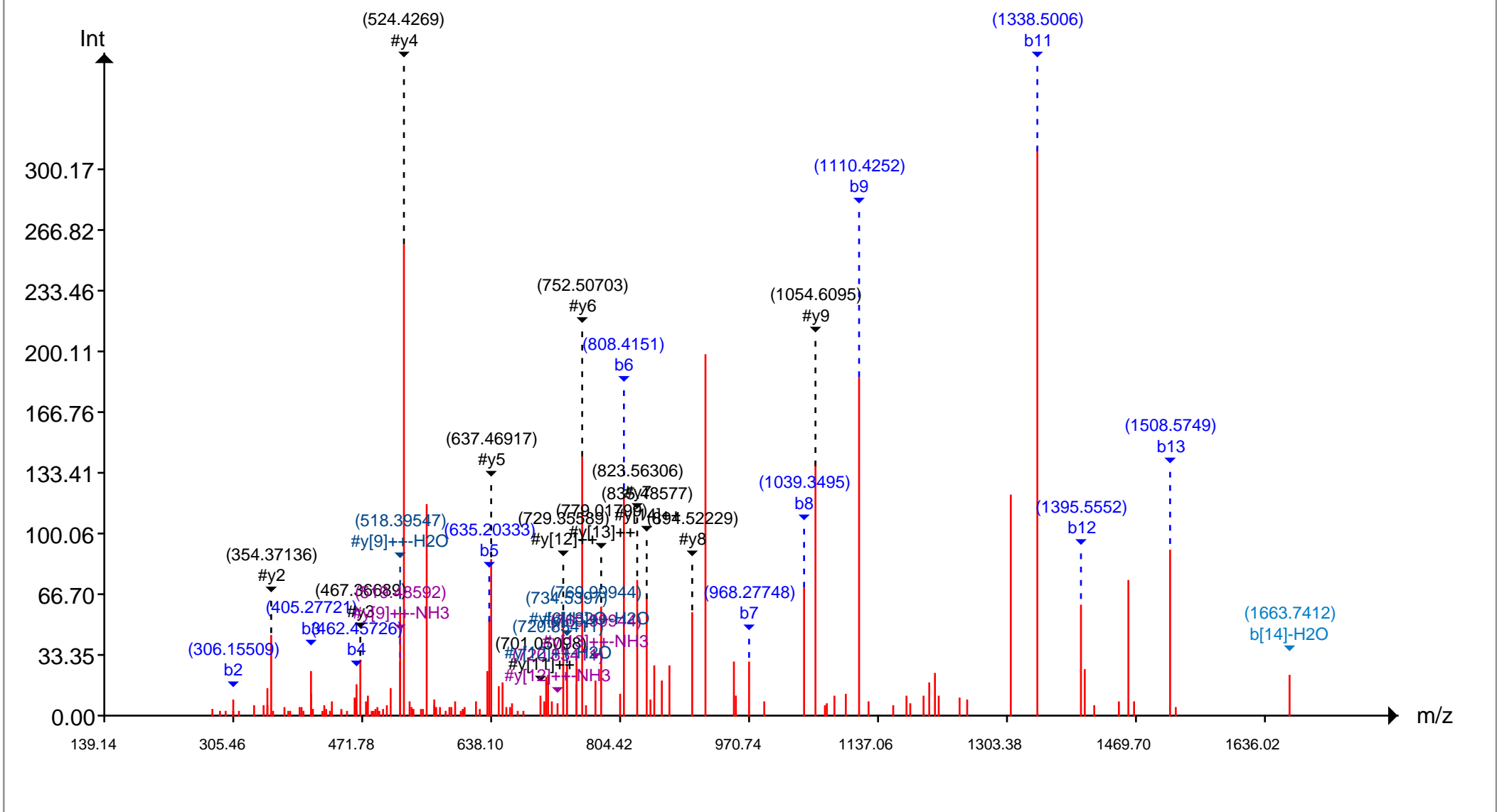
Info

Modified Sequence	AcD3-VNAIINSGPR<C13>-COOH	Ion score	61.93
Protein	P97461	Confidence level	99.00%
Precursor m/z	546.32082	Identity Threshold	46.28
Charge	2+	Mass error	1.35E-4



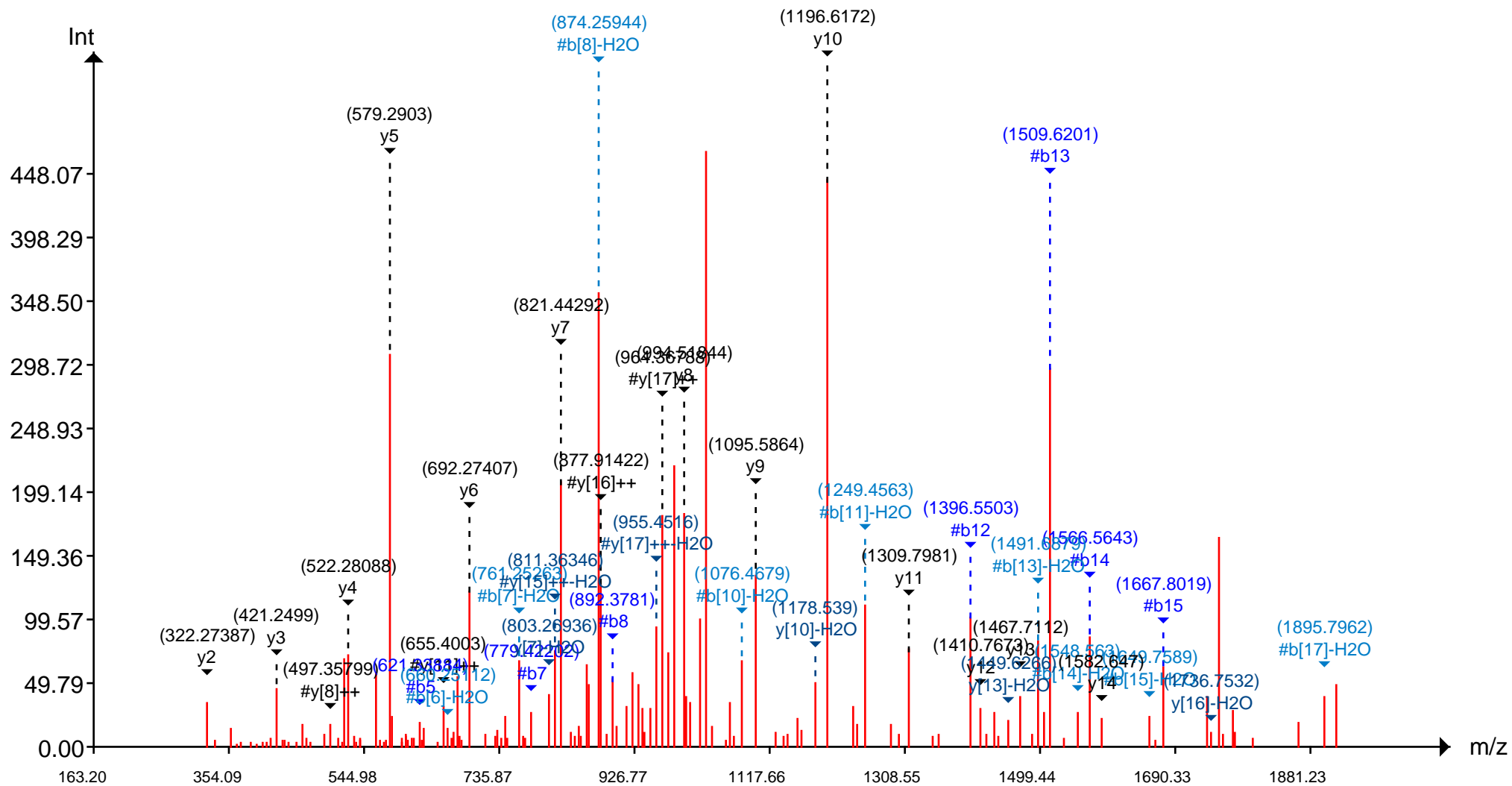
Info

Modified Sequen...	AcD3-LK<AcD3K*>TENLDDEEK<AcD3K*>LNNAK<AcD3K*>YAISM<Mox*>AR<C13>-COOH	Ion score	65.92
Prot...	Q61233	Confidence level	99.00%
Precursor m...	956.83189	Identity Threshold	47.60
Char...	3+	Mass error	0.003139



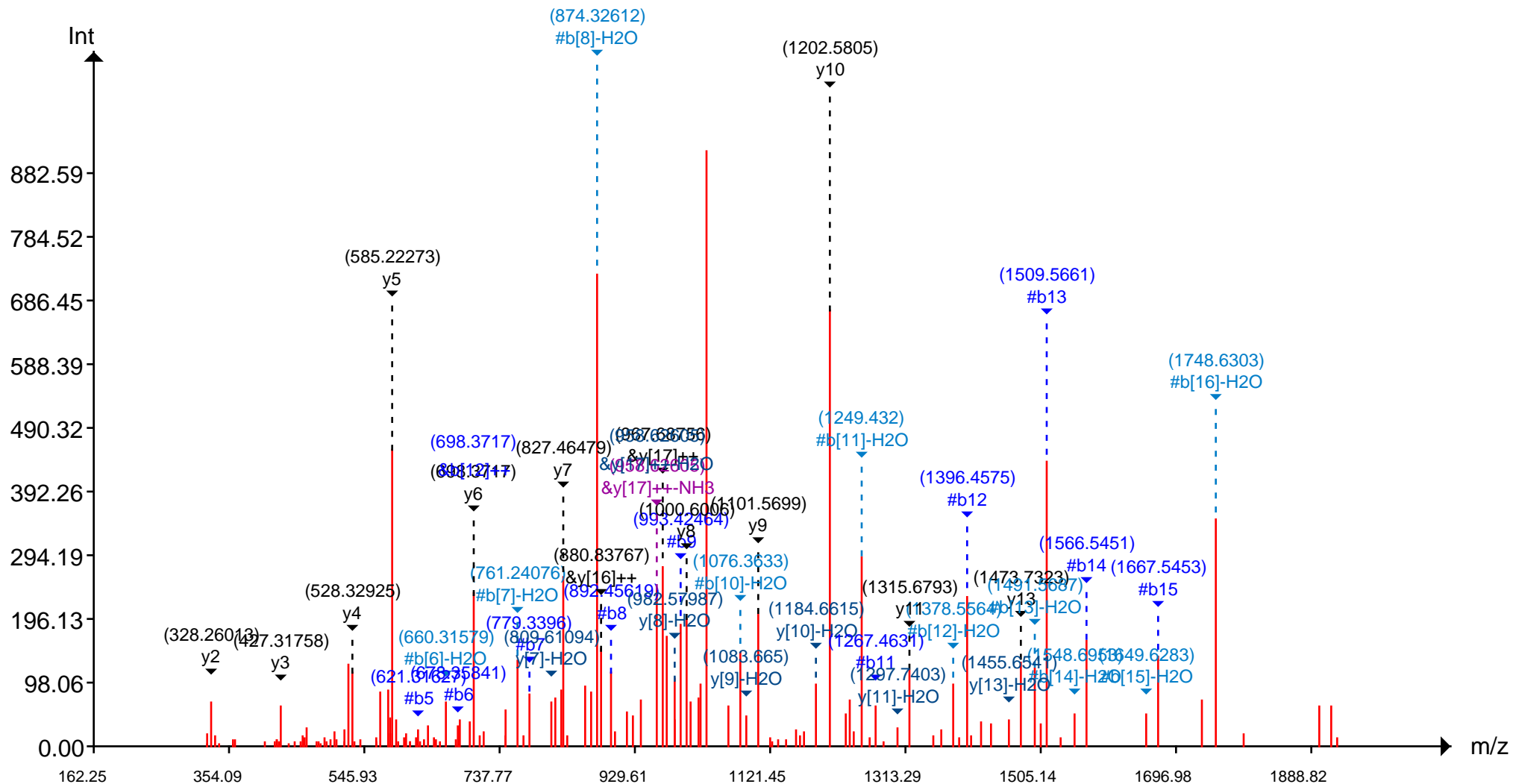
Info

Modified Sequence	AcD3-M<Mox*>IVGK<AcD3K*>K<AcD3K*>C<Cmm*>AADLGLK<AcD3K*>R<C13>-COOH	Ion score	78.56
Protein	P70349	Confidence level	99.00%
Precursor m/z	931.54046	Identity Threshold	44.74
Charge	2+	Mass error	0.001335



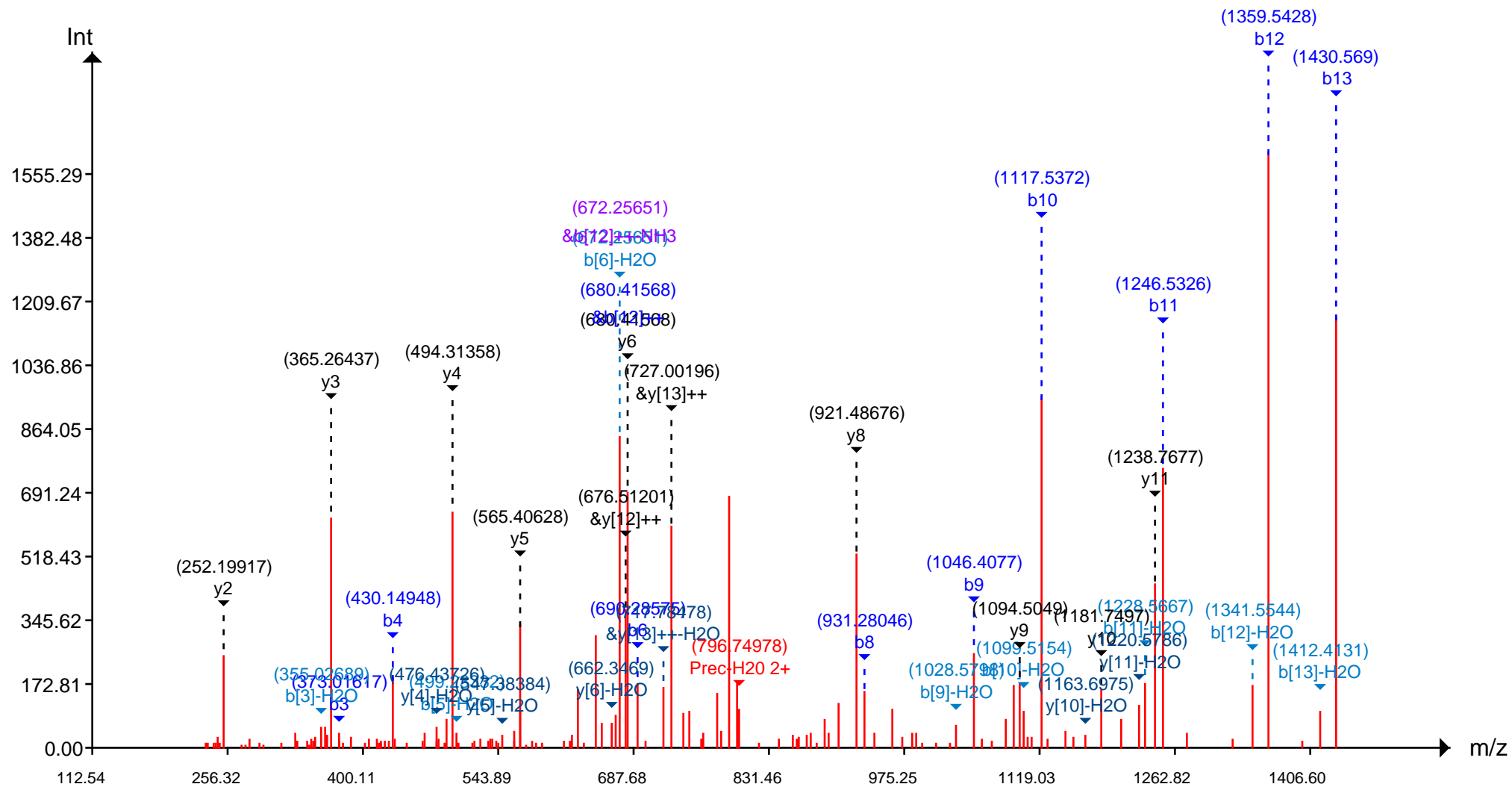
Info

Modified Sequence	AcD3-DK<AcD3K*>DGDGTITTK<AcD3K*>ELGTVM<Mox*>R-COOH	Ion score	98.04
Protein	P62204	Confidence level	99.00%
Precursor m/z	1044.528	Identity Threshold	47.75
Charge	2+	Mass error	0.001165



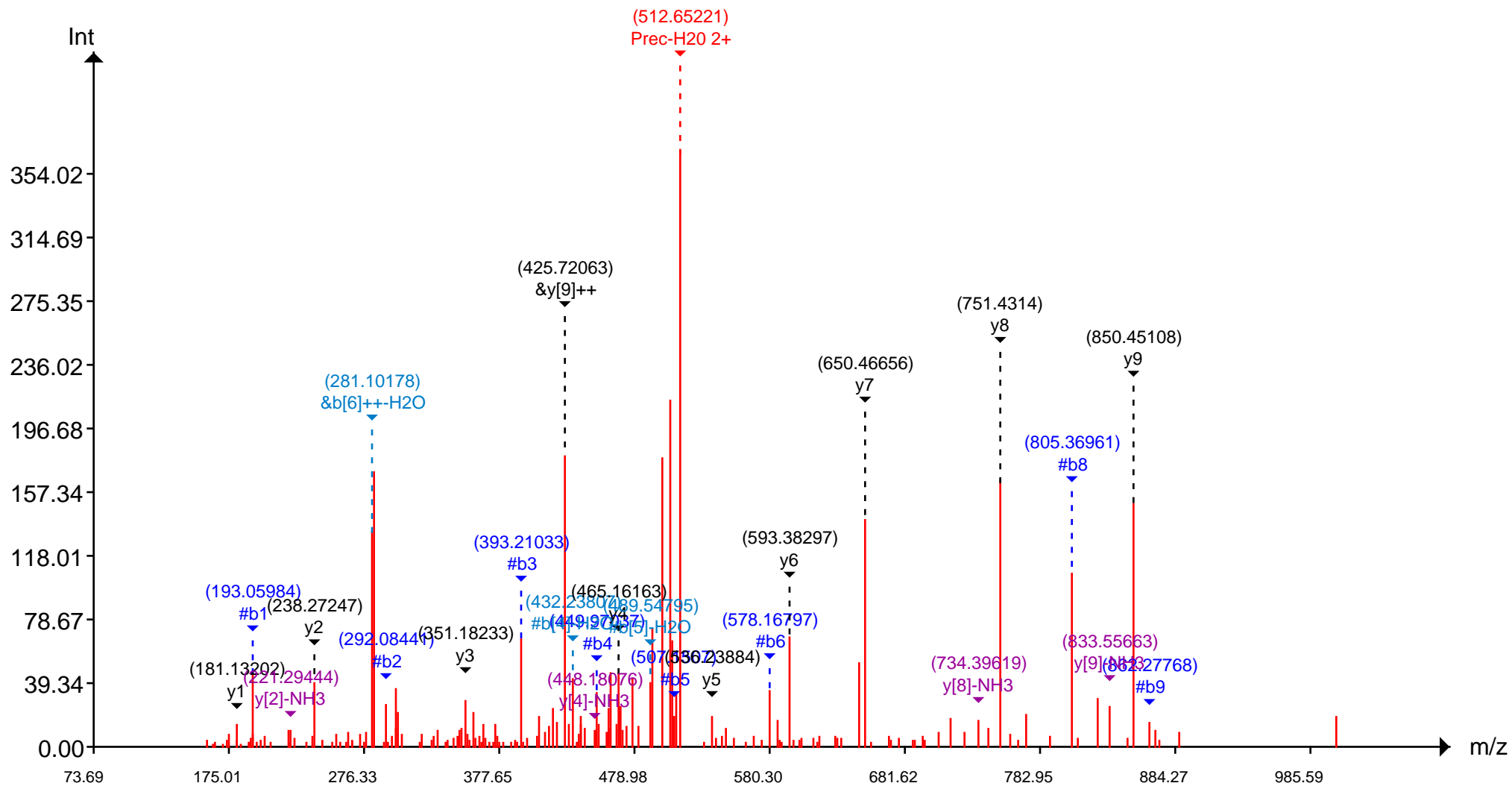
Info

Modified Sequence	AcD3-DK<AcD3K*>DGDGTITTK<AcD3K*>ELGTVM<Mox*>R<C13>-COOH	Ion score	80.53
Protein	P62204	Confidence level	99.00%
Precursor m/z	1047.5386	Identity Threshold	47.98
Charge	2+	Mass error	0.002223



Info

Modified Sequence	AcD3-LTIGSK<AcD3K*>LQDAEIAR<C13>-COOH	Ion score	67.59
Protein	P14069	Confidence level	99.00%
Precursor m/z	805.97067	Identity Threshold	45.07
Charge	2+	Mass error	0.00148



Info

Modified Sequence AcD3-M<Mox*>VTGGANLGR<C13>-COOH

Ion score 47.77

Protein P62702

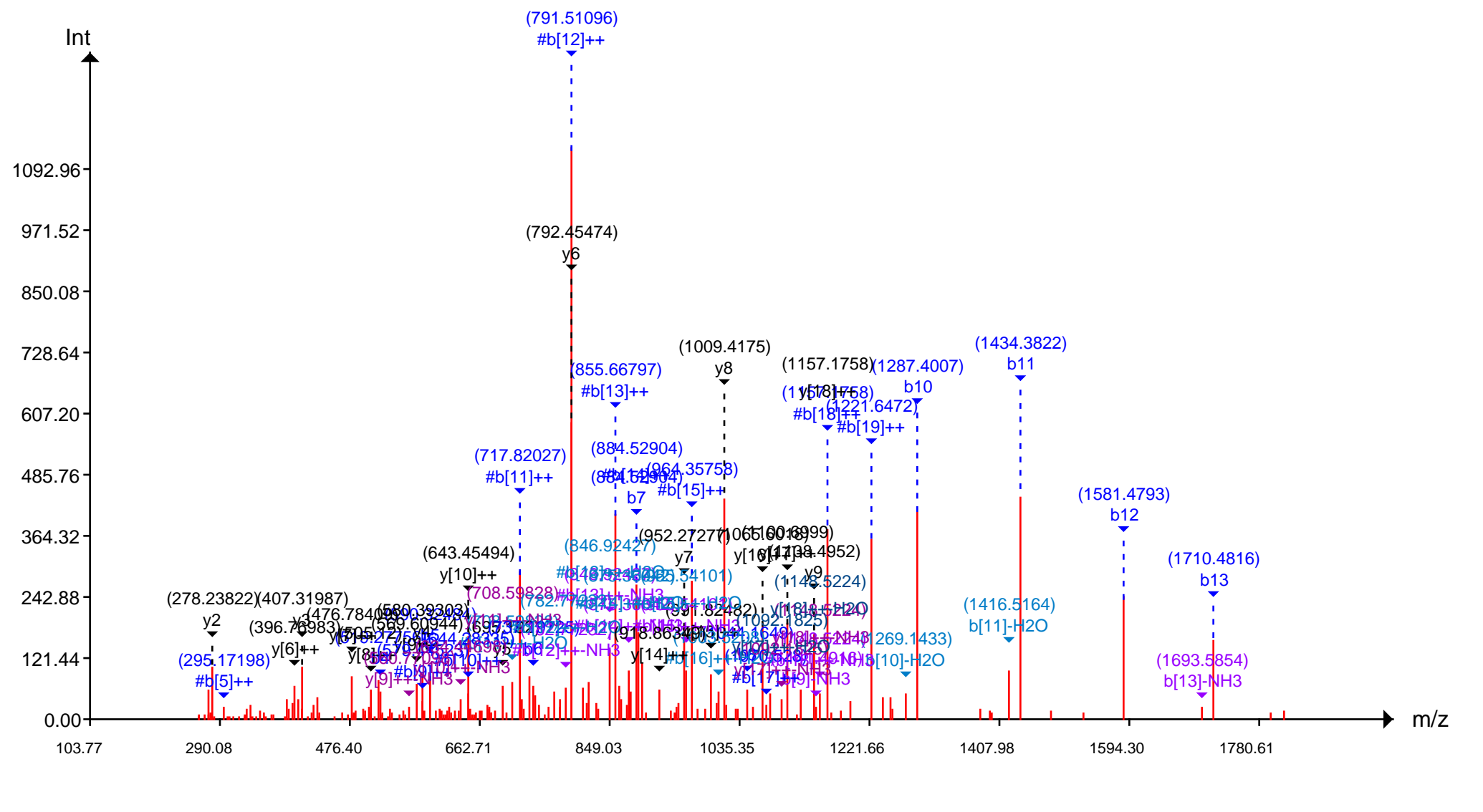
Confidence level 99.00%

Precursor m/z 521.77814

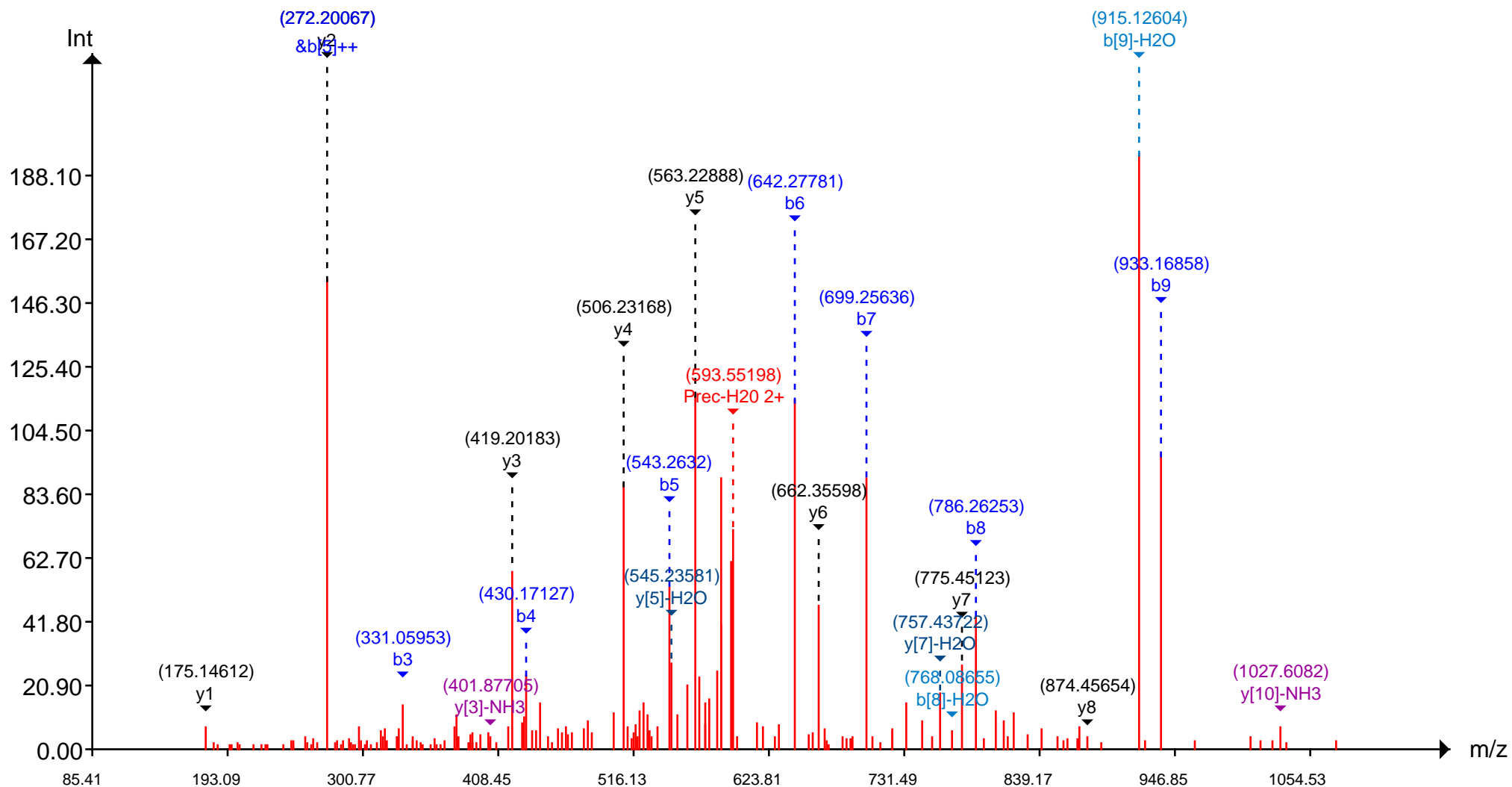
Identity Threshold 47.59

Charge 2+

Mass error 5.29E-4



Info			
Modified S...	AcD3-LSC<Cmm*>IAM<Mox*>M<Mox*>C<Cmm*>NEFFEGC<Cmm*>PDK<AcD3K*>EPR<C13>-COOH	Ion score	59
	P07091	Confidence level	99
Precu...	907.0569	Identity Threshold	39
	3+	Mass error	0.0



Info

Modified Sequence AcD3-DVAVLVGSM<Mox*>PR-COOH

Ion score 57.19

Protein P14152

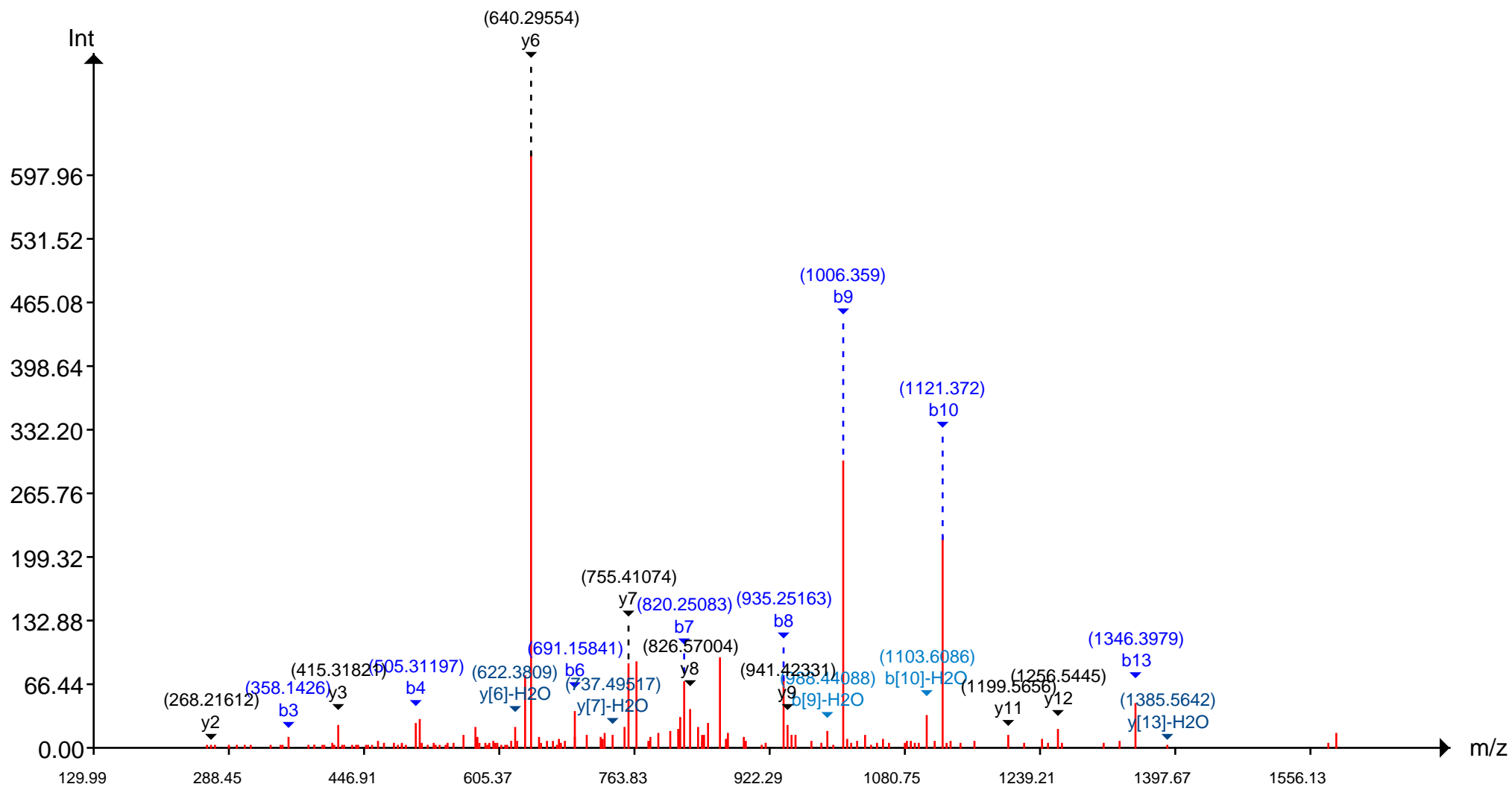
Confidence level 99.00%

Precursor m/z 602.82549

Identity Threshold 48.22

Charge 2+

Mass error 3.35E-4



Info

Modified Sequence AcD3-ALQFG EEDADPGAM<Mox*>SR<C13>-COOH

Ion score 59.47

Protein Q9Z2U1

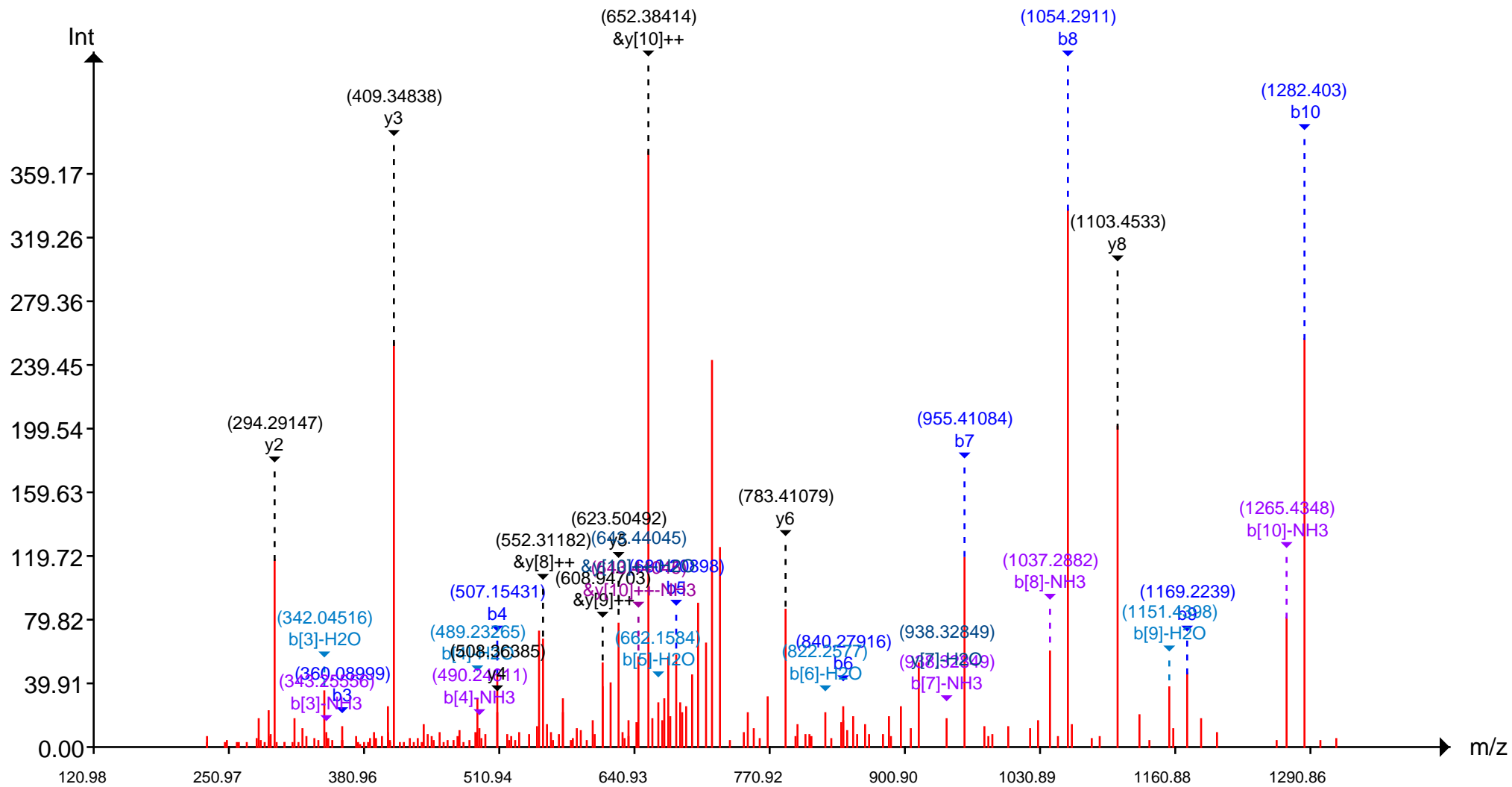
Confidence level 99.00%

Precursor m/z 880.90138

Identity Threshold 44.26

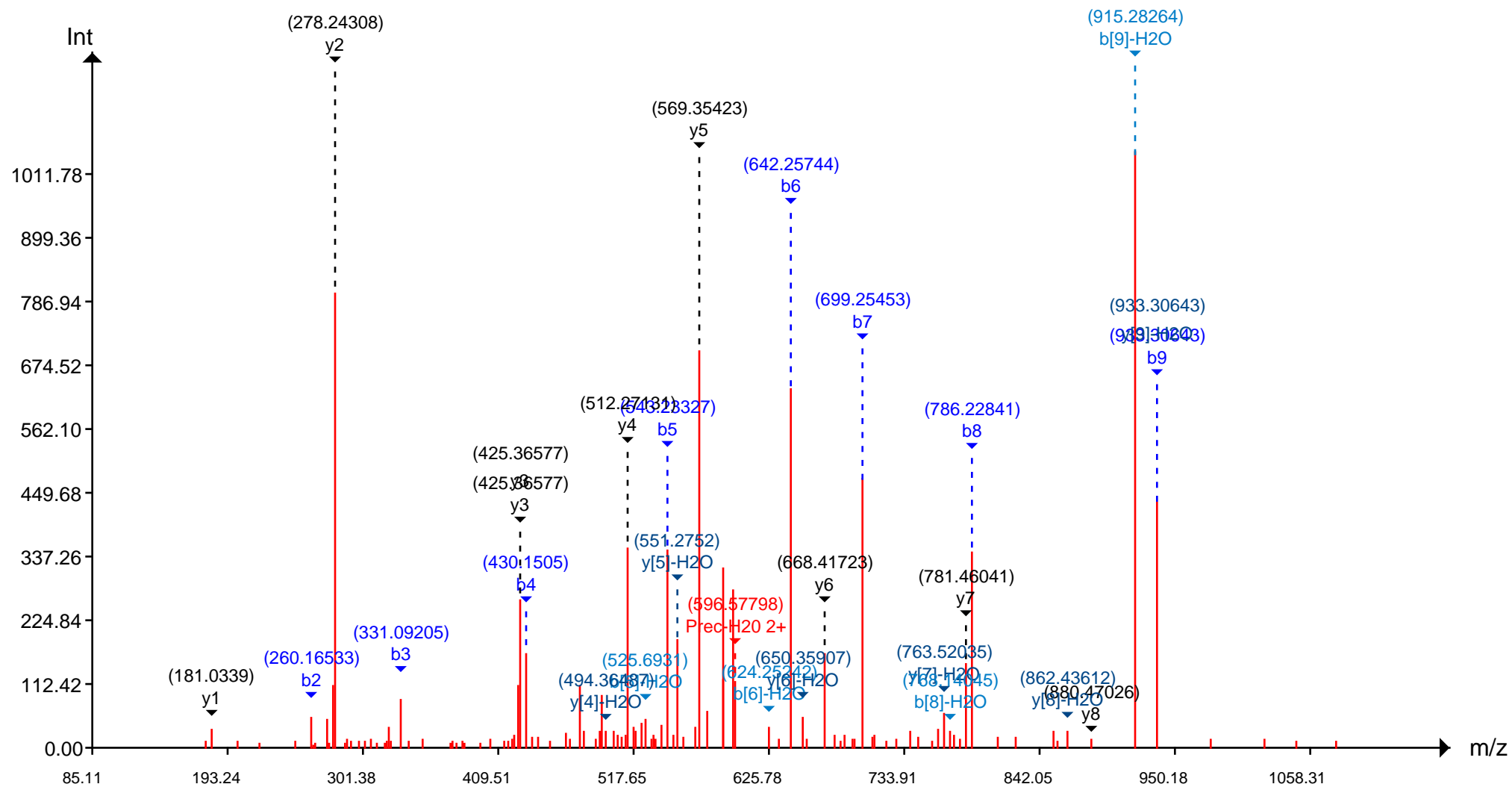
Charge 2+

Mass error 0.002426



Info

Modified Sequence	AcD3-NSIM<Mox*>K<AcD3K*>C<Cmm*>DVIDIR<C13>-COOH	Ion score	49.52
Protein	Q8BFZ3, P60710, P63260	Confidence level	99.00%
Precursor m/z	731.86637	Identity Threshold	47.59
Charge	2+	Mass error	0.001193



Info

Modified Sequence AcD3-DVAVLVGSM<Mox*>PR<C13>-COOH

Ion score 62.47

Protein P14152

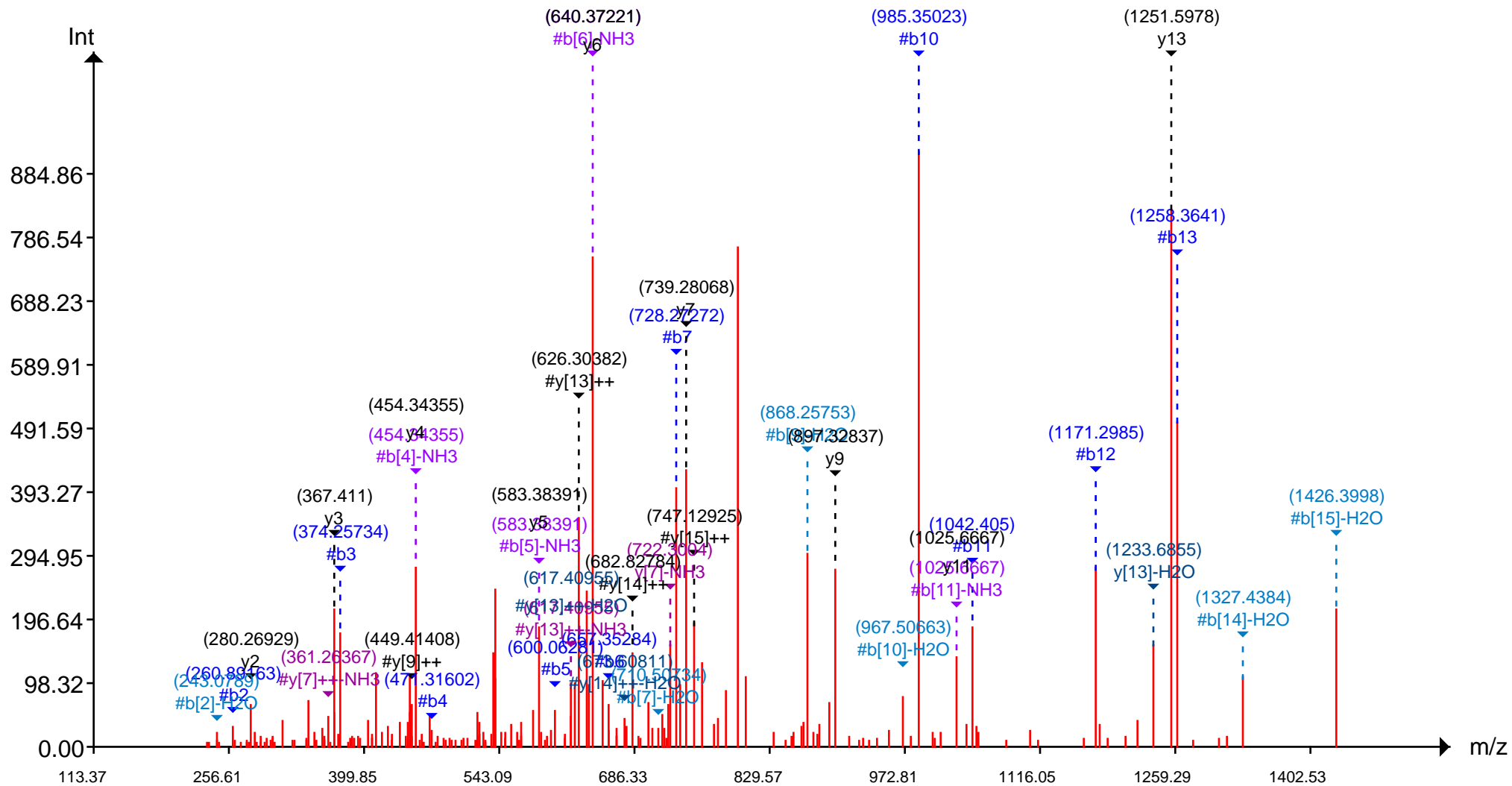
Confidence level 99.00%

Precursor m/z 605.83552

Identity Threshold 47.52

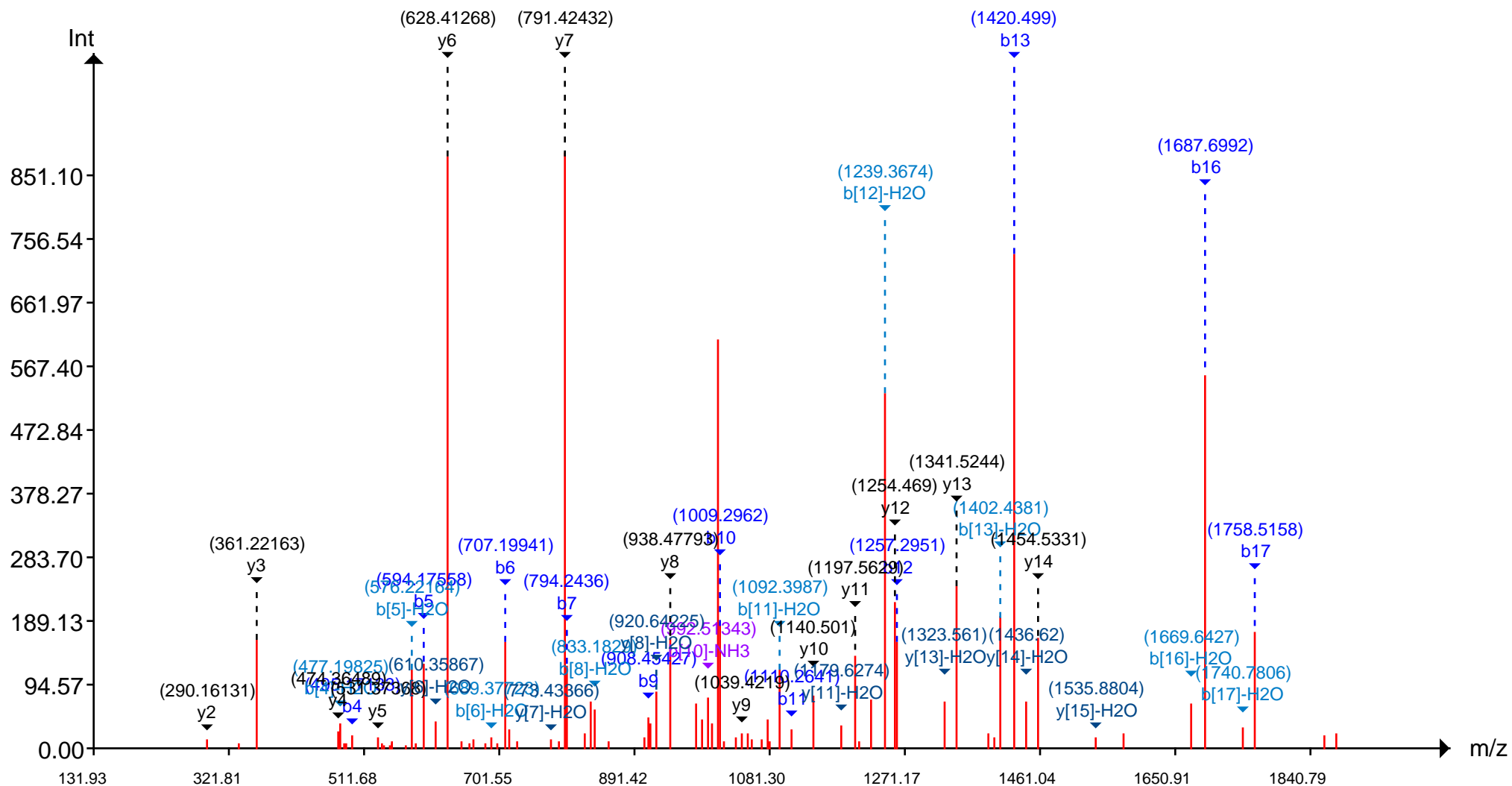
Charge 2+

Mass error 2.53E-4



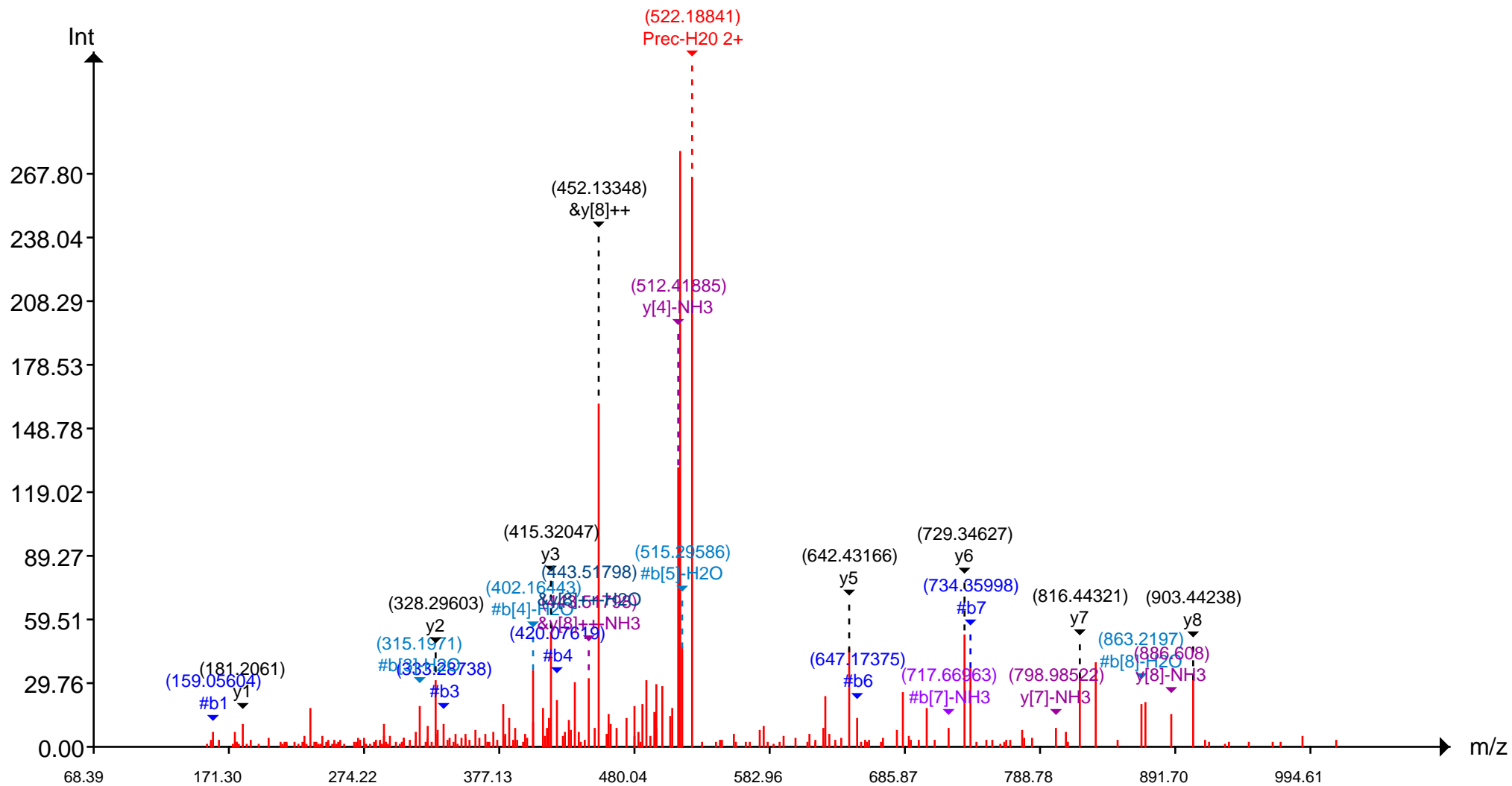
Info

Modified Sequence	AcD3-SQLPEGASAVGESSVR<C13>-COOH	Ion score	65.01
Protein	Q921W0	Confidence level	99.00%
Precursor m/z	812.9198	Identity Threshold	48.01
Charge	2+	Mass error	0.001165



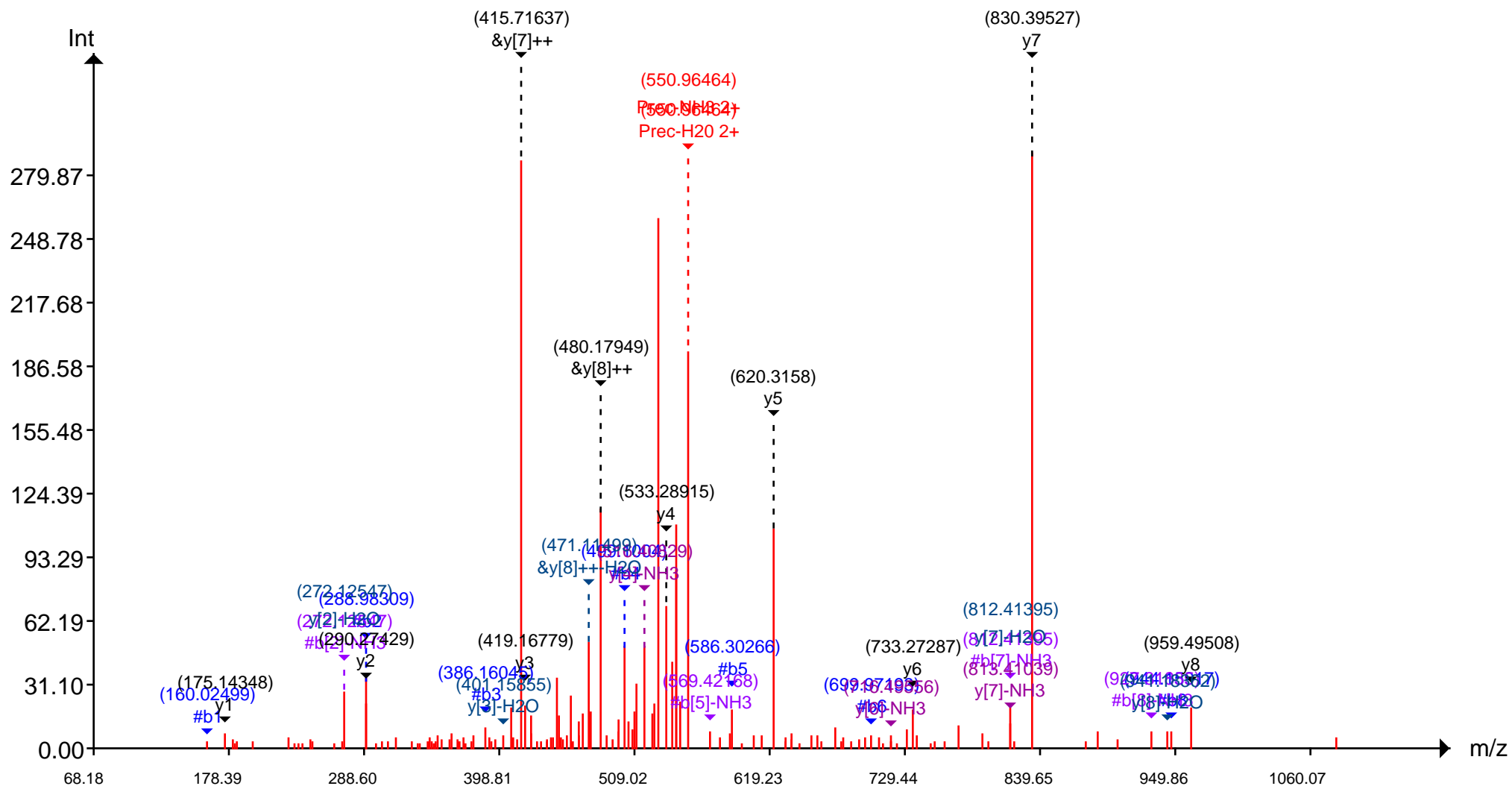
Info

Modified Sequence	AcD3-YANTVLSGGTTM<Mox*>YPGIADR-COOH	Ion score	98.86
Protein	P60710, P63260	Confidence level	99.00%
Precursor m/z	1024.4963	Identity Threshold	47.01
Charge	2+	Mass error	0.002065



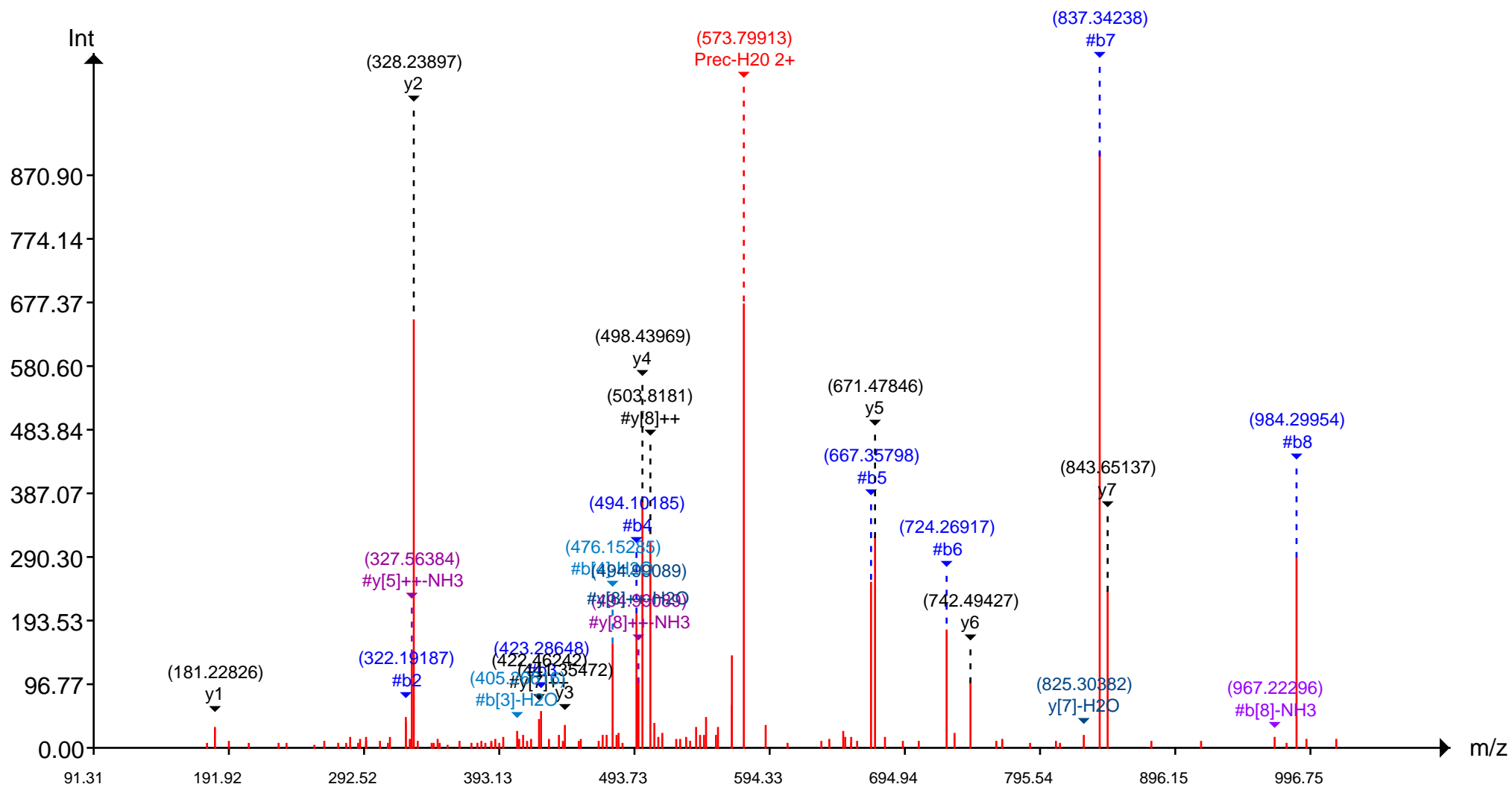
Info

Modified Sequence	AcD3-ISSSLNSFR<C13>-COOH	Ion score	51.52
Protein	Q9D1E6	Confidence level	99.00%
Precursor m/z	531.29188	Identity Threshold	48.14
Charge	2+	Mass error	4.37E-4



Info

Modified Sequence	AcD3-NEPLSNEDR-COOH	Ion score	46.16
Protein	P68510	Confidence level	99.00%
Precursor m/z	559.76111	Identity Threshold	45.49
Charge	2+	Mass error	-2.06E-4



Info

Modified Sequence AcD3-LYTAK<AcD3K*>GLFR<C13>-COOH

Ion score 52.18

Protein P17182, P17183

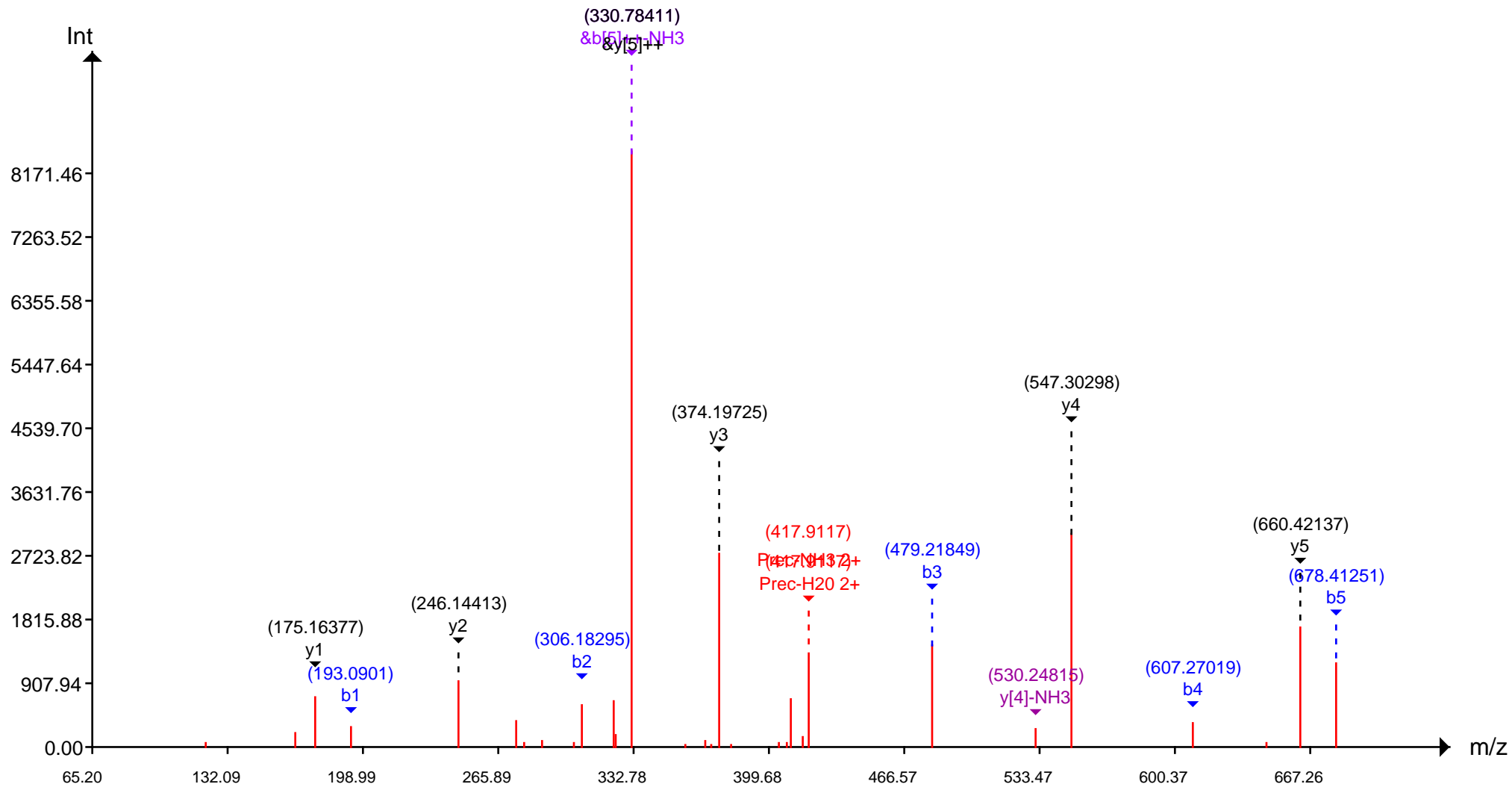
Confidence level 99.00%

Precursor m/z 582.85319

Identity Threshold 44.25

Charge 2+

Mass error 1.47E-4



Info

Modified Sequence AcD3-FLK<AcD3K*>QAR-COOH

Ion score 58.31

Protein P10107

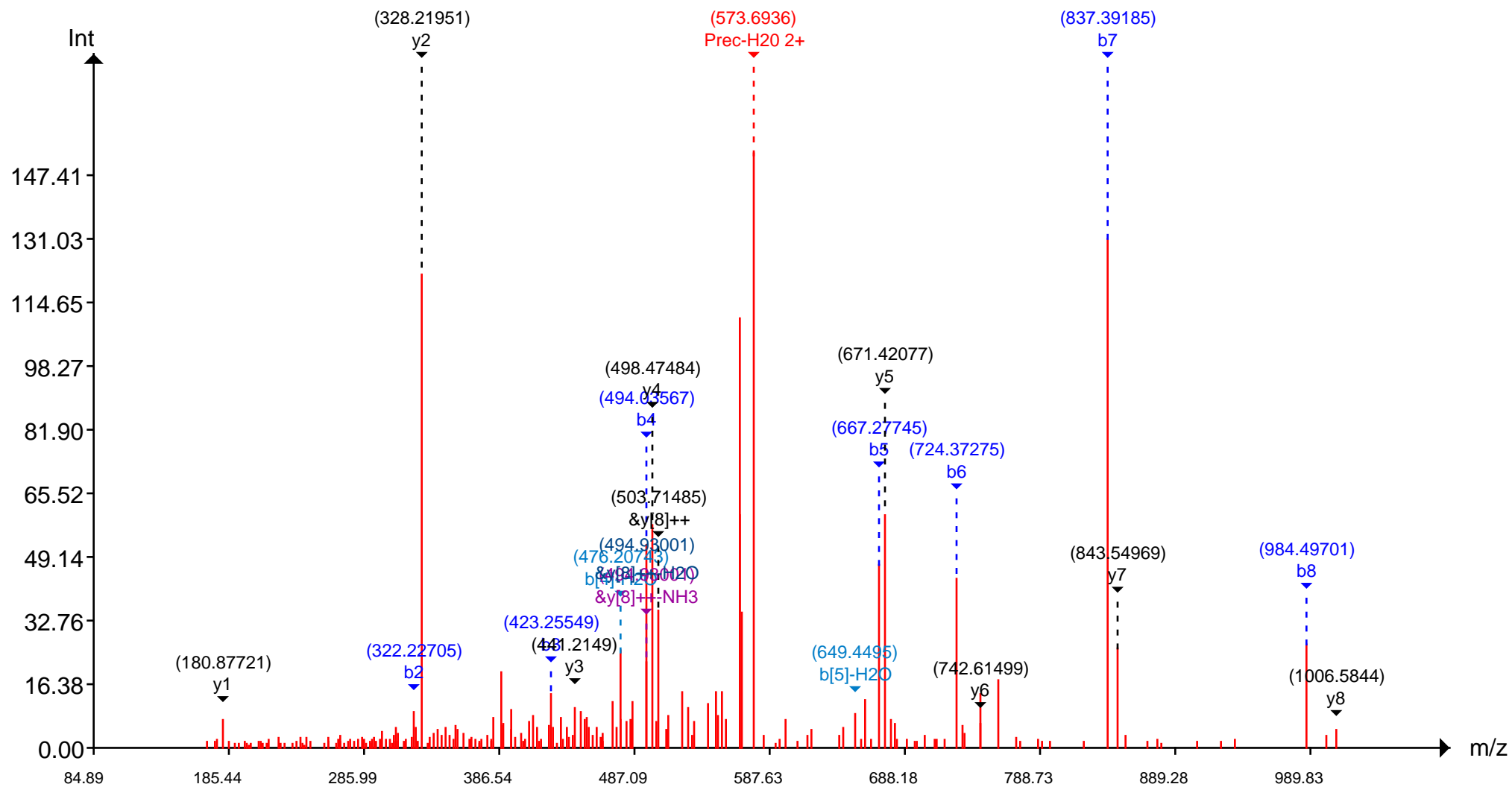
Confidence level 99.00%

Precursor m/z 426.76296

Identity Threshold 44.30

Charge 2+

Mass error -0.002212



Info

Modified Sequence AcD3-LYTAK<AcD3K*>GLFR<C13>-COOH

Ion score 46.63

Protein P17182, P17183

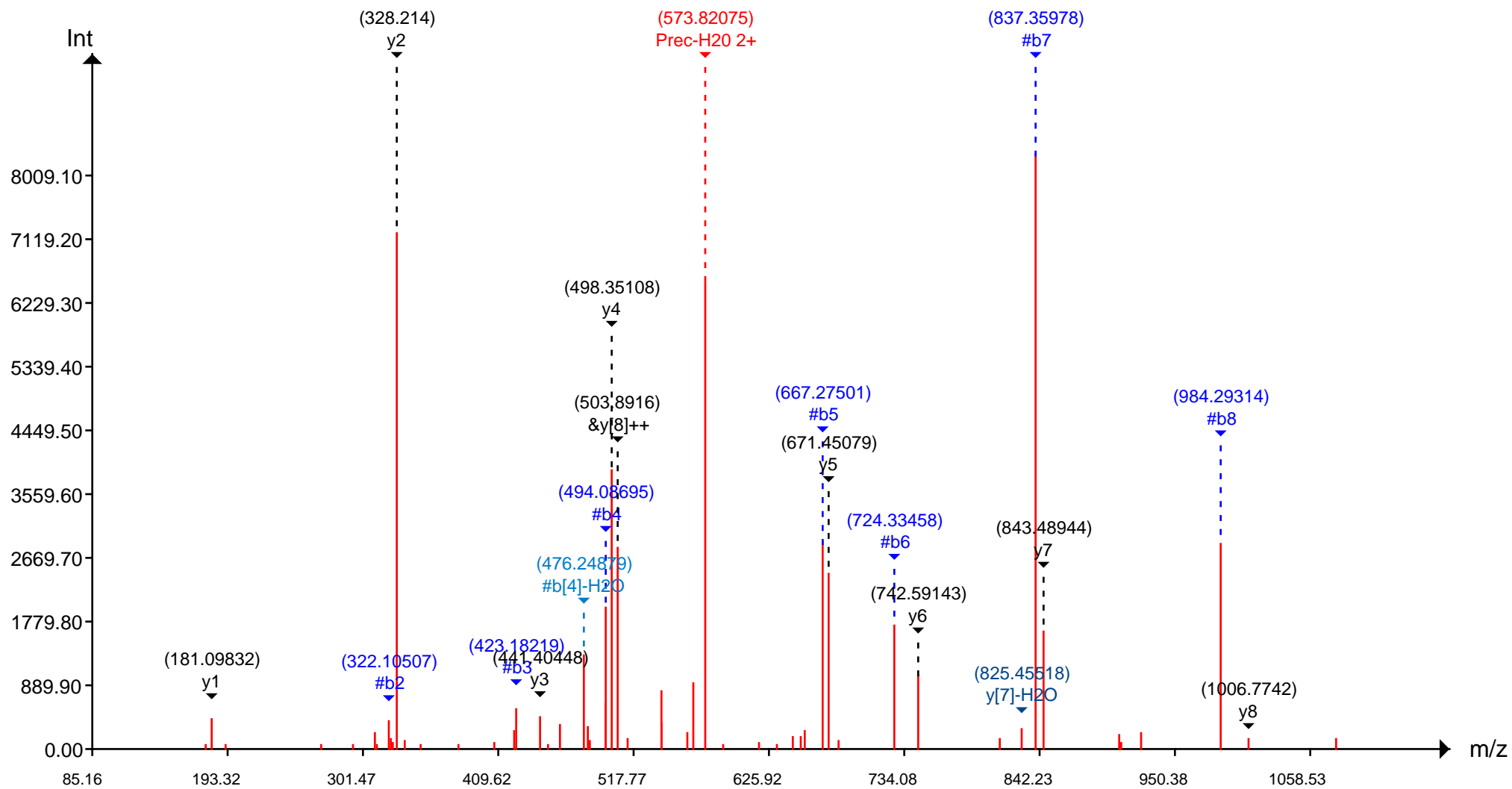
Confidence level 99.00%

Precursor m/z 582.85344

Identity Threshold 44.24

Charge 2+

Mass error 6.47E-4



Info

Modified Sequence AcD3-LYTAK<AcD3K*>GLFR<C13>-COOH

Ion score 71.43

Protein P17182, P17183

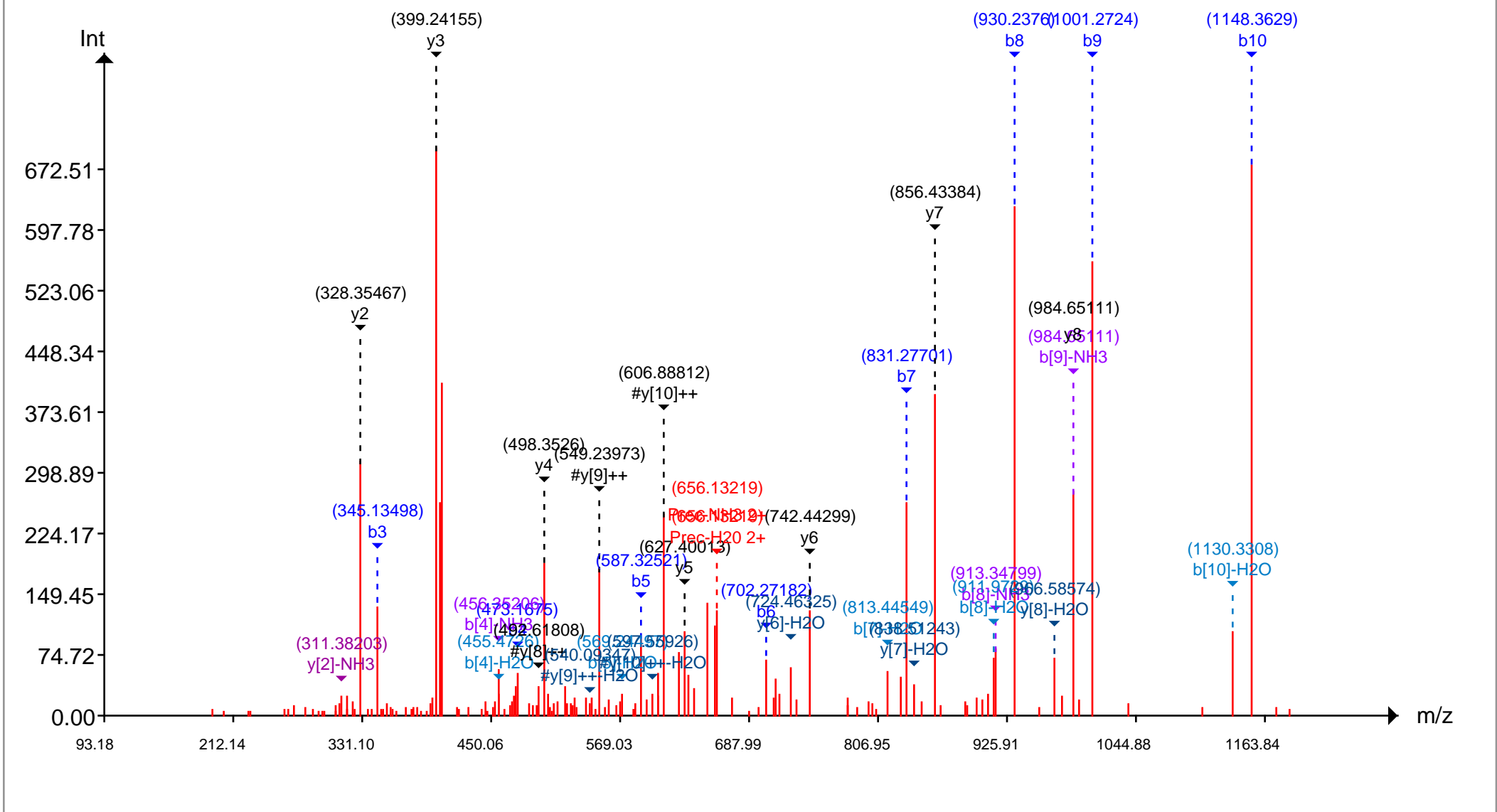
Confidence level 99.00%

Precursor m/z 582.8532

Identity Threshold 44.23

Charge 2+

Mass error 1.67E-4



Info

Modified Sequence AcD3-ADLQNDEVAFR<C13>-COOH

Ion score 58.32

Protein P97351

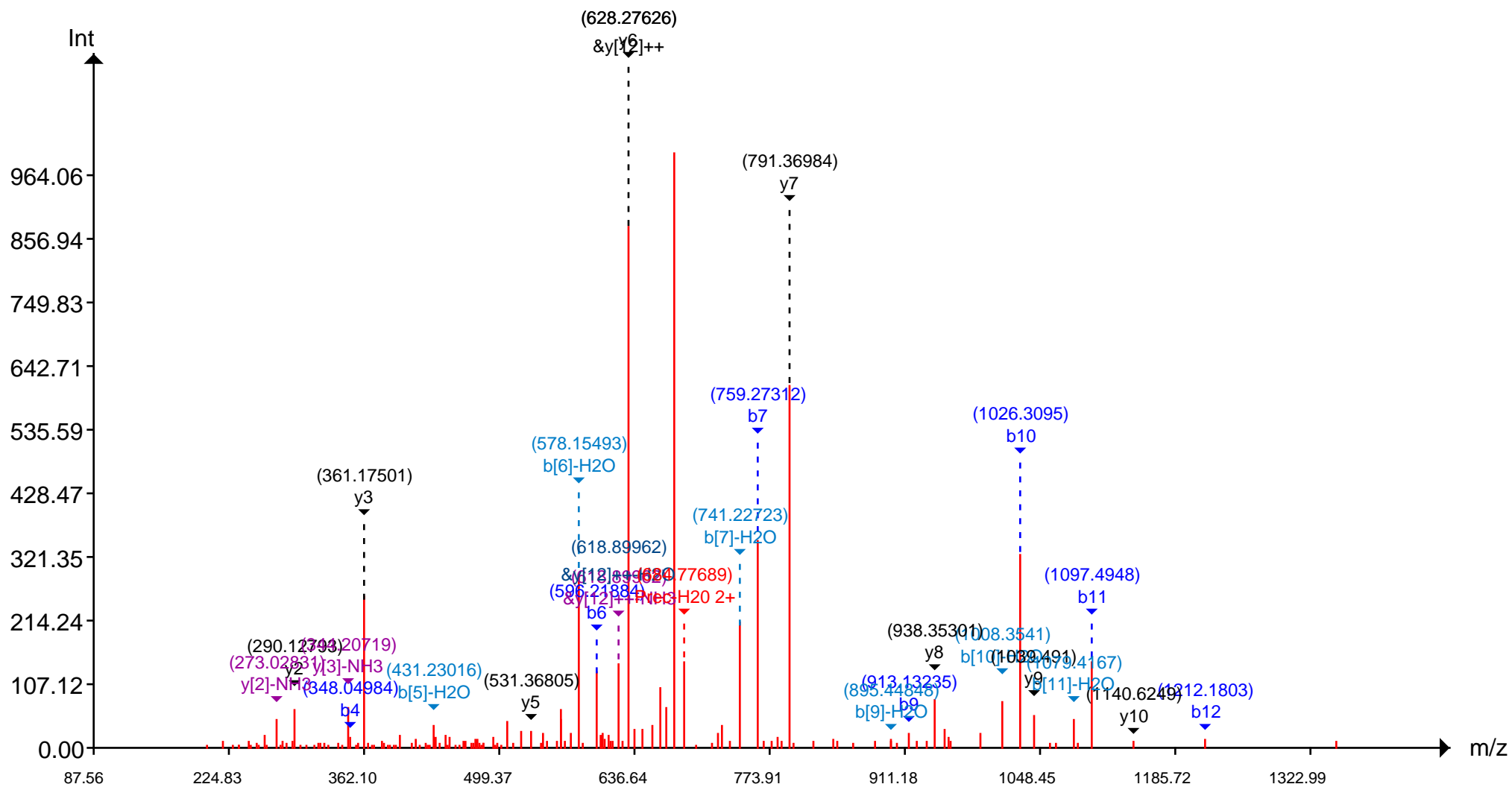
Confidence level 99.00%

Precursor m/z 664.83471

Identity Threshold 47.47

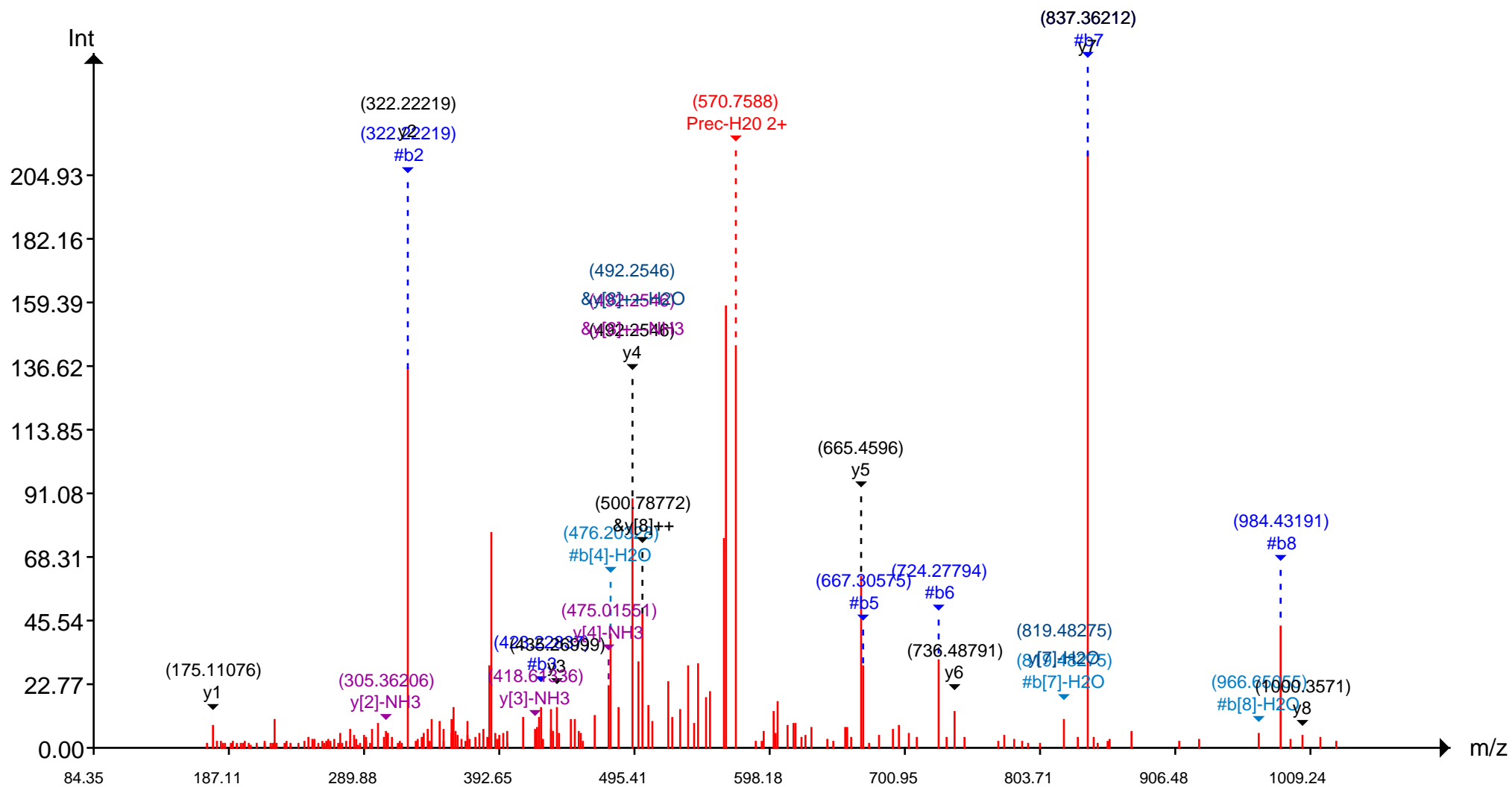
Charge 2+

Mass error 5.71E-4



Info

Modified Sequence	AcD3-SGGTTM<Mox*>YPGIADR-COOH	Ion score	50.38
Protein	P62737, P60710, P68033, P63260, P63268, P68134	Confidence level	99.00%
Precursor m/z	693.82336	Identity Threshold	45.32
Charge	2+	Mass error	-2.94E-4



Info

Modified Sequence AcD3-LYTAK<AcD3K*>GLFR-COOH

Ion score 50.86

Protein P17182, P17183

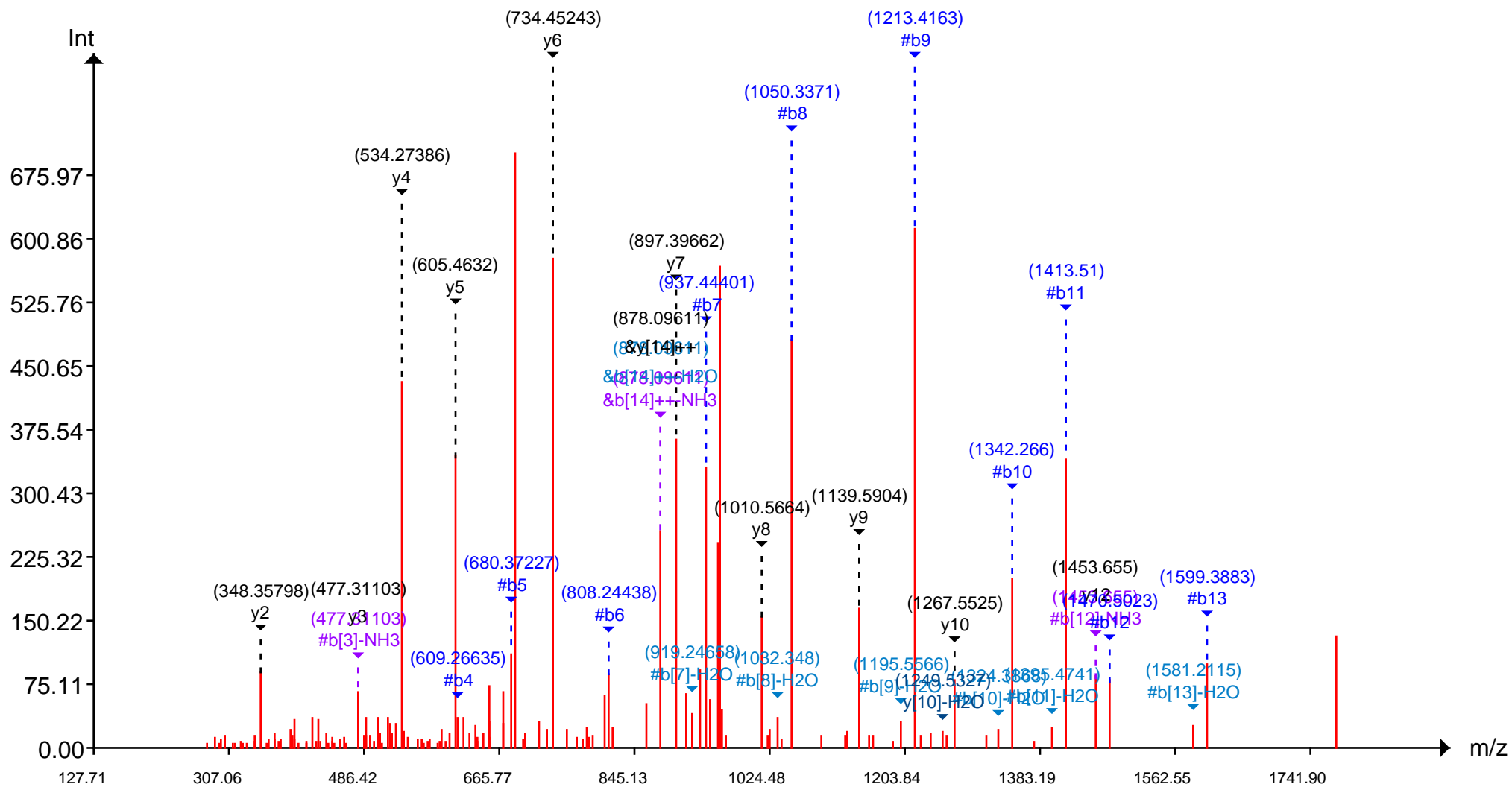
Confidence level 99.00%

Precursor m/z 579.8431

Identity Threshold 45.95

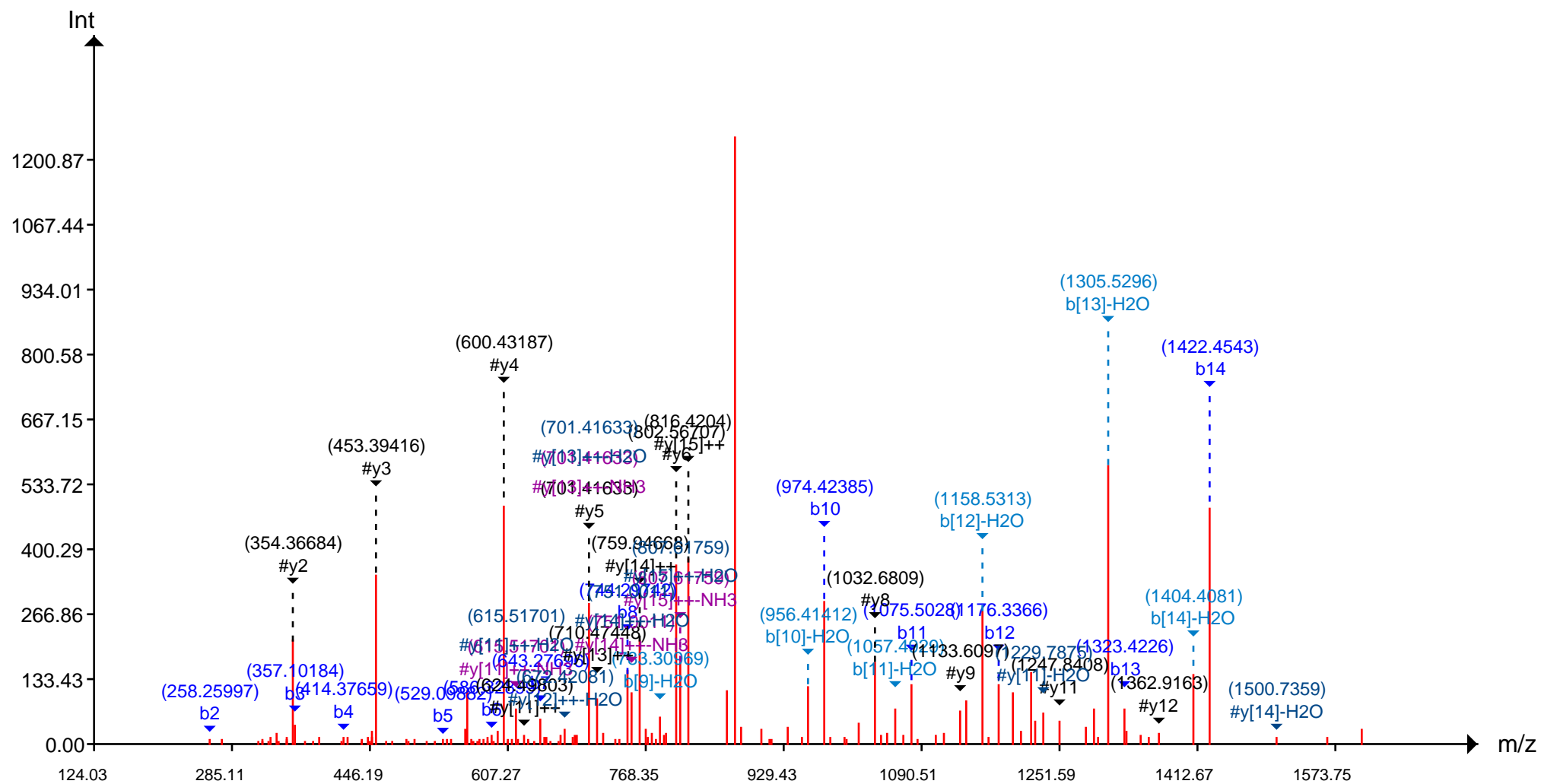
Charge 2+

Mass error 1.09E-4



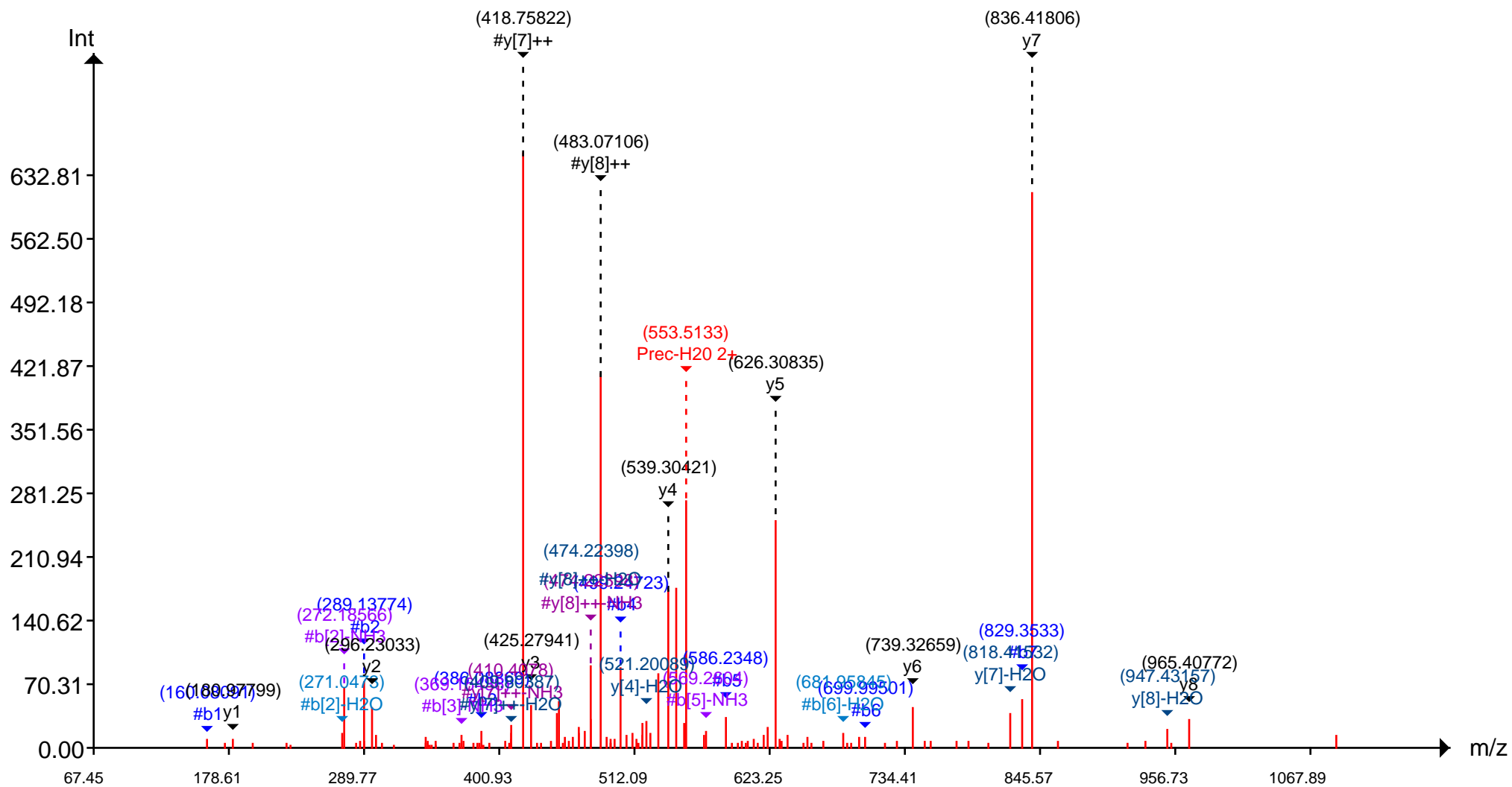
Info

Modified Sequence	AcD3-M<Mox*>K<AcD3K*>QDAQELYEAGEK<AcD3K*>R-COOH	Ion score	72.01
Protein	P97429	Confidence level	99.00%
Precursor m/z	973.97848	Identity Threshold	47.28
Charge	2+	Mass error	0.002299



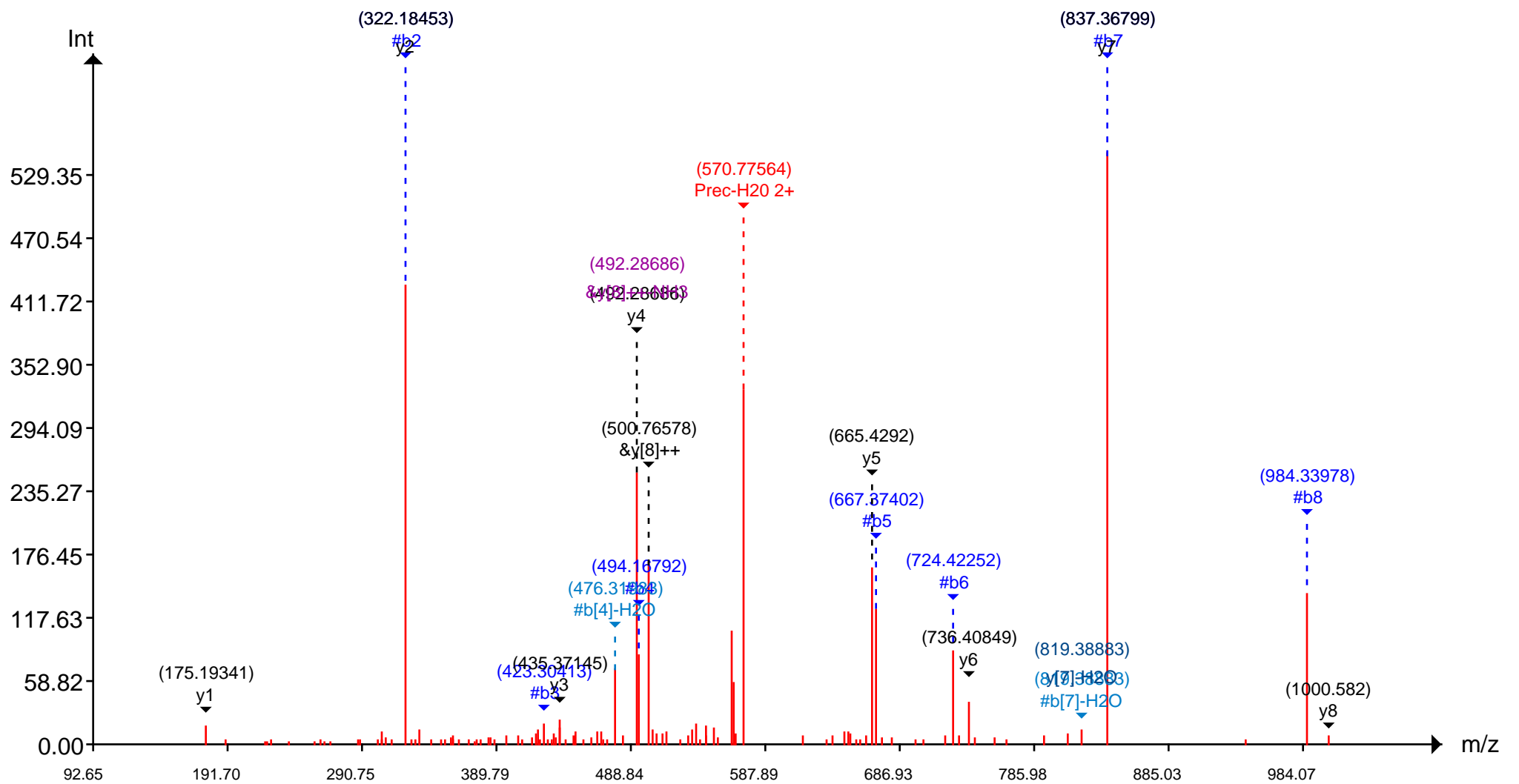
Info

Modified Sequence	AcD3-VLVGDGGTgk<AcD3K*>TTFVK<AcD3K*>R<C13>-COOH	Ion score	65.66
Protein	P62827	Confidence level	99.00%
Precursor m/z	888.5198	Identity Threshold	44.46
Charge	2+	Mass error	0.001519



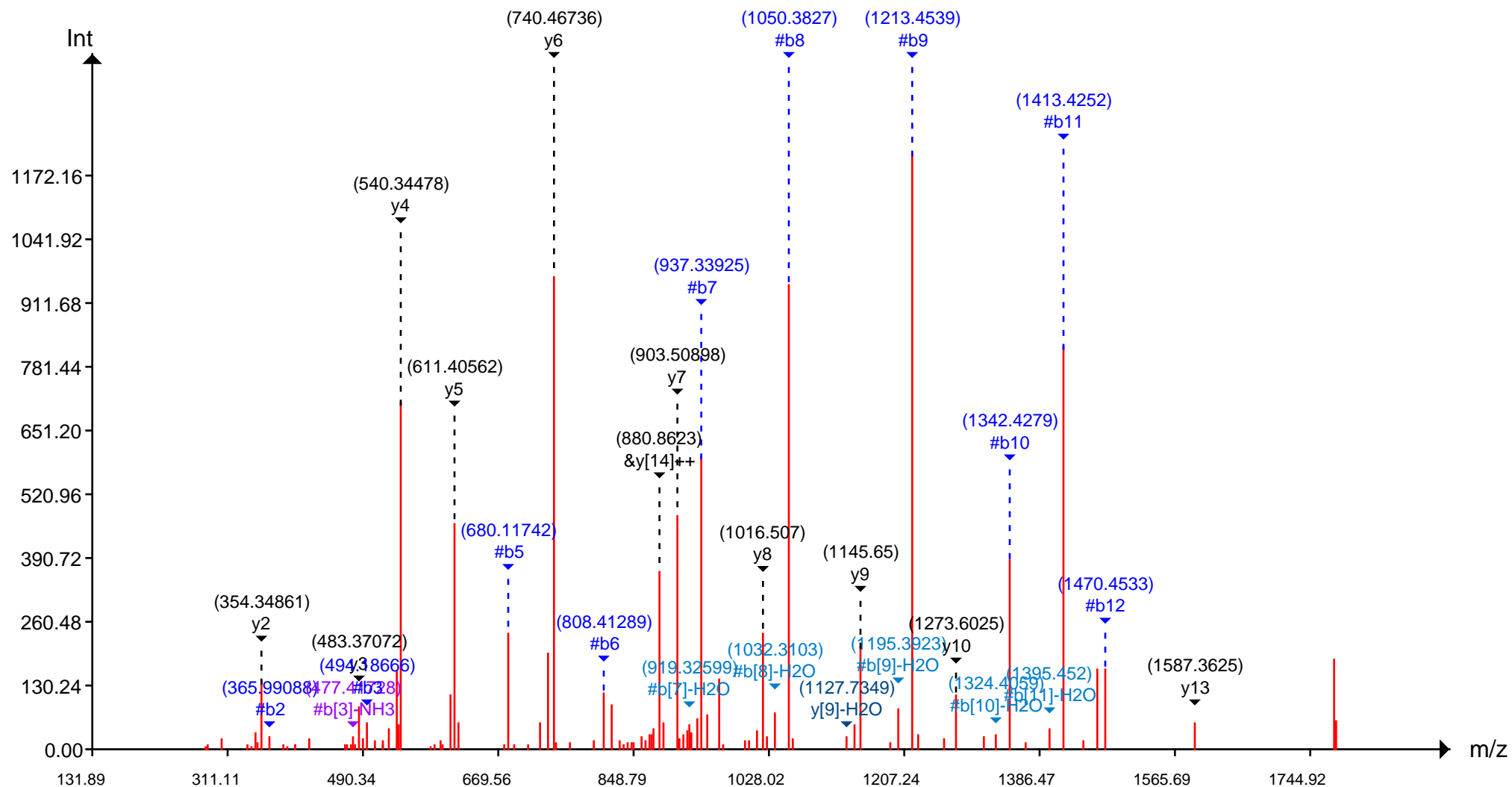
Info

Modified Sequence	AcD3-NEPLSNEDR<C13>-COOH	Ion score	60.29
Protein	P68510	Confidence level	99.00%
Precursor m/z	562.77149	Identity Threshold	46.44
Charge	2+	Mass error	4.13E-4



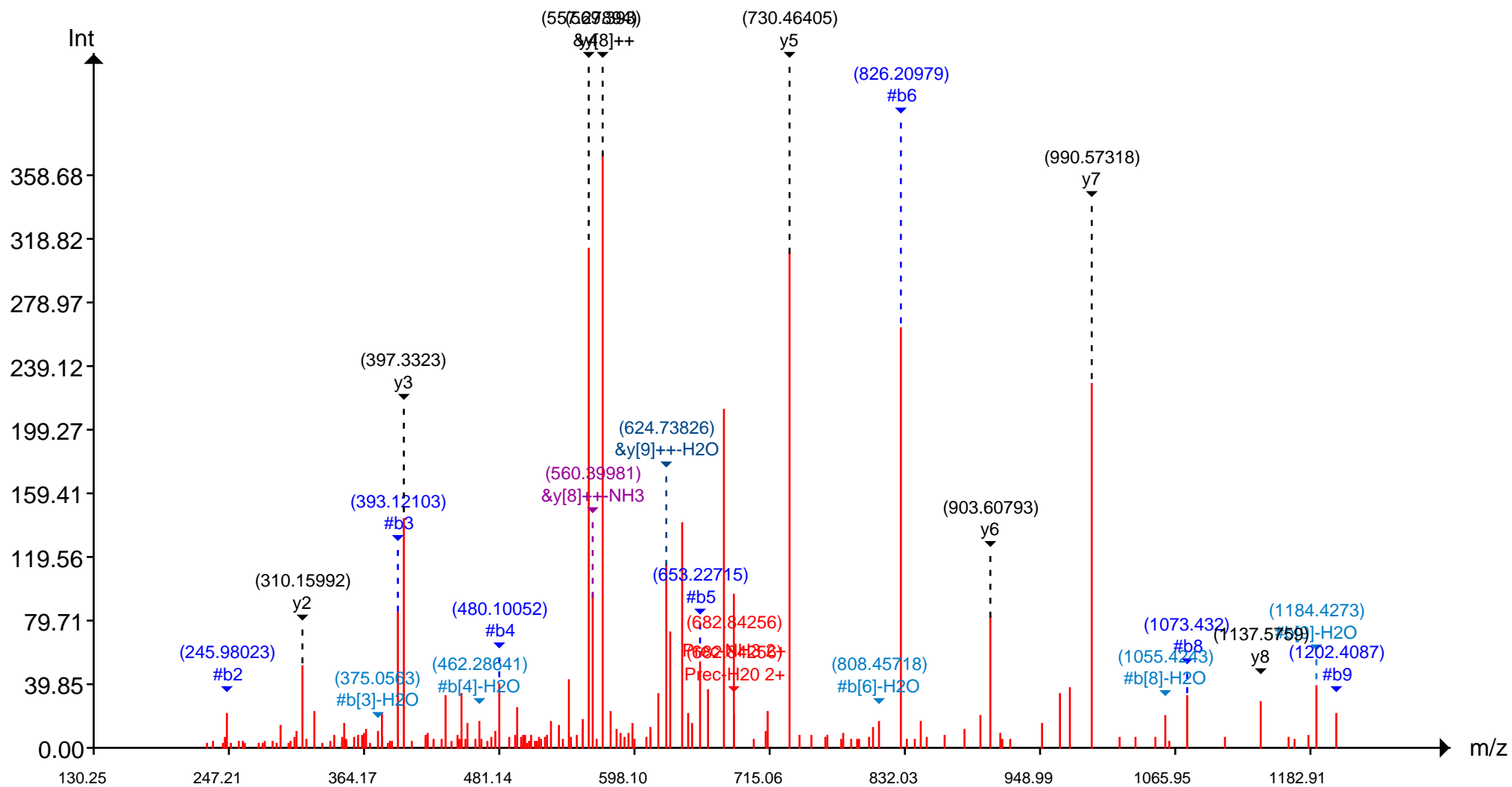
Info

Modified Sequence	AcD3-LYTAK<AcD3K*>GLFR-COOH	Ion score	73.81
Protein	P17182, P17183	Confidence level	99.00%
Precursor m/z	579.84338	Identity Threshold	45.90
Charge	2+	Mass error	6.69E-4



Info

Modified Sequence	AcD3-M<Mox*>K<AcD3K*>QDAQELYEAGEK<AcD3K*>R<C13>-COOH	Ion score	68.95
Protein	P97429	Confidence level	99.00%
Precursor m/z	976.98839	Identity Threshold	47.76
Charge	2+	Mass error	0.001977



Info

Modified Sequence AcD3-AEFSK<AcD3K*>K<AcD3K*>C<Cmm*>SER<C13>-COOH

Ion score 54.87

Protein P30681, O54879

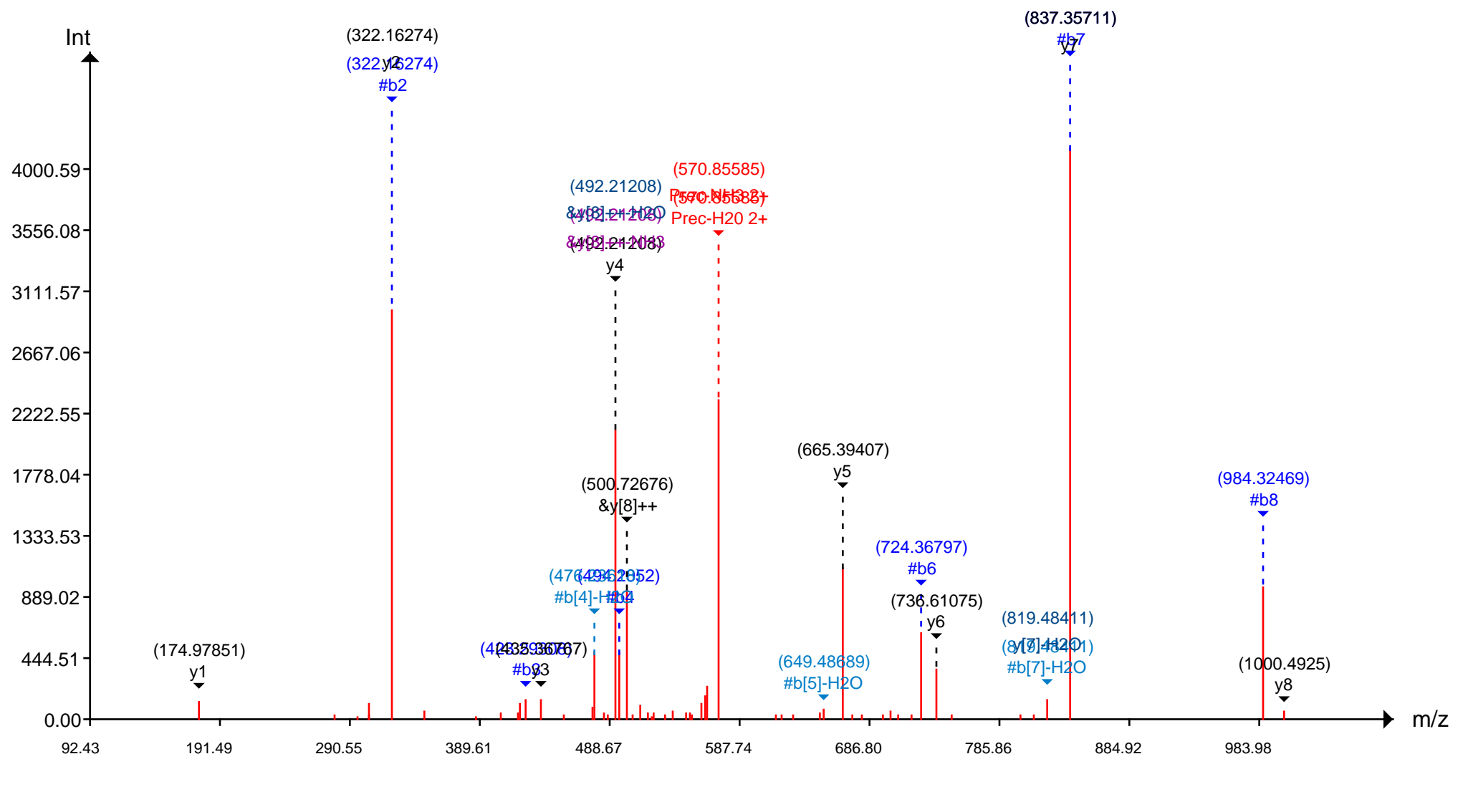
Confidence level 99.00%

Precursor m/z 691.85542

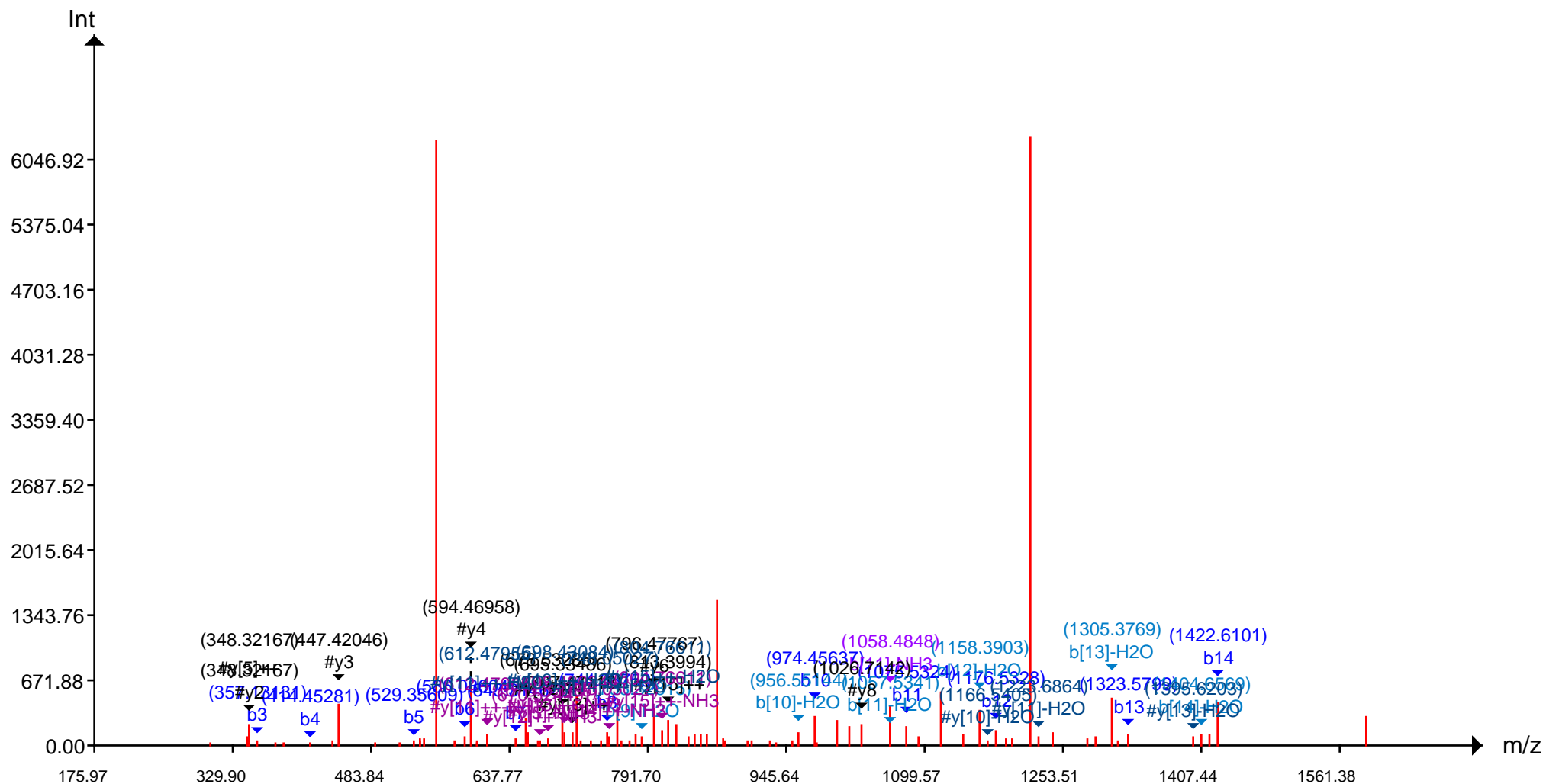
Identity Threshold 47.77

Charge 2+

Mass error 9.6E-4

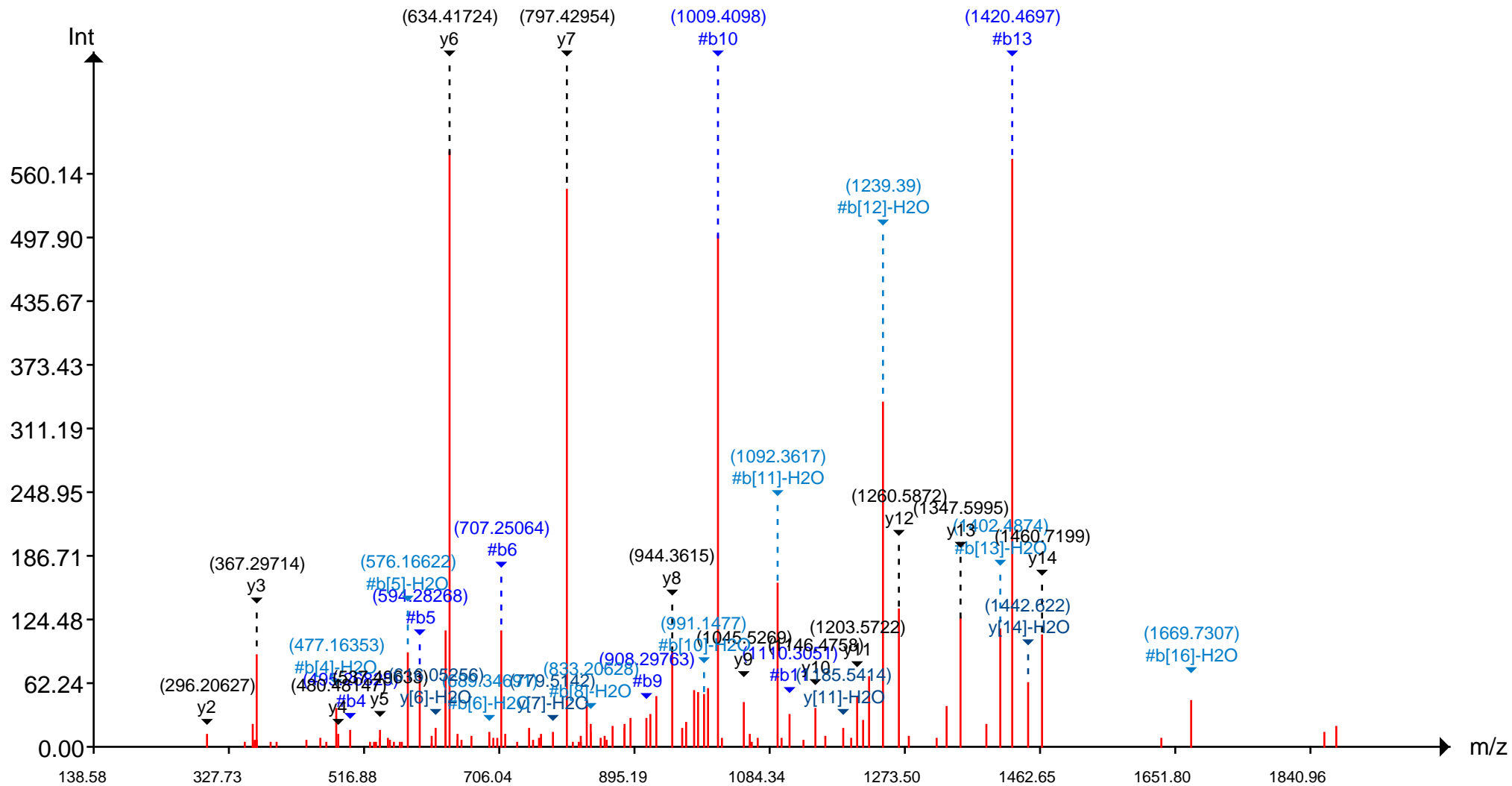


Info			
Modified Sequence	AcD3-LYTAK<AcD3K*>GLFR-COOH	Ion score	81.41
Protein	P17182, P17183	Confidence level	99.00%
Precursor m/z	579.84354	Identity Threshold	45.90
Charge	2+	Mass error	9.89E-4



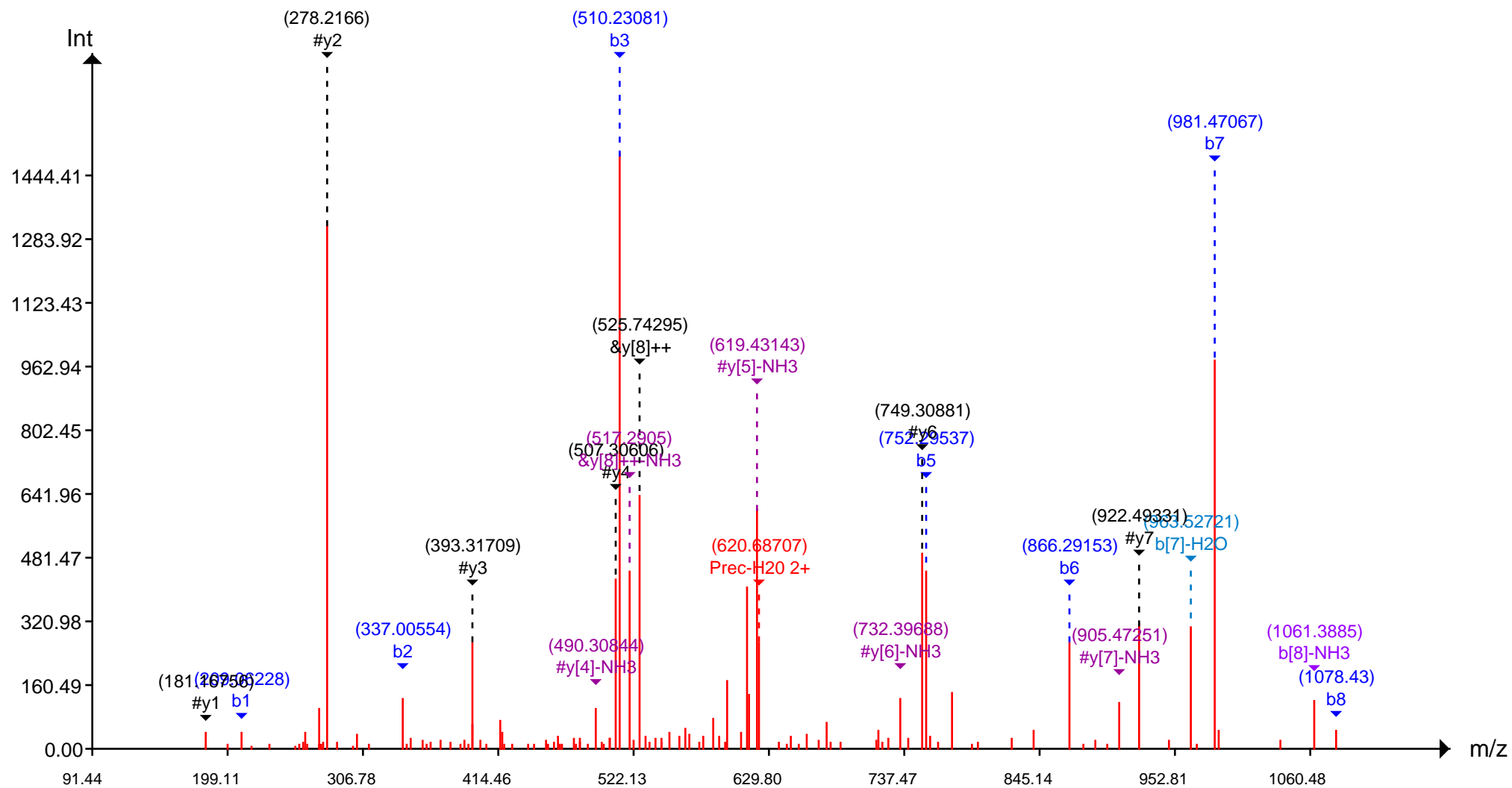
Info

Modified Sequence	AcD3-VLVGDGGTGTK<AcD3K*>TTFVK<AcD3K*>R-COOH	Ion score	53.11
Protein	P62827	Confidence level	99.00%
Precursor m/z	885.51123	Identity Threshold	45.17
Charge	2+	Mass error	0.004521



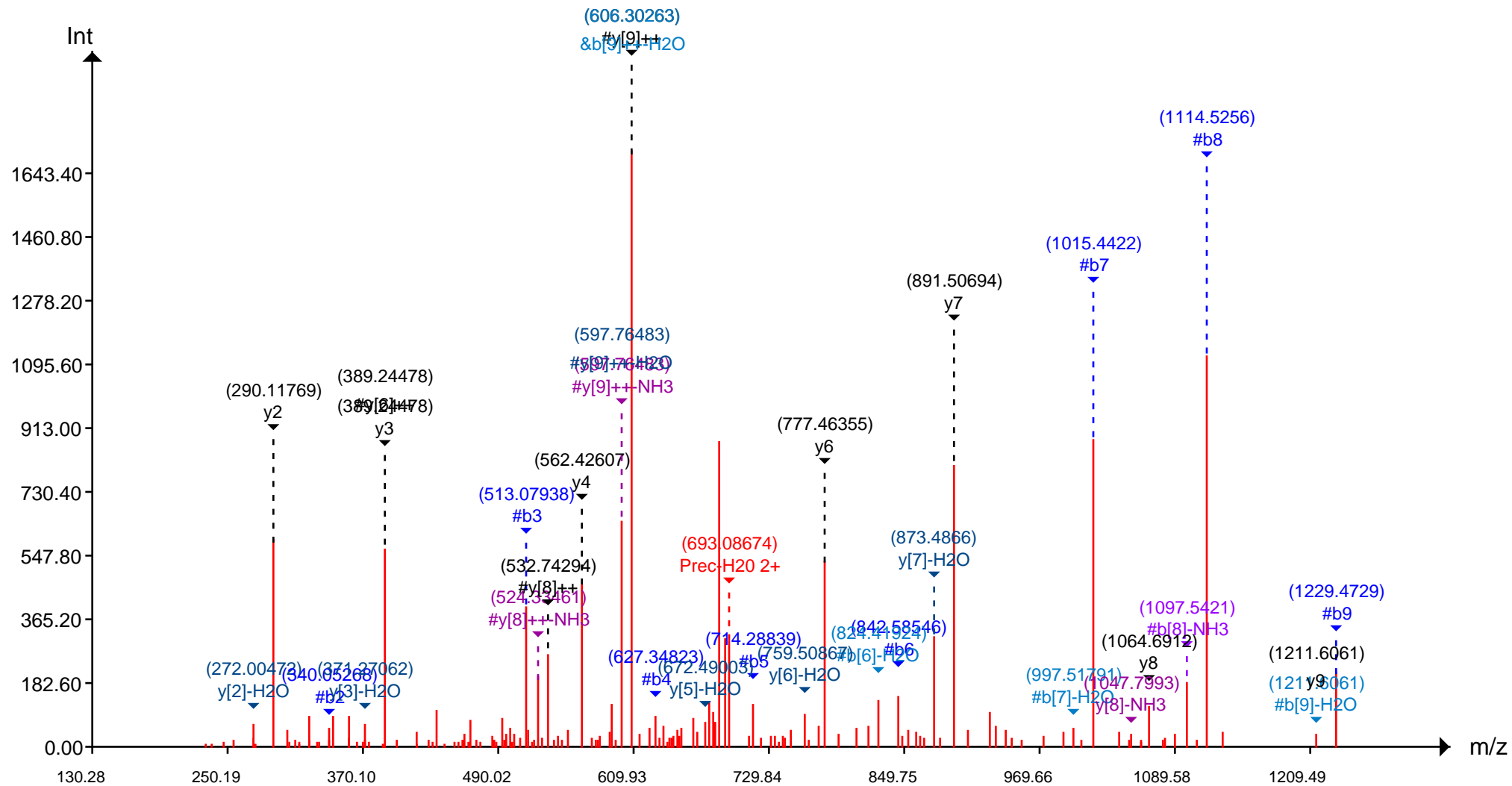
Info

Modified Sequence	AcD3-YANTVLSGGTTM<Mox*>YPGIADR<C13>-COOH	Ion score	89.49
Protein	P60710, P63260	Confidence level	99.00%
Precursor m/z	1027.5064	Identity Threshold	47.46
Charge	2+	Mass error	0.002124



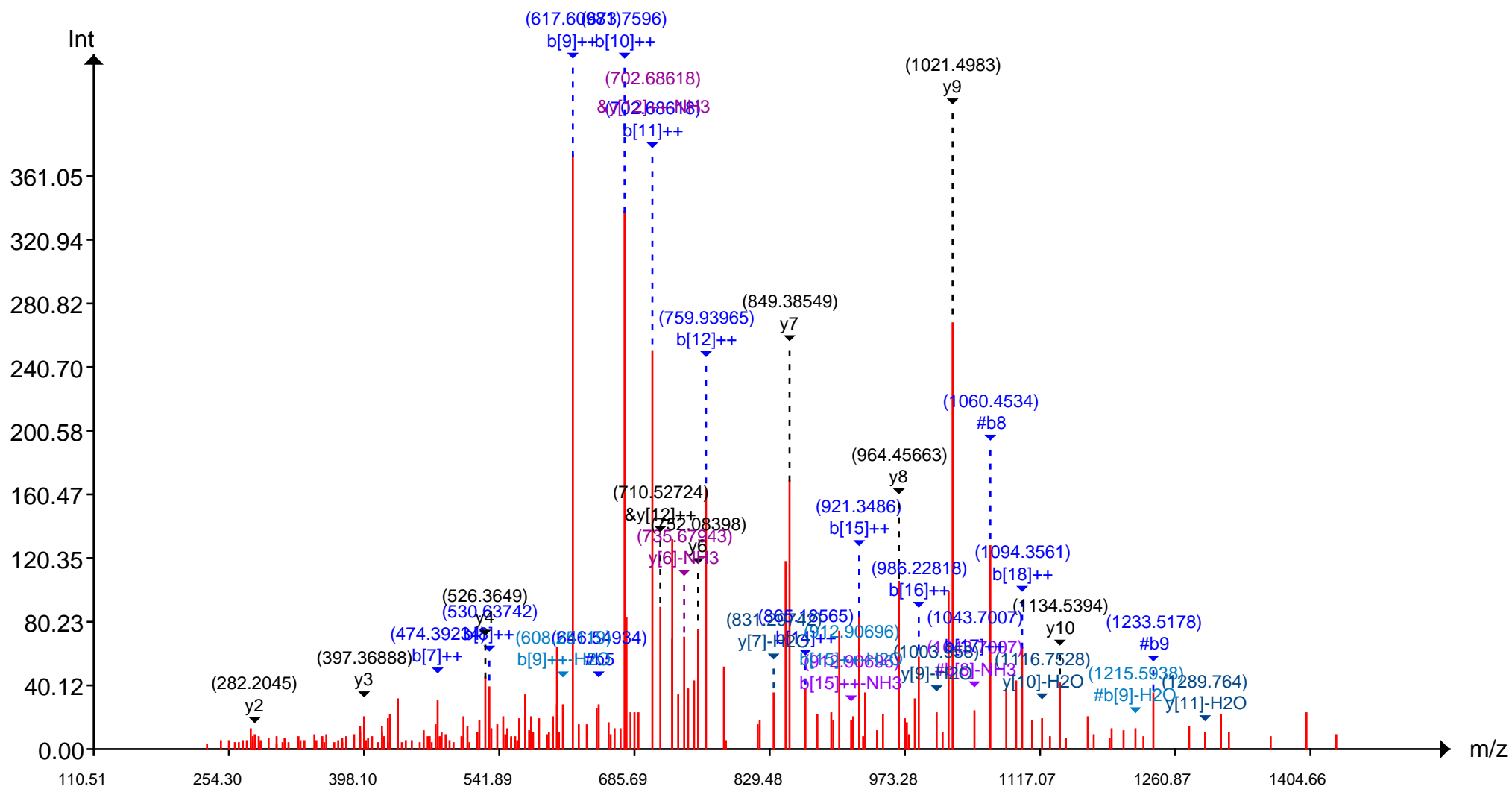
Info

Modified Sequence	AcD3-YQK<AcD3K*>LENDPR<C13>-COOH	Ion score	48.51
Protein	Q60864	Confidence level	99.00%
Precursor m/z	629.8363	Identity Threshold	47.89
Charge	2+	Mass error	0.00131



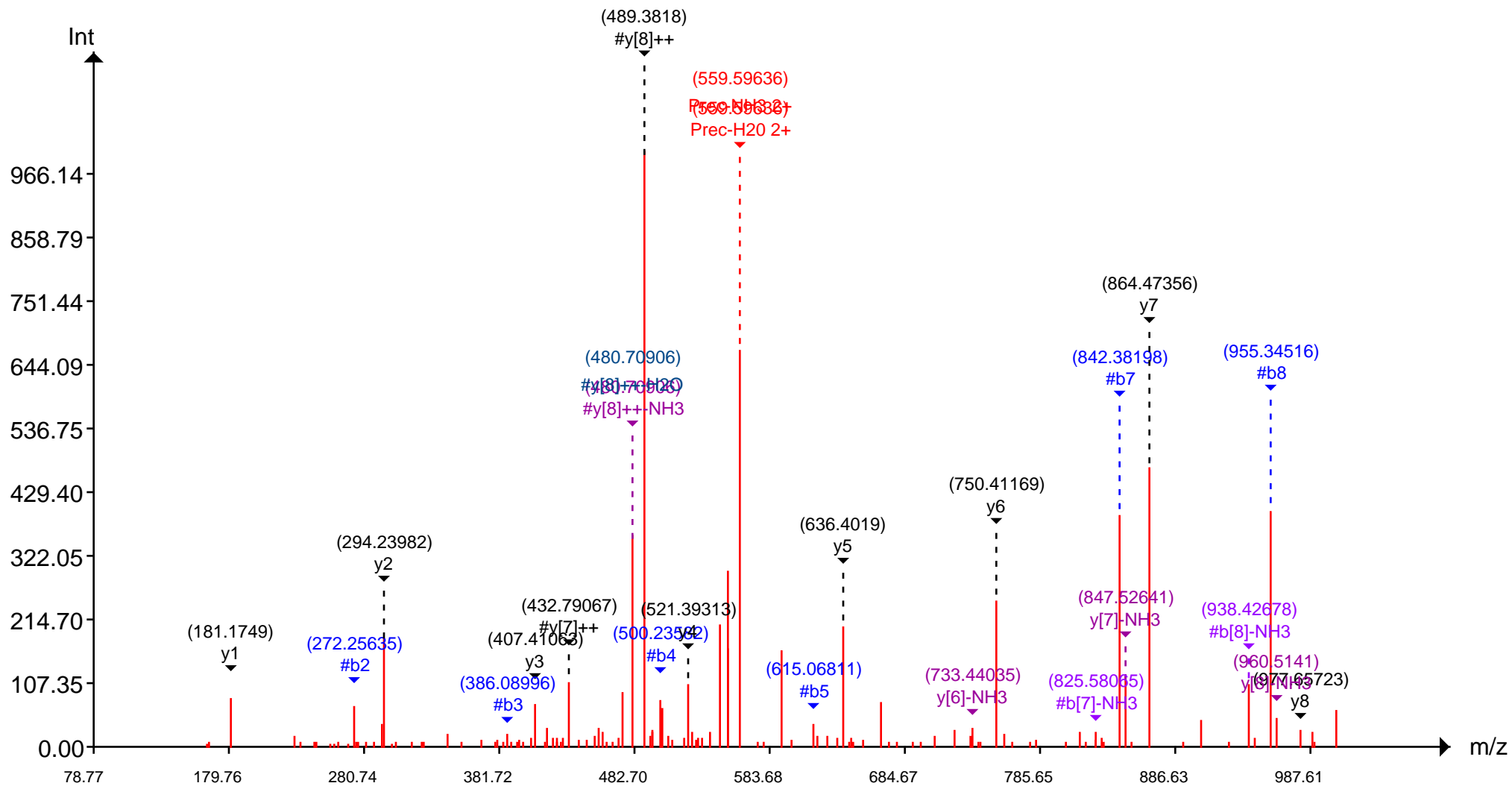
Info

Modified Sequence	AcD3-FFK<AcD3K*>NSQK<AcD3K*>VDR-COOH	Ion score	52.96
Protein	Q9CQM9	Confidence level	99.00%
Precursor m/z	702.38551	Identity Threshold	47.91
Charge	2+	Mass error	9.6E-4



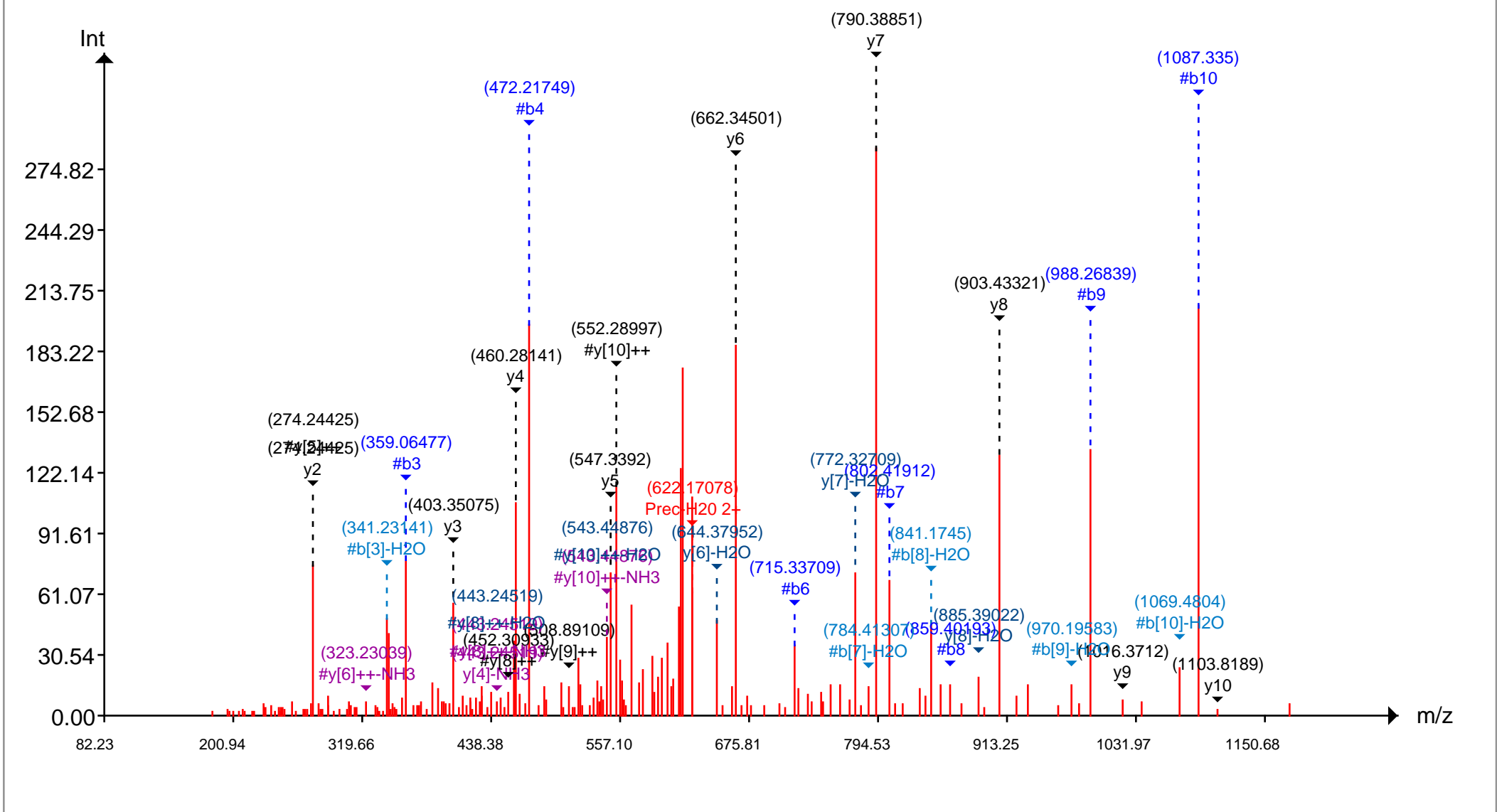
Info

Modified Sequence	AcD3-VVK<AcD3K*>QTQK<AcD3K*>IK<AcD3K*>LGDPLLEDTR<C13>-COOH	Ion score	49.62
Protein	Q9JLJ2	Confidence level	99.00%
Precursor m/z	789.80502	Identity Threshold	40.40
Charge	3+	Mass error	0.002638



Info

Modified Sequence	AcD3-LLNNDNLLR<C13>-COOH	Ion score	57.59
Protein	P47753, P47754	Confidence level	99.00%
Precursor m/z	568.33355	Identity Threshold	46.26
Charge	2+	Mass error	-6.04E-4



Info

Modified Sequence AcD3-LSLLQDSGEVR-COOH

Ion score 76.21

Protein P63242

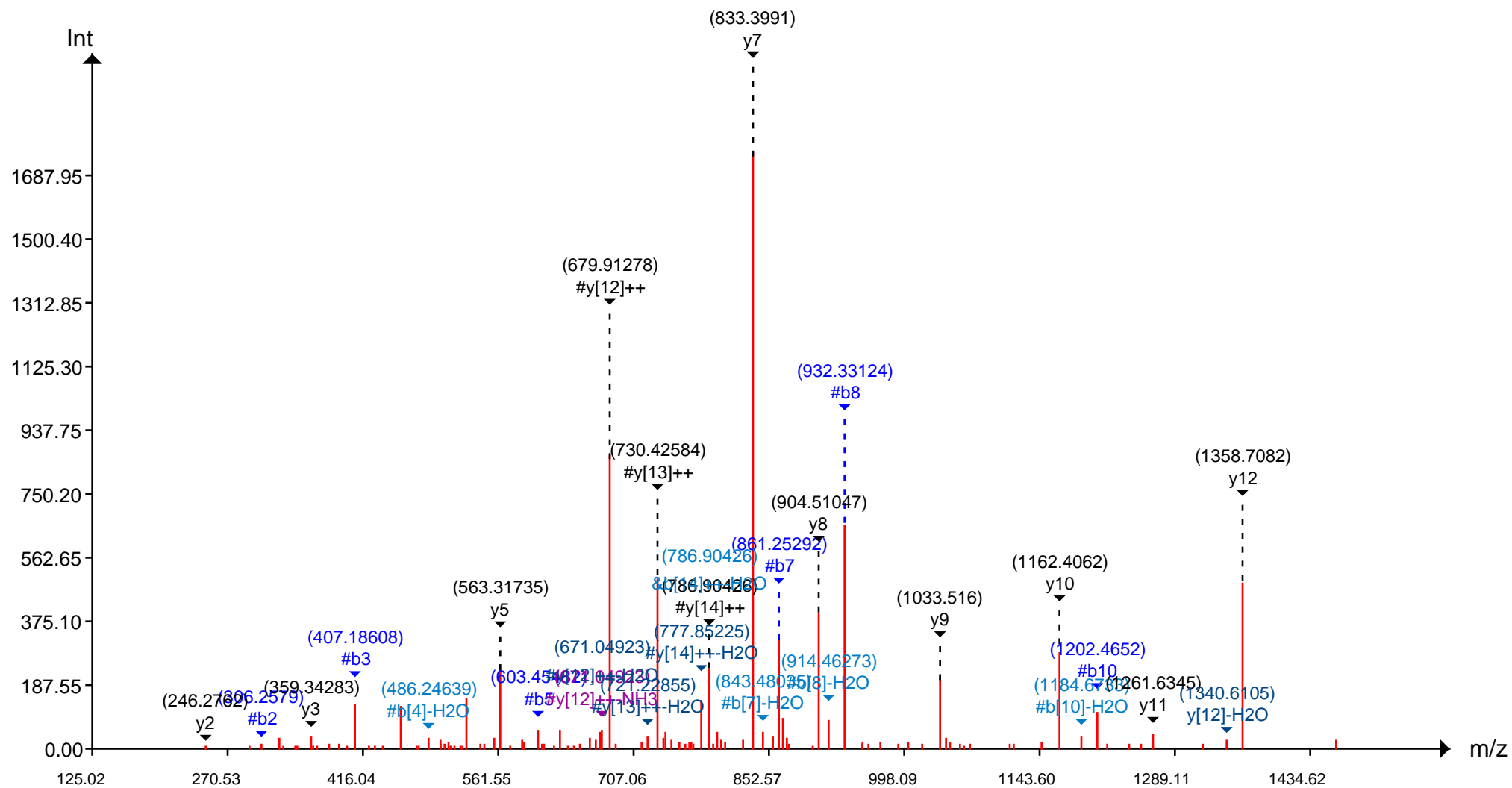
Confidence level 99.00%

Precursor m/z 631.34525

Identity Threshold 47.93

Charge 2+

Mass error 6.55E-4



Info

Modified Sequence AcD3-FLTPVEEAPK<AcD3K*>GM<MoX*>LAR-COOH

Ion score 62.27

Protein Q61599

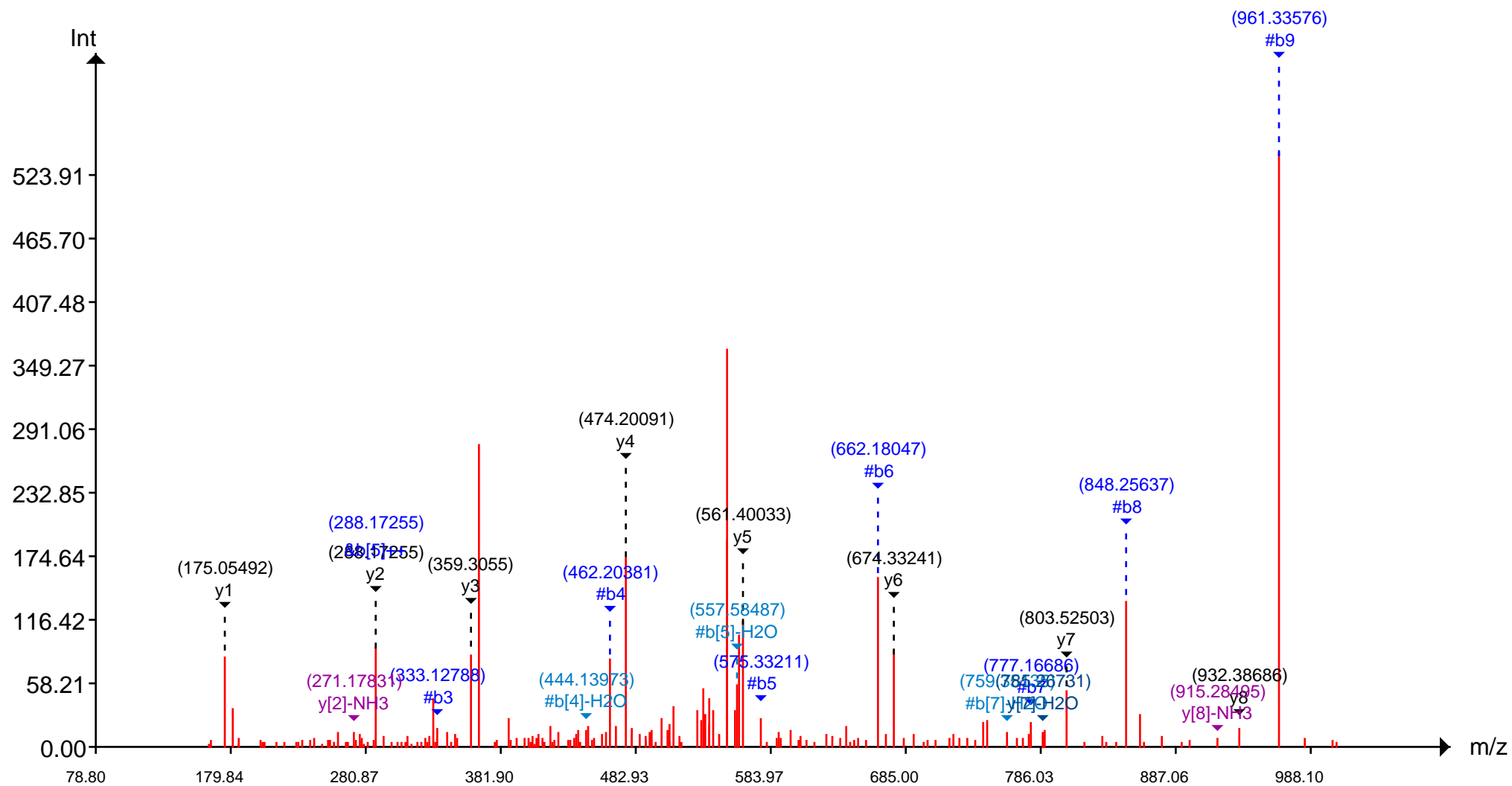
Confidence level 99.00%

Precursor m/z 882.97832

Identity Threshold 47.95

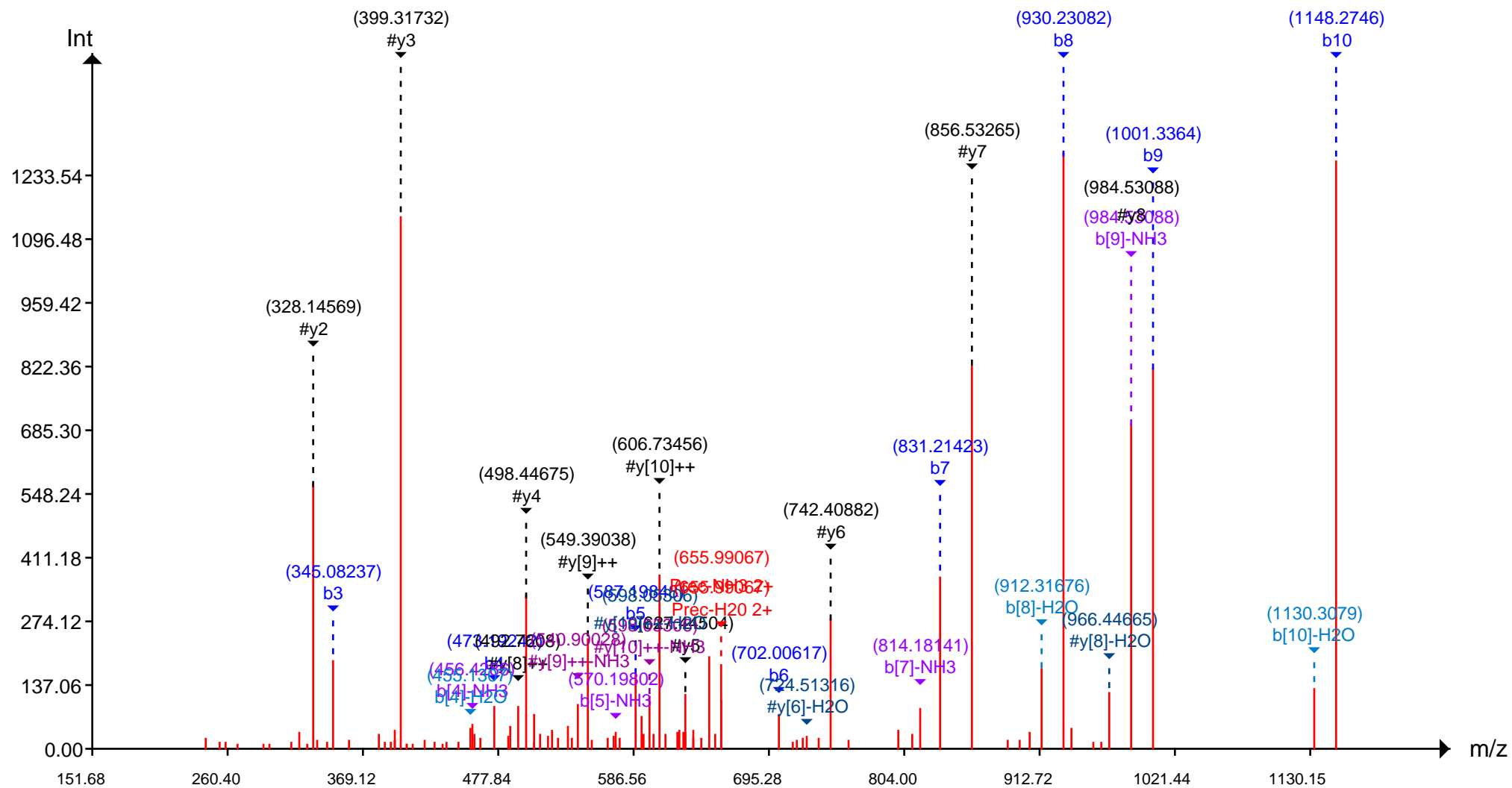
Charge 2+

Mass error 0.002238



Info

Modified Sequence	AcD3-SAEEISDAIR-COOH	Ion score	50.64
Protein	Q9JHU4	Confidence level	99.00%
Precursor m/z	568.28719	Identity Threshold	47.70
Charge	2+	Mass error	2.58E-4



Info

Modified Sequence AcD3-ADLQNDEVAFR<C13>-COOH

Ion score 62.01

Protein P97351

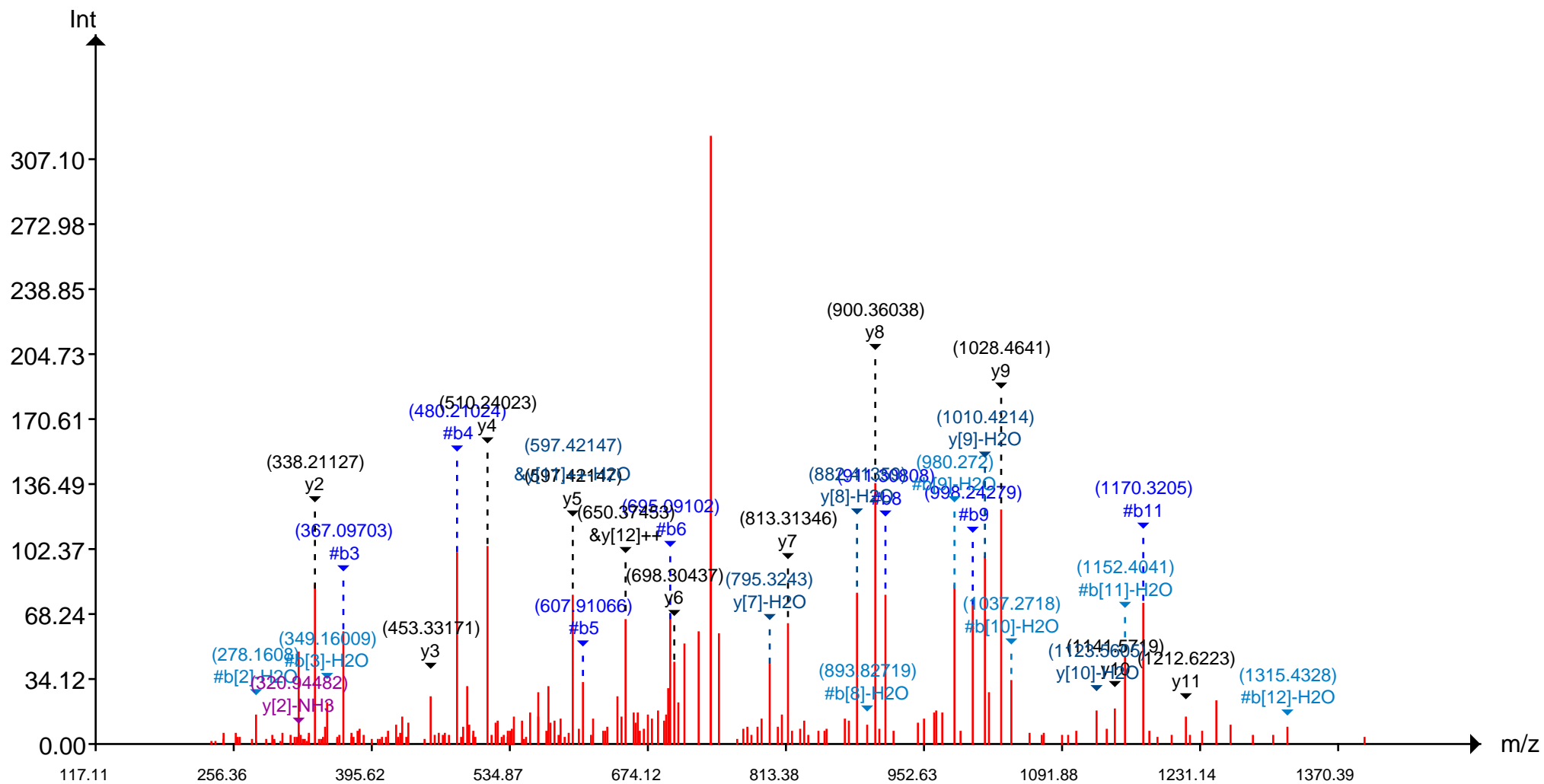
Confidence level 99.00%

Precursor m/z 664.83475

Identity Threshold 47.44

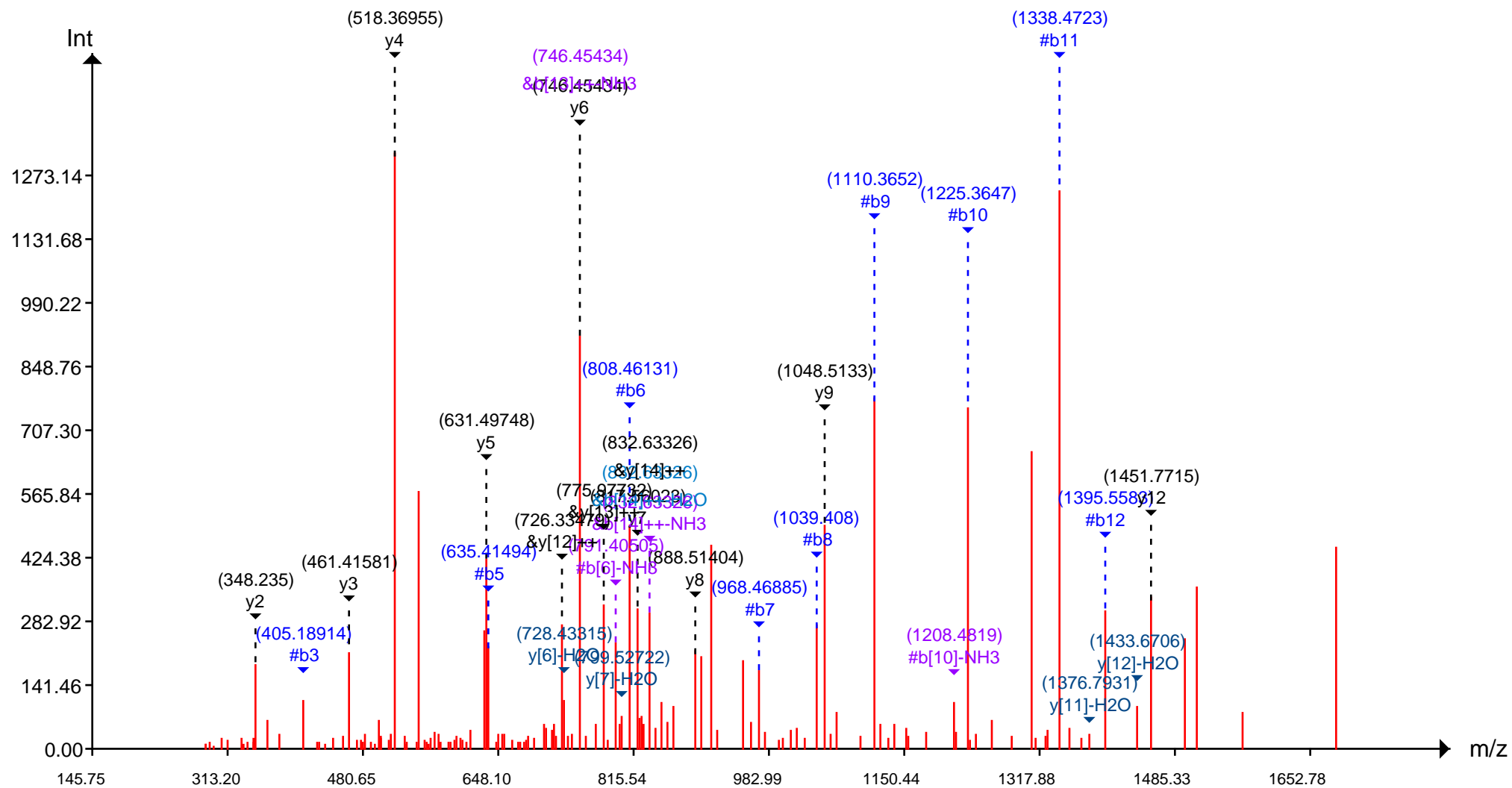
Charge 2+

Mass error 6.51E-4



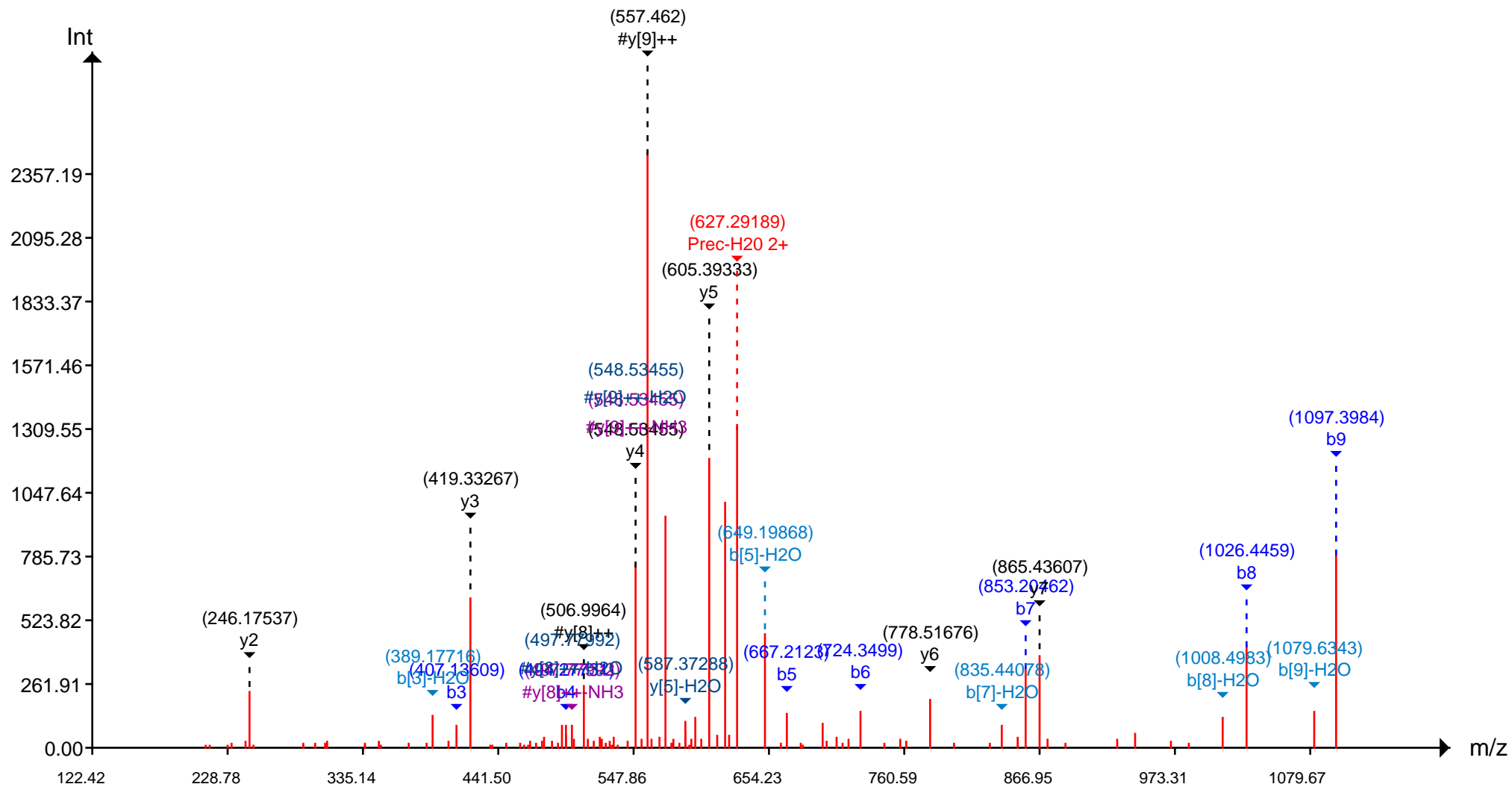
Info

Modified Sequence	AcD3-YSAIQSDTSGDYR-COOH	Ion score	63.51
Protein	O35639	Confidence level	99.00%
Precursor m/z	754.3411	Identity Threshold	44.02
Charge	2+	Mass error	0.001068



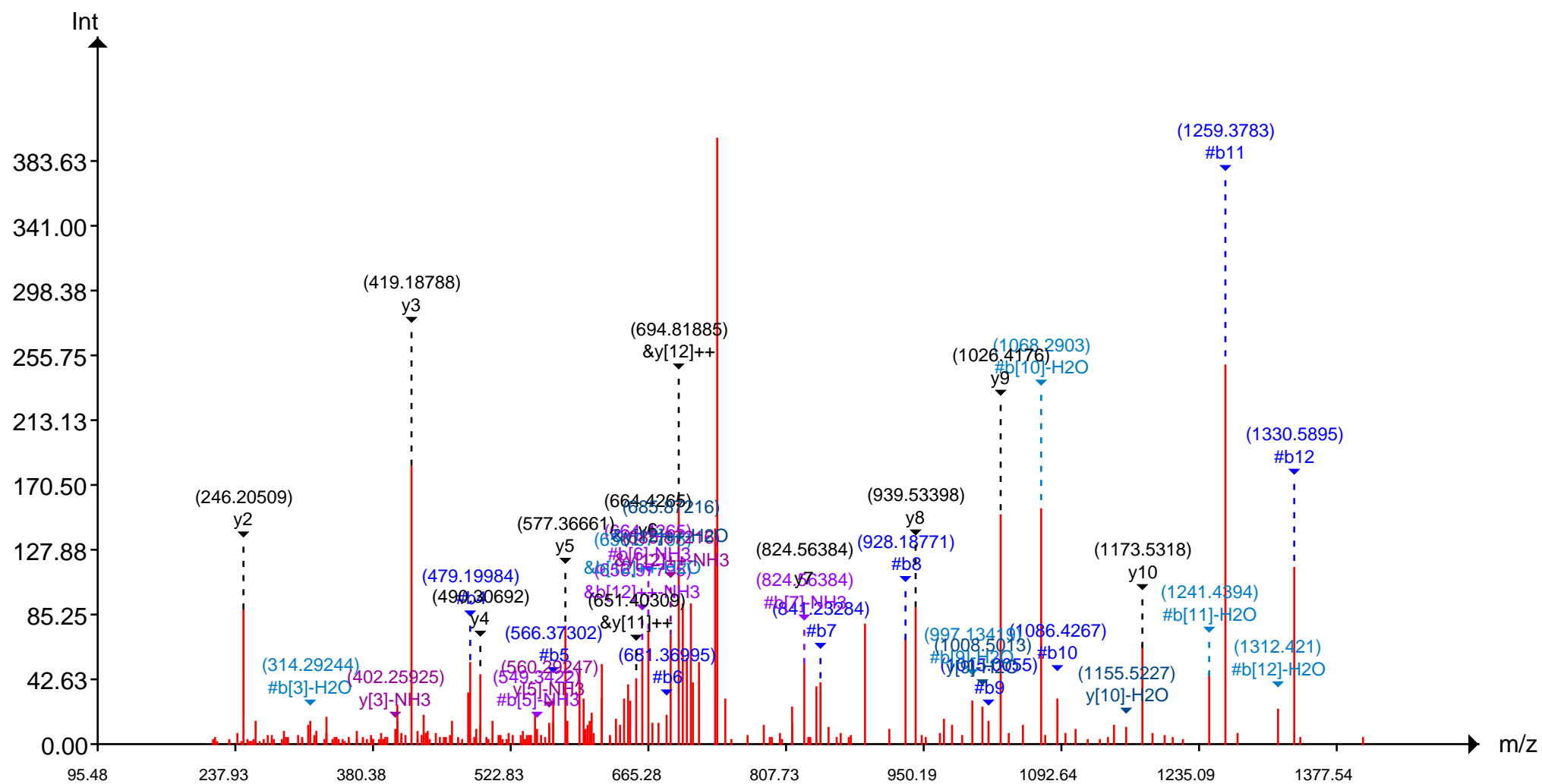
Info

Modified Sequence	AcD3-M<Mox*>IVGK<AcD3K*>K<AcD3K*>C<Cmm*>AADLGLK<AcD3K*>R-COOH	Ion score	63.14
Protein	P70349	Confidence level	99.00%
Precursor m/z	928.53049	Identity Threshold	45.52
Charge	2+	Mass error	0.001537



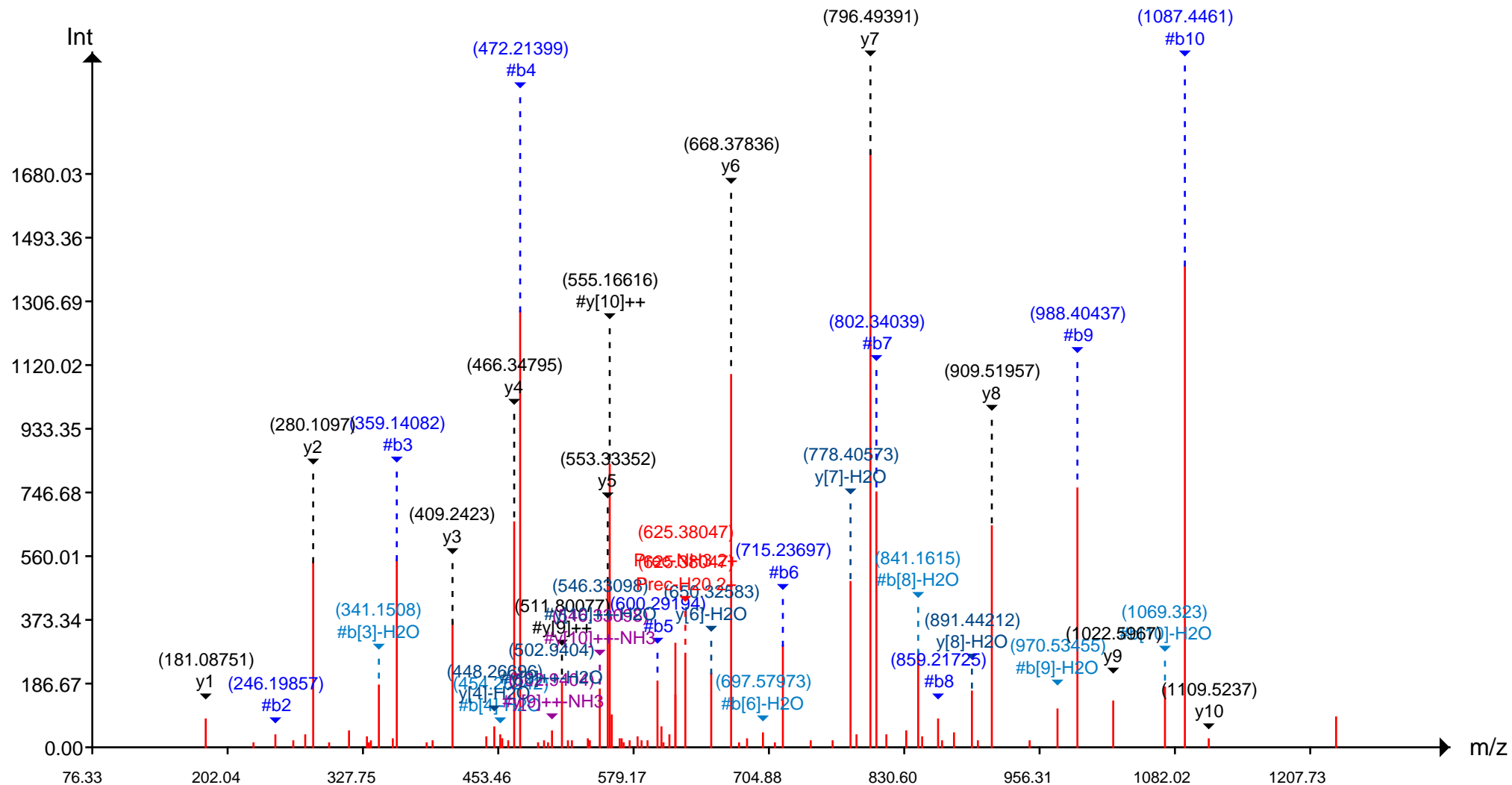
Info

Modified Sequence	AcD3-LTM<Mox*>SK<AcD3K*>GEK<AcD3K*>AR-COOH	Ion score	49.47
Protein	Q62446	Confidence level	99.00%
Precursor m/z	636.35289	Identity Threshold	47.58
Charge	2+	Mass error	0.001119



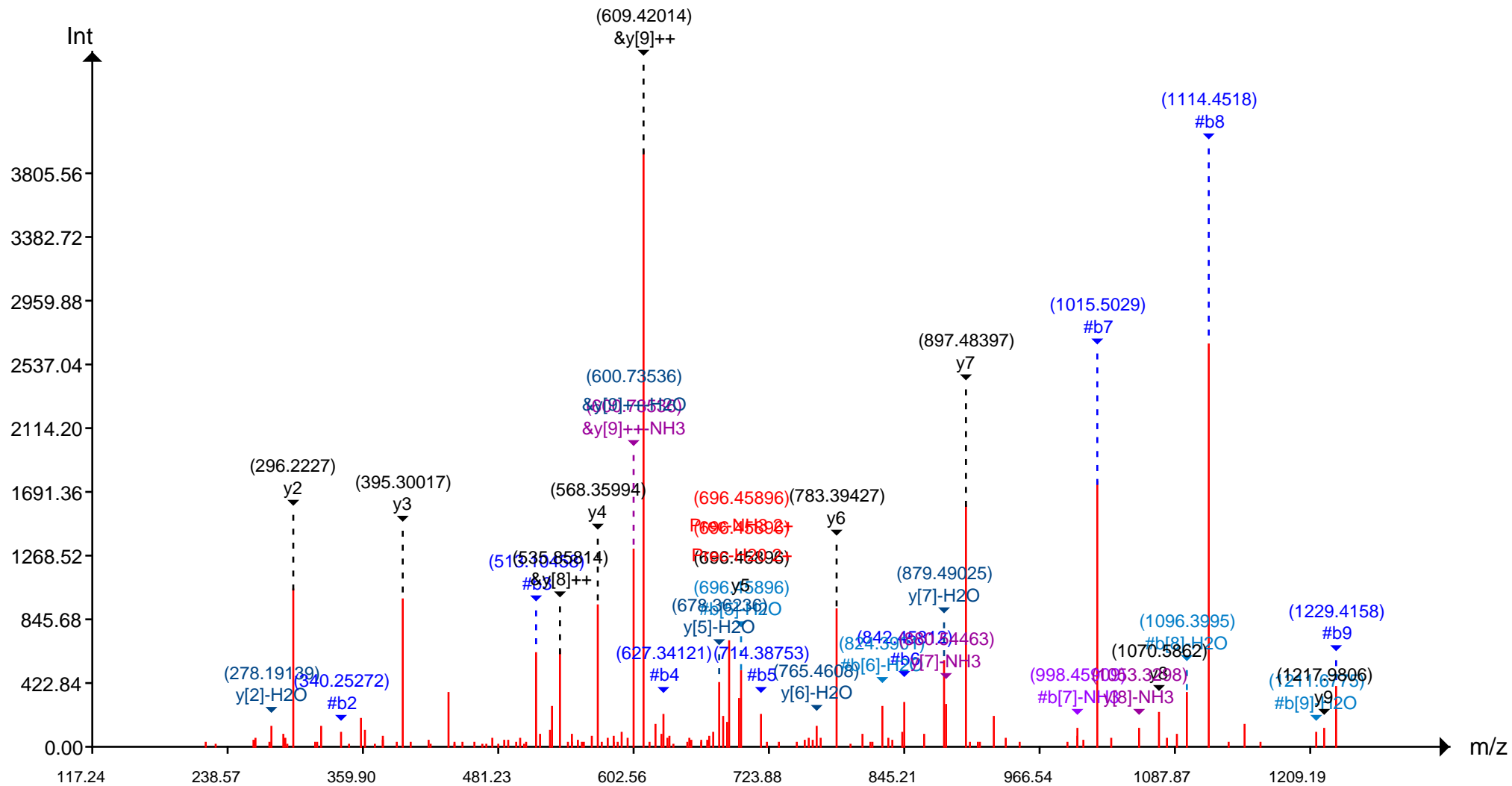
Info

Modified Sequence	AcD3-ASQFSDC<Cmm*>SSAK<AcD3K*>AR-COOH	Ion score	53.82
Protein	Q9QUR7	Confidence level	99.00%
Precursor m/z	752.85248	Identity Threshold	45.67
Charge	2+	Mass error	9.55E-4



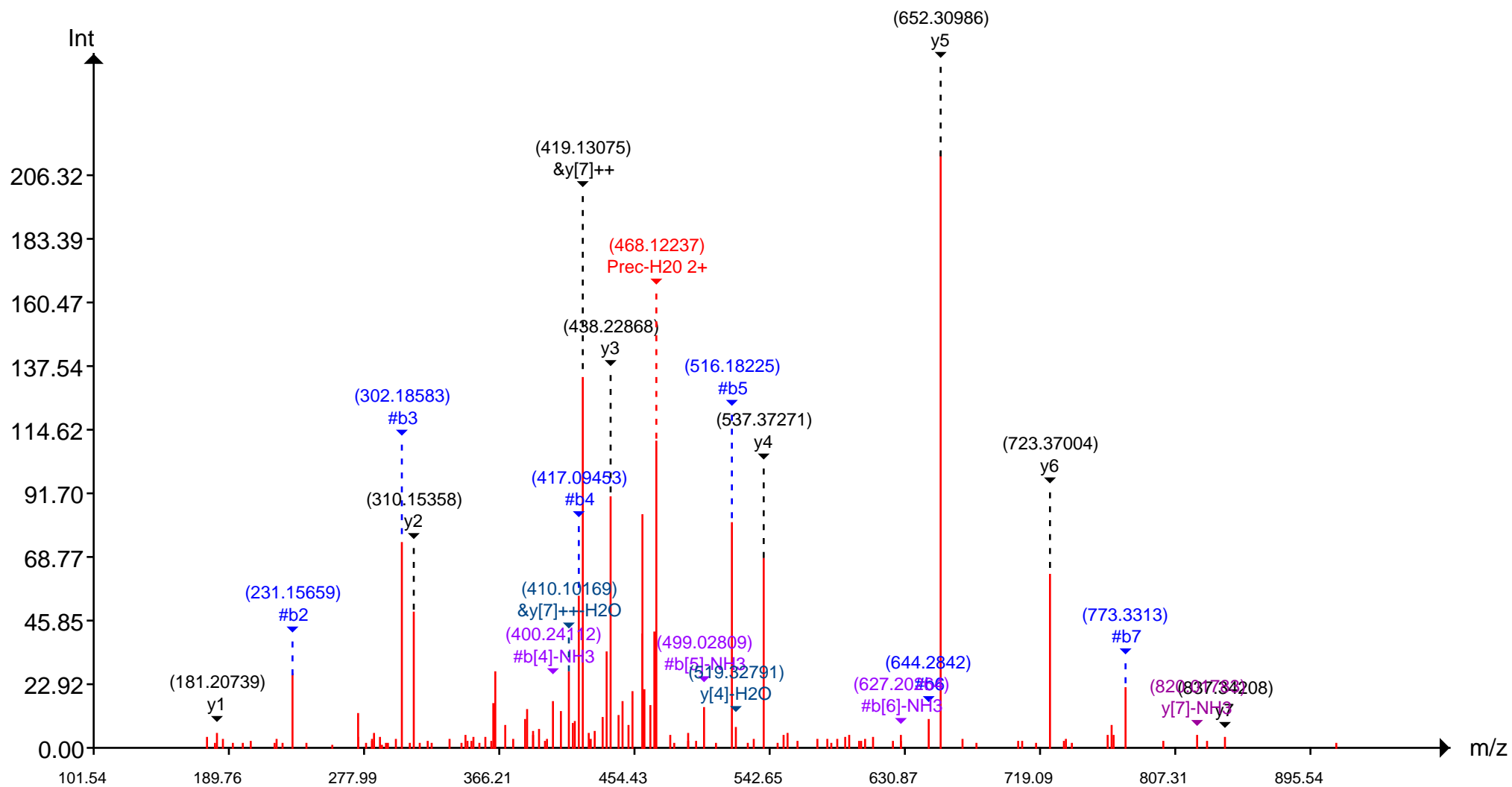
Info

Modified Sequence	AcD3-LSELLQDSGEVR<C13>-COOH	Ion score	102.33
Protein	P63242	Confidence level	99.00%
Precursor m/z	634.35529	Identity Threshold	47.19
Charge	2+	Mass error	5.93E-4



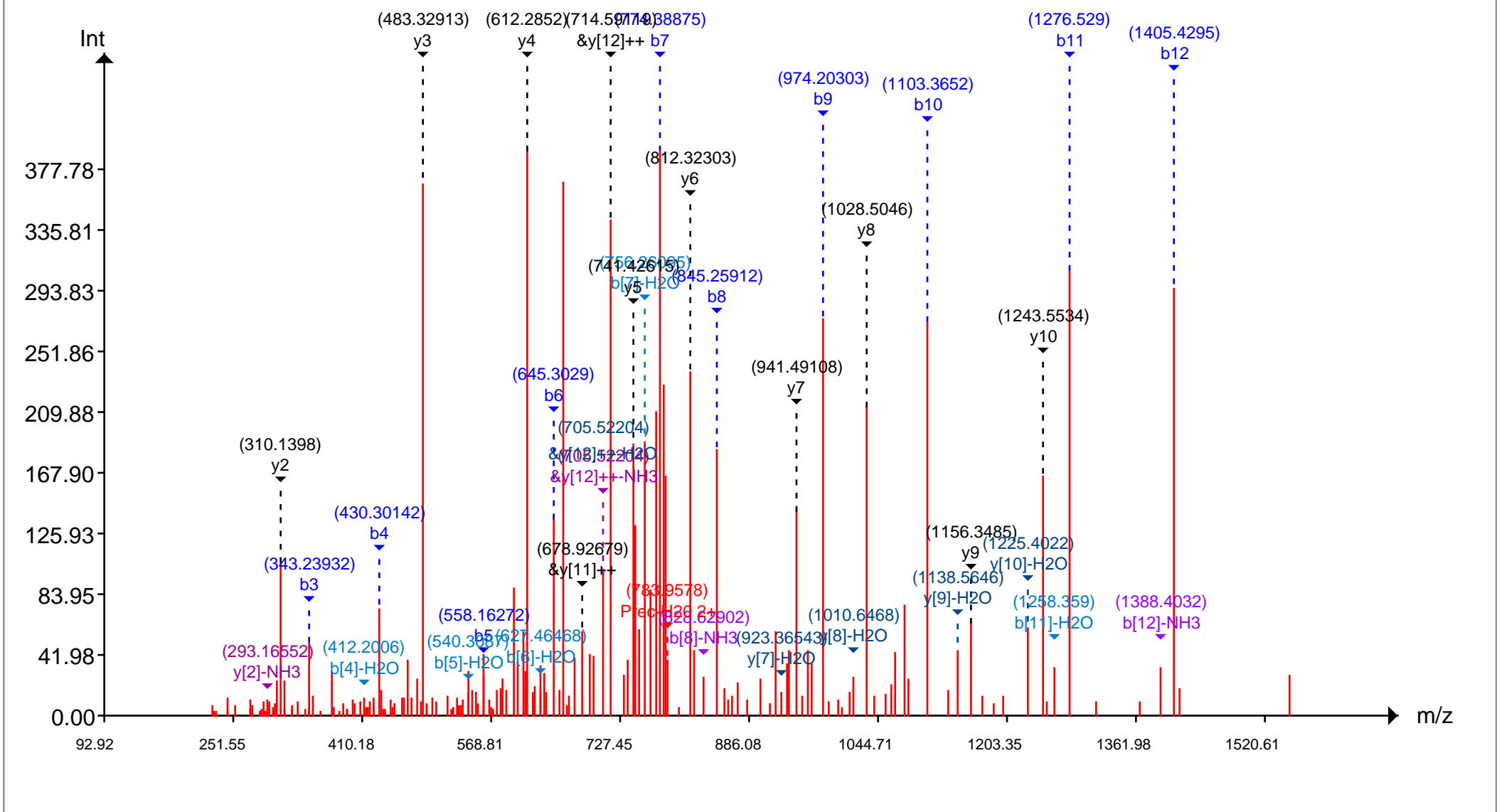
Info

Modified Sequence	AcD3-FFK<AcD3K*>NSQK<AcD3K*>VDR<C13>-COOH	Ion score	60.69
Protein	Q9CQM9	Confidence level	99.00%
Precursor m/z	705.39582	Identity Threshold	47.32
Charge	2+	Mass error	0.001438



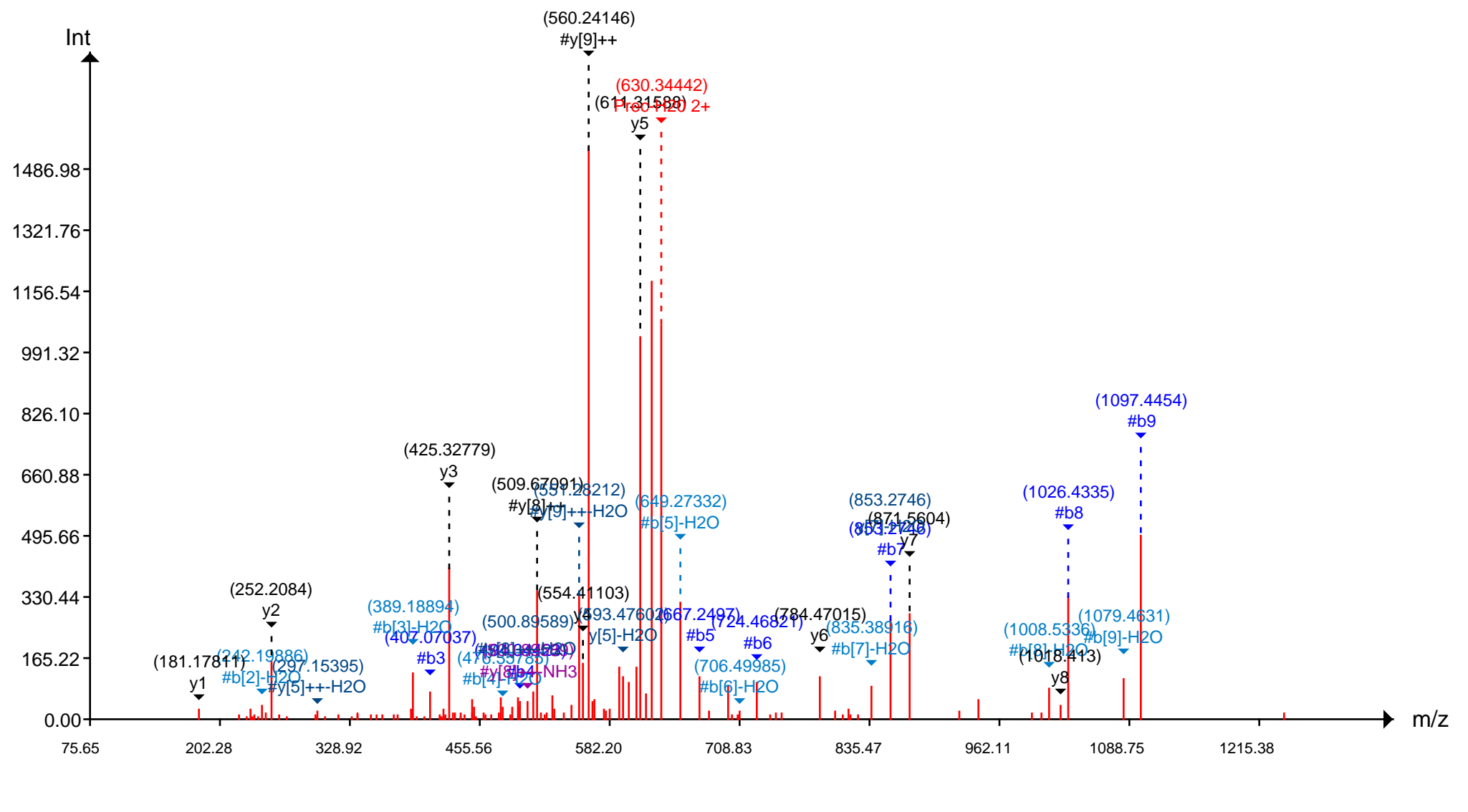
Info

Modified Sequence	AcD3-ANADVQER<C13>-COOH	Ion score	56.37
Protein	Q9JKV1	Confidence level	99.00%
Precursor m/z	477.24471	Identity Threshold	47.39
Charge	2+	Mass error	-1.0E-6

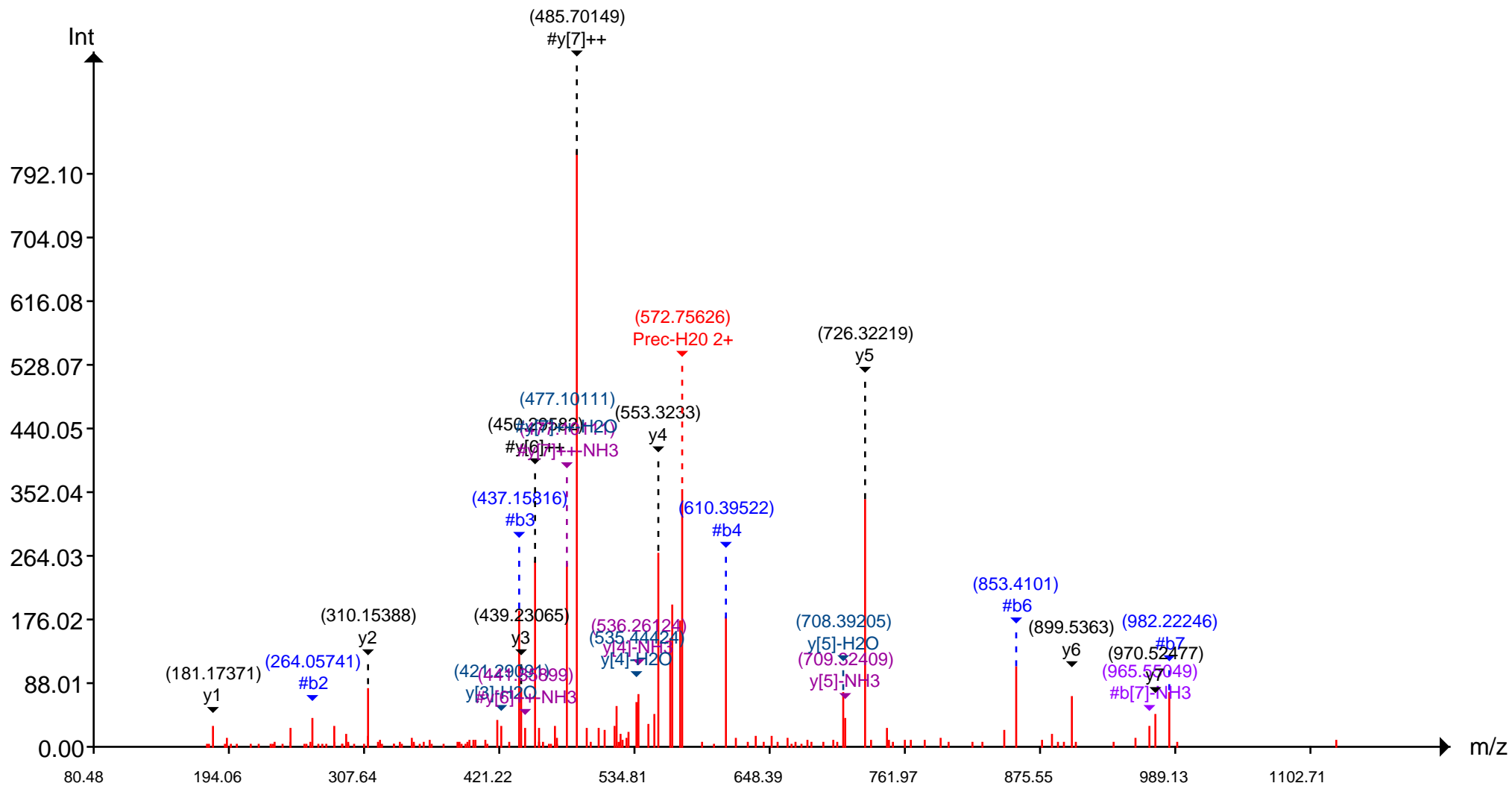


Info

Modified Sequence	AcD3-LALSQSEAEK<AcD3K*>ER<C13>-COOH	Ion score	72.06
Protein	Q99LI8	Confidence level	99.00%
Precursor m/z	793.41772	Identity Threshold	48.13
Charge	2+	Mass error	1.1E-5



Info			
Modified Sequence	AcD3-LTM<Mox*>SK<AcD3K*>GEK<AcD3K*>AR<C13>-COOH	Ion score	60.44
Protein	Q62446	Confidence level	99.00%
Precursor m/z	639.36286	Identity Threshold	47.32
Charge	2+	Mass error	9.17E-4



Info

Modified Sequence AcD3-FAK<AcD3K*>K<AcD3K*>NEER<C13>-COOH

Ion score 51.49

Protein P32067

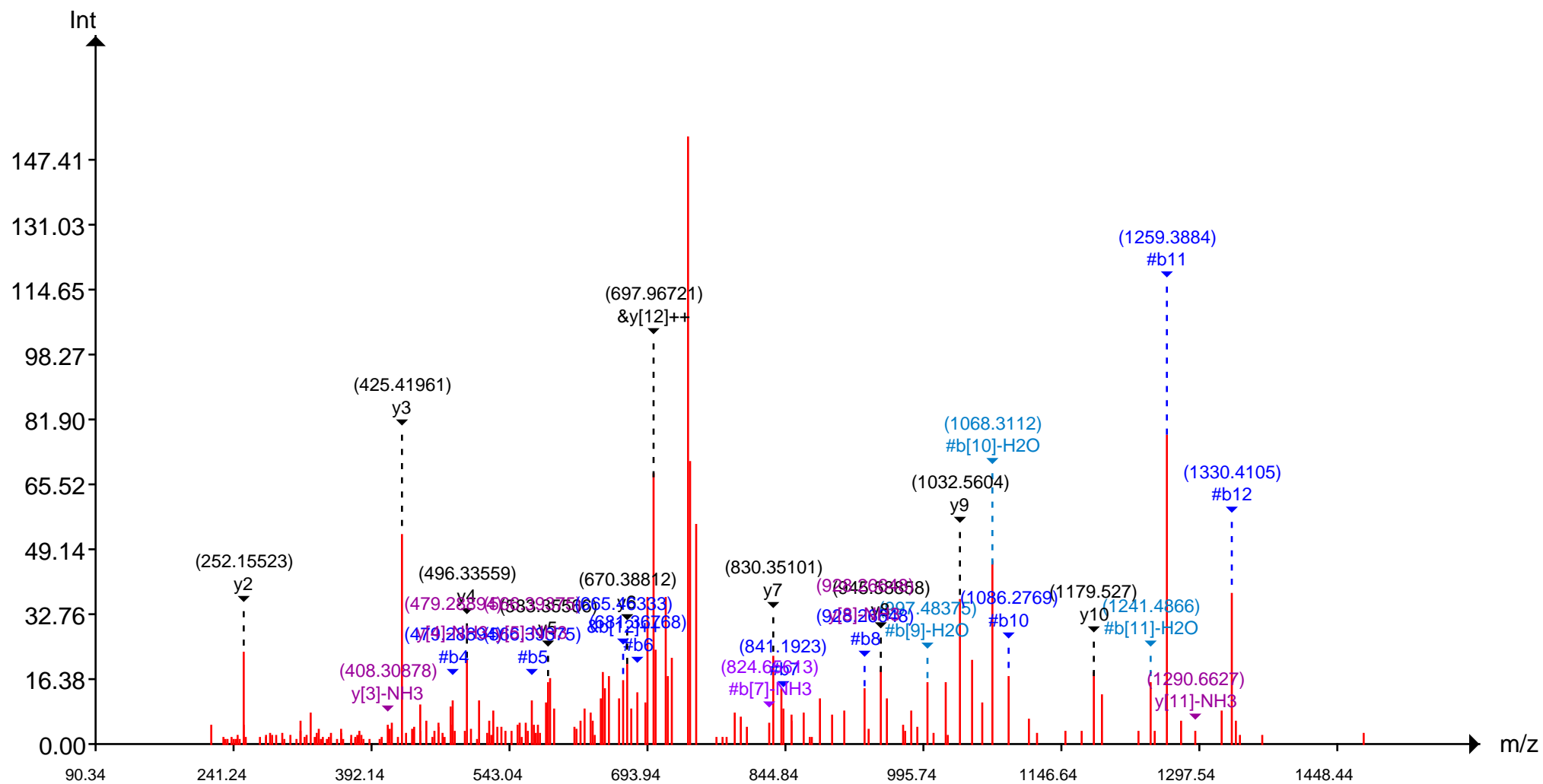
Confidence level 99.00%

Precursor m/z 581.82882

Identity Threshold 47.46

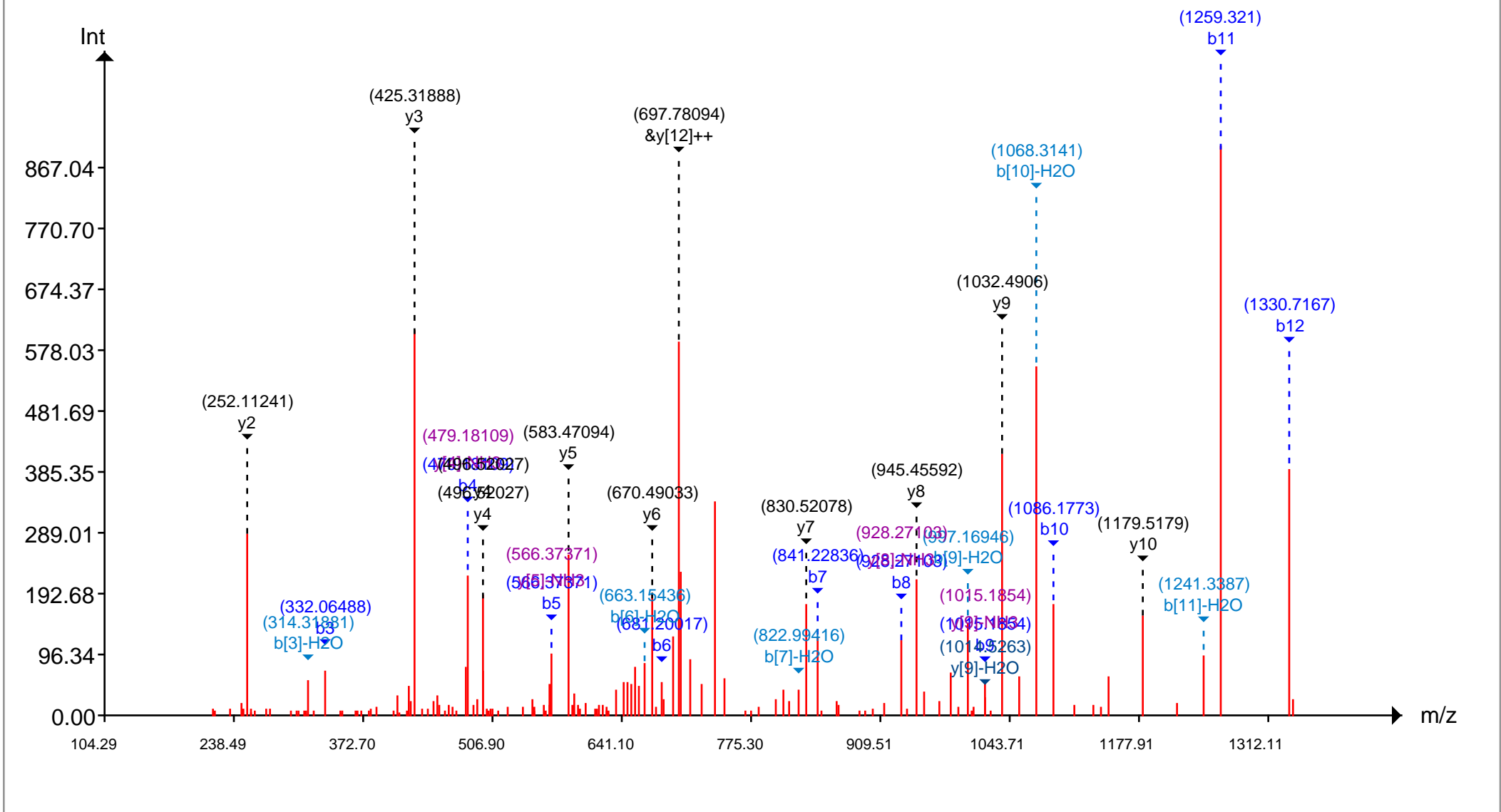
Charge 2+

Mass error -4.51E-4



Info

Modified Sequence	AcD3-ASQFSDC<Cmm*>SSAK<AcD3K*>AR<C13>-COOH	Ion score	49.87
Protein	Q9QUR7	Confidence level	99.00%
Precursor m/z	755.8619	Identity Threshold	46.45
Charge	2+	Mass error	-3.47E-4



Info

Modified Sequence AcD3-ASQFSDC<Cmm*>SSAK<AcD3K*>AR<C13>-COOH

Ion score 69.93

Protein Q9QUR7

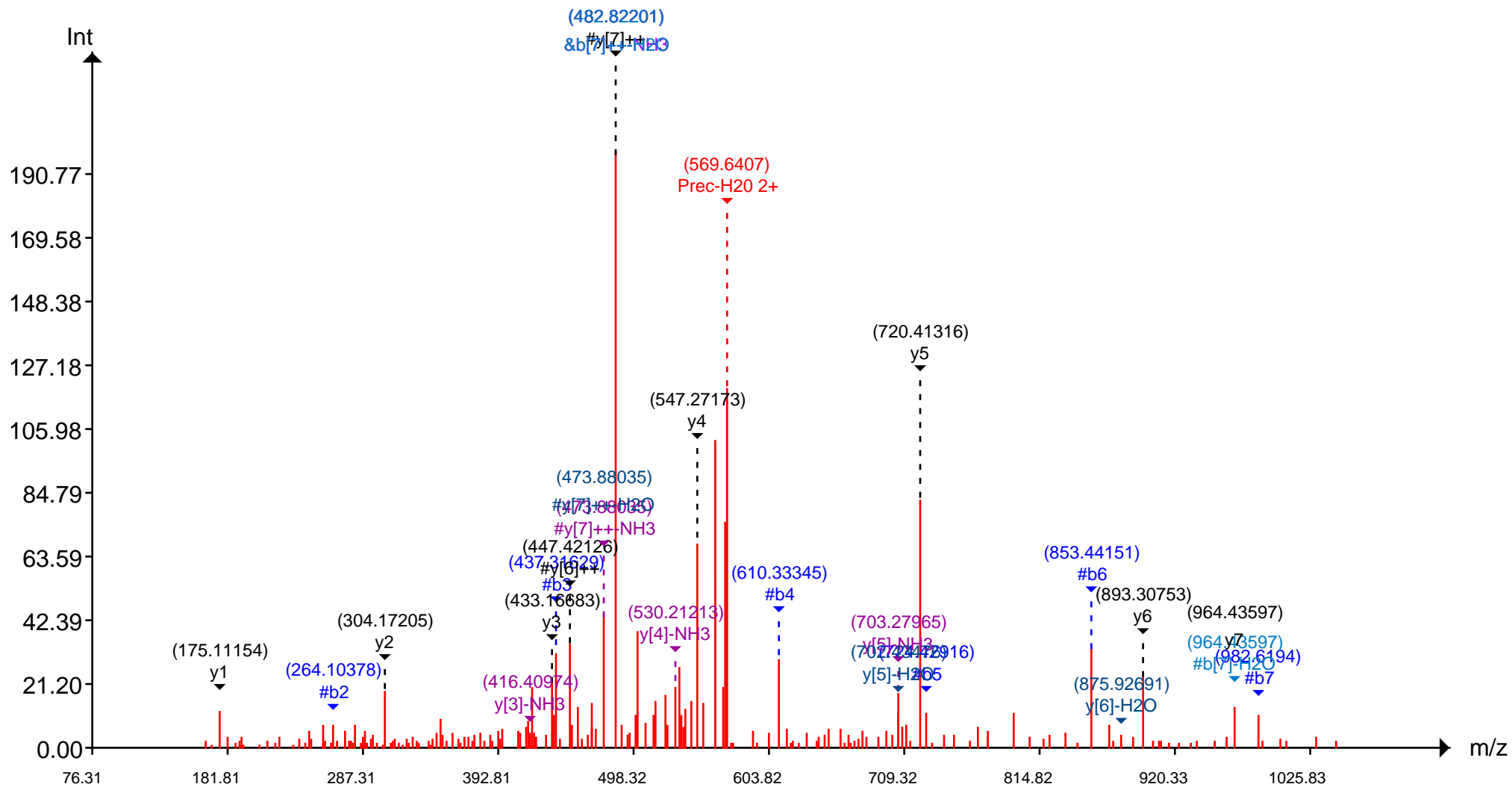
Confidence level 99.00%

Precursor m/z 755.86214

Identity Threshold 46.49

Charge 2+

Mass error 1.33E-4



Info

Modified Sequence AcD3-FAK<AcD3K*>K<AcD3K*>NEER-COOH

Ion score 48.43

Protein P32067

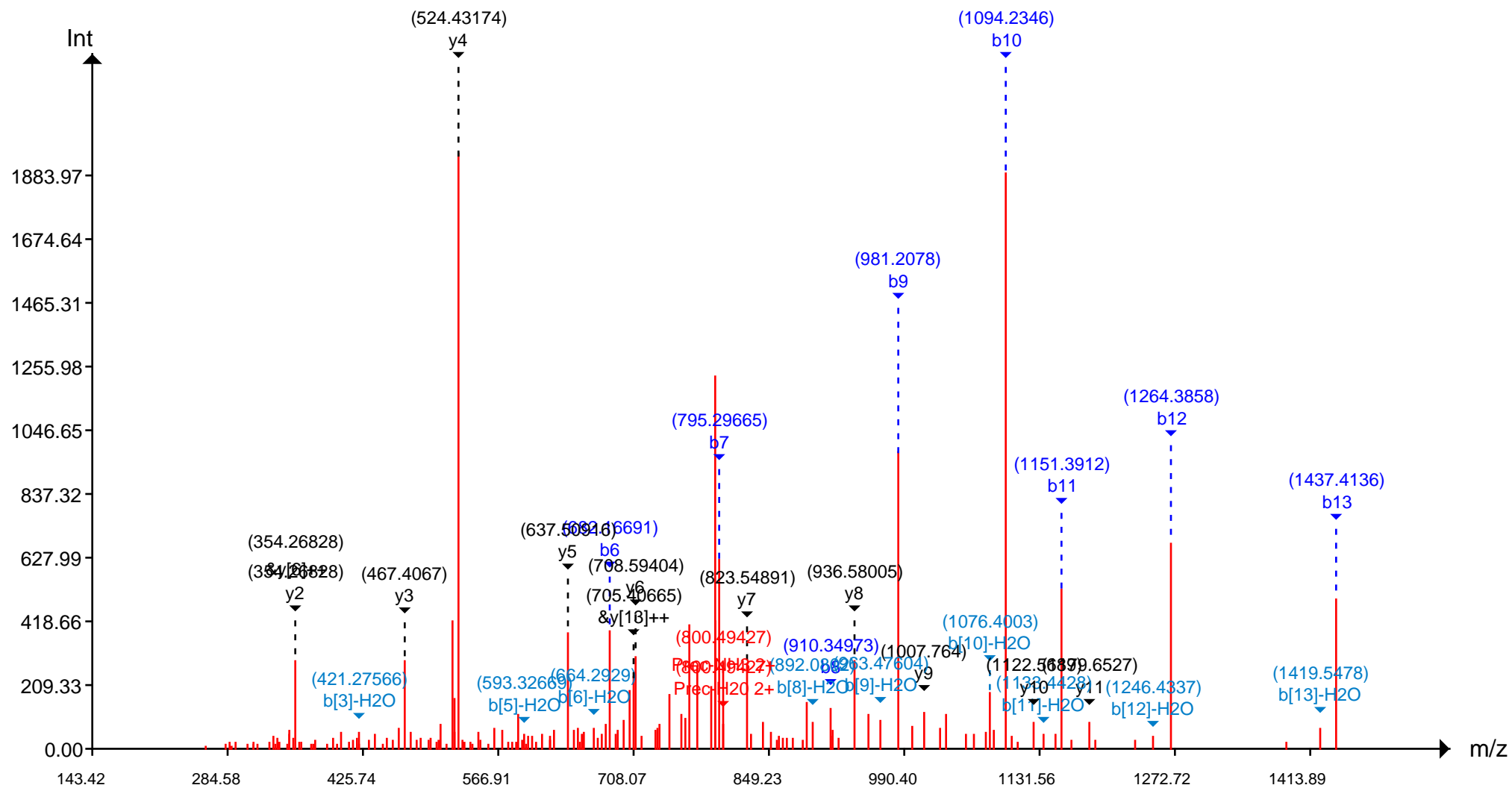
Confidence level 99.00%

Precursor m/z 578.81899

Identity Threshold 47.60

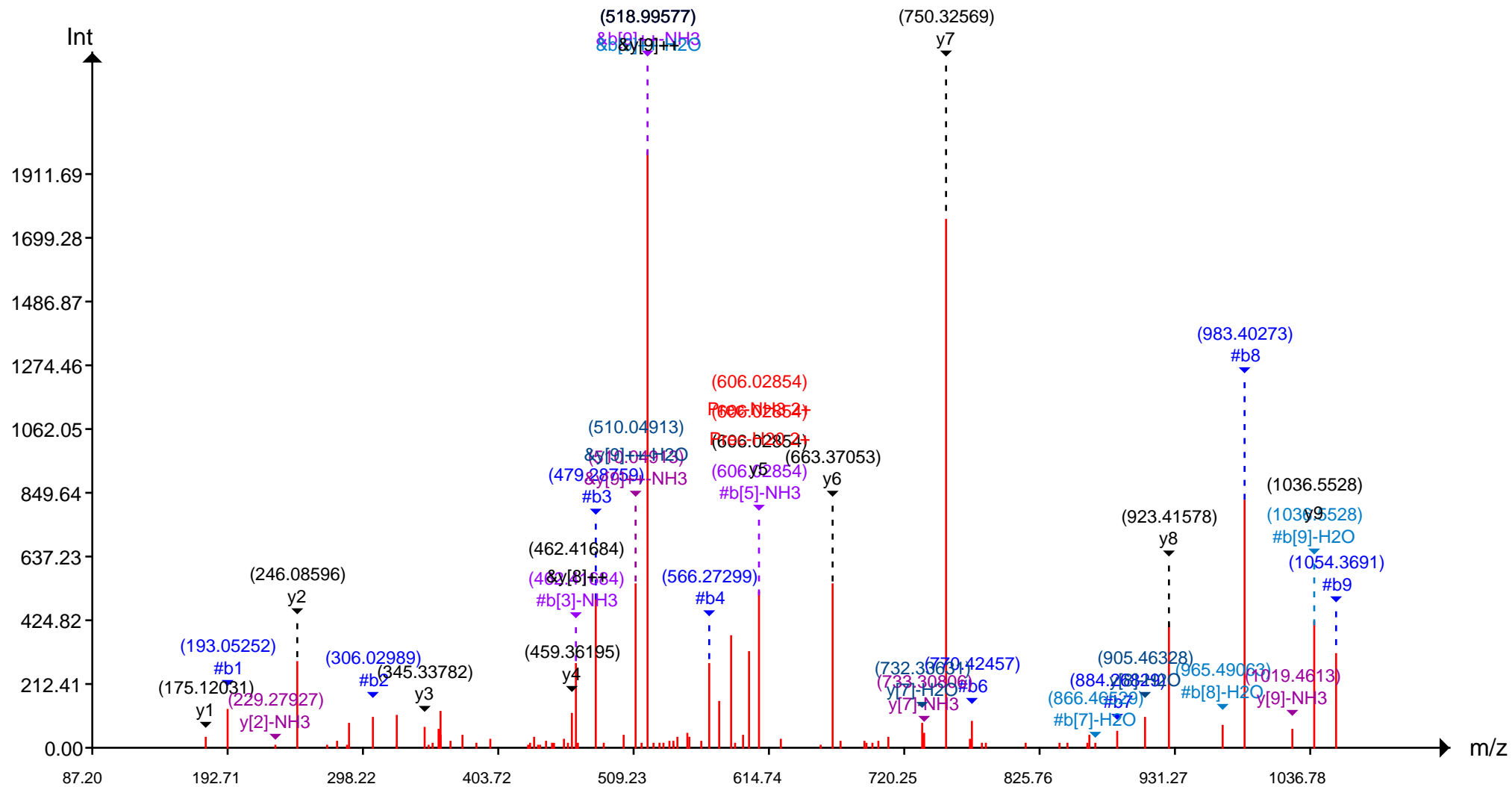
Charge 2+

Mass error 3.0E-5



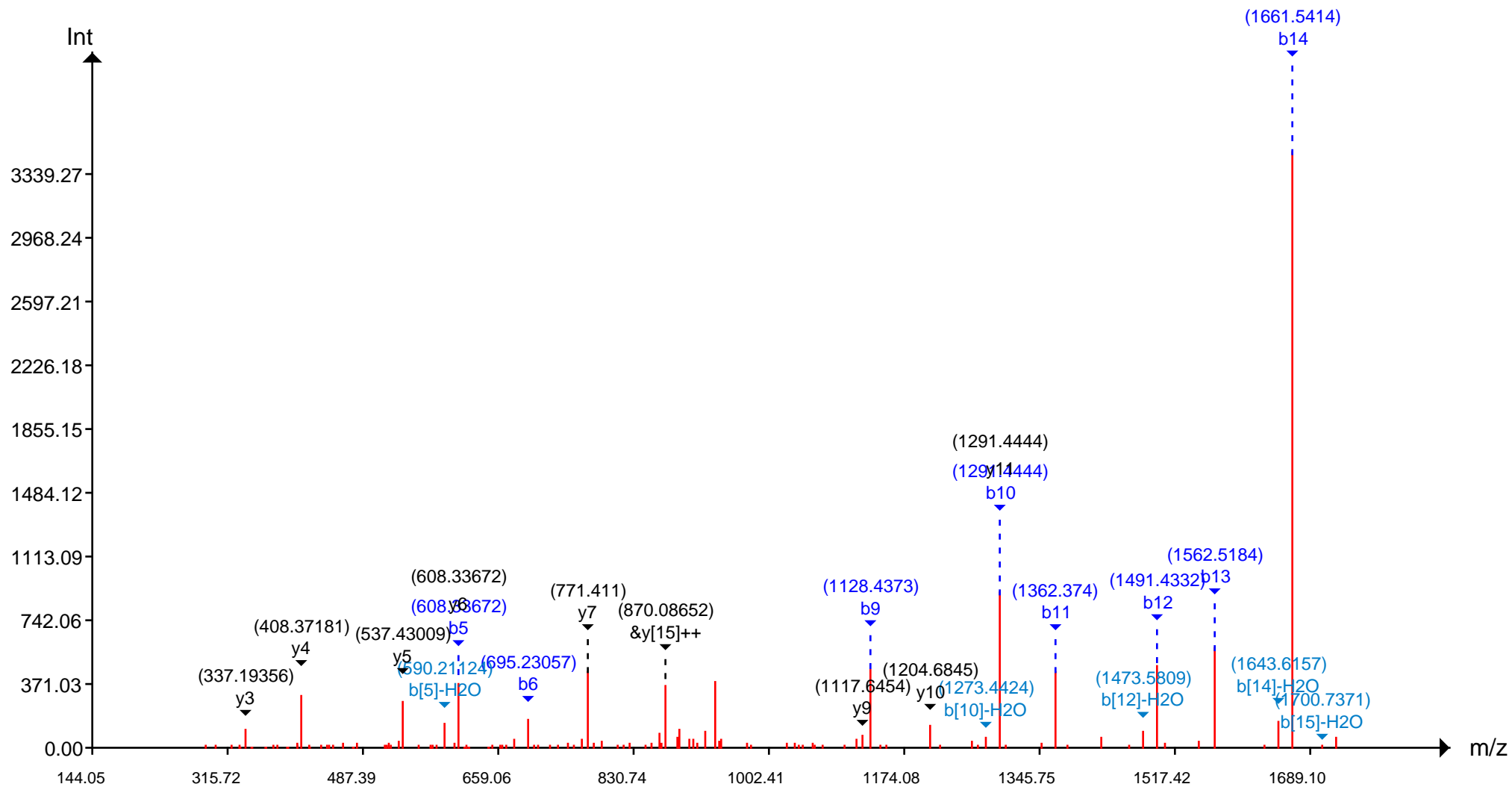
Info

Modified Sequence	AcD3-YTEGDALDALGLK<AcD3K*>R<C13>-COOH	Ion score	71.18
Protein	P62876	Confidence level	99.00%
Precursor m/z	809.43958	Identity Threshold	47.82
Charge	2+	Mass error	0.002227



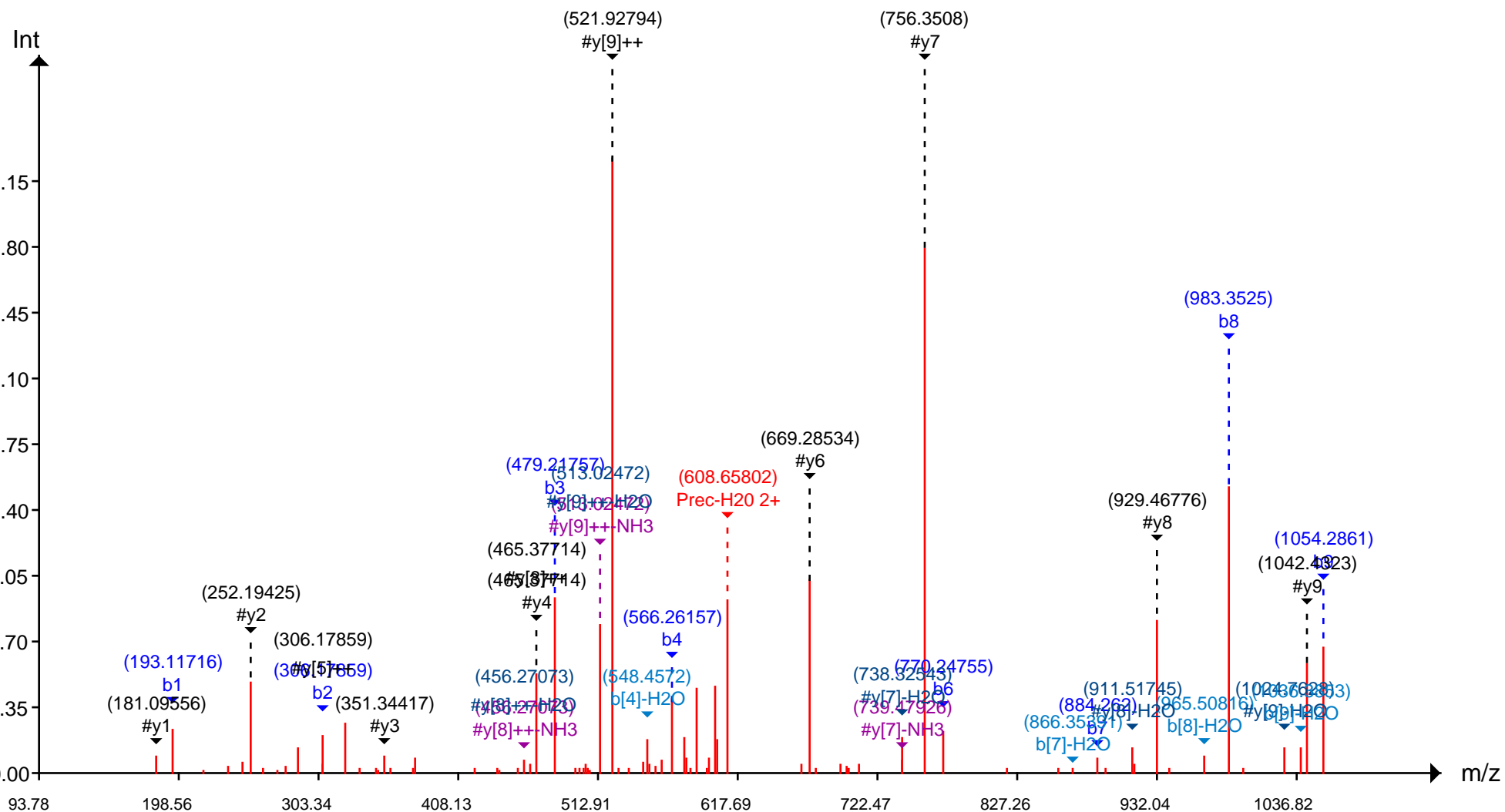
Info

Modified Sequence	AcD3-M<Mox*>IK<AcD3K*>SGM<Mox*>NVAR-COOH	Ion score	72.38
Protein	P52480	Confidence level	99.00%
Precursor m/z	614.81871	Identity Threshold	47.81
Charge	2+	Mass error	6.45E-4



Info

Modified Sequence	AcD3-DC<Cmm*>GTESSK<AcD3K*>K<AcD3K*>YAEAVGR<C13>-COOH	Ion score	59.95
Protein	P09411	Confidence level	99.00%
Precursor m/z	949.96477	Identity Threshold	47.11
Charge	2+	Mass error	0.001658



Info

Modified Sequence AcD3-M<Mox*>IK<AcD3K*>SGM<Mox*>NVAR<C13>-COOH

Ion score 63.95

Protein P52480

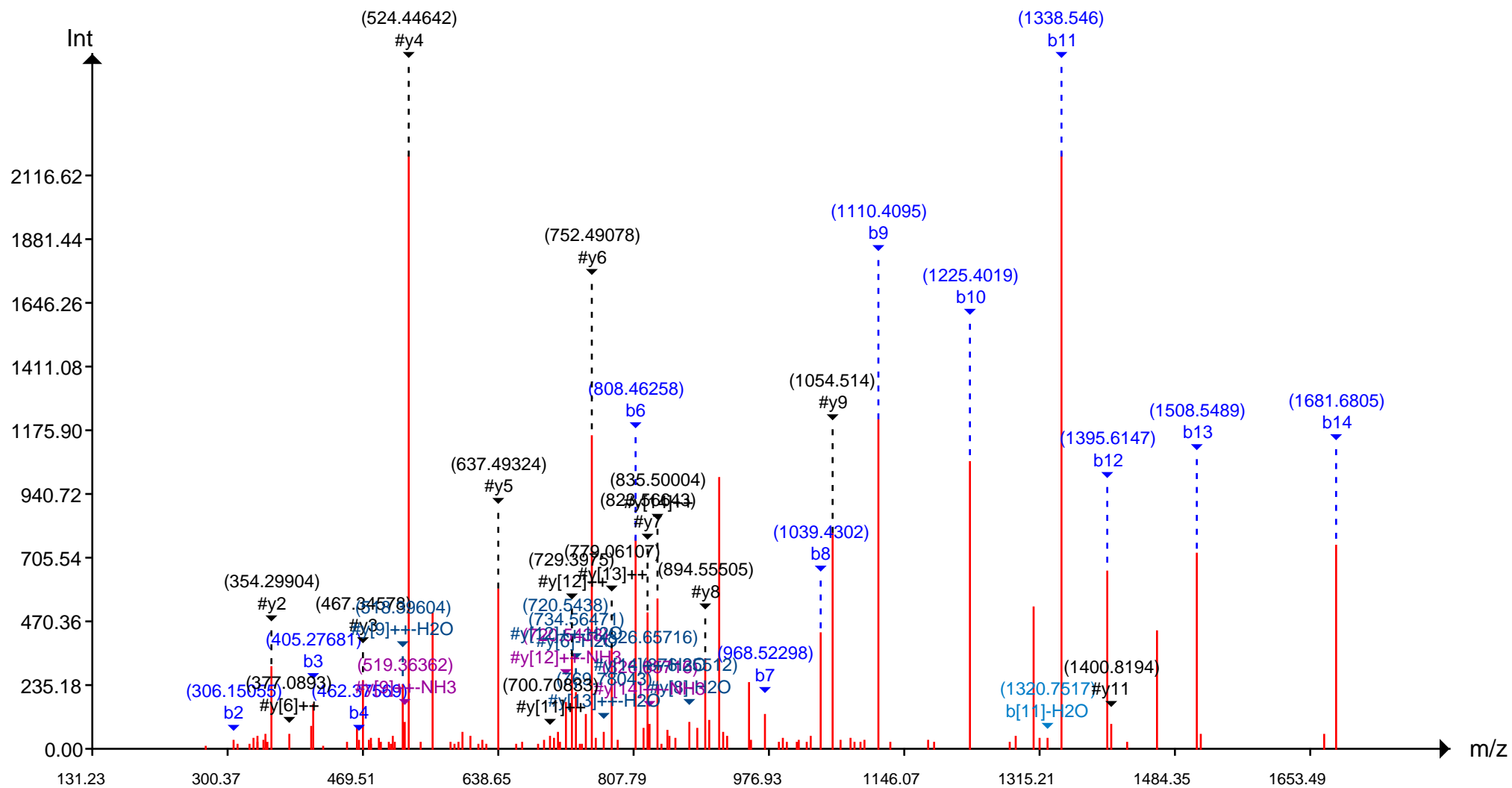
Confidence level 99.00%

Precursor m/z 617.82854

Identity Threshold 48.14

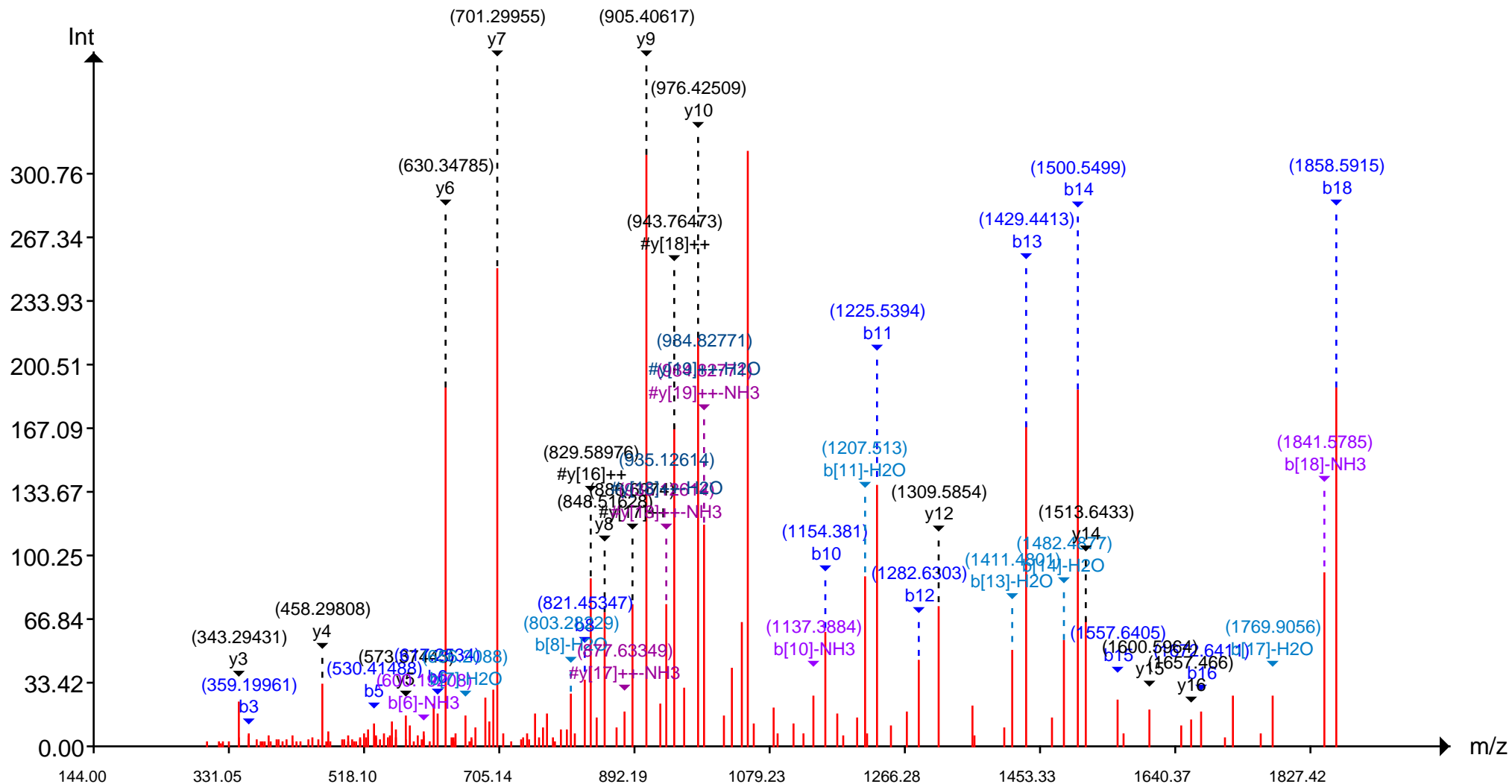
Charge 2+

Mass error 1.63E-4



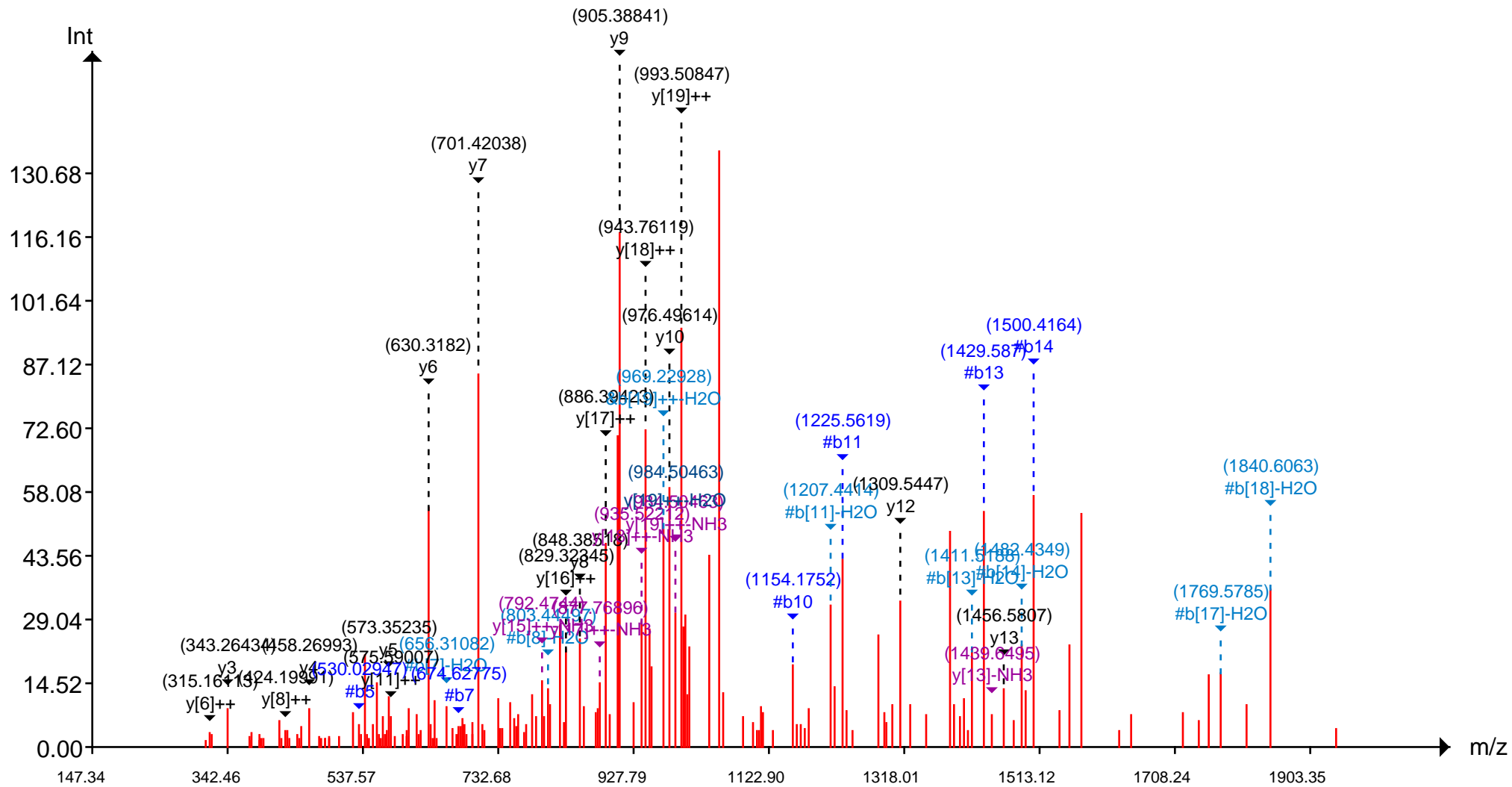
Info

Modified Sequence	AcD3-M<Mox*>IVGK<AcD3K*>K<AcD3K*>C<Cmm*>AADLGLK<AcD3K*>R<C13>-COOH	Ion score	69.33
Protein	P70349	Confidence level	99.00%
Precursor m/z	931.54029	Identity Threshold	44.73
Charge	2+	Mass error	9.95E-4



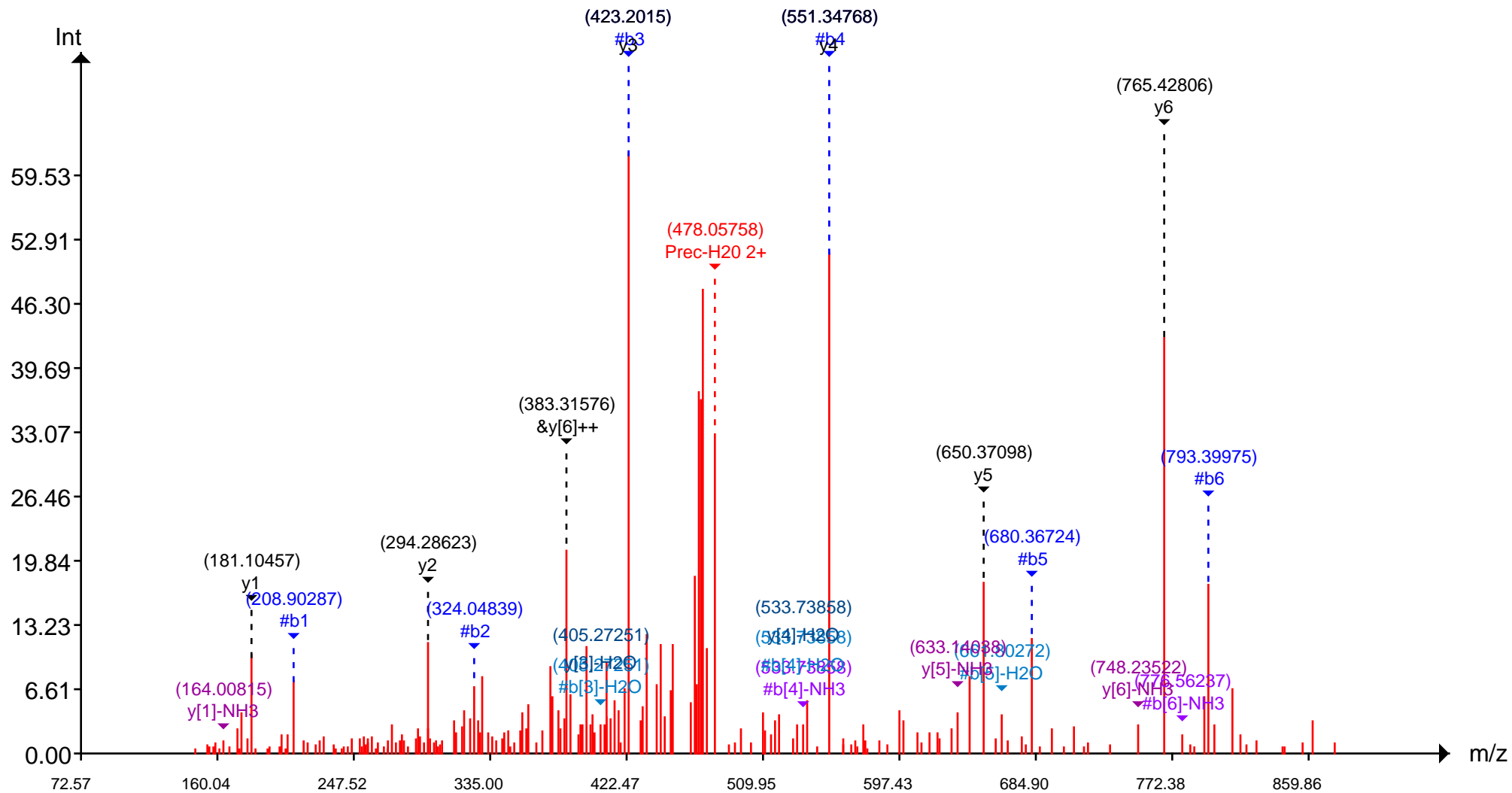
Info

Modified Sequence	AcD3-VVDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR-COOH	Ion score	92.22
Protein	P60710	Confidence level	99.00%
Precursor m/z	1065.4799	Identity Threshold	43.52
Charge	2+	Mass error	0.002758



Info

Modified Sequence	AcD3-VVDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR-COOH	Ion score	48.19
Protein	P60710	Confidence level	99.00%
Precursor m/z	1065.4794	Identity Threshold	43.45
Charge	2+	Mass error	0.001758



Info

Modified Sequence AcD3-YDVQELR<C13>-COOH

Ion score 52.39

Protein P97429

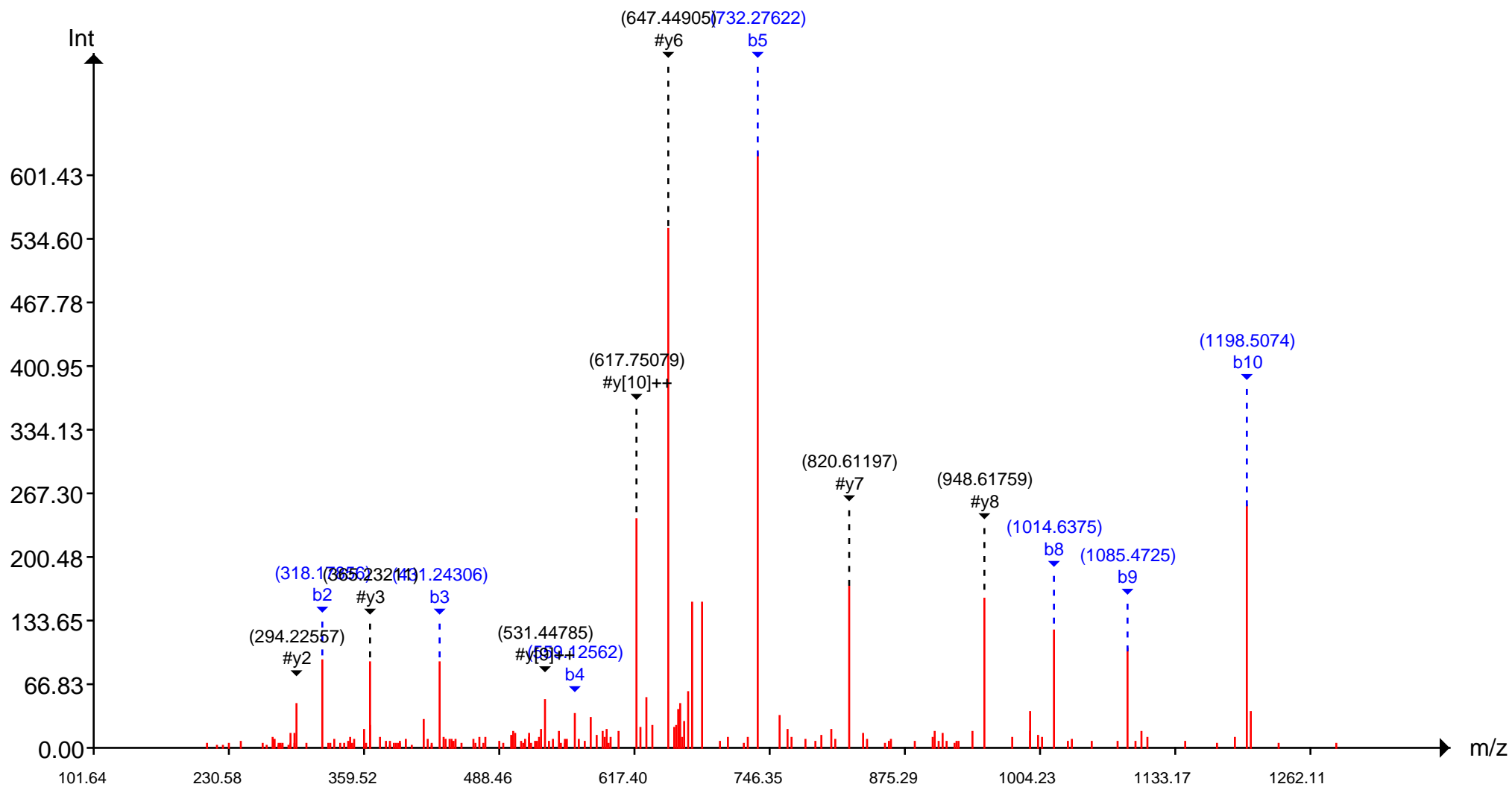
Confidence level 99.00%

Precursor m/z 487.25924

Identity Threshold 47.72

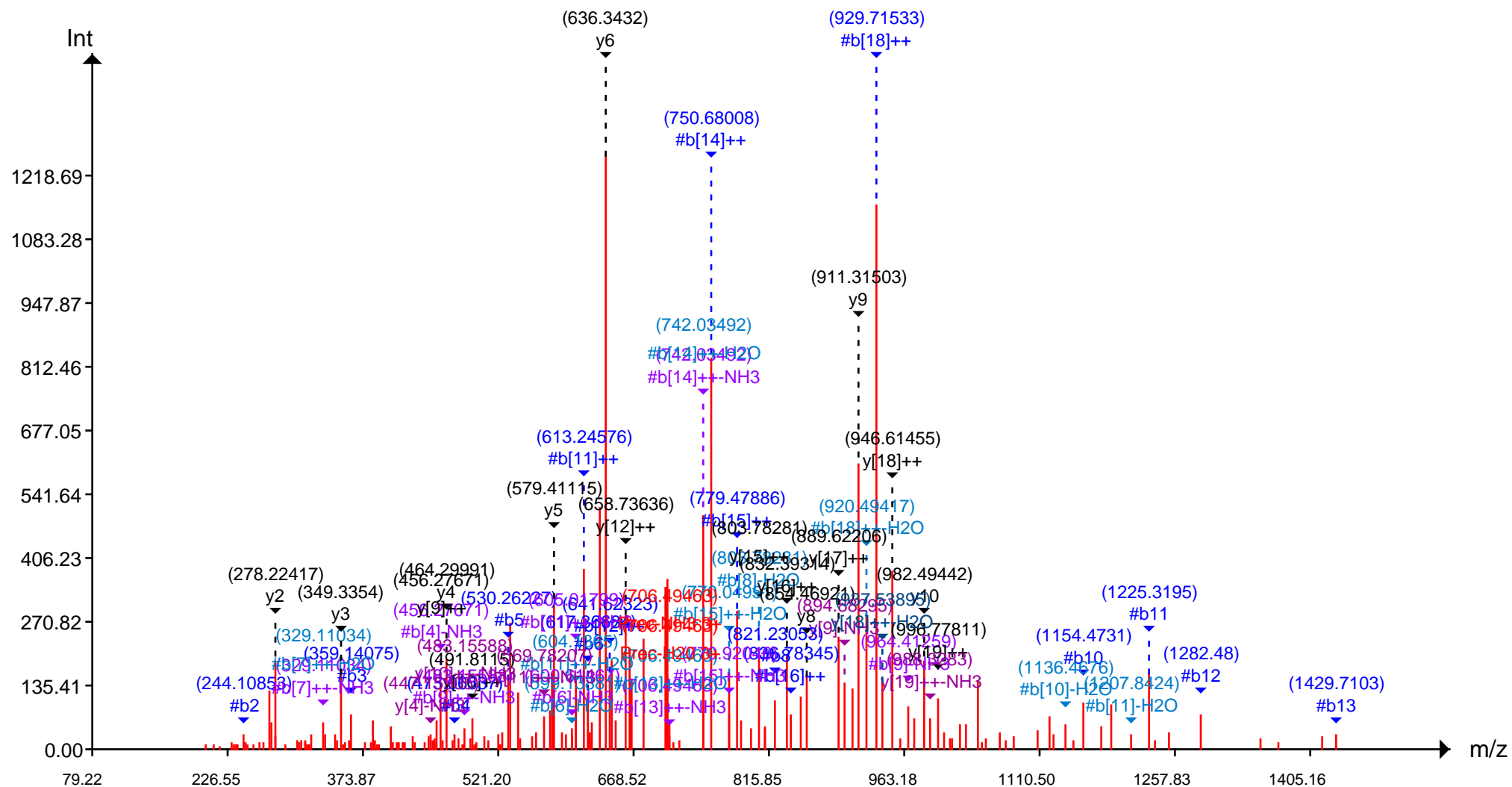
Charge 2+

Mass error -0.001184



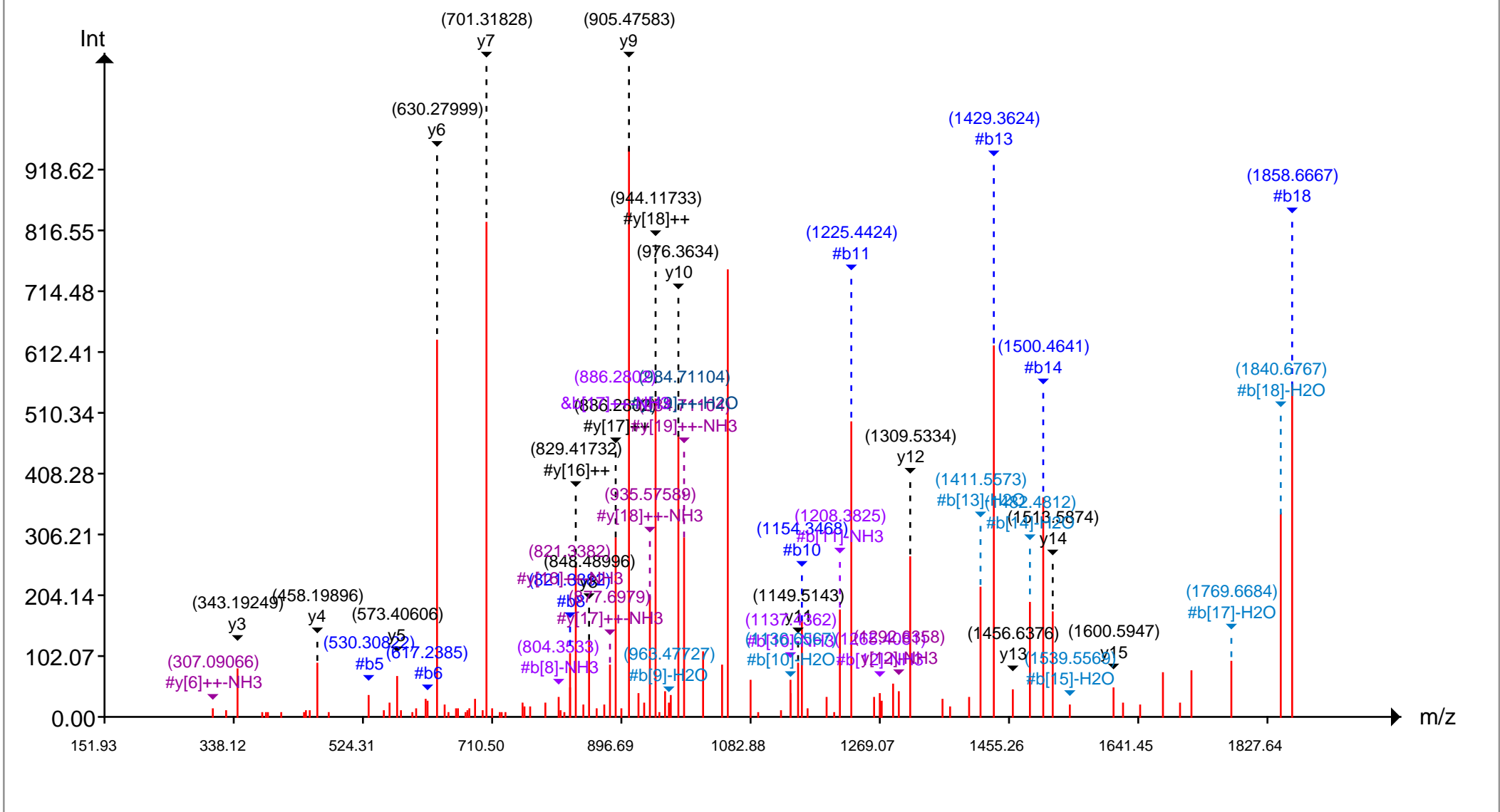
Info

Modified Sequence	AcD3-VK<AcD3K*>LQK<AcD3K*>PNAAIR<C13>-COOH	Ion score	48.62
Protein	Q99L47	Confidence level	99.00%
Precursor m/z	689.94542	Identity Threshold	37.28
Charge	2+	Mass error	0.001349



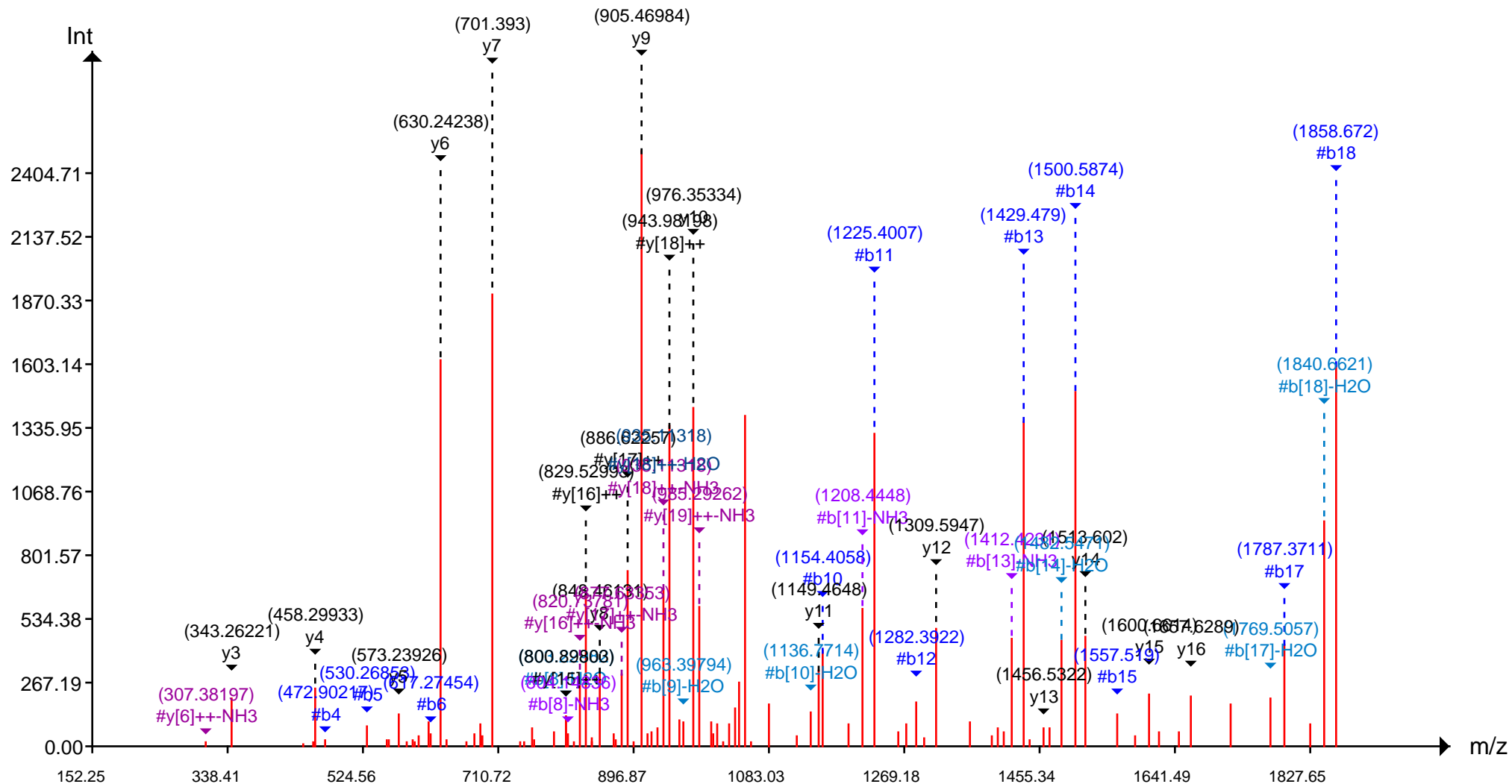
Info

Modified Sequence	AcD3-VVDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH	Ion score	49.96
Protein	P60710	Confidence level	99.00%
Precursor m/z	712.66152	Identity Threshold	44.21
Charge	3+	Mass error	1.01E-4



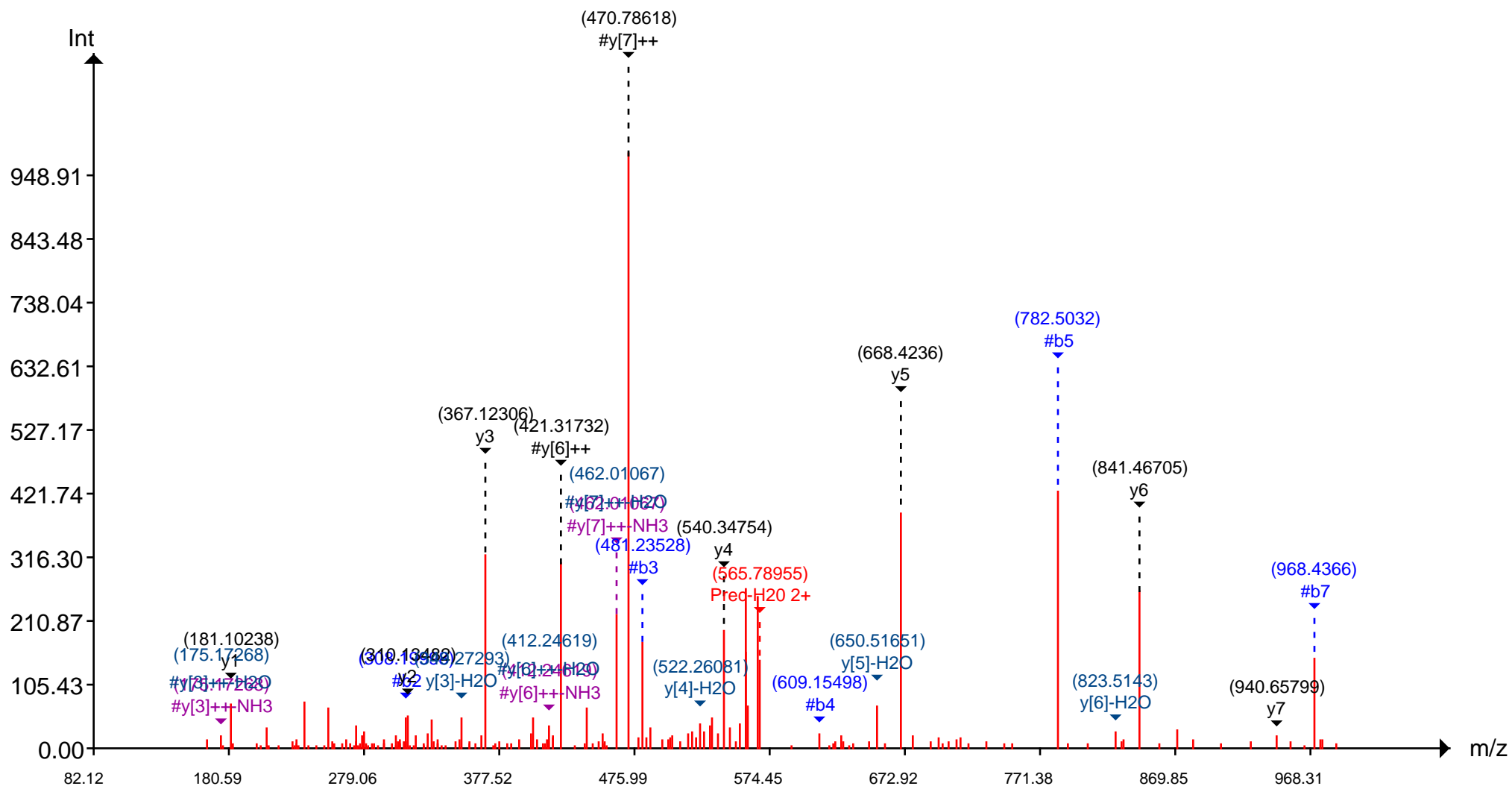
Info

Modified Sequence	AcD3-VVDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR-COOH	Ion score	88.53
Protein	P60710	Confidence level	99.00%
Precursor m/z	1065.4802	Identity Threshold	43.55
Charge	2+	Mass error	0.003358



Info

Modified Sequence	AcD3-VVDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR-COOH	Ion score	113.99
Protein	P60710	Confidence level	99.00%
Precursor m/z	1065.4803	Identity Threshold	43.54
Charge	2+	Mass error	0.003558



Info

Modified Sequence AcD3-YVK<AcD3K*>QK<AcD3K*>GER<C13>-COOH

Protein P26039

Precursor m/z 574.83927

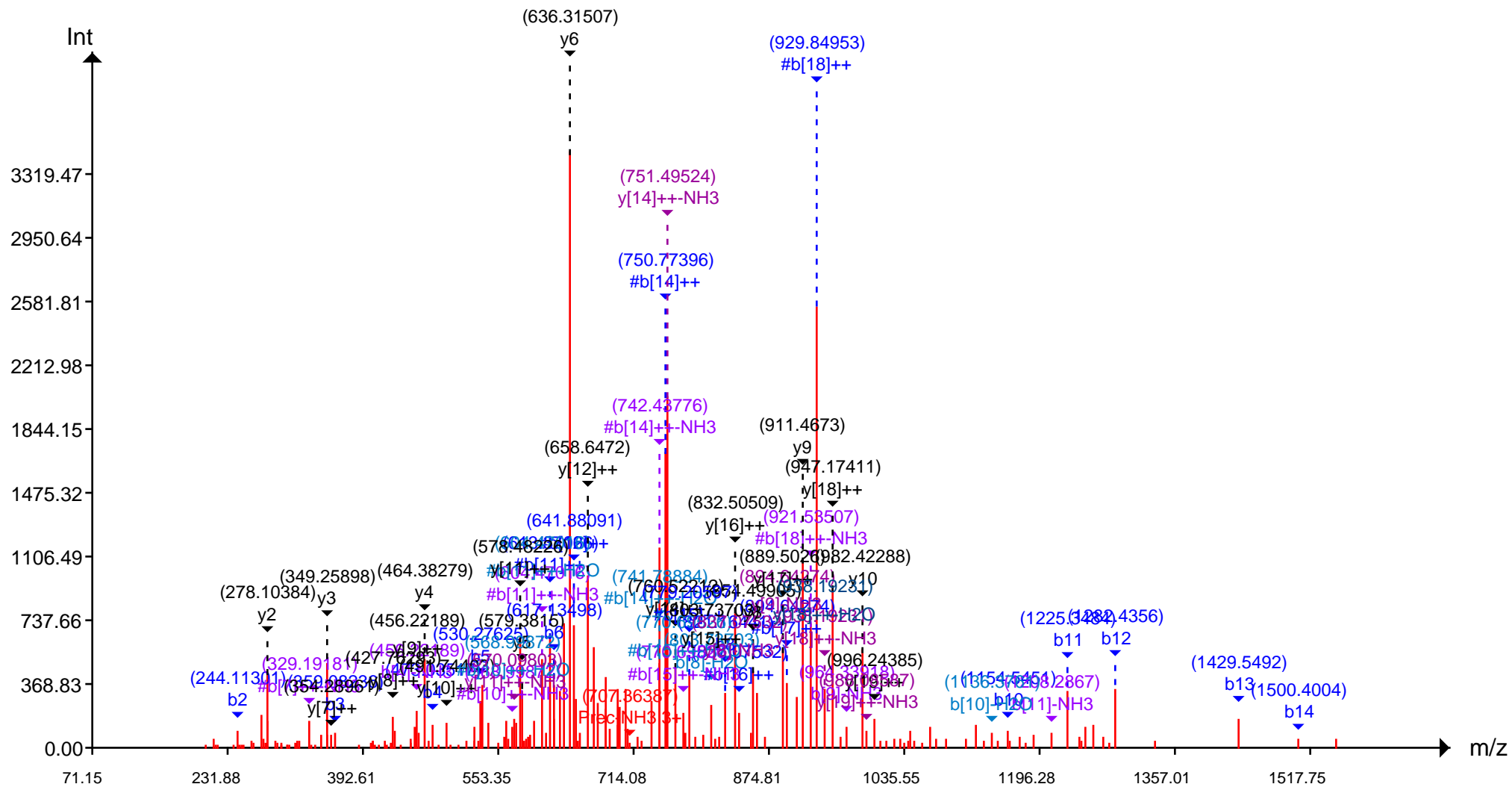
Charge 2+

Ion score 47.61

Confidence level 99.00%

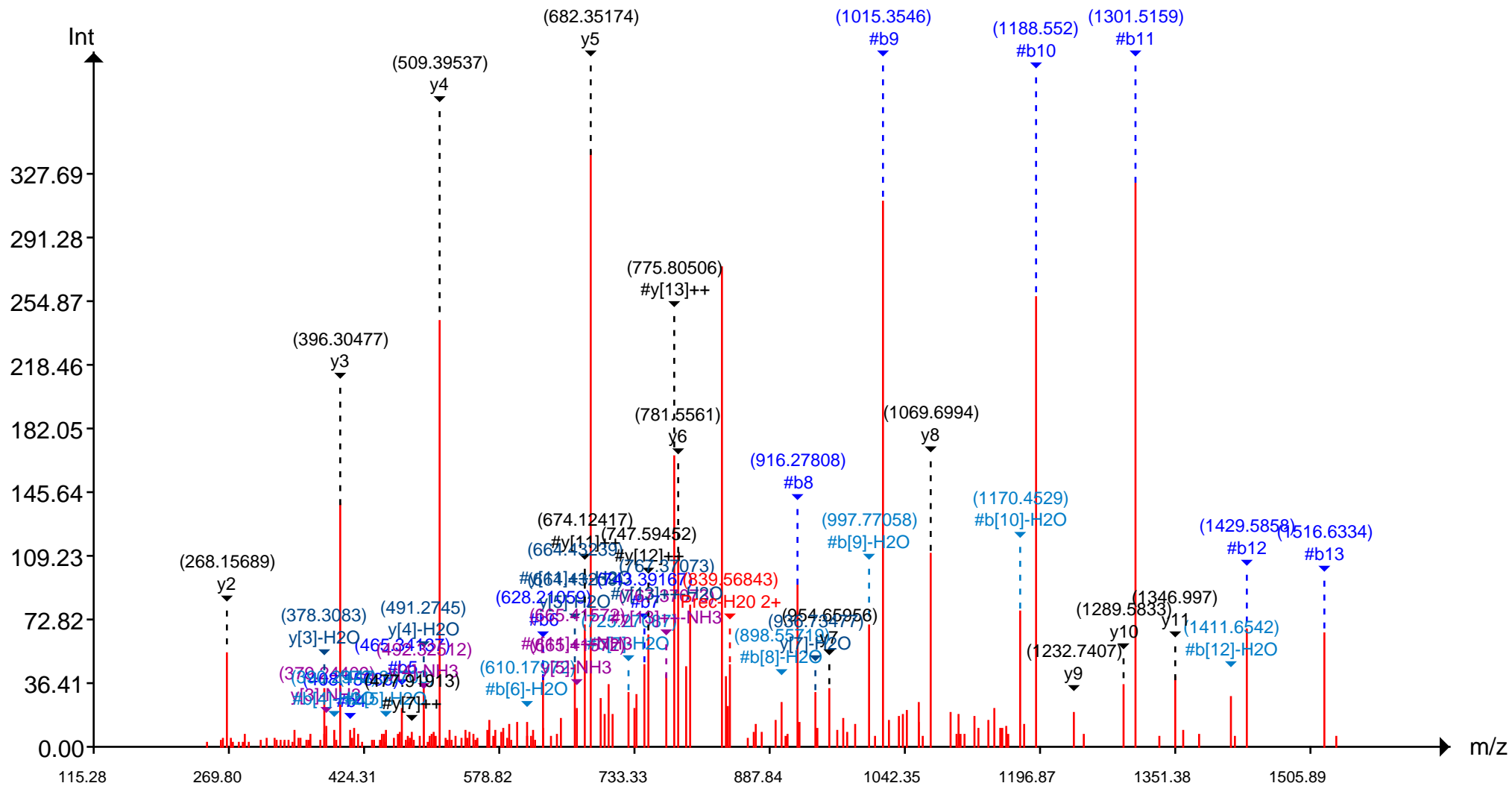
Identity Threshold 46.31

Mass error -3.03E-4



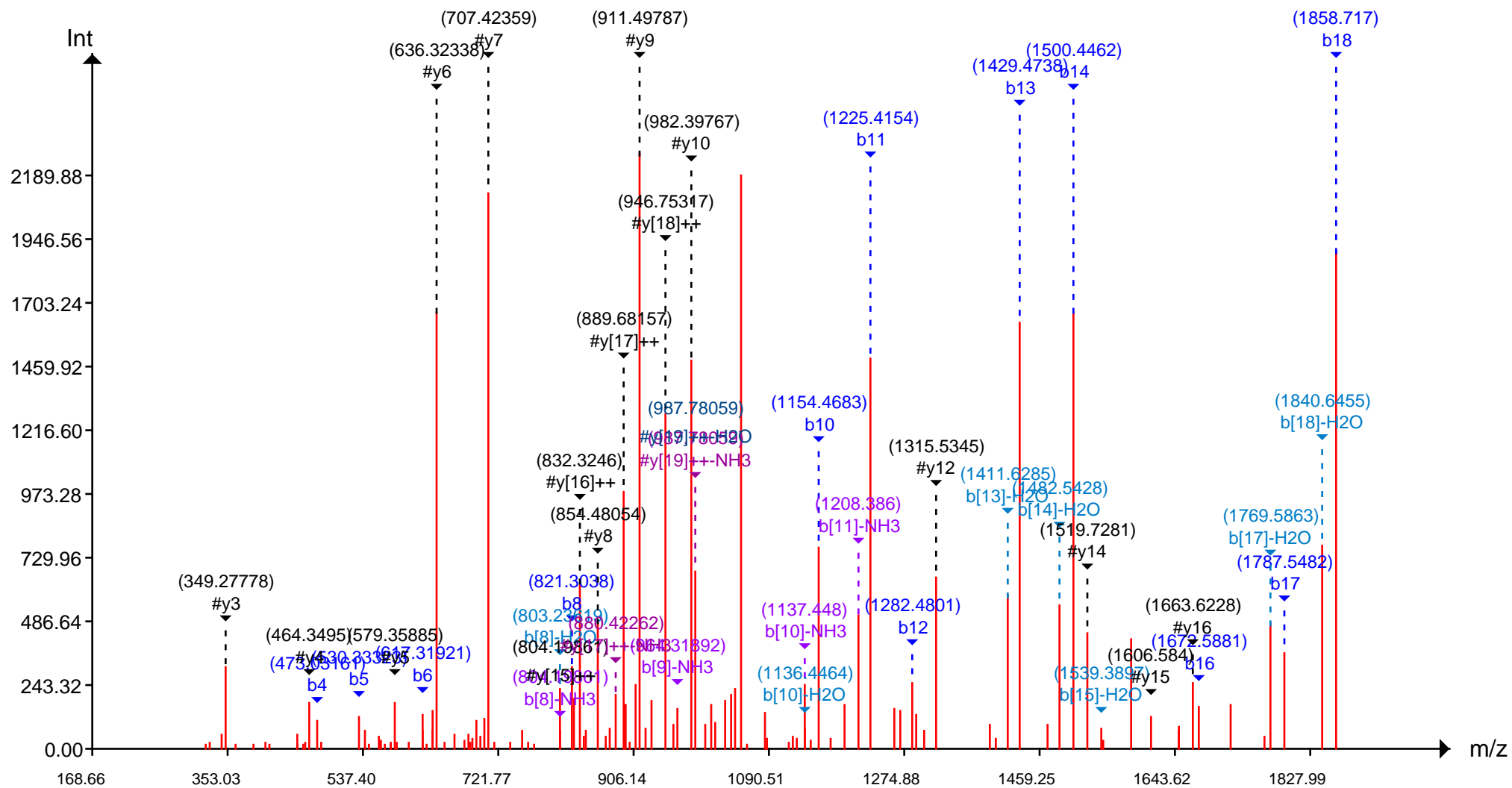
Info

Modified Sequence	AcD3-VVDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH	Ion score	74.11
Protein	P60710	Confidence level	99.00%
Precursor m/z	712.66197	Identity Threshold	44.26
Charge	3+	Mass error	0.001451



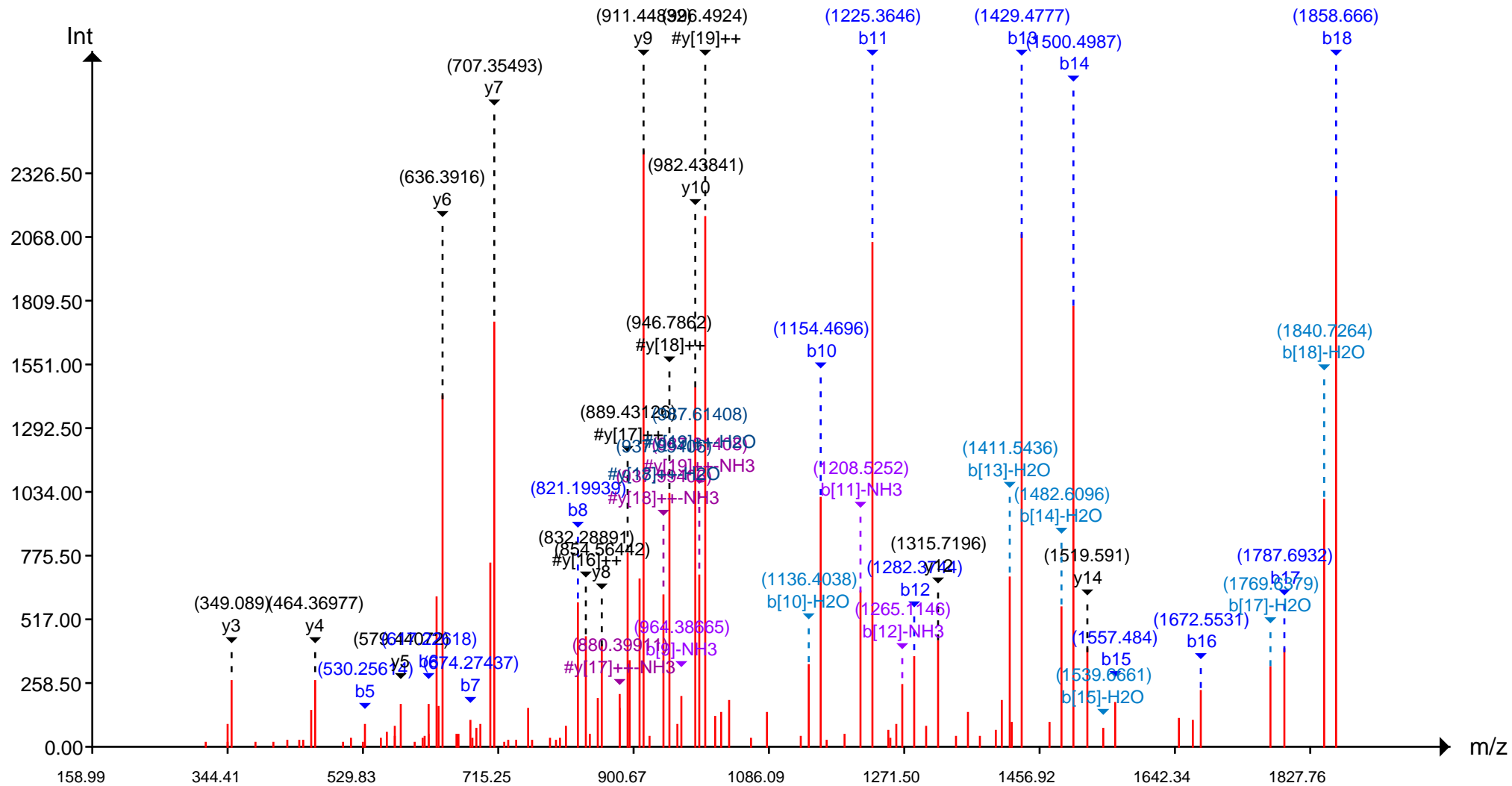
Info

Modified Sequence	AcD3-TGFGGYDK<AcD3K*>VK<AcD3K*>LQSR<C13>-COOH	Ion score	81.89
Protein	Q62465	Confidence level	99.00%
Precursor m/z	848.97029	Identity Threshold	47.34
Charge	2+	Mass error	0.002246



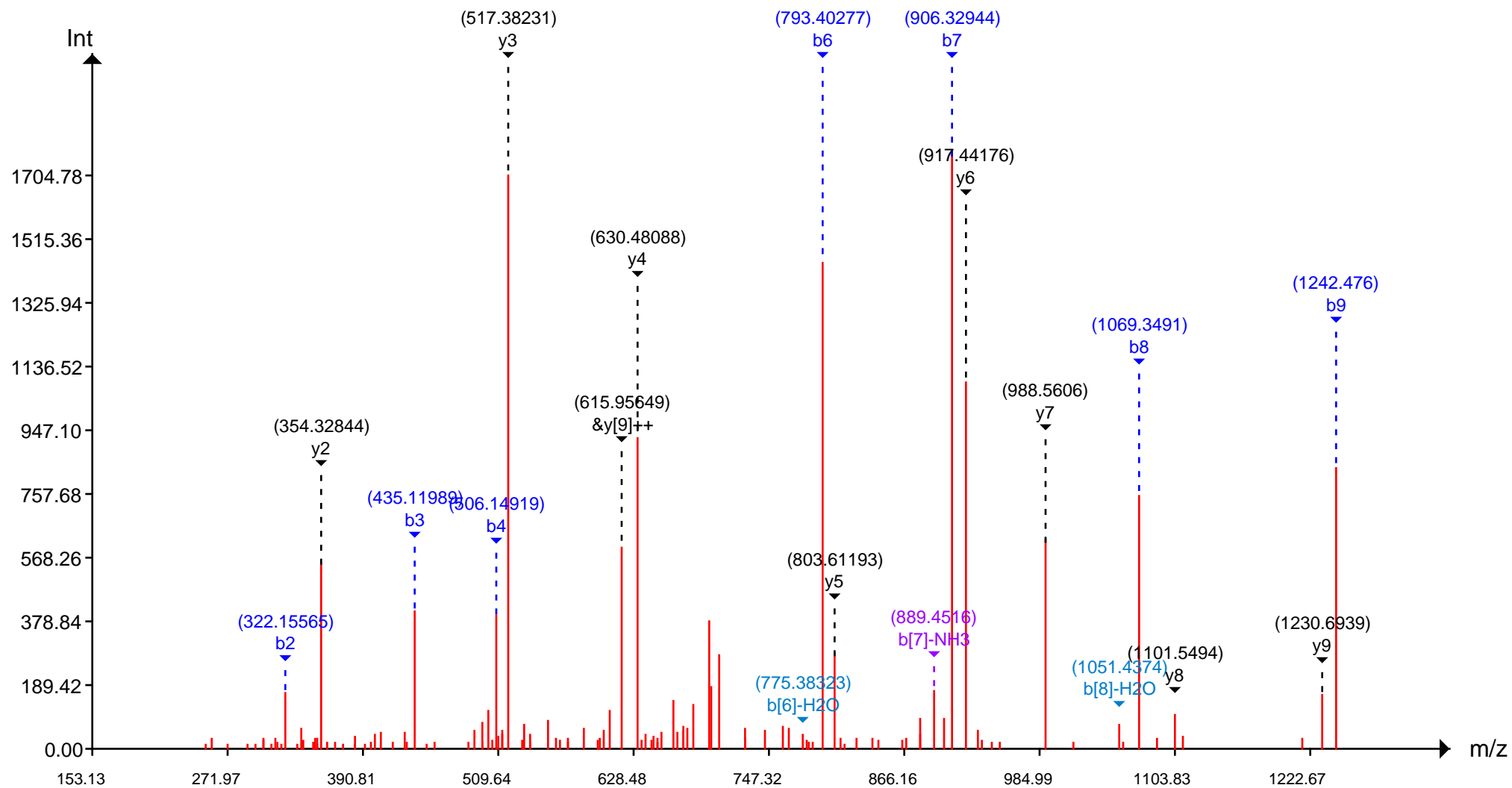
Info

Modified Sequence	AcD3-VVDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH	Ion score	95.84
Protein	P60710	Confidence level	99.00%
Precursor m/z	1068.4893	Identity Threshold	44.25
Charge	2+	Mass error	0.001417



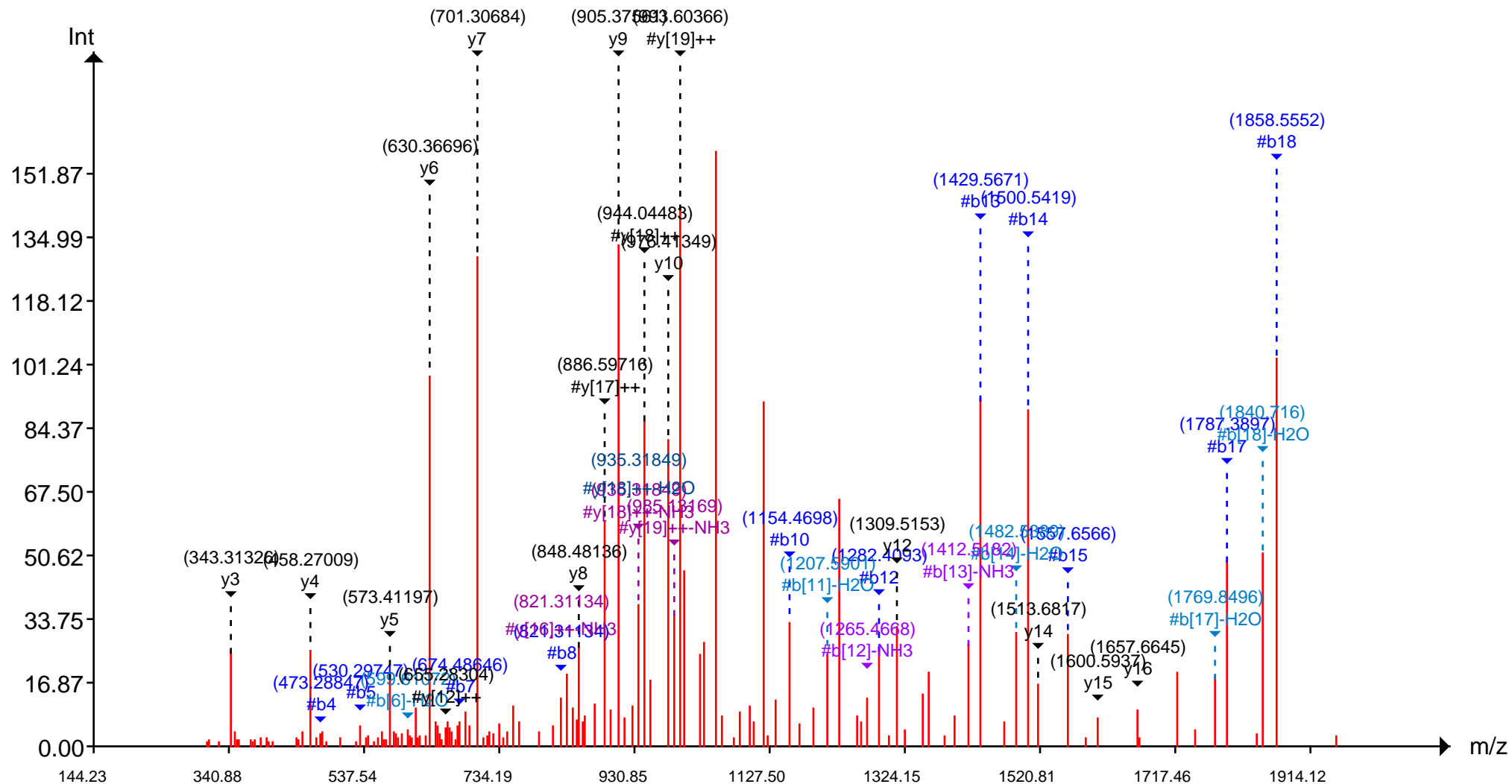
Info

Modified Sequence	AcD3-VVDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH	Ion score	84.72
Protein	P60710	Confidence level	99.00%
Precursor m/z	1068.4899	Identity Threshold	44.40
Charge	2+	Mass error	0.002617



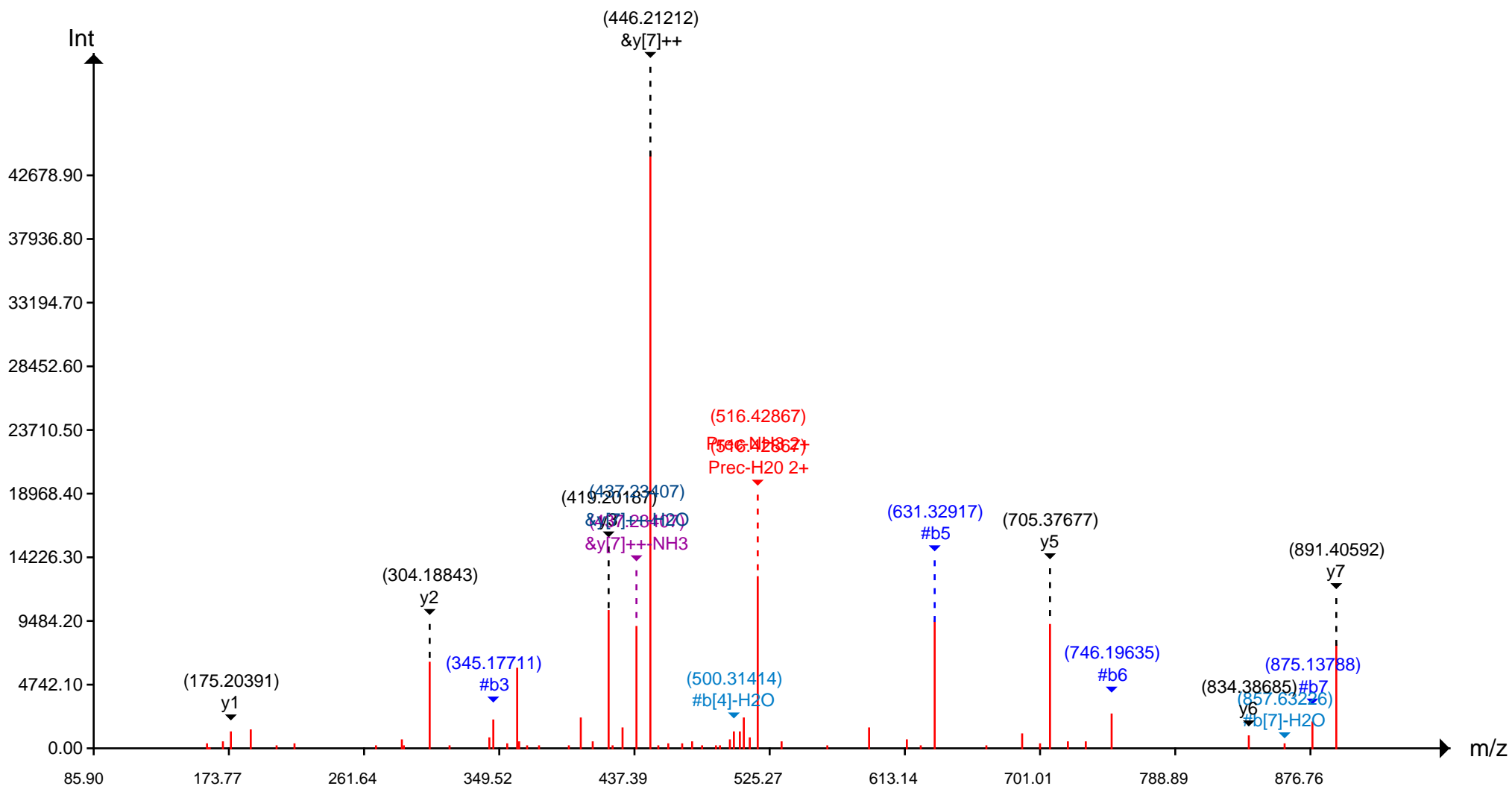
Info

Modified Sequence	AcD3-M<Mox*>ELANK<AcD3K*>IYK<AcD3K*>R<C13>-COOH	Ion score	73.51
Protein	Q920E5	Confidence level	99.00%
Precursor m/z	711.90765	Identity Threshold	46.45
Charge	2+	Mass error	0.001371



Info

Modified Sequence	AcD3-VVDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR-COOH	Ion score	81.46
Protein	P60710	Confidence level	99.00%
Precursor m/z	1065.4798	Identity Threshold	43.52
Charge	2+	Mass error	0.002558



Info

Modified Sequence AcD3-IGEK<AcD3K*>LDER-COOH

Ion score 50.57

Protein P17751

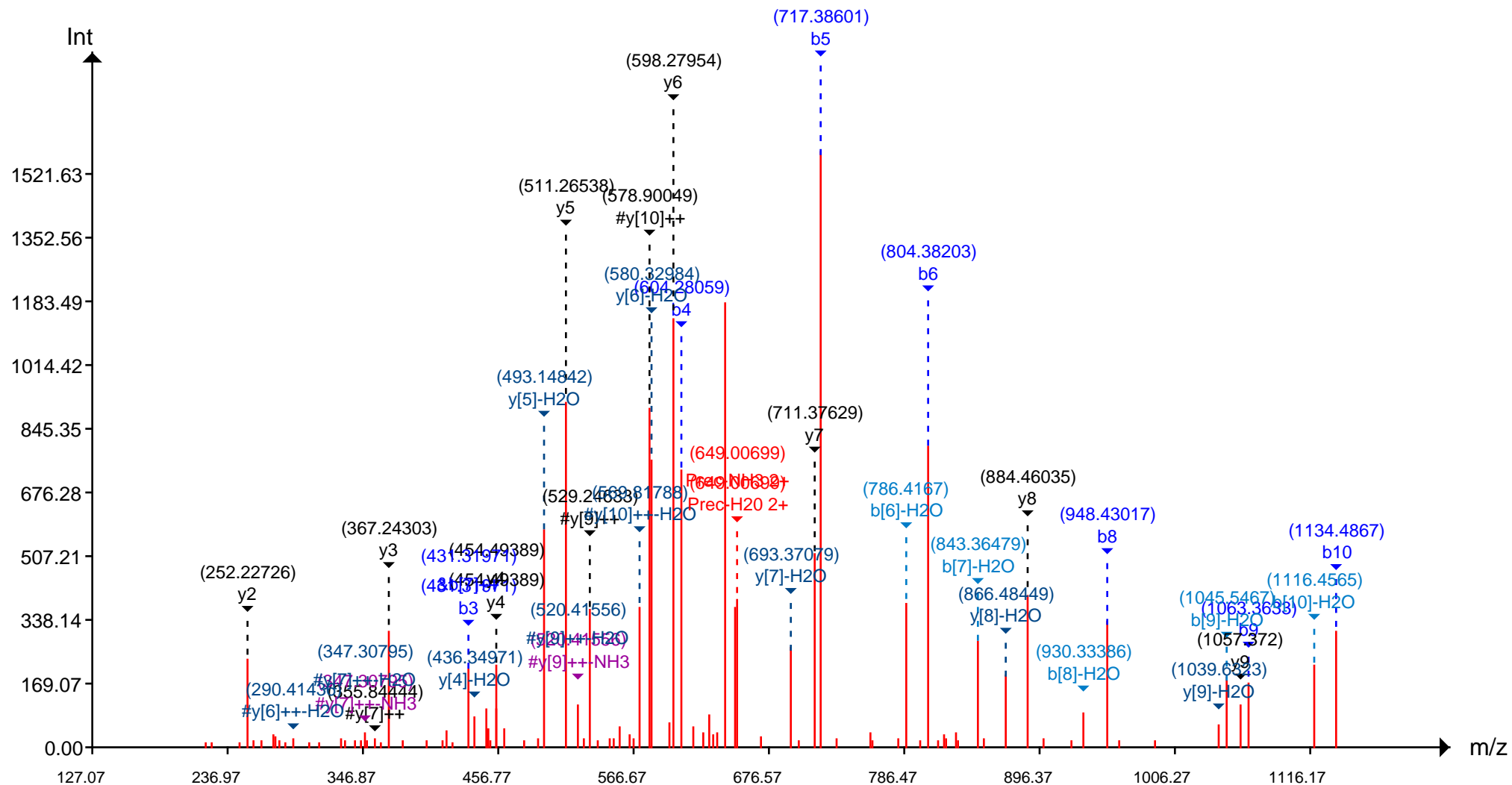
Confidence level 99.00%

Precursor m/z 525.29114

Identity Threshold 47.73

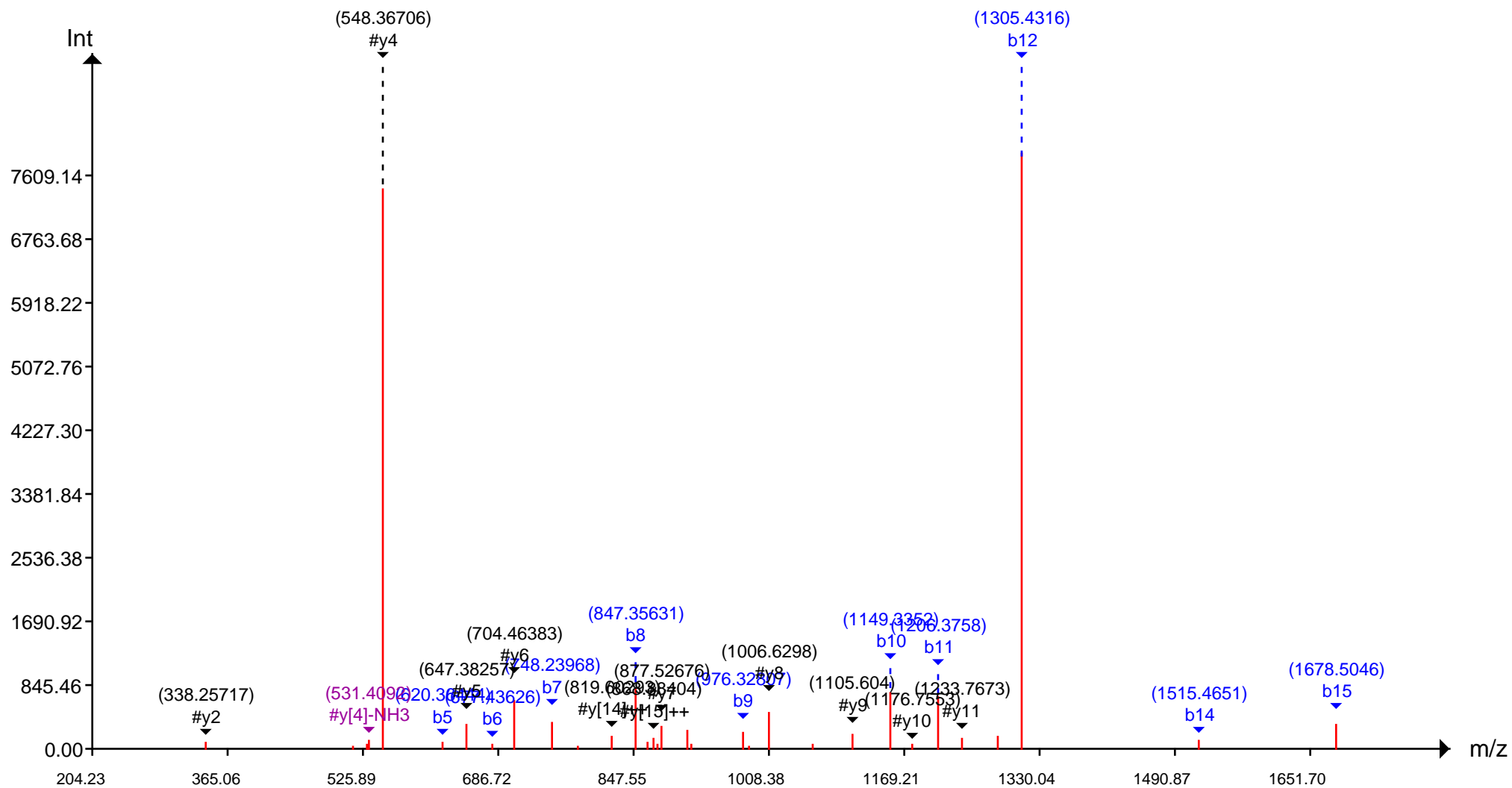
Charge 2+

Mass error 6.05E-4



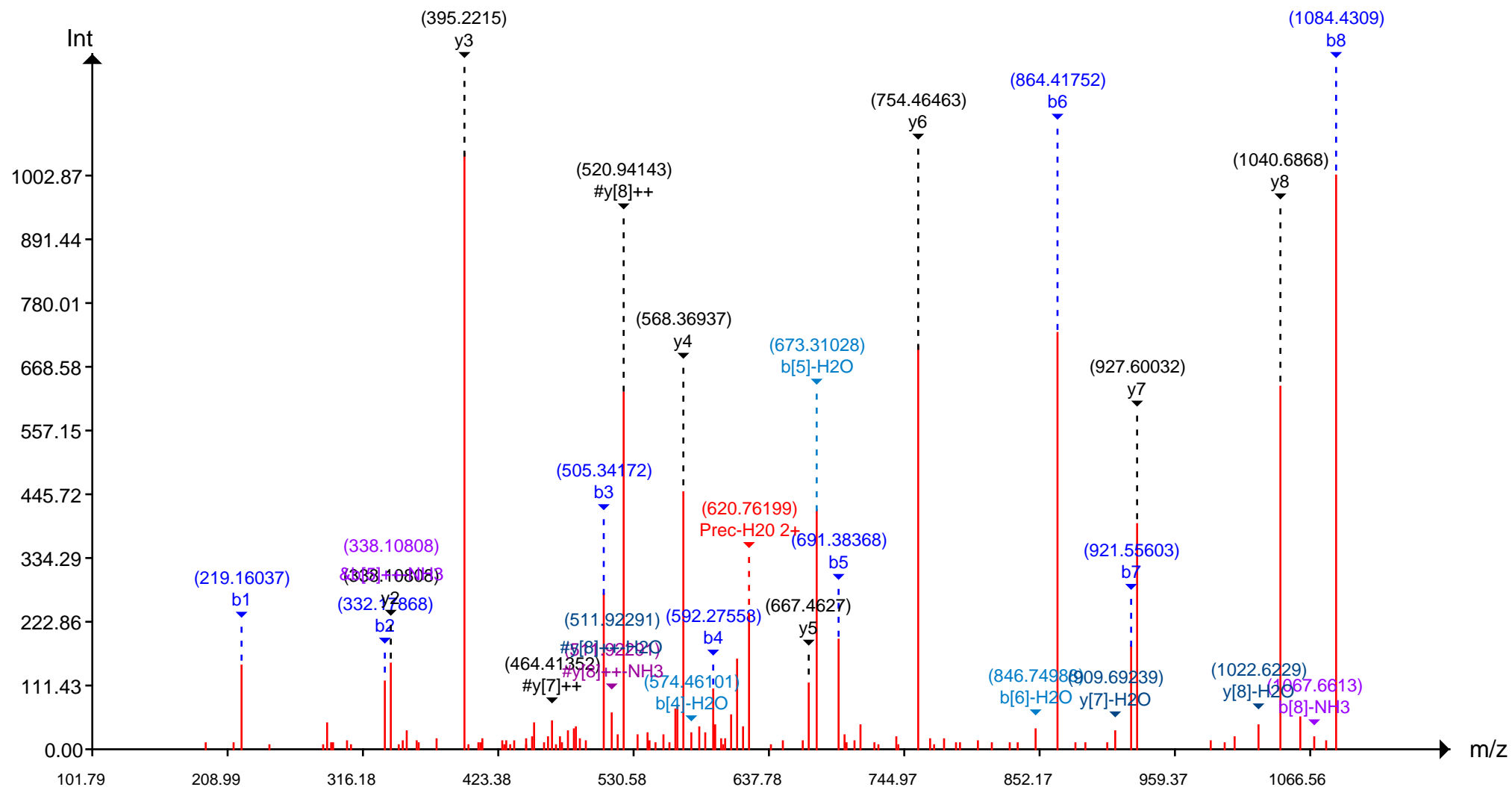
Info

Modified Sequence	AcD3-LVK<AcD3K*>K<AcD3K*>ISGSDAR<C13>-COOH	Ion score	64.74
Protein	P18654	Confidence level	99.00%
Precursor m/z	657.90565	Identity Threshold	42.87
Charge	2+	Mass error	7.12E-4



Info

Modified Sequence	AcD3-AVC<Cmm*>K<AcD3K*>AGAVEK<AcD3K*>GVPLYR-COOH	Ion score	82.21
Protein	P17182	Confidence level	99.00%
Precursor m/z	927.01964	Identity Threshold	46.51
Charge	2+	Mass error	0.002038



Info

Modified Sequence AcD3-K<AcD3K*>LK<AcD3K*>SVK<AcD3K*>GYR-COOH

Ion score 65.58

Protein P21981

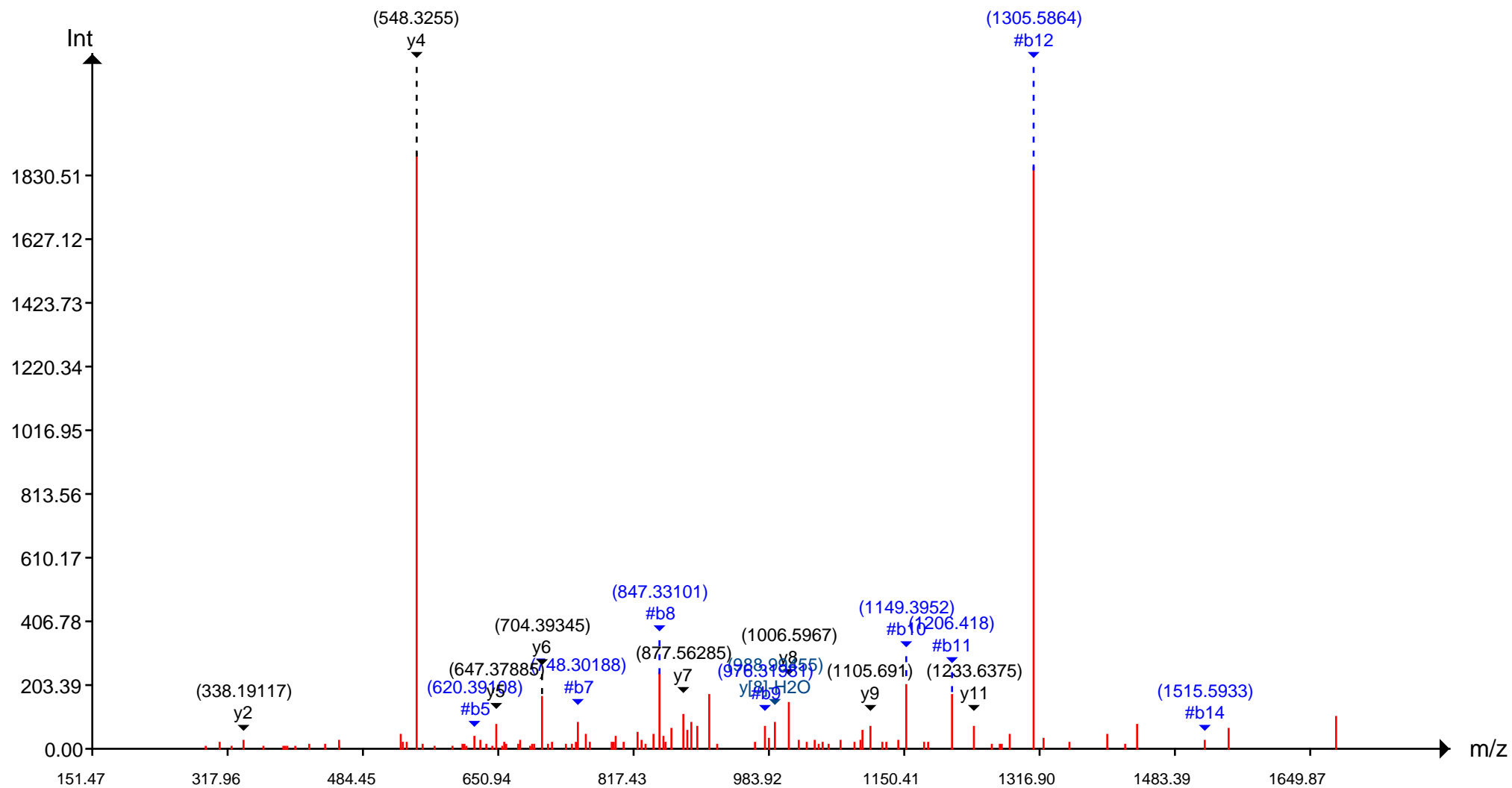
Confidence level 99.00%

Precursor m/z 629.89923

Identity Threshold 41.29

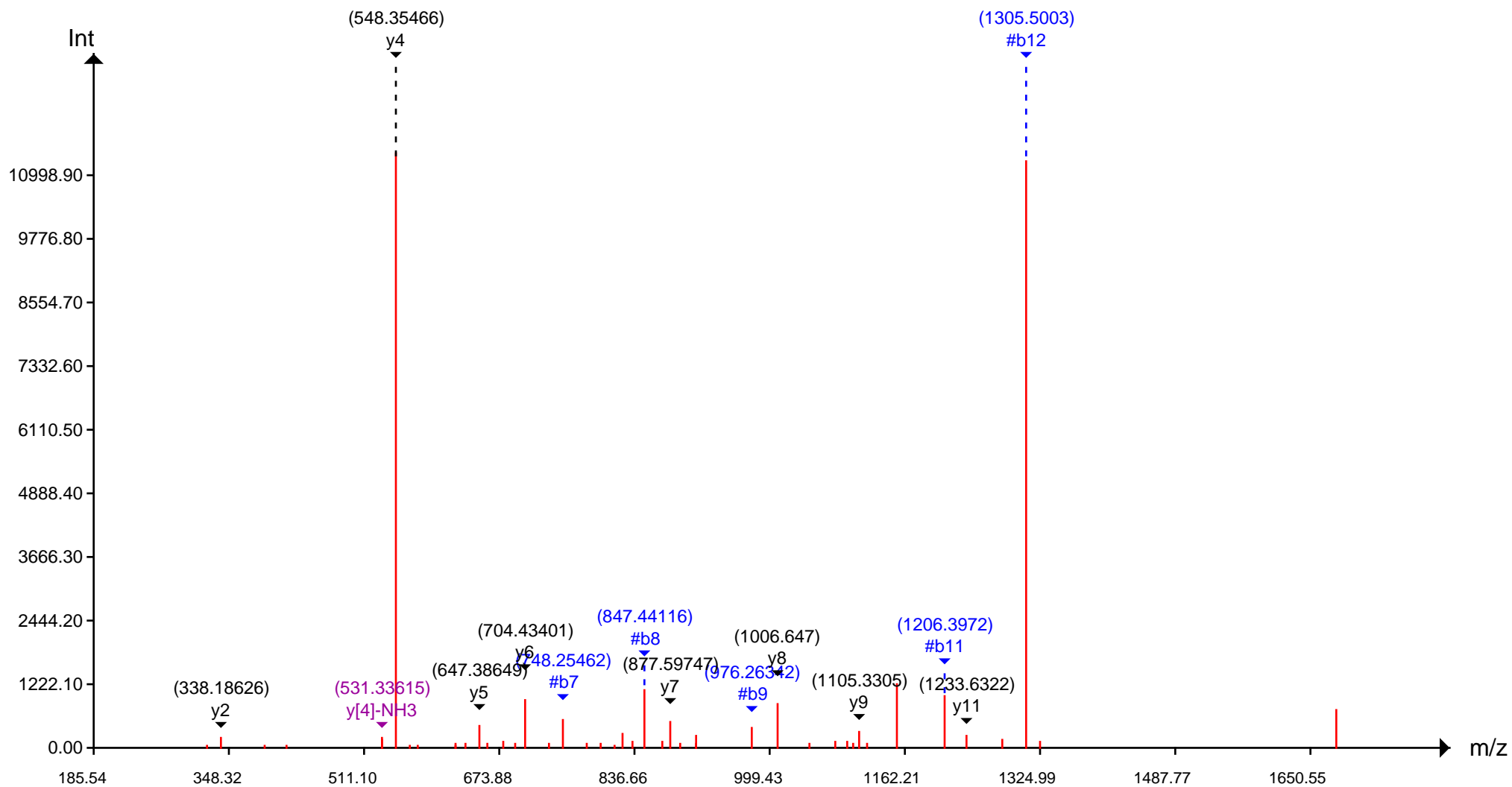
Charge 2+

Mass error 4.76E-4



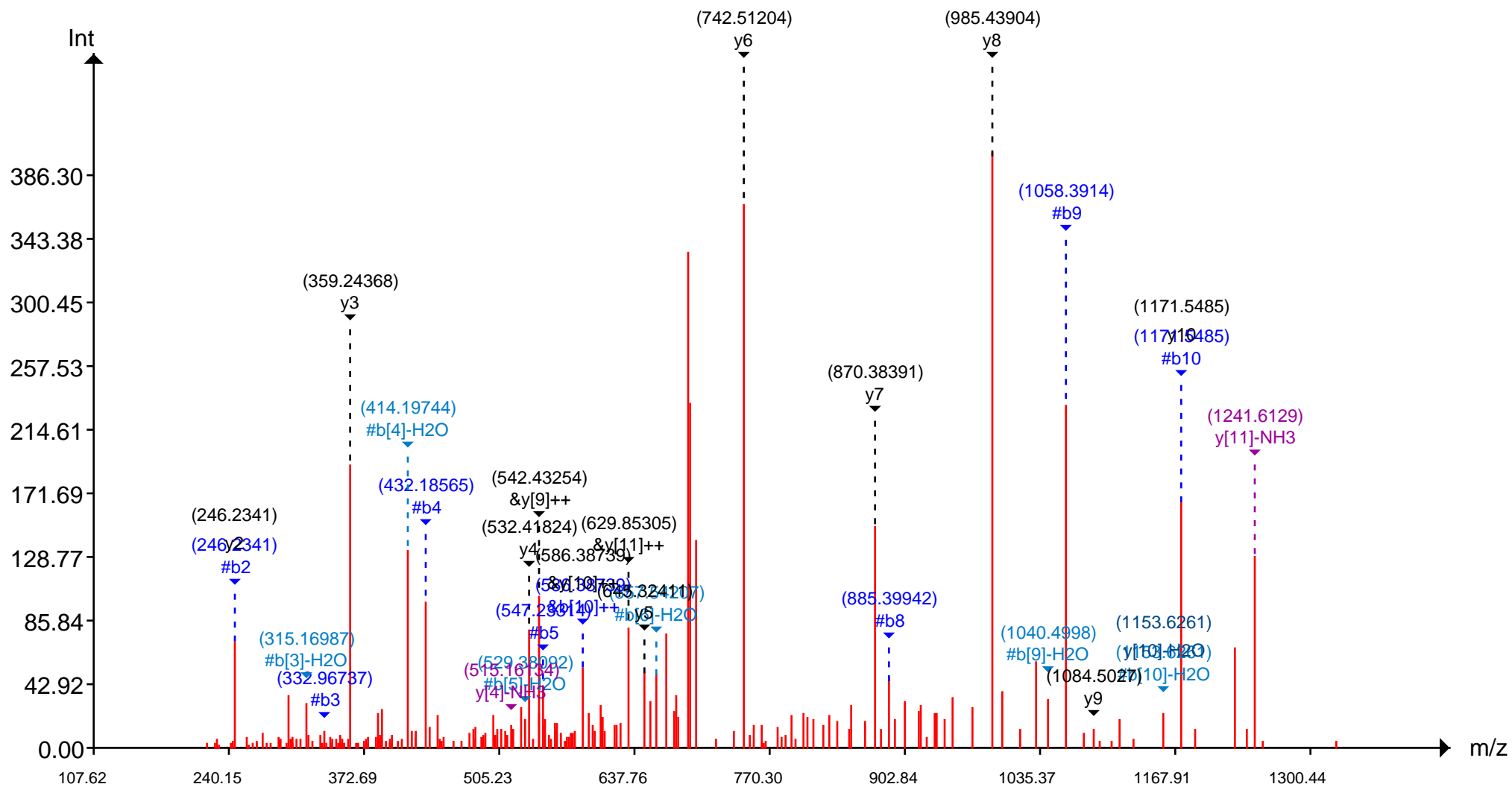
Info

Modified Sequence	AcD3-AVC<Cmm*>K<AcD3K*>AGAVEK<AcD3K*>GVPLYR-COOH	Ion score	51.84
Protein	P17182	Confidence level	99.00%
Precursor m/z	927.01989	Identity Threshold	46.54
Charge	2+	Mass error	0.002538



Info

Modified Sequence	AcD3-AVC<Cmm*>K<AcD3K*>AGAVEK<AcD3K*>GVPLYR-COOH	Ion score	59.58
Protein	P17182	Confidence level	99.00%
Precursor m/z	927.02014	Identity Threshold	46.57
Charge	2+	Mass error	0.003038



Info

Modified Sequence AcD3-LSSVDQPLK<AcD3K*>IAR-COOH

Ion score 45.96

Protein P47757

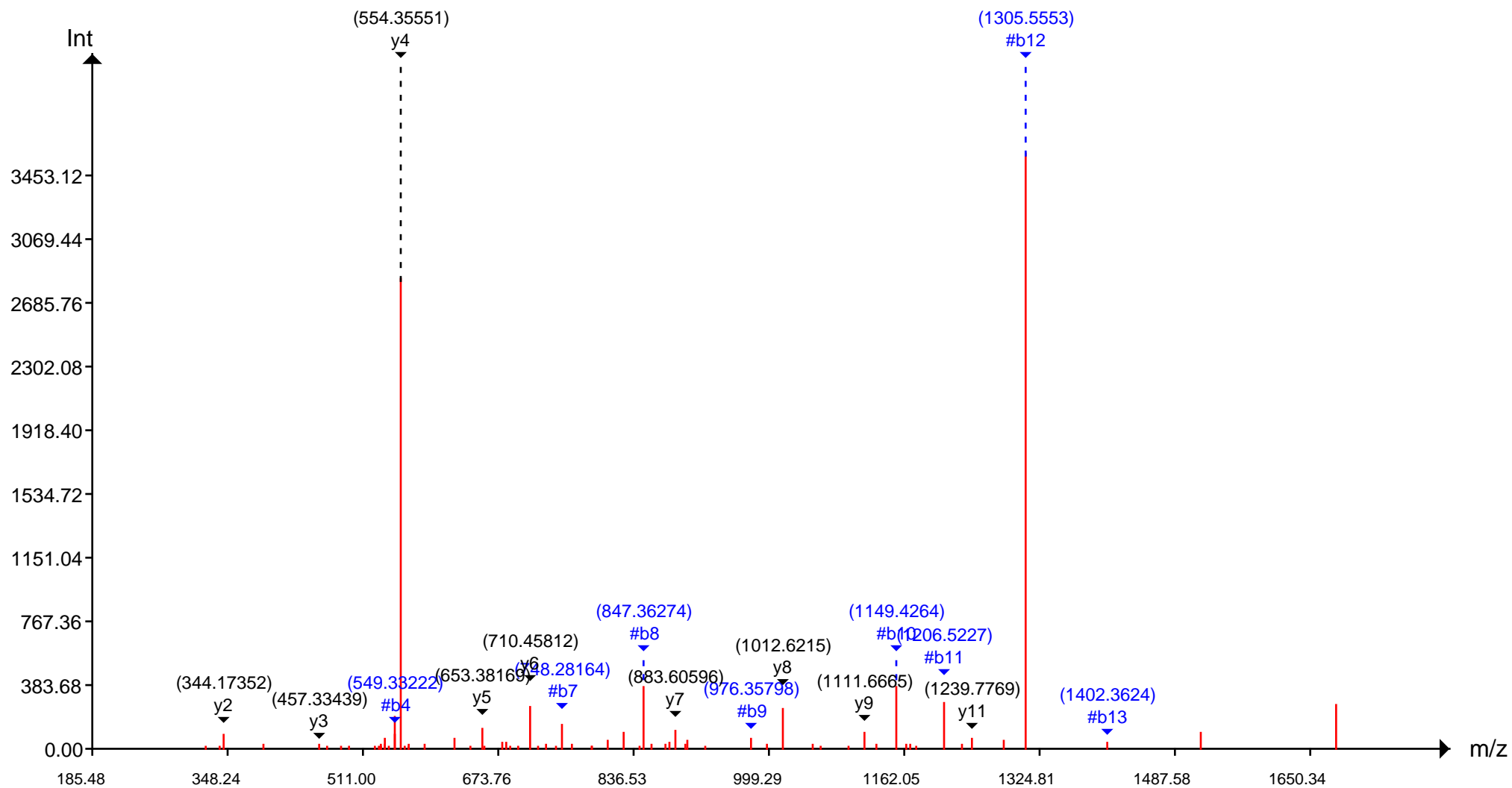
Confidence level 99.00%

Precursor m/z 708.92069

Identity Threshold 44.70

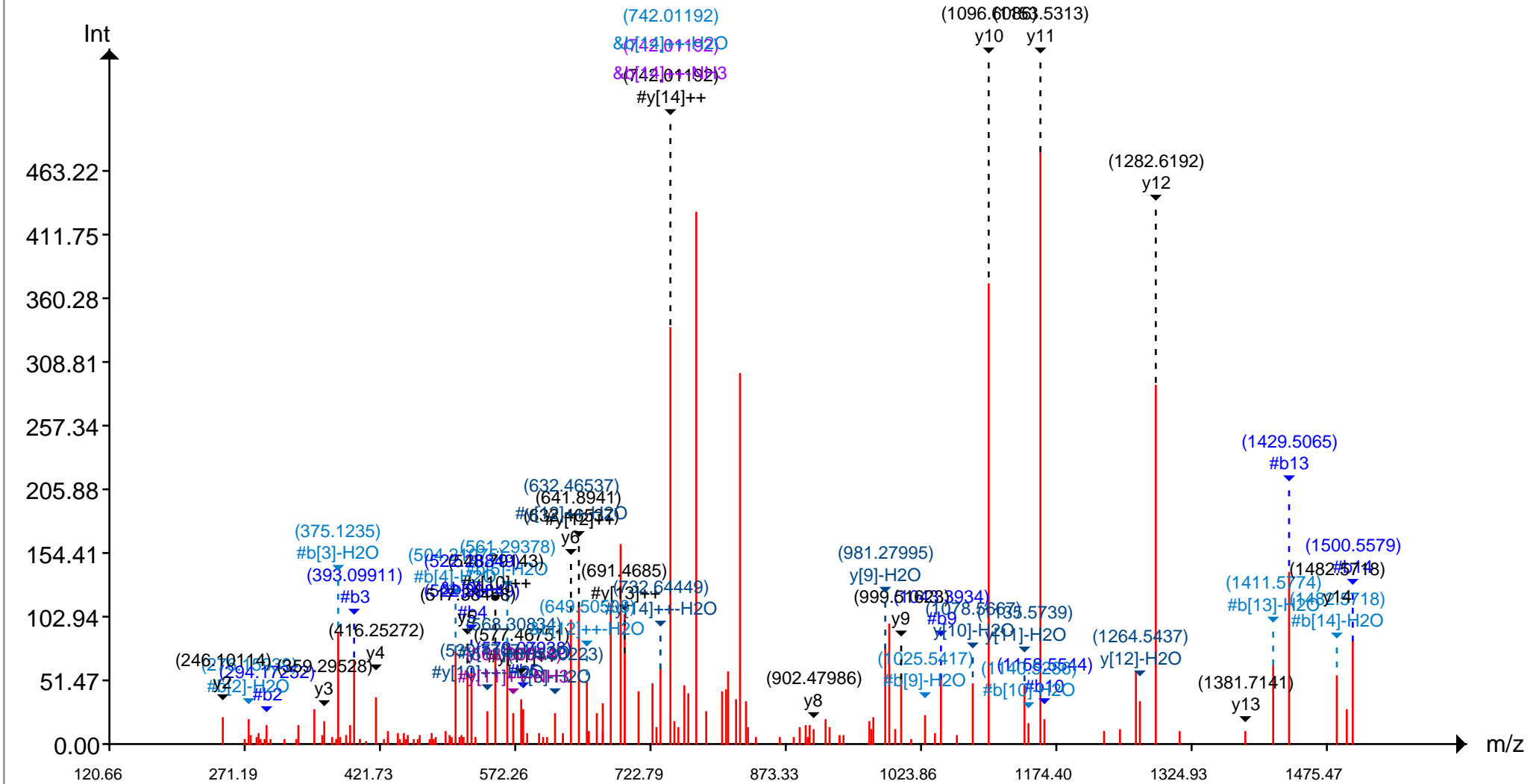
Charge 2+

Mass error 0.001358



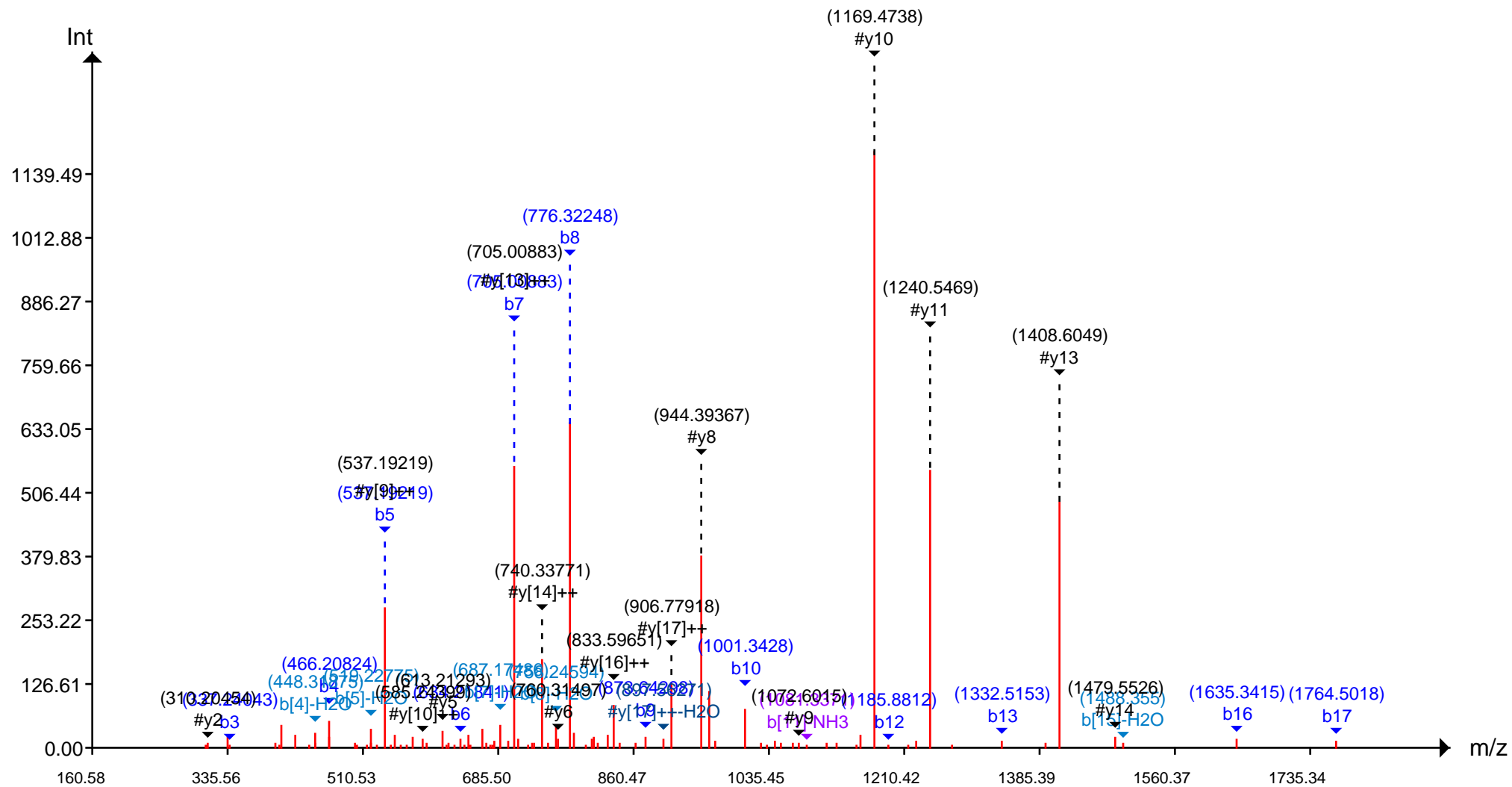
Info

Modified Sequence	AcD3-AVC<Cmm*>K<AcD3K*>AGAVEK<AcD3K*>GVPLYR<C13>-COOH	Ion score	71.34
Protein	P17182	Confidence level	99.00%
Precursor m/z	930.03009	Identity Threshold	45.69
Charge	2+	Mass error	0.002797



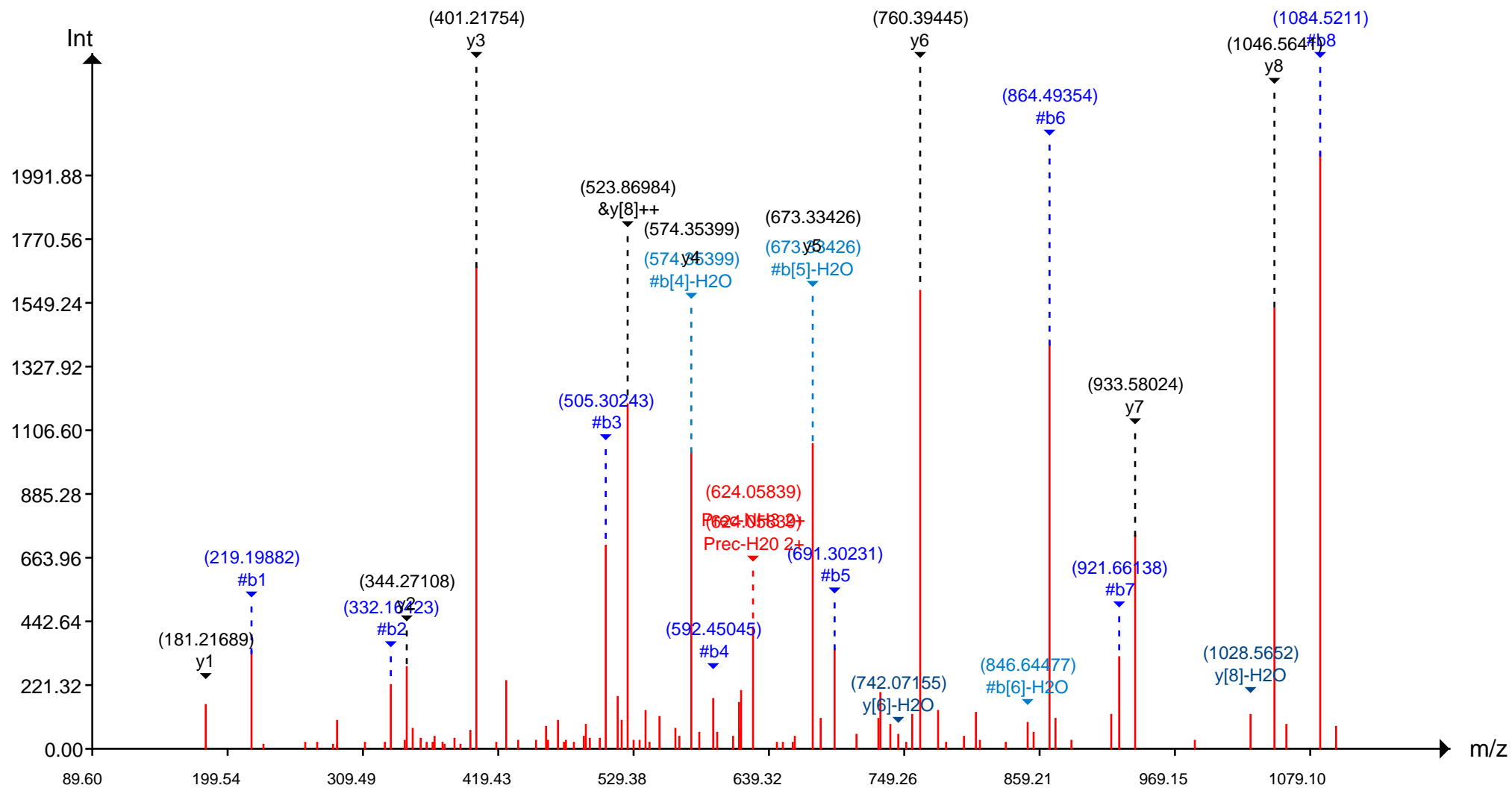
Info

Modified Sequence	AcD3-M<Mox*>TVEGPPP<AcD3K*>DTGIAR-COOH	Ion score	80.96
Protein	P27048, P63163	Confidence level	99.00%
Precursor m/z	837.93593	Identity Threshold	48.05
Charge	2+	Mass error	7.71E-4



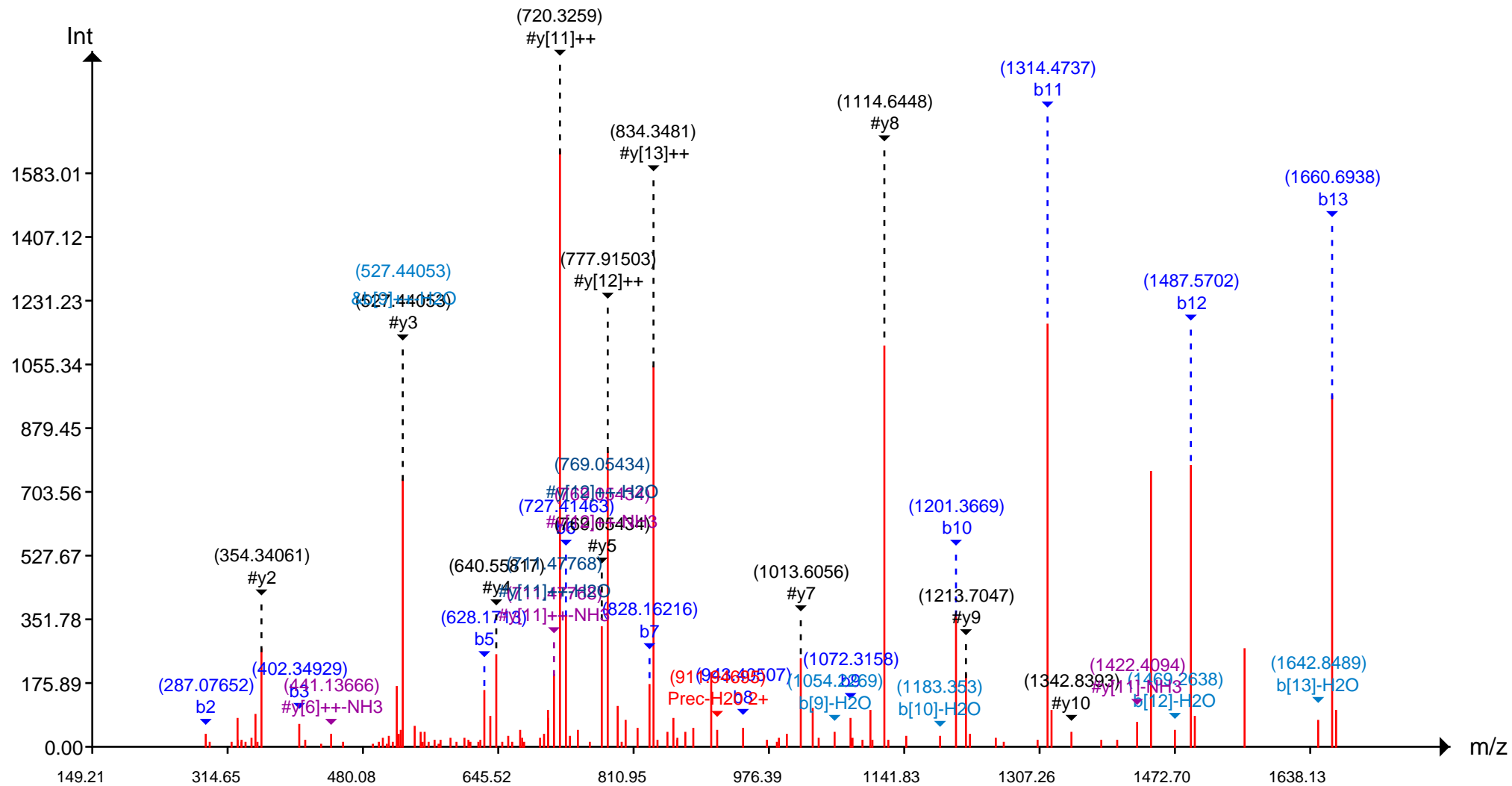
Info

Modified Sequence	AcD3-SFGEAPAAPQPSM<Mox*>SSEER<C13>-COOH	Ion score	74.2
Protein	Q9R062	Confidence level	99.00%
Precursor m/z	972.94308	Identity Threshold	44.07
Charge	2+	Mass error	0.001048



Info

Modified Sequence	AcD3-K<AcD3K*>LK<AcD3K*>SVK<AcD3K*>GYR<C13>-COOH	Ion score	69.44
Protein	P21981	Confidence level	99.00%
Precursor m/z	632.90928	Identity Threshold	38.53
Charge	2+	Mass error	4.34E-4



Info

Modified Sequence AcD3-QIDPEVTDEEIK<AcD3K*>K<AcD3K*>R<C13>-COOH

Ion score 67.52

Protein Q6NZB0

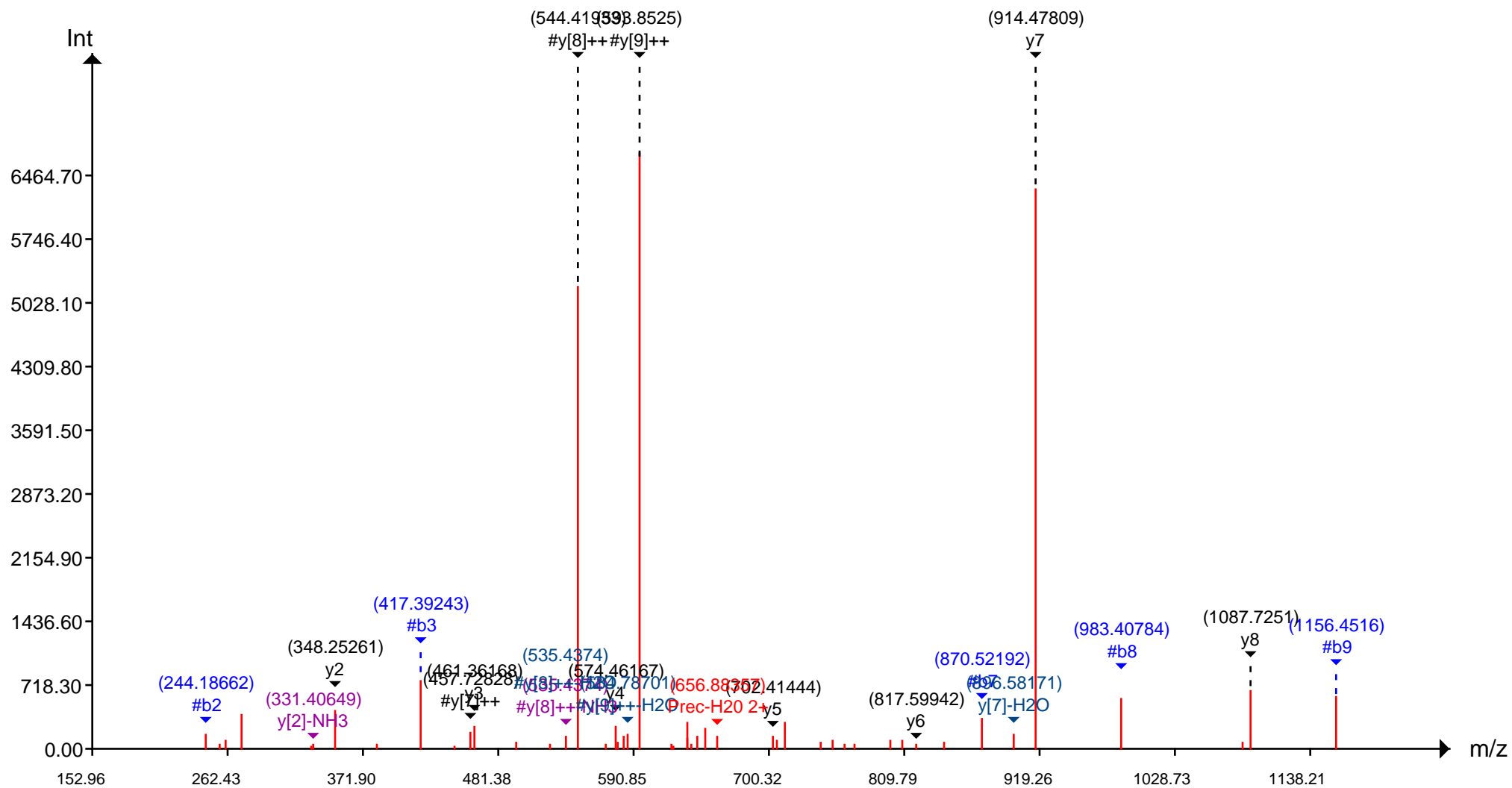
Confidence level 99.00%

Precursor m/z 921.00137

Identity Threshold 47.57

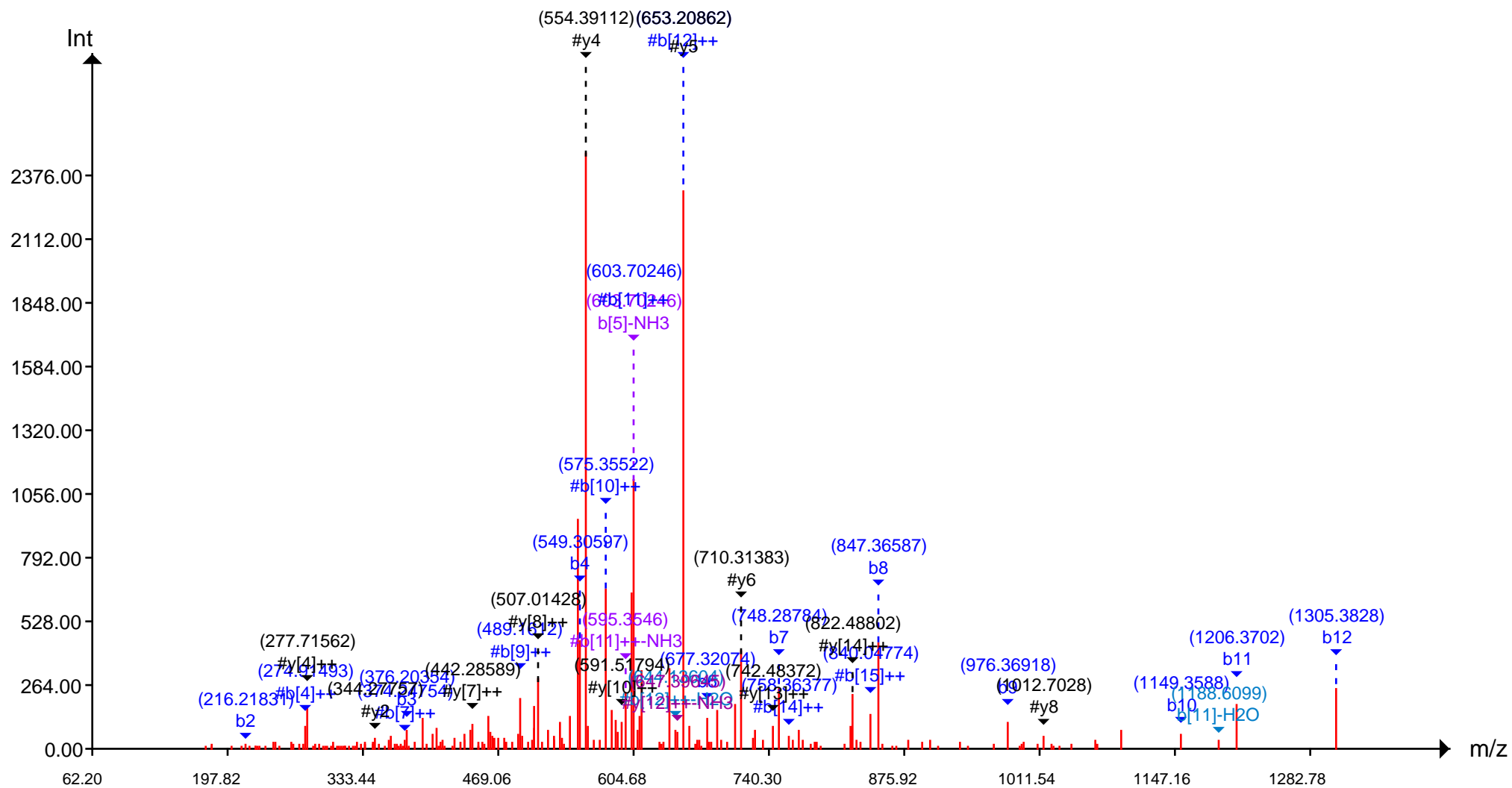
Charge 2+

Mass error 0.001051



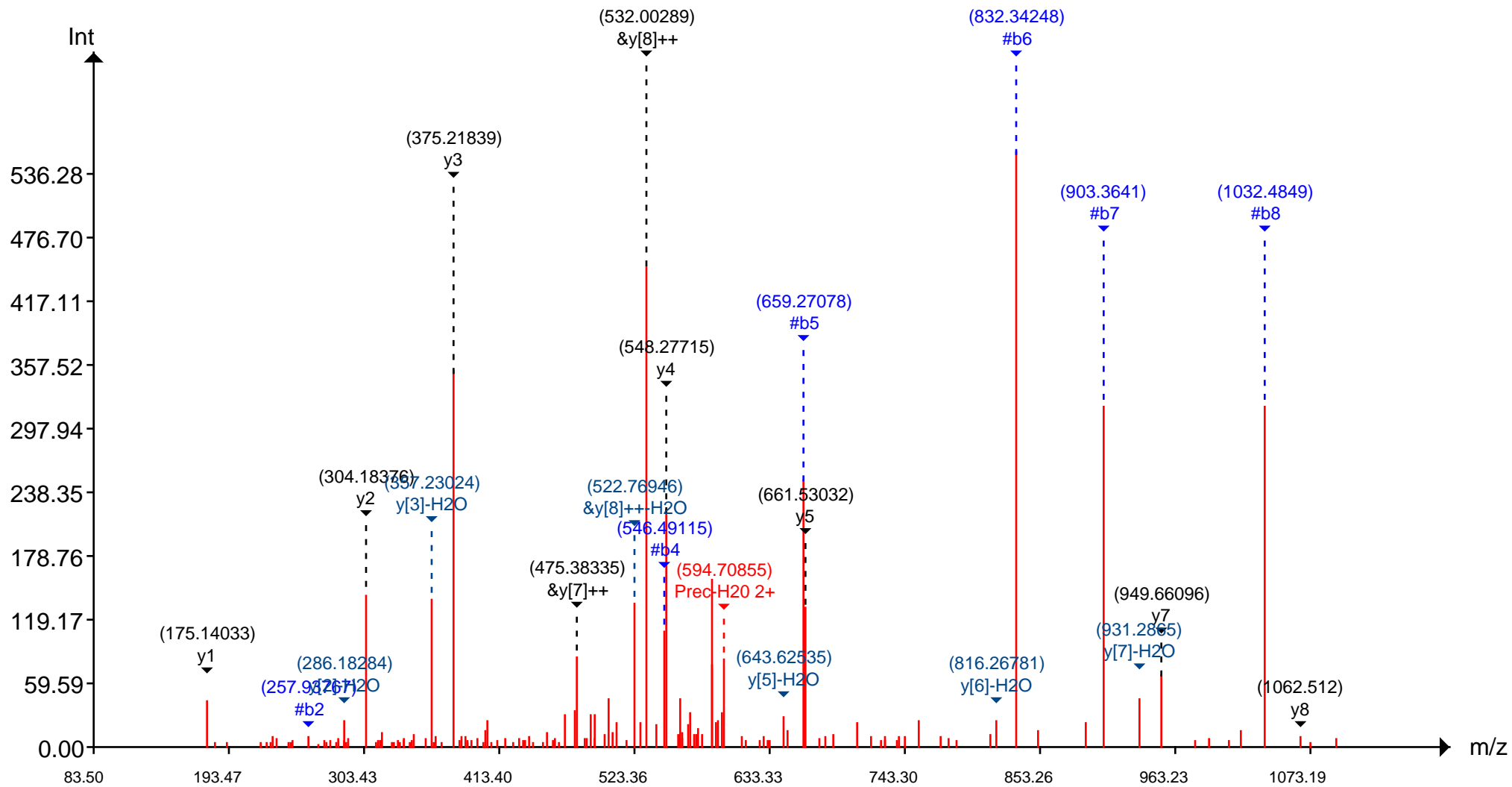
Info

Modified Sequence	AcD3-VVK<AcD3K*>PDQLIK<AcD3K*>R-COOH	Ion score	42.26
Protein	Q91V92	Confidence level	99.00%
Precursor m/z	665.92467	Identity Threshold	39.25
Charge	2+	Mass error	0.001765



Info

Modified Sequence	AcD3-AVC<Cmm*>K<AcD3K*>AGAVEK<AcD3K*>GVPLYR<C13>-COOH	Ion score	55.53
Protein	P17182	Confidence level	99.00%
Precursor m/z	620.3548	Identity Threshold	45.86
Charge	3+	Mass error	-2.59E-4



Info

Modified Sequence AcD3-VLDK<AcD3K*>LK<AcD3K*>AER-COOH

Ion score 50.26

Protein P10126, P62631

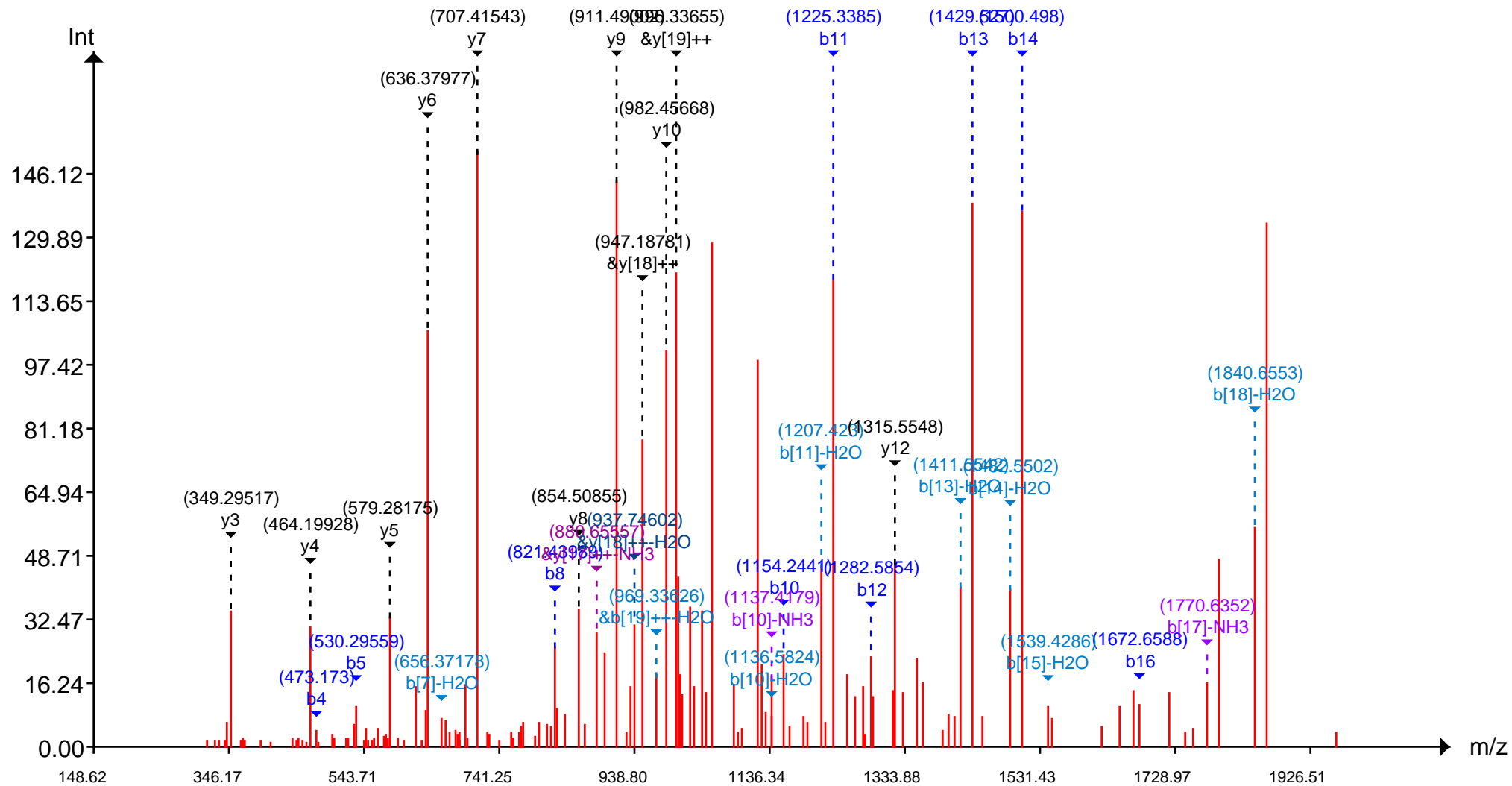
Confidence level 99.00%

Precursor m/z 603.87411

Identity Threshold 42.56

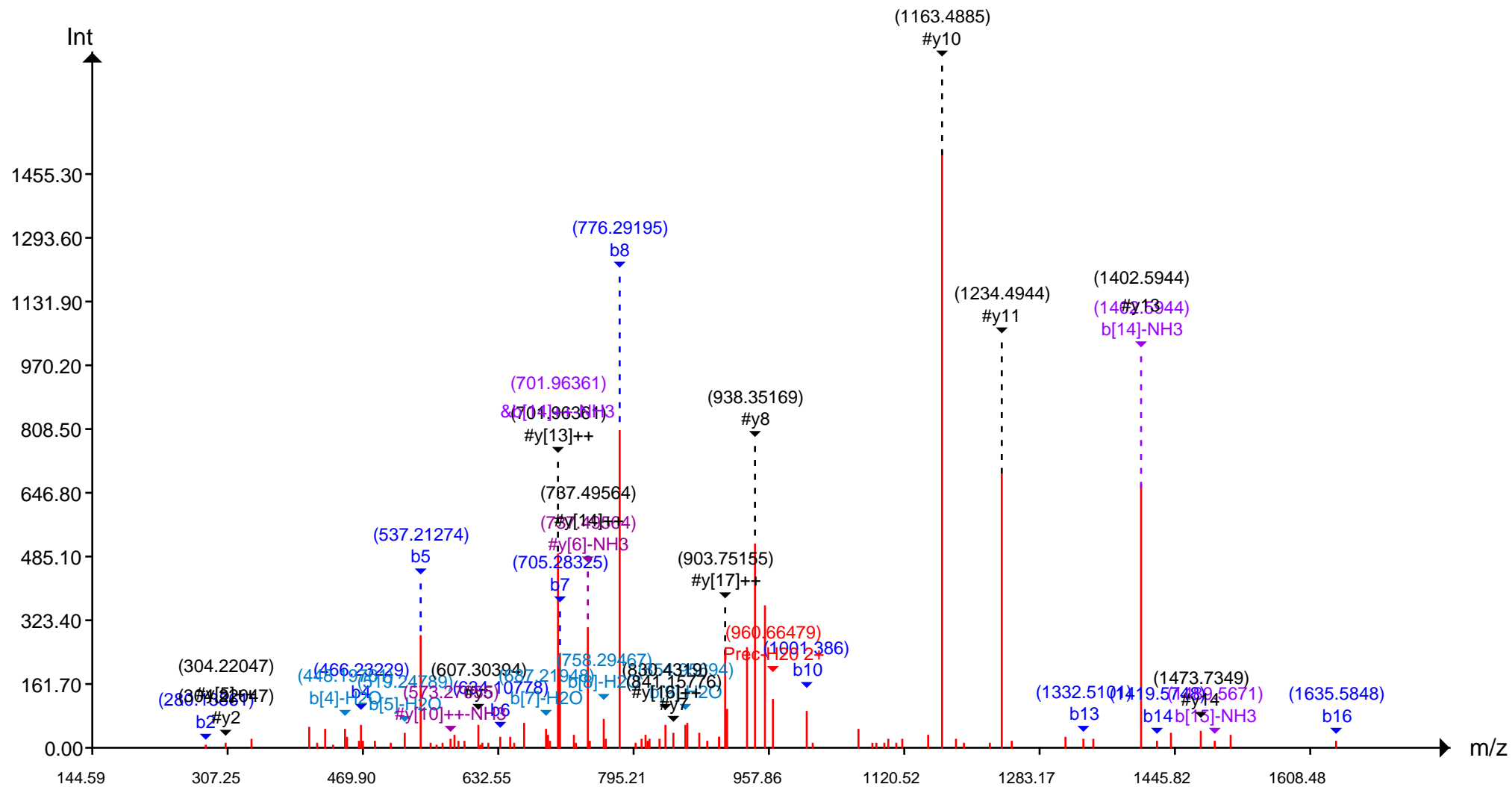
Charge 2+

Mass error 7.12E-4



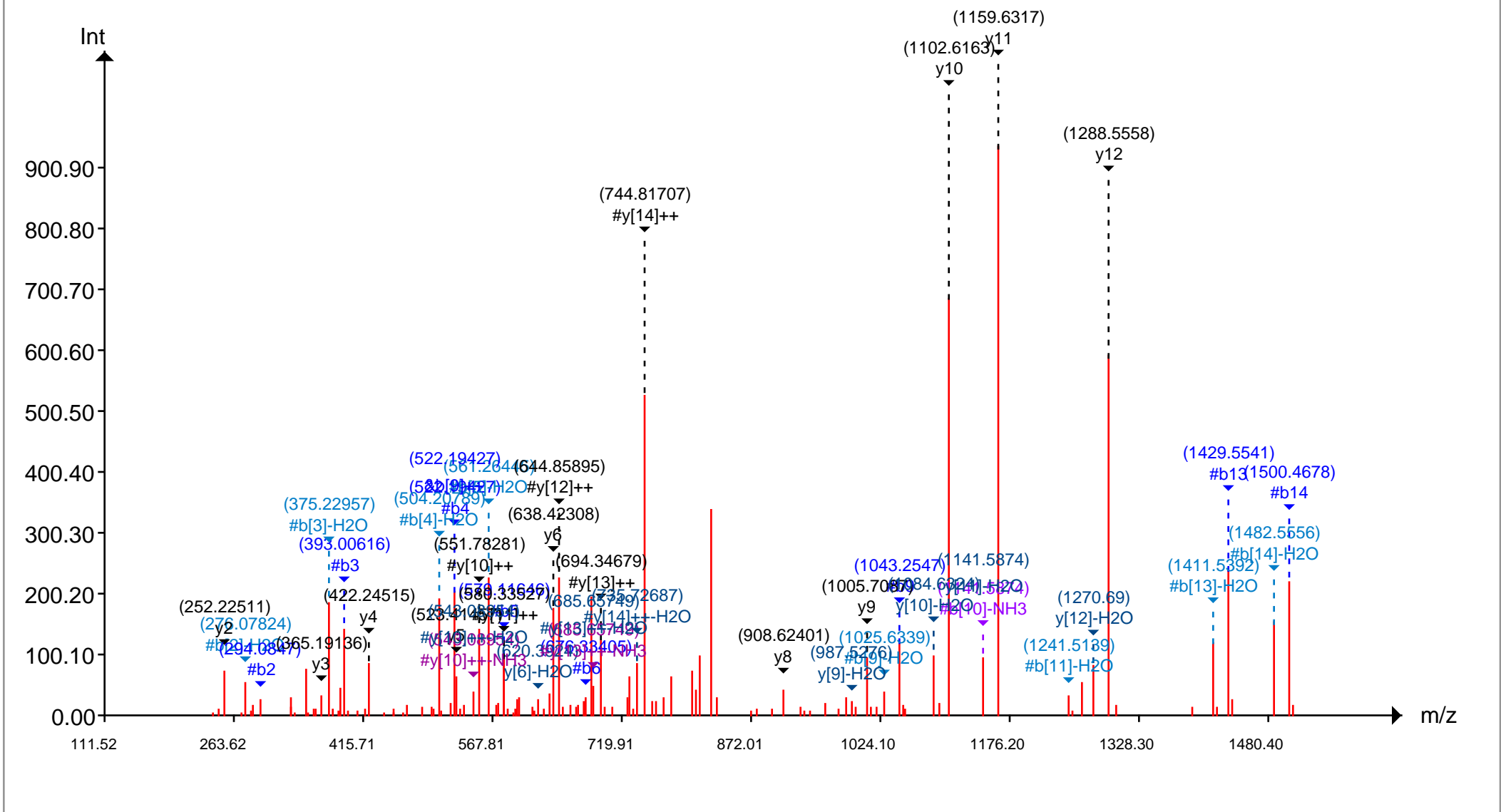
Info

Modified Sequence	AcD3-VVDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH	Ion score	60.27
Protein	P60710	Confidence level	99.00%
Precursor m/z	1068.4892	Identity Threshold	44.27
Charge	2+	Mass error	0.001217



Info

Modified Sequence	AcD3-SFGEAPAAPQPSM<Mox*>SSEER-COOH	Ion score	60.27
Protein	Q9R062	Confidence level	99.00%
Precursor m/z	969.93326	Identity Threshold	43.12
Charge	2+	Mass error	0.00155



Info

Modified Sequence AcD3-M<Mox*>TVEGPPPK<AcD3K*>DTGIAR<C13>-COOH

Ion score 72.56

Protein P27048, P63163

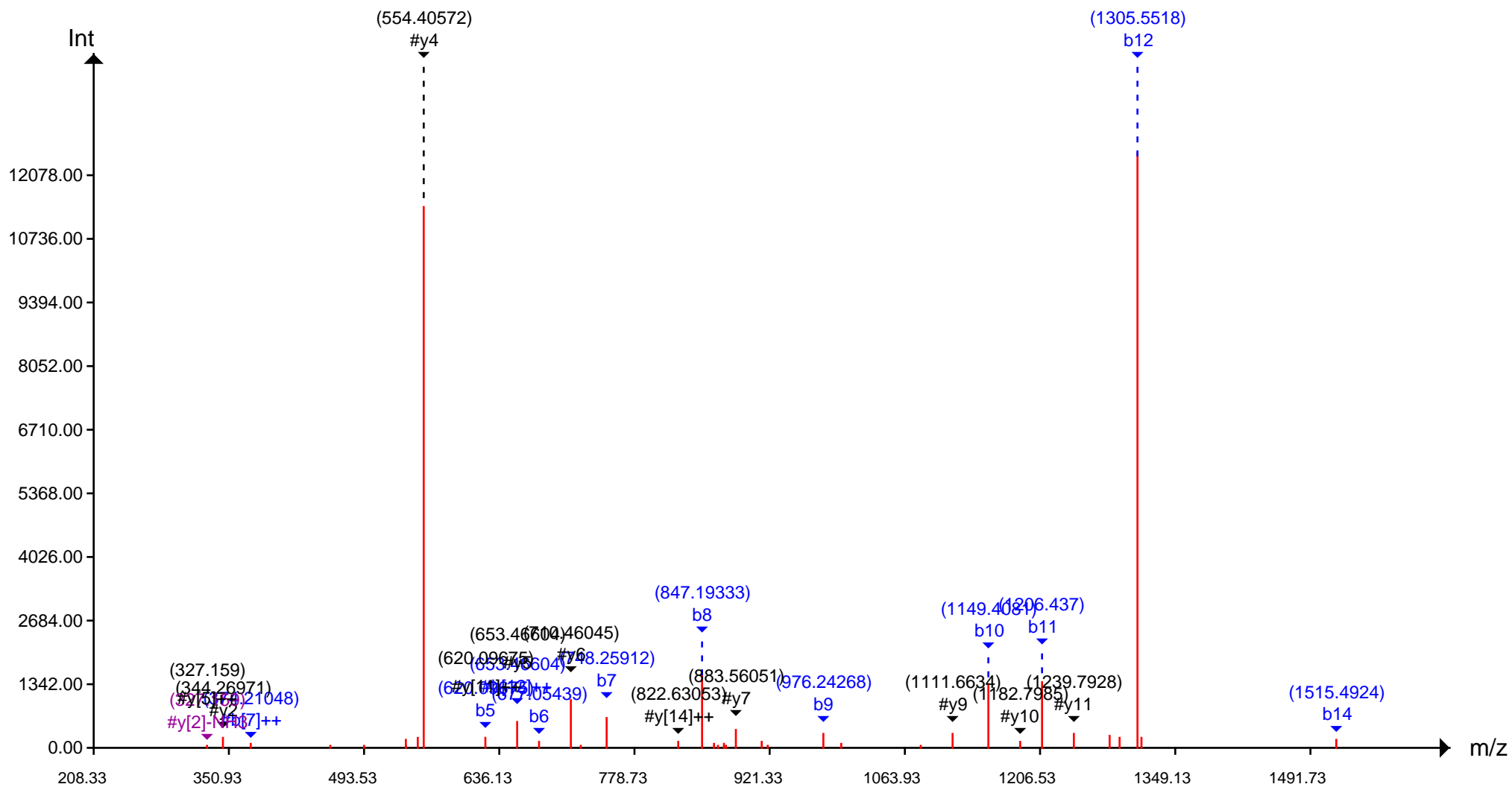
Confidence level 99.00%

Precursor m/z 840.94601

Identity Threshold 48.06

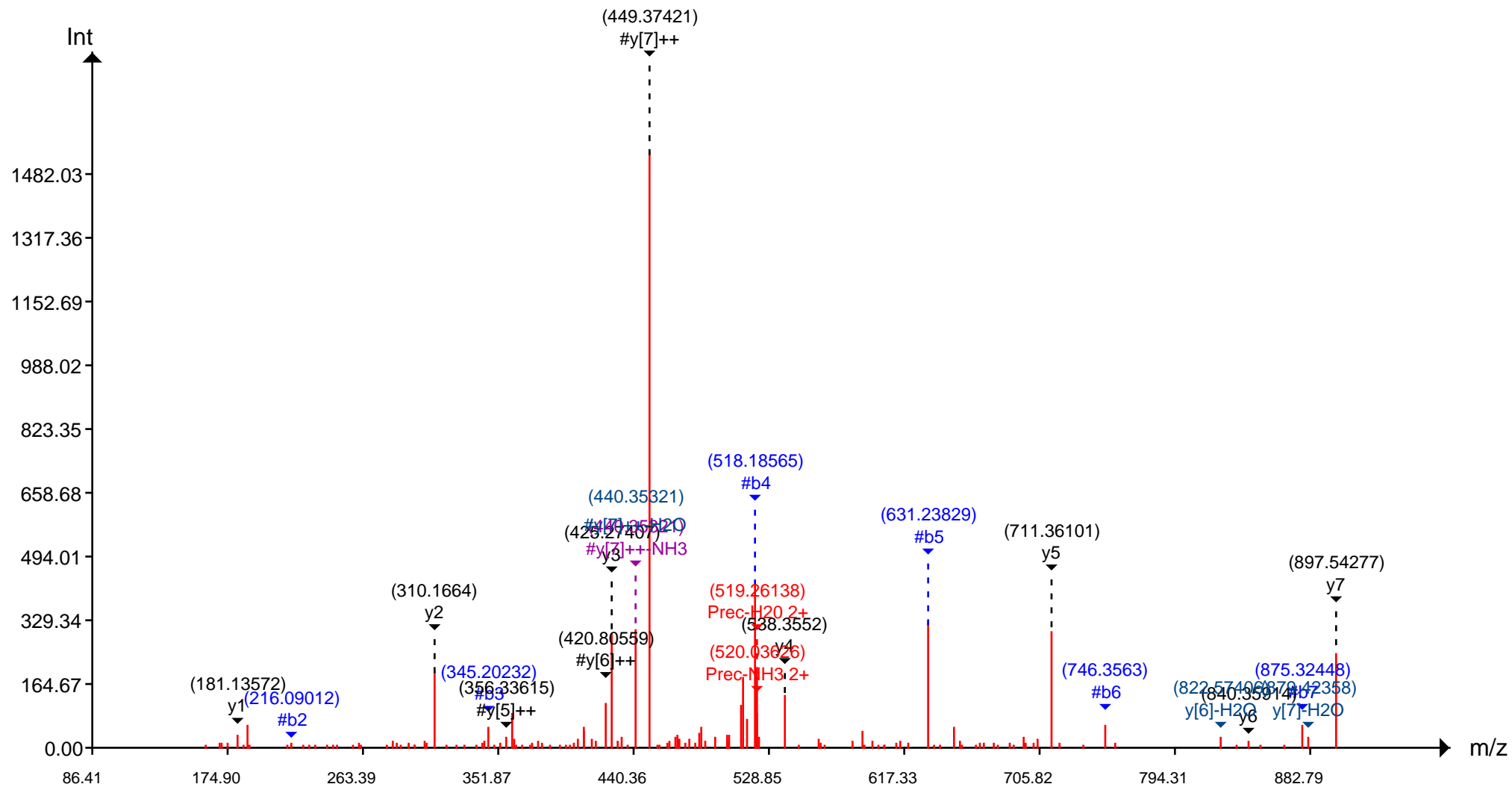
Charge 2+

Mass error 7.9E-4



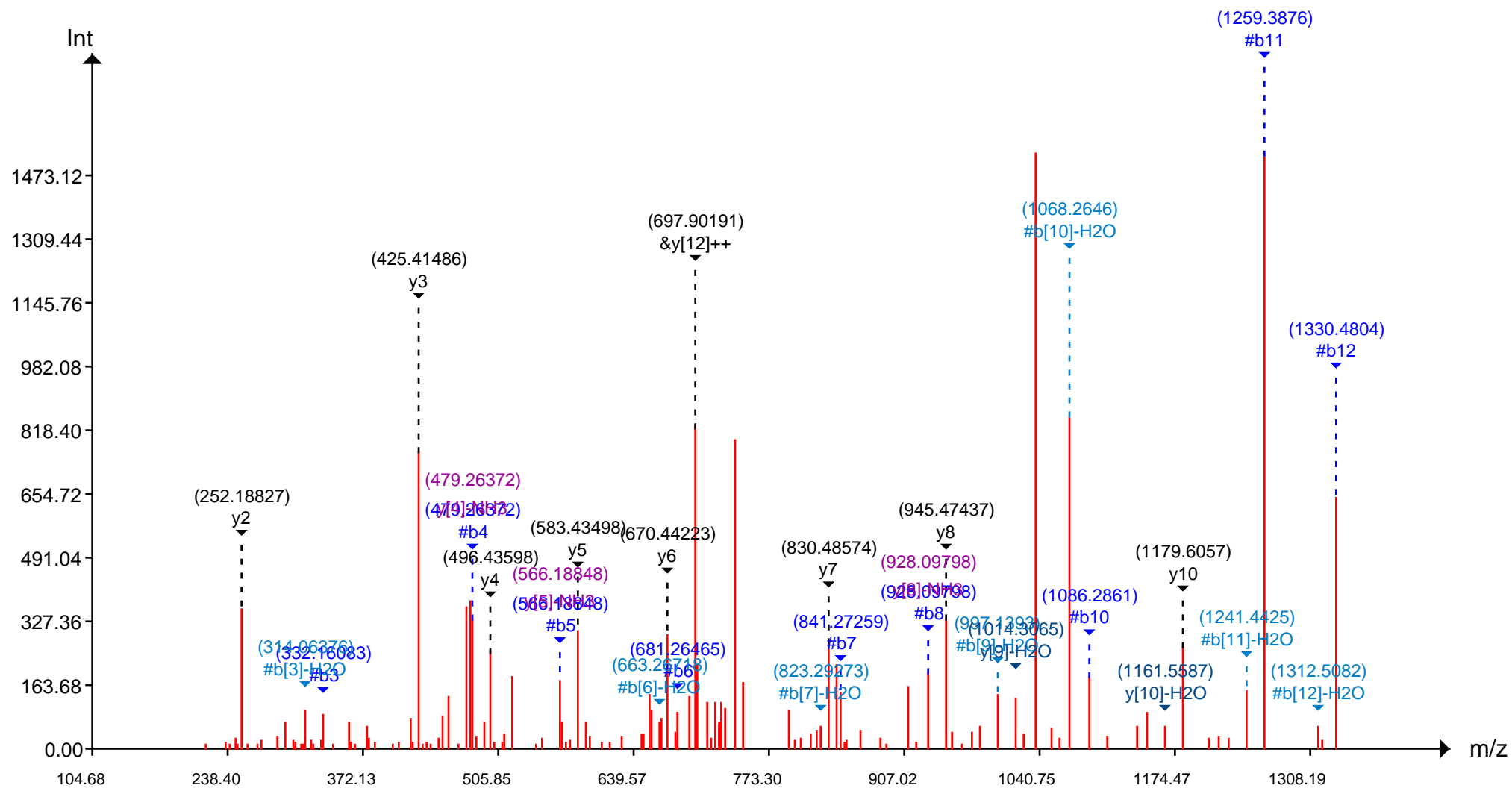
Info

Modified Sequence	AcD3-AVC<Cmm*>K<AcD3K*>AGAVEK<AcD3K*>GVPLYR<C13>-COOH	Ion score	70.17
Protein	P17182	Confidence level	99.00%
Precursor m/z	930.02947	Identity Threshold	45.80
Charge	2+	Mass error	0.001557



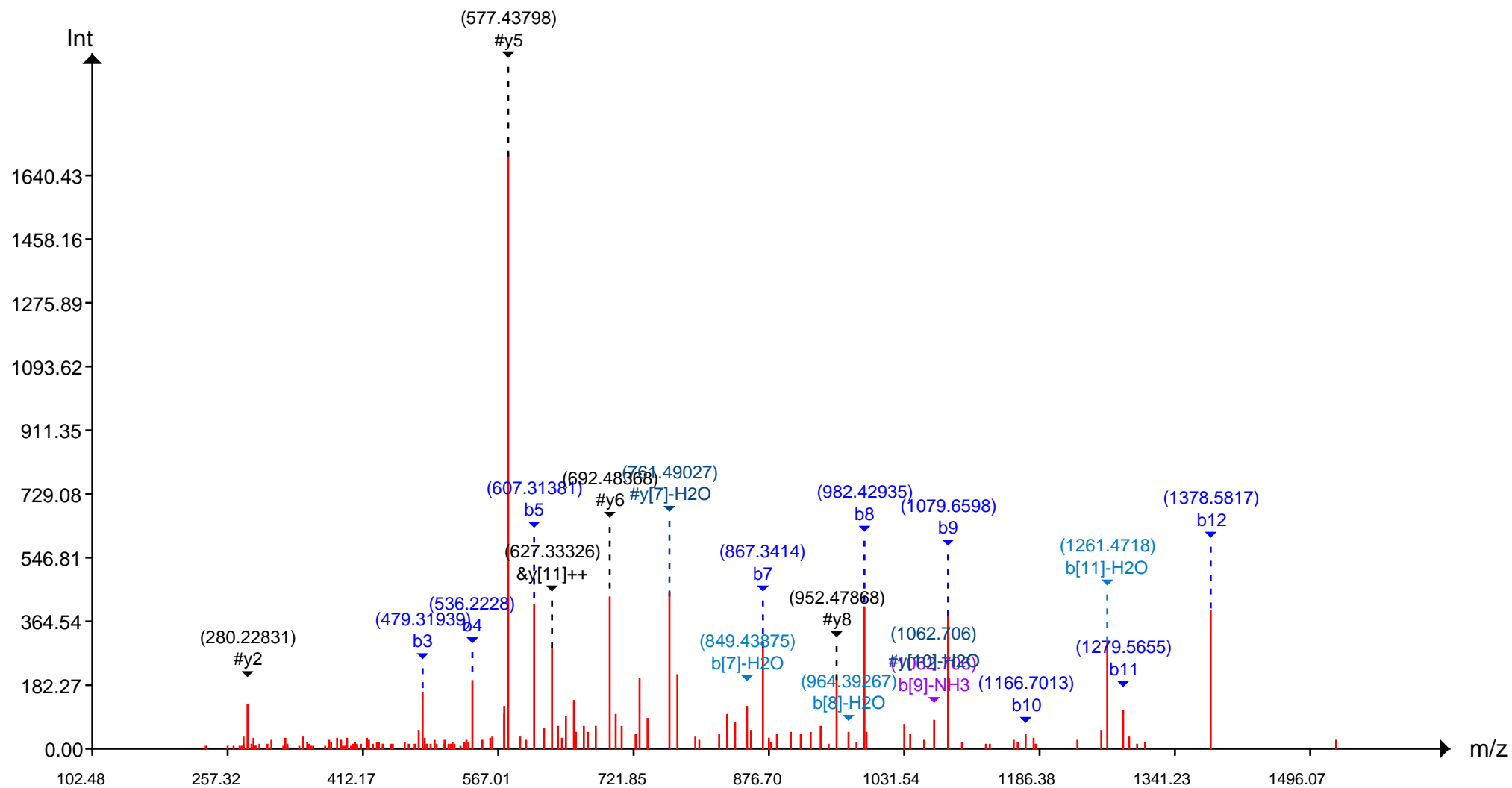
Info

Modified Sequence	AcD3-IGEK<AcD3K*>LDER<C13>-COOH	Ion score	52.09
Protein	P17751	Confidence level	99.00%
Precursor m/z	528.30097	Identity Threshold	47.18
Charge	2+	Mass error	1.23E-4



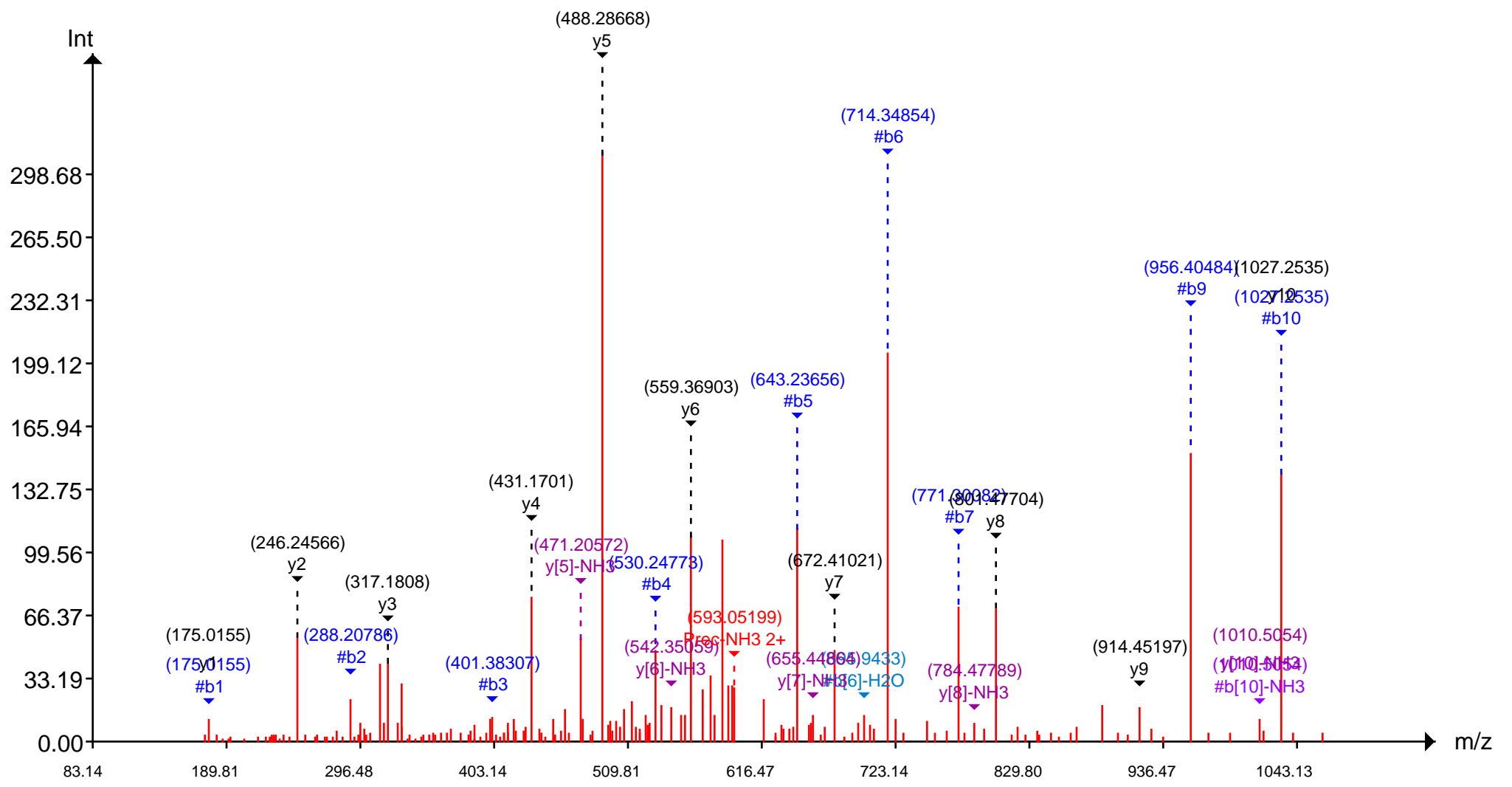
Info

Modified Sequence	AcD3-ASQFSDC<Cmm*>SSAK<AcD3K*>AR<C13>-COOH	Ion score	47.93
Protein	Q9QUR7	Confidence level	99.00%
Precursor m/z	755.8623	Identity Threshold	46.49
Charge	2+	Mass error	4.53E-4

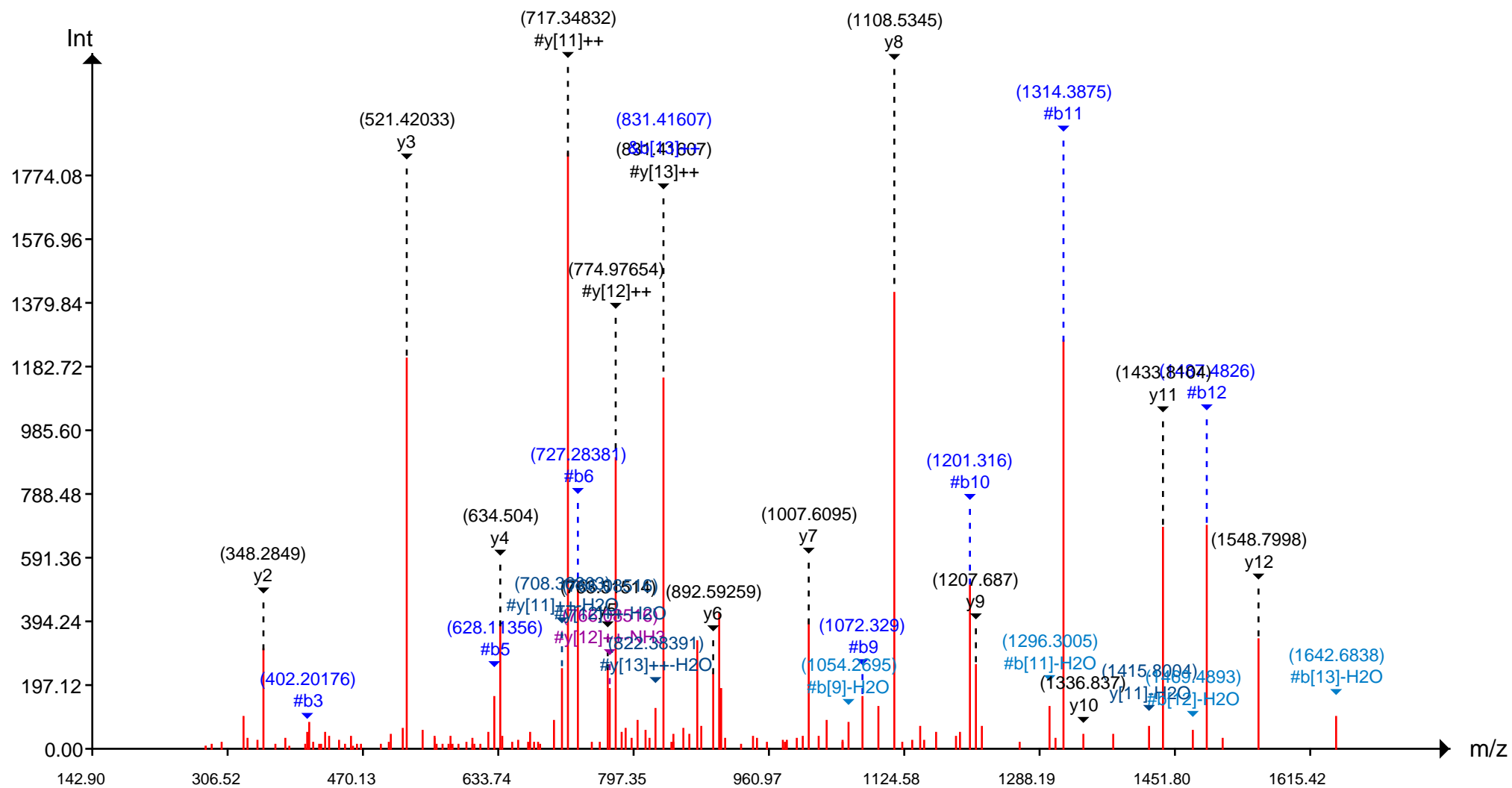


Info

Modified Sequence	AcD3-FIK<AcD3K*>GAK<AcD3K*>SDPSIVR<C13>-COOH	Ion score	56.8
Protein	P62748	Confidence level	99.00%
Precursor m/z	779.96691	Identity Threshold	43.61
Charge	2+	Mass error	0.002047



Info	
Modi...	AcD3-EILELAGNAAR-COOH
	Q8CGP5, Q8CGP6, Q8CGP7, P22752, Q6GSS7, Q64522, Q64523, Q8BFU2, Q8R1M2, Q8CCK0, P27661, Q9QZQ8
	601.3352
	2+
	Ion score
	Confidence level
	Identity Threshold
	Mass error



Info

Modified Sequence AcD3-QIDPEVTDEEIK<AcD3K*>K<AcD3K*>R-COOH

Ion score 77.21

Protein Q6NZB0

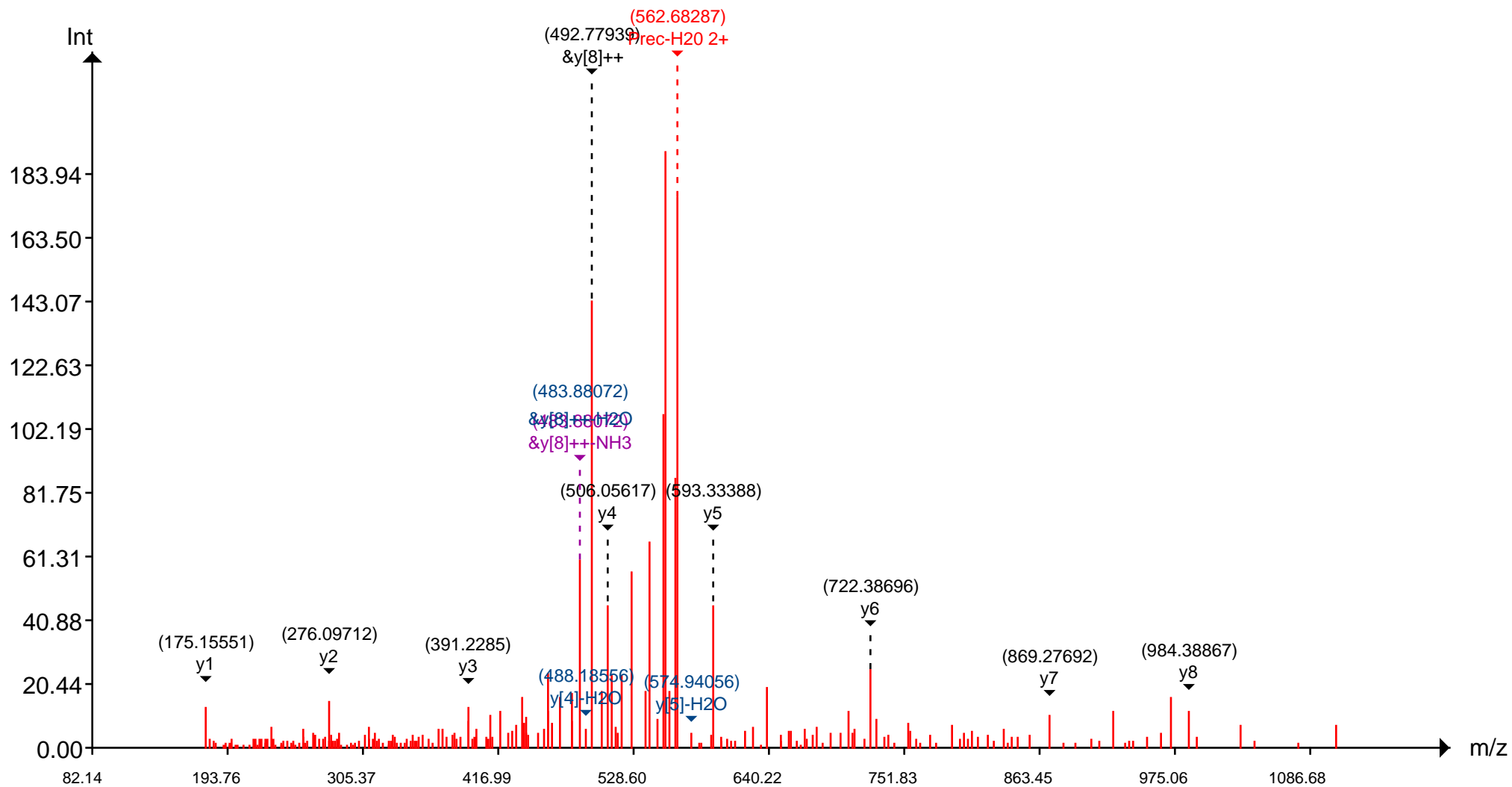
Confidence level 99.00%

Precursor m/z 917.9916

Identity Threshold 47.98

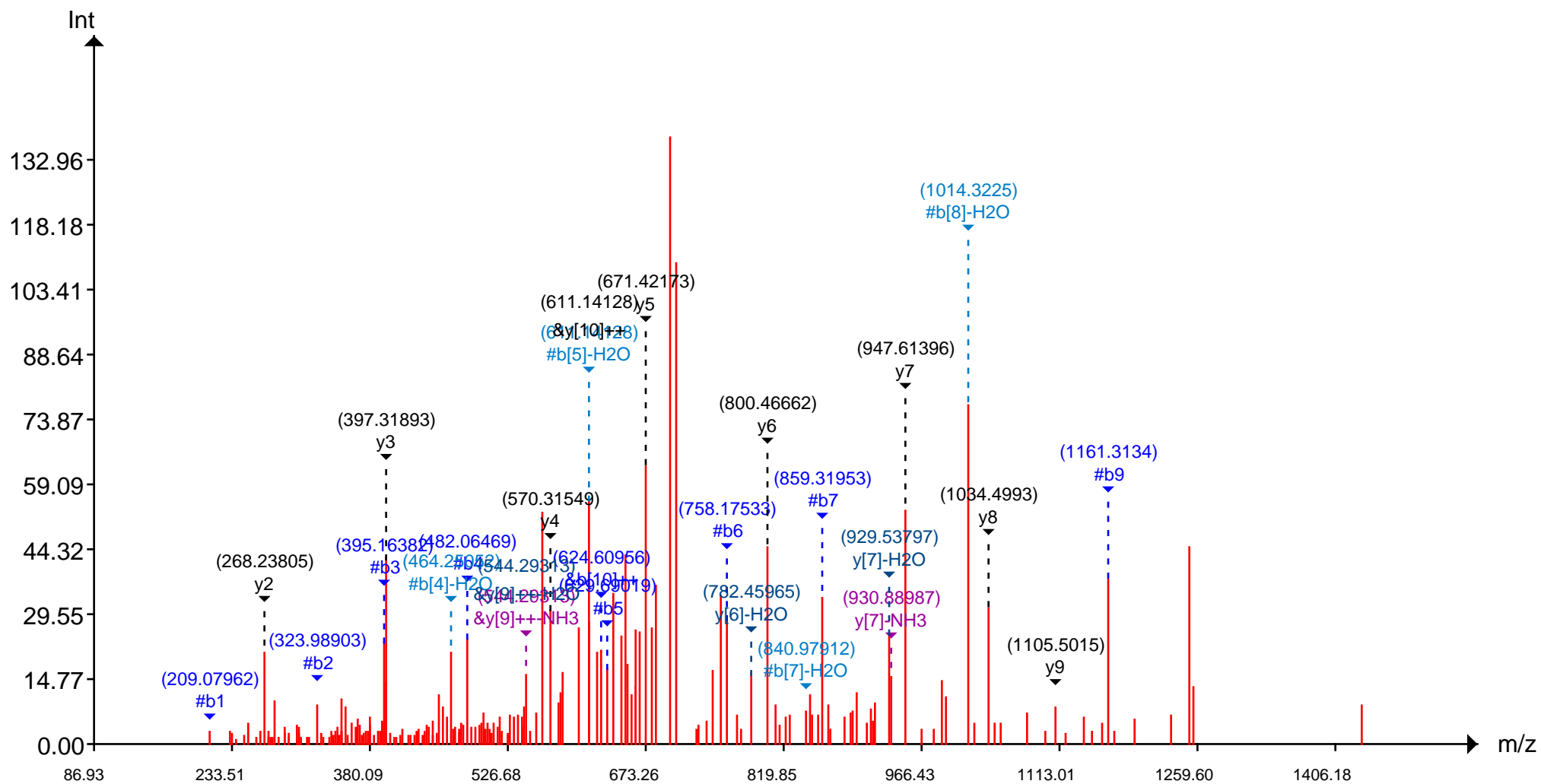
Charge 2+

Mass error 0.001653



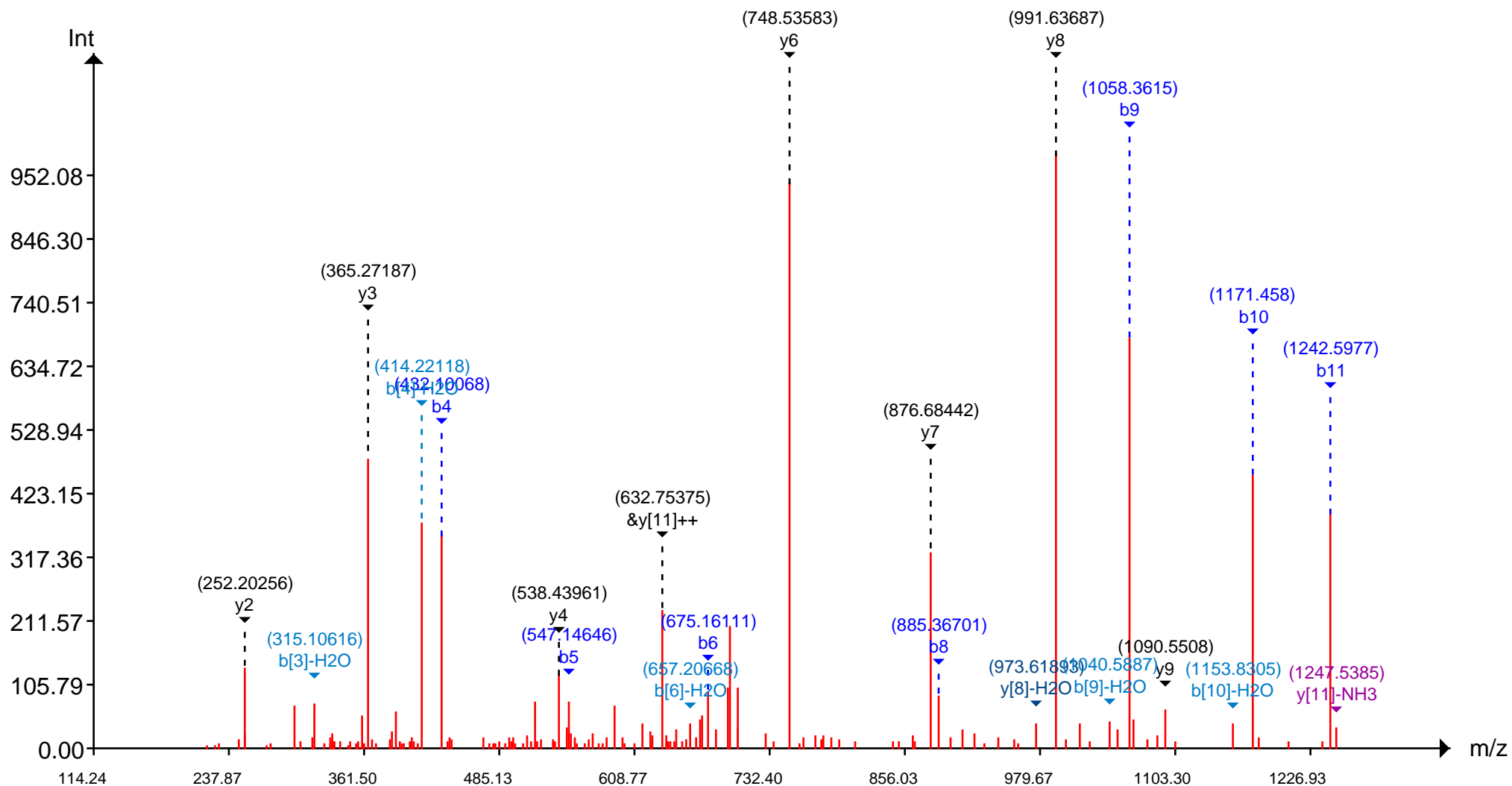
Info

Modified Sequence	AcD3-LDM<Mox*>ESDDTR-COOH	Ion score	40.85
Protein	Q62084	Confidence level	99.00%
Precursor m/z	571.73927	Identity Threshold	40.28
Charge	2+	Mass error	3.34E-4



Info

Modified Sequence	AcD3-YDASFETK<AcD3K*>ESR<C13>-COOH	Ion score	49.7
Protein	Q9R0P5	Confidence level	99.00%
Precursor m/z	714.84694	Identity Threshold	46.77
Charge	2+	Mass error	0.00106



Info

Modified Sequence AcD3-LSSVDQPLK<AcD3K*>IAR<C13>-COOH

Ion score 64.92

Protein P47757

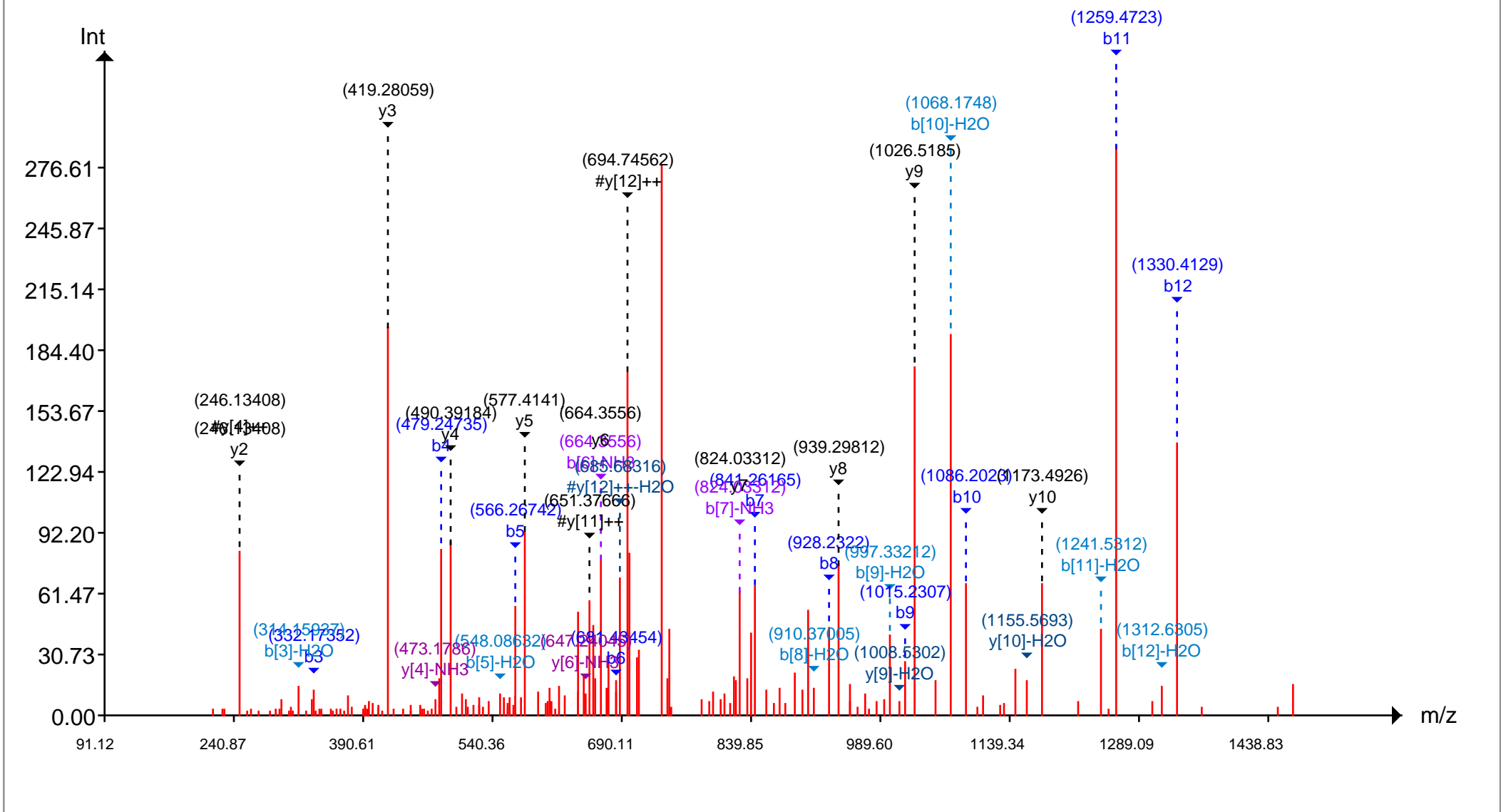
Confidence level 99.00%

Precursor m/z 711.93065

Identity Threshold 43.34

Charge 2+

Mass error 0.001136



Info

Modified Sequence AcD3-ASQFSDC<Cmm*>SSAK<AcD3K*>AR-COOH

Ion score 66.19

Protein Q9QUR7

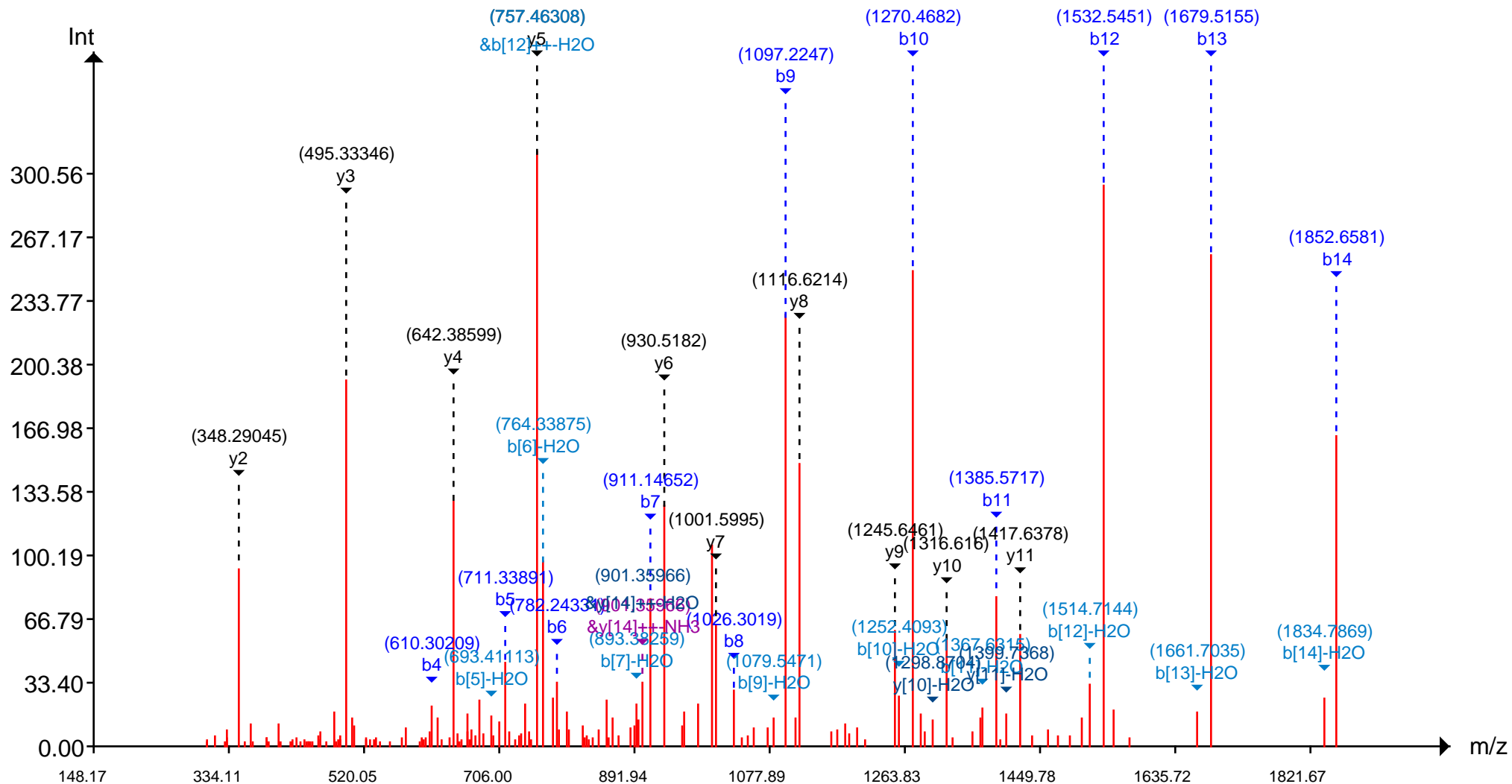
Confidence level 99.00%

Precursor m/z 752.85248

Identity Threshold 45.67

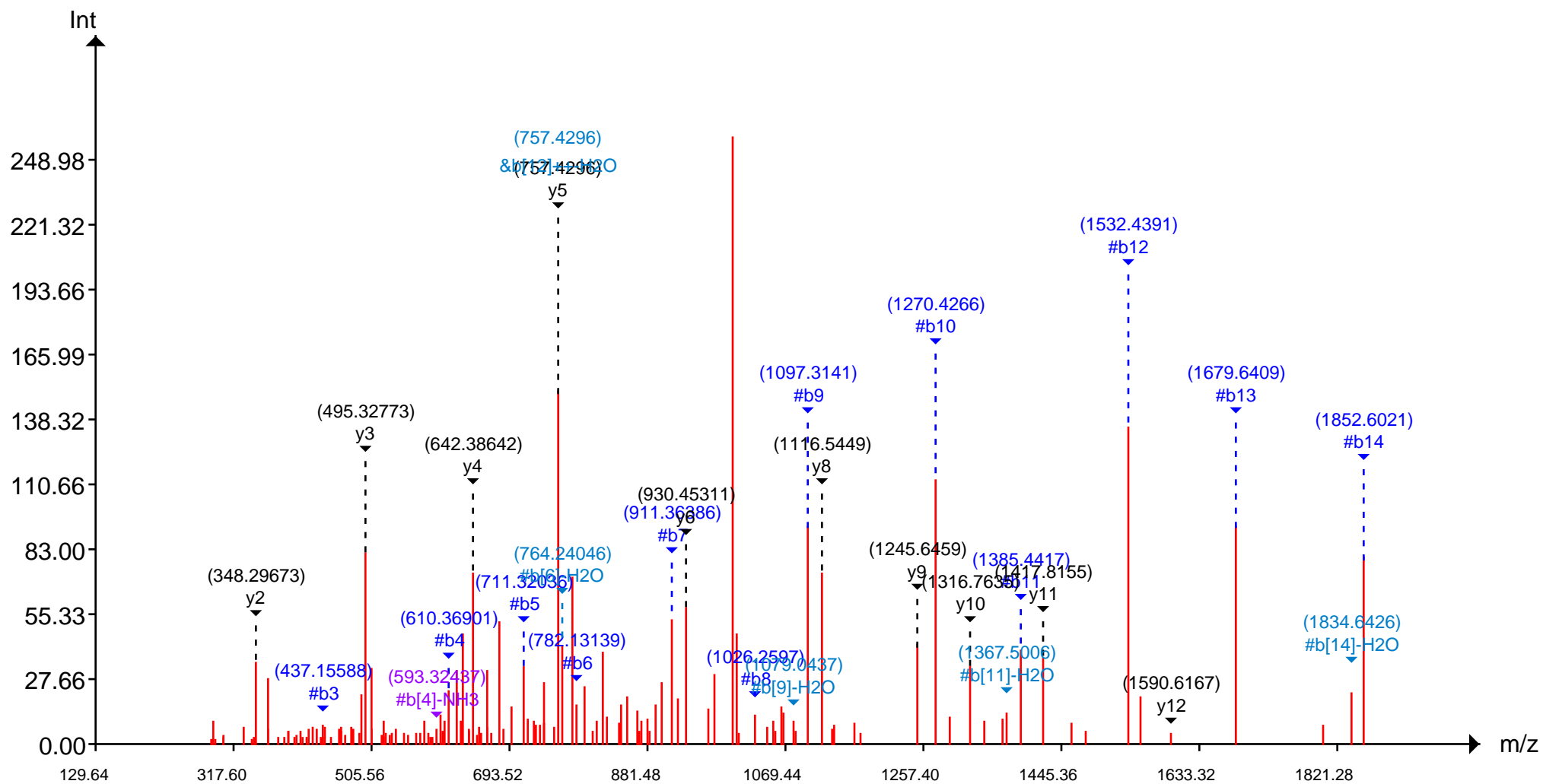
Charge 2+

Mass error 9.55E-4



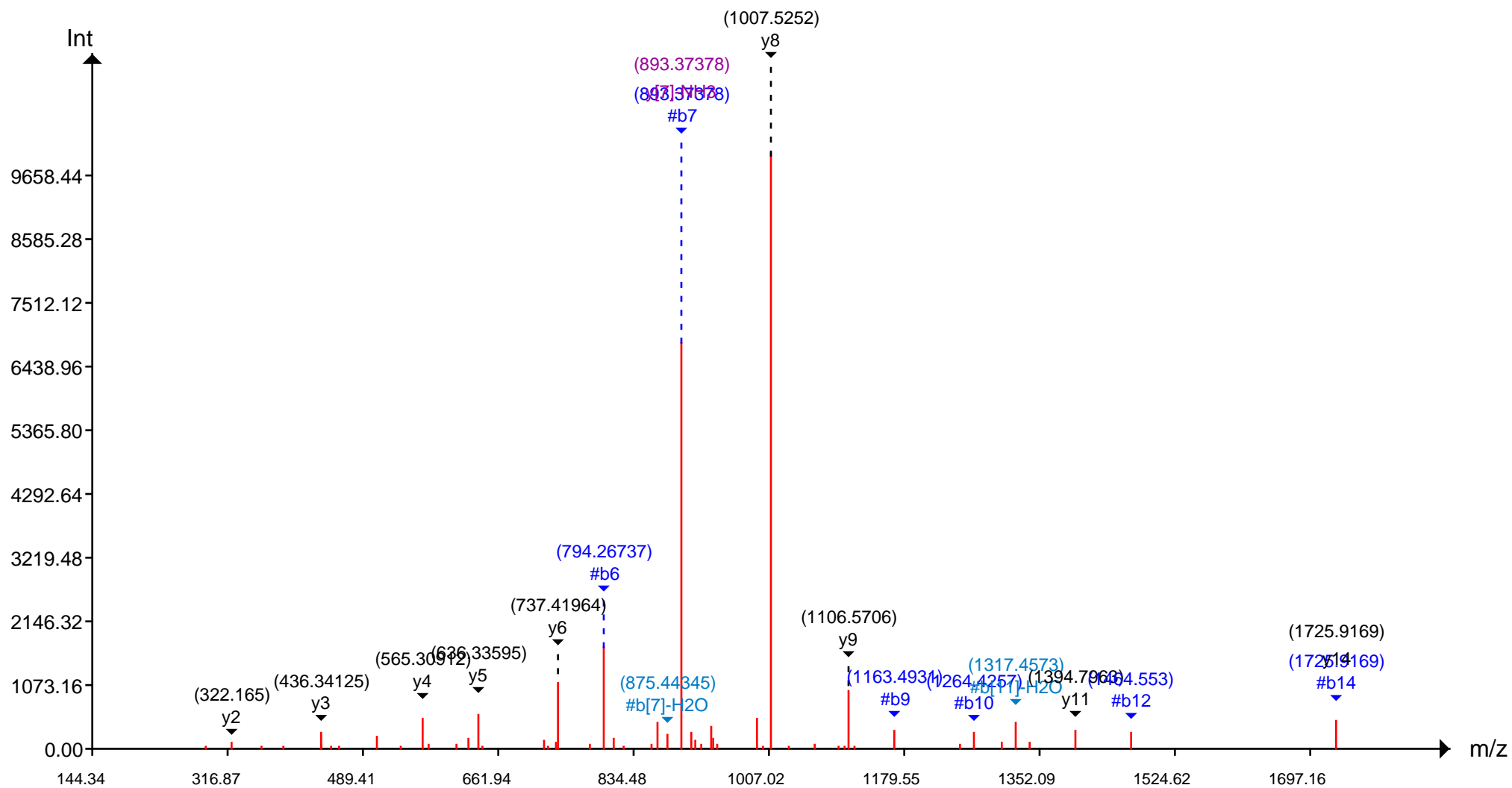
Info

Modified Sequence	AcD3-YVEK<AcD3K*>TAEDAK<AcD3K*>DFFK<AcD3K*>R-COOH	Ion score	73.85
Protein	Q9WU28	Confidence level	99.00%
Precursor m/z	1014.0309	Identity Threshold	48.09
Charge	2+	Mass error	0.003562



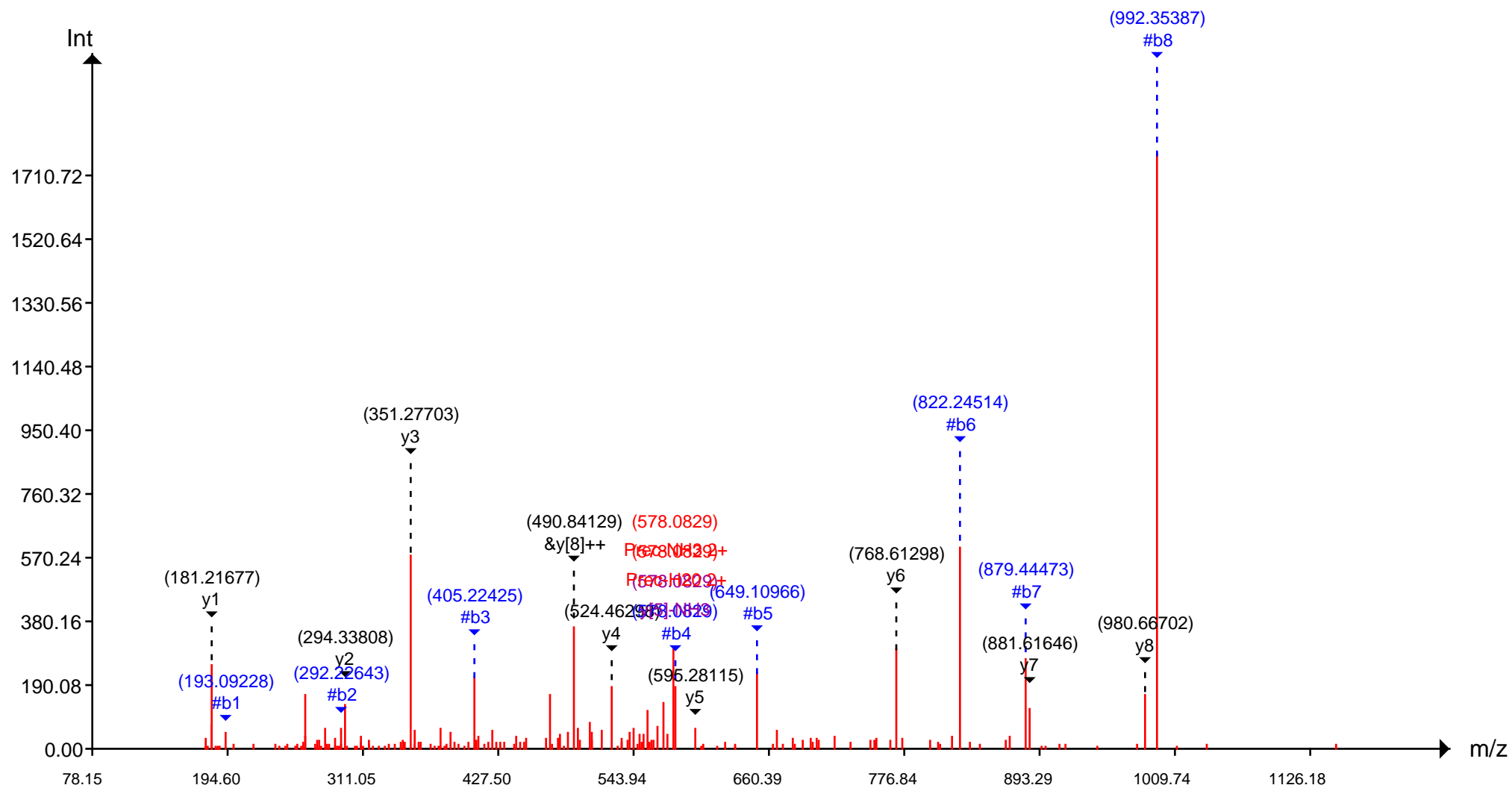
Info

Modified Sequence	AcD3-YVEK<AcD3K*>TAEDAK<AcD3K*>DFFK<AcD3K*>R-COOH	Ion score	66.43
Protein	Q9WU28	Confidence level	99.00%
Precursor m/z	1014.0316	Identity Threshold	48.09
Charge	2+	Mass error	0.004962



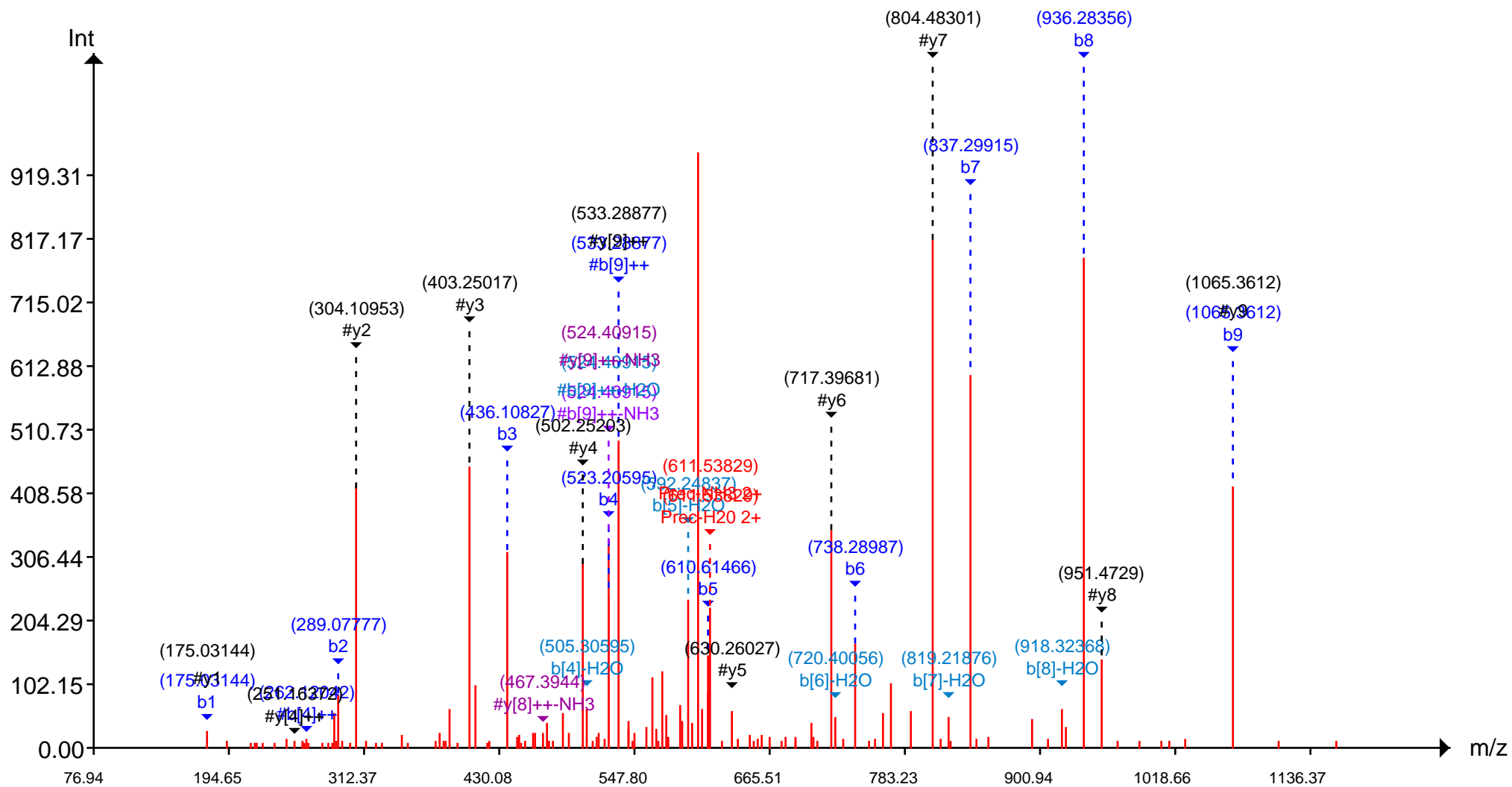
Info

Modified Sequence	AcD3-ELFADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH	Ion score	93.26
Protein	P17742	Confidence level	99.00%
Precursor m/z	950.51328	Identity Threshold	47.69
Charge	2+	Mass error	0.003204



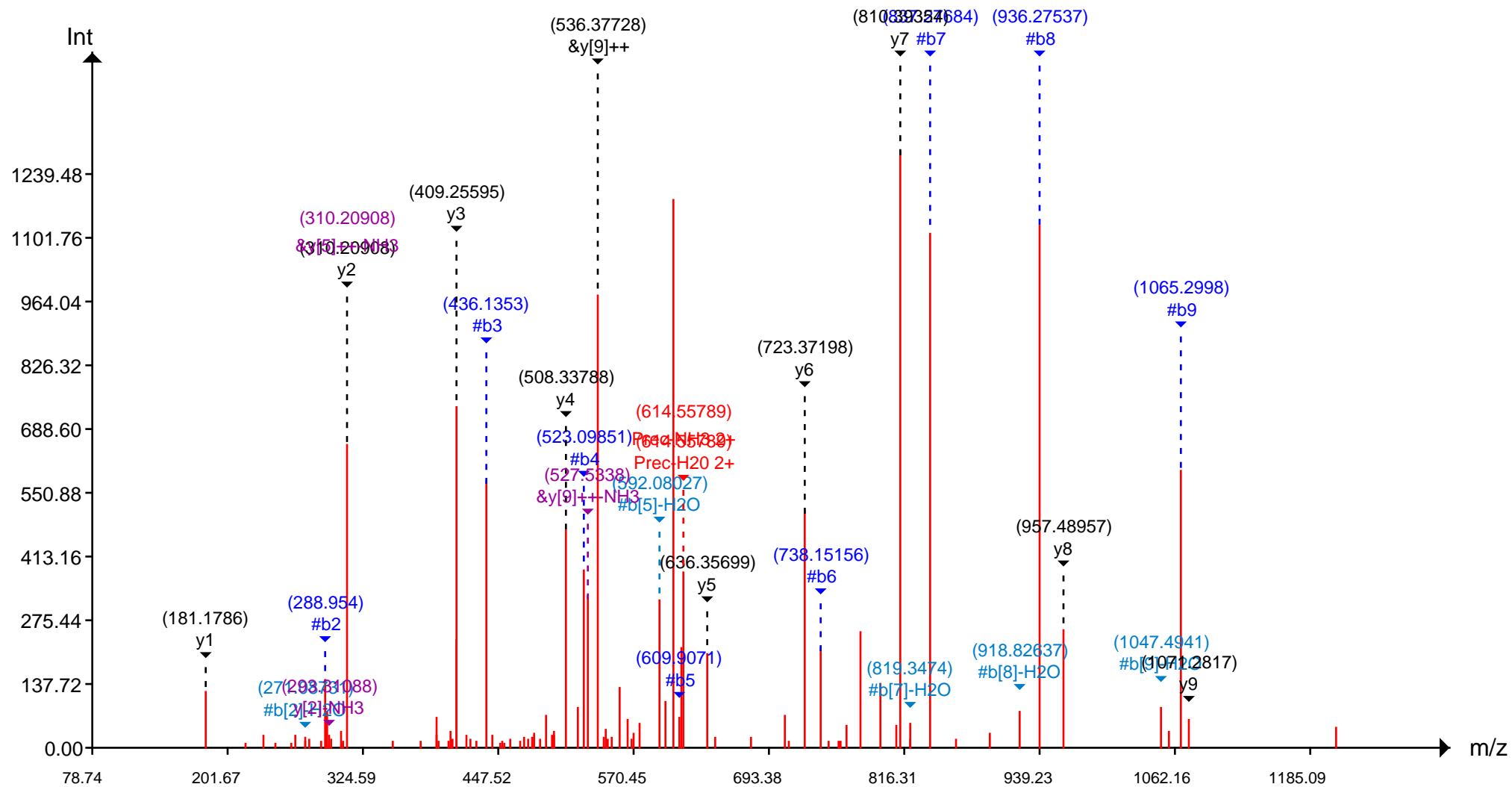
Info

Modified Sequence	AcD3-M<Mox*>VLK<AcD3K*>AK<AcD3K*>GIR<C13>-COOH	Ion score	52.86
Protein	O09131	Confidence level	99.00%
Precursor m/z	586.87739	Identity Threshold	40.35
Charge	2+	Mass error	-2.05E-4



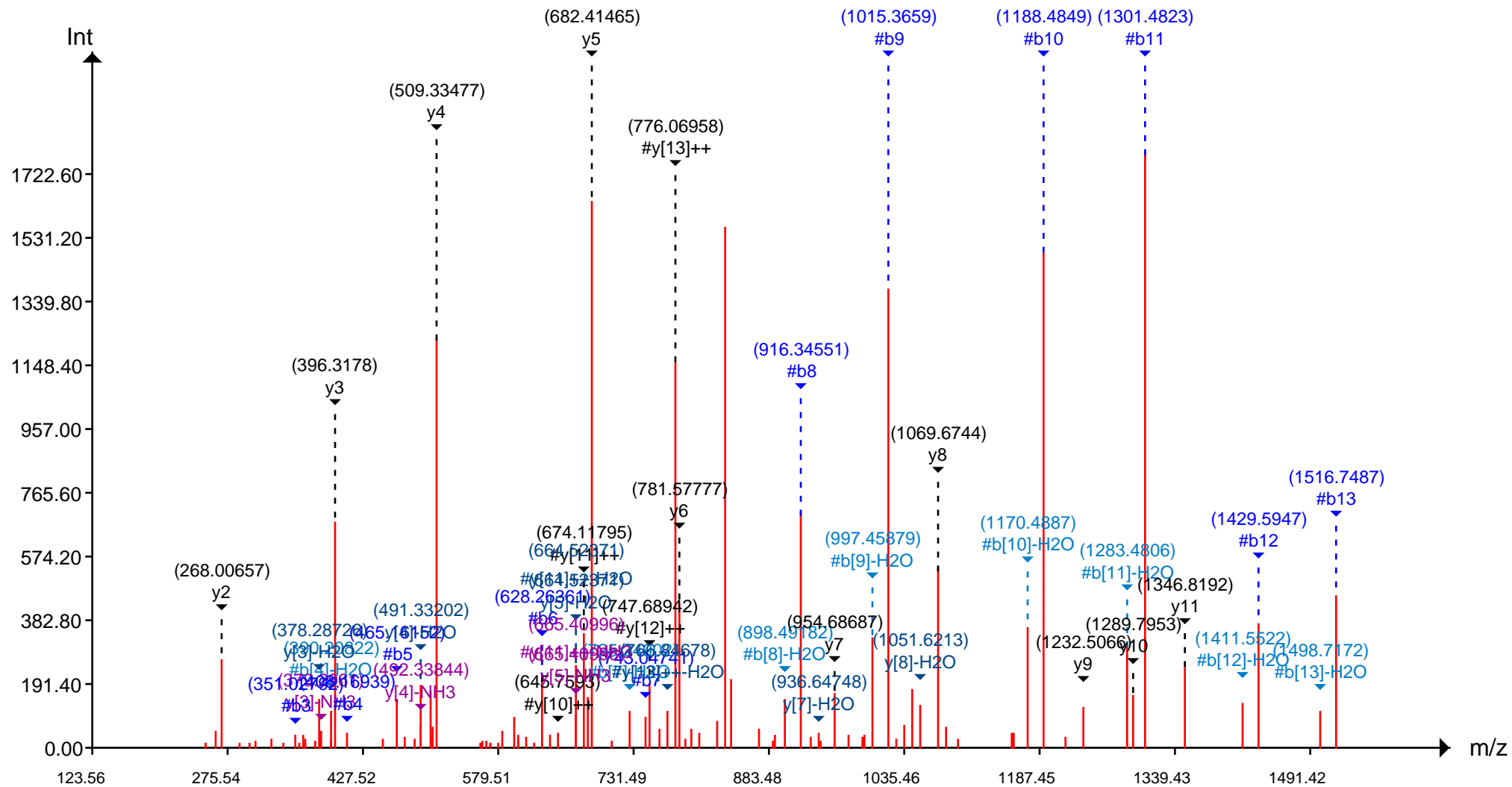
Info

Modified Sequence	AcD3-ENFSSQVVER-COOH	Ion score	72.83
Protein	P59999	Confidence level	99.00%
Precursor m/z	620.3059	Identity Threshold	47.22
Charge	2+	Mass error	2.02E-4



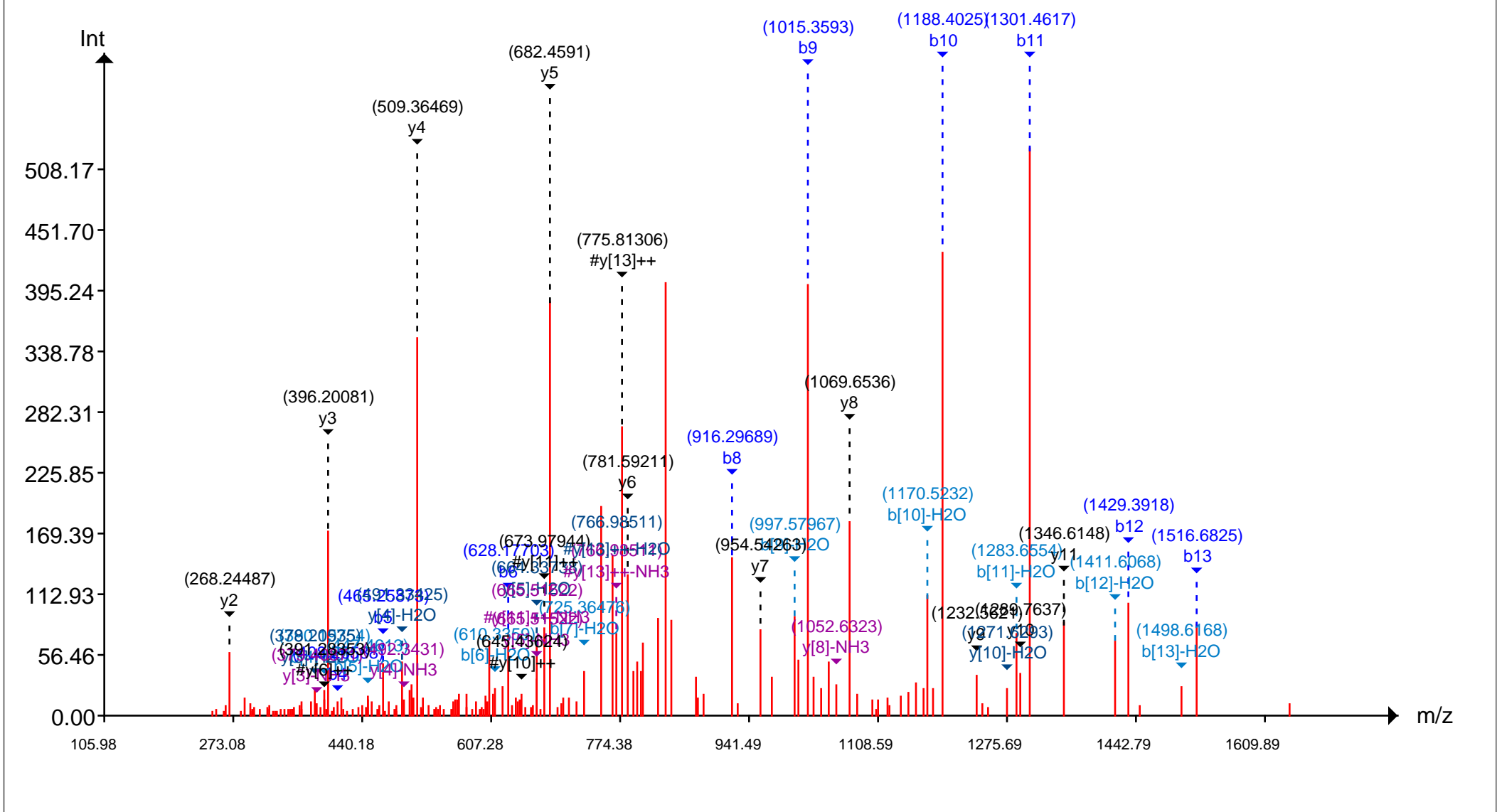
Info

Modified Sequence	AcD3-ENFSSQVVER<C13>-COOH	Ion score	67.61
Protein	P59999	Confidence level	99.00%
Precursor m/z	623.31596	Identity Threshold	47.52
Charge	2+	Mass error	1.8E-4



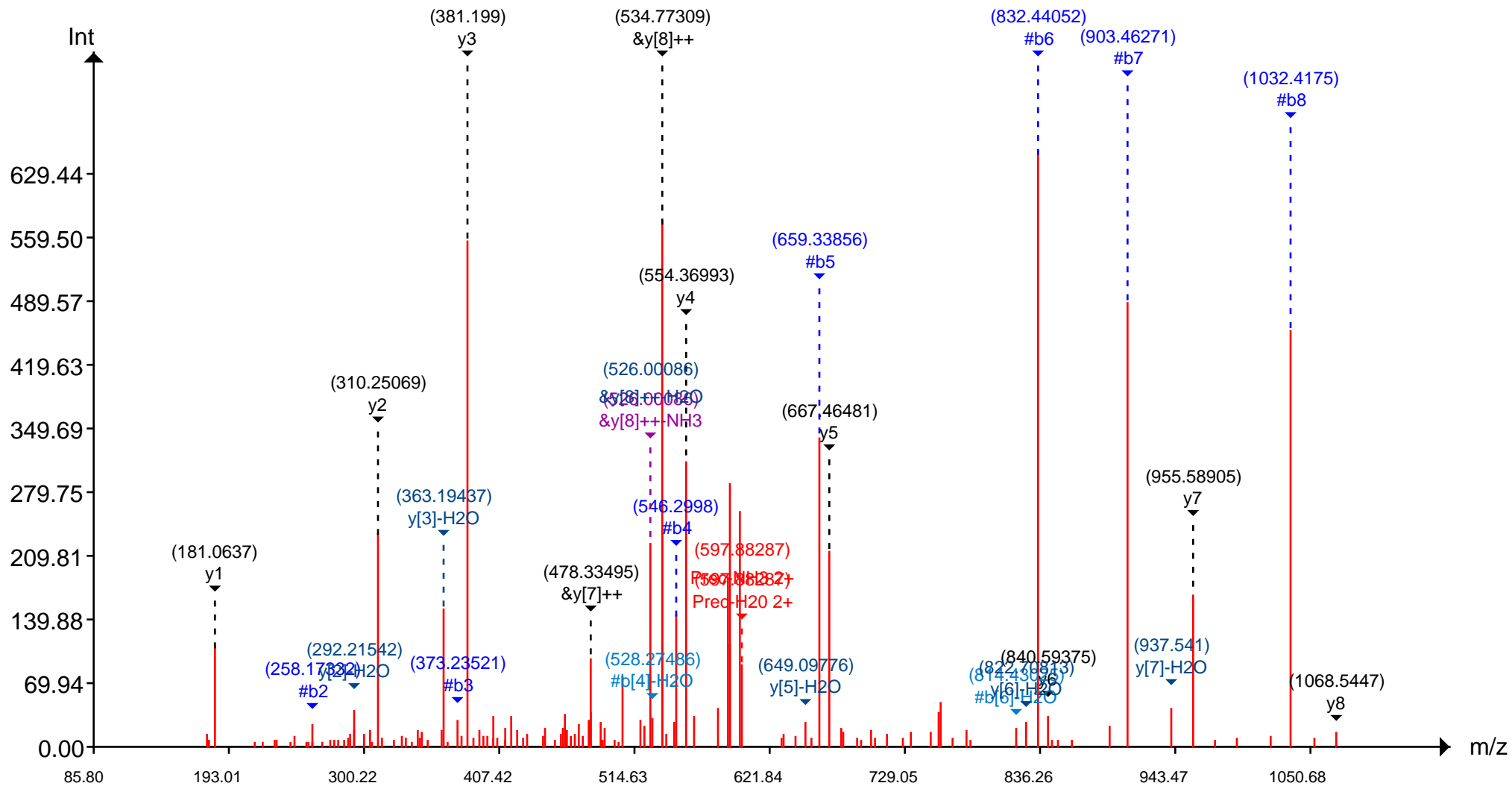
Info

Modified Sequence	AcD3-TGFGGYDK<AcD3K*>VK<AcD3K*>LQSR<C13>-COOH	Ion score	78.01
Protein	Q62465	Confidence level	99.00%
Precursor m/z	848.97006	Identity Threshold	47.25
Charge	2+	Mass error	0.001786



Info

Modified Sequence	AcD3-TGFGGYDK<AcD3K*>VK<AcD3K*>LQSR<C13>-COOH	Ion score	74.46
Protein	Q62465	Confidence level	99.00%
Precursor m/z	848.97005	Identity Threshold	47.25
Charge	2+	Mass error	0.001766



Info

Modified Sequence AcD3-VLDK<AcD3K*>LK<AcD3K*>AER<C13>-COOH

Ion score 57.41

Protein P10126, P62631

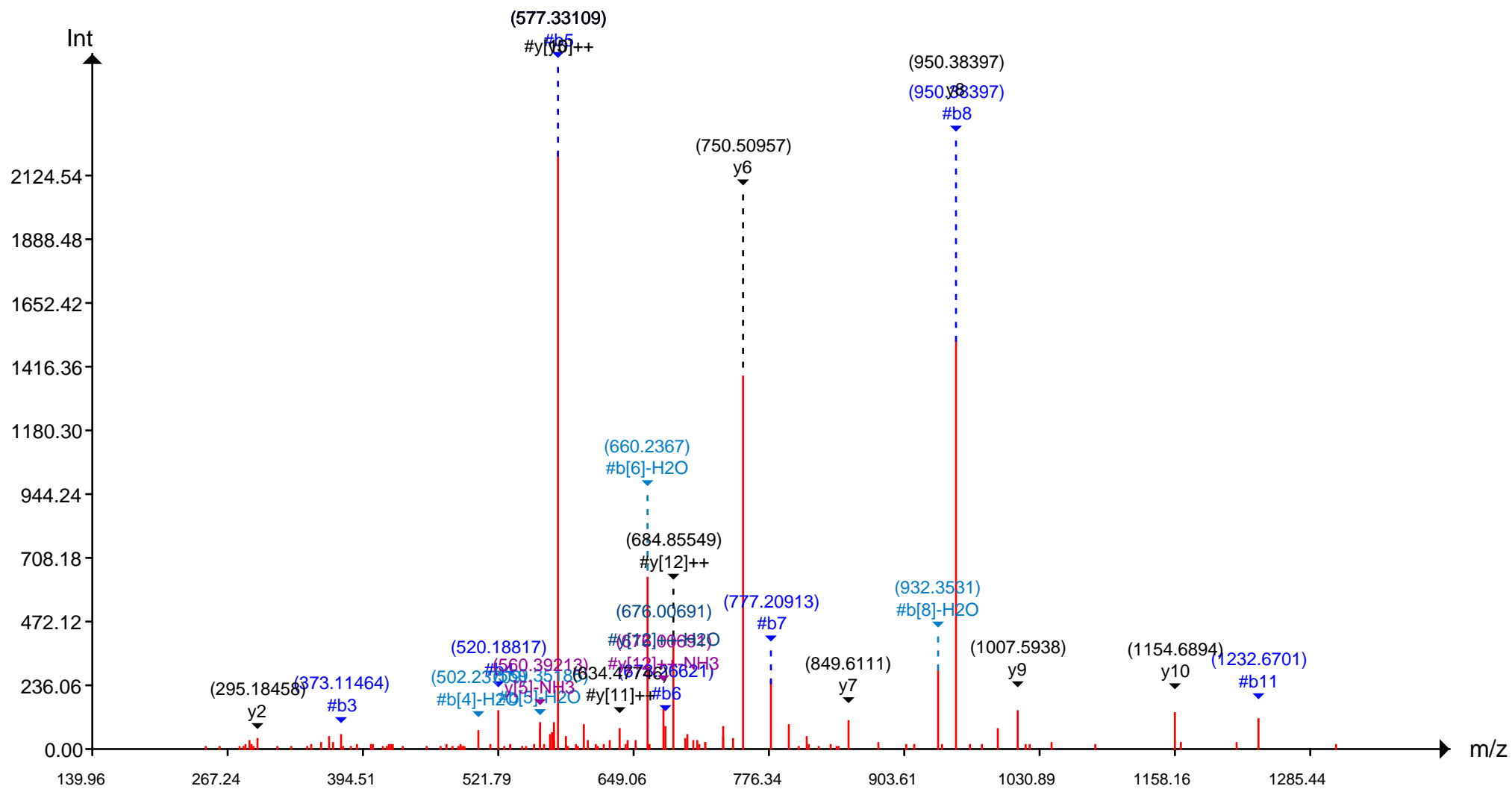
Confidence level 99.00%

Precursor m/z 606.88422

Identity Threshold 41.09

Charge 2+

Mass error 7.9E-4



Info

Modified Sequence AcD3-ITIM<Mox*>GTVK<AcD3K*>PNANR<C13>-COOH

Ion score 57.4

Protein P16110

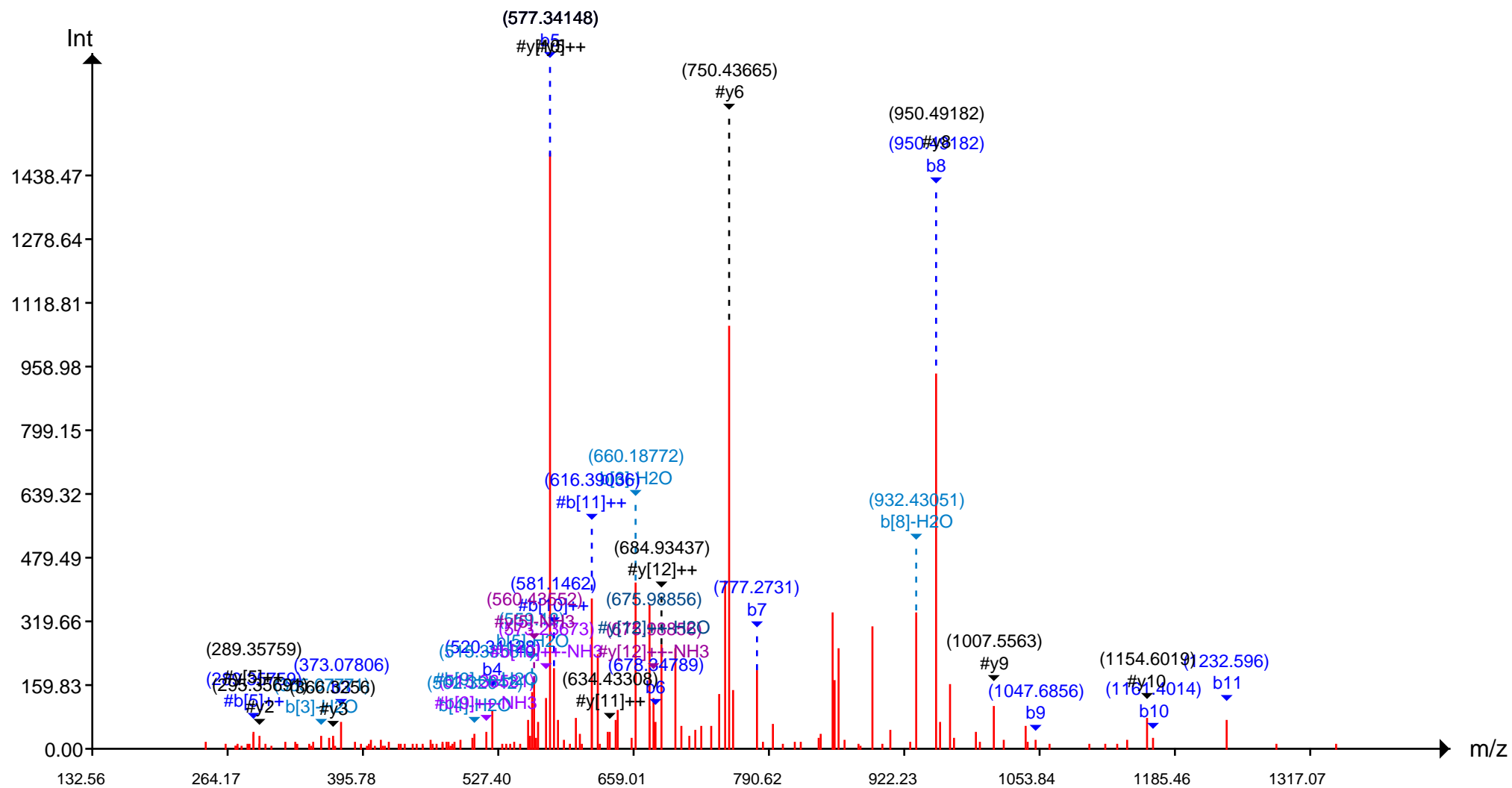
Confidence level 99.00%

Precursor m/z 763.93261

Identity Threshold 46.76

Charge 2+

Mass error 6.16E-4



Info

Modified Sequence AcD3-ITIM<Mox*>GTVK<AcD3K*>PNANR<C13>-COOH

Ion score 58.02

Protein P16110

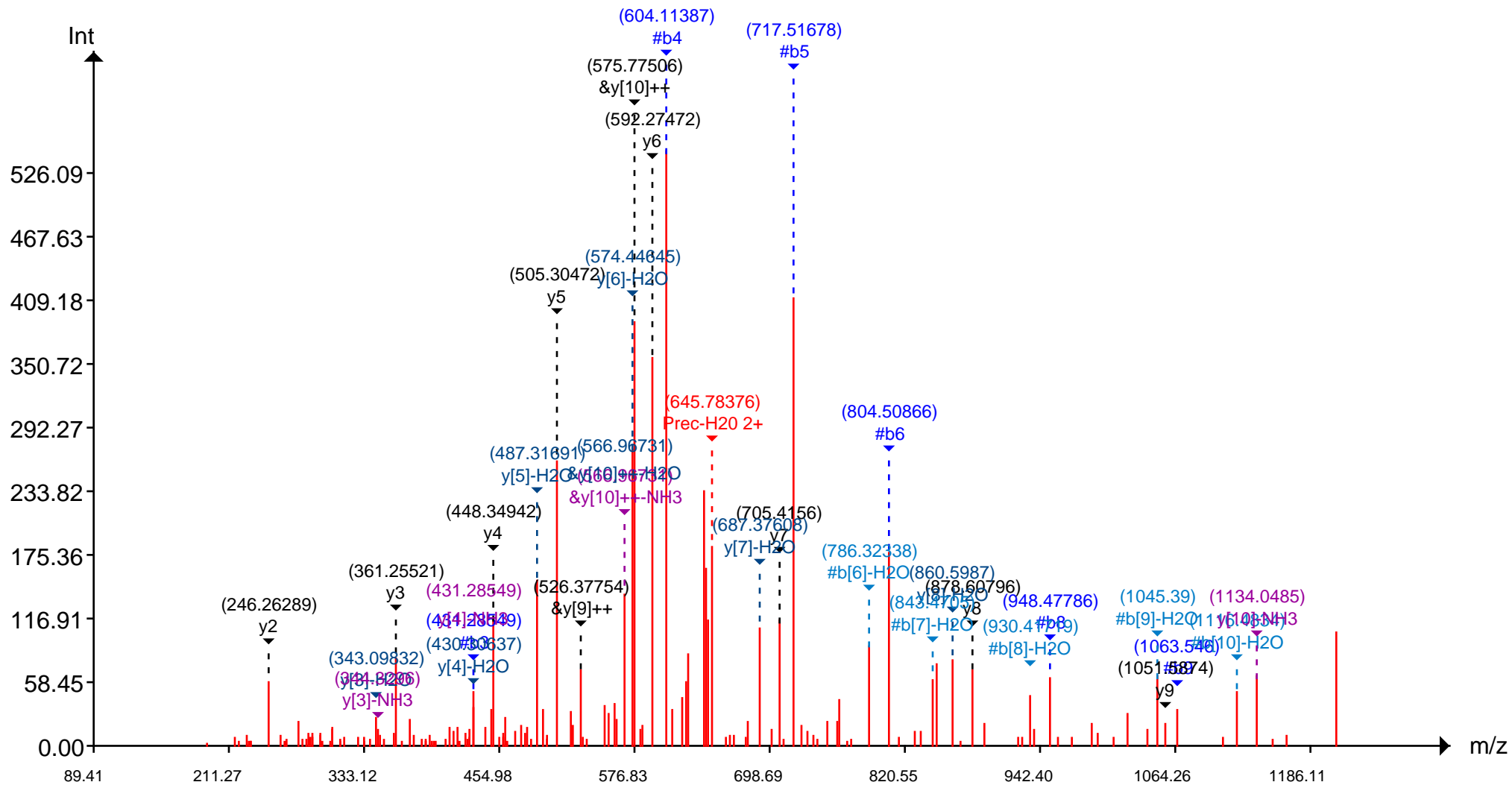
Confidence level 99.00%

Precursor m/z 763.93322

Identity Threshold 46.72

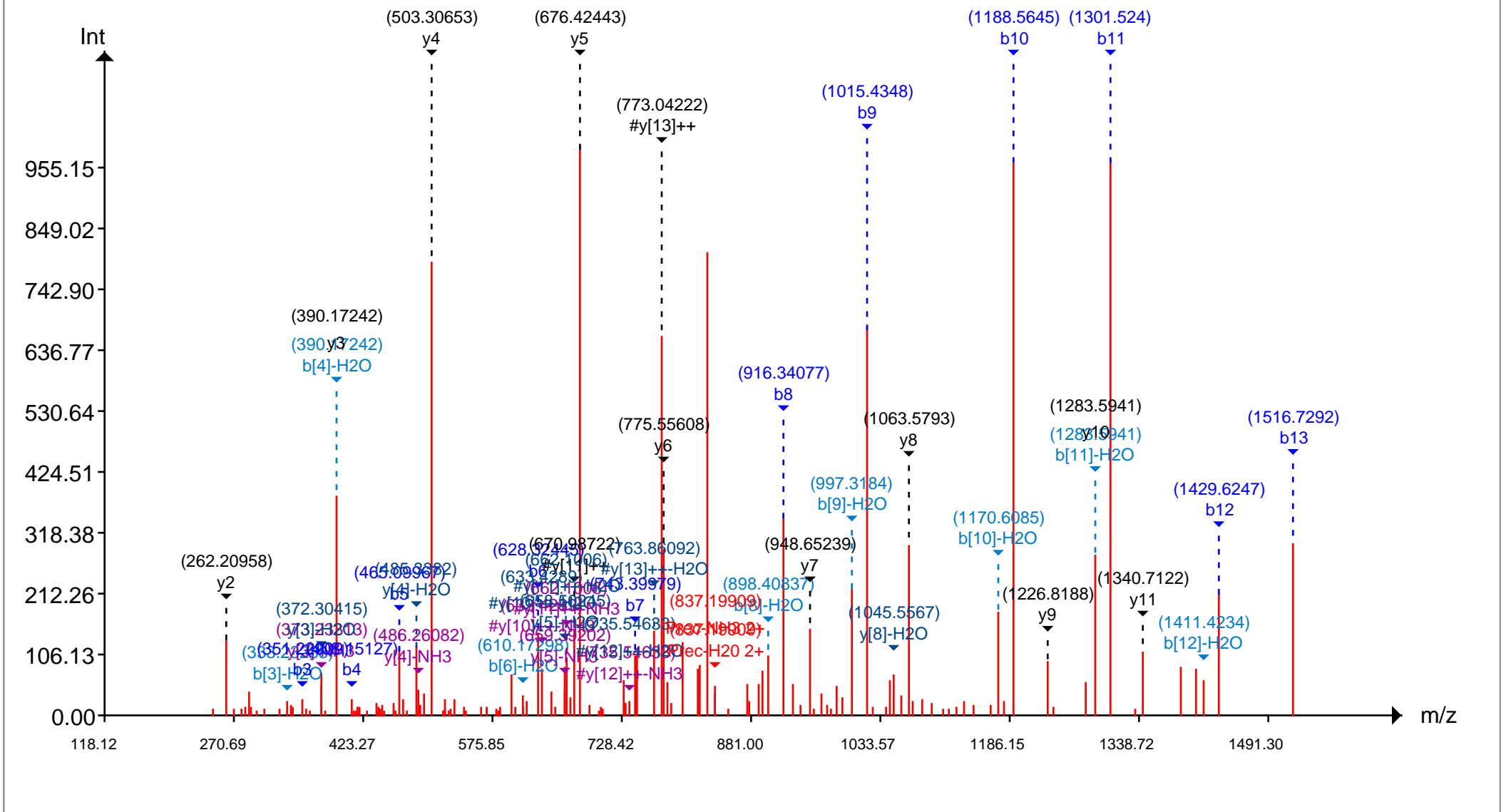
Charge 2+

Mass error 0.001836



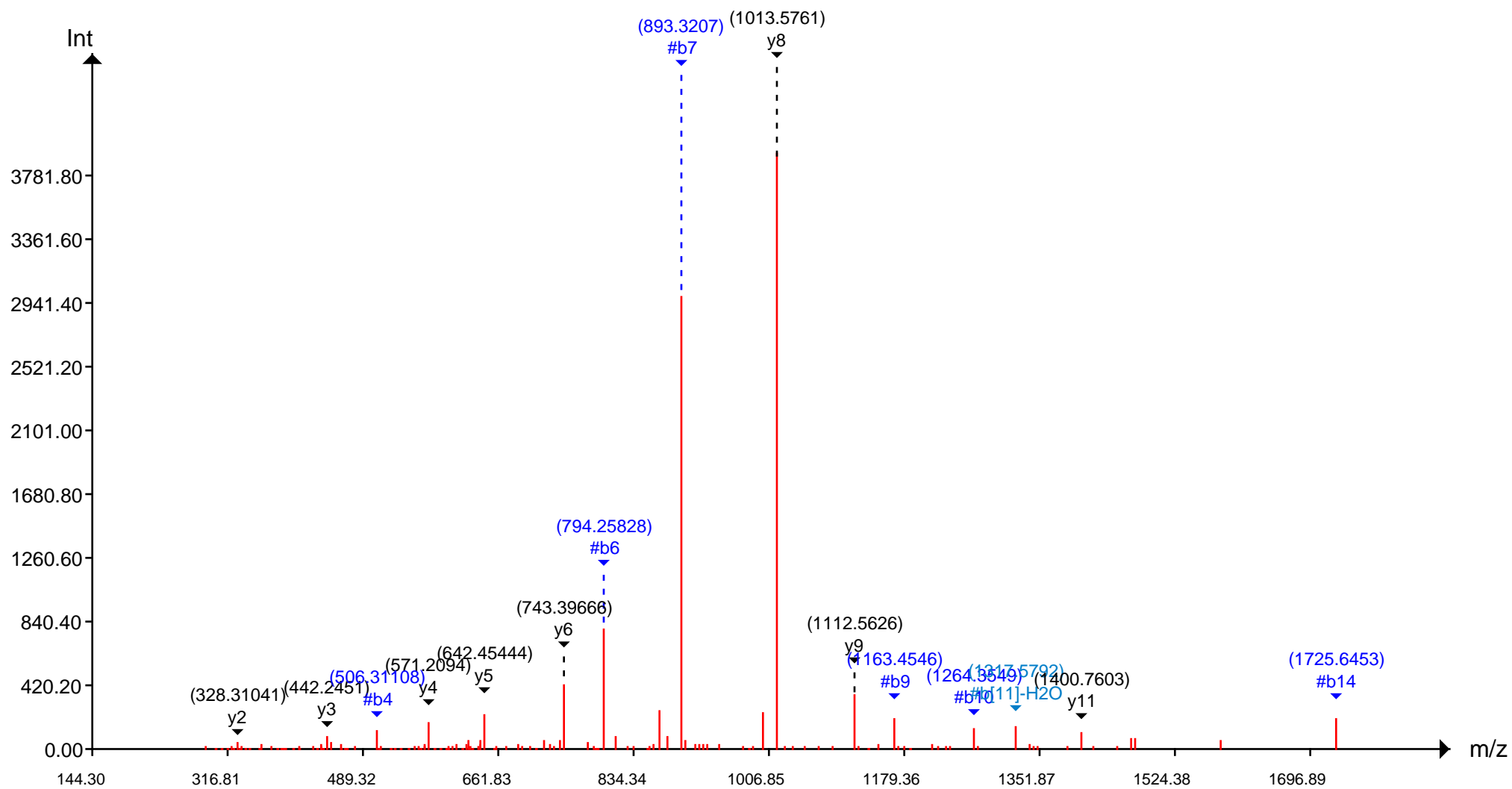
Info

Modified Sequence	AcD3-LVK<AcD3K*>K<AcD3K*>ISGSDAR-COOH	Ion score	54.62
Protein	P18654	Confidence level	99.00%
Precursor m/z	654.89589	Identity Threshold	43.94
Charge	2+	Mass error	0.001334



Info

Modified Sequence	AcD3-TGFGGYDK<AcD3K*>VK<AcD3K*>LQSR-COOH	Ion score	89.42
Protein	Q62465	Confidence level	99.00%
Precursor m/z	845.9605	Identity Threshold	47.63
Charge	2+	Mass error	0.002807



Info

Modified Sequence AcD3-ELFADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH

Ion score 74.6

Protein P17742

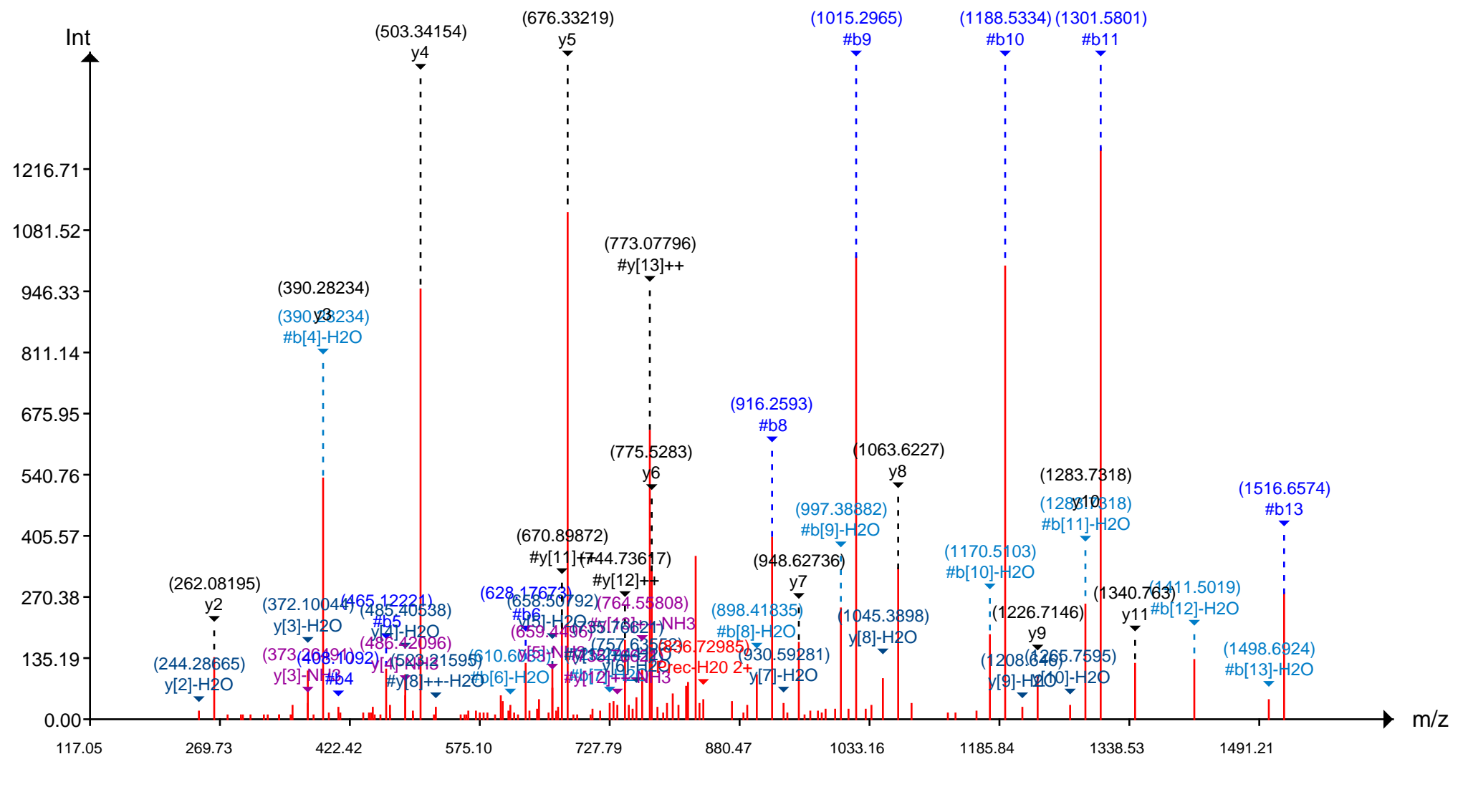
Confidence level 99.00%

Precursor m/z 953.52363

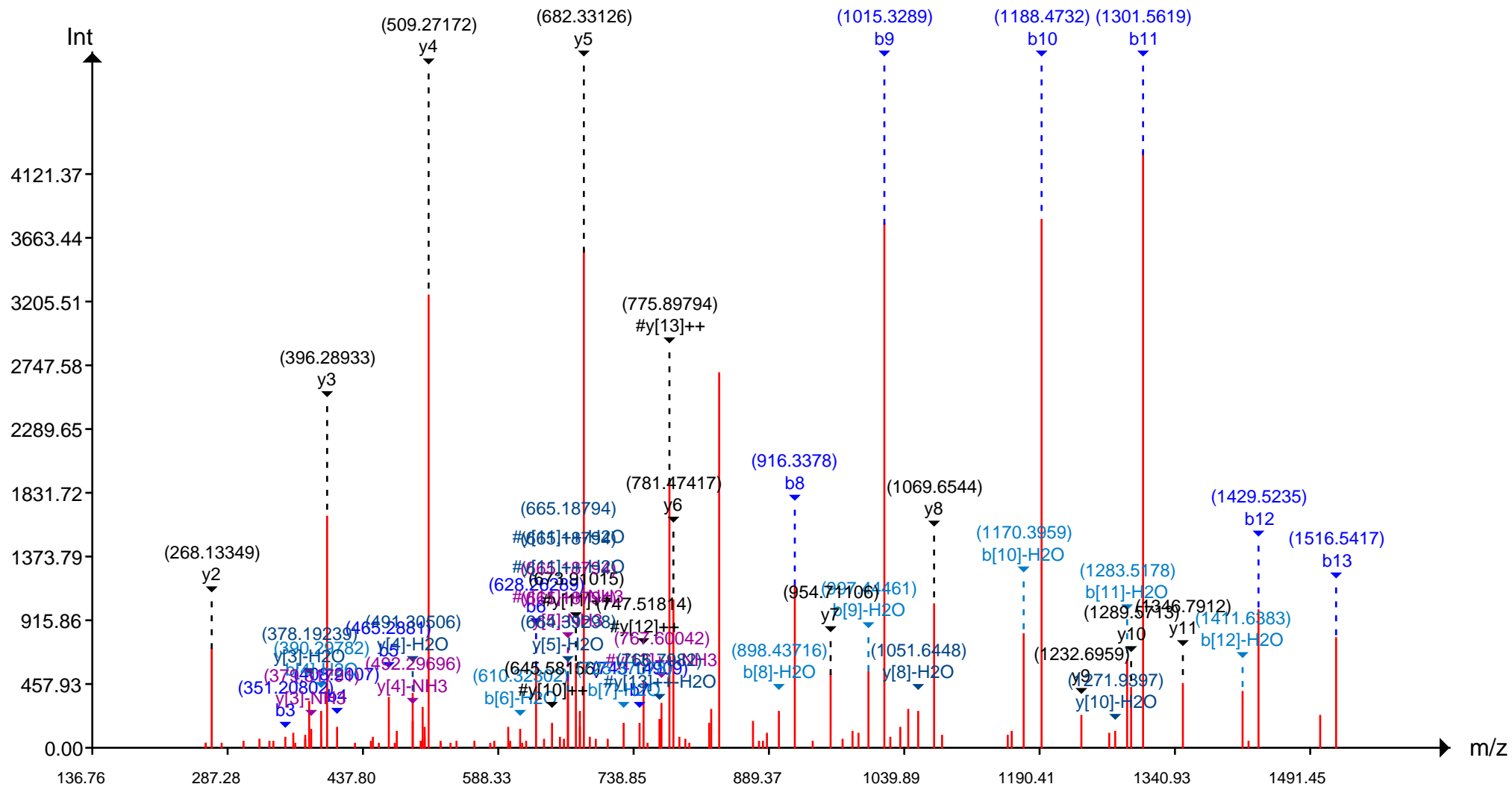
Identity Threshold 47.23

Charge 2+

Mass error 0.003762

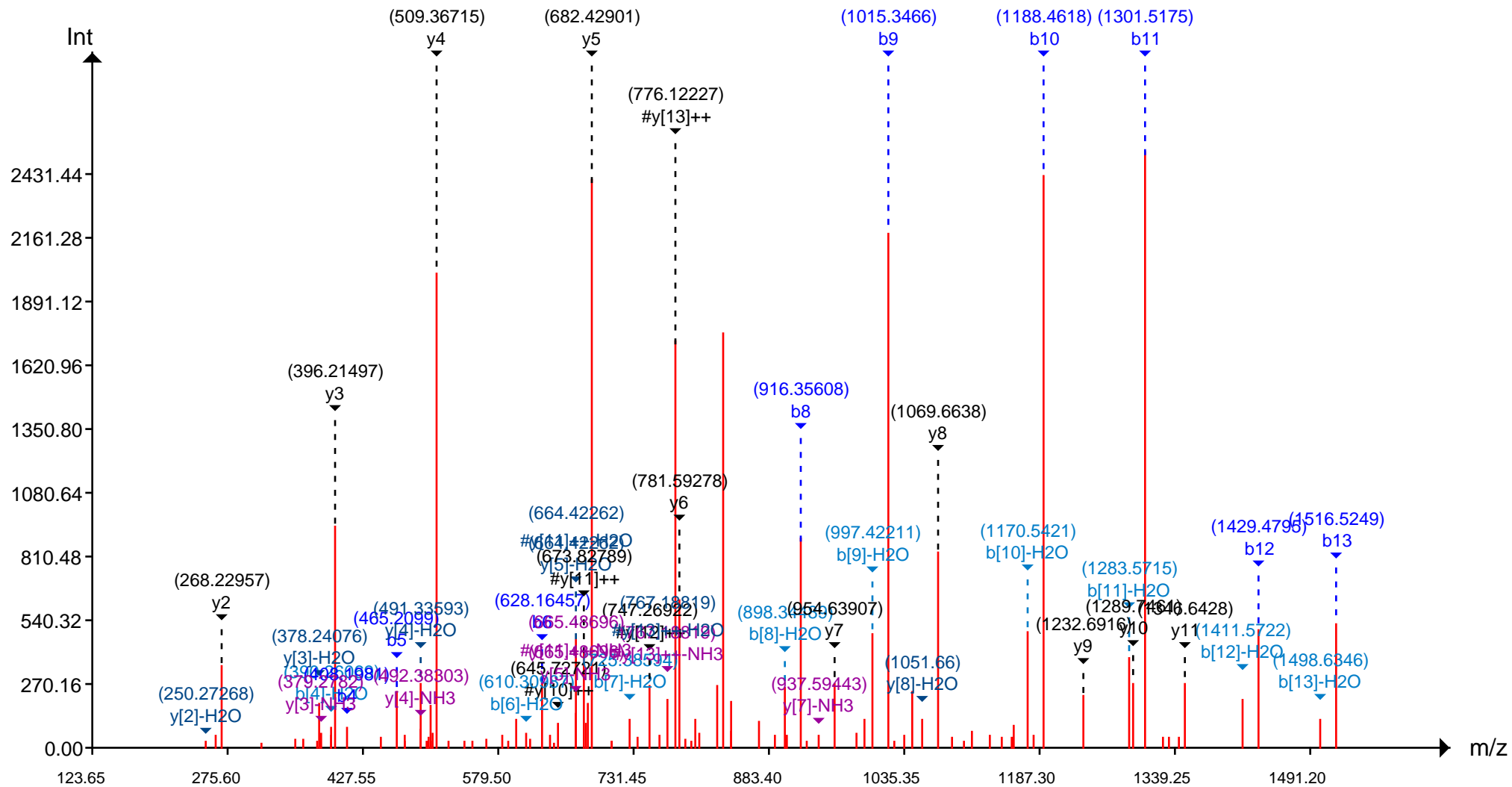


Info			
Modified Sequence	AcD3-TGFGGYDK<AcD3K*>VK<AcD3K*>LQSR-COOH	Ion score	89.93
Protein	Q62465	Confidence level	99.00%
Precursor m/z	845.96018	Identity Threshold	47.66
Charge	2+	Mass error	0.002167



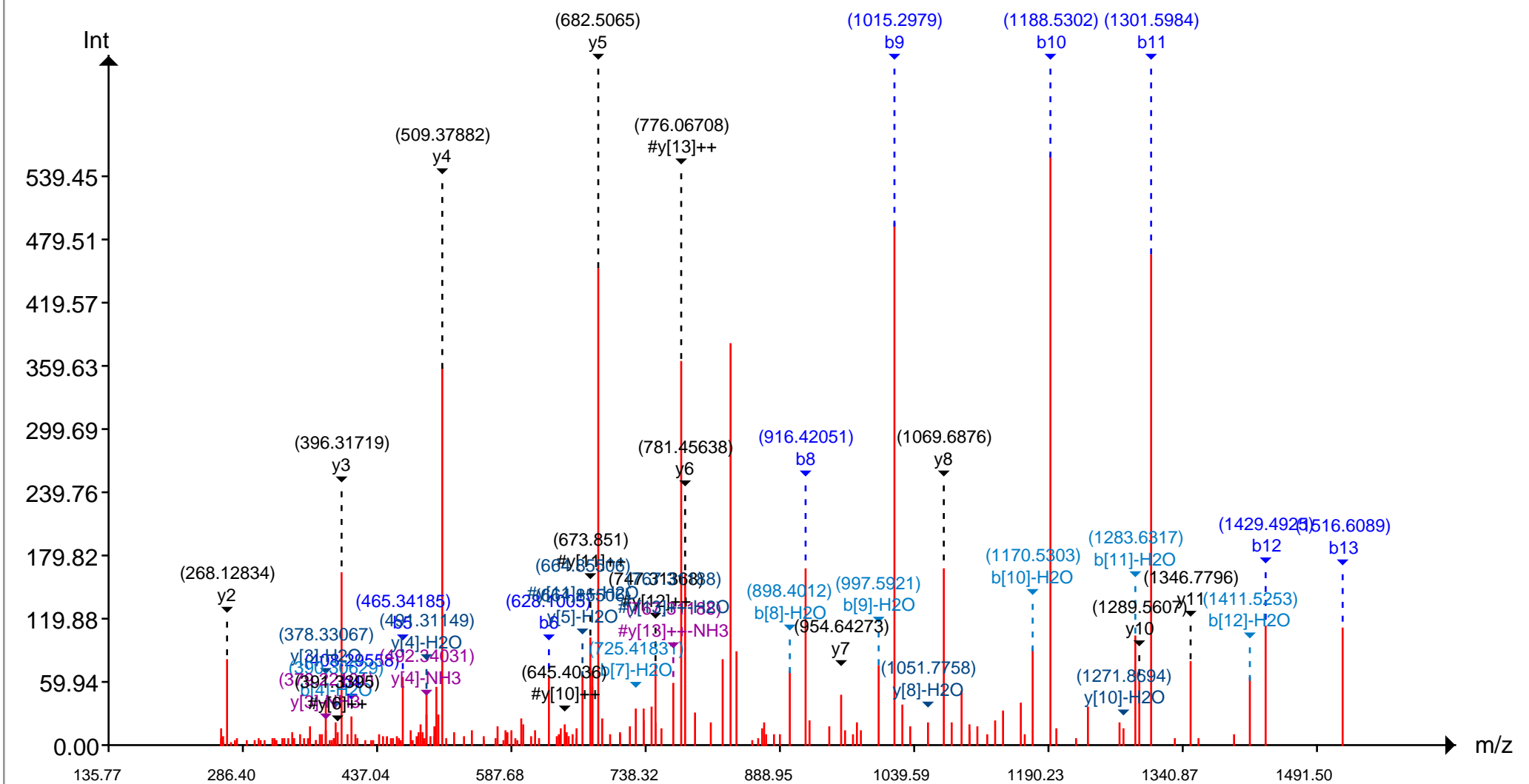
Info

Modified Sequence	AcD3-TGFGGYDK<AcD3K*>VK<AcD3K*>LQSR<C13>-COOH	Ion score	81.99
Protein	Q62465	Confidence level	99.00%
Precursor m/z	848.97008	Identity Threshold	47.25
Charge	2+	Mass error	0.001826



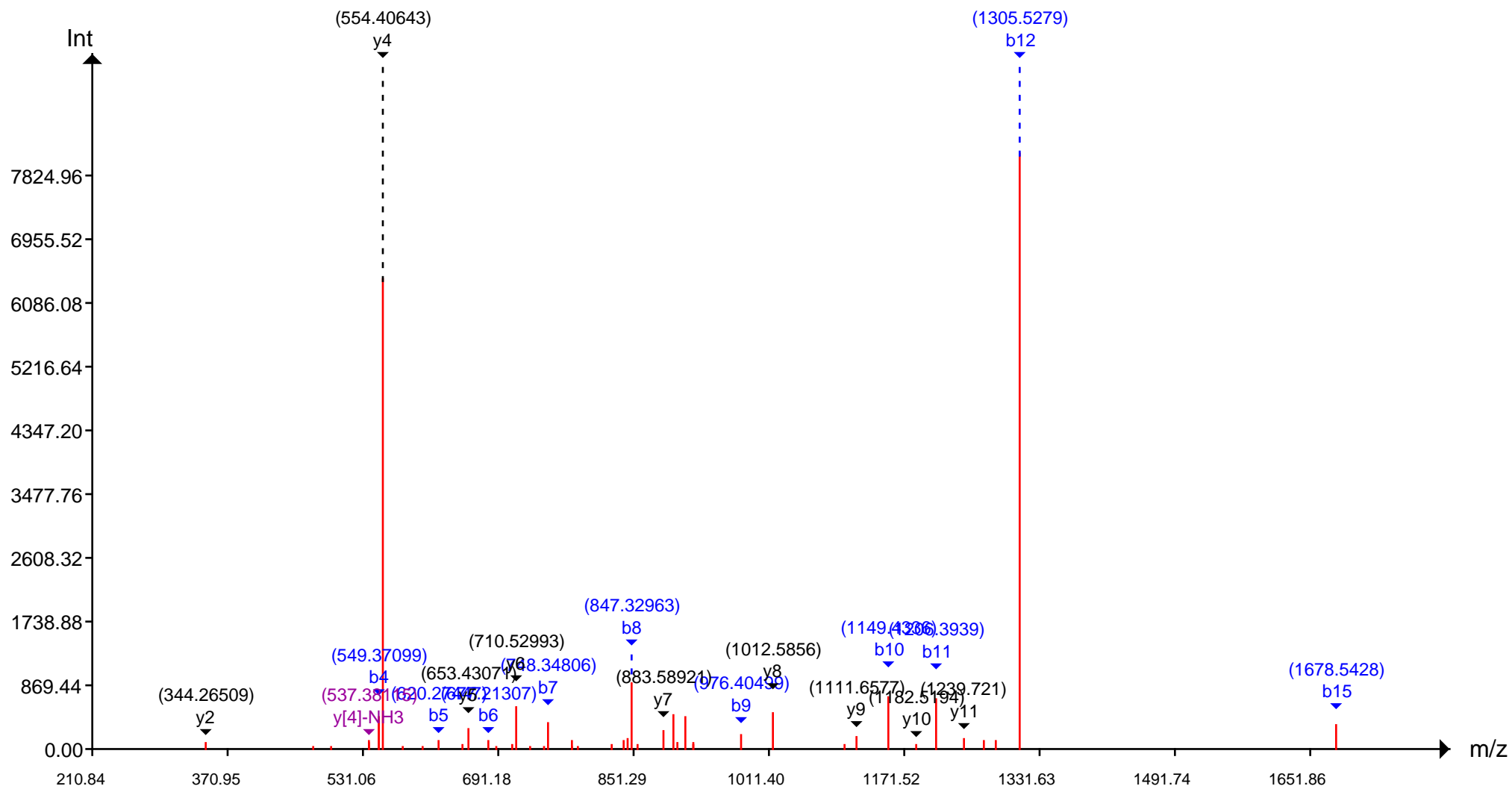
Info

Modified Sequence	AcD3-TGFGGYDK<AcD3K*>VK<AcD3K*>LQSR<C13>-COOH	Ion score	83.66
Protein	Q62465	Confidence level	99.00%
Precursor m/z	848.96973	Identity Threshold	47.30
Charge	2+	Mass error	0.001126



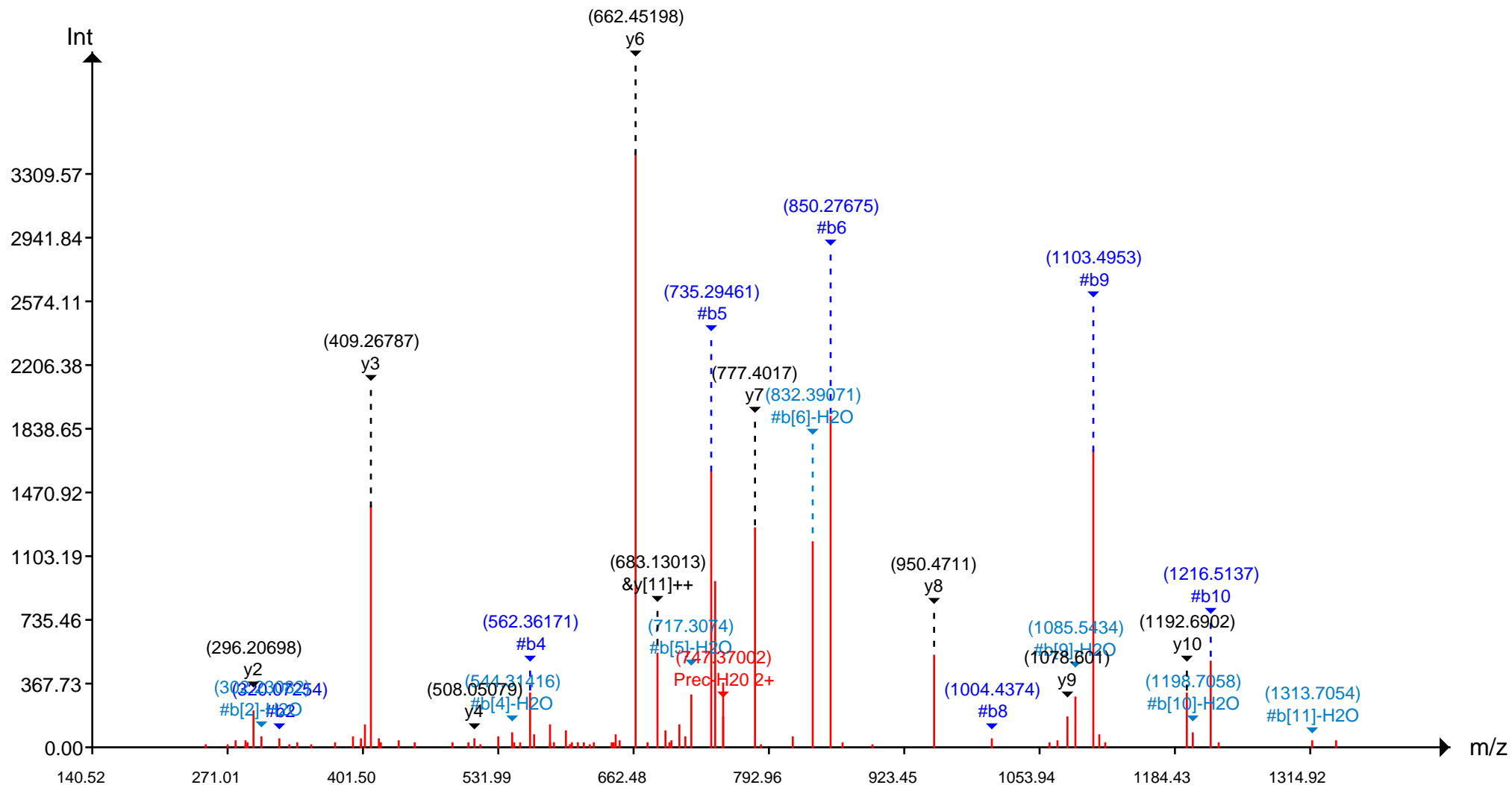
Info

Modified Sequence	AcD3-TGFGGYDK<AcD3K*>VK<AcD3K*>LQSR<C13>-COOH	Ion score	72.31
Protein	Q62465	Confidence level	99.00%
Precursor m/z	848.97071	Identity Threshold	47.28
Charge	2+	Mass error	0.003086



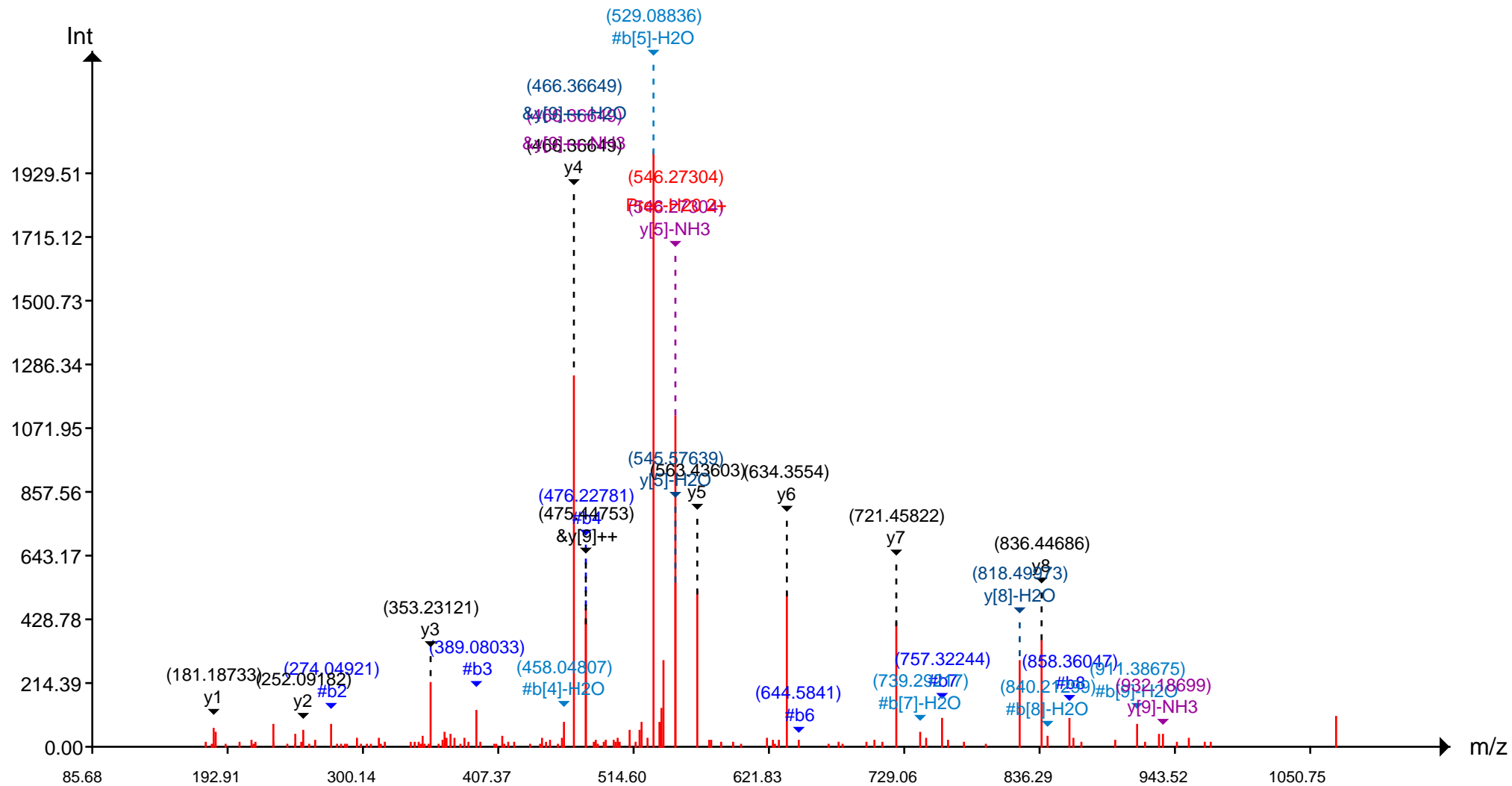
Info

Modified Sequence	AcD3-AVC<Cmm*>K<AcD3K*>AGAVEK<AcD3K*>GVPLYR<C13>-COOH	Ion score	70.78
Protein	P17182	Confidence level	99.00%
Precursor m/z	930.02982	Identity Threshold	45.68
Charge	2+	Mass error	0.002257



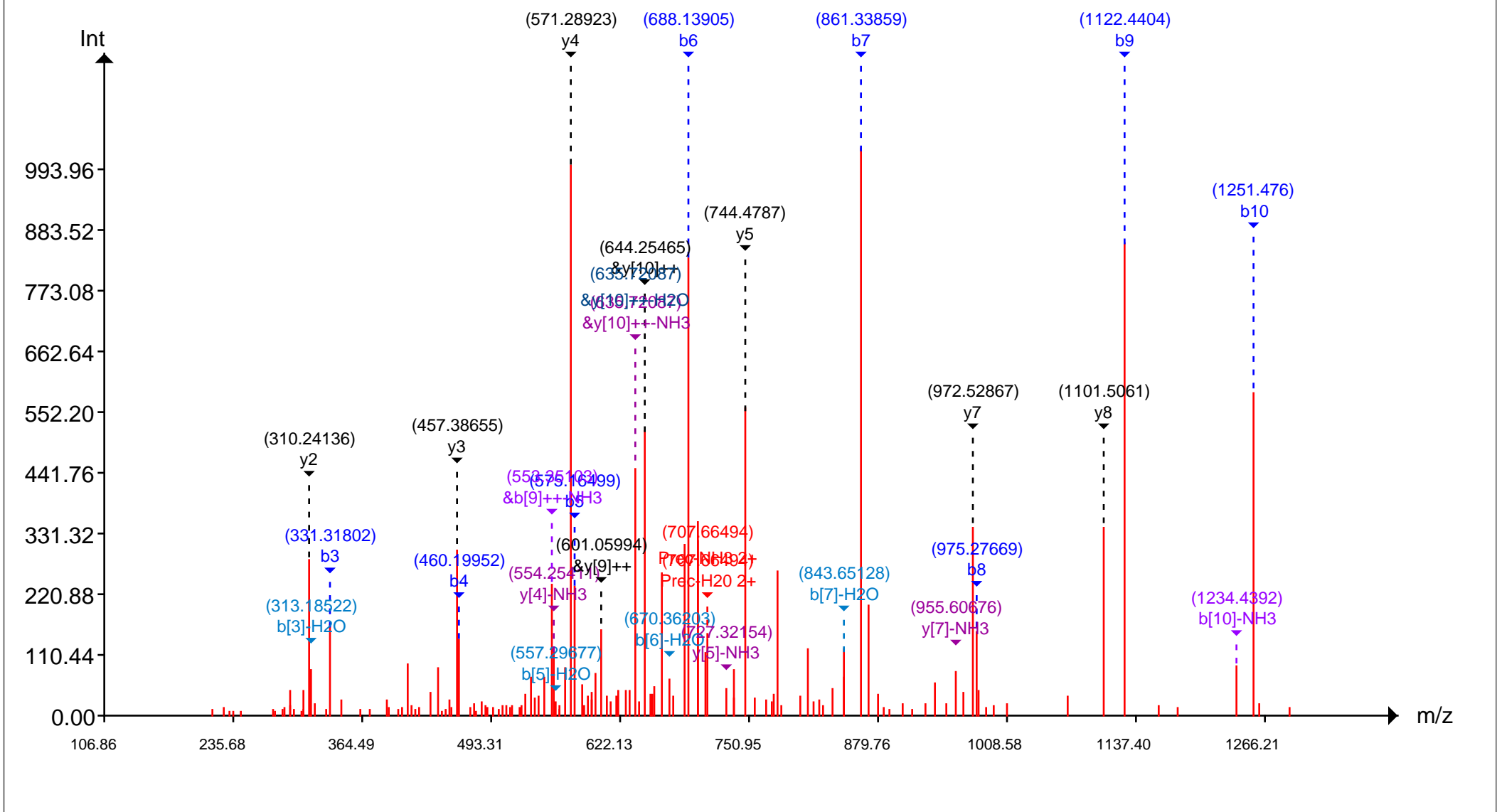
Info

Modified Sequence	AcD3-TK<AcD3K*>NQK<AcD3K*>DPGVLDLDR<C13>-COOH	Ion score	63.62
Protein	P50543	Confidence level	99.00%
Precursor m/z	756.42724	Identity Threshold	46.81
Charge	2+	Mass error	2.22E-4



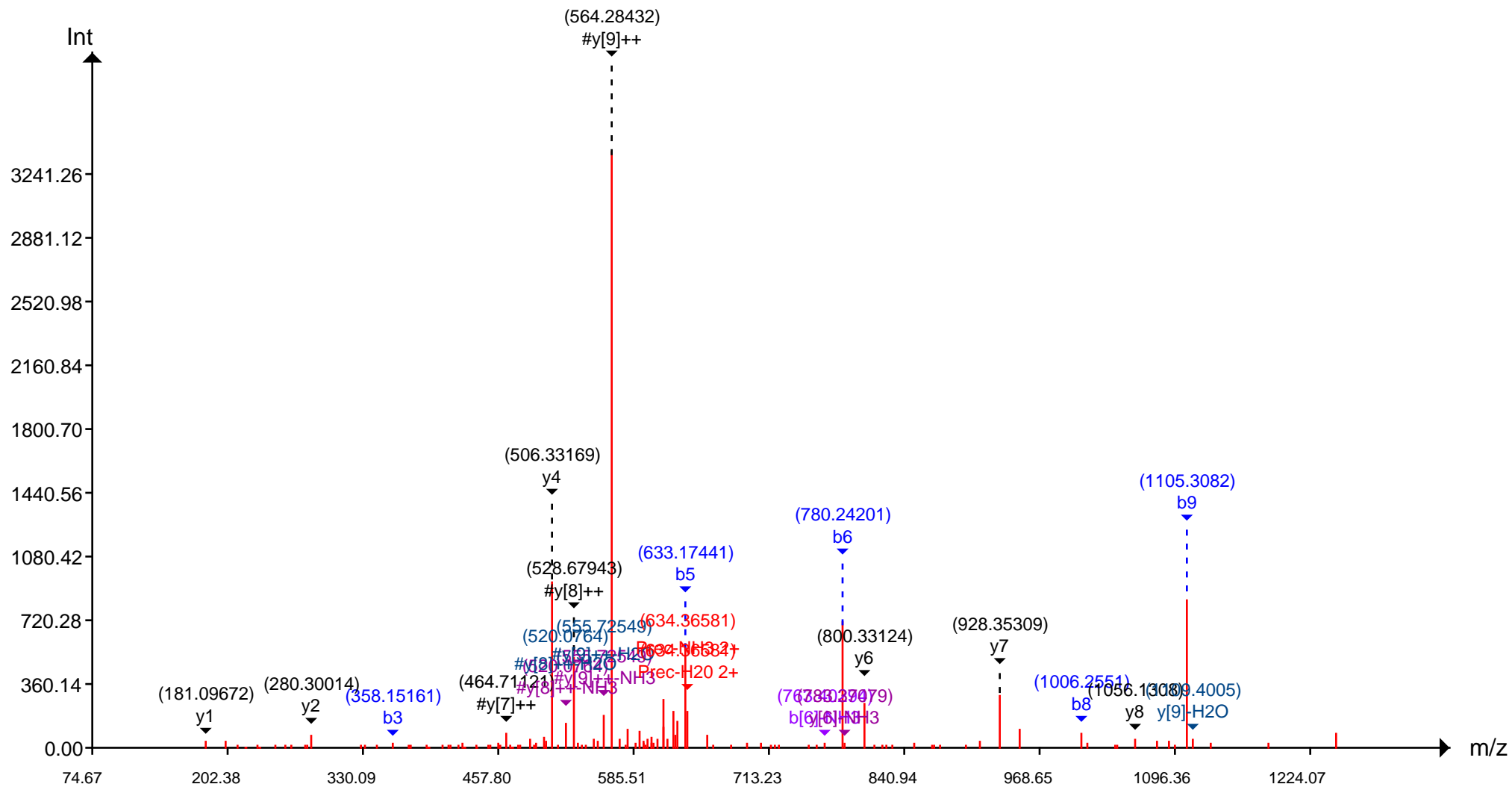
Info

Modified Sequence	AcD3-DIDSAPITAR<C13>-COOH	Ion score	67.96
Protein	P52480	Confidence level	99.00%
Precursor m/z	555.3021	Identity Threshold	47.83
Charge	2+	Mass error	-2.57E-4



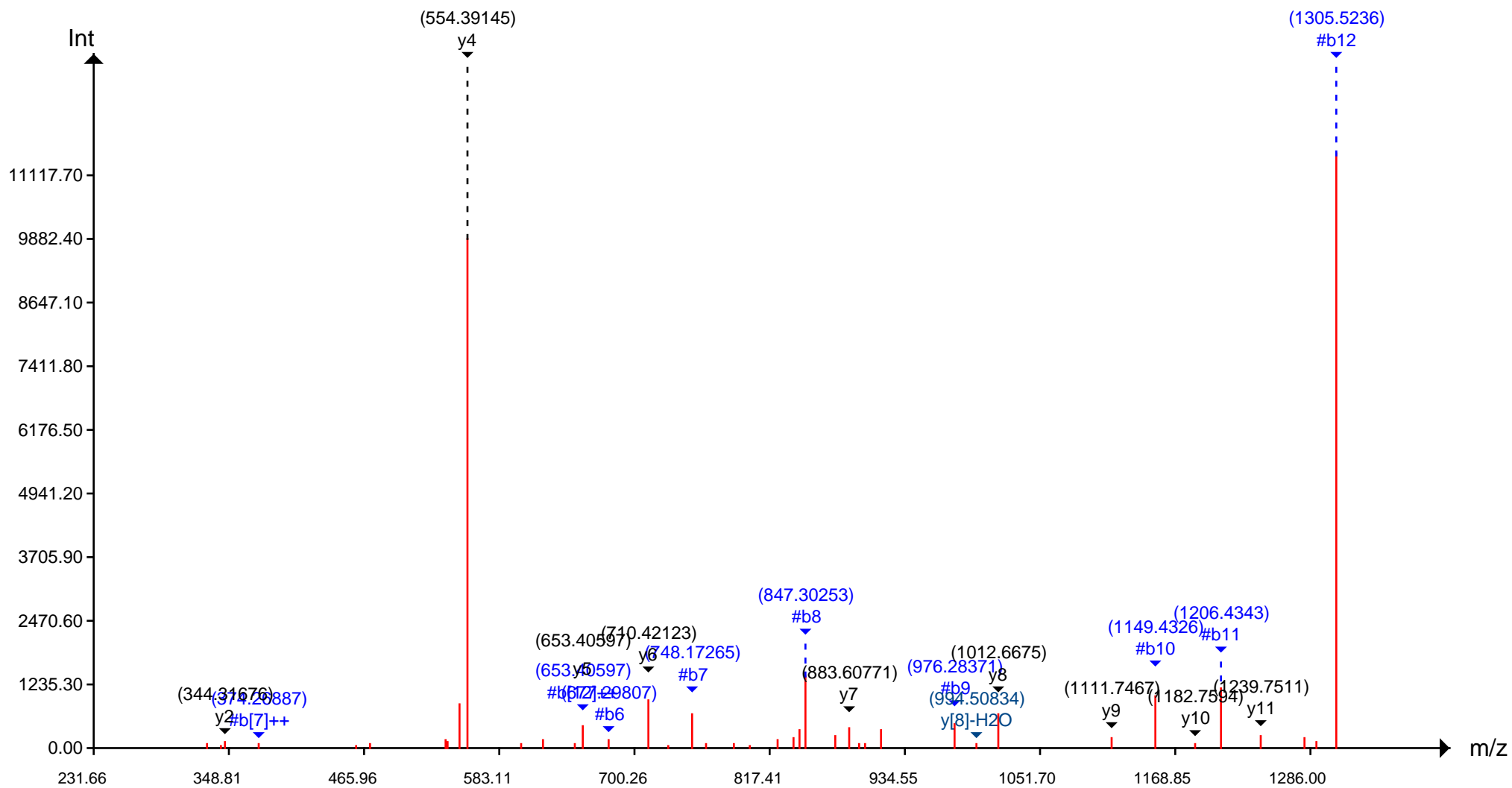
Info

Modified Sequence	AcD3-VSVEDLK<AcD3K*>NFER<C13>-COOH	Ion score	64.28
Protein	Q9CQ92	Confidence level	99.00%
Precursor m/z	716.38877	Identity Threshold	47.84
Charge	2+	Mass error	0.00104



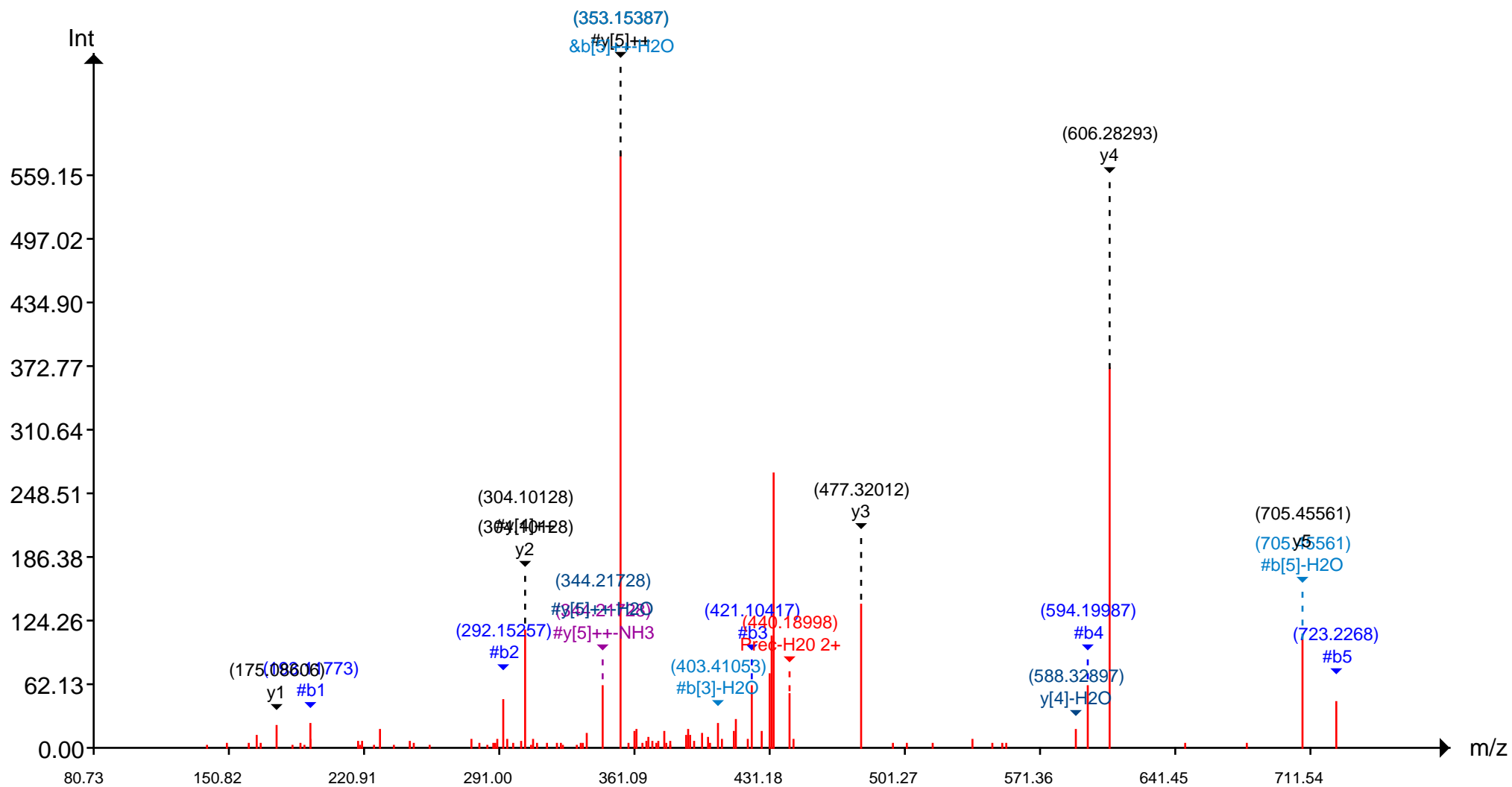
Info

Modified Sequence	AcD3-IAQQM<Mox*>M<Mox*>PEVR<C13>-COOH	Ion score	48.34
Protein	P50518	Confidence level	99.00%
Precursor m/z	643.32432	Identity Threshold	47.44
Charge	2+	Mass error	-6.0E-6



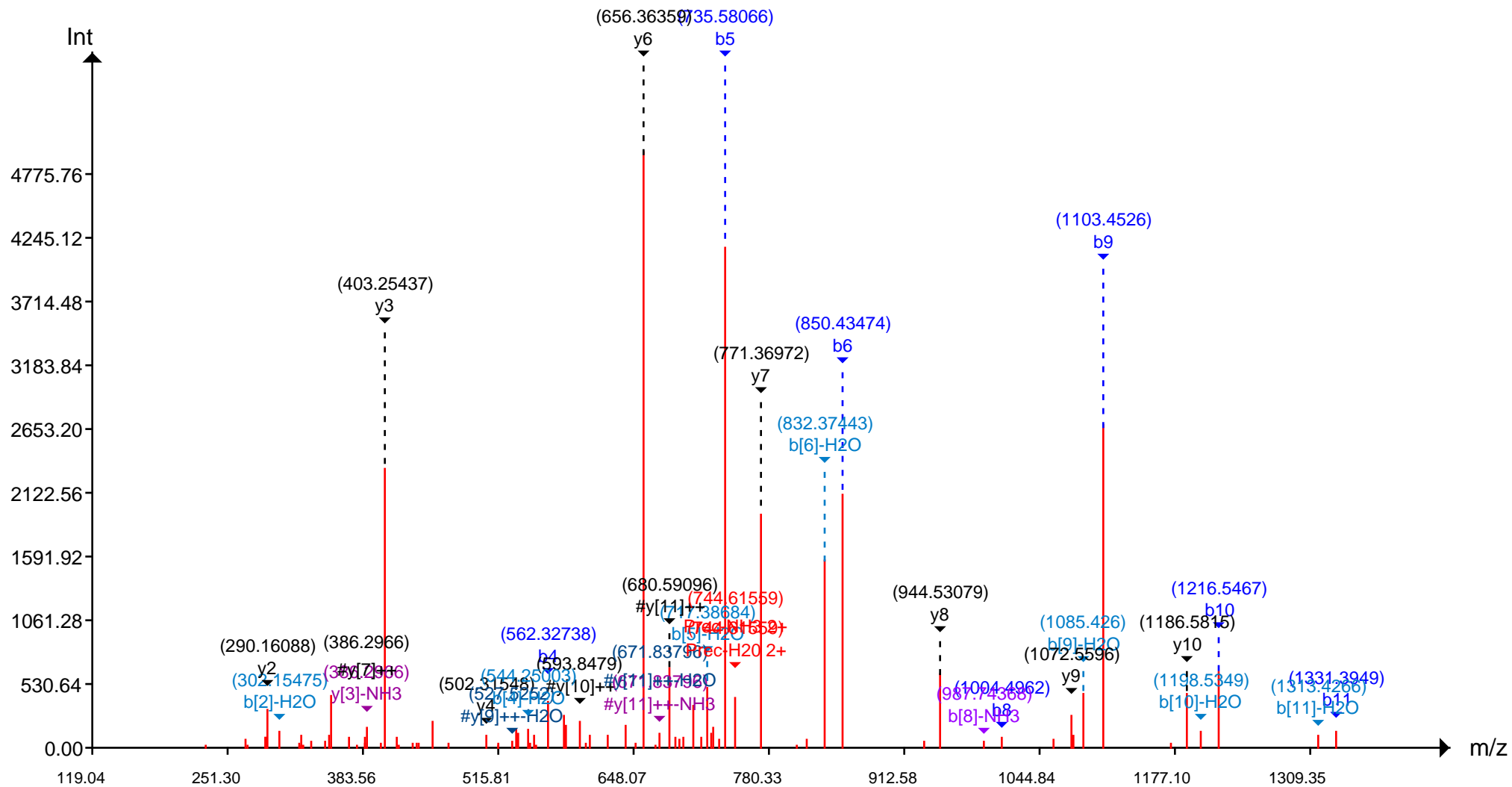
Info

Modified Sequence	AcD3-AVC<Cmm*>K<AcD3K*>AGAVEK<AcD3K*>GVPLYR<C13>-COOH	Ion score	65.11
Protein	P17182	Confidence level	99.00%
Precursor m/z	930.02842	Identity Threshold	45.93
Charge	2+	Mass error	-5.43E-4



Info

Modified Sequence	AcD3-FVEK<AcD3K*>ER-COOH	Ion score	48.79
Protein	P07901, Q9WU62	Confidence level	99.00%
Precursor m/z	449.251	Identity Threshold	47.61
Charge	2+	Mass error	2.1E-5



Info

Modified Sequence AcD3-TK<AcD3K*>NQK<AcD3K*>DPGVLDLDR-COOH

Ion score 70.72

Protein P50543

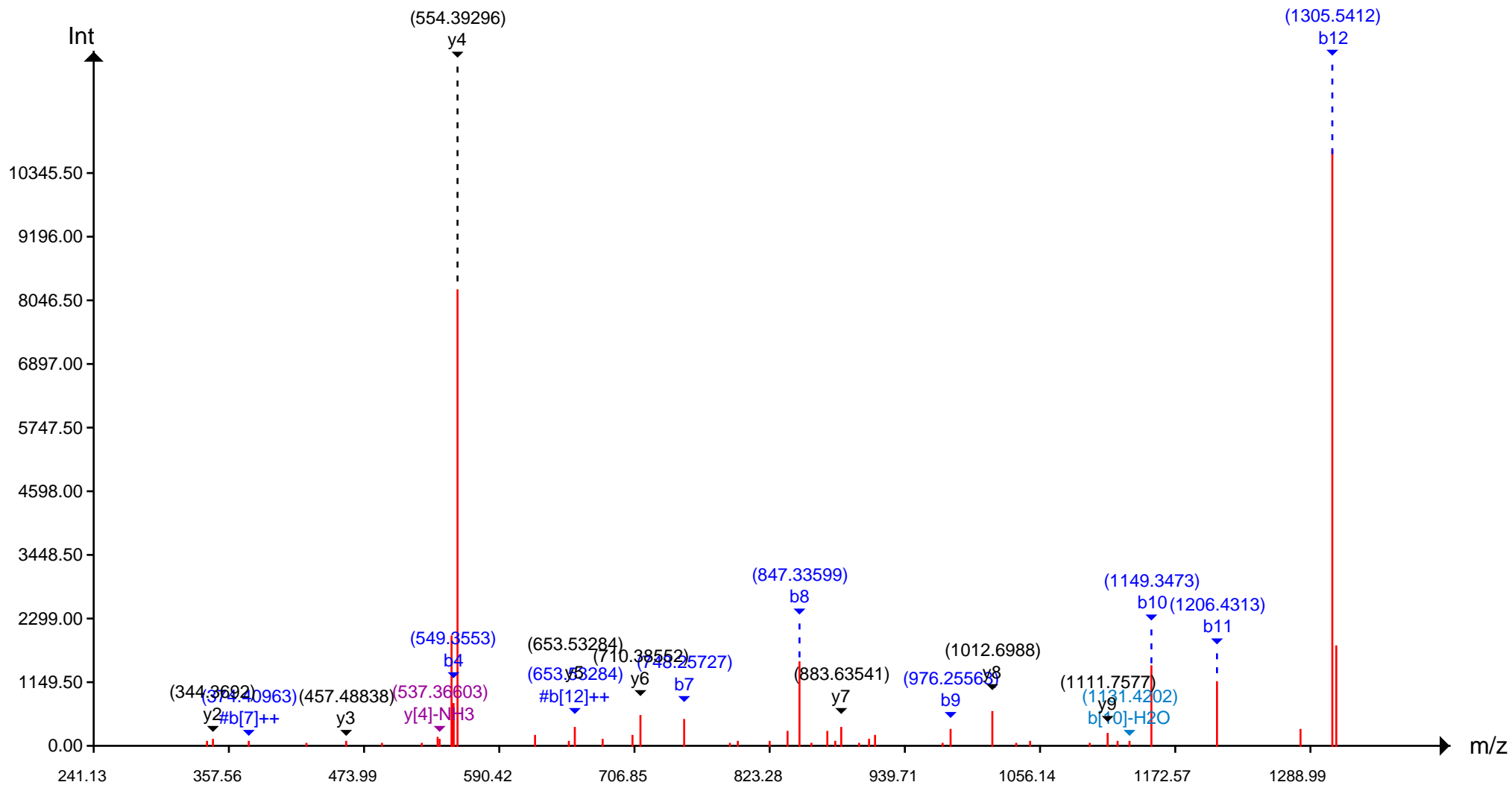
Confidence level 99.00%

Precursor m/z 753.41527

Identity Threshold 47.61

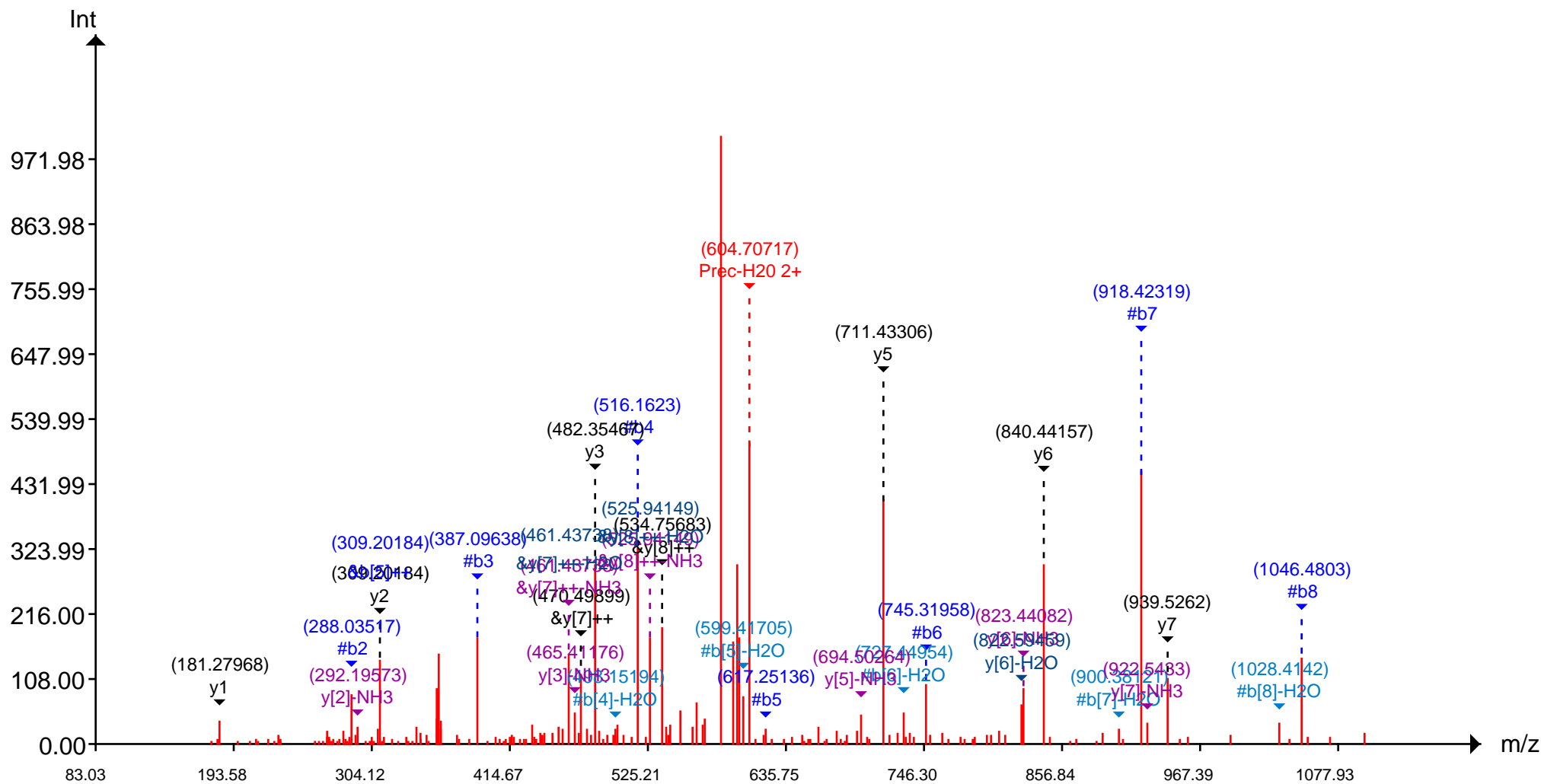
Charge 2+

Mass error -0.003577



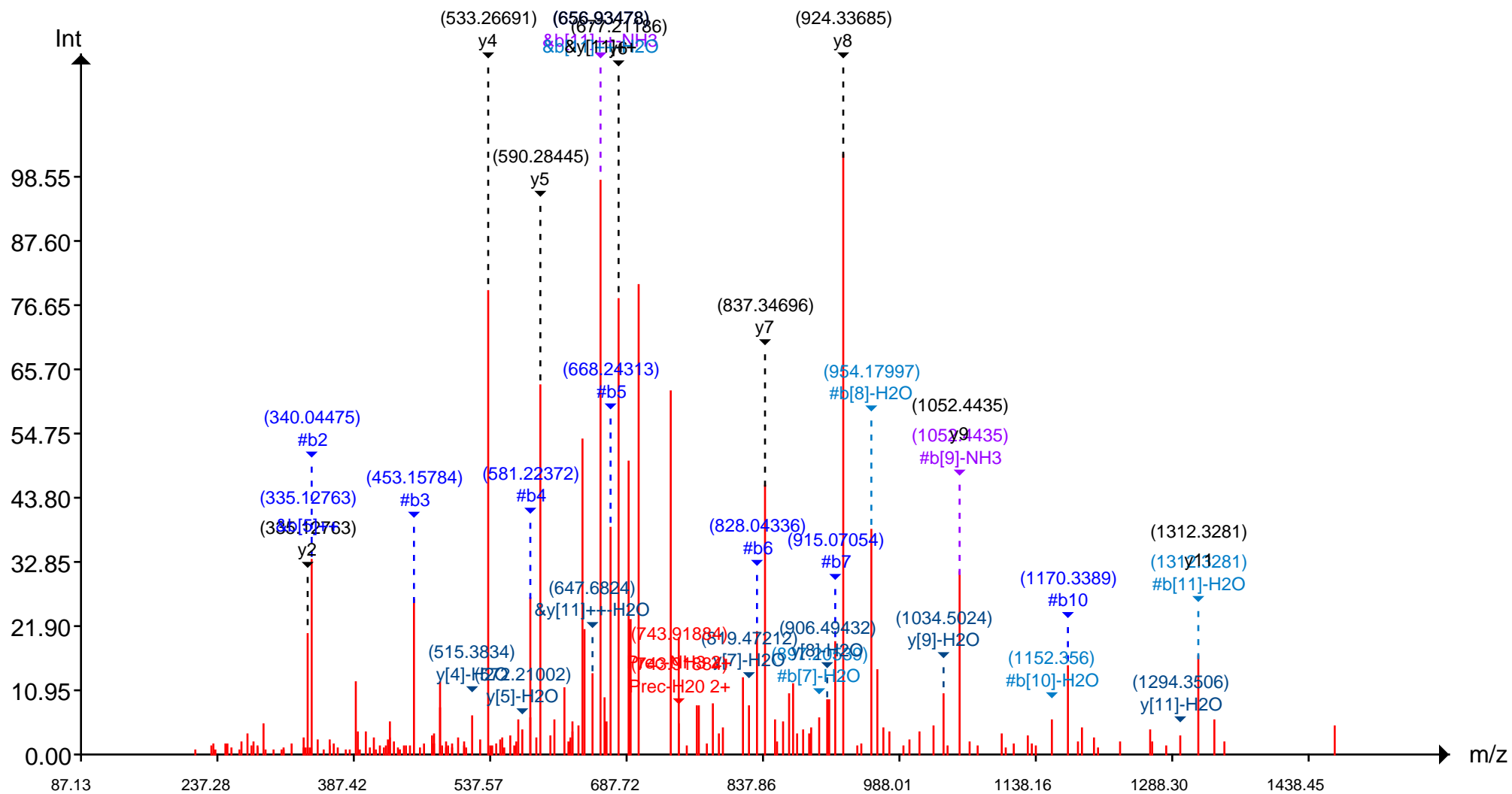
Info

Modified Sequence	AcD3-AVC<Cmm*>K<AcD3K*>AGAVEK<AcD3K*>GVPLYR<C13>-COOH	Ion score	53.15
Protein	P17182	Confidence level	99.00%
Precursor m/z	930.02882	Identity Threshold	45.79
Charge	2+	Mass error	2.57E-4



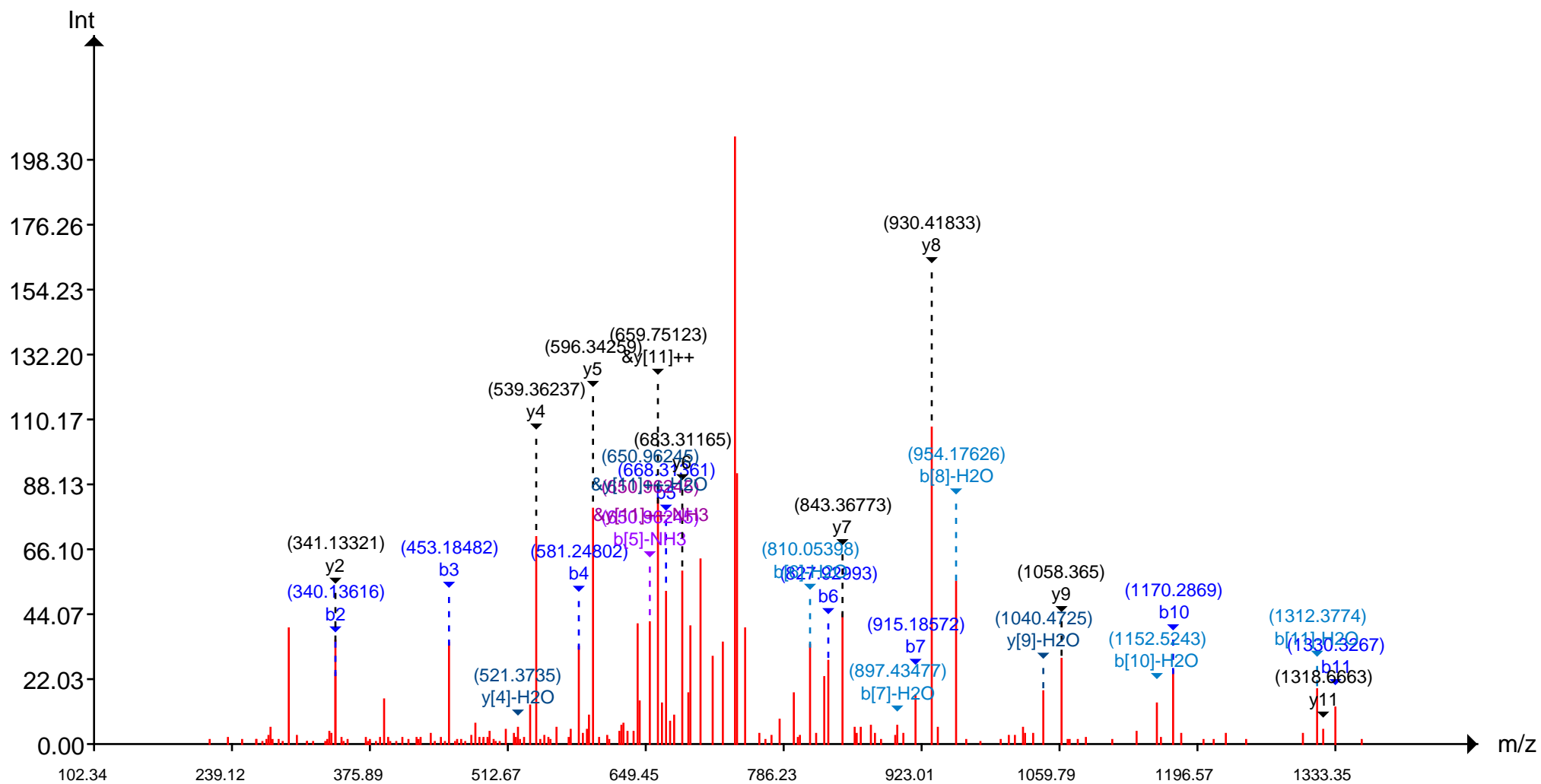
Info

Modified Sequence	AcD3-IEVETQK<AcD3K*>QR<C13>-COOH	Ion score	47.24
Protein	Q9WVA2	Confidence level	99.00%
Precursor m/z	613.8512	Identity Threshold	46.73
Charge	2+	Mass error	-2.01E-4



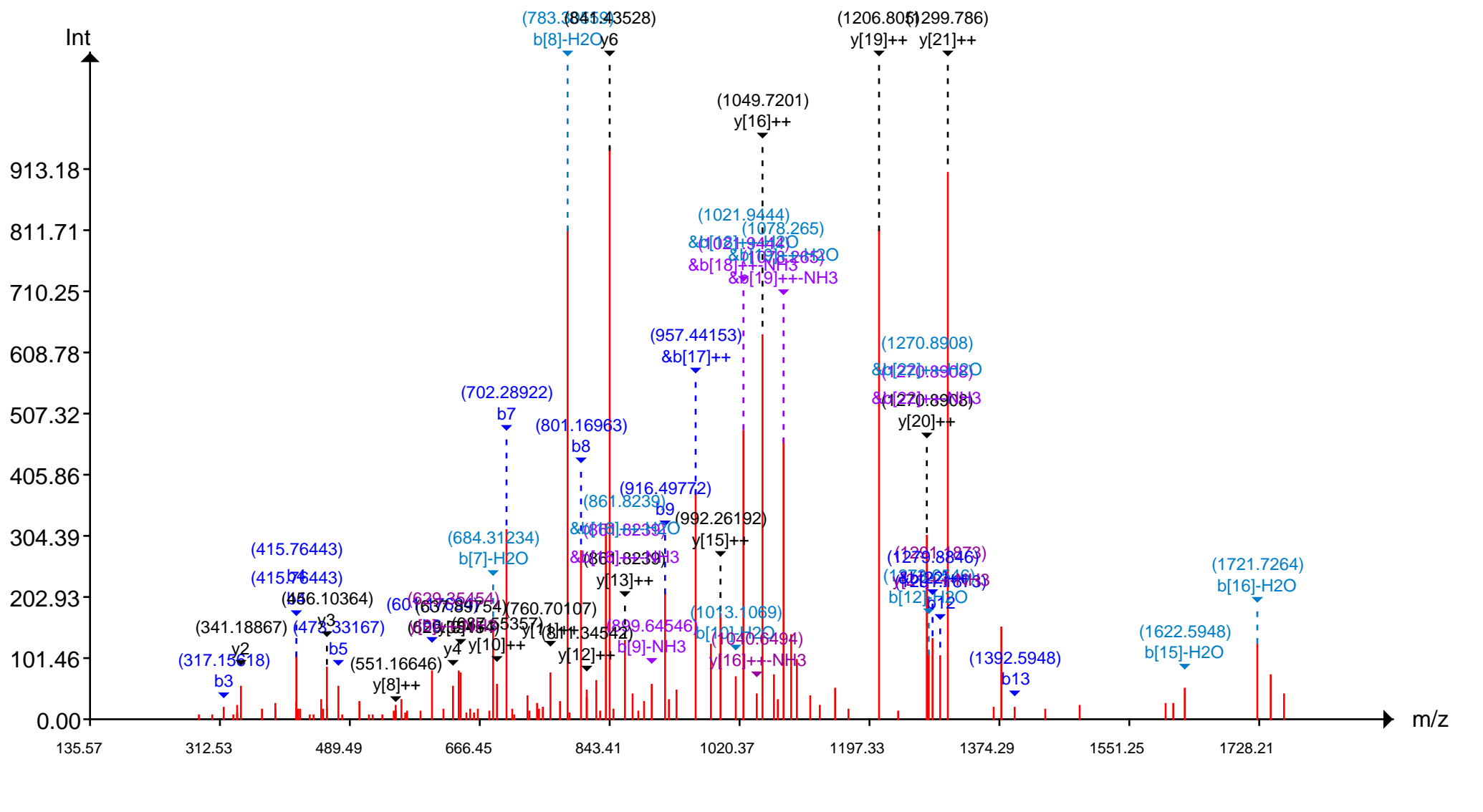
Info

Modified Sequence	AcD3-M<Mox*>M<Mox*>LQSC<Cmm*>SGPTC<Cmm*>R-COOH	Ion score	46.78
Protein	Q8BFR4	Confidence level	99.00%
Precursor m/z	752.80852	Identity Threshold	38.12
Charge	2+	Mass error	7.12E-4

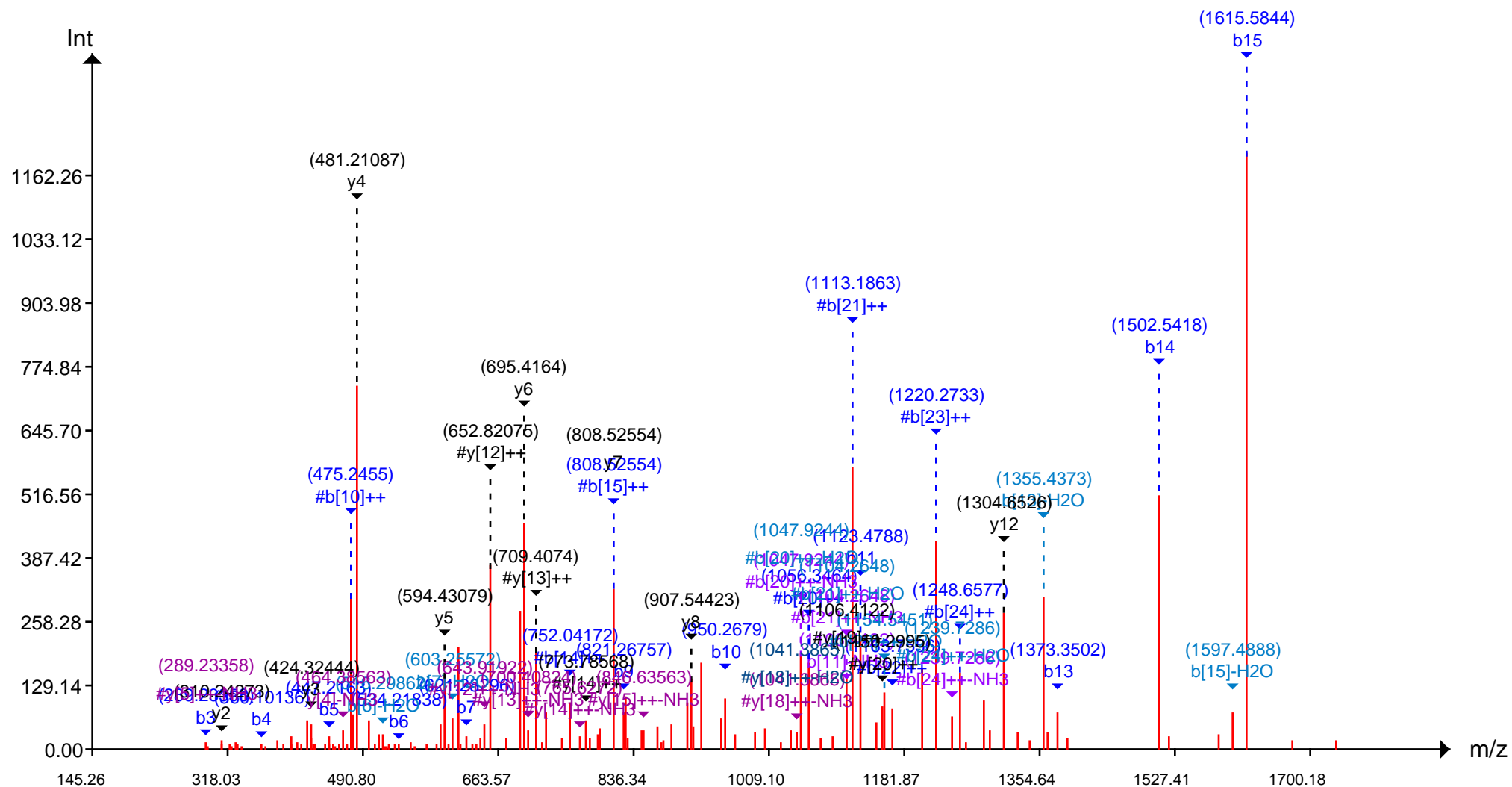


Info

Modified Sequence	AcD3-M<Mox*>M<Mox*>LQSC<Cmm*>SGPTC<Cmm*>R<C13>-COOH	Ion score	56.25
Protein	Q8BFR4	Confidence level	99.00%
Precursor m/z	755.8189	Identity Threshold	40.50
Charge	2+	Mass error	0.00133

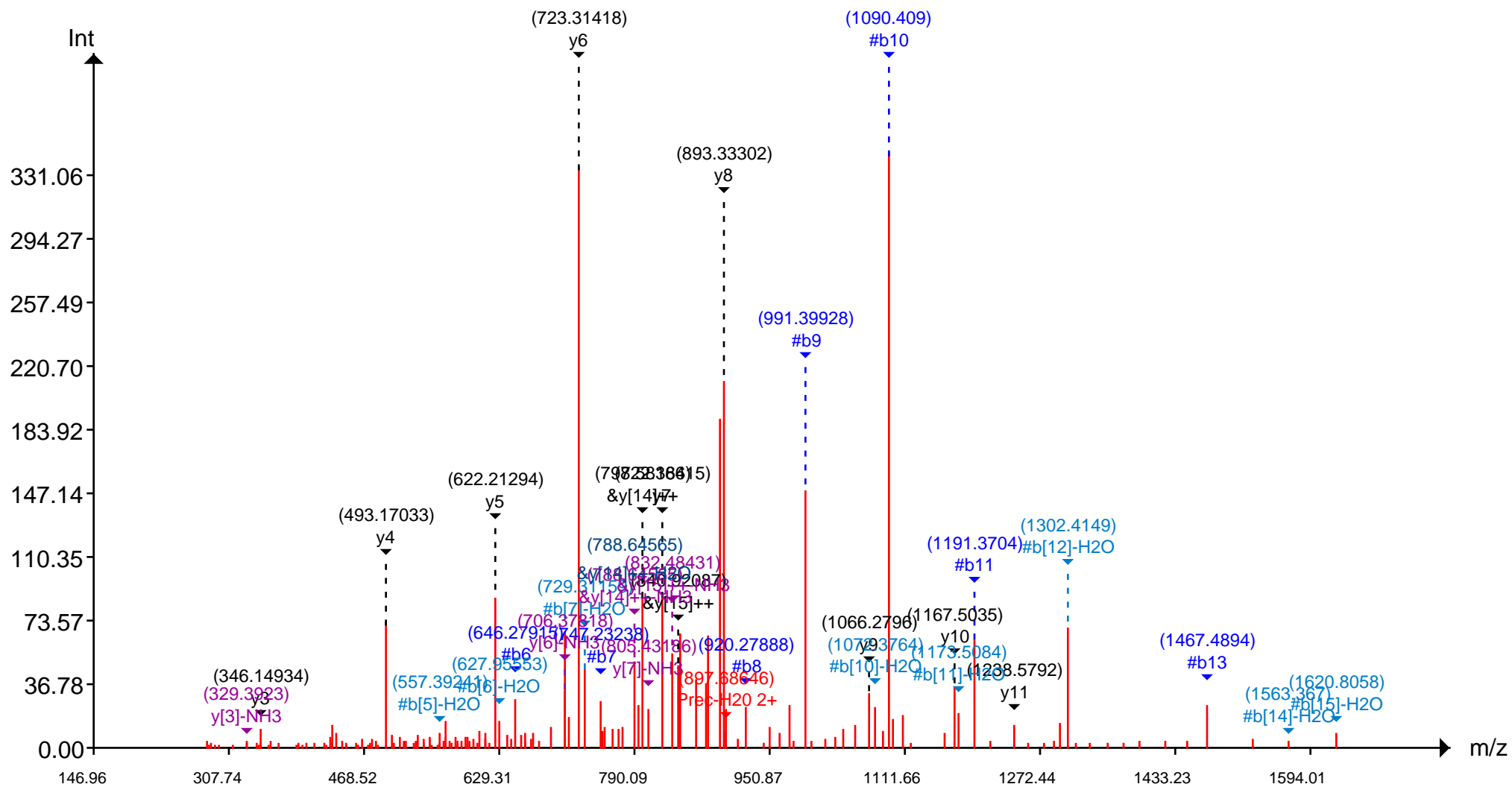


Info			
Modified Sequ...	AcD3-VGDVGQTVDDPYTTFFVK<AcD3K*>M<Mox*>LPDK<AcD3K*>DC<Cmm*>R<C13>-COOH	Ion score	54.81
Pro...	P18760	Confidence level	99.00%
Precursor ...	1005.159	Identity Threshold	46.74
Ch...	3+	Mass error	0.00488



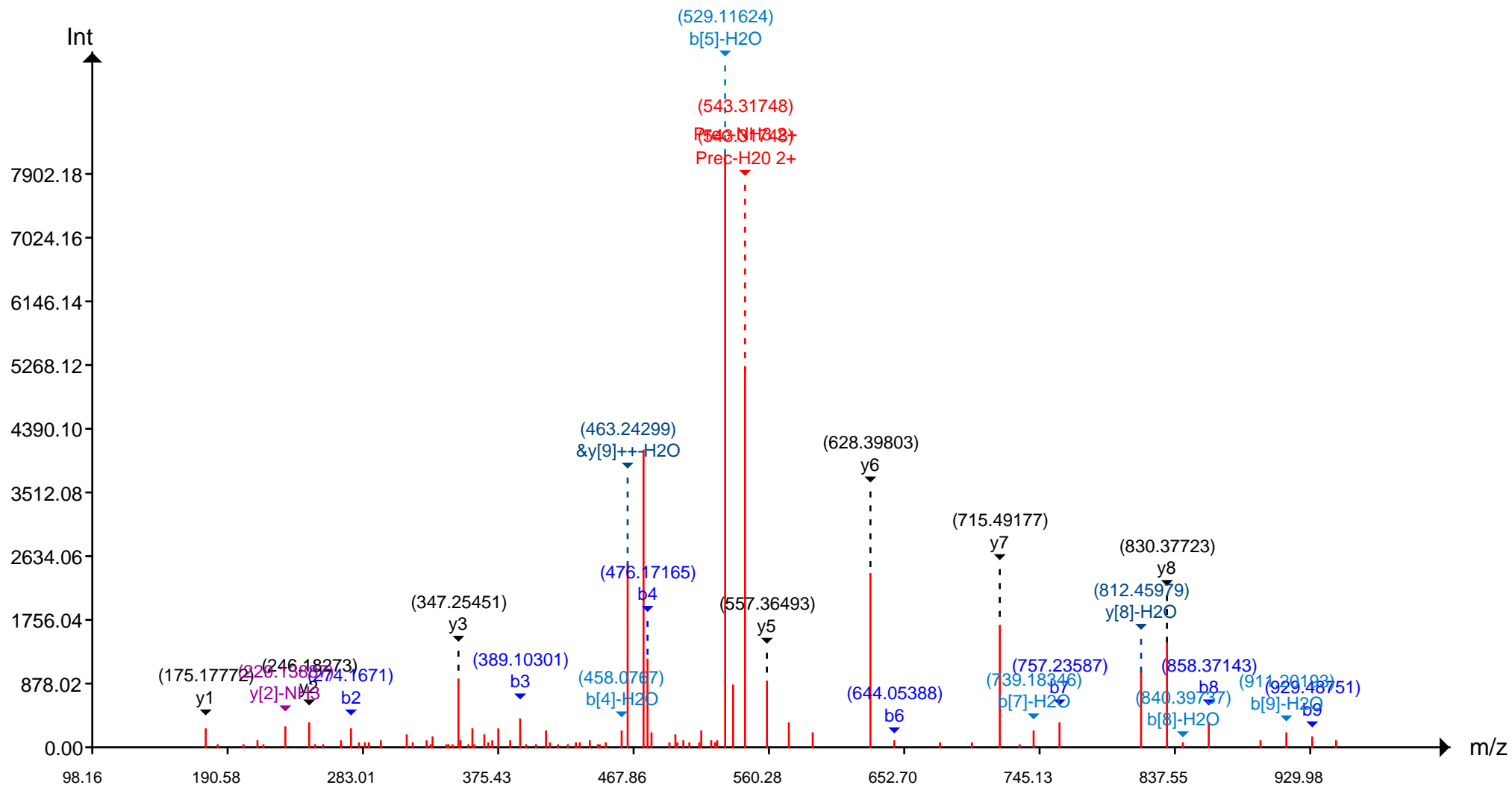
Info

Modified Sequence	AcD3-ATAASSSSLEK<AcD3K*>SYELPDGQVITIGNER<C13>-COOH	Ion score	47.89
Protein	P62737, P60710, P68033, P63260, P63268, P68134	Confidence level	99.00%
Precursor m/z	973.83213	Identity Threshold	47.63
Charge	3+	Mass error	0.002272



Info

Modified Sequence	AcD3-SSPEATK<AcD3K*>AVTEM<Mox*>NGR-COOH	Ion score	68.54
Protein	P29341	Confidence level	99.00%
Precursor m/z	906.93239	Identity Threshold	46.45
Charge	2+	Mass error	0.002434



Info

Modified Sequence AcD3-DIDSAPITAR-COOH

Ion score 58.8

Protein P52480

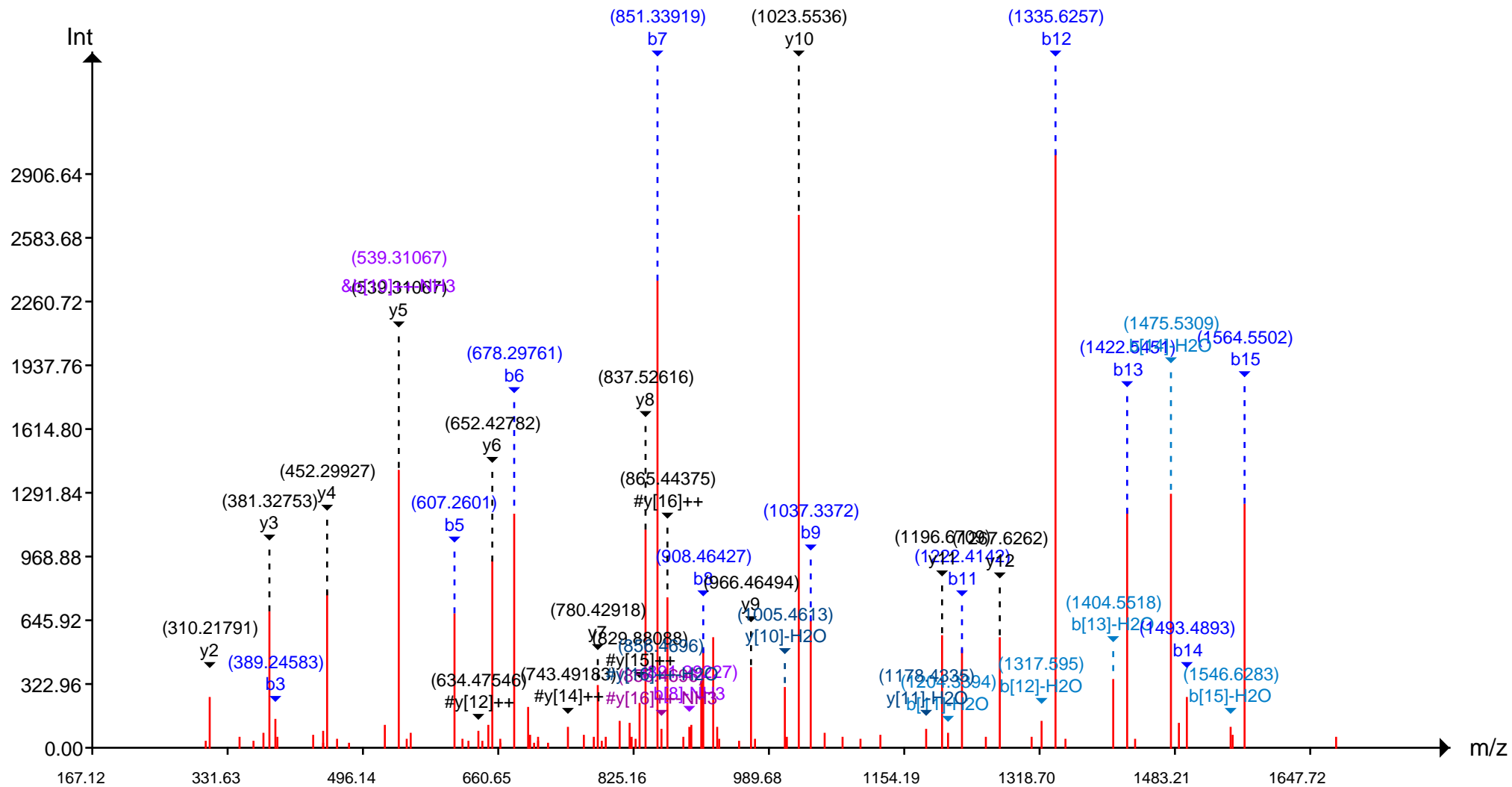
Confidence level 99.00%

Precursor m/z 552.29251

Identity Threshold 48.07

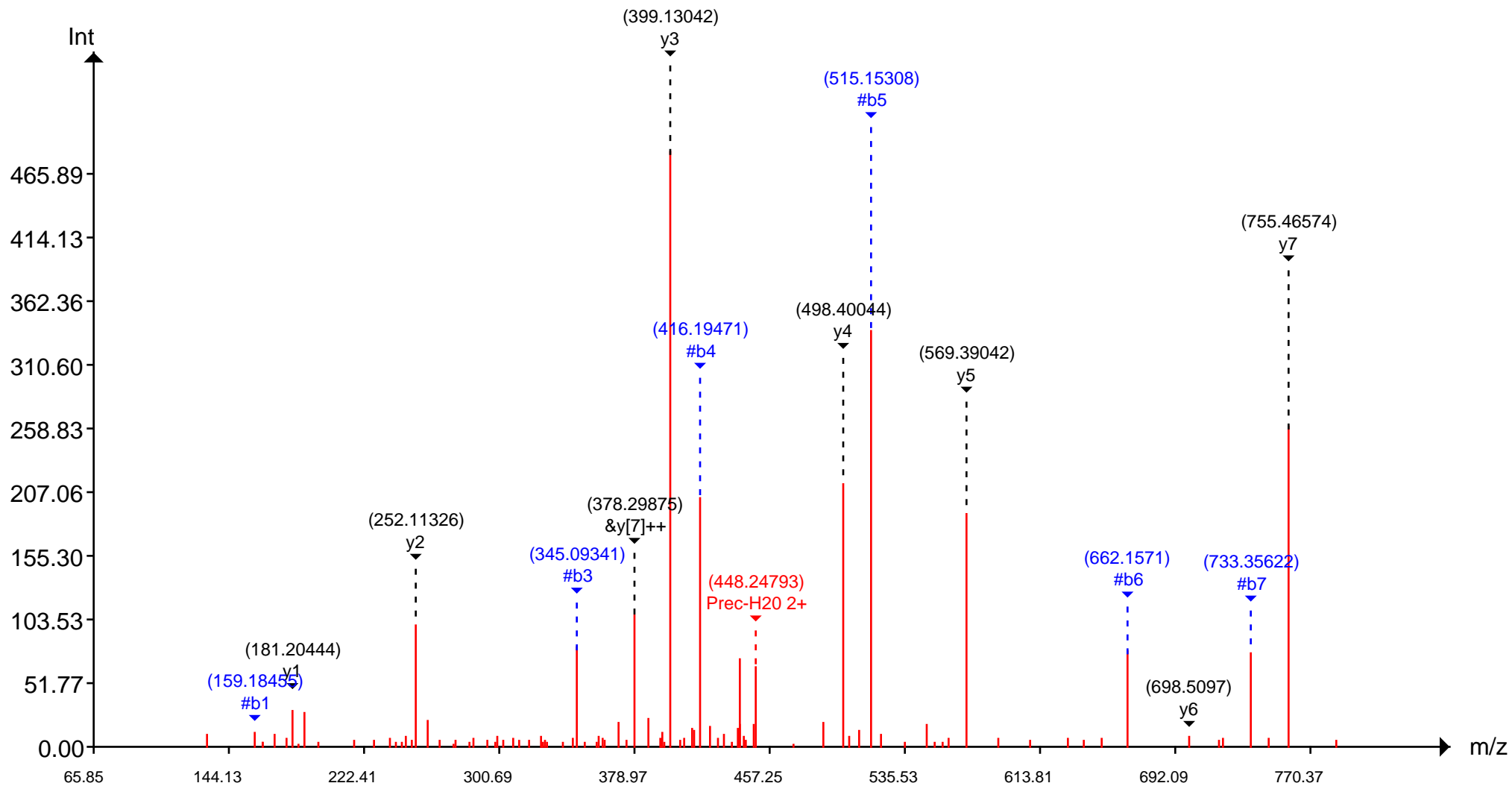
Charge 2+

Mass error 7.05E-4



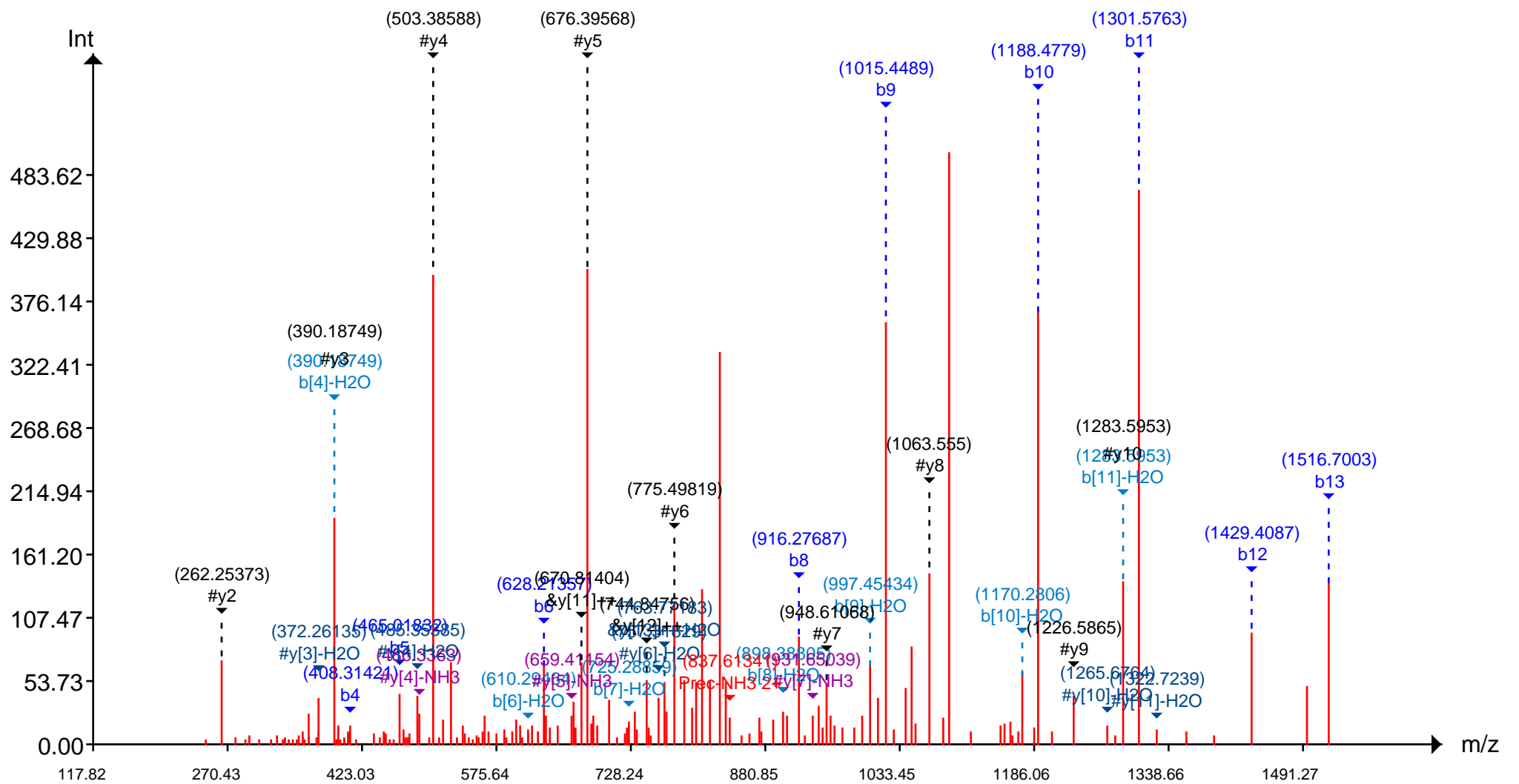
Info

Modified Sequence	AcD3-VAK<AcD3K*>FAAK<AcD3K*>GEGQLSAAER<C13>-COOH	Ion score	104.29
Protein	P58252	Confidence level	99.00%
Precursor m/z	937.52575	Identity Threshold	46.44
Charge	2+	Mass error	0.001868

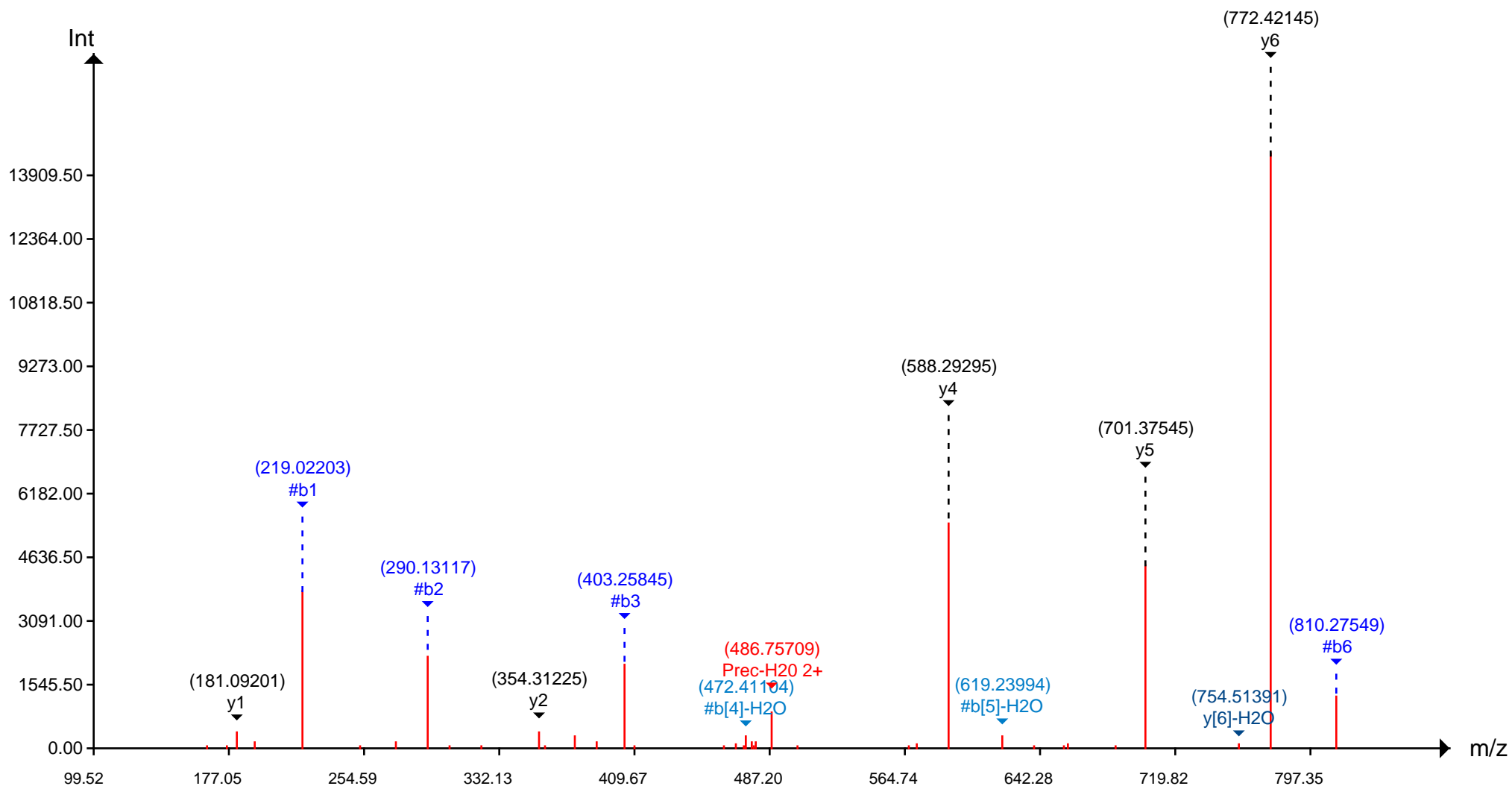


Info

Modified Sequence	AcD3-IGEA VFAR<C13>-COOH	Ion score	68.01
Protein	Q9DCD0	Confidence level	99.00%
Precursor m/z	457.26744	Identity Threshold	46.71
Charge	2+	Mass error	-4.2E-5



Info			
Modified Sequence	AcD3-TGFGGYDK<AcD3K*>VK<AcD3K*>LQSR-COOH	Ion score	62.67
Protein	Q62465	Confidence level	99.00%
Precursor m/z	845.96089	Identity Threshold	47.58
Charge	2+	Mass error	0.003587



Info

Modified Sequence AcD3-K<AcD3K*>AISFK<AcD3K*>R<C13>-COOH

Ion score 46.44

Protein Q62426

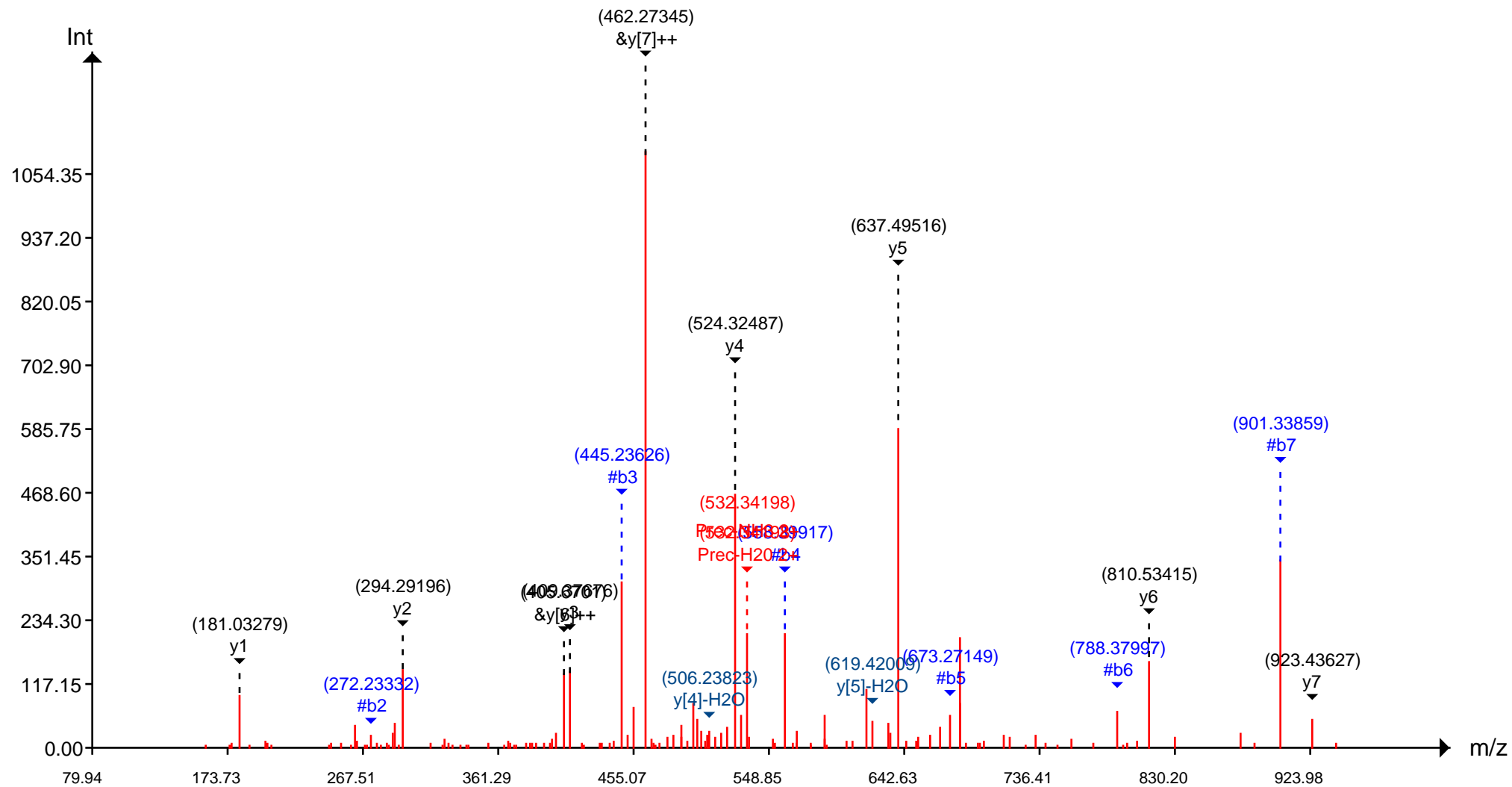
Confidence level 99.00%

Precursor m/z 495.82282

Identity Threshold 41.59

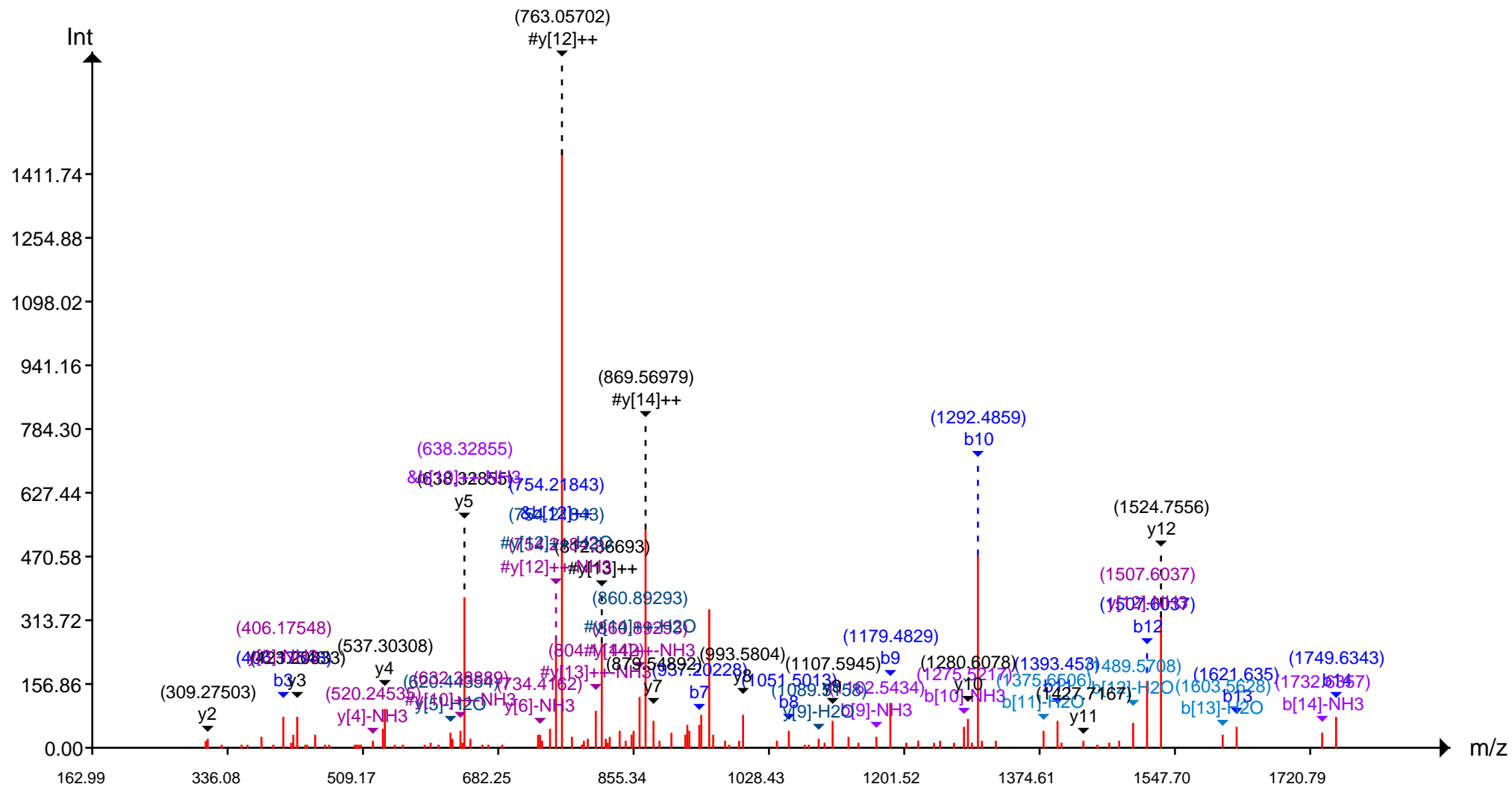
Charge 2+

Mass error -4.43E-4



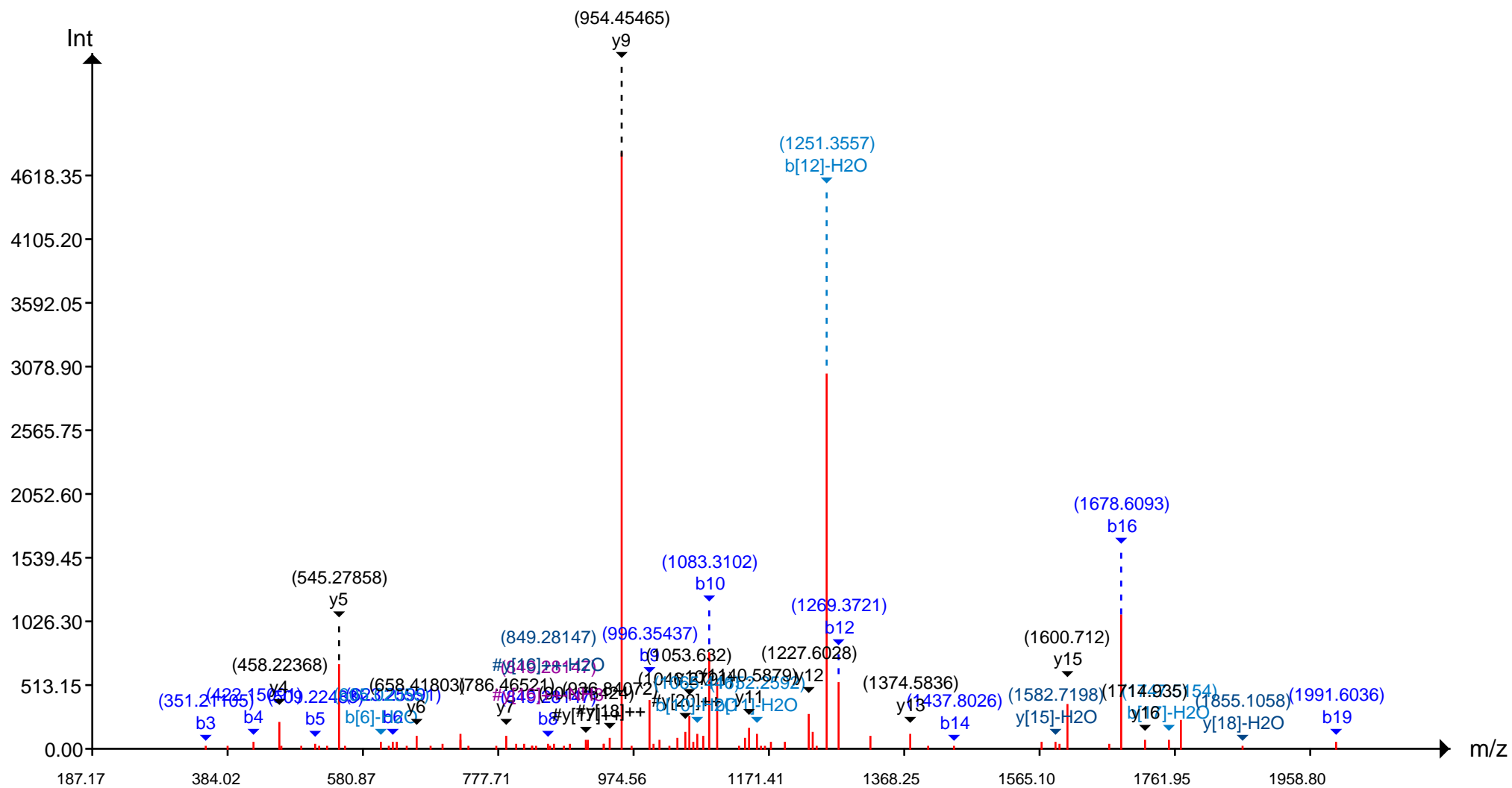
Info

Modified Sequence	AcD3-ILK<AcD3K*>IDDIR<C13>-COOH	Ion score	59.18
Protein	P80316	Confidence level	99.00%
Precursor m/z	541.3452	Identity Threshold	40.85
Charge	2+	Mass error	1.58E-4



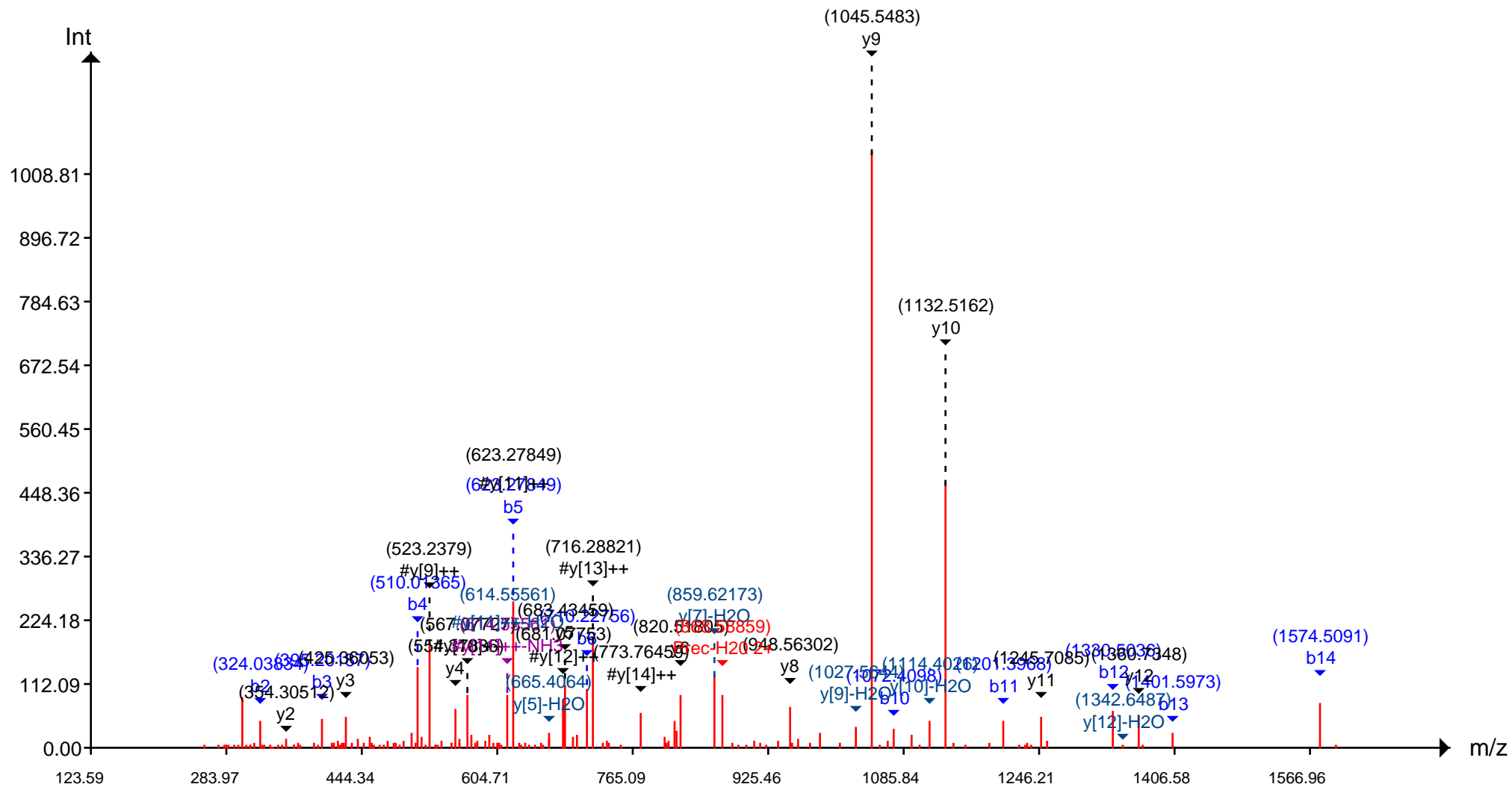
Info

Modified Sequence	AcD3-FNVPM<Mox*>K<AcD3K*>NNQITNNQR<C13>-COOH	Ion score	77.23
Protein	P09411, P09041	Confidence level	99.00%
Precursor m/z	965.49537	Identity Threshold	48.02
Charge	2+	Mass error	0.002113



Info

Modified Sequence	AcD3-SAFASNPEFSSVPAQISPEGR-COOH	Ion score	91.71
Protein	P26039	Confidence level	99.00%
Precursor m/z	1112.0437	Identity Threshold	47.20
Charge	2+	Mass error	0.004565



Info

Modified Sequence AcD3-YDADLSPAGHEEAK<AcD3K*>R<C13>-COOH

Ion score 92.72

Protein Q9DBJ1

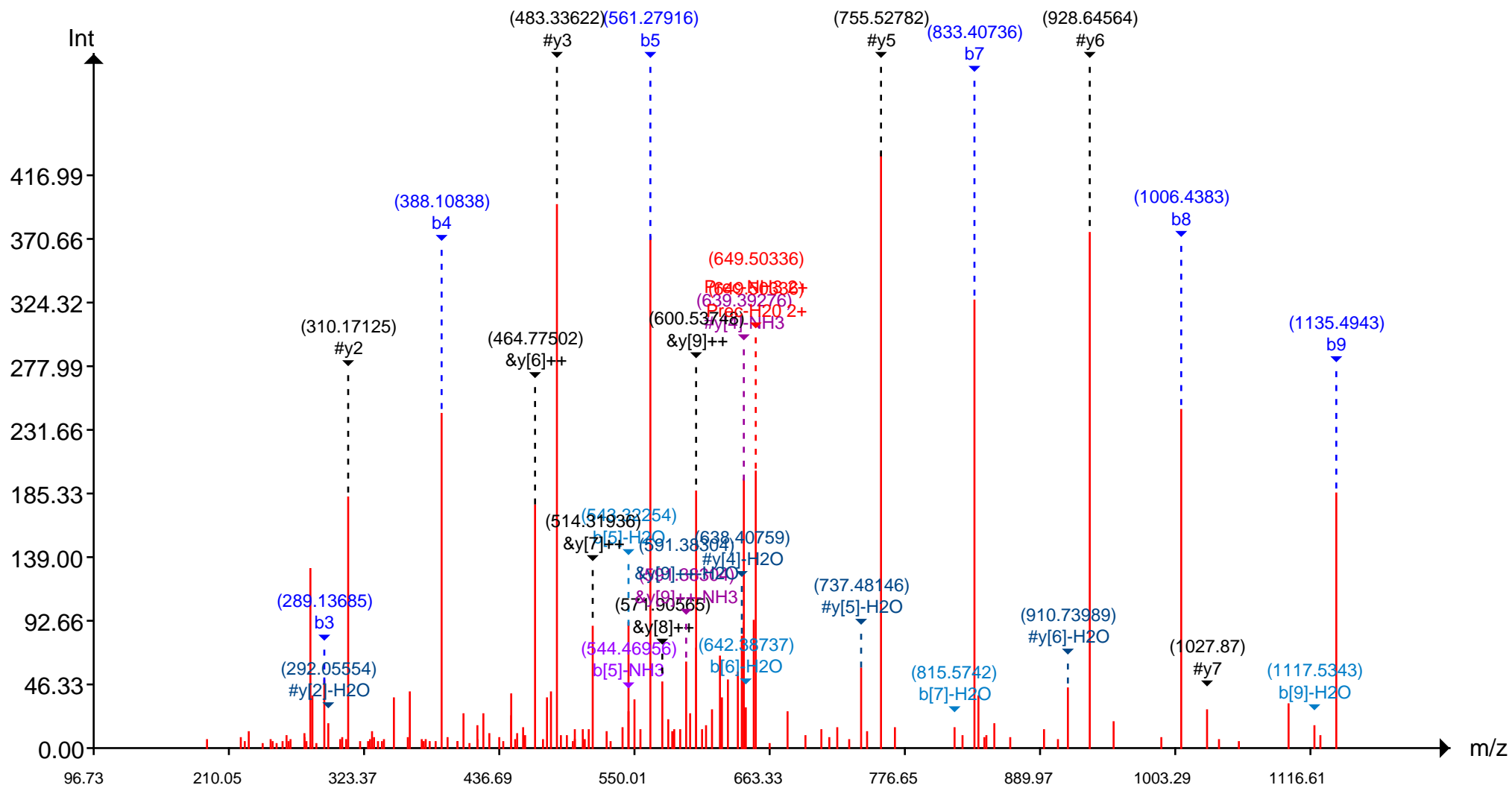
Confidence level 99.00%

Precursor m/z 877.93231

Identity Threshold 47.27

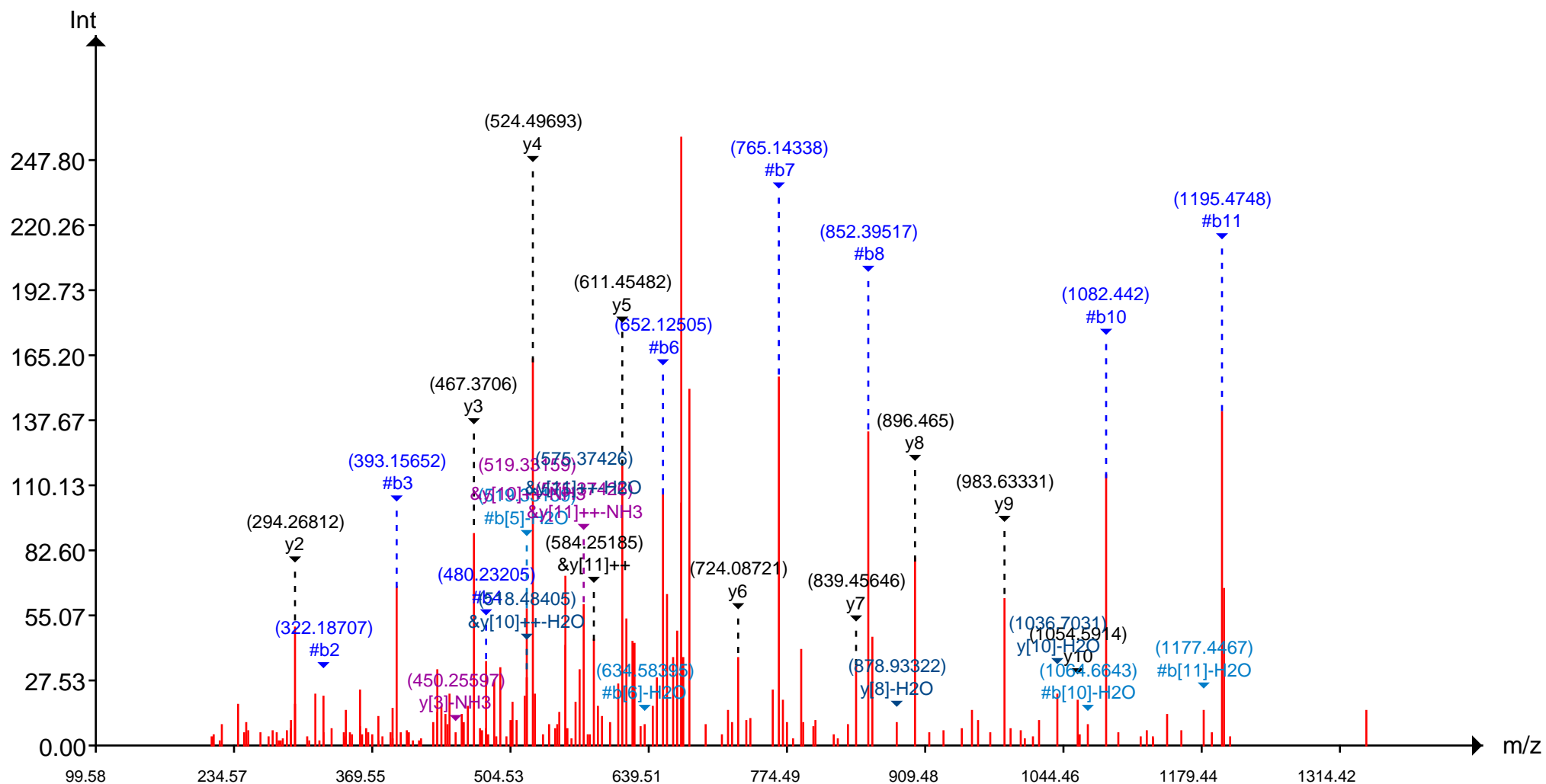
Charge 2+

Mass error 0.001557



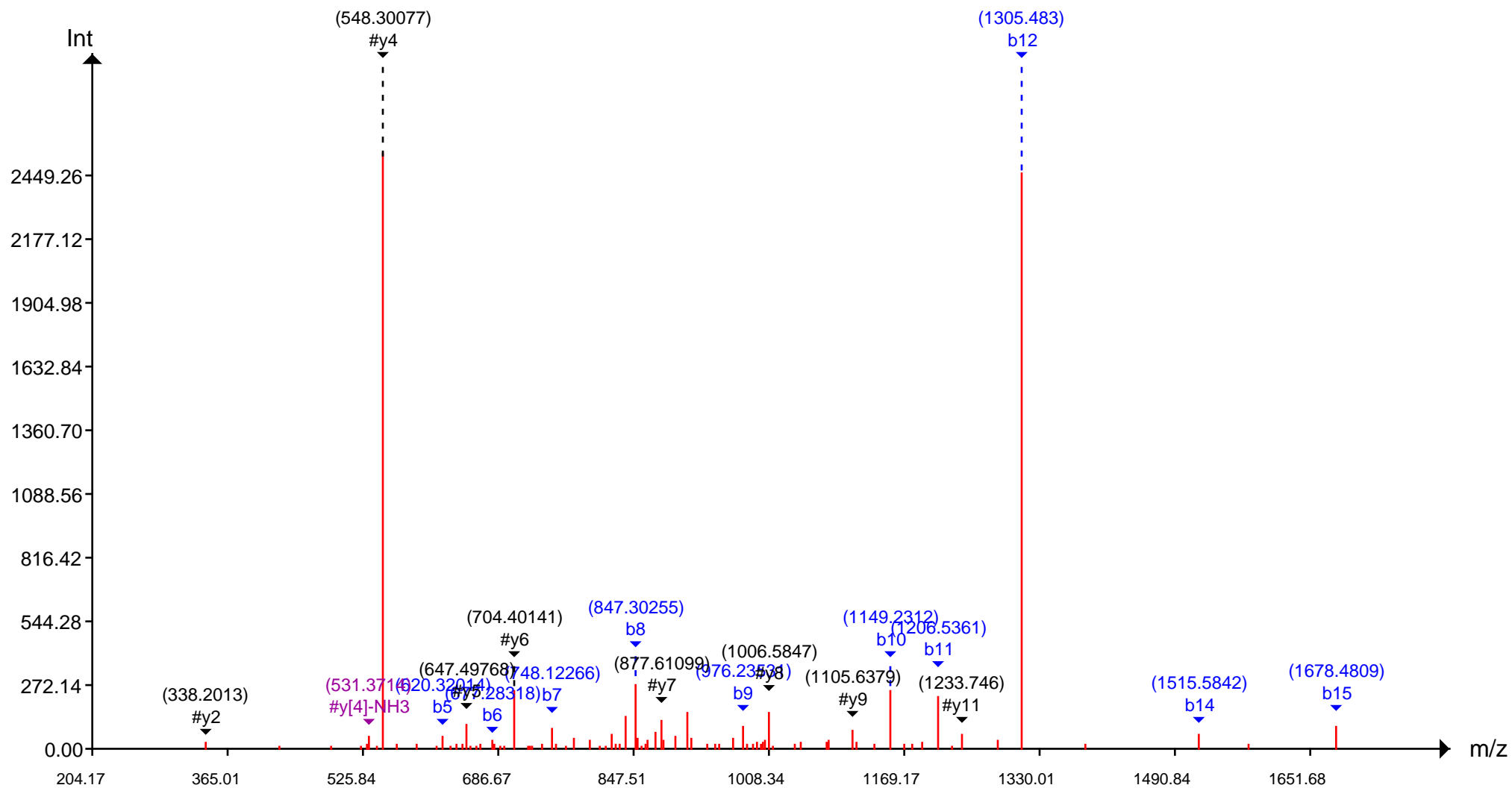
Info

Modified Sequence	AcD3-AGDVK<AcD3K*>VK<AcD3K*>K<AcD3K*>ER<C13>-COOH	Ion score	45.44
Protein	Q8K3X4	Confidence level	99.00%
Precursor m/z	658.40791	Identity Threshold	42.37
Charge	2+	Mass error	0.002043



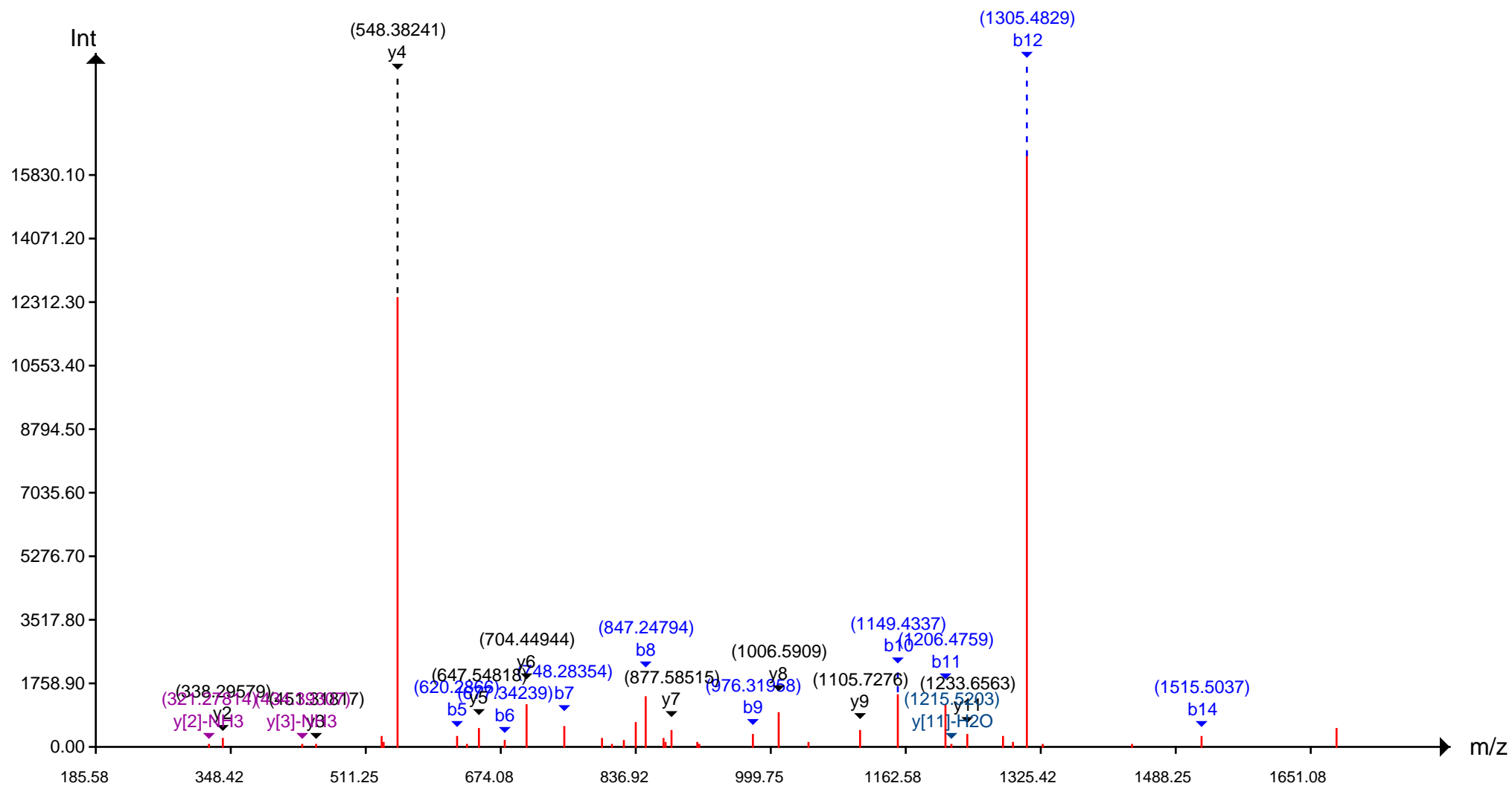
Info

Modified Sequence	AcD3-YIASGDISGK<AcD3K*>LR<C13>-COOH	Ion score	54.05
Protein	O88342	Confidence level	99.00%
Precursor m/z	688.39397	Identity Threshold	46.74
Charge	2+	Mass error	0.001278



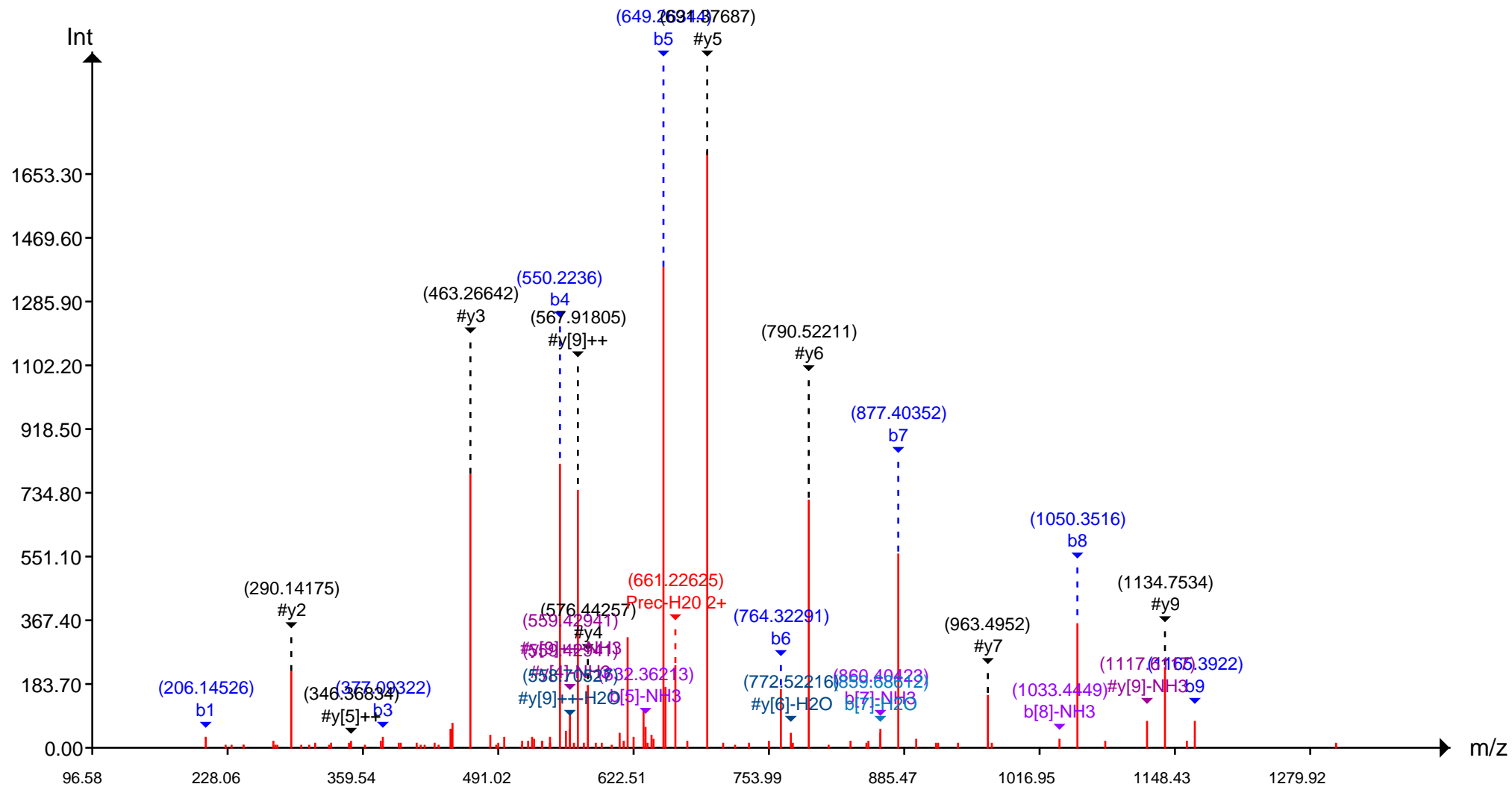
Info

Modified Sequence	AcD3-AVC<Cmm*>K<AcD3K*>AGAVEK<AcD3K*>GVPLYR-COOH	Ion score	70.65
Protein	P17182	Confidence level	99.00%
Precursor m/z	927.01976	Identity Threshold	46.50
Charge	2+	Mass error	0.002278



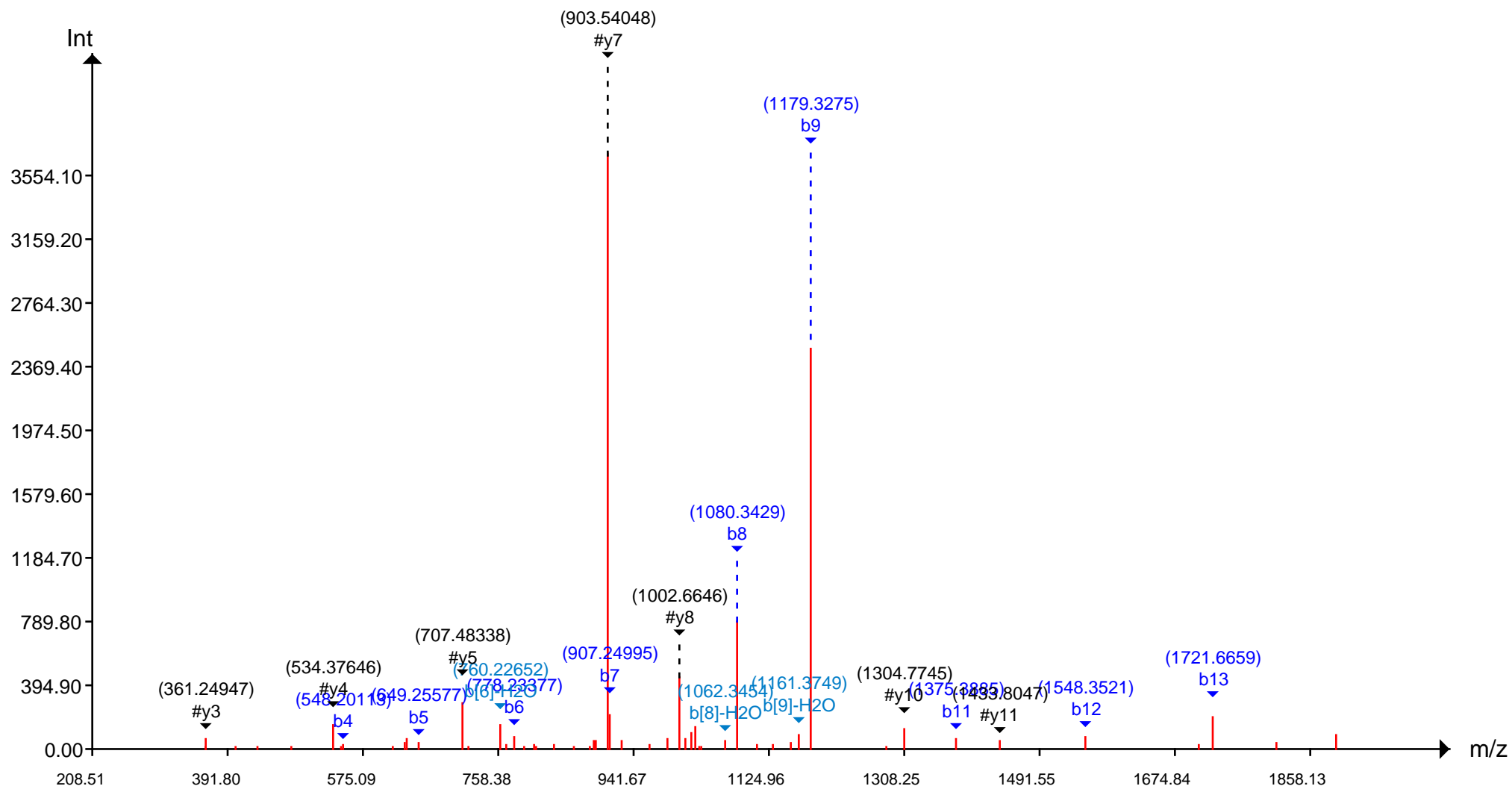
Info

Modified Sequence	AcD3-AVC<Cmm*>K<AcD3K*>AGAVEK<AcD3K*>GVPLYR-COOH	Ion score	71.88
Protein	P17182	Confidence level	99.00%
Precursor m/z	927.01938	Identity Threshold	46.59
Charge	2+	Mass error	0.001518



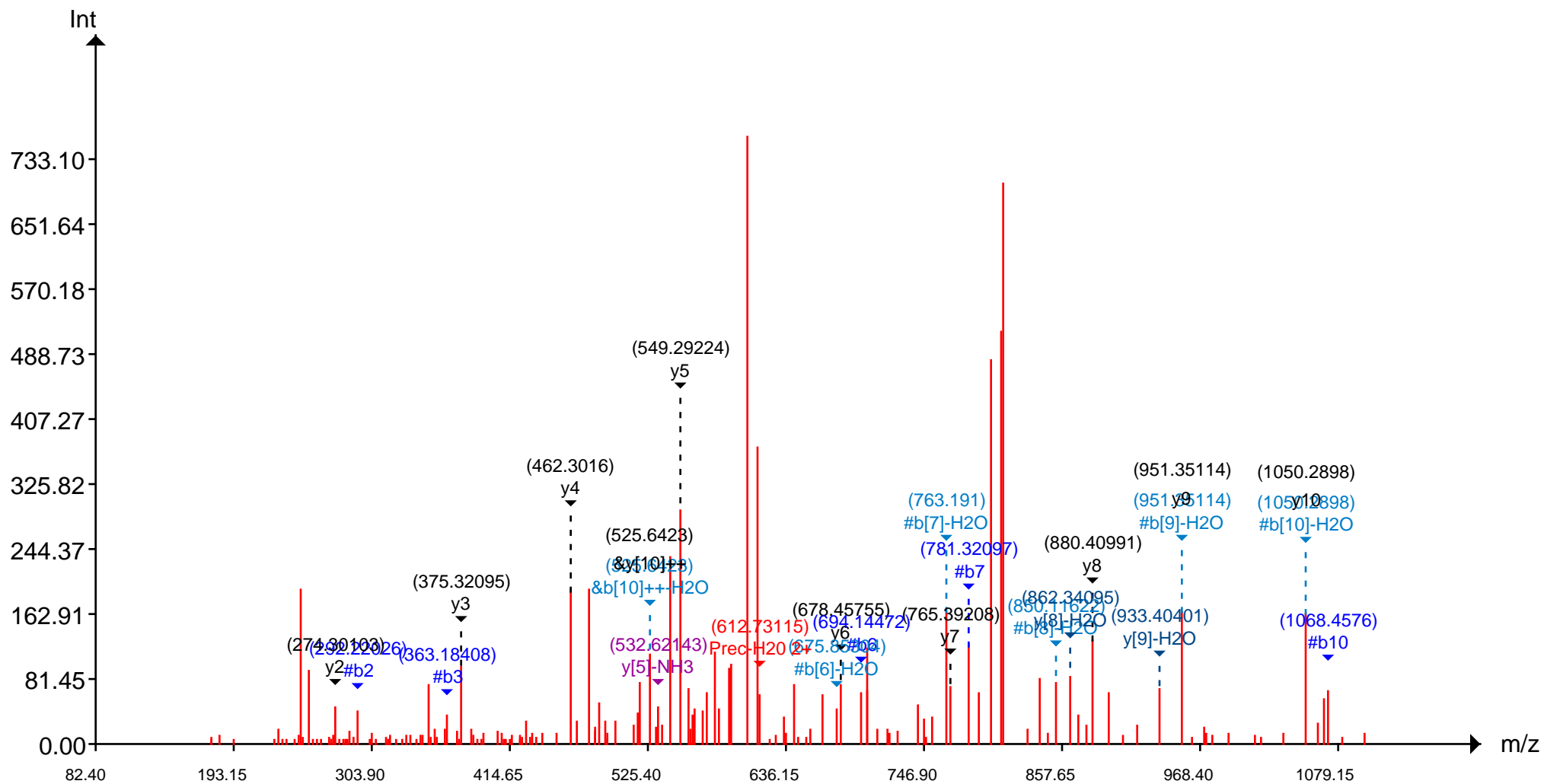
Info

Modified Sequence	AcD3-C<Cmm*>GNK<AcD3K*>VDIK<AcD3K*>DR-COOH	Ion score	77.8
Protein	P62827	Confidence level	99.00%
Precursor m/z	670.3533	Identity Threshold	48.05
Charge	2+	Mass error	8.55E-4



Info

Modified Sequence	AcD3-DEEETEEK<AcD3K*>VPVK<AcD3K*>K<AcD3K*>SVR-COOH	Ion score	64.88
Protein	Q61937	Confidence level	99.00%
Precursor m/z	1041.5544	Identity Threshold	47.86
Charge	2+	Mass error	0.002497



Info

Modified Sequence AcD3-FVADESSTVR-COOH

Protein Q8BZW8

Precursor m/z 621.80545

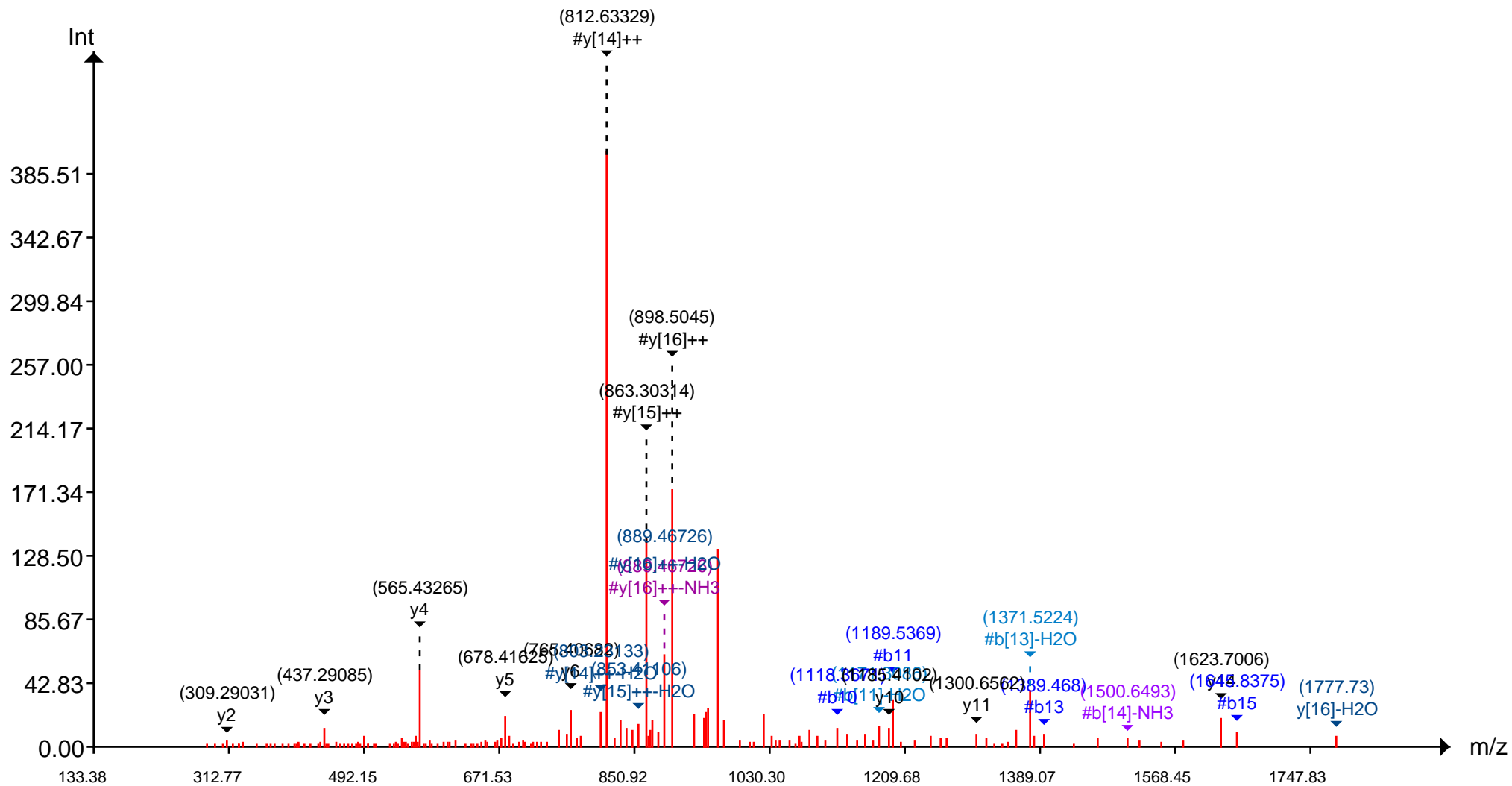
Charge 2+

Ion score 55.34

Confidence level 99.00%

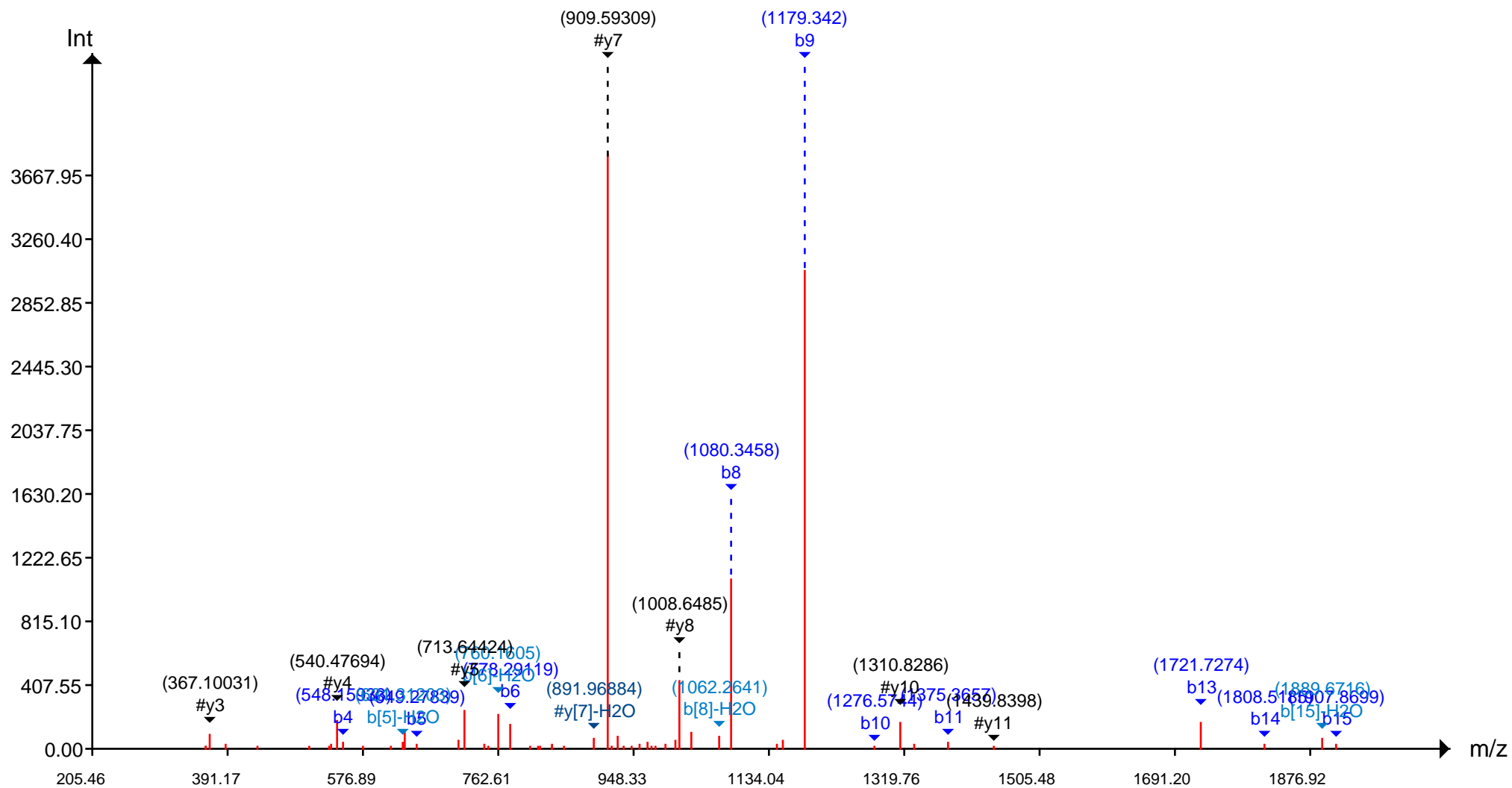
Identity Threshold 46.78

Mass error -3.77E-4



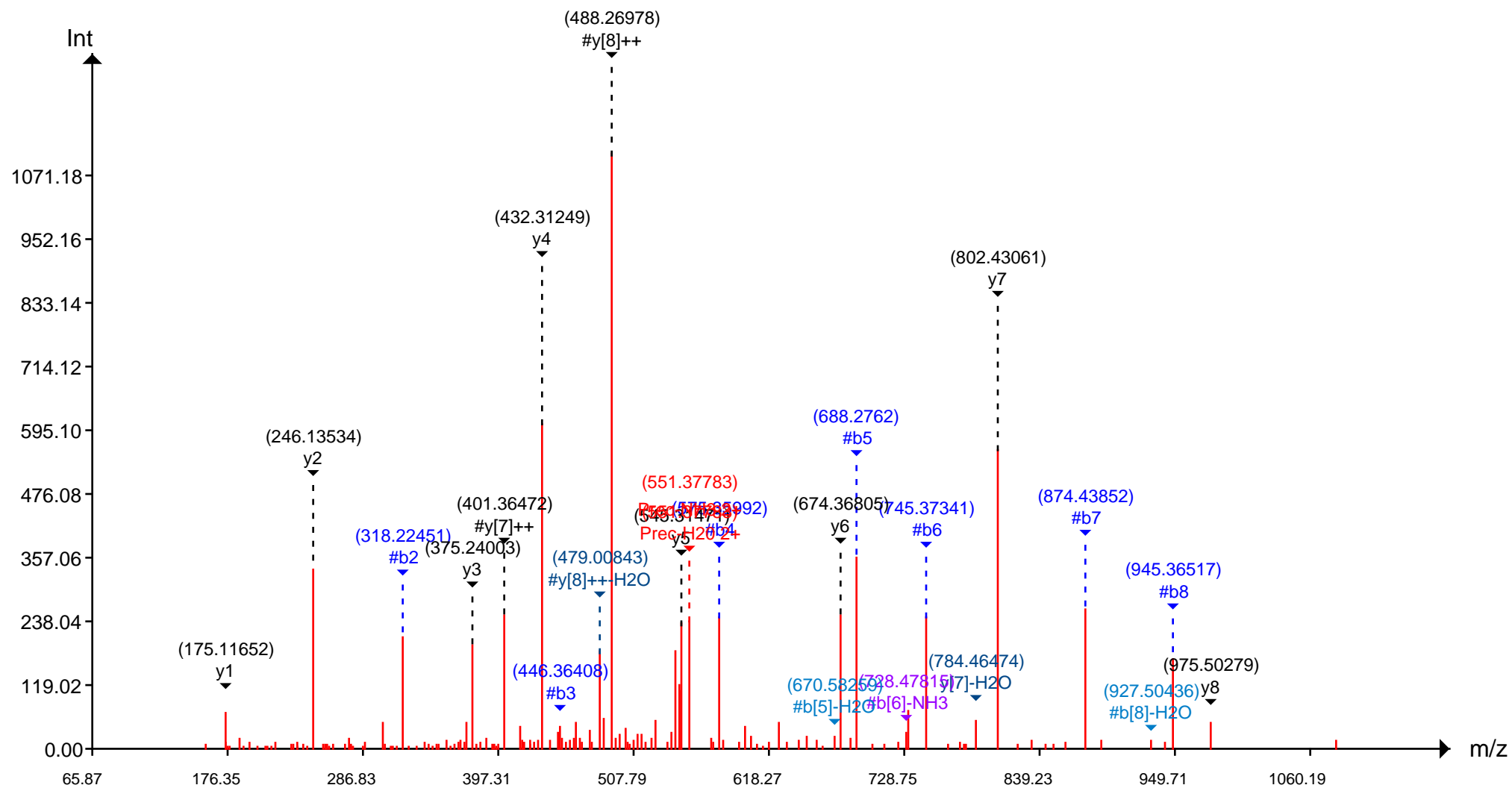
Info

Modified Sequence	AcD3-IATPPEDSDM<Mox*>ASLQQQR<C13>-COOH	Ion score	49.42
Protein	Q9DBG5	Confidence level	99.00%
Precursor m/z	977.4726	Identity Threshold	46.91
Charge	2+	Mass error	0.002227



Info

Modified Sequence	AcD3-DEEETEEK<AcD3K*>VPVK<AcD3K*>K<AcD3K*>SVR<C13>-COOH	Ion score	68.67
Protein	Q61937	Confidence level	99.00%
Precursor m/z	1044.5642	Identity Threshold	47.62
Charge	2+	Mass error	0.001955



Info

Modified Sequence AcD3-VK<AcD3K*>QELGEAR-COOH

Protein Q03958

Precursor m/z 560.31664

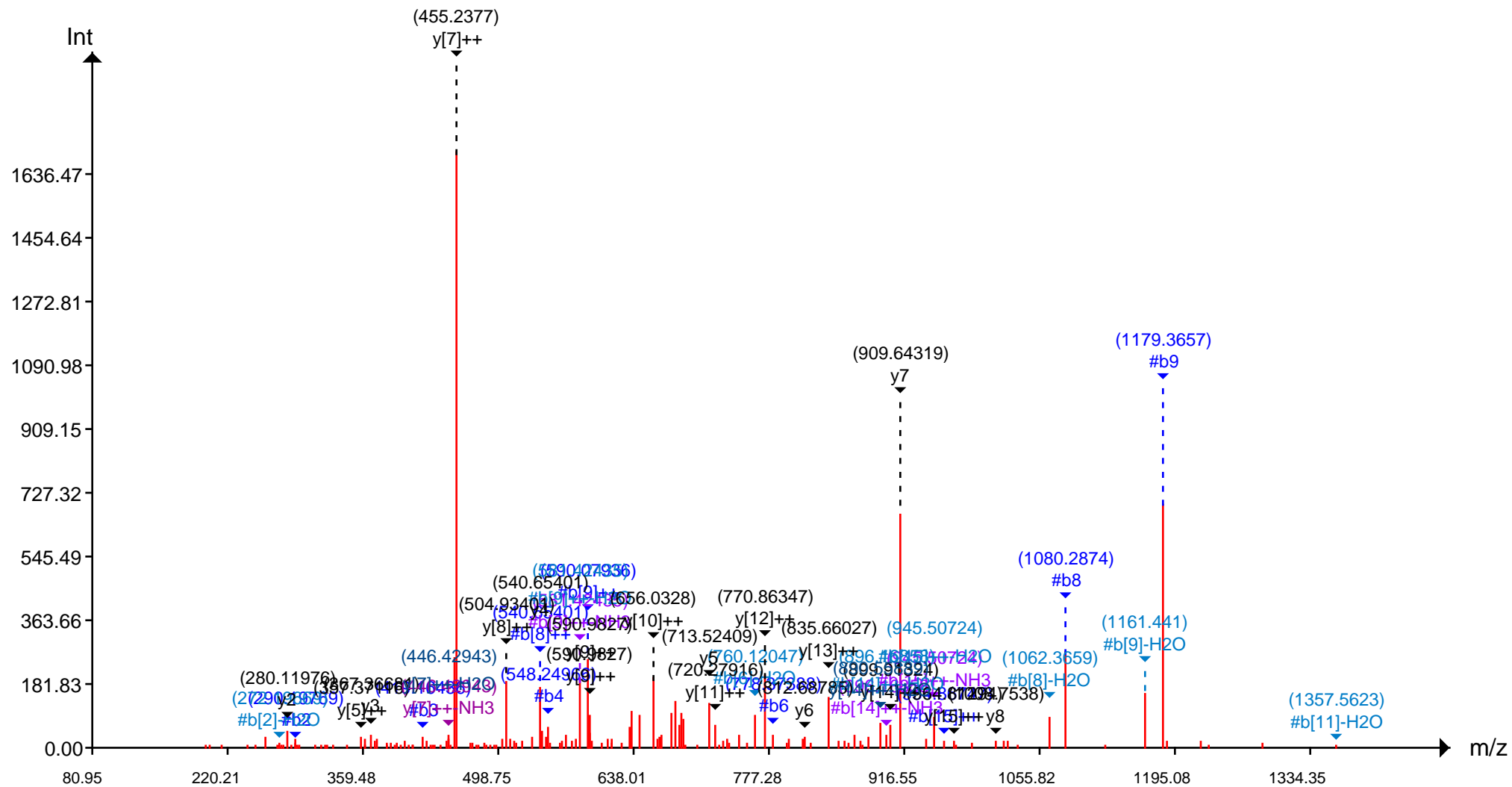
Charge 2+

Ion score 65.74

Confidence level 99.00%

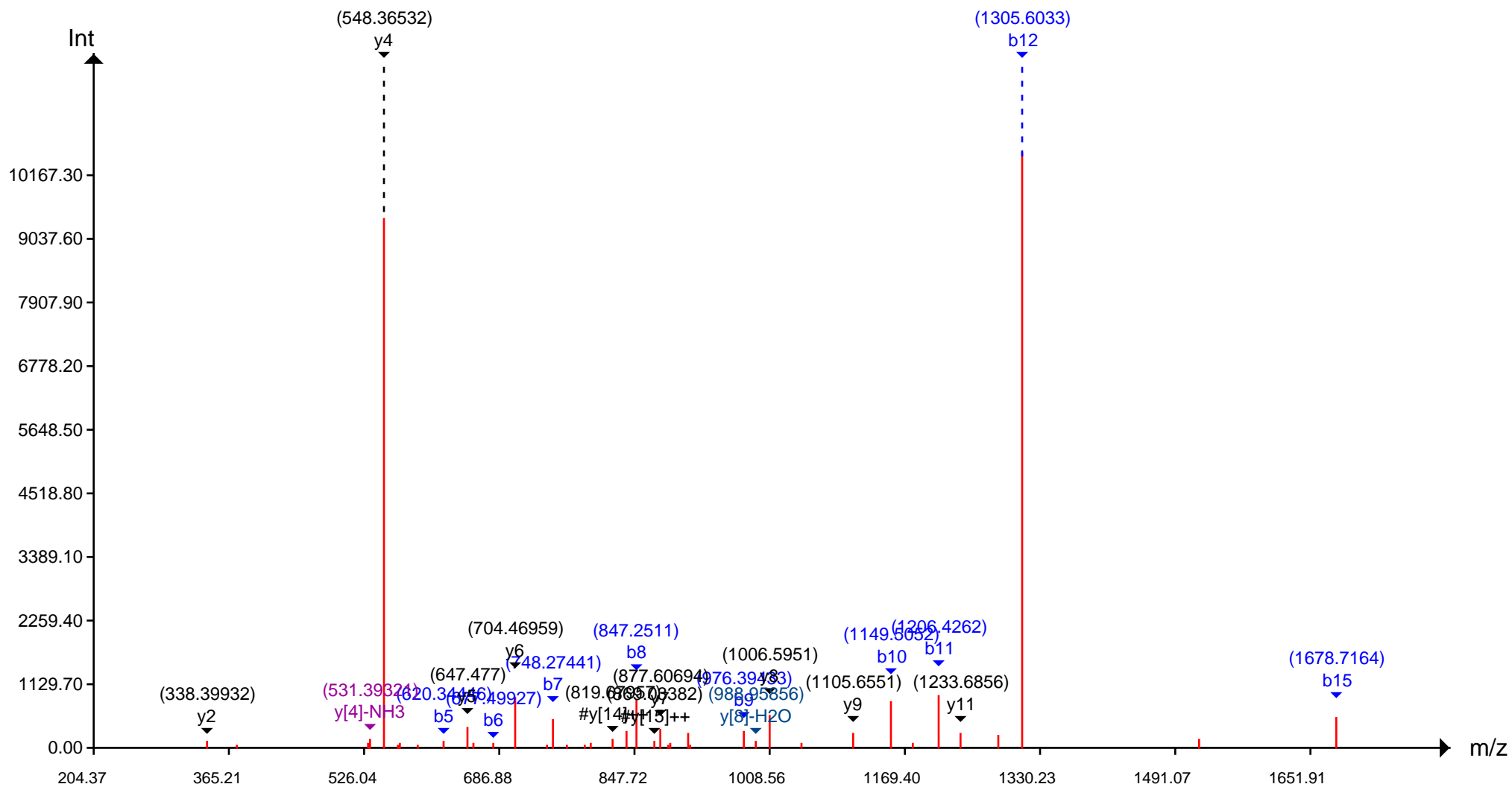
Identity Threshold 47.35

Mass error -0.001496



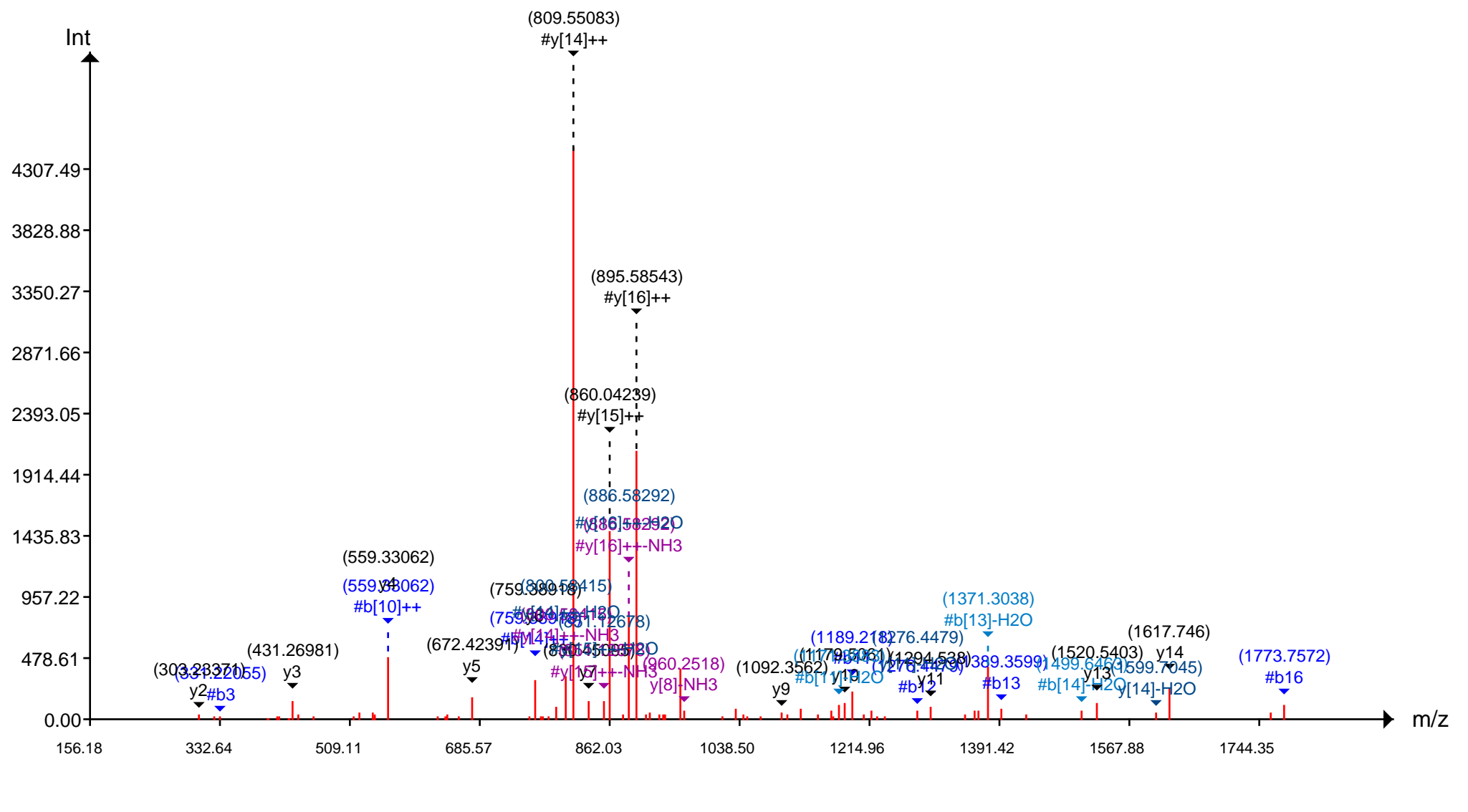
Info

Modified Sequence	AcD3-DEEETEEK<AcD3K*>VPVK<AcD3K*>K<AcD3K*>SVR<C13>-COOH	Ion score	62.47
Protein	Q61937	Confidence level	99.00%
Precursor m/z	696.71154	Identity Threshold	47.64
Charge	3+	Mass error	8.99E-4

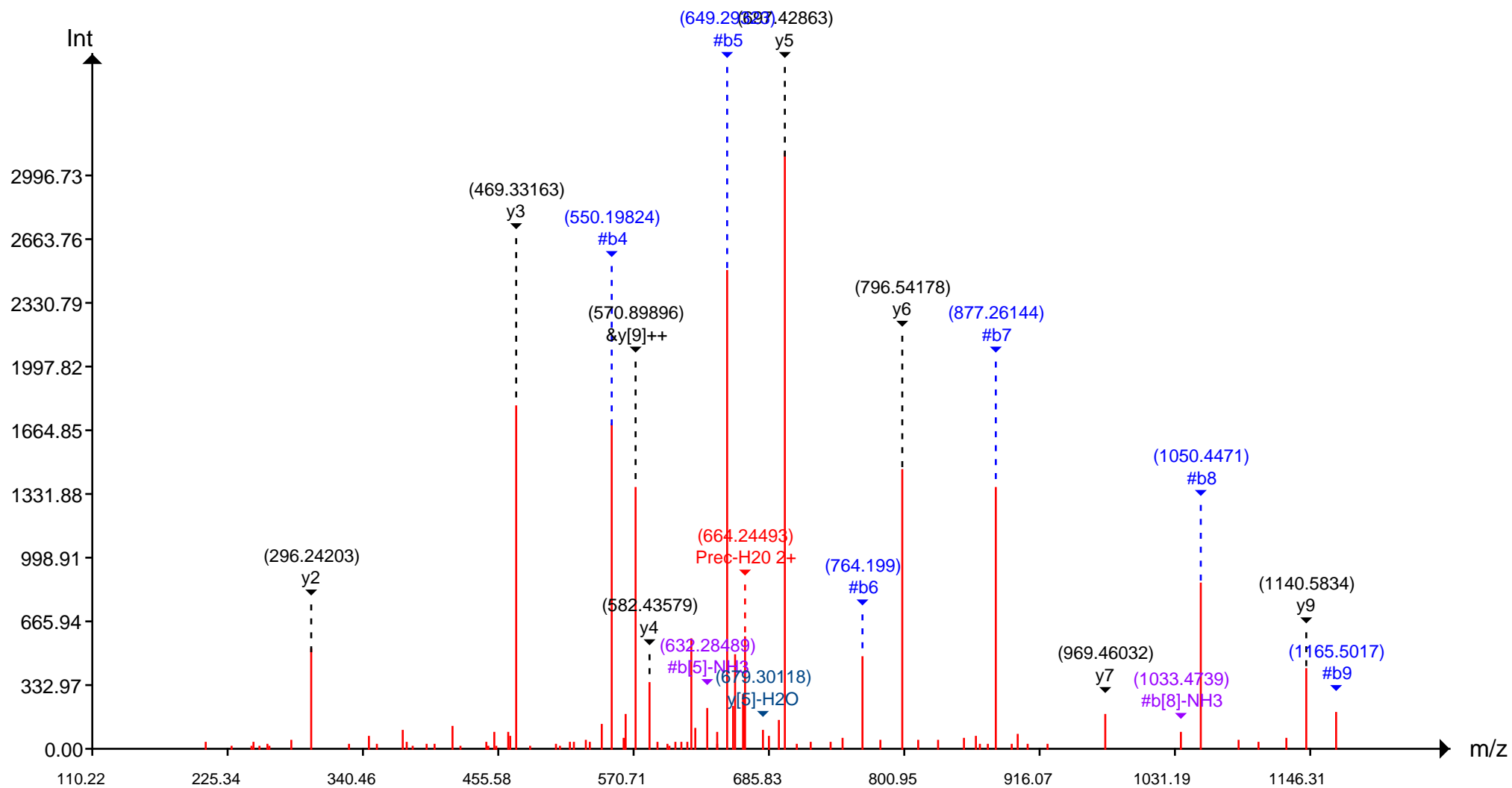


Info

Modified Sequence	AcD3-AVC<Cmm*>K<AcD3K*>AGAVEK<AcD3K*>GVPLYR-COOH	Ion score	66.99
Protein	P17182	Confidence level	99.00%
Precursor m/z	927.01885	Identity Threshold	46.58
Charge	2+	Mass error	4.58E-4

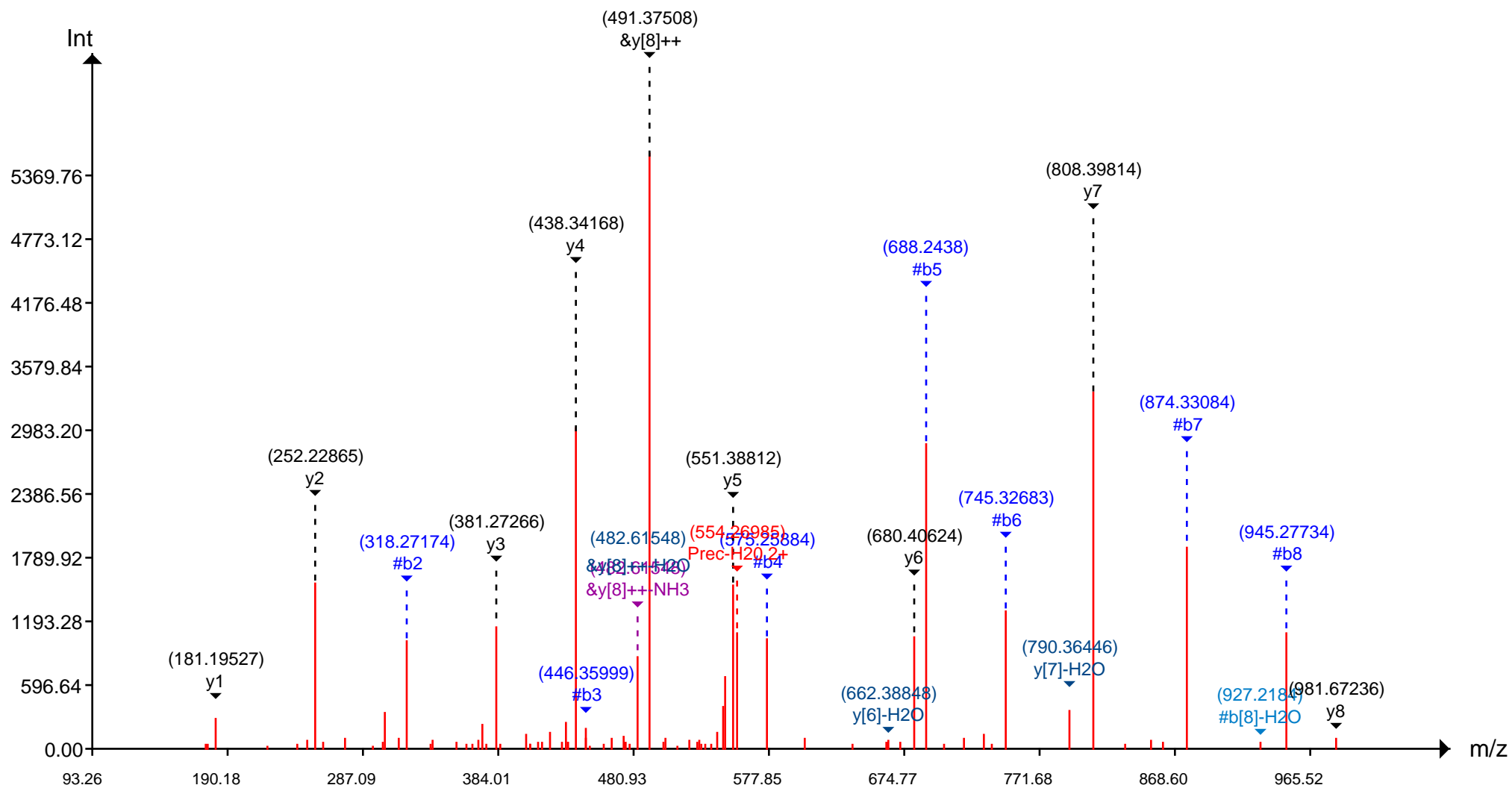


Info			
Modified Sequence	AcD3-IATPPEDSDM<Mox*>ASLQQQR-COOH	Ion score	82.04
Protein	Q9DBG5	Confidence level	99.00%
Precursor m/z	974.46256	Identity Threshold	46.09
Charge	2+	Mass error	0.002289



Info

Modified Sequence	AcD3-C<Cmm*>GNK<AcD3K*>VDIK<AcD3K*>DR<C13>-COOH	Ion score	59.64
Protein	P62827	Confidence level	99.00%
Precursor m/z	673.36311	Identity Threshold	47.87
Charge	2+	Mass error	3.34E-4



Info

Modified Sequence AcD3-VK<AcD3K*>QELGEAR<C13>-COOH

Ion score 65.85

Protein Q03958

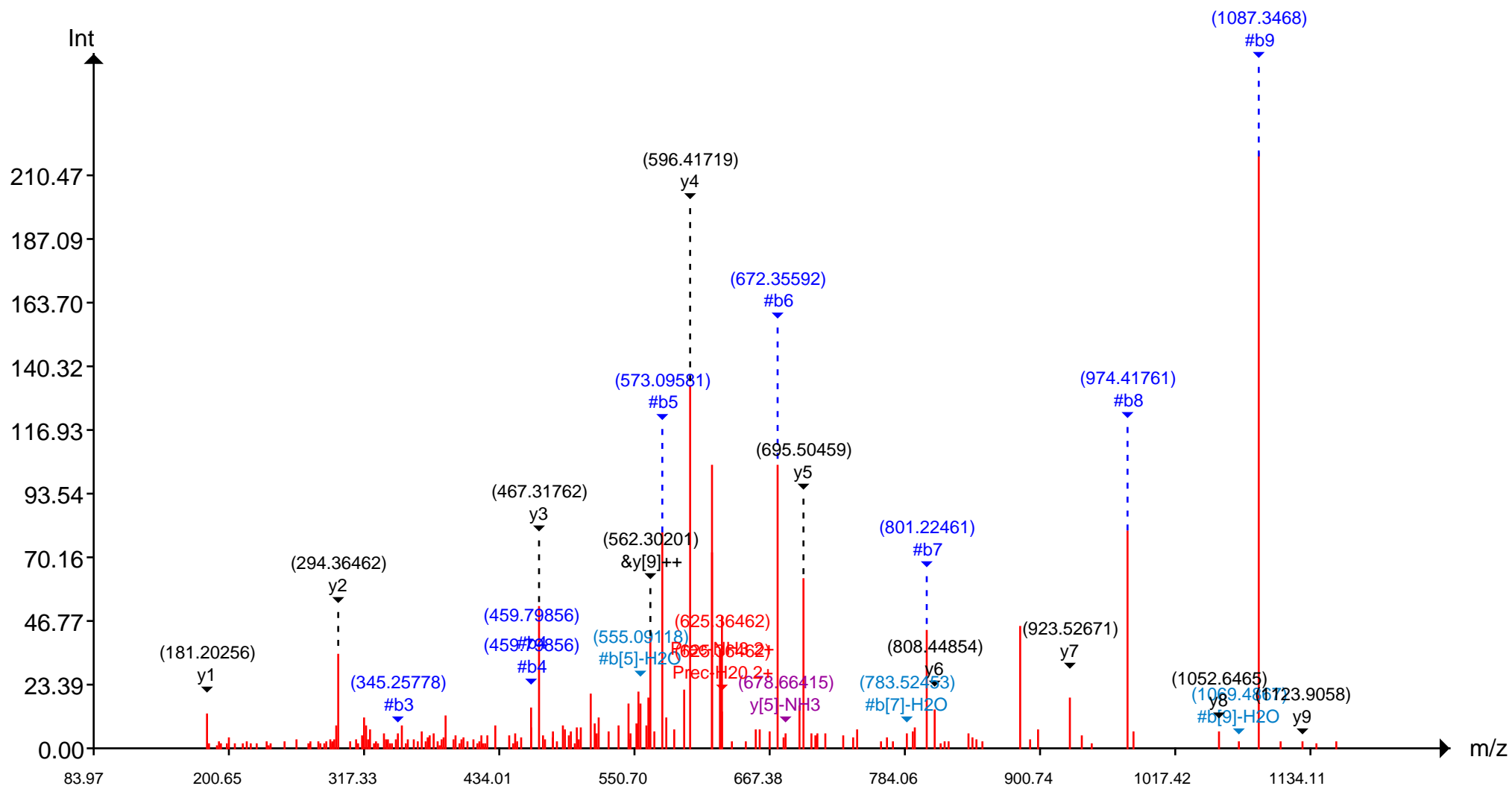
Confidence level 99.00%

Precursor m/z 563.32623

Identity Threshold 46.61

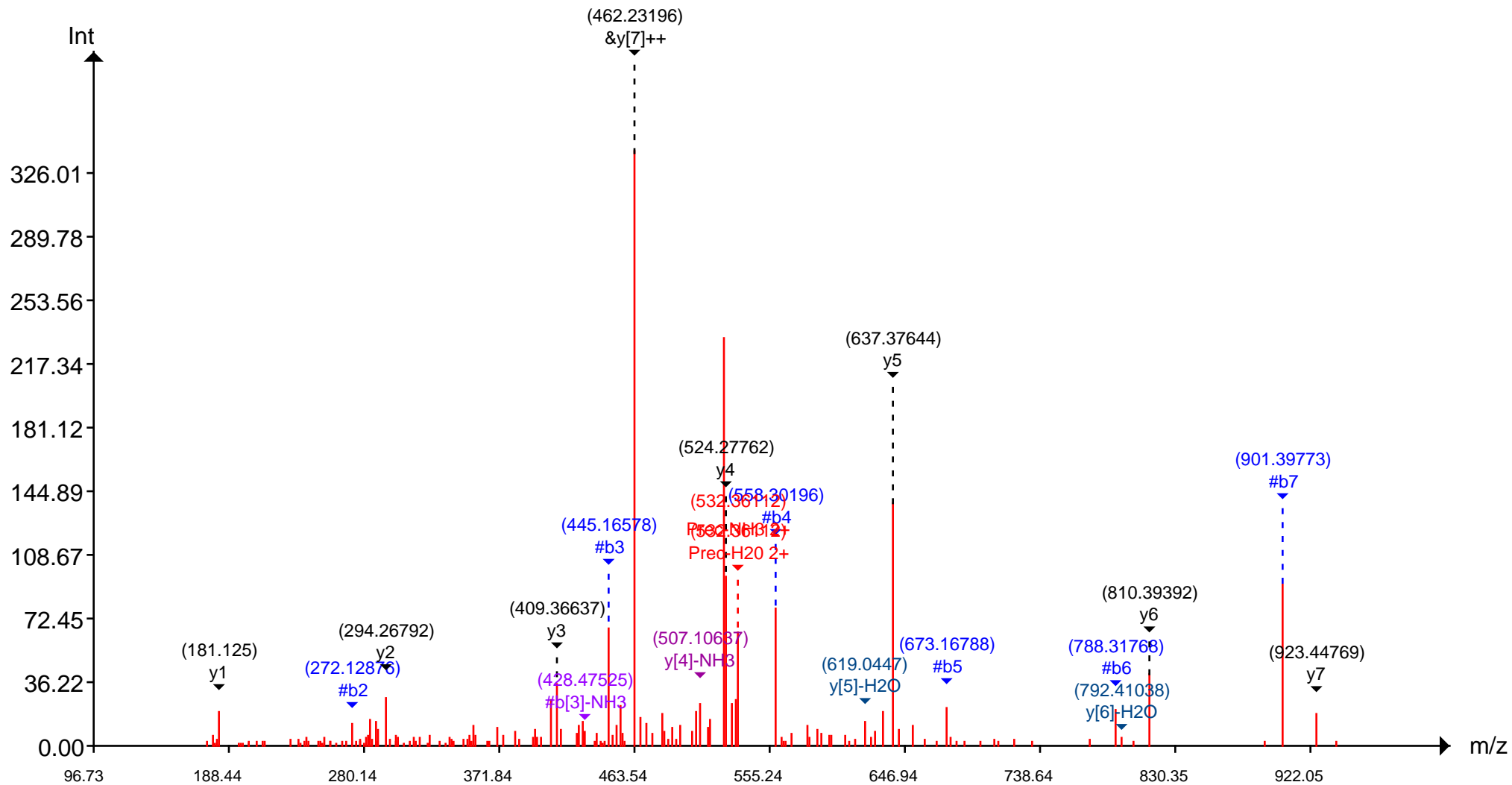
Charge 2+

Mass error -0.002458



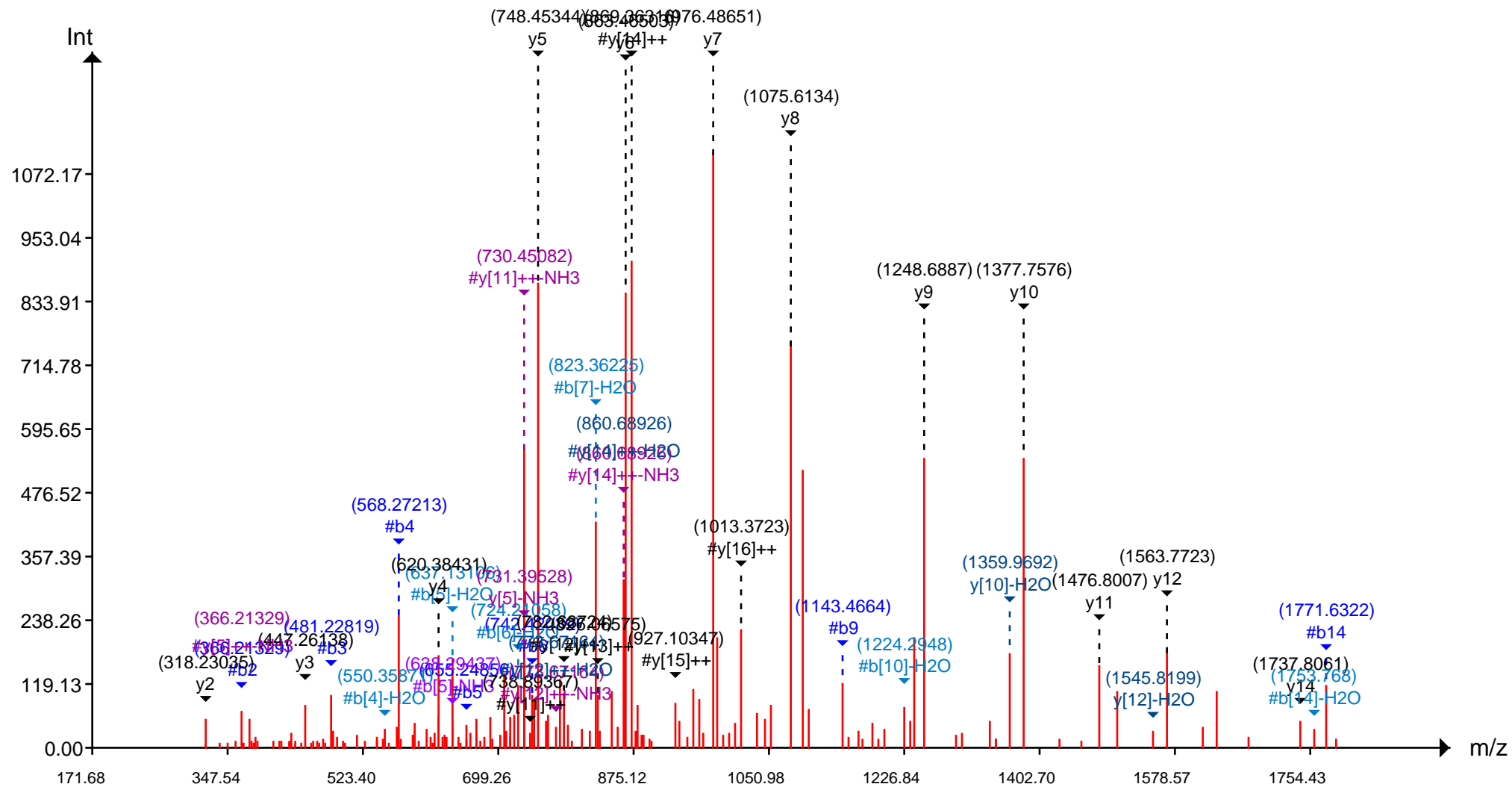
Info

Modified Sequence	AcD3-VAEDLVEK<AcD3K*>LR<C13>-COOH	Ion score	69.14
Protein	Q9CQI3	Confidence level	99.00%
Precursor m/z	634.37766	Identity Threshold	45.28
Charge	2+	Mass error	0.001022



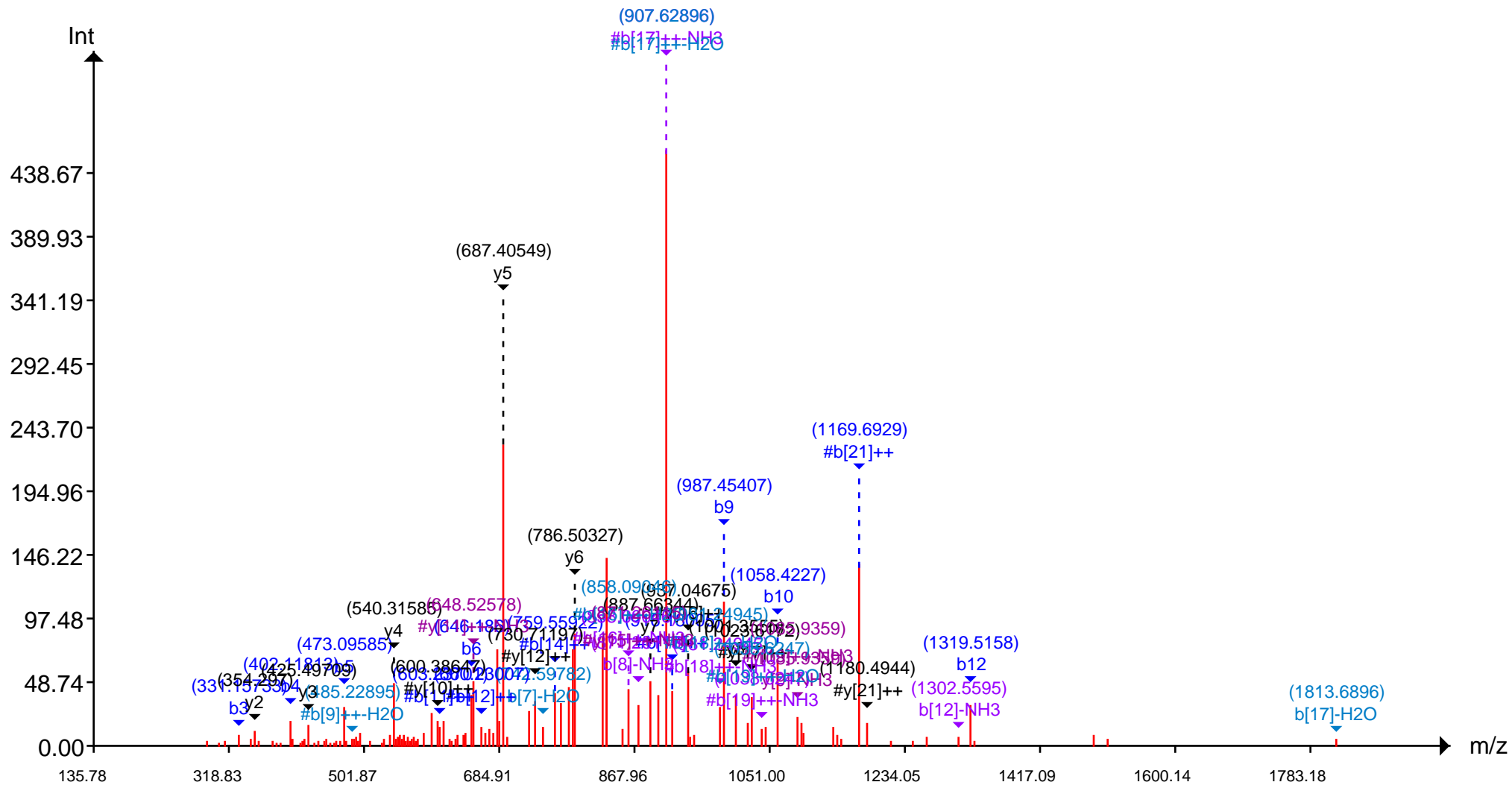
Info

Modified Sequence	AcD3-ILK<AcD3K*>IDDIR<C13>-COOH	Ion score	54.89
Protein	P80316	Confidence level	99.00%
Precursor m/z	541.34493	Identity Threshold	40.84
Charge	2+	Mass error	-3.82E-4



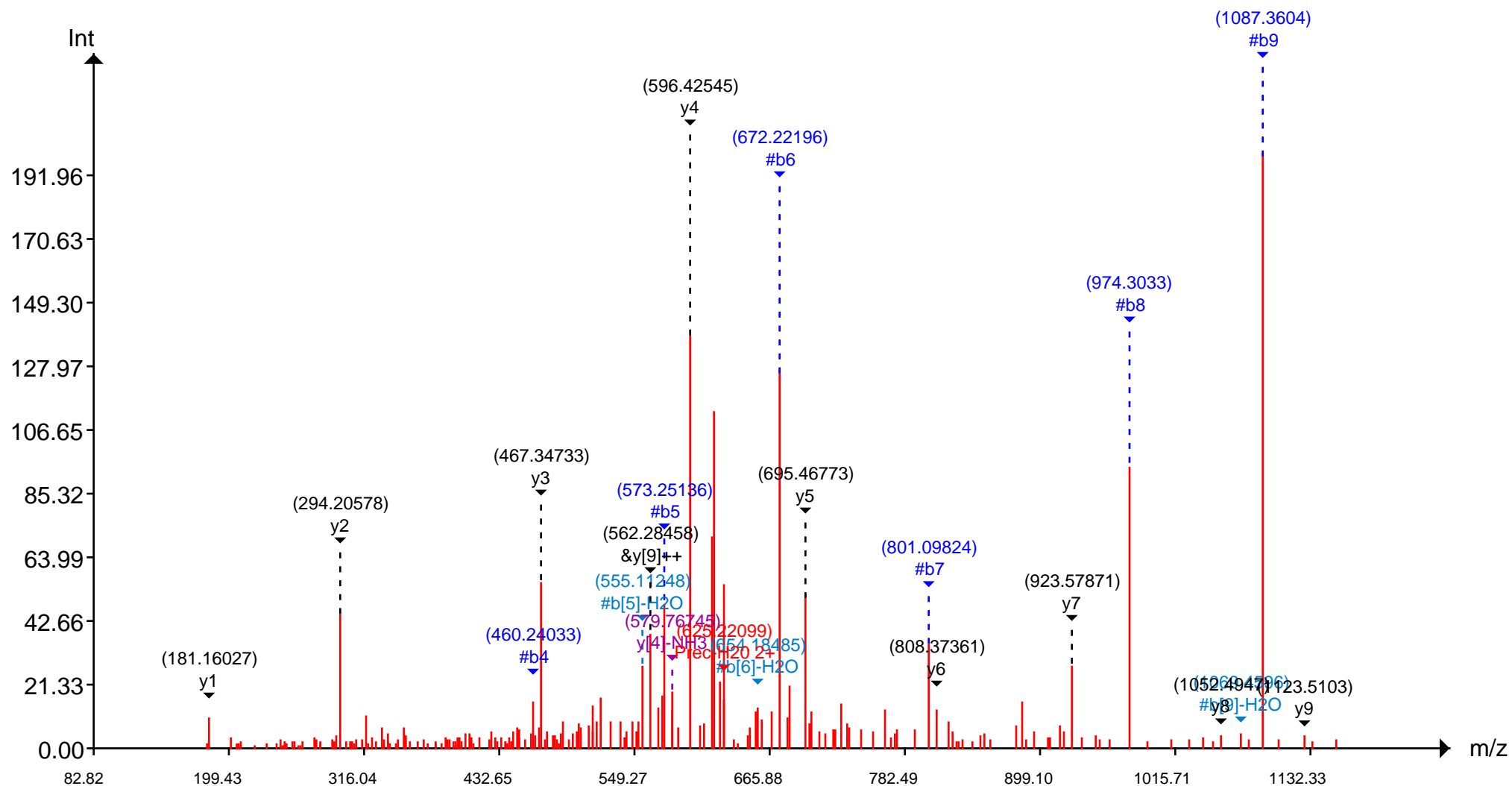
Info

Modified Sequence	AcD3-FK<AcD3K*>DSSSVEK<AcD3K*>VLDQK<AcD3K*>EHR<C13>-COOH	Ion score	84.77
Protein	Q99020	Confidence level	99.00%
Precursor m/z	1109.5962	Identity Threshold	47.59
Charge	2+	Mass error	0.001609



Info

Modified Sequence	AcD3-IGDAAK<AcD3K*>NQVAM<Mox*>NPTNTVFDAAK<AcD3K*>R<C13>-COOH	Ion score	54.47
Protein	P63017	Confidence level	99.00%
Precursor m/z	840.44023	Identity Threshold	47.86
Charge	3+	Mass error	1.004947



Info

Modified Sequence AcD3-VAEDLVEK<AcD3K*>LR<C13>-COOH

Ion score 84.36

Protein Q9CQI3

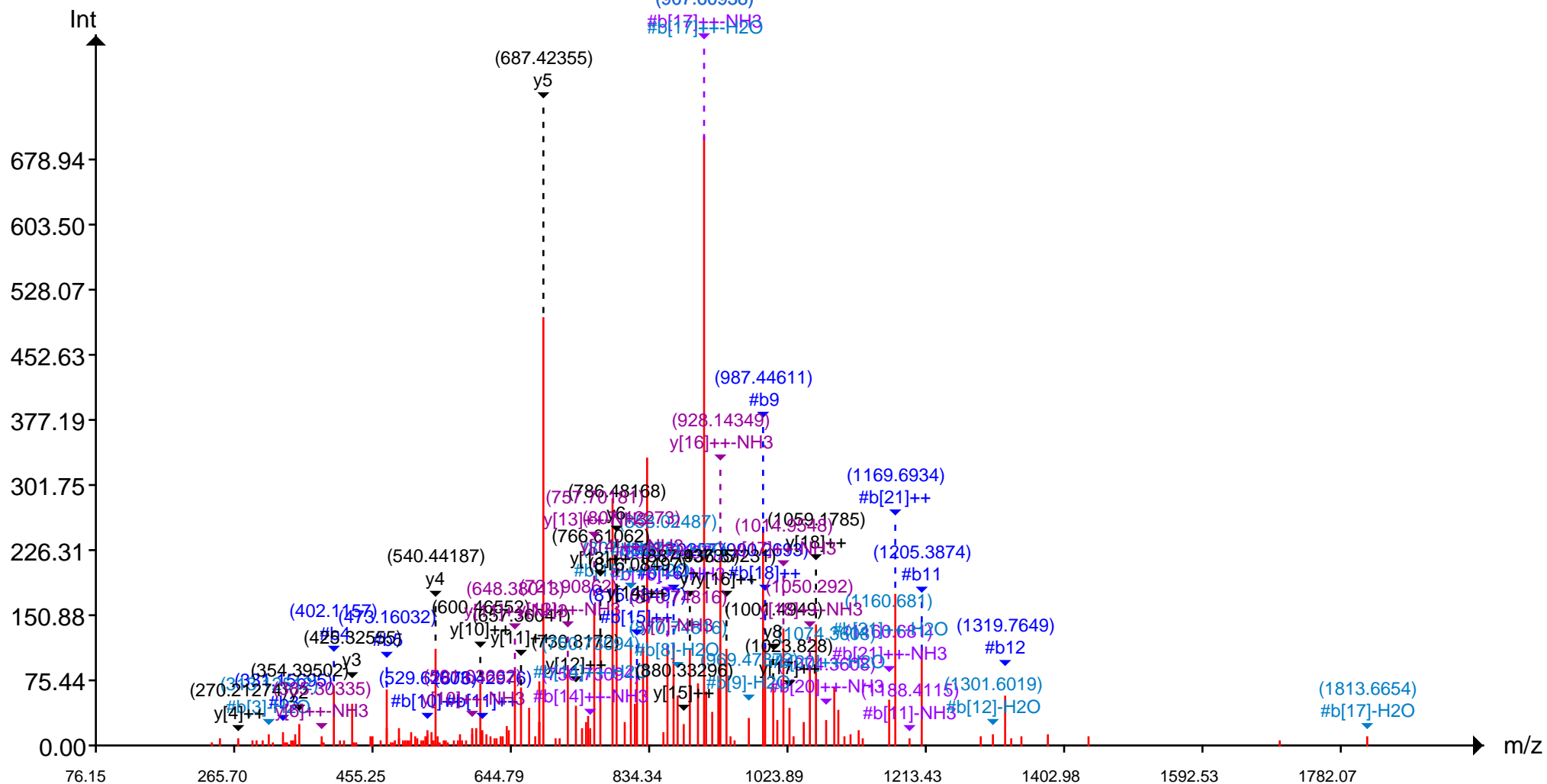
Confidence level 99.00%

Precursor m/z 634.37741

Identity Threshold 45.45

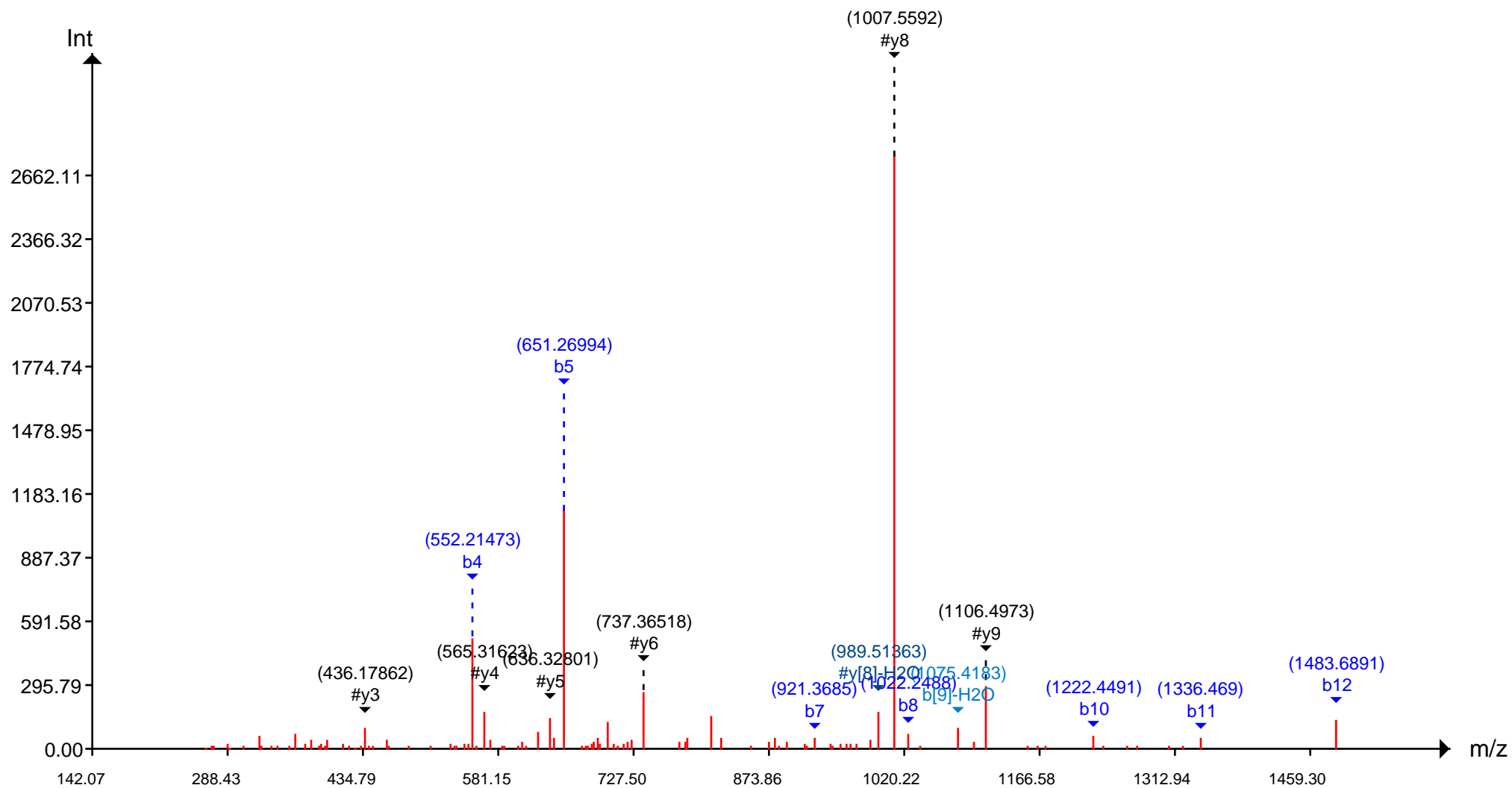
Charge 2+

Mass error 5.22E-4



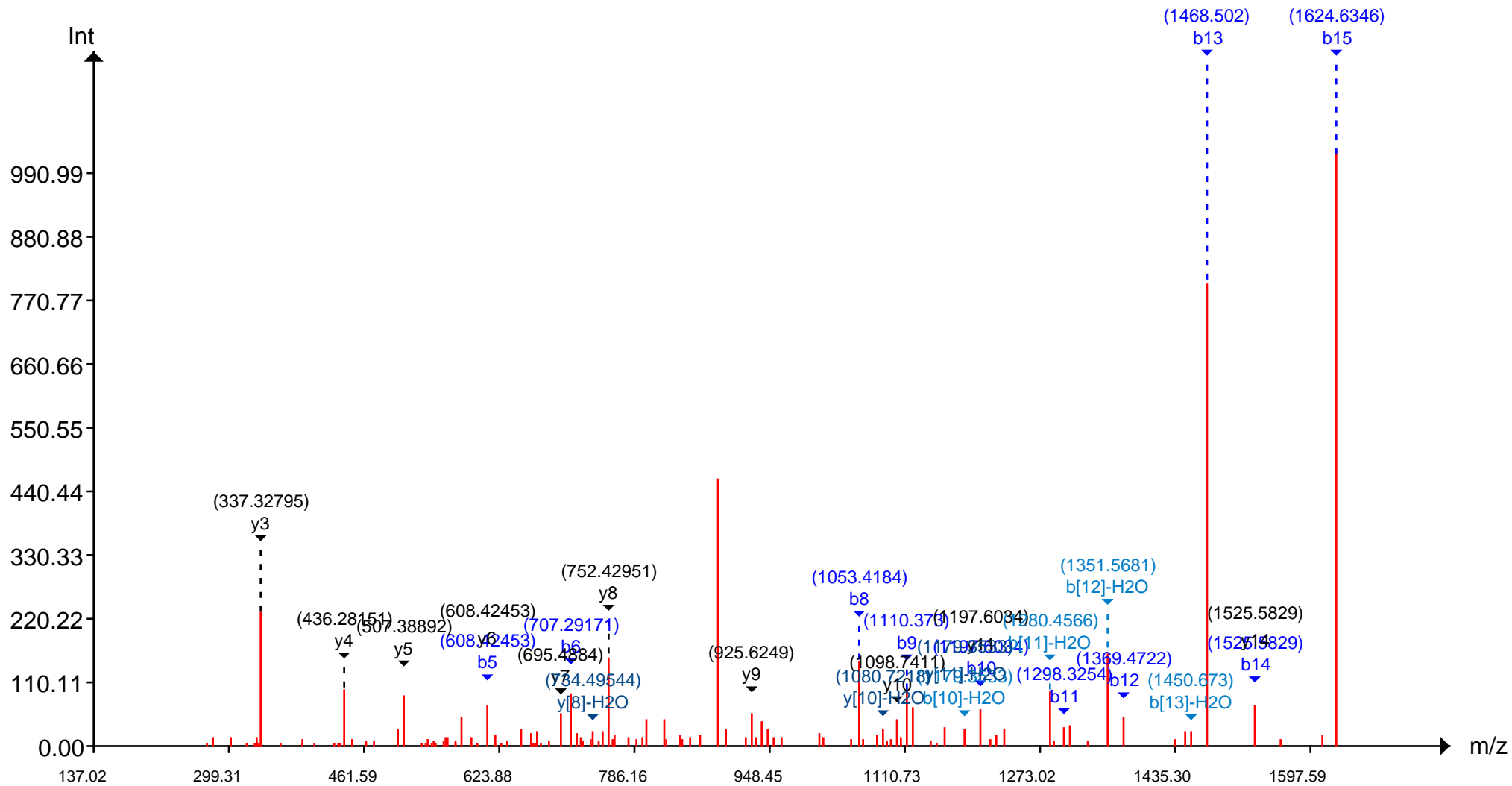
Info

Modified Sequence	AcD3-IGDAAK<AcD3K*>NQVAM<Mox*>NPTNTVFDAAK<AcD3K*>R<C13>-COOH	Ion score	54.75
Protein	P63017	Confidence level	99.00%
Precursor m/z	840.10612	Identity Threshold	47.88
Charge	3+	Mass error	0.002617



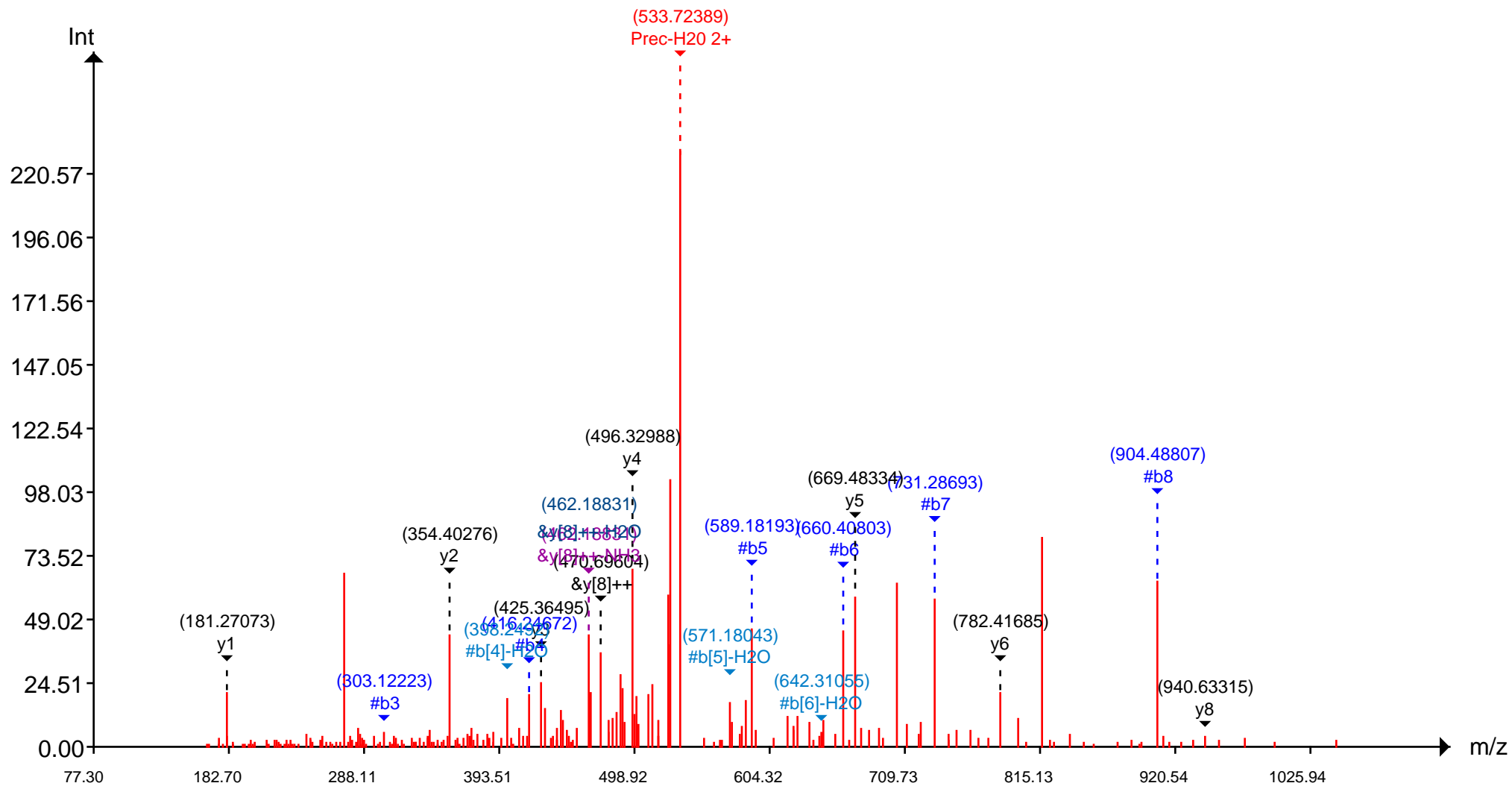
Info

Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH	Ion score	53.51
Protein	P17742	Confidence level	99.00%
Precursor m/z	829.44892	Identity Threshold	47.81
Charge	2+	Mass error	0.001132



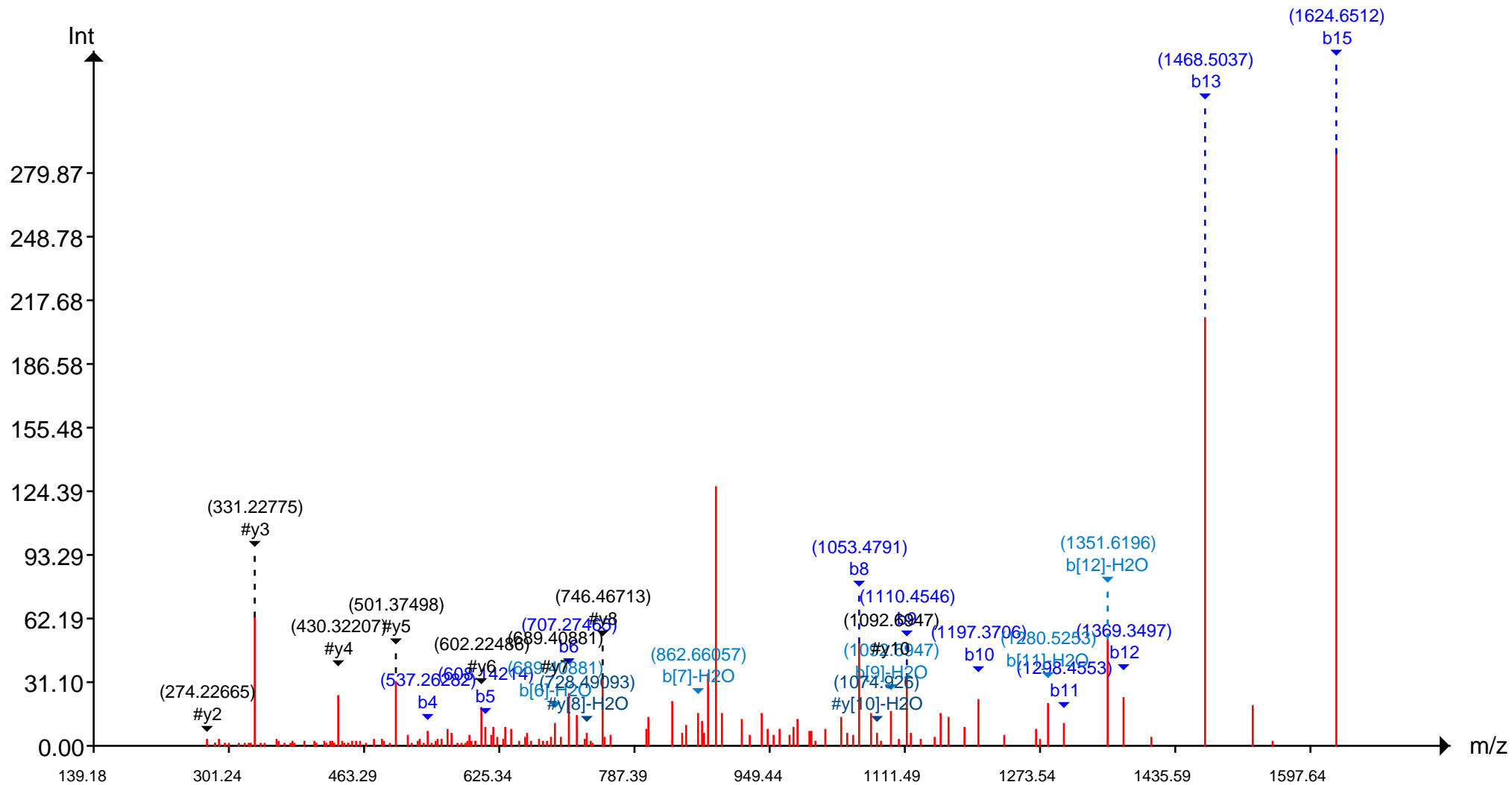
Info

Modified Sequence	AcD3-YAQEAVK<AcD3K*>K<AcD3K*>GSTAVGVR<C13>-COOH	Ion score	76.72
Protein	Q9Z2U0	Confidence level	99.00%
Precursor m/z	903.01551	Identity Threshold	45.72
Charge	2+	Mass error	0.002827



Info

Modified Sequence	AcD3-VGTLK<AcD3K*>AAK<AcD3K*>R<C13>-COOH	Ion score	43.1
Protein	Q4KML4	Confidence level	99.00%
Precursor m/z	542.86017	Identity Threshold	38.40
Charge	2+	Mass error	2.3E-5



Info

Modified Sequence AcD3-YAQEAVK<AcD3K*>K<AcD3K*>GSTAVGVR-COOH

Ion score 51.08

Protein Q9Z2U0

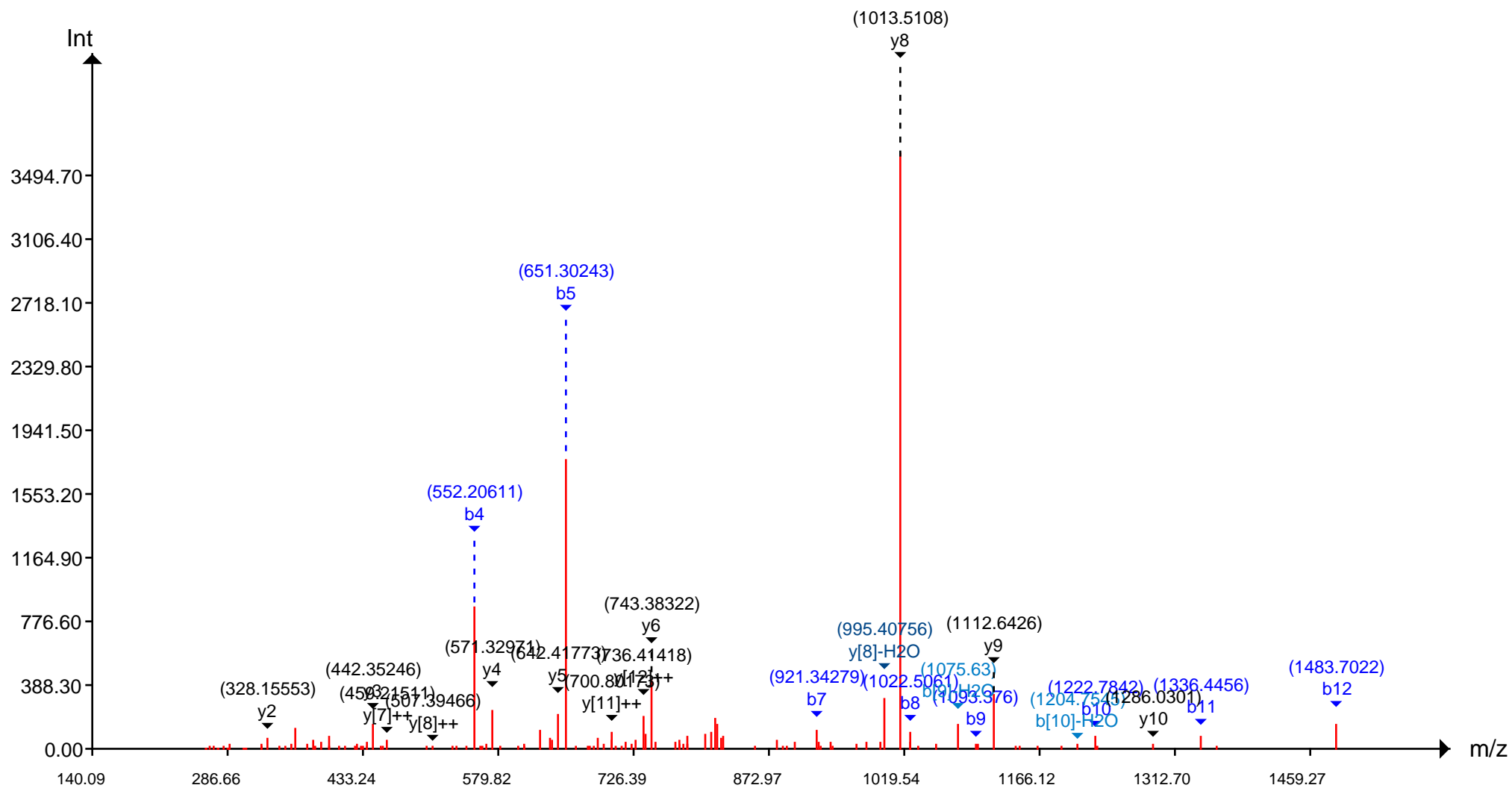
Confidence level 99.00%

Precursor m/z 900.00529

Identity Threshold 46.60

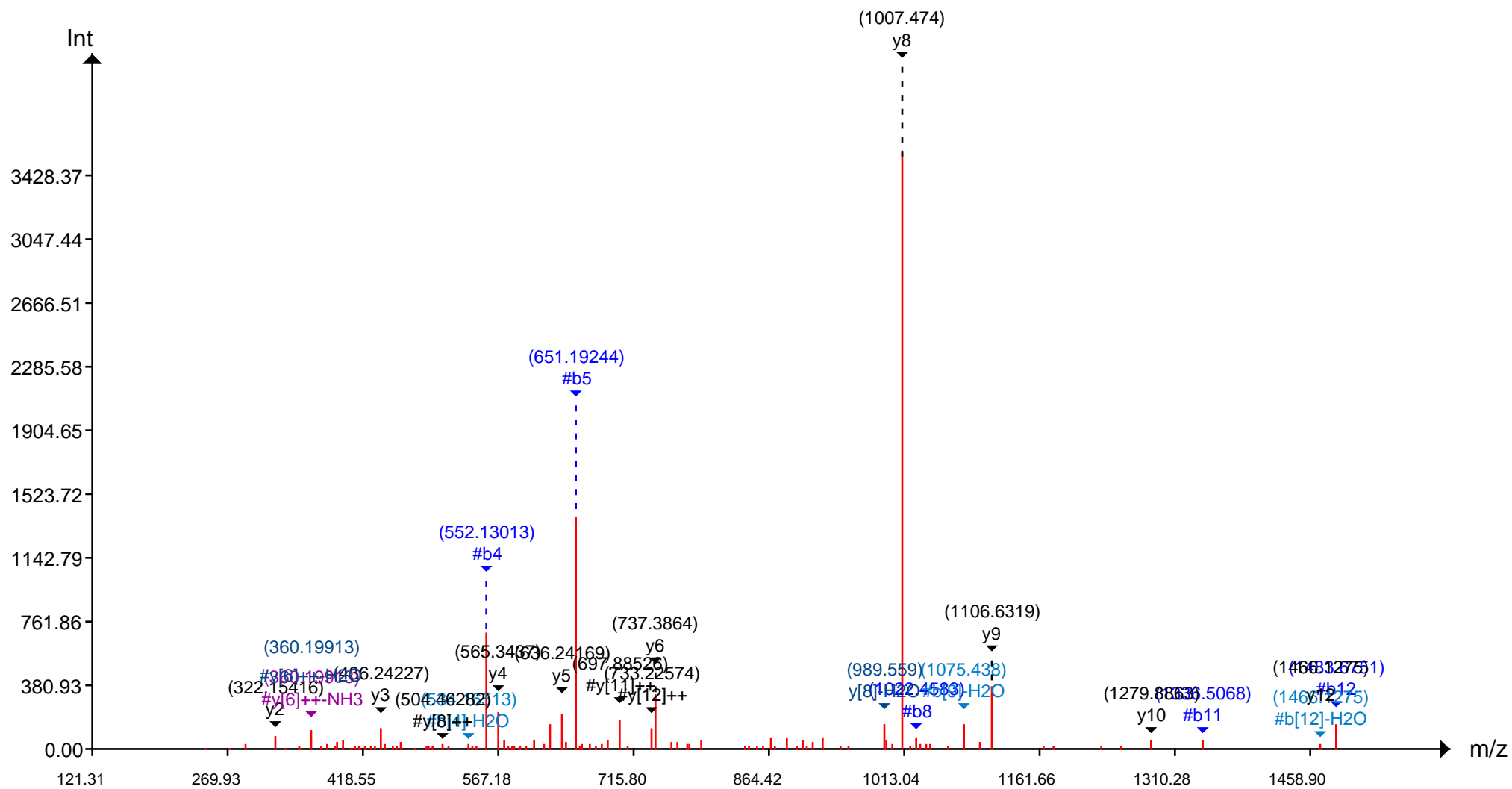
Charge 2+

Mass error 0.002528



Info

Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH	Ion score	62.59
Protein	P17742	Confidence level	99.00%
Precursor m/z	832.45937	Identity Threshold	47.48
Charge	2+	Mass error	0.00189



Info

Modified Sequence AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH

Ion score 58.52

Protein P17742

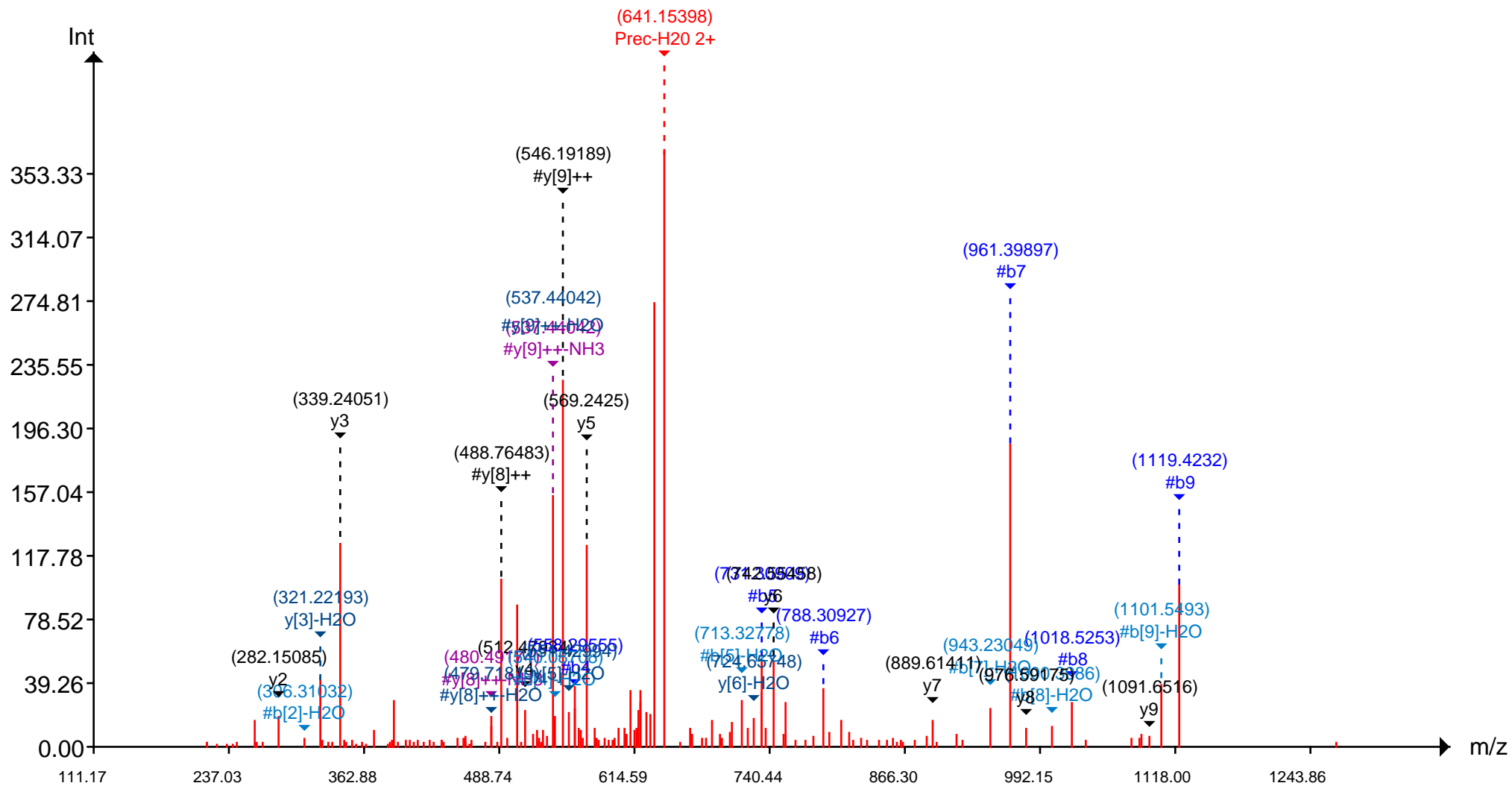
Confidence level 99.00%

Precursor m/z 829.44962

Identity Threshold 47.77

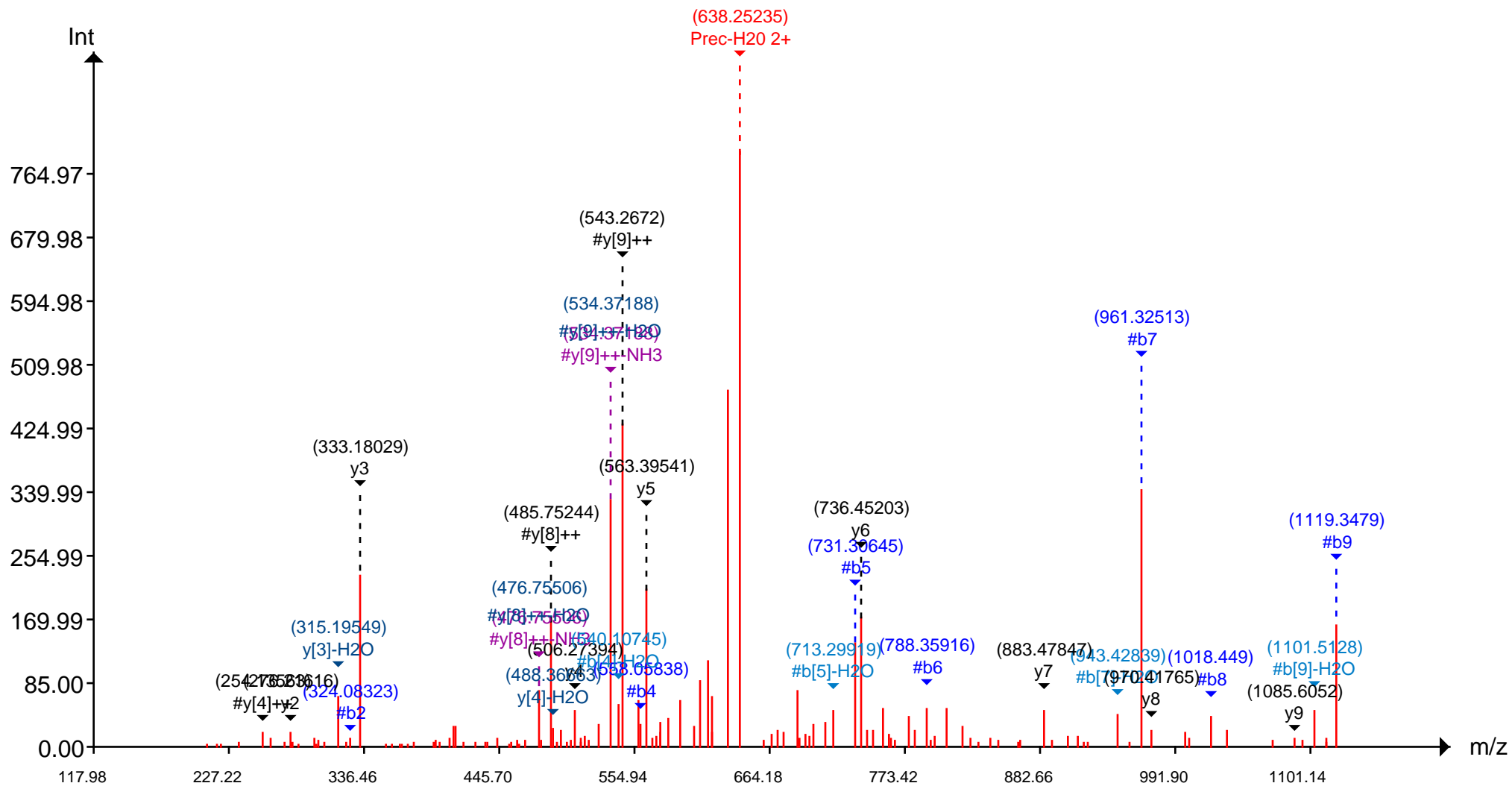
Charge 2+

Mass error 0.002532



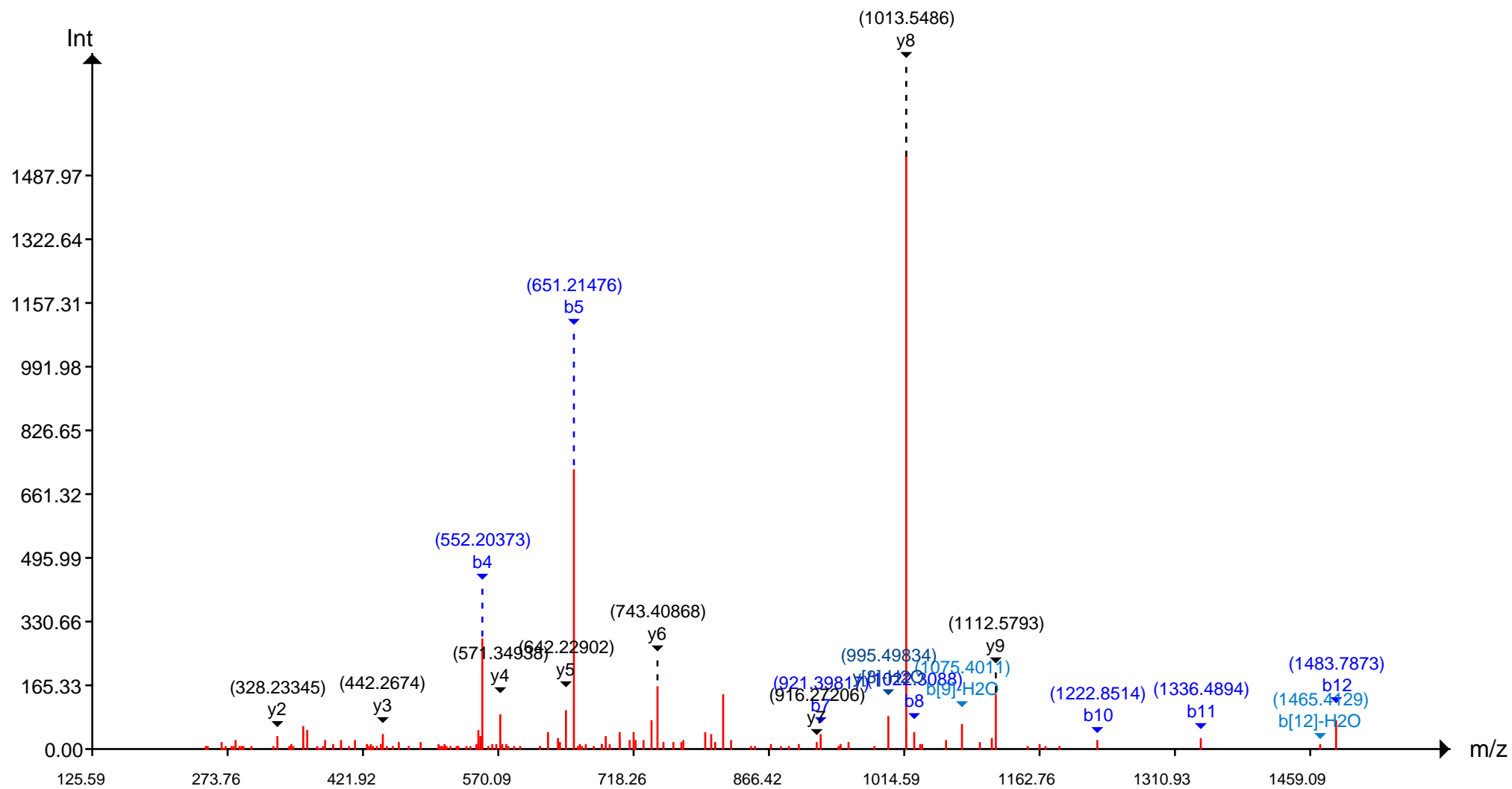
Info

Modified Sequence	AcD3-YDSM<Mox*>K<AcD3K*>GK<AcD3K*>GTR<C13>-COOH	Ion score	58.78
Protein	P07356	Confidence level	99.00%
Precursor m/z	650.33673	Identity Threshold	47.96
Charge	2+	Mass error	6.74E-4



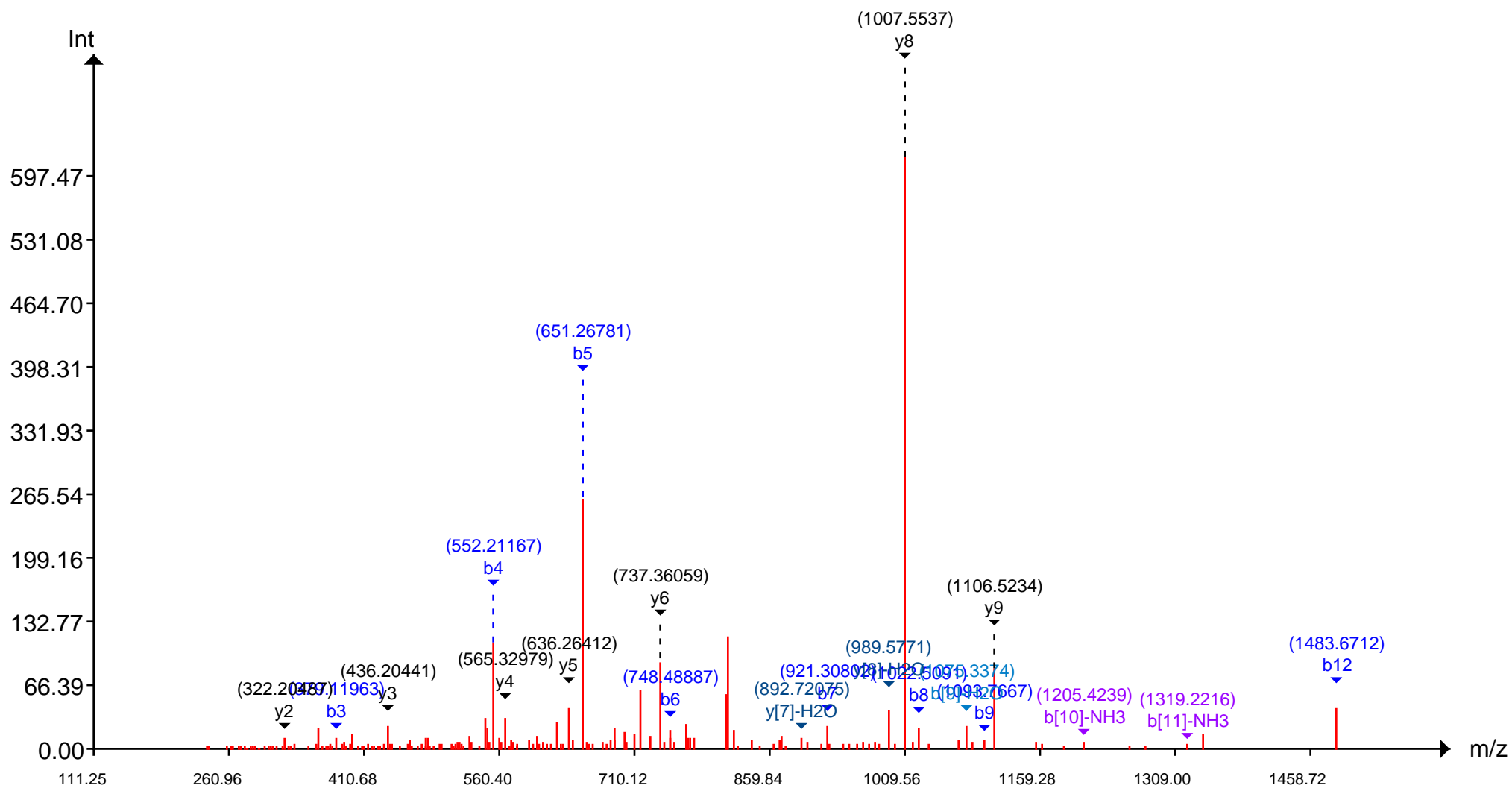
Info

Modified Sequence	AcD3-YDSM<Mox*>K<AcD3K*>GK<AcD3K*>GTR-COOH	Ion score	52.81
Protein	P07356	Confidence level	99.00%
Precursor m/z	647.3269	Identity Threshold	47.80
Charge	2+	Mass error	0.001156



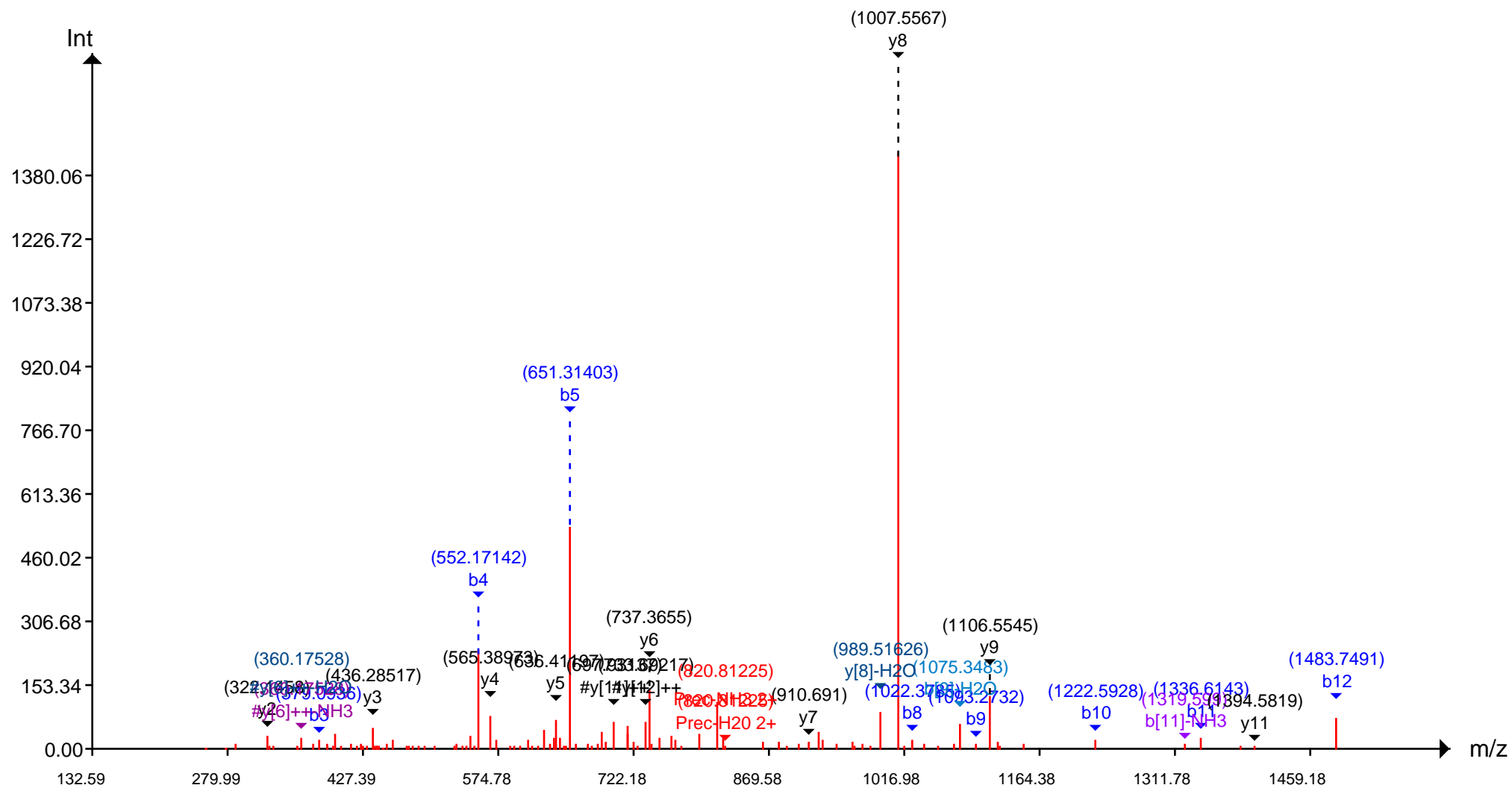
Info

Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH	Ion score	50.46
Protein	P17742	Confidence level	99.00%
Precursor m/z	832.45877	Identity Threshold	47.57
Charge	2+	Mass error	6.9E-4



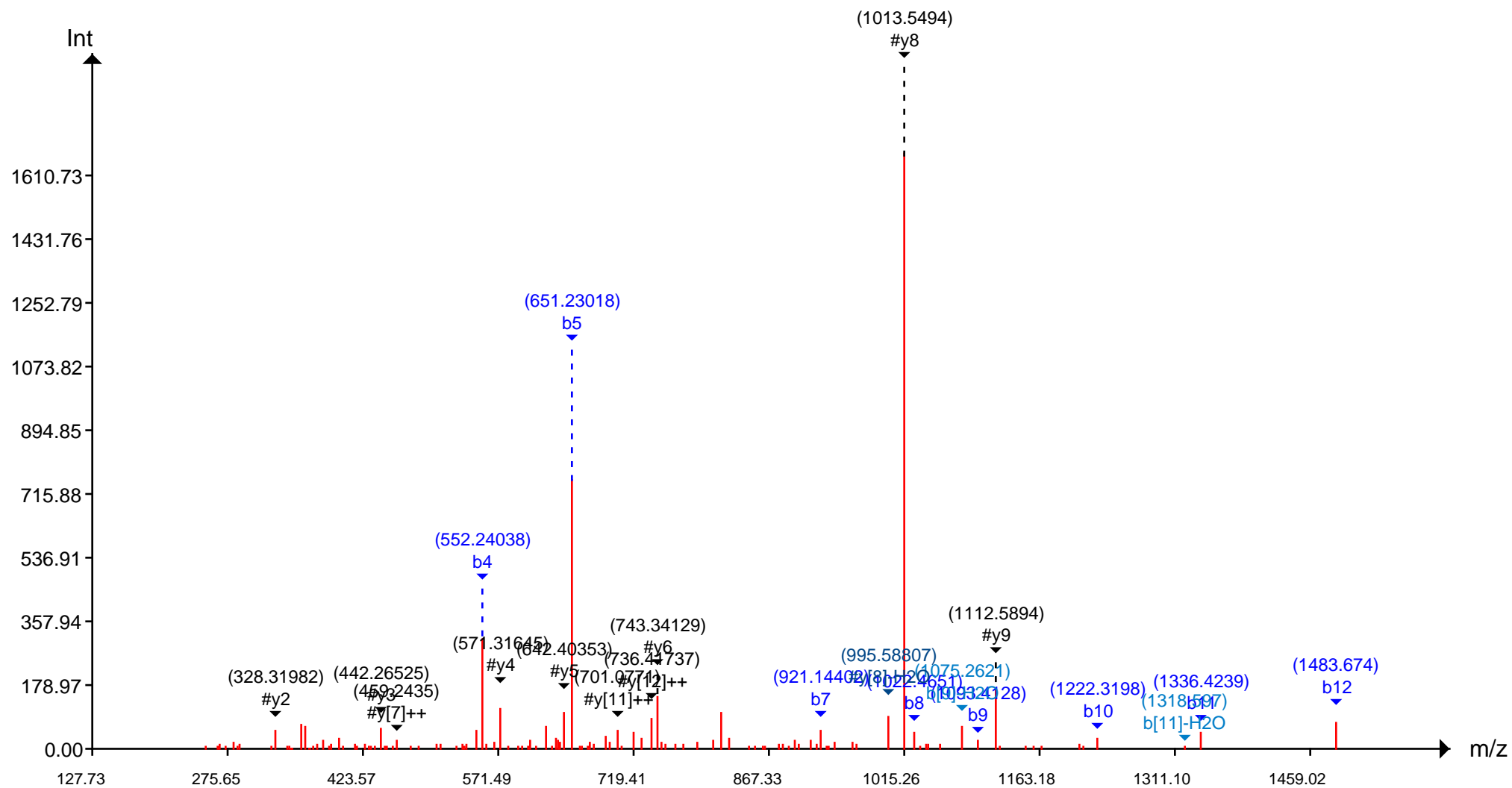
Info

Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH	Ion score	49.12
Protein	P17742	Confidence level	99.00%
Precursor m/z	829.44892	Identity Threshold	47.81
Charge	2+	Mass error	0.001132



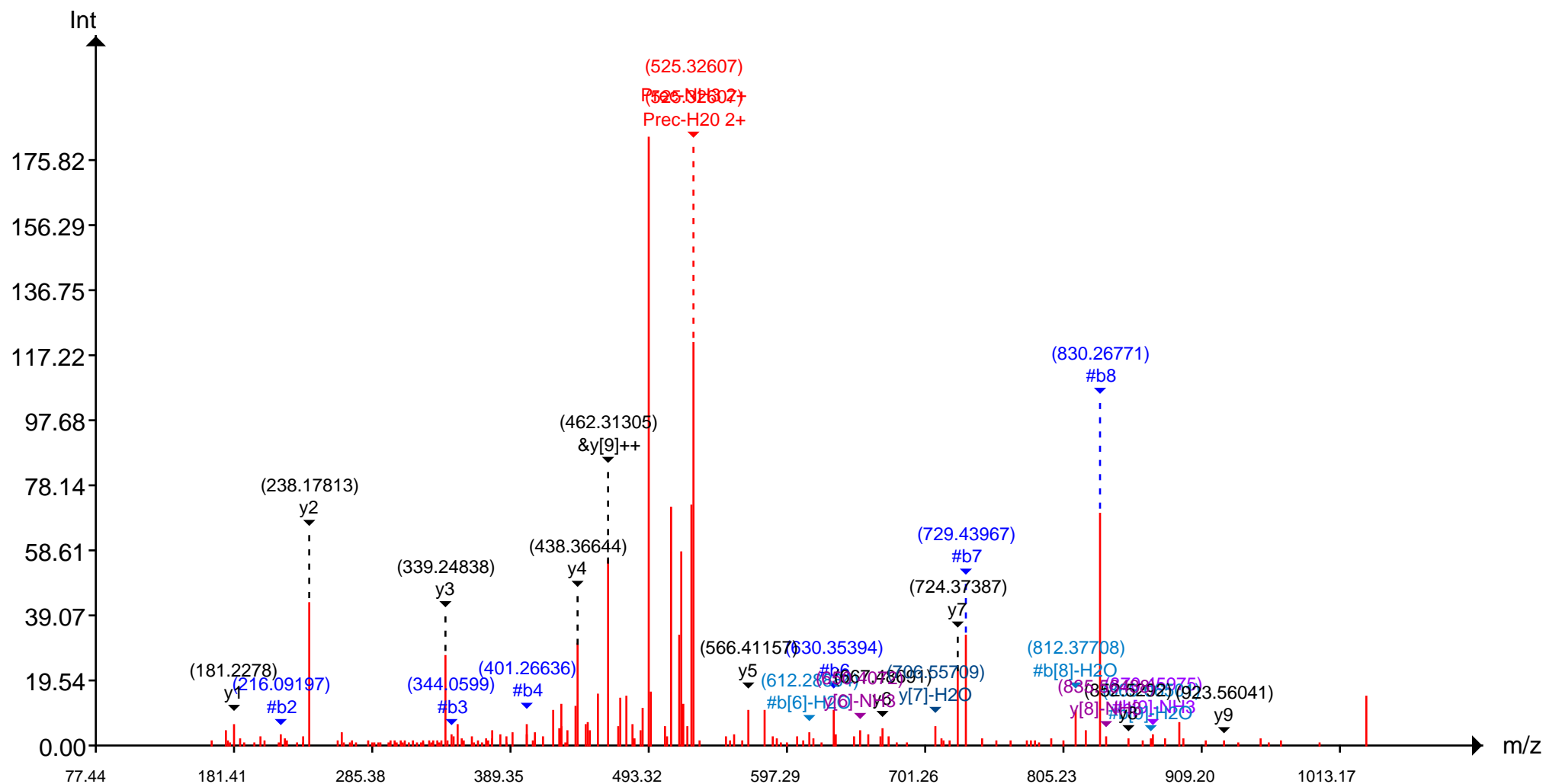
Info

Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH	Ion score	60.79
Protein	P17742	Confidence level	99.00%
Precursor m/z	829.44935	Identity Threshold	47.77
Charge	2+	Mass error	0.001992



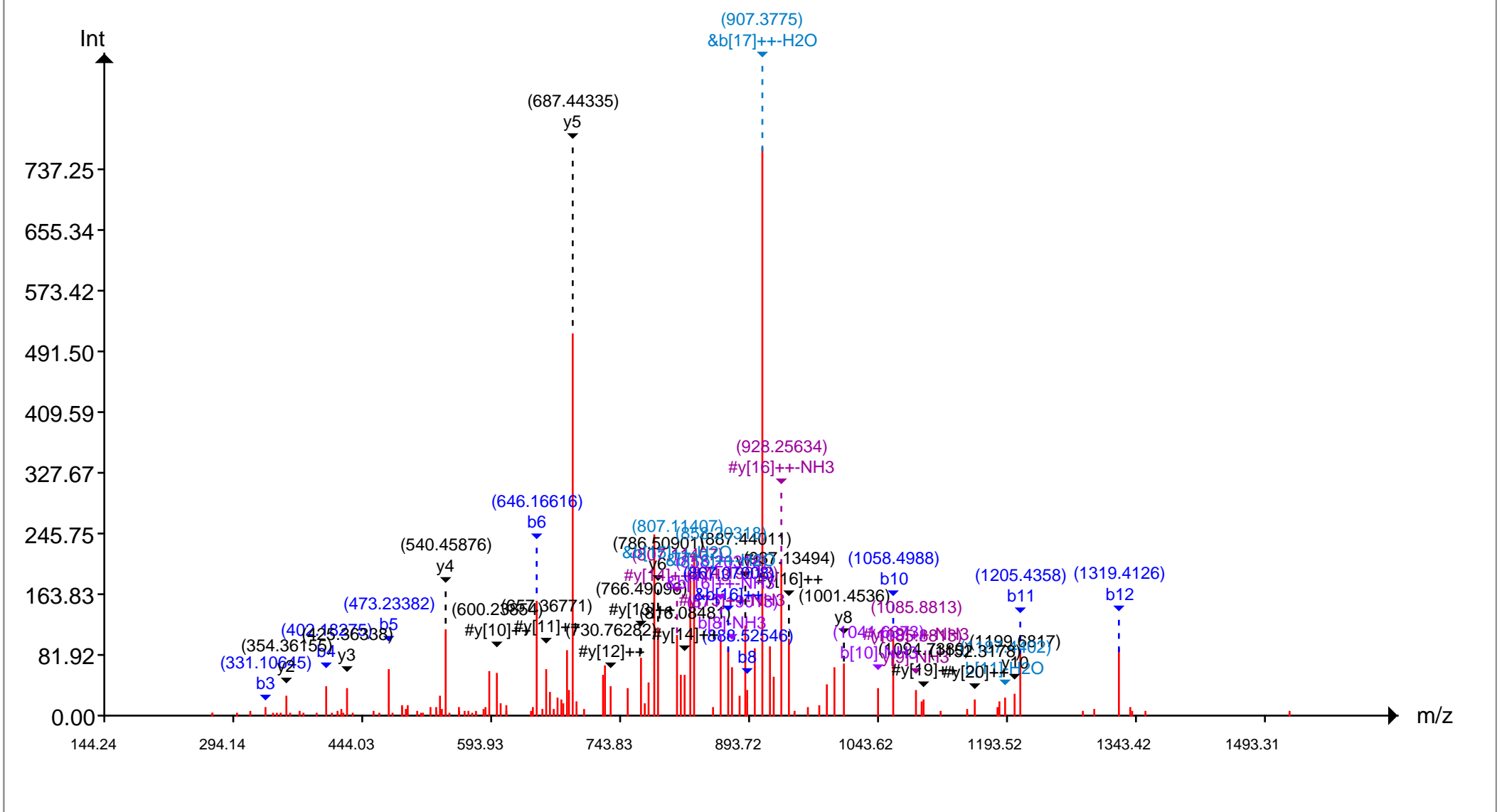
Info

Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH	Ion score	50.92
Protein	P17742	Confidence level	99.00%
Precursor m/z	832.45916	Identity Threshold	47.45
Charge	2+	Mass error	0.00147



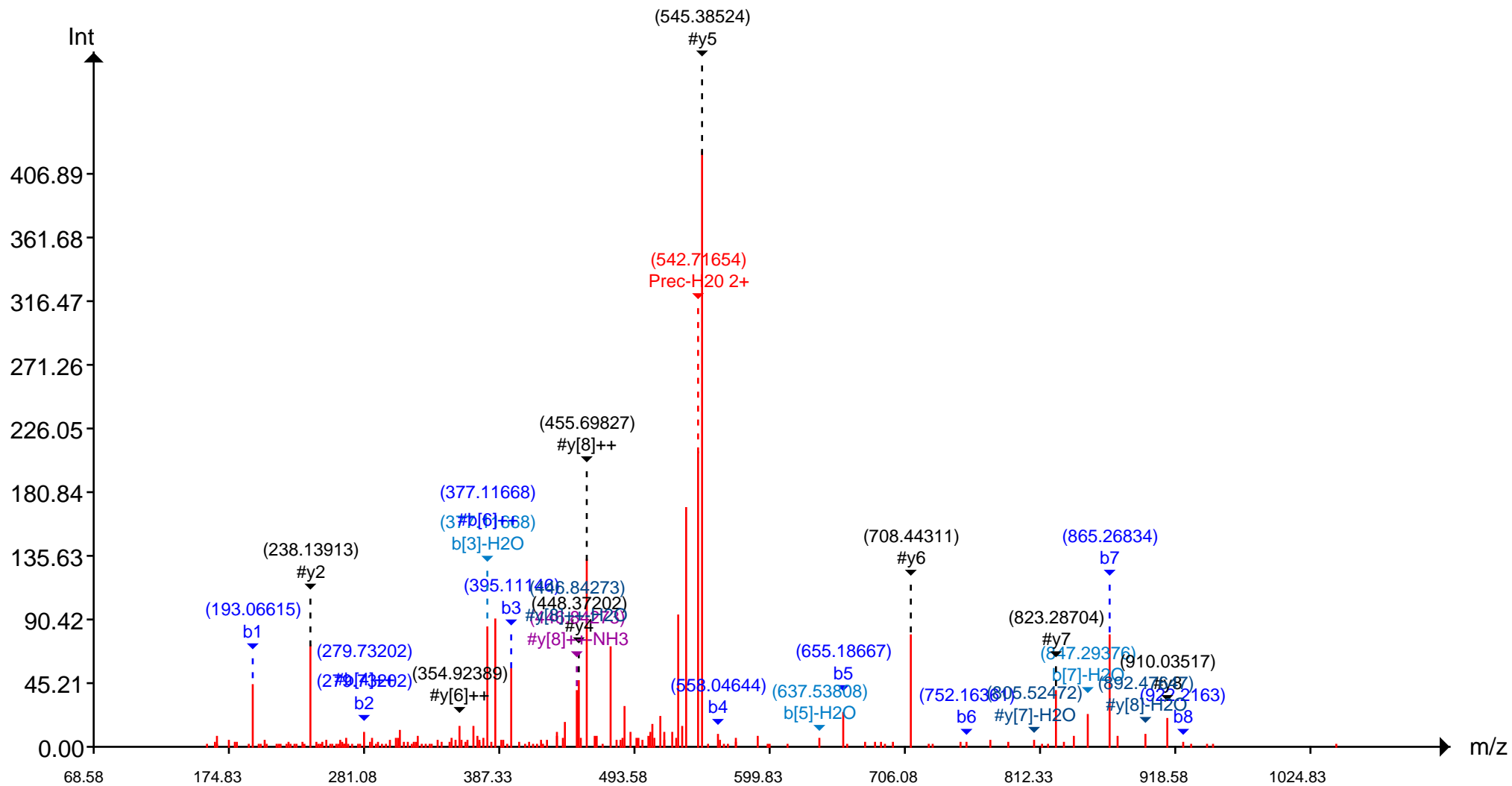
Info

Modified Sequence	AcD3-VAQGTQVTGR<C13>-COOH	Ion score	50.89
Protein	P50580	Confidence level	99.00%
Precursor m/z	534.30244	Identity Threshold	47.28
Charge	2+	Mass error	-2.79E-4



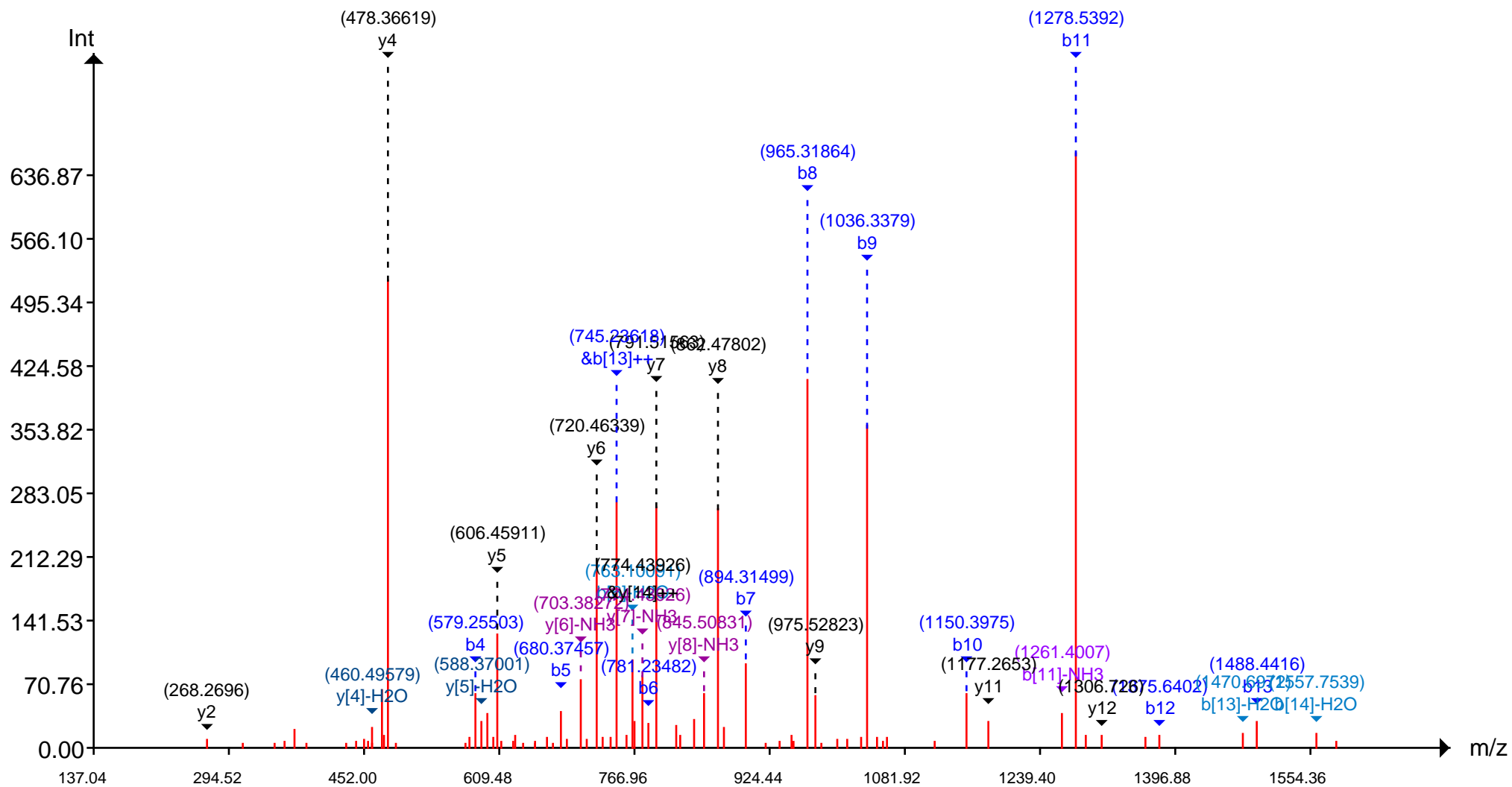
Info

Modified Sequence	AcD3-IGDAAK<AcD3K*>NQVAM<Mox*>NPTNTVFDAAK<AcD3K*>R<C13>-COOH	Ion score	63.0
Protein	P63017	Confidence level	99.00%
Precursor m/z	840.43978	Identity Threshold	47.87
Charge	3+	Mass error	1.003597



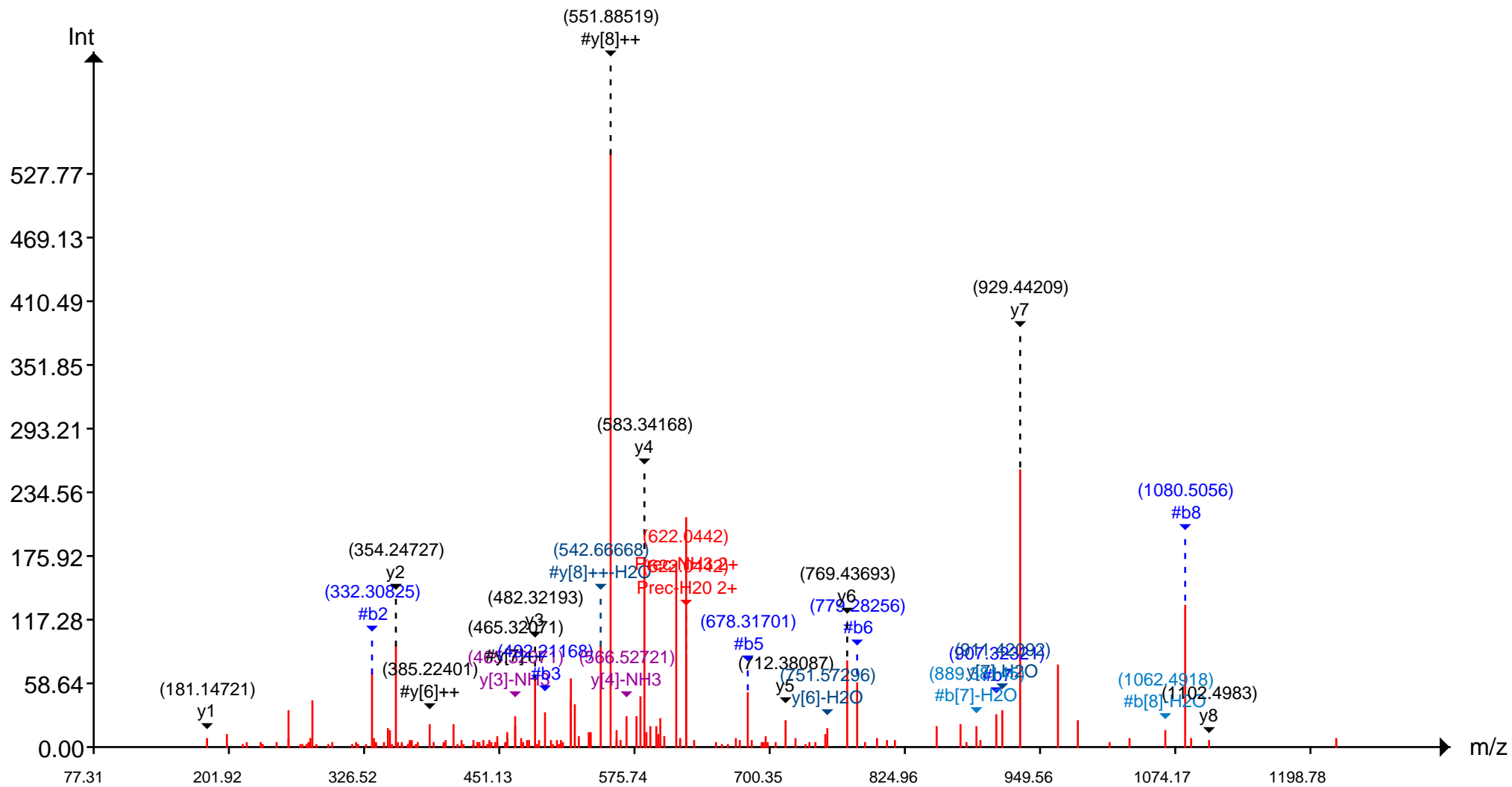
Info

Modified Sequence	AcD3-FSDYPPLGR<C13>-COOH	Ion score	55.29
Protein	P10126	Confidence level	99.00%
Precursor m/z	551.78874	Identity Threshold	47.66
Charge	2+	Mass error	-4.5E-5



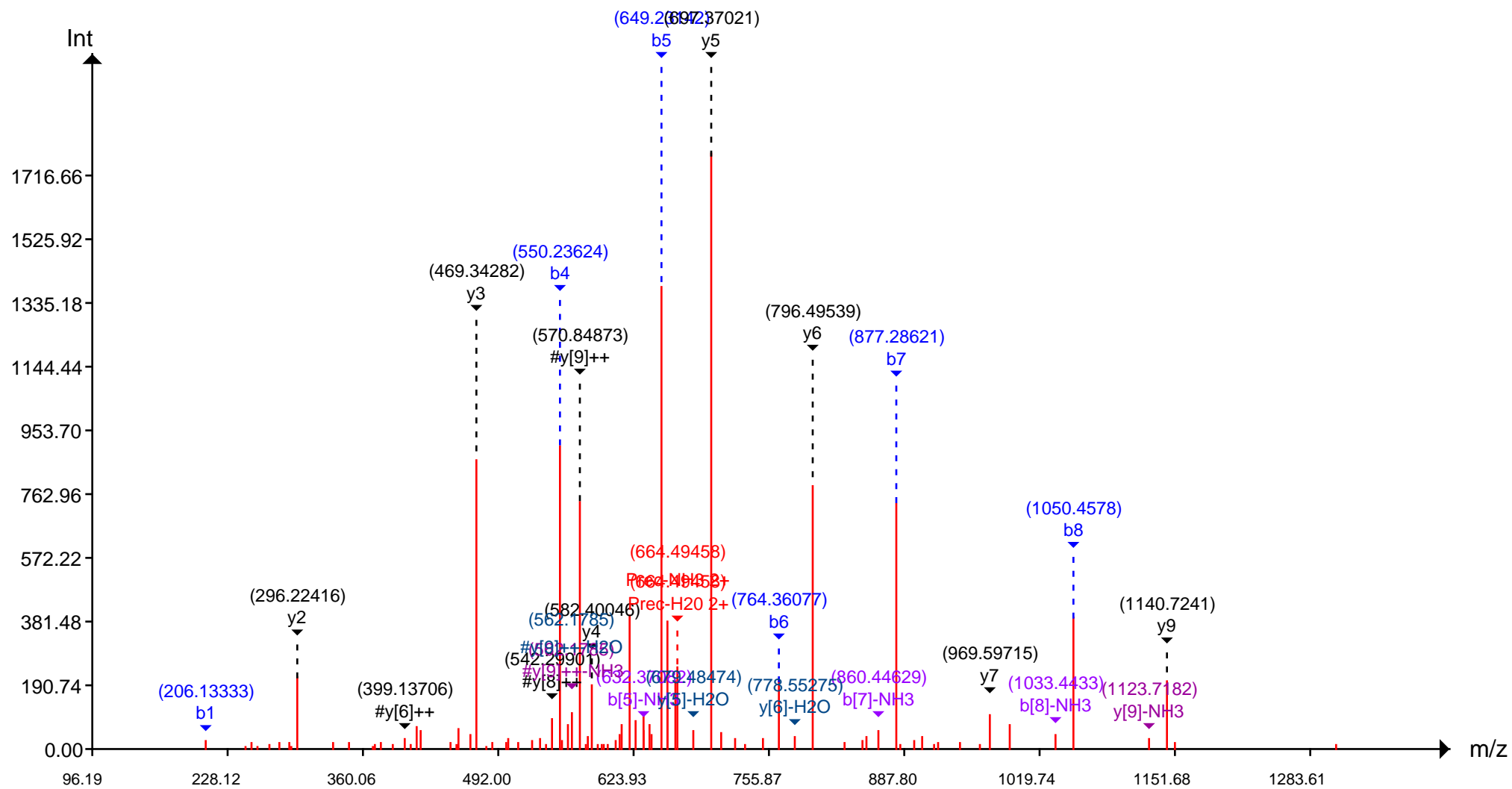
Info

Modified Sequence	AcD3-YLQETTLAANQPLSR<C13>-COOH	Ion score	76.19
Protein	O09159	Confidence level	99.00%
Precursor m/z	878.47506	Identity Threshold	47.83
Charge	2+	Mass error	0.001822



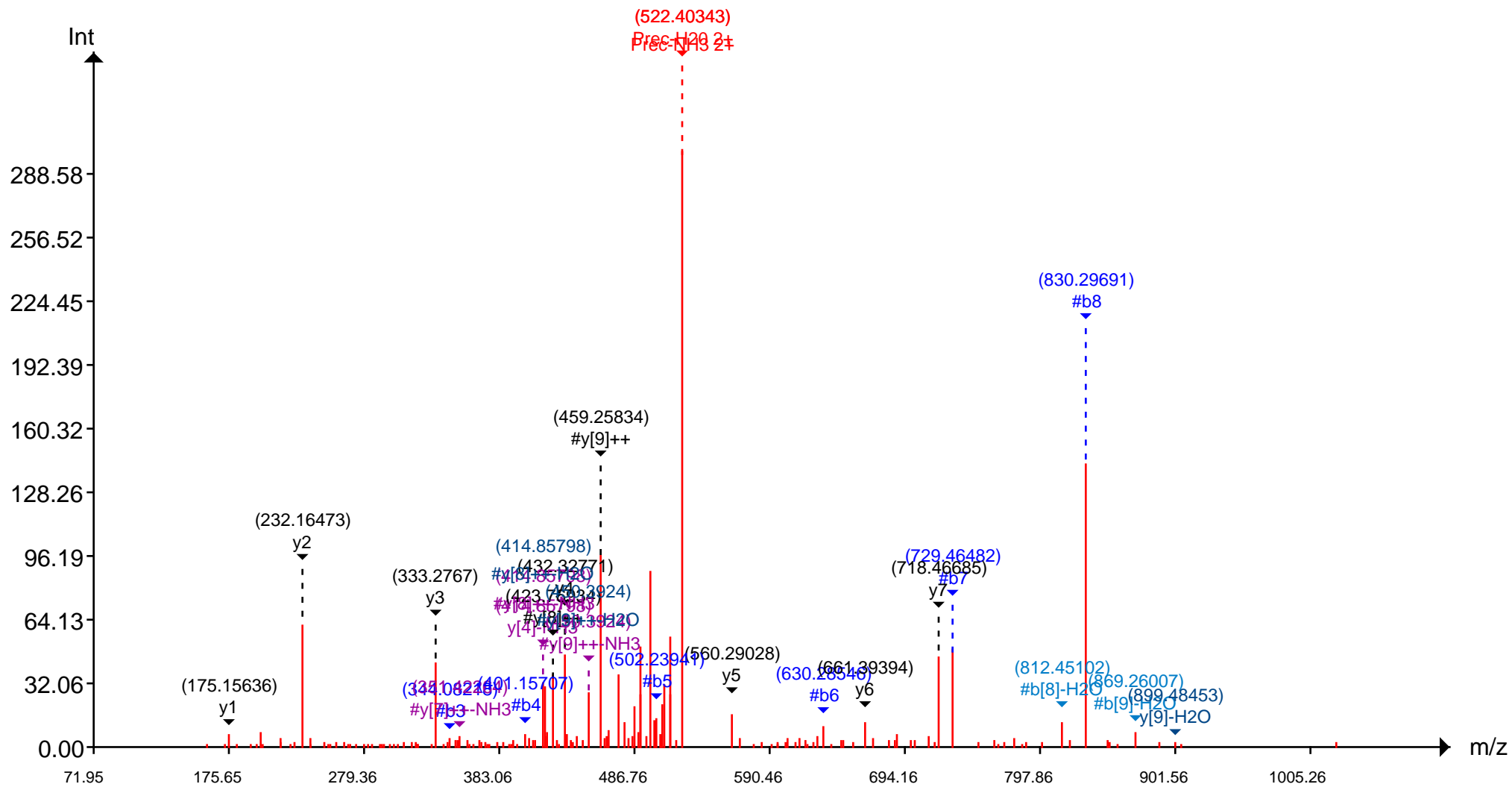
Info

Modified Sequence	AcD3-IK<AcD3K*>C<Cmm*>GETQK<AcD3K*>R<C13>-COOH	Ion score	47.82
Protein	Q9JK48	Confidence level	99.00%
Precursor m/z	630.85521	Identity Threshold	47.33
Charge	2+	Mass error	9.22E-4



Info

Modified Sequence	AcD3-C<Cmm*>GNK<AcD3K*>VDIK<AcD3K*>DR<C13>-COOH	Ion score	53.35
Protein	P62827	Confidence level	99.00%
Precursor m/z	673.36228	Identity Threshold	47.97
Charge	2+	Mass error	-0.001326



Info

Modified Sequence AcD3-VAQGTQVTGR-COOH

Protein P50580

Precursor m/z 531.2924

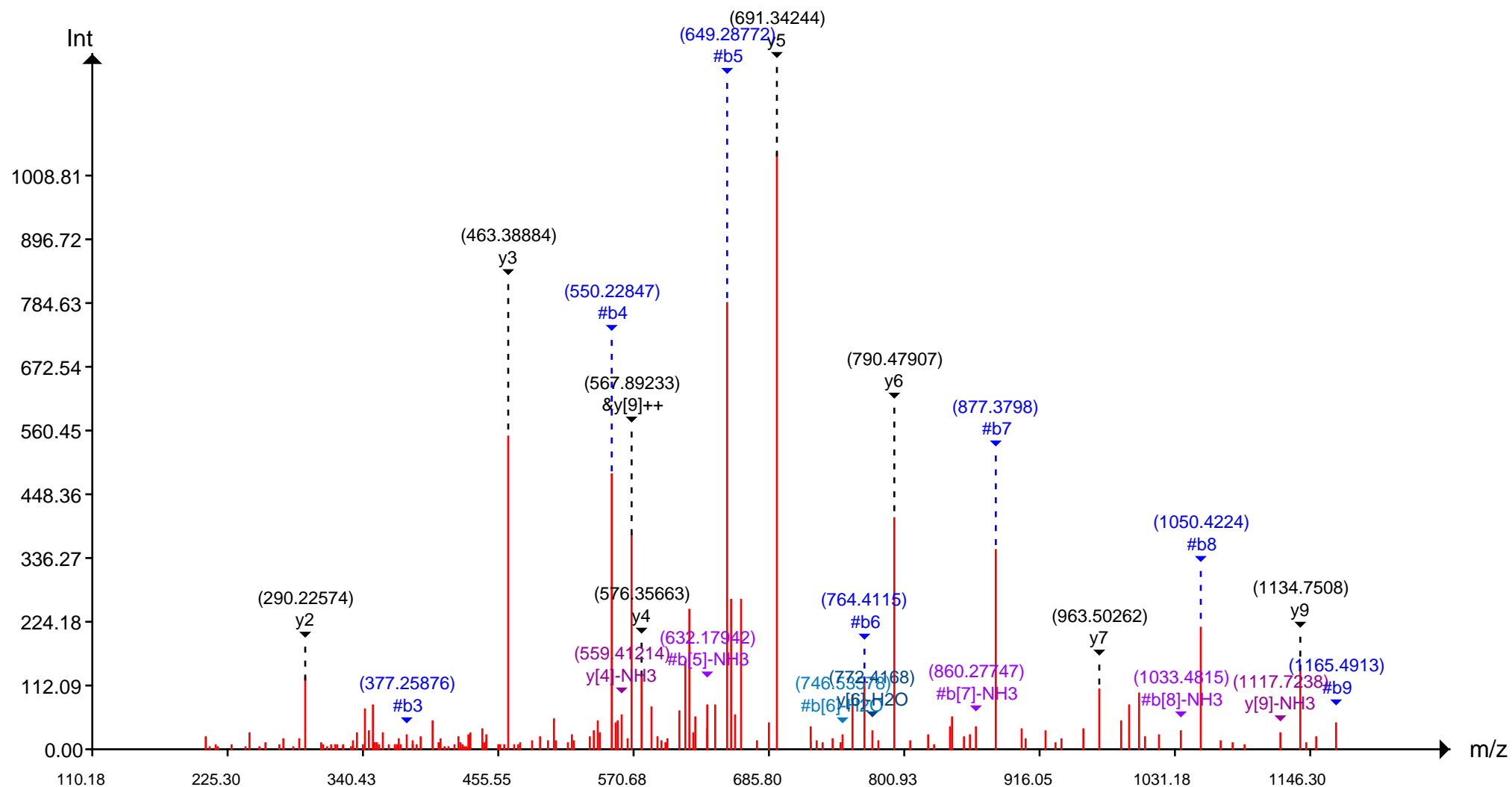
Charge 2+

Ion score 56.79

Confidence level 99.00%

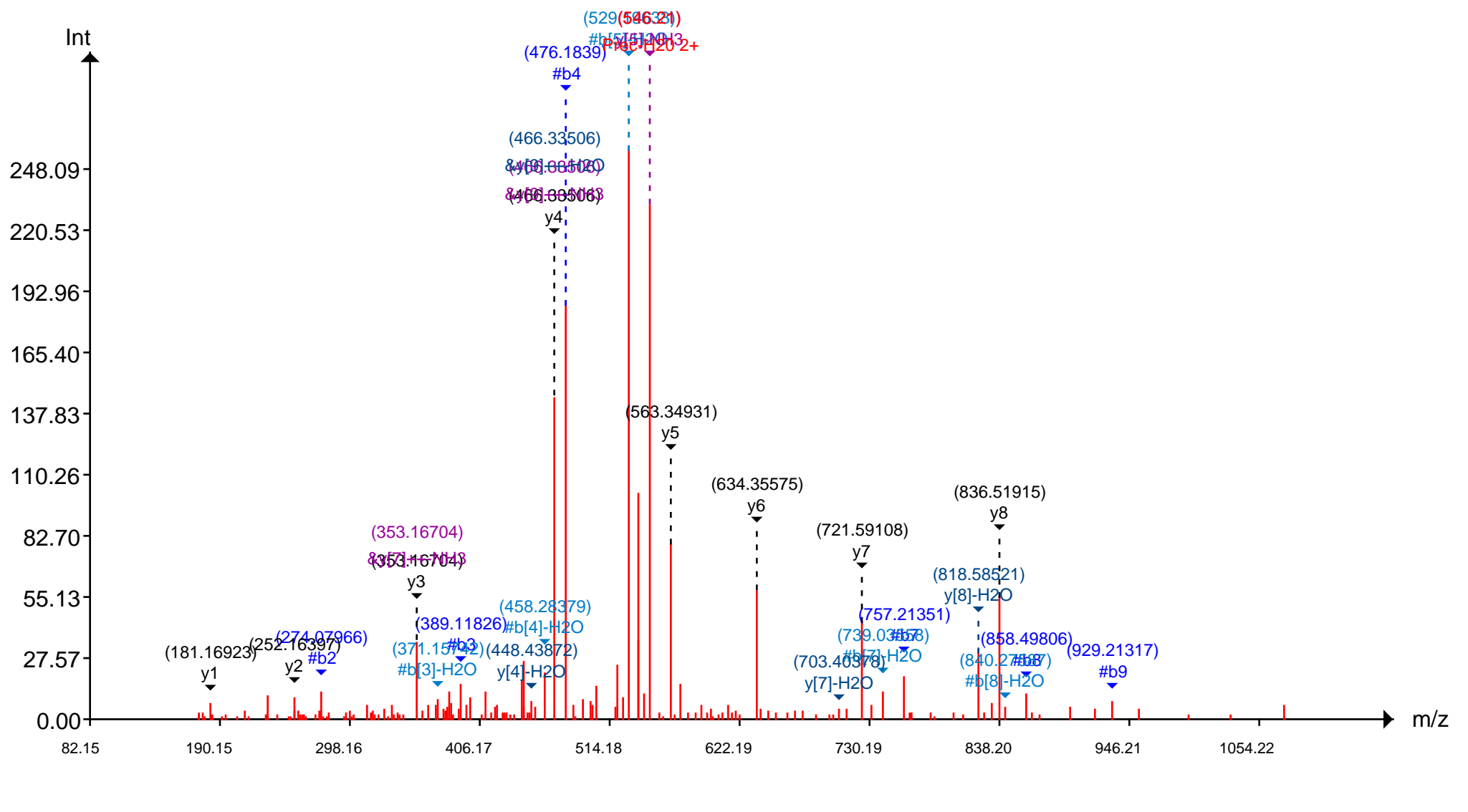
Identity Threshold 47.85

Mass error -2.17E-4

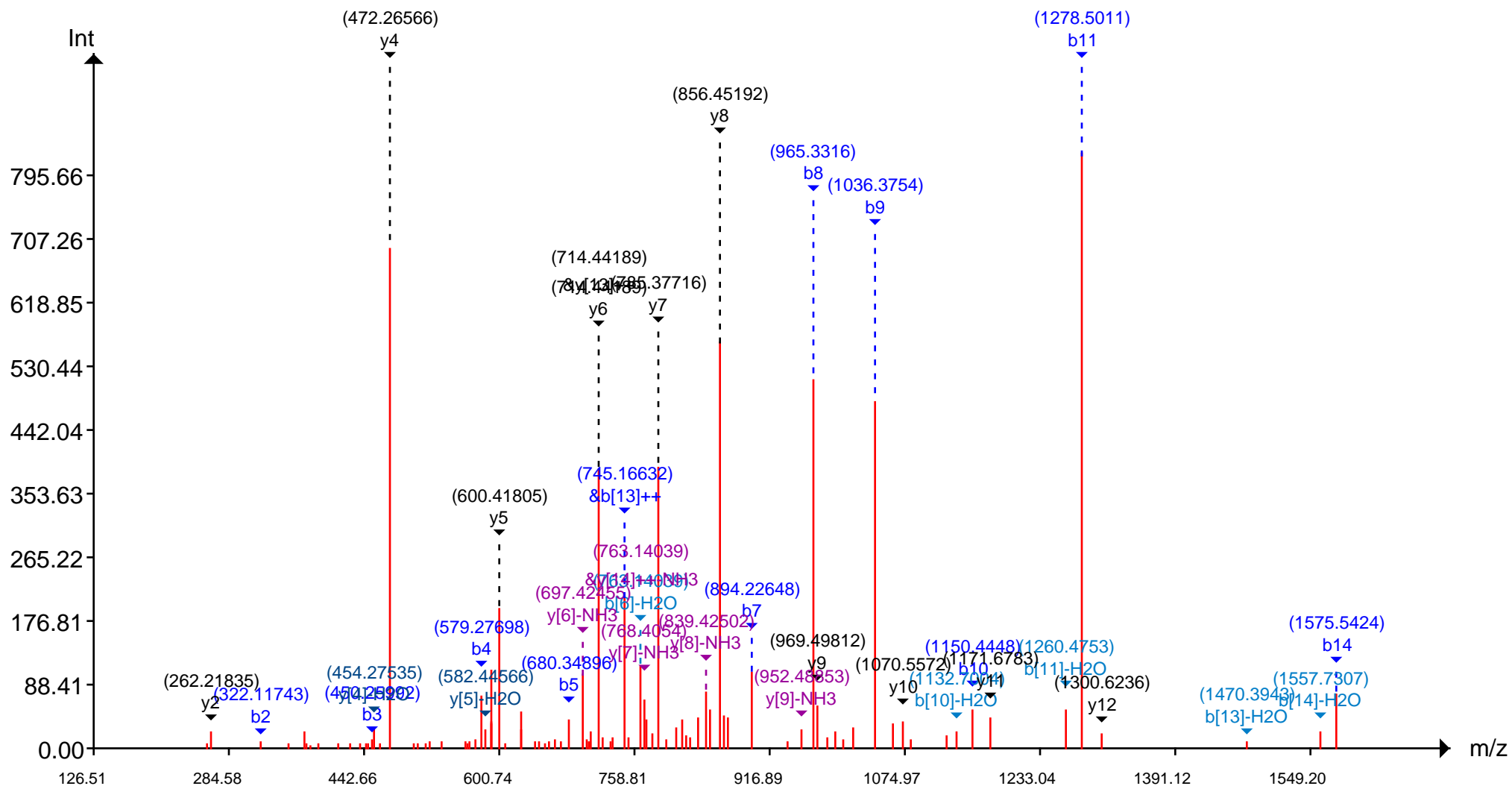


Info

Modified Sequence	AcD3-C<Cmm*>GNK<AcD3K*>VDIK<AcD3K*>DR-COOH	Ion score	53.09
Protein	P62827	Confidence level	99.00%
Precursor m/z	670.35291	Identity Threshold	47.91
Charge	2+	Mass error	7.5E-5

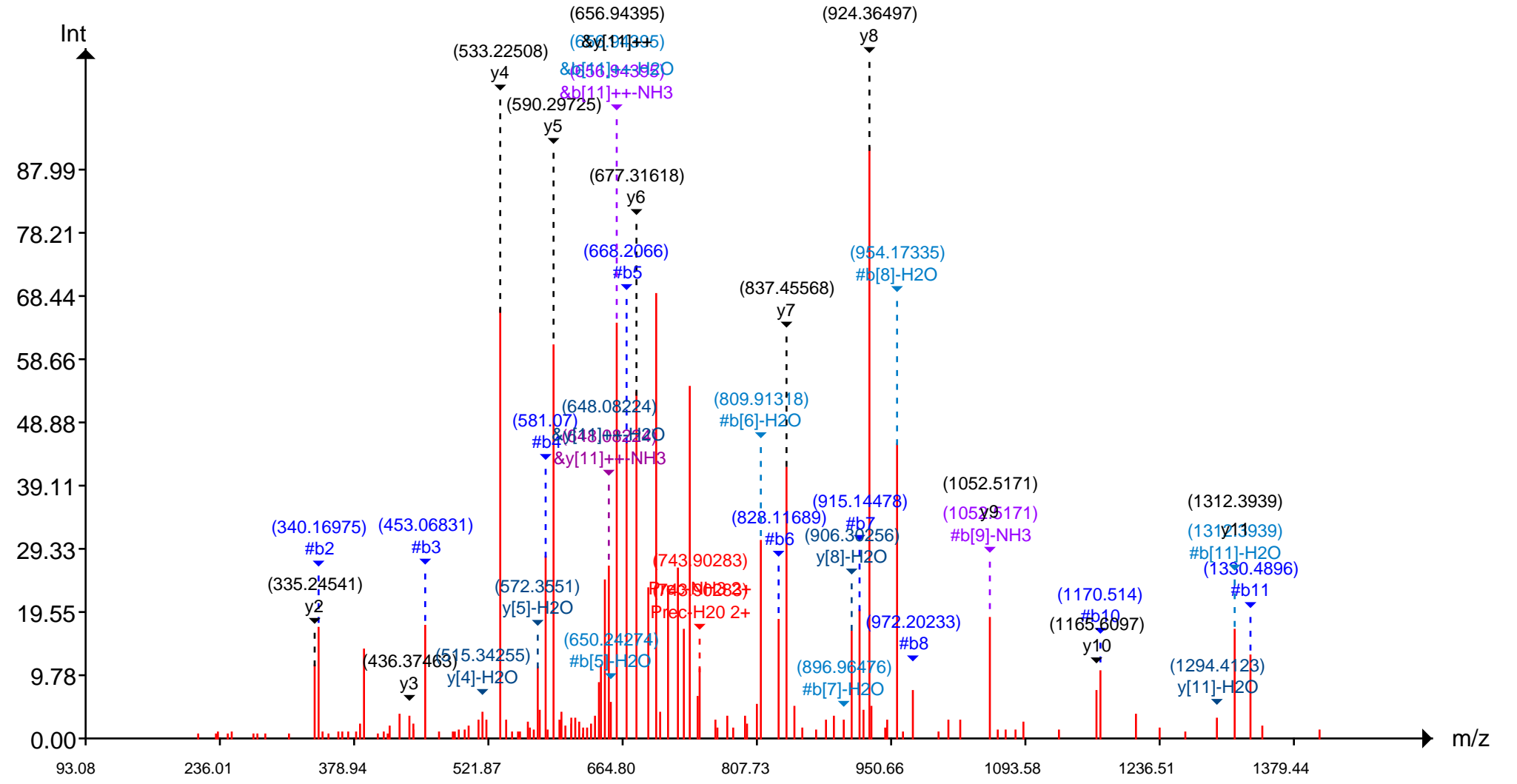


Info			
Modified Sequence	AcD3-DIDSAPITAR<C13>-COOH	Ion score	56.02
Protein	P52480	Confidence level	99.00%
Precursor m/z	555.30224	Identity Threshold	47.79
Charge	2+	Mass error	2.3E-5



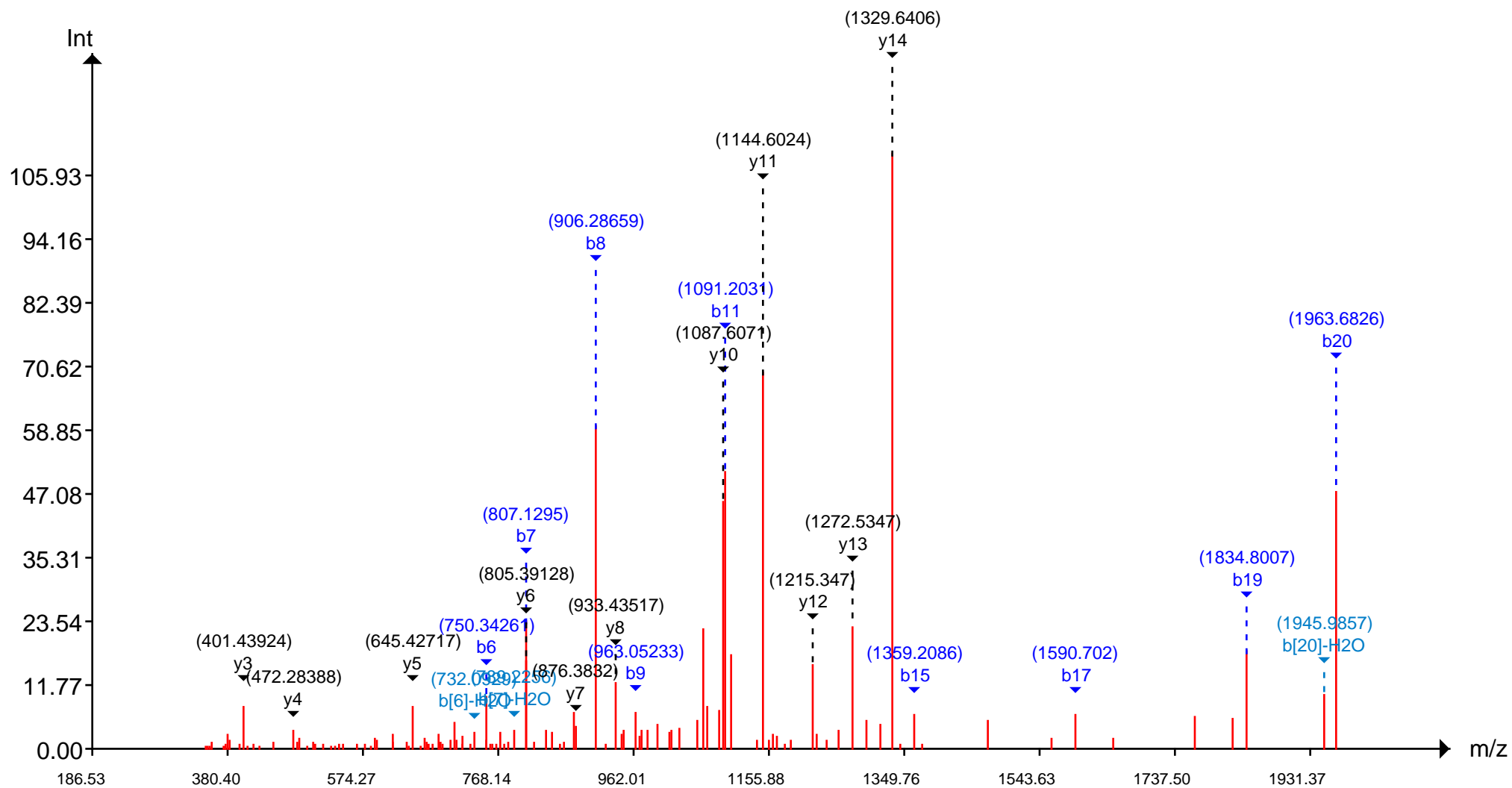
Info

Modified Sequence	AcD3-YLQETTLAANQPLSR-COOH	Ion score	88.85
Protein	O09159	Confidence level	99.00%
Precursor m/z	875.46499	Identity Threshold	48.12
Charge	2+	Mass error	0.001823



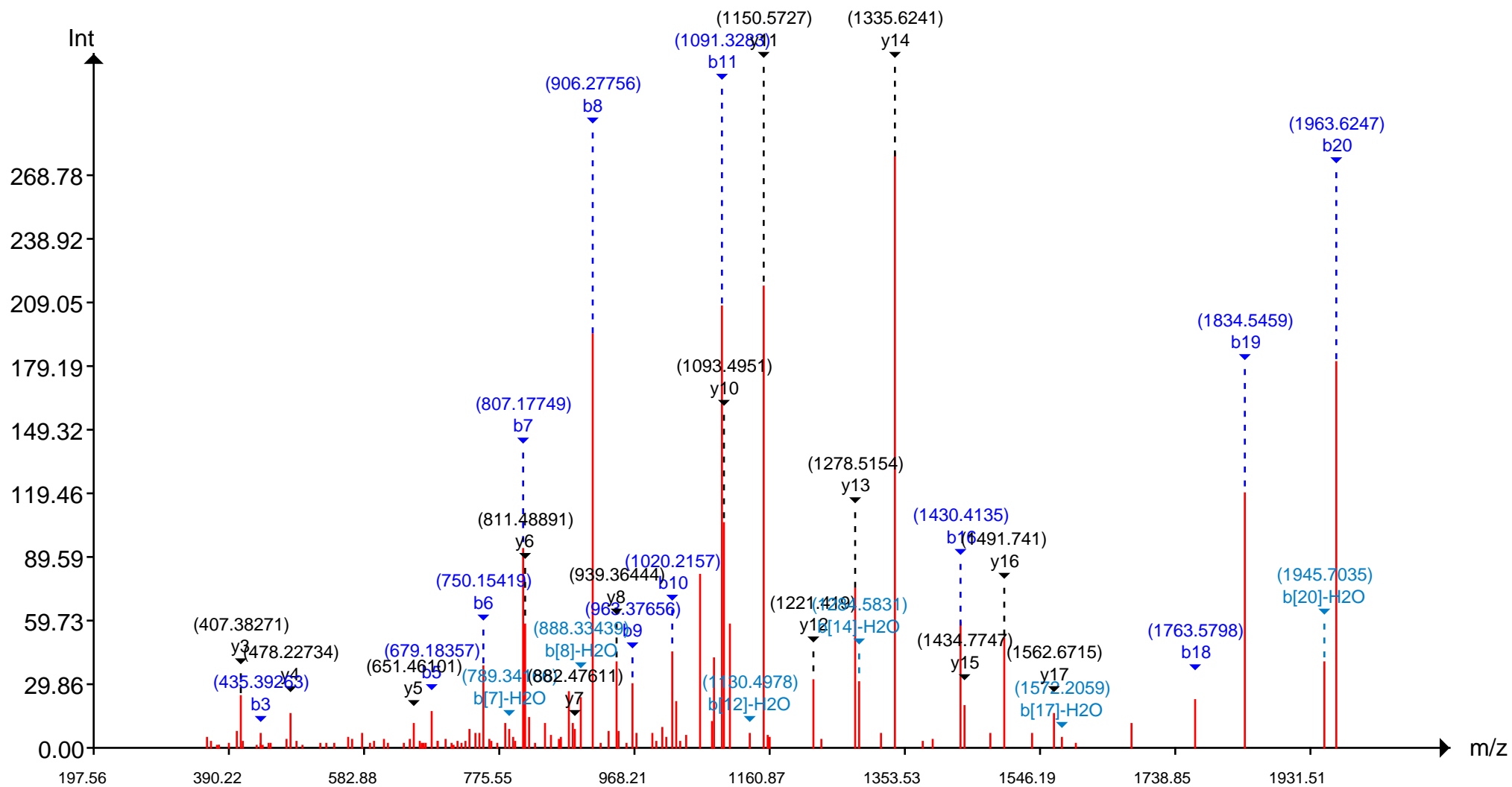
Info

Modified Sequence	AcD3-M<Mox*>M<Mox*>LQSC<Cmm*>SGPTC<Cmm*>R-COOH	Ion score	76.65
Protein	Q8BFR4	Confidence level	99.00%
Precursor m/z	752.80908	Identity Threshold	38.18
Charge	2+	Mass error	0.001832



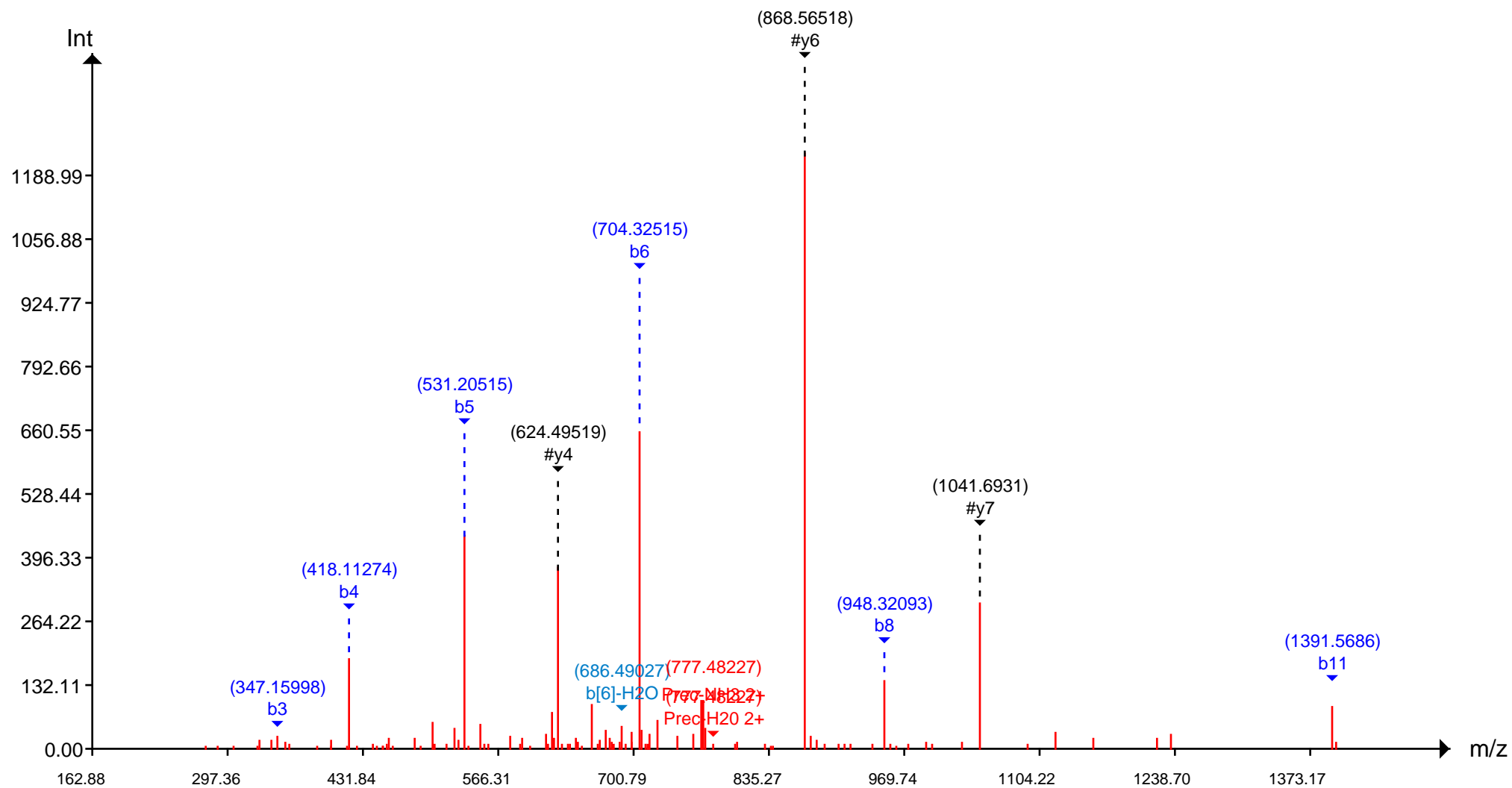
Info

Modified Sequence	AcD3-M<Mox*>LEDEAGVGGAGPGGAC<Cmm*>K<AcD3K*>AEPR-COOH	Ion score	70.72
Protein	Q00PI9	Confidence level	99.00%
Precursor m/z	1118.0197	Identity Threshold	45.11
Charge	2+	Mass error	0.003409



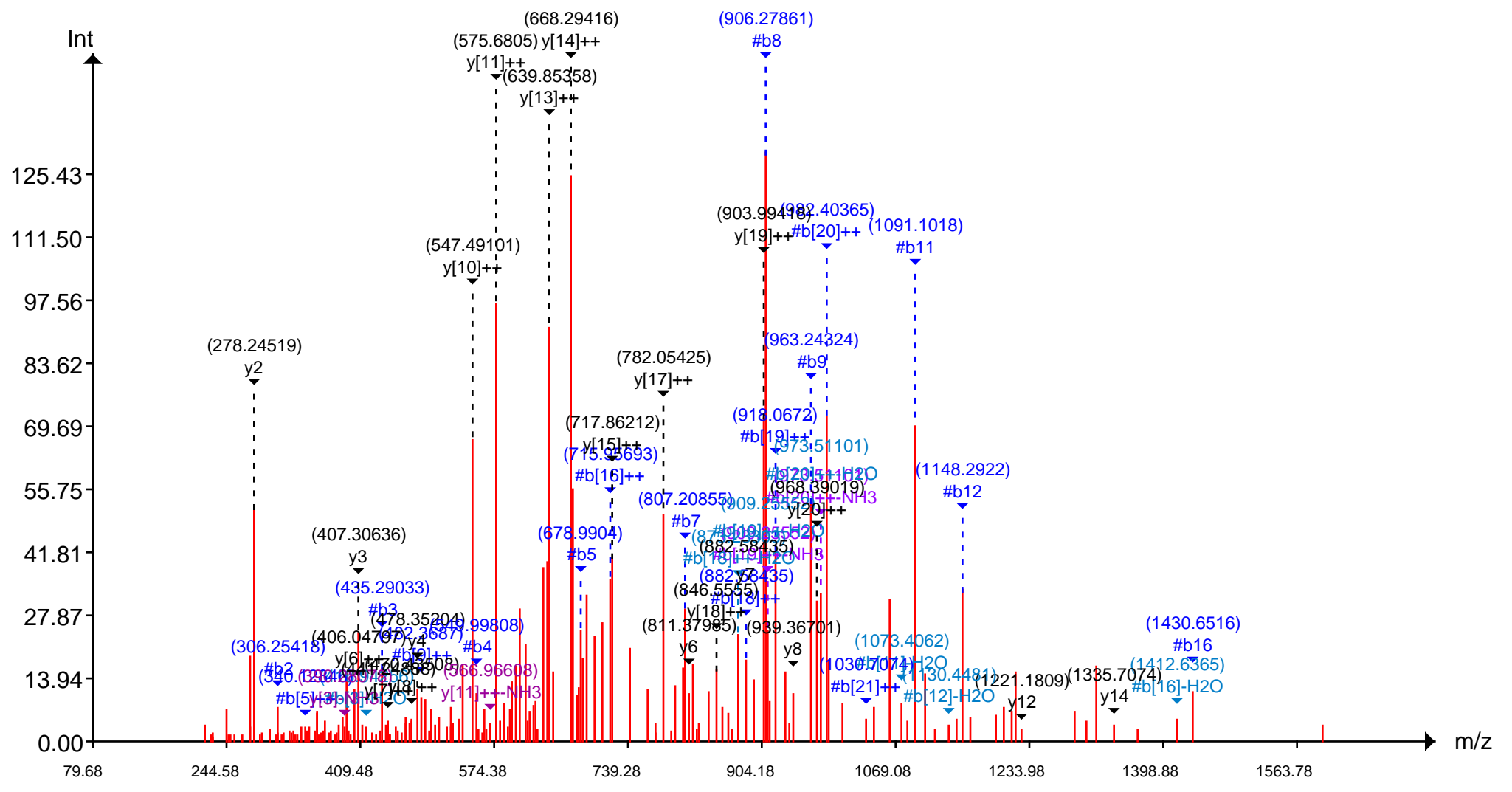
Info

Modified Sequence	AcD3-M<Mox*>LEDEAGVGGAGPGGAC<Cmm*>K<AcD3K*>AEPR<C13>-COOH	Ion score	104.75
Protein	Q00PI9	Confidence level	99.00%
Precursor m/z	1121.0287	Identity Threshold	45.91
Charge	2+	Mass error	0.001268

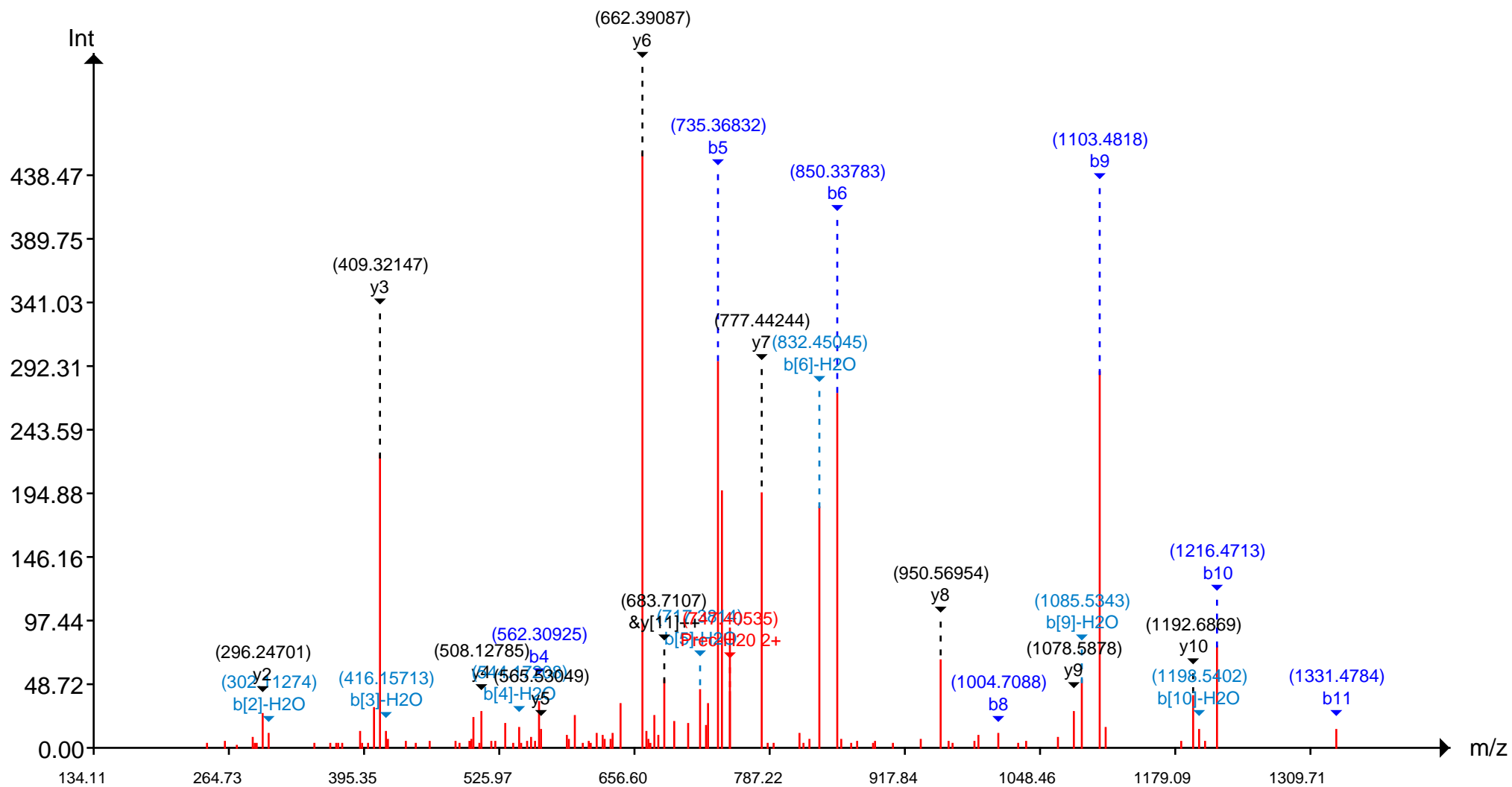


Info

Modified Sequence	AcD3-AATEAIK<AcD3K*>PFPK<AcD3K*>K<AcD3K*>R<C13>-COOH	Ion score	49.23
Protein	Q6PGL7	Confidence level	99.00%
Precursor m/z	786.48588	Identity Threshold	40.78
Charge	2+	Mass error	4.36E-4

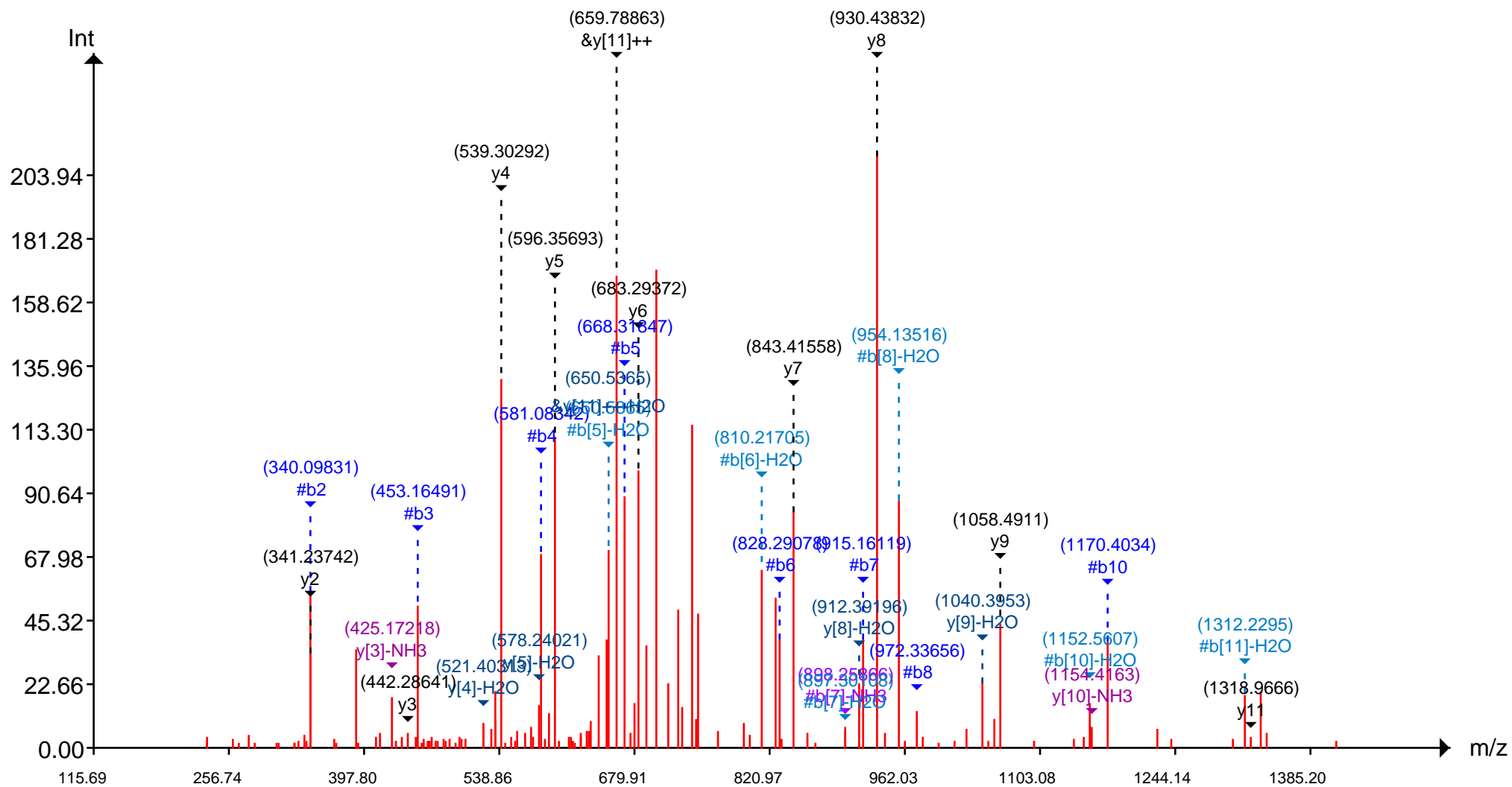


Info			
Modified Sequence	AcD3-M<Mox*>LEDEAGVGGAGPGGAC<Cmm*>K<AcD3K*>AEPR<C13>-COOH	Ion score	49.43
Protein	Q00PI9	Confidence level	99.00%
Precursor m/z	747.68813	Identity Threshold	45.85
Charge	3+	Mass error	9.82E-4



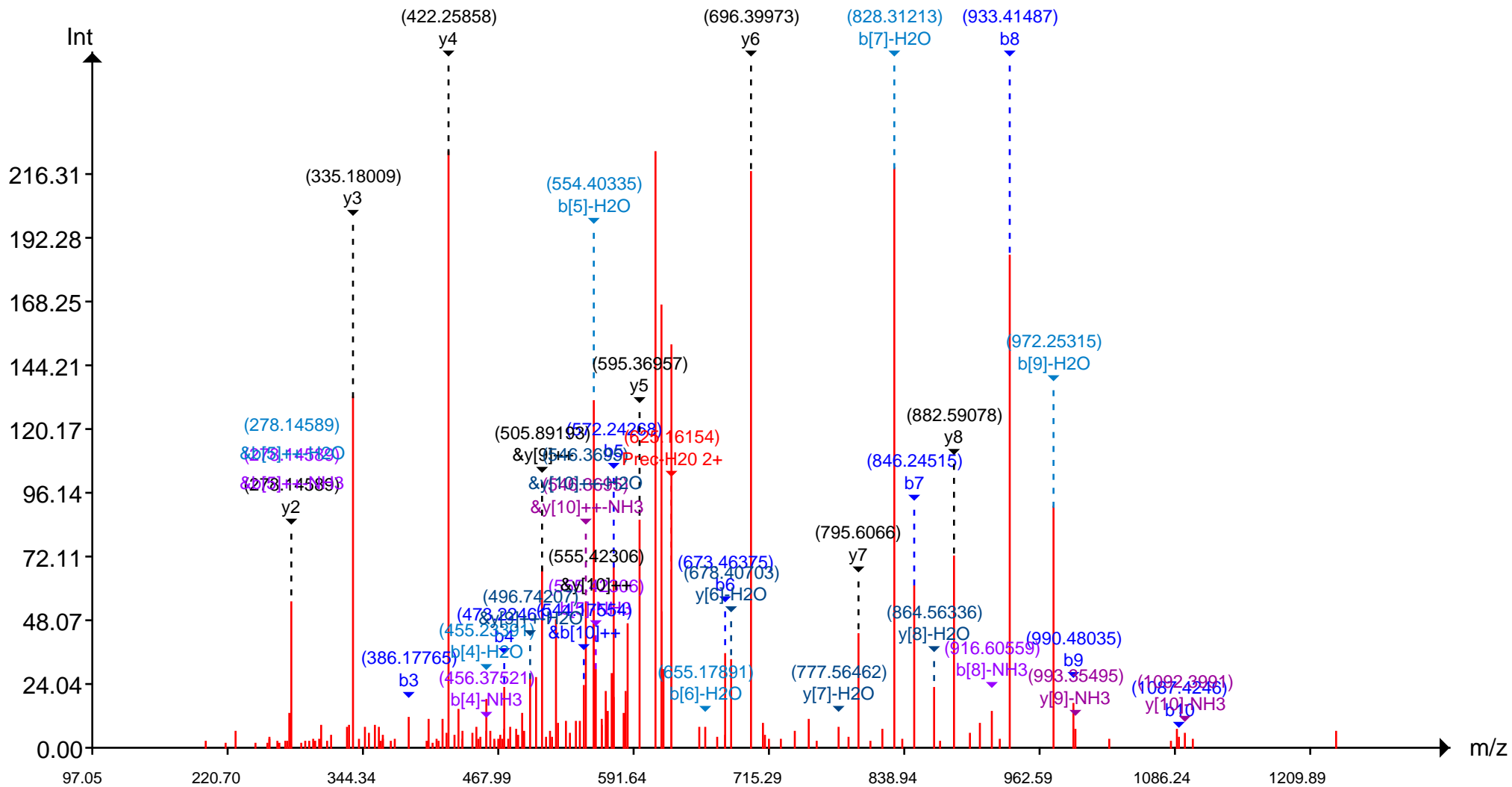
Info

Modified Sequence	AcD3-TK<AcD3K*>NQK<AcD3K*>DPGVLDK<C13>-COOH	Ion score	63.64
Protein	P50543	Confidence level	99.00%
Precursor m/z	756.42784	Identity Threshold	46.81
Charge	2+	Mass error	0.001422



Info

Modified Sequence	AcD3-M<Mox*>M<Mox*>LQSC<Cmm*>SGPTC<Cmm*>R<C13>-COOH	Ion score	55.23
Protein	Q8BFR4	Confidence level	99.00%
Precursor m/z	755.81897	Identity Threshold	40.45
Charge	2+	Mass error	0.00147



Info

Modified Sequence AcD3-LVQSVTK<AcD3K*>SGPR<C13>-COOH

Ion score 55.62

Protein Q6P9Q4

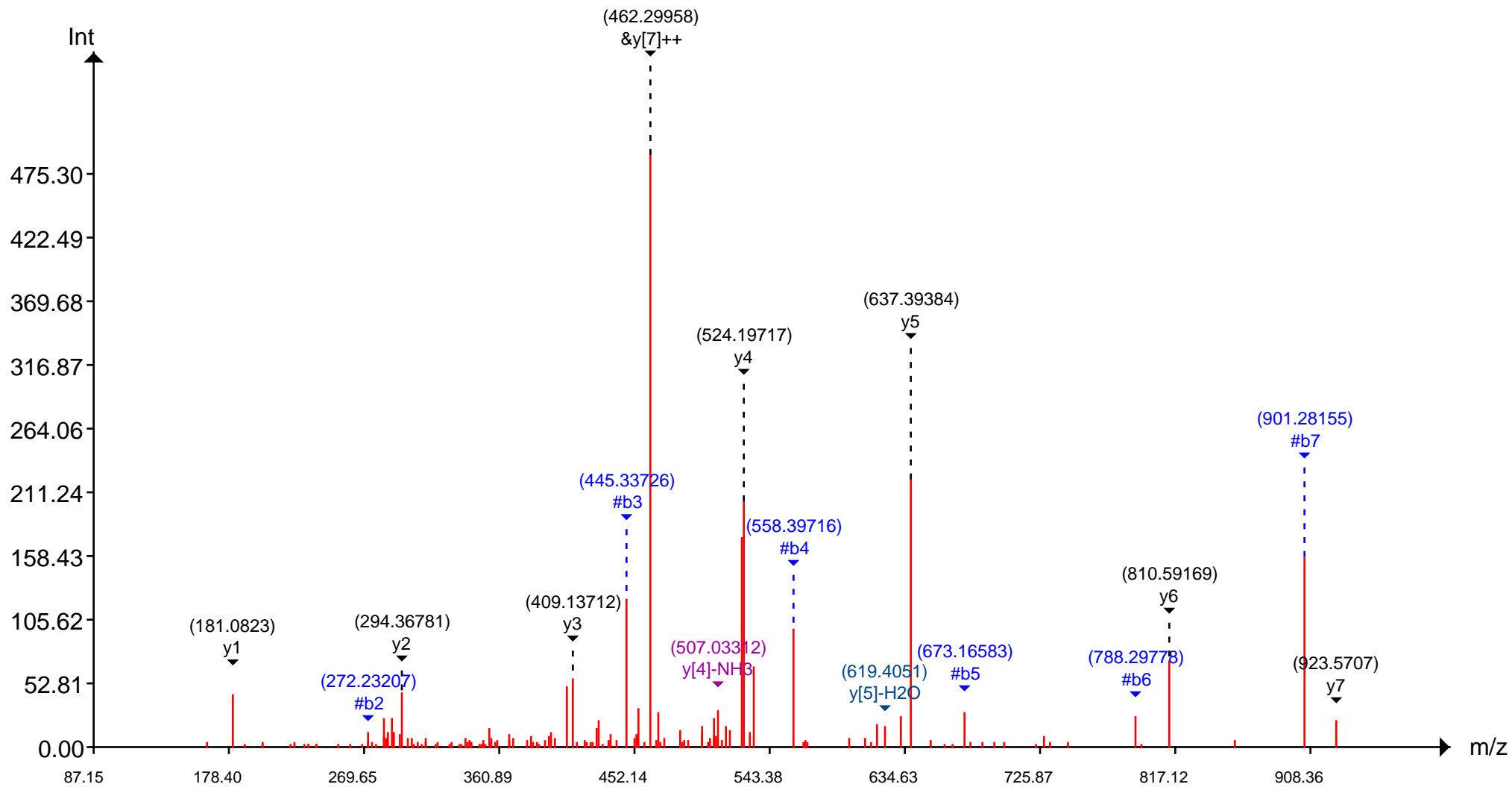
Confidence level 99.00%

Precursor m/z 634.38314

Identity Threshold 44.35

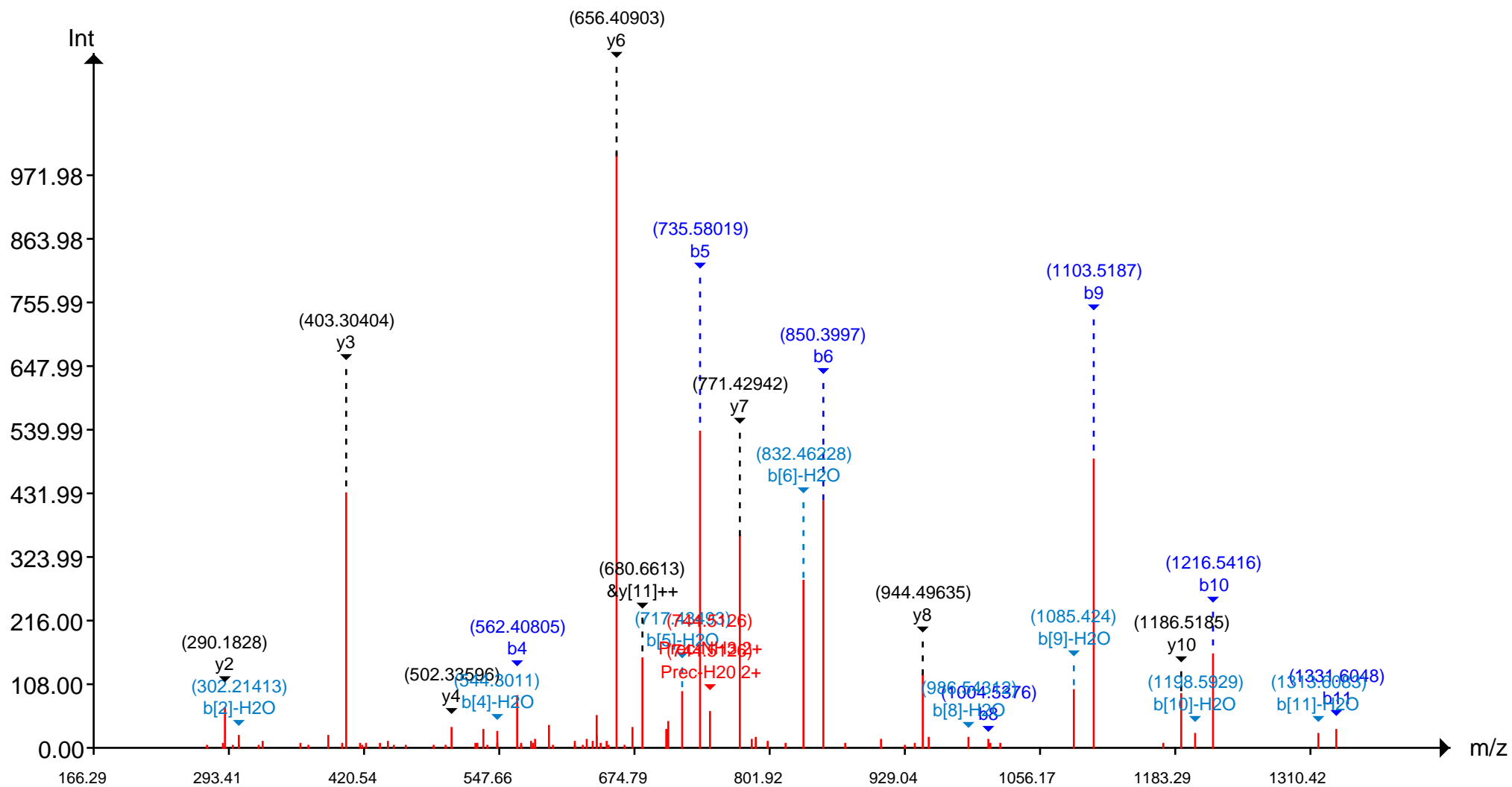
Charge 2+

Mass error 7.21E-4



Info

Modified Sequence	AcD3-ILK<AcD3K*>IDDIR<C13>-COOH	Ion score	50.53
Protein	P80316	Confidence level	99.00%
Precursor m/z	541.34499	Identity Threshold	40.84
Charge	2+	Mass error	-2.62E-4



Info

Modified Sequence AcD3-TK<AcD3K*>NQK<AcD3K*>DPGVLDLDR-COOH

Ion score 76.6

Protein P50543

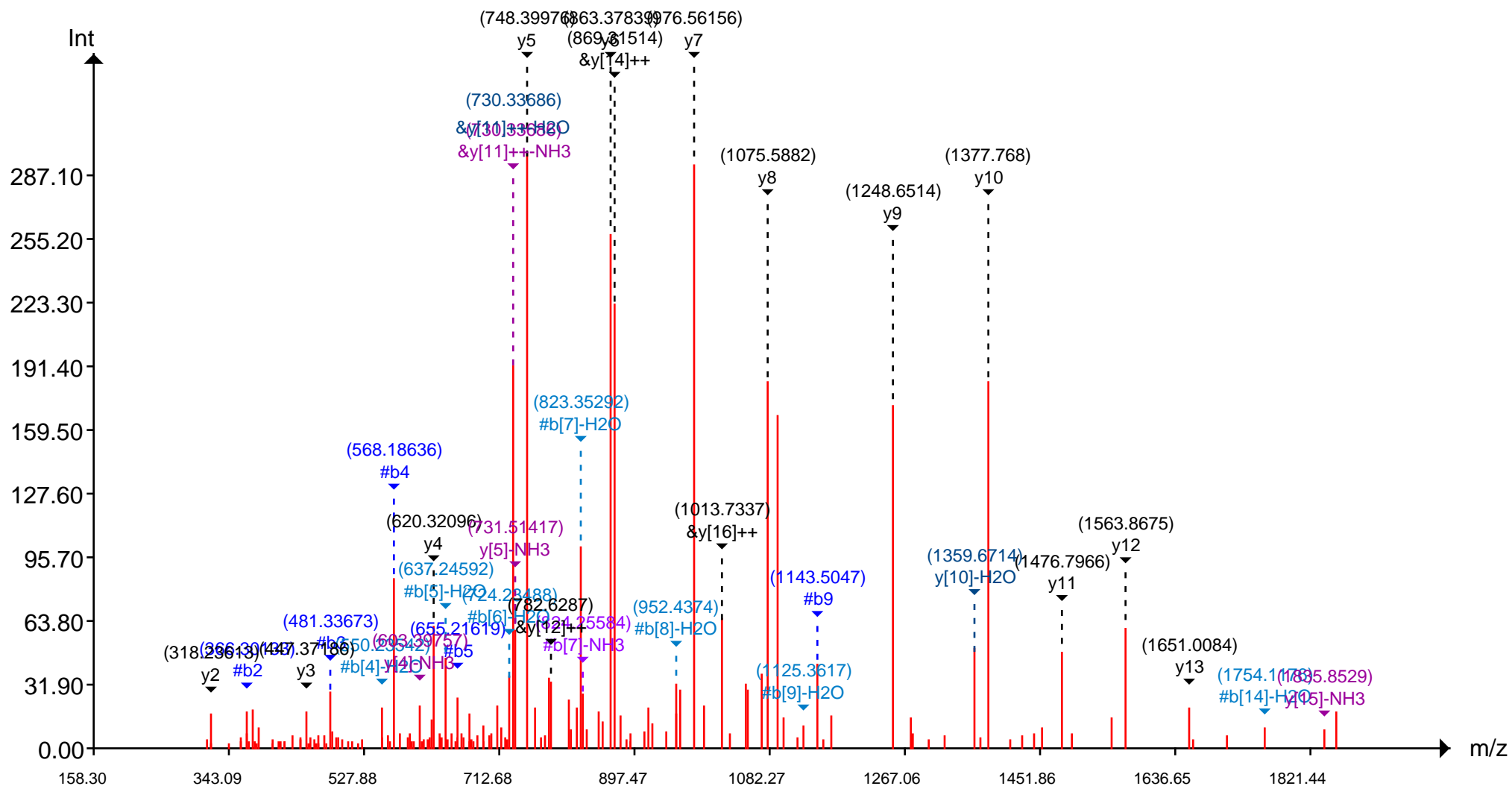
Confidence level 99.00%

Precursor m/z 753.41761

Identity Threshold 47.50

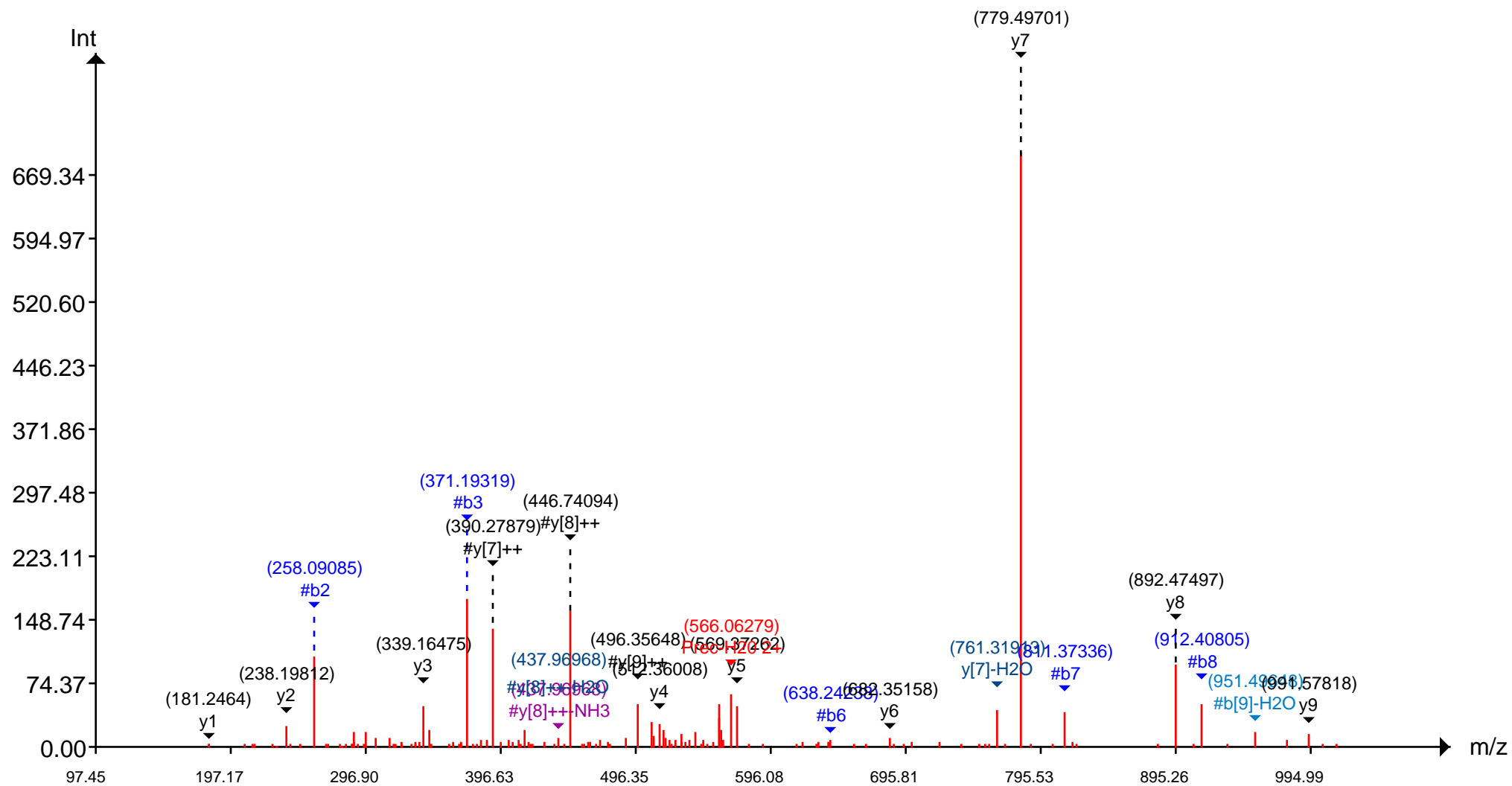
Charge 2+

Mass error 0.001103



Info

Modified Sequence	AcD3-FK<AcD3K*>DSSSVEK<AcD3K*>VLDQK<AcD3K*>EHR<C13>-COOH	Ion score	89.35
Protein	Q99020	Confidence level	99.00%
Precursor m/z	1109.5967	Identity Threshold	47.54
Charge	2+	Mass error	0.002609



Info

Modified Sequence AcD3-LVLPLGK<AcD3K*>TGR<C13>-COOH

Ion score 54.03

Protein O89053

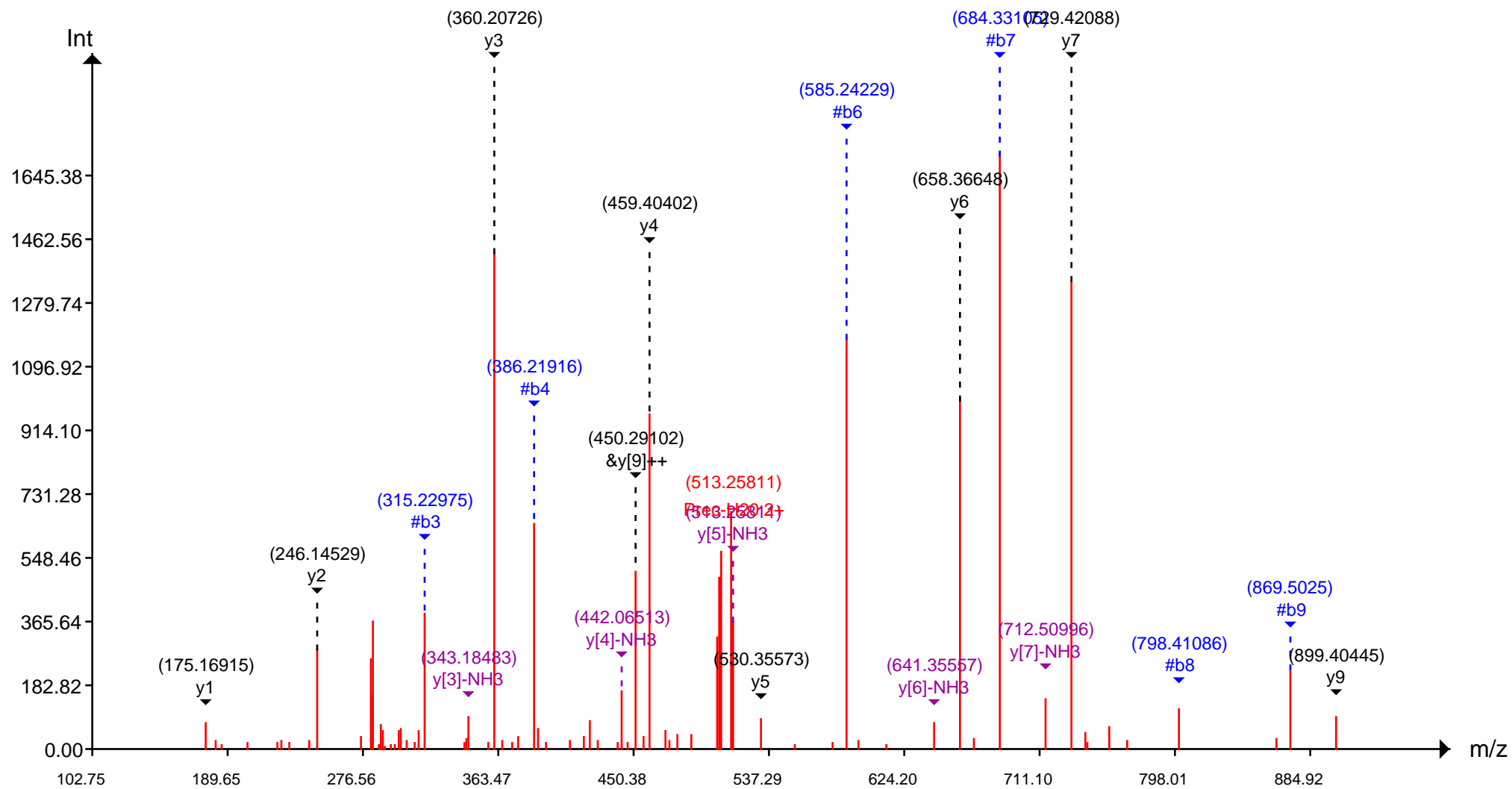
Confidence level 99.00%

Precursor m/z 575.38206

Identity Threshold 36.20

Charge 2+

Mass error 2.6E-5



Info

Modified Sequence AcD3-VGLAQAVNAR-COOH

Ion score 73.43

Protein Q02053

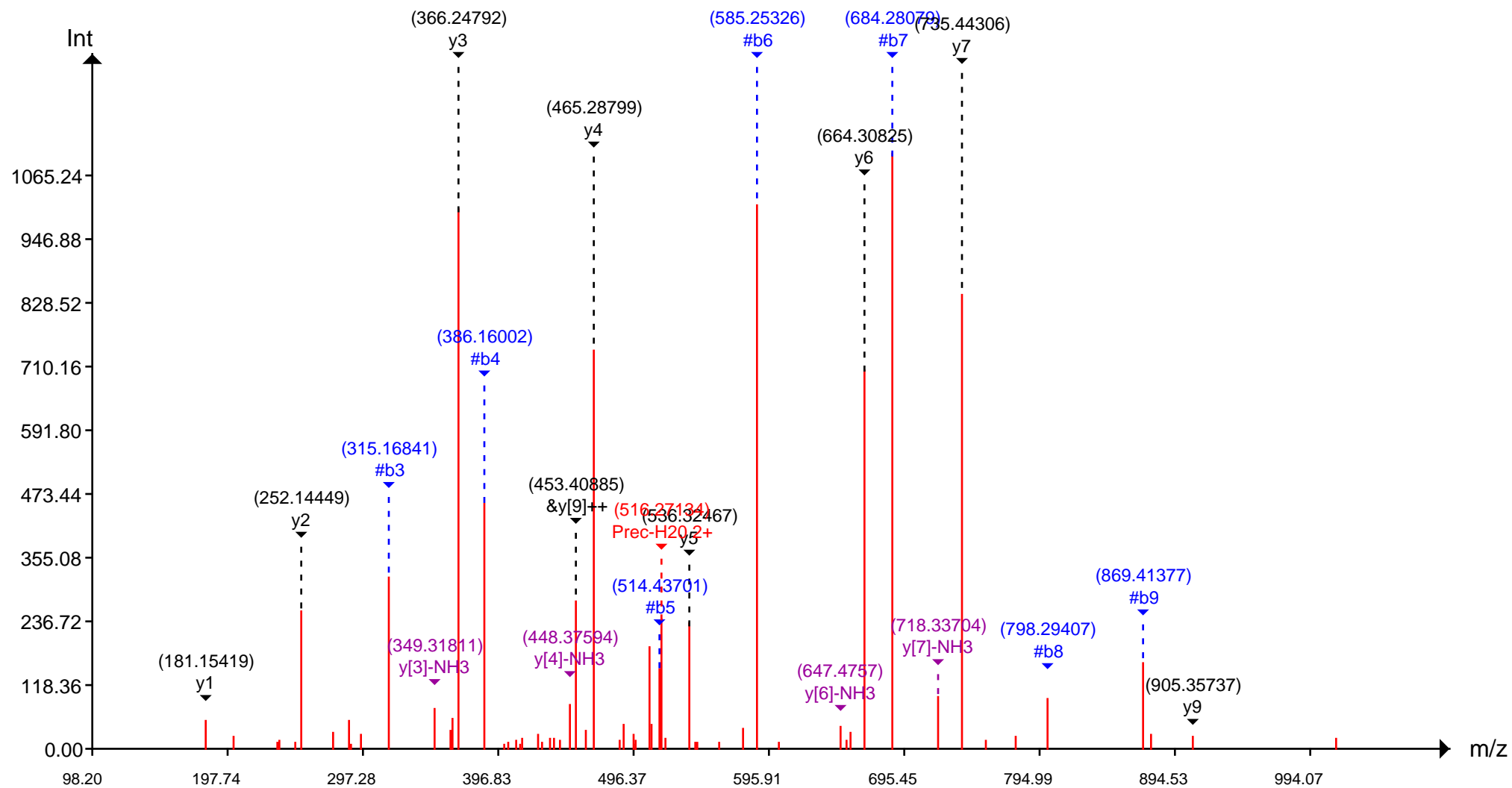
Confidence level 99.00%

Precursor m/z 522.30538

Identity Threshold 46.69

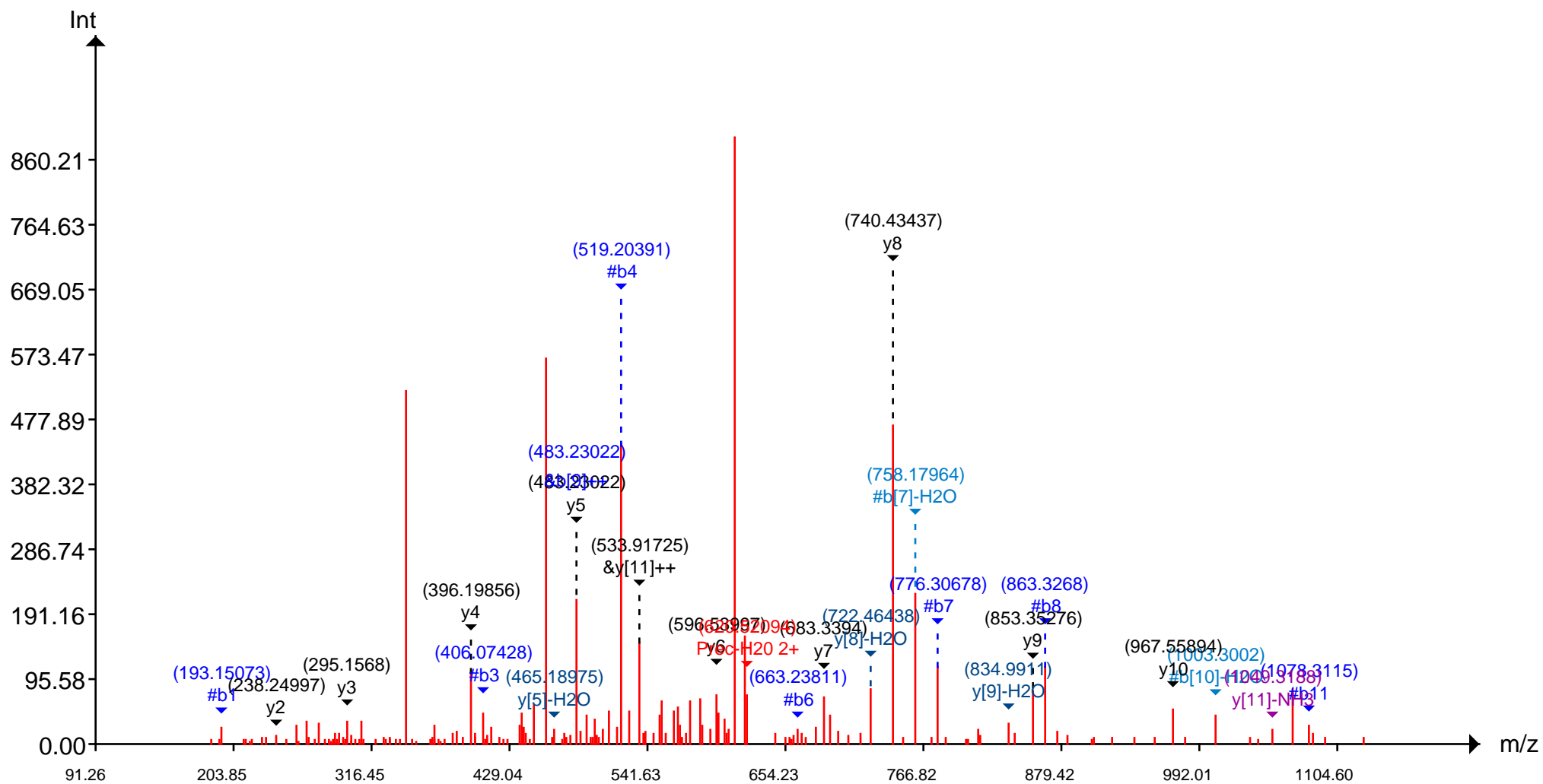
Charge 2+

Mass error -4.4E-5



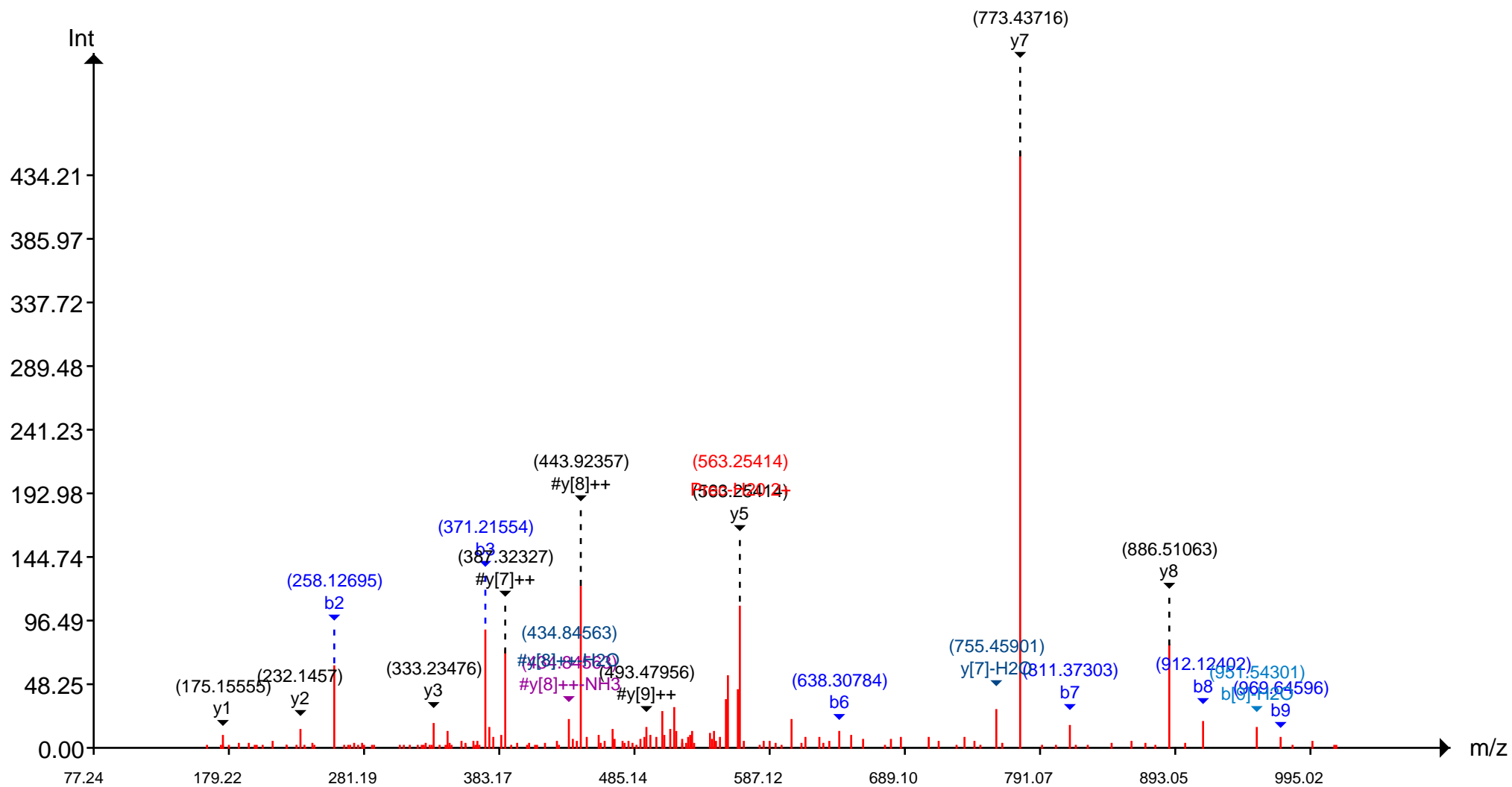
Info

Modified Sequence	AcD3-VGLAQAVNAR<C13>-COOH	Ion score	84.29
Protein	Q02053	Confidence level	99.00%
Precursor m/z	525.31539	Identity Threshold	45.47
Charge	2+	Mass error	-1.66E-4



Info

Modified Sequence	AcD3-M<Mox*>VNIGSLSTGGR<C13>-COOH	Ion score	49.5
Protein	Q9Z0N1, Q9Z0N2	Confidence level	99.00%
Precursor m/z	629.83377	Identity Threshold	47.91
Charge	2+	Mass error	7.82E-4



Info

Modified Sequence AcD3-LVLPLGK<AcD3K*>TGR-COOH

Ion score 43.4

Protein O89053

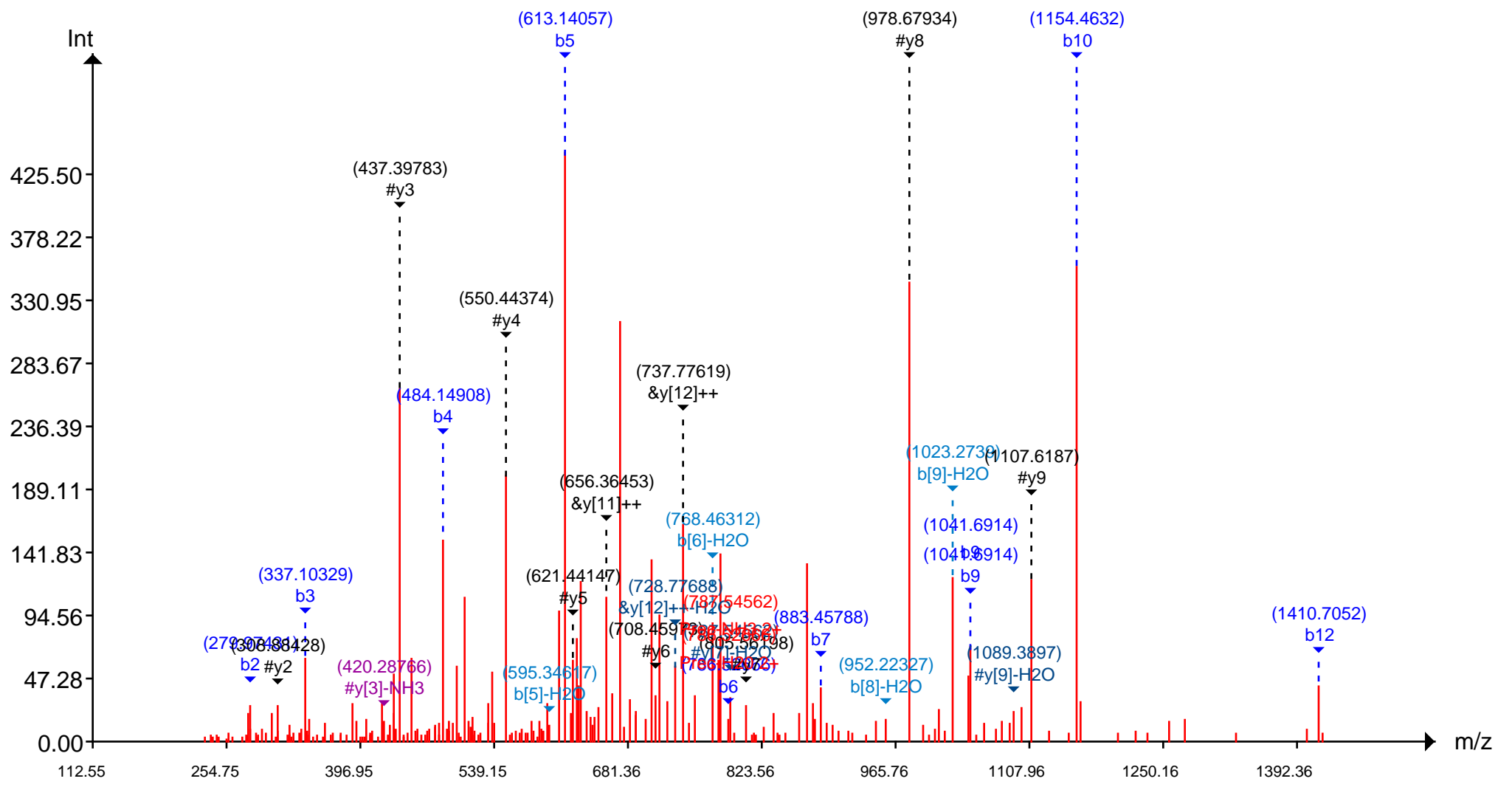
Confidence level 99.00%

Precursor m/z 572.3722

Identity Threshold 38.76

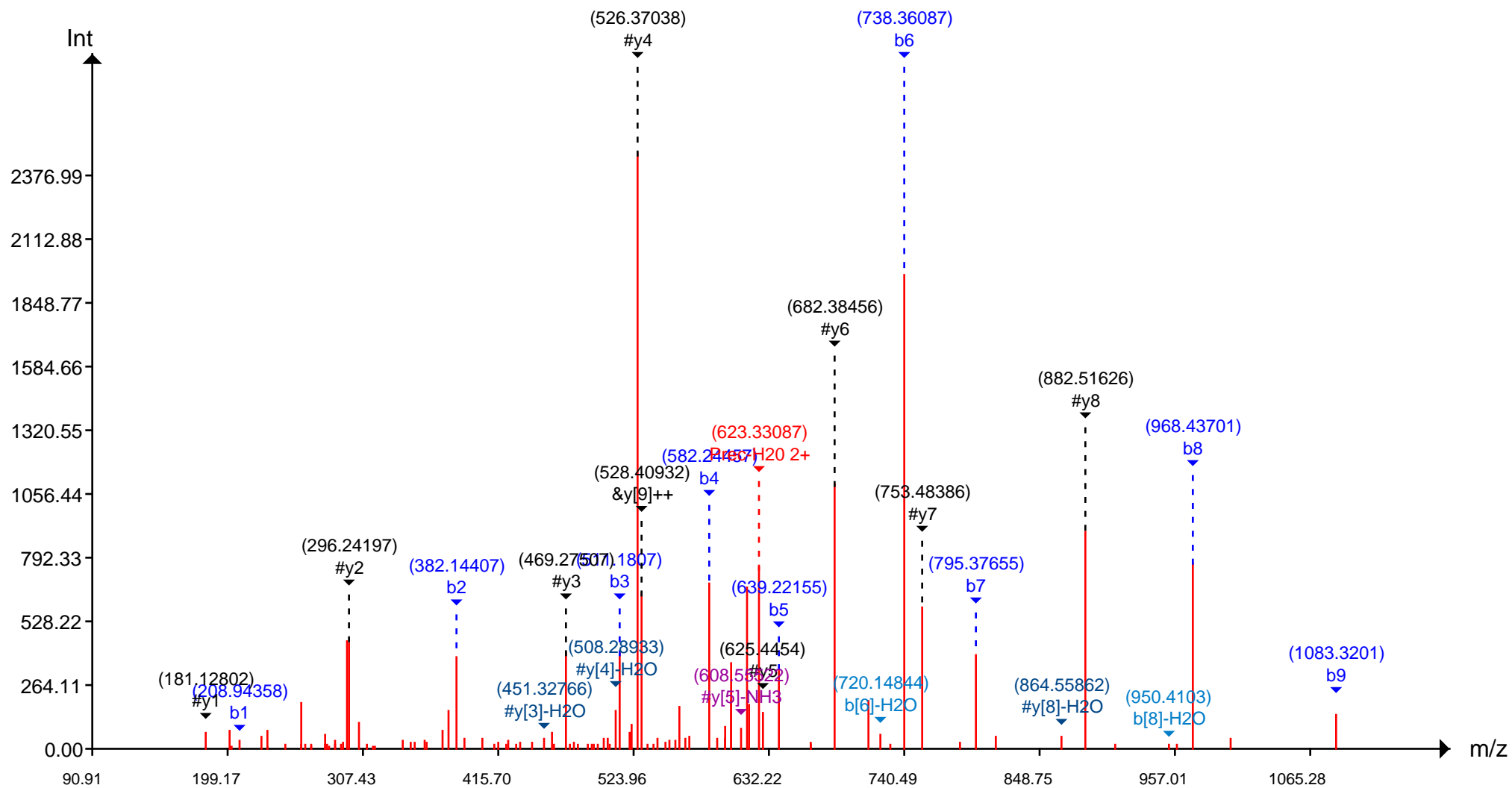
Charge 2+

Mass error 4.47E-4



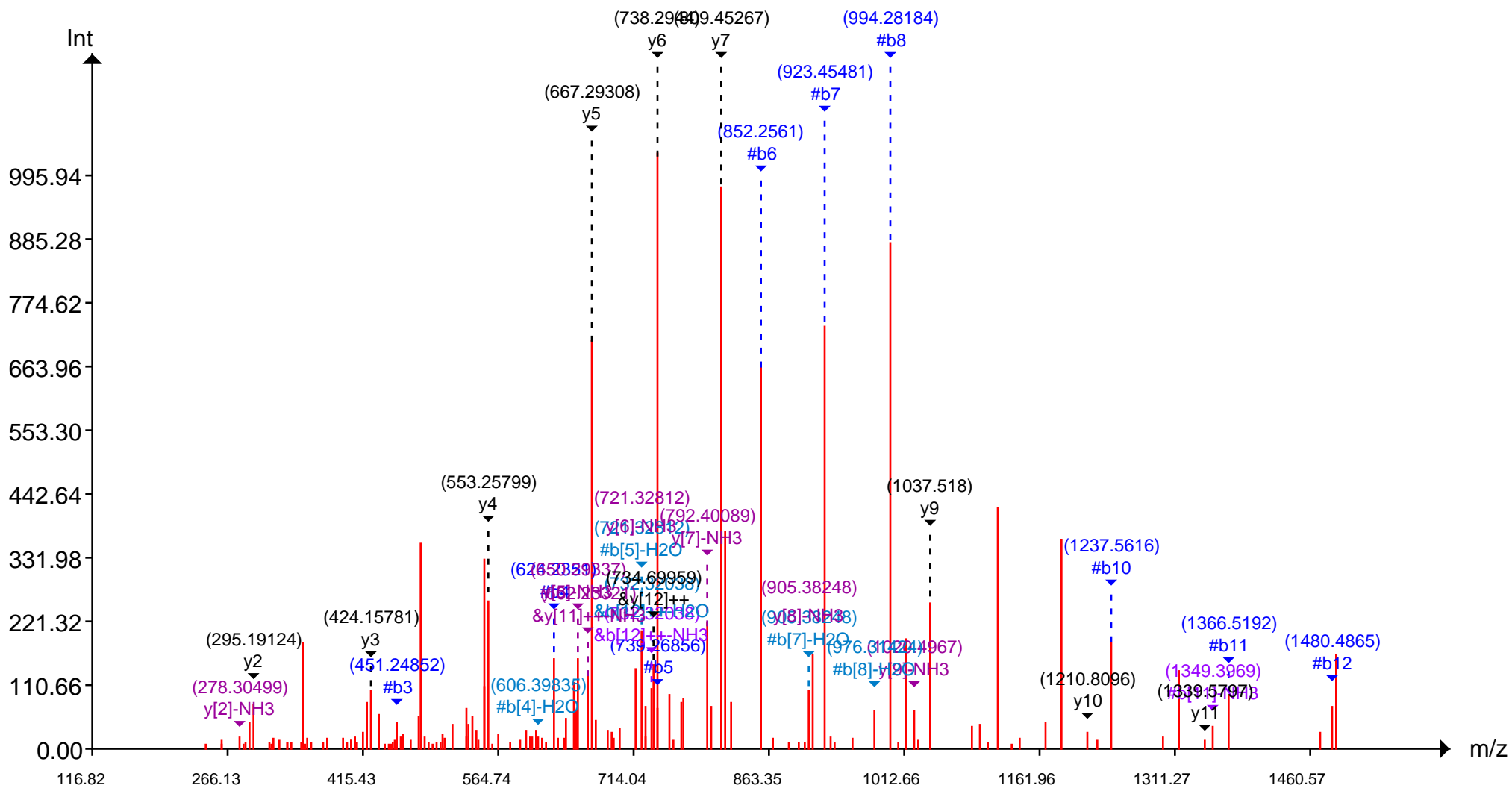
Info

Modified Sequence	AcD3-AYGF EK<AcD3K*>PSAIQQR<C13>-COOH	Ion score	50.95
Protein	P60843, P10630, Q91VC3	Confidence level	99.00%
Precursor m/z	795.92816	Identity Threshold	47.90
Charge	2+	Mass error	1.85E-4



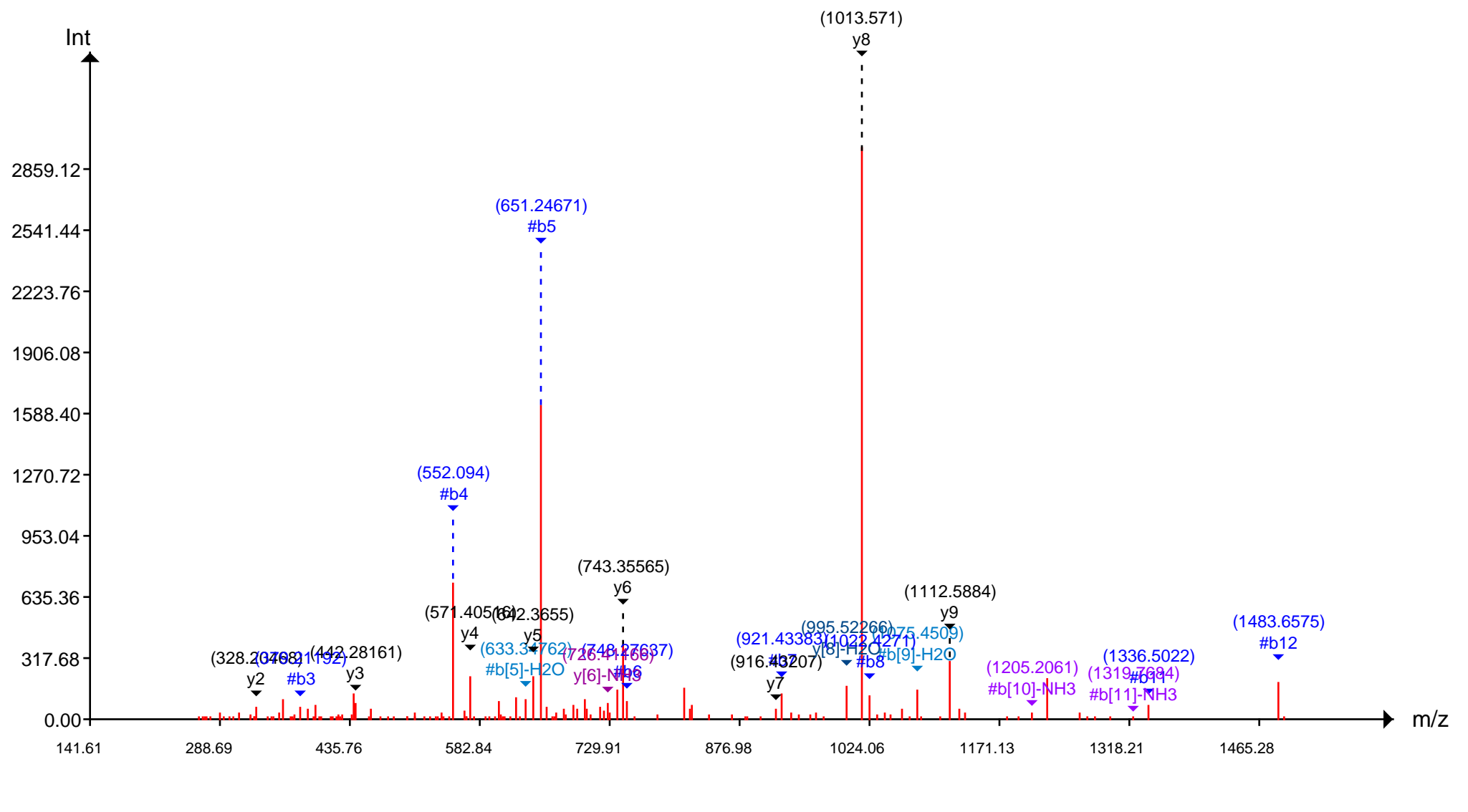
Info

Modified Sequence	AcD3-YK<AcD3K*>EAGVGK<AcD3K*>DR<C13>-COOH	Ion score	53.36
Protein	Q93092	Confidence level	99.00%
Precursor m/z	632.35335	Identity Threshold	47.66
Charge	2+	Mass error	9.1E-4

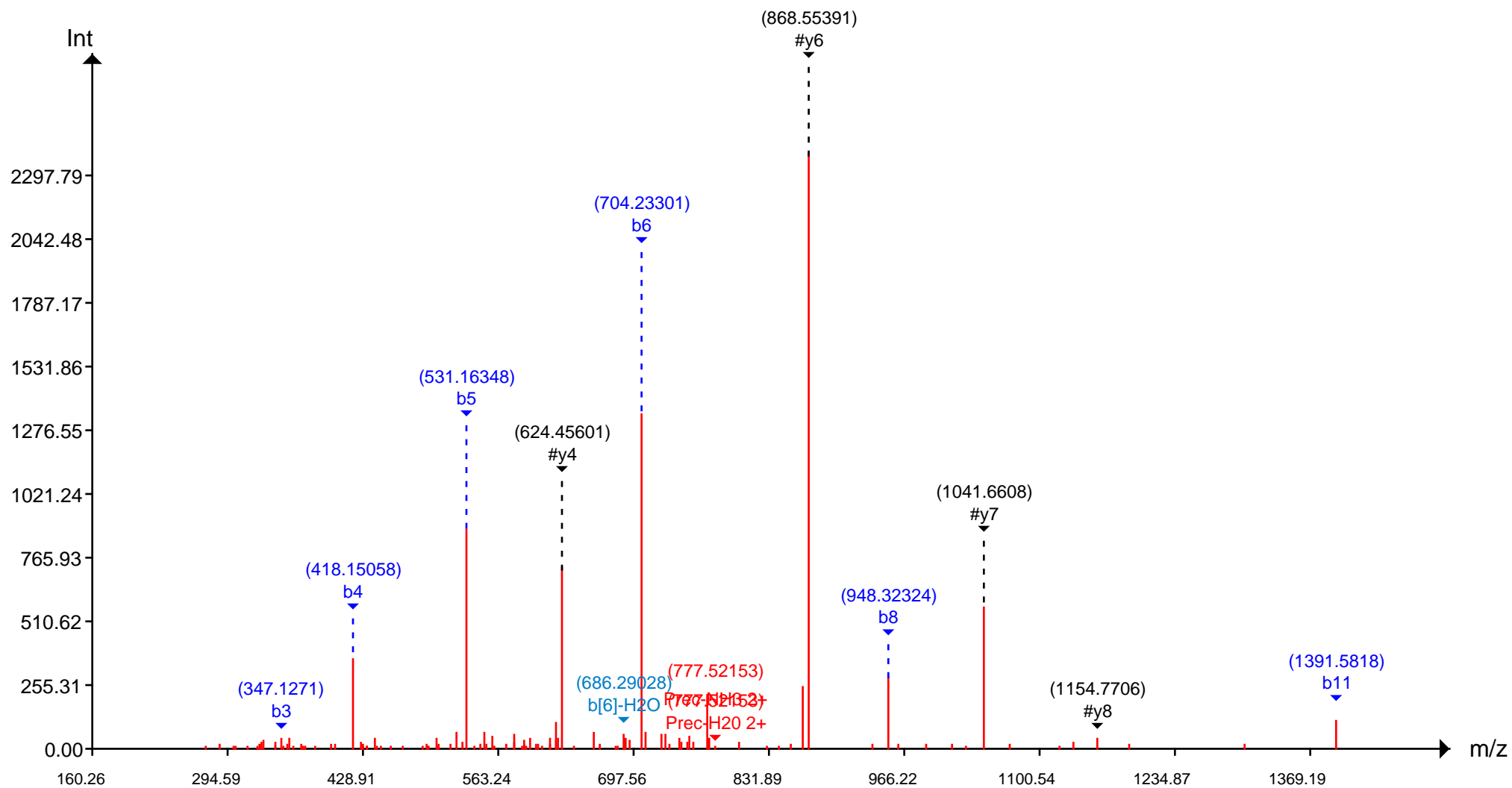


Info

Modified Sequence	AcD3-FEEK<AcD3K*>DIAANEENR<C13>-COOH	Ion score	51.12
Protein	Q9JJU8	Confidence level	99.00%
Precursor m/z	830.90588	Identity Threshold	47.13
Charge	2+	Mass error	0.001813

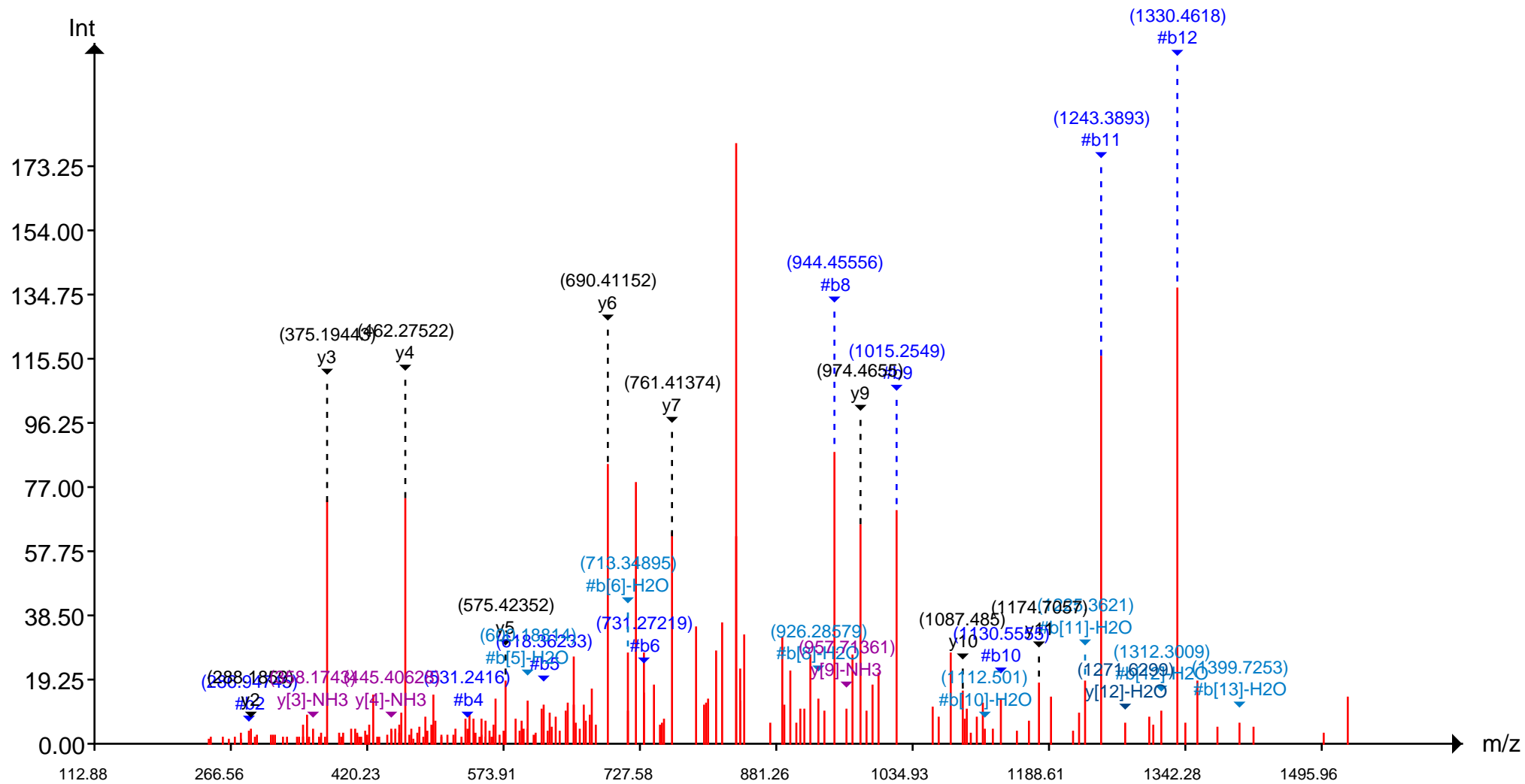


Info			
Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH	Ion score	49.09
Protein	P17742	Confidence level	99.00%
Precursor m/z	832.4593	Identity Threshold	47.48
Charge	2+	Mass error	0.00175



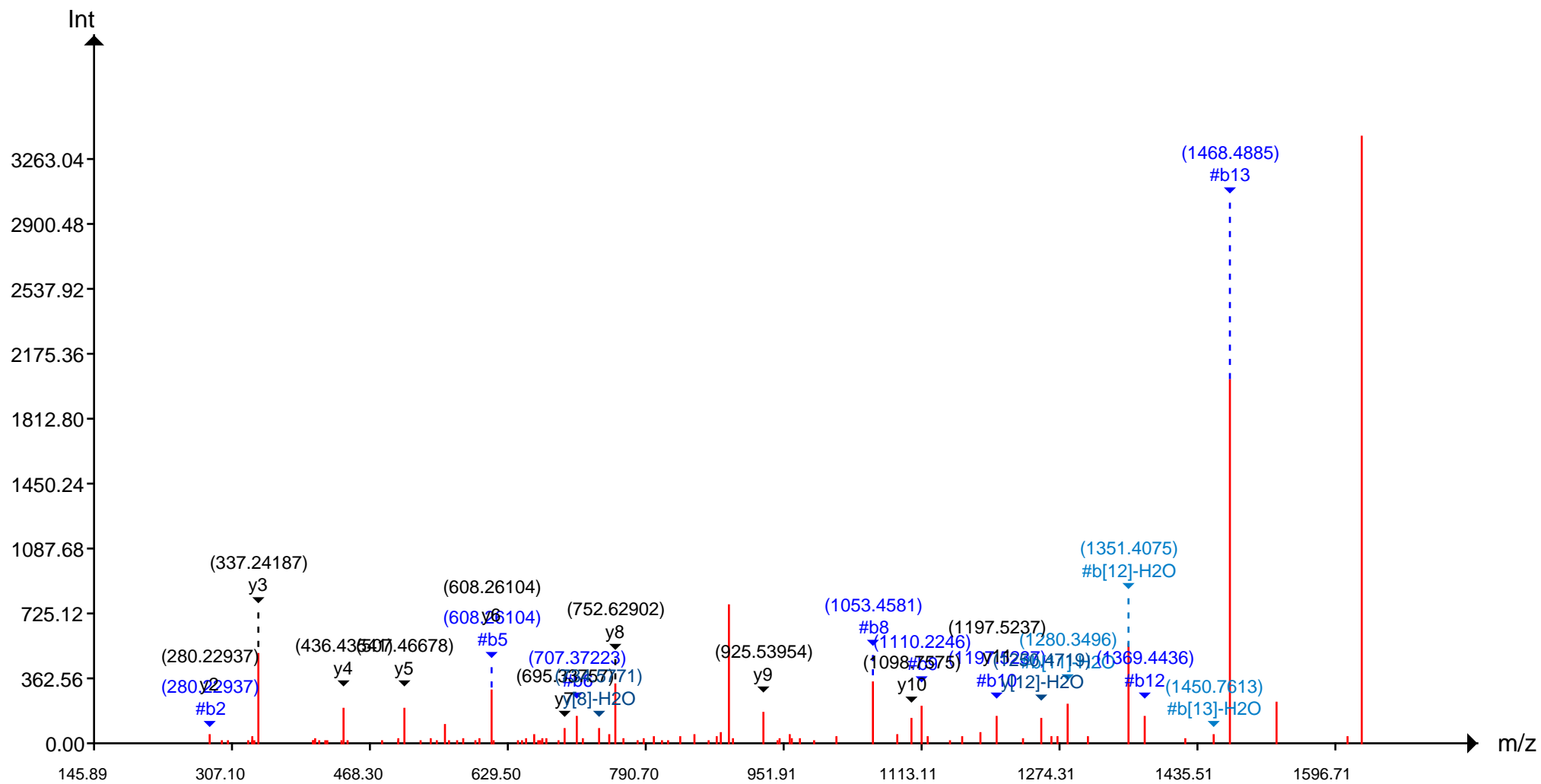
Info

Modified Sequence	AcD3-AATEAIK<AcD3K*>PFPK<AcD3K*>K<AcD3K*>R<C13>-COOH	Ion score	49.79
Protein	Q6PGL7	Confidence level	99.00%
Precursor m/z	786.48652	Identity Threshold	40.47
Charge	2+	Mass error	0.001716



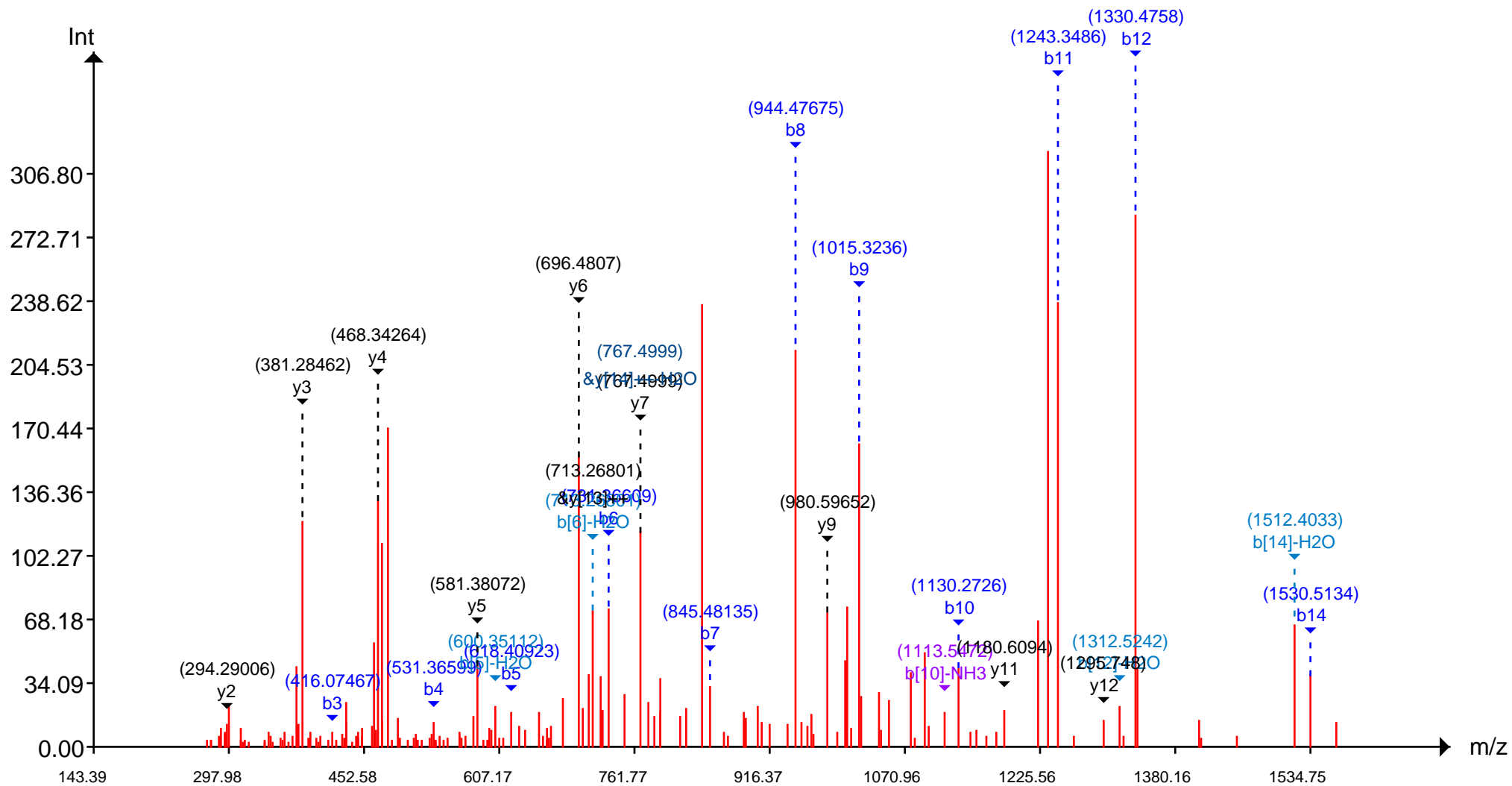
Info

Modified Sequence	AcD3-LQEDSLNVADLSSLR-COOH	Ion score	52.69
Protein	P97372	Confidence level	99.00%
Precursor m/z	852.94731	Identity Threshold	48.21
Charge	2+	Mass error	0.003176



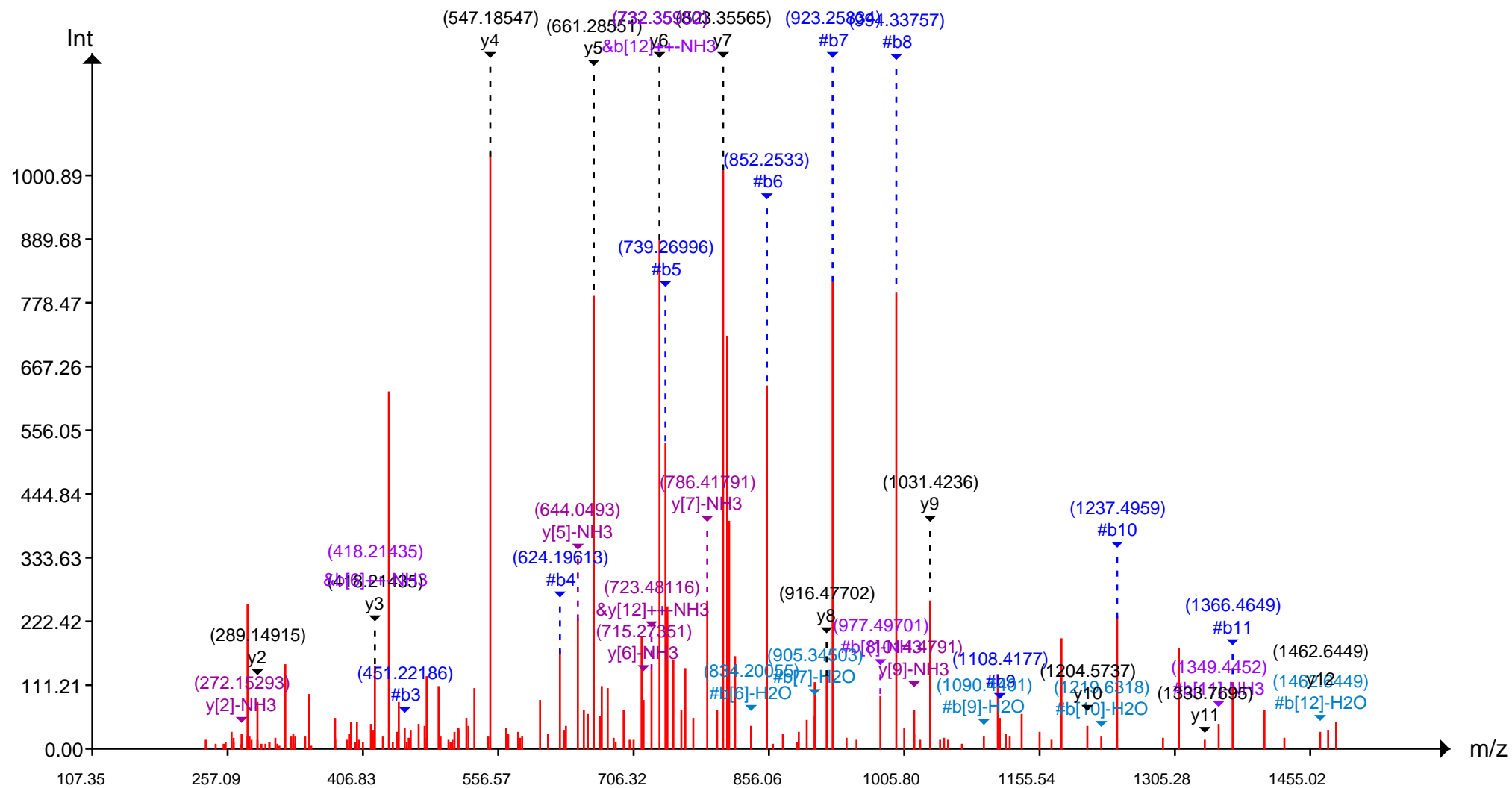
Info

Modified Sequence	AcD3-YAQEAVK<AcD3K*>K<AcD3K*>GSTAVGVR<C13>-COOH	Ion score	62.93
Protein	Q9Z2U0	Confidence level	99.00%
Precursor m/z	903.01508	Identity Threshold	45.80
Charge	2+	Mass error	0.001967



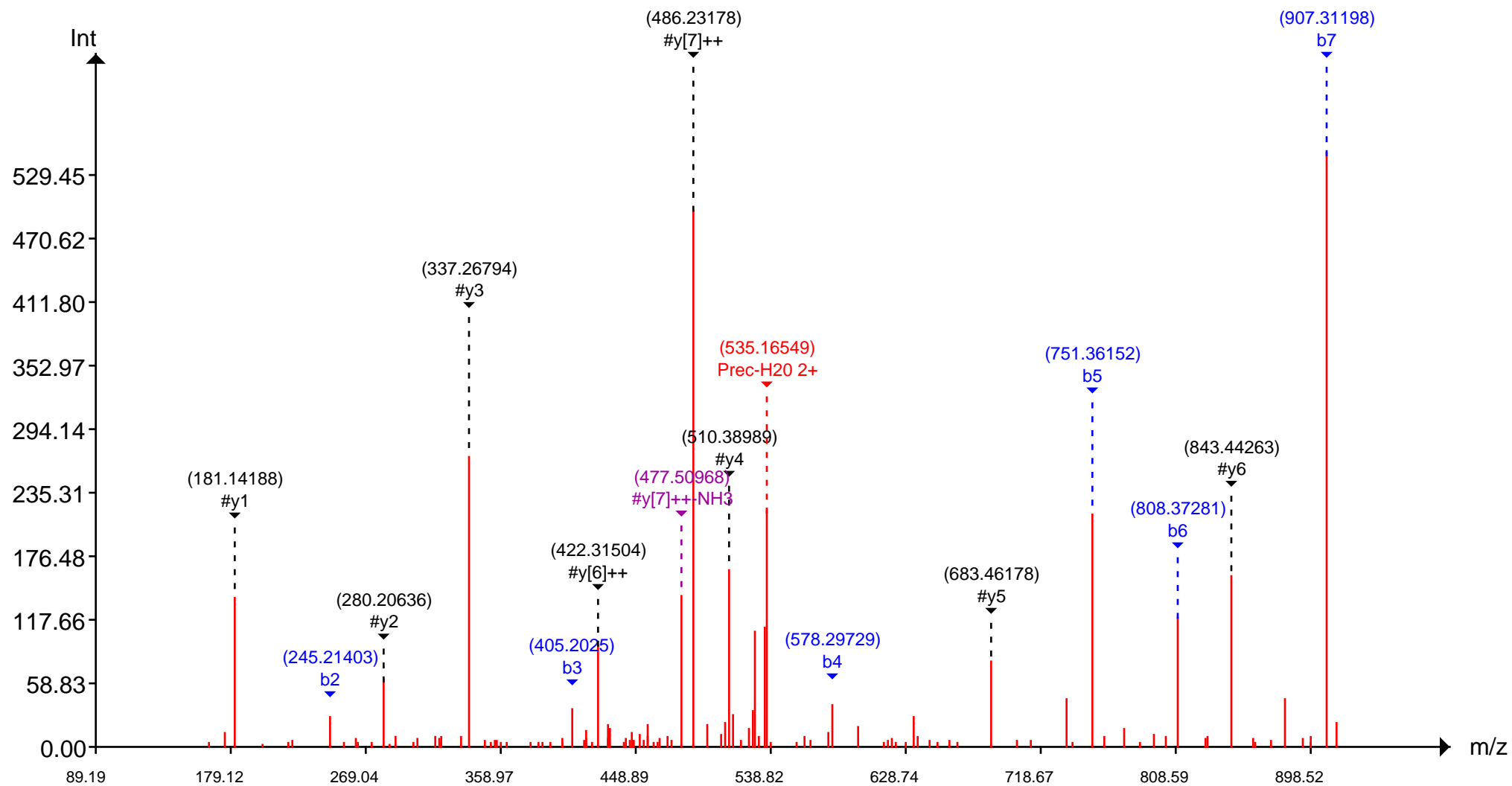
Info

Modified Sequence	AcD3-LQEDSLNVADLSSLR<C13>-COOH	Ion score	60.73
Protein	P97372	Confidence level	99.00%
Precursor m/z	855.95672	Identity Threshold	47.93
Charge	2+	Mass error	0.001854



Info

Modified Sequence	AcD3-FEEK<AcD3K*>DIAANEENR-COOH	Ion score	67.83
Protein	Q9JJU8	Confidence level	99.00%
Precursor m/z	827.89588	Identity Threshold	46.41
Charge	2+	Mass error	0.001955



Info

Modified Sequence AcD3-AQC<Cmm*>K<AcD3K*>K<AcD3K*>GVR<C13>-COOH

Ion score 52.94

Protein Q61753

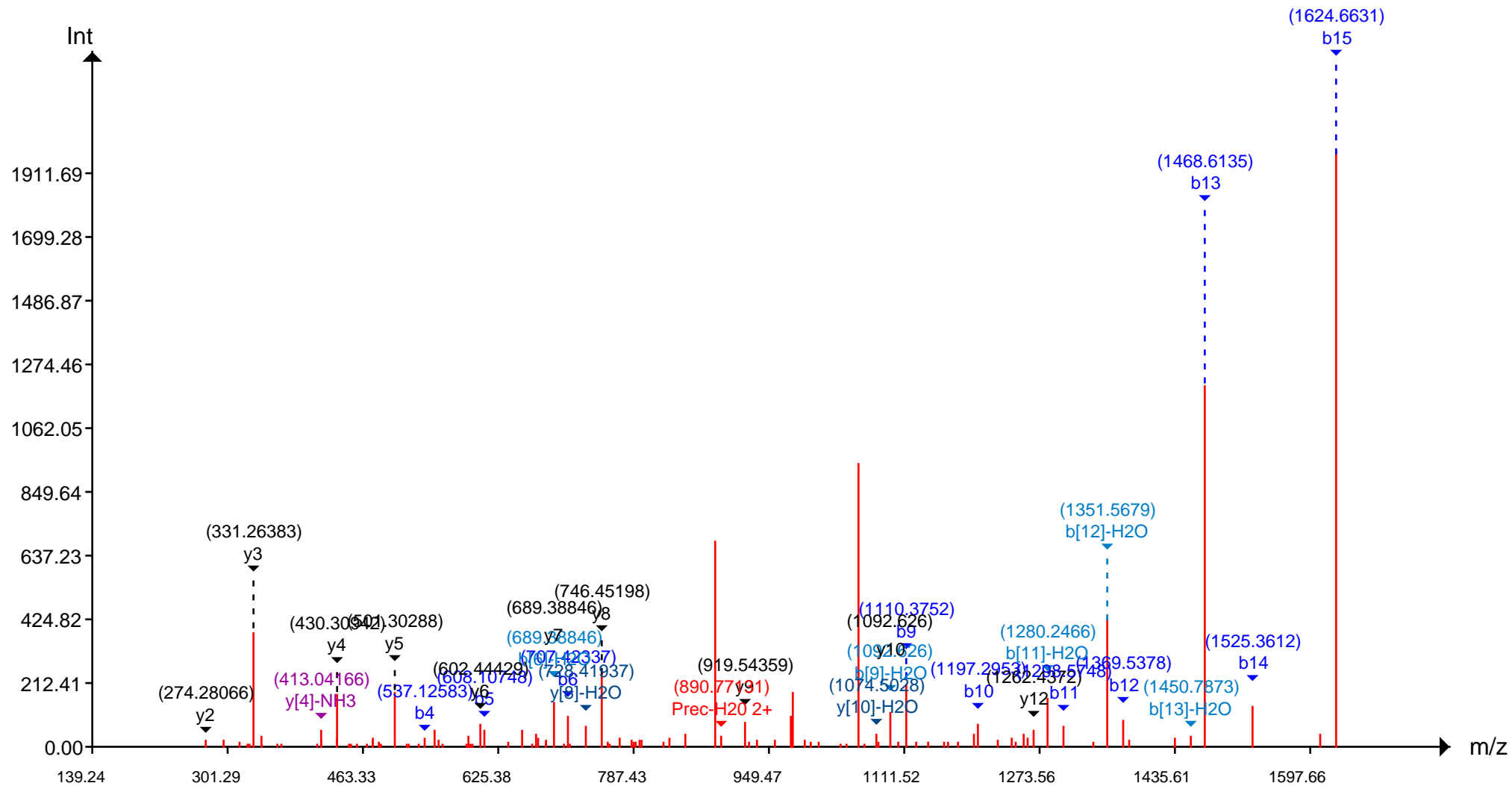
Confidence level 99.00%

Precursor m/z 544.32044

Identity Threshold 46.24

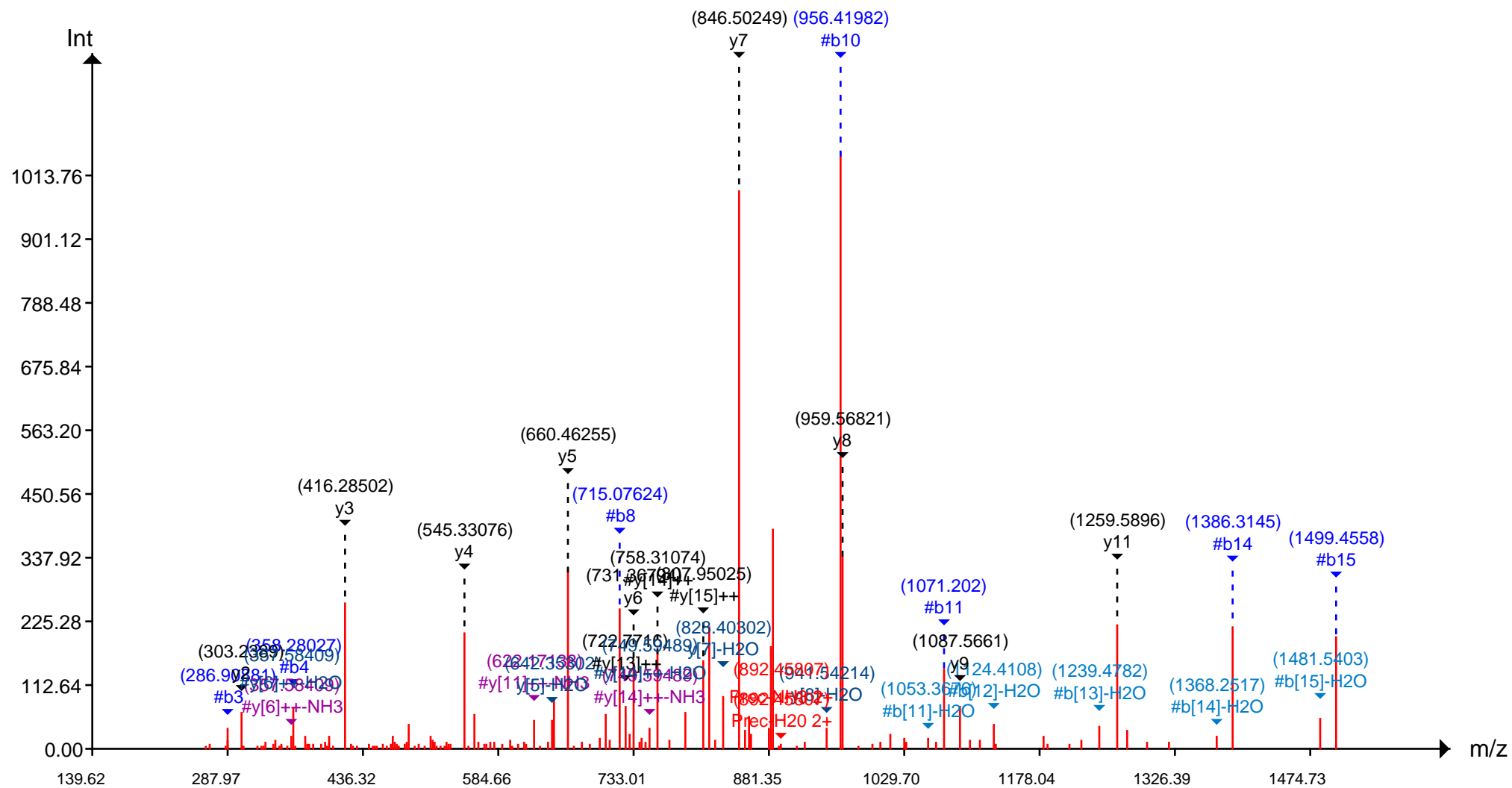
Charge 2+

Mass error 1.84E-4



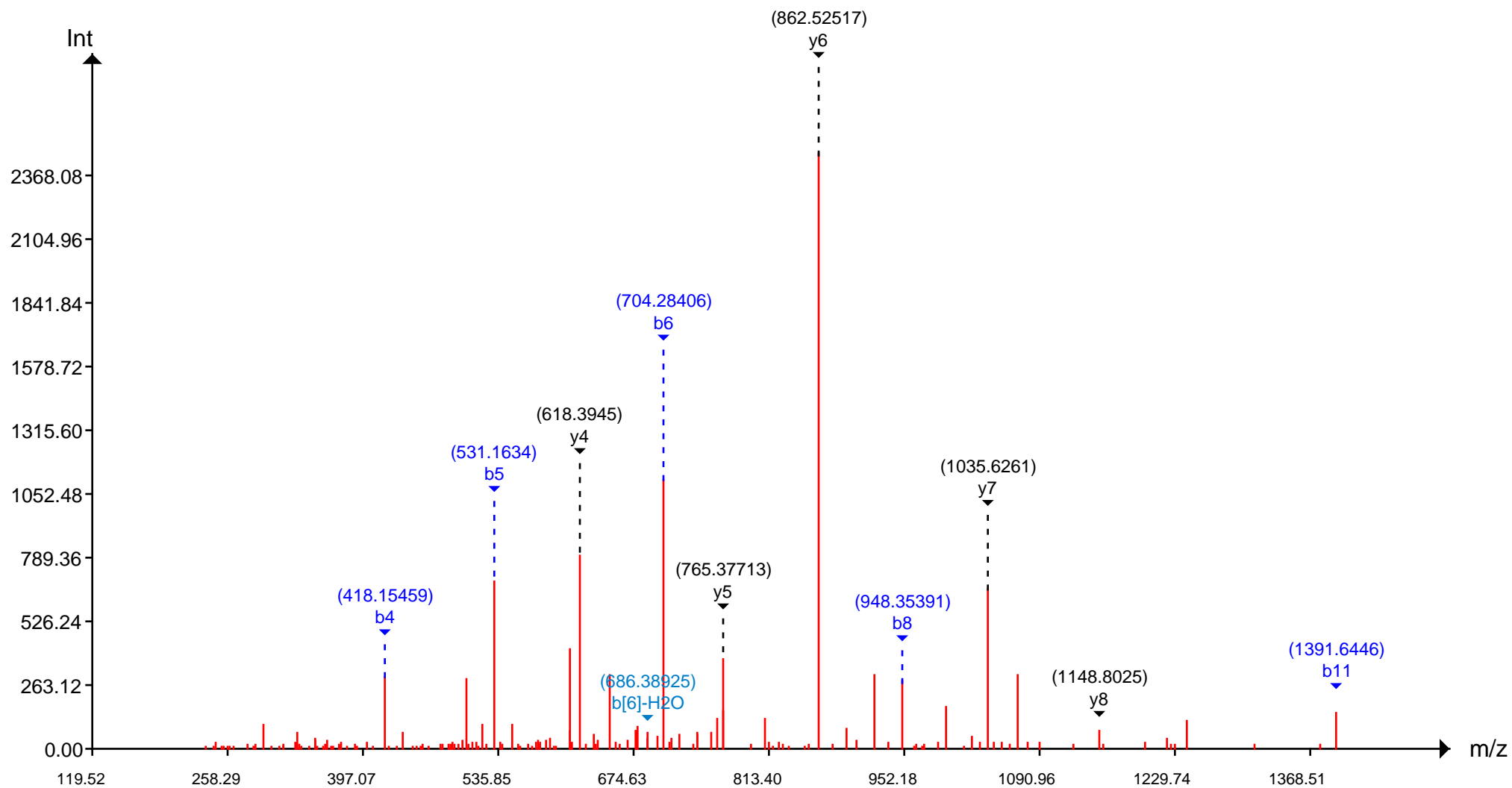
Info

Modified Sequence	AcD3-YAQEAVK<AcD3K*>K<AcD3K*>GSTAVGVR-COOH	Ion score	73.94
Protein	Q9Z2U0	Confidence level	99.00%
Precursor m/z	900.00512	Identity Threshold	46.61
Charge	2+	Mass error	0.002188



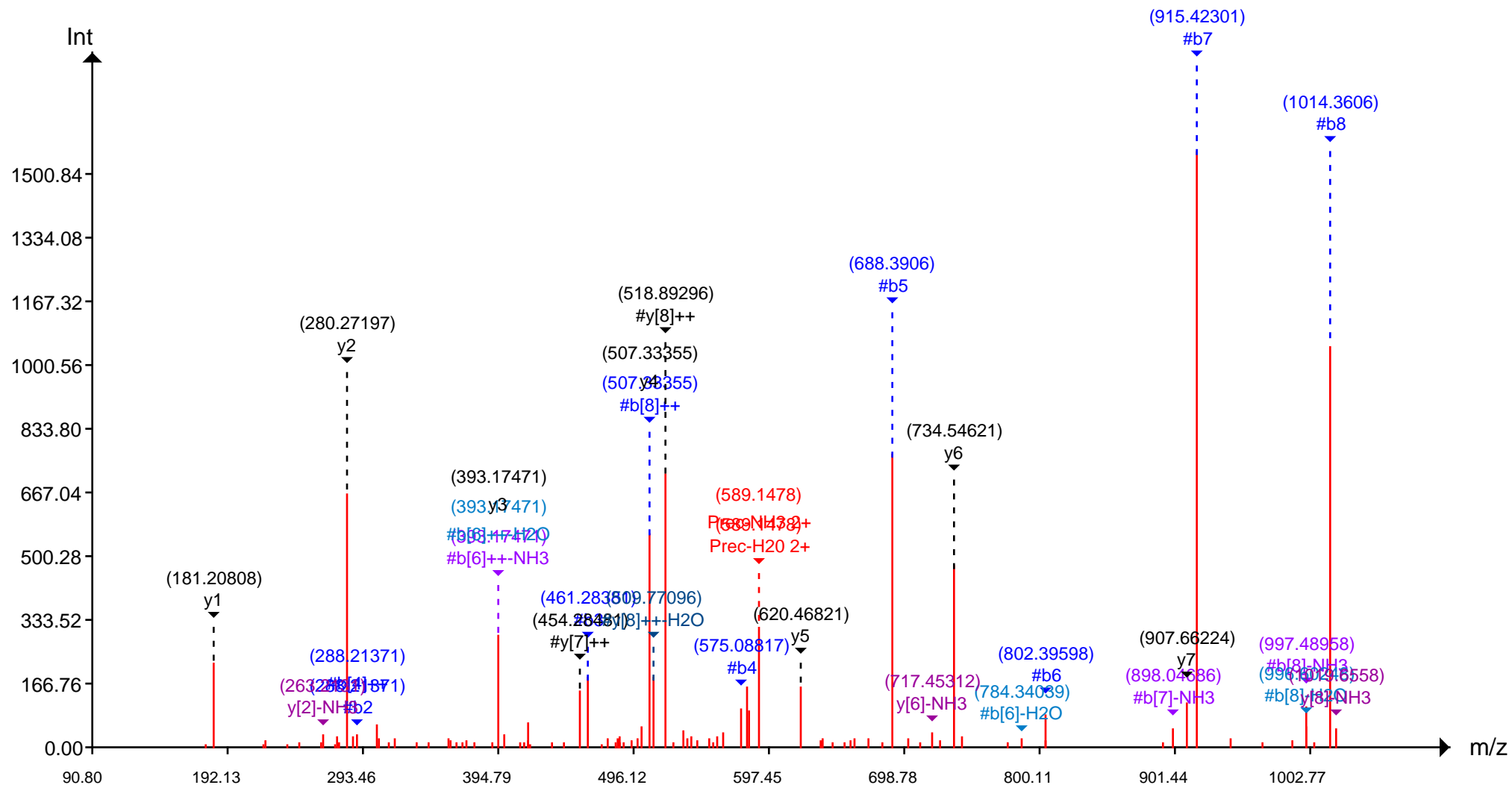
Info

Modified Sequence	AcD3-AAVAGQDGQIDADELQR-COOH	Ion score	61.01
Protein	Q6P069	Confidence level	99.00%
Precursor m/z	901.44238	Identity Threshold	47.25
Charge	2+	Mass error	0.002074



Info

Modified Sequence	AcD3-AATEAIK<AcD3K*>PFPK<AcD3K*>K<AcD3K*>R-COOH	Ion score	45.11
Protein	Q6PGL7	Confidence level	99.00%
Precursor m/z	783.47661	Identity Threshold	42.21
Charge	2+	Mass error	0.002038



Info

Modified Sequence AcD3-LEK<AcD3K*>NINIVR<C13>-COOH

Ion score 66.06

Protein Q99K10

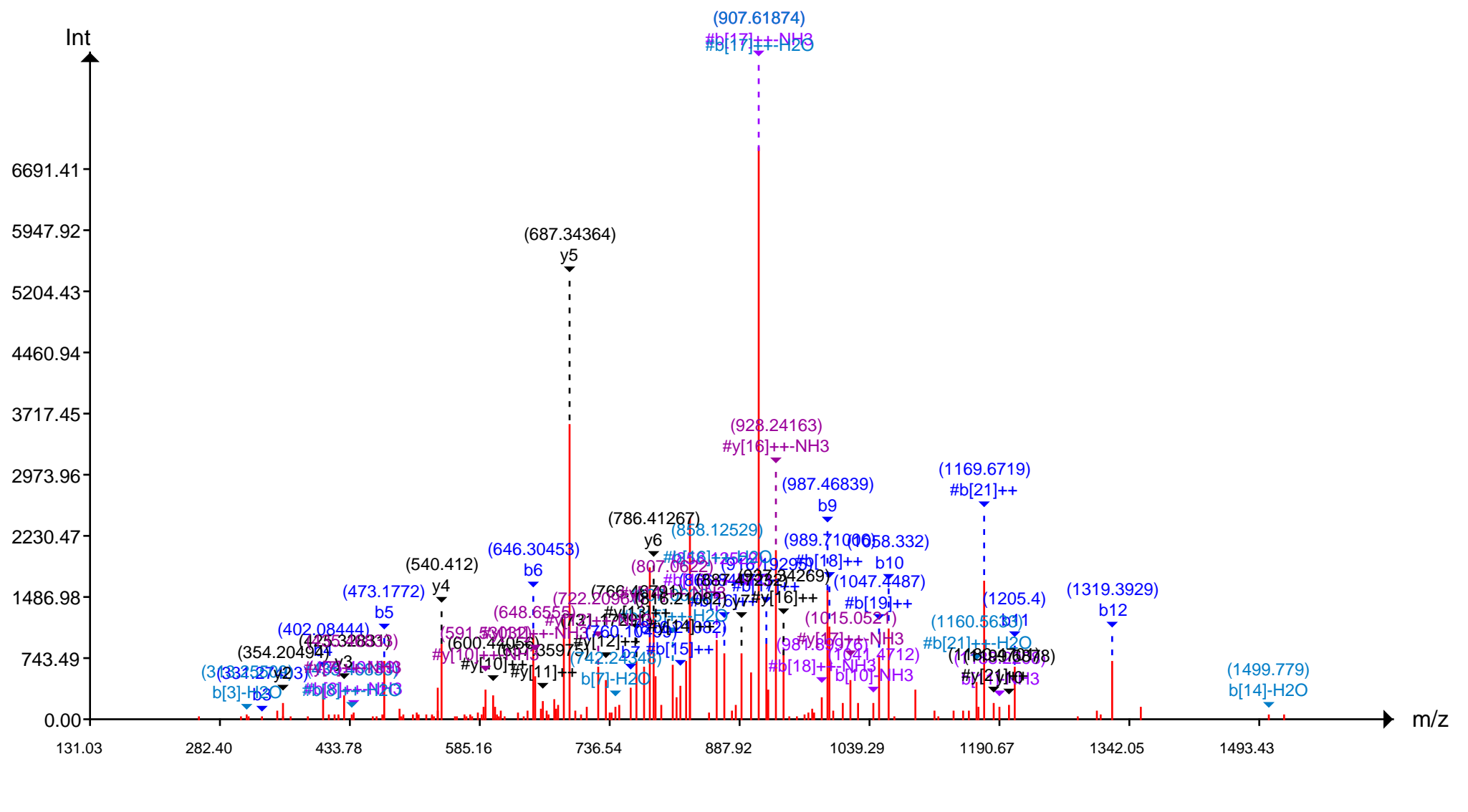
Confidence level 99.00%

Precursor m/z 597.87466

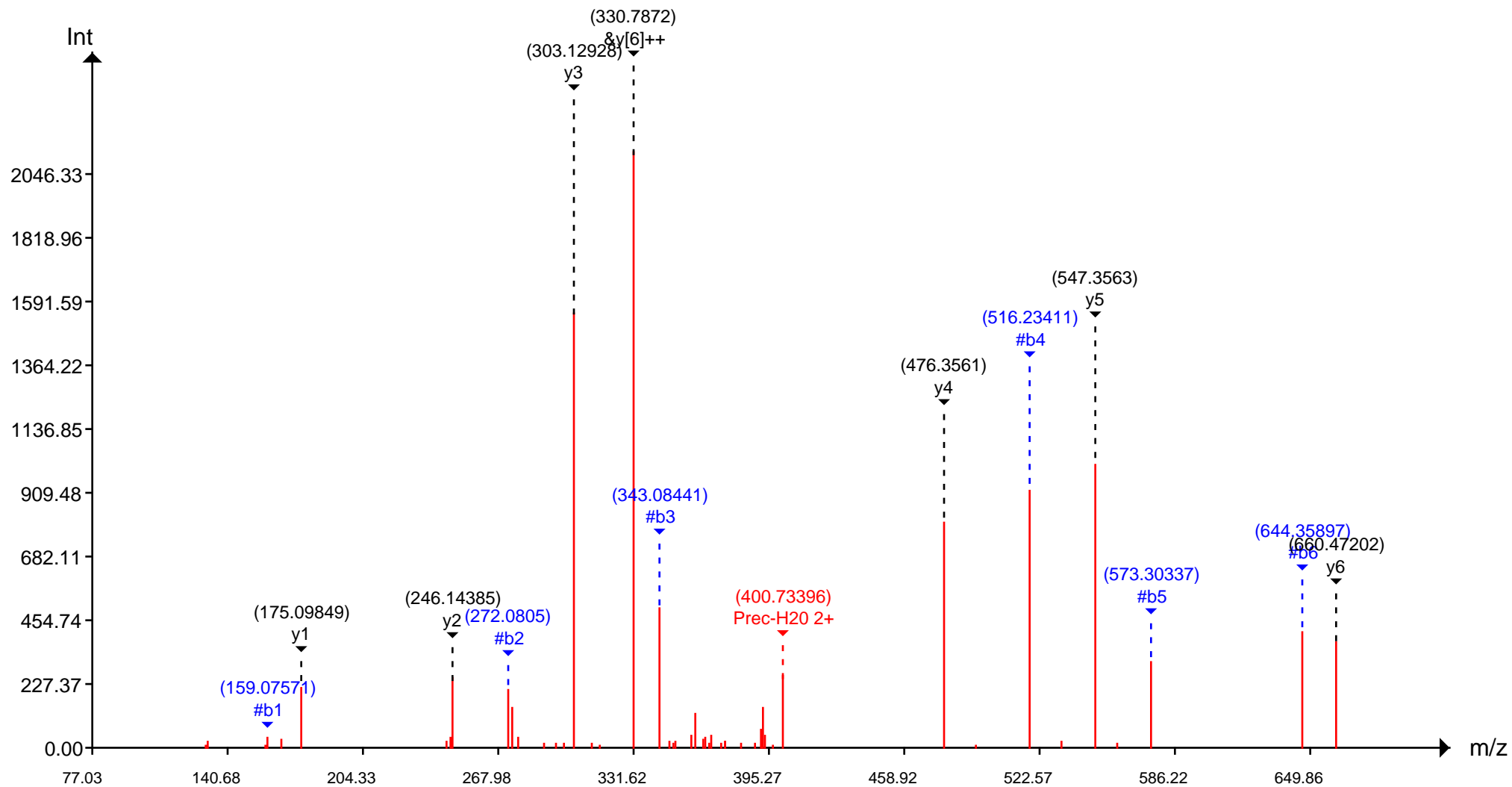
Identity Threshold 41.81

Charge 2+

Mass error 1.79E-4



Info			
Modified Sequence	AcD3-IGDAAK<AcD3K*>NQVAM<Mox*>NPTNTVFDAAK<AcD3K*>R<C13>-COOH	Ion score	47.89
Protein	P63017	Confidence level	99.00%
Precursor m/z	840.10585	Identity Threshold	47.87
Charge	3+	Mass error	0.001807



Info

Modified Sequence AcD3-LLAK<AcD3K*>GAR-COOH

Ion score 64.78

Protein Q62422

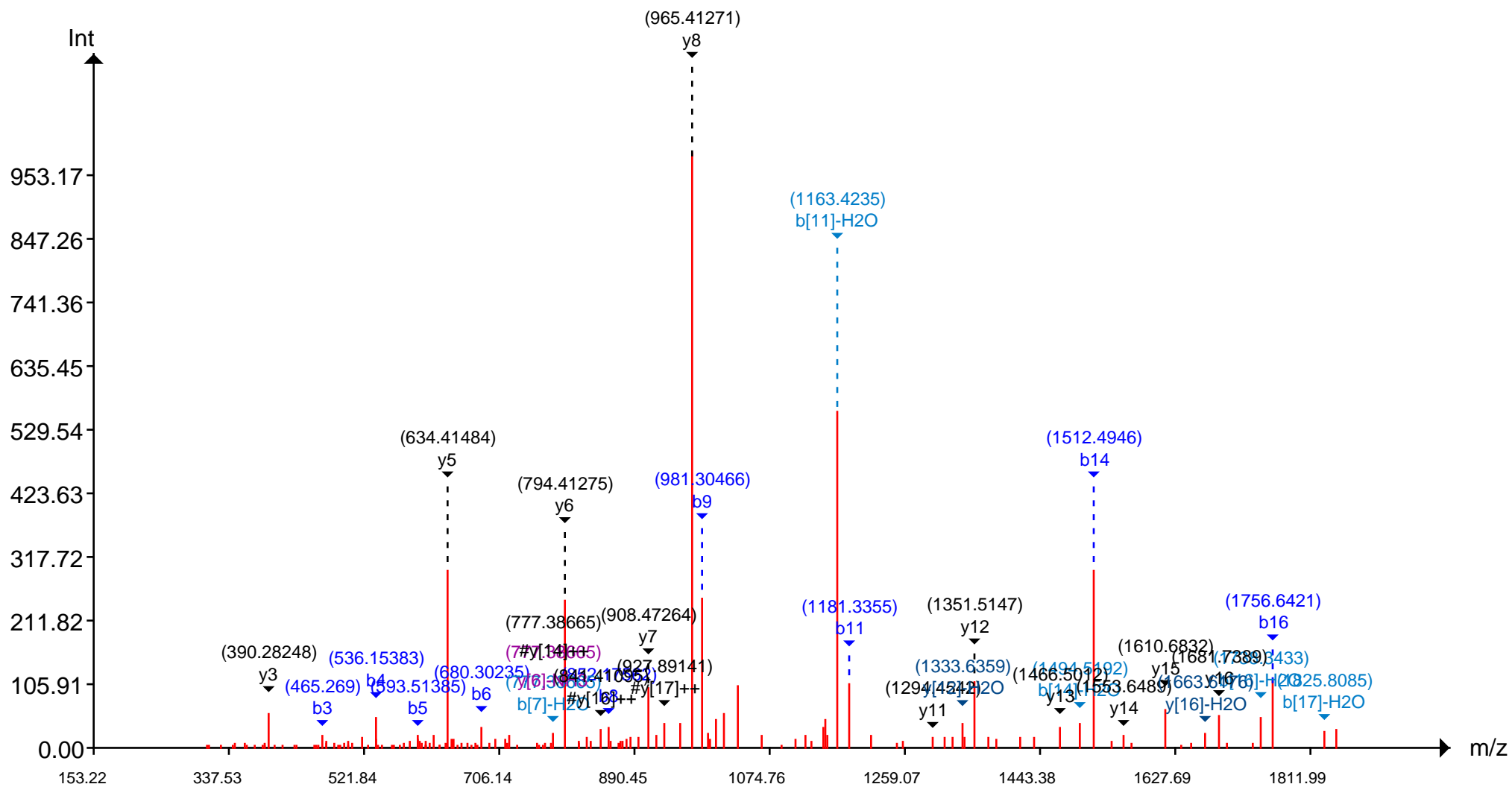
Confidence level 99.00%

Precursor m/z 409.7719

Identity Threshold 39.57

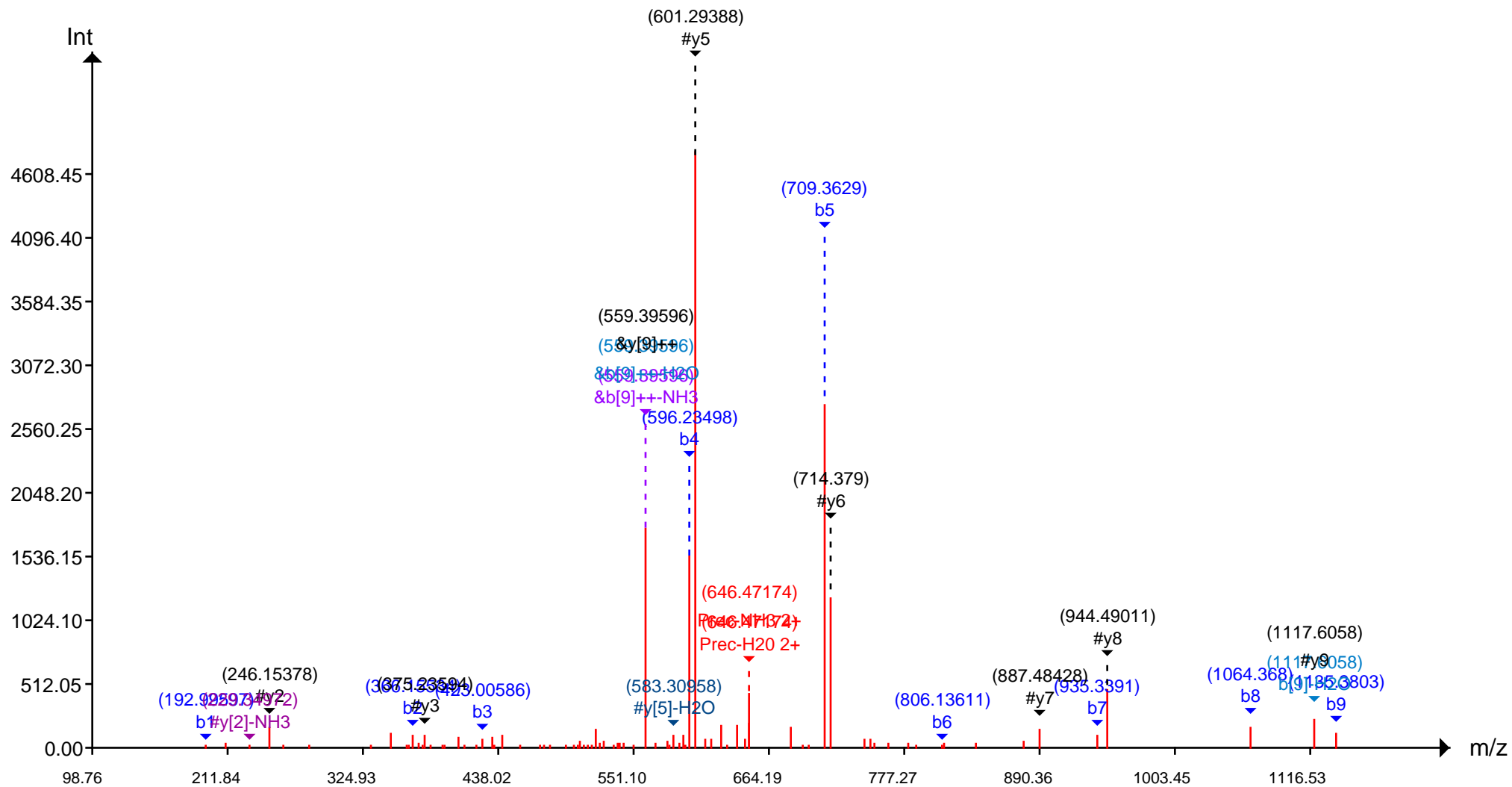
Charge 2+

Mass error 2.7E-5



Info

Modified Sequence	AcD3-FVK<AcD3K*>AGSDGESIGNC<Cmm*>PFSQR-COOH	Ion score	83.75
Protein	Q9QYB1	Confidence level	99.00%
Precursor m/z	1073.511	Identity Threshold	46.23
Charge	2+	Mass error	6.58E-4



Info

Modified Sequence AcD3-M<Mox*>K<AcD3K*>GK<AcD3K*>LPPEAR-COOH

Ion score 60.18

Protein P06745

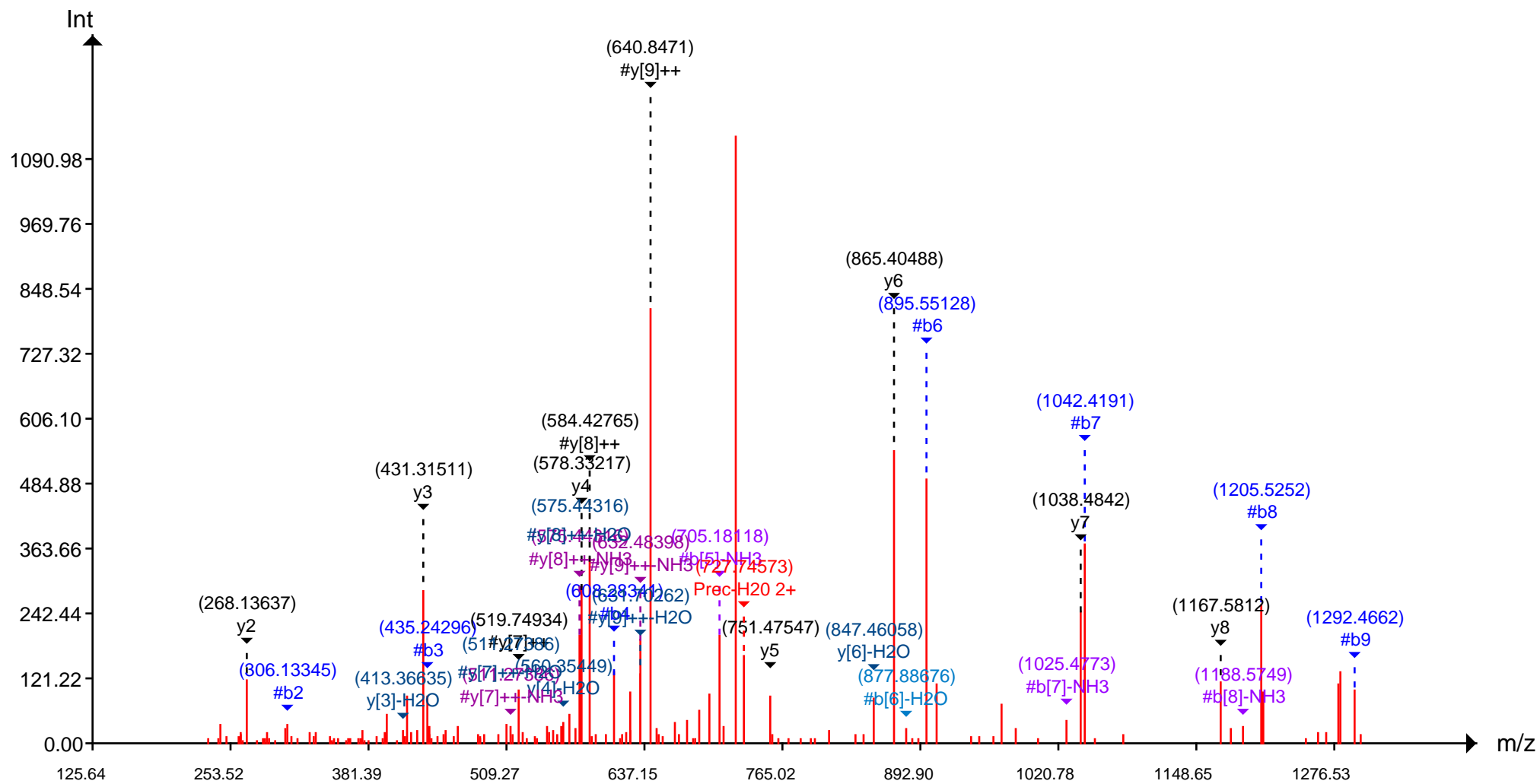
Confidence level 99.00%

Precursor m/z 655.3608

Identity Threshold 47.84

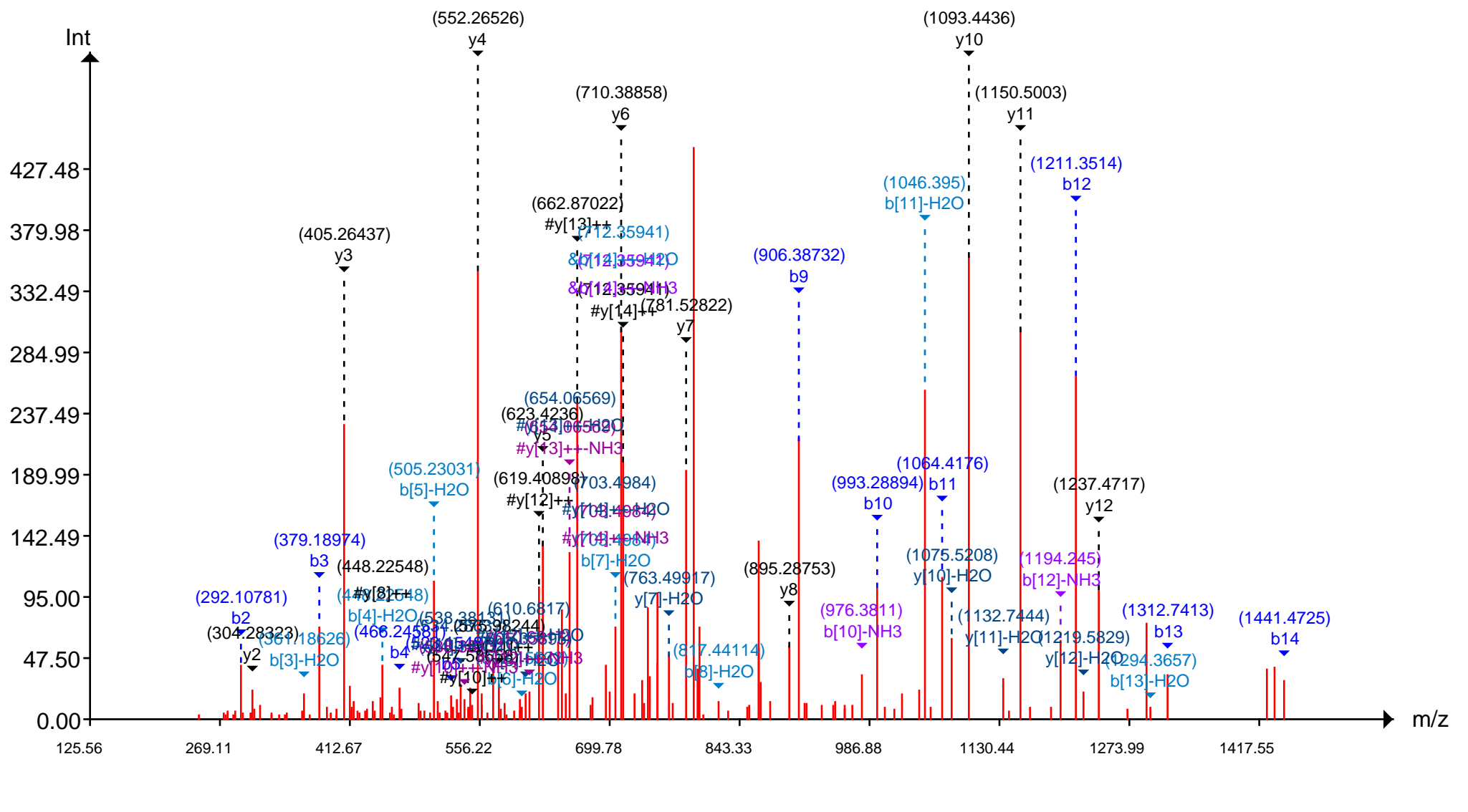
Charge 2+

Mass error 0.001298

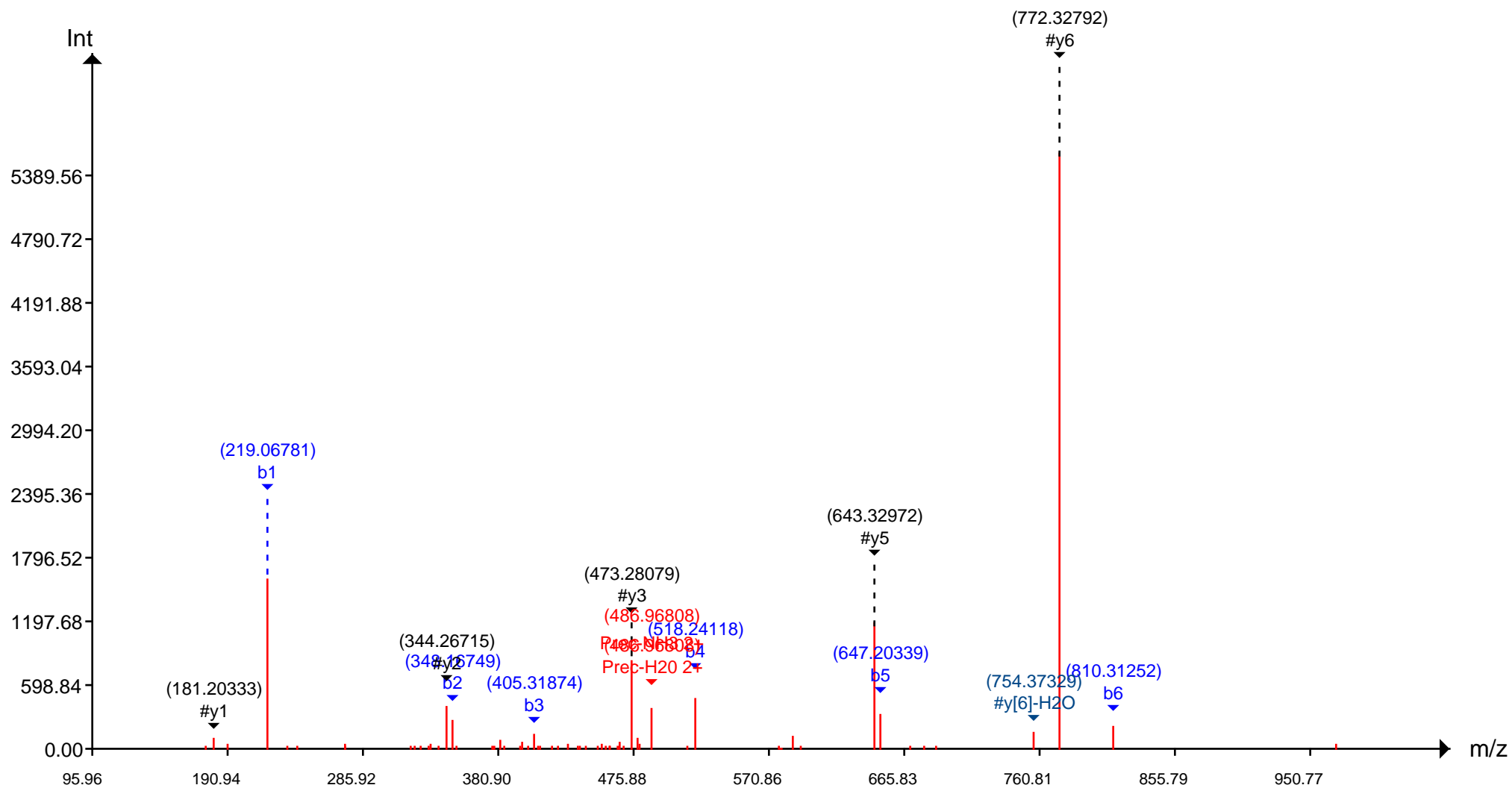


Info

Modified Sequence	AcD3-FLEK<AcD3K*>NK<AcD3K*>M<Mox*>YSR<C13>-COOH	Ion score	51.6
Protein	Q9D7A6	Confidence level	99.00%
Precursor m/z	736.89715	Identity Threshold	48.04
Charge	2+	Mass error	0.001092



Info			
Modified Sequence	AcD3-FVSSGPTNASAFter-COOH	Ion score	86.86
Protein	P27546	Confidence level	99.00%
Precursor m/z	808.3944	Identity Threshold	47.03
Charge	2+	Mass error	0.002519



Info

Modified Sequence AcD3-K<AcD3K*>EGIEYR<C13>-COOH

Ion score 59.05

Protein Q61599

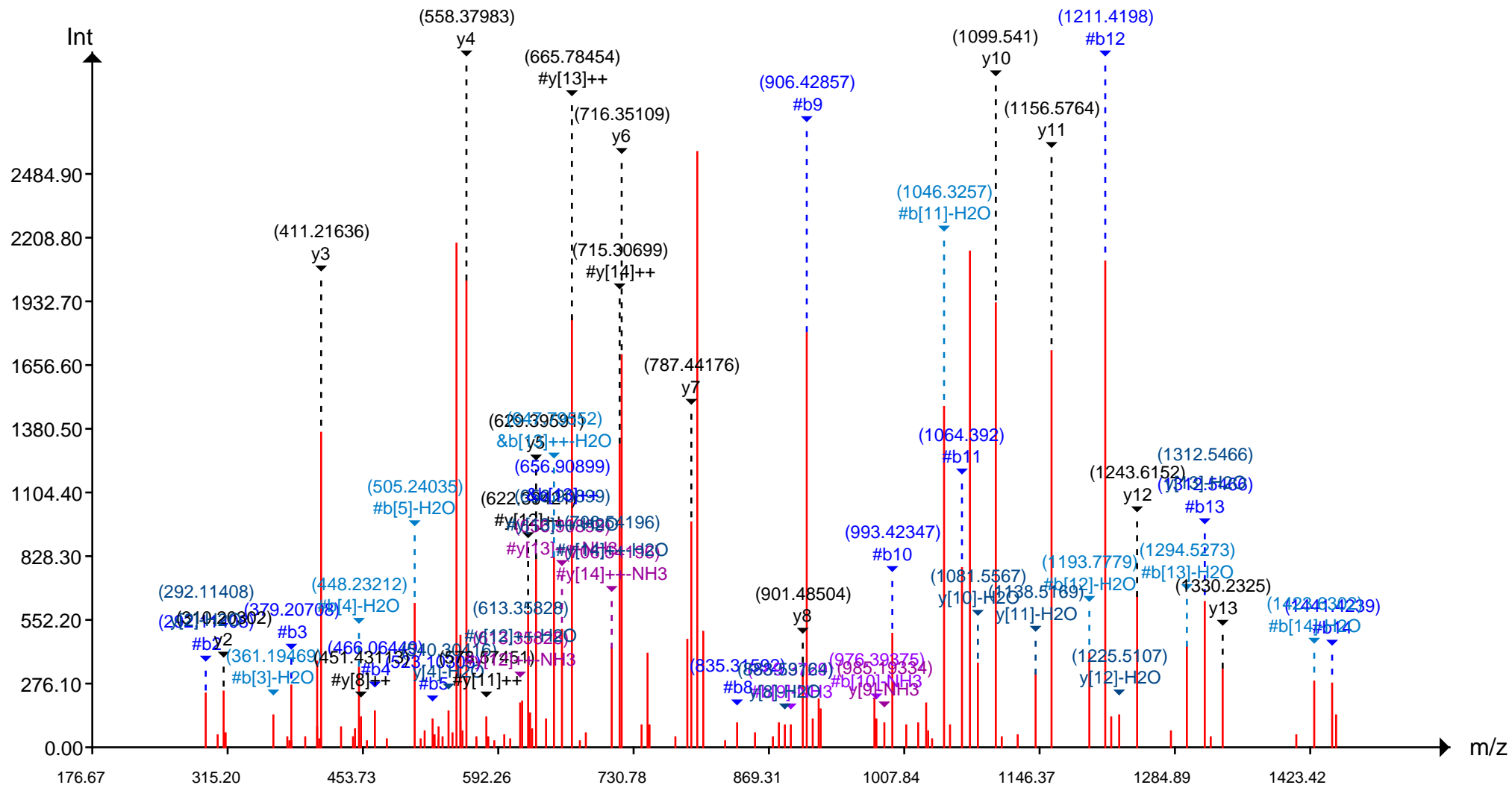
Confidence level 99.00%

Precursor m/z 495.77679

Identity Threshold 48.30

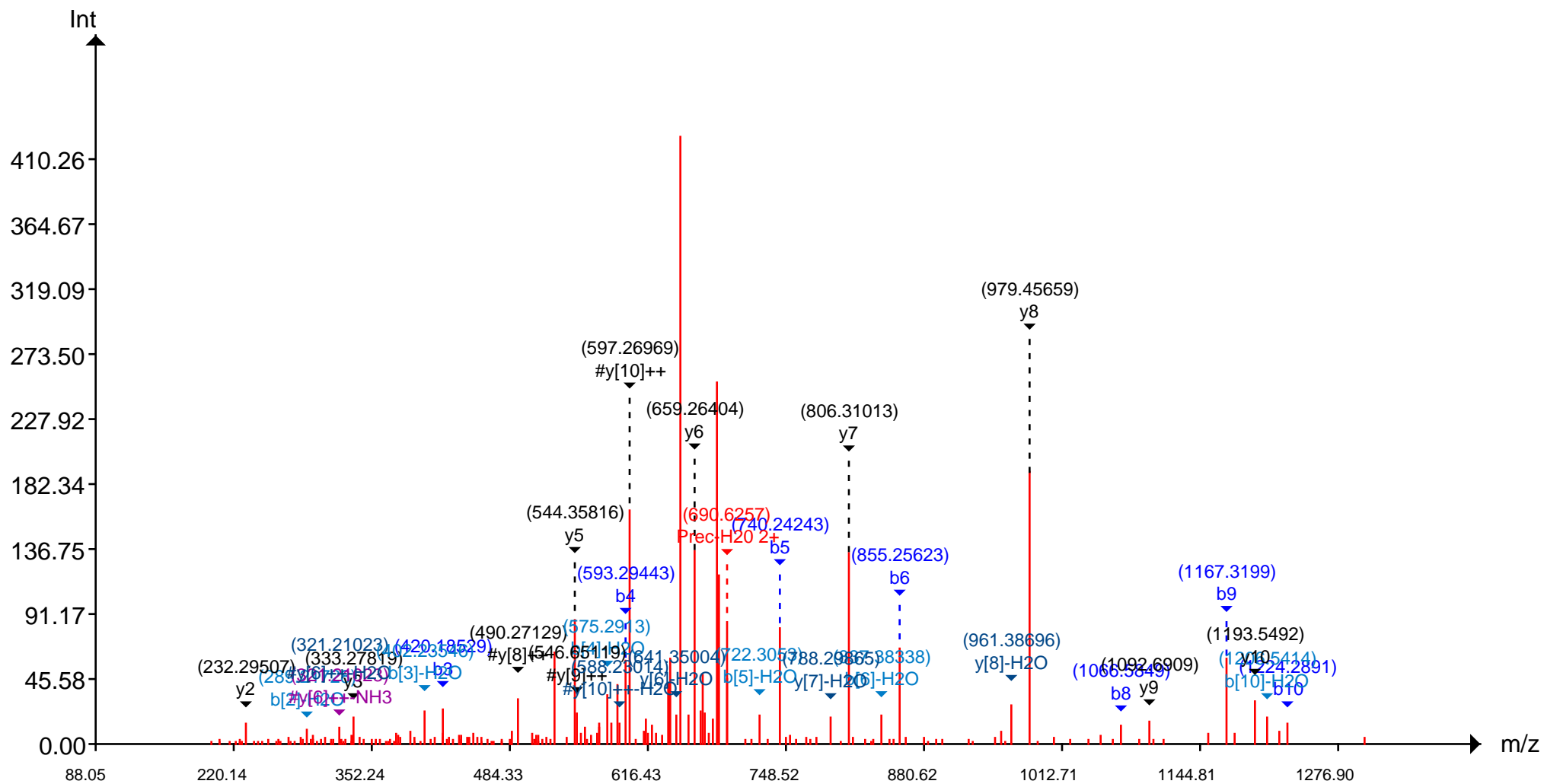
Charge 2+

Mass error -5.53E-4



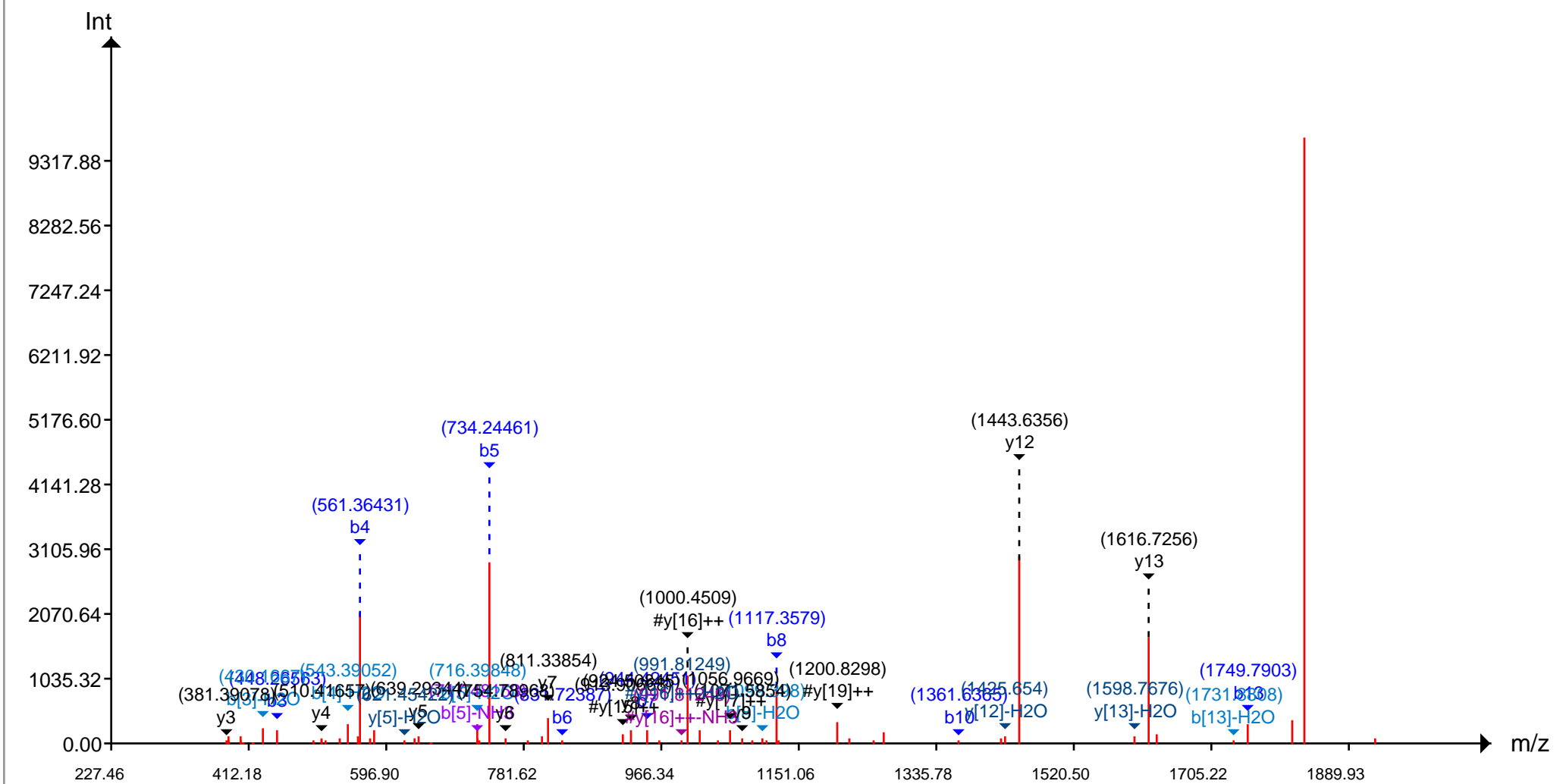
Info

Modified Sequence	AcD3-FVSSGPTNASAFter<C13>-COOH	Ion score	101.35
Protein	P27546	Confidence level	99.00%
Precursor m/z	811.40383	Identity Threshold	47.52
Charge	2+	Mass error	0.001238



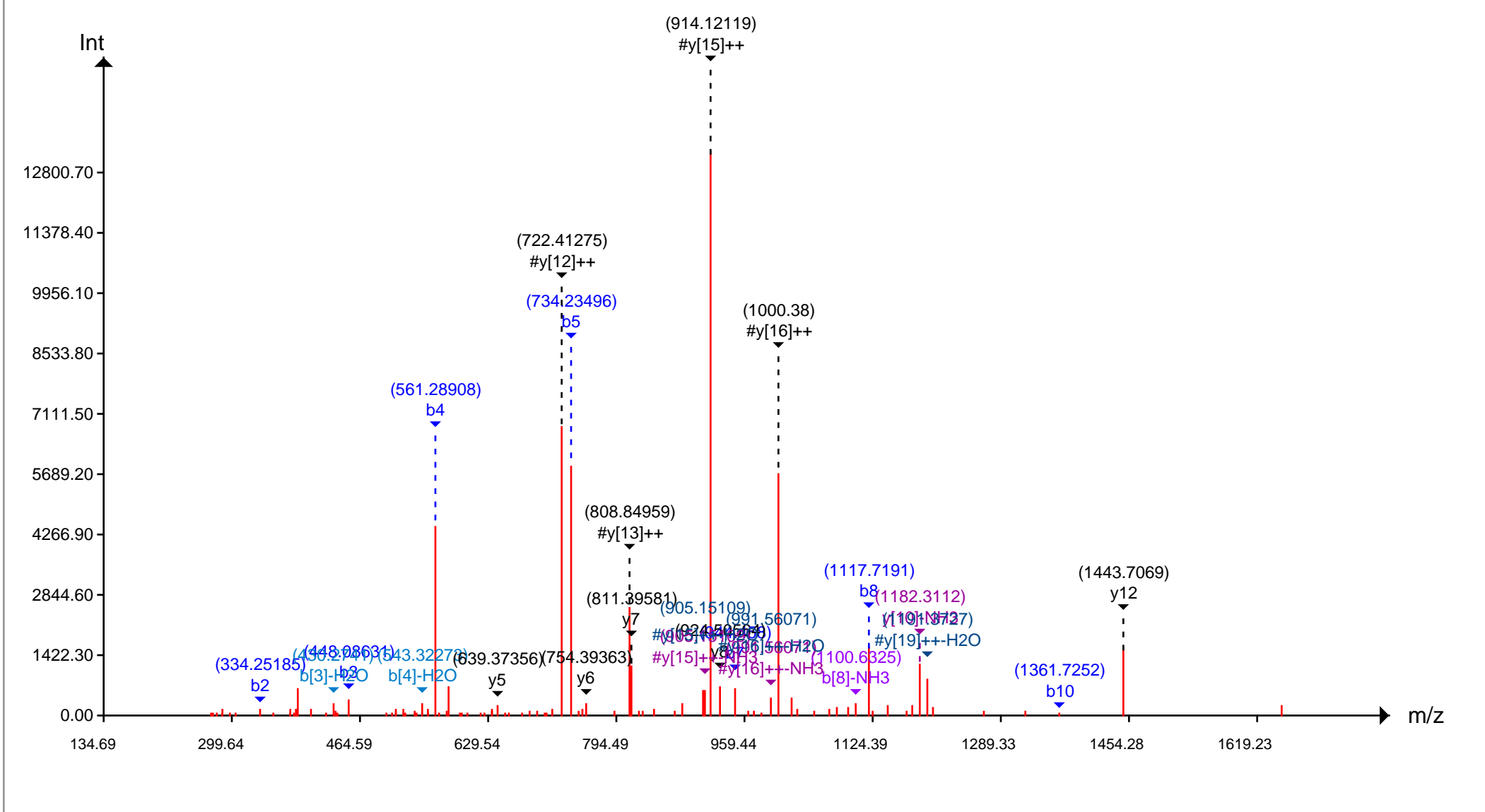
Info

Modified Sequence	AcD3-C<Cmm*>TIK<AcD3K*>M<Mox*>DPNTGR-COOH	Ion score	71.38
Protein	Q99020	Confidence level	99.00%
Precursor m/z	699.83589	Identity Threshold	46.44
Charge	2+	Mass error	0.002229

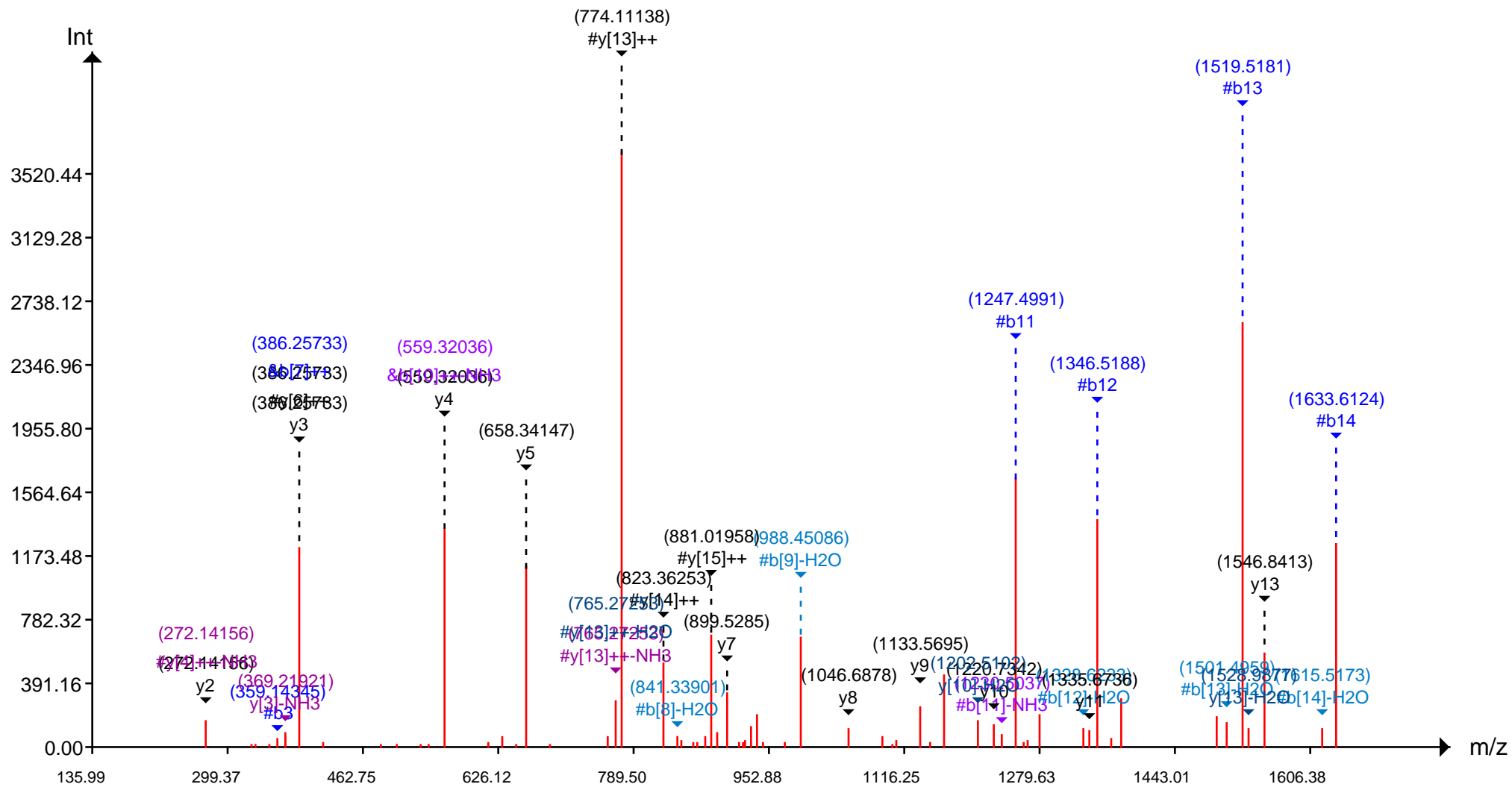


Info

Modified Sequence	AcD3-DK<AcD3K*>NLK<AcD3K*>PIK<AcD3K*>PM<Mox*>QFLGDEETVR<C13>-COOH	Ion score	60.02
Protein	Q9DBJ1	Confidence level	99.00%
Precursor m/z	1280.6953	Identity Threshold	46.76
Charge	2+	Mass error	0.002101

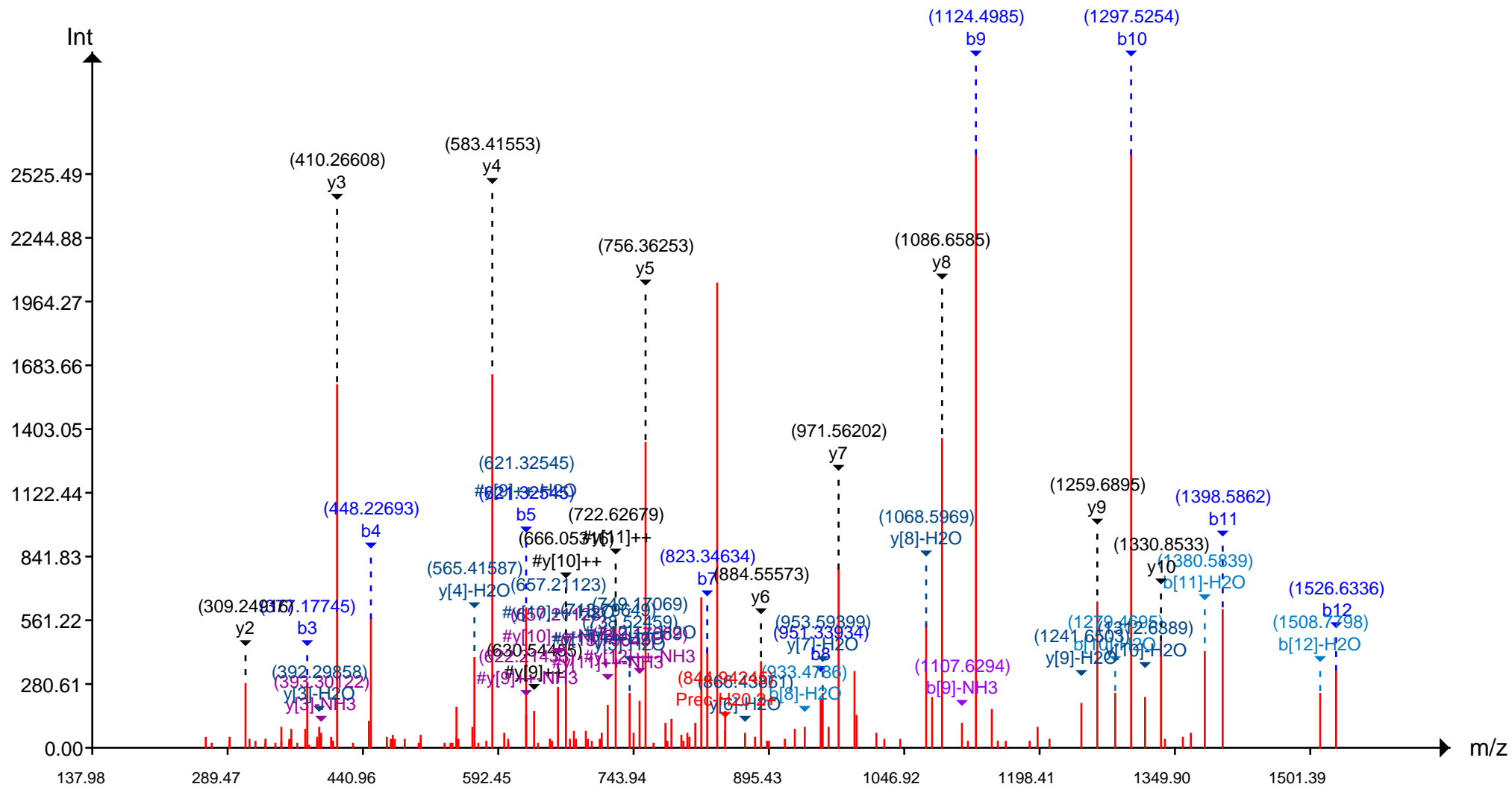


Info			
Modified Sequence	AcD3-DK<AcD3K*>NLK<AcD3K*>PIK<AcD3K*>PM<Mox*>QFLGDEETVR<C13>-COOH	Ion score	46.97
Protein	Q9DBJ1	Confidence level	99.00%
Precursor m/z	854.13244	Identity Threshold	46.77
Charge	3+	Mass error	0.001545



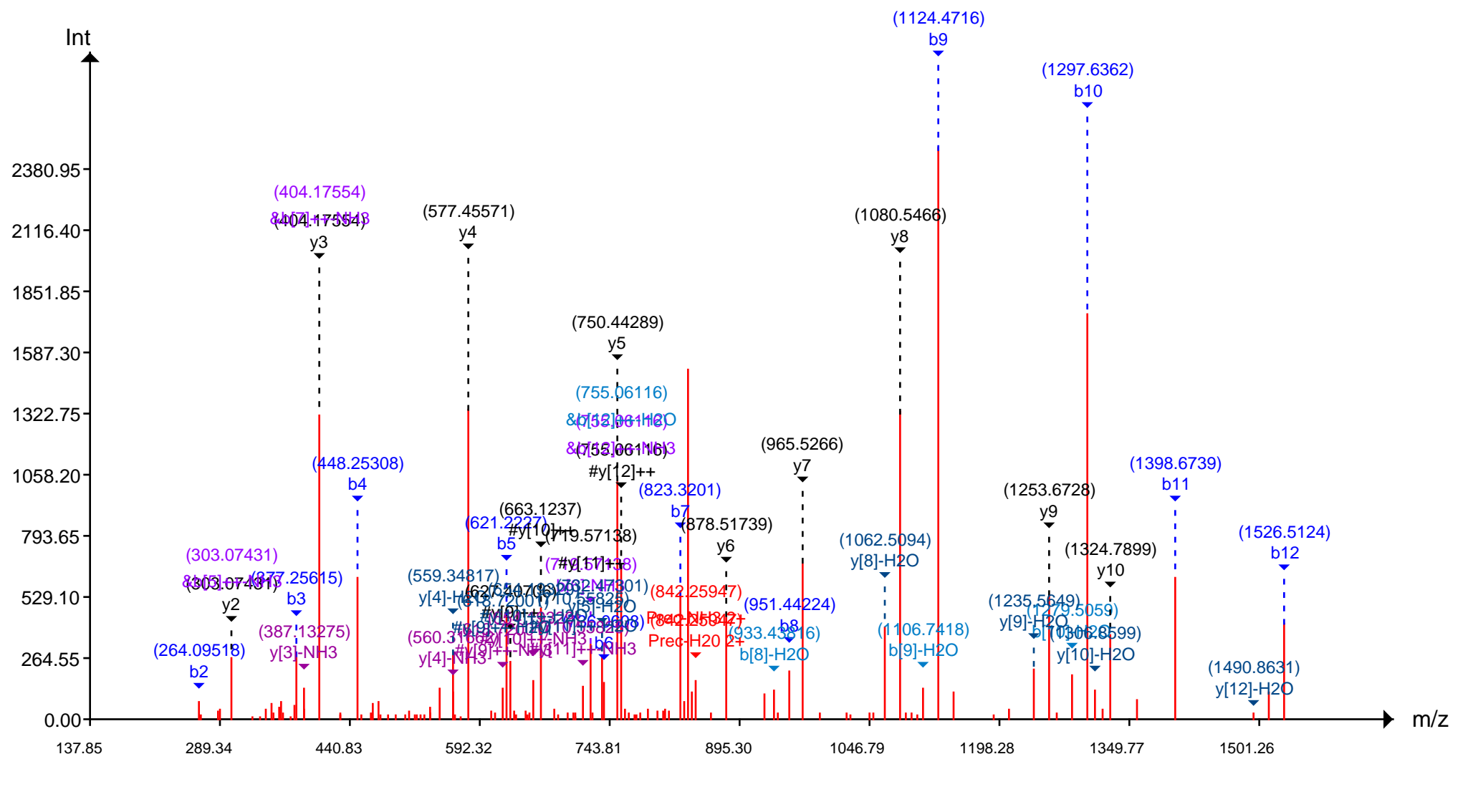
Info

Modified Sequence	AcD3-VDVPNDSSFQIVK<AcD3K*>NPR-COOH	Ion score	62.89
Protein	Q61176	Confidence level	99.00%
Precursor m/z	953.00335	Identity Threshold	48.17
Charge	2+	Mass error	0.00109

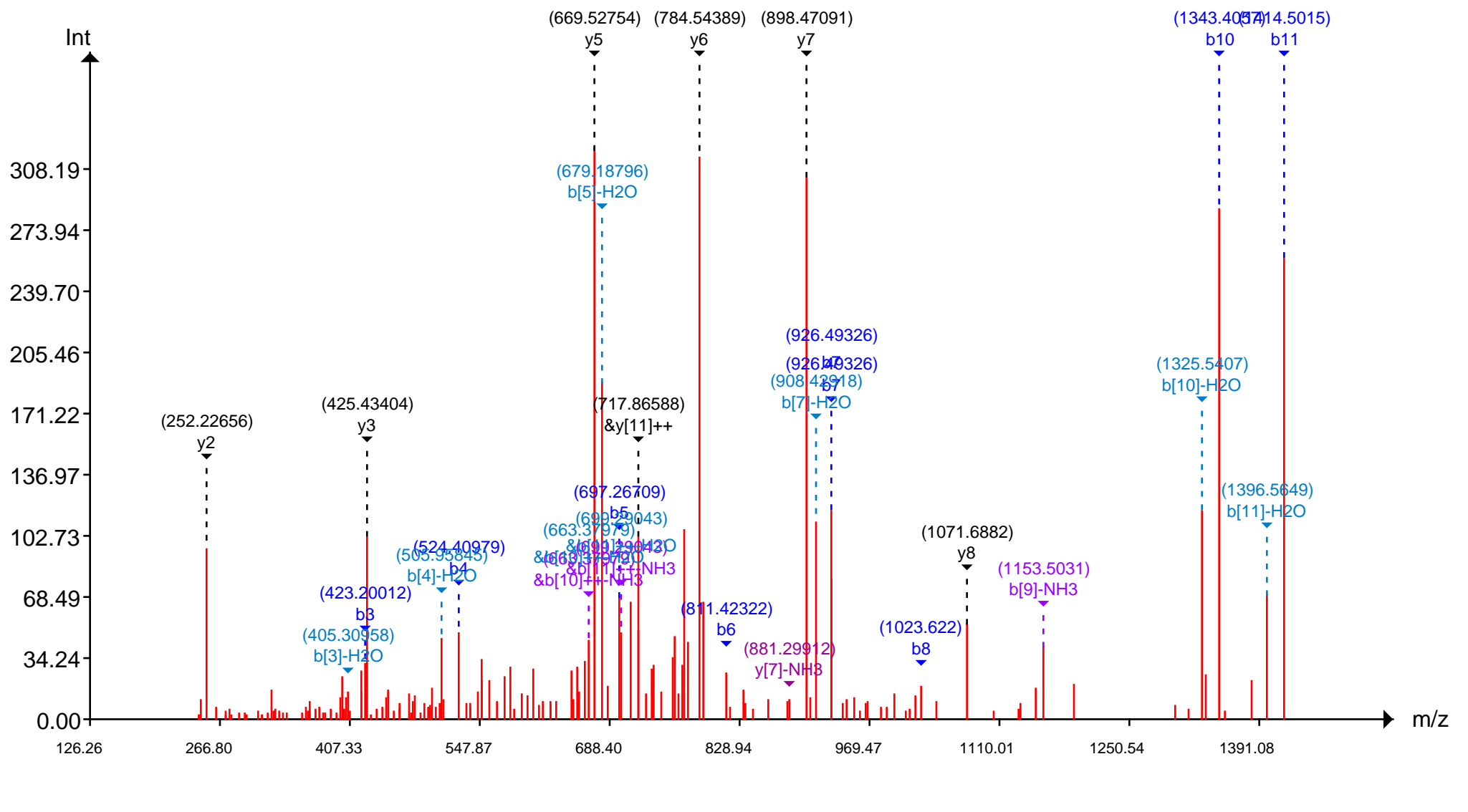


Info

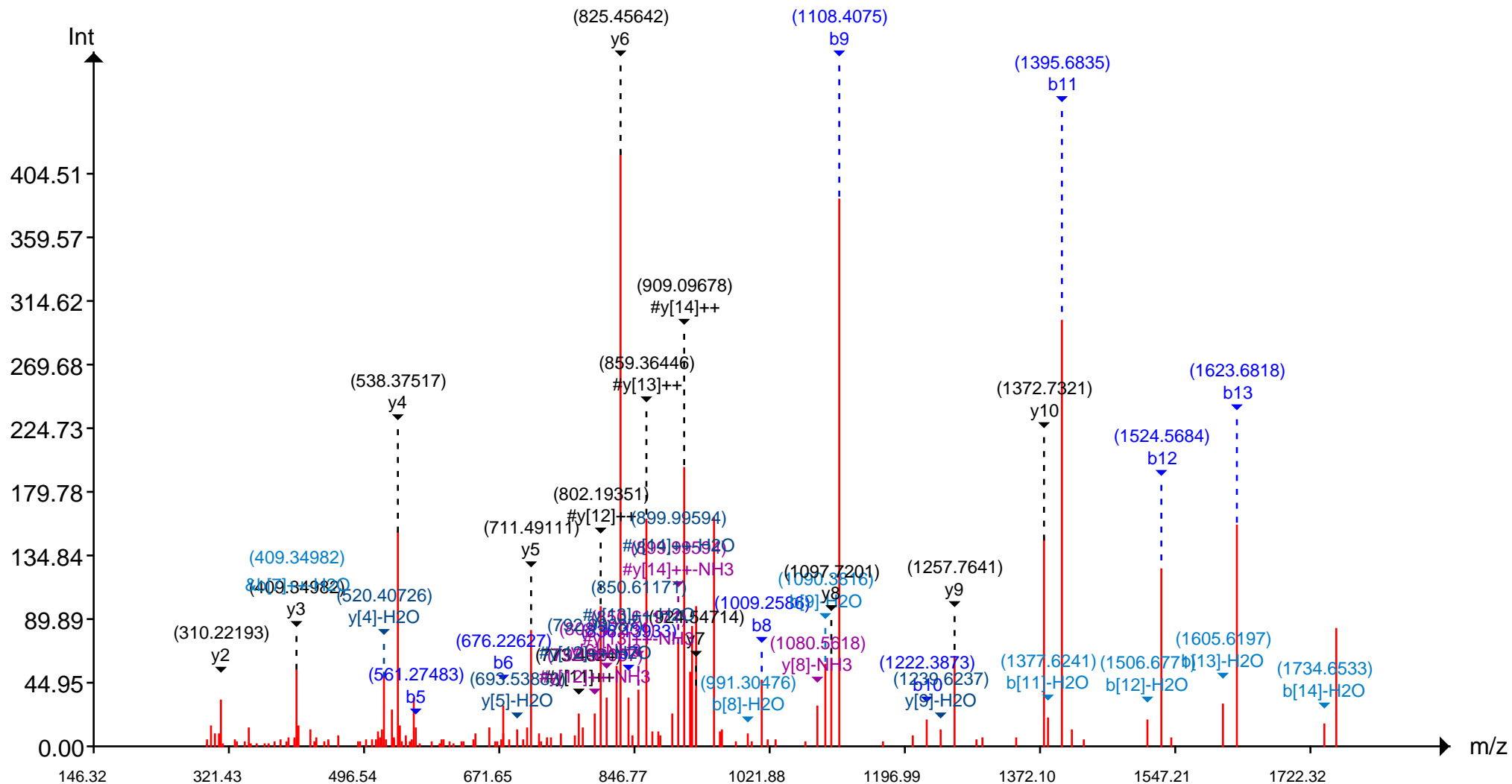
Modified Sequence	AcD3-FAIAK<AcD3K*>DSQK<AcD3K*>K<AcD3K*>TQR<C13>-COOH	Ion score	75.69
Protein	Q6PGL7	Confidence level	99.00%
Precursor m/z	854.00046	Identity Threshold	44.49
Charge	2+	Mass error	0.001581



Info			
Modified Sequence	AcD3-FAIAK<AcD3K*>DSQK<AcD3K*>K<AcD3K*>TQR-COOH	Ion score	92.82
Protein	Q6PGL7	Confidence level	99.00%
Precursor m/z	850.99068	Identity Threshold	45.43
Charge	2+	Mass error	0.002163

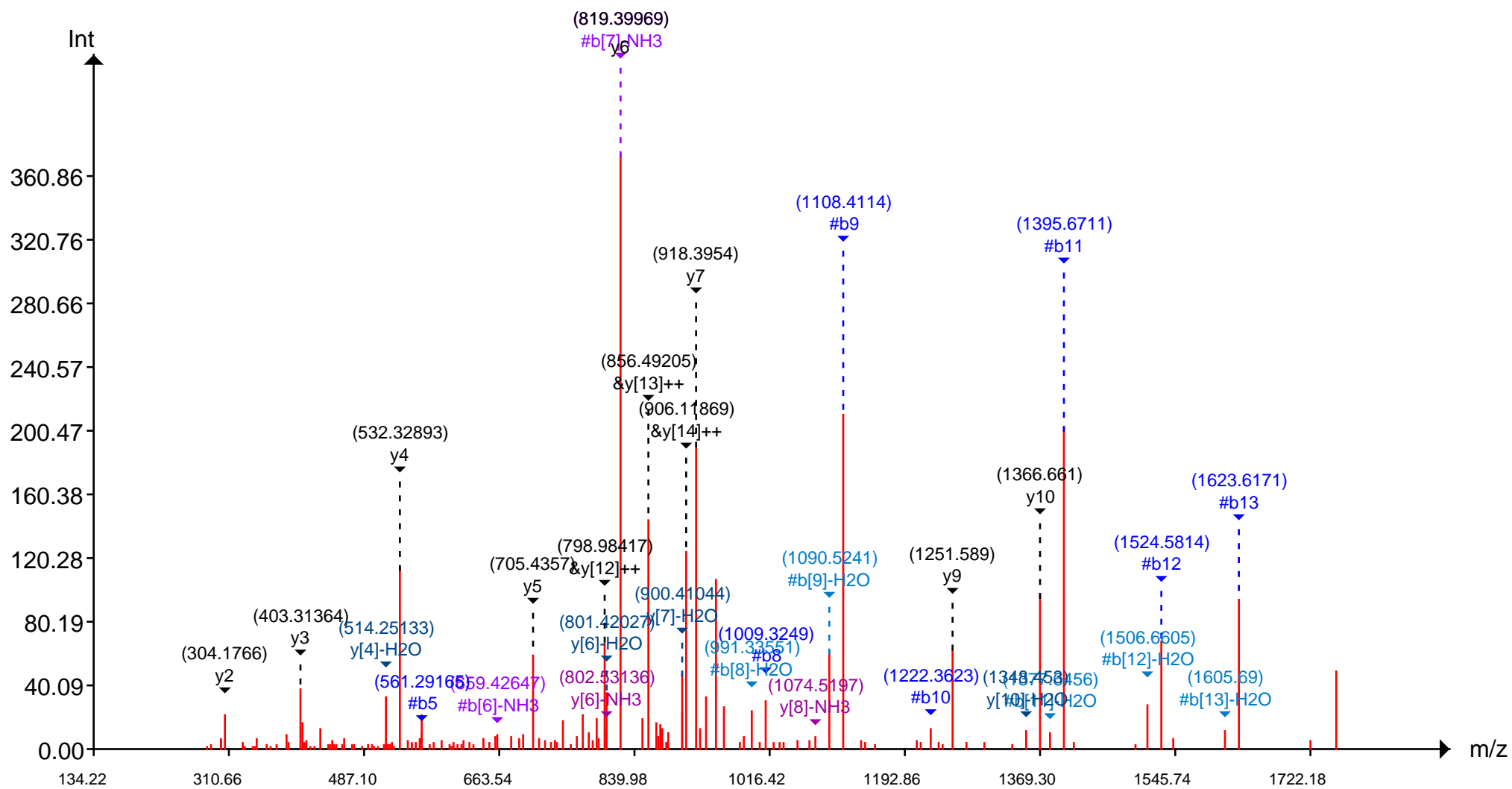


Info			
Modified Sequence	AcD3-DDM<Mox*>TK<AcD3K*>NDPFK<AcD3K*>AR<C13>-COOH	Ion score	51.58
Protein	Q62523	Confidence level	99.00%
Precursor m/z	797.89572	Identity Threshold	47.50
Charge	2+	Mass error	0.001833



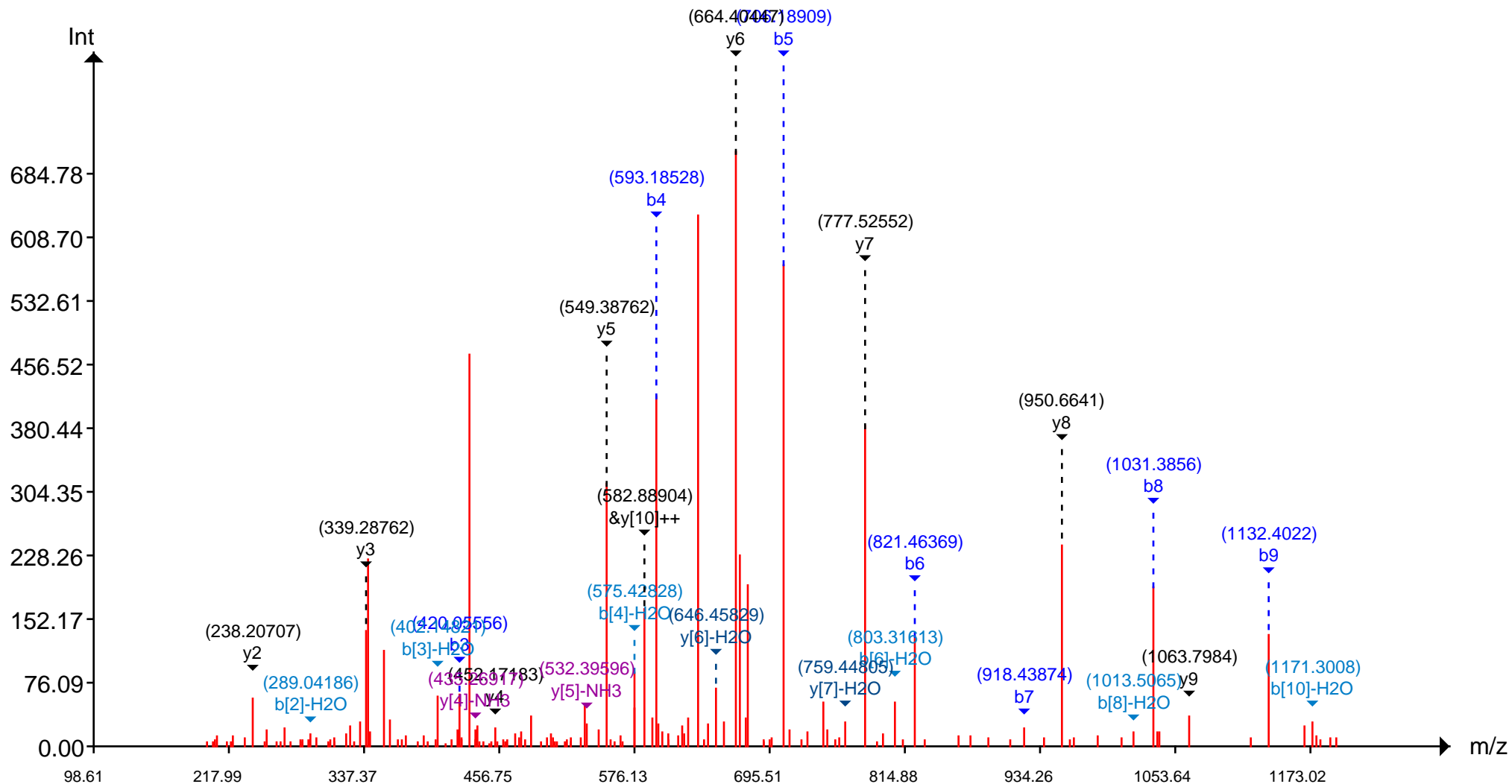
Info

Modified Sequence	AcD3-AVDGK<AcD3K*>DC<Cmm*>K<AcD3K*>VNK<AcD3K*>EVER<C13>-COOH	Ion score	62.53
Protein	Q9D172	Confidence level	99.00%
Precursor m/z	967.01364	Identity Threshold	48.02
Charge	2+	Mass error	0.001971



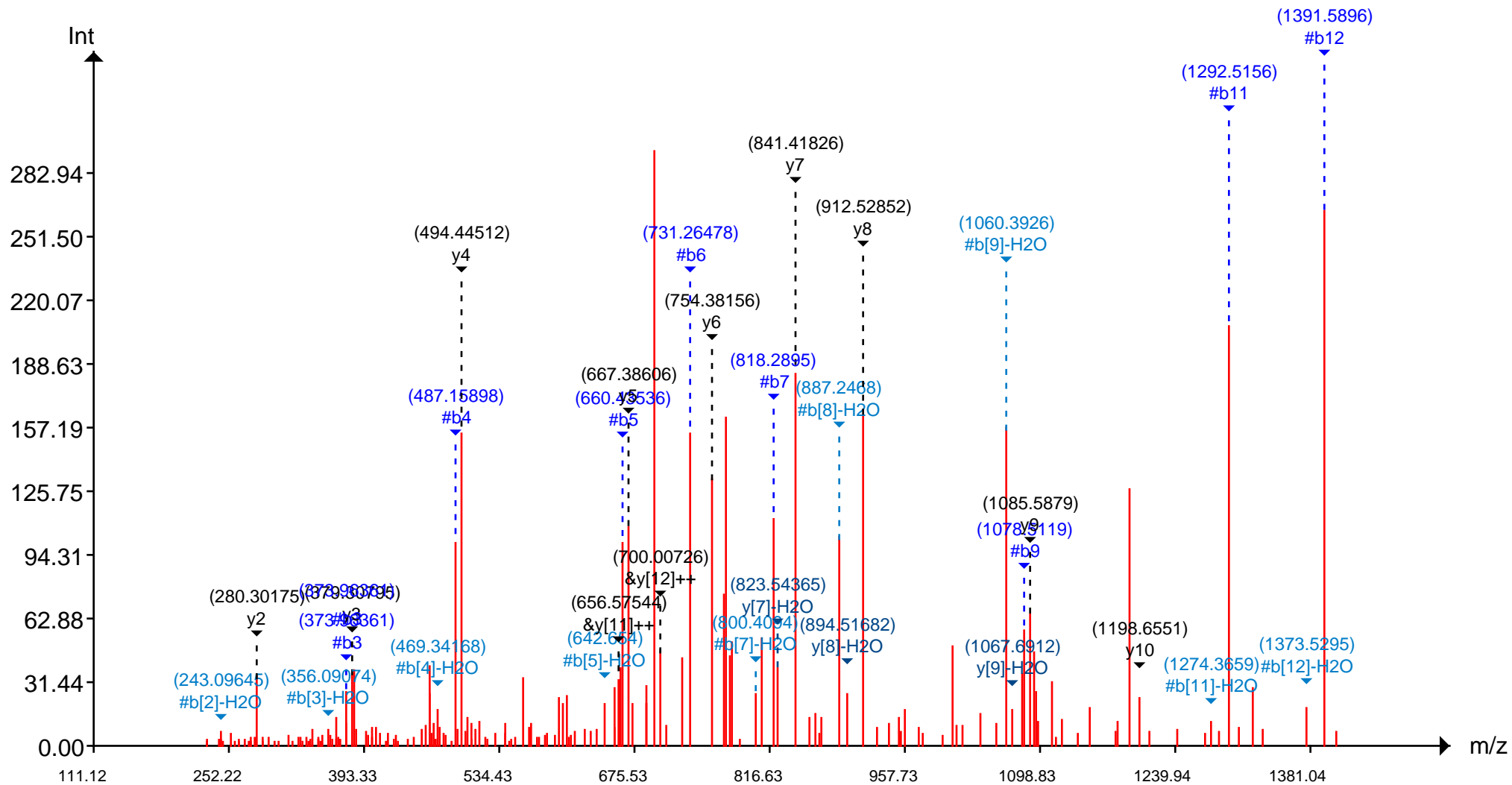
Info

Modified Sequence	AcD3-AVDGK<AcD3K*>DC<Cmm*>K<AcD3K*>VNK<AcD3K*>EVER-COOH	Ion score	72.99
Protein	Q9D172	Confidence level	99.00%
Precursor m/z	964.00357	Identity Threshold	48.13
Charge	2+	Mass error	0.001972



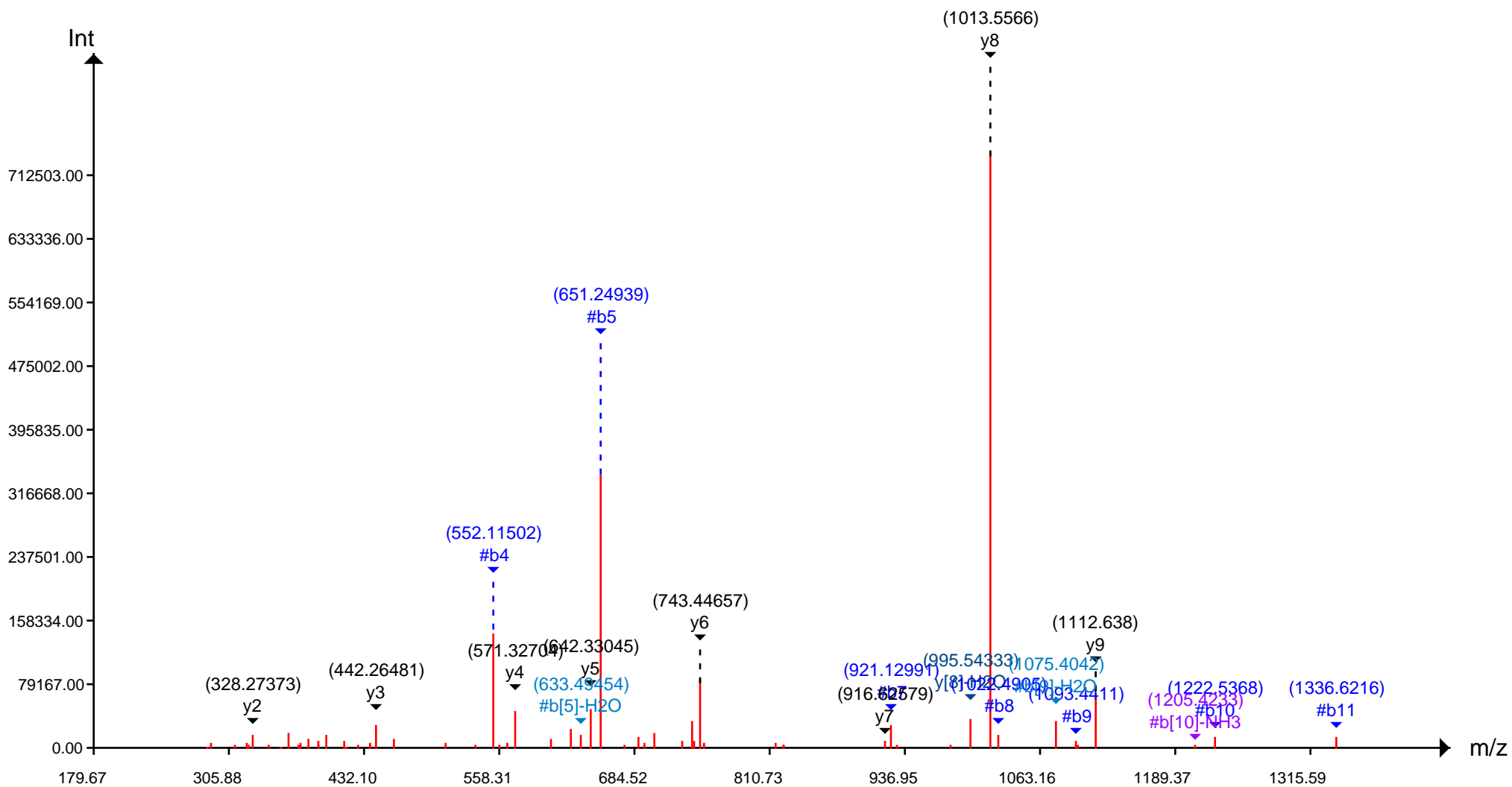
Info

Modified Sequence	AcD3-C<Cmm*>TLK<AcD3K*>LDPITGR<C13>-COOH	Ion score	66.78
Protein	Q60668	Confidence level	99.00%
Precursor m/z	685.39015	Identity Threshold	46.81
Charge	2+	Mass error	8.1E-4



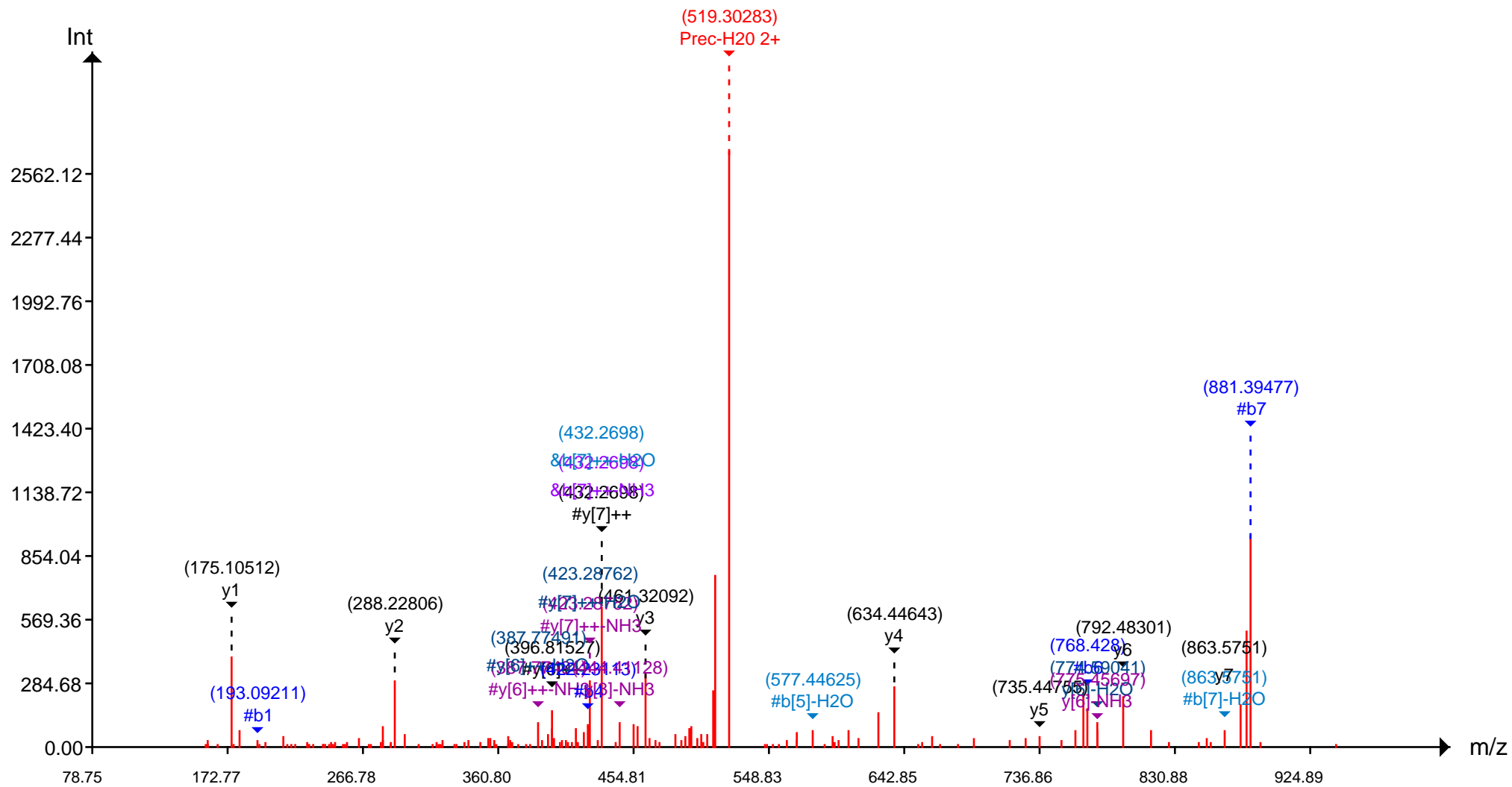
Info

Modified Sequence	AcD3-QSLLK<AcD3K*>ASSK<AcD3K*>DVVR<C13>-COOH	Ion score	58.74
Protein	Q8K2Q0	Confidence level	99.00%
Precursor m/z	786.47499	Identity Threshold	42.90
Charge	2+	Mass error	0.001834



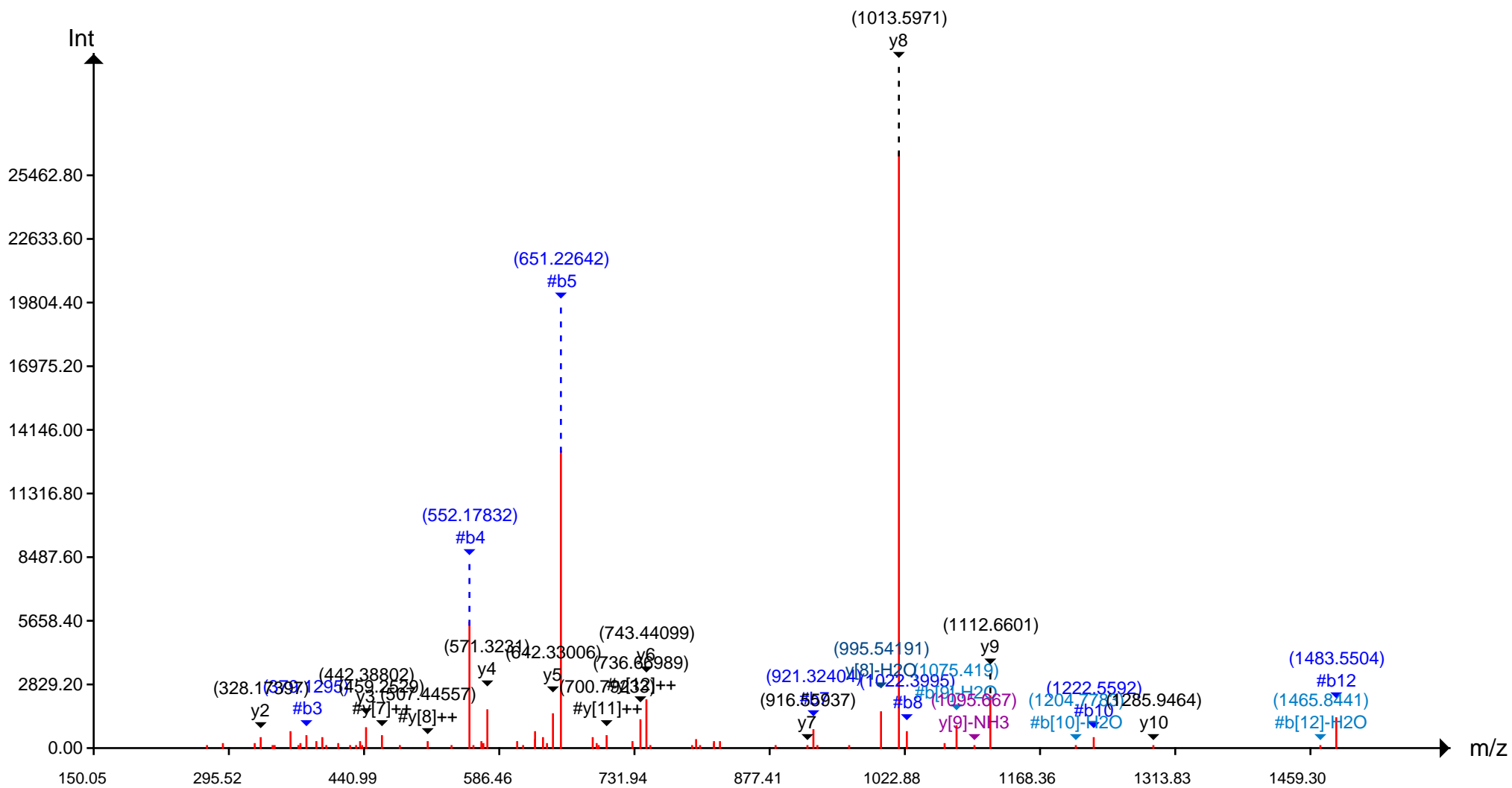
Info

Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH	Ion score	68.11
Protein	P17742	Confidence level	99.00%
Precursor m/z	832.45809	Identity Threshold	47.55
Charge	2+	Mass error	-6.7E-4



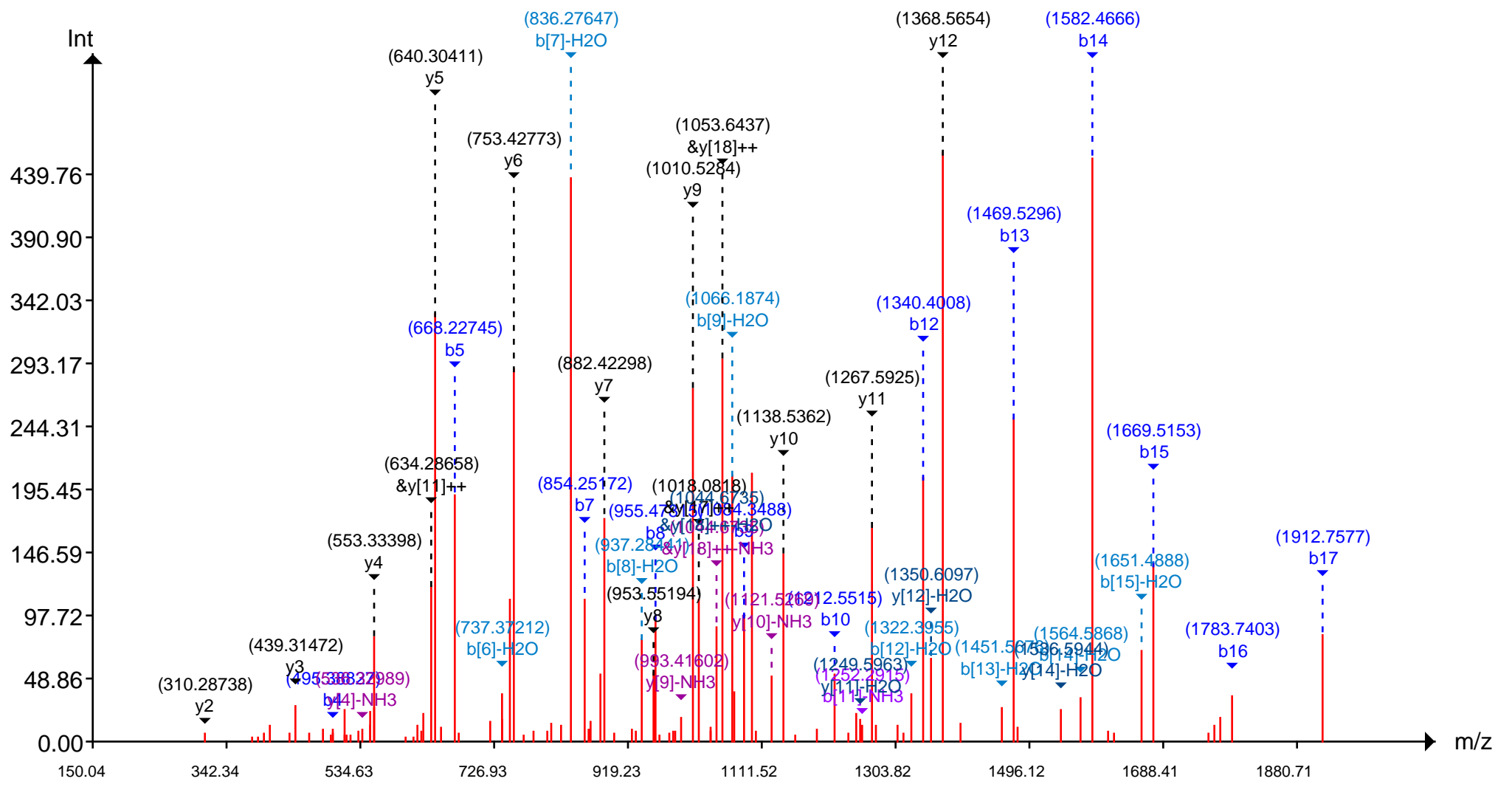
Info

Modified Sequence	AcD3-FAGTK<AcD3K*>K<AcD3K*>IR-COOH	Ion score	45.09
Protein	Q6PGL7	Confidence level	99.00%
Precursor m/z	528.33186	Identity Threshold	42.64
Charge	2+	Mass error	6.54E-4

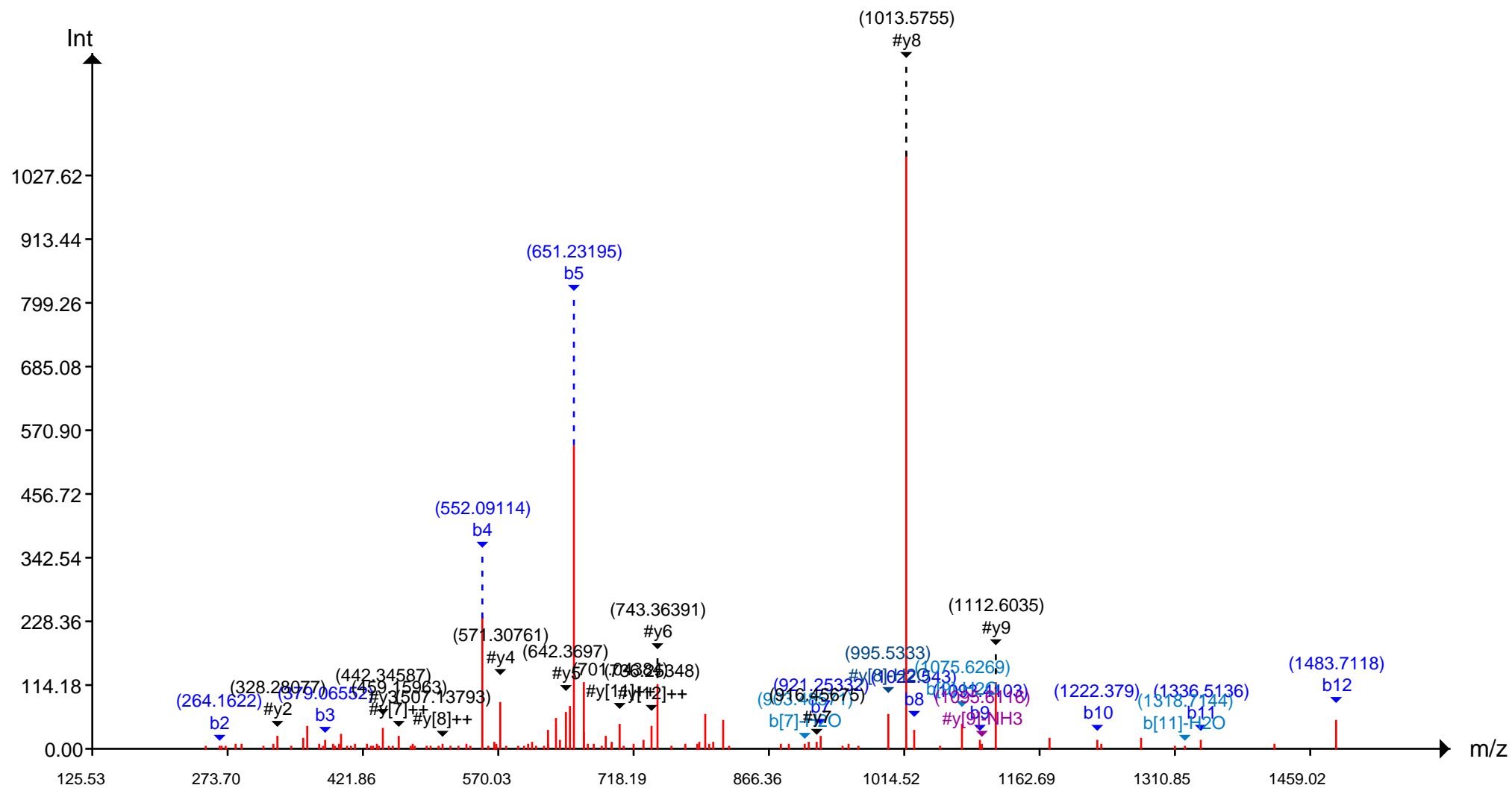


Info

Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH	Ion score	76.43
Protein	P17742	Confidence level	99.00%
Precursor m/z	832.45862	Identity Threshold	47.56
Charge	2+	Mass error	3.9E-4

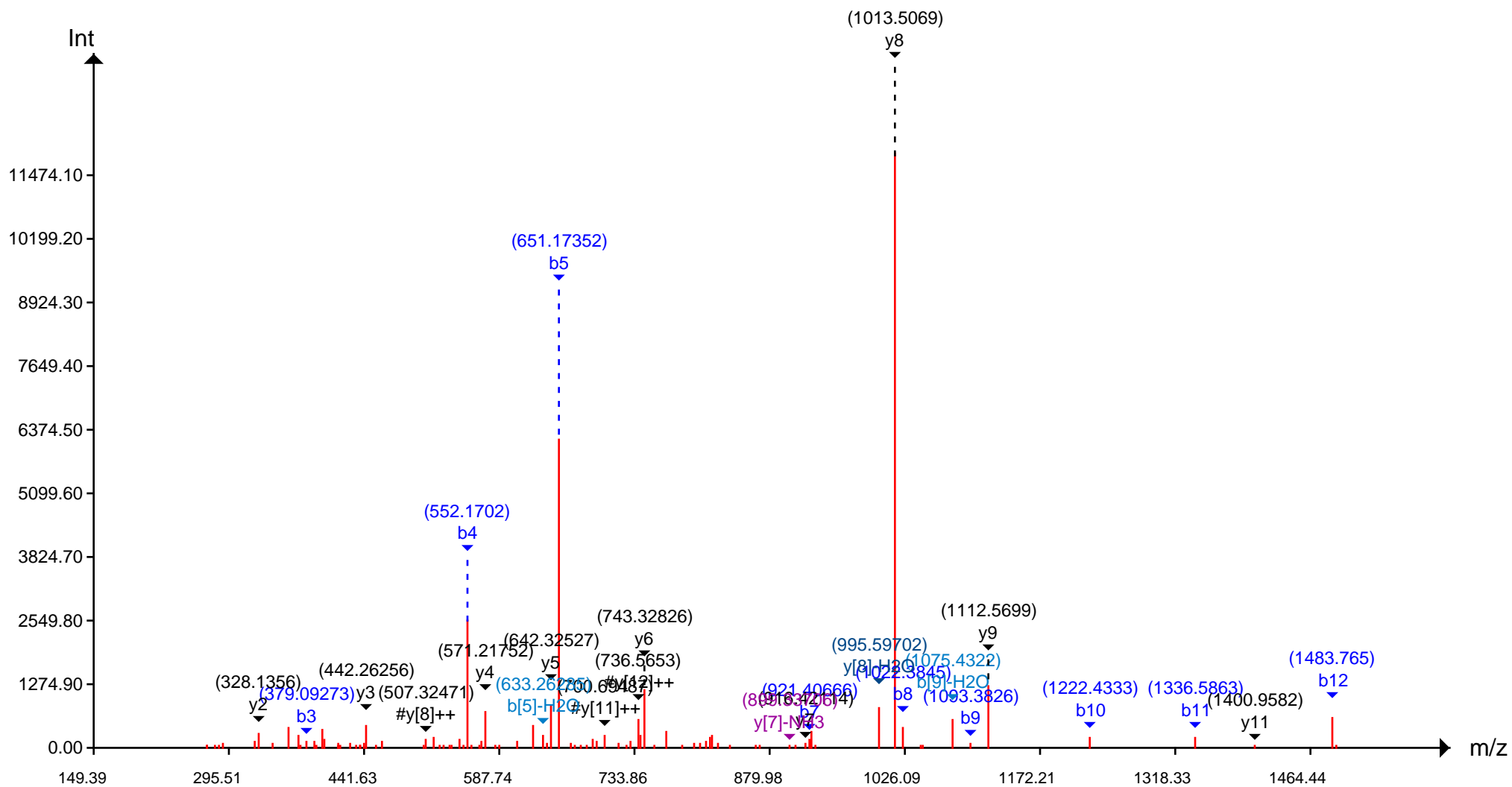


Info			
Modified Sequence	AcD3-AAC<Cmm*>M<Mox*>K<AcD3K*>SVTEQGAELSNEER<C13>-COOH	Ion score	84.1
Protein	P63101	Confidence level	99.00%
Precursor m/z	1111.5186	Identity Threshold	45.40
Charge	2+	Mass error	0.002171



Info

Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH	Ion score	54.97
Protein	P17742	Confidence level	99.00%
Precursor m/z	832.45899	Identity Threshold	47.47
Charge	2+	Mass error	0.00113



Info

Modified Sequence AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH

Ion score 70.17

Protein P17742

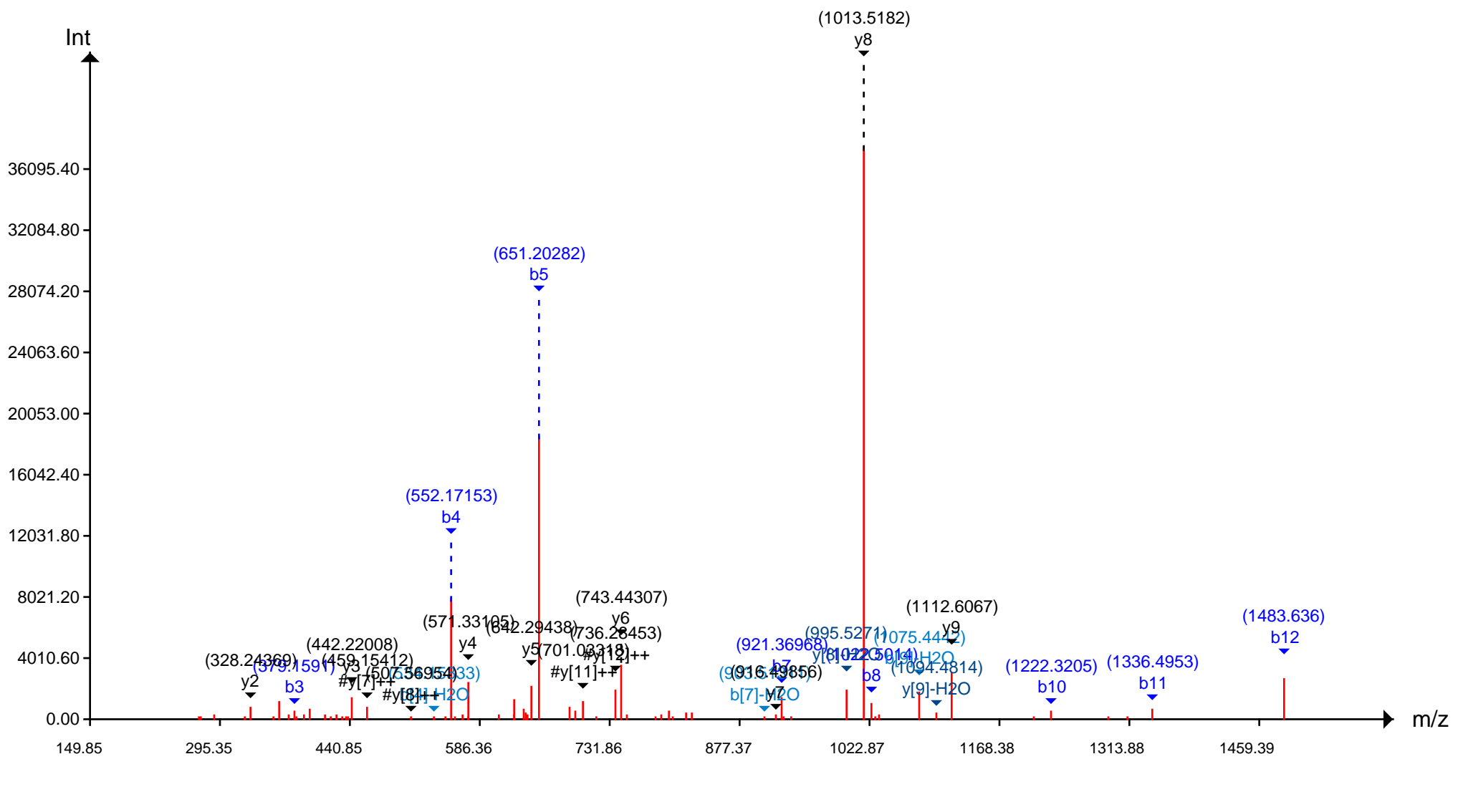
Confidence level 99.00%

Precursor m/z 832.45912

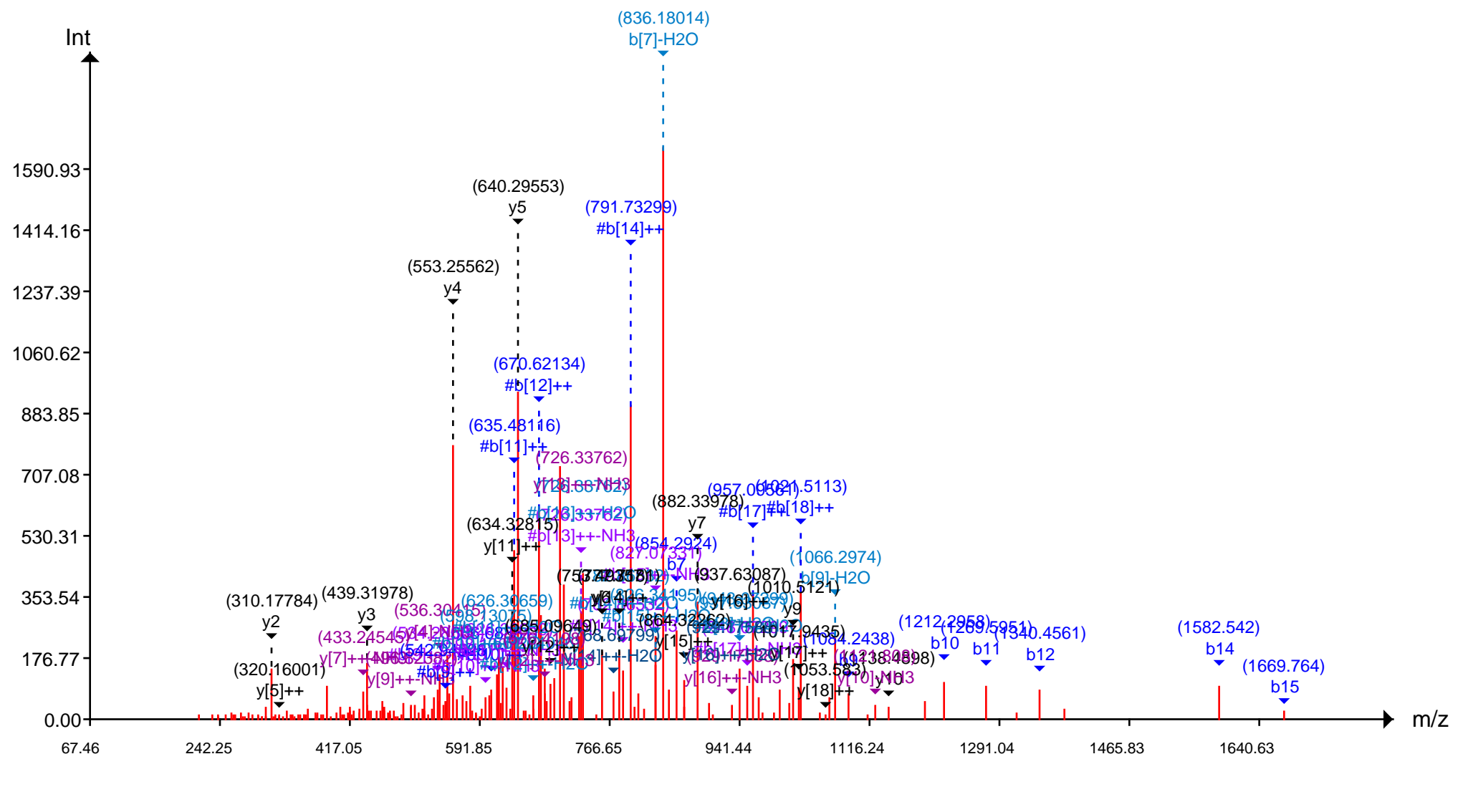
Identity Threshold 47.45

Charge 2+

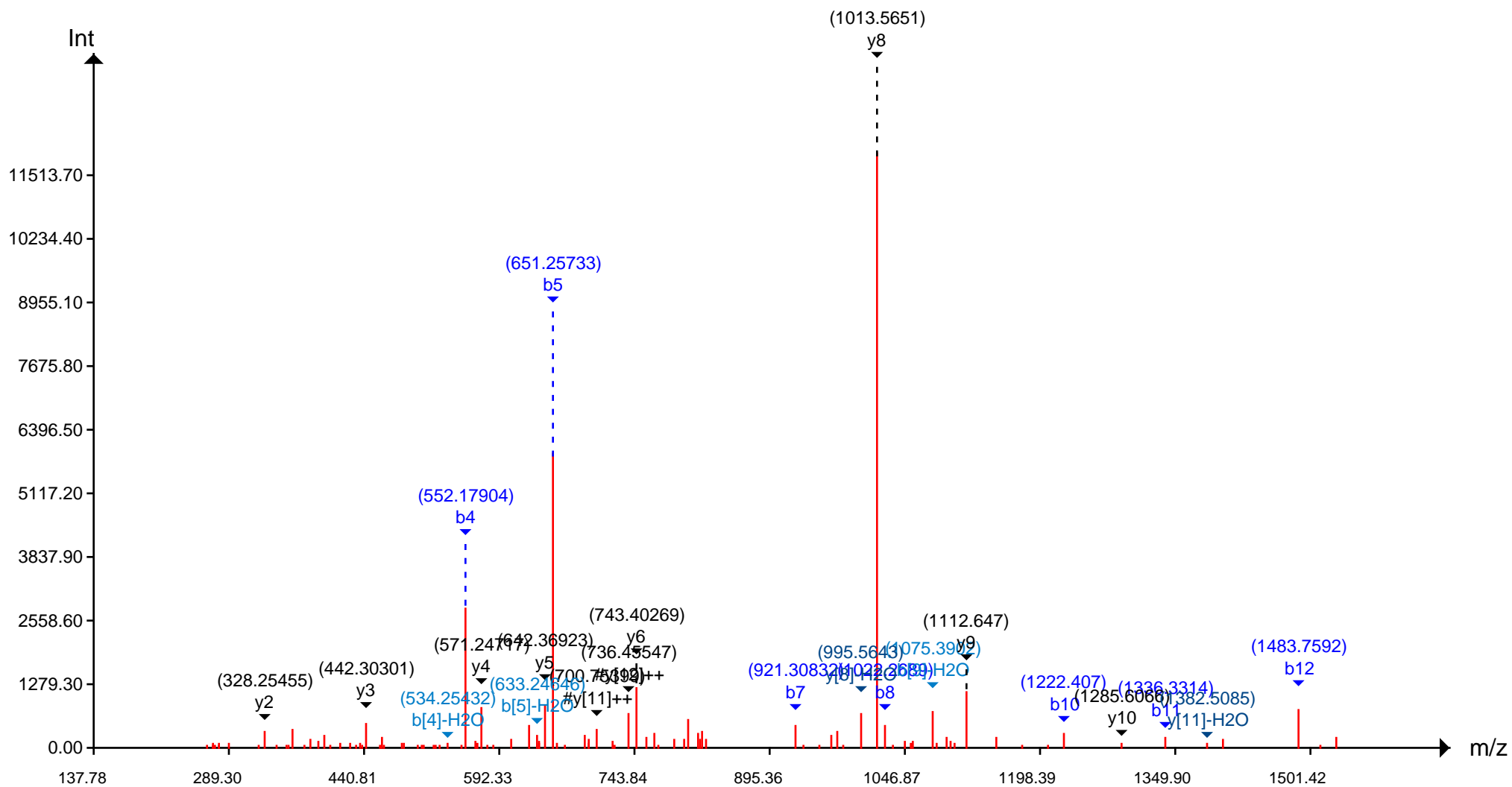
Mass error 0.00139



Info			
Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH	Ion score	58.4
Protein	P17742	Confidence level	99.00%
Precursor m/z	832.45939	Identity Threshold	47.48
Charge	2+	Mass error	0.00193



Info			
Modified Sequence	AcD3-AAC<Cmm*>M<Mox*>K<AcD3K*>SVTEQGAELSNEER<C13>-COOH	Ion score	47.5
Protein	P63101	Confidence level	99.00%
Precursor m/z	741.34763	Identity Threshold	45.23
Charge	3+	Mass error	5.85E-4



Info

Modified Sequence AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH

Ion score 48.42

Protein P17742

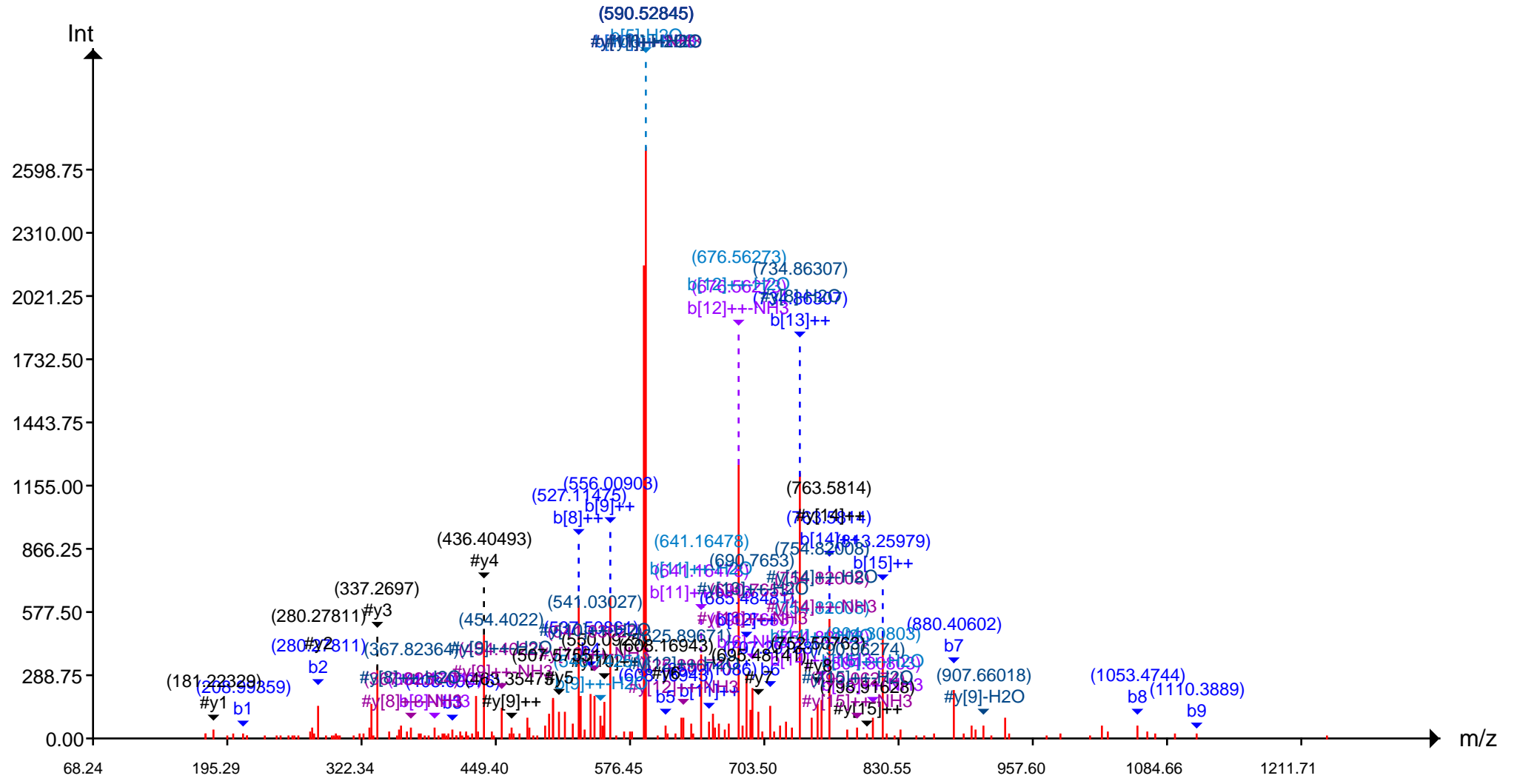
Confidence level 99.00%

Precursor m/z 832.4594

Identity Threshold 47.48

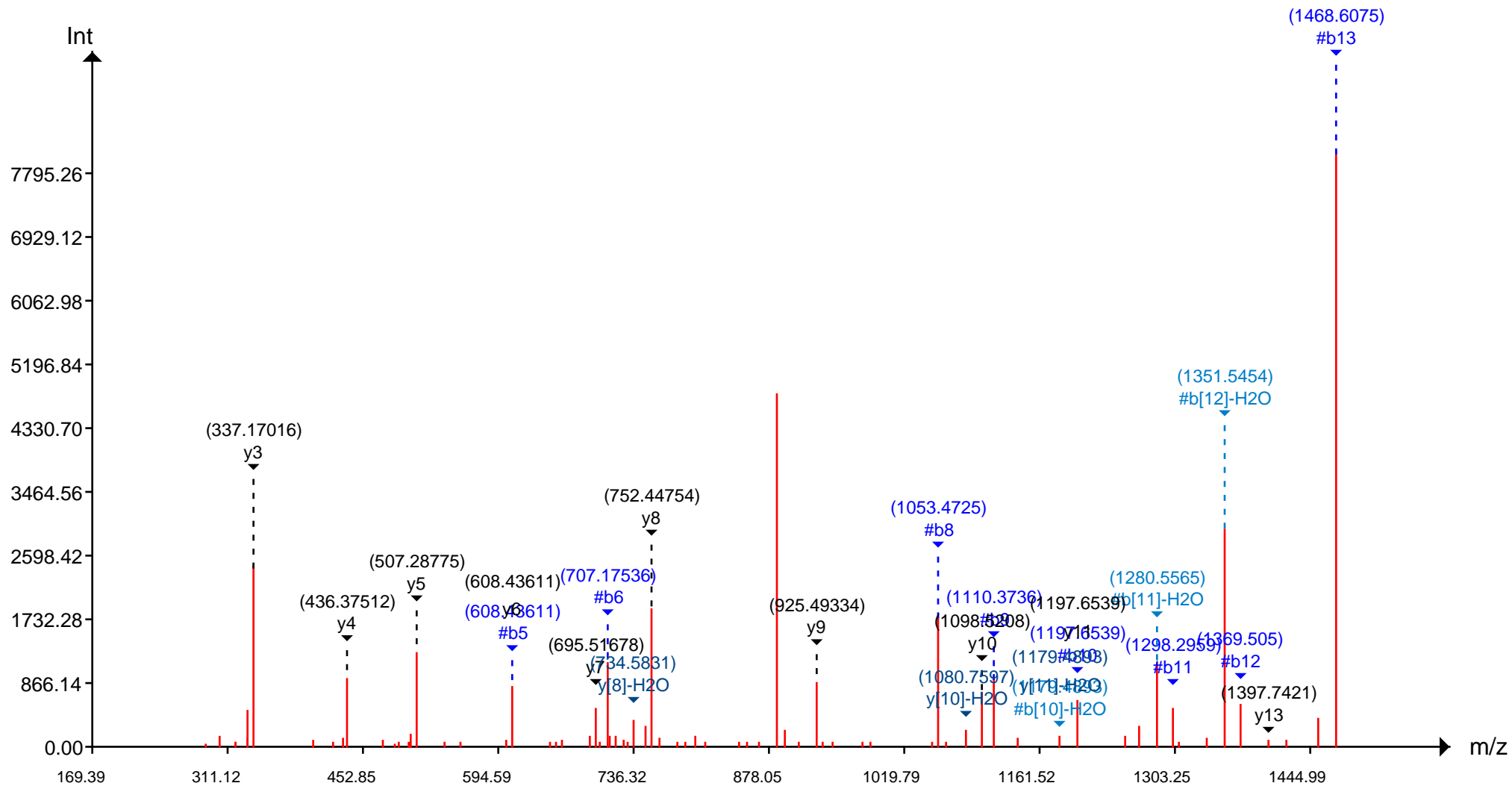
Charge 2+

Mass error 0.00195



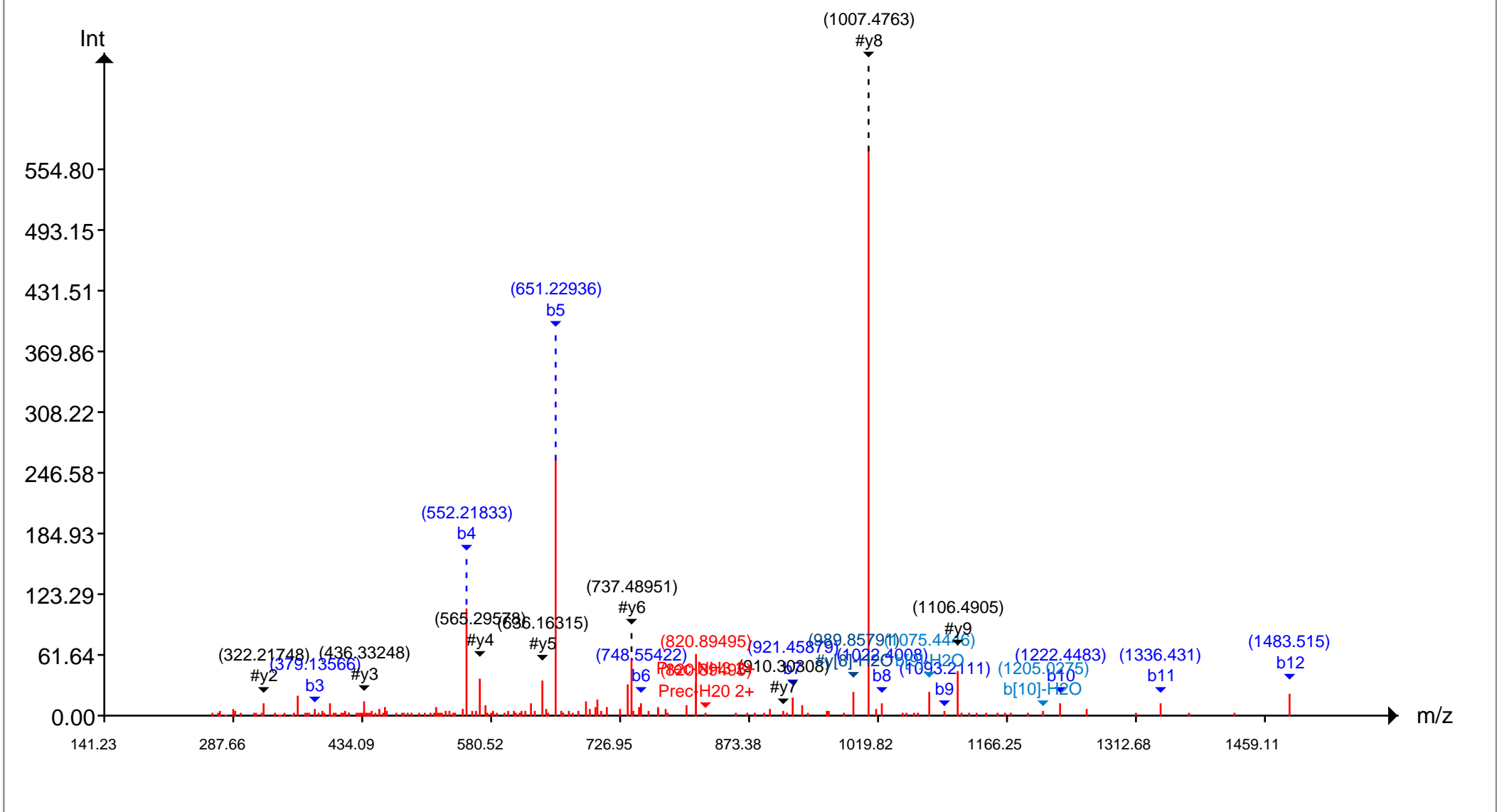
Info

Modified Sequence	AcD3-YAQEAVK<AcD3K*>K<AcD3K*>GSTAVGVR<C13>-COOH	Ion score	46.32
Protein	Q9Z2U0	Confidence level	99.00%
Precursor m/z	602.34506	Identity Threshold	45.95
Charge	3+	Mass error	-2.89E-4



Info

Modified Sequence	AcD3-YAQEAVK<AcD3K*>K<AcD3K*>GSTAVGVR<C13>-COOH	Ion score	66.14
Protein	Q9Z2U0	Confidence level	99.00%
Precursor m/z	903.01433	Identity Threshold	45.92
Charge	2+	Mass error	4.67E-4



Info

Modified Sequence AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH

Ion score 54.0

Protein P17742

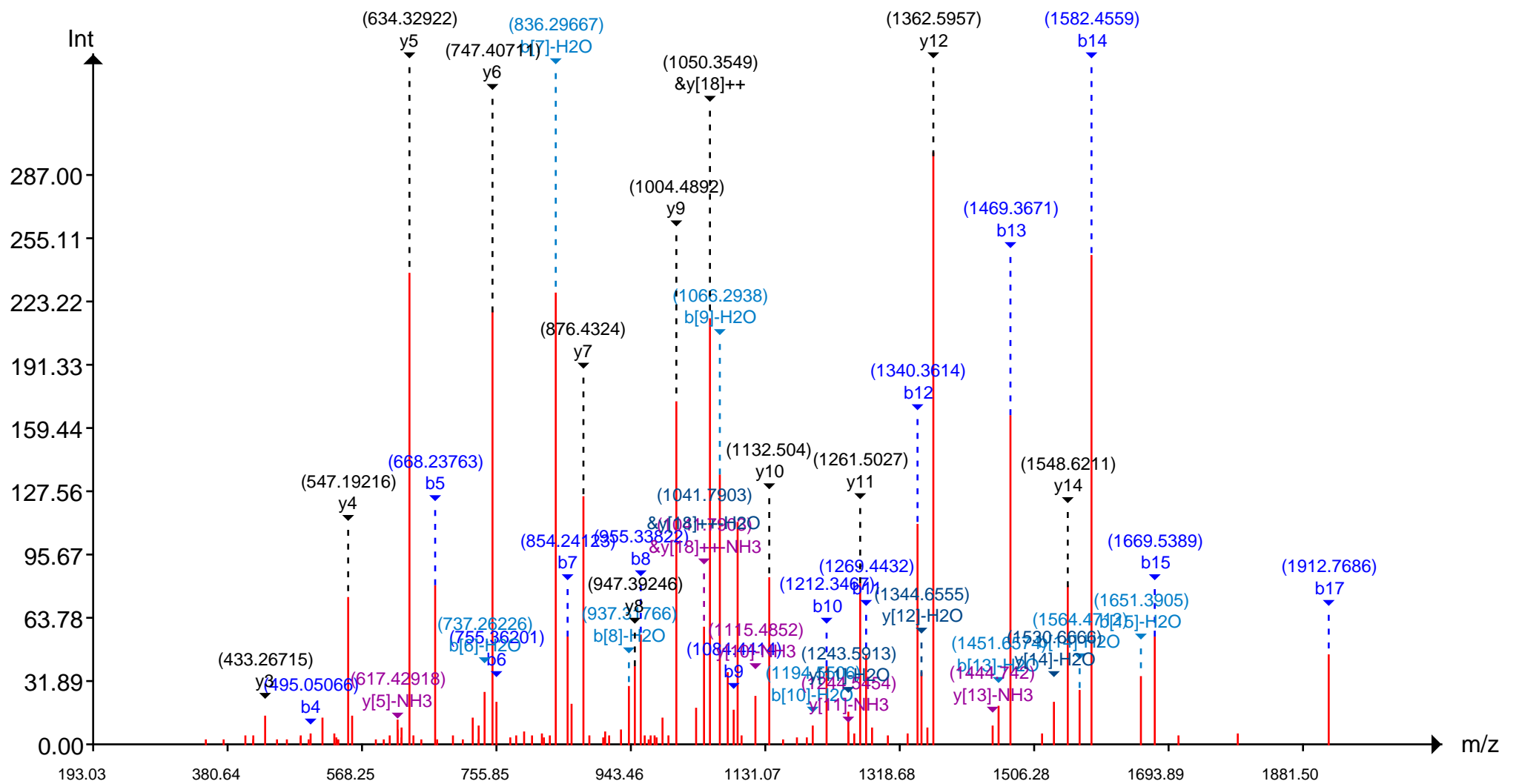
Confidence level 99.00%

Precursor m/z 829.44962

Identity Threshold 47.77

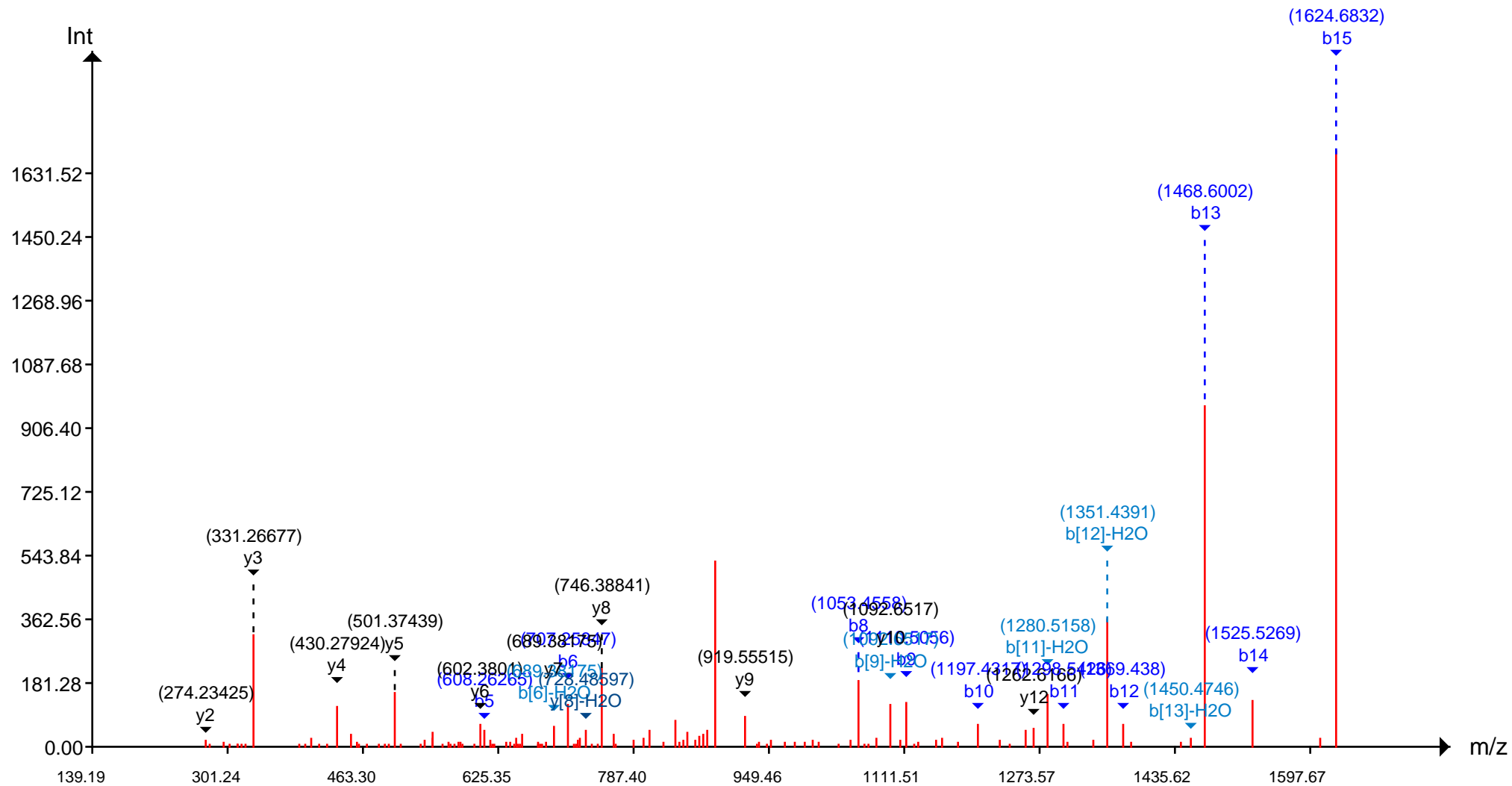
Charge 2+

Mass error 0.002532



Info

Modified Sequence	AcD3-AAC<Cmm*>M<Mox*>K<AcD3K*>SVTEQGAELSNEER-COOH	Ion score	95.05
Protein	P63101	Confidence level	99.00%
Precursor m/z	1108.5089	Identity Threshold	44.48
Charge	2+	Mass error	0.002912



Info

Modified Sequence AcD3-YAQEAVK<AcD3K*>K<AcD3K*>GSTAVGVR-COOH

Ion score 87.78

Protein Q9Z2U0

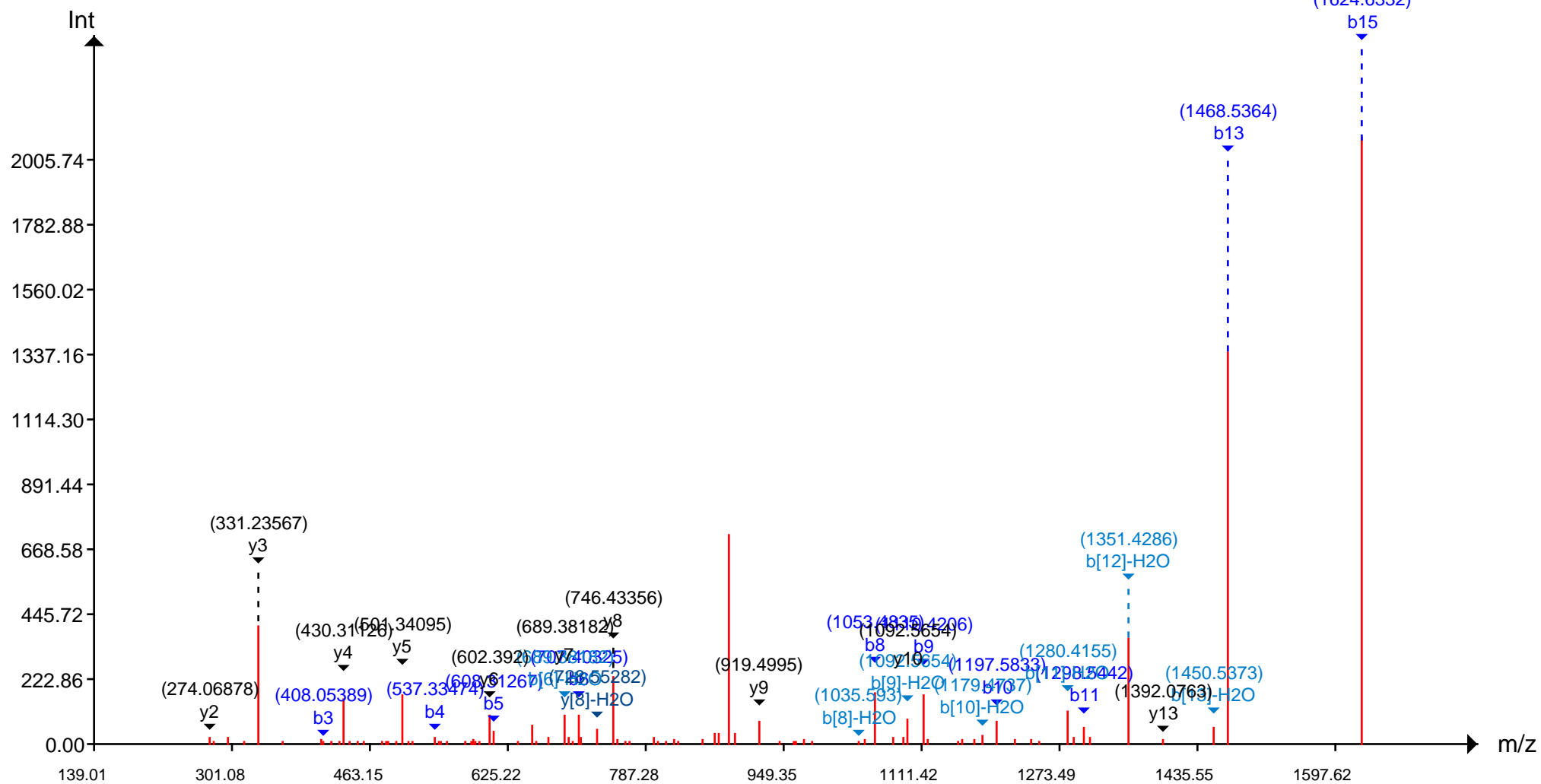
Confidence level 99.00%

Precursor m/z 900.00544

Identity Threshold 46.61

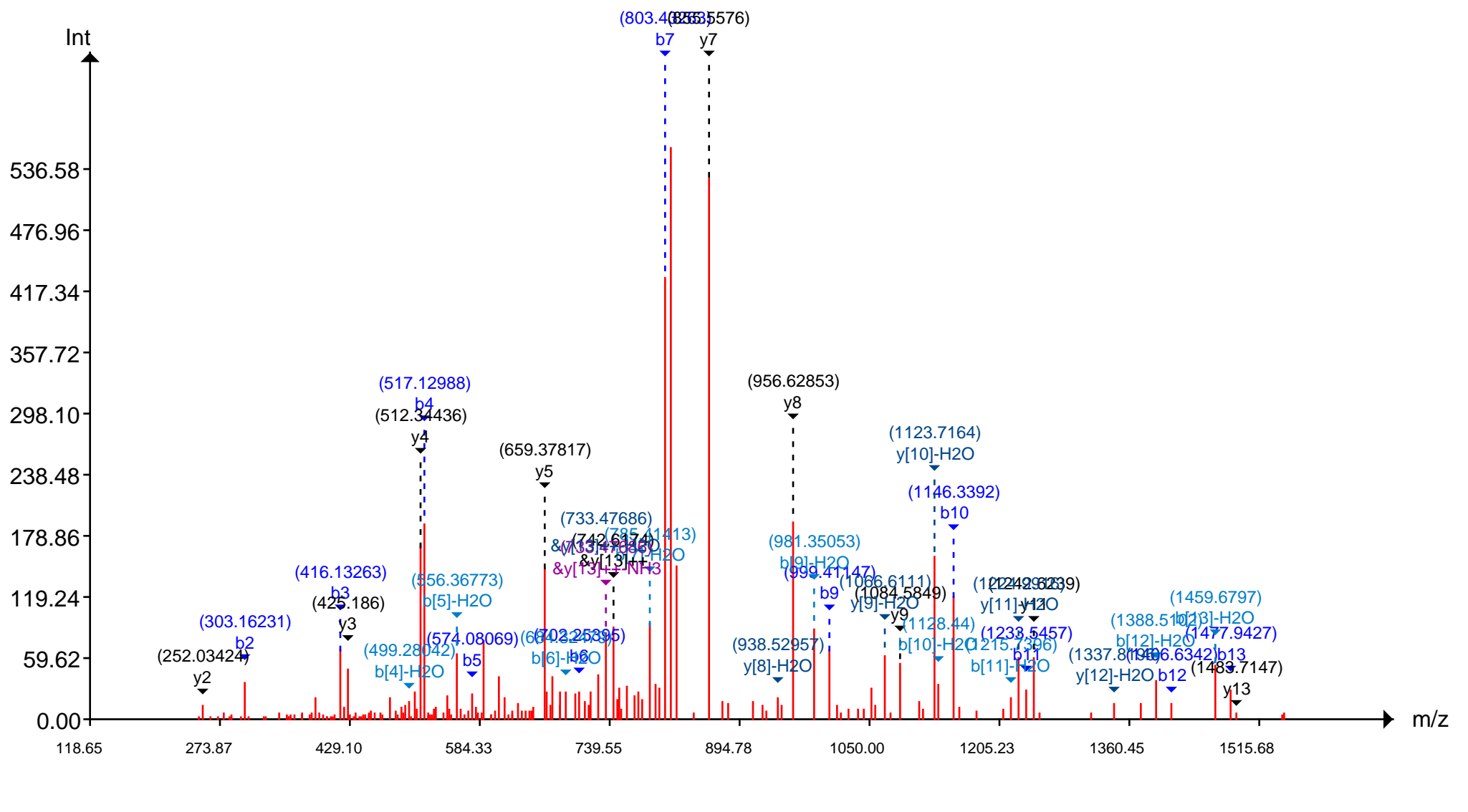
Charge 2+

Mass error 0.002828

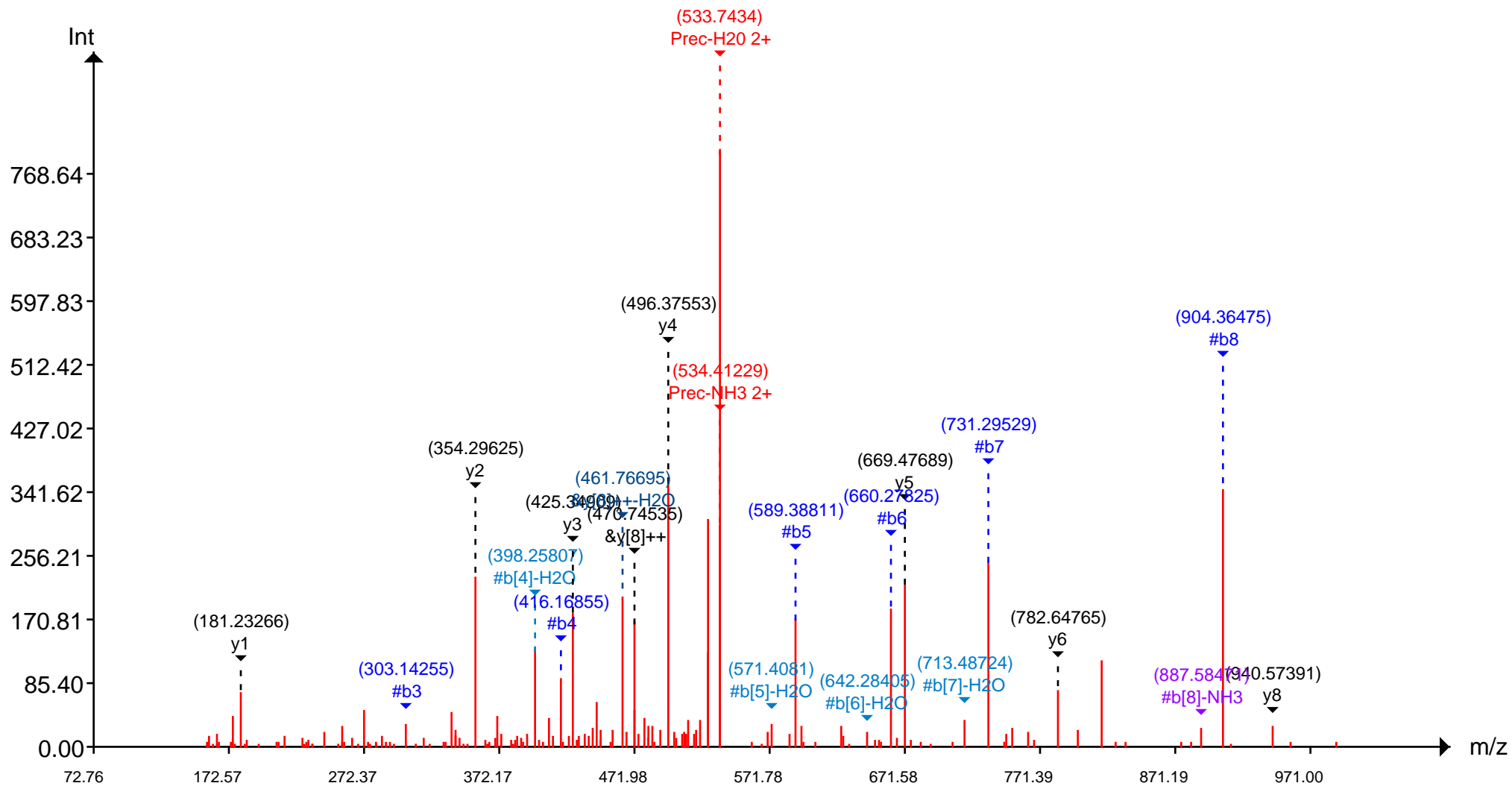


Info

Modified Sequence	AcD3-YAQEAVK<AcD3K*>K<AcD3K*>GSTAVGVR-COOH	Ion score	91.52
Protein	Q9Z2U0	Confidence level	99.00%
Precursor m/z	900.00544	Identity Threshold	46.61
Charge	2+	Mass error	0.002828

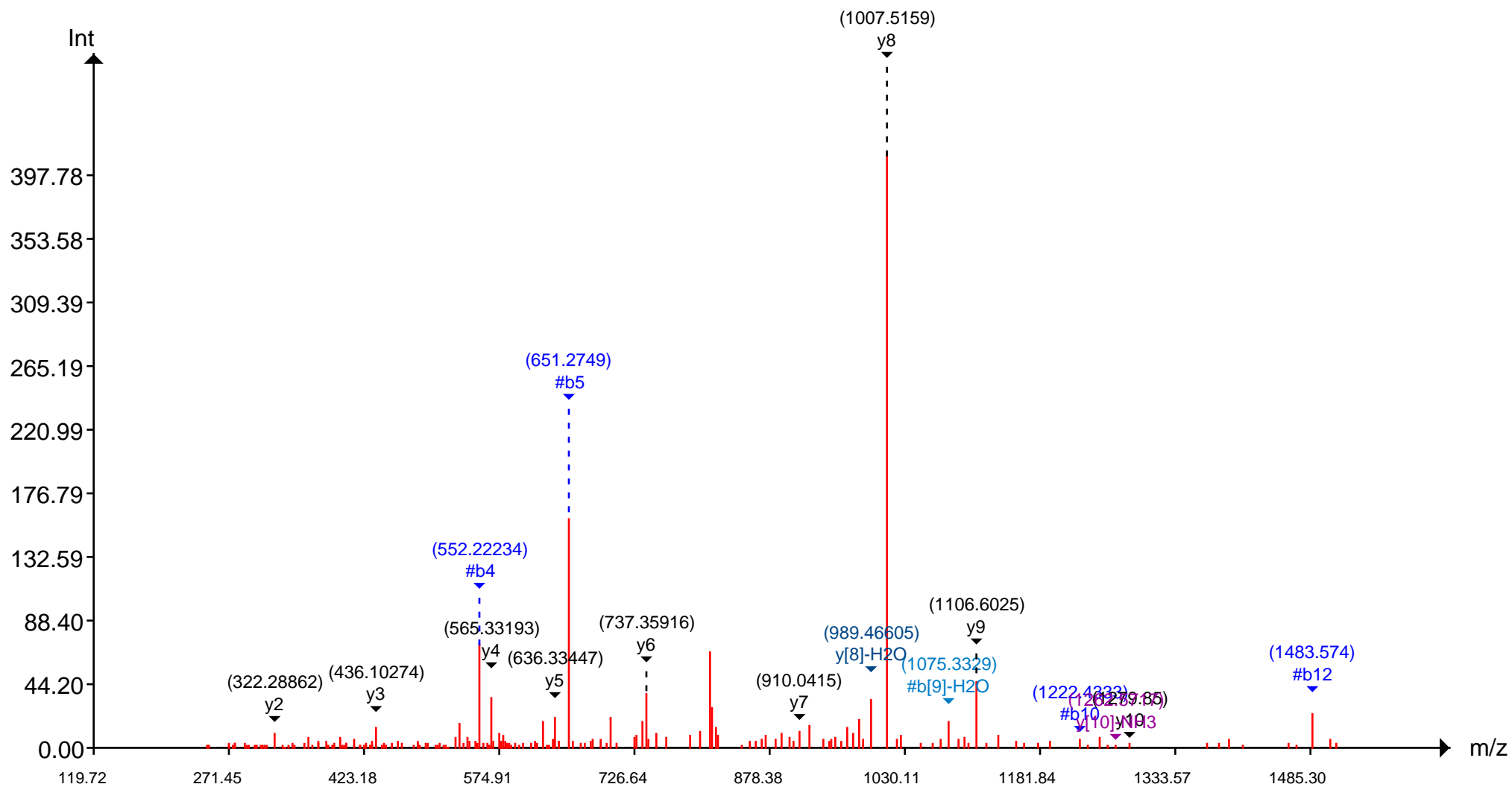


Info			
Modified Sequence	AcD3-EQLTGQTPVFSK<AcD3K*>AR<C13>-COOH	Ion score	76.19
Protein	Q9CXW4	Confidence level	99.00%
Precursor m/z	829.46059	Identity Threshold	47.26
Charge	2+	Mass error	0.00169



Info

Modified Sequence	AcD3-VGTLK<AcD3K*>AAK<AcD3K*>R<C13>-COOH	Ion score	58.72
Protein	Q4KML4	Confidence level	99.00%
Precursor m/z	542.86017	Identity Threshold	38.40
Charge	2+	Mass error	2.3E-5



Info

Modified Sequence AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH

Ion score 55.2

Protein P17742

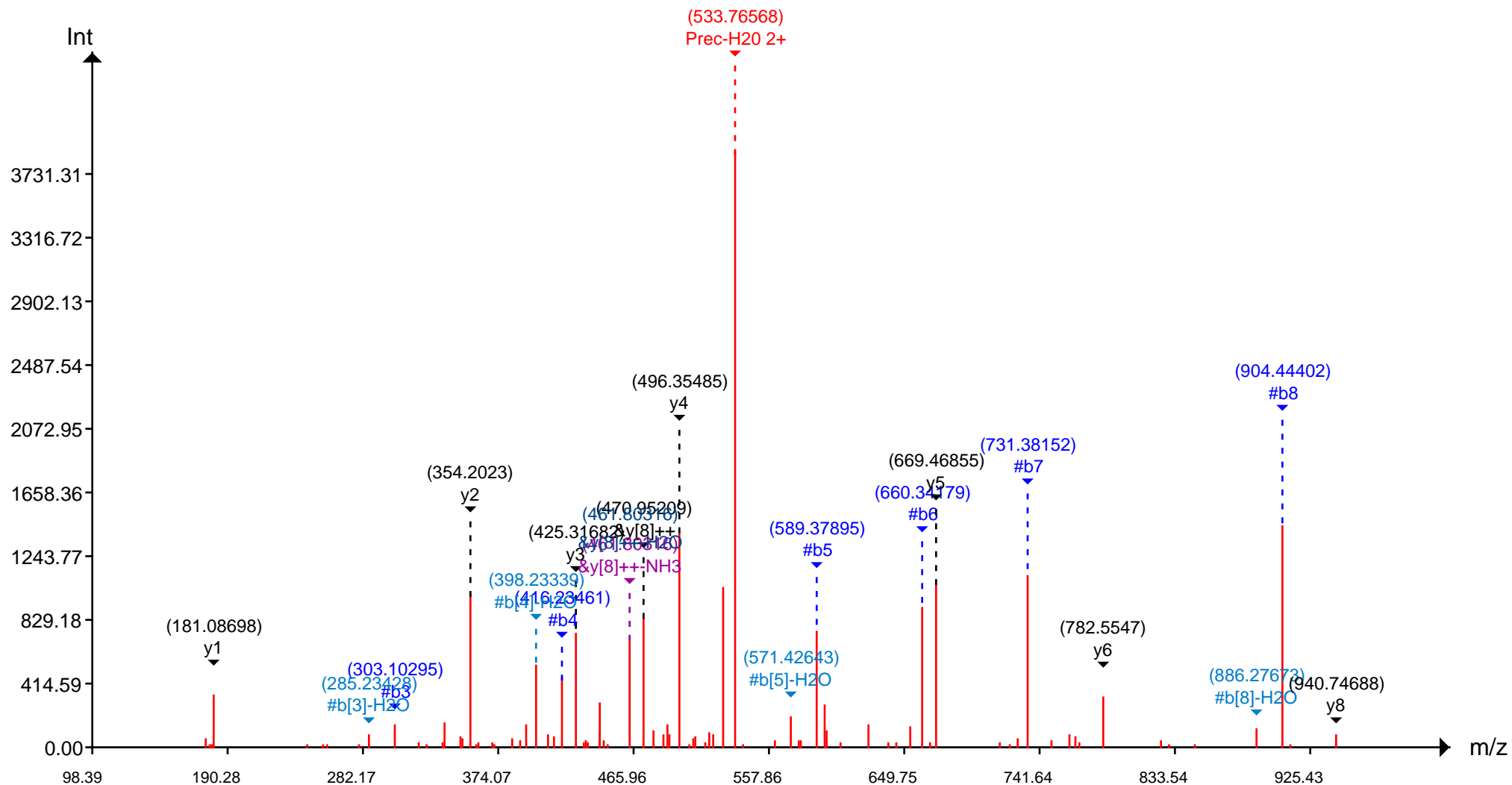
Confidence level 99.00%

Precursor m/z 829.44906

Identity Threshold 47.80

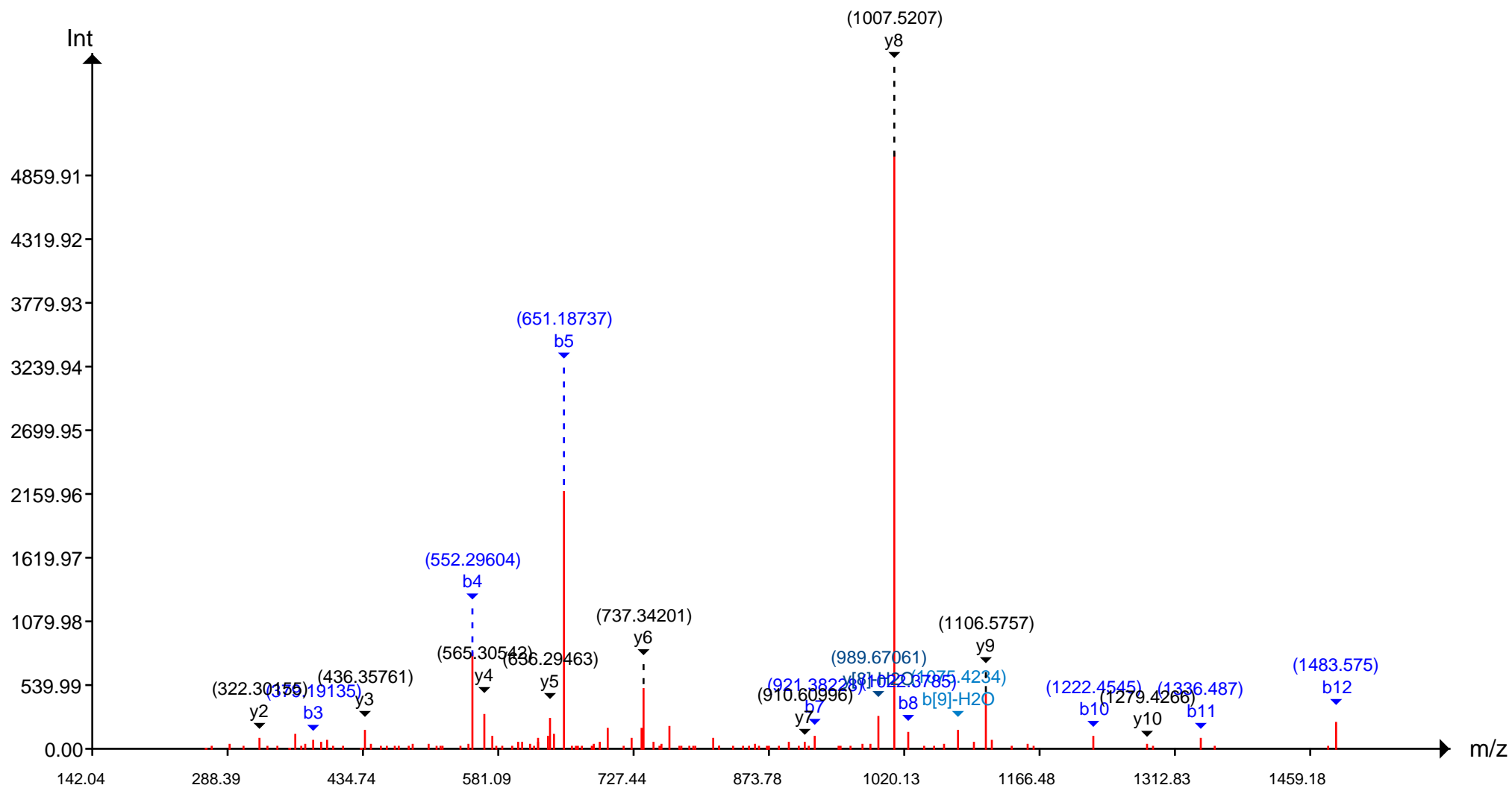
Charge 2+

Mass error 0.001412



Info

Modified Sequence	AcD3-VGTLK<AcD3K*>AAK<AcD3K*>R<C13>-COOH	Ion score	53.33
Protein	Q4KML4	Confidence level	99.00%
Precursor m/z	542.85991	Identity Threshold	38.40
Charge	2+	Mass error	-4.97E-4



Info

Modified Sequence AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH

Ion score 71.41

Protein P17742

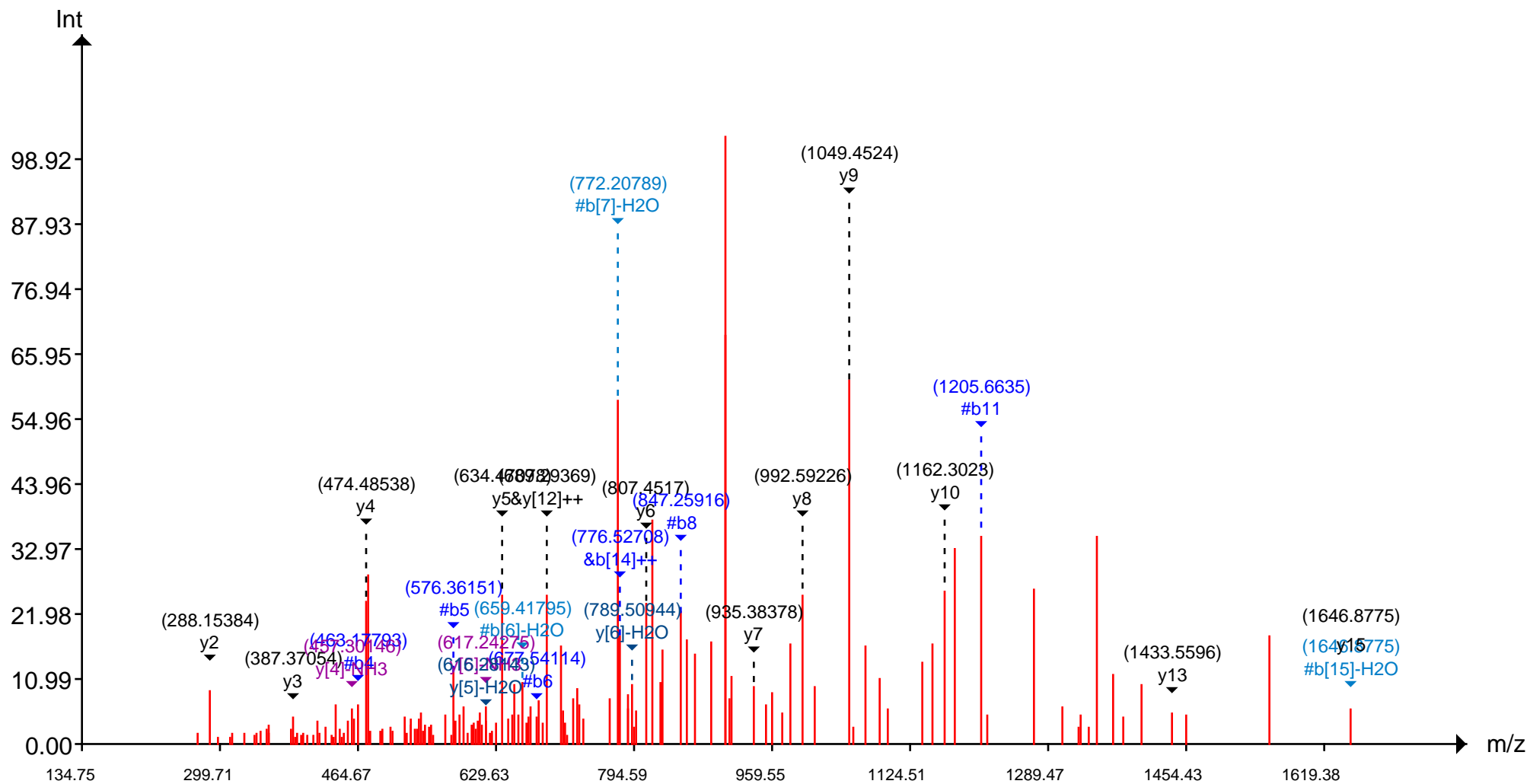
Confidence level 99.00%

Precursor m/z 829.44927

Identity Threshold 47.76

Charge 2+

Mass error 0.001832



Info

Modified Sequence AcD3-FVNGTLGGQK<AcD3K*>C<Cmm*>SVIR-COOH

Ion score 53.45

Protein P62962

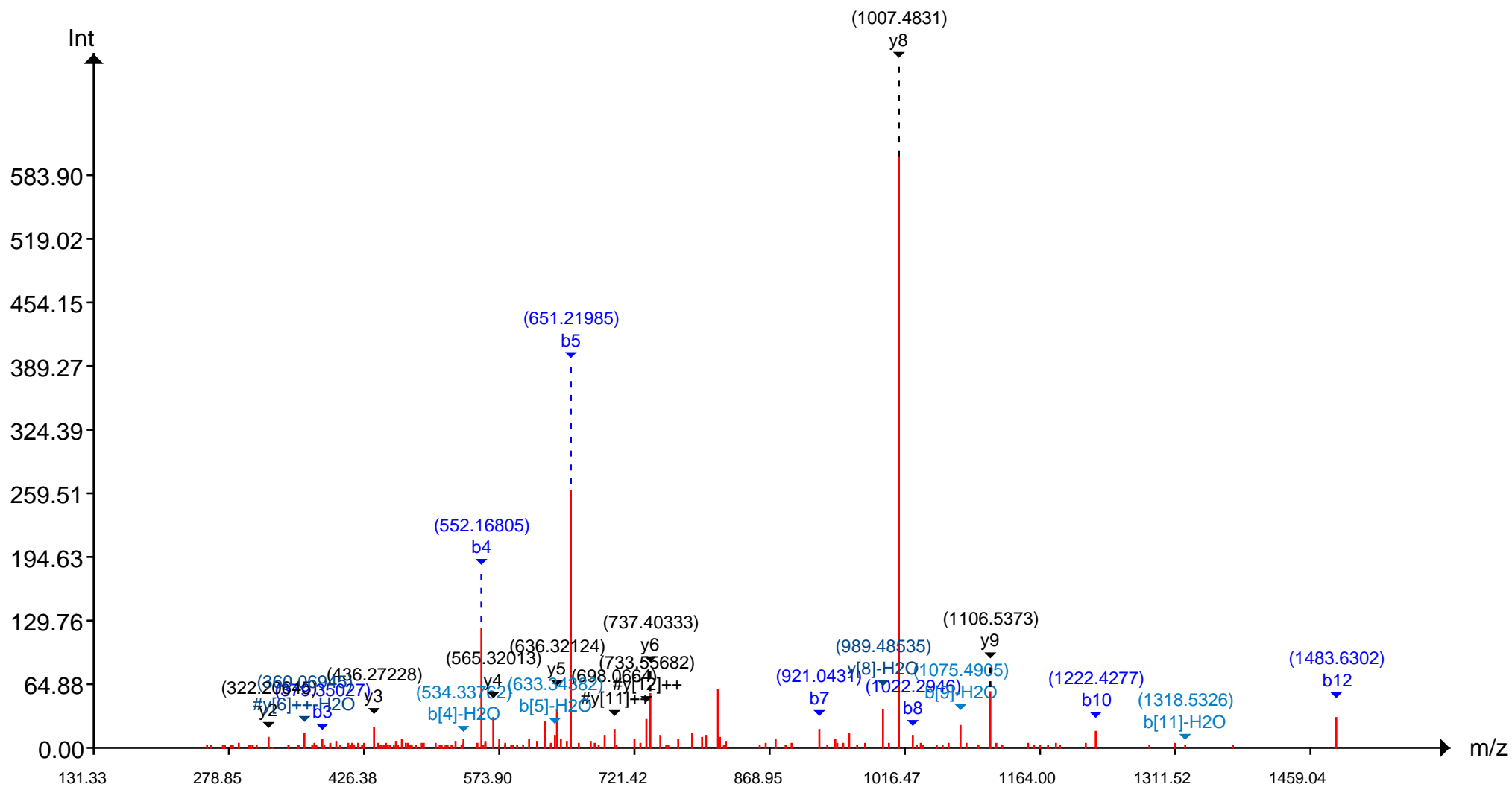
Confidence level 99.00%

Precursor m/z 920.00711

Identity Threshold 47.19

Charge 2+

Mass error 5.38E-4



Info

Modified Sequence AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH

Ion score 49.74

Protein P17742

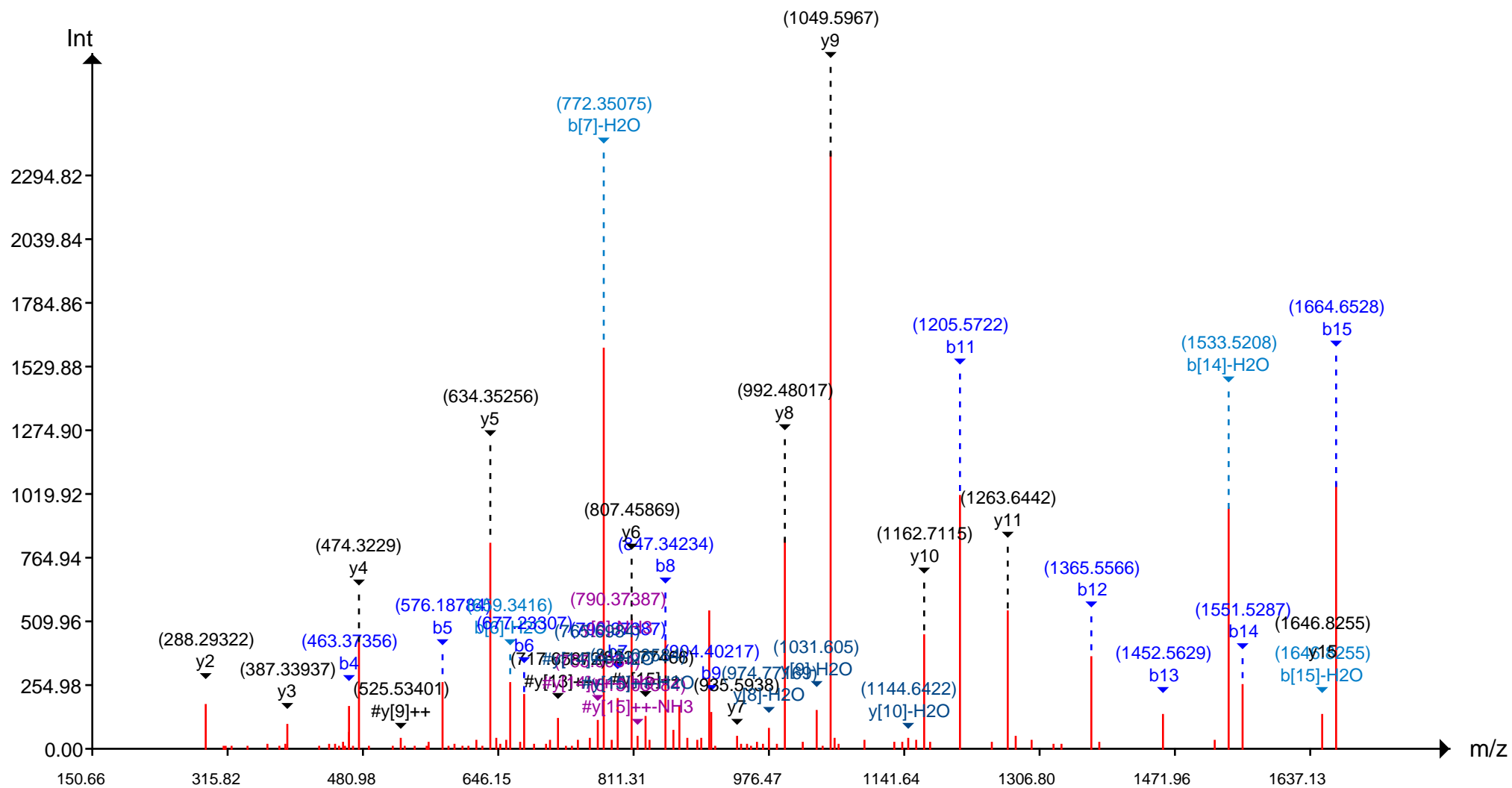
Confidence level 99.00%

Precursor m/z 829.44925

Identity Threshold 47.76

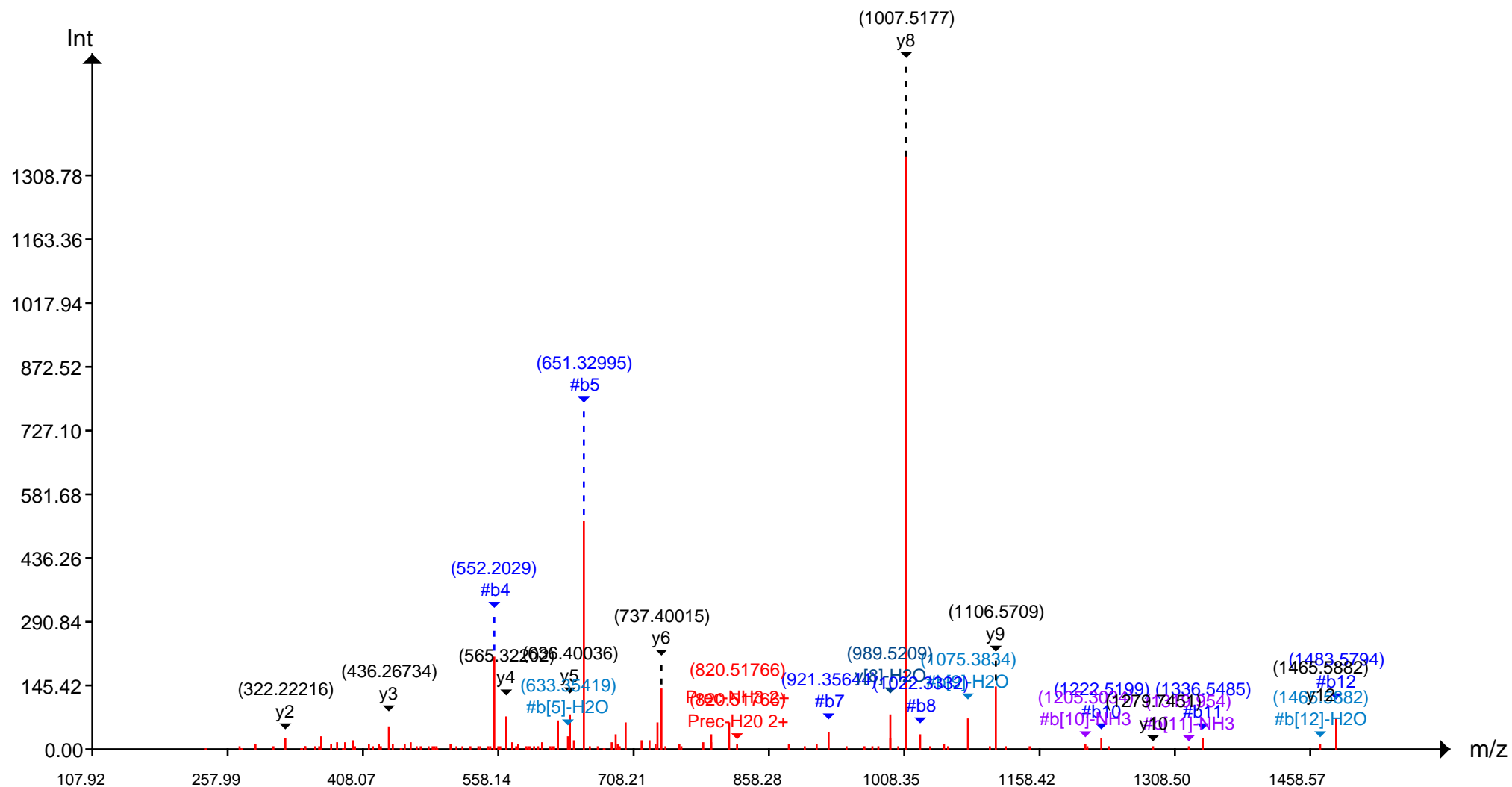
Charge 2+

Mass error 0.001792



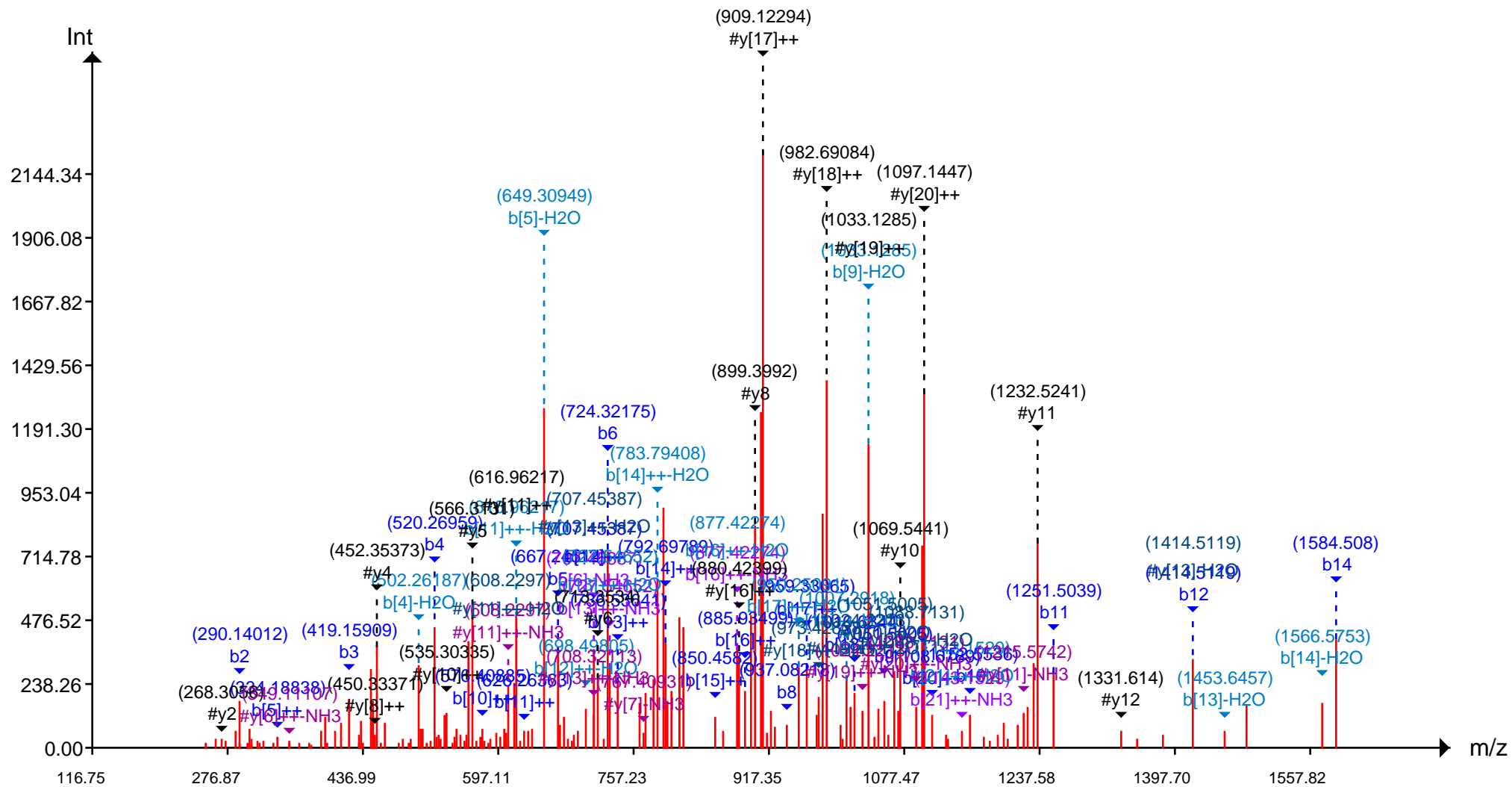
Info

Modified Sequence	AcD3-FVNGTLGGQK<AcD3K*>C<Cmm*>SVIR-COOH	Ion score	119.01
Protein	P62962	Confidence level	99.00%
Precursor m/z	920.00767	Identity Threshold	47.18
Charge	2+	Mass error	0.001658



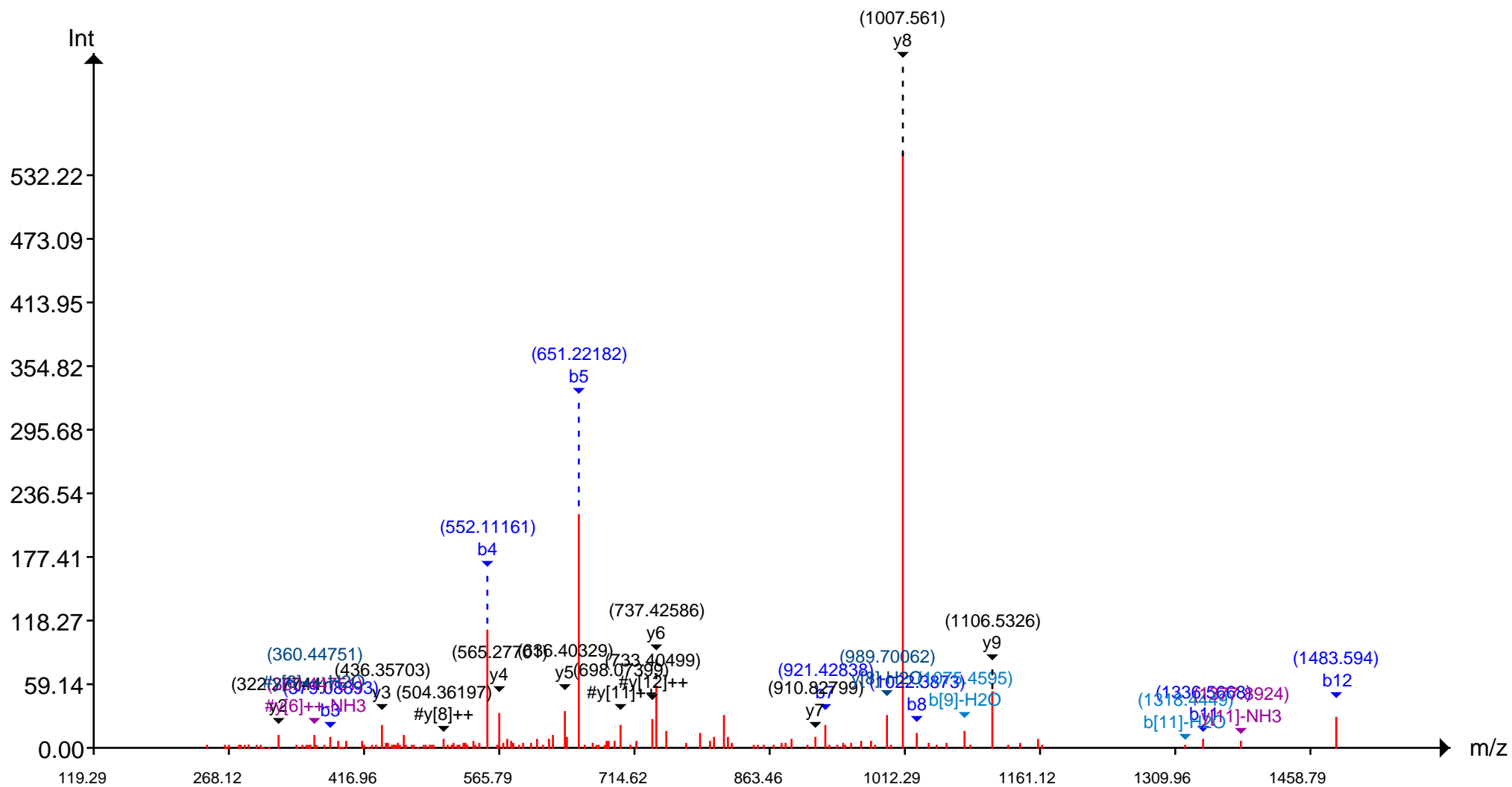
Info

Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH	Ion score	80.58
Protein	P17742	Confidence level	99.00%
Precursor m/z	829.44922	Identity Threshold	47.76
Charge	2+	Mass error	0.001732



Info

Modified Sequence	AcD3-AK<AcD3K*>ETFGVNN TVY GIDAM<Mox*>NPSSR<C13>-COOH	Ion score	62.26
Protein	Q66JS6	Confidence level	99.00%
Precursor m/z	828.40842	Identity Threshold	47.12
Charge	3+	Mass error	0.002184



Info

Modified Sequence AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH

Ion score 62.37

Protein P17742

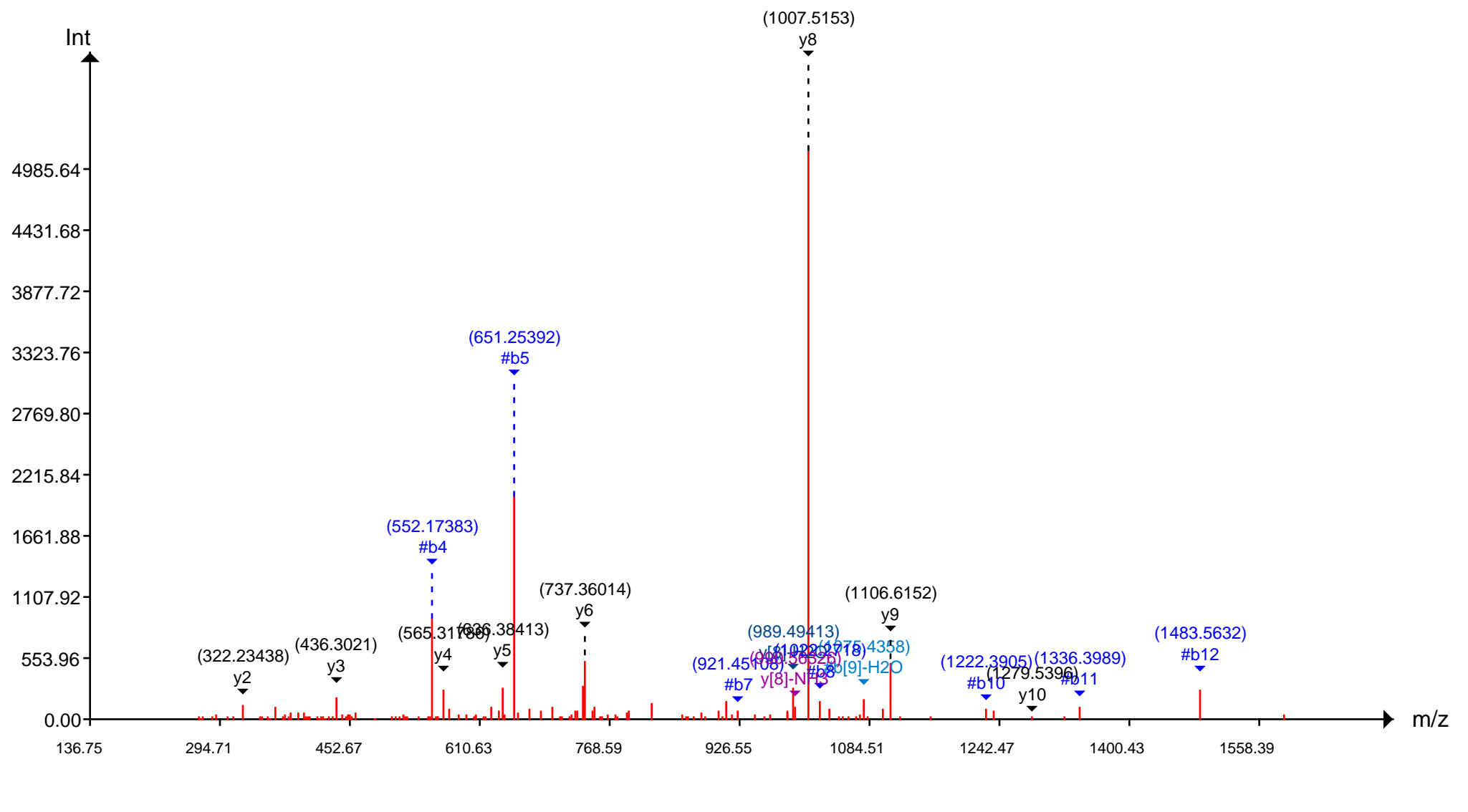
Confidence level 99.00%

Precursor m/z 829.44941

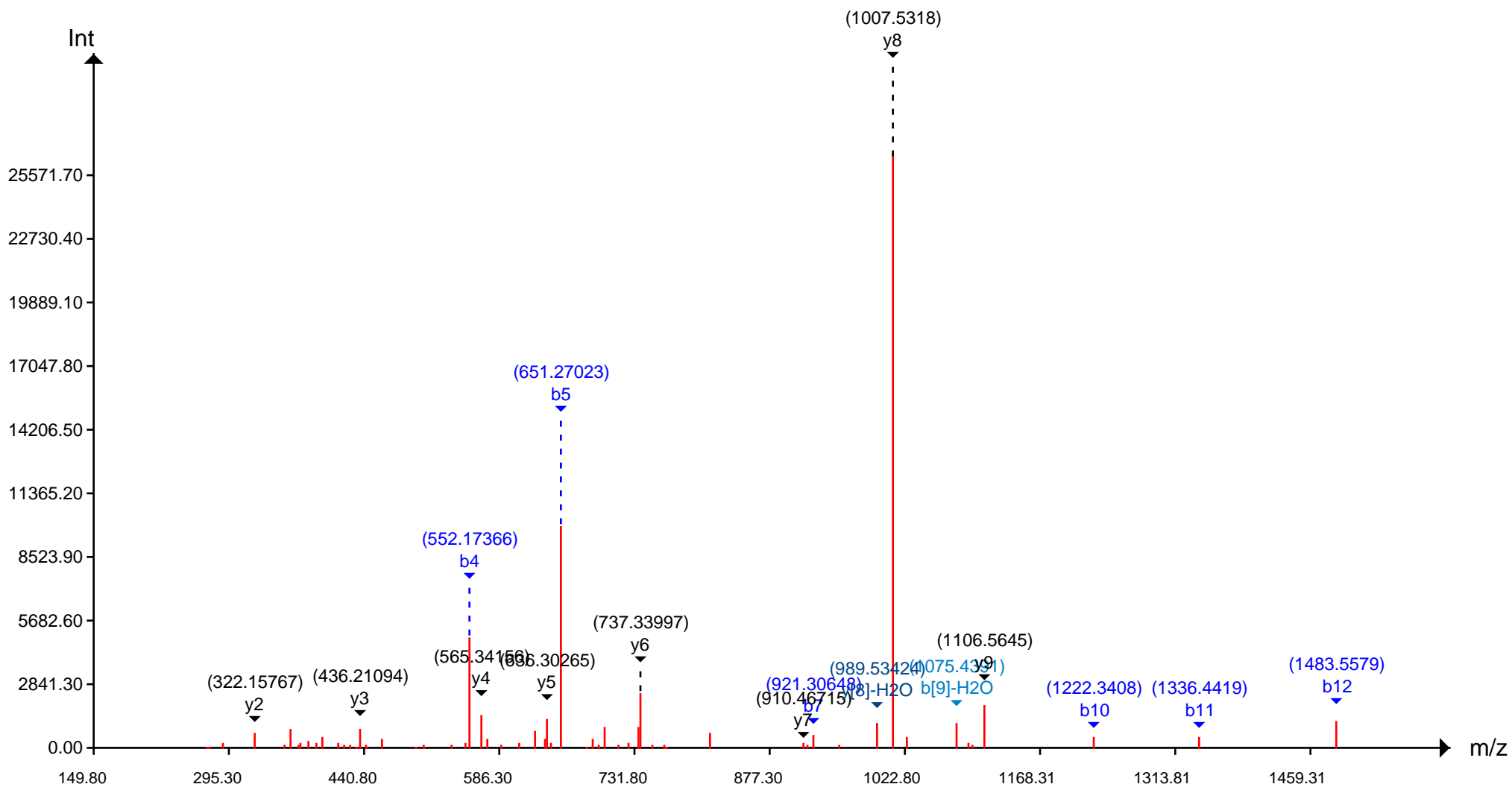
Identity Threshold 47.78

Charge 2+

Mass error 0.002112

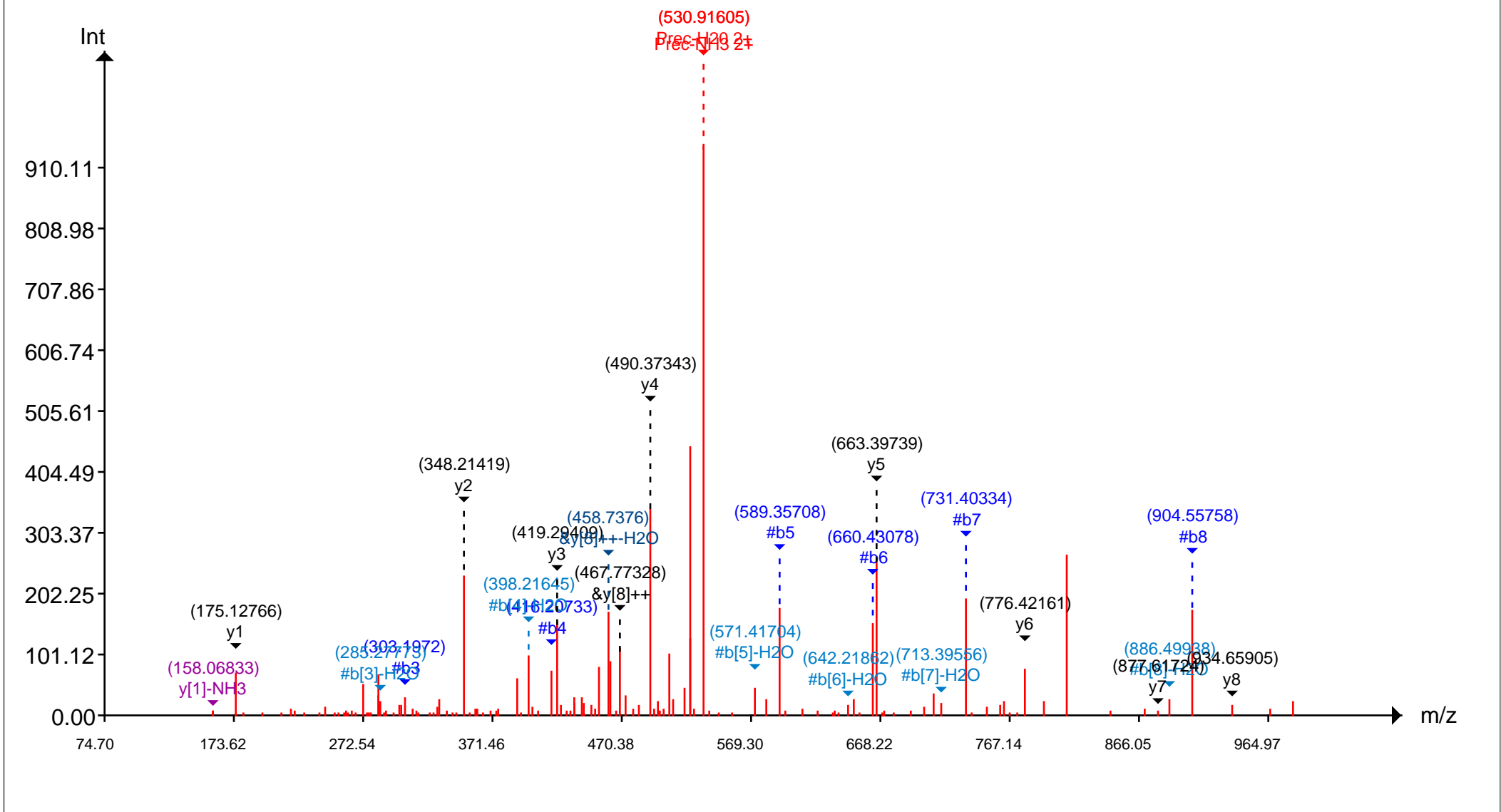


Info			
Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH	Ion score	62.77
Protein	P17742	Confidence level	99.00%
Precursor m/z	829.44961	Identity Threshold	47.77
Charge	2+	Mass error	0.002512



Info

Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH	Ion score	80.34
Protein	P17742	Confidence level	99.00%
Precursor m/z	829.44946	Identity Threshold	47.78
Charge	2+	Mass error	0.002212



Info

Modified Sequence AcD3-VGTLK<AcD3K*>AAK<AcD3K*>R-COOH

Ion score 52.82

Protein Q4KML4

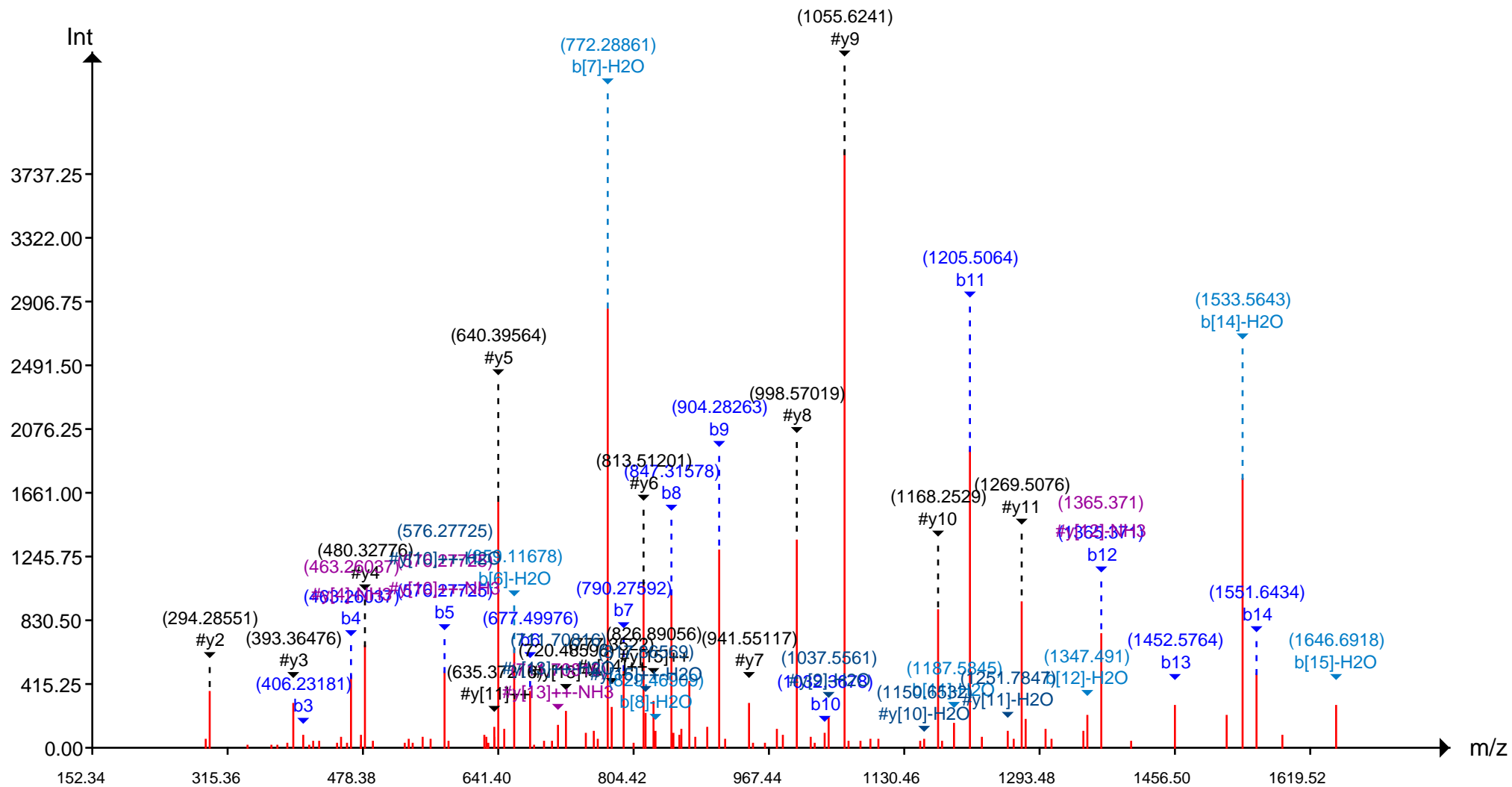
Confidence level 99.00%

Precursor m/z 539.8502

Identity Threshold 39.67

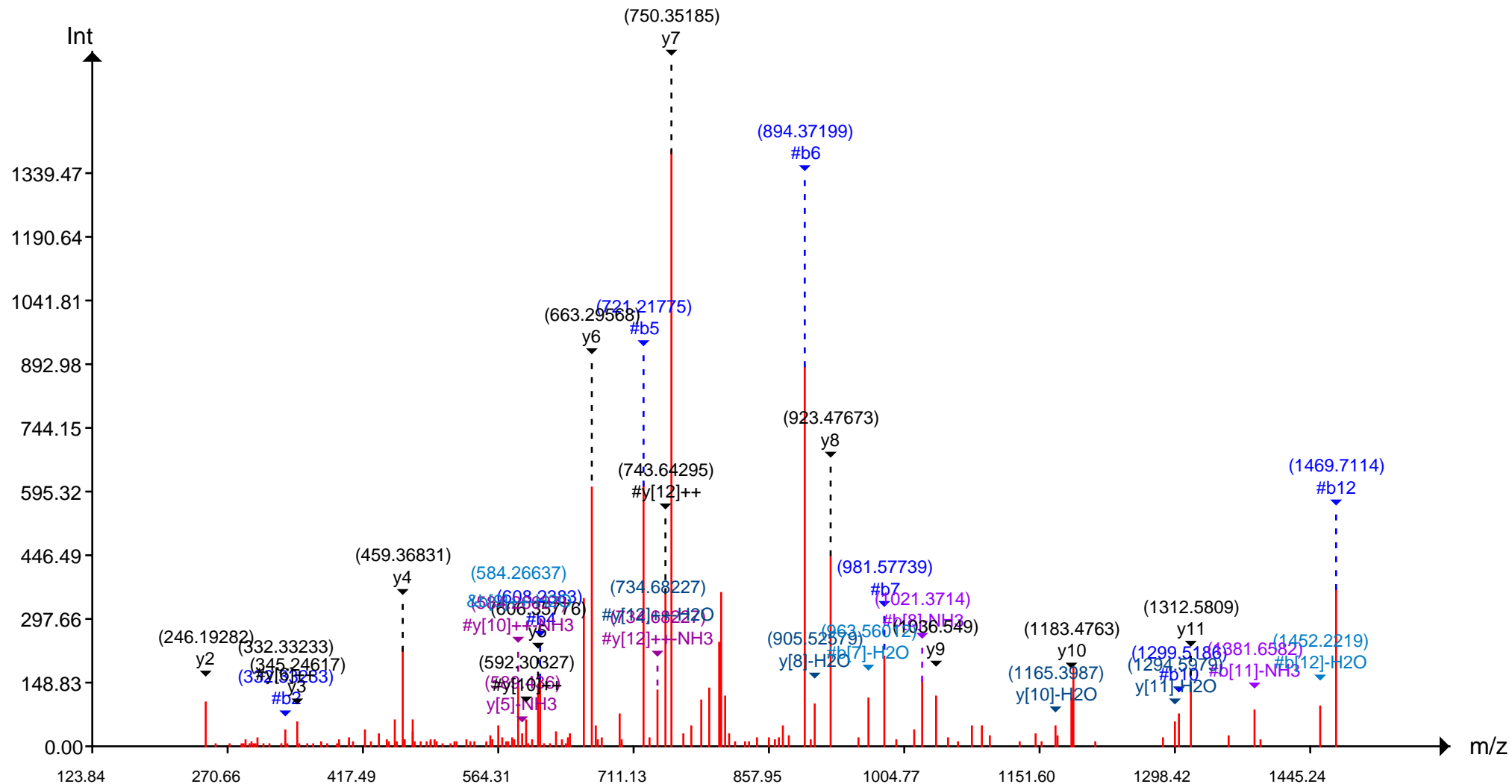
Charge 2+

Mass error 2.25E-4



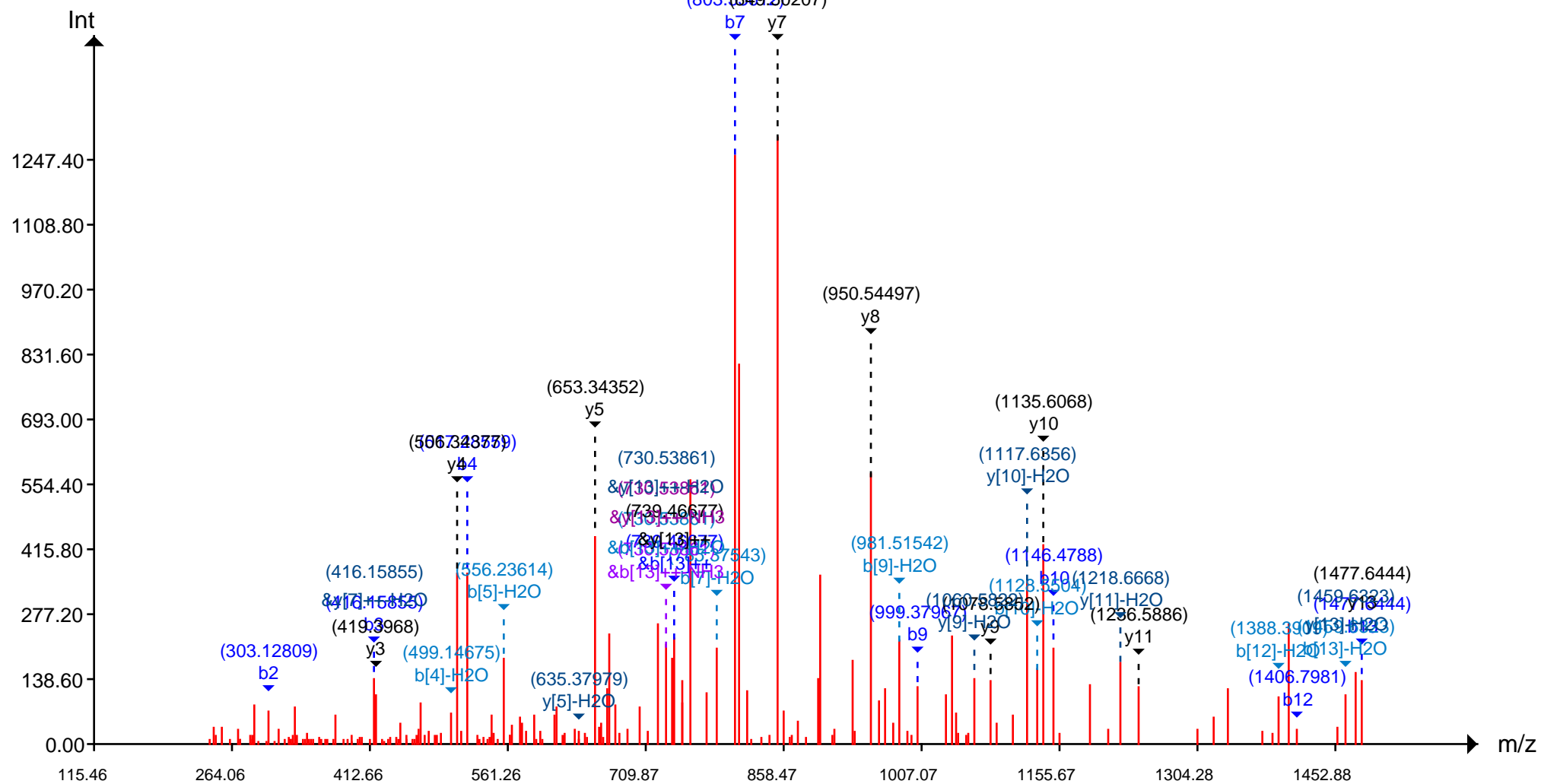
Info

Modified Sequence	AcD3-FVNGTLGGQK<AcD3K*>C<Cmm*>SVIR<C13>-COOH	Ion score	92.92
Protein	P62962	Confidence level	99.00%
Precursor m/z	923.01772	Identity Threshold	46.68
Charge	2+	Mass error	0.001616



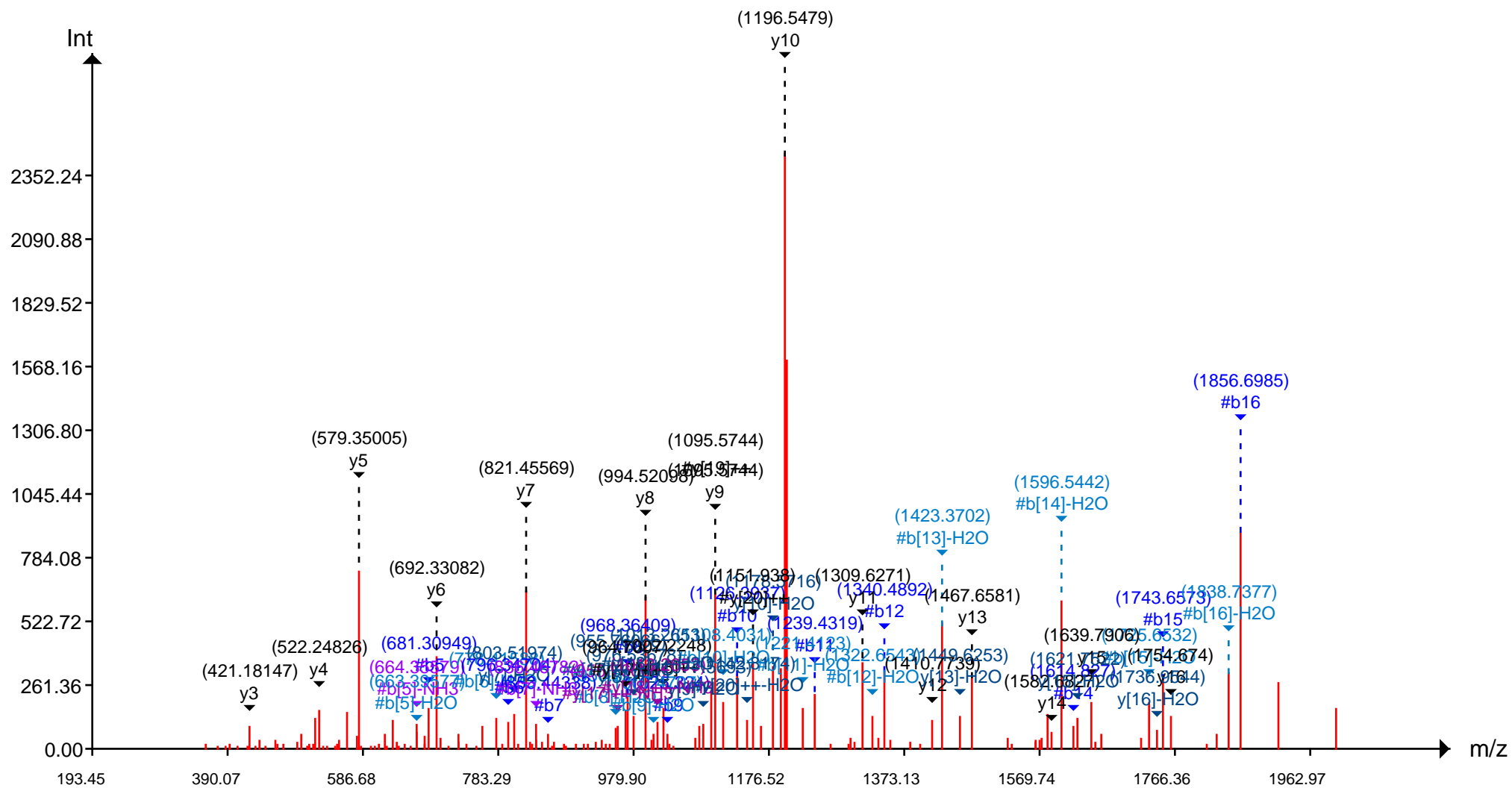
Info

Modified Sequence	AcD3-LK<AcD3K*>EM<Mox*>IK<AcD3K*>SGM<Mox*>NVAR-COOH	Ion score	75.48
Protein	P52480	Confidence level	99.00%
Precursor m/z	822.44483	Identity Threshold	47.89
Charge	2+	Mass error	0.001878



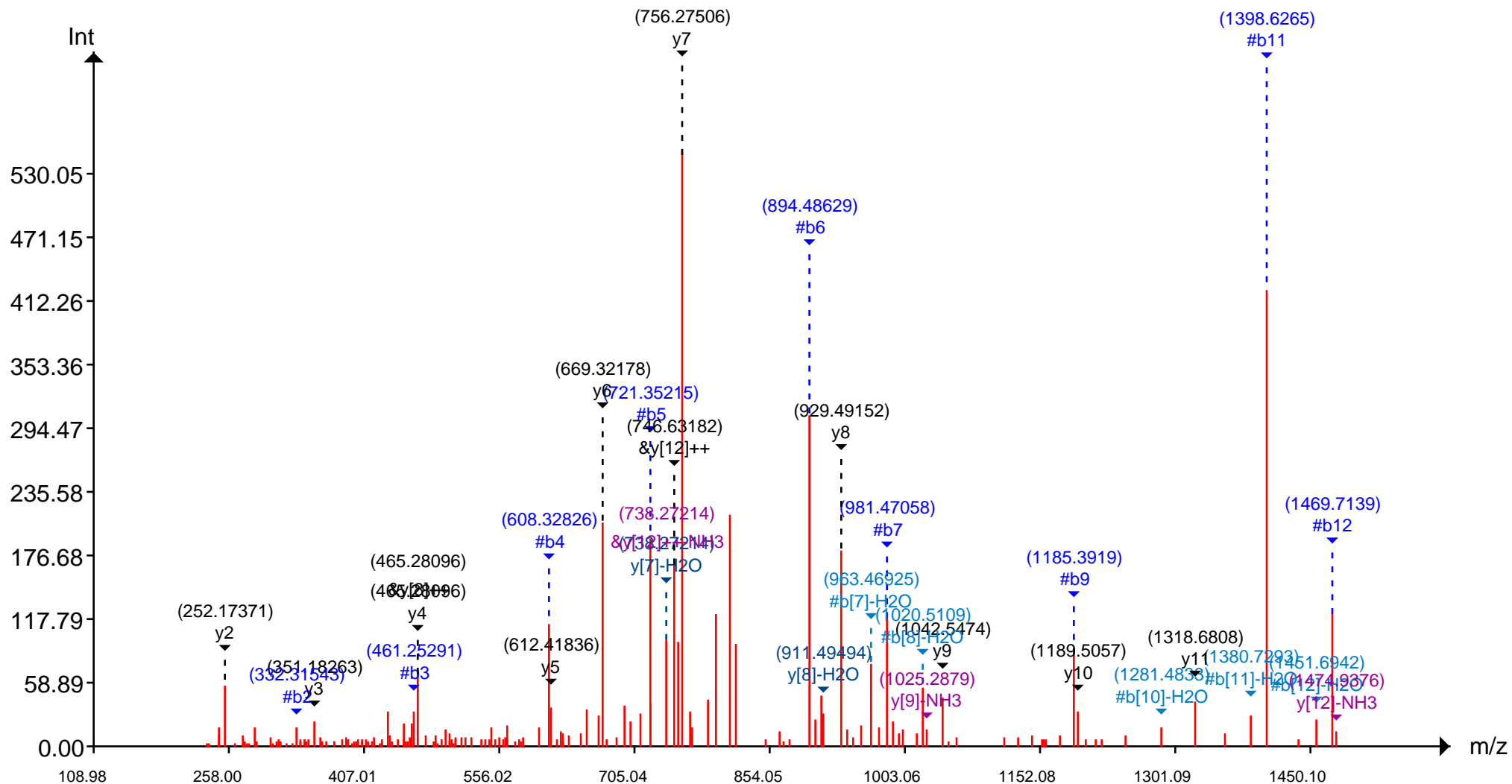
Info

Modified Sequence	AcD3-EQLTGQTPVFSK<AcD3K*>AR-COOH	Ion score	48.33
Protein	Q9CXW4	Confidence level	99.00%
Precursor m/z	826.4507	Identity Threshold	47.65
Charge	2+	Mass error	0.002052



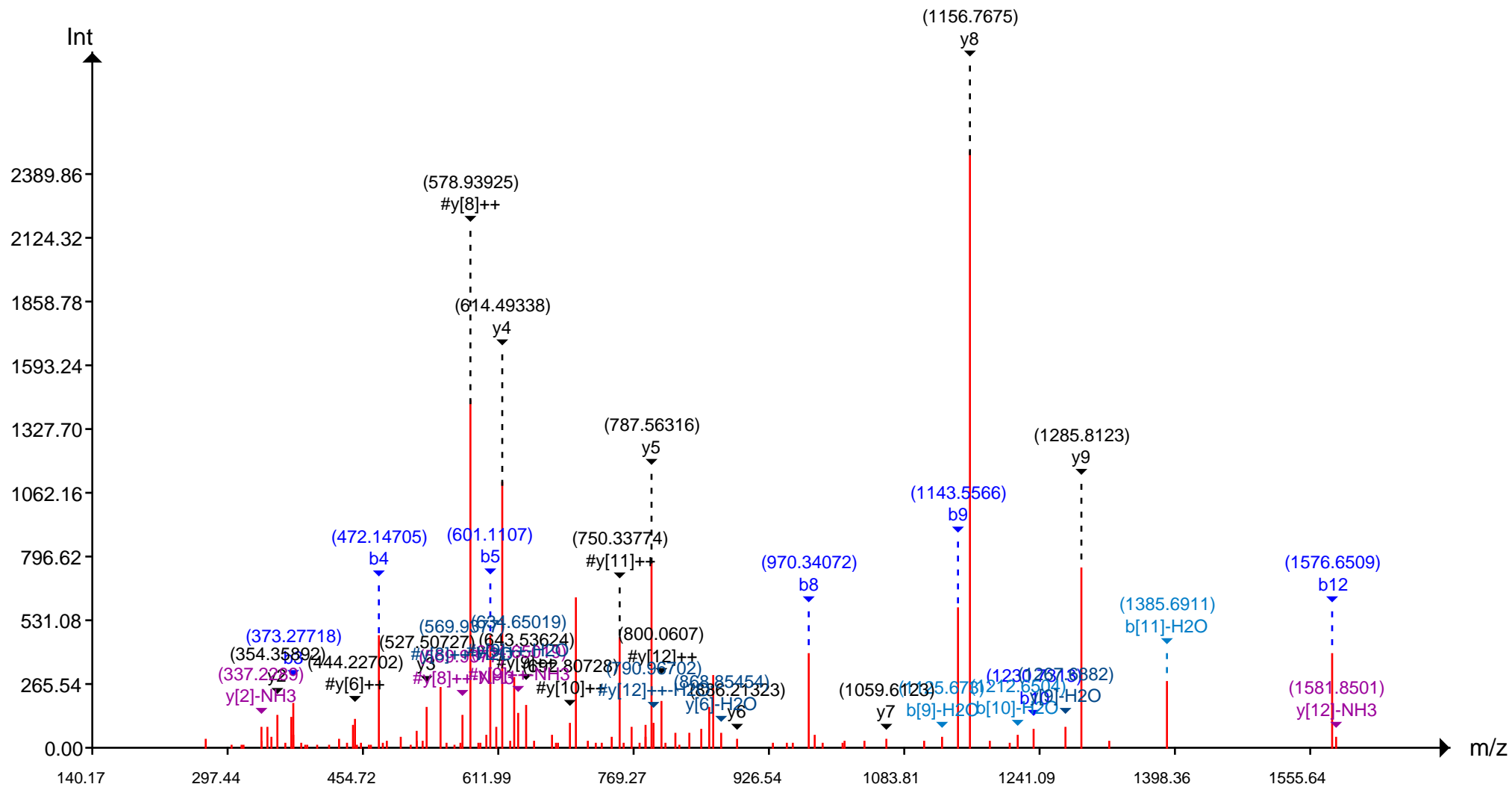
Info

Modified Sequence	AcD3-SLFDK<AcD3K*>DGDGTITTK<AcD3K*>ELGTVM<Mox*>R-COOH	Ion score	99.2
Protein	P62204	Confidence level	99.00%
Precursor m/z	1218.1198	Identity Threshold	47.89
Charge	2+	Mass error	2.56E-4



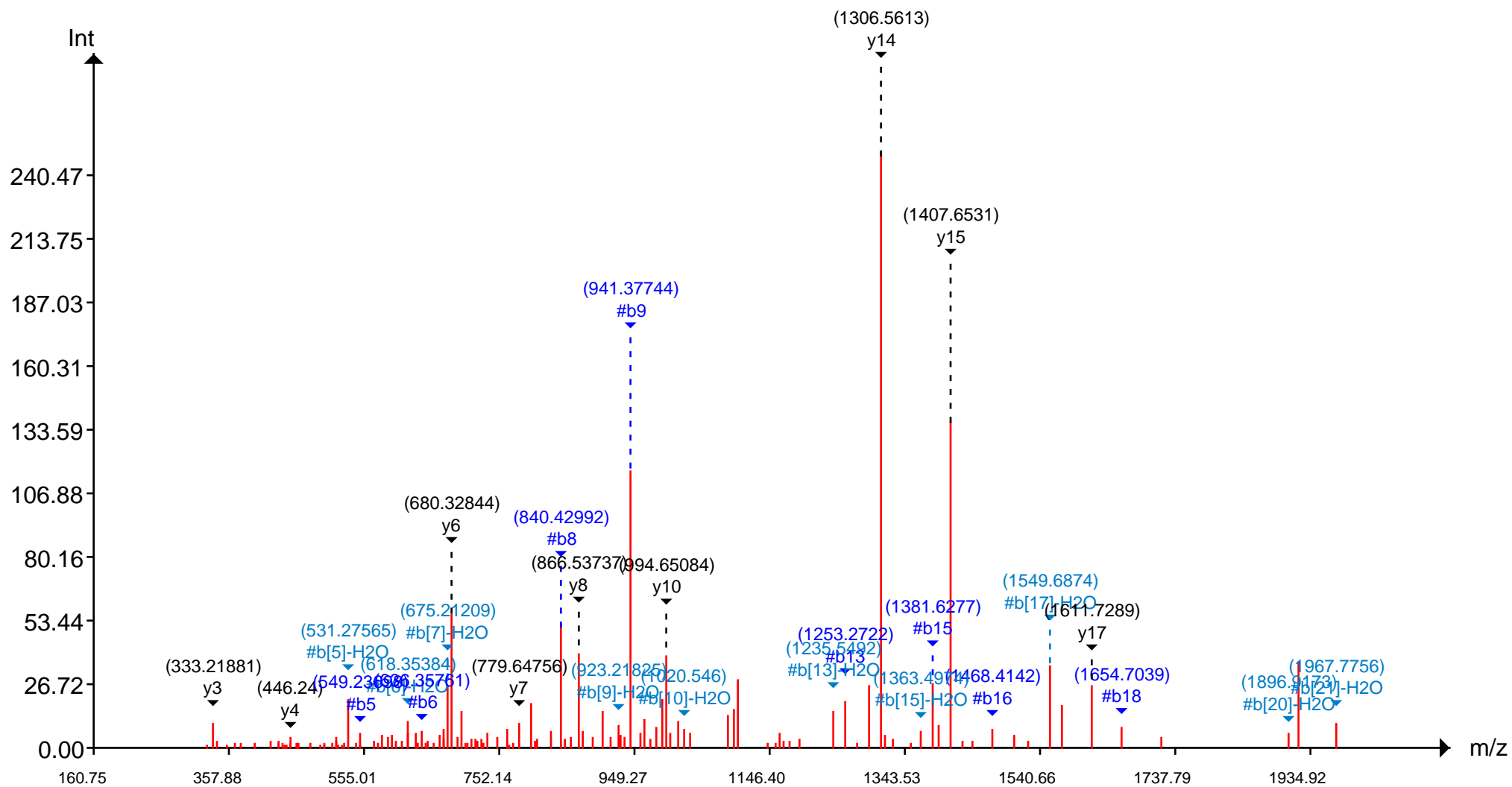
Info

Modified Sequence	AcD3-LK<AcD3K*>EM<Mox*>IK<AcD3K*>SGM<Mox*>NVAR<C13>-COOH	Ion score	96.15
Protein	P52480	Confidence level	99.00%
Precursor m/z	825.45459	Identity Threshold	47.54
Charge	2+	Mass error	0.001256



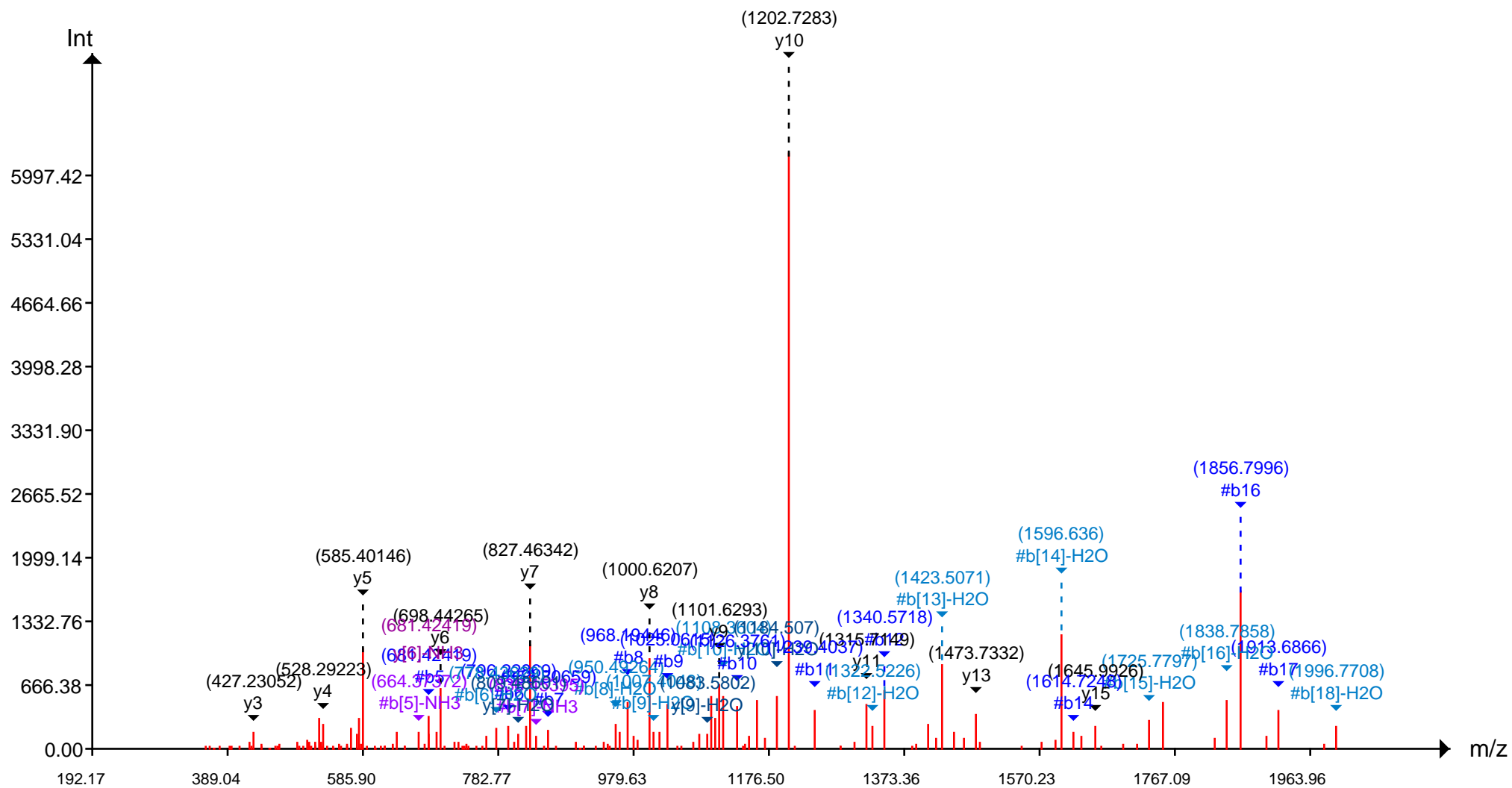
Info

Modified Sequence	AcD3-LVDVEPK<AcD3K*>VK<AcD3K*>SK<AcD3K*>K<AcD3K*>R<C13>-COOH	Ion score	63.01
Protein	P62315	Confidence level	99.00%
Precursor m/z	879.05933	Identity Threshold	35.93
Charge	2+	Mass error	0.001828



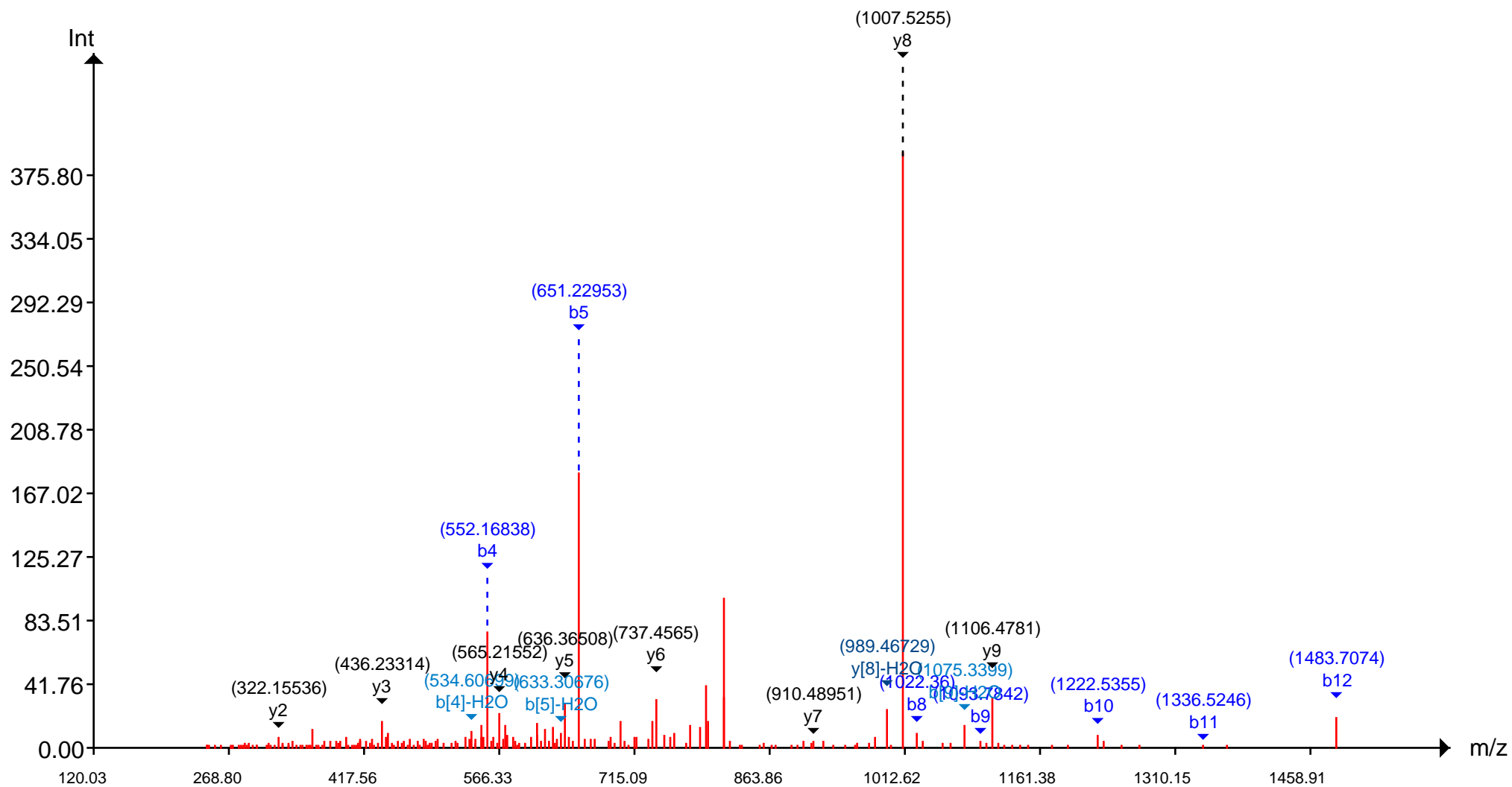
Info

Modified Sequence	AcD3-VESSTSGFTPSGAGASVSM<Mox*>IASR-COOH	Ion score	58.24
Protein	Q99MD9	Confidence level	99.00%
Precursor m/z	1124.0369	Identity Threshold	46.29
Charge	2+	Mass error	0.002821



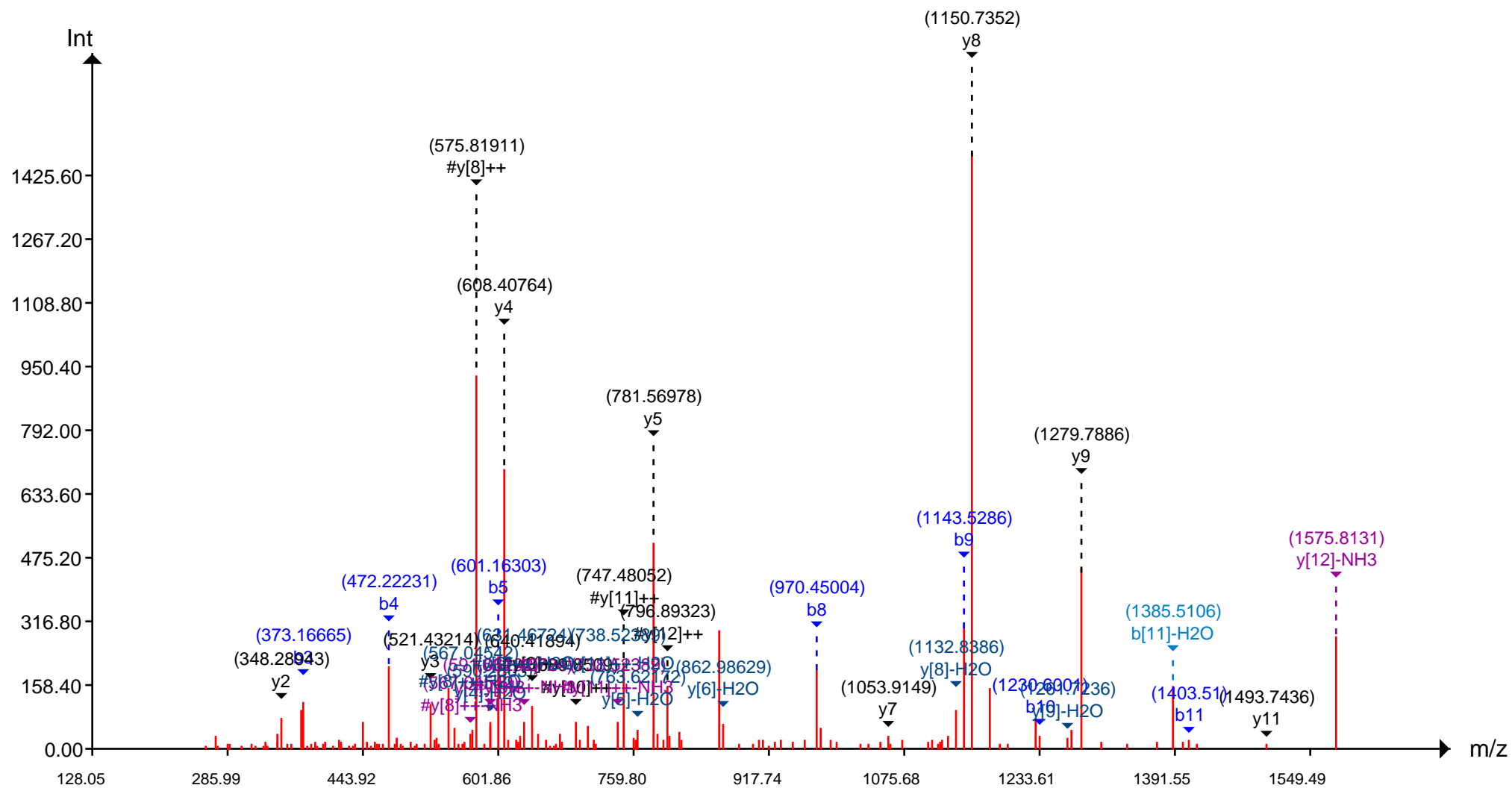
Info

Modified Sequence	AcD3-SLFDK<AcD3K*>DGDGTITTK<AcD3K*>ELGTVM<Mox*>R<C13>-COOH	Ion score	80.54
Protein	P62204	Confidence level	99.00%
Precursor m/z	1221.1296	Identity Threshold	47.90
Charge	2+	Mass error	-2.86E-4



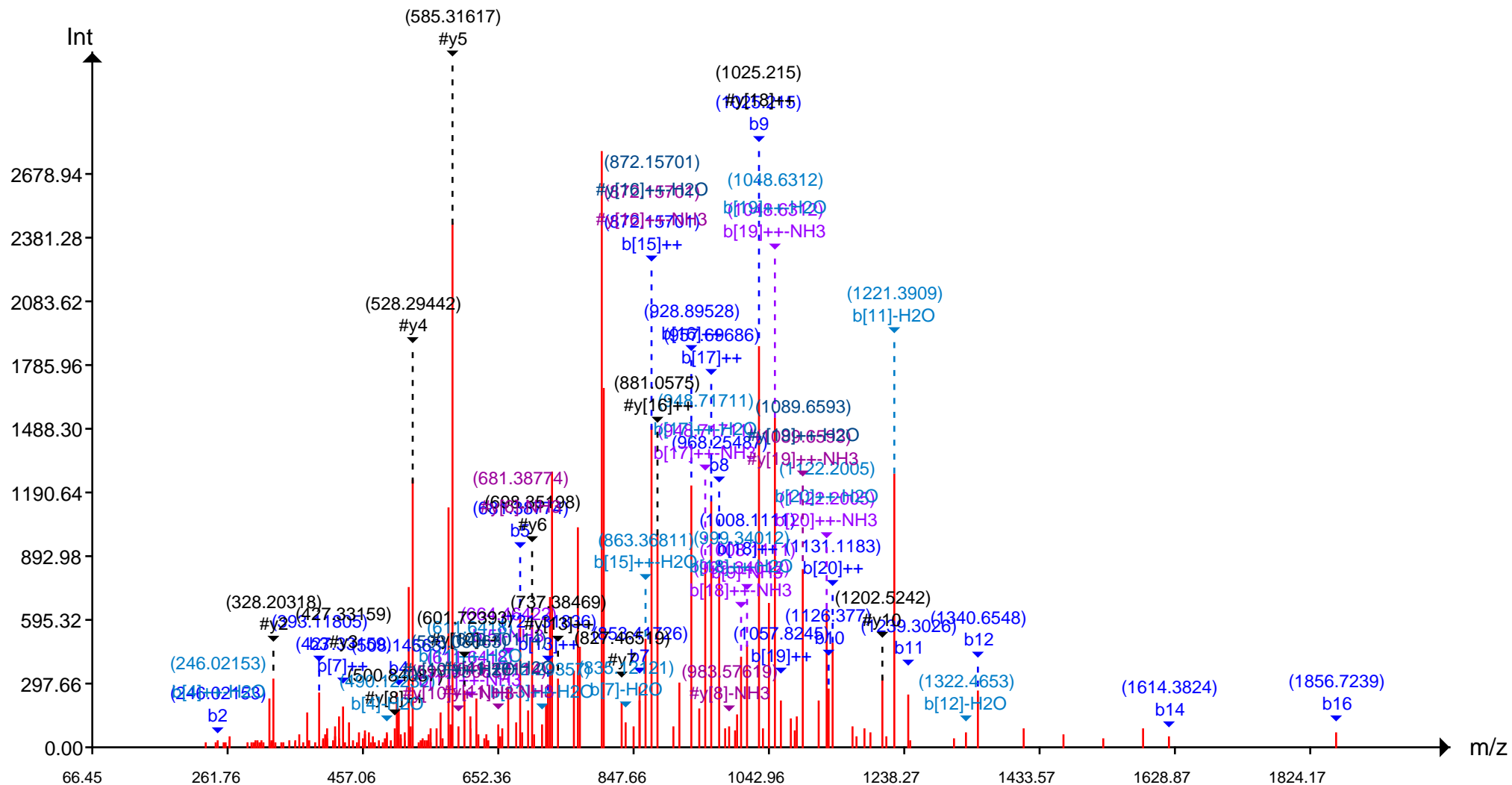
Info

Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH	Ion score	48.64
Protein	P17742	Confidence level	99.00%
Precursor m/z	829.44887	Identity Threshold	47.81
Charge	2+	Mass error	0.001032



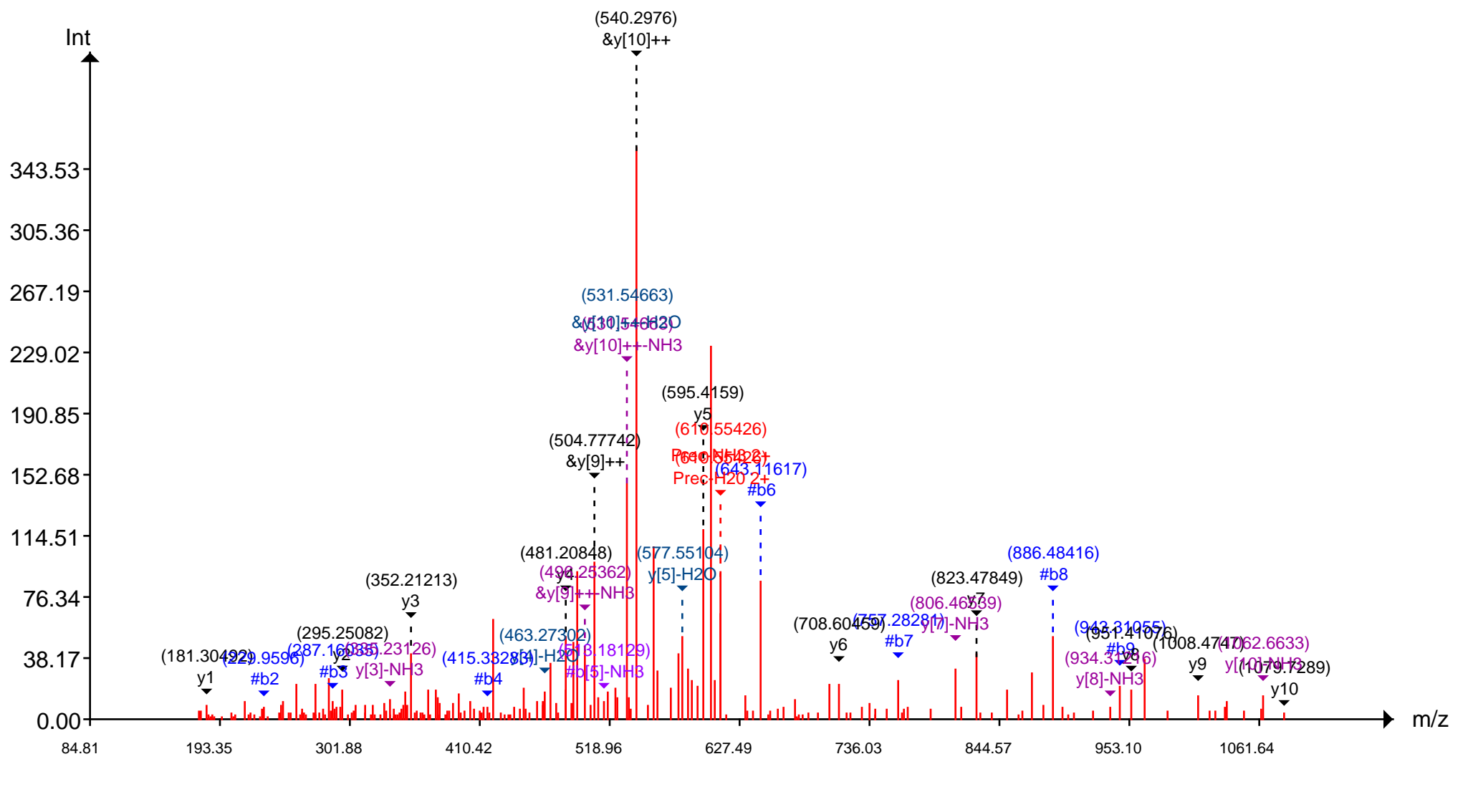
Info

Modified Sequence	AcD3-LVDVEPK<AcD3K*>VK<AcD3K*>SK<AcD3K*>K<AcD3K*>R-COOH	Ion score	51.9
Protein	P62315	Confidence level	99.00%
Precursor m/z	876.04945	Identity Threshold	37.68
Charge	2+	Mass error	0.00221

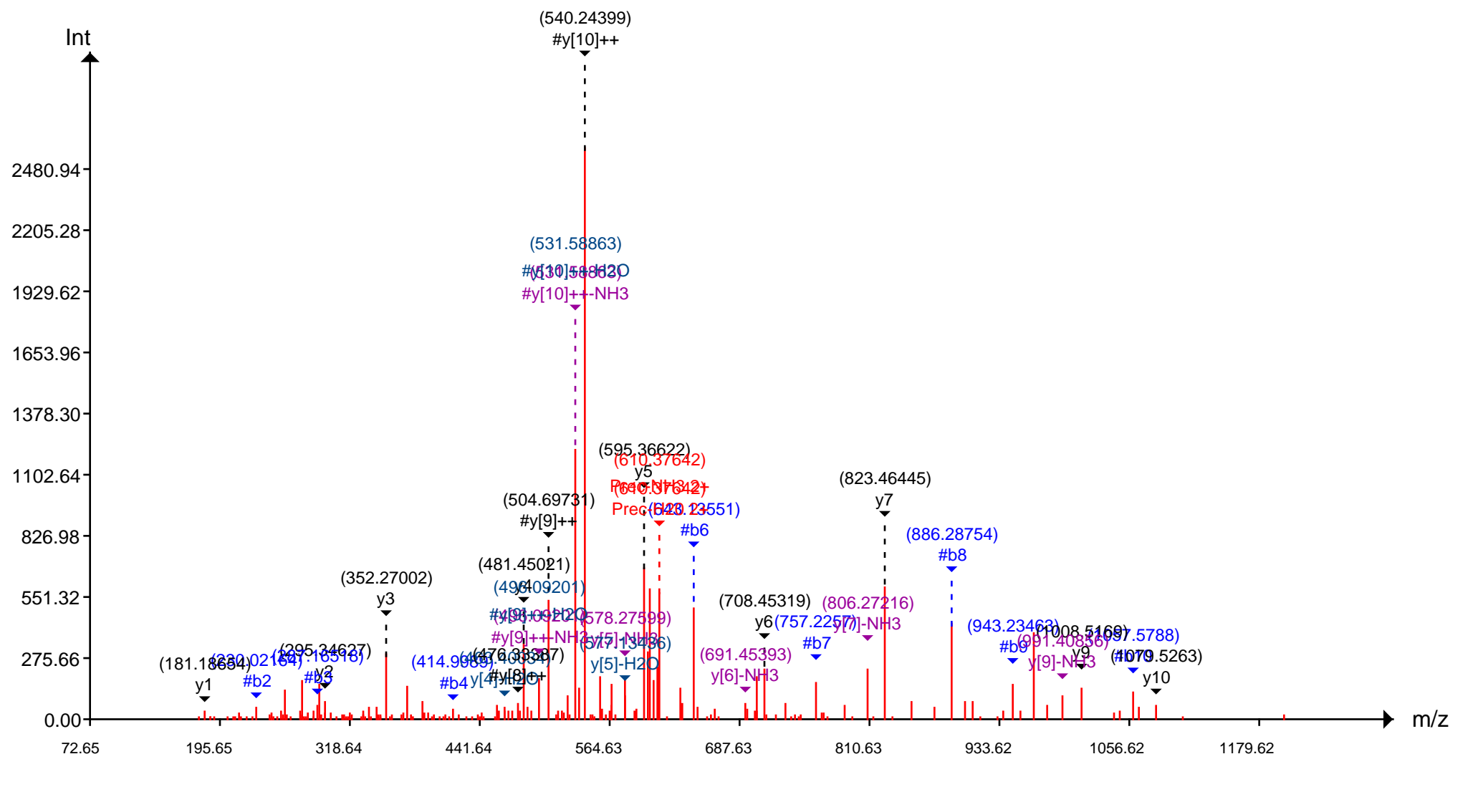


Info

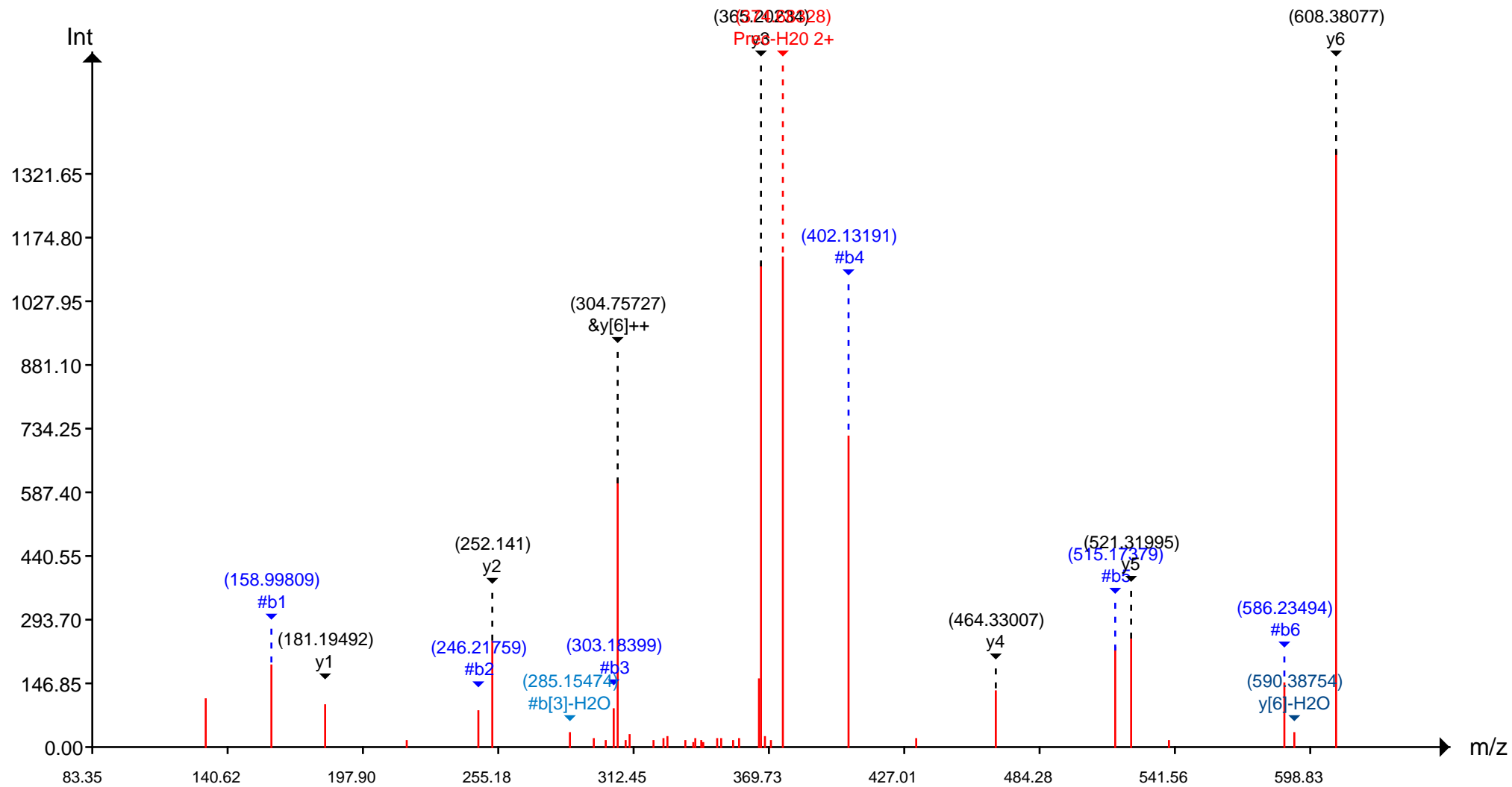
Modified Sequence	AcD3-SLFDK<AcD3K*>DGDGTITTK<AcD3K*>ELGTVM<Mox*>R<C13>-COOH	Ion score	66.34
Protein	P62204	Confidence level	99.00%
Precursor m/z	814.42275	Identity Threshold	47.91
Charge	3+	Mass error	0.001488



Info			
Modified Sequence	AcD3-IAGQDLNEGNR<C13>-COOH	Ion score	57.44
Protein	Q61233	Confidence level	99.00%
Precursor m/z	619.31919	Identity Threshold	47.85
Charge	2+	Mass error	5.06E-4

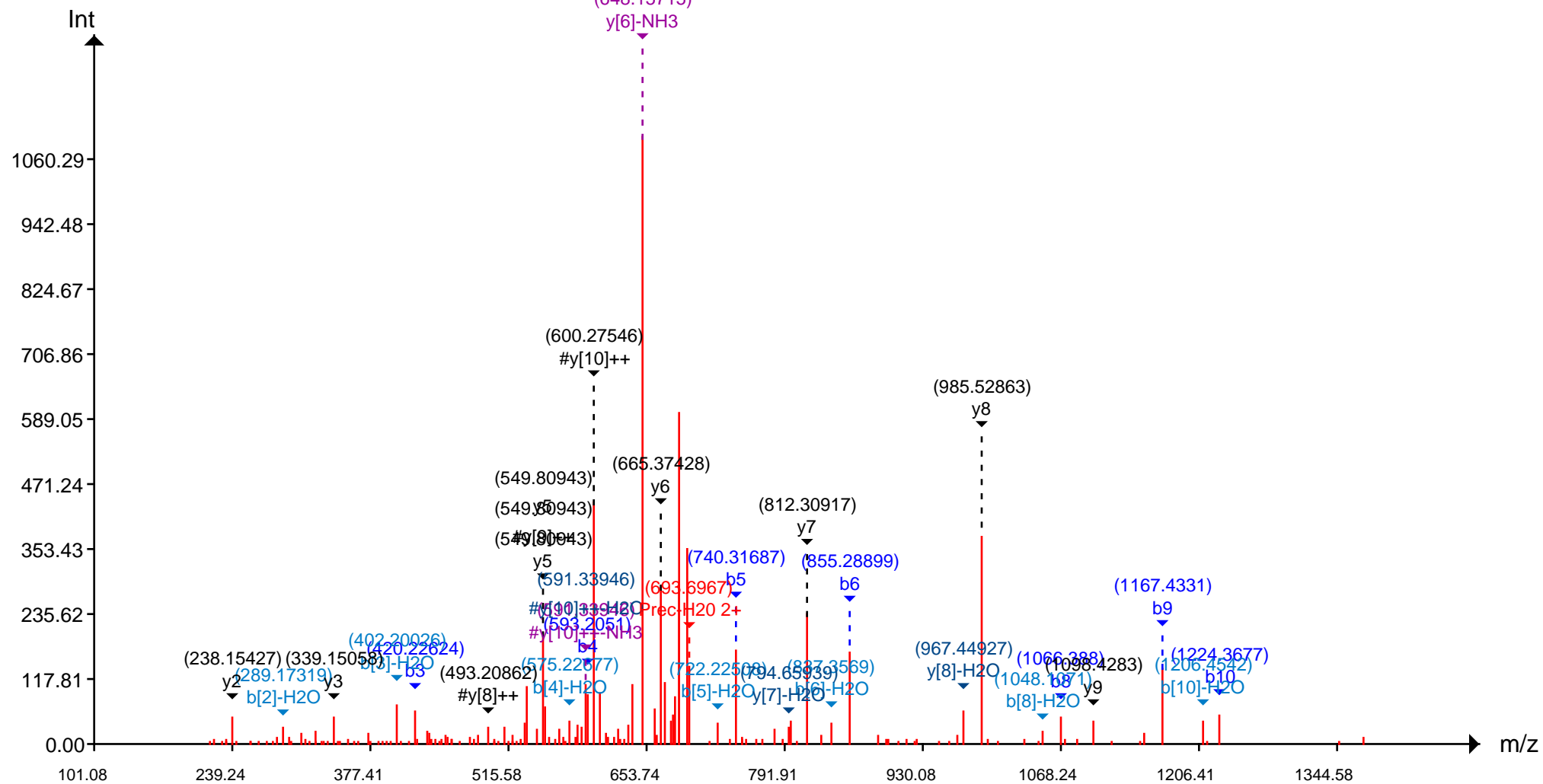


Info			
Modified Sequence	AcD3-IAGQDLNEGNR<C13>-COOH	Ion score	66.97
Protein	Q61233	Confidence level	99.00%
Precursor m/z	619.31981	Identity Threshold	47.97
Charge	2+	Mass error	0.001746



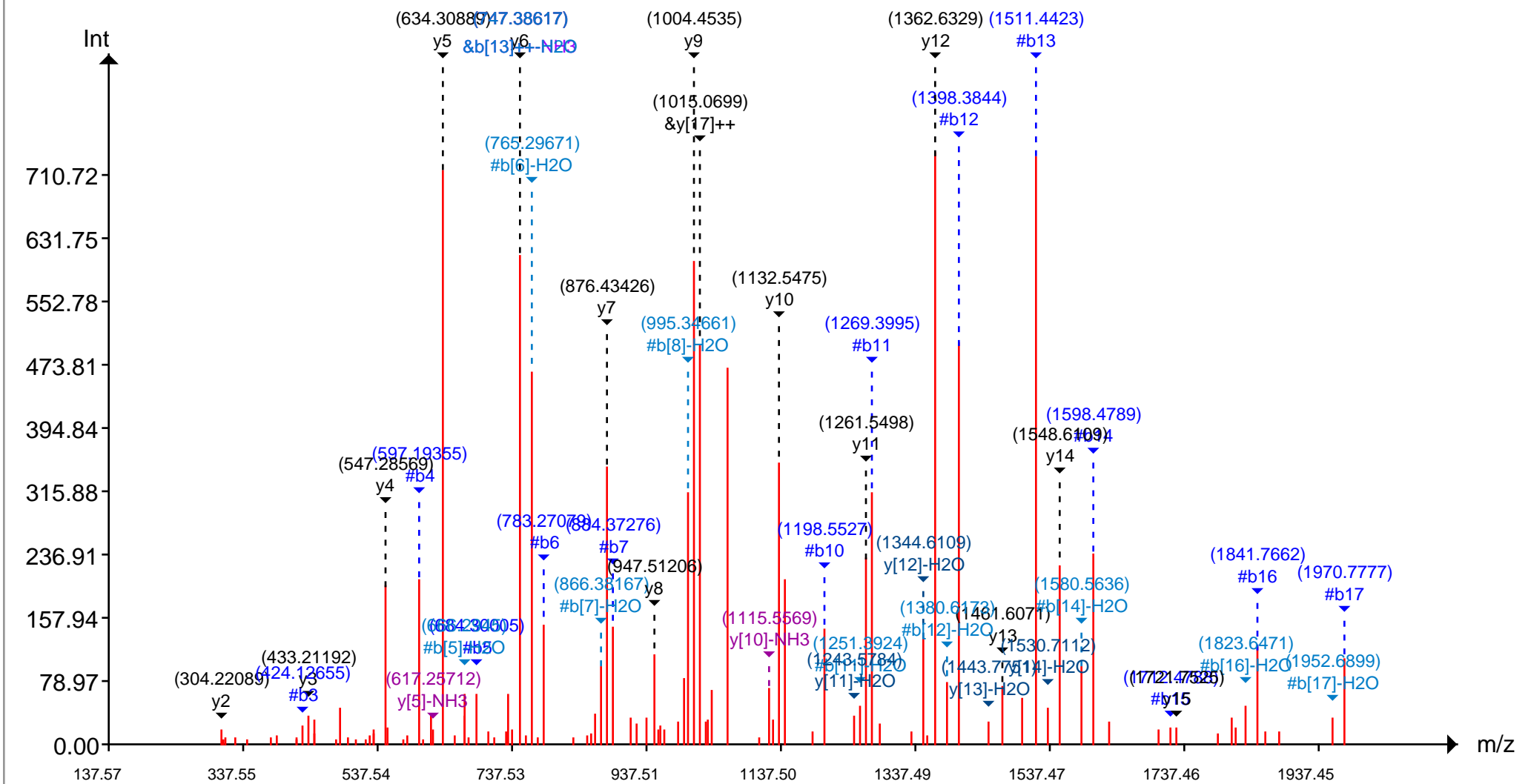
Info

Modified Sequence	AcD3-ISGVIAR<C13>-COOH	Ion score	60.96
Protein	Q91XC8	Confidence level	99.00%
Precursor m/z	383.75148	Identity Threshold	43.01
Charge	2+	Mass error	6.6E-5



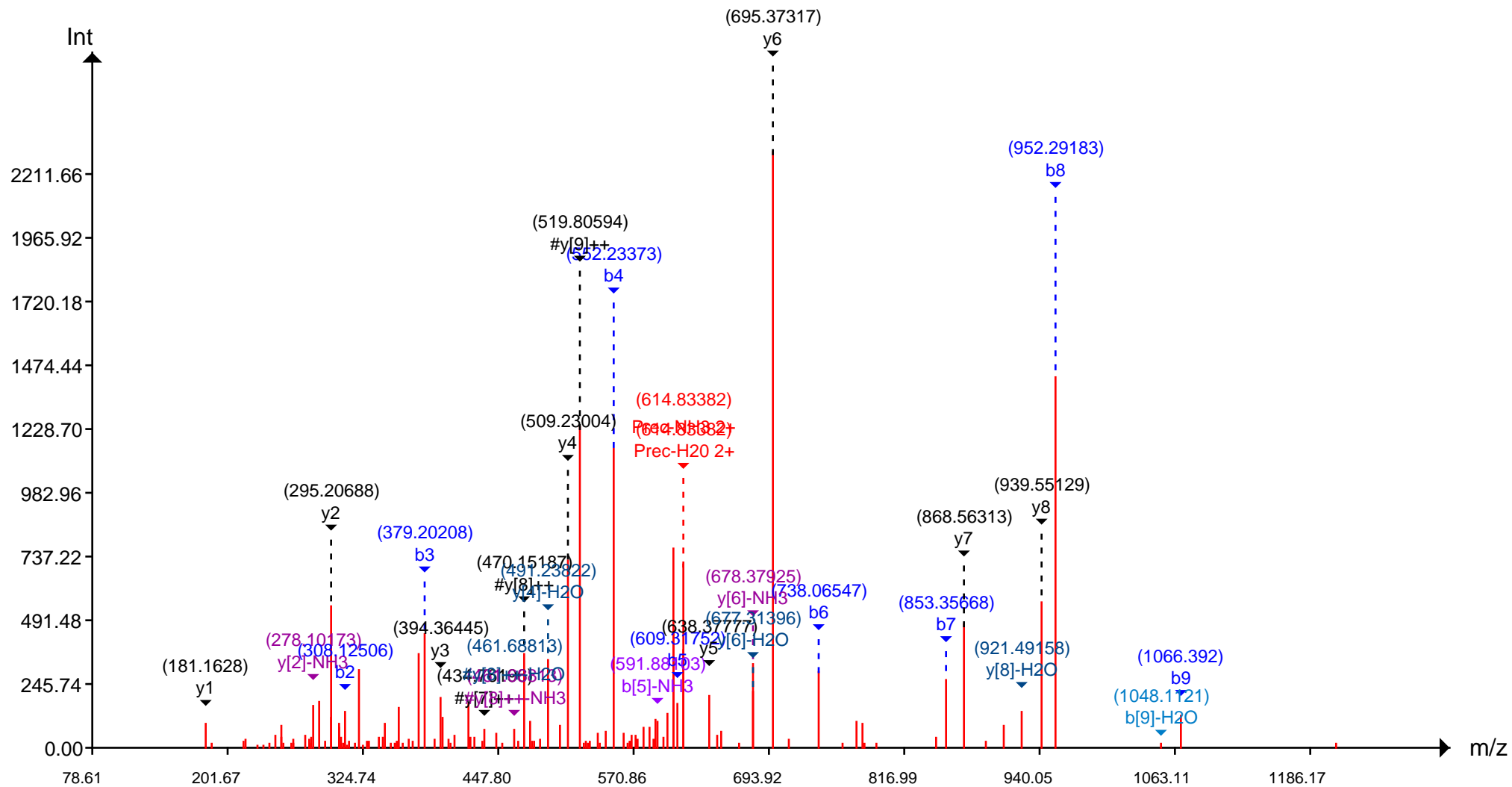
Info

Modified Sequence	AcD3-C<Cmm*>TIK<AcD3K*>M<Mox*>DPNTGR<C13>-COOH	Ion score	60.11
Protein	Q99020	Confidence level	99.00%
Precursor m/z	702.84469	Identity Threshold	47.11
Charge	2+	Mass error	-3.12E-4



Info

Modified Sequence	AcD3-AC<Cmm*>M<Mox*>K<AcD3K*>SVTEQGAELSNEER-COOH	Ion score	113.99
Protein	P63101	Confidence level	99.00%
Precursor m/z	1072.9902	Identity Threshold	44.28
Charge	2+	Mass error	0.002622



Info

Modified Sequence AcD3-YVAK<AcD3K*>GEDVNR<C13>-COOH

Ion score 60.27

Protein P62774

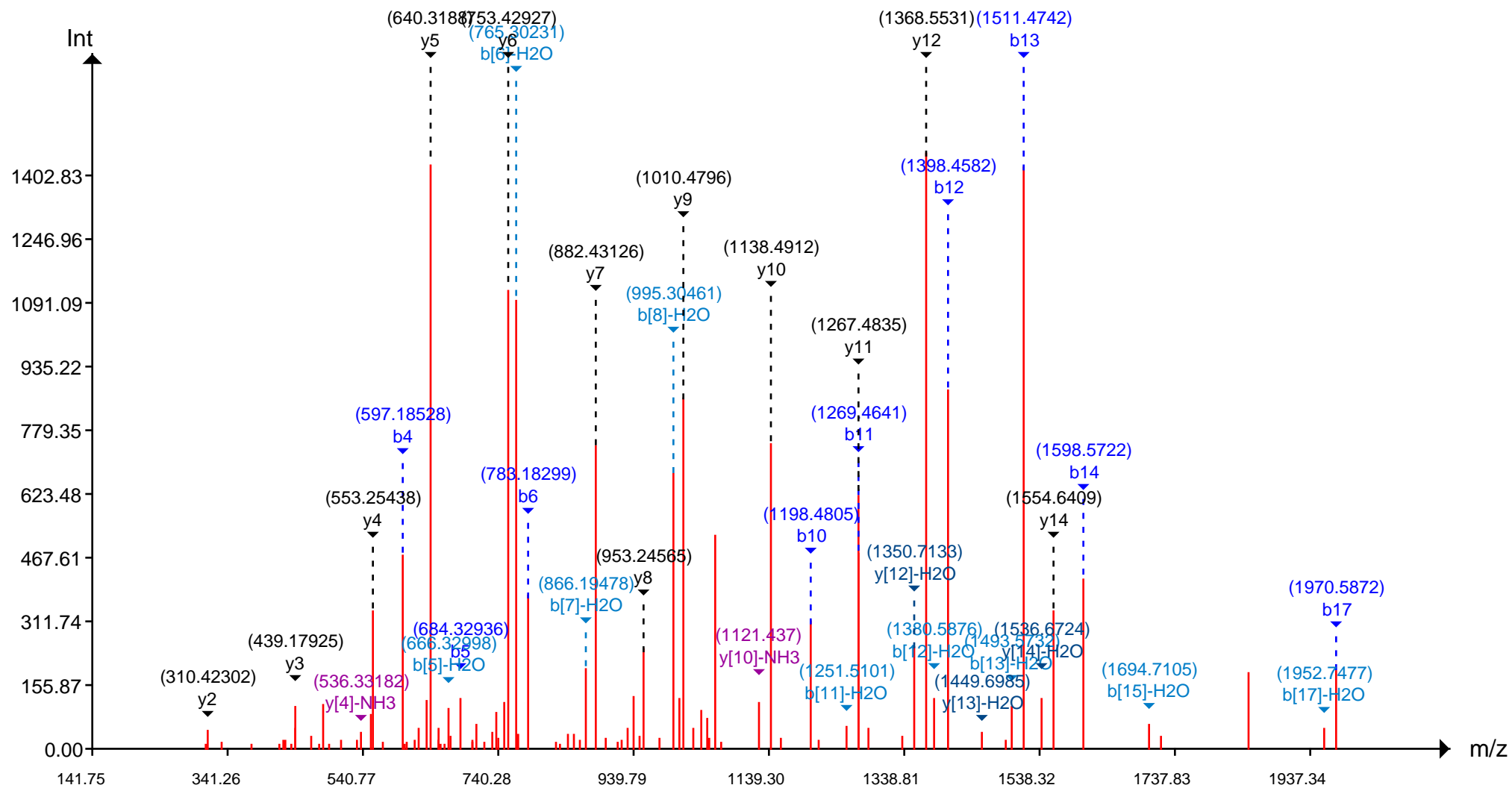
Confidence level 99.00%

Precursor m/z 623.83619

Identity Threshold 48.07

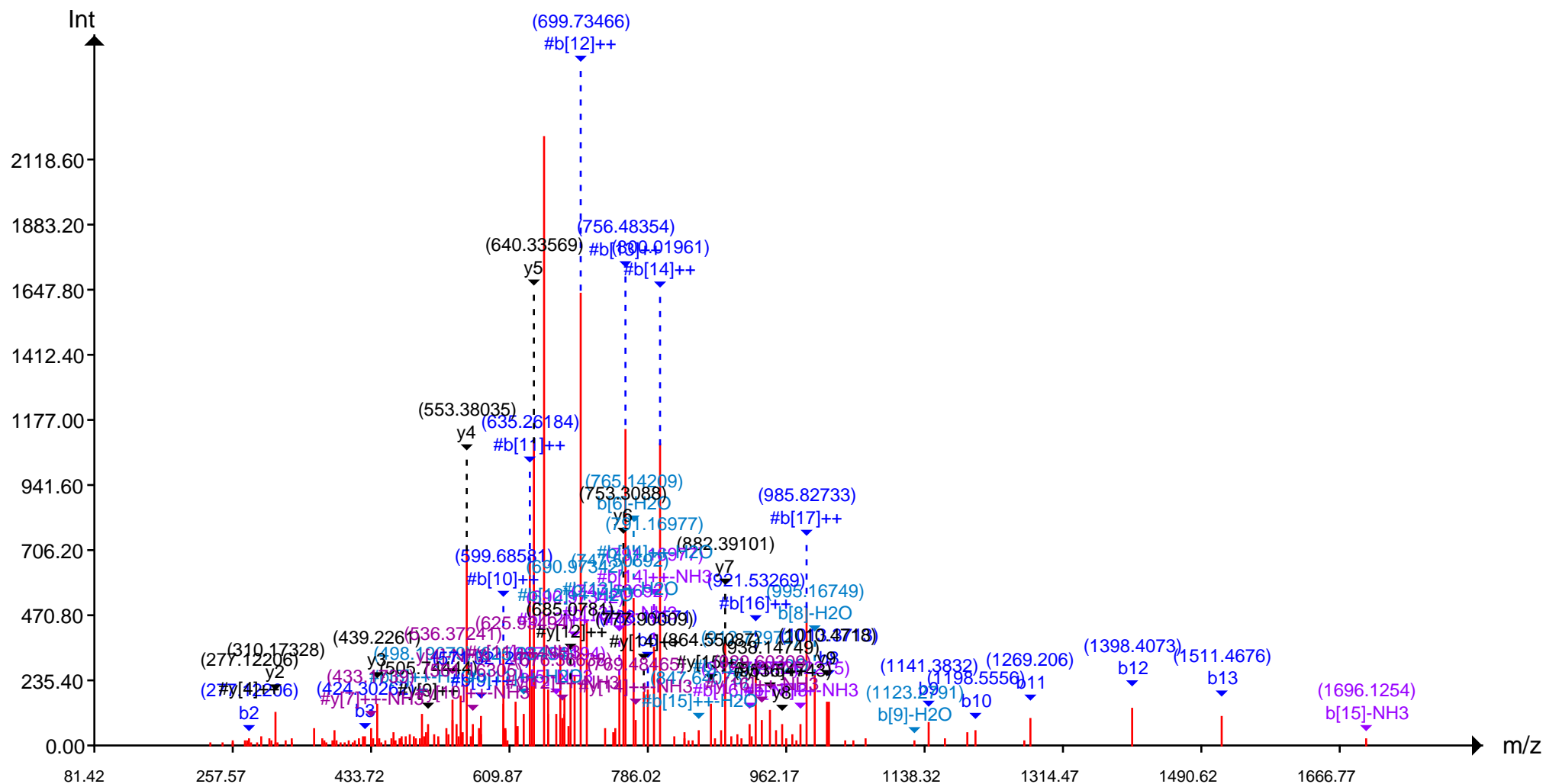
Charge 2+

Mass error 0.001074



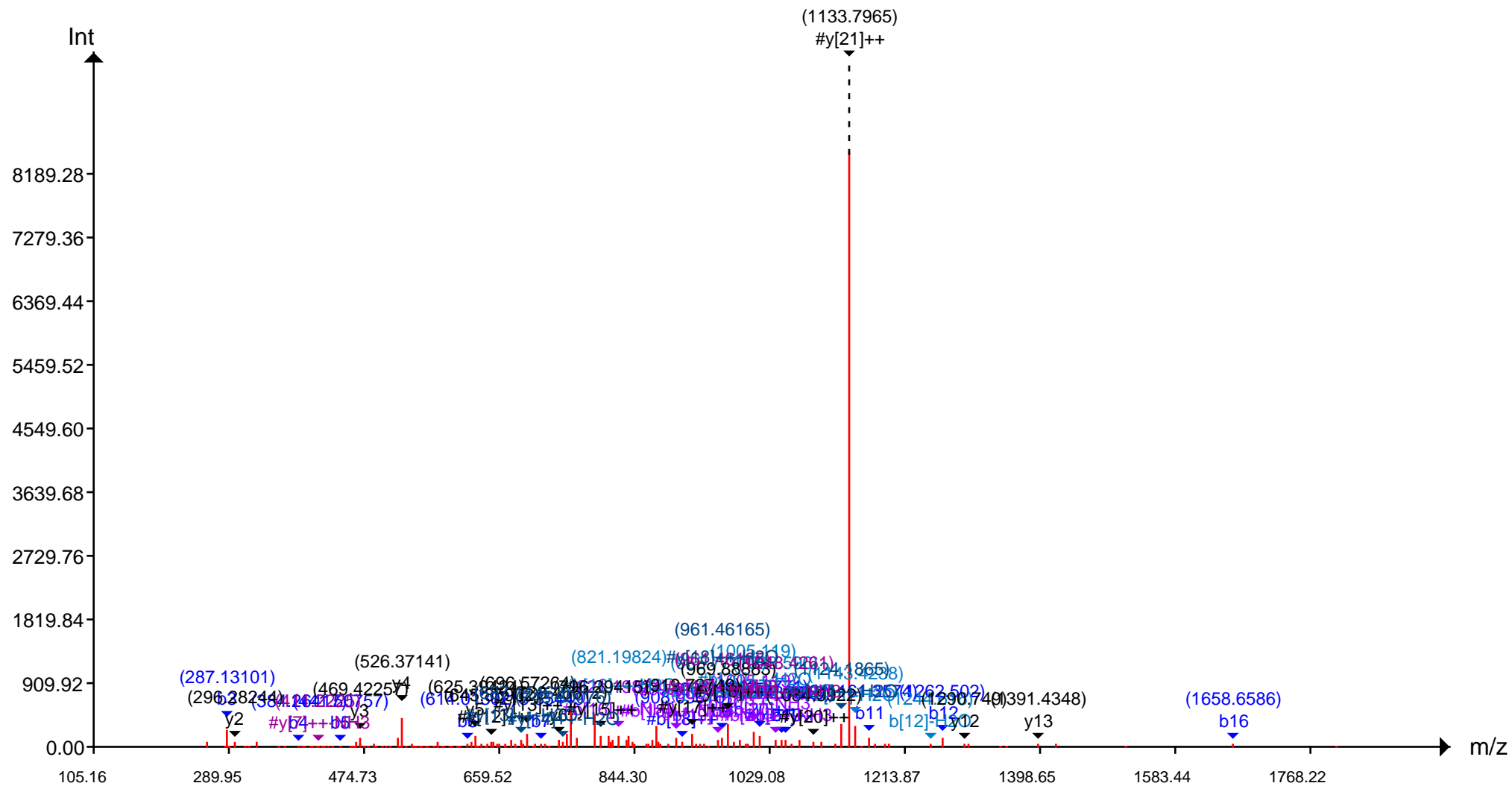
Info

Modified Sequence	AcD3-AC<Cmm*>M<Mox*>K<AcD3K*>SVTEQGAELSNEER<C13>-COOH	Ion score	109.04
Protein	P63101	Confidence level	99.00%
Precursor m/z	1076.0001	Identity Threshold	45.24
Charge	2+	Mass error	0.00228



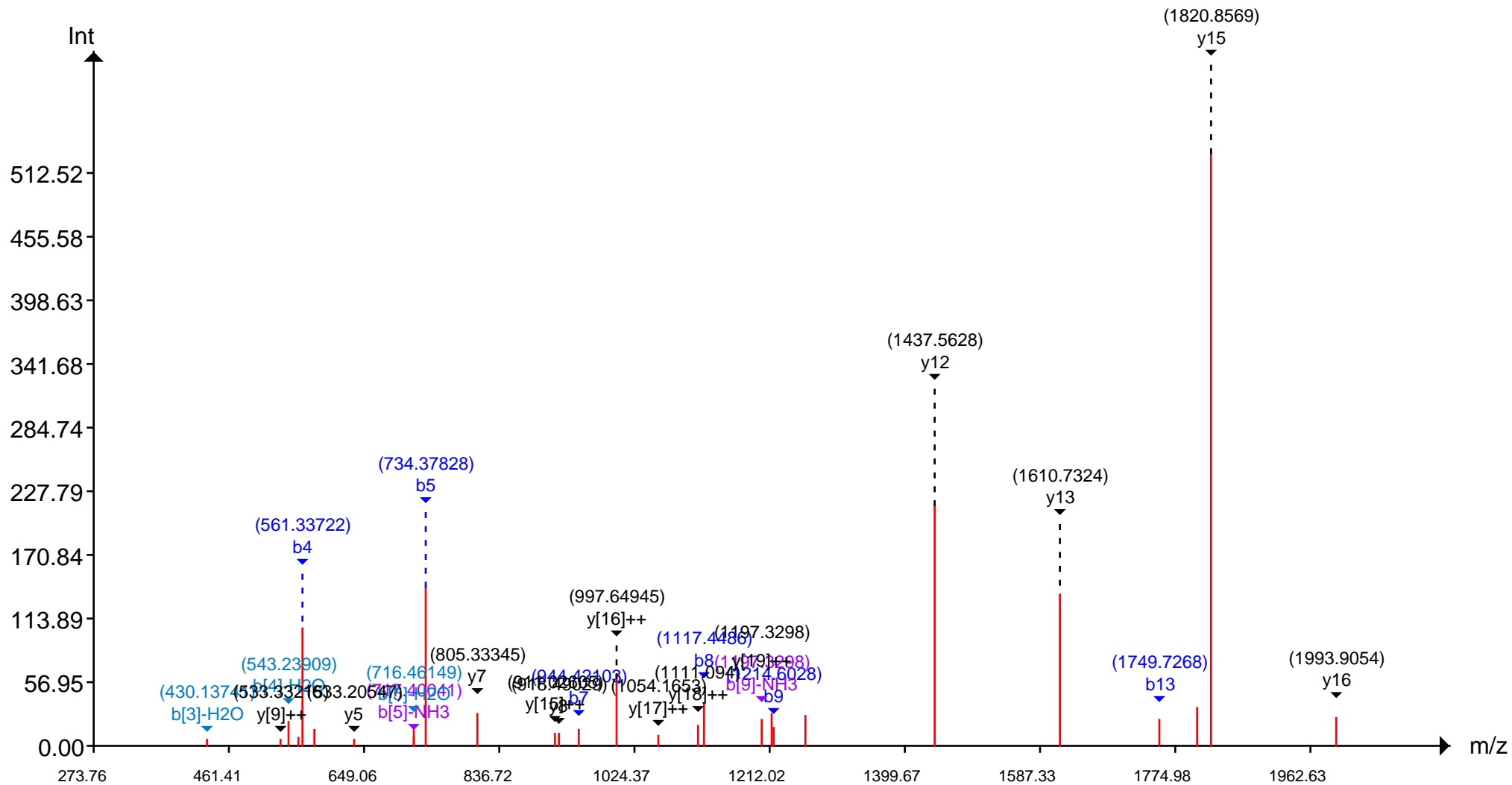
Info

Modified Sequence	AcD3-AC<Cmm*>M<Mox*>K<AcD3K*>SVTEQGAELSNEER<C13>-COOH	Ion score	61.14
Protein	P63101	Confidence level	99.00%
Precursor m/z	717.66818	Identity Threshold	45.07
Charge	3+	Mass error	-6.56E-4



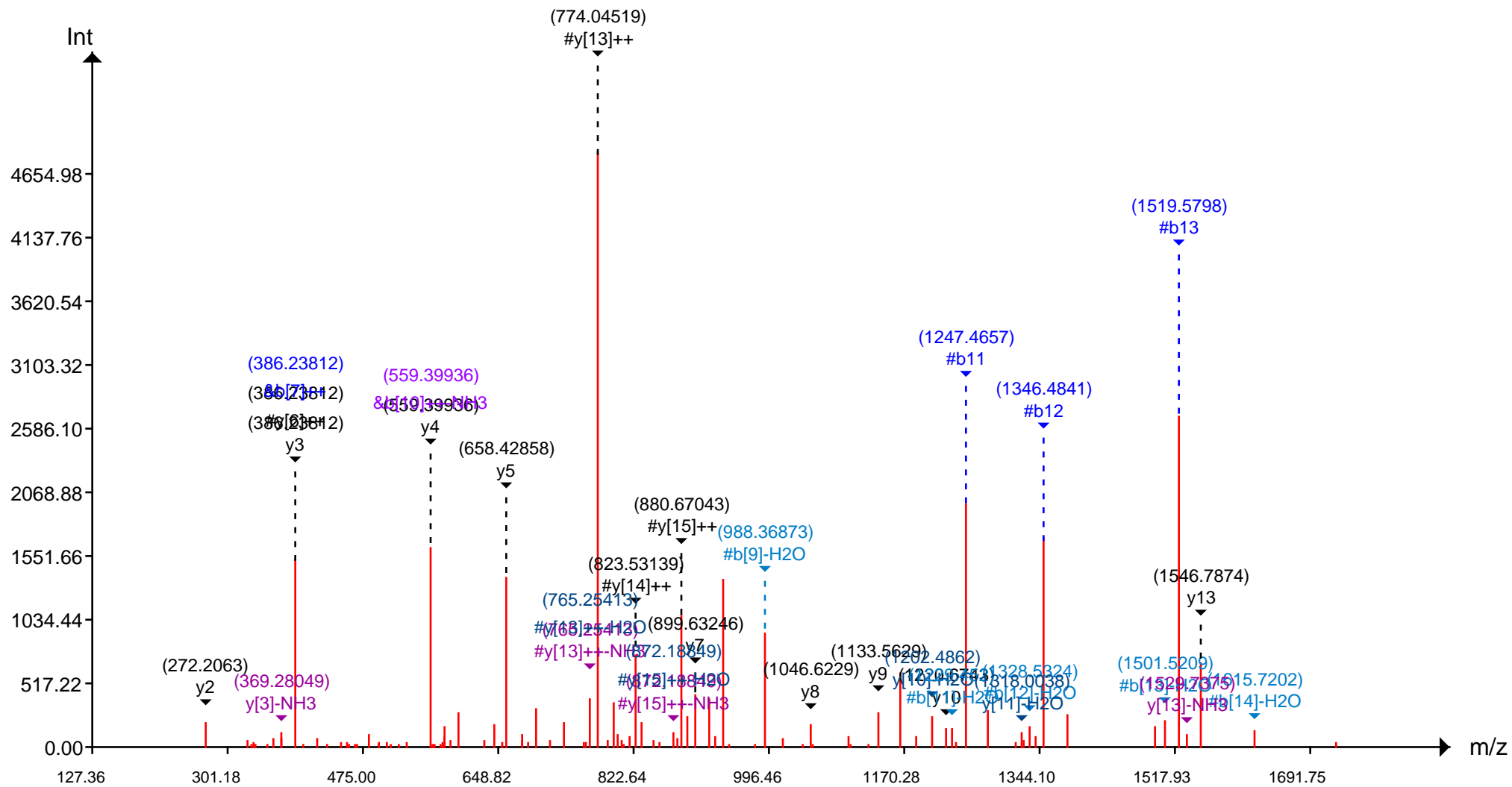
Info

Modified Sequence	AcD3-AAVPGK<AcD3K*>TFVSITPAEVGVLVGK<AcD3K*>DR<C13>-COOH	Ion score	44.47
Protein	P62962	Confidence level	99.00%
Precursor m/z	851.49705	Identity Threshold	42.20
Charge	3+	Mass error	0.002571



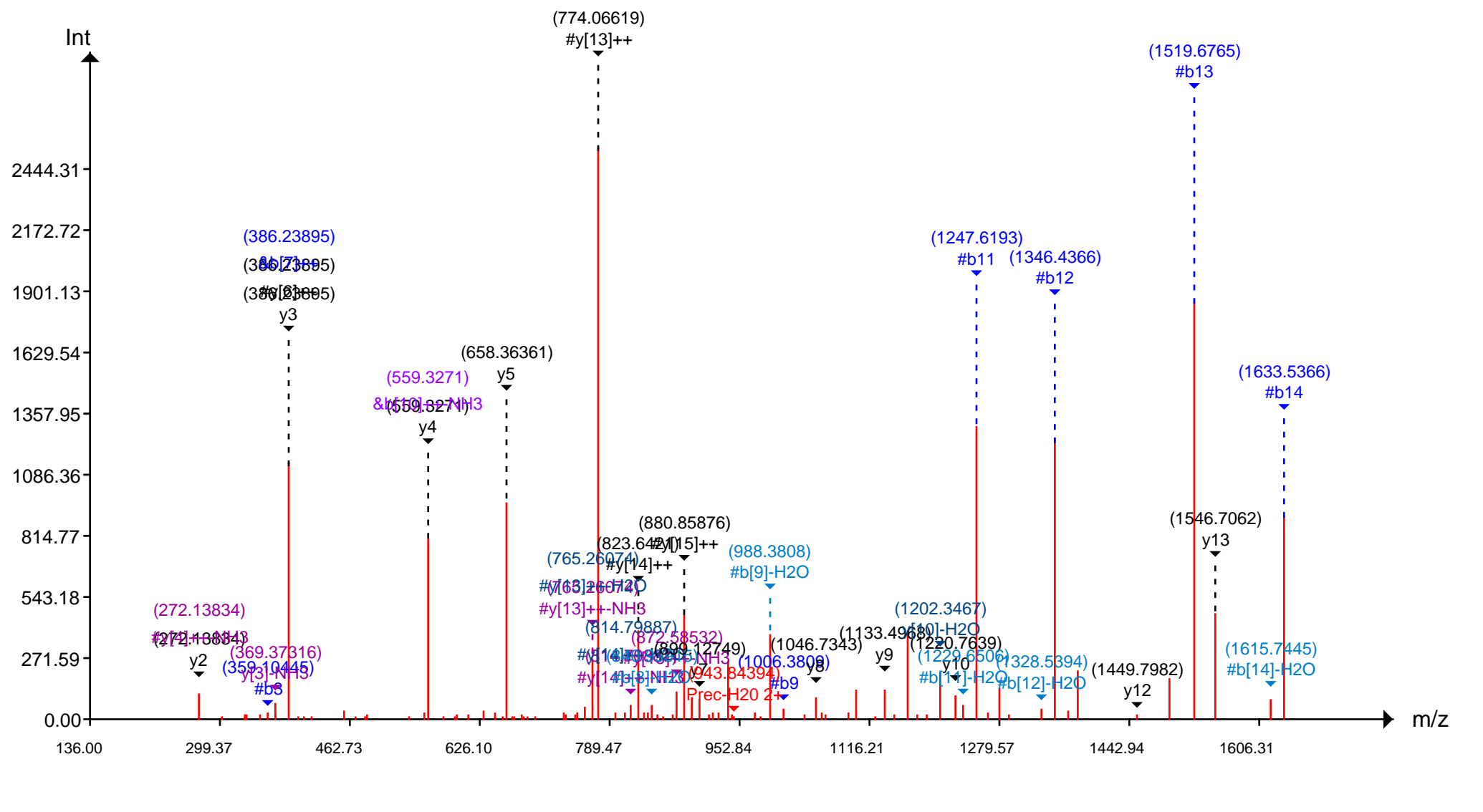
Info

Modified Sequence	AcD3-DK<AcD3K*>NLK<AcD3K*>PIK<AcD3K*>PM<Mox*>QFLGDEETVR-COOH	Ion score	62.45
Protein	Q9DBJ1	Confidence level	99.00%
Precursor m/z	1277.6861	Identity Threshold	47.20
Charge	2+	Mass error	0.003842

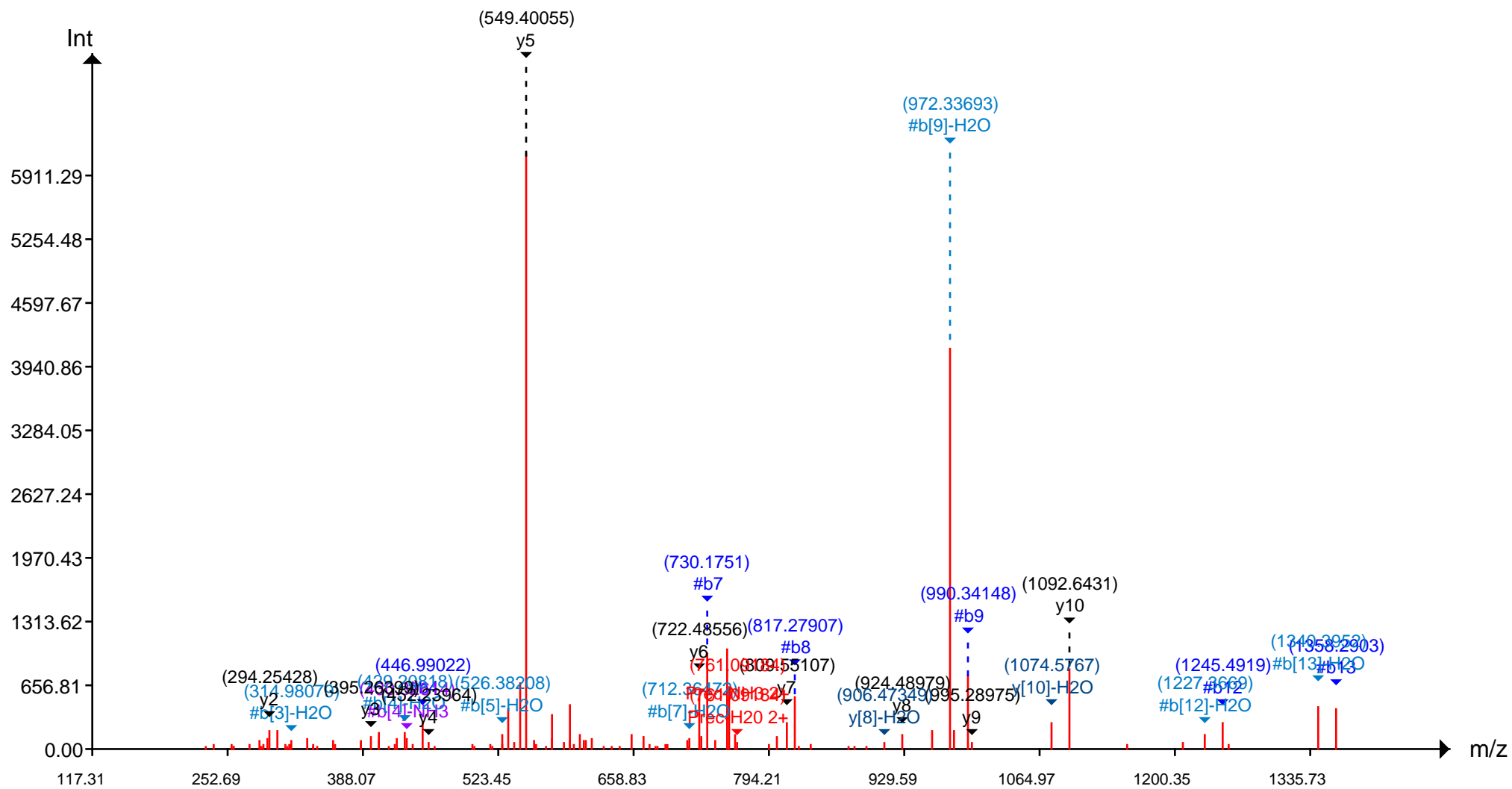


Info

Modified Sequence	AcD3-VDVPNDSSFQIVK<AcD3K*>NPR-COOH	Ion score	53.41
Protein	Q61176	Confidence level	99.00%
Precursor m/z	953.00435	Identity Threshold	48.16
Charge	2+	Mass error	0.00309

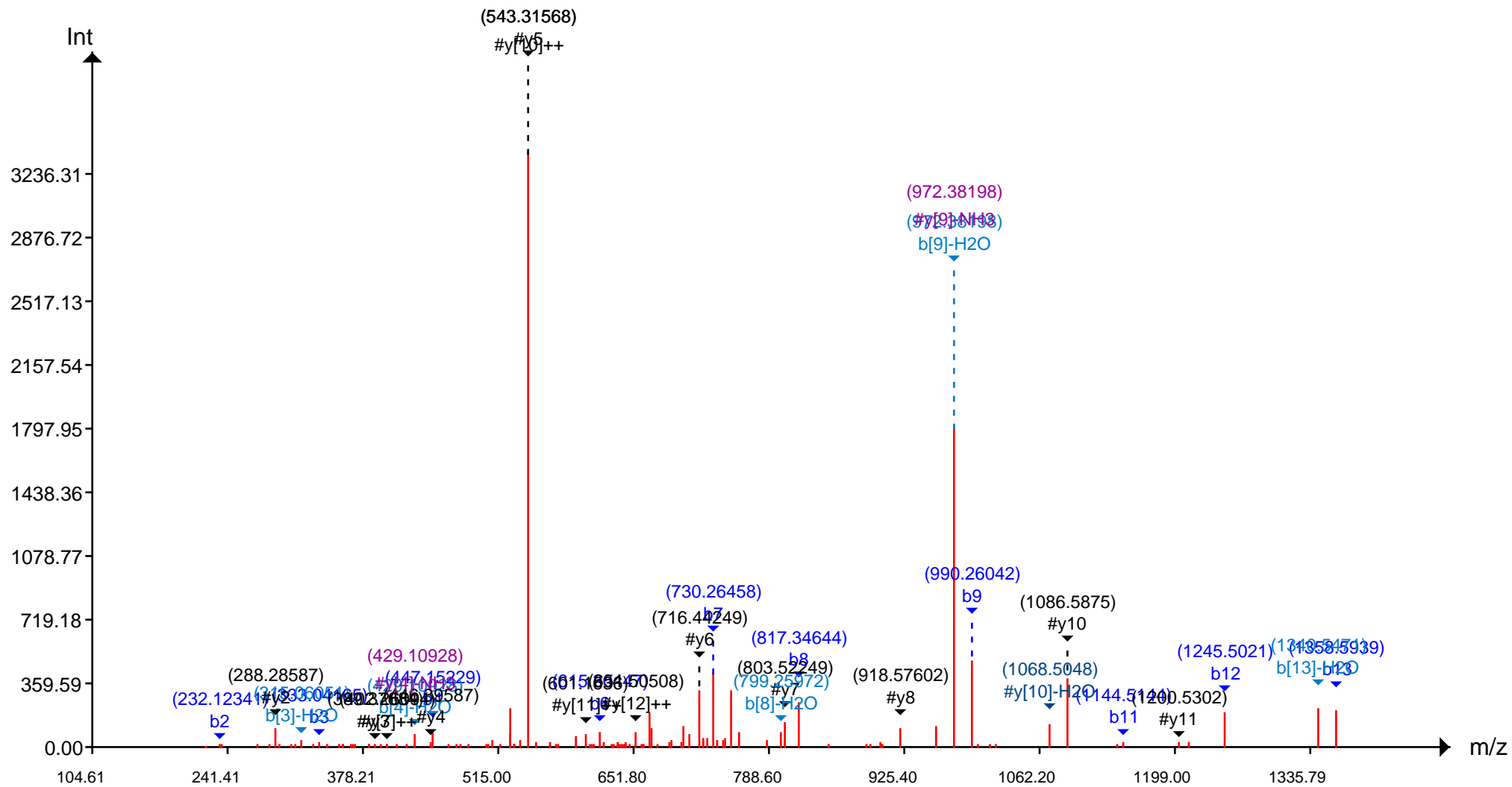


Info			
Modified Sequence	AcD3-VDVPNDSSFQIVK<AcD3K*>NPR-COOH	Ion score	59.2
Protein	Q61176	Confidence level	99.00%
Precursor m/z	953.00402	Identity Threshold	48.15
Charge	2+	Mass error	0.00243



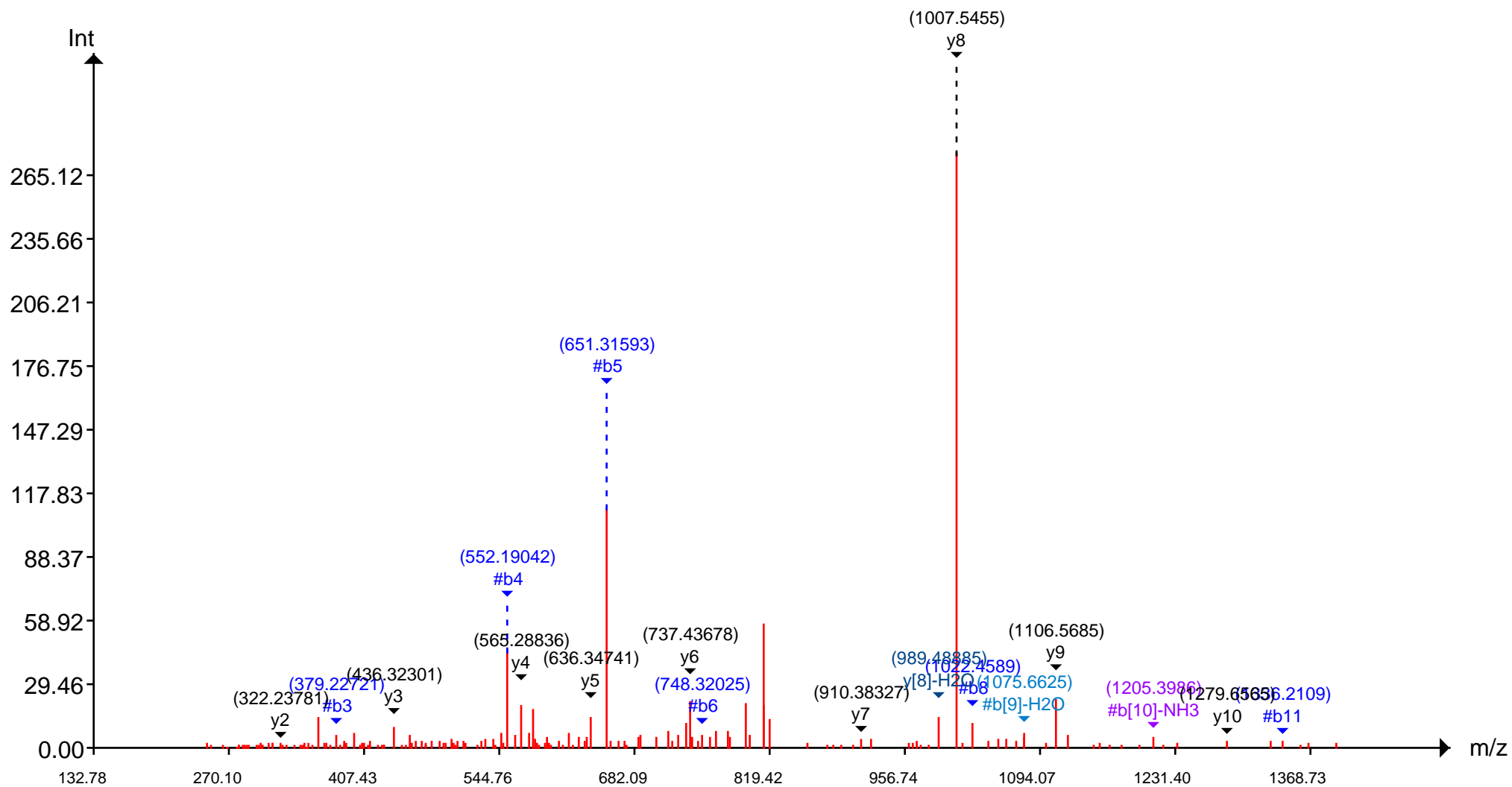
Info

Modified Sequence	AcD3-GETNPADSK<AcD3K*>PGTIR<C13>-COOH	Ion score	54.09
Protein	P15532, Q01768	Confidence level	99.00%
Precursor m/z	769.90473	Identity Threshold	48.13
Charge	2+	Mass error	-1.36E-4



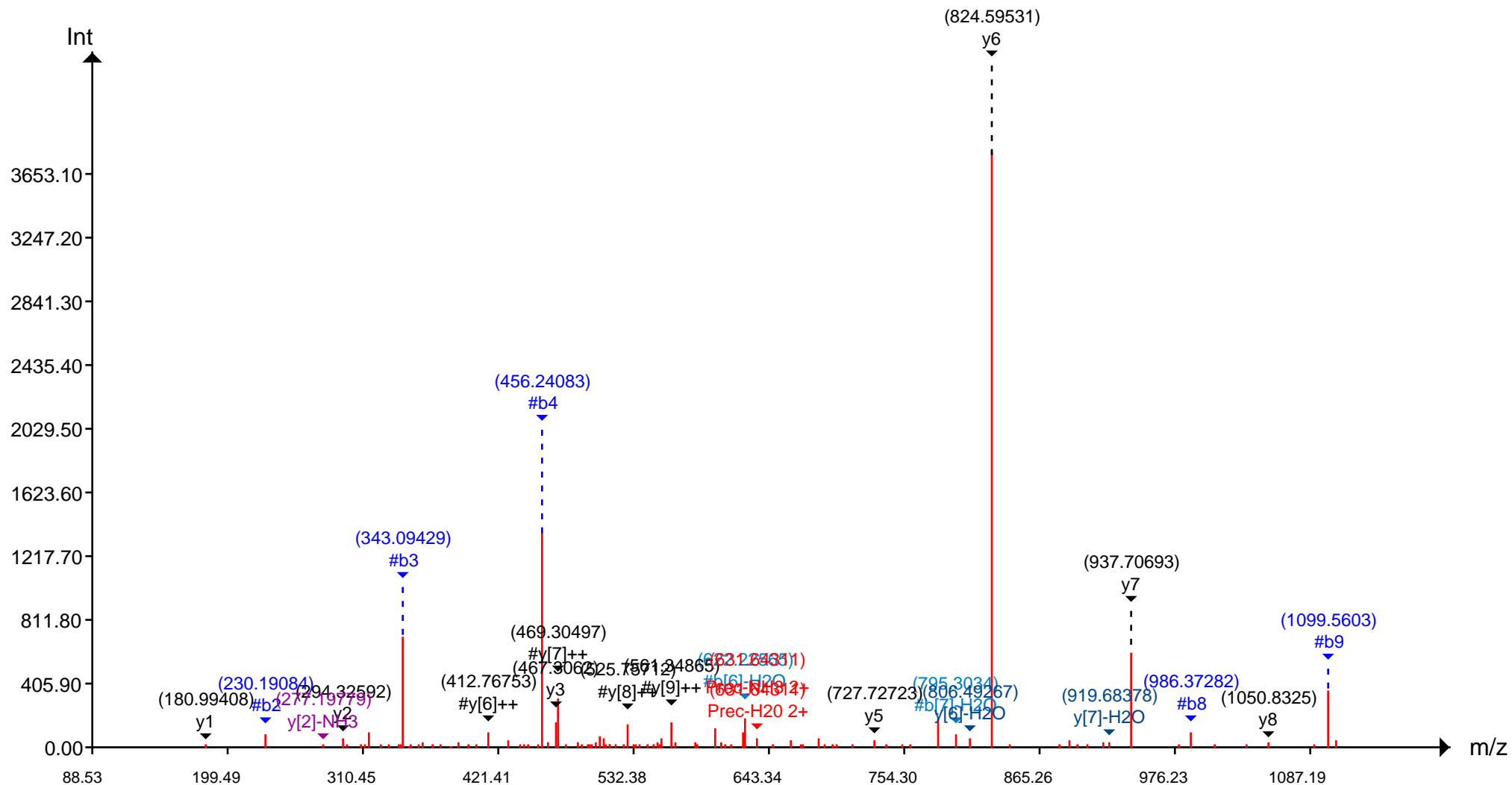
Info

Modified Sequence	AcD3-GETNPADSK<AcD3K*>PGTIR-COOH	Ion score	77.24
Protein	P15532, Q01768	Confidence level	99.00%
Precursor m/z	766.89491	Identity Threshold	48.07
Charge	2+	Mass error	3.66E-4



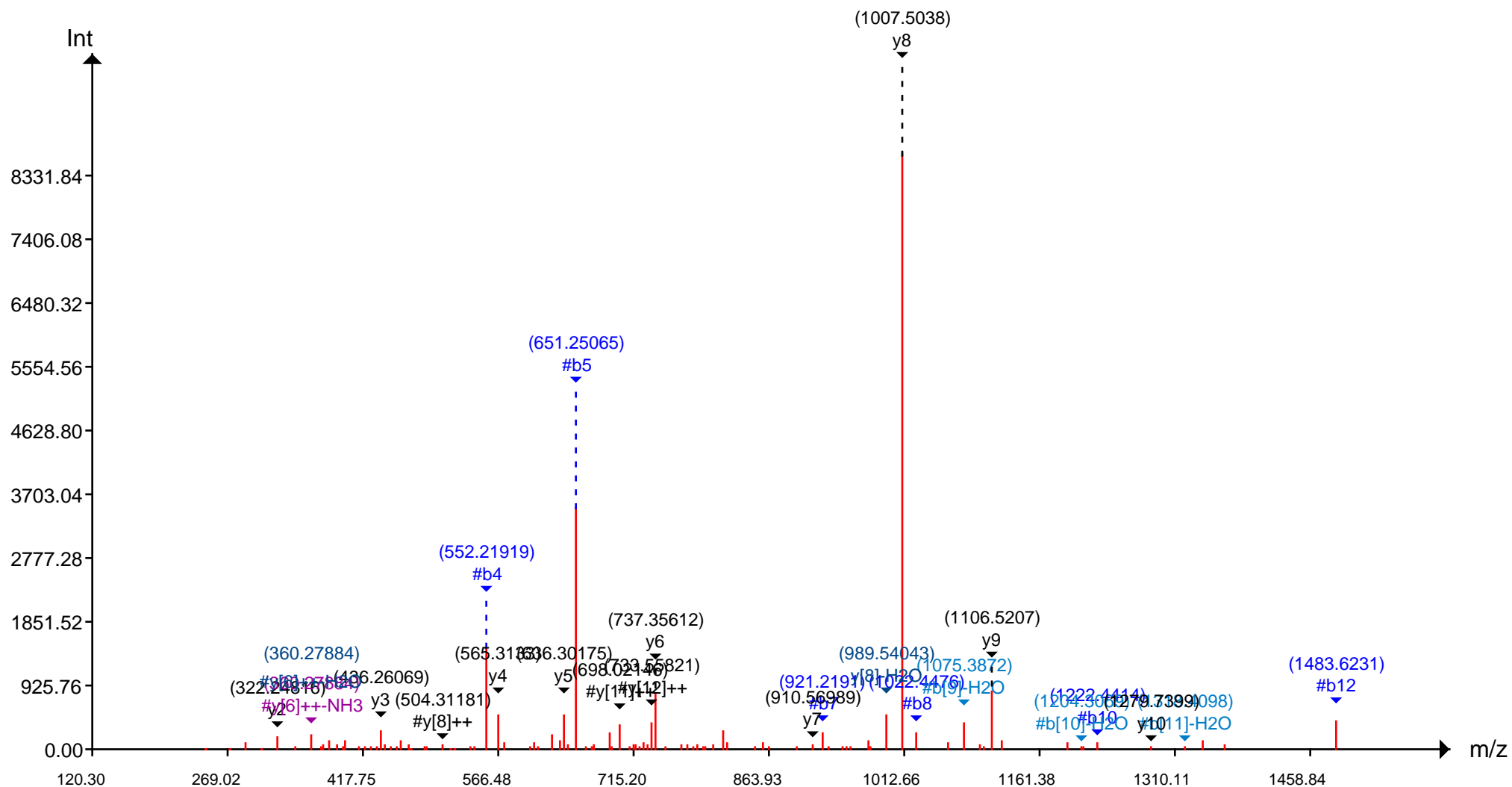
Info

Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH	Ion score	49.36
Protein	P17742	Confidence level	99.00%
Precursor m/z	829.44951	Identity Threshold	47.80
Charge	2+	Mass error	0.002312



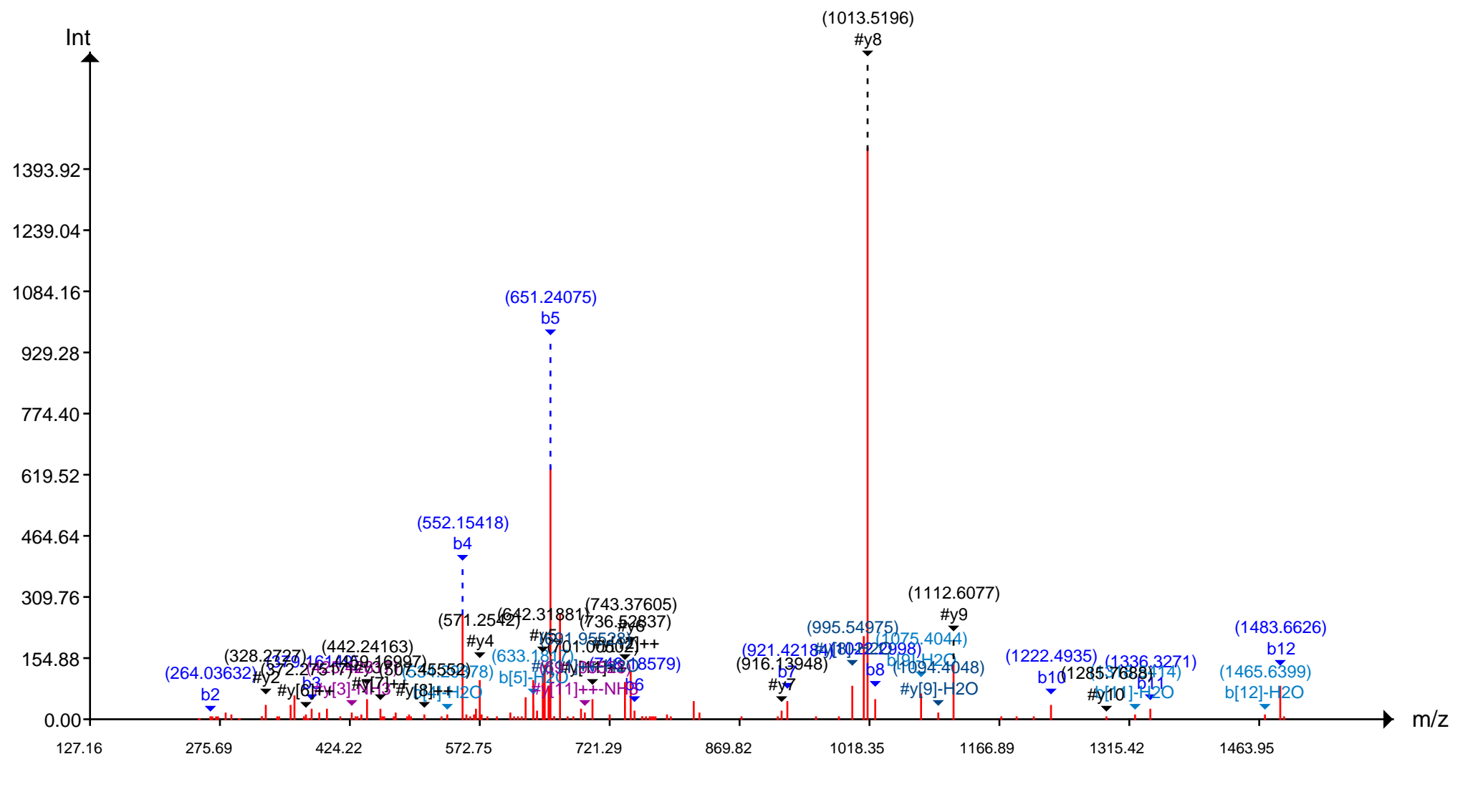
Info

Modified Sequence	AcD3-IAIIPSK<AcD3K*>K<AcD3K*>LR<C13>-COOH	Ion score	49.72
Protein	P63276	Confidence level	99.00%
Precursor m/z	640.44143	Identity Threshold	29.55
Charge	2+	Mass error	2.51E-4

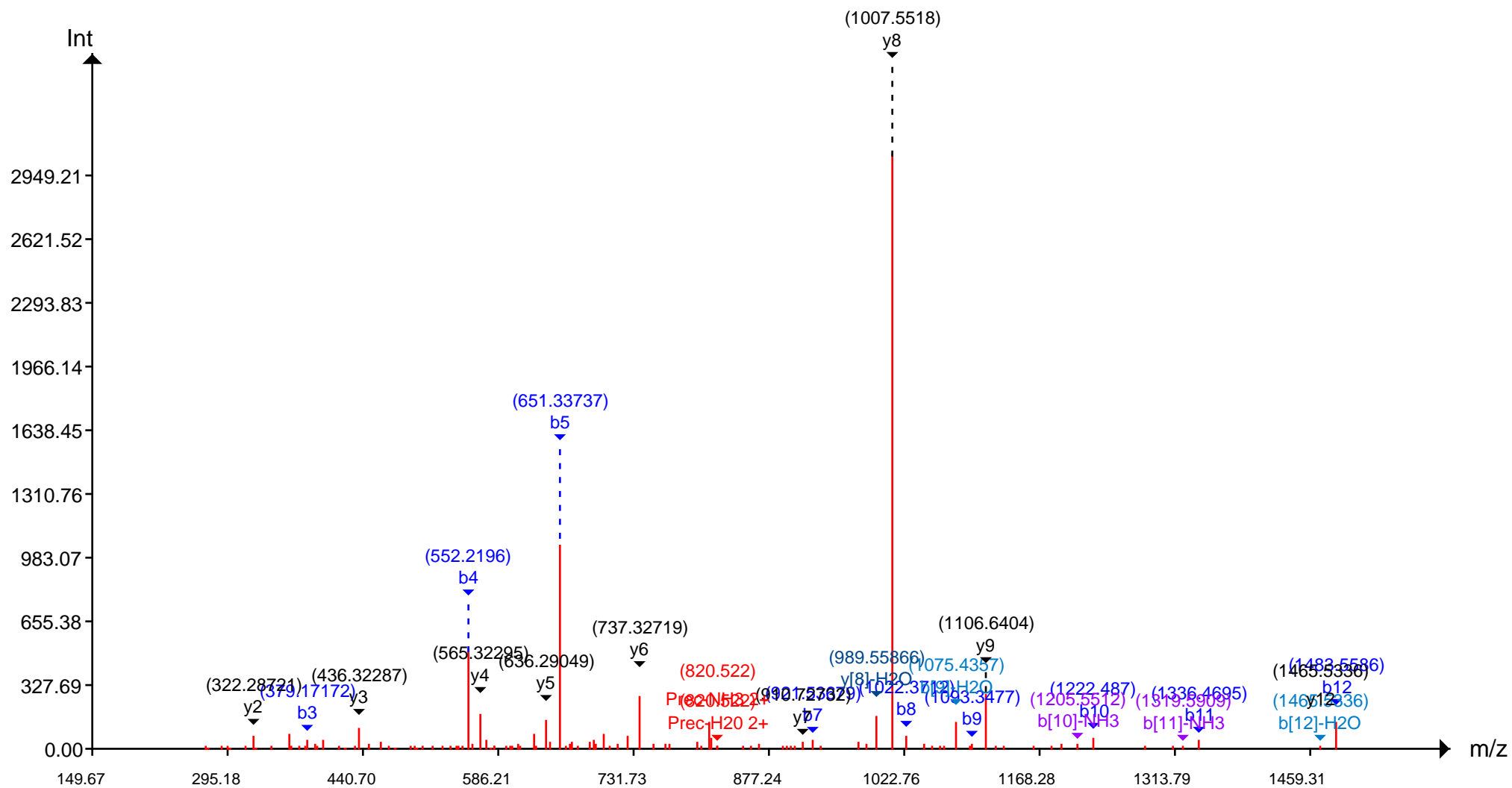


Info

Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH	Ion score	64.05
Protein	P17742	Confidence level	99.00%
Precursor m/z	829.44957	Identity Threshold	47.76
Charge	2+	Mass error	0.002432



Info			
Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH	Ion score	51.59
Protein	P17742	Confidence level	99.00%
Precursor m/z	832.45906	Identity Threshold	47.46
Charge	2+	Mass error	0.00127



Info

Modified Sequence AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH

Ion score 79.8

Protein P17742

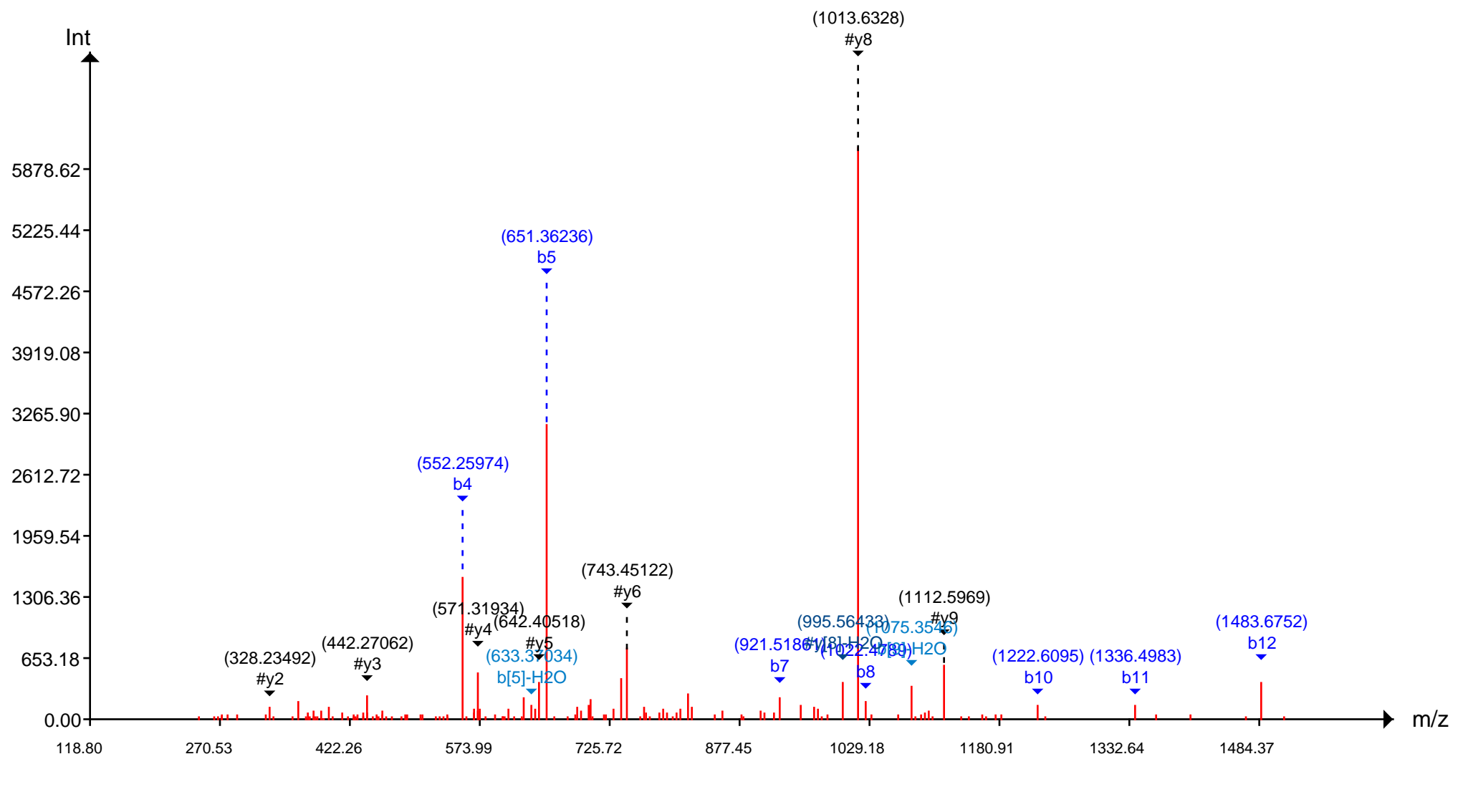
Confidence level 99.00%

Precursor m/z 829.4496

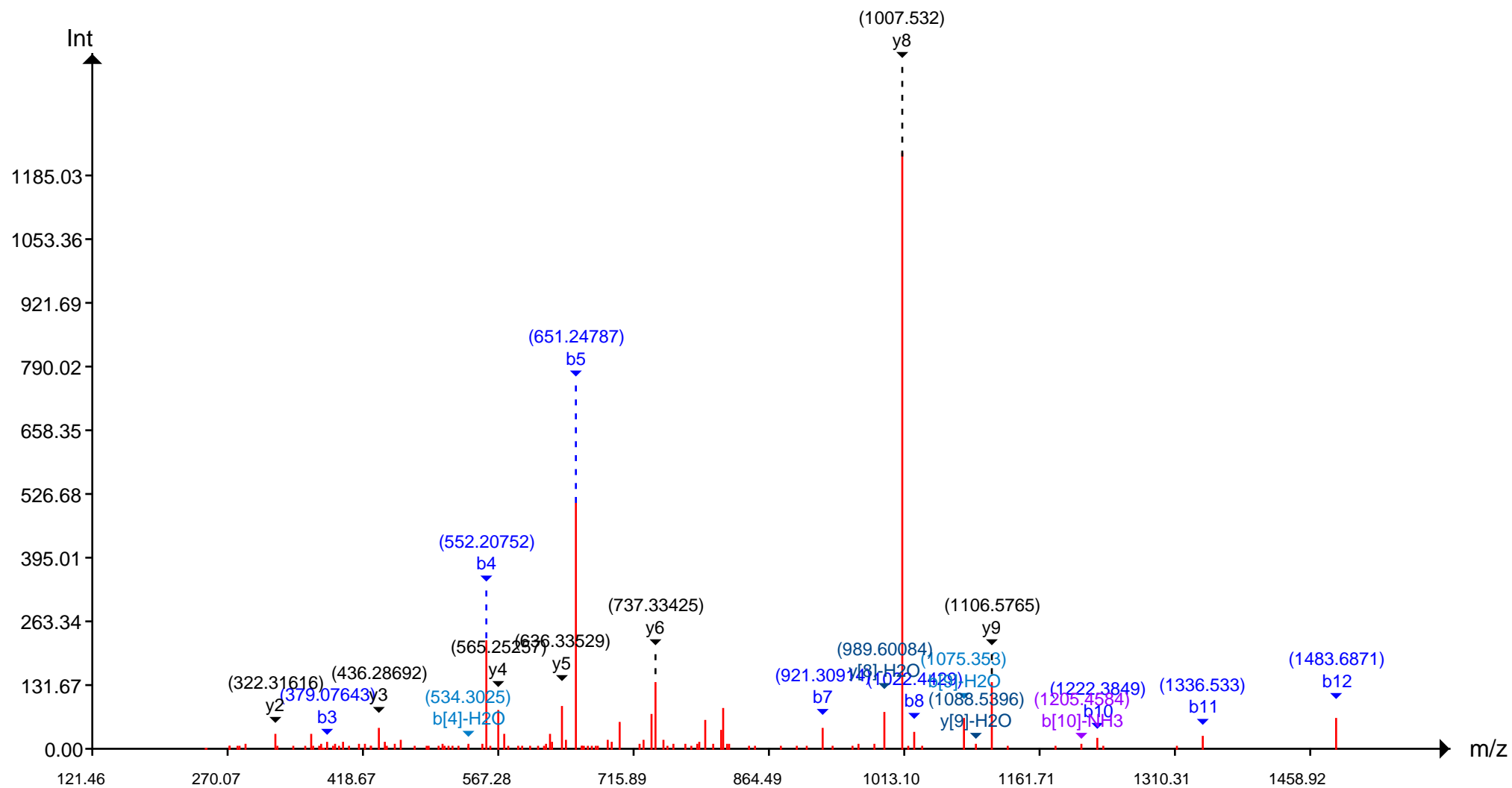
Identity Threshold 47.77

Charge 2+

Mass error 0.002492



Info			
Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH	Ion score	49.07
Protein	P17742	Confidence level	99.00%
Precursor m/z	832.45911	Identity Threshold	47.45
Charge	2+	Mass error	0.00137



Info

Modified Sequence AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH

Ion score 58.11

Protein P17742

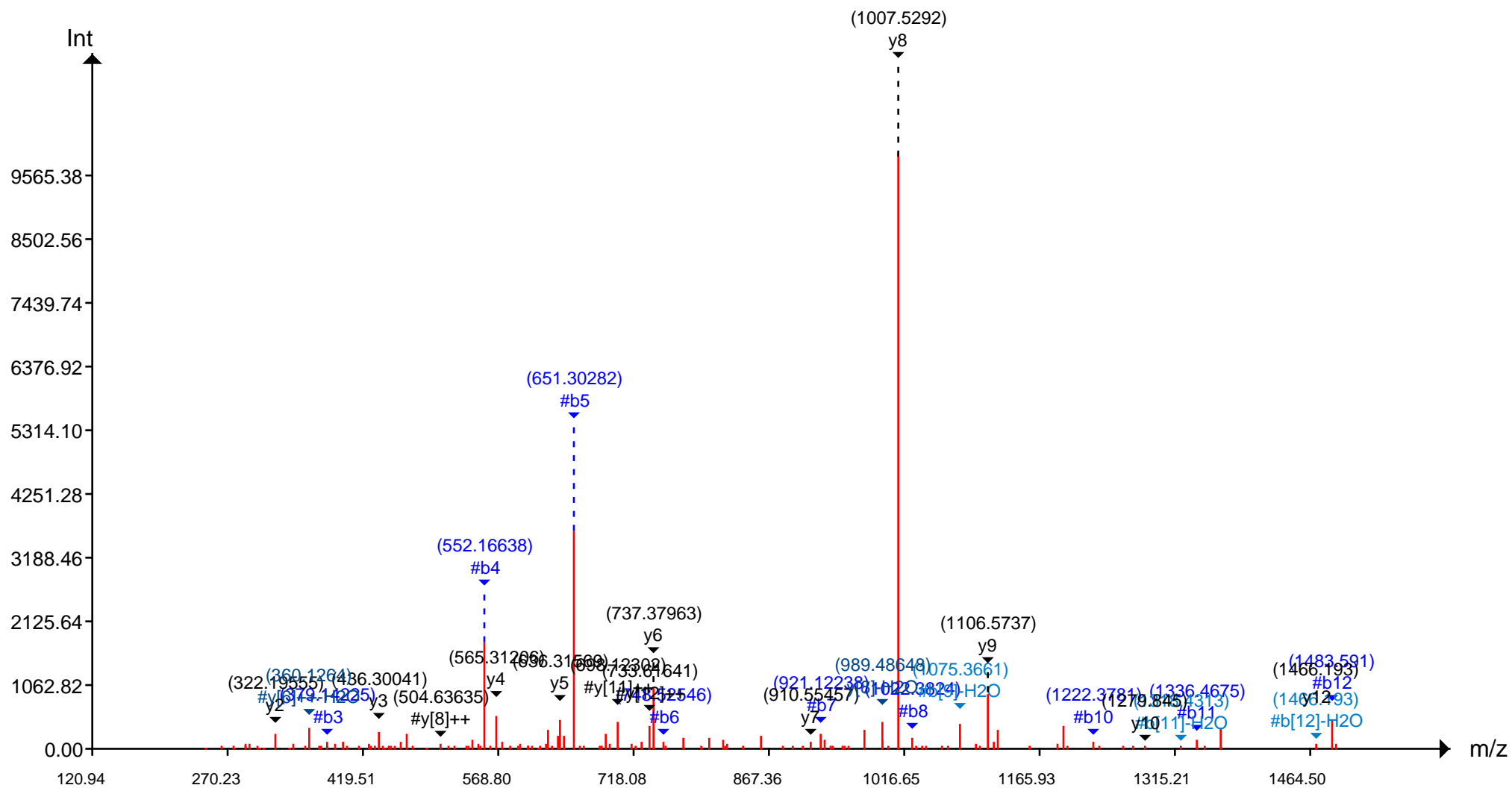
Confidence level 99.00%

Precursor m/z 829.44932

Identity Threshold 47.76

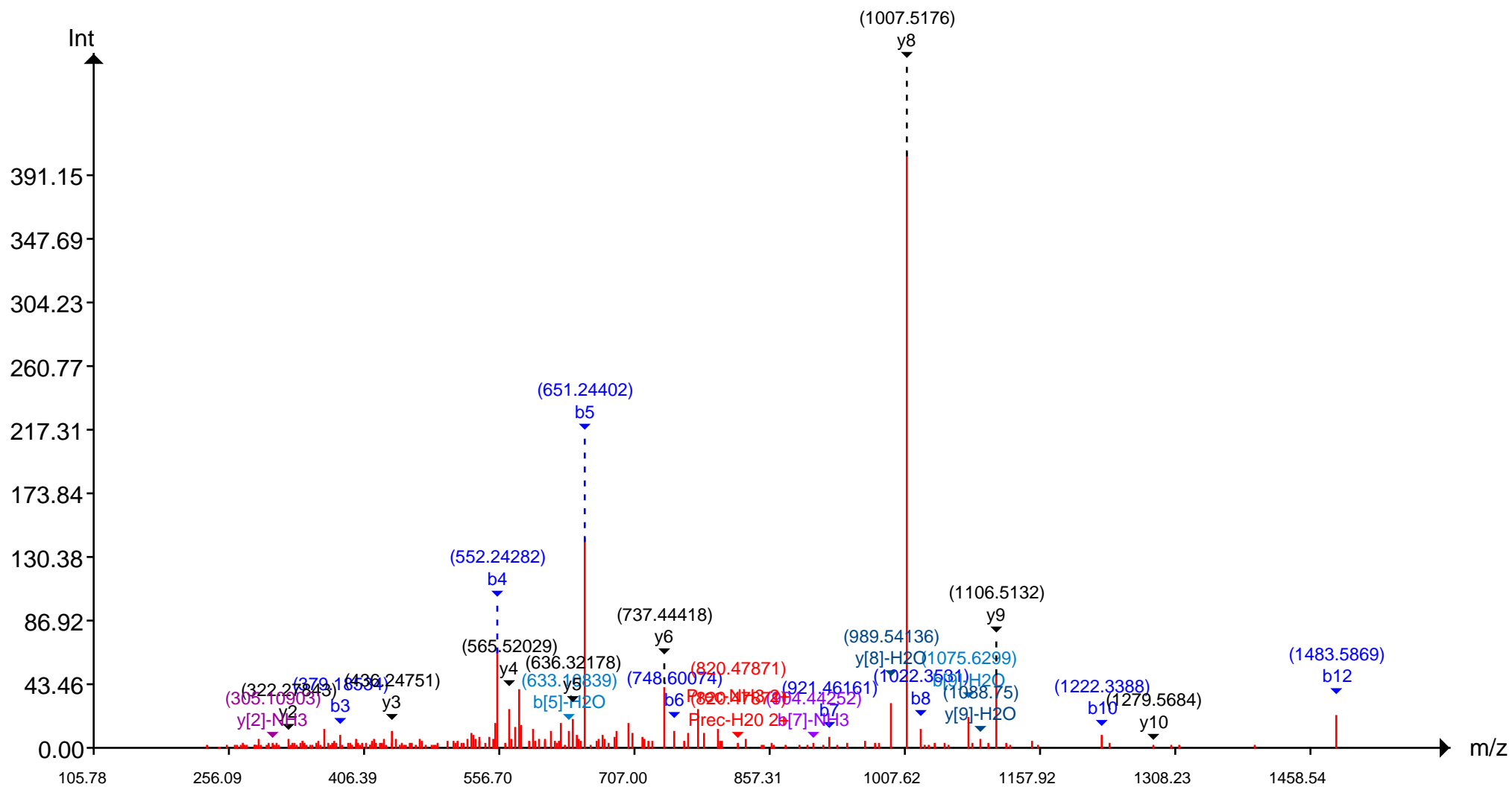
Charge 2+

Mass error 0.001932



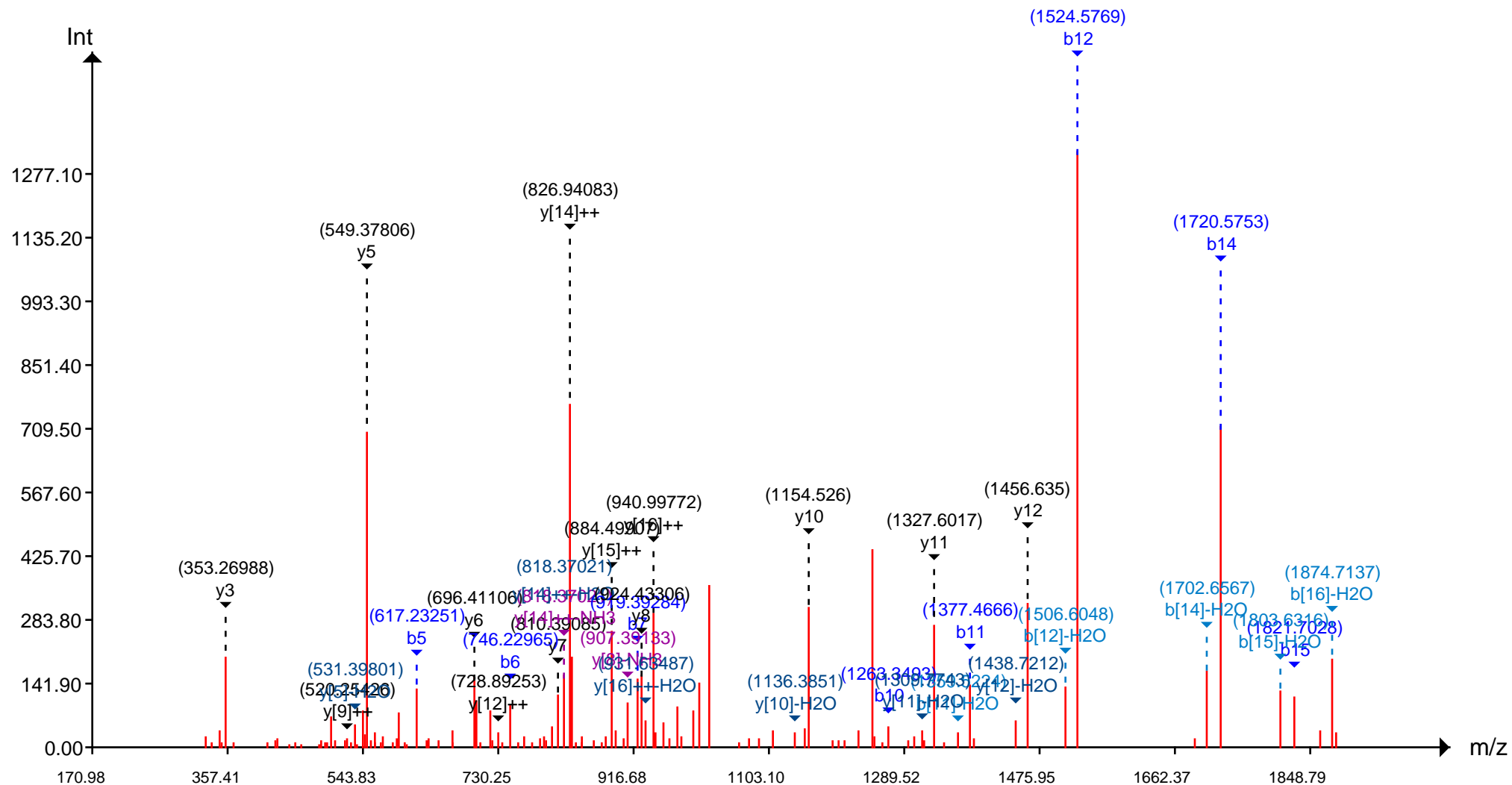
Info

Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH	Ion score	63.65
Protein	P17742	Confidence level	99.00%
Precursor m/z	829.44968	Identity Threshold	47.78
Charge	2+	Mass error	0.002652



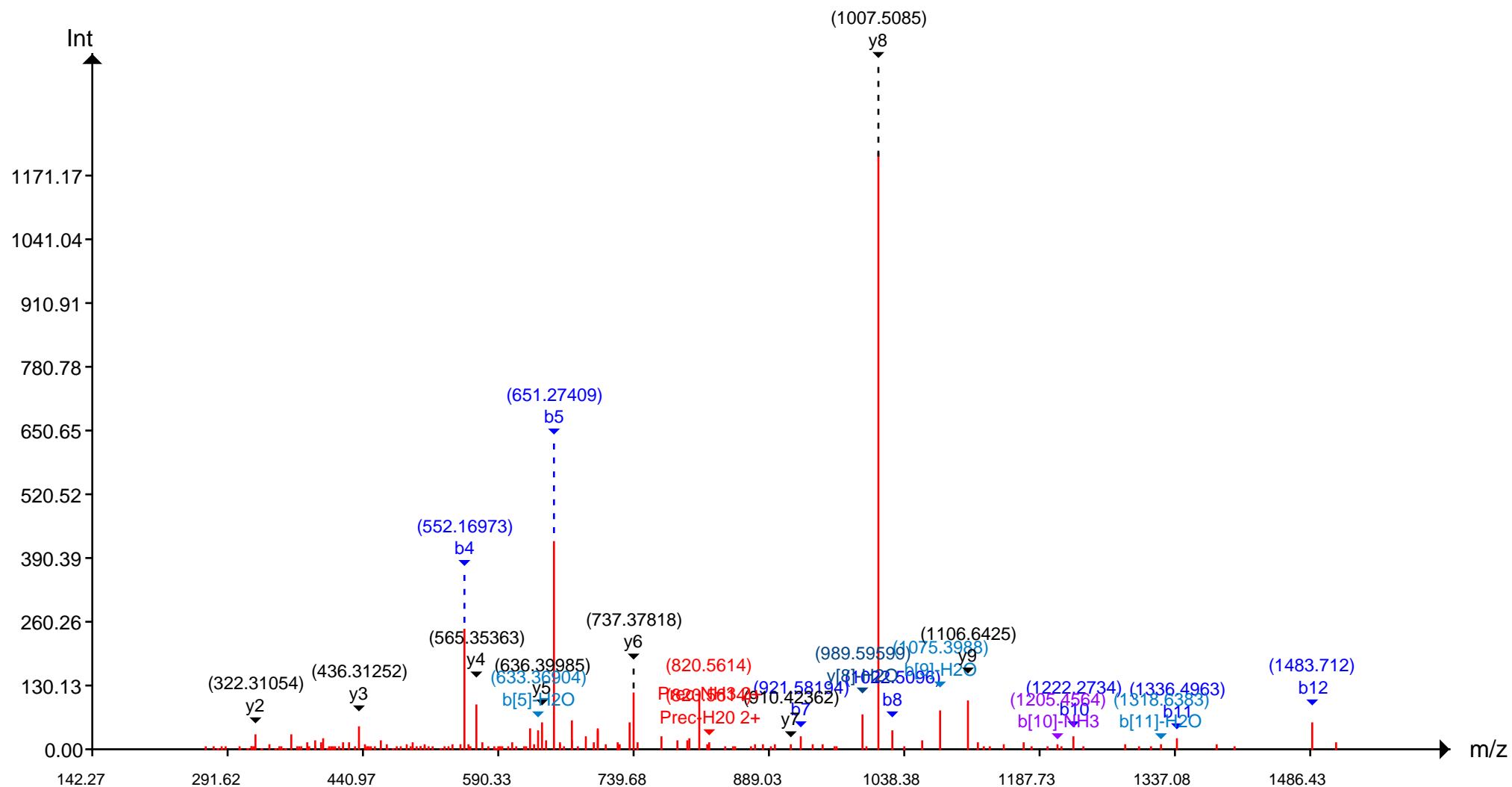
Info

Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH	Ion score	49.57
Protein	P17742	Confidence level	99.00%
Precursor m/z	829.44981	Identity Threshold	47.76
Charge	2+	Mass error	0.002912



Info

Modified Sequence	AcD3-M<Mox*>LDPVEK<AcD3K*>DDNNM<Mox*>PVTAR<C13>-COOH	Ion score	53.22
Protein	O08709	Confidence level	99.00%
Precursor m/z	1036.9961	Identity Threshold	46.44
Charge	2+	Mass error	7.96E-4



Info

Modified Sequence AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH

Ion score 49.74

Protein P17742

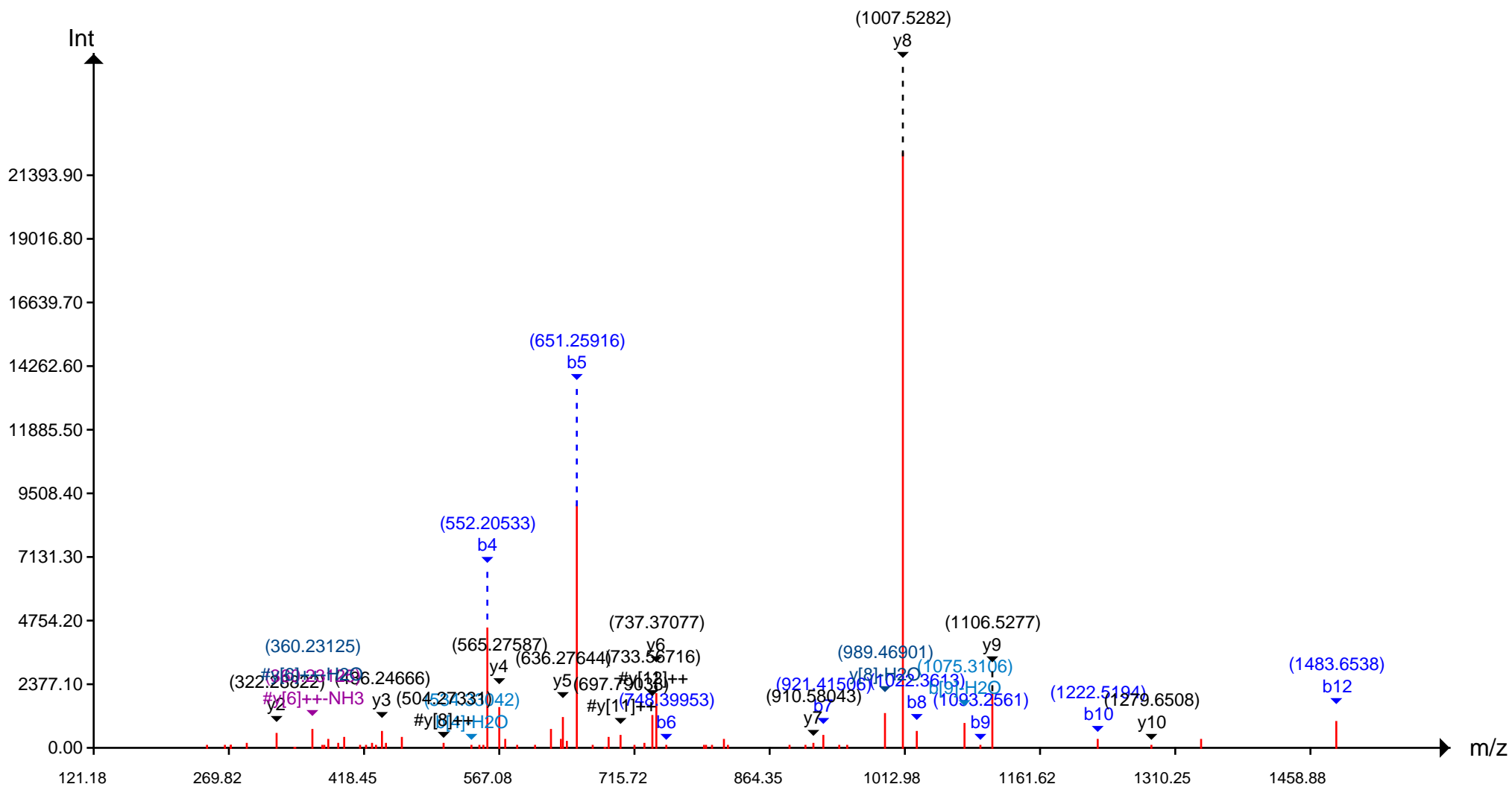
Confidence level 99.00%

Precursor m/z 829.4499

Identity Threshold 47.74

Charge 2+

Mass error 0.003092



Info

Modified Sequence AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH

Ion score 76.03

Protein P17742

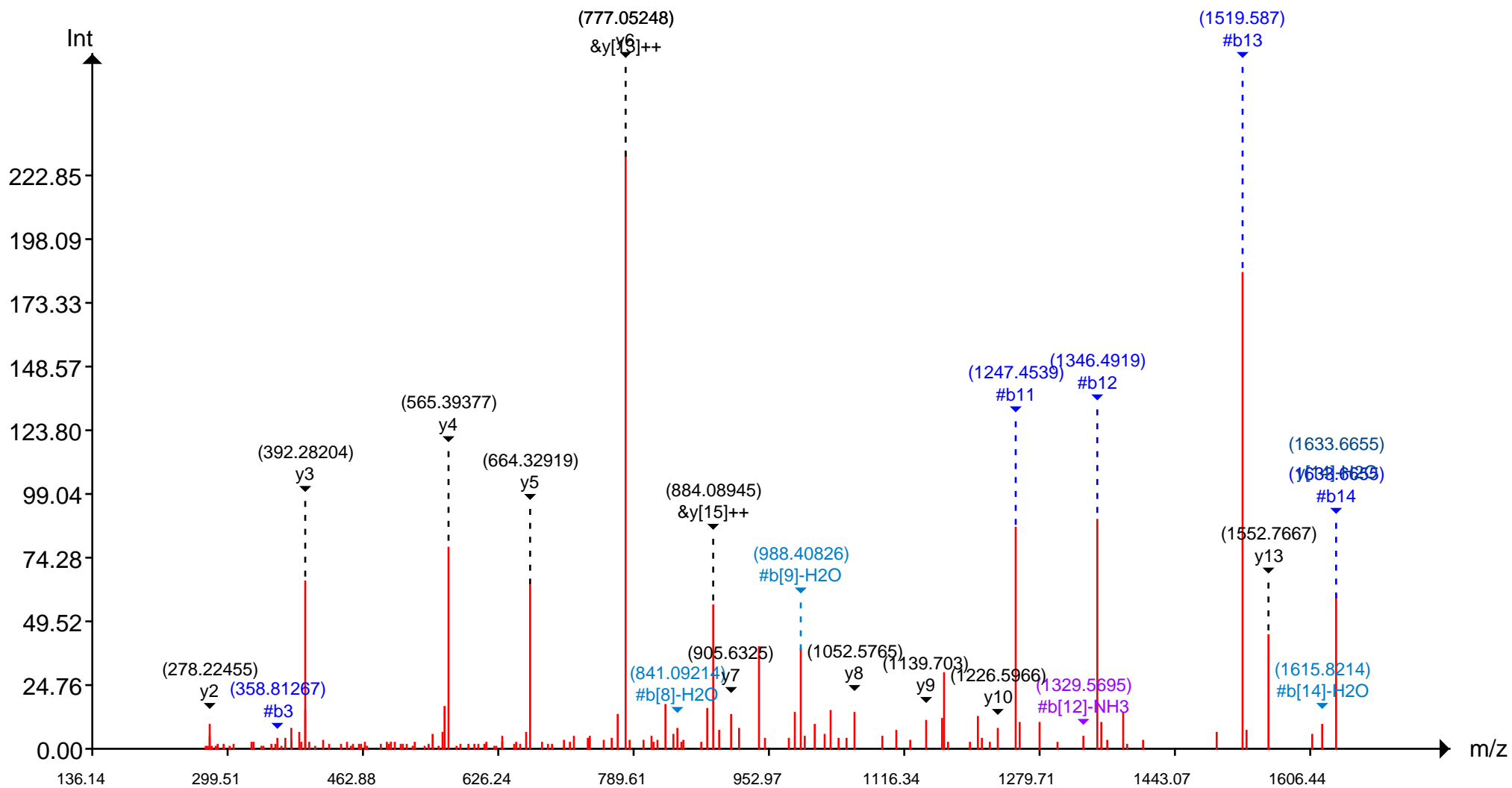
Confidence level 99.00%

Precursor m/z 829.45032

Identity Threshold 47.76

Charge 2+

Mass error 0.003932



Info

Modified Sequence AcD3-VDVPNDSSFQIVK<AcD3K*>NPR<C13>-COOH

Ion score 49.5

Protein Q61176

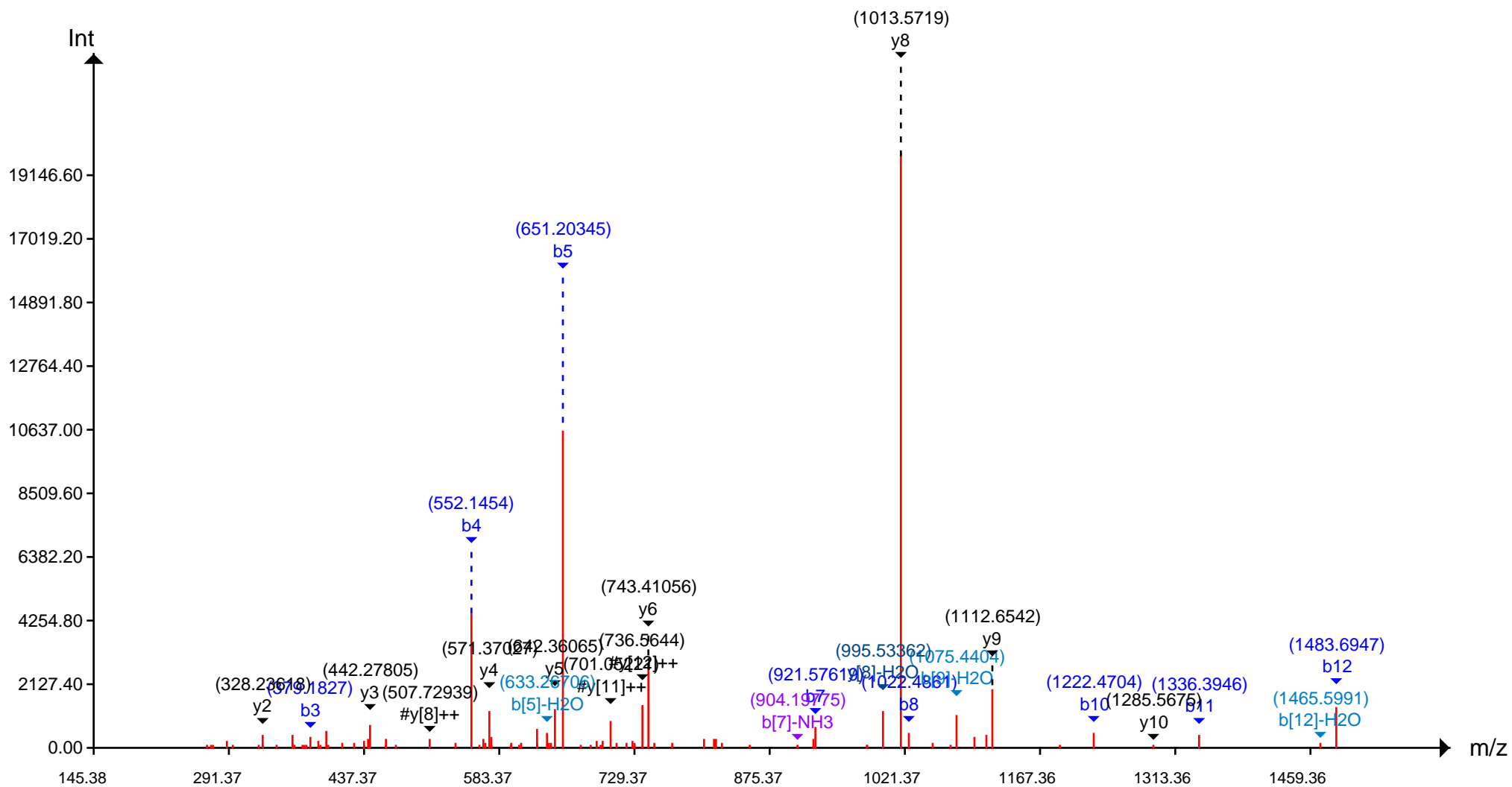
Confidence level 99.00%

Precursor m/z 956.01561

Identity Threshold 47.74

Charge 2+

Mass error 0.005468



Info

Modified Sequence AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH

Ion score 63.43

Protein P17742

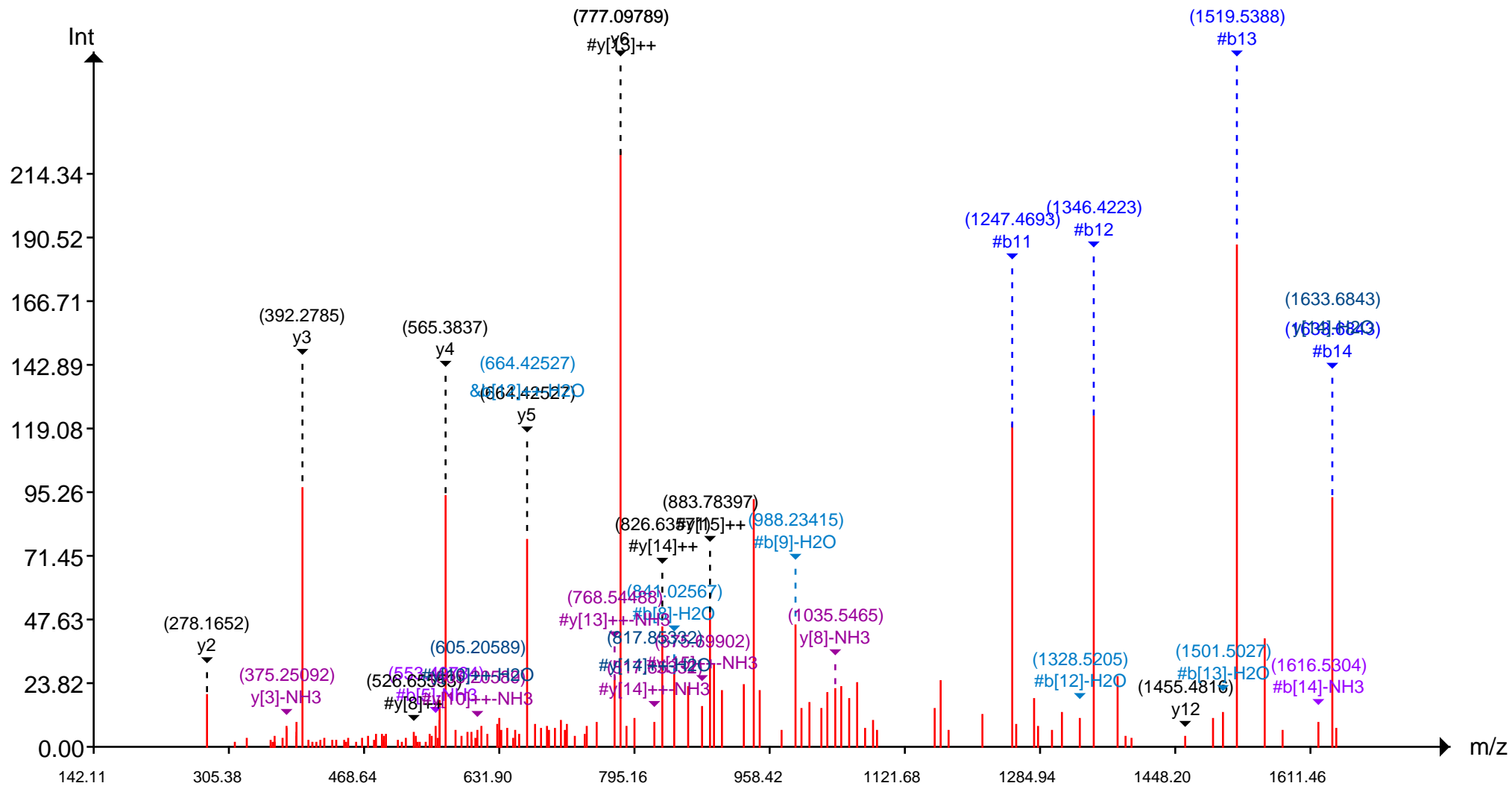
Confidence level 99.00%

Precursor m/z 832.45928

Identity Threshold 47.47

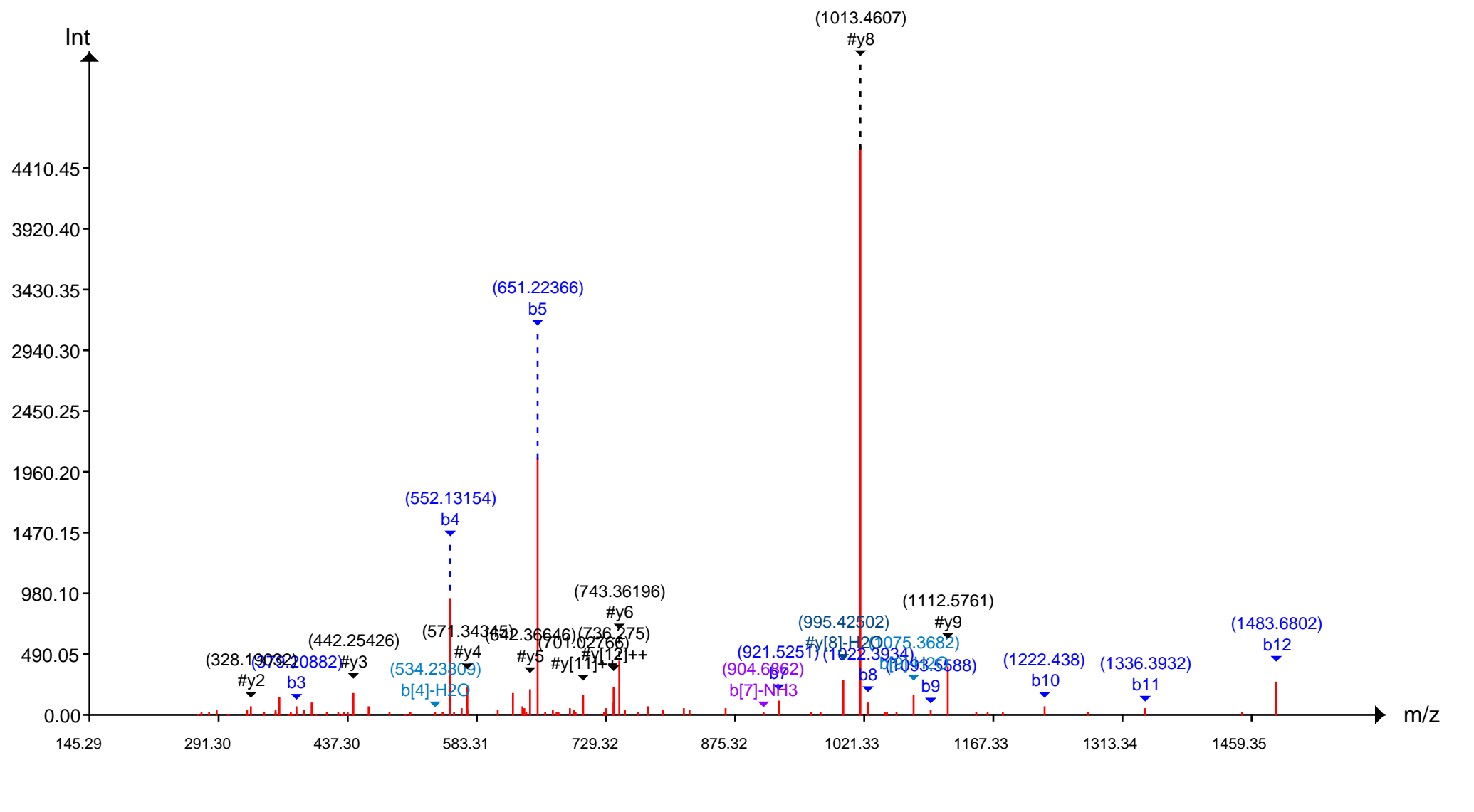
Charge 2+

Mass error 0.00171



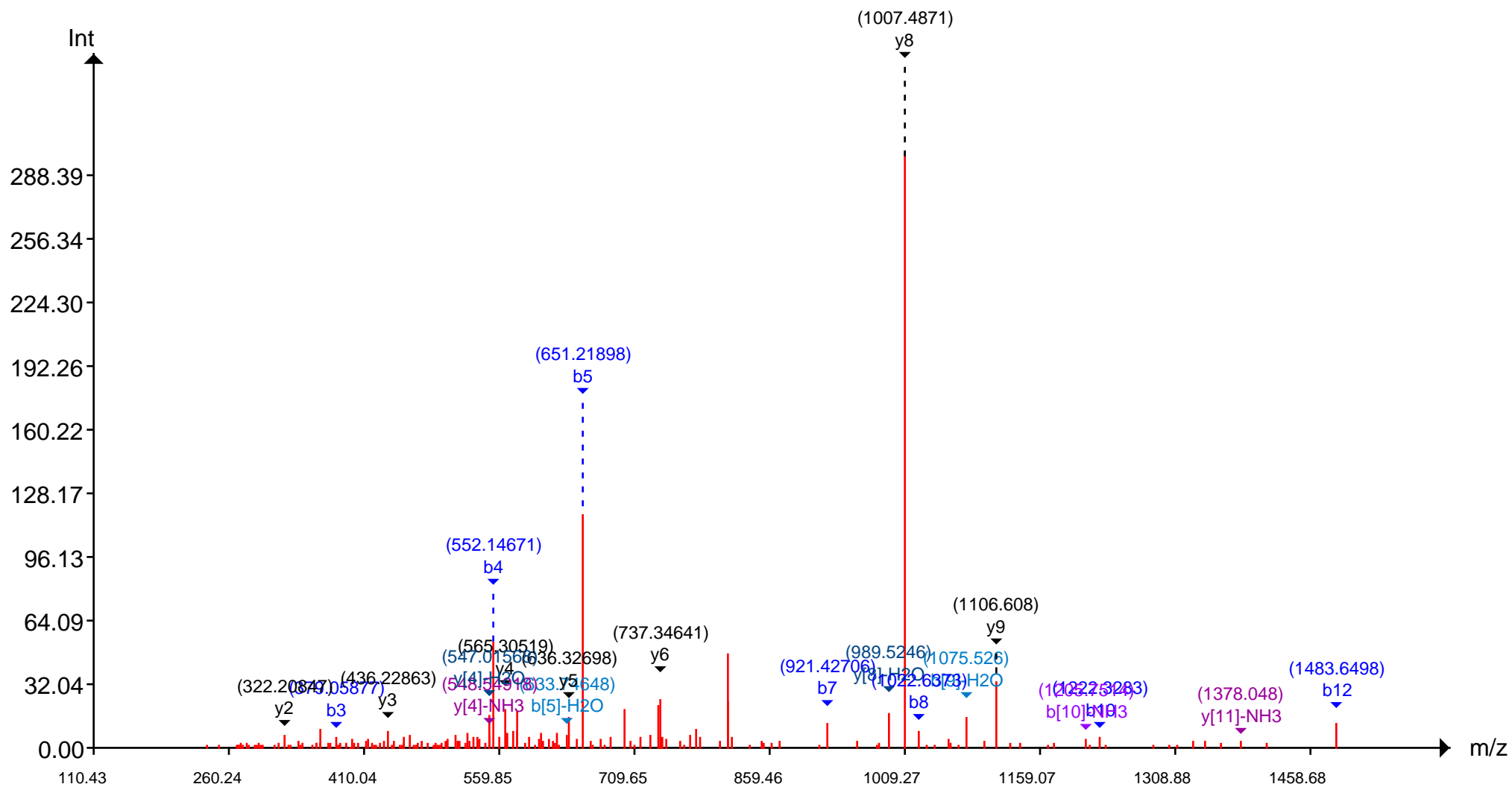
Info

Modified Sequence	AcD3-VDVPNDSSFQIVK<AcD3K*>NPR<C13>-COOH	Ion score	50.23
Protein	Q61176	Confidence level	99.00%
Precursor m/z	956.01433	Identity Threshold	47.81
Charge	2+	Mass error	0.002908



Info

Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH	Ion score	67.58
Protein	P17742	Confidence level	99.00%
Precursor m/z	832.45933	Identity Threshold	47.48
Charge	2+	Mass error	0.00181



Info

Modified Sequence AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH

Ion score 50.55

Protein P17742

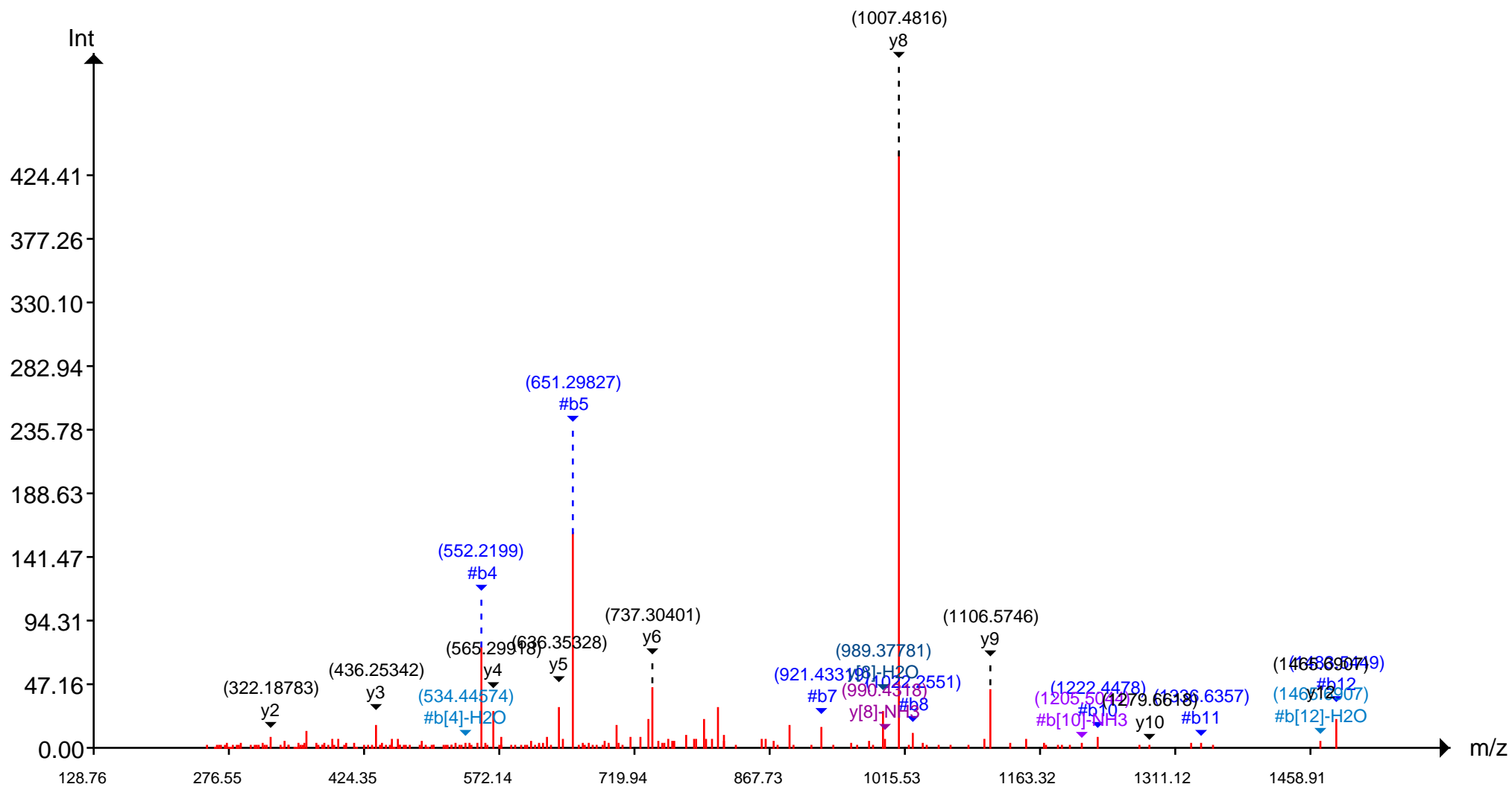
Confidence level 99.00%

Precursor m/z 829.44901

Identity Threshold 47.80

Charge 2+

Mass error 0.001312



Info

Modified Sequence AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH

Ion score 49.94

Protein P17742

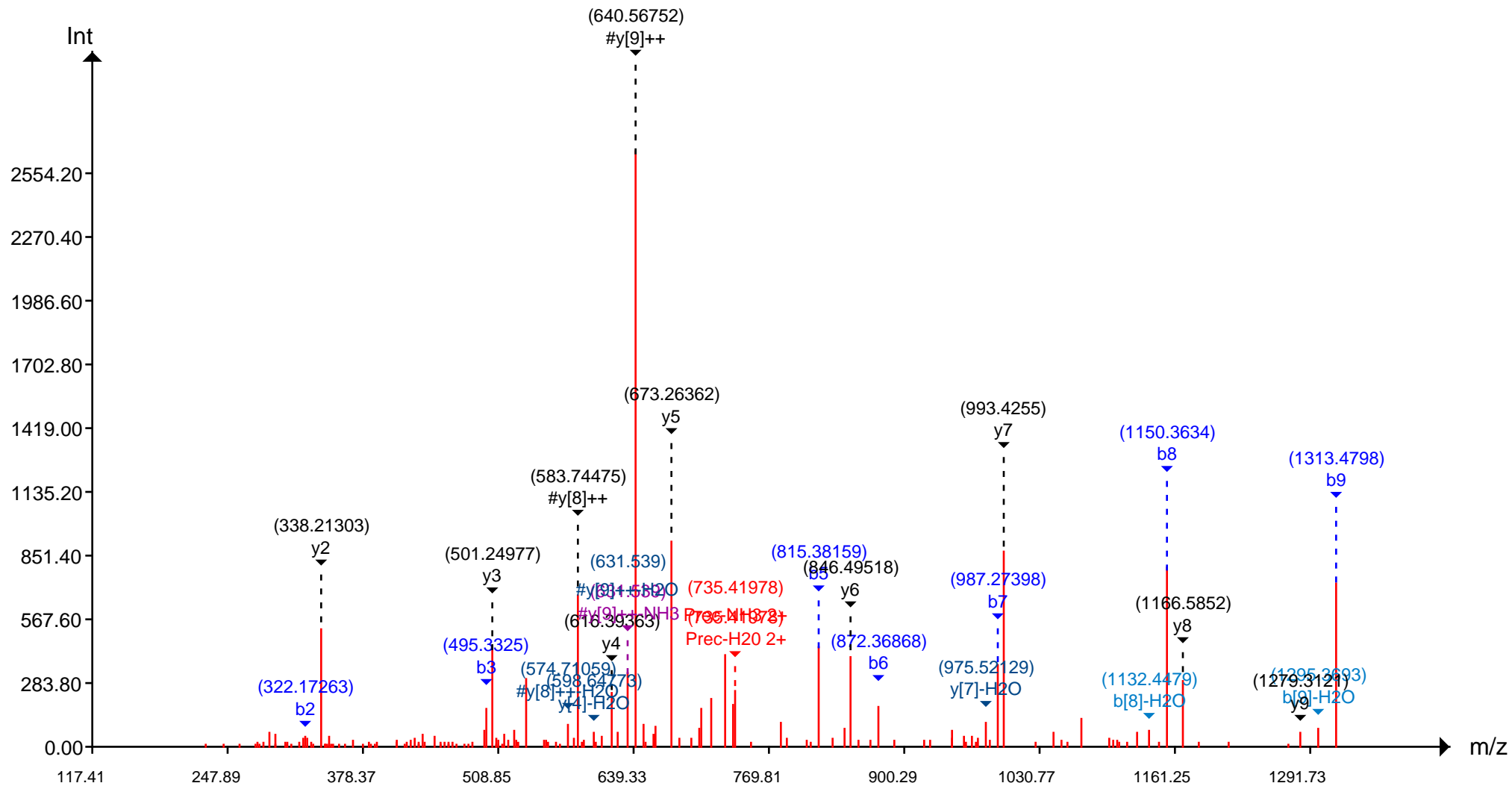
Confidence level 99.00%

Precursor m/z 829.44932

Identity Threshold 47.76

Charge 2+

Mass error 0.001932



Info

Modified Sequence AcD3-YLK<AcD3K*>M<Mox*>K<AcD3K*>GDYYR-COOH

Ion score 55.01

Protein P68510, P61982, O70456, P63101

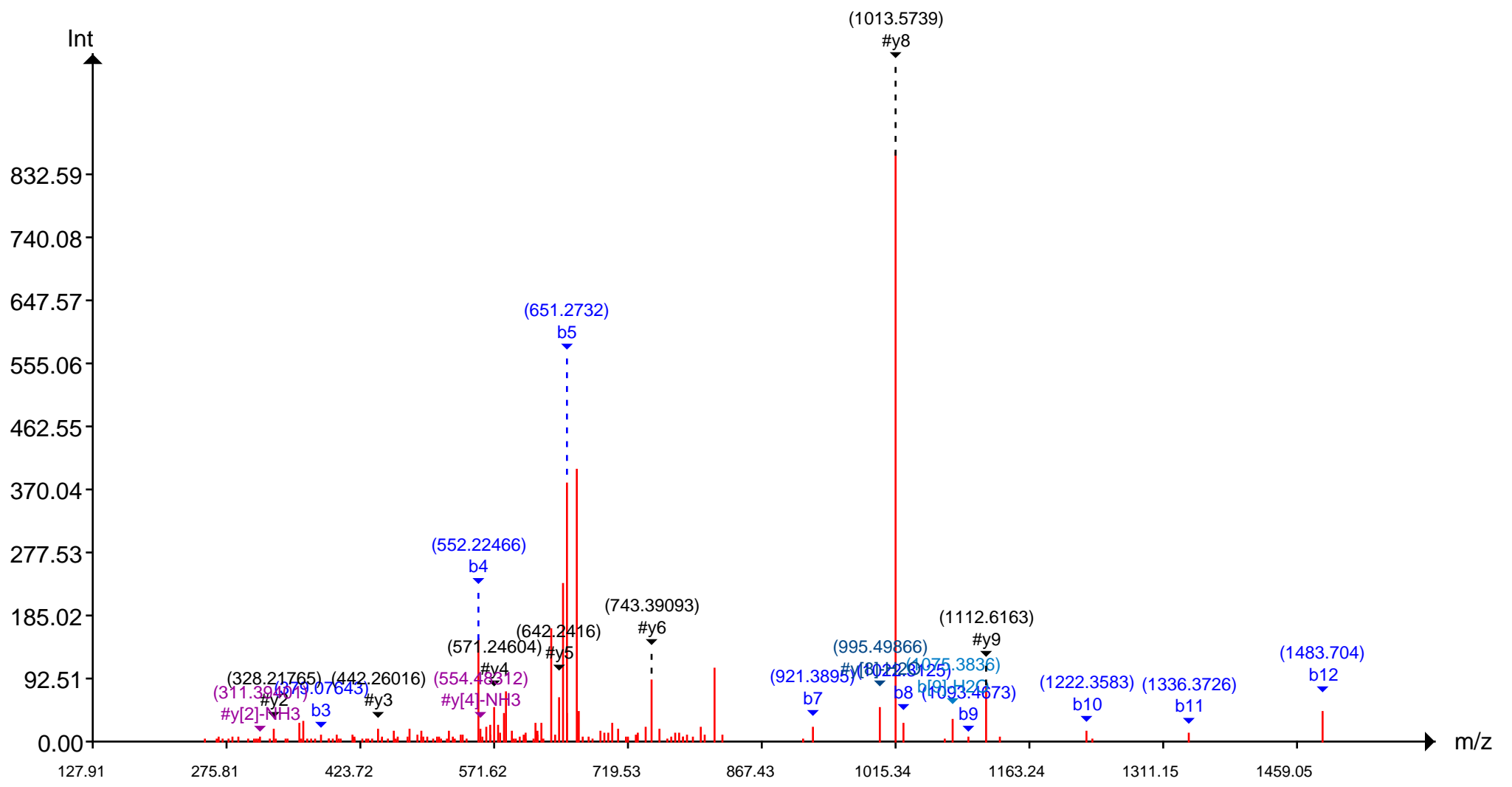
Confidence level 99.00%

Precursor m/z 744.38145

Identity Threshold 47.99

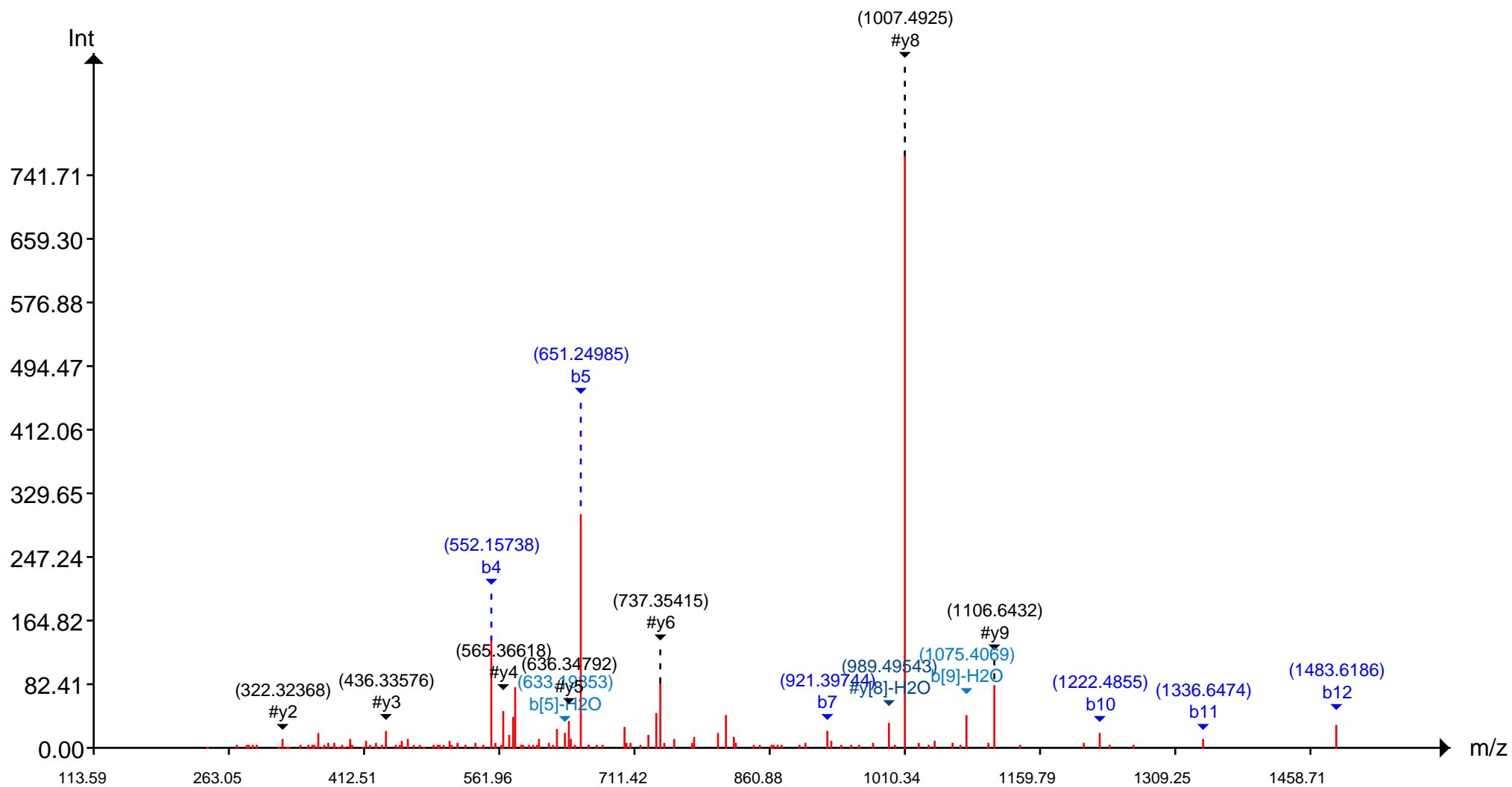
Charge 2+

Mass error 7.28E-4



Info

Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH	Ion score	50.64
Protein	P17742	Confidence level	99.00%
Precursor m/z	832.45936	Identity Threshold	47.48
Charge	2+	Mass error	0.00187



Info

Modified Sequence AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH

Ion score 53.06

Protein P17742

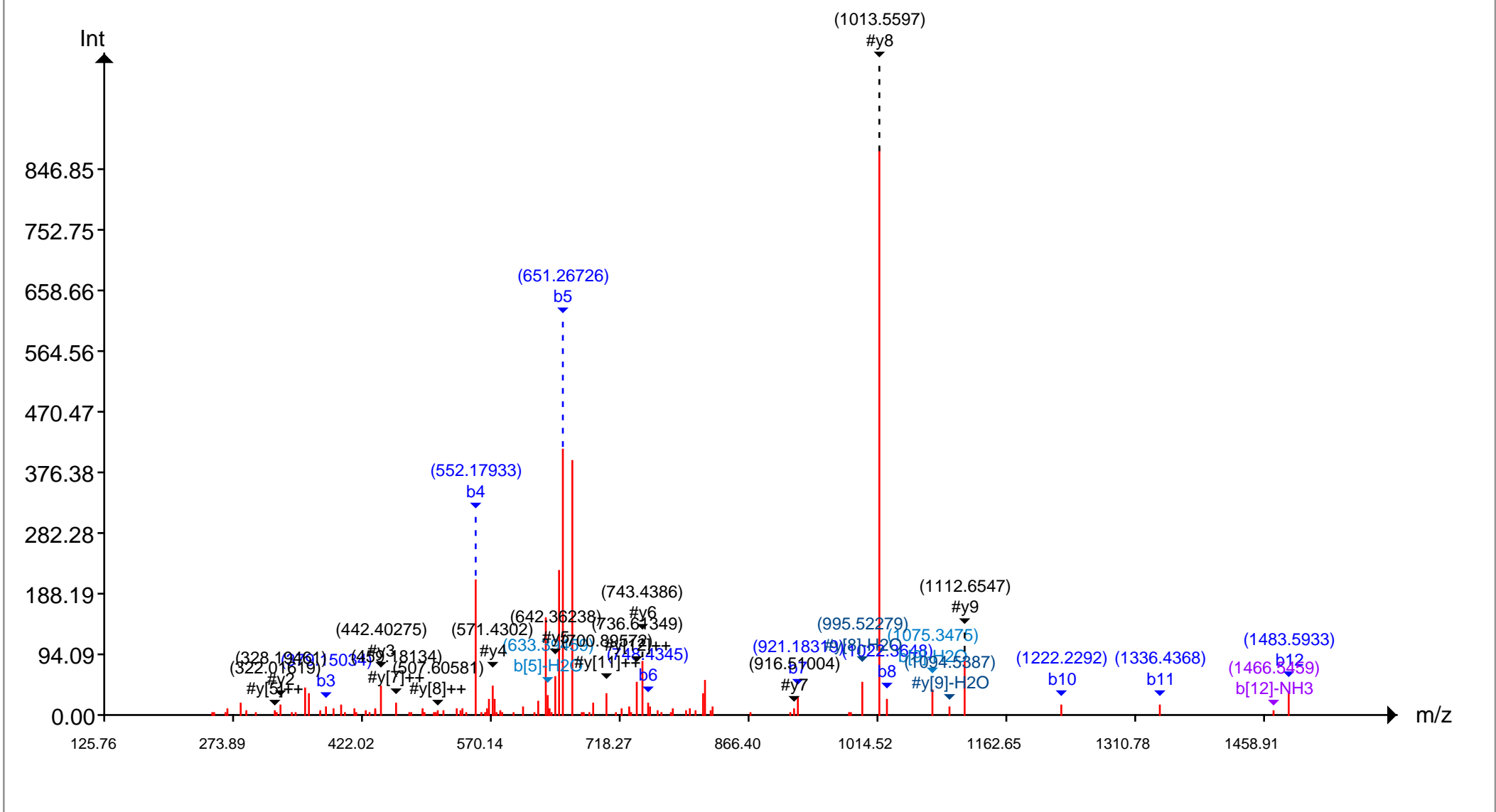
Confidence level 99.00%

Precursor m/z 829.44931

Identity Threshold 47.76

Charge 2+

Mass error 0.001912



Info

Modified Sequence AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH

Ion score 50.09

Protein P17742

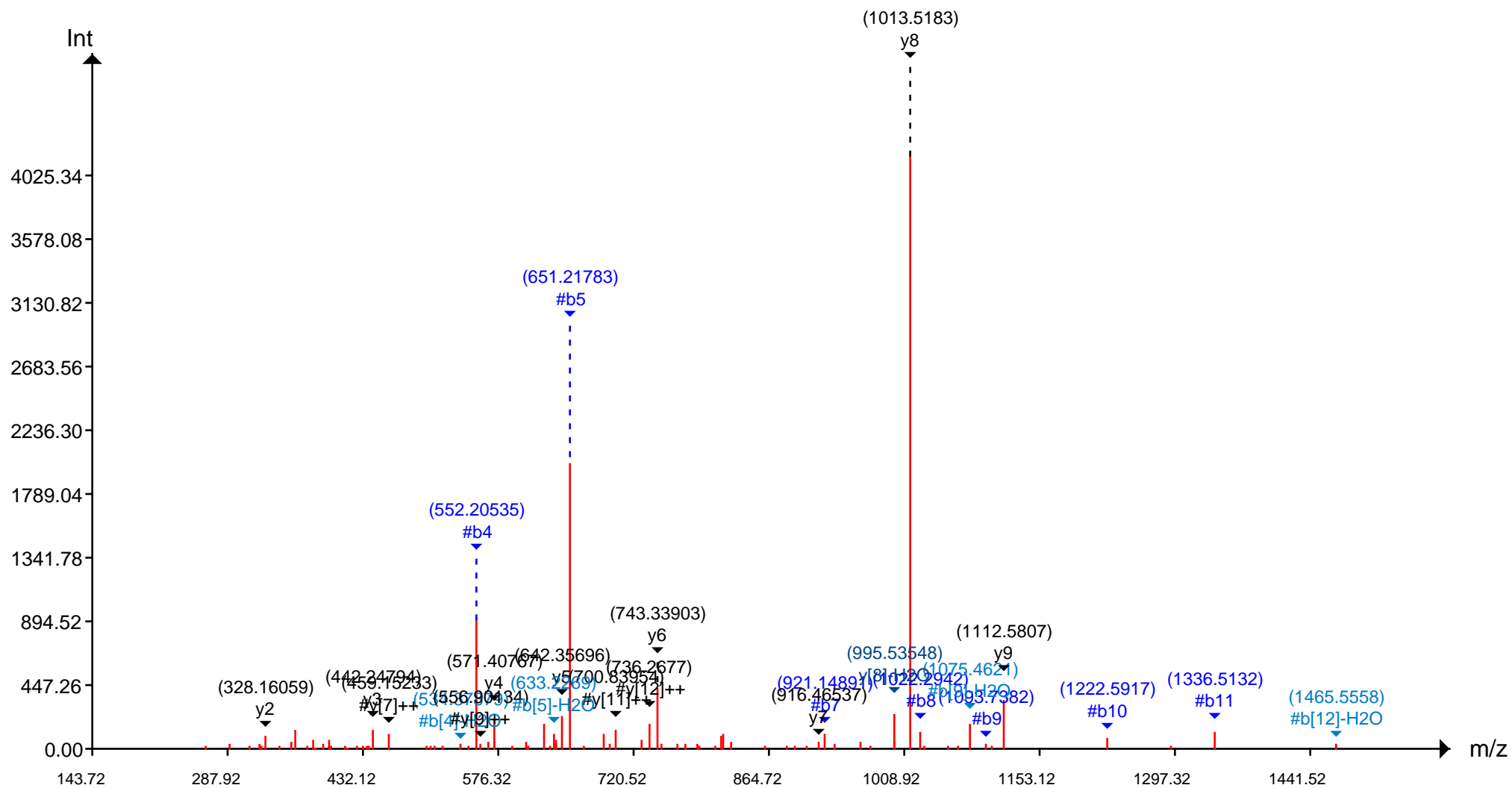
Confidence level 99.00%

Precursor m/z 832.45916

Identity Threshold 47.45

Charge 2+

Mass error 0.00147



Info

Modified Sequence AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH

Ion score 62.13

Protein P17742

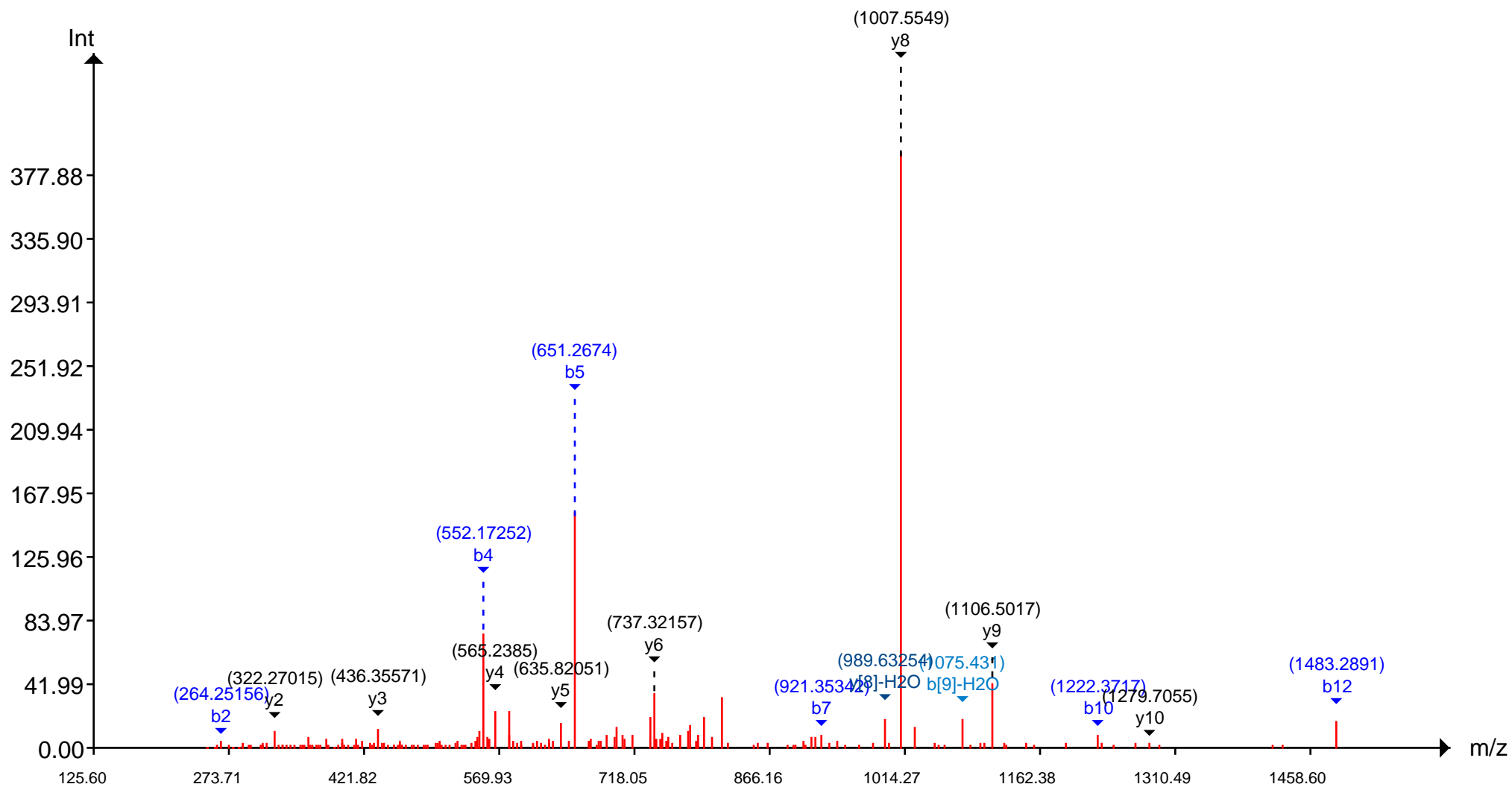
Confidence level 99.00%

Precursor m/z 832.45916

Identity Threshold 47.45

Charge 2+

Mass error 0.00147



Info

Modified Sequence AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH

Ion score 55.8

Protein P17742

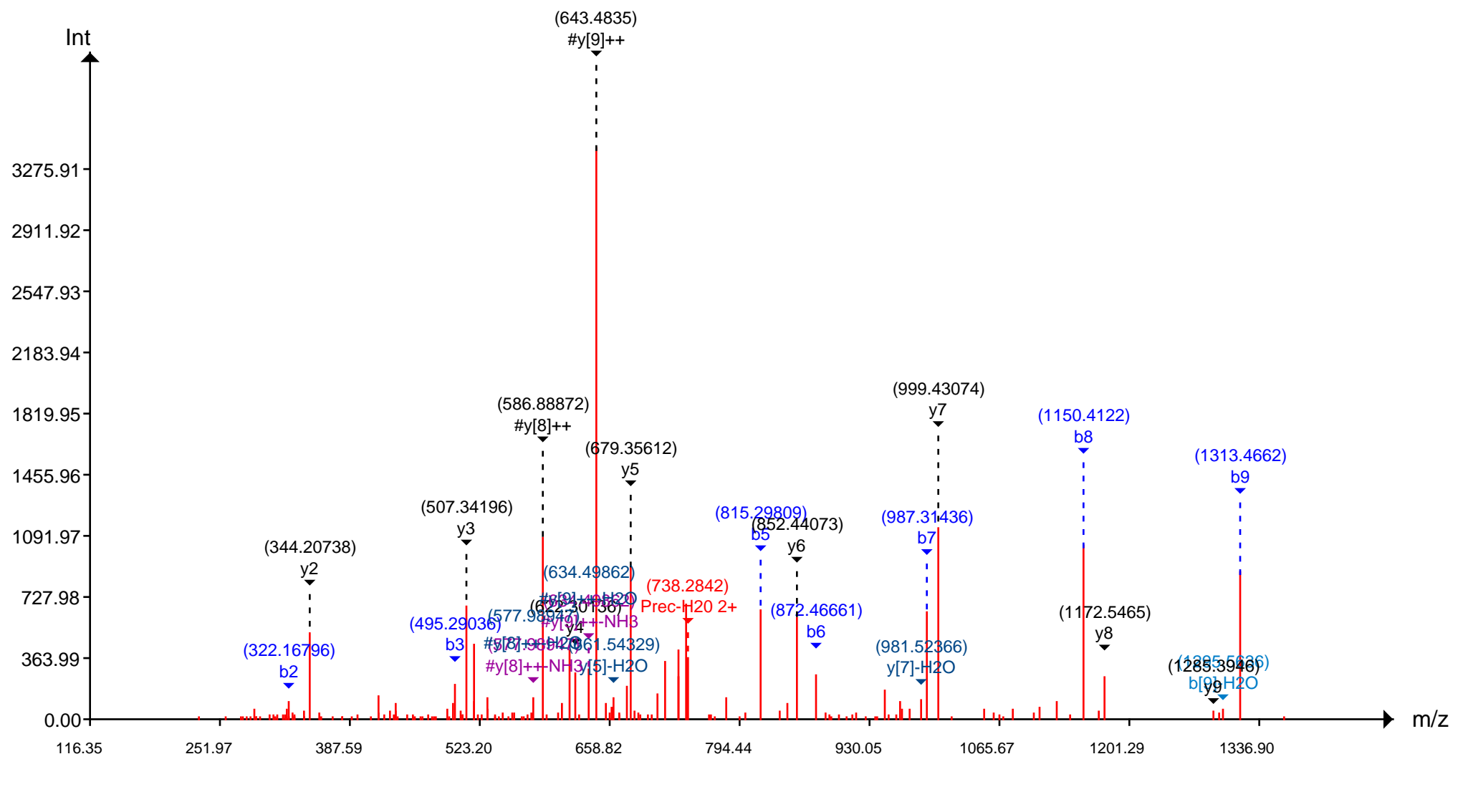
Confidence level 99.00%

Precursor m/z 829.44854

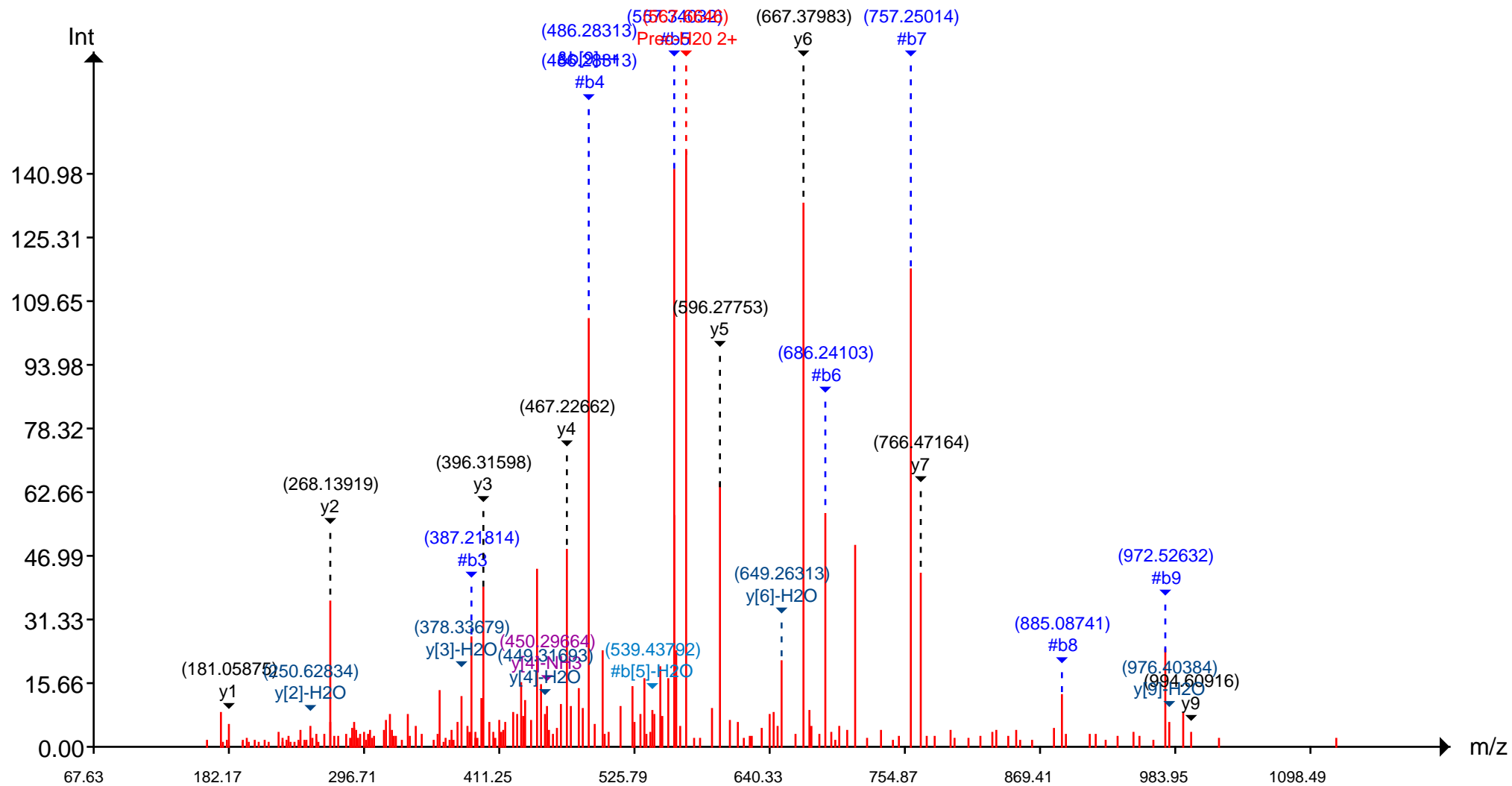
Identity Threshold 47.89

Charge 2+

Mass error 3.72E-4

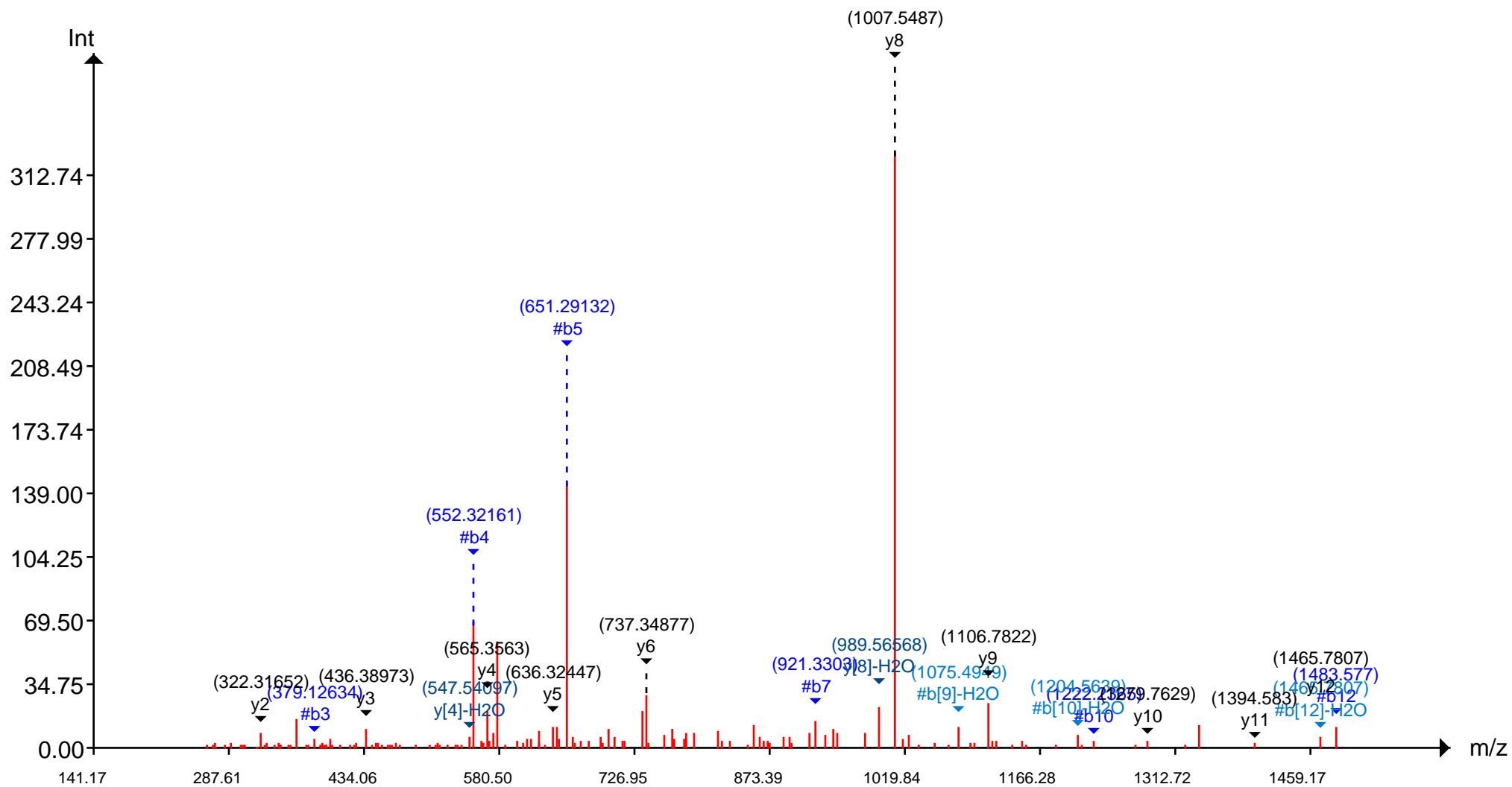


Info			
Modified Sequence	AcD3-YLK<AcD3K*>M<Mox*>K<AcD3K*>GDYYR<C13>-COOH	Ion score	53.19
Protein	P68510, P61982, O70456, P63101	Confidence level	99.00%
Precursor m/z	747.3917	Identity Threshold	48.22
Charge	2+	Mass error	0.001087



Info

Modified Sequence	AcD3-LDLVAEAQSR<C13>-COOH	Ion score	52.46
Protein	Q3U1Z5	Confidence level	99.00%
Precursor m/z	576.82302	Identity Threshold	47.35
Charge	2+	Mass error	-6.07E-4



Info

Modified Sequence AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH

Ion score 62.09

Protein P17742

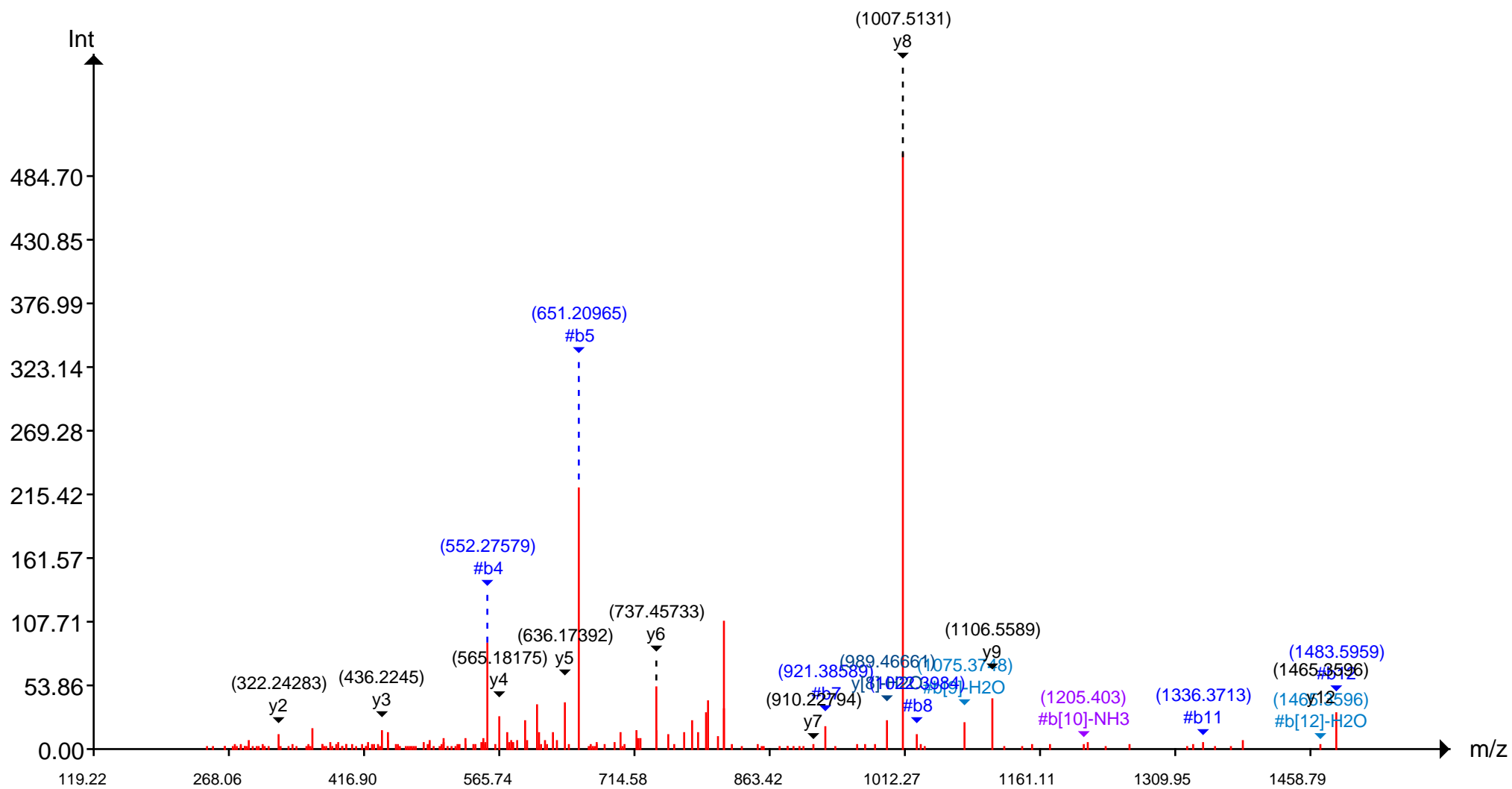
Confidence level 99.00%

Precursor m/z 829.44866

Identity Threshold 47.89

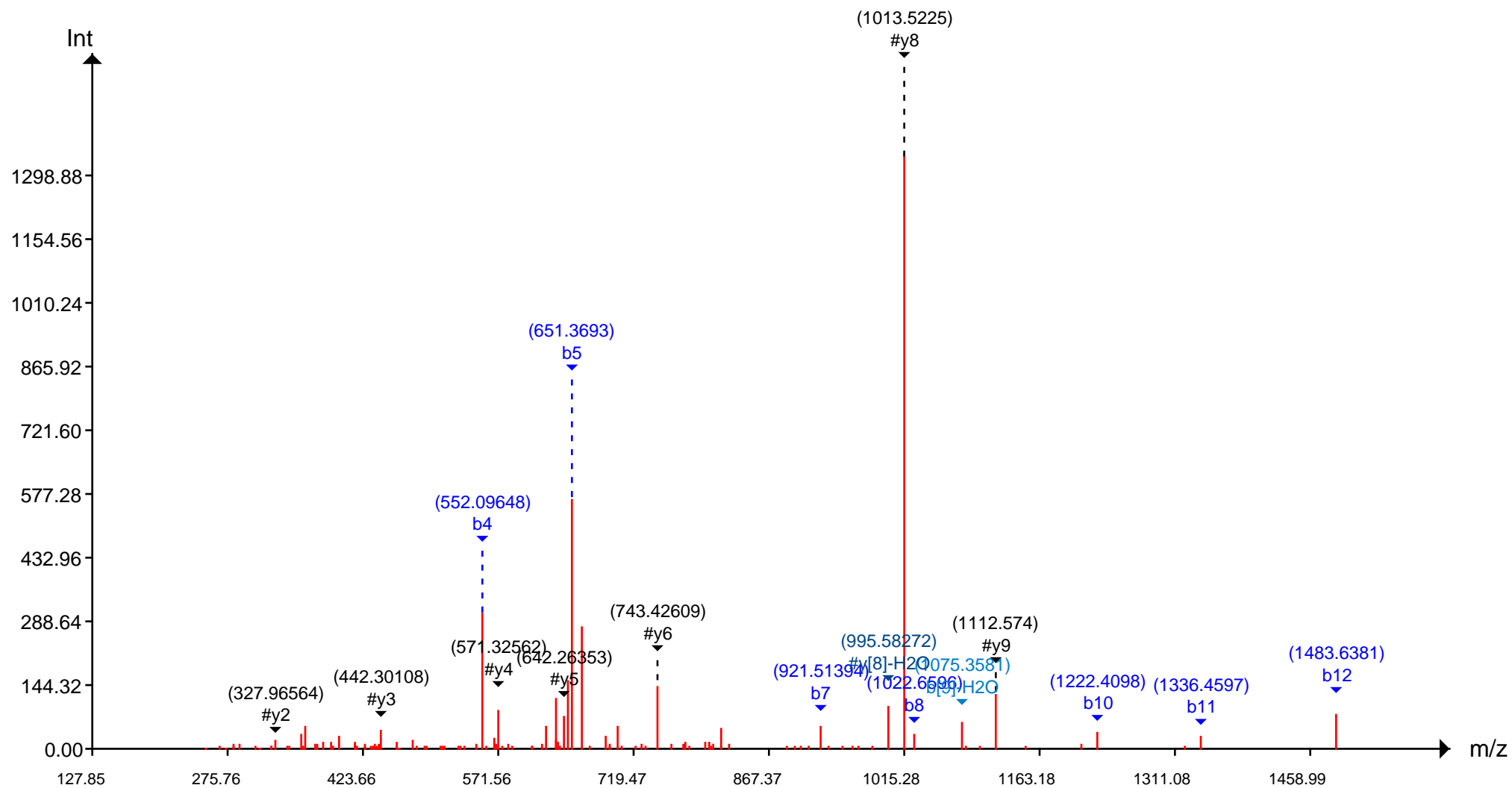
Charge 2+

Mass error 6.12E-4



Info

Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH	Ion score	55.31
Protein	P17742	Confidence level	99.00%
Precursor m/z	829.44864	Identity Threshold	47.89
Charge	2+	Mass error	5.72E-4



Info

Modified Sequence AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH

Ion score 48.78

Protein P17742

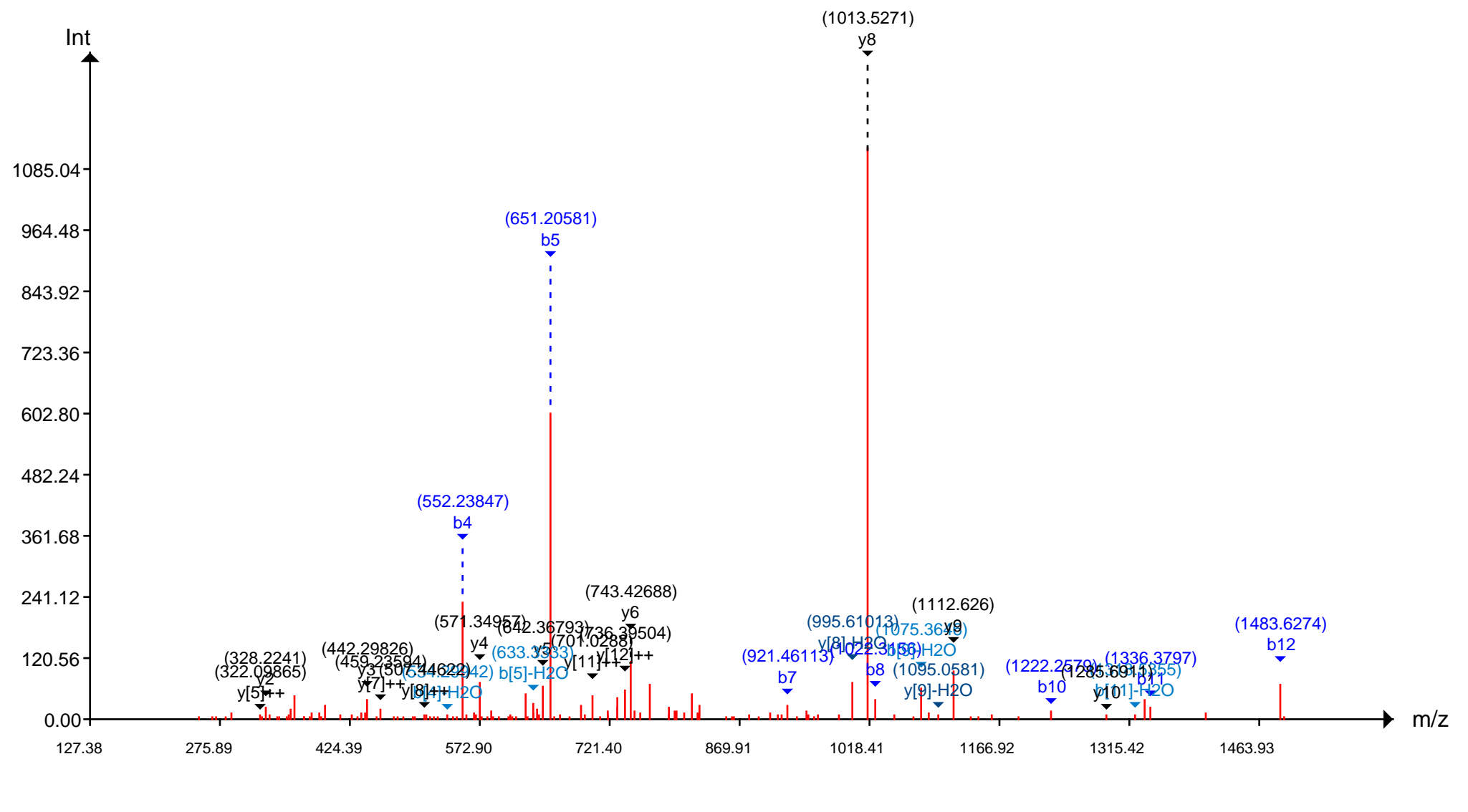
Confidence level 99.00%

Precursor m/z 832.45919

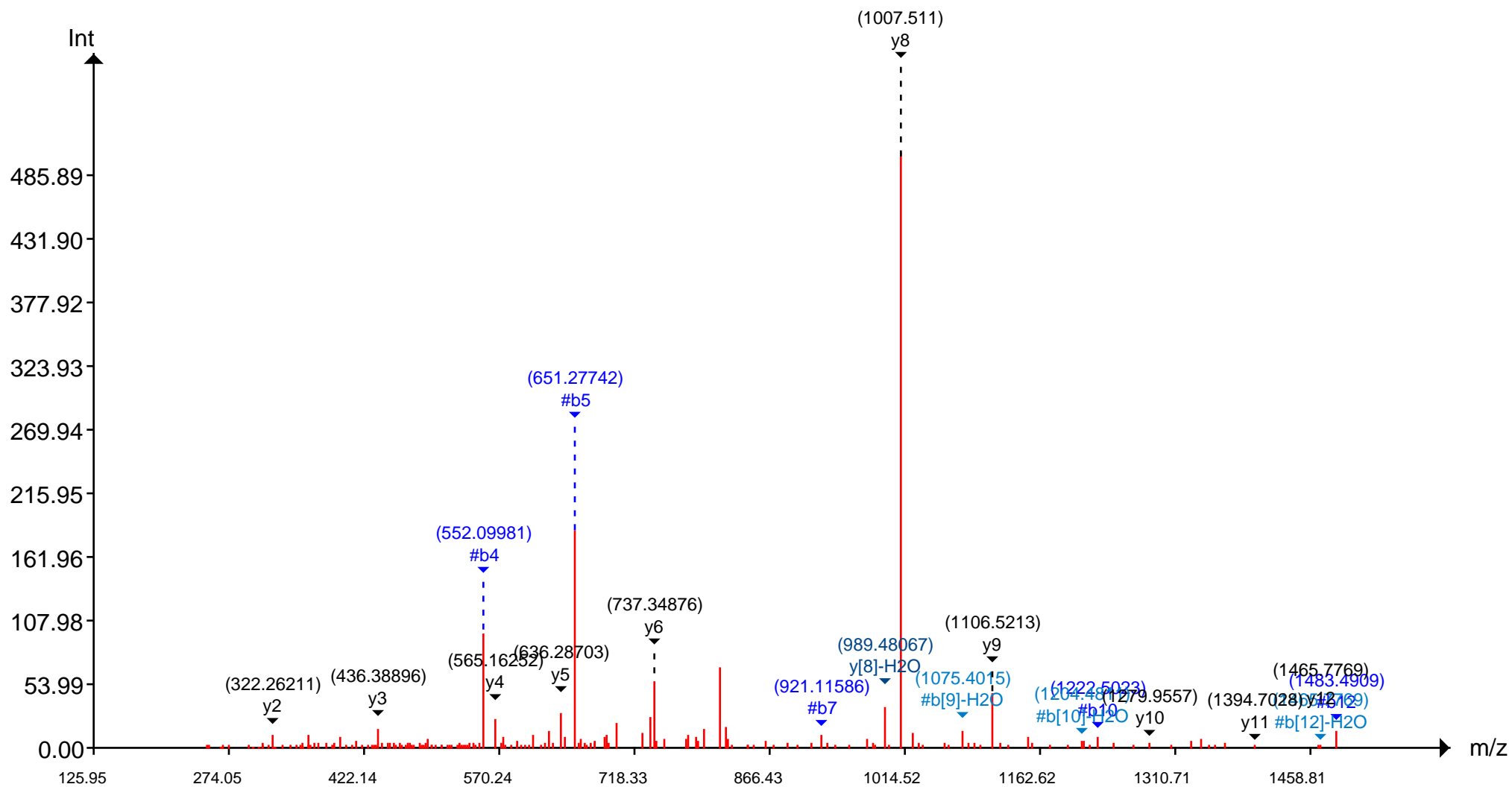
Identity Threshold 47.46

Charge 2+

Mass error 0.00153



Info			
Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH	Ion score	49.11
Protein	P17742	Confidence level	99.00%
Precursor m/z	832.45927	Identity Threshold	47.47
Charge	2+	Mass error	0.00169



Info

Modified Sequence AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR-COOH

Ion score 66.38

Protein P17742

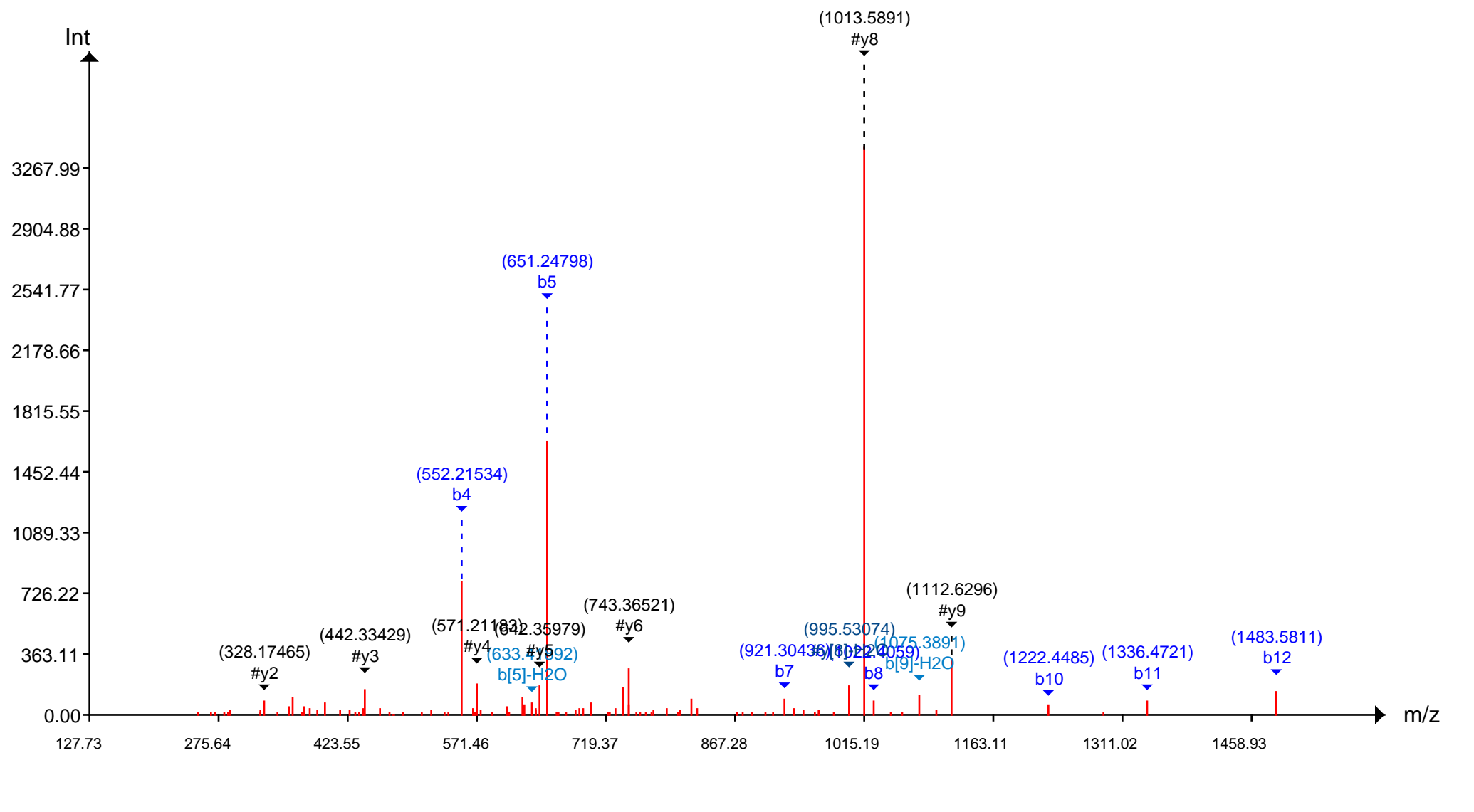
Confidence level 99.00%

Precursor m/z 829.44883

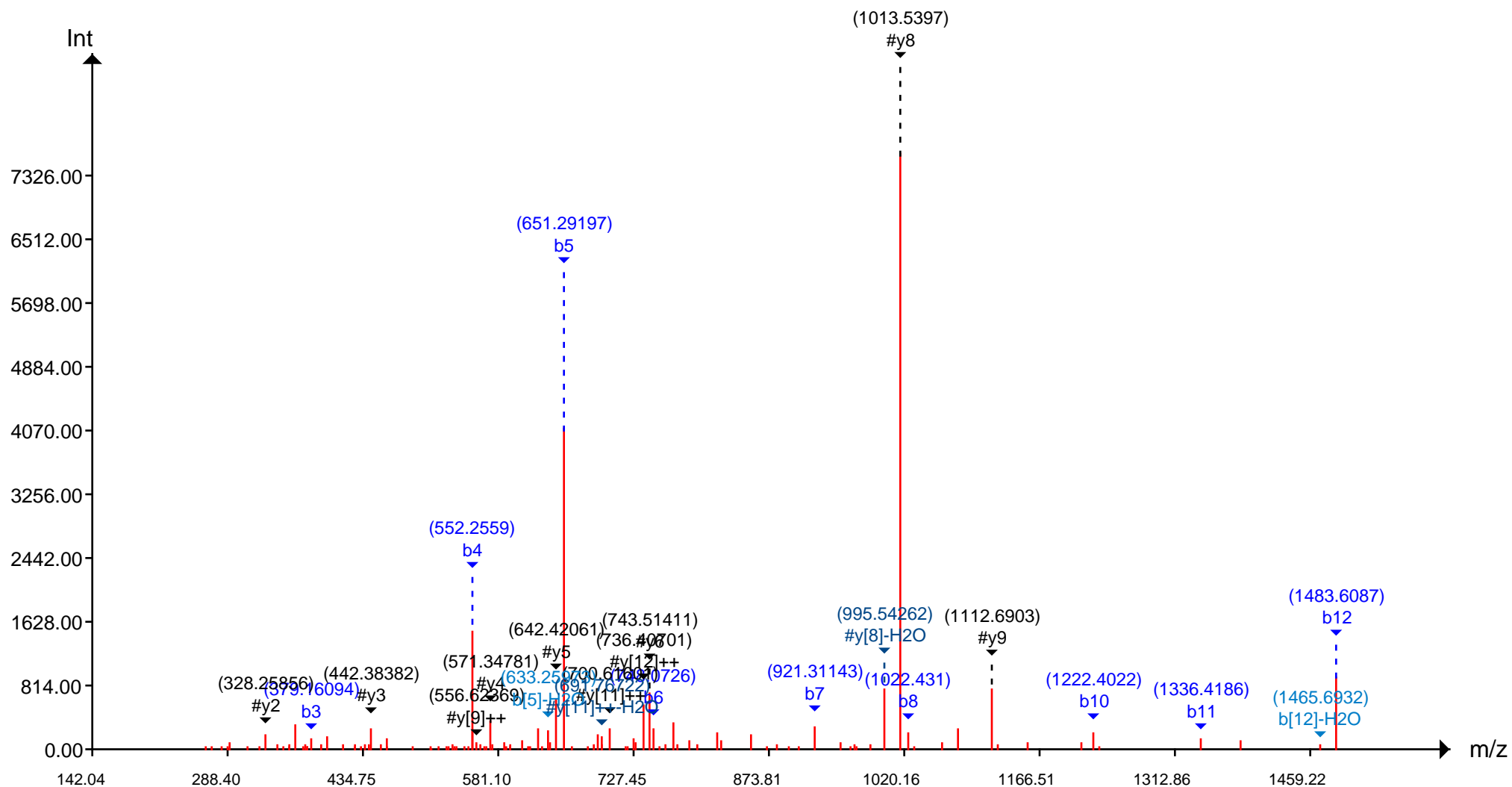
Identity Threshold 47.83

Charge 2+

Mass error 9.52E-4

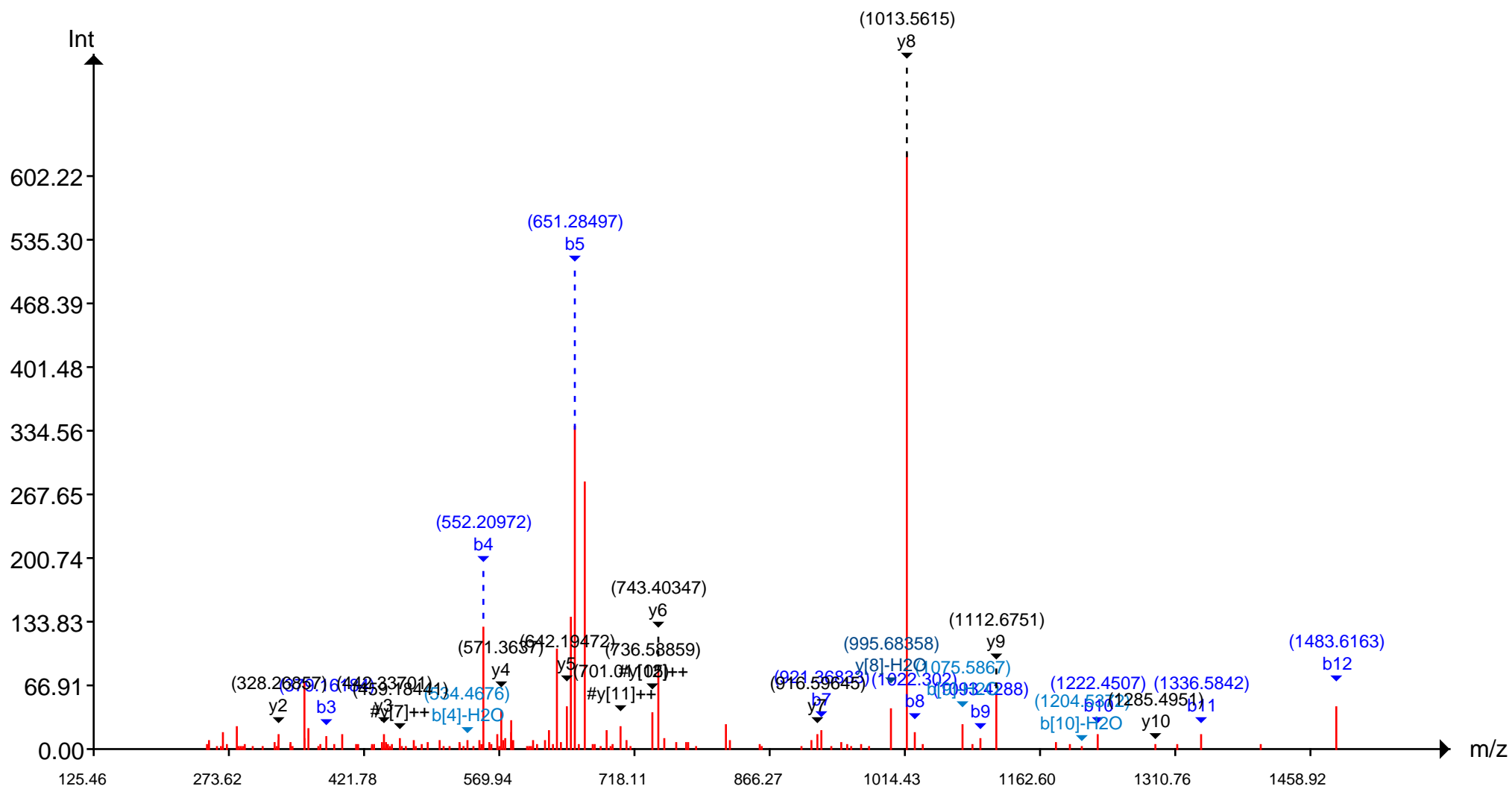


Info			
Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH	Ion score	53.54
Protein	P17742	Confidence level	99.00%
Precursor m/z	832.45924	Identity Threshold	47.46
Charge	2+	Mass error	0.00163



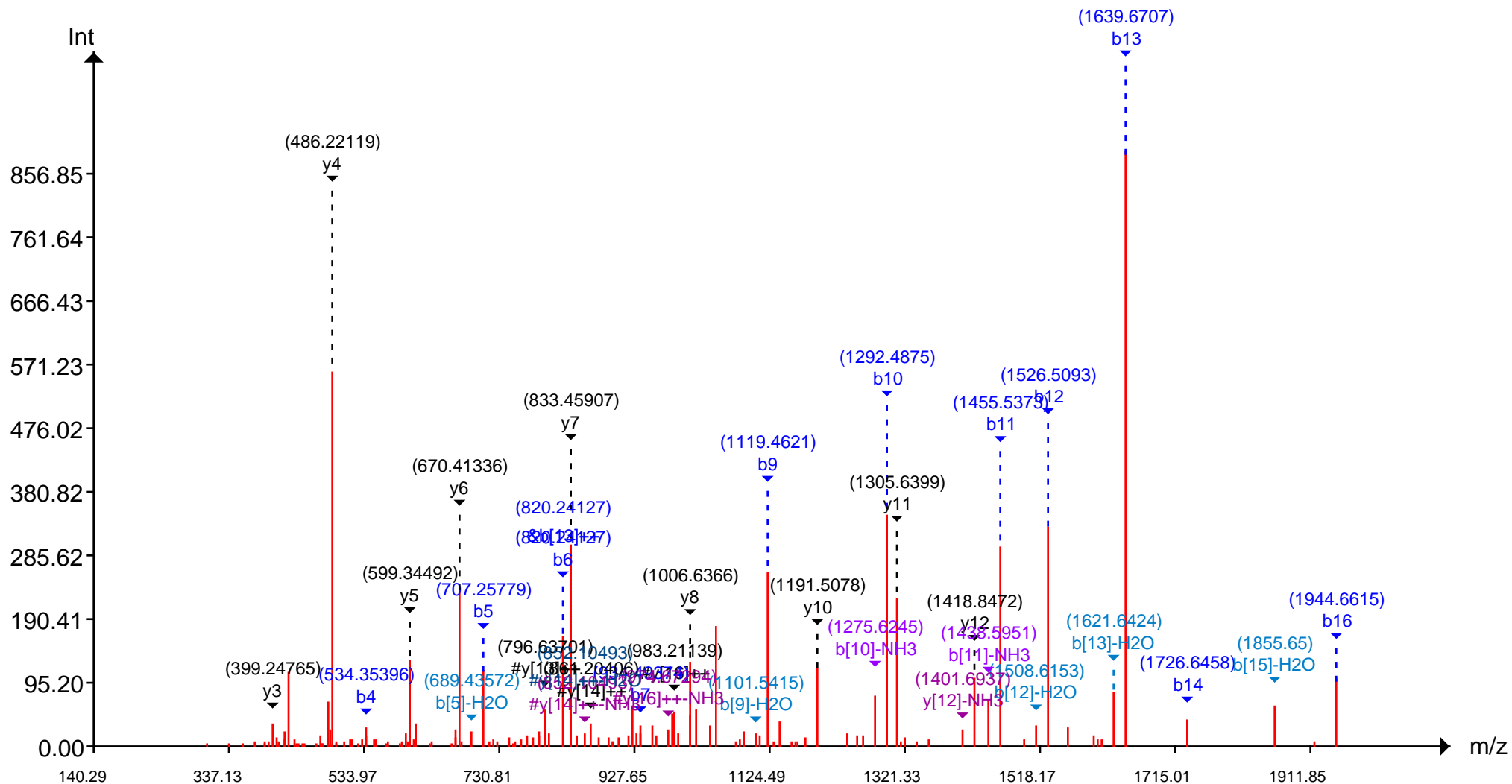
Info

Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH	Ion score	51.88
Protein	P17742	Confidence level	99.00%
Precursor m/z	832.45982	Identity Threshold	47.49
Charge	2+	Mass error	0.00279



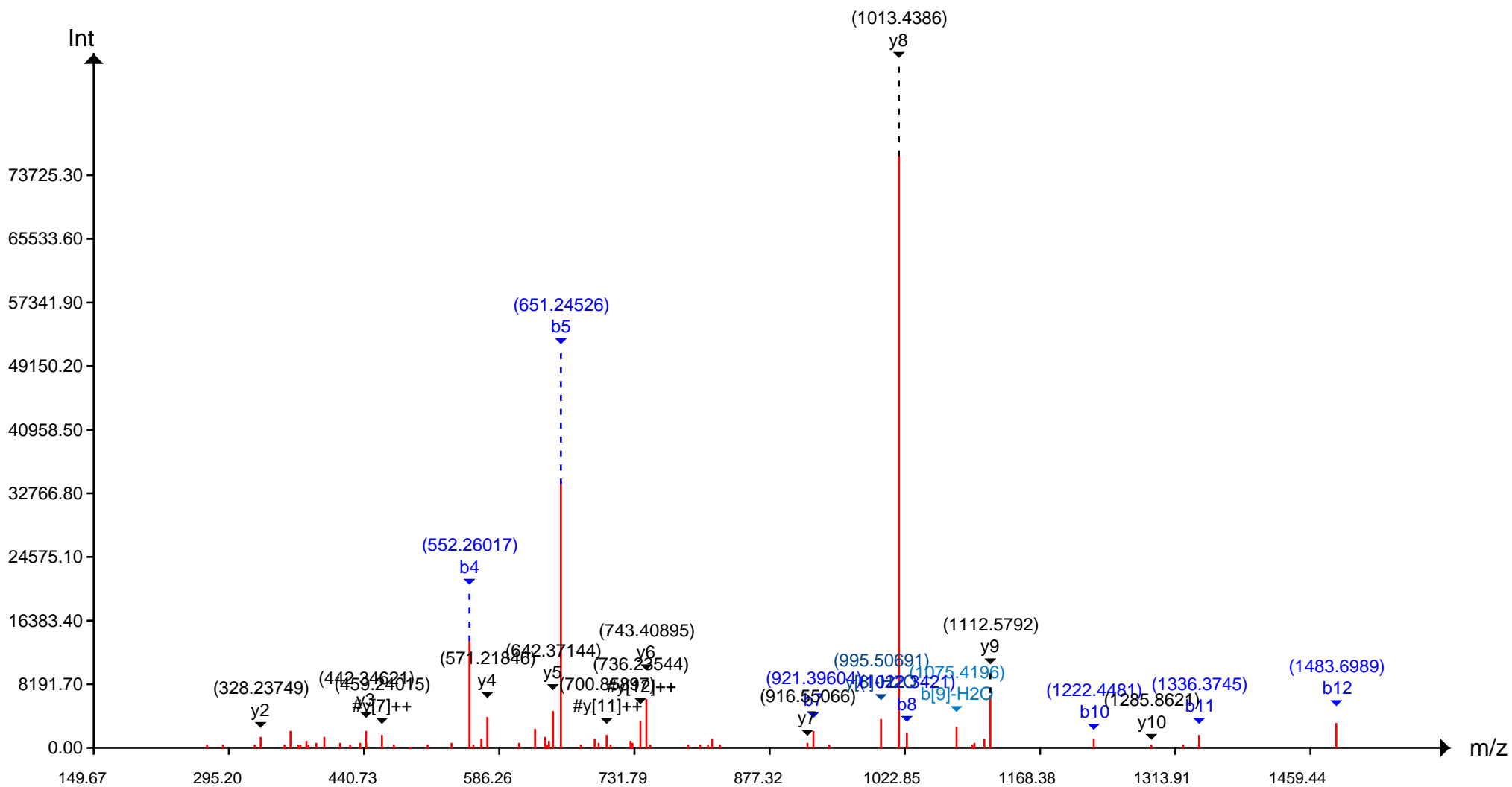
Info

Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH	Ion score	49.73
Protein	P17742	Confidence level	99.00%
Precursor m/z	832.4599	Identity Threshold	47.48
Charge	2+	Mass error	0.00295



Info

Modified Sequence	AcD3-DDEEK<AcD3K*>LNNAK<AcD3K*>YAISM<Mox*>AR<C13>-COOH	Ion score	83.8
Protein	Q61233	Confidence level	99.00%
Precursor m/z	1063.0315	Identity Threshold	47.73
Charge	2+	Mass error	0.003419



Info

Modified Sequence AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH

Ion score 75.42

Protein P17742

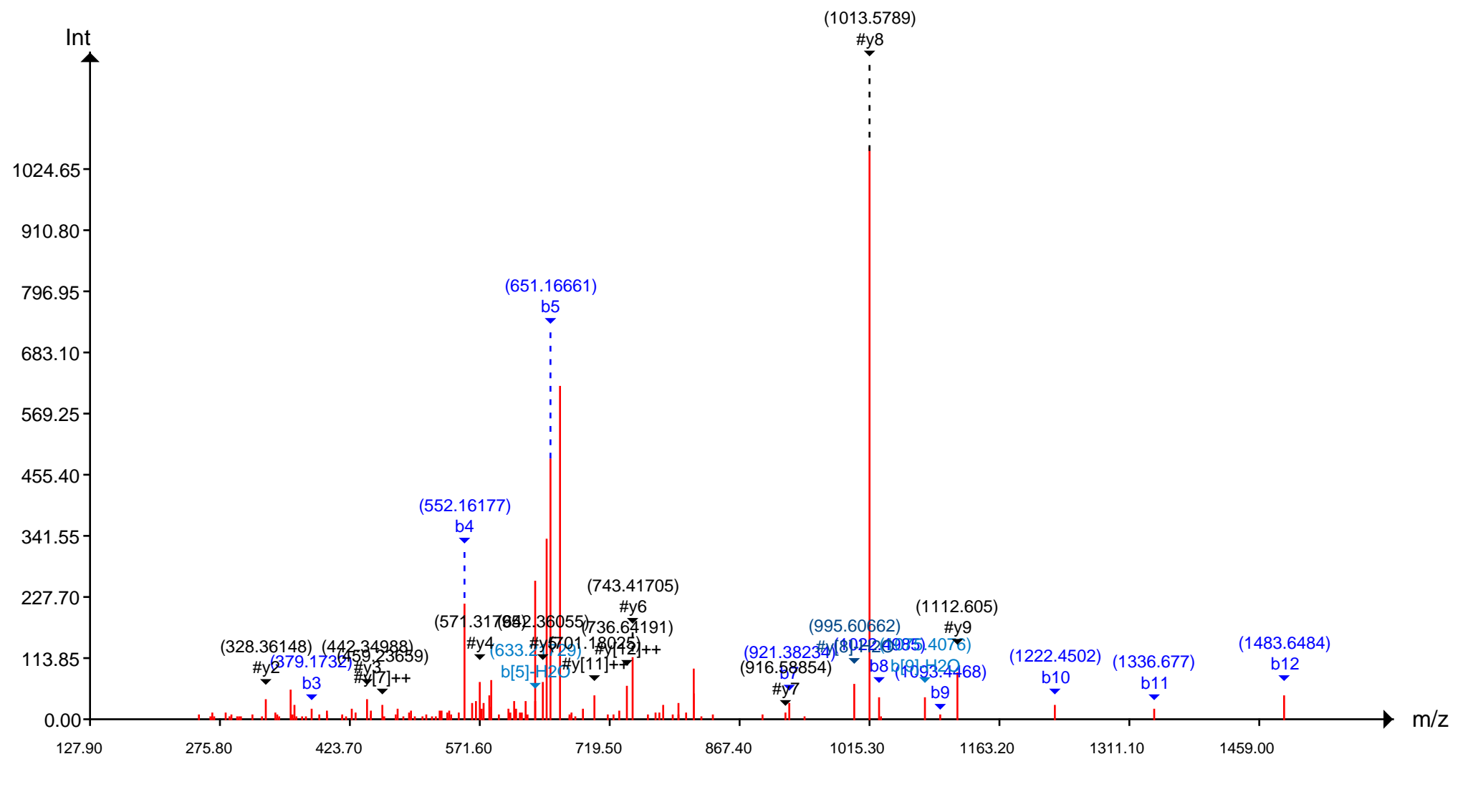
Confidence level 99.00%

Precursor m/z 832.45977

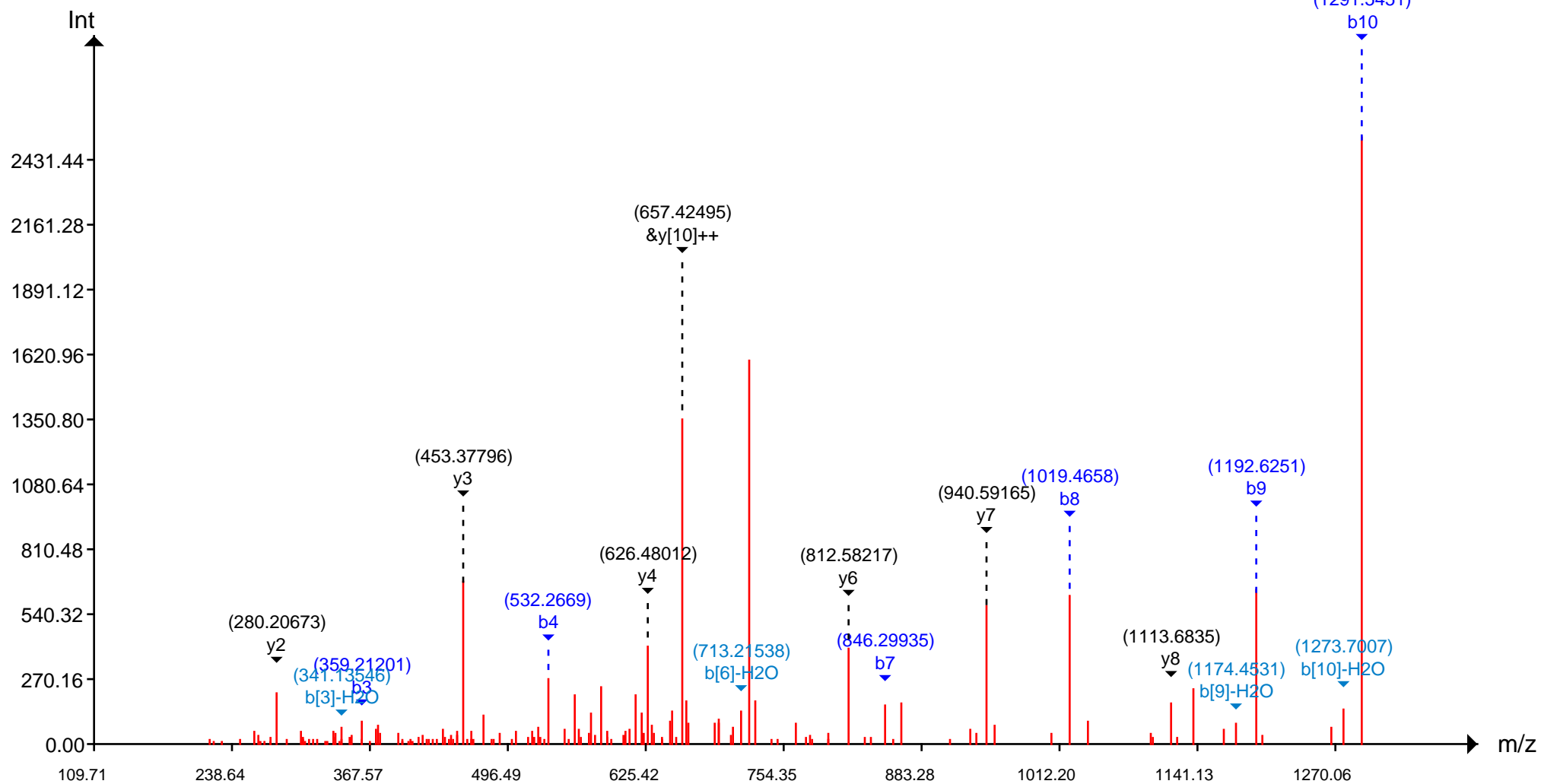
Identity Threshold 47.51

Charge 2+

Mass error 0.00269

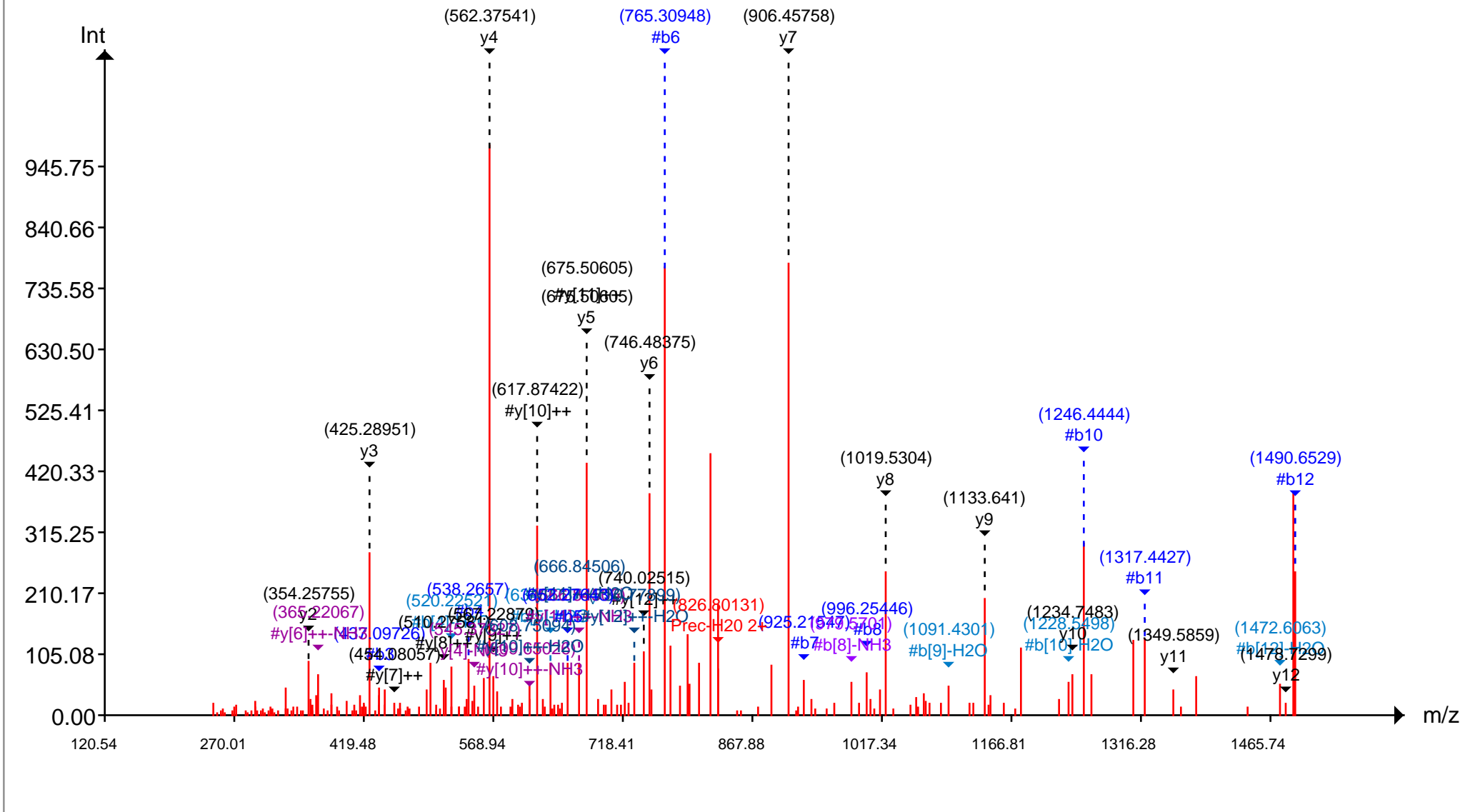


Info			
Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH	Ion score	54.32
Protein	P17742	Confidence level	99.00%
Precursor m/z	832.45991	Identity Threshold	47.48
Charge	2+	Mass error	0.00297



Info

Modified Sequence	AcD3-ISLK<AcD3K*>QADK<AcD3K*>K<AcD3K*>VR<C13>-COOH	Ion score	42.77
Protein	Q9DCL9	Confidence level	99.00%
Precursor m/z	736.47049	Identity Threshold	38.20
Charge	2+	Mass error	9.52E-4



Info

Modified Sequence AcD3-FEDTNLC<Cmm*>AIHAK<AcD3K*>R<C13>-COOH

Ion score 75.98

Protein P68433, P84228, P84244

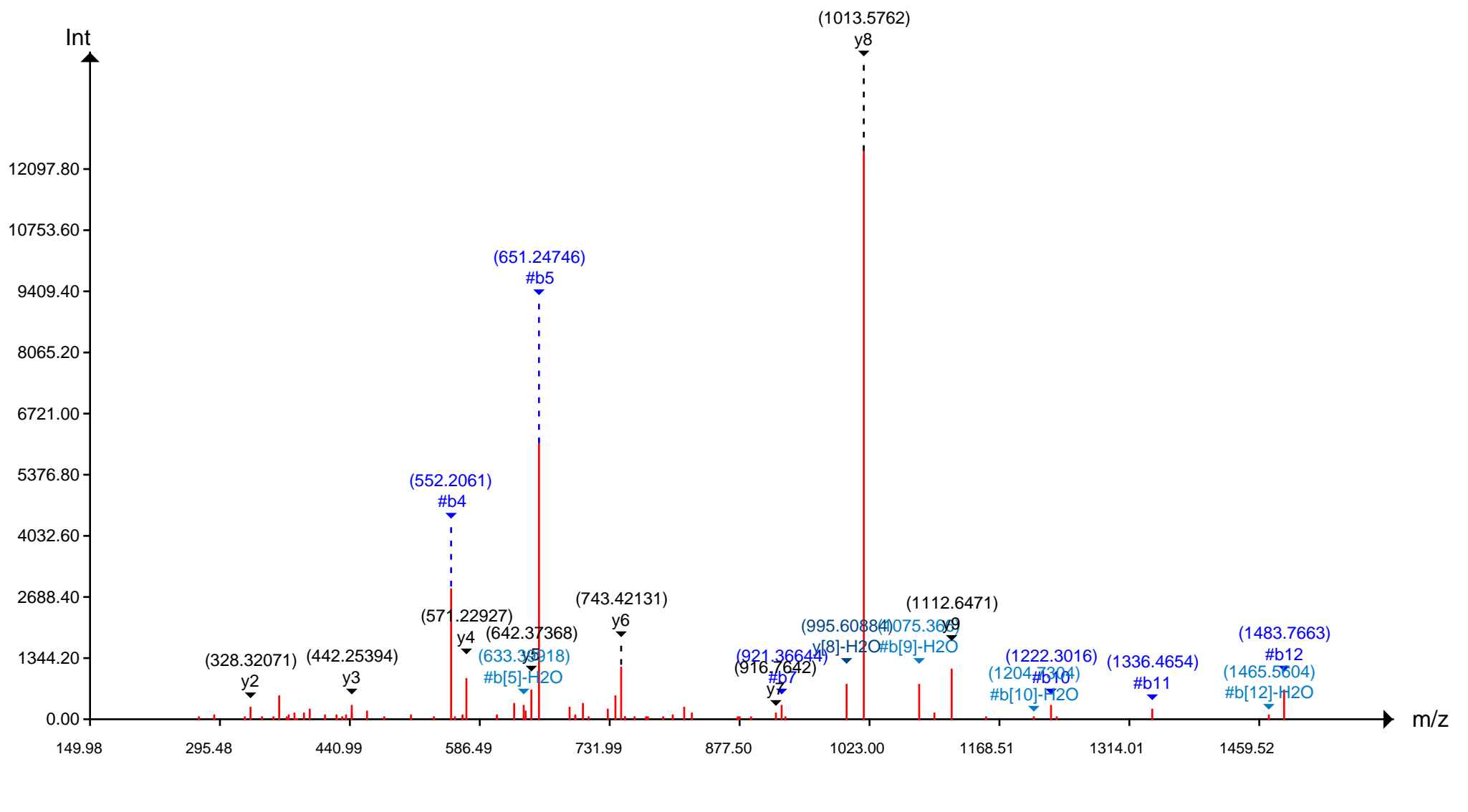
Confidence level 99.00%

Precursor m/z 835.93109

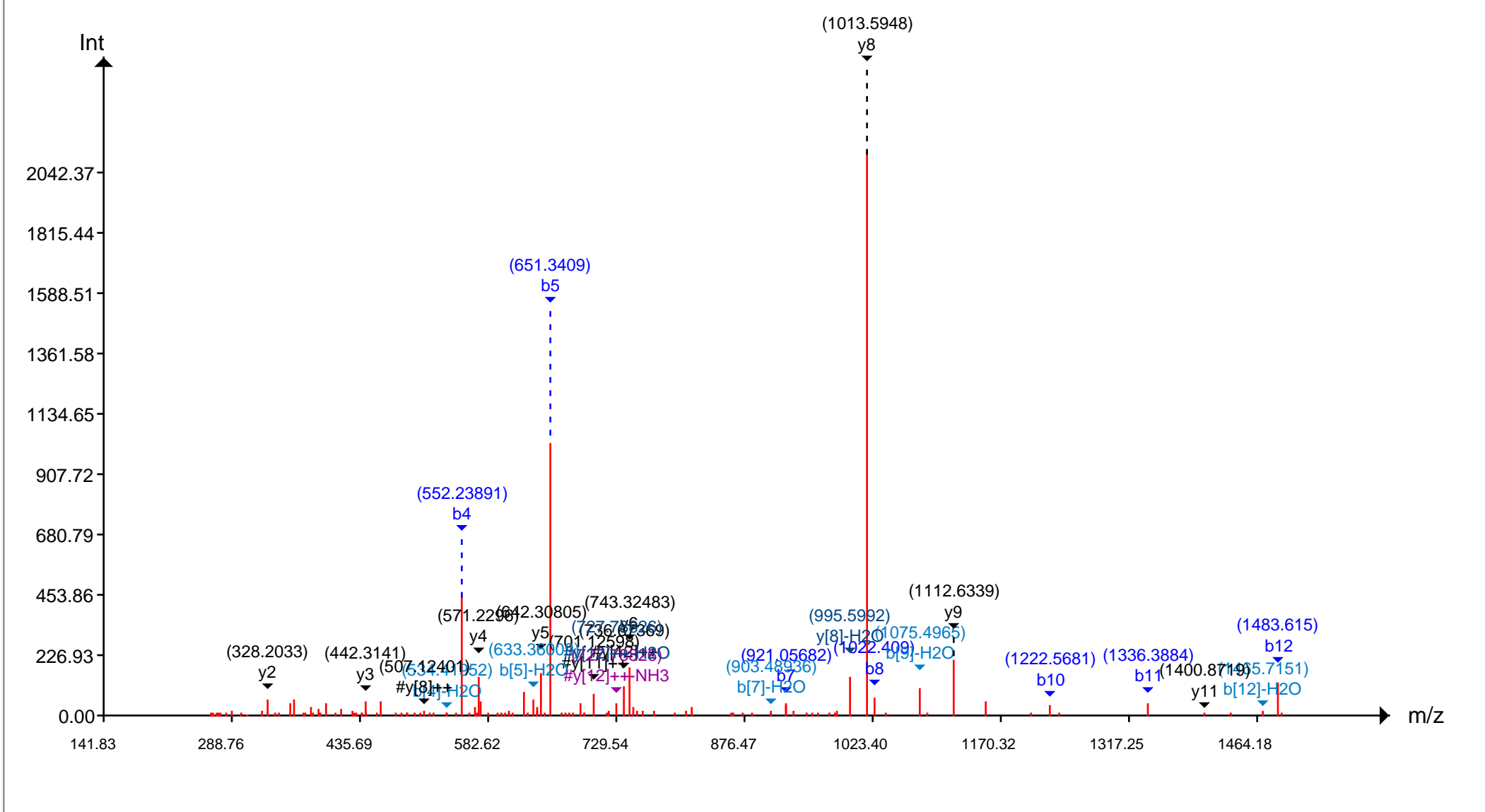
Identity Threshold 48.14

Charge 2+

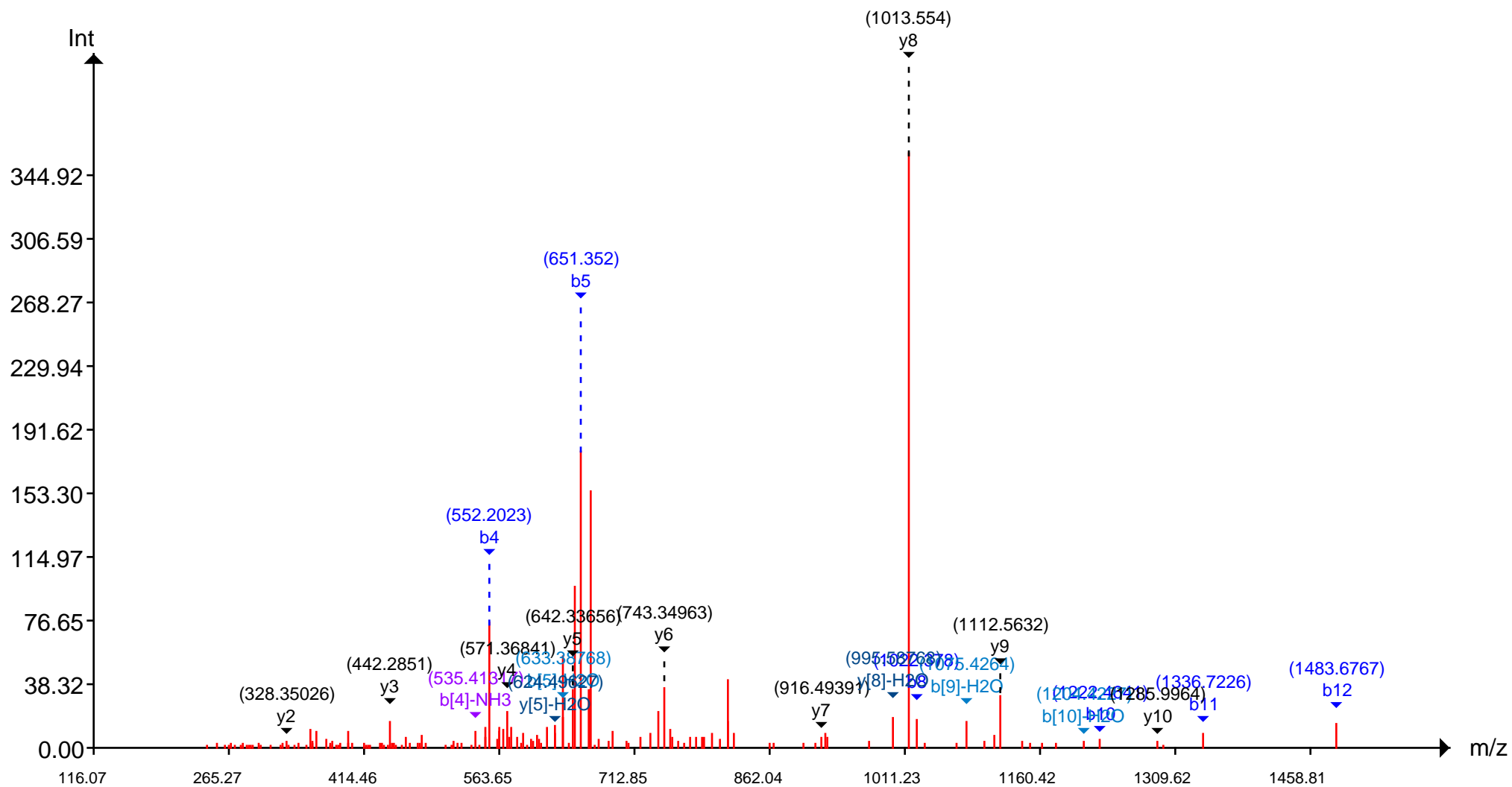
Mass error 0.001604



Info			
Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH	Ion score	68.46
Protein	P17742	Confidence level	99.00%
Precursor m/z	832.45946	Identity Threshold	47.48
Charge	2+	Mass error	0.00207

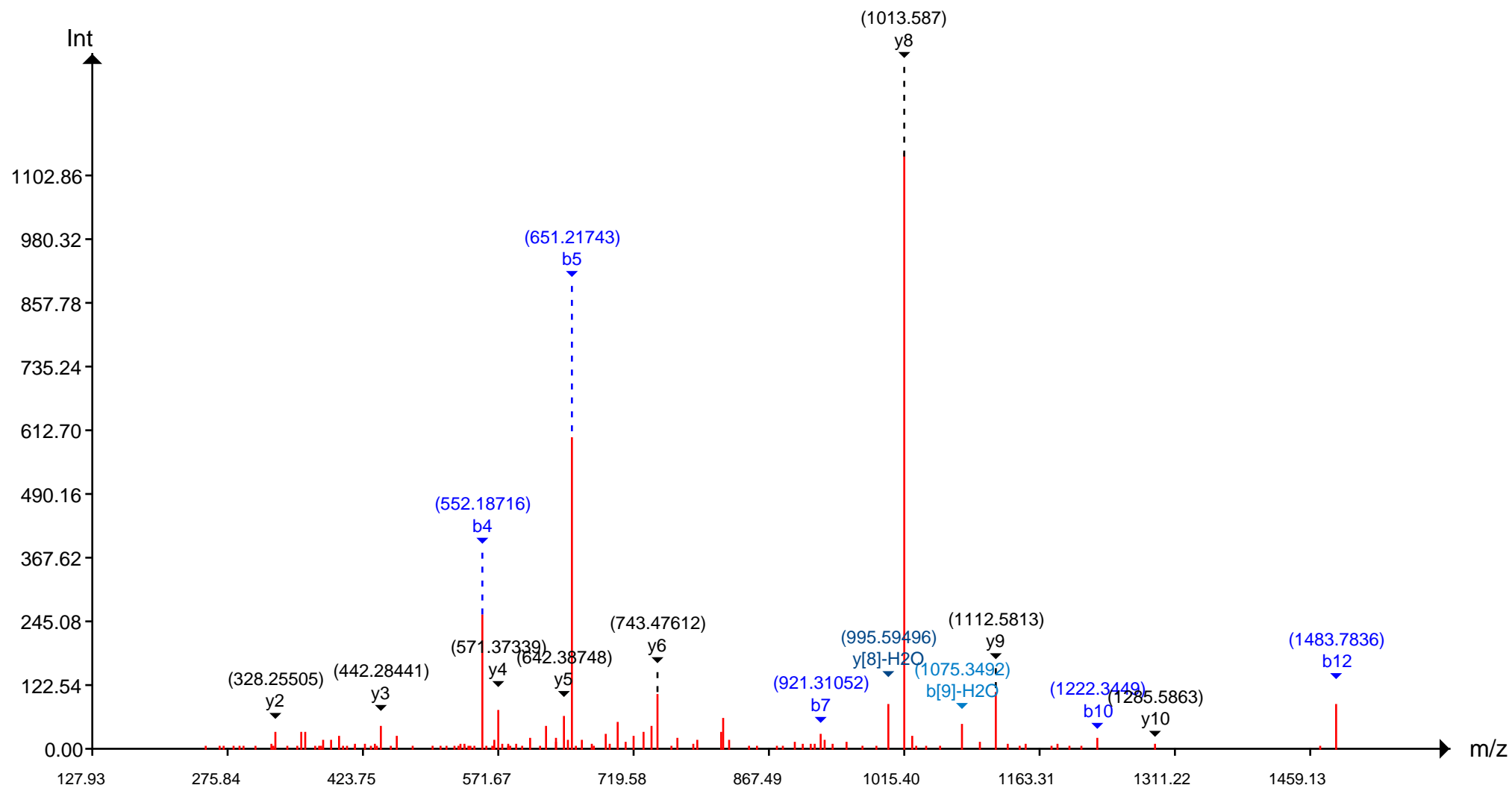


Info			
Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH	Ion score	54.89
Protein	P17742	Confidence level	99.00%
Precursor m/z	832.4595	Identity Threshold	47.49
Charge	2+	Mass error	0.00215



Info

Modified Sequence	AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH	Ion score	50.59
Protein	P17742	Confidence level	99.00%
Precursor m/z	832.4594	Identity Threshold	47.48
Charge	2+	Mass error	0.00195



Info

Modified Sequence AcD3-FADK<AcD3K*>VPK<AcD3K*>TAENFR<C13>-COOH

Ion score 53.58

Protein P17742

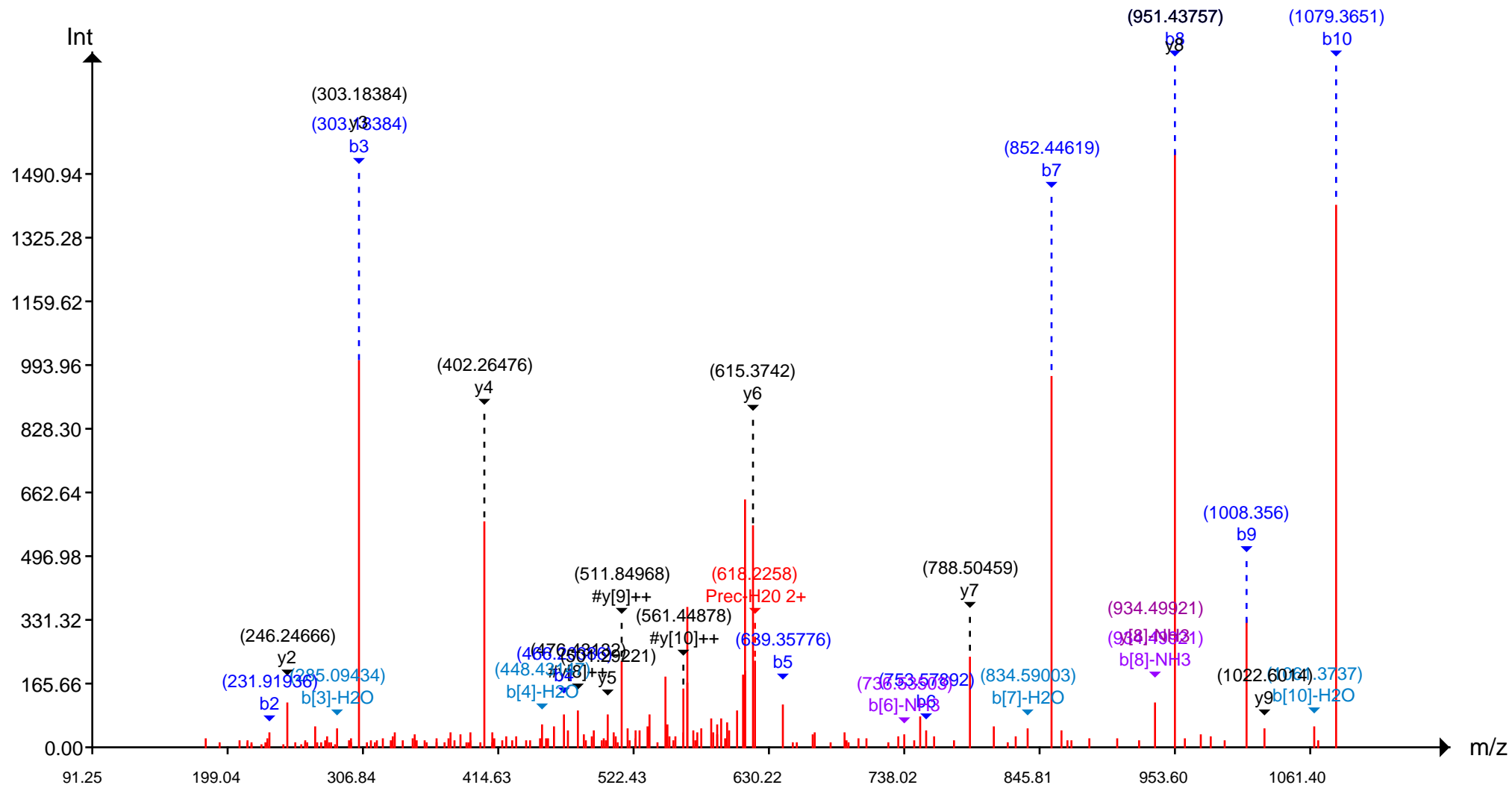
Confidence level 99.00%

Precursor m/z 832.45965

Identity Threshold 47.56

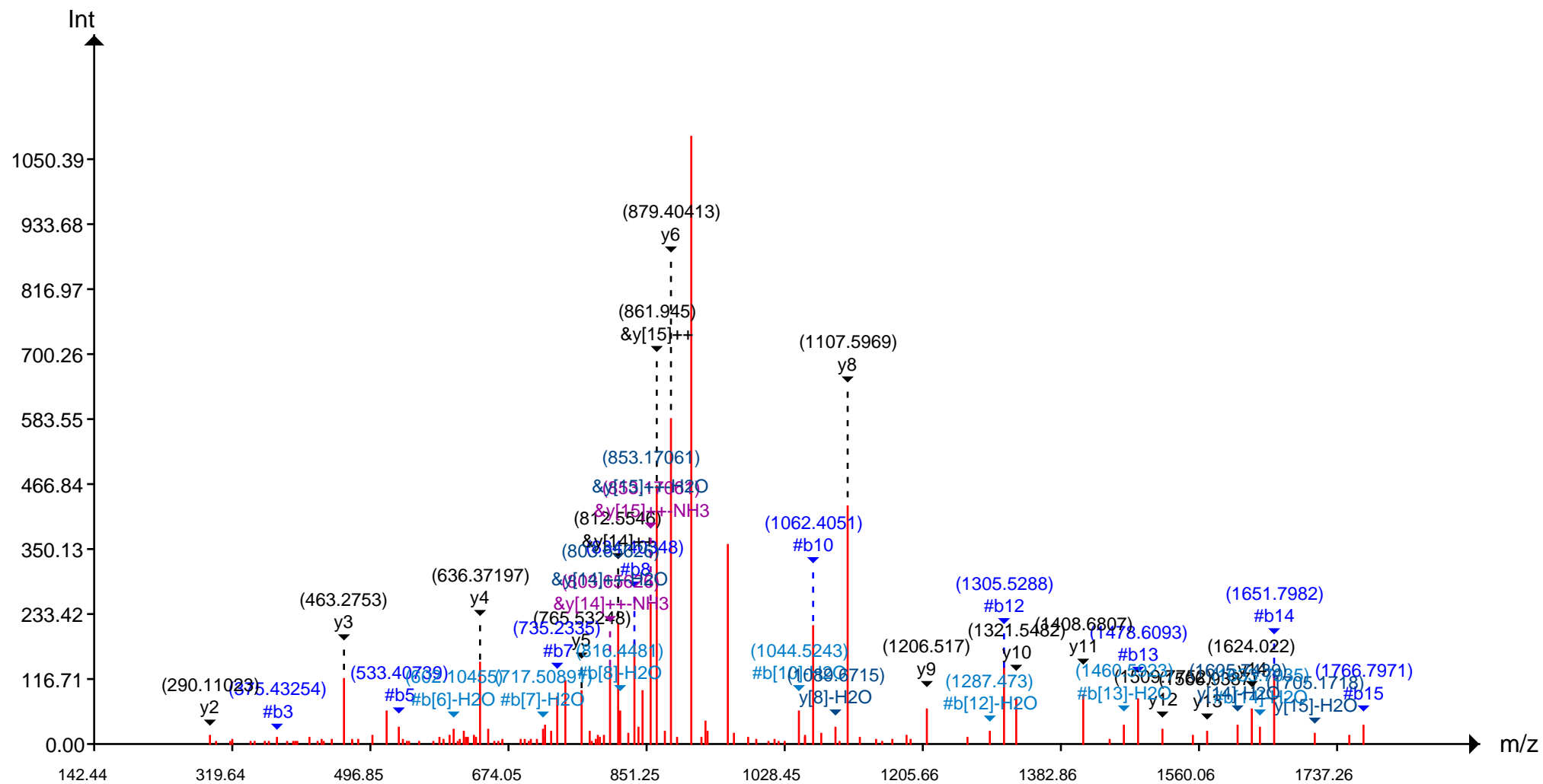
Charge 2+

Mass error 0.00245



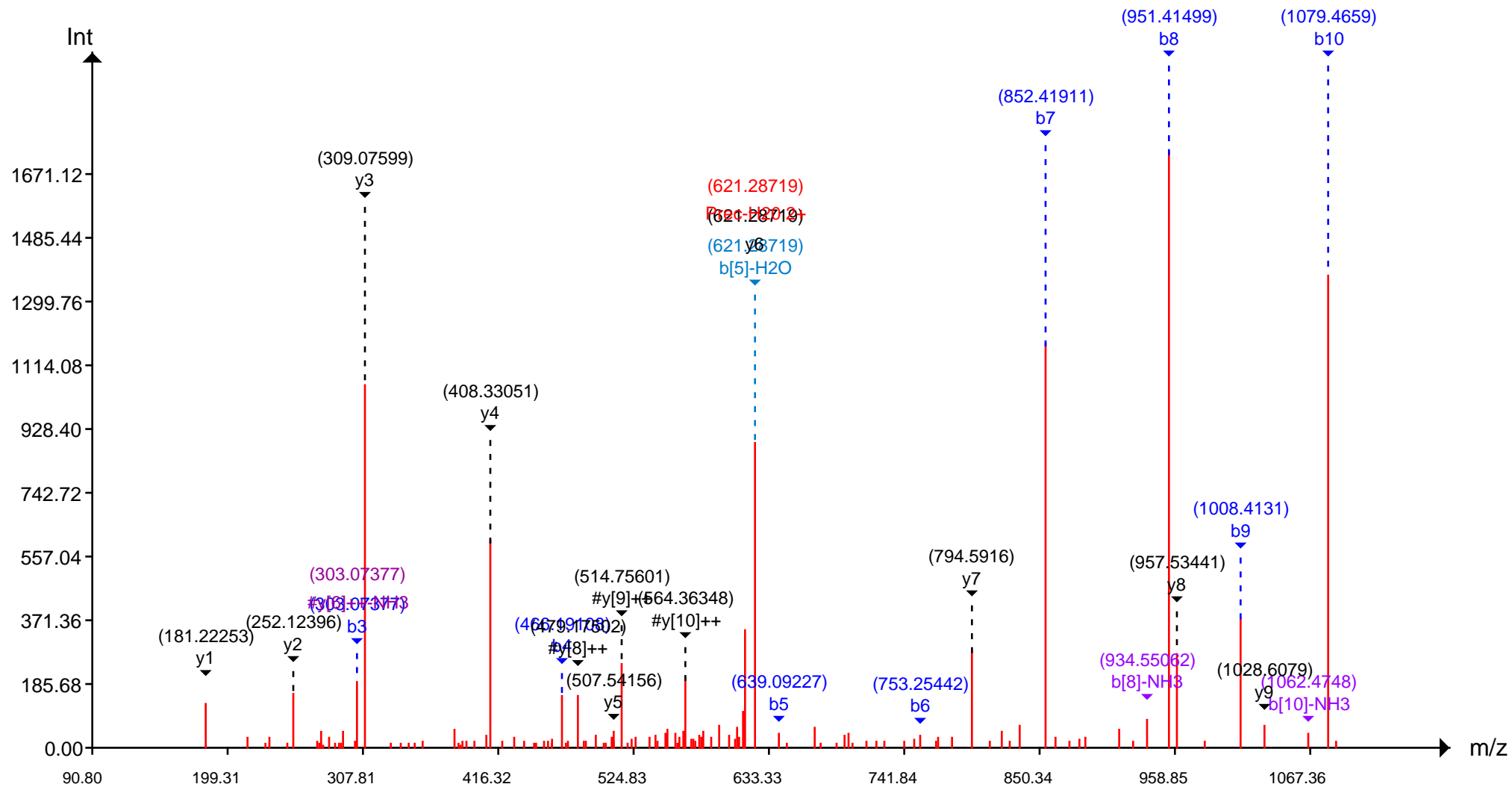
Info

Modified Sequence	AcD3-SVAYK<AcD3K*>NVVGAR-COOH	Ion score	61.49
Protein	Q9CQV8, P68510, P61982, P63101	Confidence level	99.00%
Precursor m/z	627.35983	Identity Threshold	47.06
Charge	2+	Mass error	4.72E-4



Info

Modified Sequence	AcD3-K<AcD3K*>VGGTSDVEVNEK<AcD3K*>K<AcD3K*>DR-COOH	Ion score	108.74
Protein	P63038	Confidence level	99.00%
Precursor m/z	971.02023	Identity Threshold	47.69
Charge	2+	Mass error	0.001875



Info

Modified Sequence AcD3-SVAYK<AcD3K*>NVVGAR<C13>-COOH

Ion score 69.22

Protein Q9CQV8, P68510, P61982, P63101

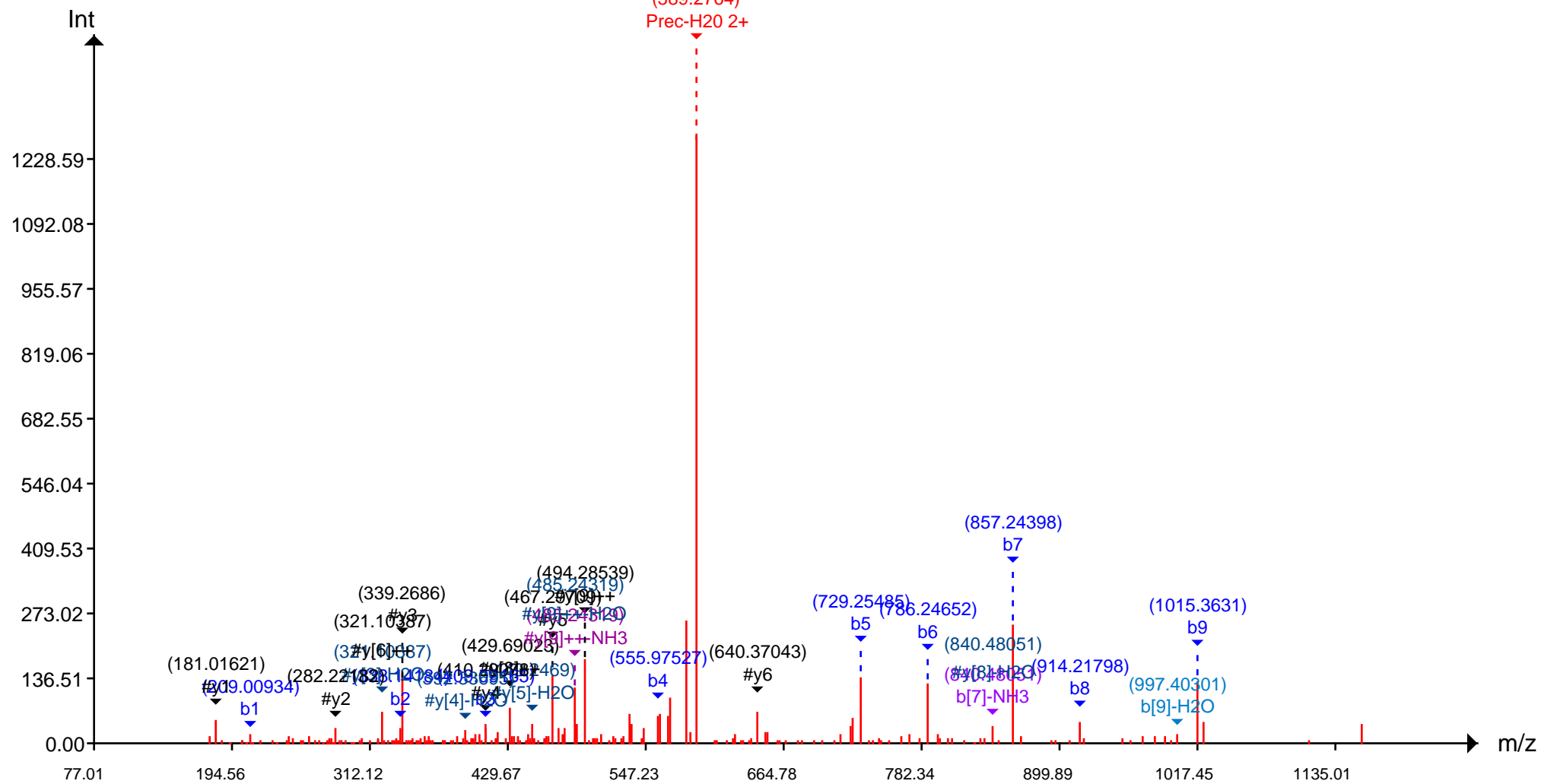
Confidence level 99.00%

Precursor m/z 630.36973

Identity Threshold 46.17

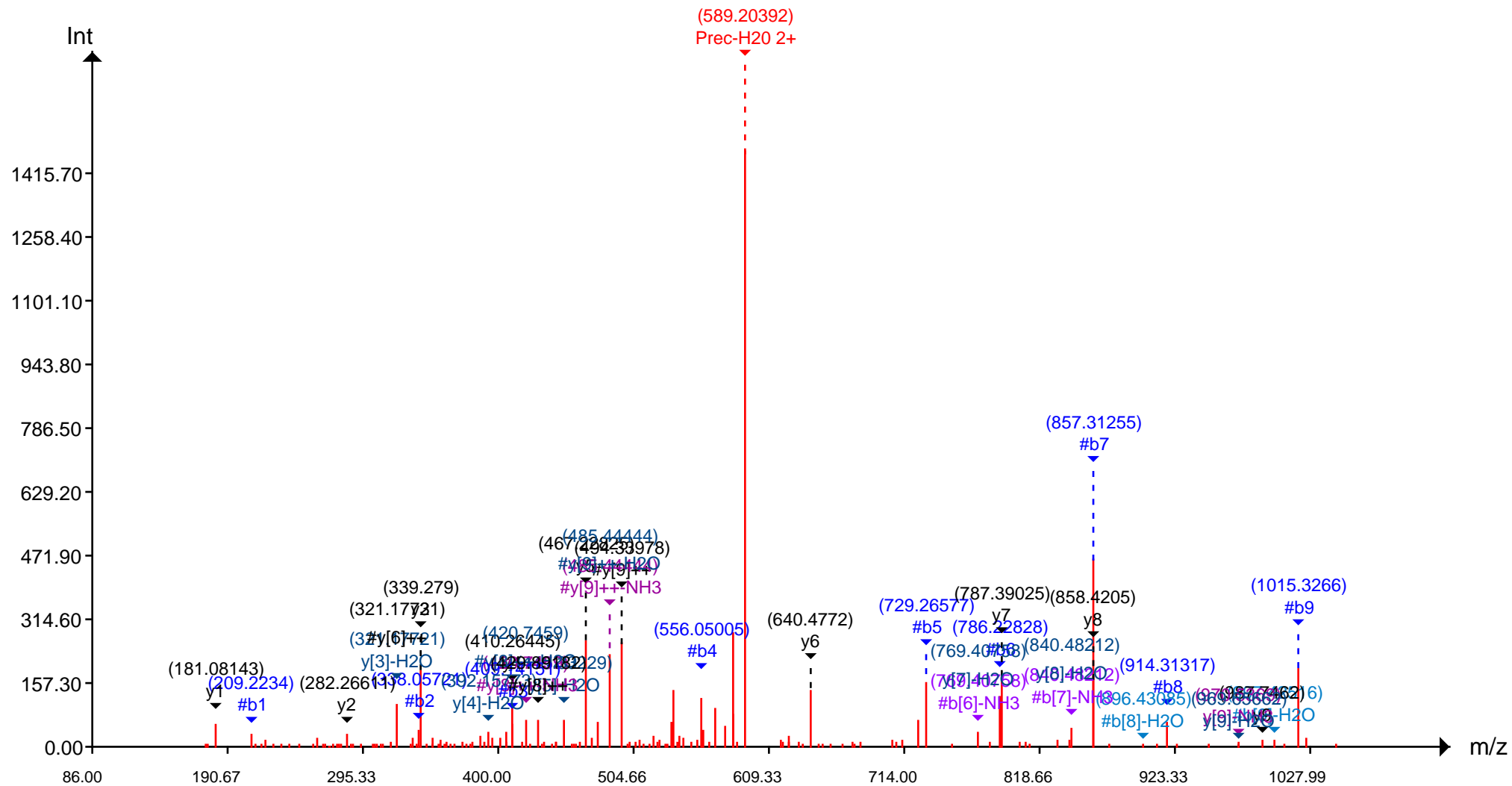
Charge 2+

Mass error 1.31E-4



Info

Modified Sequence	AcD3-YEAM<Mox*>K<AcD3K*>GAGTR<C13>-COOH	Ion score	52.41
Protein	P10107	Confidence level	99.00%
Precursor m/z	598.30319	Identity Threshold	47.79
Charge	2+	Mass error	1.23E-4



Info

Modified Sequence AcD3-YEAM<Mox*>K<AcD3K*>GAGTR<C13>-COOH

Ion score 81.11

Protein P10107

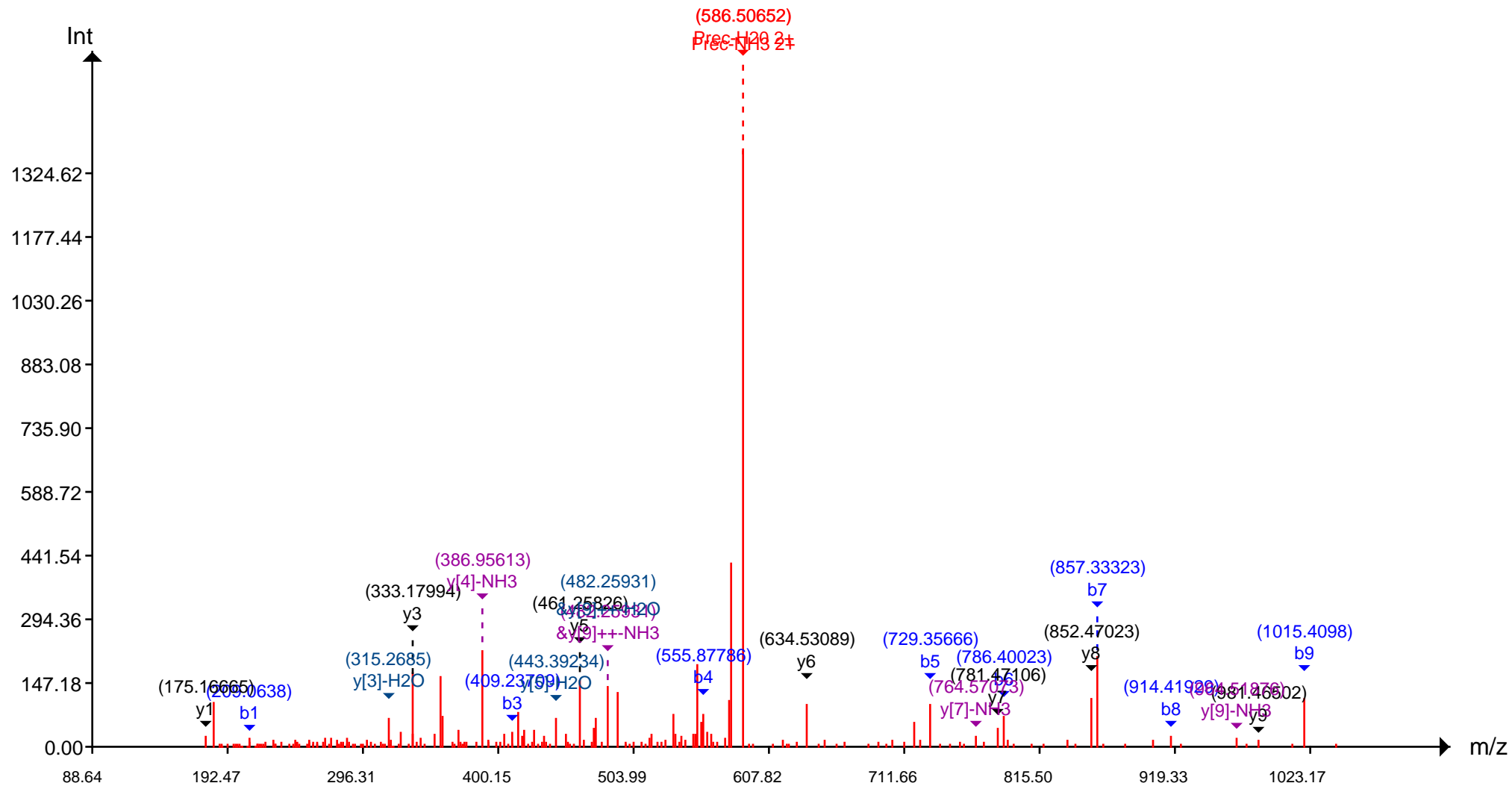
Confidence level 99.00%

Precursor m/z 598.30255

Identity Threshold 47.72

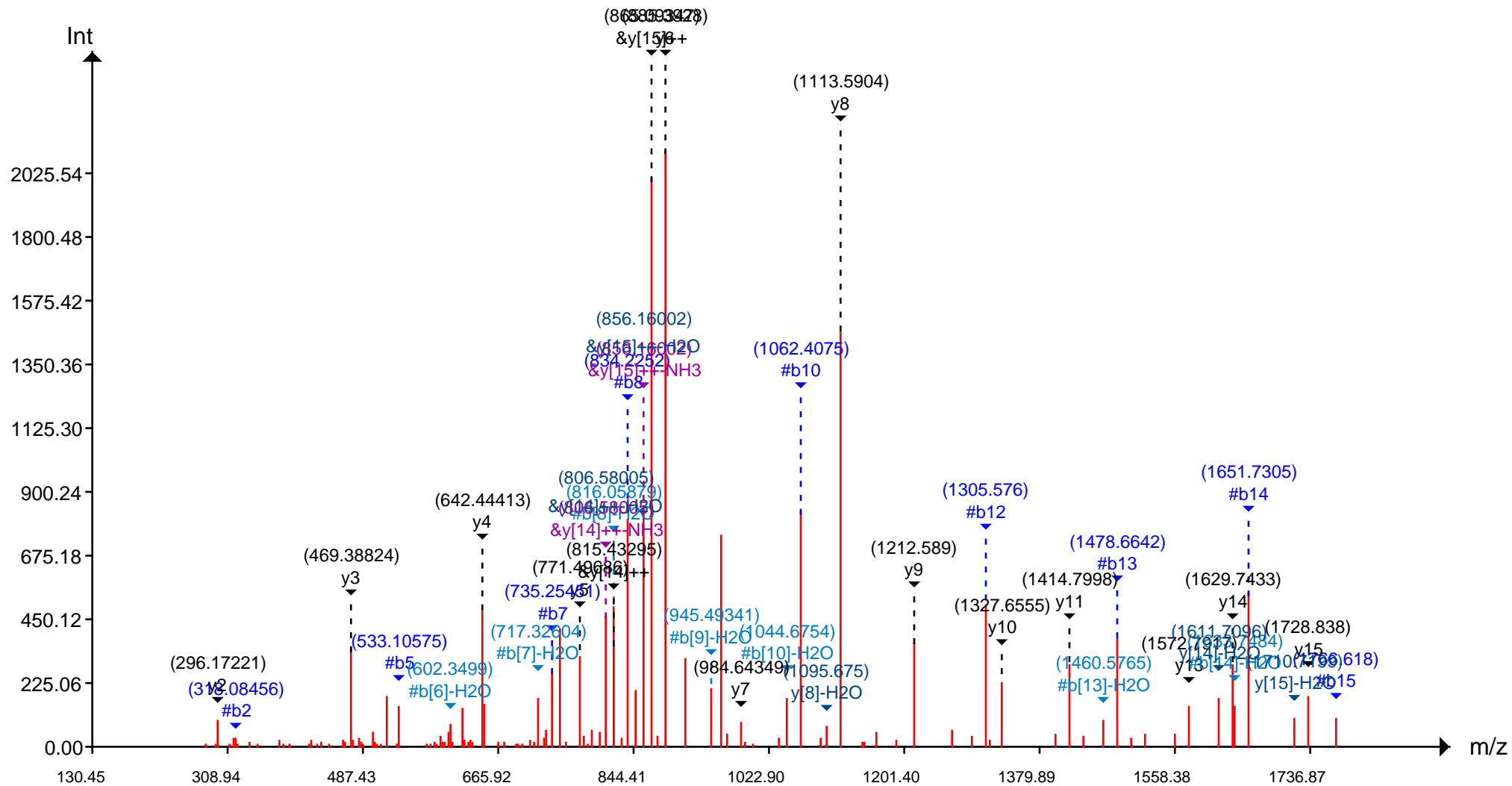
Charge 2+

Mass error -0.001157



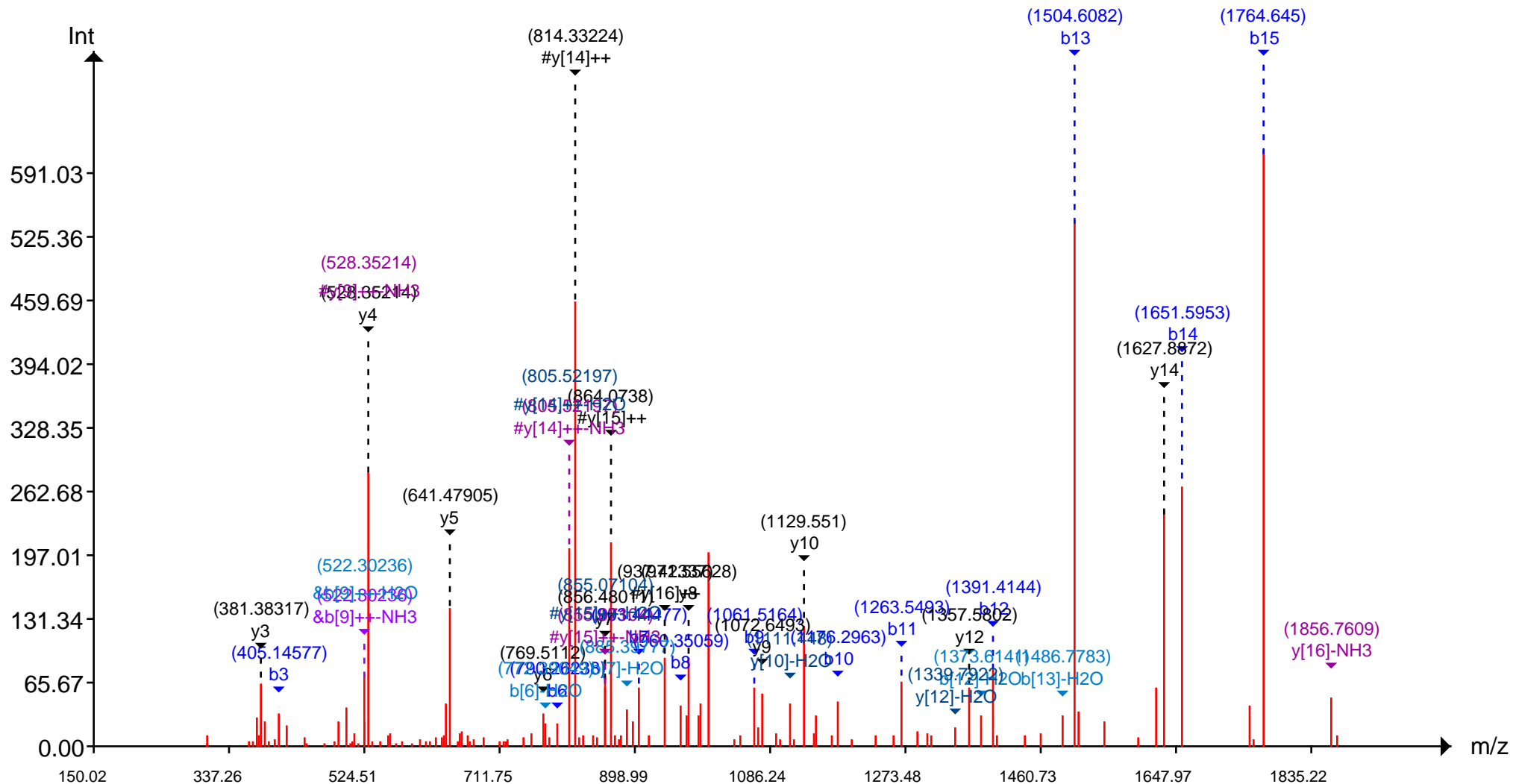
Info

Modified Sequence	AcD3-YEAM<Mox*>K<AcD3K*>GAGTR-COOH	Ion score	48.17
Protein	P10107	Confidence level	99.00%
Precursor m/z	595.29334	Identity Threshold	47.05
Charge	2+	Mass error	5.64E-4



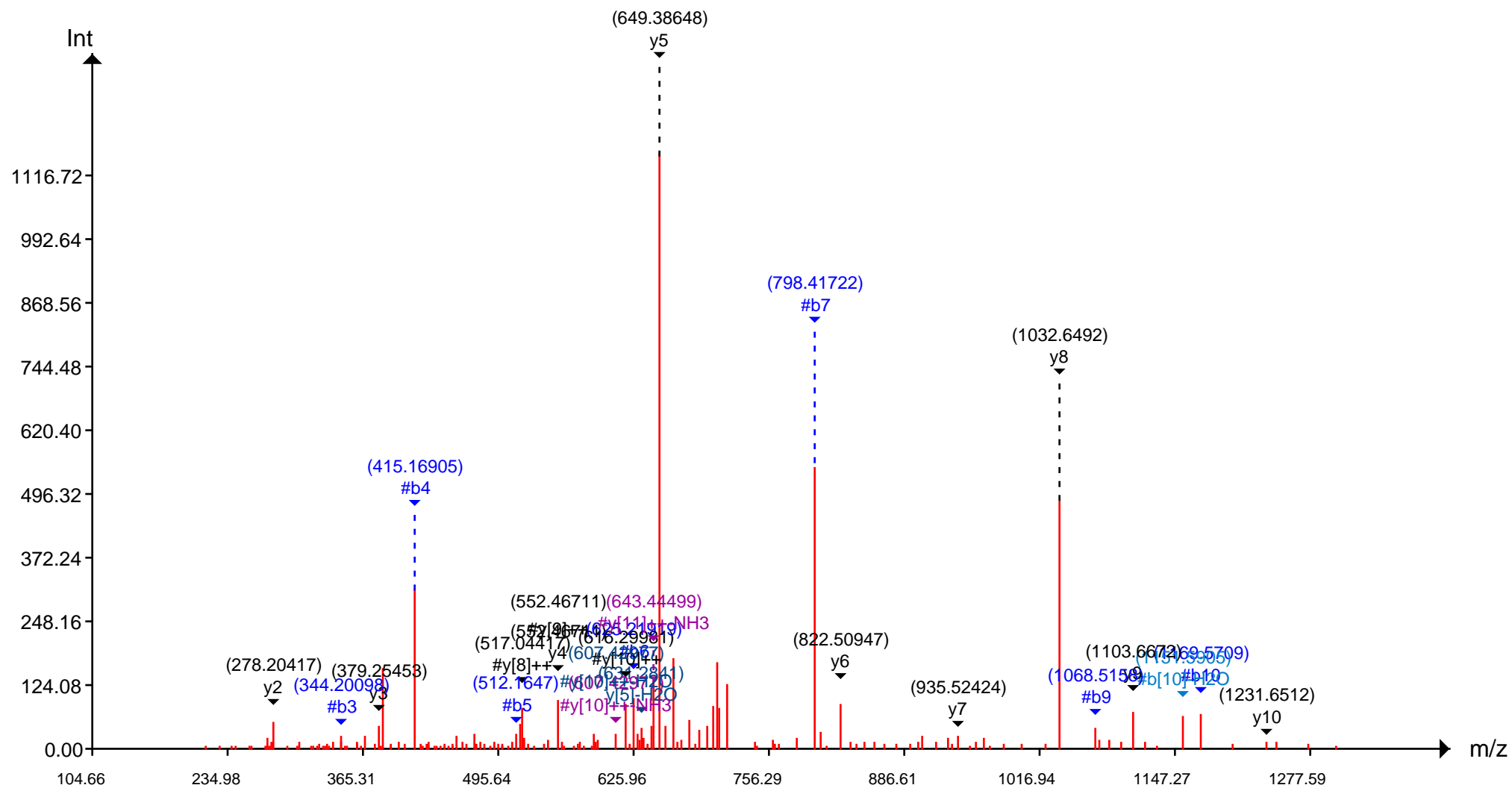
Info

Modified Sequence	AcD3-K<AcD3K*>VGGTSDVEVNEK<AcD3K*>K<AcD3K*>DR<C13>-COOH	Ion score	103.48
Protein	P63038	Confidence level	99.00%
Precursor m/z	974.02999	Identity Threshold	47.55
Charge	2+	Mass error	0.001254



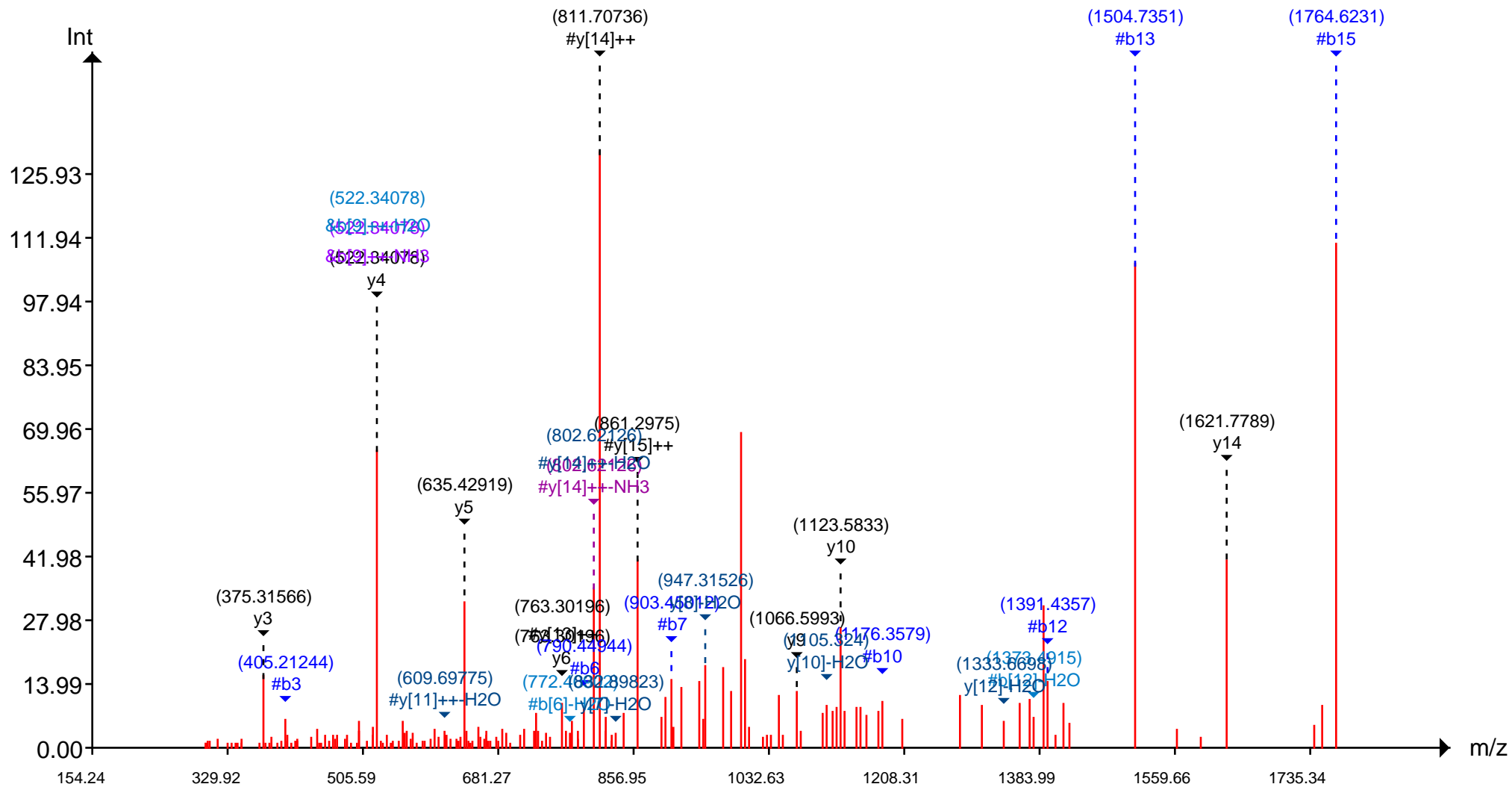
Info

Modified Sequence	AcD3-LFVPK<AcD3K*>DLGTDSQIFISR<C13>-COOH	Ion score	69.63
Protein	Q61598	Confidence level	99.00%
Precursor m/z	1016.5717	Identity Threshold	46.08
Charge	2+	Mass error	0.003344



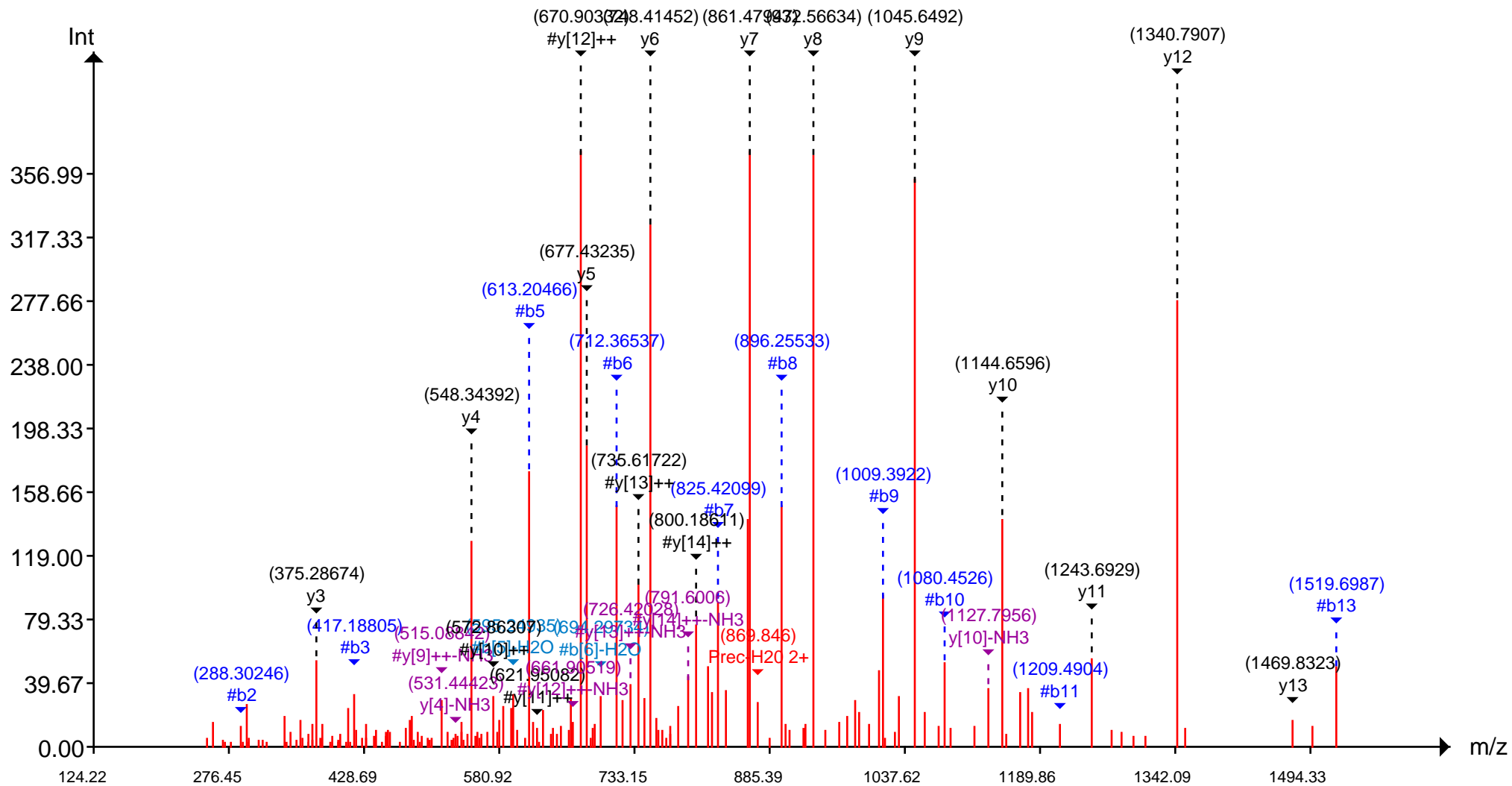
Info

Modified Sequence	AcD3-VAQAPLK<AcD3K*>PK<AcD3K*>TPR<C13>-COOH	Ion score	68.74
Protein	Q9R190	Confidence level	99.00%
Precursor m/z	723.95835	Identity Threshold	39.22
Charge	2+	Mass error	9.79E-4



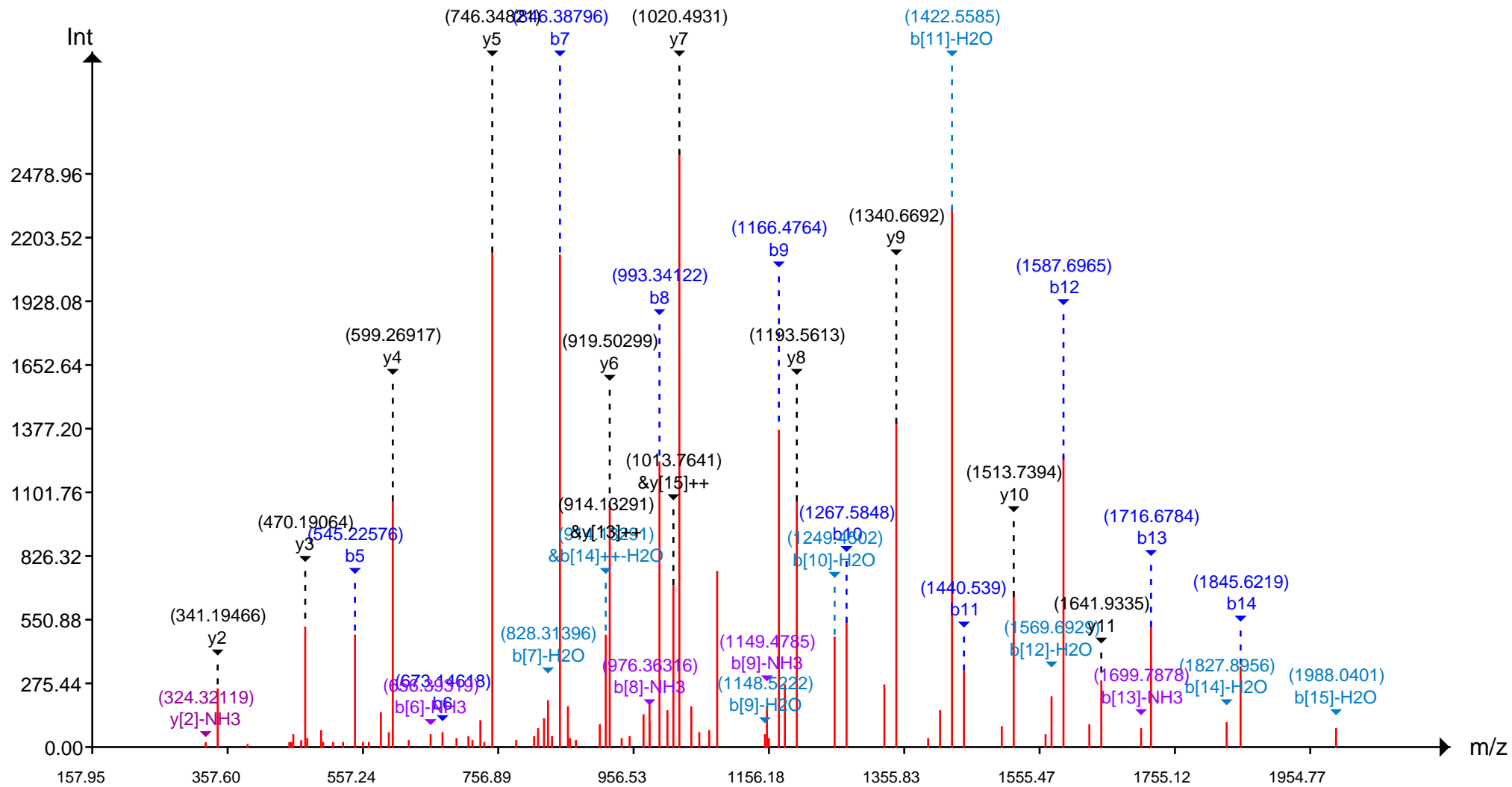
Info

Modified Sequence	AcD3-LFVPK<AcD3K*>DLGTDSQIFISR-COOH	Ion score	48.54
Protein	Q61598	Confidence level	99.00%
Precursor m/z	1014.0574	Identity Threshold	47.06
Charge	2+	Mass error	0.994886



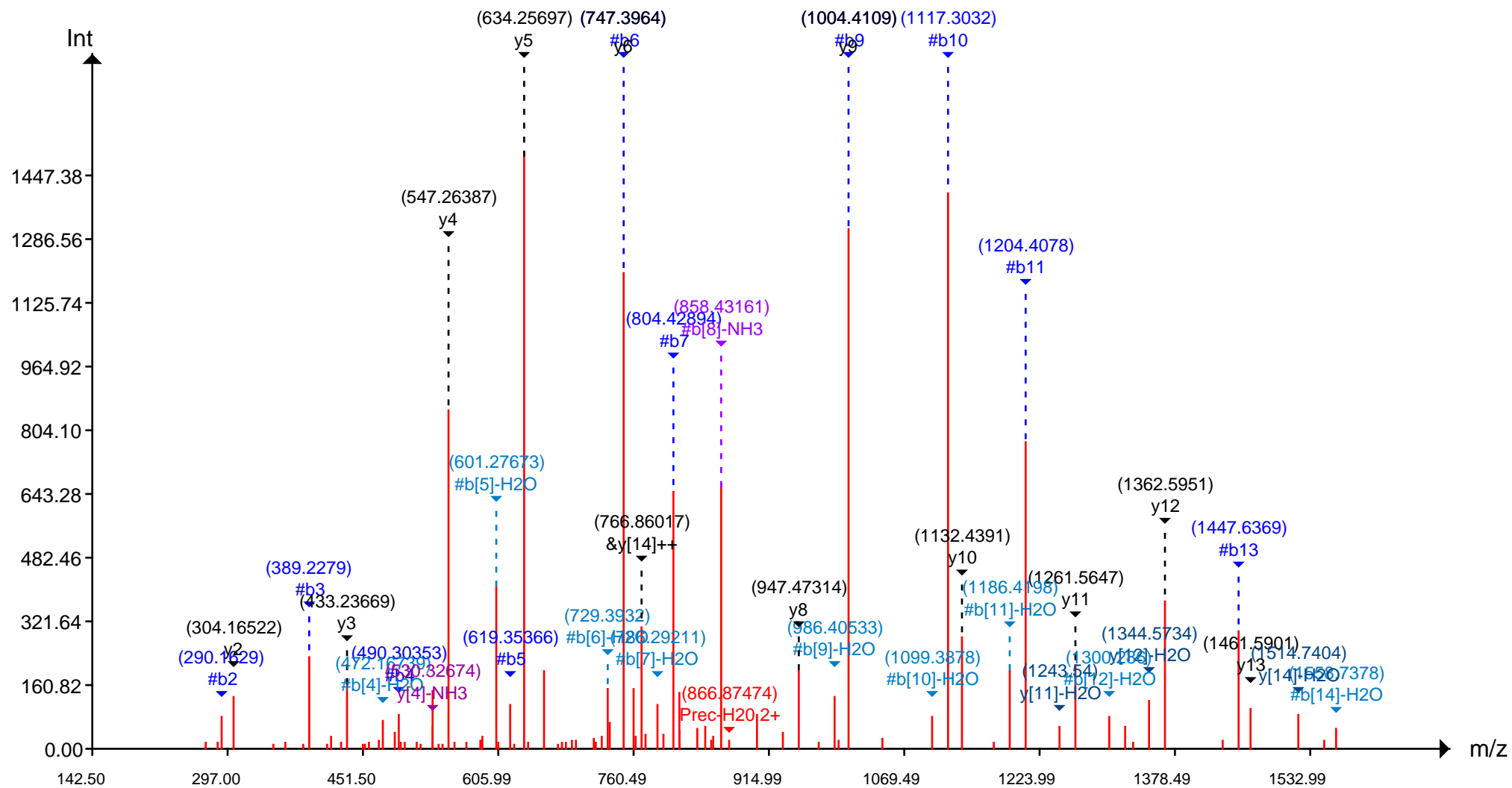
Info

Modified Sequence	AcD3-LEEPVVLALAEK<AcD3K*>HGR<C13>-COOH	Ion score	89.94
Protein	Q9JII6	Confidence level	99.00%
Precursor m/z	879.01351	Identity Threshold	44.49
Charge	2+	Mass error	0.002763



Info

Modified Sequence	AcD3-NAENAQK<AcD3K*>FK<AcD3K*>TK<AcD3K*>FEEC<Cmm*>R<C13>-COOH	Ion score	94.47
Protein	P34022	Confidence level	99.00%
Precursor m/z	1093.5566	Identity Threshold	47.93
Charge	2+	Mass error	0.002792



Info

Modified Sequence AcD3-K<AcD3K*>AVTEQGAELSNEER-COOH

Ion score 112.74

Protein P68254

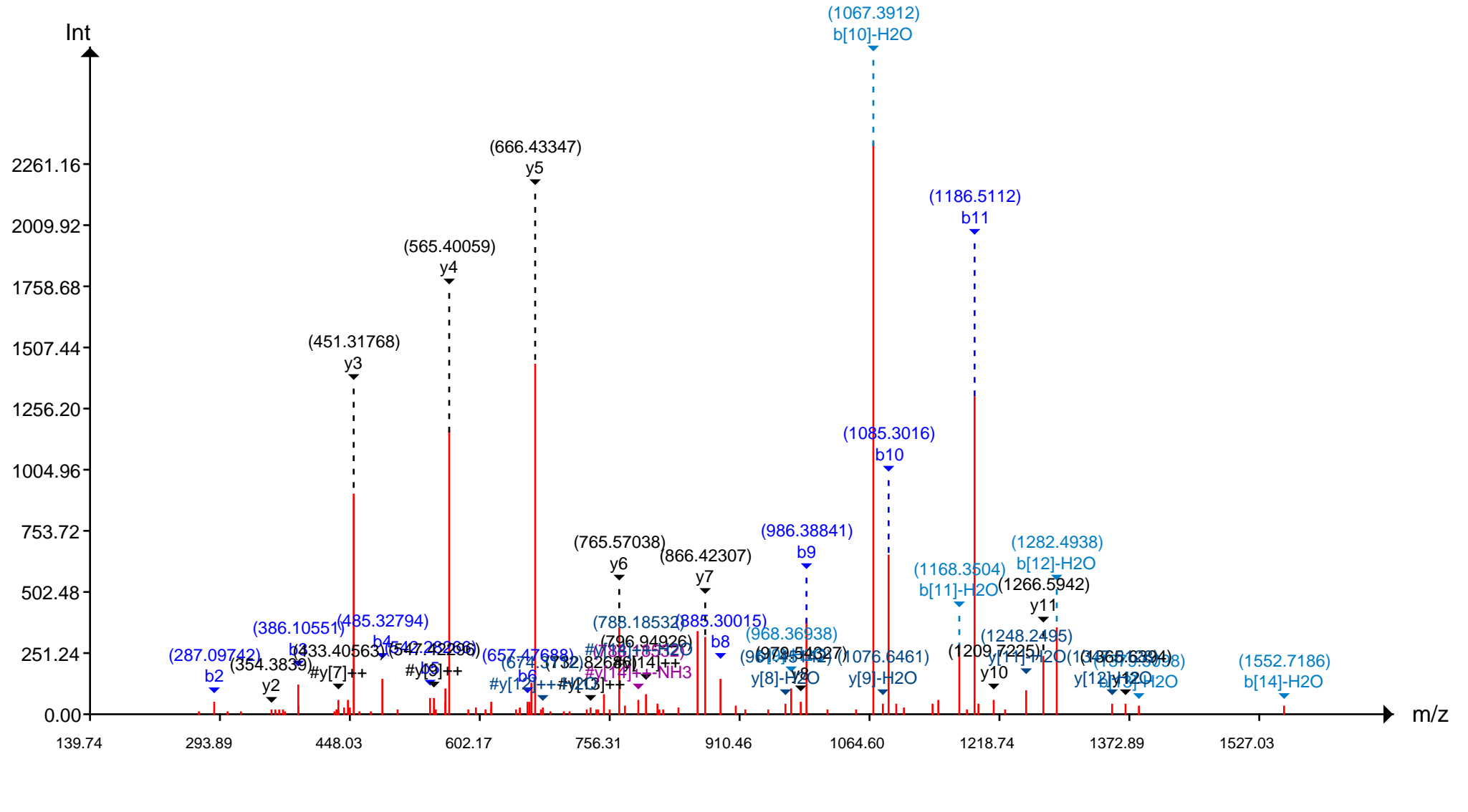
Confidence level 99.00%

Precursor m/z 875.94098

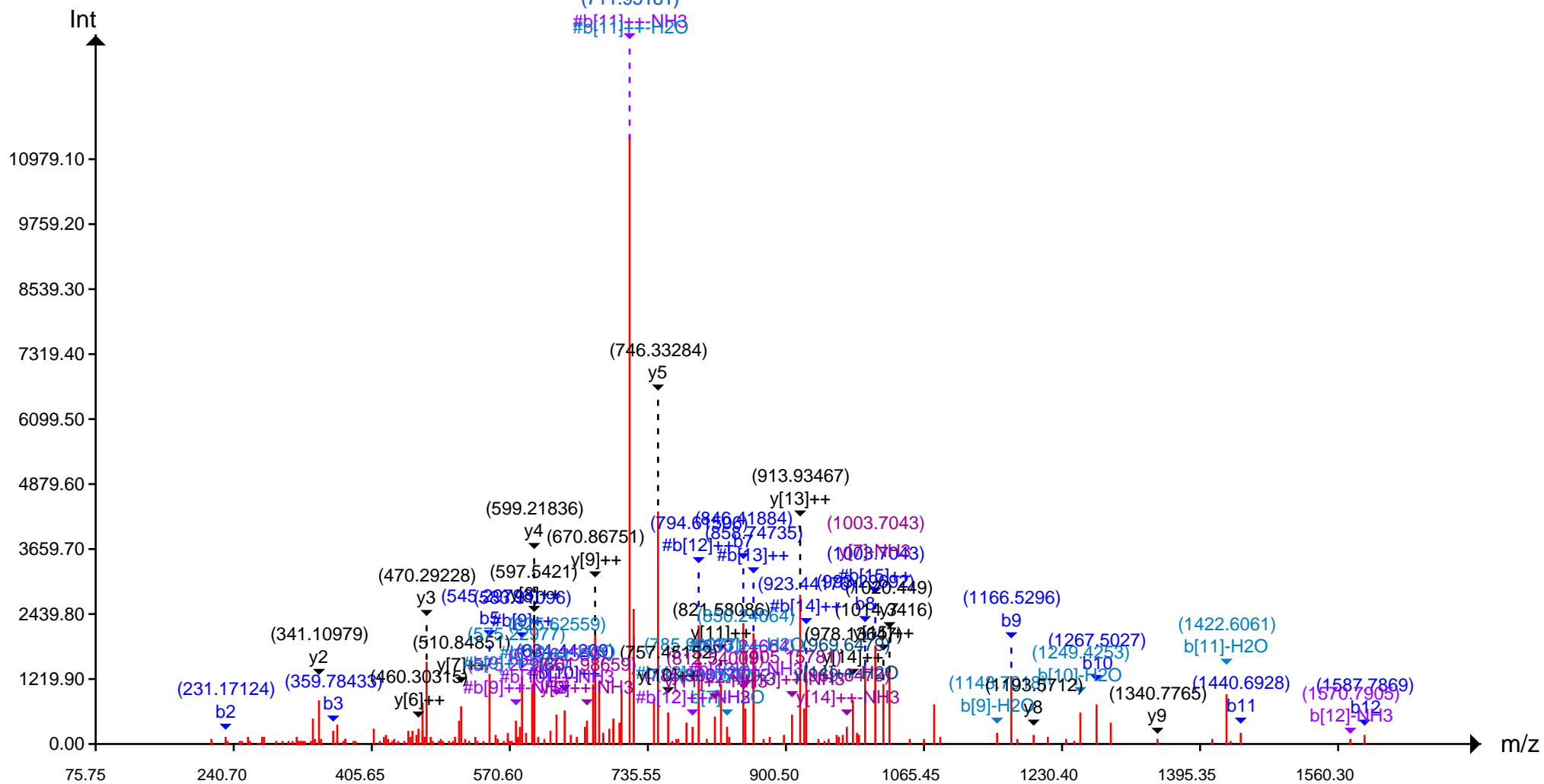
Identity Threshold 47.90

Charge 2+

Mass error 0.002265

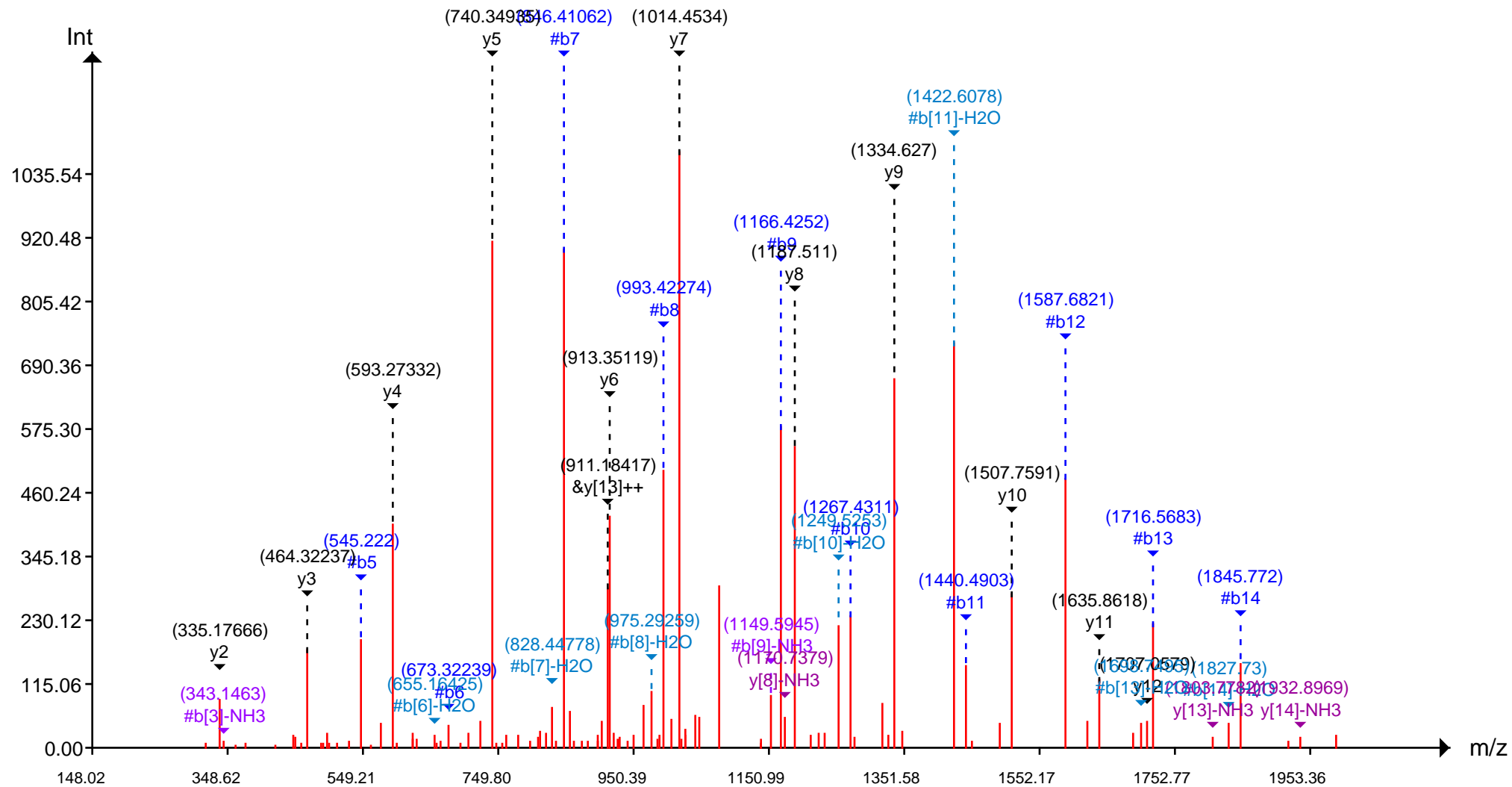


Info			
Modified Sequence	AcD3-IQVVGDDLTVTNPK<AcD3K*>R<C13>-COOH	Ion score	72.73
Protein	P17182	Confidence level	99.00%
Precursor m/z	876.00048	Identity Threshold	45.78
Charge	2+	Mass error	0.002476



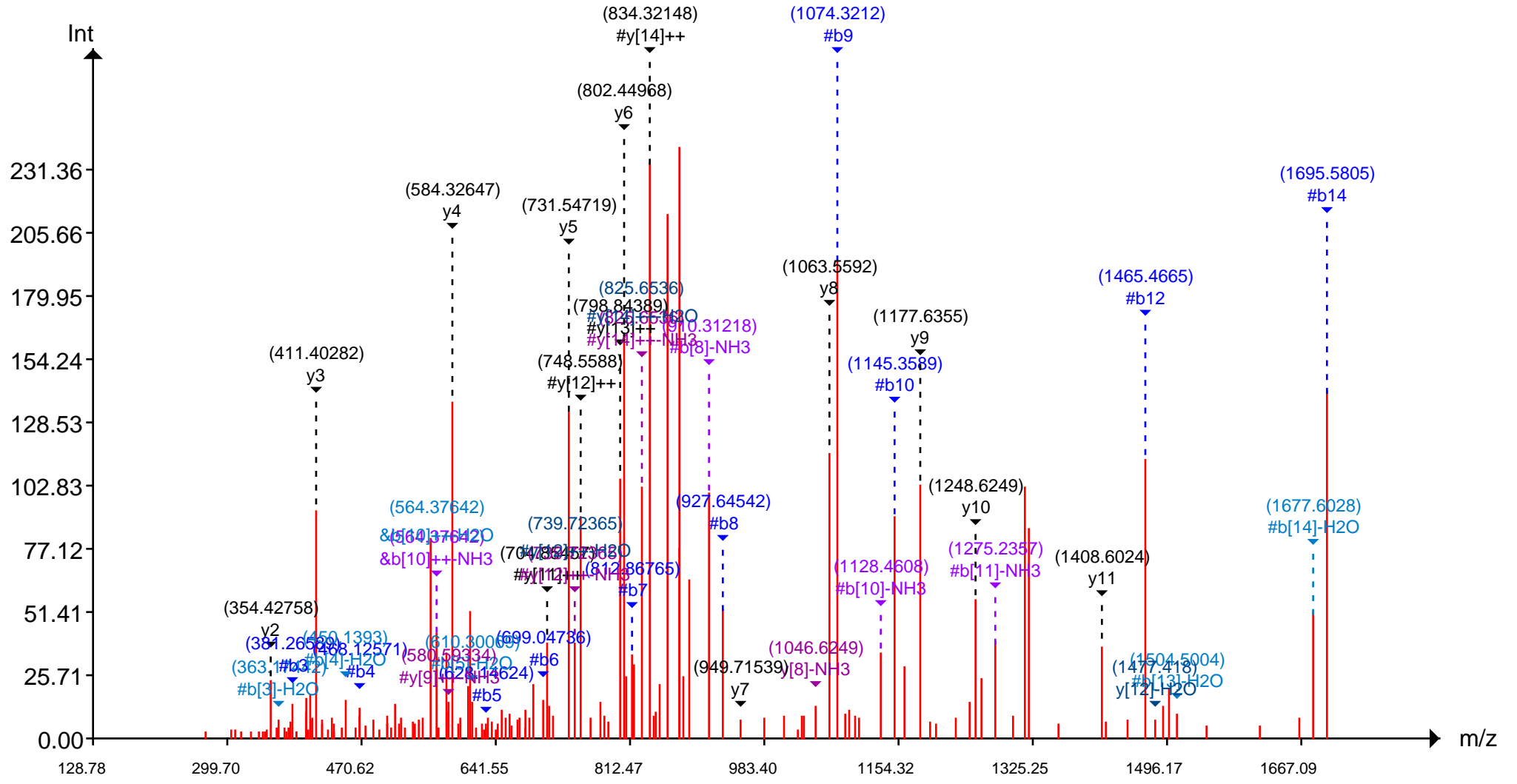
Info

Modified Sequence	AcD3-NAENAQK<AcD3K*>FK<AcD3K*>TK<AcD3K*>FEEC<Cmm*>R<C13>-COOH	Ion score	58.45
Protein	P34022	Confidence level	99.00%
Precursor m/z	729.37309	Identity Threshold	47.94
Charge	3+	Mass error	0.001586



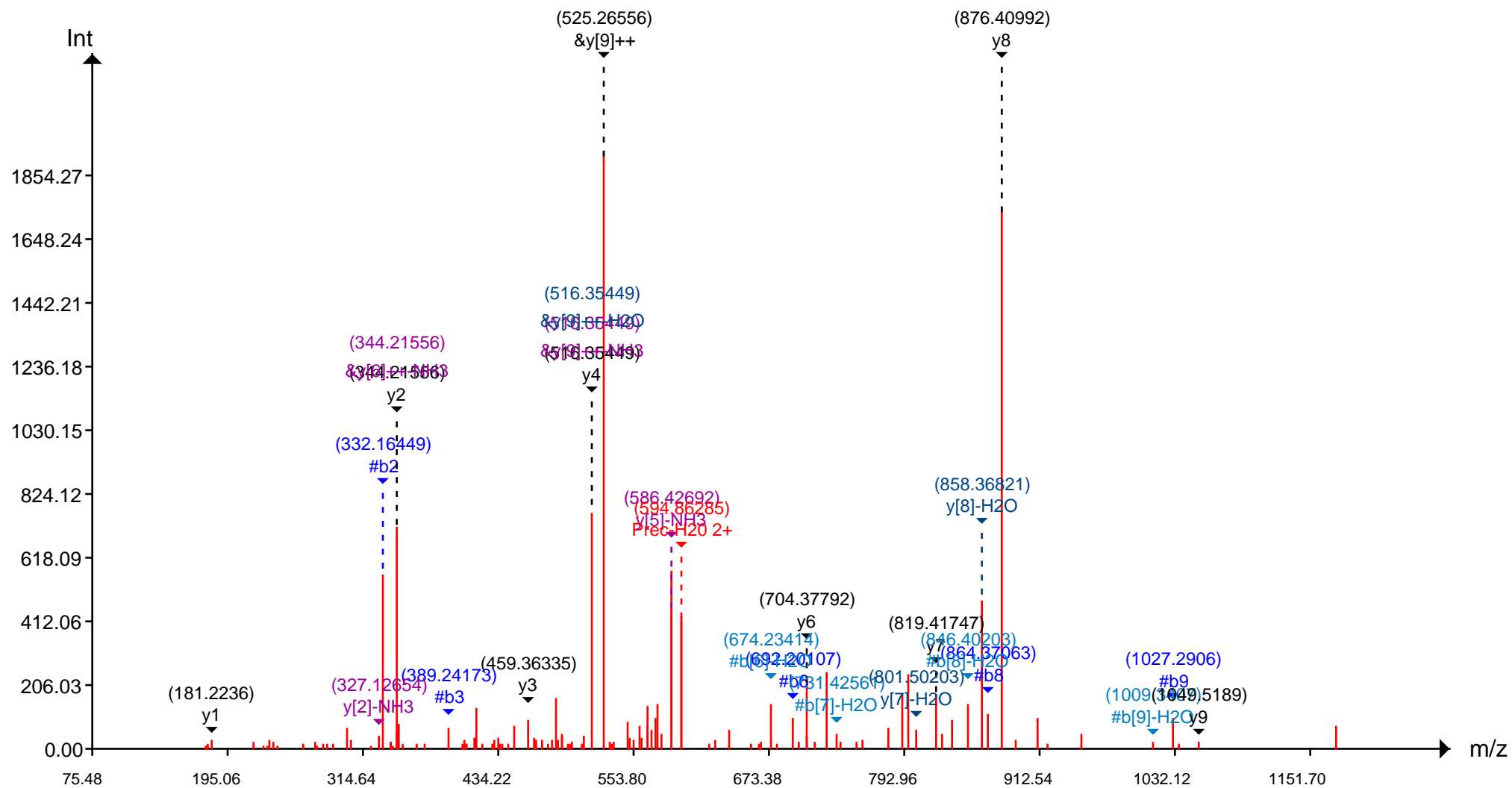
Info

Modified Sequence	AcD3-NAENAQK<AcD3K*>FK<AcD3K*>TK<AcD3K*>FEEC<Cmm*>R-COOH	Ion score	98.2
Protein	P34022	Confidence level	99.00%
Precursor m/z	1090.5467	Identity Threshold	47.74
Charge	2+	Mass error	0.003134



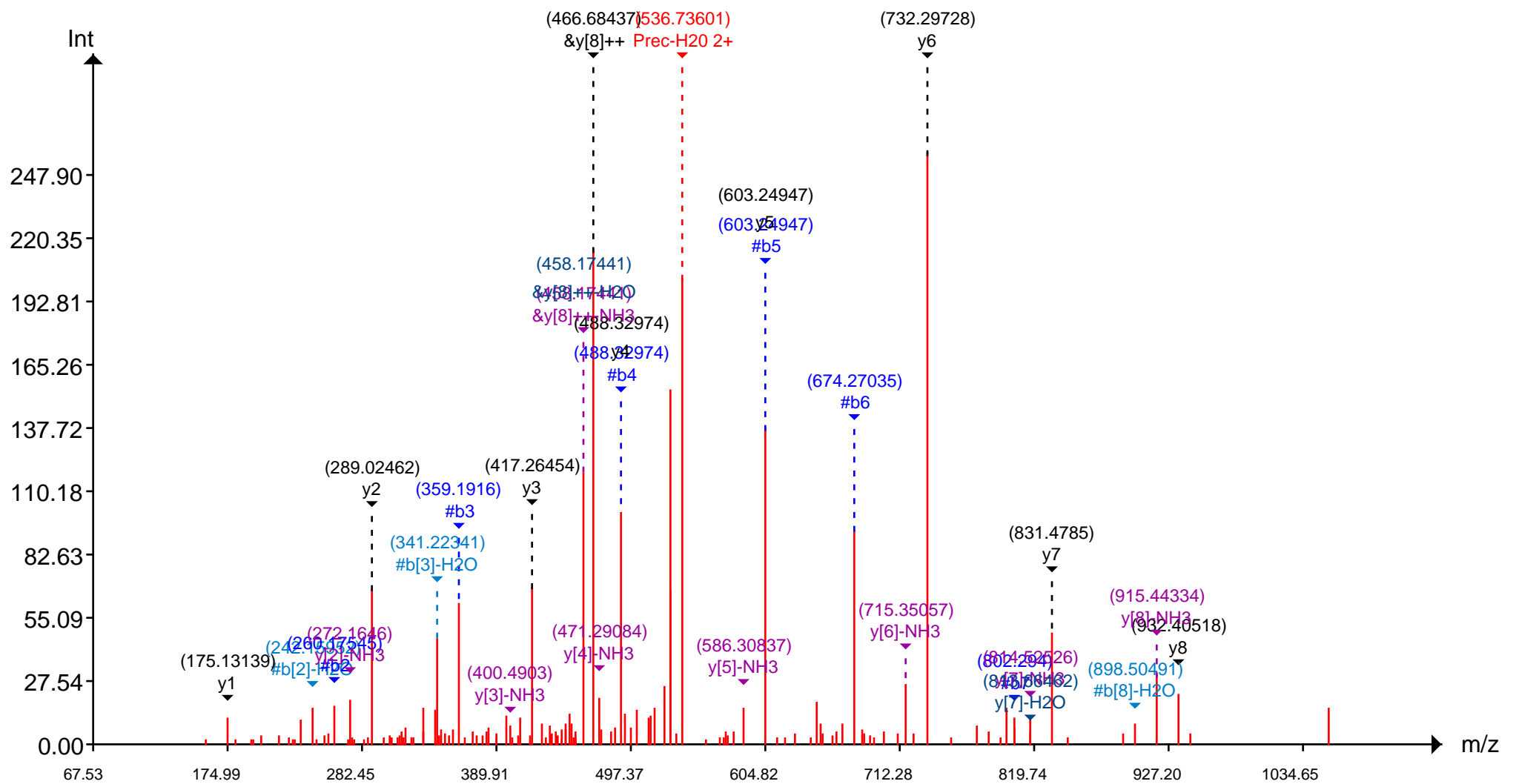
Info

Modified Sequence	AcD3-YATSC<Cmm*>ANNFAM<Mox*>K<AcD3K*>GK<AcD3K*>R<C13>-COOH	Ion score	48.54
Protein	Q9JKP7	Confidence level	99.00%
Precursor m/z	938.46143	Identity Threshold	47.23
Charge	2+	Mass error	0.002241



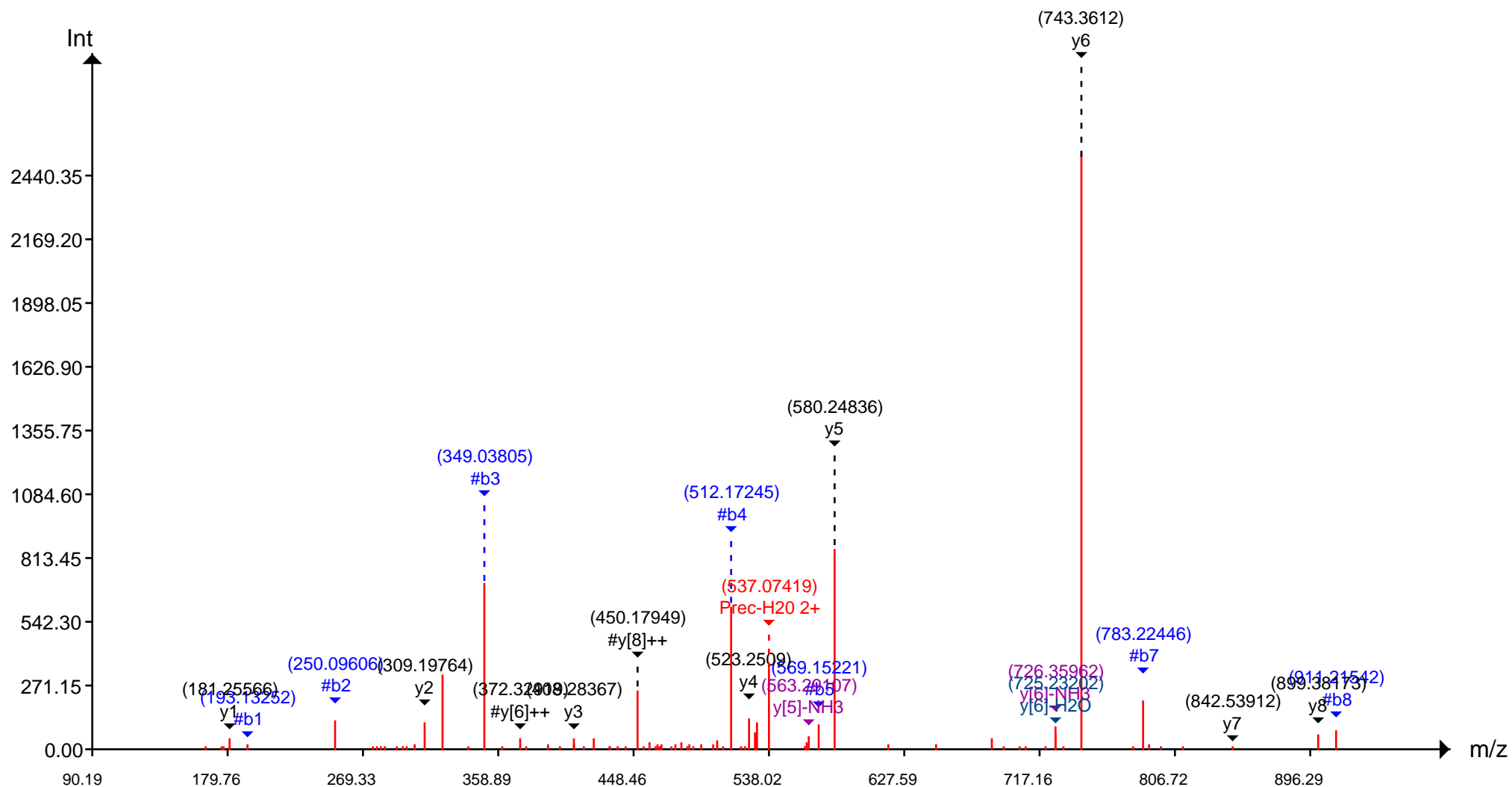
Info

Modified Sequence	AcD3-IK<AcD3K*>GDTSGDYR<C13>-COOH	Ion score	57.54
Protein	P97429	Confidence level	99.00%
Precursor m/z	604.31199	Identity Threshold	47.92
Charge	2+	Mass error	-5.4E-5



Info

Modified Sequence	AcD3-ITVEDAQNR-COOH	Ion score	59.04
Protein	Q9QYS9	Confidence level	99.00%
Precursor m/z	545.78202	Identity Threshold	47.56
Charge	2+	Mass error	1.26E-4



Info

Modified Sequence AcD3-FGVYGDVQR<C13>-COOH

Ion score 78.54

Protein P17225, Q91Z31

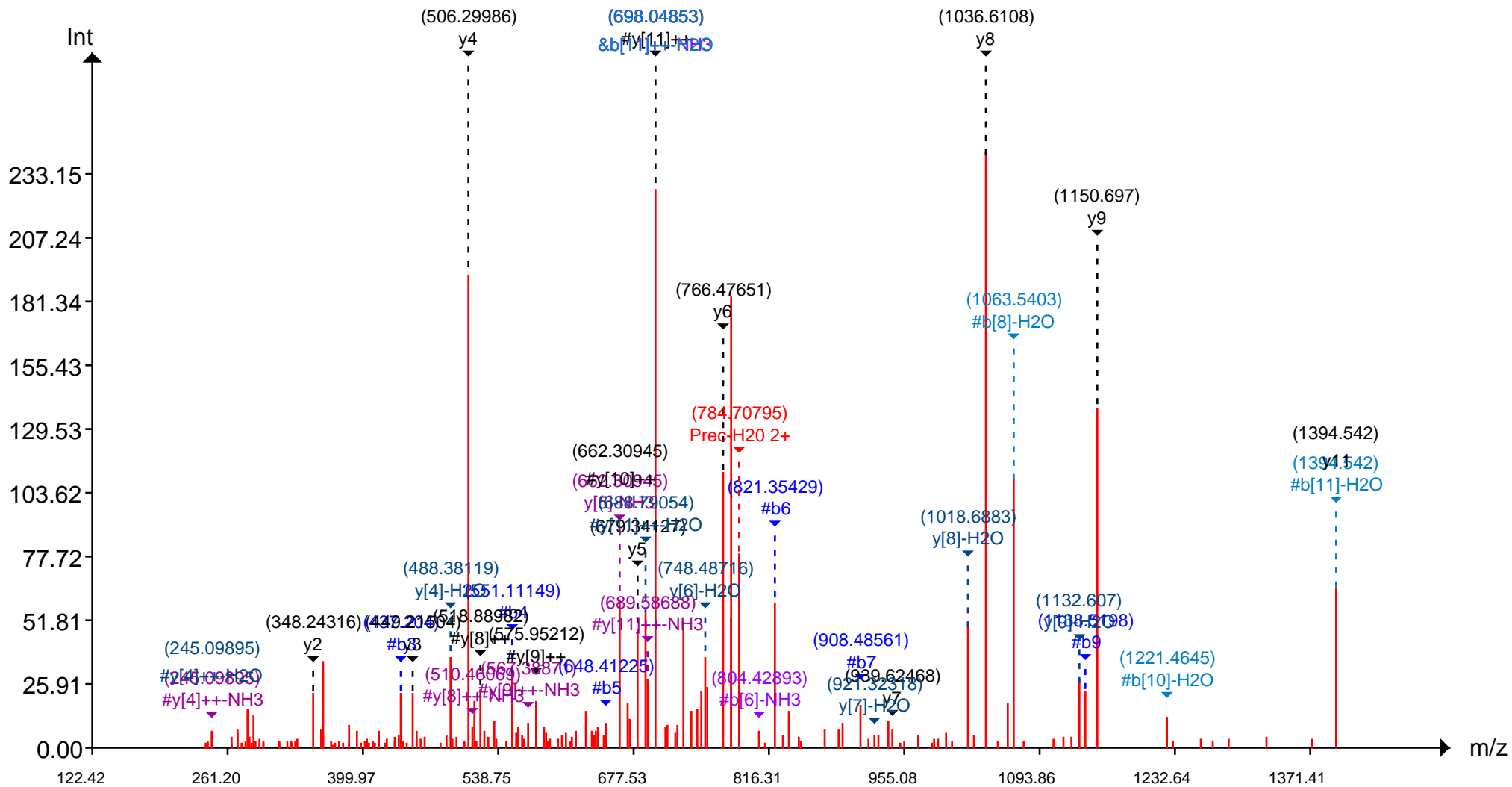
Confidence level 99.00%

Precursor m/z 546.28668

Identity Threshold 47.94

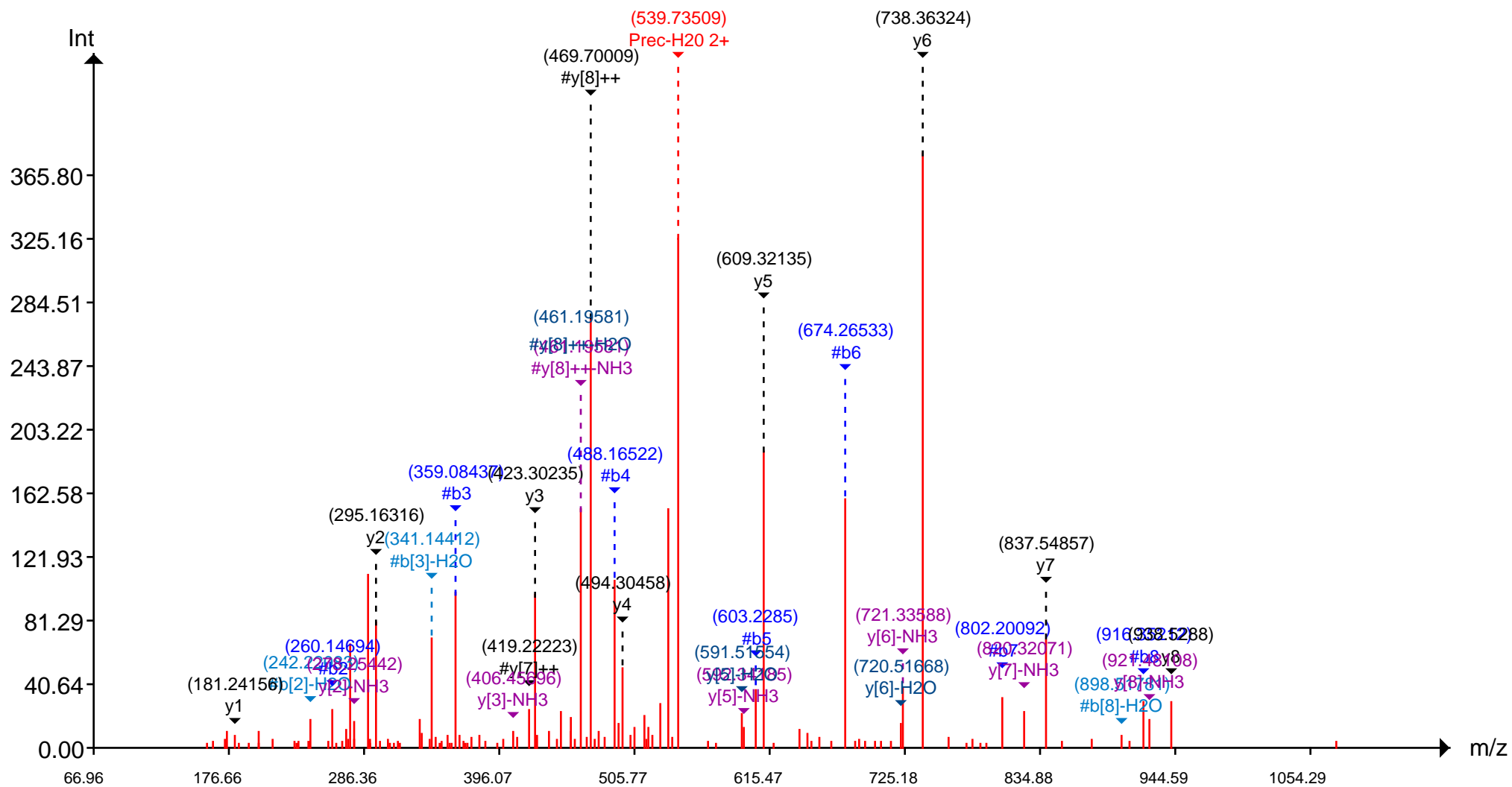
Charge 2+

Mass error 5.65E-4



Info

Modified Sequence	AcD3-FAK<AcD3K*>NPK<AcD3K*>SK<AcD3K*>GTK<AcD3K*>R-COOH	Ion score	48.13
Protein	Q9D554	Confidence level	99.00%
Precursor m/z	793.9786	Identity Threshold	42.99
Charge	2+	Mass error	0.001761



Info

Modified Sequence AcD3-ITVEDAQNR<C13>-COOH

Ion score 57.11

Protein Q9QYS9

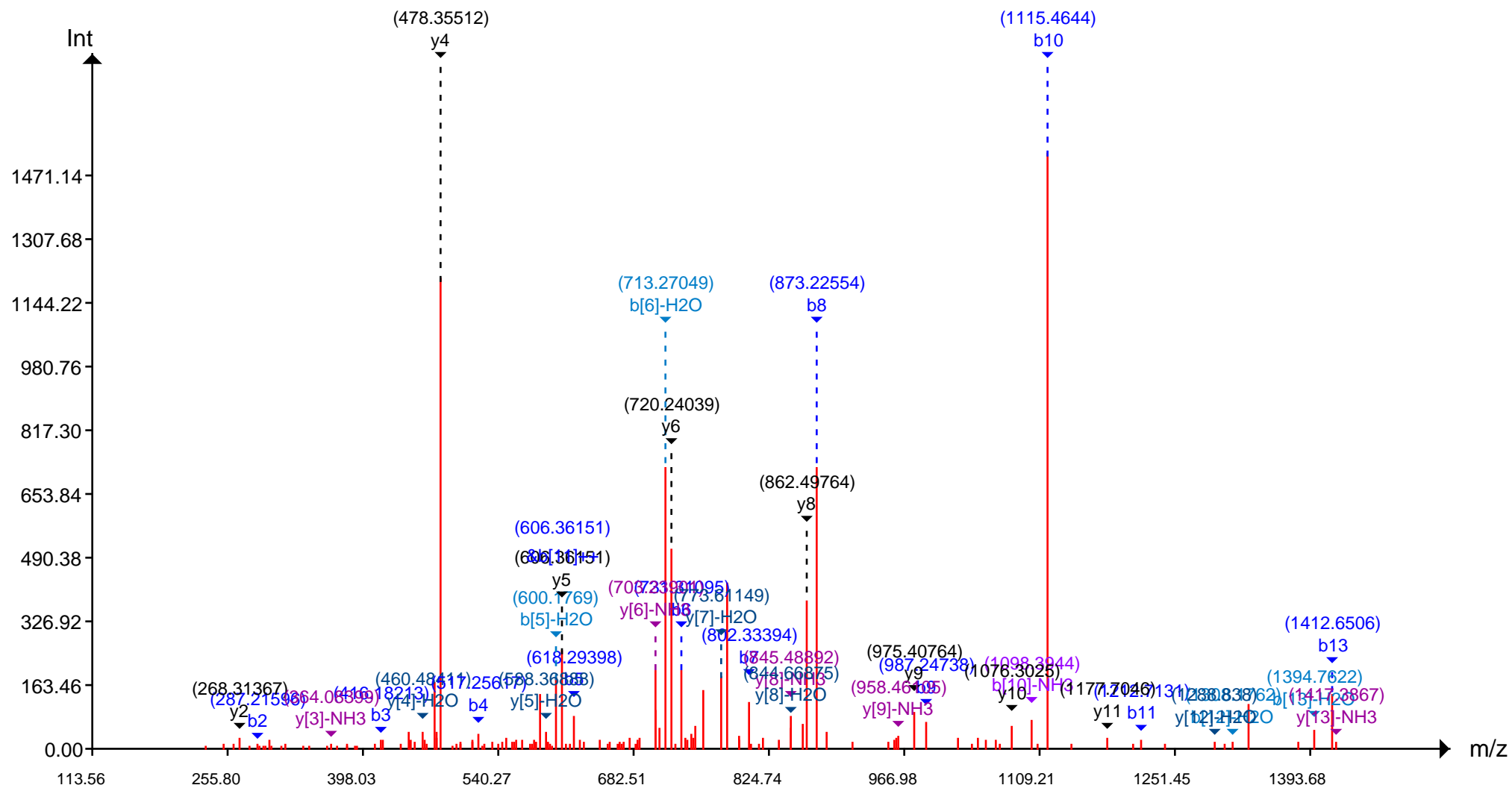
Confidence level 99.00%

Precursor m/z 548.79211

Identity Threshold 47.54

Charge 2+

Mass error 1.64E-4



Info

Modified Sequence AcD3-LQETTLAANQPLSR<C13>-COOH

Ion score 73.29

Protein O09159

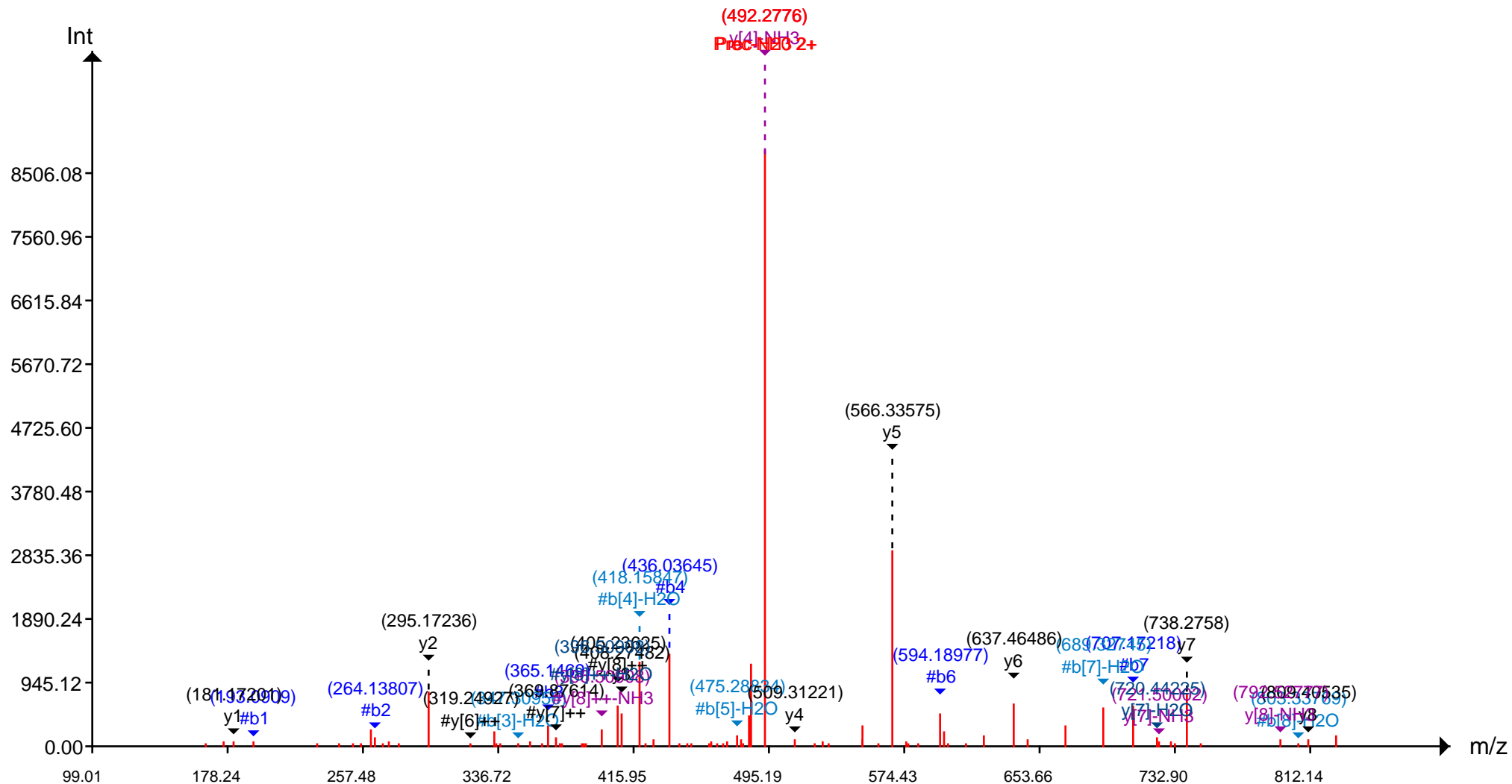
Confidence level 99.00%

Precursor m/z 796.94356

Identity Threshold 47.35

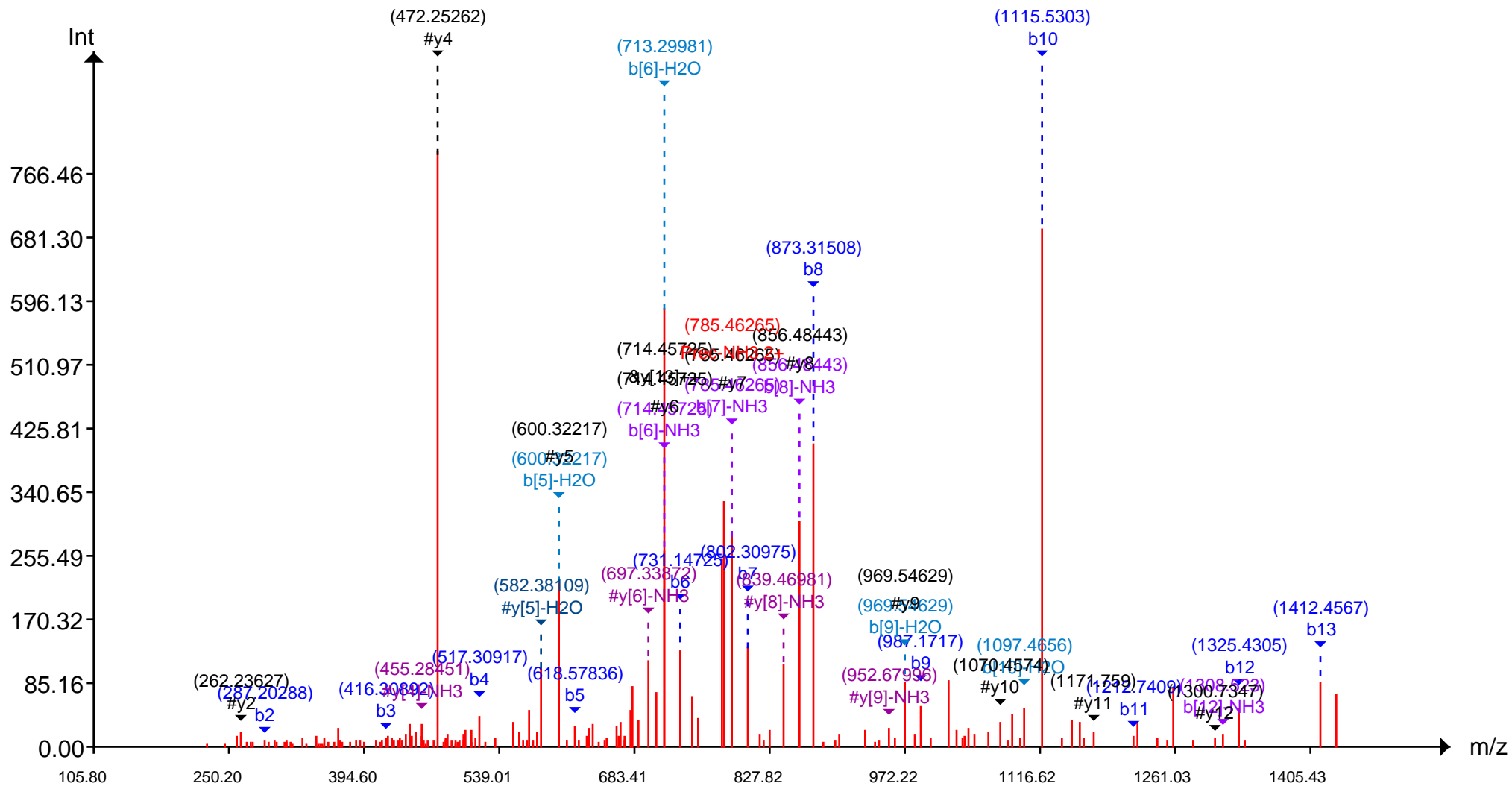
Charge 2+

Mass error 0.002146



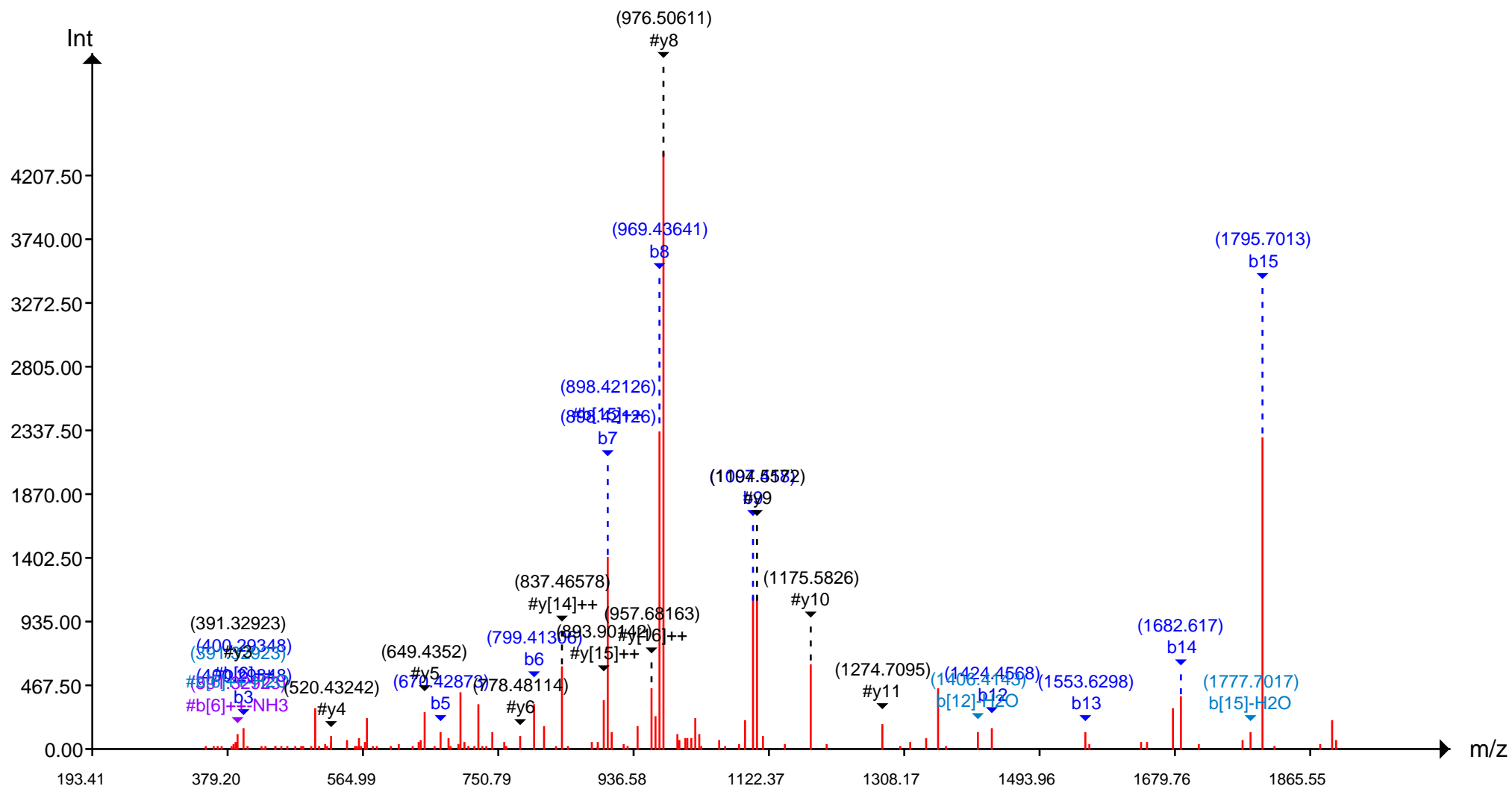
Info

Modified Sequence	AcD3-FATAGTLNR<C13>-COOH	Ion score	60.16
Protein	P26039	Confidence level	99.00%
Precursor m/z	501.28124	Identity Threshold	47.99
Charge	2+	Mass error	2.75E-4



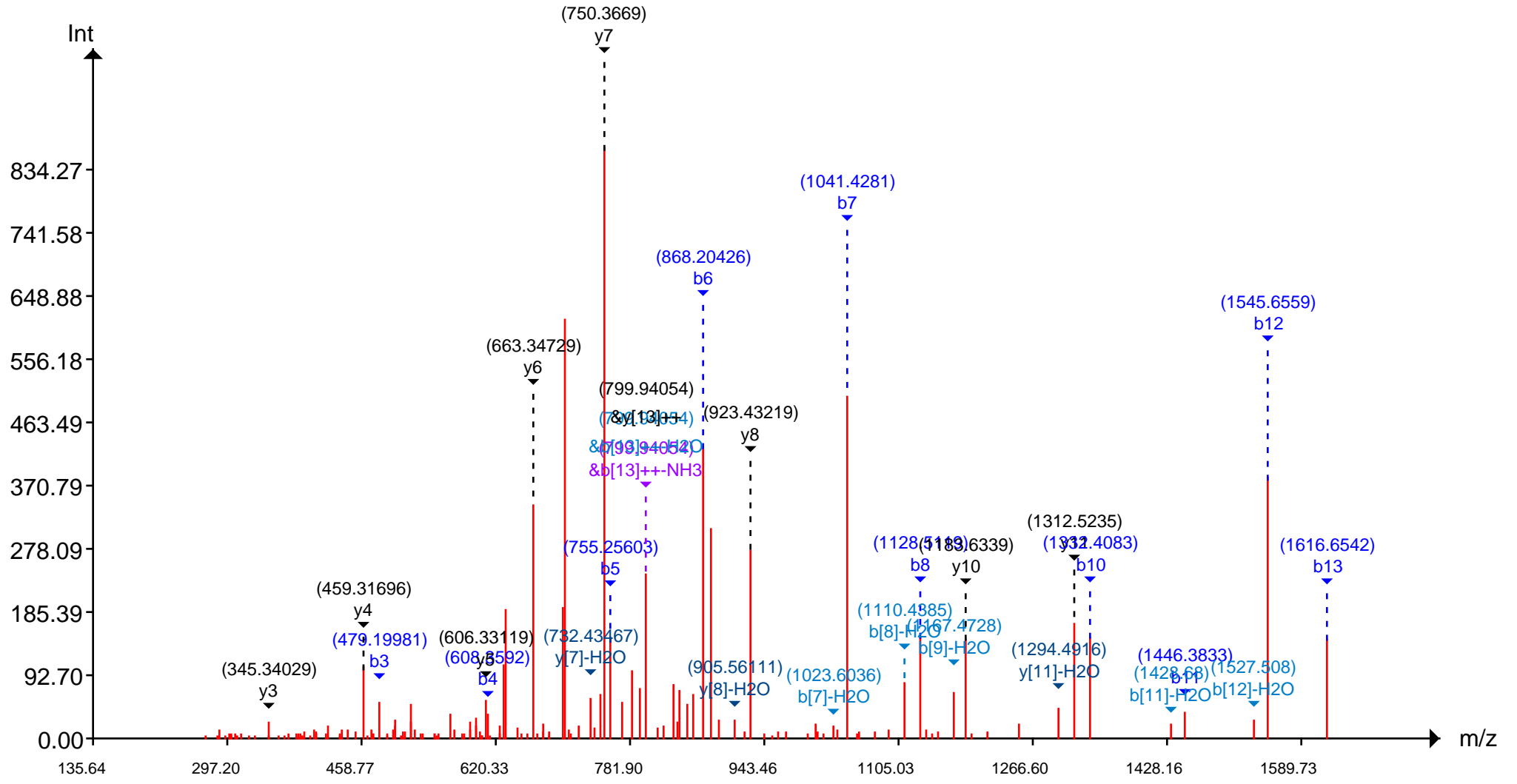
Info

Modified Sequence	AcD3-LQETTLAANQPLSR-COOH	Ion score	81.66
Protein	O09159	Confidence level	99.00%
Precursor m/z	793.93339	Identity Threshold	47.81
Charge	2+	Mass error	0.001947



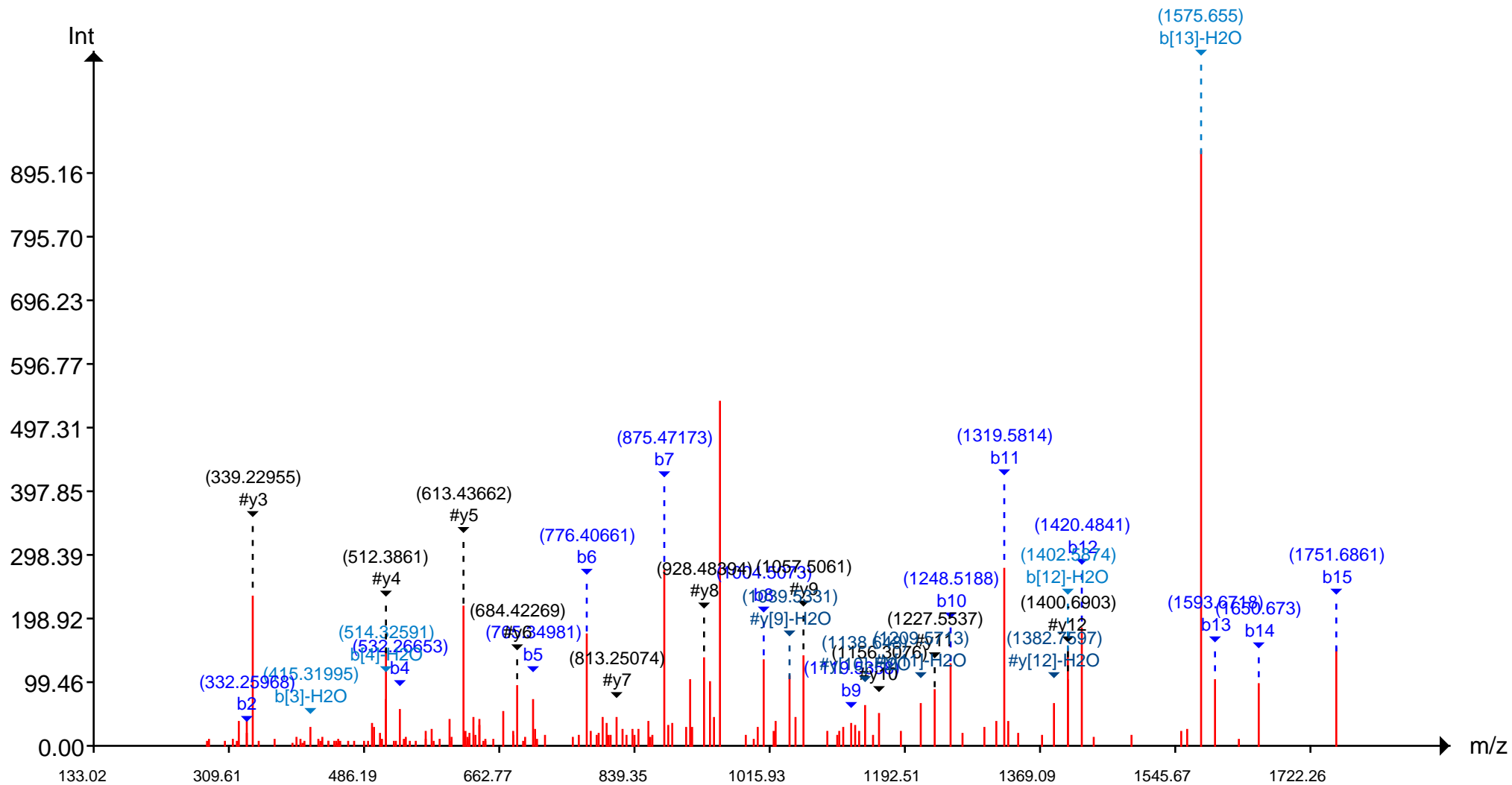
Info

Modified Sequence	AcD3-LQLPK<AcD3K*>EVAQPTEEELPR<C13>-COOH	Ion score	56.07
Protein	Q9ERD8	Confidence level	99.00%
Precursor m/z	1037.0769	Identity Threshold	46.54
Charge	2+	Mass error	0.002499



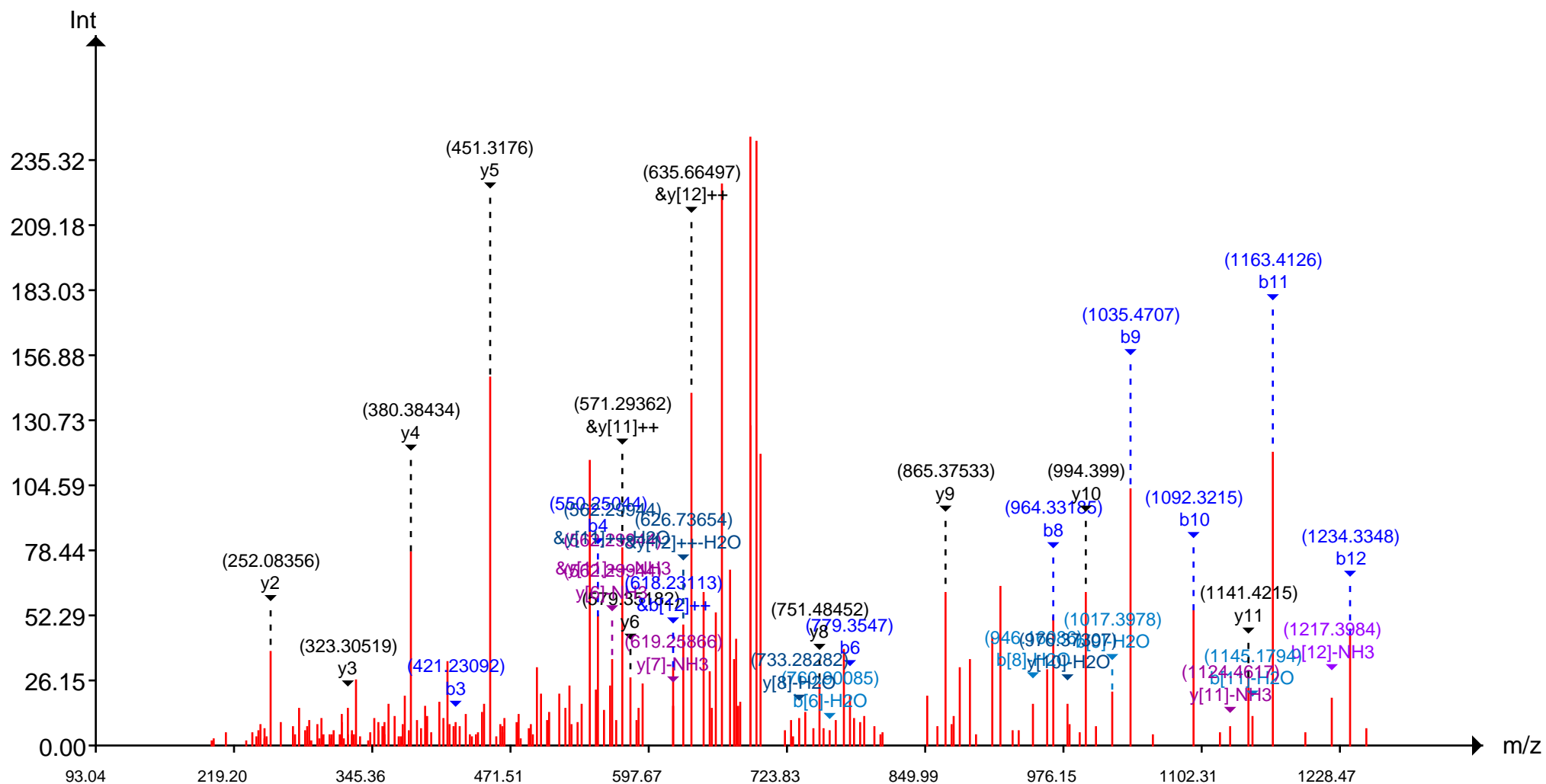
Info

Modified Sequence	AcD3-M<Mox*>LK<AcD3K*>EM<Mox*>IK<AcD3K*>SGM<Mox*>NVAR-COOH	Ion score	77.89
Protein	P52480	Confidence level	99.00%
Precursor m/z	895.96239	Identity Threshold	48.14
Charge	2+	Mass error	0.001598



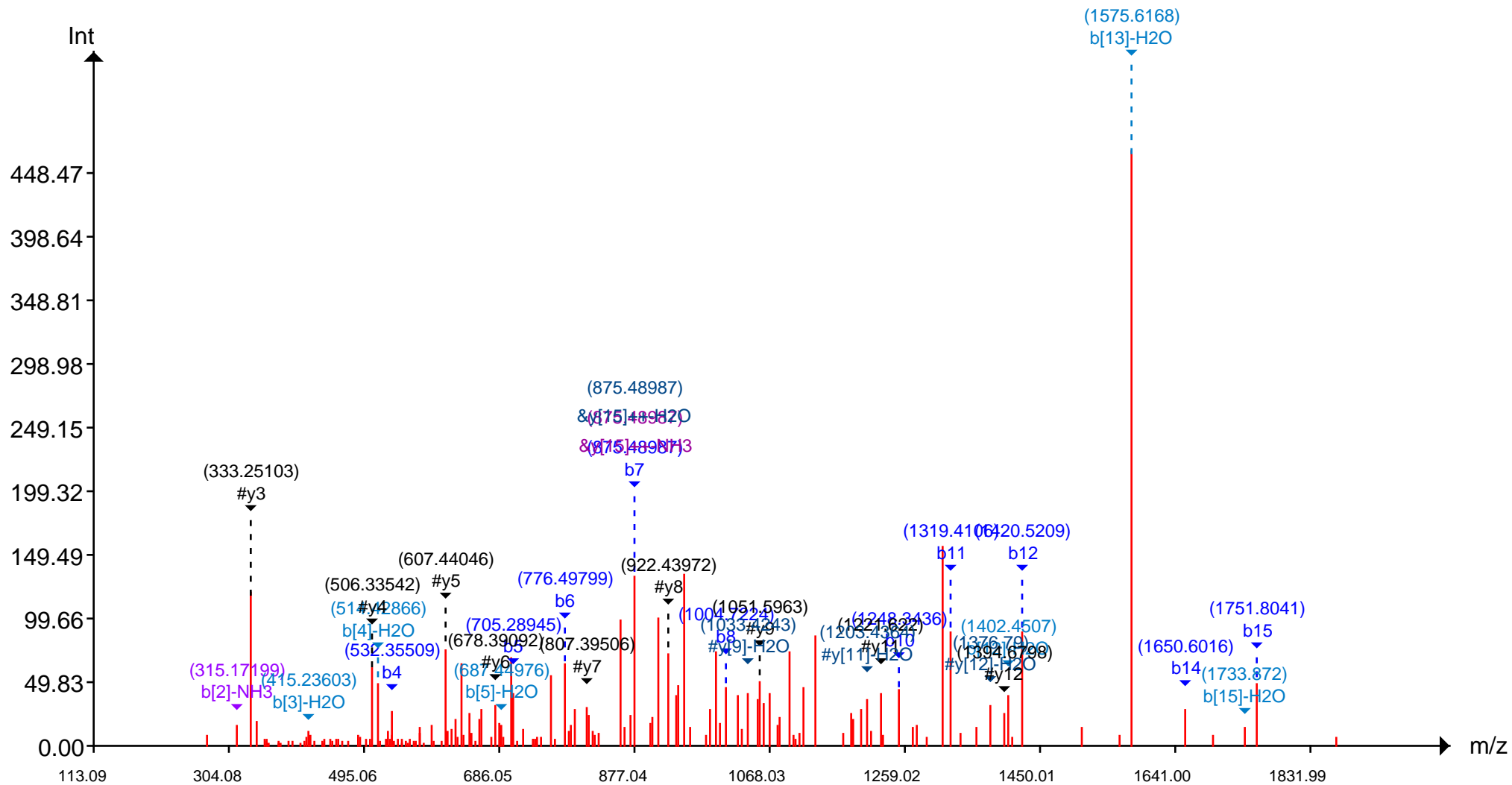
Info

Modified Sequence	AcD3-LK<AcD3K*>TVK<AcD3K*>AVEDEATK<AcD3K*>GTR<C13>-COOH	Ion score	106.18
Protein	P26039	Confidence level	99.00%
Precursor m/z	966.56146	Identity Threshold	44.42
Charge	2+	Mass error	0.002335



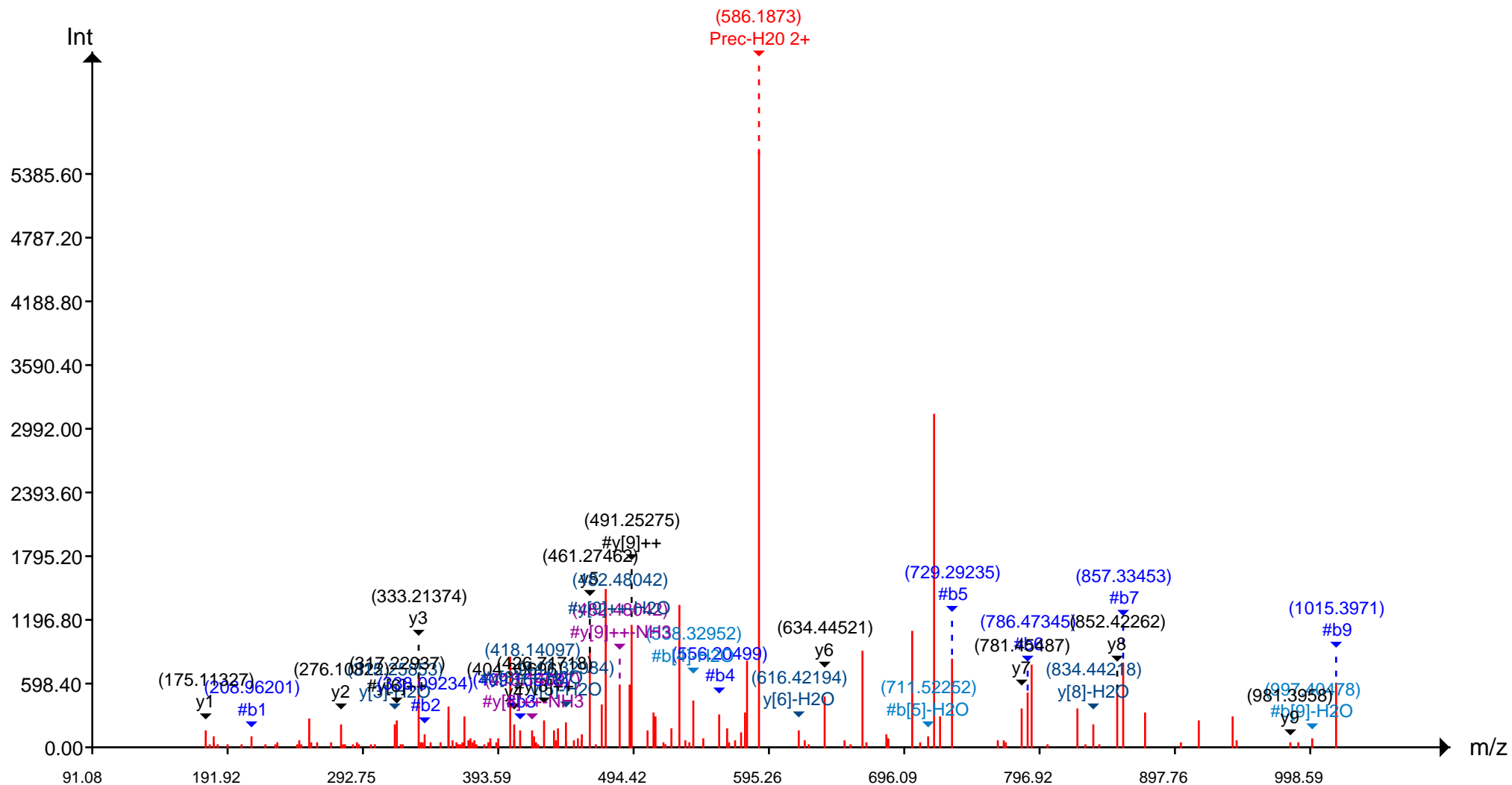
Info

Modified Sequence	AcD3-VEFENDGQAGAAR<C13>-COOH	Ion score	47.3
Protein	Q9CQI7	Confidence level	99.00%
Precursor m/z	707.84073	Identity Threshold	46.29
Charge	2+	Mass error	9.84E-4



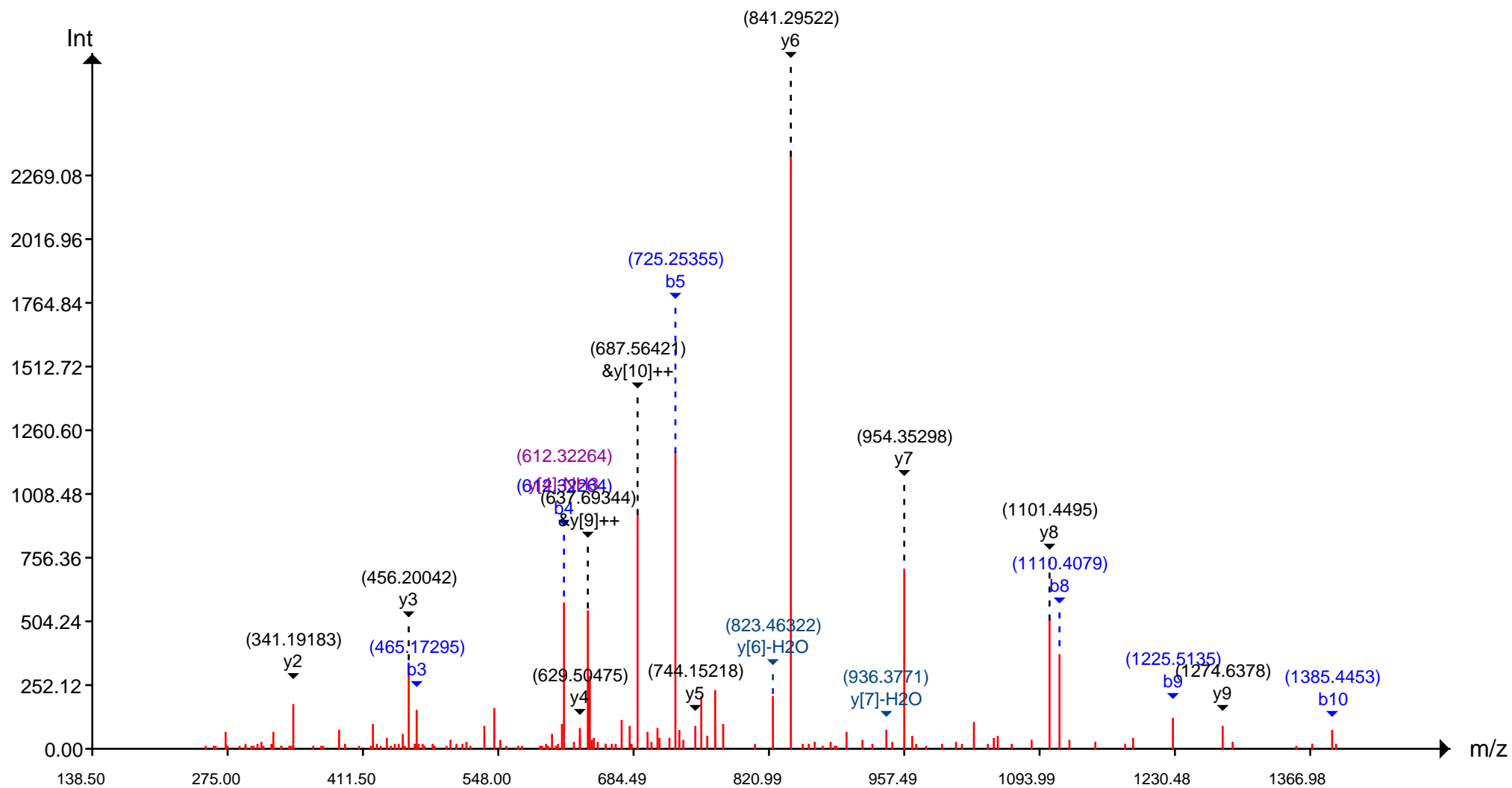
Info

Modified Sequence	AcD3-LK<AcD3K*>TVK<AcD3K*>AVEDEATK<AcD3K*>GTR-COOH	Ion score	61.13
Protein	P26039	Confidence level	99.00%
Precursor m/z	963.55097	Identity Threshold	45.40
Charge	2+	Mass error	0.001496



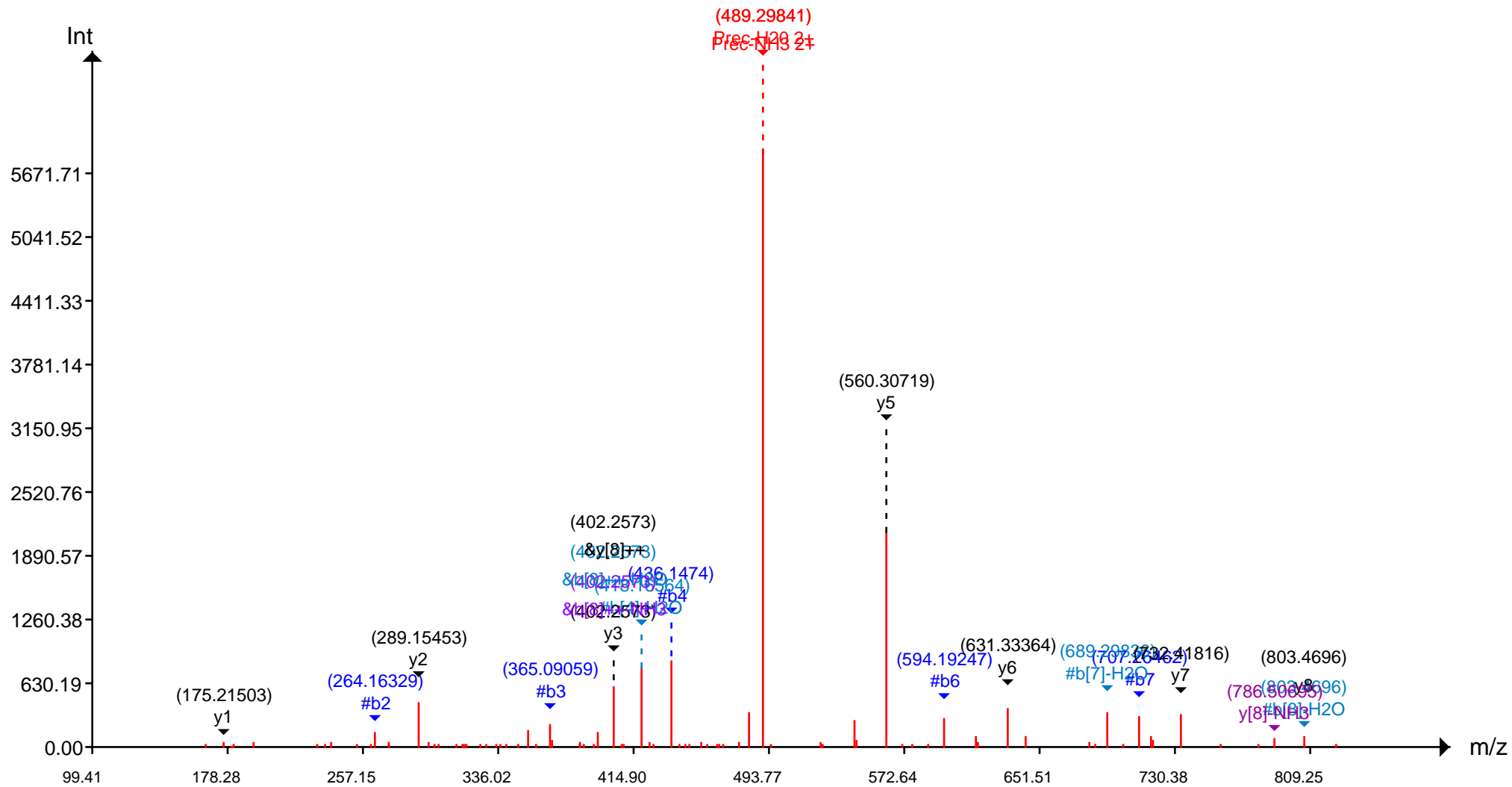
Info

Modified Sequence	AcD3-YEAM<Mox*>K<AcD3K*>GAGTR-COOH	Ion score	52.52
Protein	P10107	Confidence level	99.00%
Precursor m/z	595.29583	Identity Threshold	47.19
Charge	2+	Mass error	0.005544



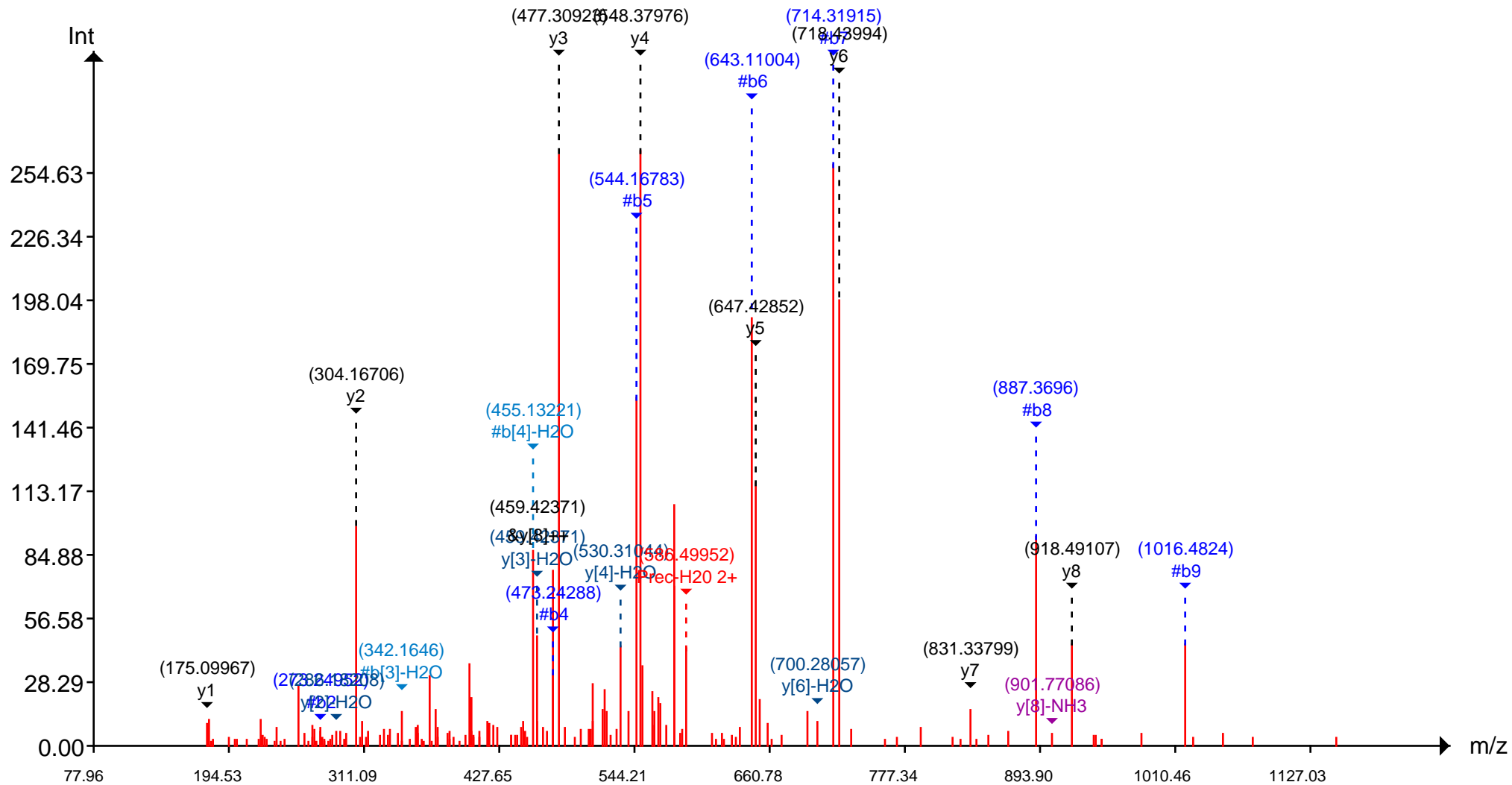
Info

Modified Sequence	AcD3-FVK<AcD3K*>M<Mox*>LPDK<AcD3K*>DC<Cmm*>R<C13>-COOH	Ion score	52.54
Protein	P18760	Confidence level	99.00%
Precursor m/z	783.41005	Identity Threshold	48.04
Charge	2+	Mass error	0.00202



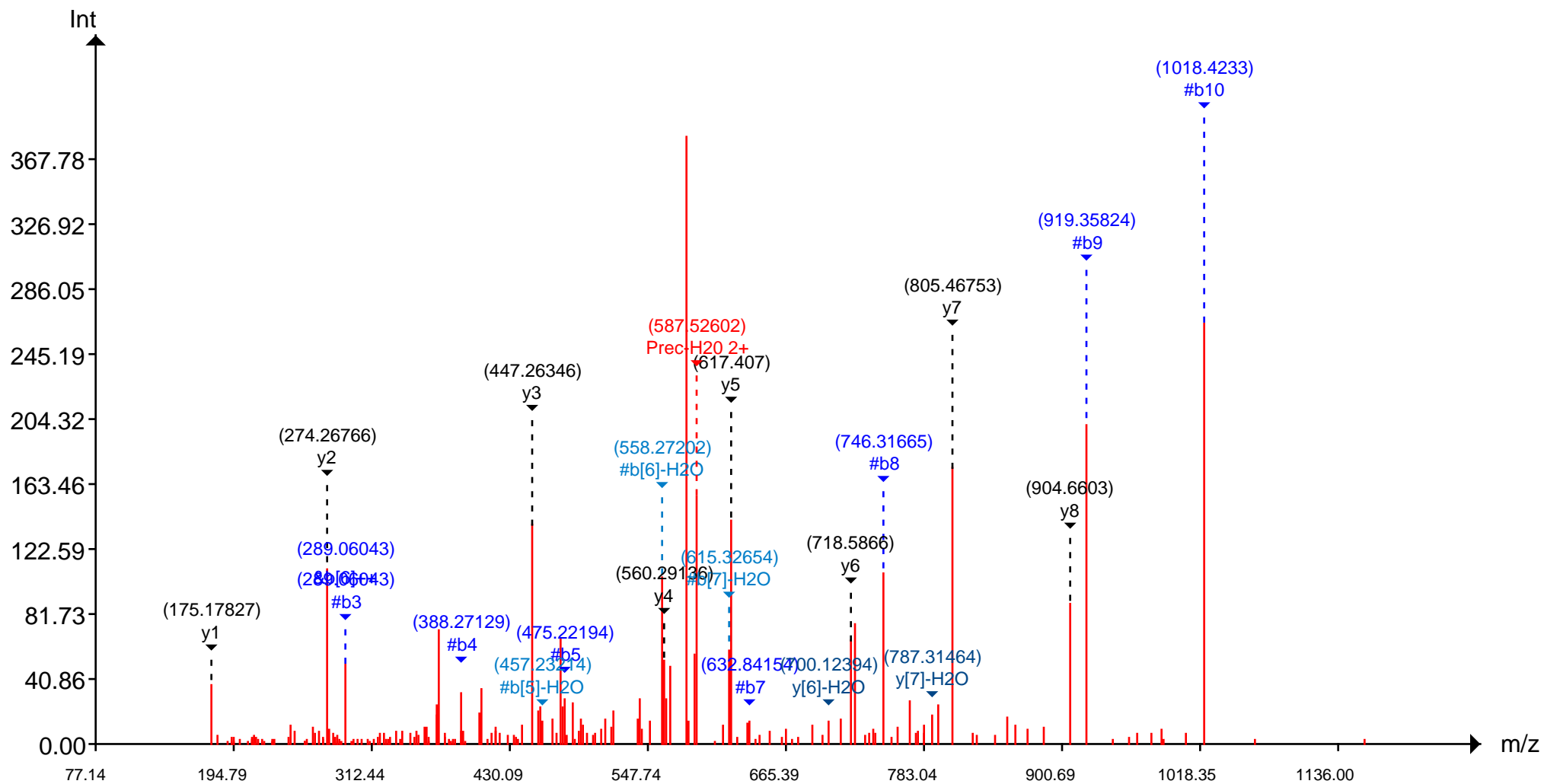
Info

Modified Sequence	AcD3-FATAGTLNR-COOH	Ion score	67.16
Protein	P26039	Confidence level	99.00%
Precursor m/z	498.27128	Identity Threshold	47.68
Charge	2+	Mass error	4.96E-4



Info

Modified Sequence	AcD3-NISLAVAK<AcD3K*>ER-COOH	Ion score	52.72
Protein	P70441	Confidence level	99.00%
Precursor m/z	595.85444	Identity Threshold	45.80
Charge	2+	Mass error	6.18E-4



Info

Modified Sequence AcD3-SGVVSTGLK<AcD3K*>VR-COOH

Ion score 57.36

Protein P58252

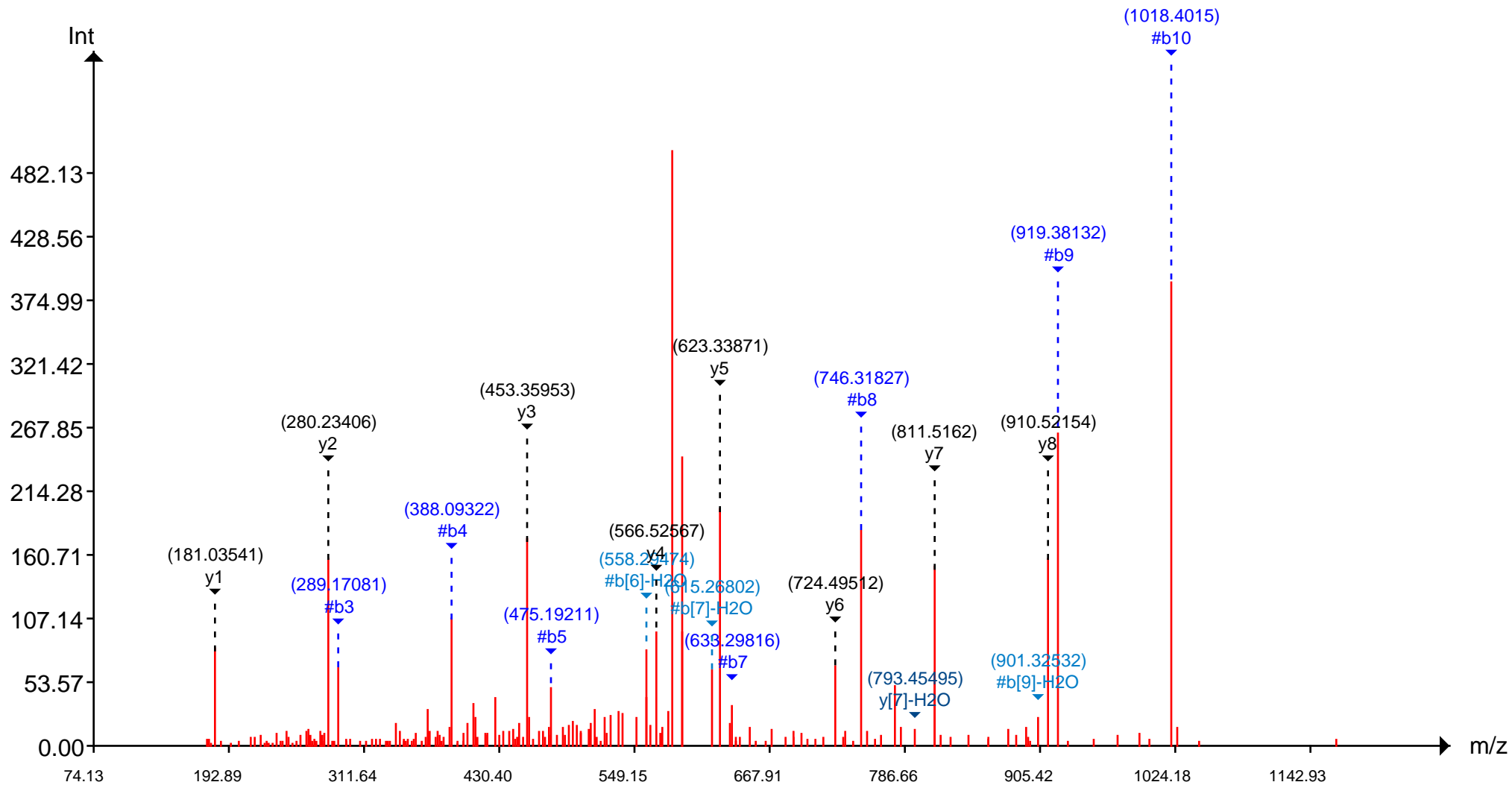
Confidence level 99.00%

Precursor m/z 596.86166

Identity Threshold 44.68

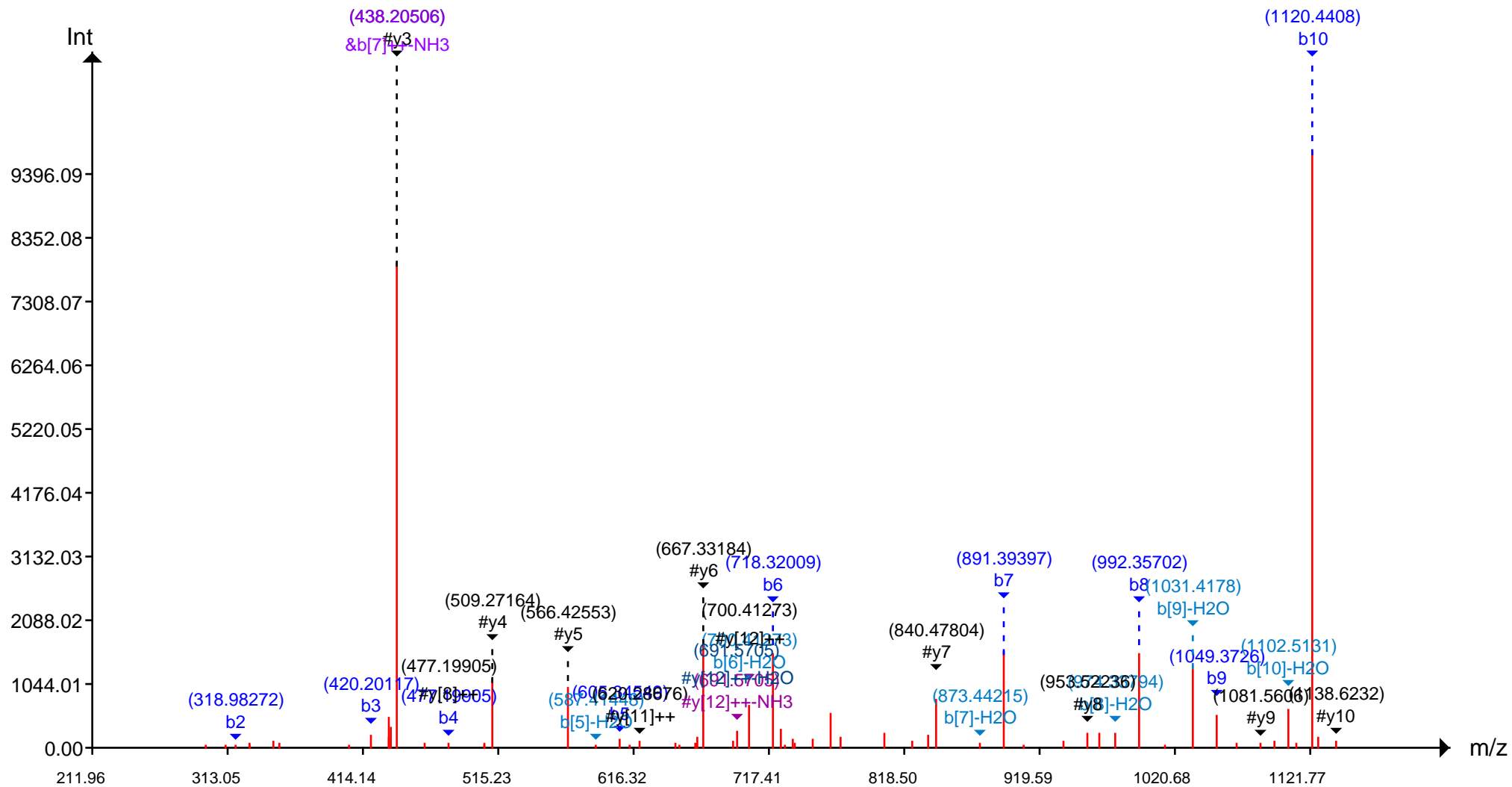
Charge 2+

Mass error -6.44E-4



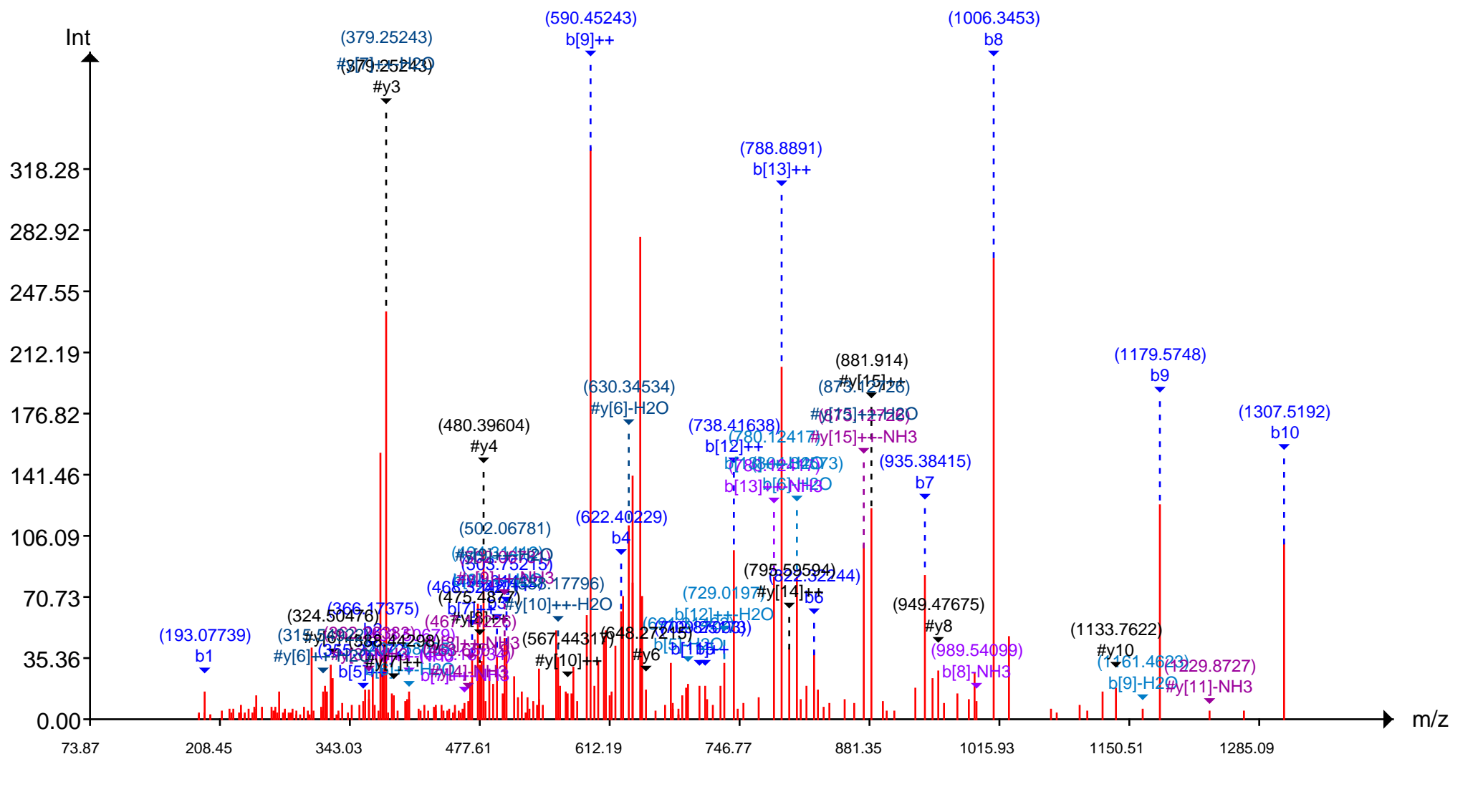
Info

Modified Sequence	AcD3-SGVVSTGLK<AcD3K*>VR<C13>-COOH	Ion score	61.39
Protein	P58252	Confidence level	99.00%
Precursor m/z	599.87195	Identity Threshold	42.67
Charge	2+	Mass error	-2.05E-4

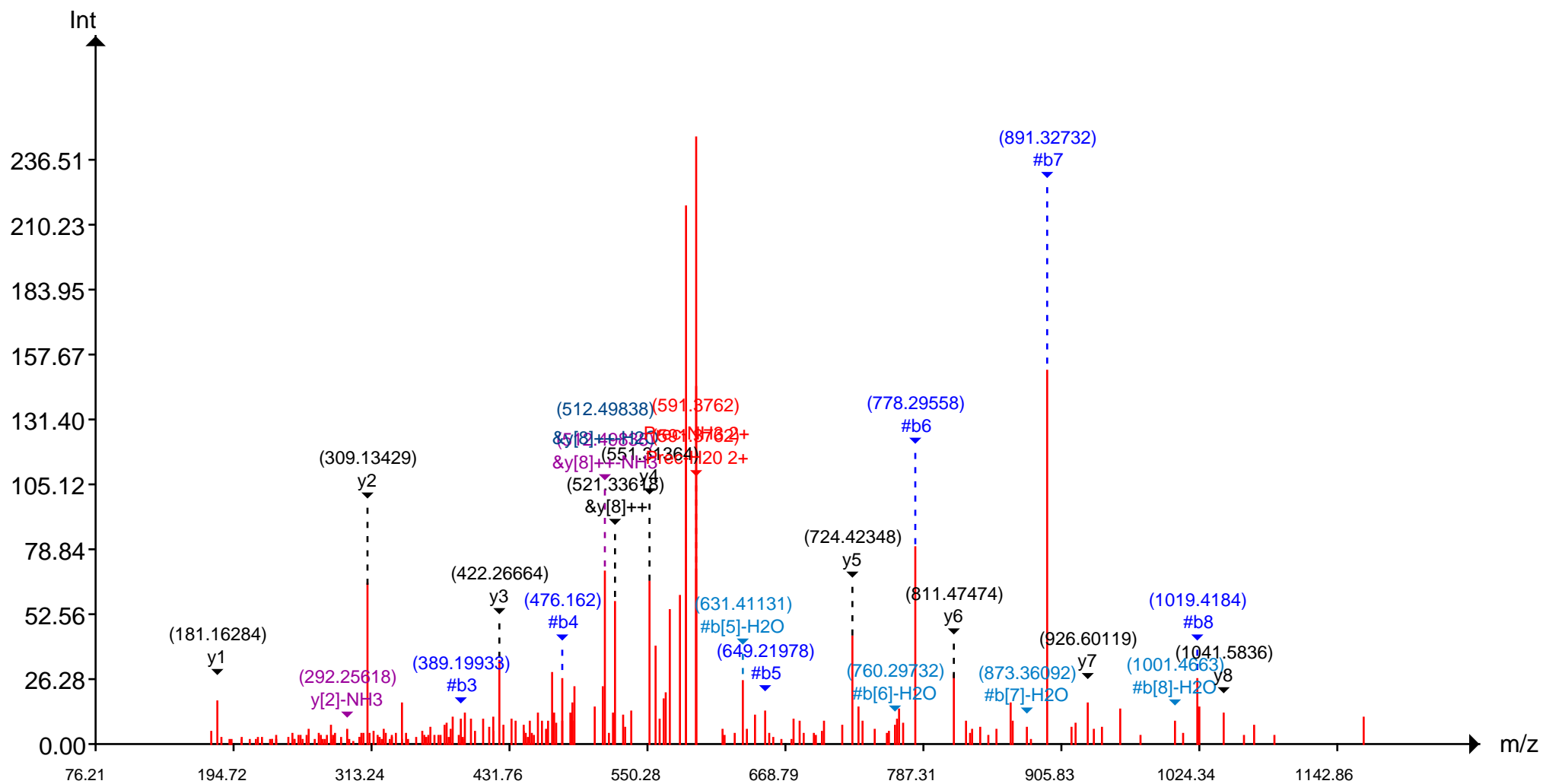


Info

Modified Sequence	AcD3-LC<Cmm*>TGQIK<AcD3K*>TGAPC<Cmm*>R<C13>-COOH	Ion score	55.18
Protein	P17182, P21550, P17183	Confidence level	99.00%
Precursor m/z	779.40842	Identity Threshold	48.08
Charge	2+	Mass error	5.45E-4

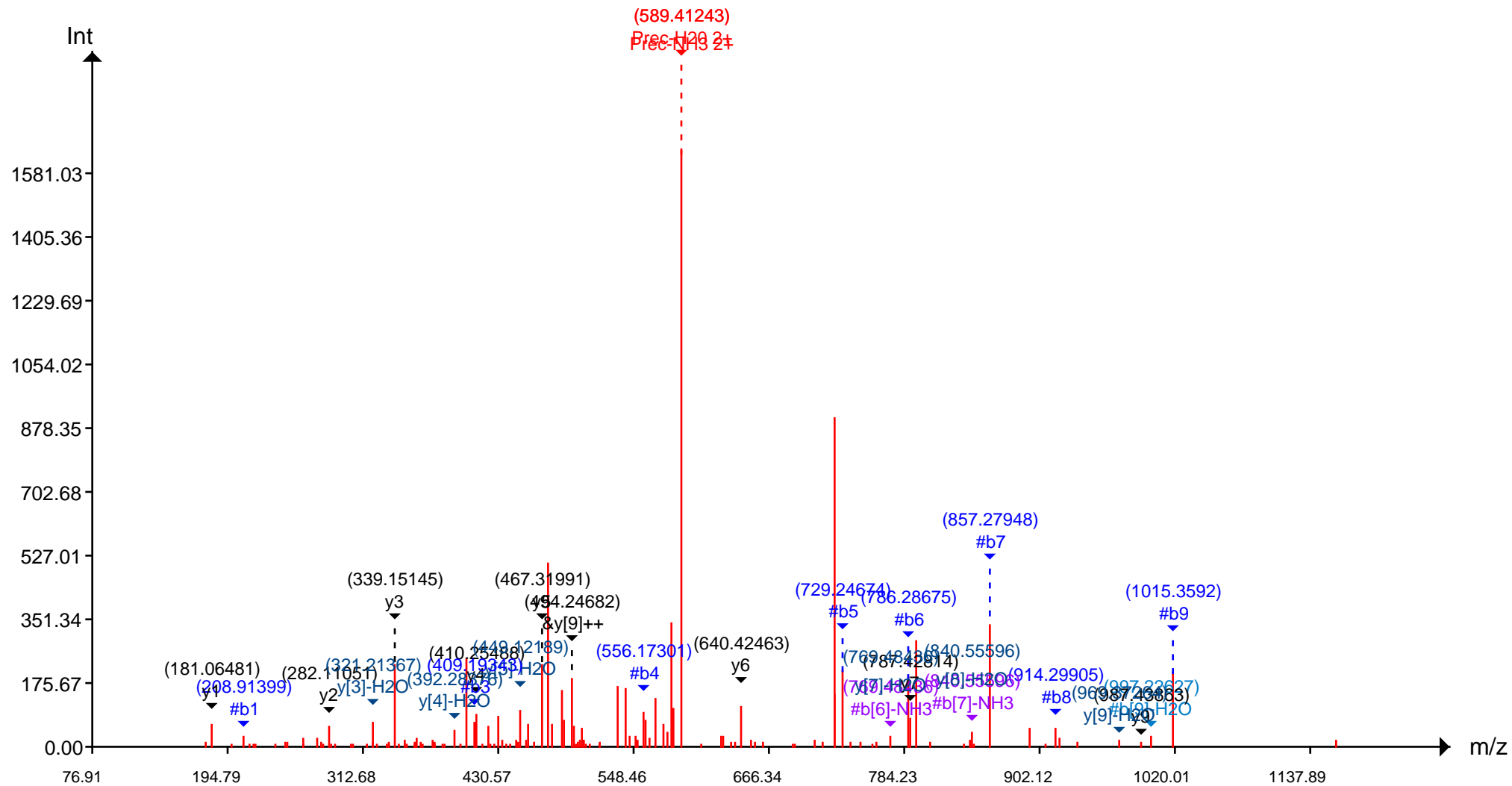


Info			
Modified Sequence	AcD3-M<Mox*>K<AcD3K*>QQSLLAK<AcD3K*>QPATPTR<C13>-COOH	Ion score	46.7
Protein	Q6PDL0	Confidence level	99.00%
Precursor m/z	652.37365	Identity Threshold	45.57
Charge	3+	Mass error	0.002809



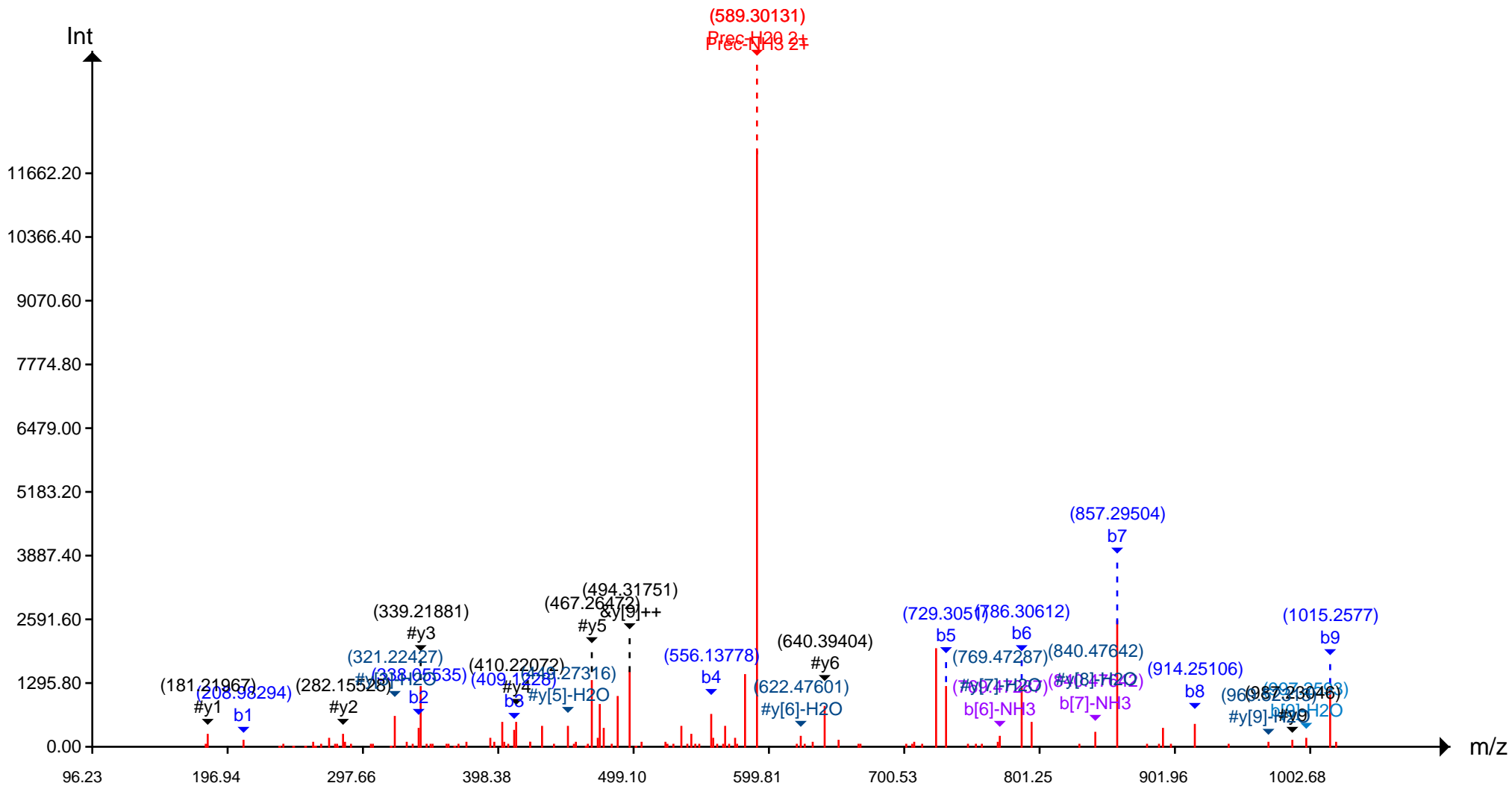
Info

Modified Sequence	AcD3-LDDSK<AcD3K*>ELQR<C13>-COOH	Ion score	48.42
Protein	Q7TQI3	Confidence level	99.00%
Precursor m/z	600.32786	Identity Threshold	47.67
Charge	2+	Mass error	4.06E-4



Info

Modified Sequence	AcD3-YEAM<Mox*>K<AcD3K*>GAGTR<C13>-COOH	Ion score	48.46
Protein	P10107	Confidence level	99.00%
Precursor m/z	598.30251	Identity Threshold	47.72
Charge	2+	Mass error	-0.001237



Info

Modified Sequence AcD3-YEAM<Mox*>K<AcD3K*>GAGTR<C13>-COOH

Ion score 58.97

Protein P10107

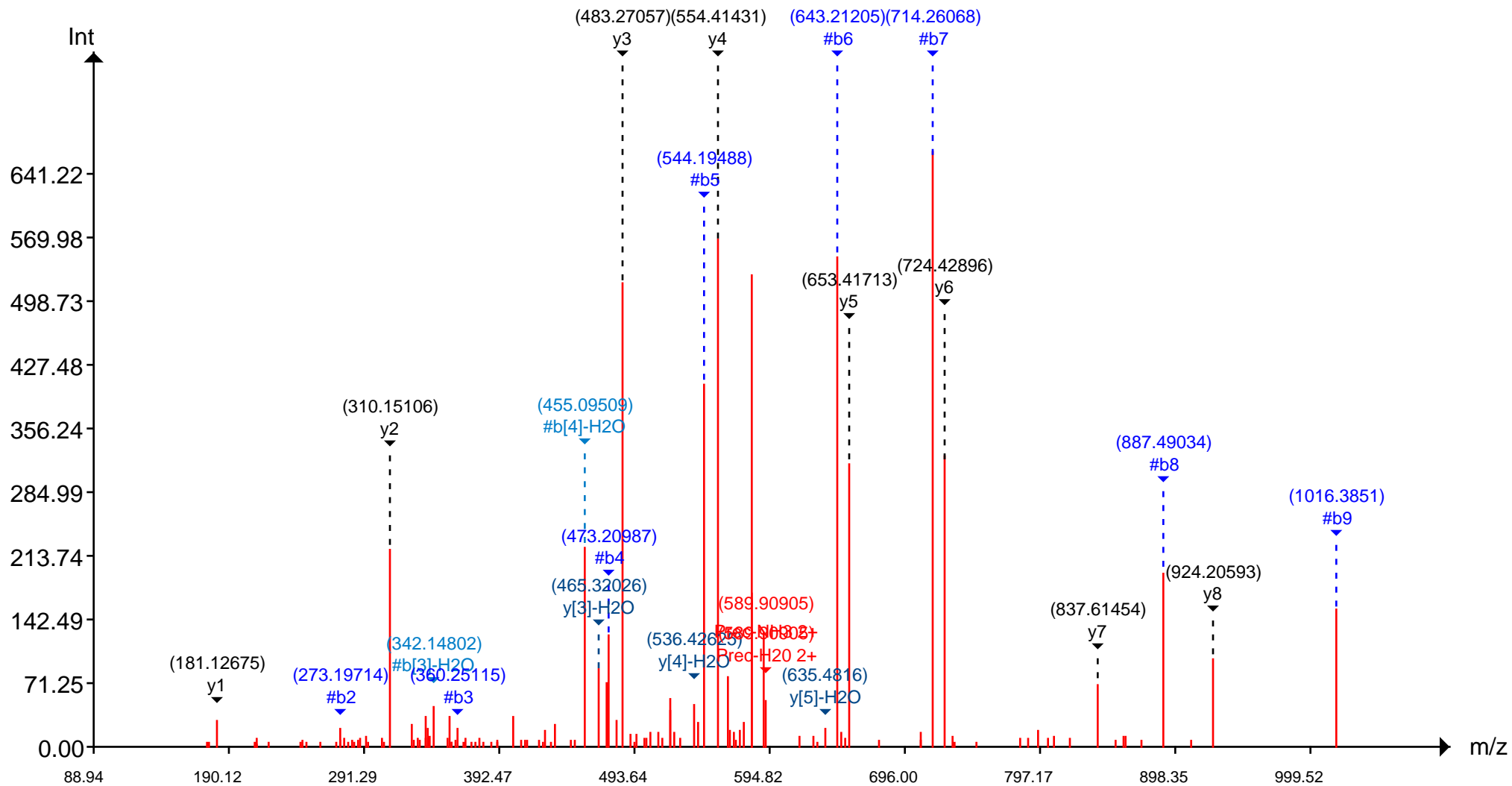
Confidence level 99.00%

Precursor m/z 598.30222

Identity Threshold 47.90

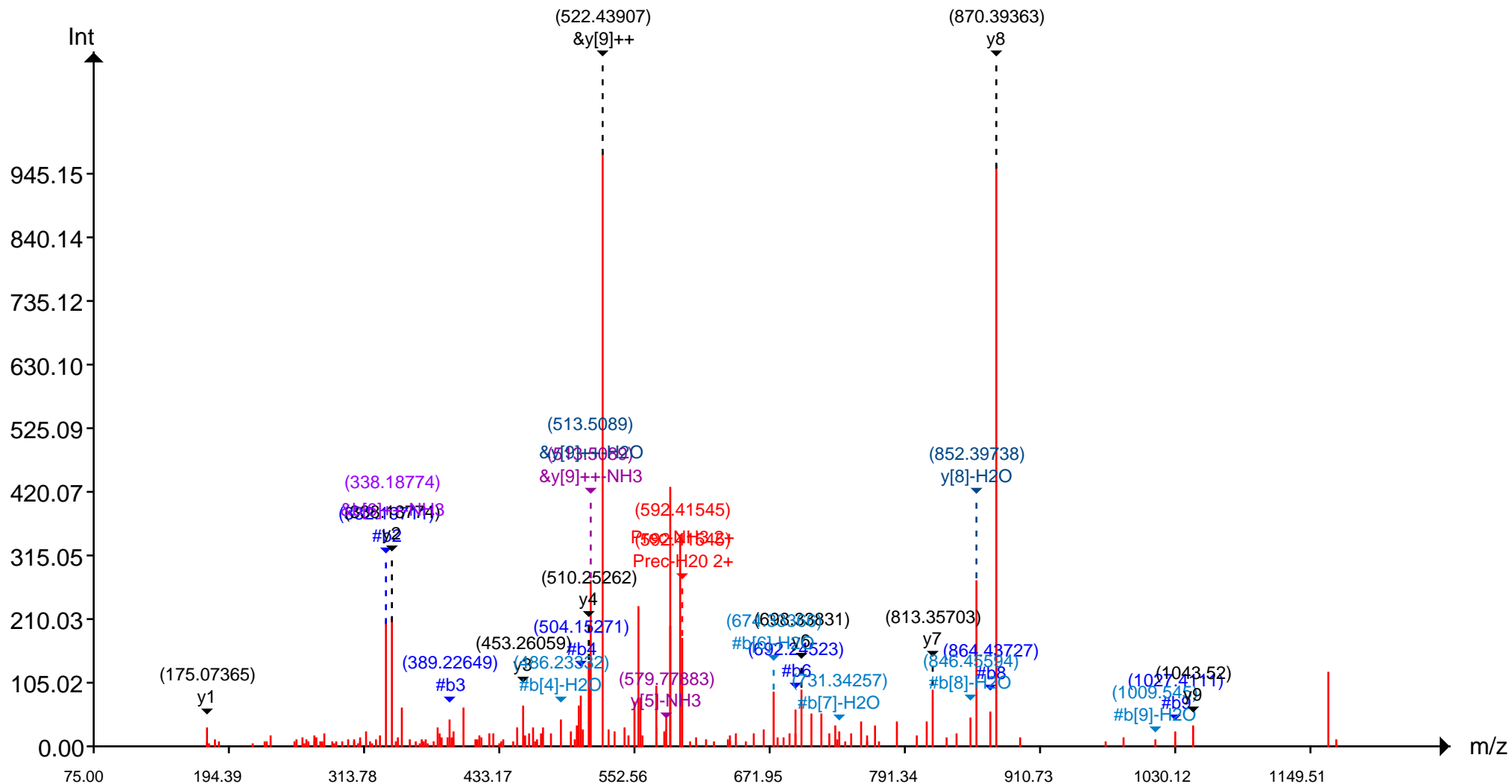
Charge 2+

Mass error -0.001817



Info

Modified Sequence	AcD3-NISLAVAK<AcD3K*>ER<C13>-COOH	Ion score	71.47
Protein	P70441	Confidence level	99.00%
Precursor m/z	598.86464	Identity Threshold	43.77
Charge	2+	Mass error	8.76E-4



Info

Modified Sequence AcD3-IK<AcD3K*>GDTSGDYR-COOH

Protein P97429

Precursor m/z 601.30221

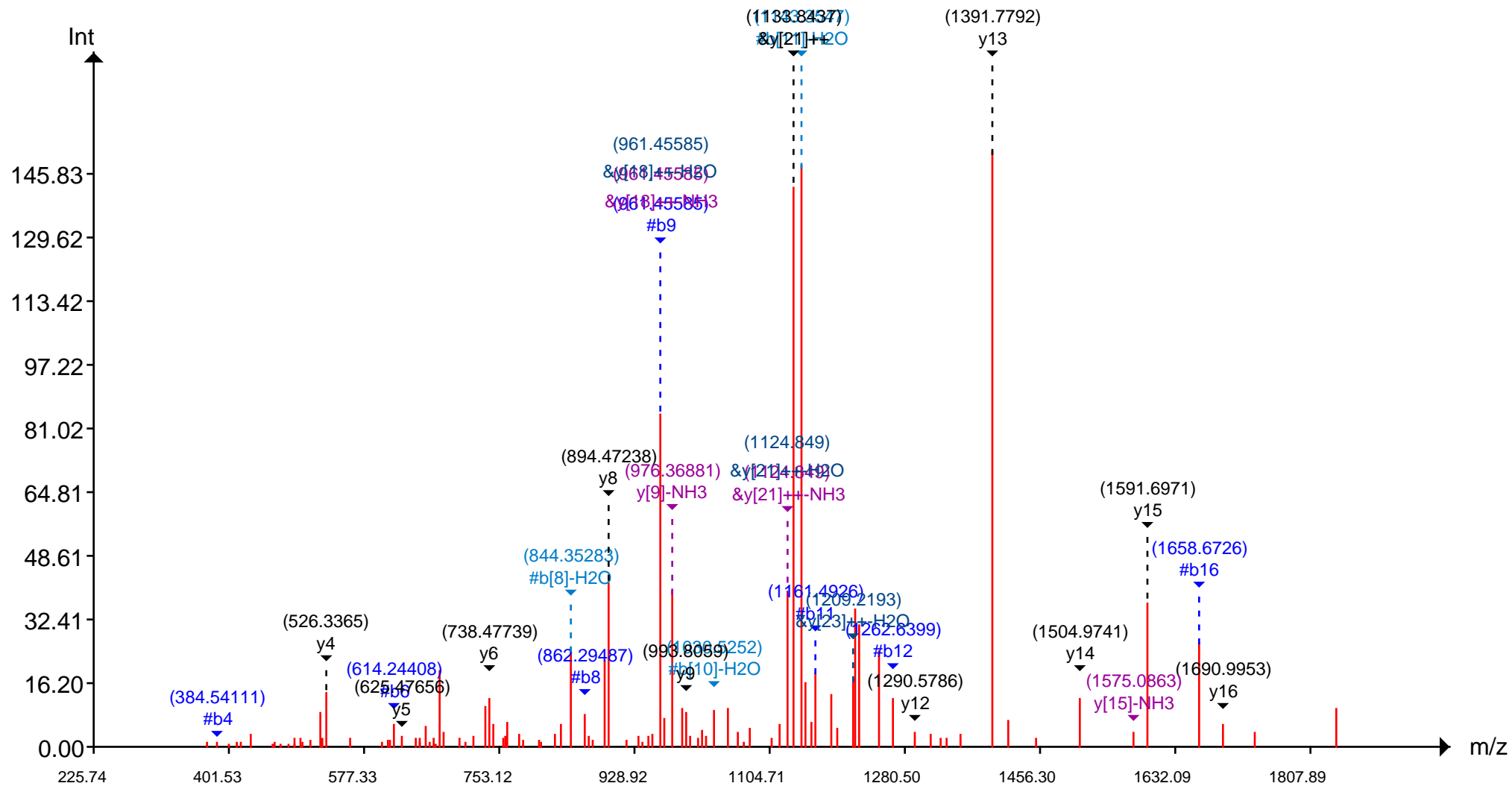
Charge 2+

Ion score 48.47

Confidence level 99.00%

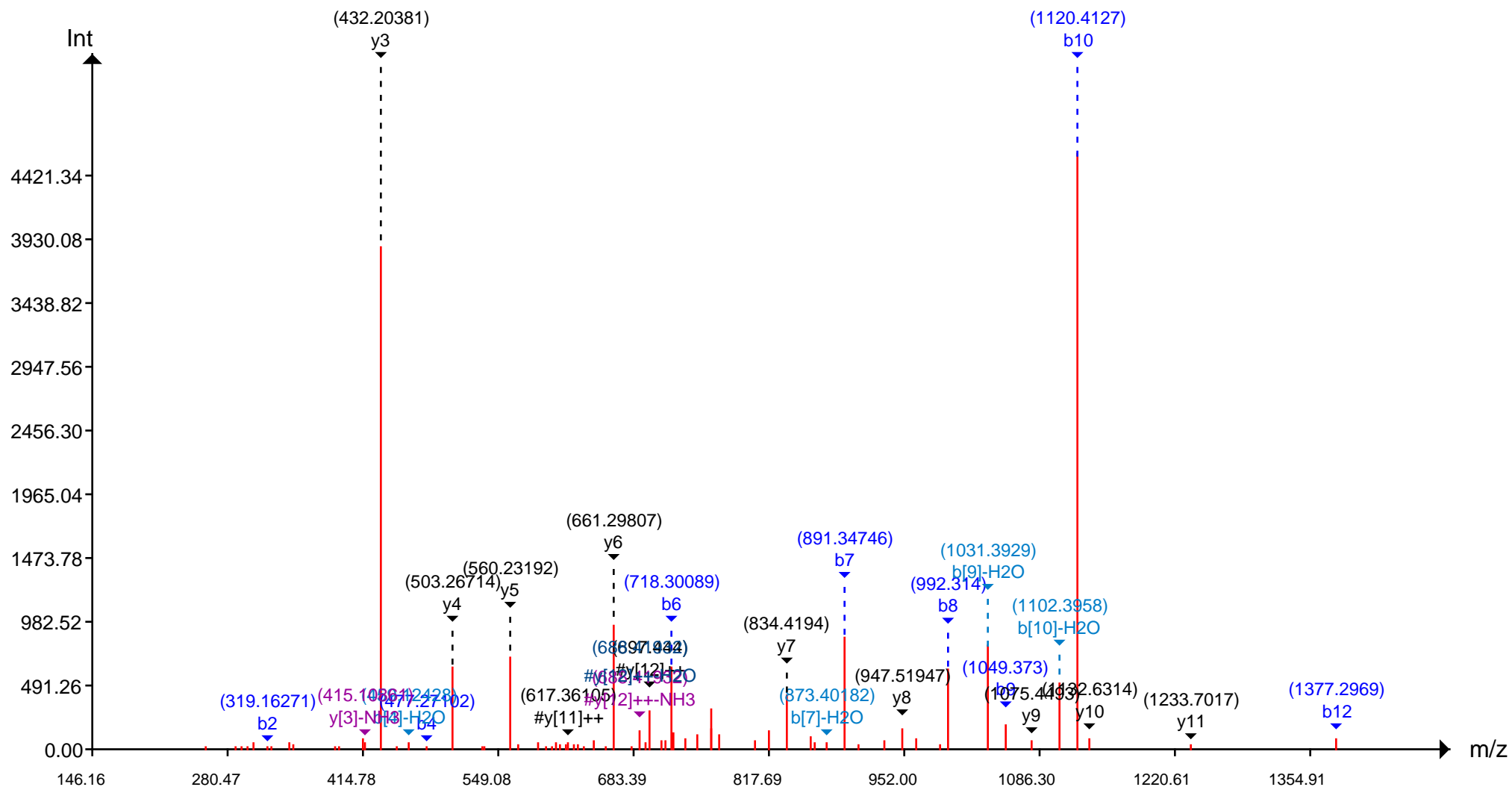
Identity Threshold 47.20

Mass error 5.28E-4



Info

Modified Sequence	AcD3-AAVPGK<AcD3K*>TFVSITPAEVGVLVGK<AcD3K*>DR<C13>-COOH	Ion score	42.78
Protein	P62962	Confidence level	99.00%
Precursor m/z	1276.7423	Identity Threshold	42.20
Charge	2+	Mass error	0.003297



Info

Modified Sequence AcD3-LC<Cmm*>TGQIK<AcD3K*>TGAPC<Cmm*>R-COOH

Ion score 75.94

Protein P17182, P21550, P17183

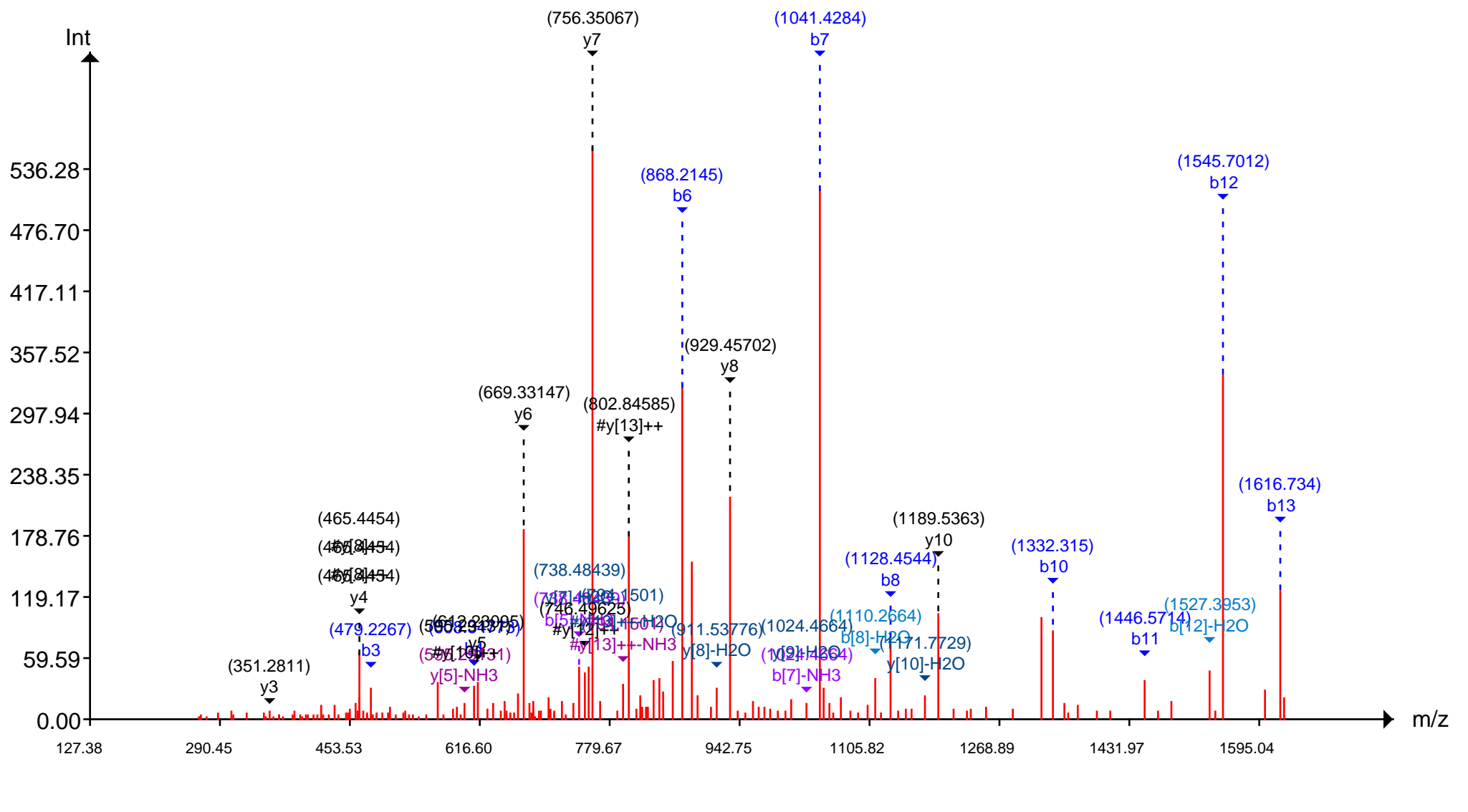
Confidence level 99.00%

Precursor m/z 776.39876

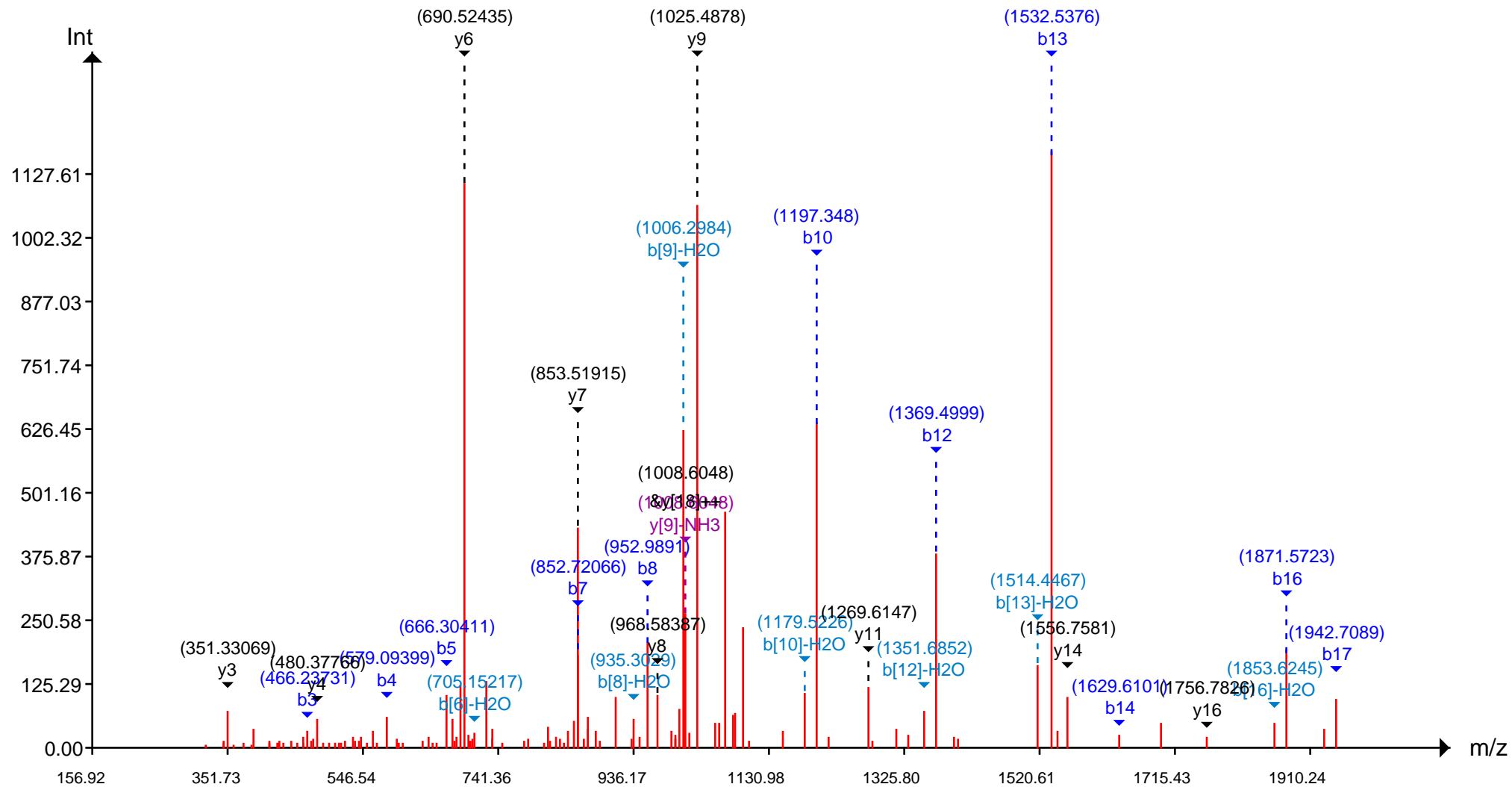
Identity Threshold 47.95

Charge 2+

Mass error 0.001367

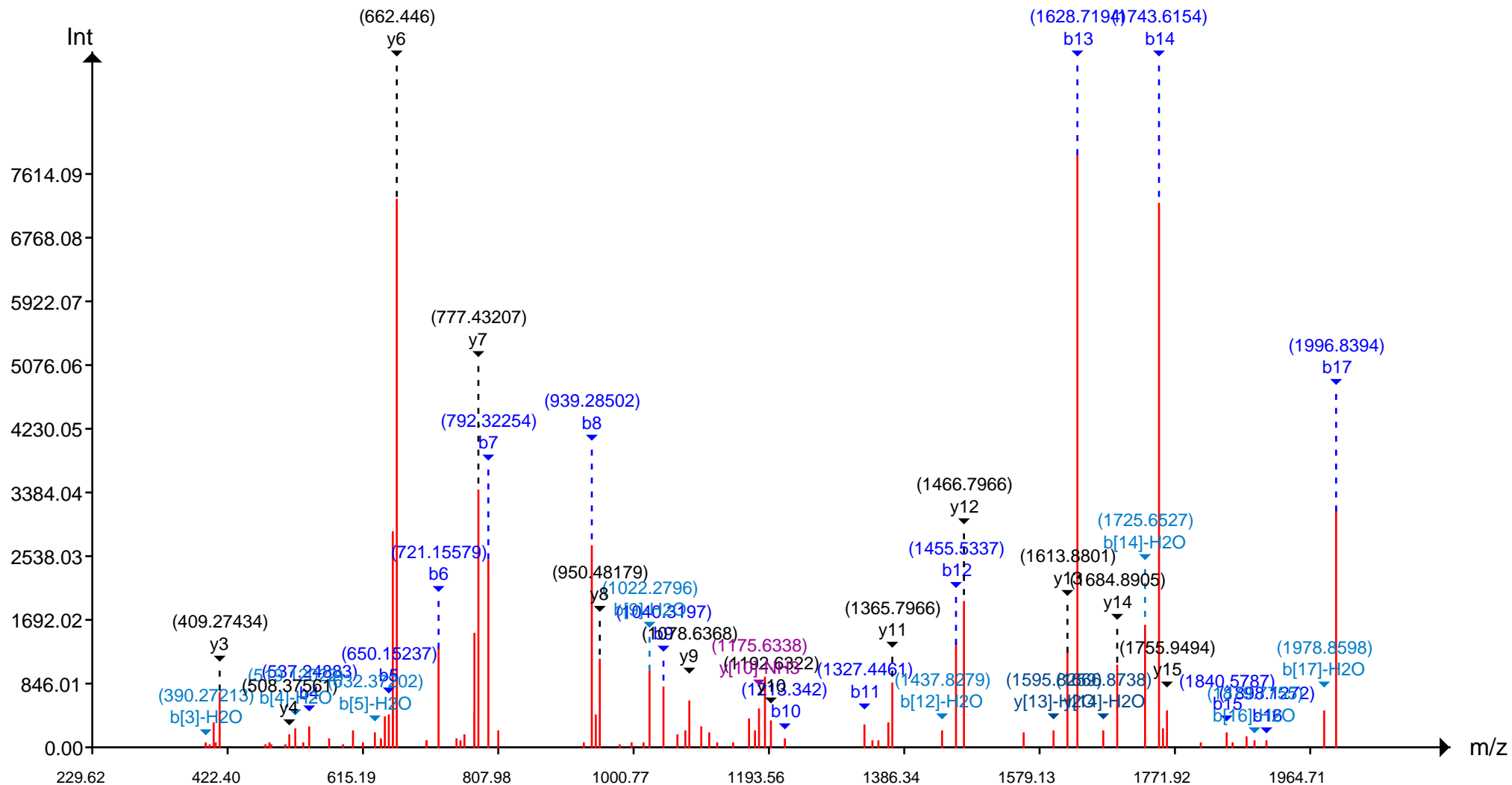


Info			
Modified Sequence	AcD3-M<Mox*>LK<AcD3K*>EM<Mox*>IK<AcD3K*>SGM<Mox*>NVAR<C13>-COOH	Ion score	60.1
Protein	P52480	Confidence level	99.00%
Precursor m/z	898.97238	Identity Threshold	48.03
Charge	2+	Mass error	0.001436



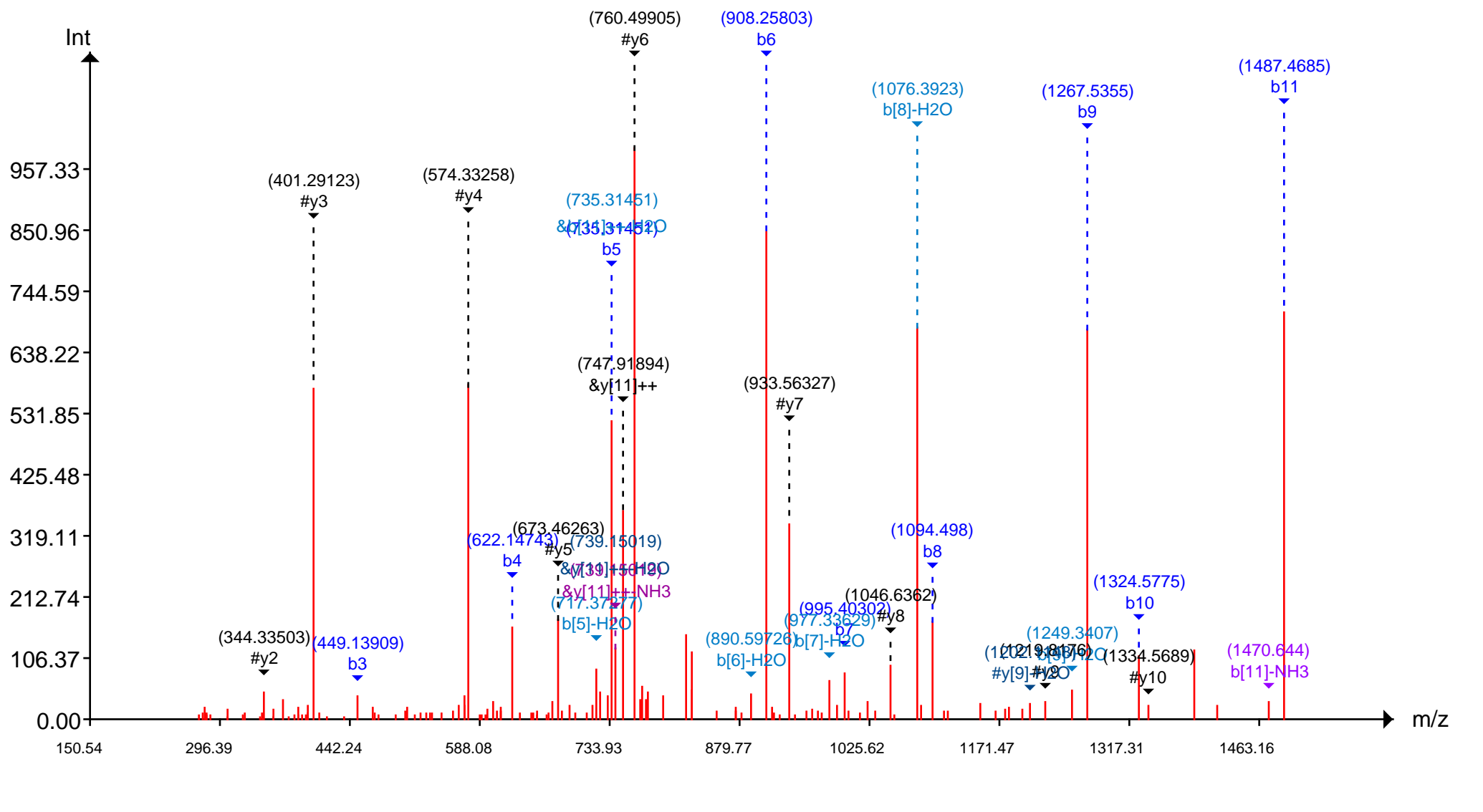
Info

Modified Sequence	AcD3-C<Cmm*>IM<Mox*>LSGETAK<AcD3K*>GDYPLEAVR<C13>-COOH	Ion score	71.61
Protein	P52480	Confidence level	99.00%
Precursor m/z	1111.5577	Identity Threshold	47.73
Charge	2+	Mass error	0.003558

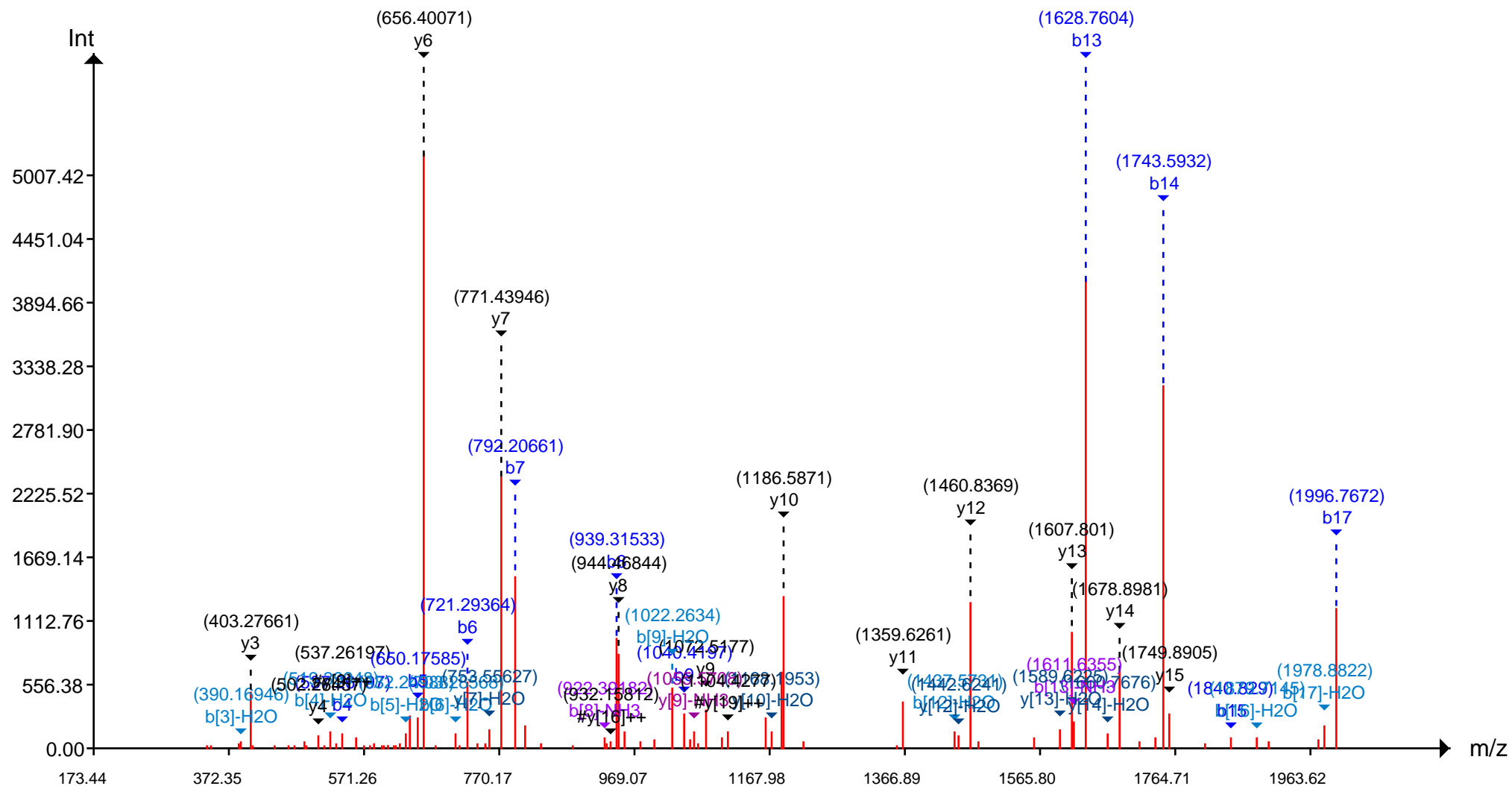


Info

Modified Sequence	AcD3-M<Mox*>NTELAFTK<AcD3K*>NQK<AcD3K*>DPGVLDR<C13>-COOH	Ion score	109.27
Protein	P50543	Confidence level	99.00%
Precursor m/z	1203.1261	Identity Threshold	47.86
Charge	2+	Mass error	0.002647

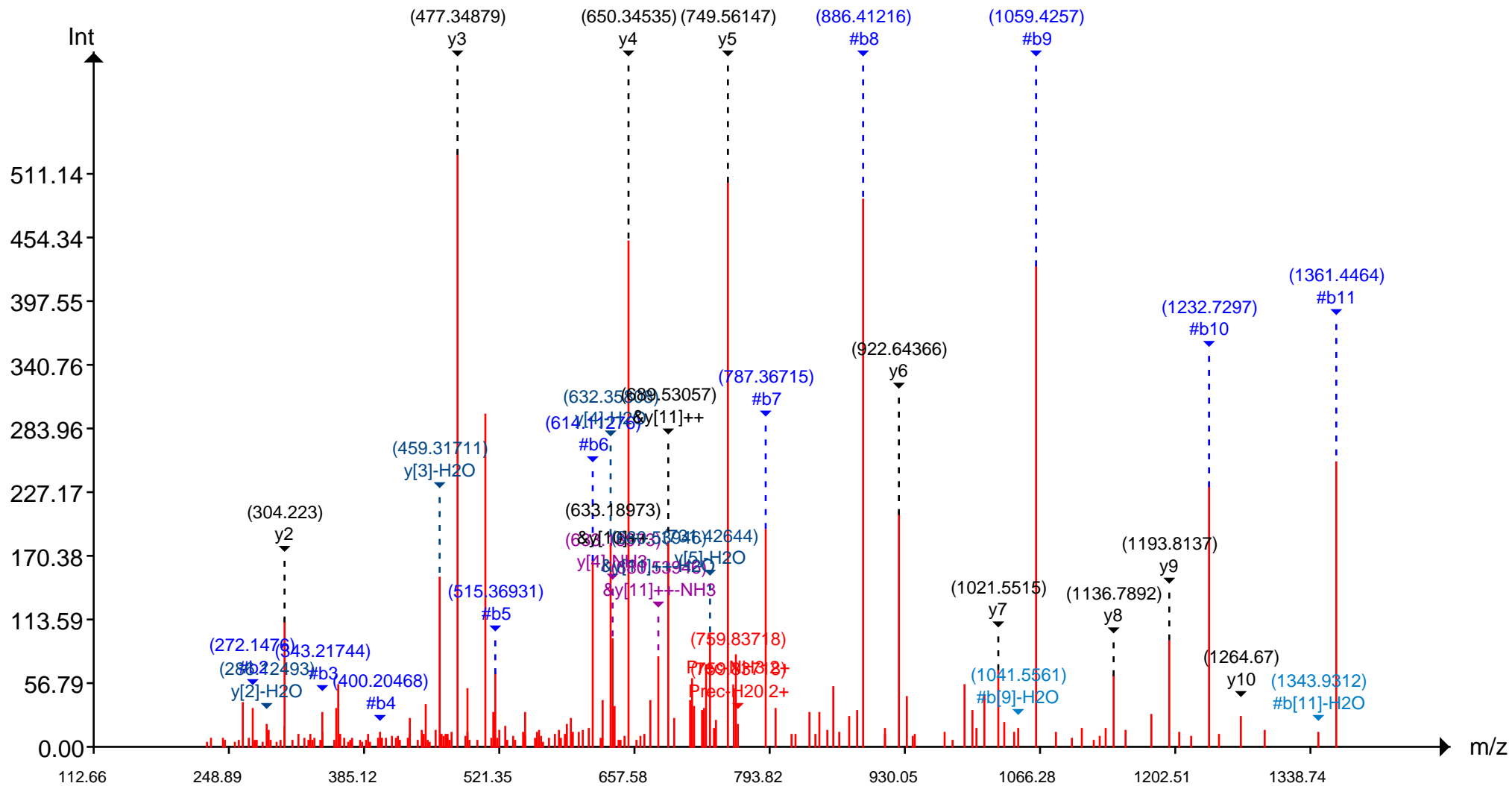


Info			
Modified Sequence	AcD3-QC<Cmm*>DK<AcD3K*>LK<AcD3K*>SVK<AcD3K*>GYR<C13>-COOH	Ion score	72.65
Protein	P21981	Confidence level	99.00%
Precursor m/z	834.4691	Identity Threshold	46.65
Charge	2+	Mass error	0.003894



Info

Modified Sequence	AcD3-M<Mox*>NTELAAF TK<AcD3K*>NQK<AcD3K*>DPGVLDR-COOH	Ion score	118.83
Protein	P50543	Confidence level	99.00%
Precursor m/z	1200.1159	Identity Threshold	47.90
Charge	2+	Mass error	0.002389



Info

Modified Sequence AcD3-ILAGDVK<AcD3K*>VK<AcD3K*>K<AcD3K*>ER-COOH

Ion score 72.26

Protein Q8K3X4

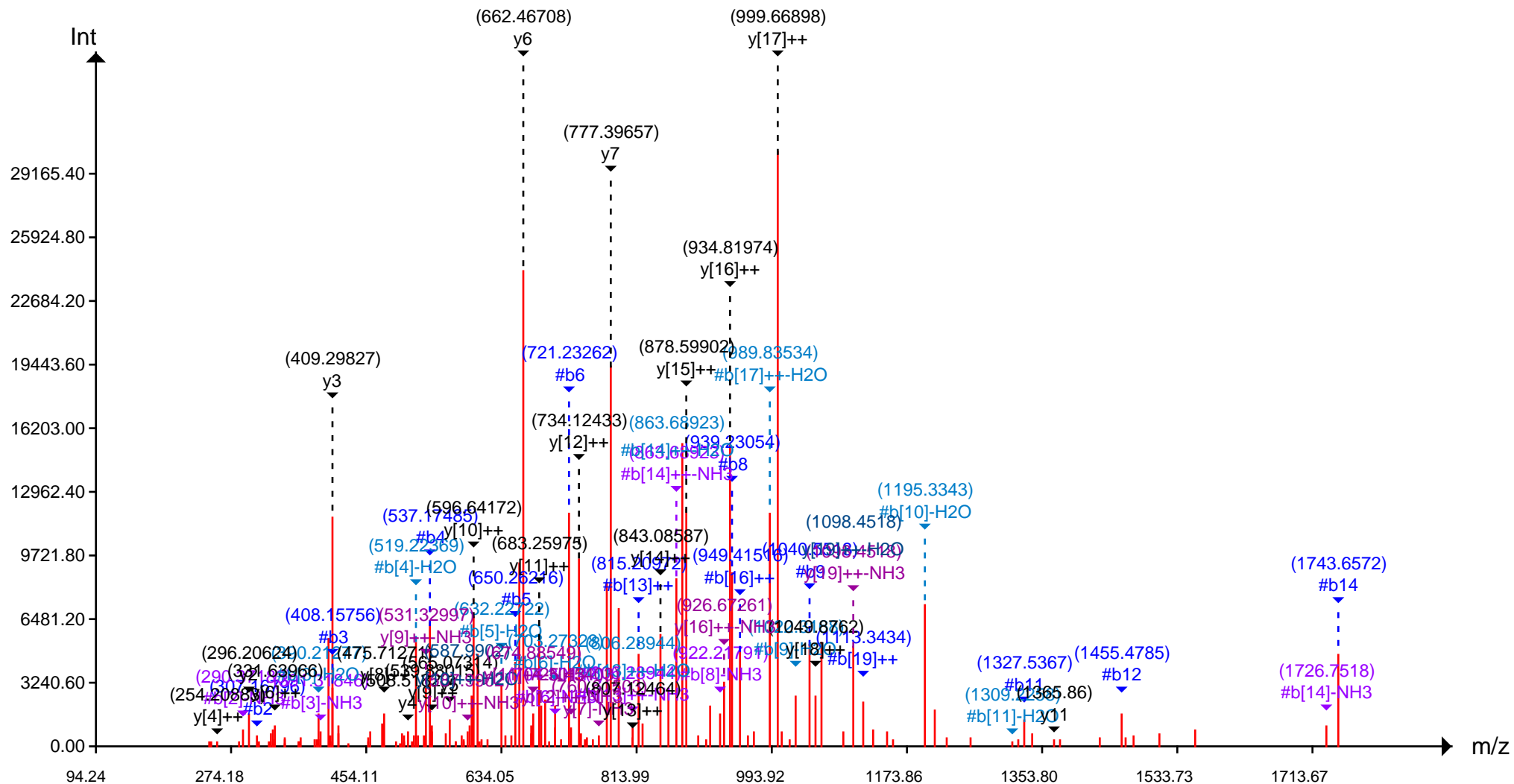
Confidence level 99.00%

Precursor m/z 768.48159

Identity Threshold 39.38

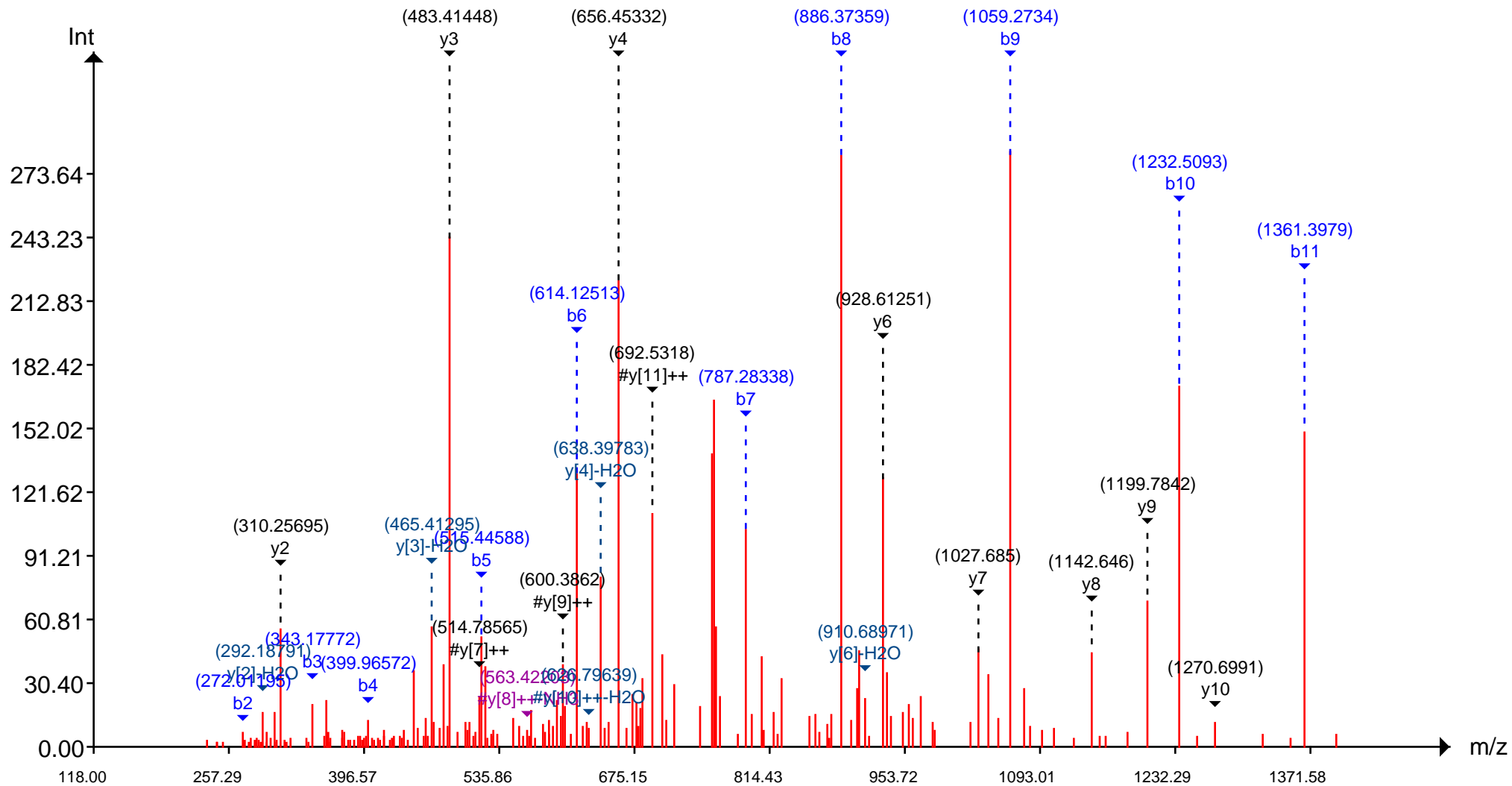
Charge 2+

Mass error 0.001423



Info

Modified Sequence	AcD3-M<Mox*>NTELAFTK<AcD3K*>NQK<AcD3K*>DPGVLDK<C13>-COOH	Ion score	66.35
Protein	P50543	Confidence level	99.00%
Precursor m/z	802.41931	Identity Threshold	47.88
Charge	3+	Mass error	0.001101



Info

Modified Sequence AcD3-ILAGDVK<AcD3K*>VK<AcD3K*>K<AcD3K*>ER<C13>-COOH

Ion score 61.69

Protein Q8K3X4

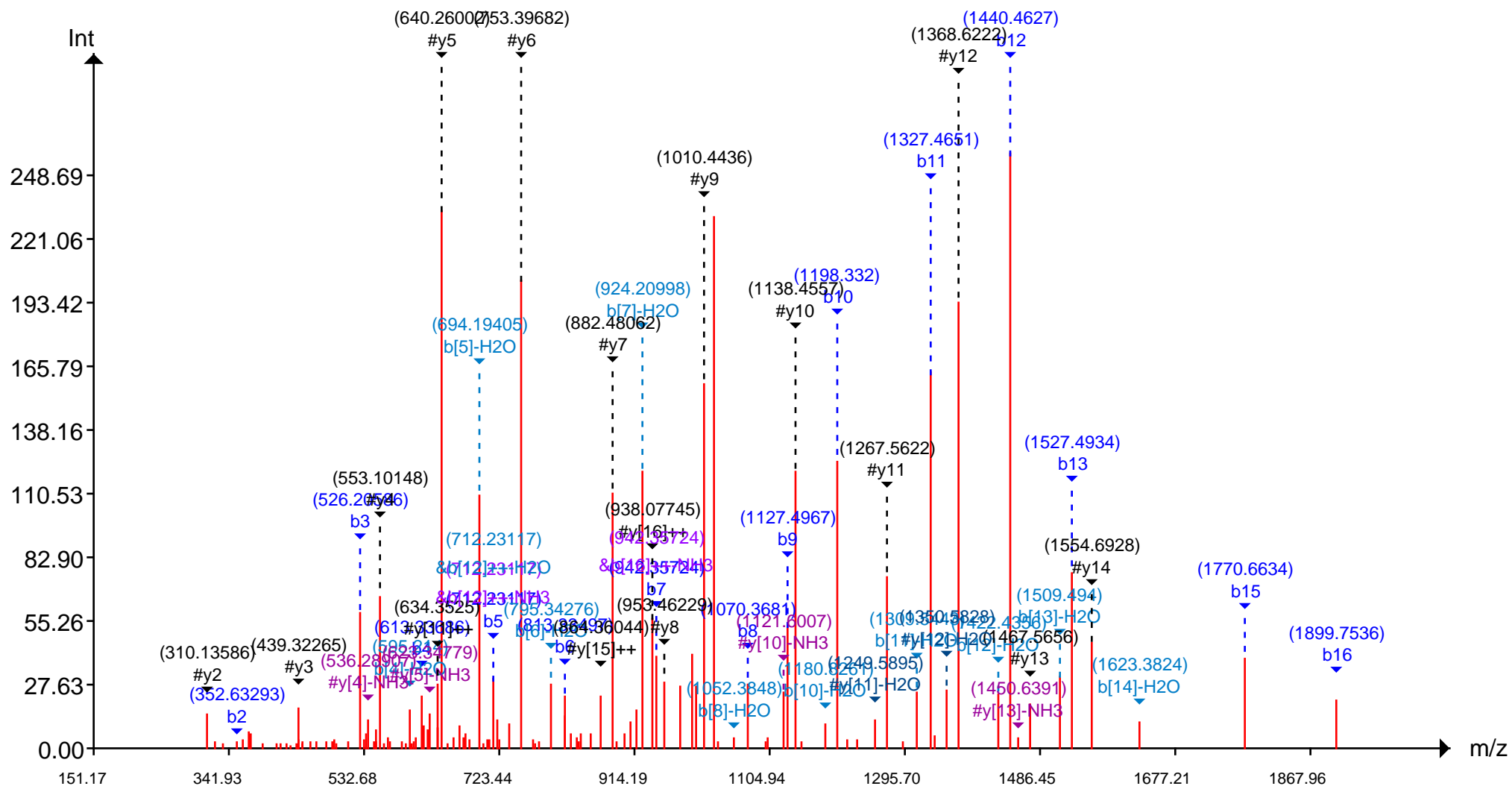
Confidence level 99.00%

Precursor m/z 771.49185

Identity Threshold 37.68

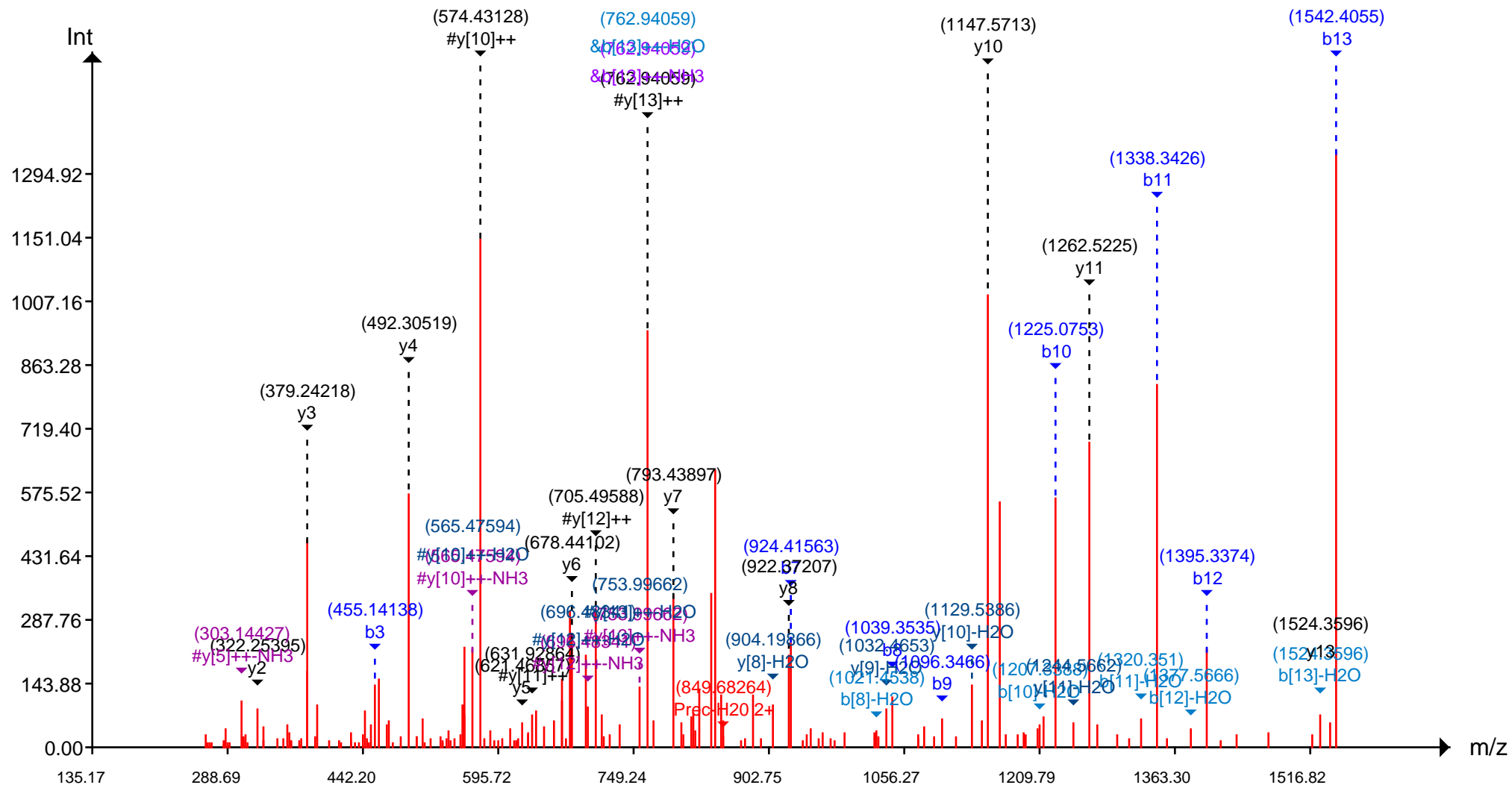
Charge 2+

Mass error 0.001802



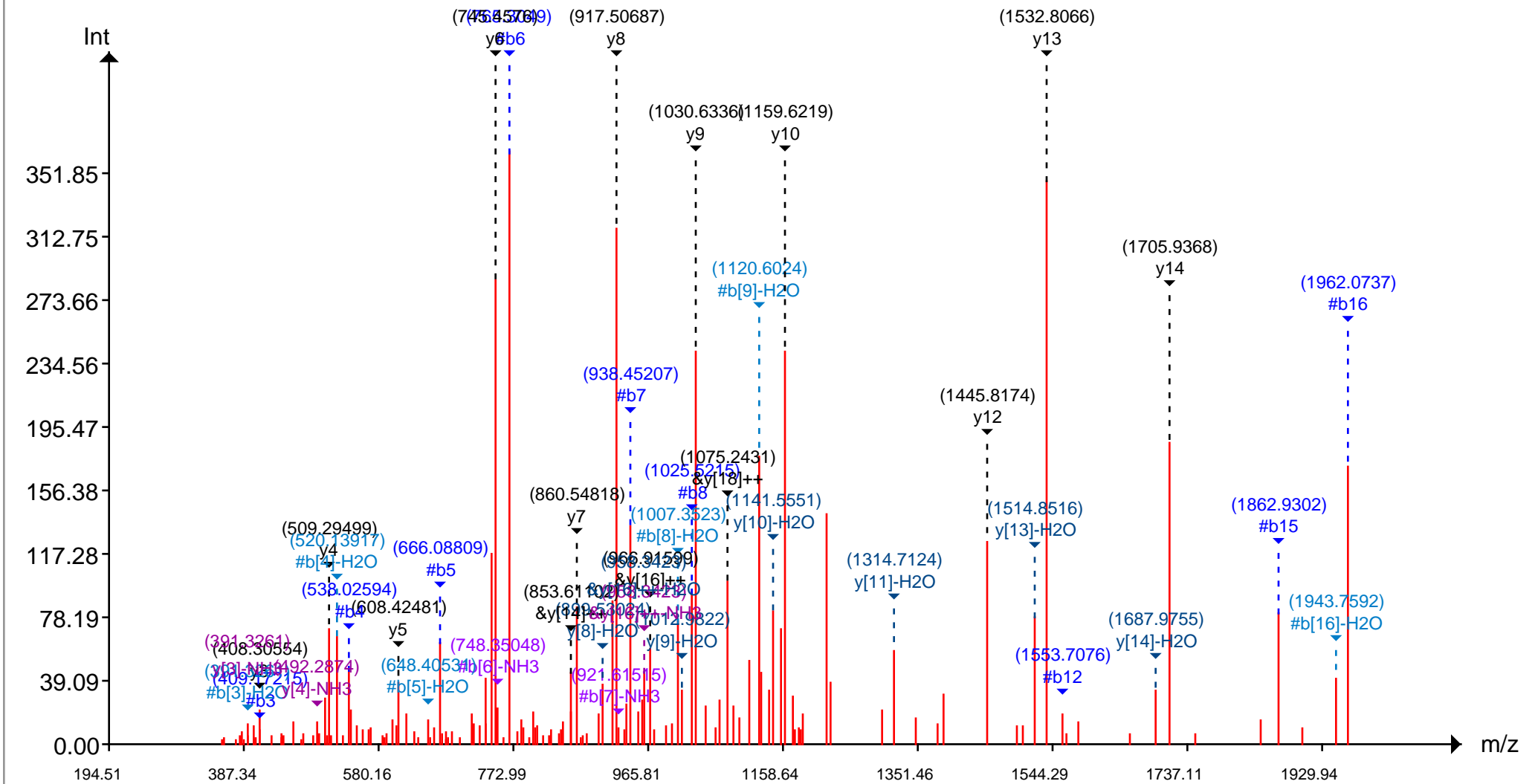
Info

Modified Sequence	AcD3-C<Cmm*>M<Mox*>K<AcD3K*>SVTEQGAELSNEER<C13>-COOH	Ion score	115.29
Protein	P63101	Confidence level	99.00%
Precursor m/z	1040.4812	Identity Threshold	44.94
Charge	2+	Mass error	0.001589



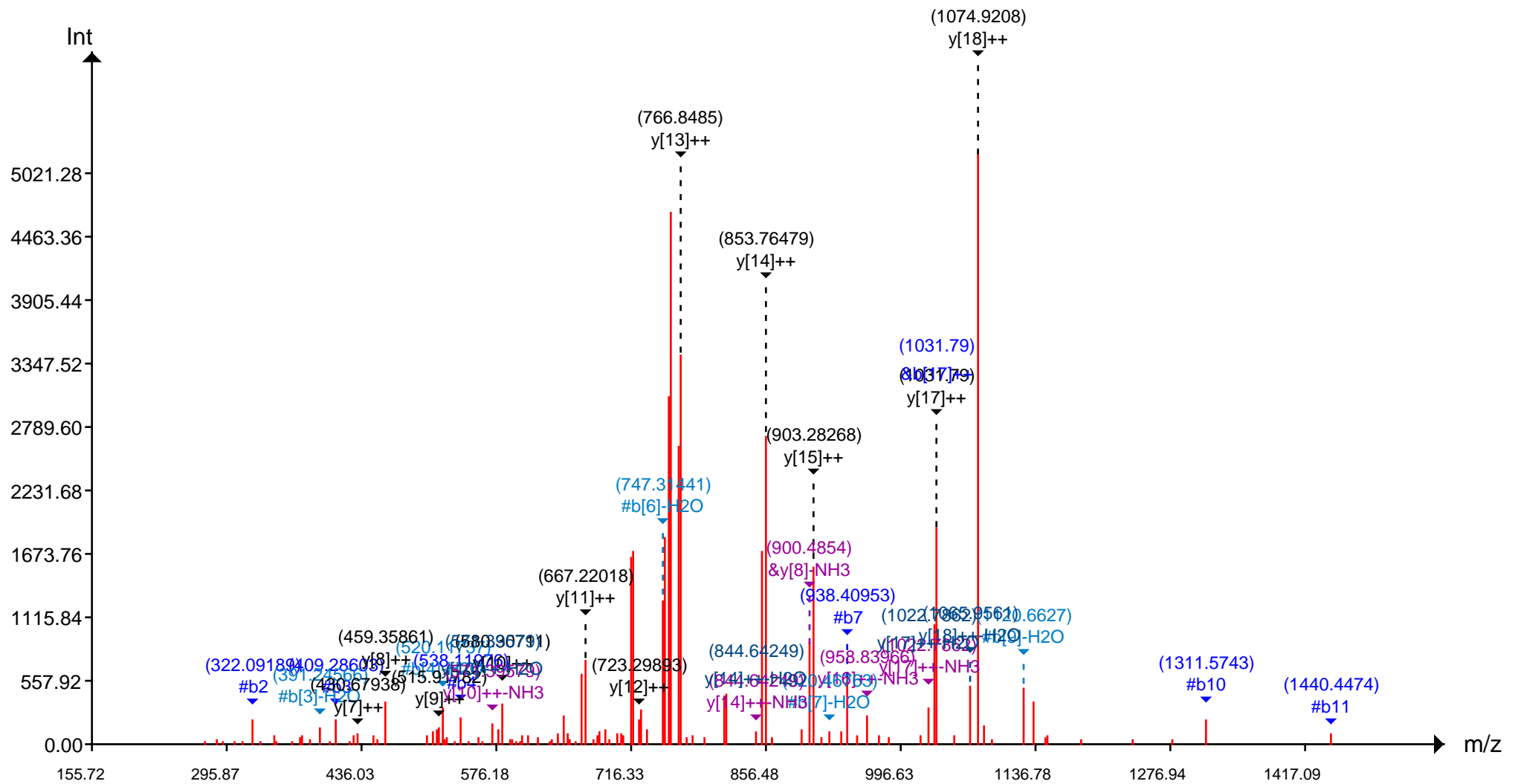
Info

Modified Sequence	AcD3-FDFDPQEDGELGFR-COOH	Ion score	66.55
Protein	Q60631	Confidence level	99.00%
Precursor m/z	858.88407	Identity Threshold	43.28
Charge	2+	Mass error	0.002917



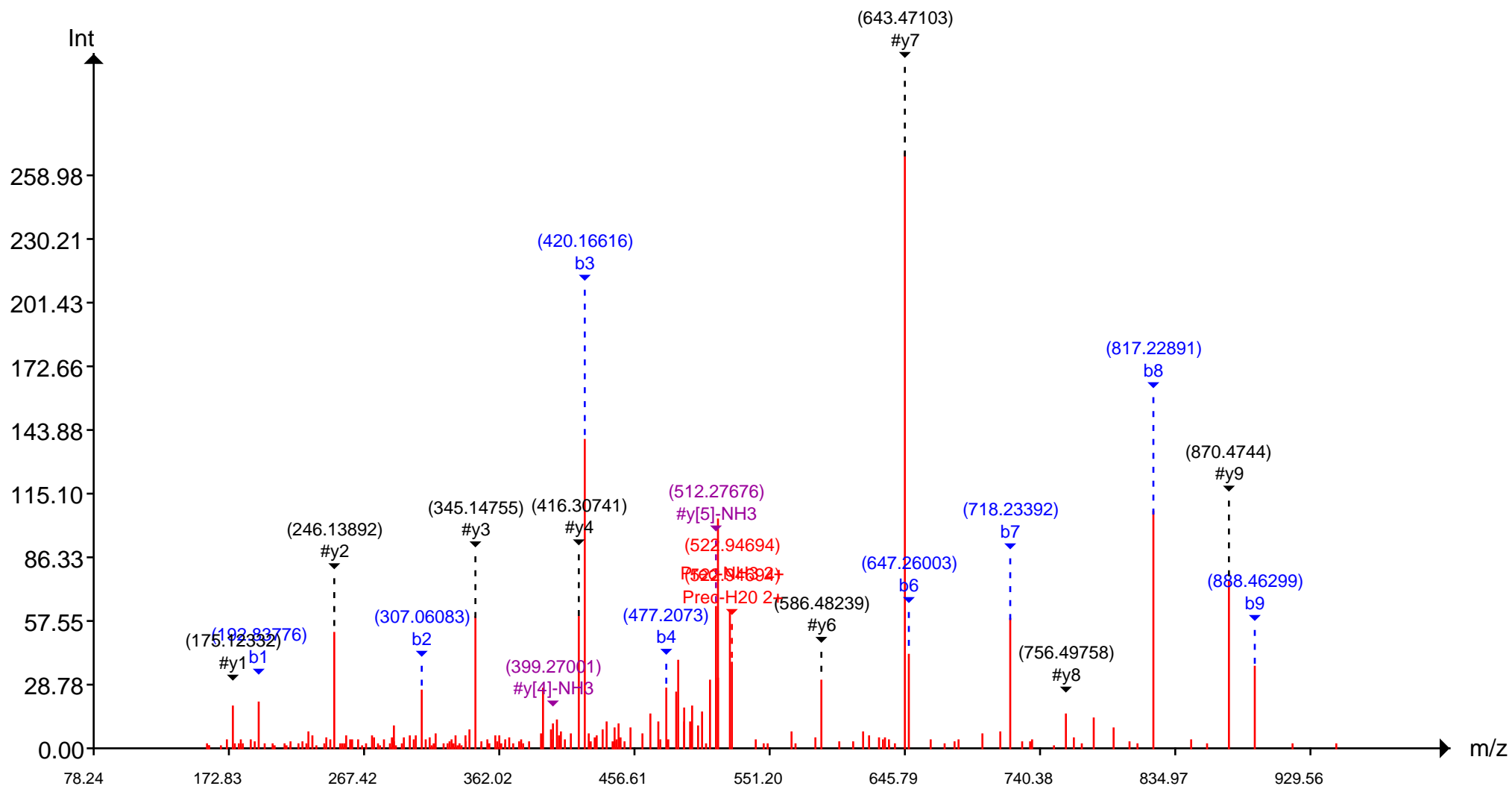
Info

Modified Sequence	AcD3-YLSEQVK<AcD3K*>SIK<AcD3K*>ELGDHVTNLR<C13>-COOH	Ion score	86.24
Protein	P09528	Confidence level	99.00%
Precursor m/z	1235.685	Identity Threshold	45.99
Charge	2+	Mass error	0.003382



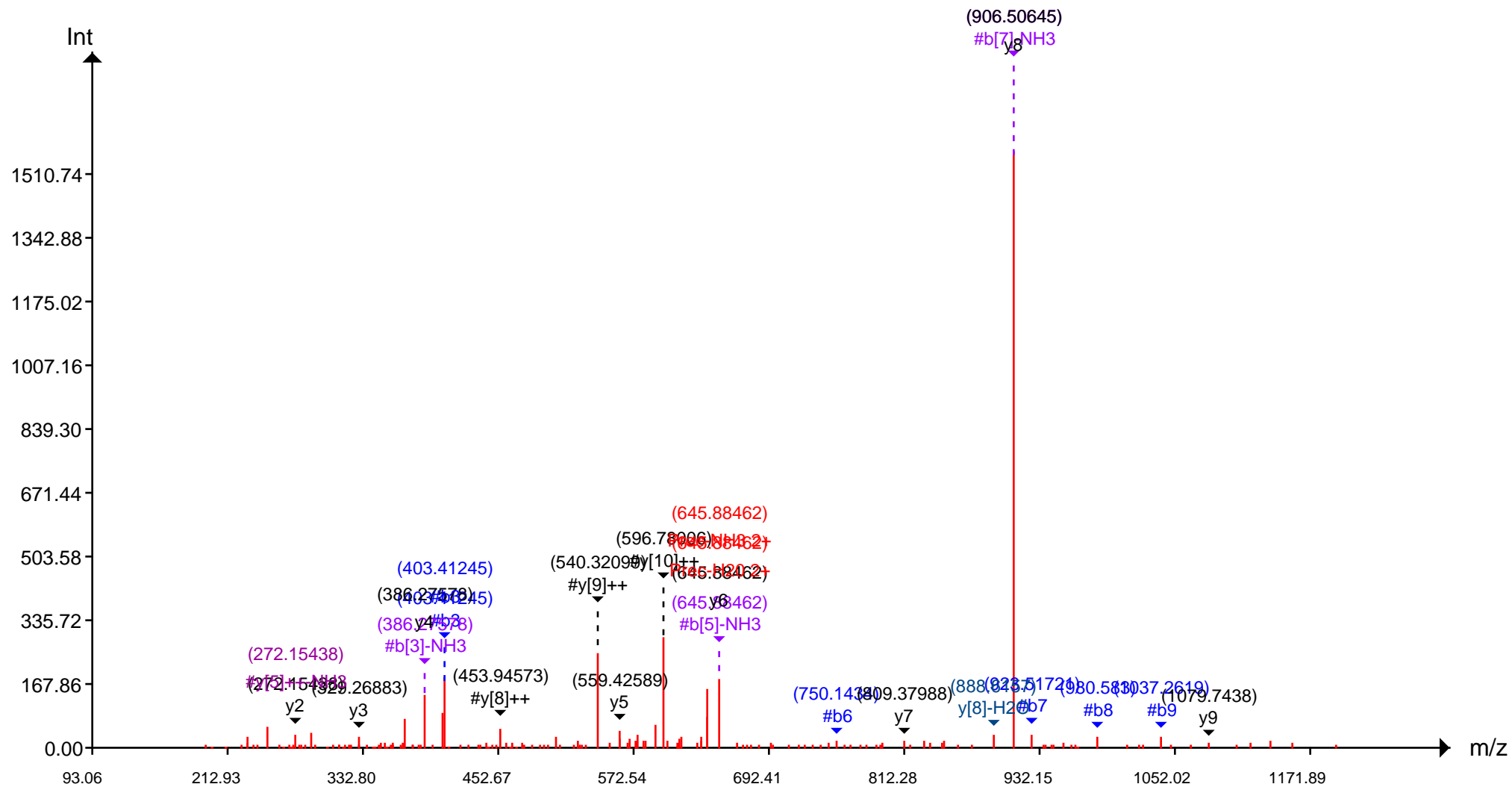
Info

Modified Sequence	AcD3-YLSEQVK<AcD3K*>SIK<AcD3K*>ELGDHVTNLR<C13>-COOH	Ion score	49.05
Protein	P09528	Confidence level	99.00%
Precursor m/z	824.12568	Identity Threshold	46.02
Charge	3+	Mass error	0.003146



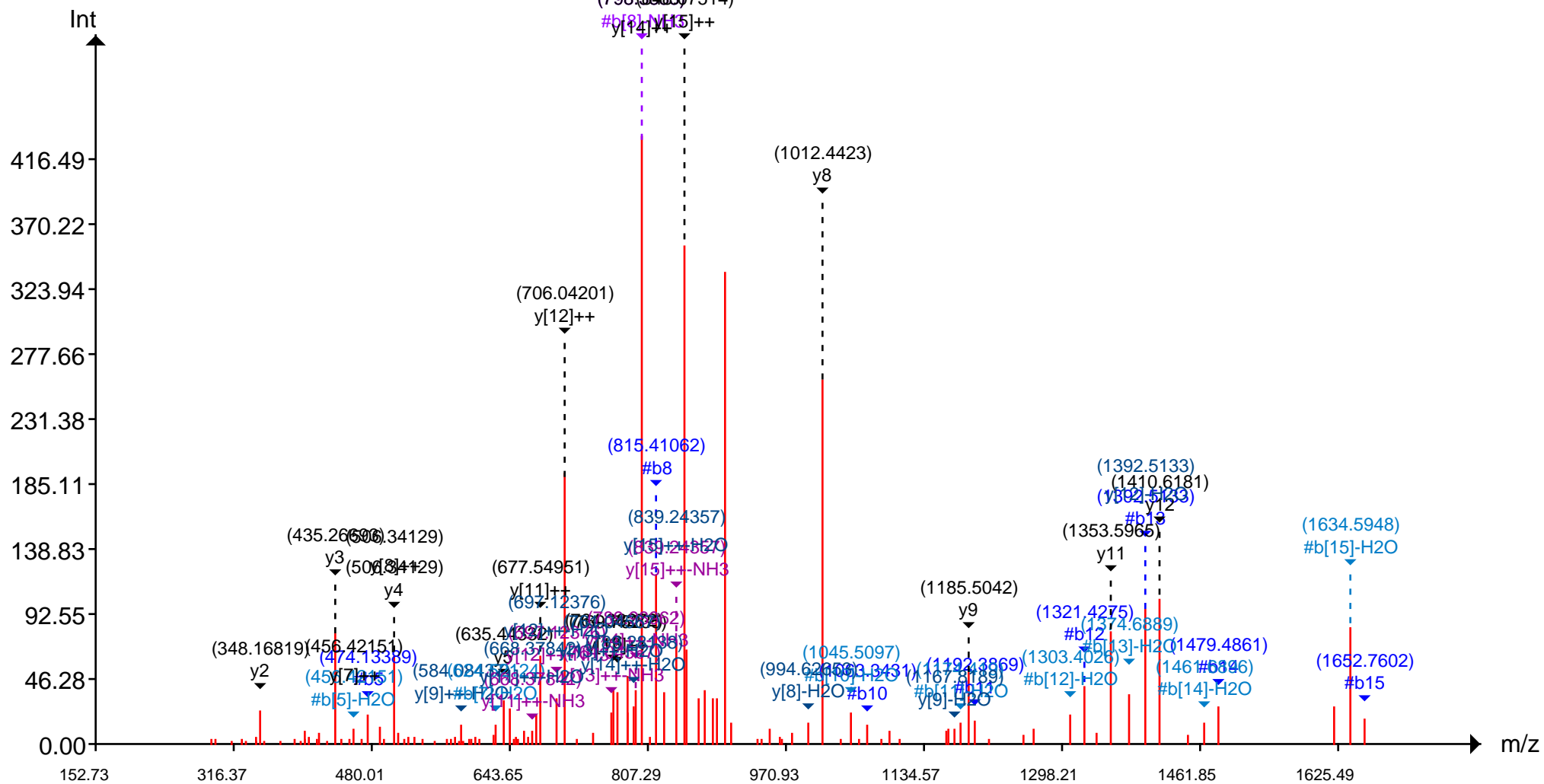
Info

Modified Sequence	AcD3-M<Mox*>NLGGLAVAR-COOH	Ion score	68.68
Protein	Q9WVA4	Confidence level	99.00%
Precursor m/z	531.79373	Identity Threshold	48.03
Charge	2+	Mass error	-1.66E-4



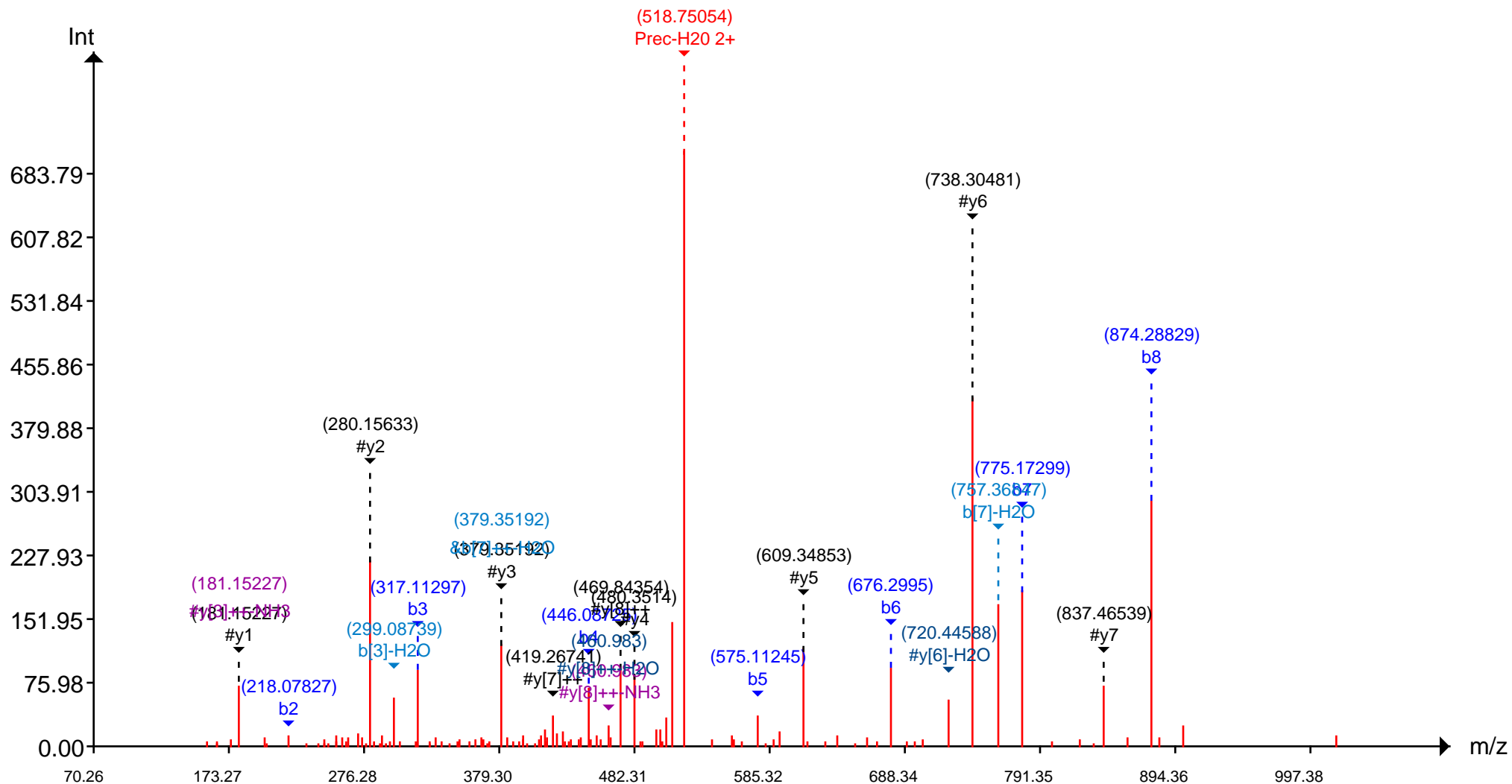
Info

Modified Sequence	AcD3-ALK<AcD3K*>PYSK<AcD3K*>GGPR-COOH	Ion score	56.41
Protein	P06745	Confidence level	99.00%
Precursor m/z	654.88531	Identity Threshold	45.32
Charge	2+	Mass error	0.001307



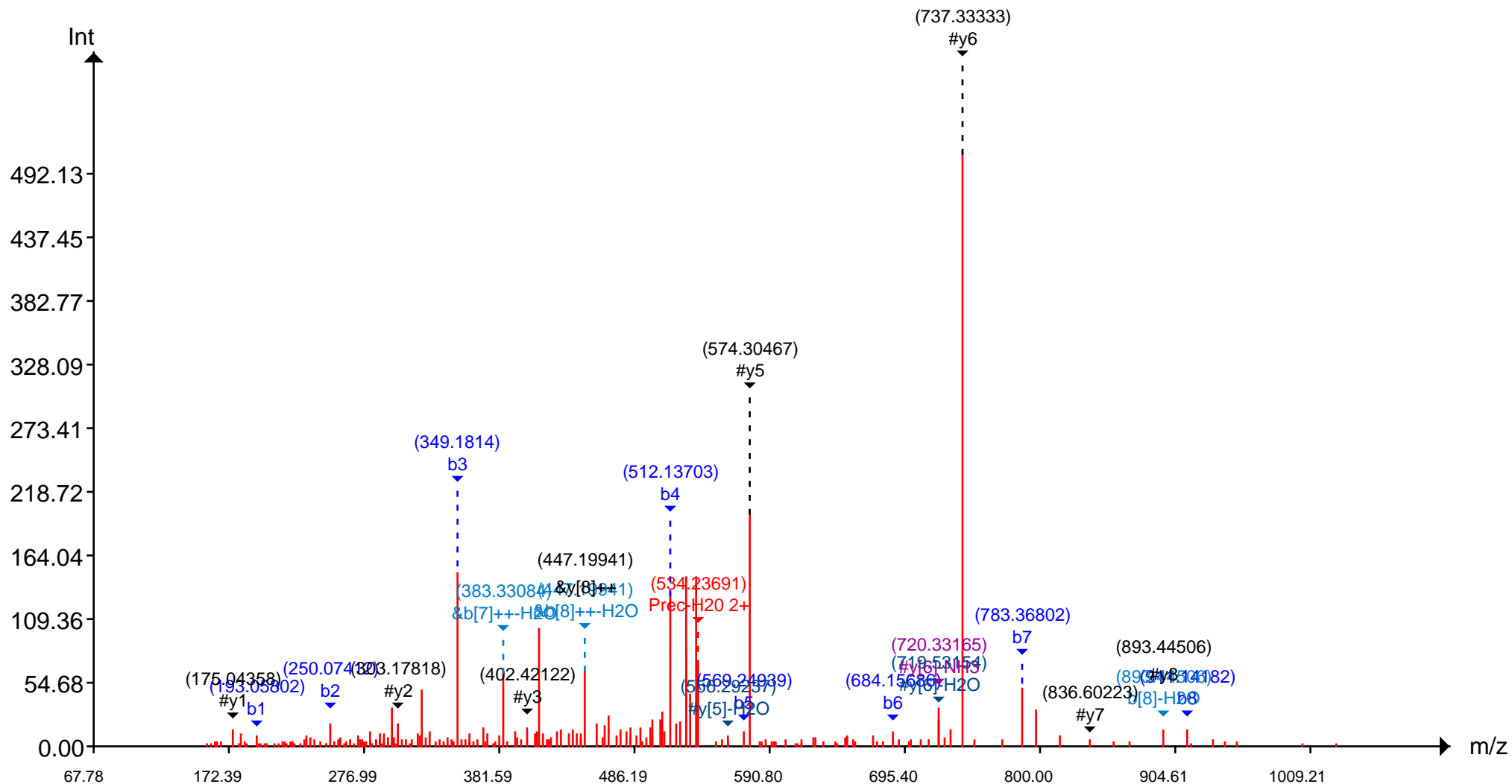
Info

Modified Sequence	AcD3-SVGQGPAK<AcD3K*>TM<Mox*>EEASK<AcD3K*>R-COOH	Ion score	71.21
Protein	O70310	Confidence level	99.00%
Precursor m/z	913.96755	Identity Threshold	48.03
Charge	2+	Mass error	0.001542



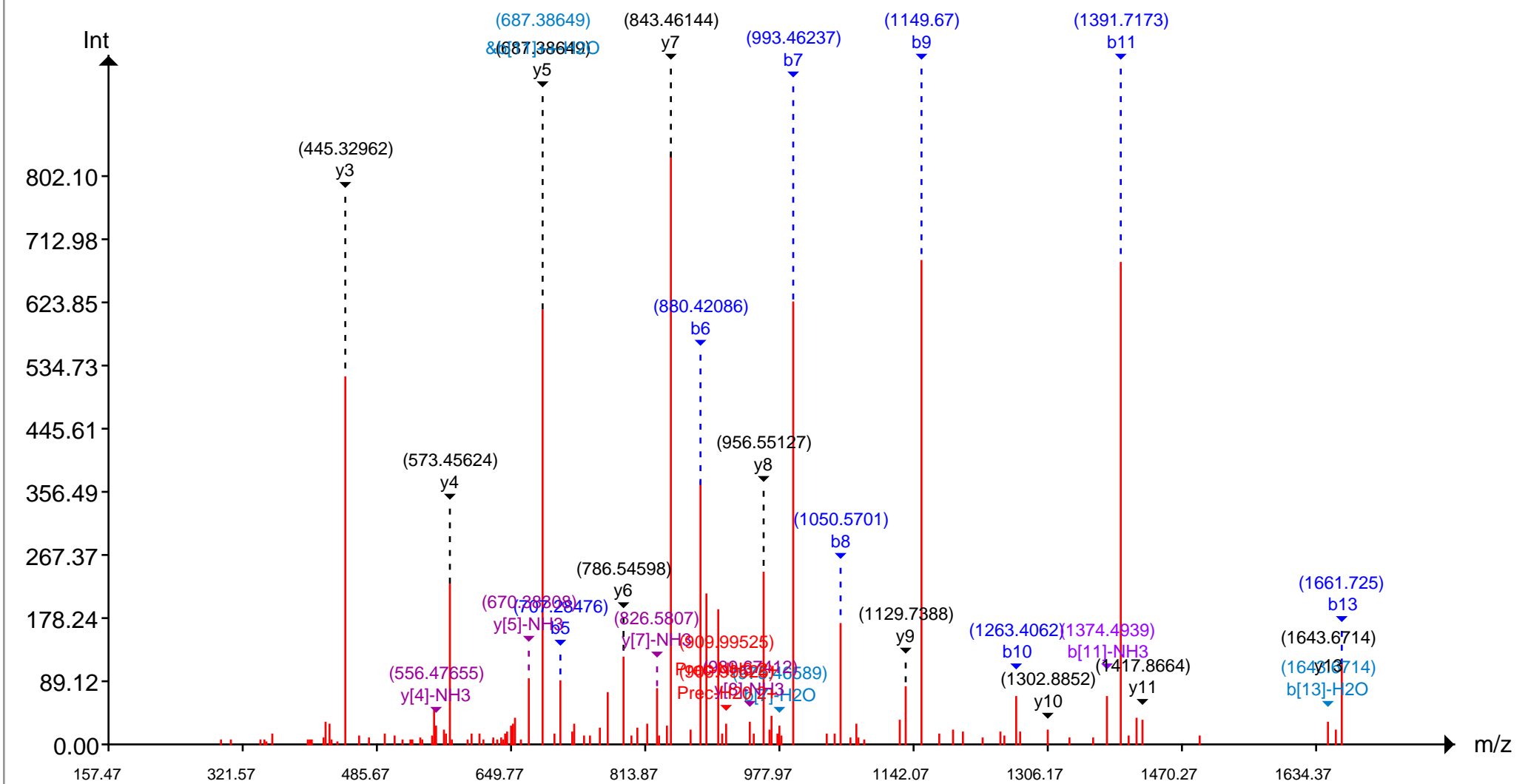
Info

Modified Sequence	AcD3-ATVEETVVR<C13>-COOH	Ion score	62.58
Protein	Q76MZ3, Q7TNP2	Confidence level	99.00%
Precursor m/z	527.79916	Identity Threshold	46.78
Charge	2+	Mass error	-3.38E-4



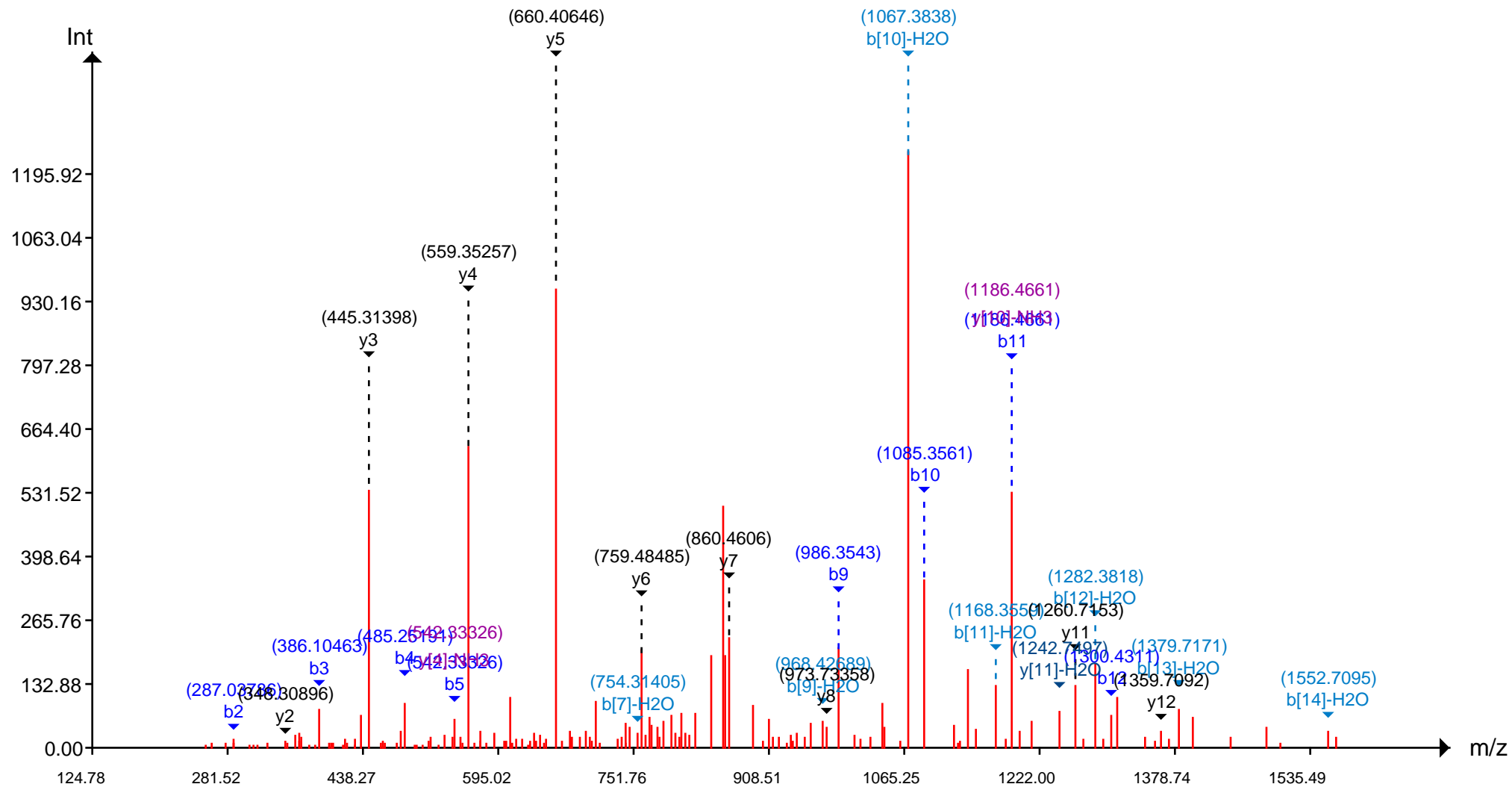
Info

Modified Sequence	AcD3-FGVYGDVQR-COOH	Ion score	50.77
Protein	P17225, Q91Z31	Confidence level	99.00%
Precursor m/z	543.27675	Identity Threshold	47.43
Charge	2+	Mass error	8.47E-4



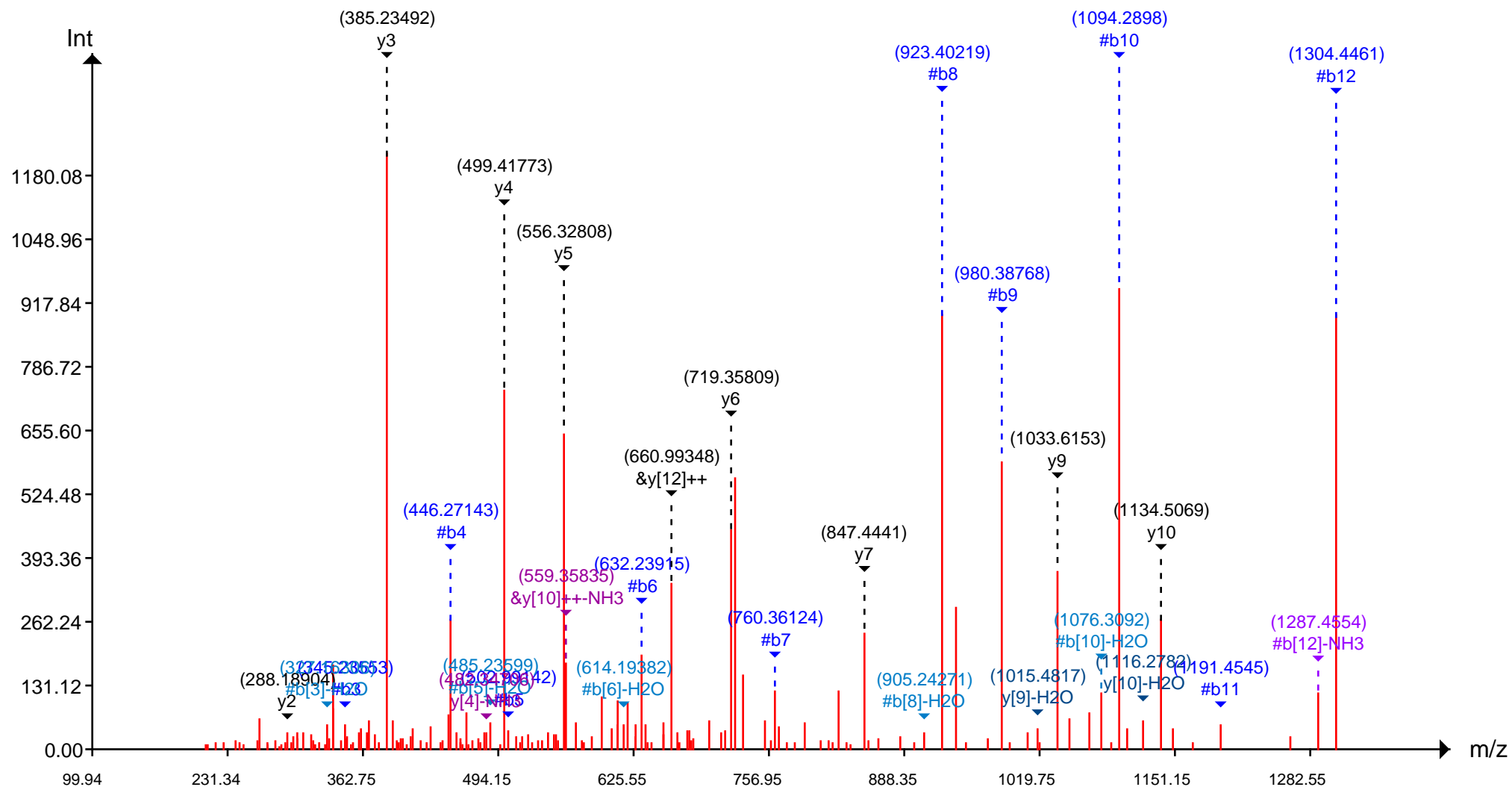
Info

Modified Sequence	AcD3-FLLDK<AcD3K*>K<AcD3K*>IGVNQPK<AcD3K*>R-COOH	Ion score	72.03
Protein	P80314	Confidence level	99.00%
Precursor m/z	918.56105	Identity Threshold	40.24
Charge	2+	Mass error	0.001713



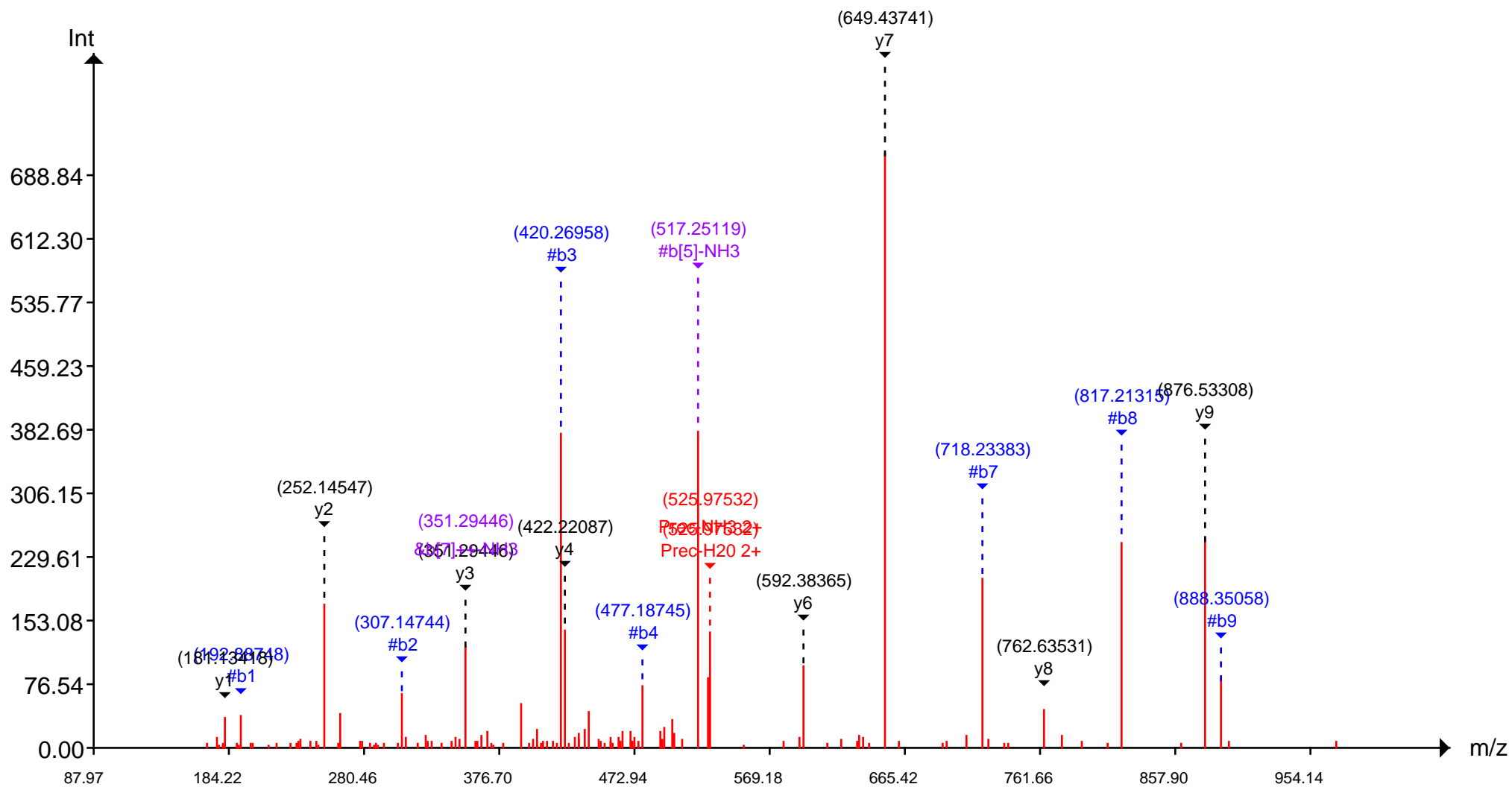
Info

Modified Sequence	AcD3-IQVVGDDLTVTNPK<AcD3K*>R-COOH	Ion score	59.68
Protein	P17182	Confidence level	99.00%
Precursor m/z	872.99031	Identity Threshold	46.76
Charge	2+	Mass error	0.002277



Info

Modified Sequence	AcD3-LSVTGEQYGNPIR-COOH	Ion score	65.8
Protein	Q9ESP1	Confidence level	99.00%
Precursor m/z	739.88854	Identity Threshold	48.06
Charge	2+	Mass error	0.002106



Info

Modified Sequence AcD3-M<Mox*>NLGGLAVAR<C13>-COOH

Ion score 63.29

Protein Q9WVA4

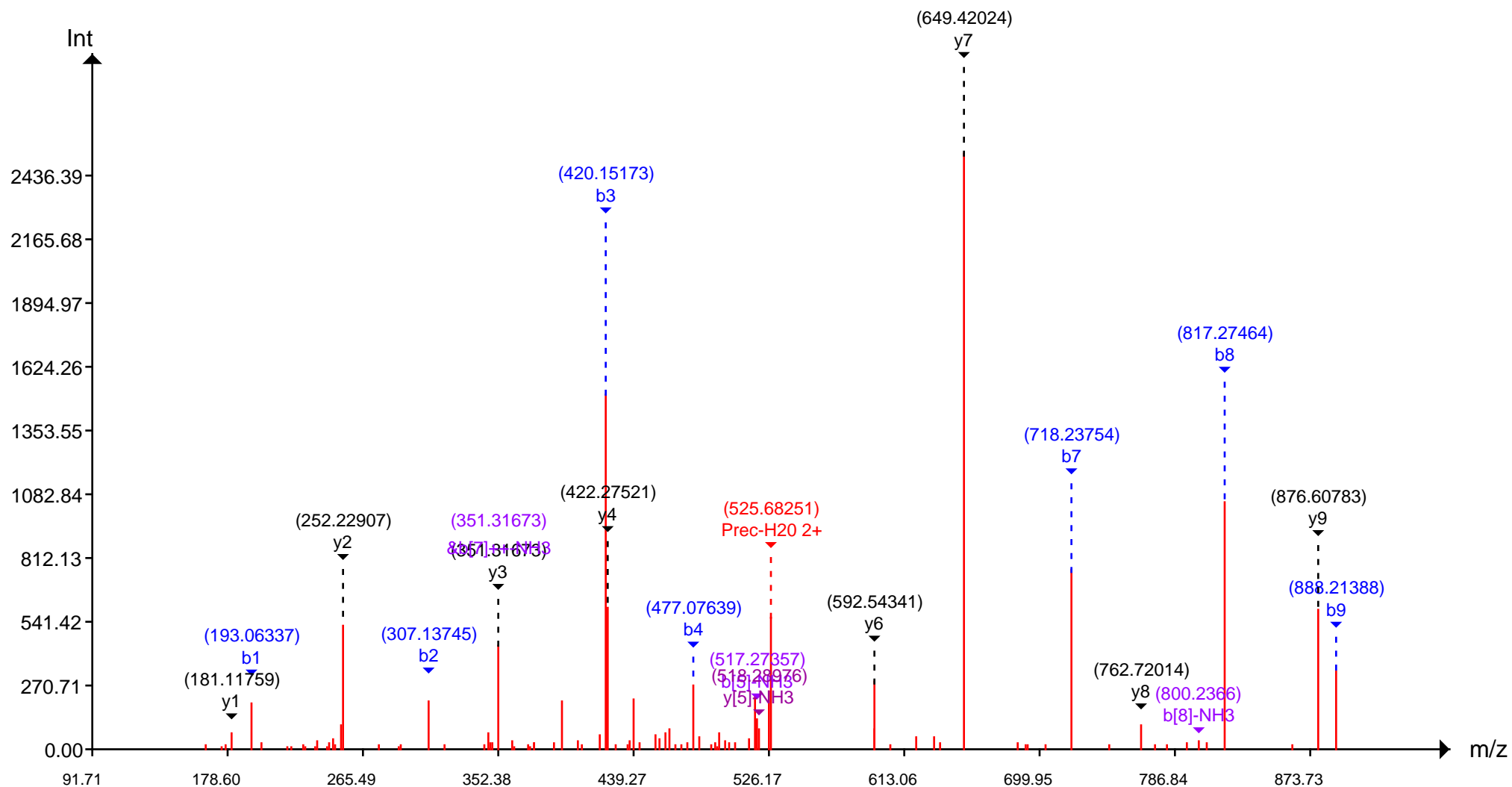
Confidence level 99.00%

Precursor m/z 534.80378

Identity Threshold 47.18

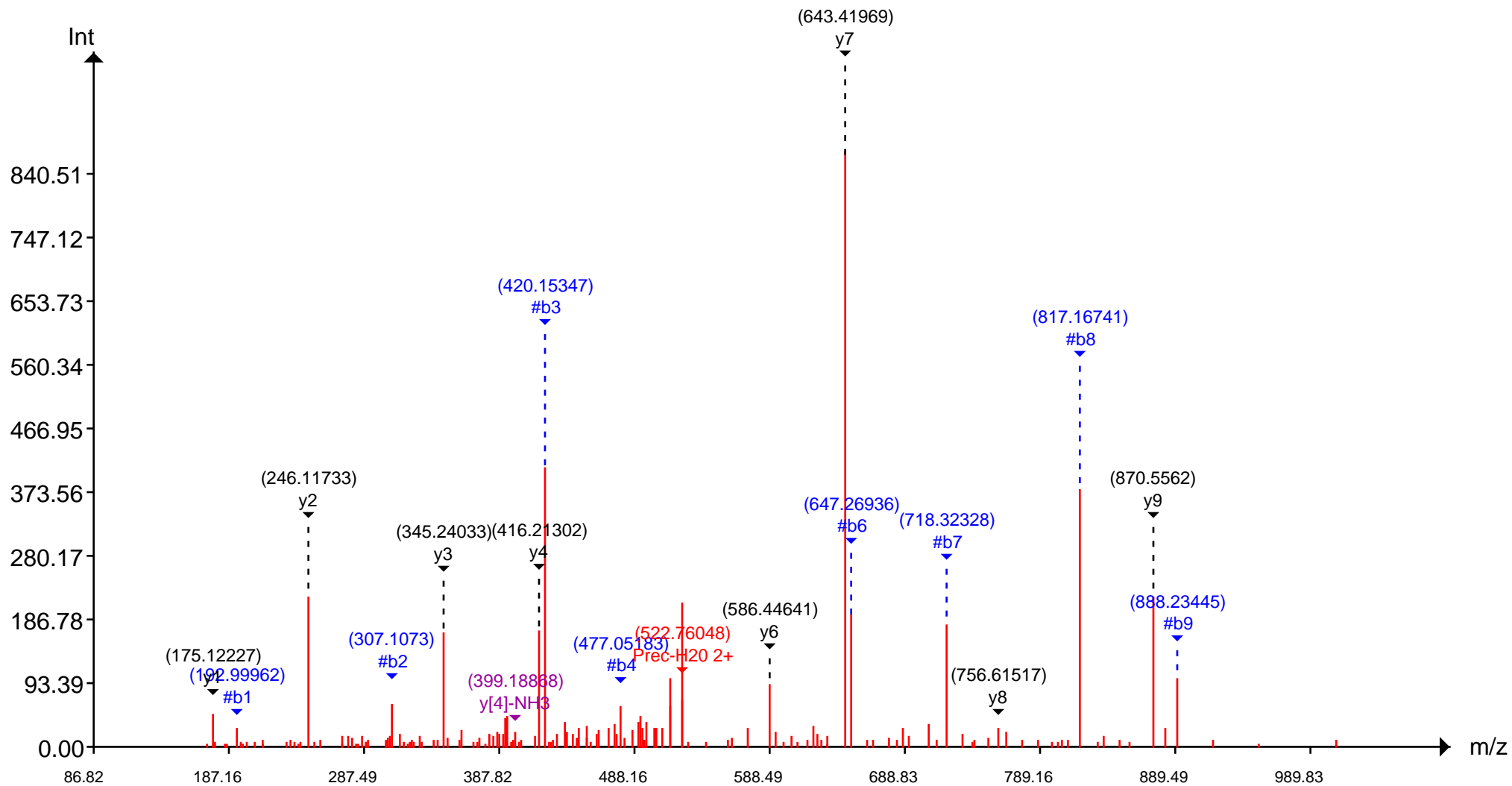
Charge 2+

Mass error -2.08E-4



Info

Modified Sequence	AcD3-M<Mox*>NLGGLAVAR<C13>-COOH	Ion score	81.94
Protein	Q9WVA4	Confidence level	99.00%
Precursor m/z	534.80419	Identity Threshold	46.81
Charge	2+	Mass error	6.12E-4



Info

Modified Sequence AcD3-M<Mox*>NLGGLAVAR-COOH

Ion score 65.49

Protein Q9WVA4

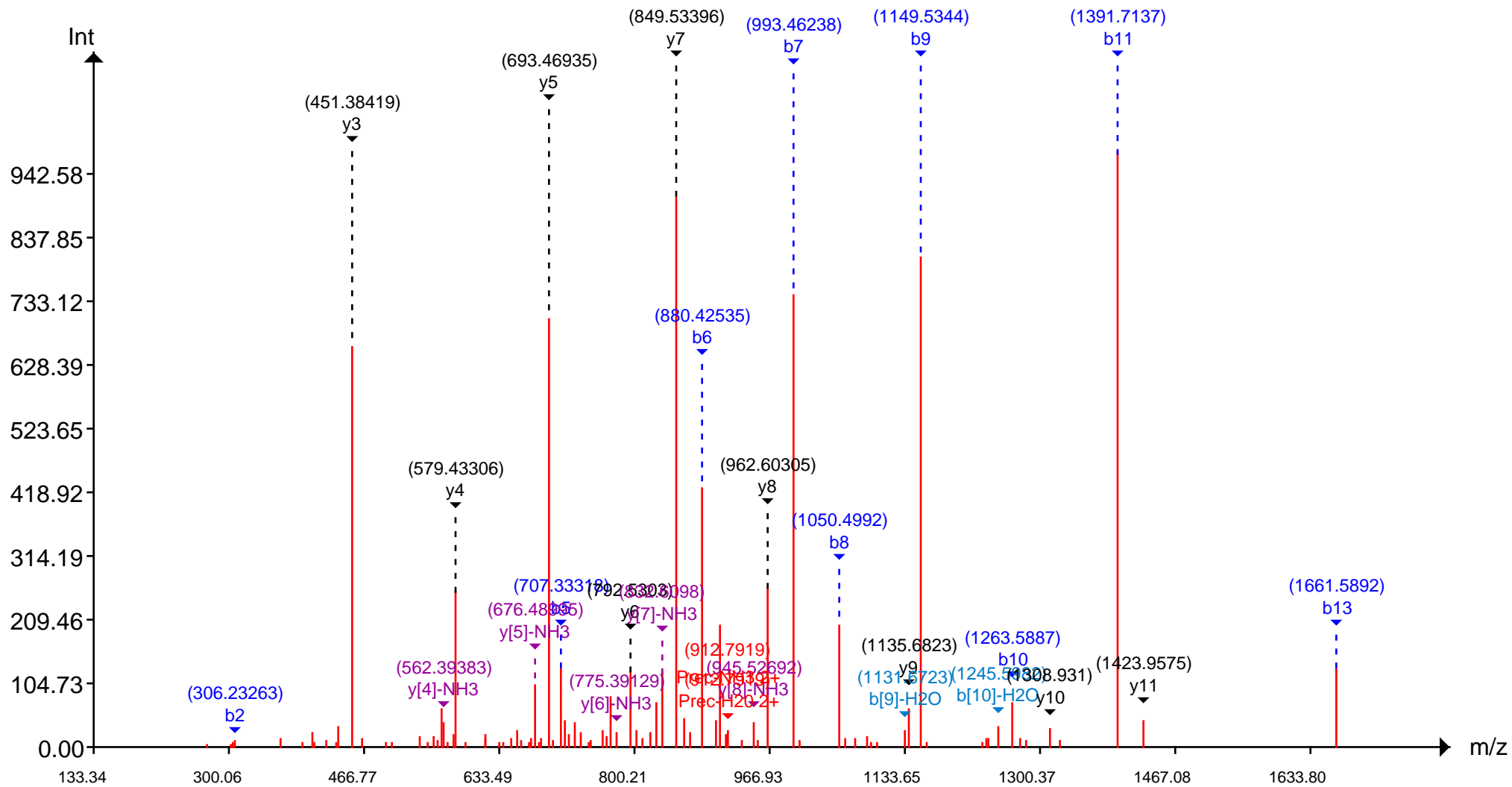
Confidence level 99.00%

Precursor m/z 531.79395

Identity Threshold 47.77

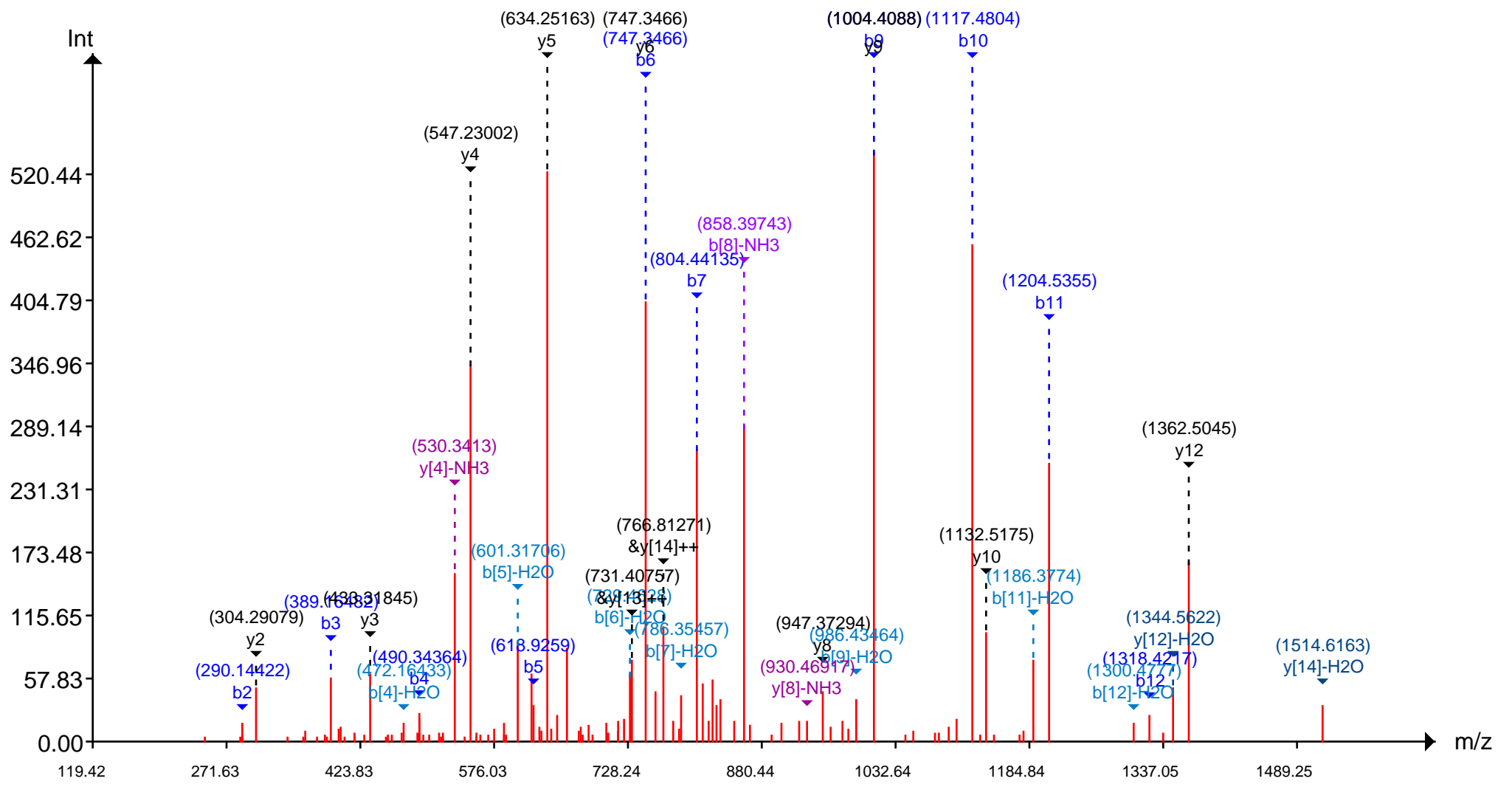
Charge 2+

Mass error 2.74E-4



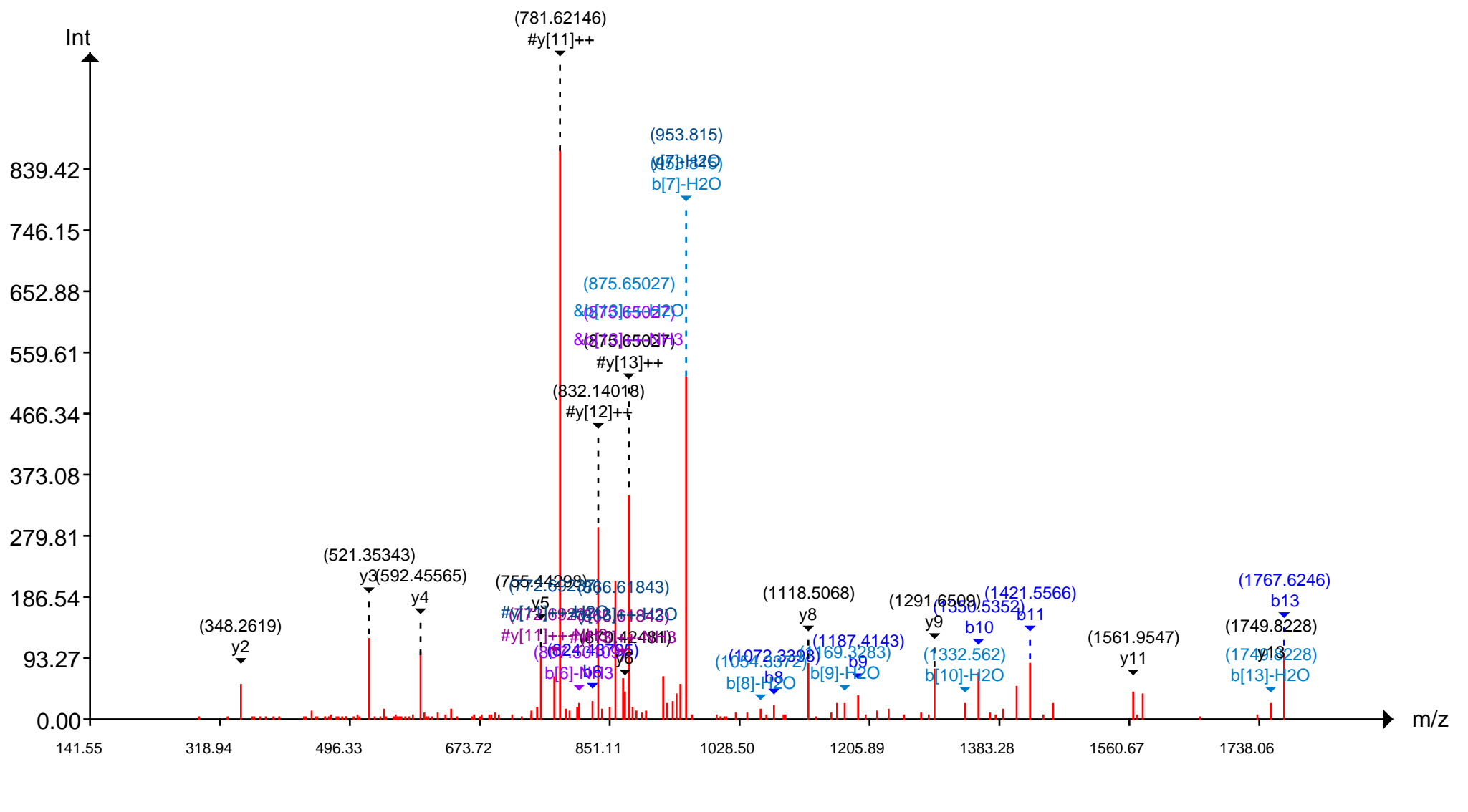
Info

Modified Sequence	AcD3-FLLDK<AcD3K*>K<AcD3K*>IGVNQPK<AcD3K*>R<C13>-COOH	Ion score	86.29
Protein	P80314	Confidence level	99.00%
Precursor m/z	921.57105	Identity Threshold	38.37
Charge	2+	Mass error	0.001571

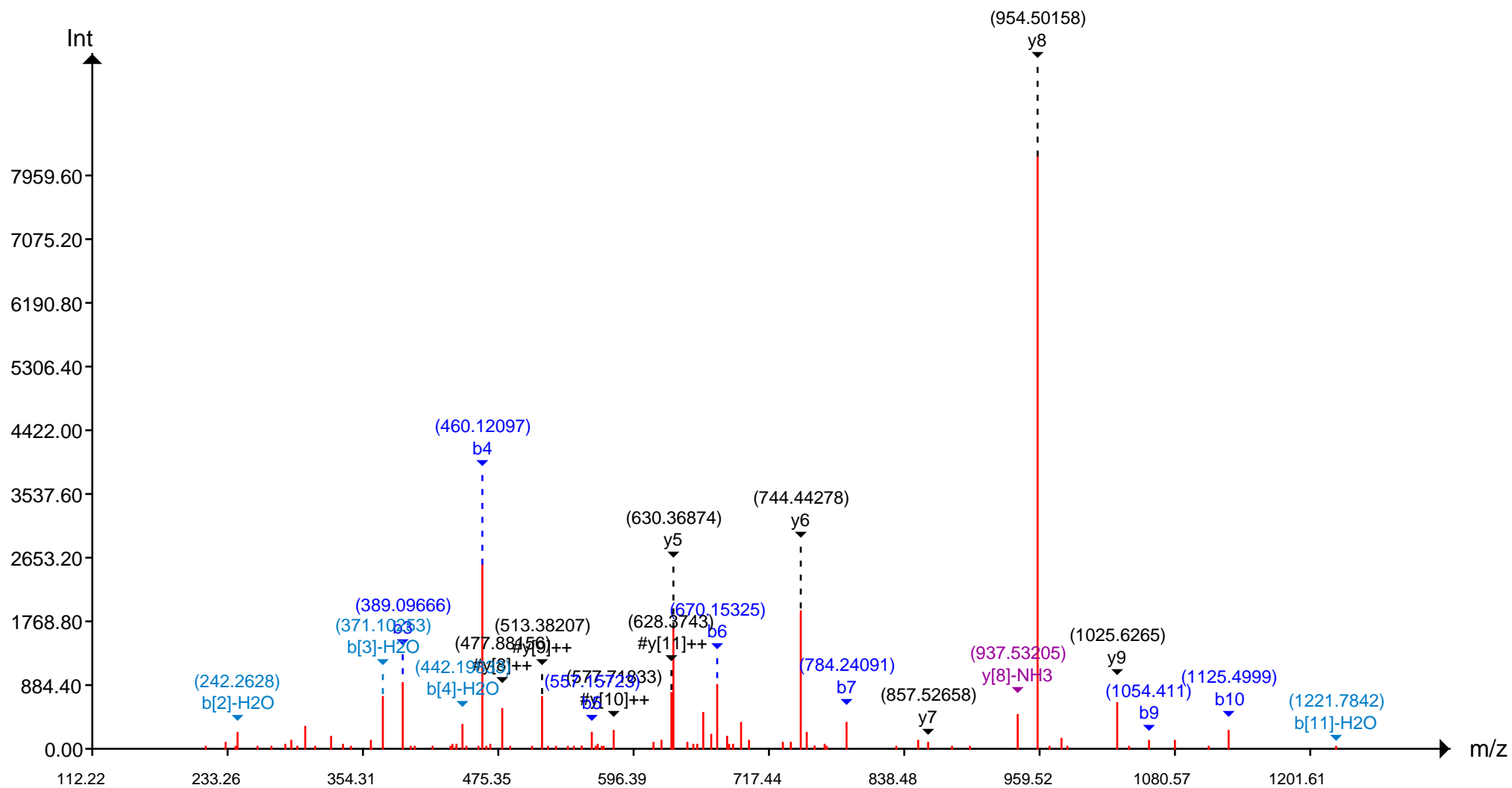


Info

Modified Sequence	AcD3-K<AcD3K*>AVTEQGAELSNEER-COOH	Ion score	94.86
Protein	P68254	Confidence level	99.00%
Precursor m/z	875.94086	Identity Threshold	47.93
Charge	2+	Mass error	0.002025

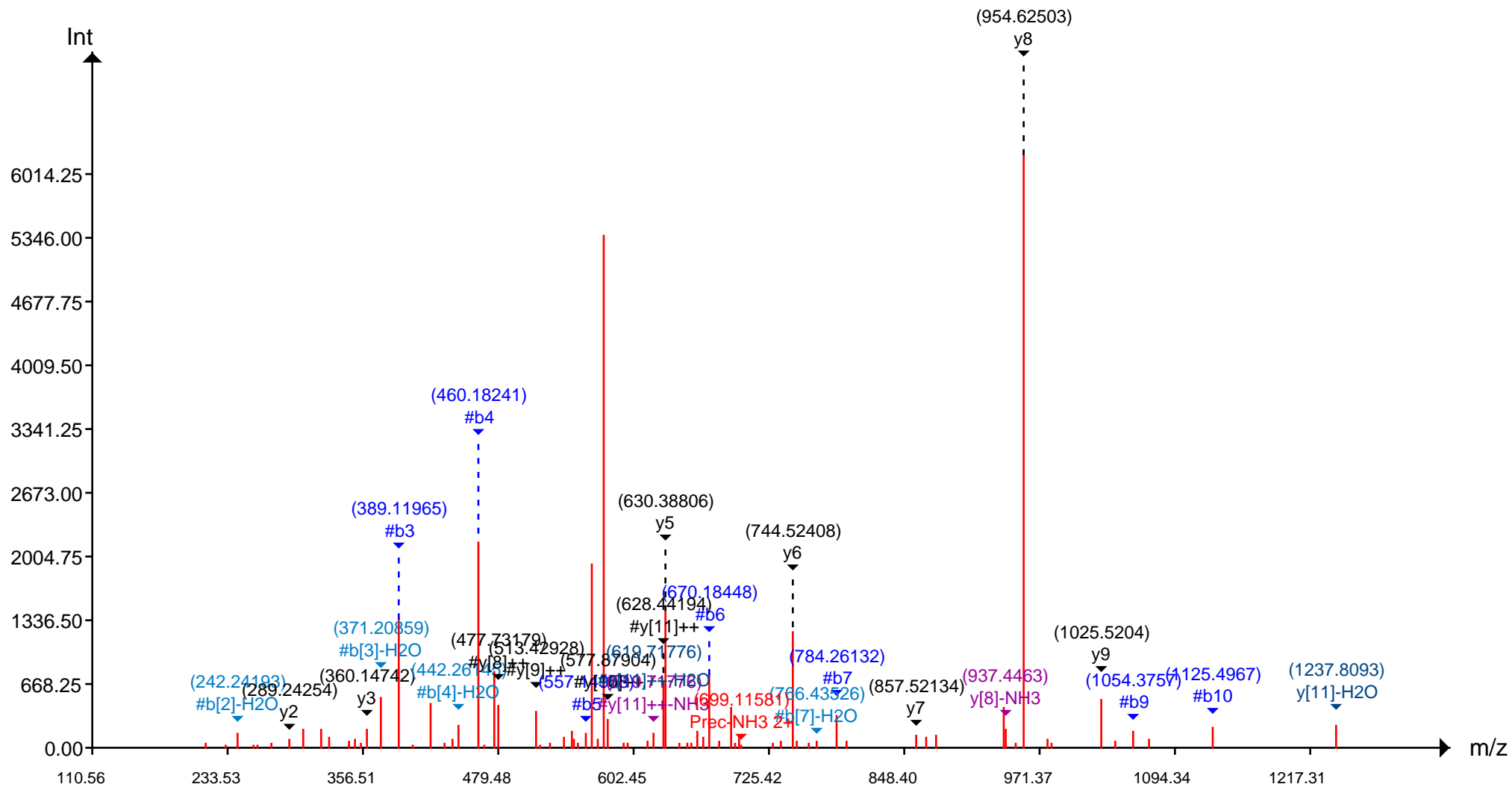


Info			
Modified Sequence	AcD3-FSTPK<AcD3K*>K<AcD3K*>M<Mox*>TDYAK<AcD3K*>K<AcD3K*>R-COOH	Ion score	49.52
Protein	Q9CX56	Confidence level	99.00%
Precursor m/z	971.53319	Identity Threshold	47.27
Charge	2+	Mass error	0.001978



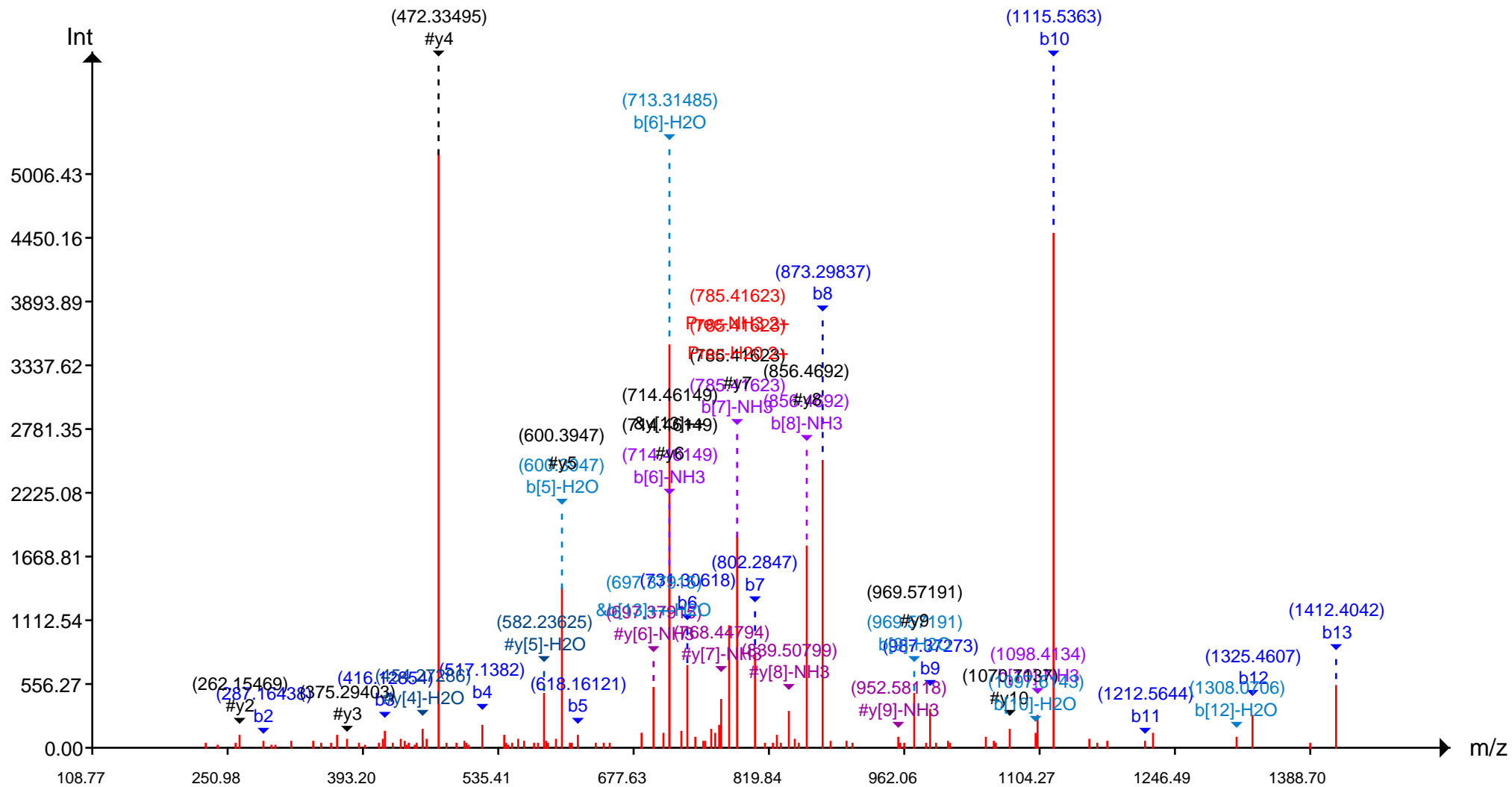
Info

Modified Sequence	AcD3-LTEAPLNPK<AcD3K*>ANR-COOH	Ion score	56.33
Protein	P62737, P60710, P68033, P63260, P63268, P68134	Confidence level	99.00%
Precursor m/z	707.40276	Identity Threshold	46.89
Charge	2+	Mass error	0.00157



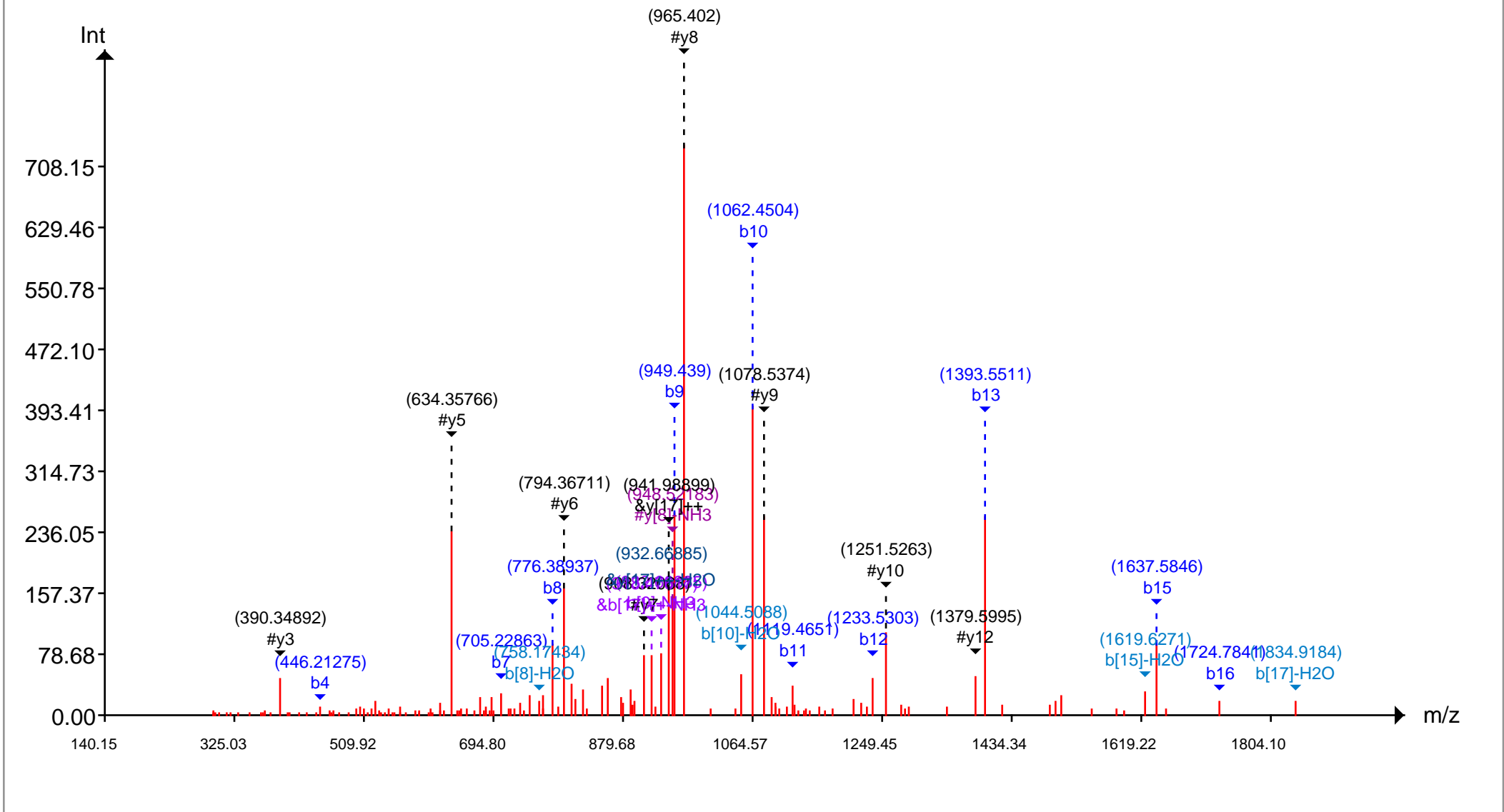
Info

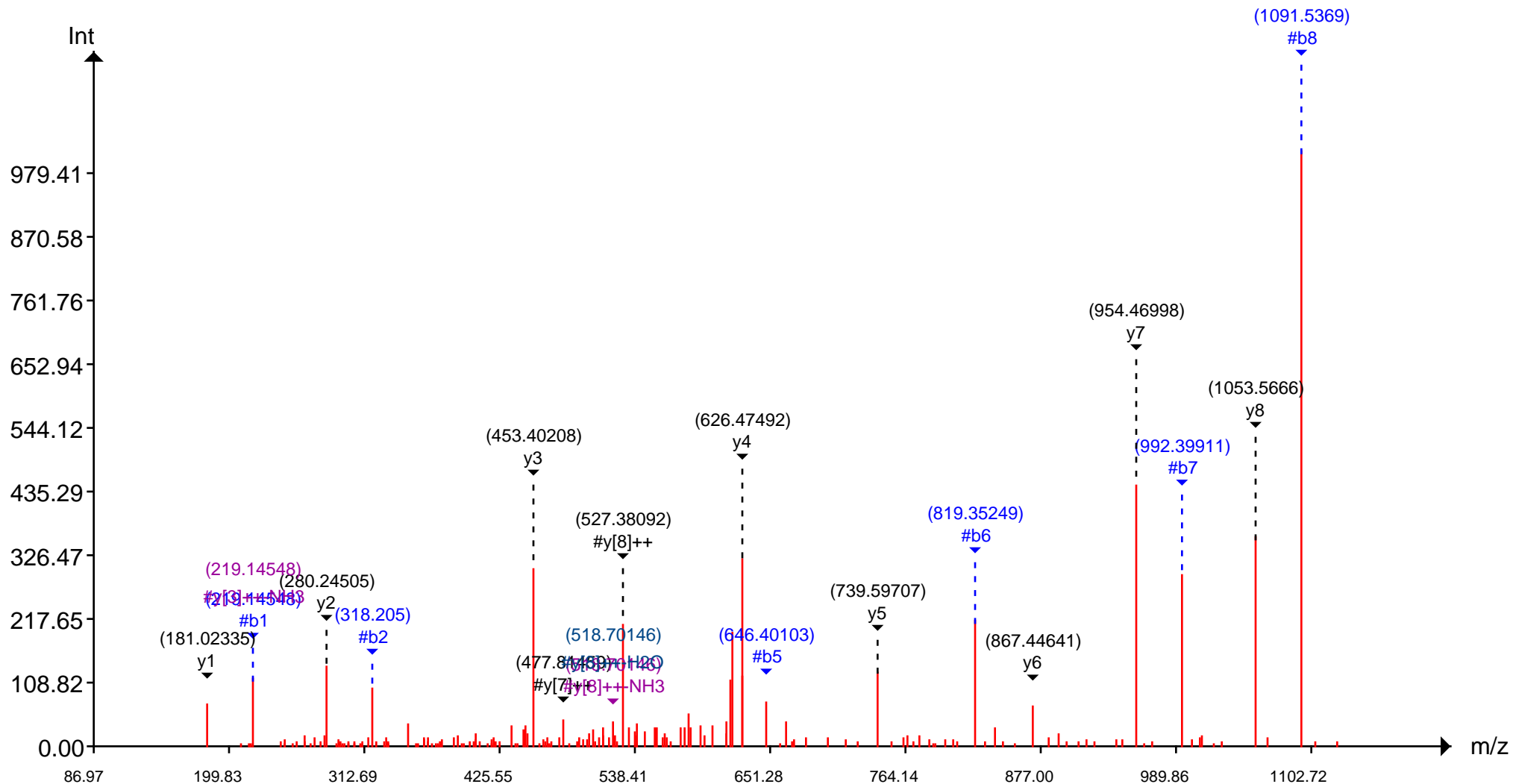
Modified Sequence	AcD3-LTEAPLNPK<AcD3K*>ANR-COOH	Ion score	47.96
Protein	P62737, P60710, P68033, P63260, P63268, P68134	Confidence level	99.00%
Precursor m/z	707.40254	Identity Threshold	46.72
Charge	2+	Mass error	0.00113



Info

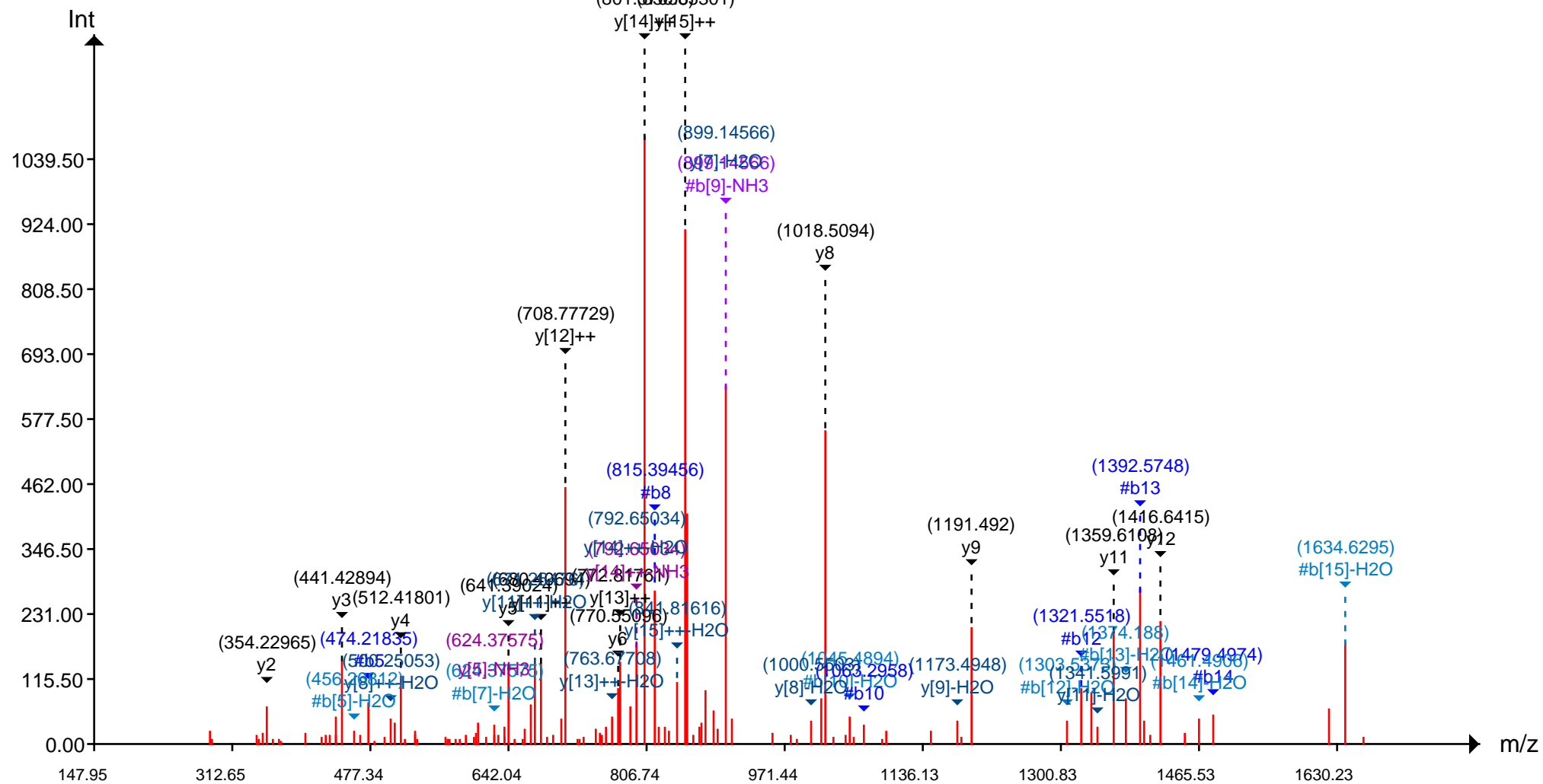
Modified Sequence	AcD3-LQETTLAANQPLSR-COOH	Ion score	105.13
Protein	O09159	Confidence level	99.00%
Precursor m/z	793.93269	Identity Threshold	47.74
Charge	2+	Mass error	5.47E-4





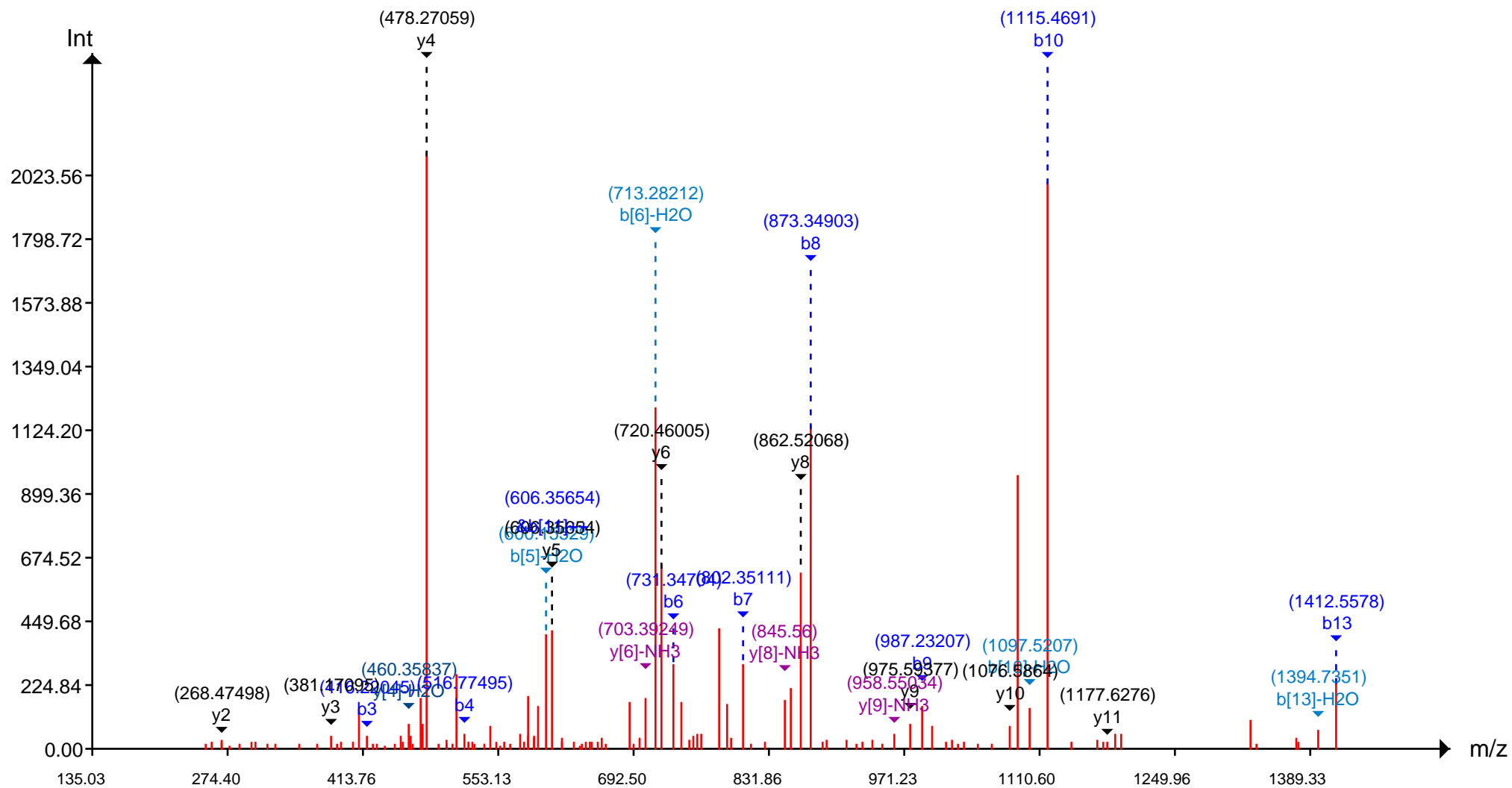
Info

Modified Sequence	AcD3-K<AcD3K*>VSQIK<AcD3K*>K<AcD3K*>VR<C13>-COOH	Ion score	59.83
Protein	P26638	Confidence level	99.00%
Precursor m/z	636.43025	Identity Threshold	33.36
Charge	2+	Mass error	1.69E-4



Info

Modified Sequence	AcD3-SVGQGPAK<AcD3K*>TM<Mox*>EEASK<AcD3K*>R<C13>-COOH	Ion score	62.6
Protein	O70310	Confidence level	99.00%
Precursor m/z	916.97767	Identity Threshold	48.17
Charge	2+	Mass error	0.00164



Info

Modified Sequence AcD3-LQETTLAANQPLSR<C13>-COOH

Ion score 55.88

Protein O09159

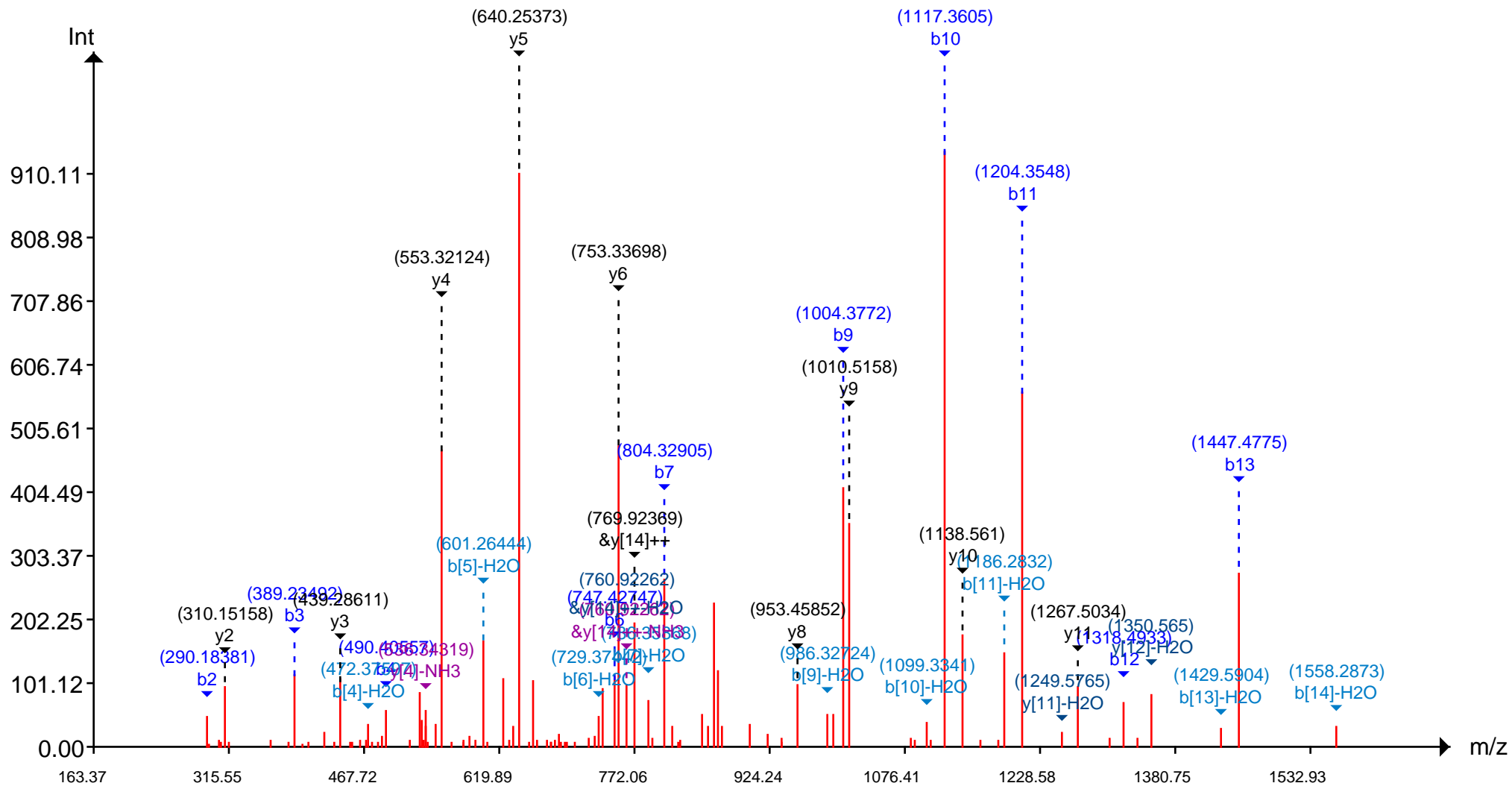
Confidence level 99.00%

Precursor m/z 796.94379

Identity Threshold 47.28

Charge 2+

Mass error 0.002606



Info

Modified Sequence AcD3-K<AcD3K*>AVTEQGAELSNEER<C13>-COOH

Ion score 92.61

Protein P68254

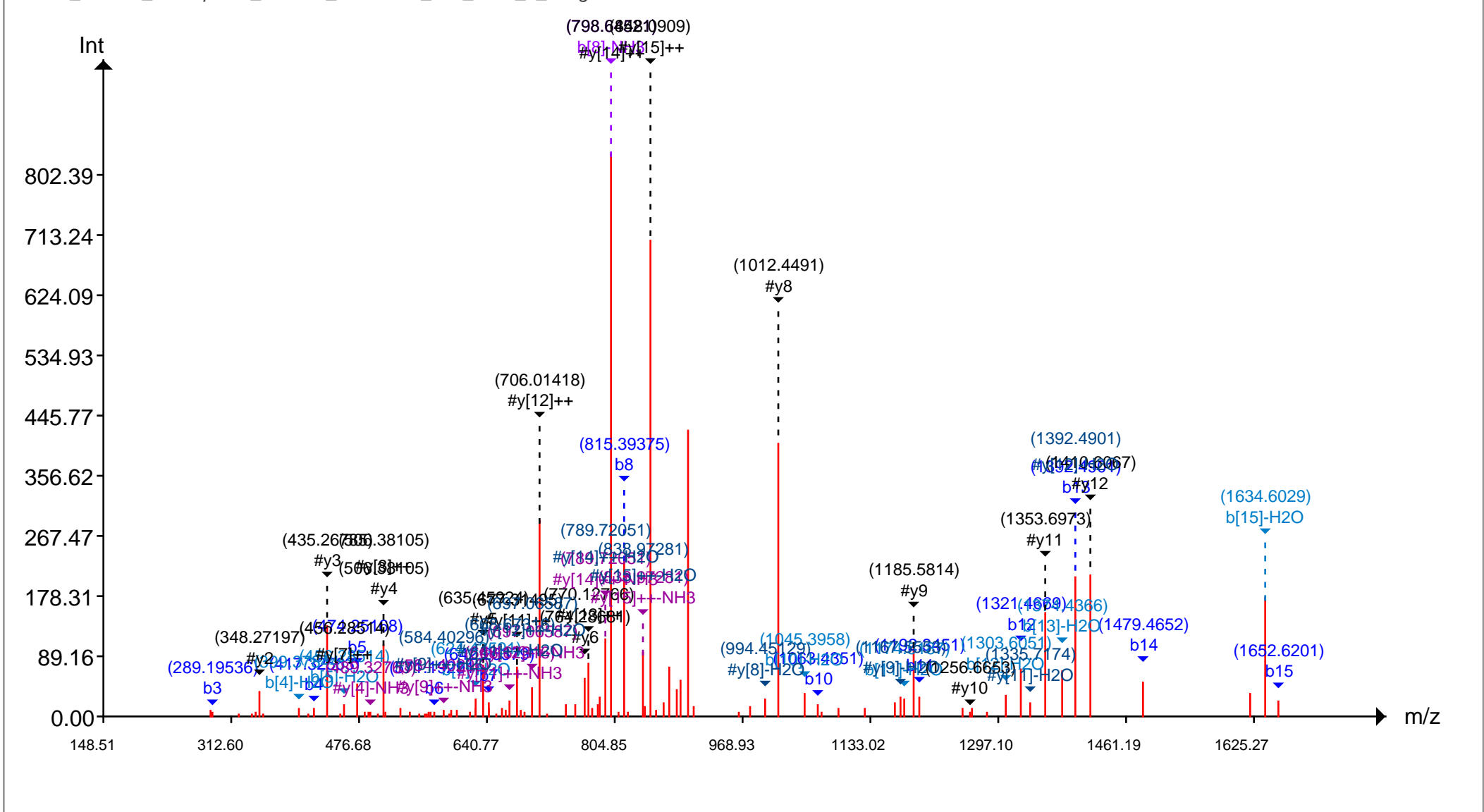
Confidence level 99.00%

Precursor m/z 878.95074

Identity Threshold 48.09

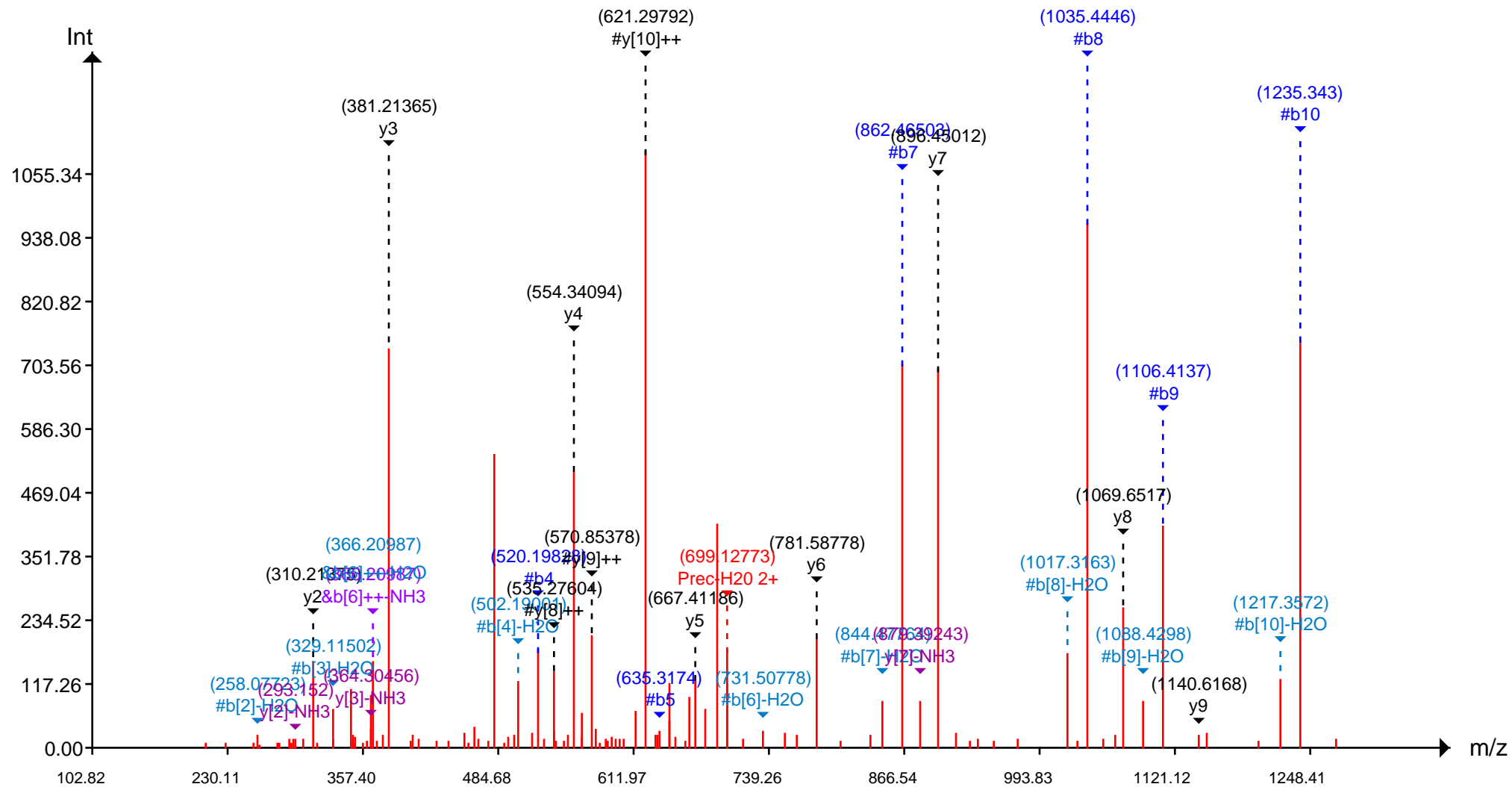
Charge 2+

Mass error 0.001644



Info

Modified Sequence	AcD3-SVGQGPAK<AcD3K*>TM<Mox*>EEASK<AcD3K*>R-COOH	Ion score	93.13
Protein	O70310	Confidence level	99.00%
Precursor m/z	913.96799	Identity Threshold	48.03
Charge	2+	Mass error	0.002422



Info

Modified Sequence AcD3-ETAK<AcD3K*>DNLK<AcD3K*>AER<C13>-COOH

Ion score 53.25

Protein Q9CVD2

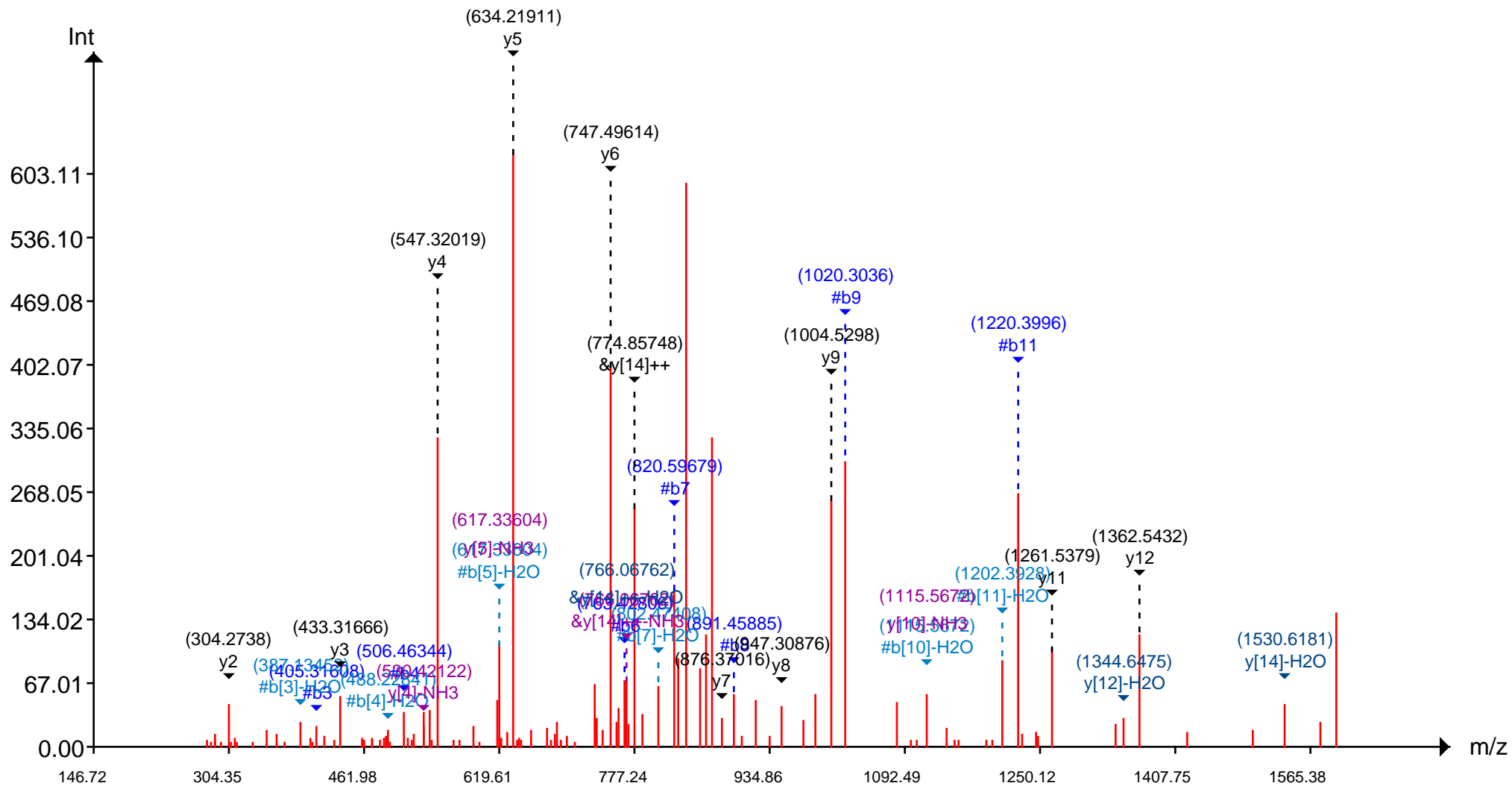
Confidence level 99.00%

Precursor m/z 708.39342

Identity Threshold 47.67

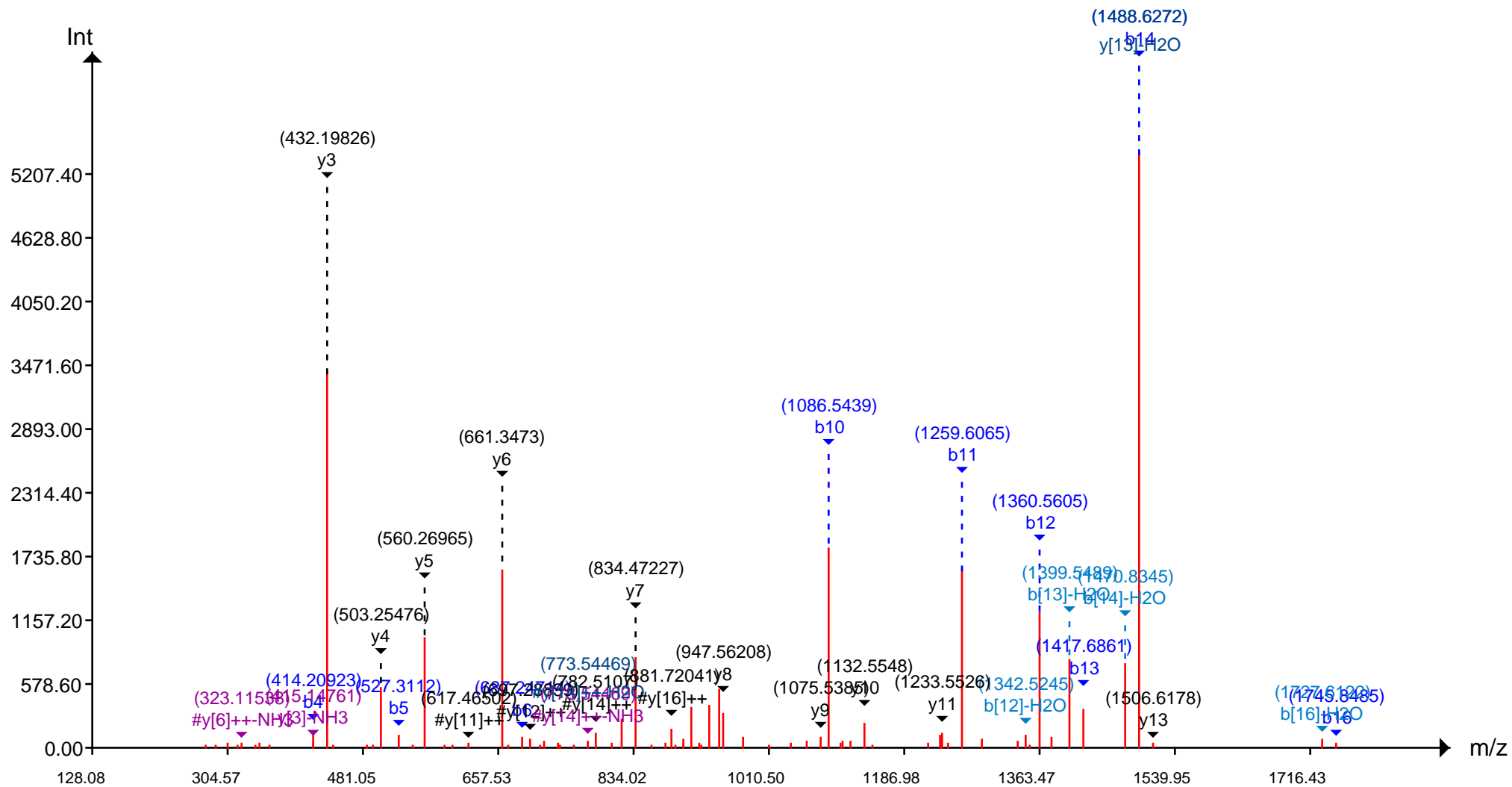
Charge 2+

Mass error 0.001368



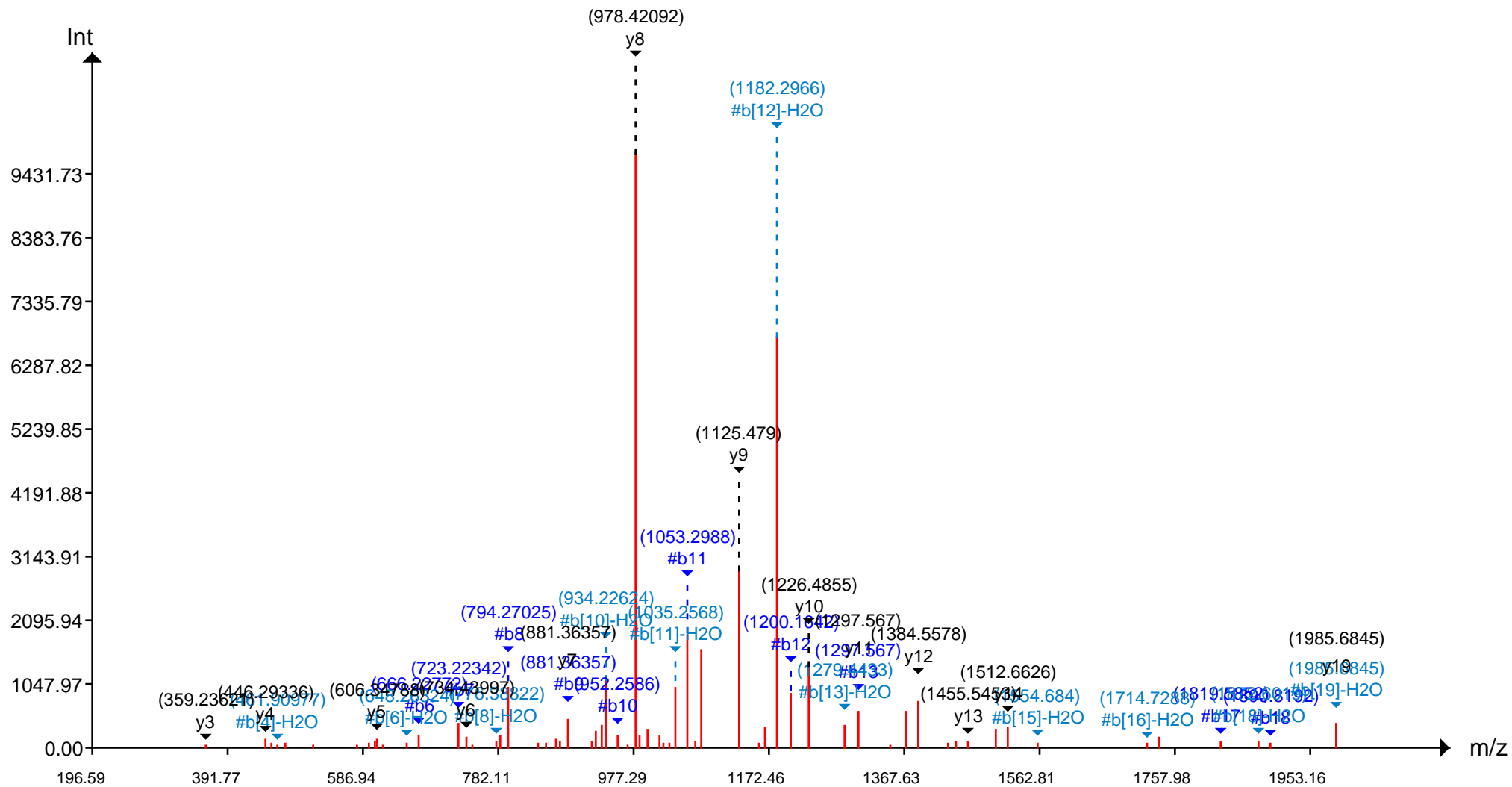
Info

Modified Sequence	AcD3-K<AcD3K*>SVTEQGAELSNEER-COOH	Ion score	60.33
Protein	P63101	Confidence level	99.00%
Precursor m/z	883.93833	Identity Threshold	47.65
Charge	2+	Mass error	0.002046



Info

Modified Sequence	AcD3-LVVGLC<Cmm*>TGQIK<AcD3K*>TGAPC<Cmm*>R-COOH	Ion score	78.75
Protein	P17182, P21550, P17183	Confidence level	99.00%
Precursor m/z	960.5204	Identity Threshold	47.57
Charge	2+	Mass error	0.002276



Info

Modified Sequence AcD3-FETGDAGASATFPM<Mox*>QC<Cmm*>SALR-COOH

Ion score 112.8

Protein P63242

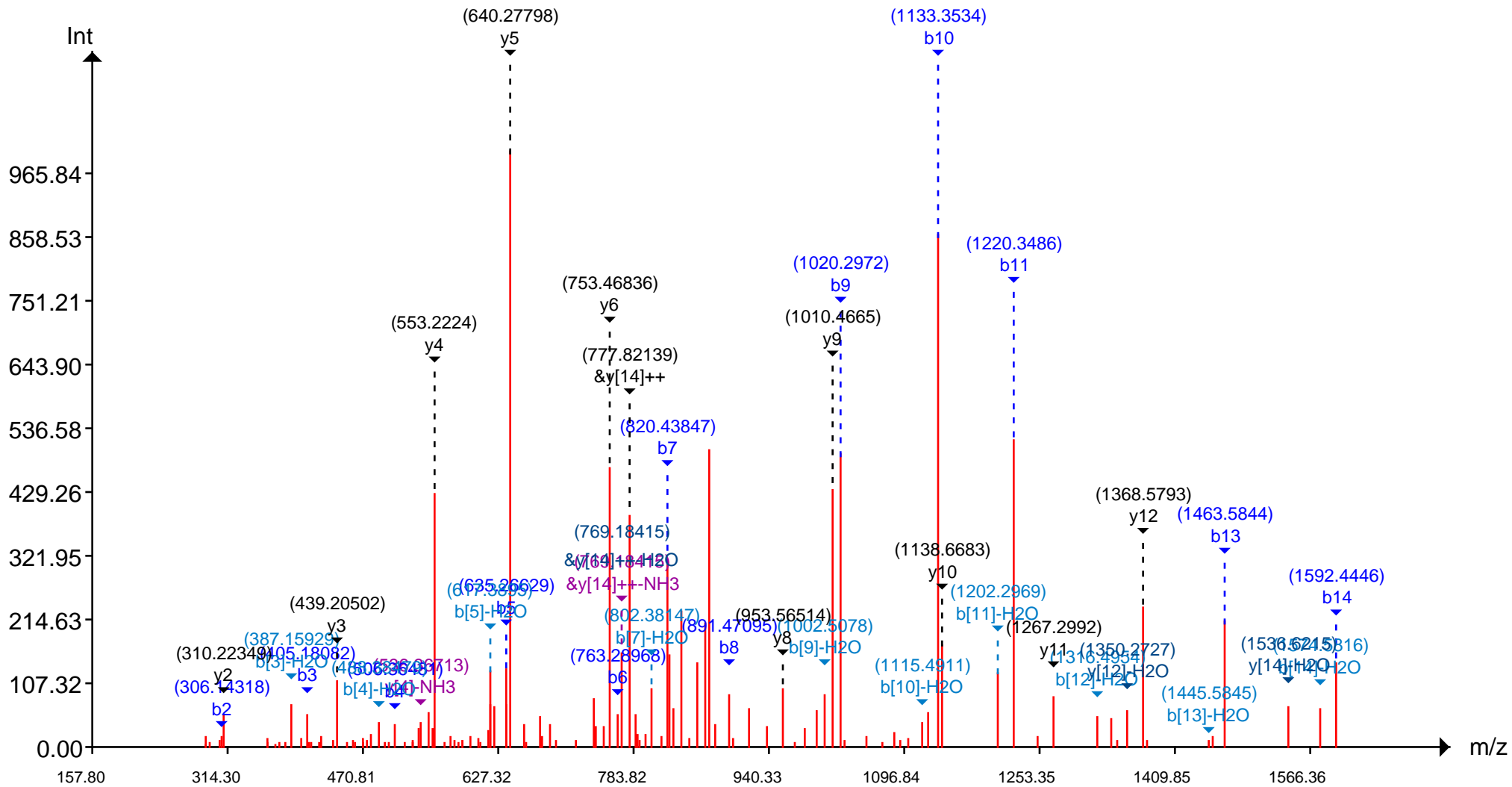
Confidence level 99.00%

Precursor m/z 1089.4891

Identity Threshold 43.25

Charge 2+

Mass error 0.003962



Info

Modified Sequence AcD3-K<AcD3K*>SVTEQGAELSNEER<C13>-COOH

Ion score 104.01

Protein P63101

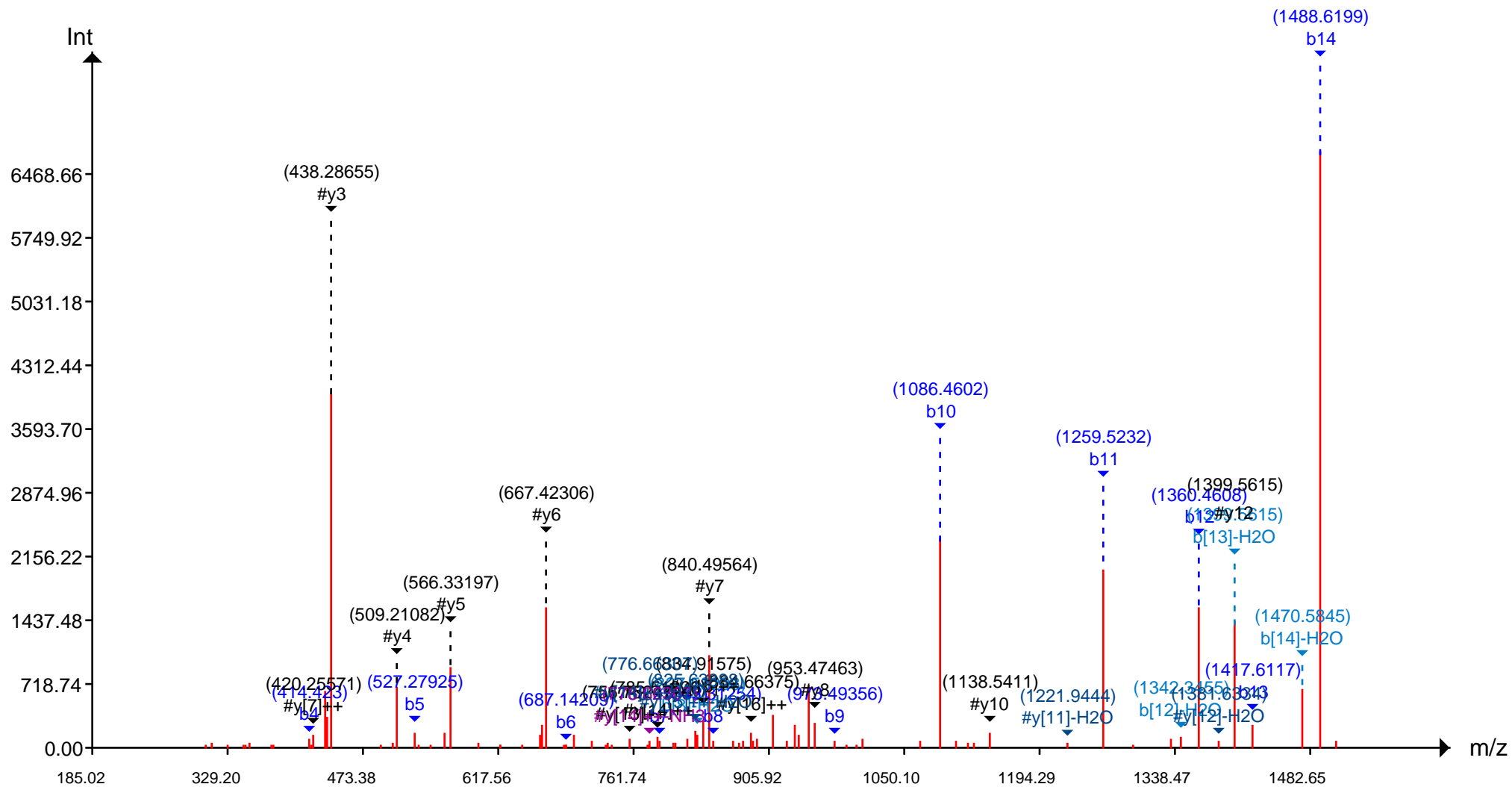
Confidence level 99.00%

Precursor m/z 886.94826

Identity Threshold 47.94

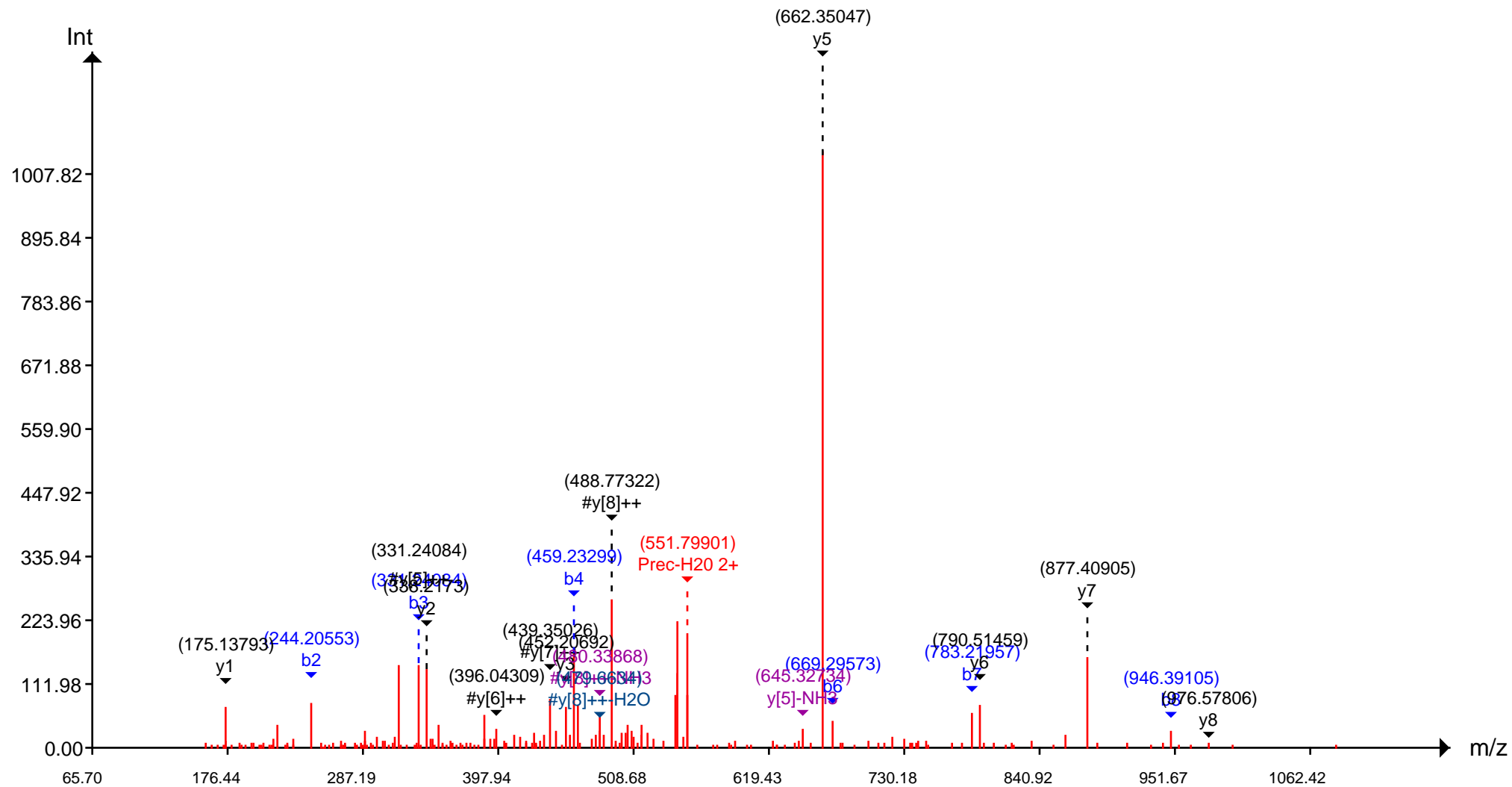
Charge 2+

Mass error 0.001765



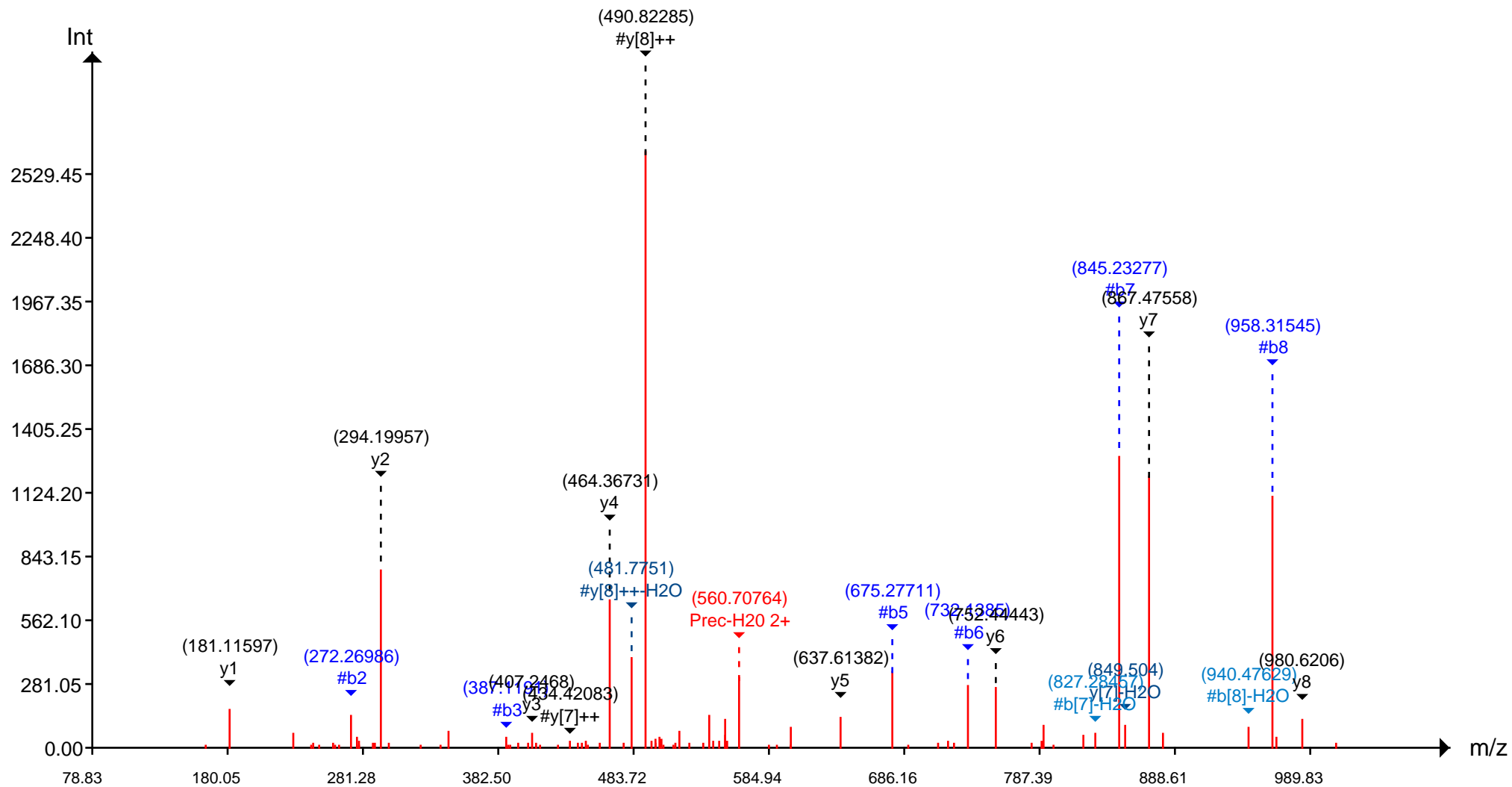
Info

Modified Sequence	AcD3-LVVGLC<Cmm*>TGQIK<AcD3K*>TGAPC<Cmm*>R<C13>-COOH	Ion score	56.47
Protein	P17182, P21550, P17183	Confidence level	99.00%
Precursor m/z	963.53016	Identity Threshold	47.17
Charge	2+	Mass error	0.001655



Info

Modified Sequence	AcD3-VVSQPLNYR-COOH	Ion score	50.61
Protein	Q9JLJ2	Confidence level	99.00%
Precursor m/z	560.81308	Identity Threshold	47.33
Charge	2+	Mass error	3.6E-5



Info

Modified Sequence AcD3-IIDDK<AcD3K*>GILR<C13>-COOH

Protein P35700

Precursor m/z 569.85547

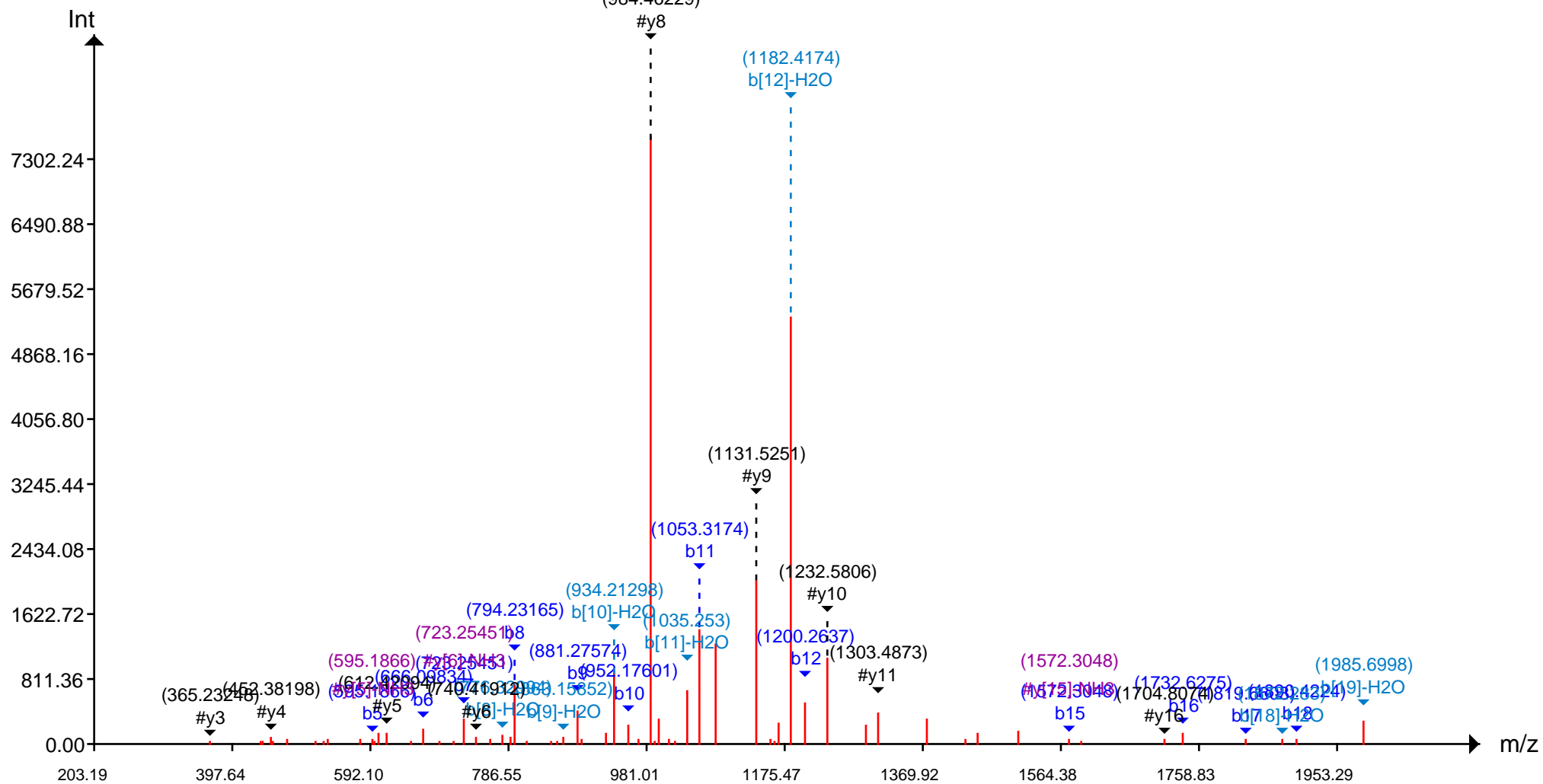
Charge 2+

Ion score 68.47

Confidence level 99.00%

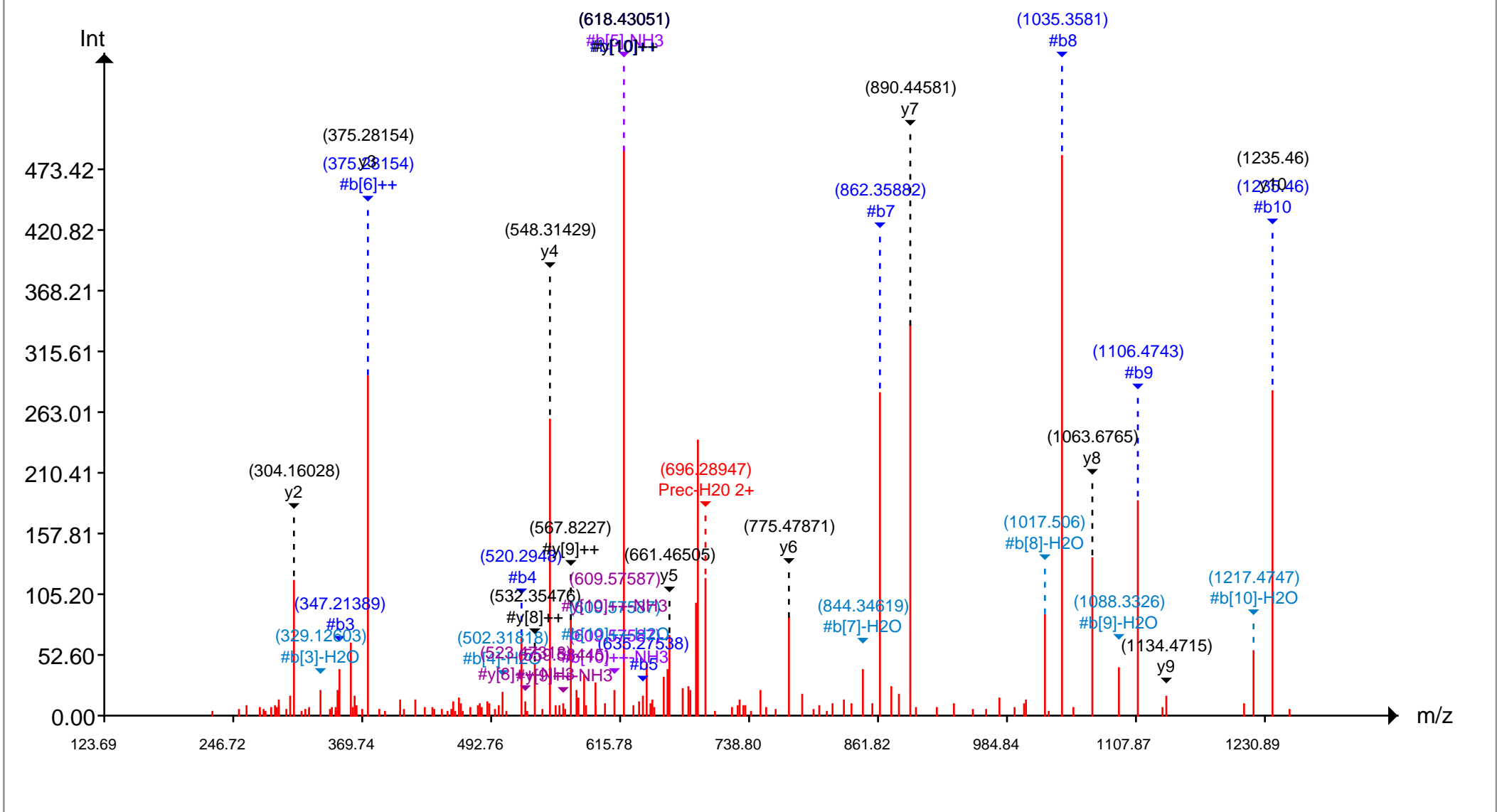
Identity Threshold 42.64

Mass error -7.71E-4



Info

Modified Sequence	AcD3-FETGDAGASATFPM<Mox*>QC<Cmm*>SALR<C13>-COOH	Ion score	100.39
Protein	P63242	Confidence level	99.00%
Precursor m/z	1092.4986	Identity Threshold	44.22
Charge	2+	Mass error	0.00282



Info

Modified Sequence AcD3-ETAK<AcD3K*>DNLK<AcD3K*>AER-COOH

Ion score 62.91

Protein Q9CVD2

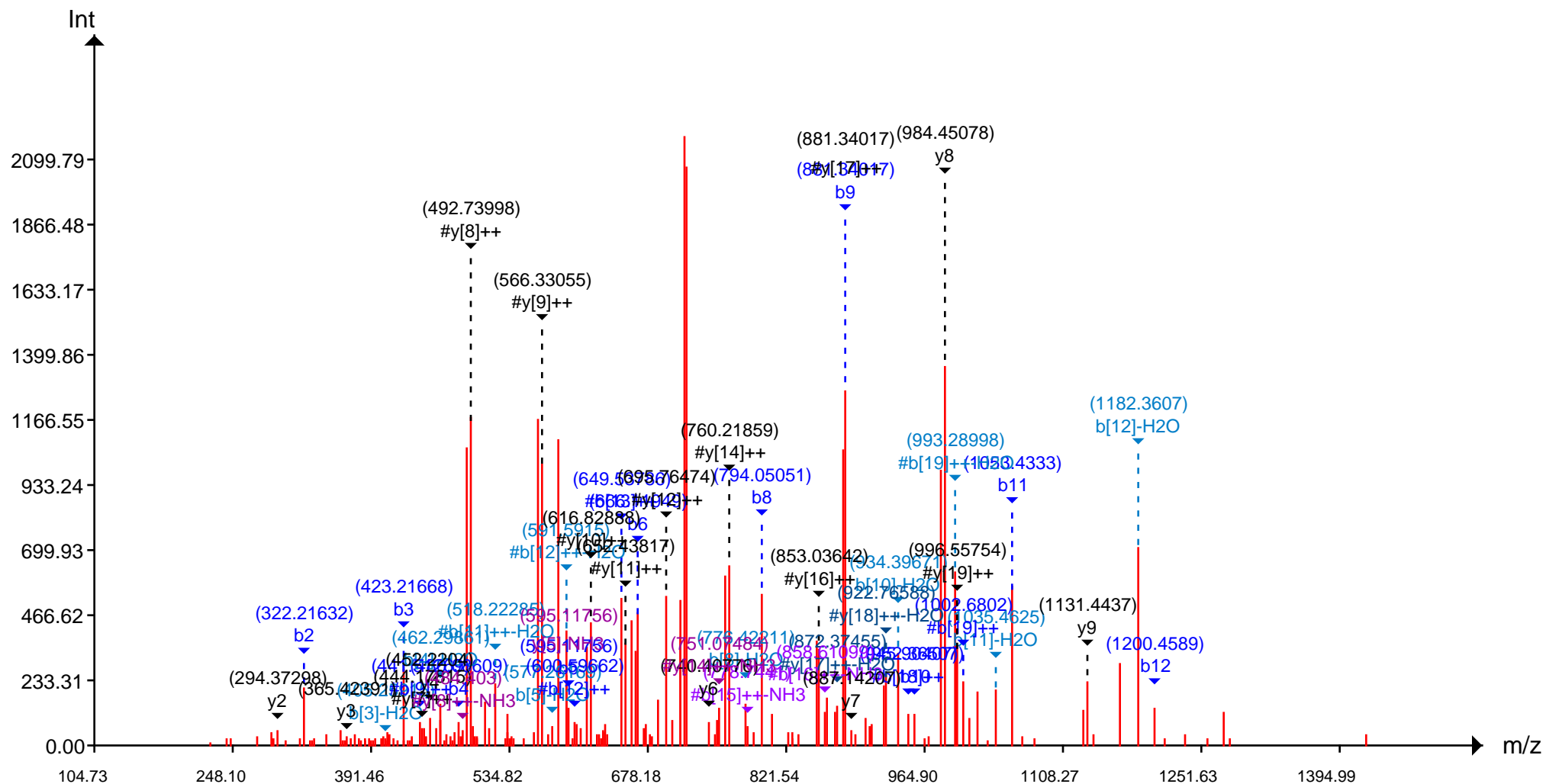
Confidence level 99.00%

Precursor m/z 705.38325

Identity Threshold 47.90

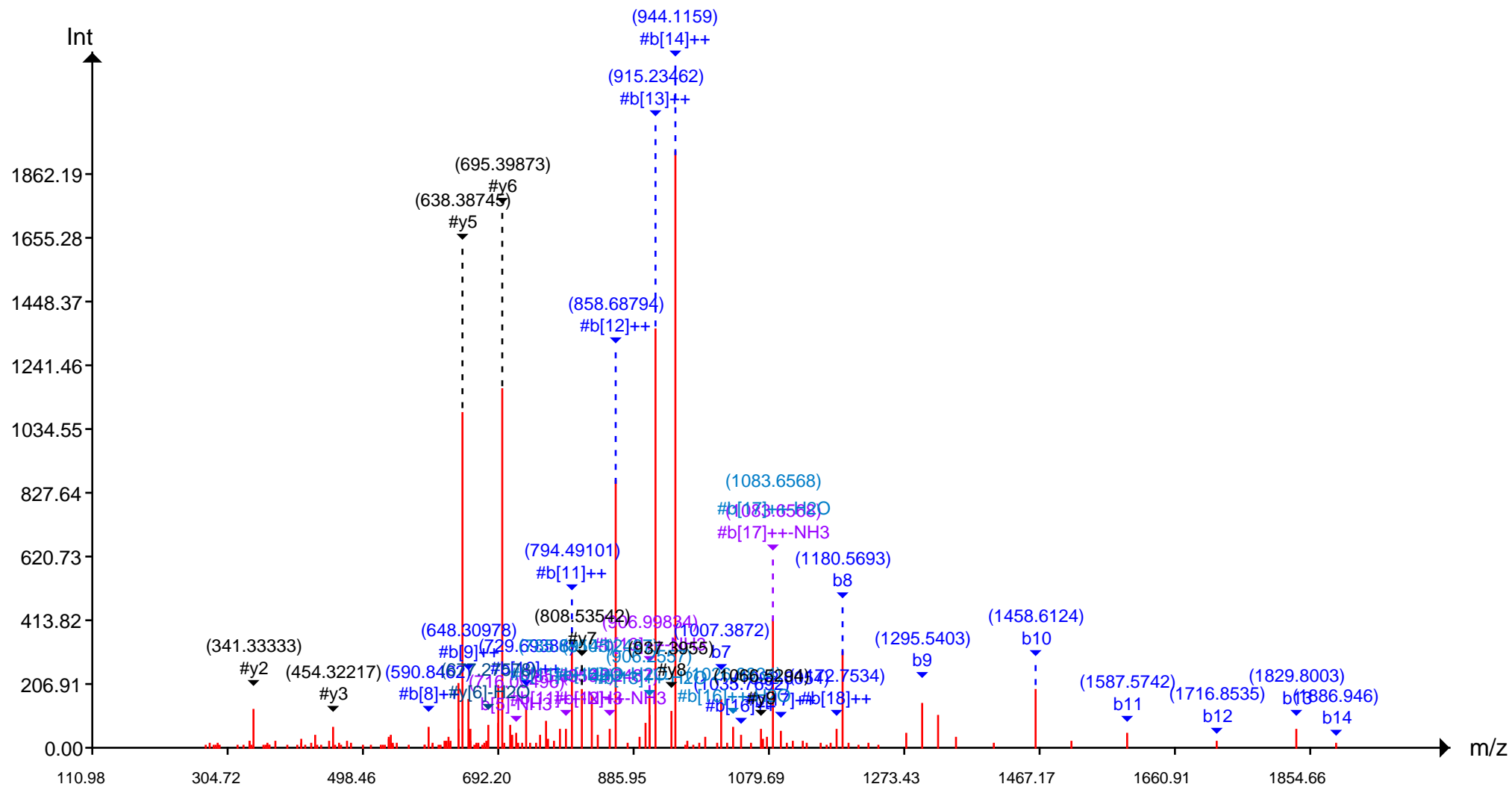
Charge 2+

Mass error 0.00117



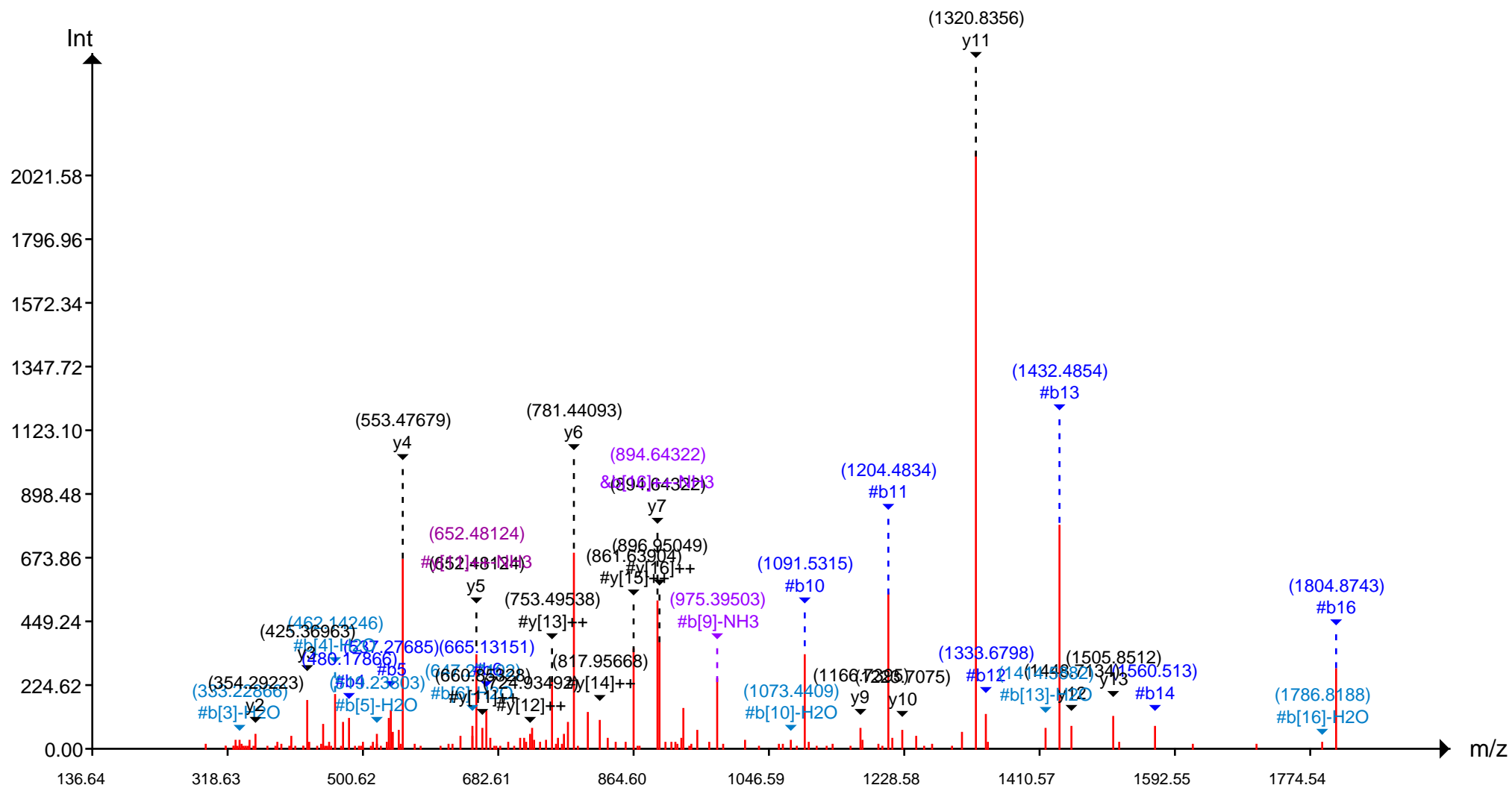
Info

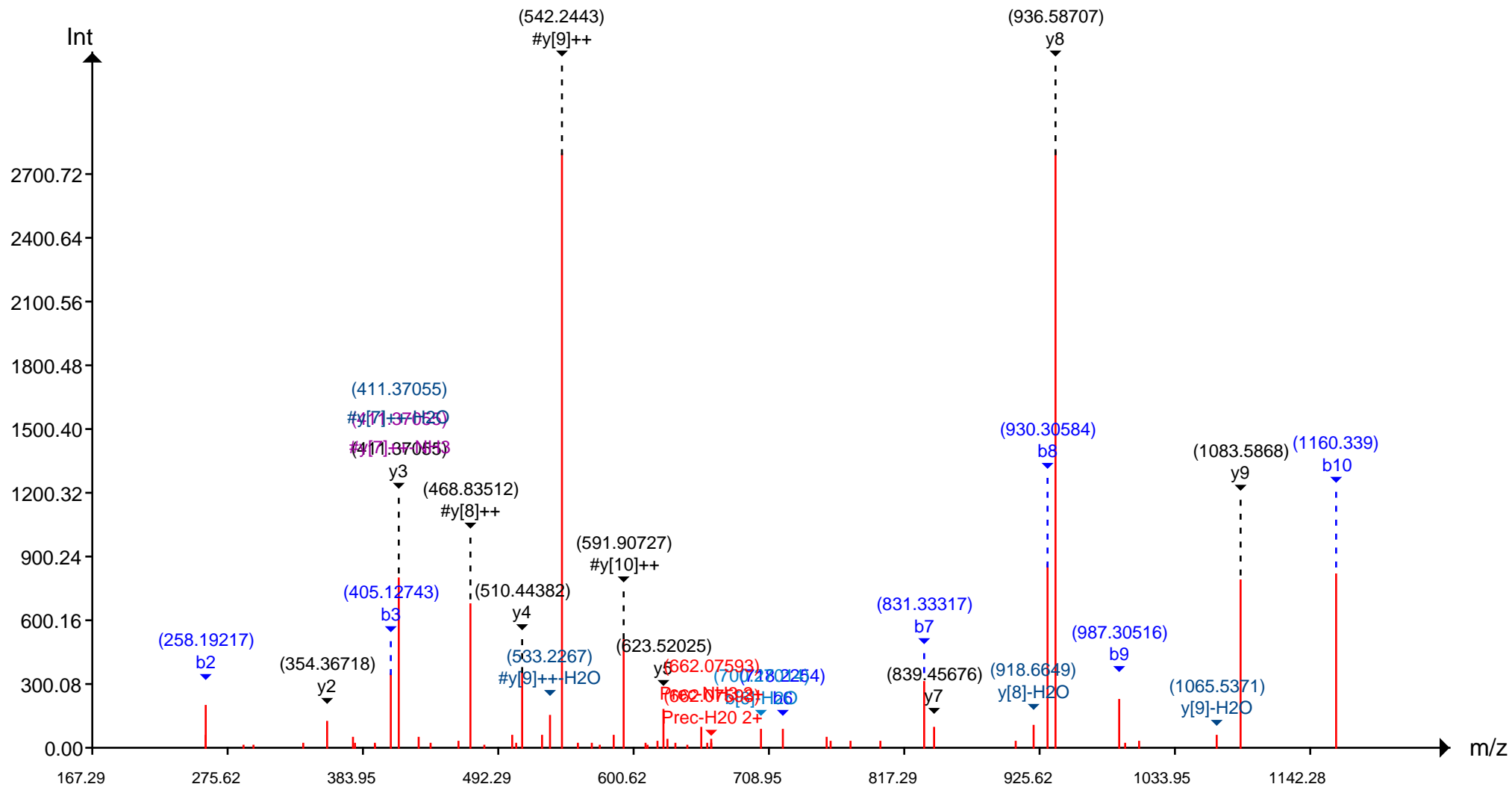
Modified Sequence	AcD3-FETGDAGASATFPM<Mox*>QC<Cmm*>SALR<C13>-COOH	Ion score	44.64
Protein	P63242	Confidence level	99.00%
Precursor m/z	728.6671	Identity Threshold	43.99
Charge	3+	Mass error	-3.56E-4



Info

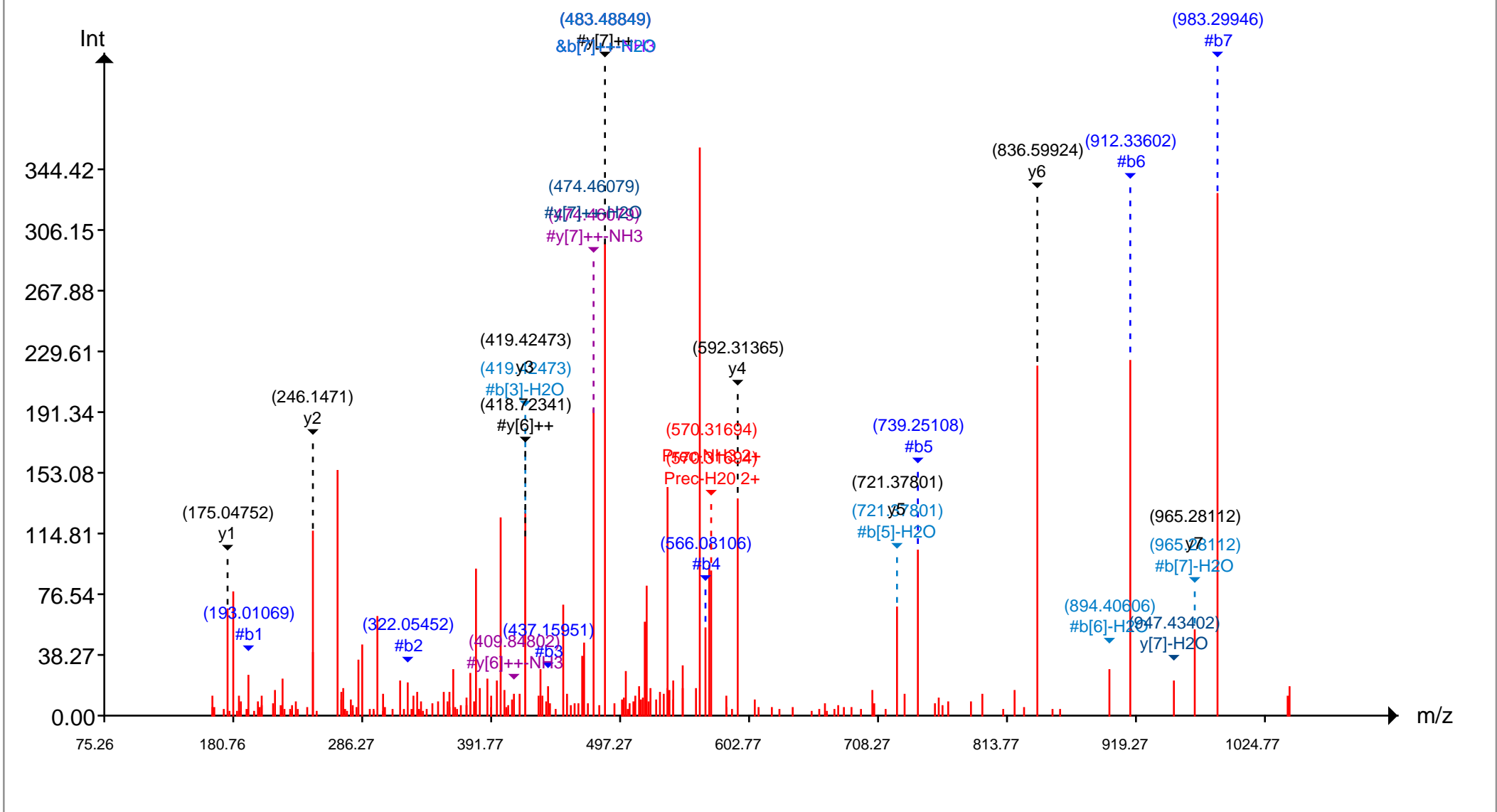
Modified Sequence	AcD3-YQVC<Cmm*>HTK<AcD3K*>K<AcD3K*>DYEEIGPSIC<Cmm*>R<C13>-COOH	Ion score	64.19
Protein	Q99JY9	Confidence level	99.00%
Precursor m/z	842.08116	Identity Threshold	47.09
Charge	3+	Mass error	0.00368





Info

Modified Sequence	AcD3-LVFPSEIVGK<AcD3K*>R<C13>-COOH	Ion score	73.75
Protein	P62082	Confidence level	99.00%
Precursor m/z	670.91184	Identity Threshold	42.50
Charge	2+	Mass error	0.001343



Info

Modified Sequence AcD3-FEDEK<AcD3K*>K<AcD3K*>AR-COOH

Ion score 49.08

Protein P50580

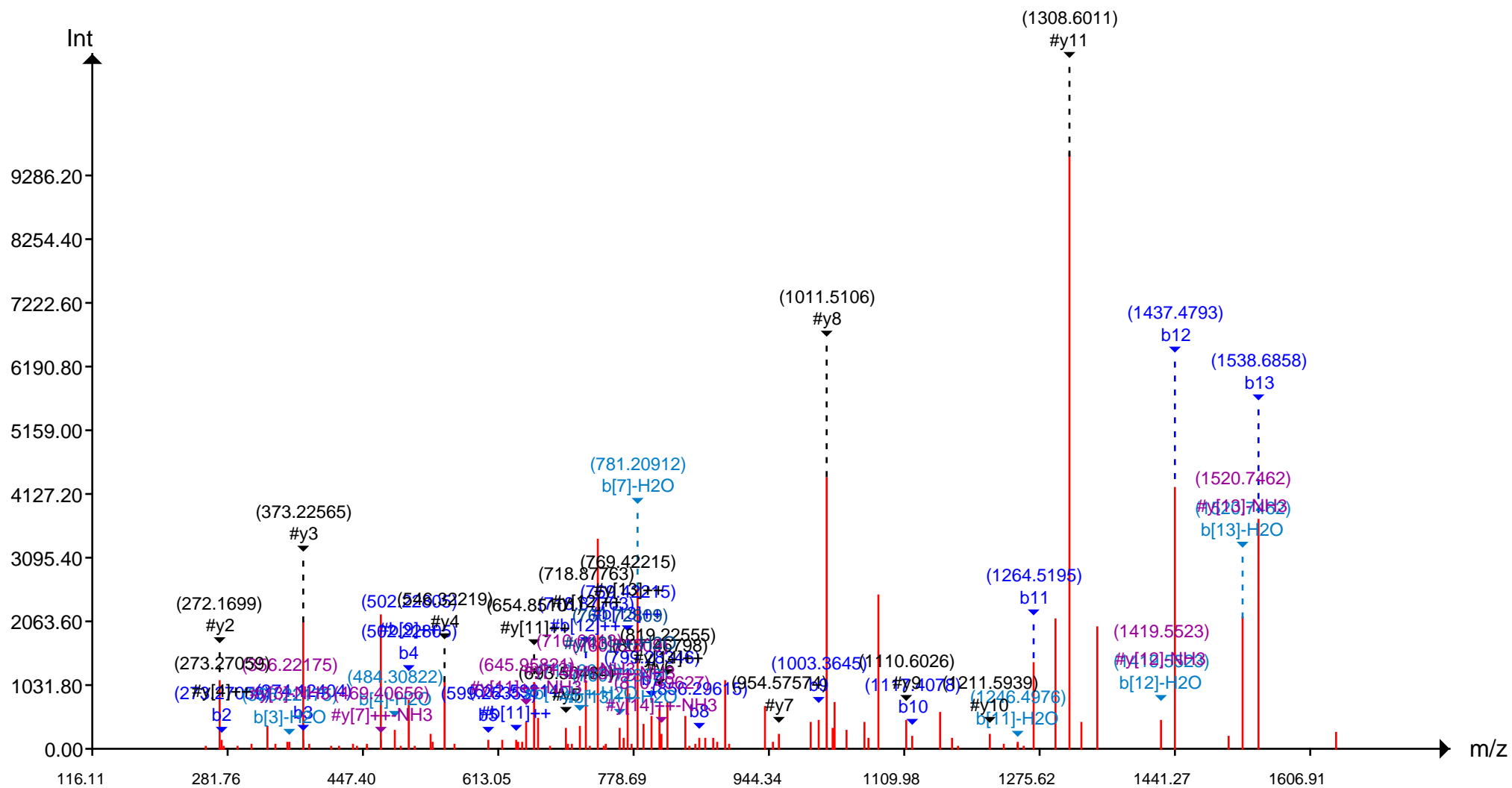
Confidence level 99.00%

Precursor m/z 579.3114

Identity Threshold 47.86

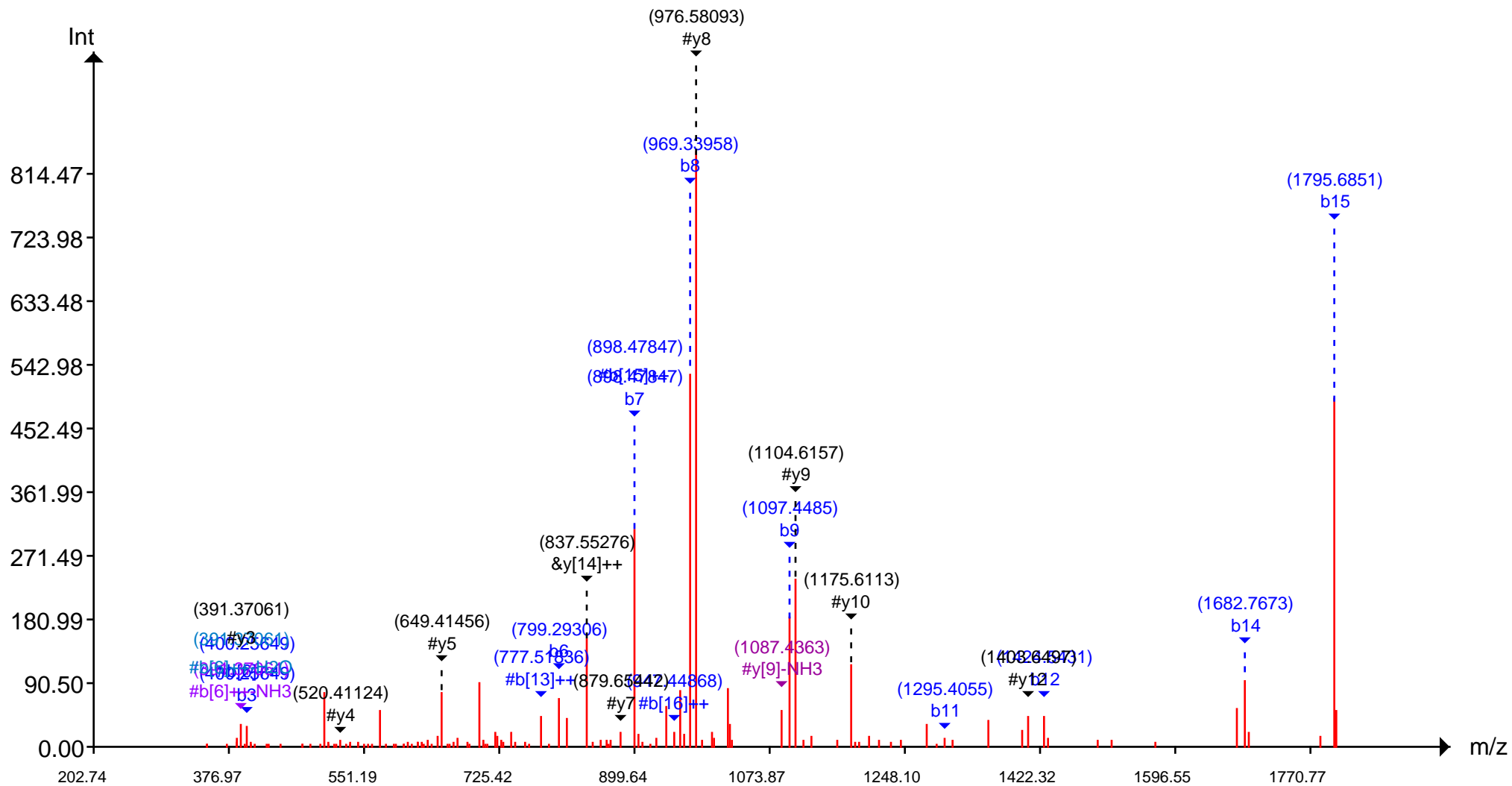
Charge 2+

Mass error 8.26E-4



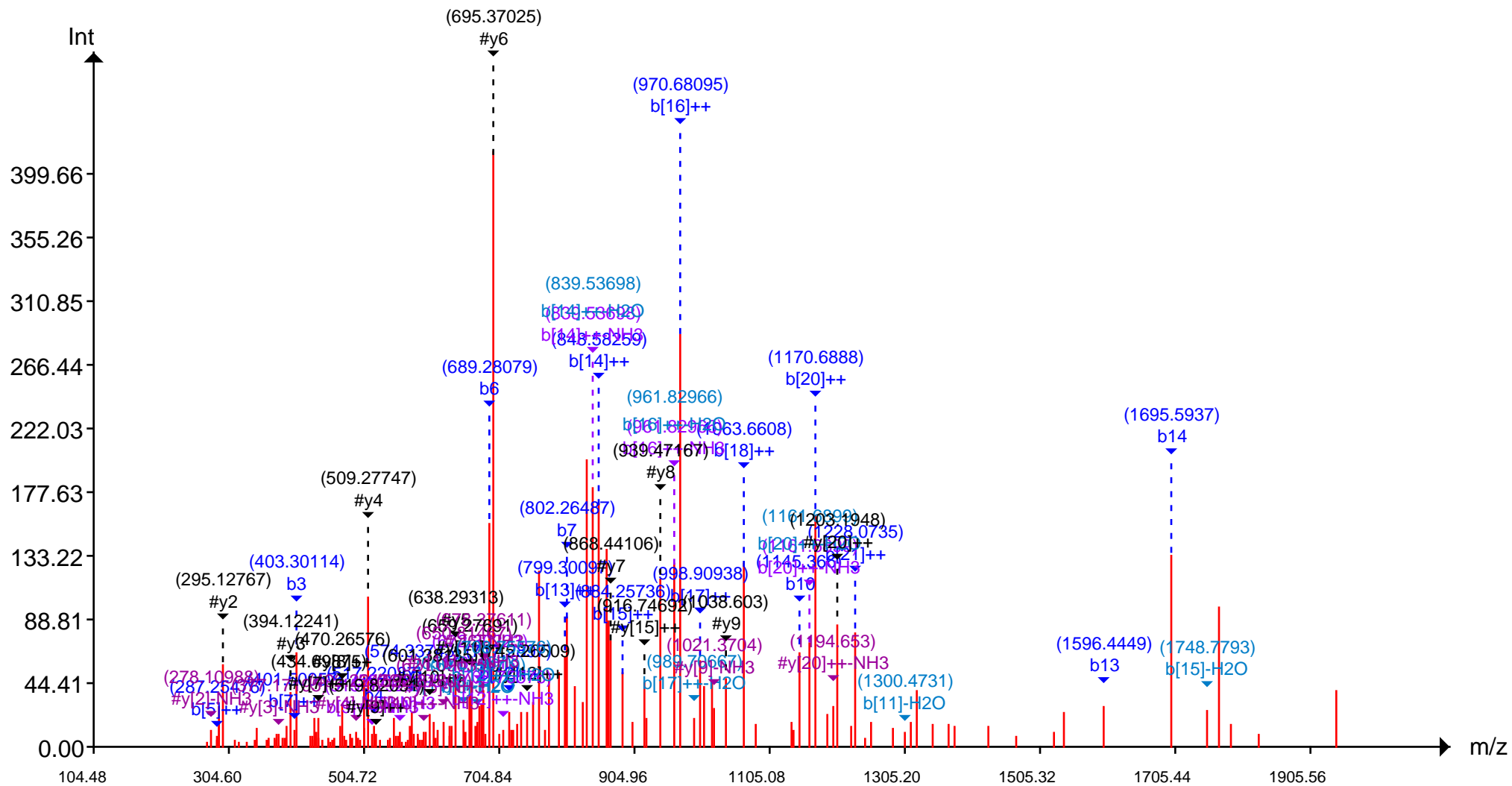
Info

Modified Sequence	AcD3-QVTQPTVGM<Mox*>NFK<AcD3K*>TPR-COOH	Ion score	64.64
Protein	P63276	Confidence level	99.00%
Precursor m/z	905.47666	Identity Threshold	48.05
Charge	2+	Mass error	0.002565



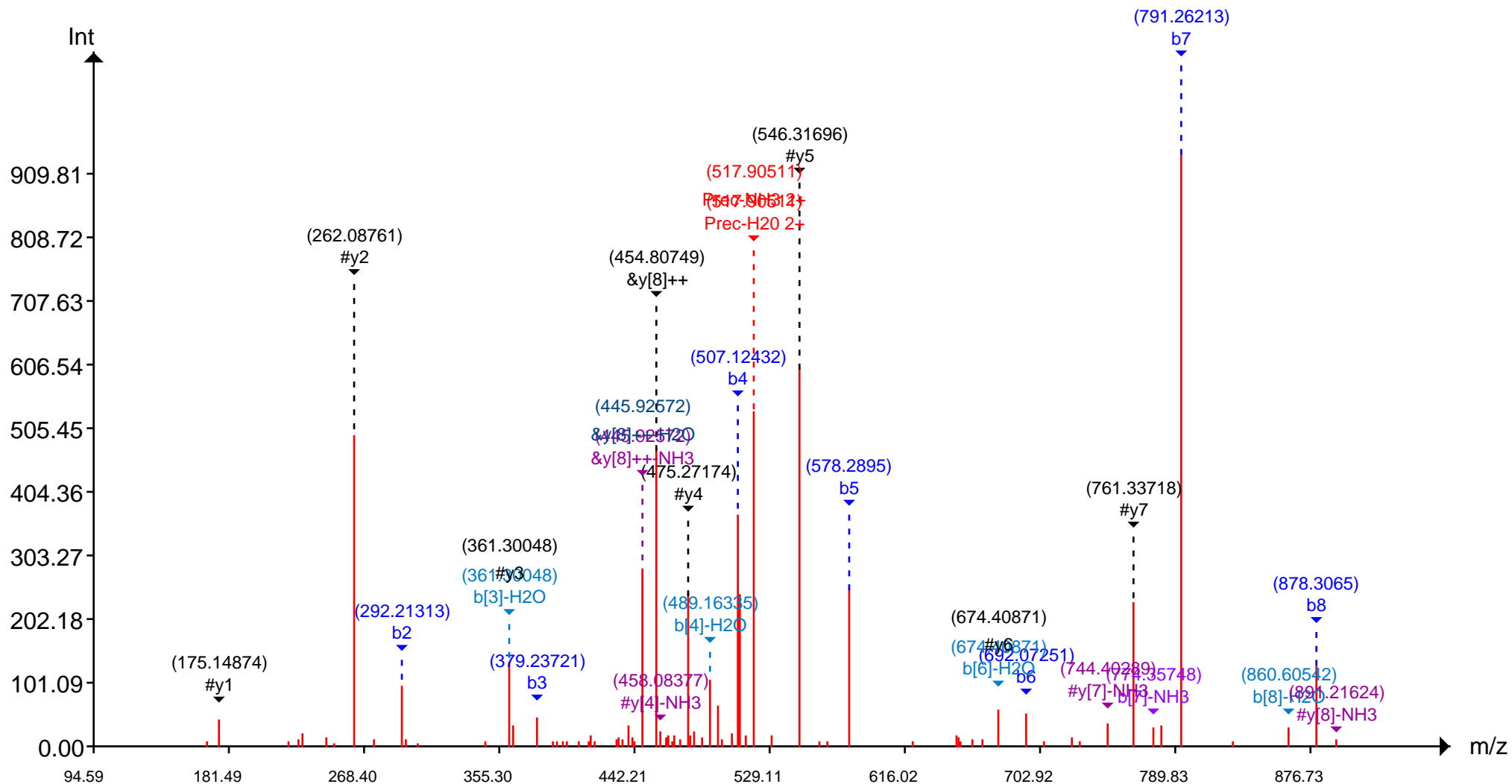
Info

Modified Sequence	AcD3-LQLPK<AcD3K*>EVAQPTEEELPR<C13>-COOH	Ion score	53.61
Protein	Q9ERD8	Confidence level	99.00%
Precursor m/z	1037.0766	Identity Threshold	46.53
Charge	2+	Mass error	0.001899



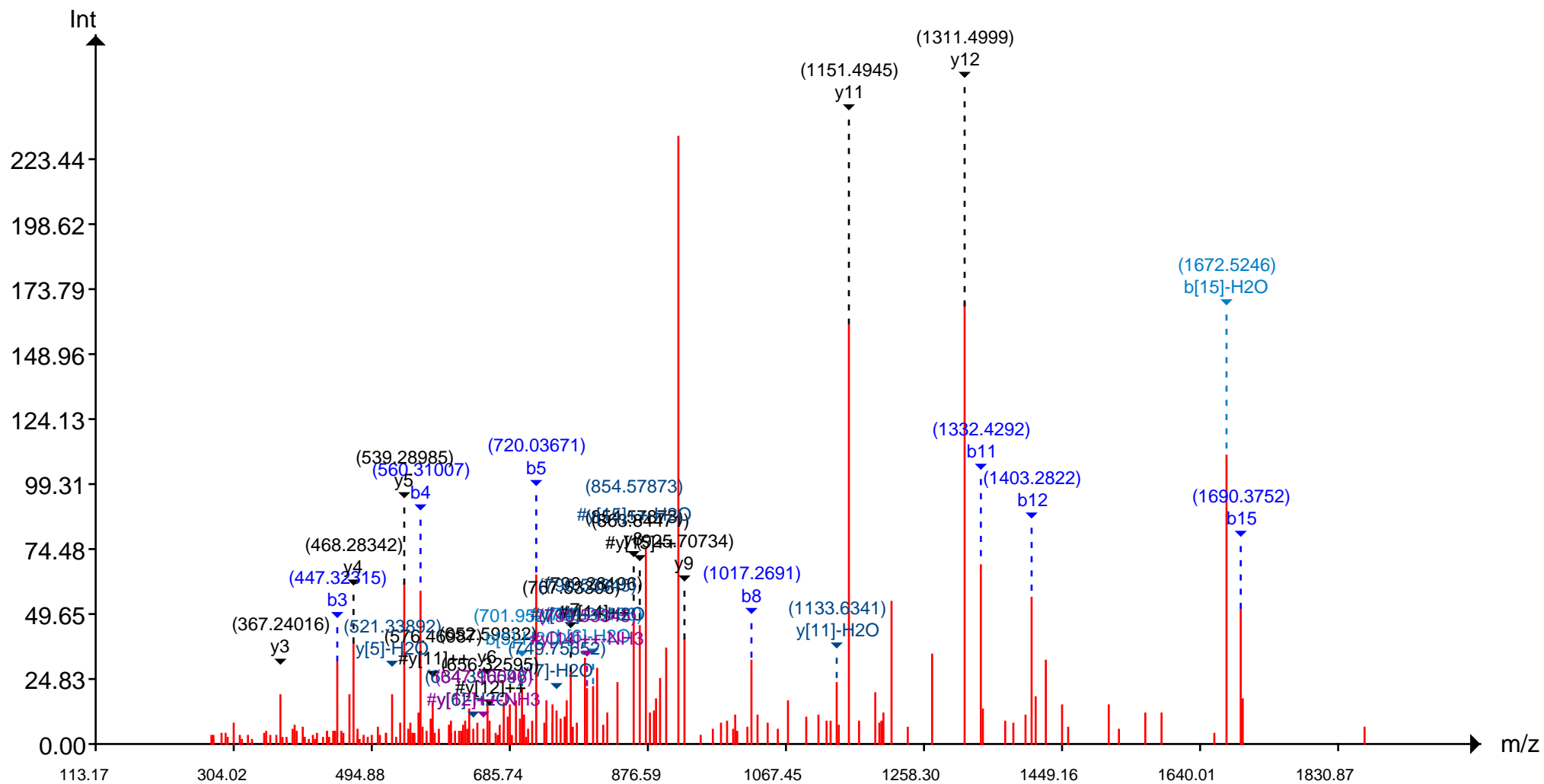
Info

Modified Sequence	AcD3-ALK<AcD3K*>NGDLDEVK<AcD3K*>DYVAK<AcD3K*>GEDVNR<C13>-COOH	Ion score	49.86
Protein	P62774	Confidence level	99.00%
Precursor m/z	878.79718	Identity Threshold	47.70
Charge	3+	Mass error	0.002769



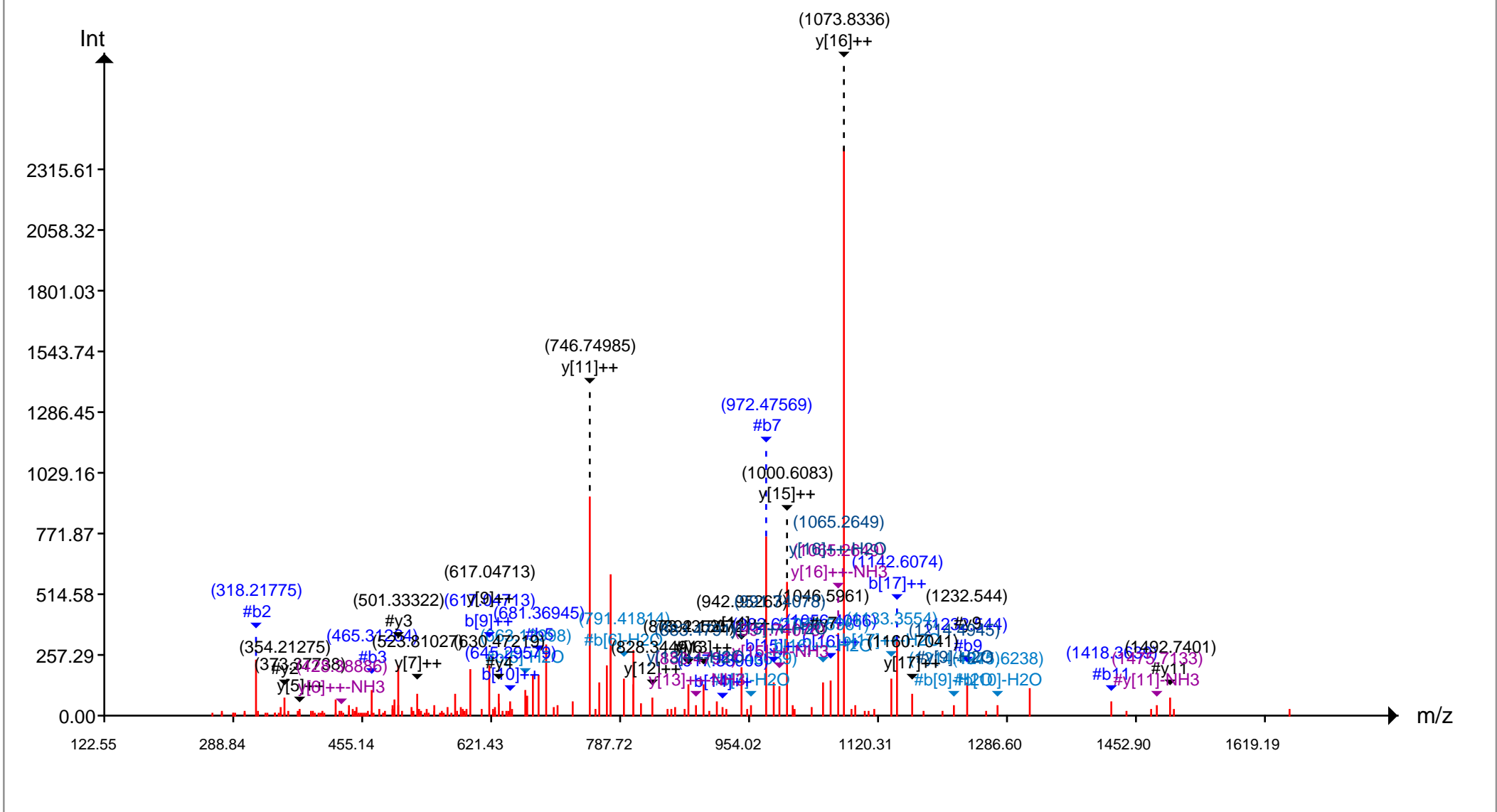
Info

Modified Sequence	AcD3-VM<Mox*>SQANVSR-COOH	Ion score	57.35
Protein	P70670, Q60817	Confidence level	99.00%
Precursor m/z	526.76547	Identity Threshold	47.53
Charge	2+	Mass error	4.27E-4



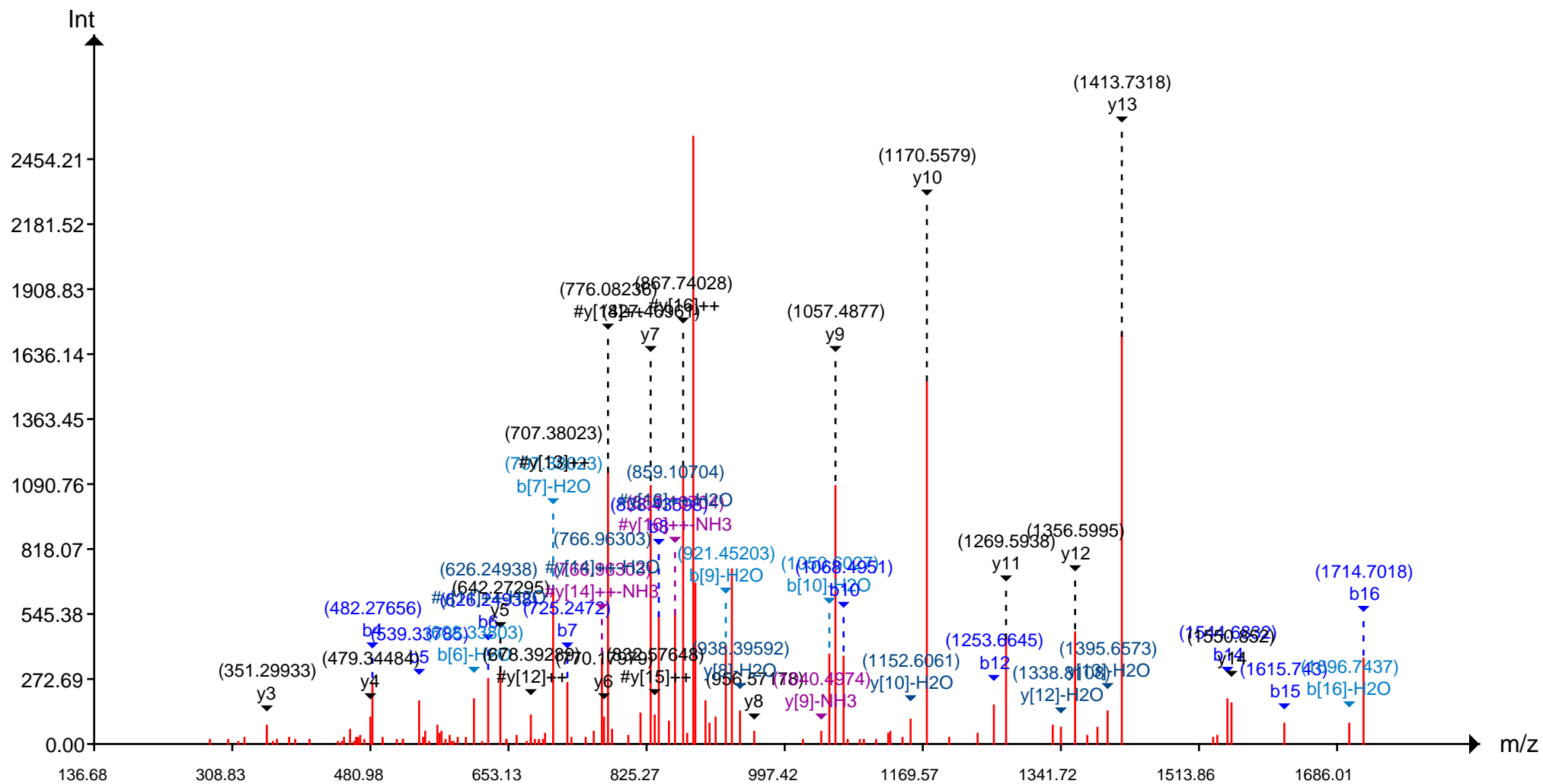
Info

Modified Sequence	AcD3-VEK<AcD3K*>LC<Cmm*>PEASDIATSVR<C13>-COOH	Ion score	52.85
Protein	Q8BIJ7	Confidence level	99.00%
Precursor m/z	935.99513	Identity Threshold	48.00
Charge	2+	Mass error	0.003693



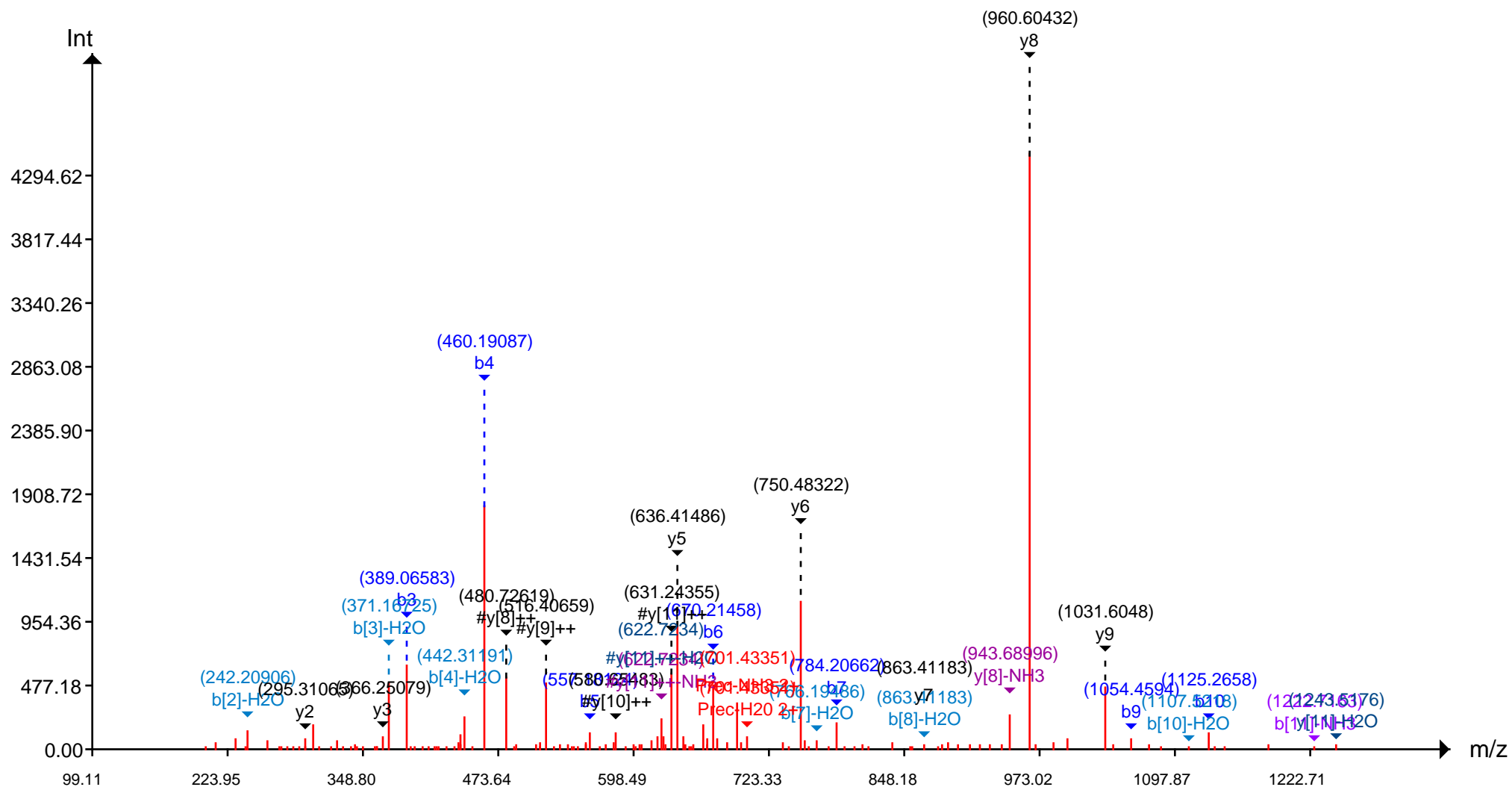
Info

Modified Sequence	AcD3-VK<AcD3K*>FDTQYPYGEK<AcD3K*>QDEFK<AcD3K*>R<C13>-COOH	Ion score	49.74
Protein	P57759	Confidence level	99.00%
Precursor m/z	822.42362	Identity Threshold	47.97
Charge	3+	Mass error	1.004708



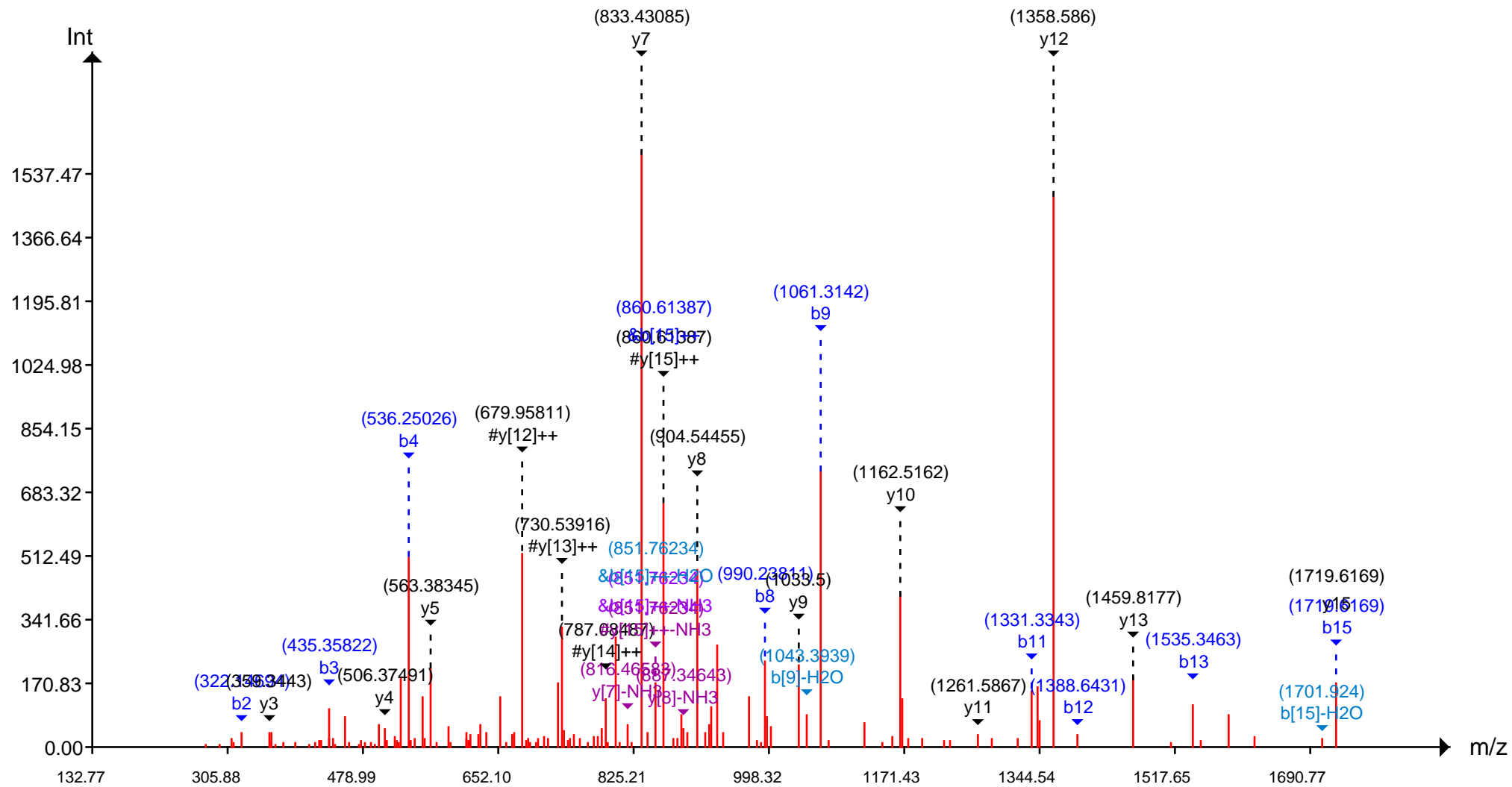
Info

Modified Sequence	AcD3-DALHGSLVLTGQYQAVR<C13>-COOH	Ion score	88.77
Protein	Q9EPB4	Confidence level	99.00%
Precursor m/z	947.99414	Identity Threshold	48.02
Charge	2+	Mass error	0.001774



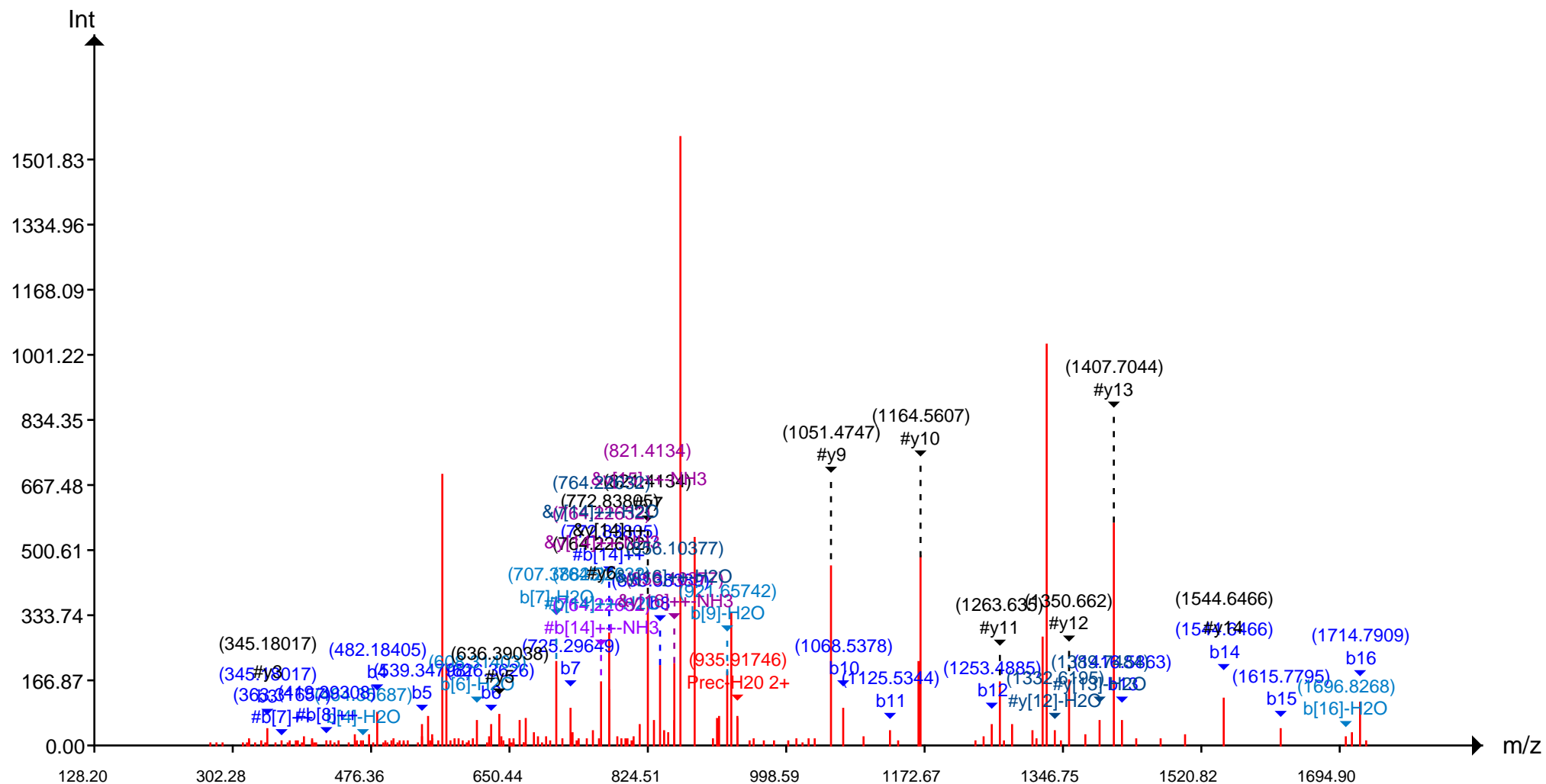
Info

Modified Sequence	AcD3-LTEAPLNPK<AcD3K*>ANR<C13>-COOH	Ion score	56.73
Protein	P62737, P60710, P68033, P63260, P63268, P68134	Confidence level	99.00%
Precursor m/z	710.4127	Identity Threshold	45.92
Charge	2+	Mass error	0.001308



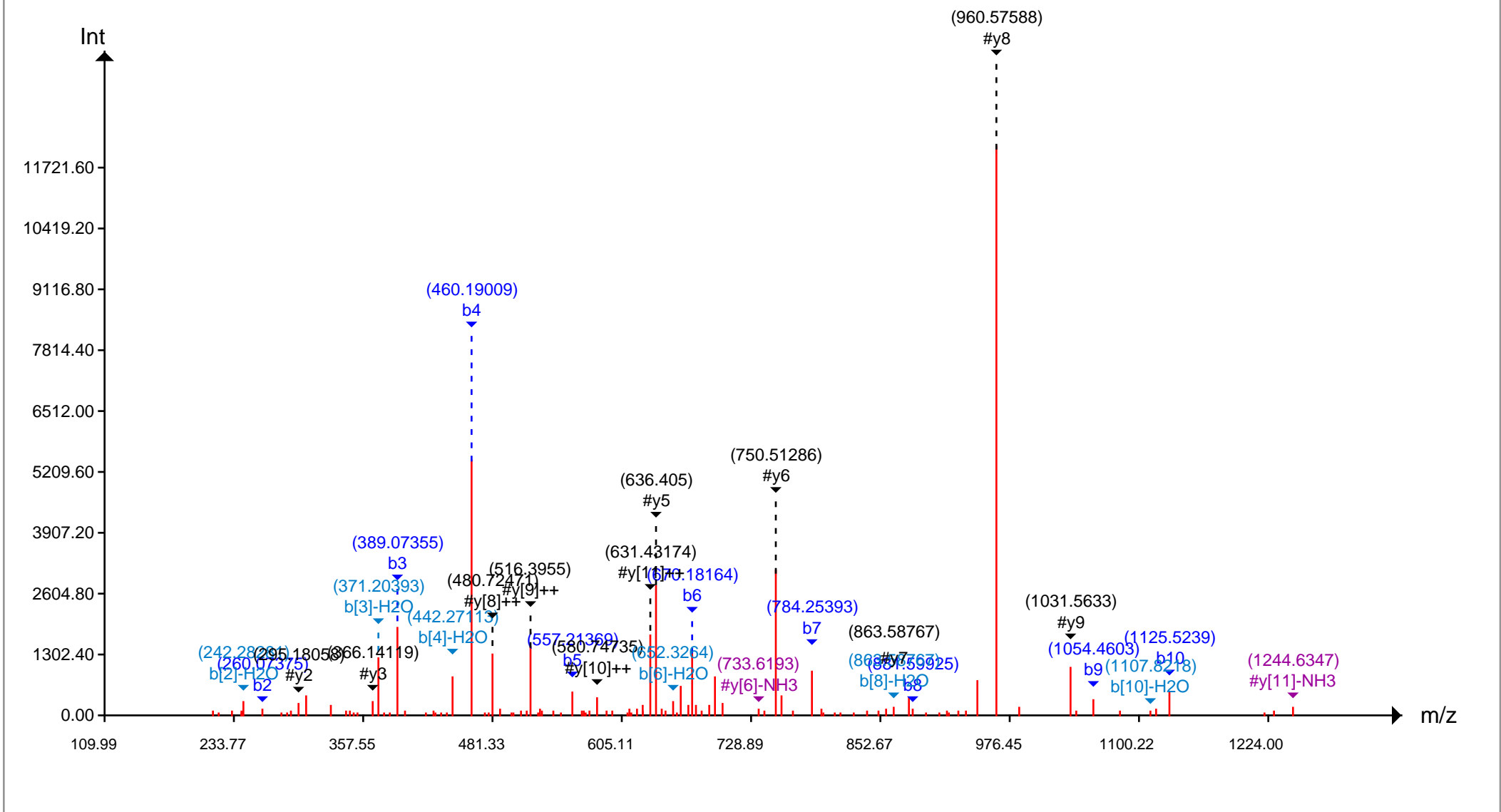
Info

Modified Sequence	AcD3-EFLTPVEEAPK<AcD3K*>GM<Mox*>LAR-COOH	Ion score	63.94
Protein	Q61599	Confidence level	99.00%
Precursor m/z	947.50038	Identity Threshold	48.03
Charge	2+	Mass error	0.003771



Info

Modified Sequence	AcD3-DALHGSLVTEGQYQAVR-COOH	Ion score	71.13
Protein	Q9EPB4	Confidence level	99.00%
Precursor m/z	944.98412	Identity Threshold	48.00
Charge	2+	Mass error	0.001875



Info

Modified Sequence AcD3-LTEAPLNPK<AcD3K*>ANR<C13>-COOH

Ion score 66.0

Protein P62737, P60710, P68033, P63260, P63268, P68134

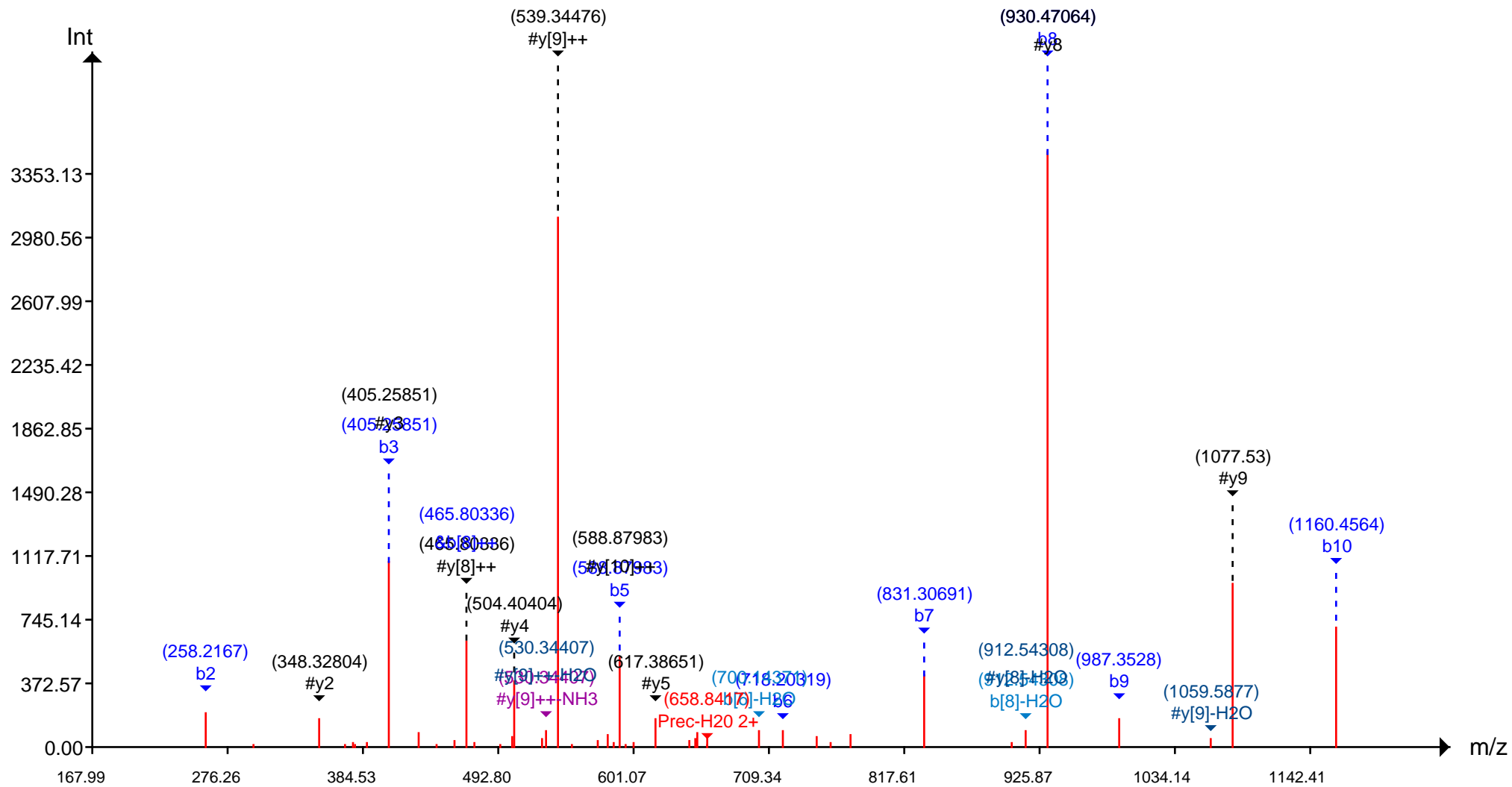
Confidence level 99.00%

Precursor m/z 710.41256

Identity Threshold 45.88

Charge 2+

Mass error 0.001028



Info

Modified Sequence AcD3-LVFPSEIVGK<AcD3K*>R-COOH

Ion score 78.05

Protein P62082

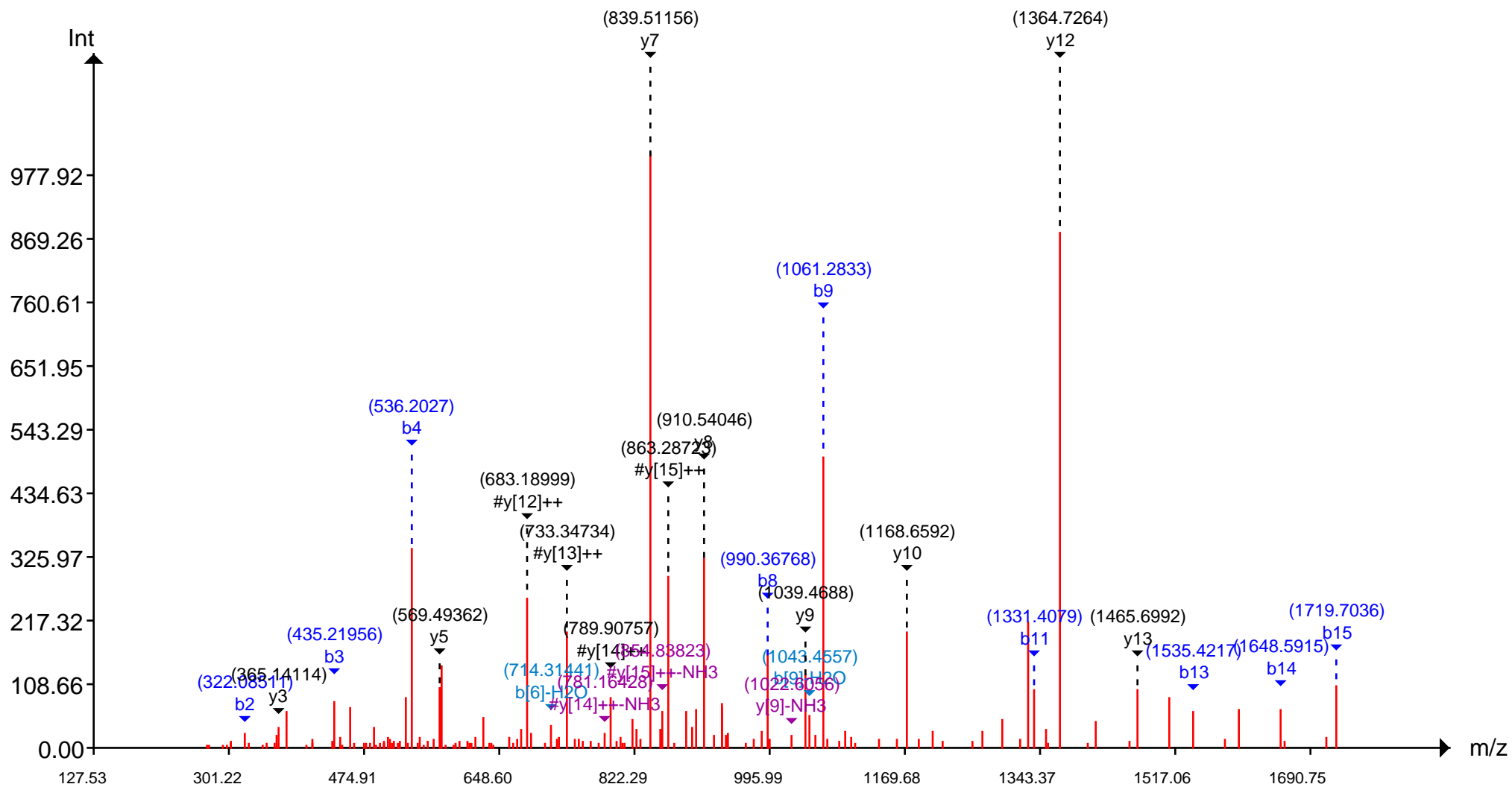
Confidence level 99.00%

Precursor m/z 667.90185

Identity Threshold 44.08

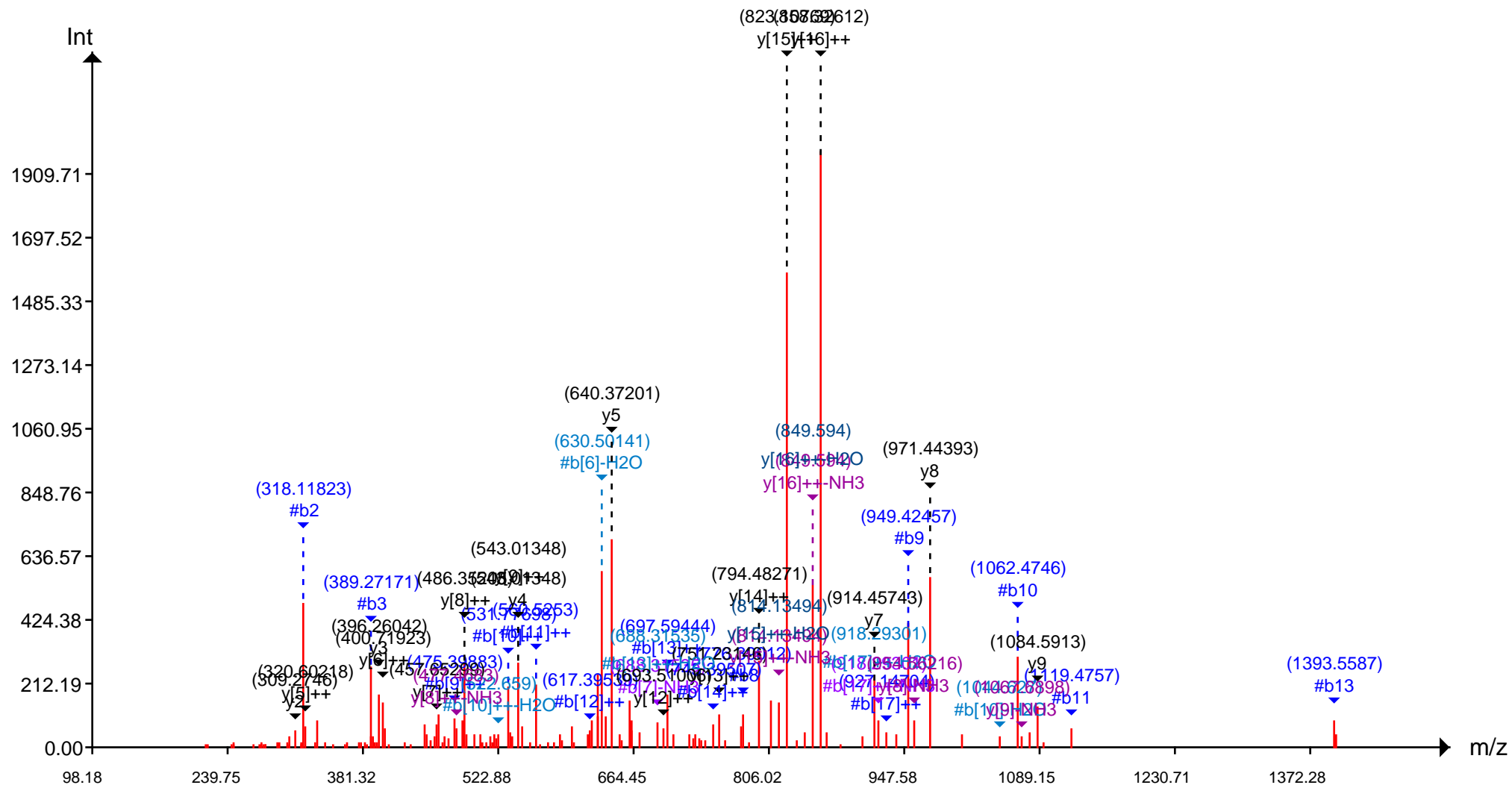
Charge 2+

Mass error 0.001505



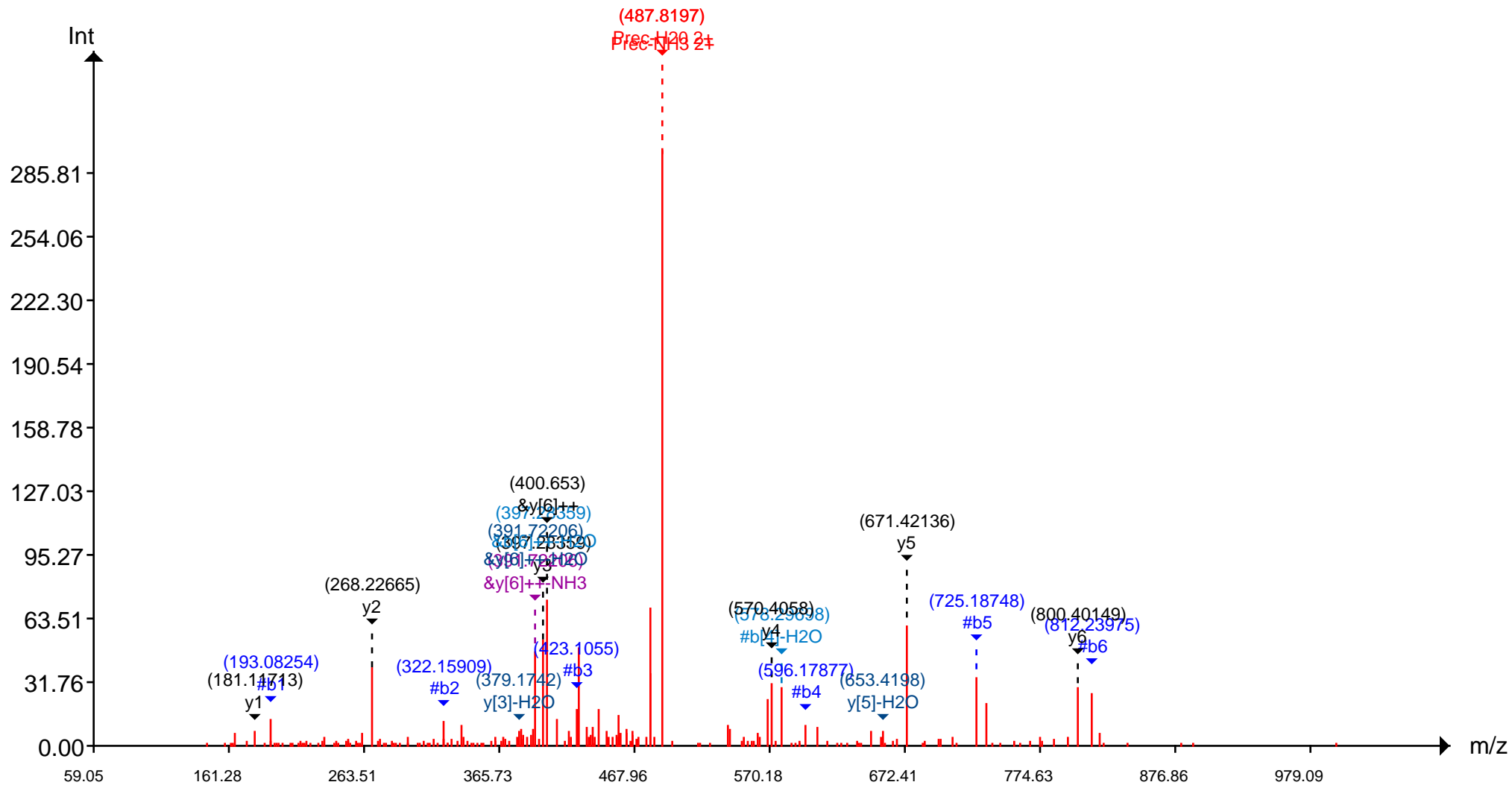
Info

Modified Sequence	AcD3-EFLTPVEEAPK<AcD3K*>GM<Mox*>LAR<C13>-COOH	Ion score	55.22
Protein	Q61599	Confidence level	99.00%
Precursor m/z	950.50927	Identity Threshold	47.86
Charge	2+	Mass error	0.001409



Info

Modified Sequence	AcD3-VK<AcD3K*>AGSDGAK<AcD3K*>IGNC<Cmm*>PFSQR<C13>-COOH	Ion score	55.66
Protein	Q9Z1Q5	Confidence level	99.00%
Precursor m/z	678.35628	Identity Threshold	48.12
Charge	3+	Mass error	0.001648



Info

Modified Sequence AcD3-FETK<AcD3K*>ESR<C13>-COOH

Ion score 51.52

Protein Q9R0P5

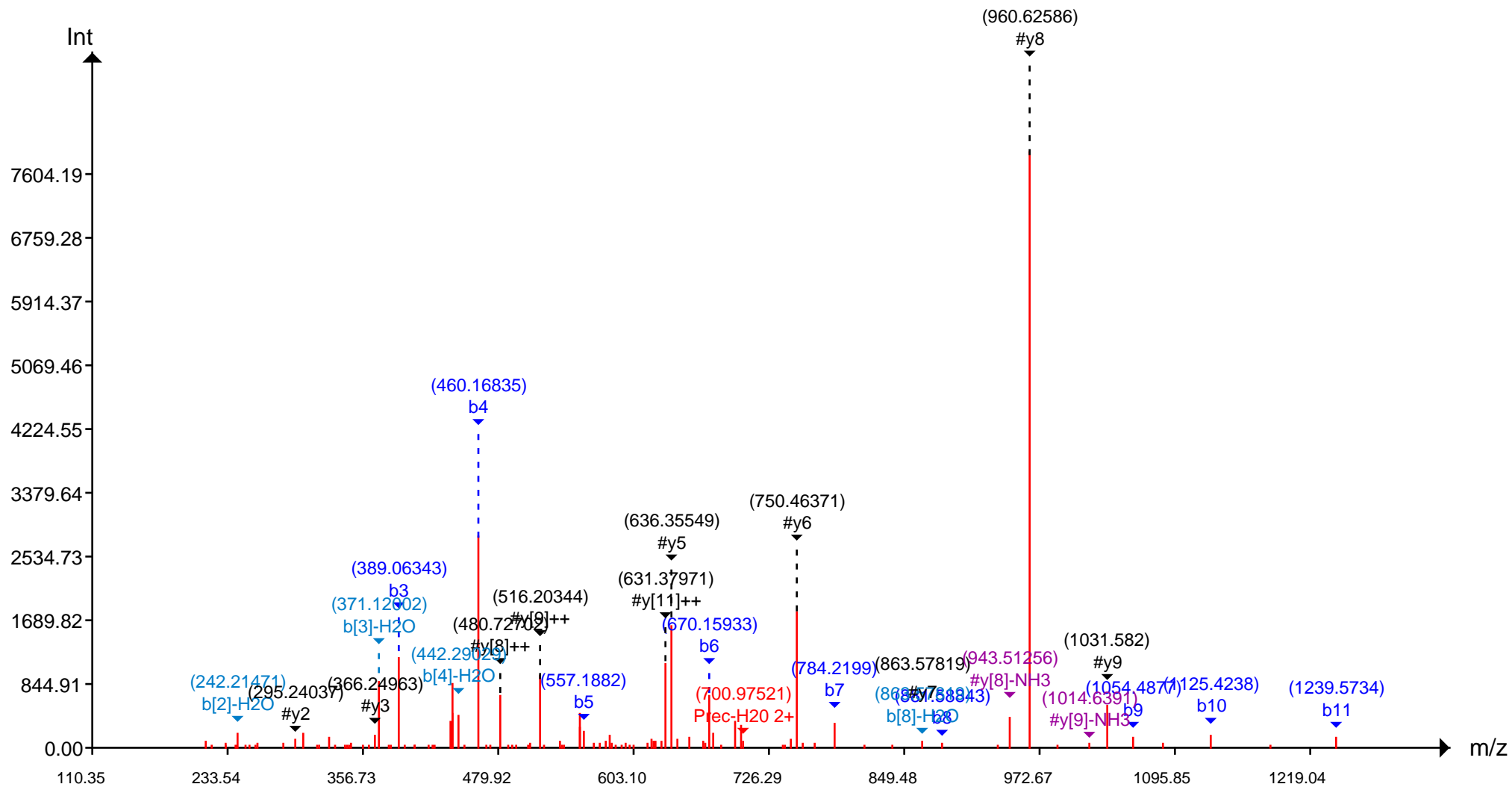
Confidence level 99.00%

Precursor m/z 496.76672

Identity Threshold 48.05

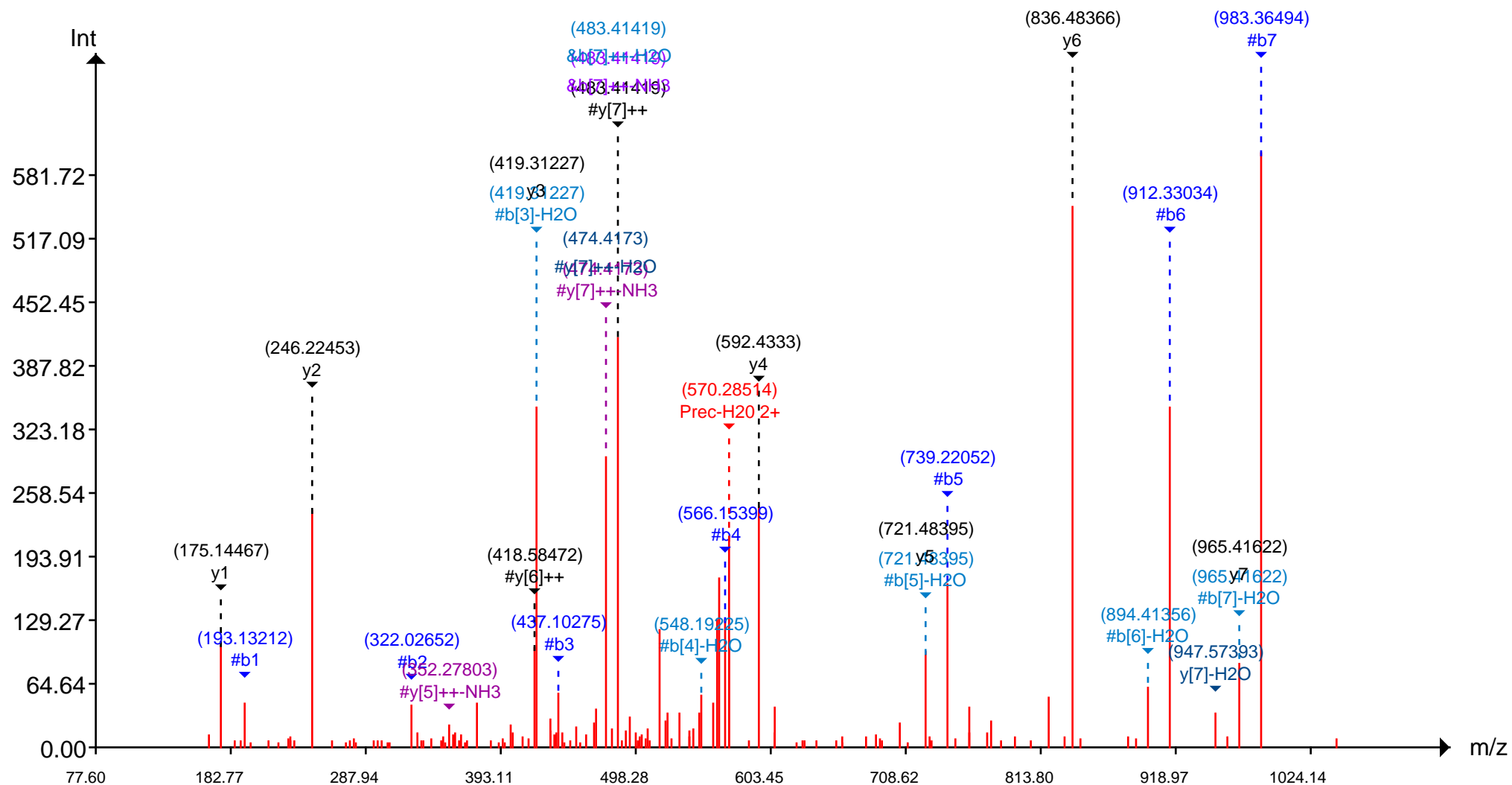
Charge 2+

Mass error 2.8E-5



Info

Modified Sequence	AcD3-LTEAPLNPK<AcD3K*>ANR<C13>-COOH	Ion score	65.01
Protein	P62737, P60710, P68033, P63260, P63268, P68134	Confidence level	99.00%
Precursor m/z	710.41166	Identity Threshold	46.01
Charge	2+	Mass error	-7.72E-4



Info

Modified Sequence AcD3-FEDEK<AcD3K*>K<AcD3K*>AR-COOH

Ion score 71.71

Protein P50580

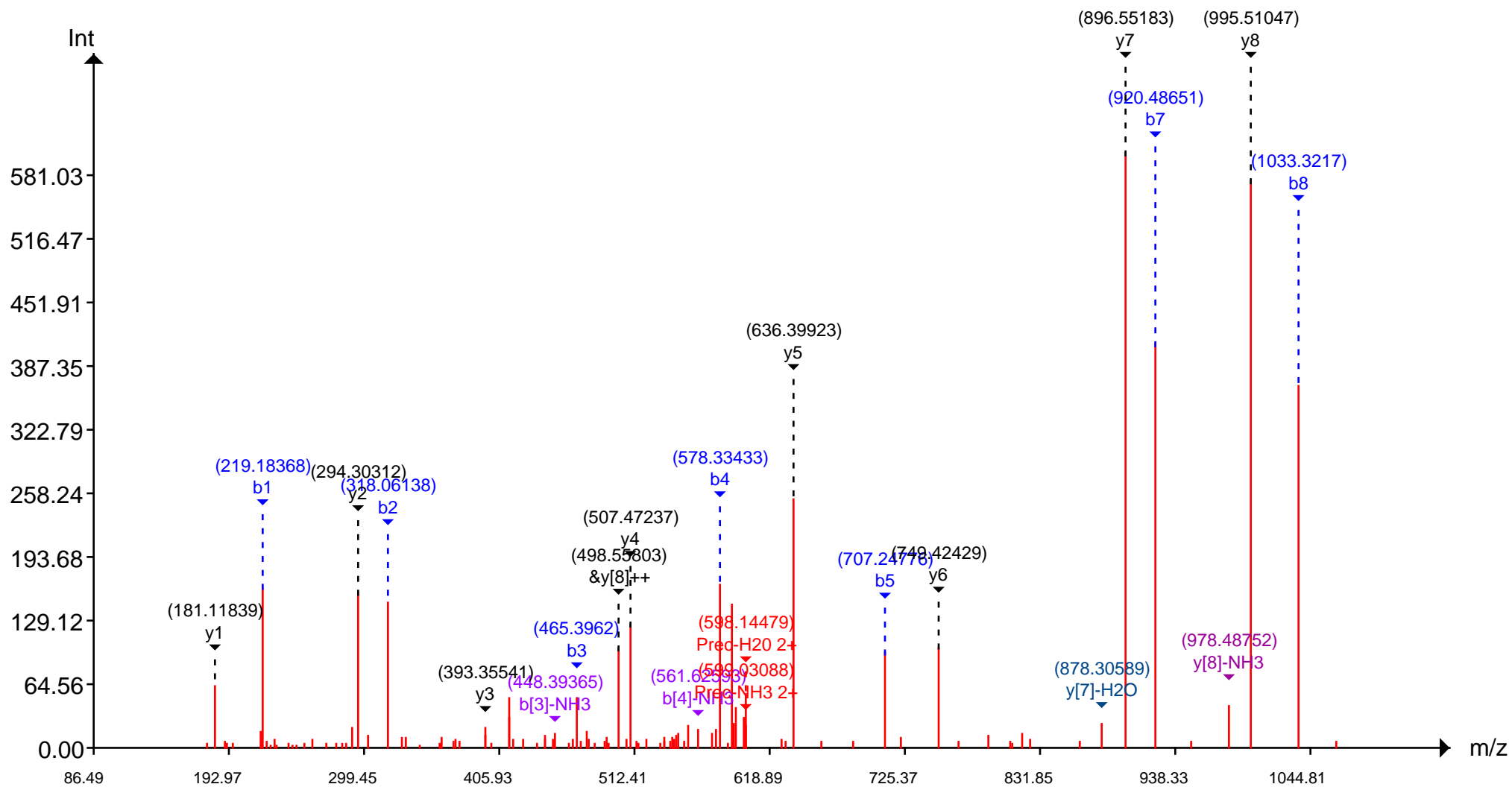
Confidence level 99.00%

Precursor m/z 579.3113

Identity Threshold 47.84

Charge 2+

Mass error 6.26E-4



Info

Modified Sequence AcD3-K<AcD3K*>VFLENVIR<C13>-COOH

Ion score 62.81

Protein P62806

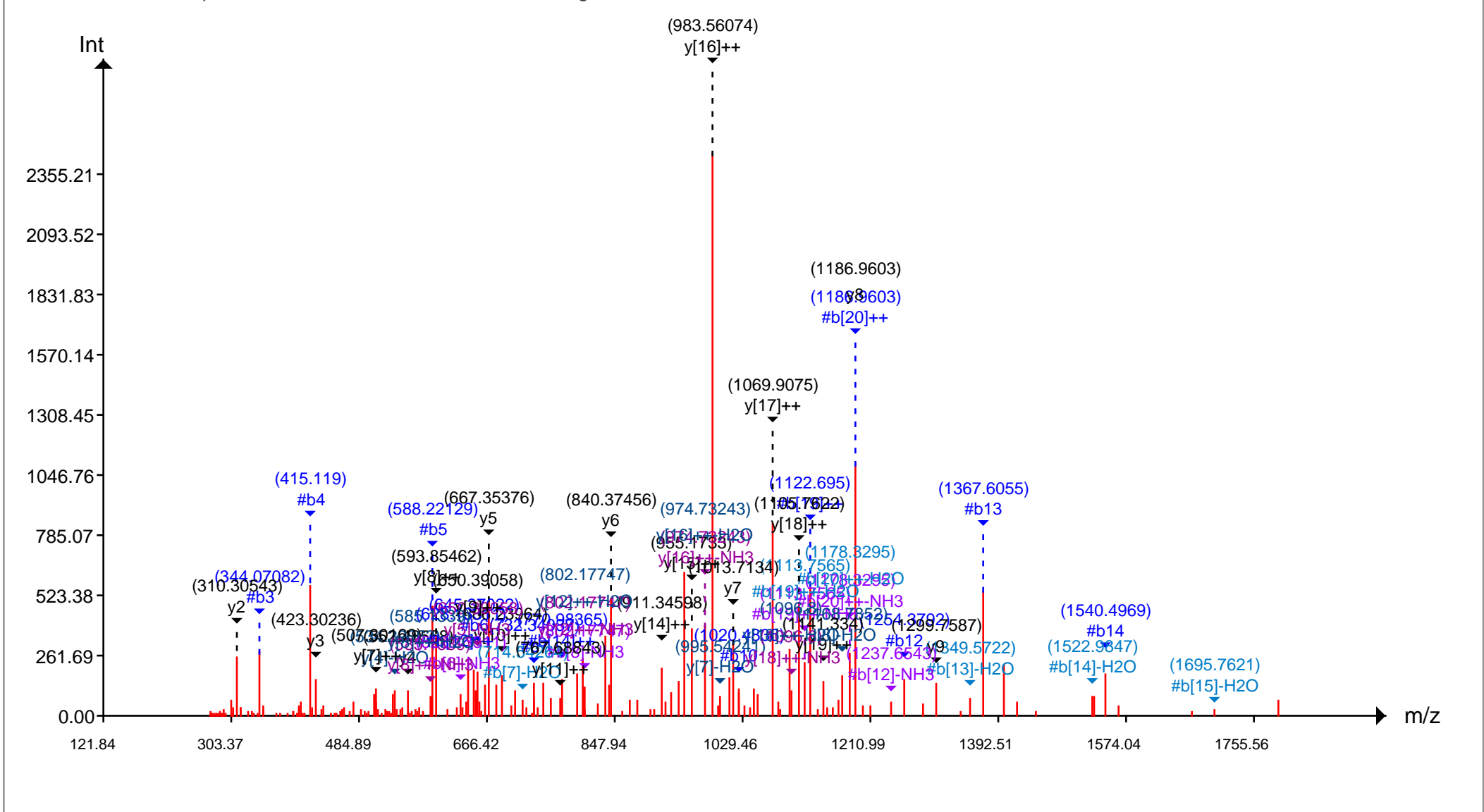
Confidence level 99.00%

Precursor m/z 607.37973

Identity Threshold 42.00

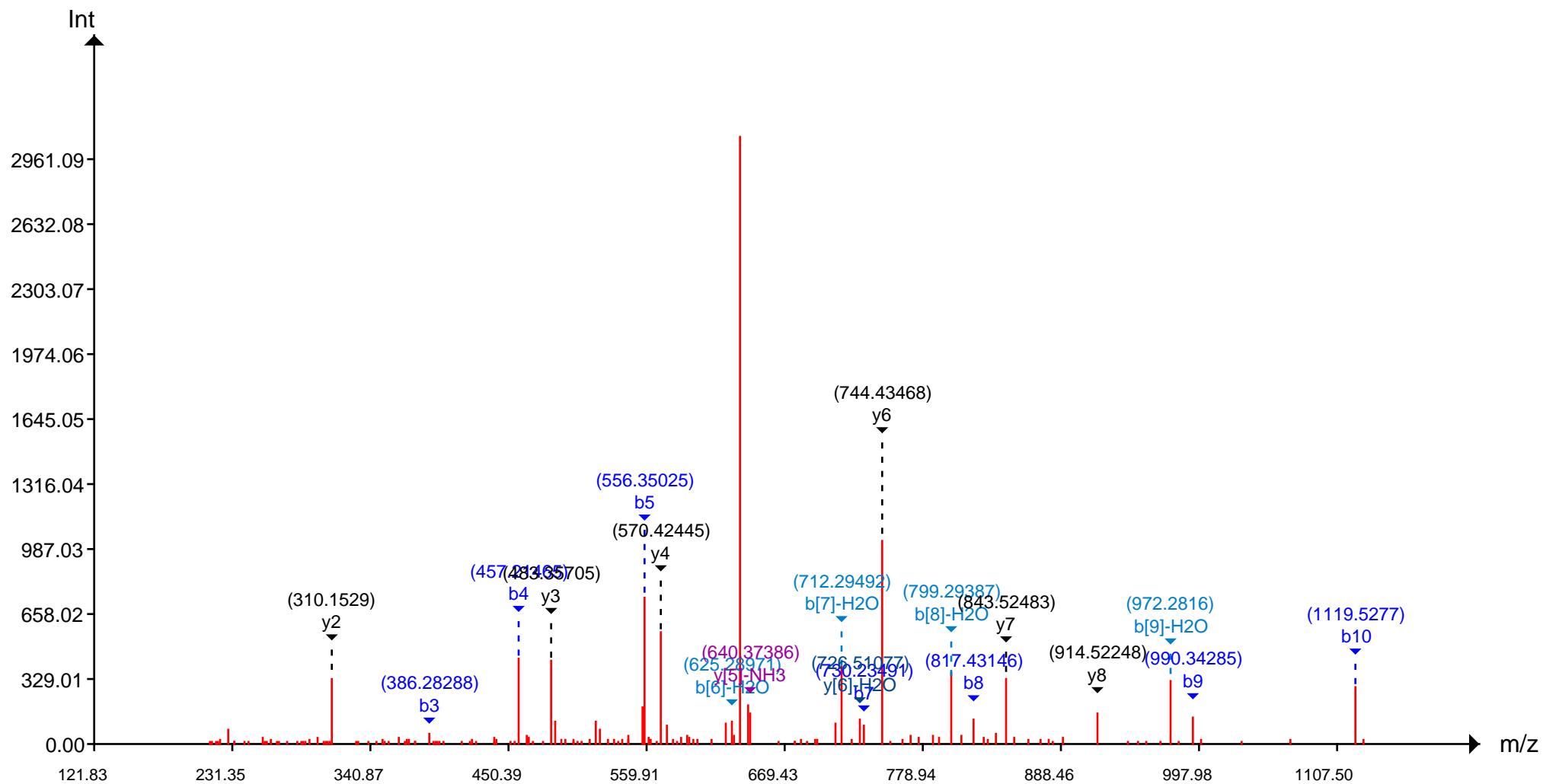
Charge 2+

Mass error 4.62E-4



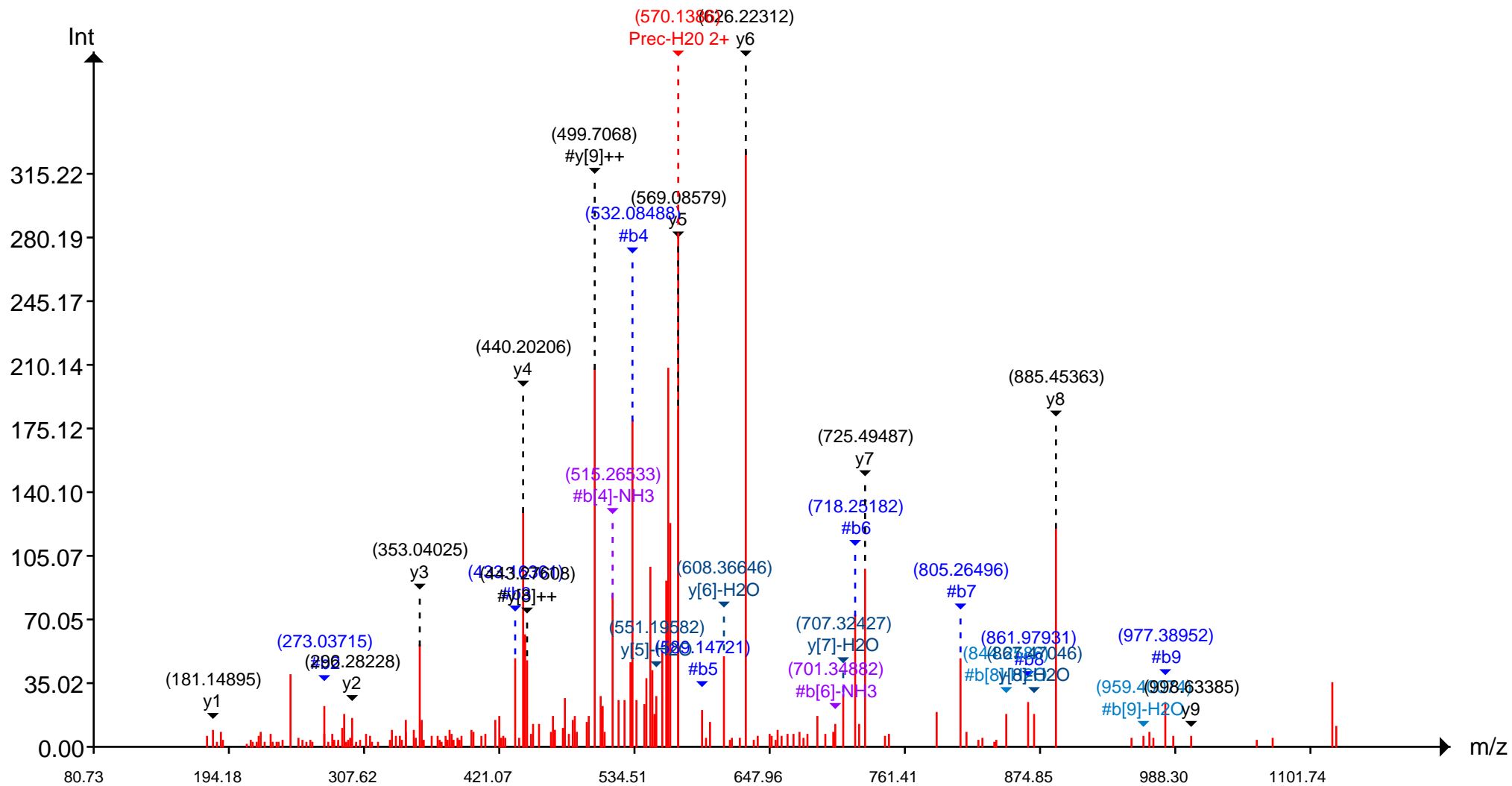
Info

Modified Sequen...	AcD3-INAAK<AcD3K*>GSSNSYAIK<AcD3K*>K<AcD3K*>K<AcD3K*>DELER<C13>-COOH	Ion score	64.63
Prote...	P97461	Confidence level	99.00%
Precursor m...	851.80833	Identity Threshold	45.80
Char...	3+	Mass error	0.002176



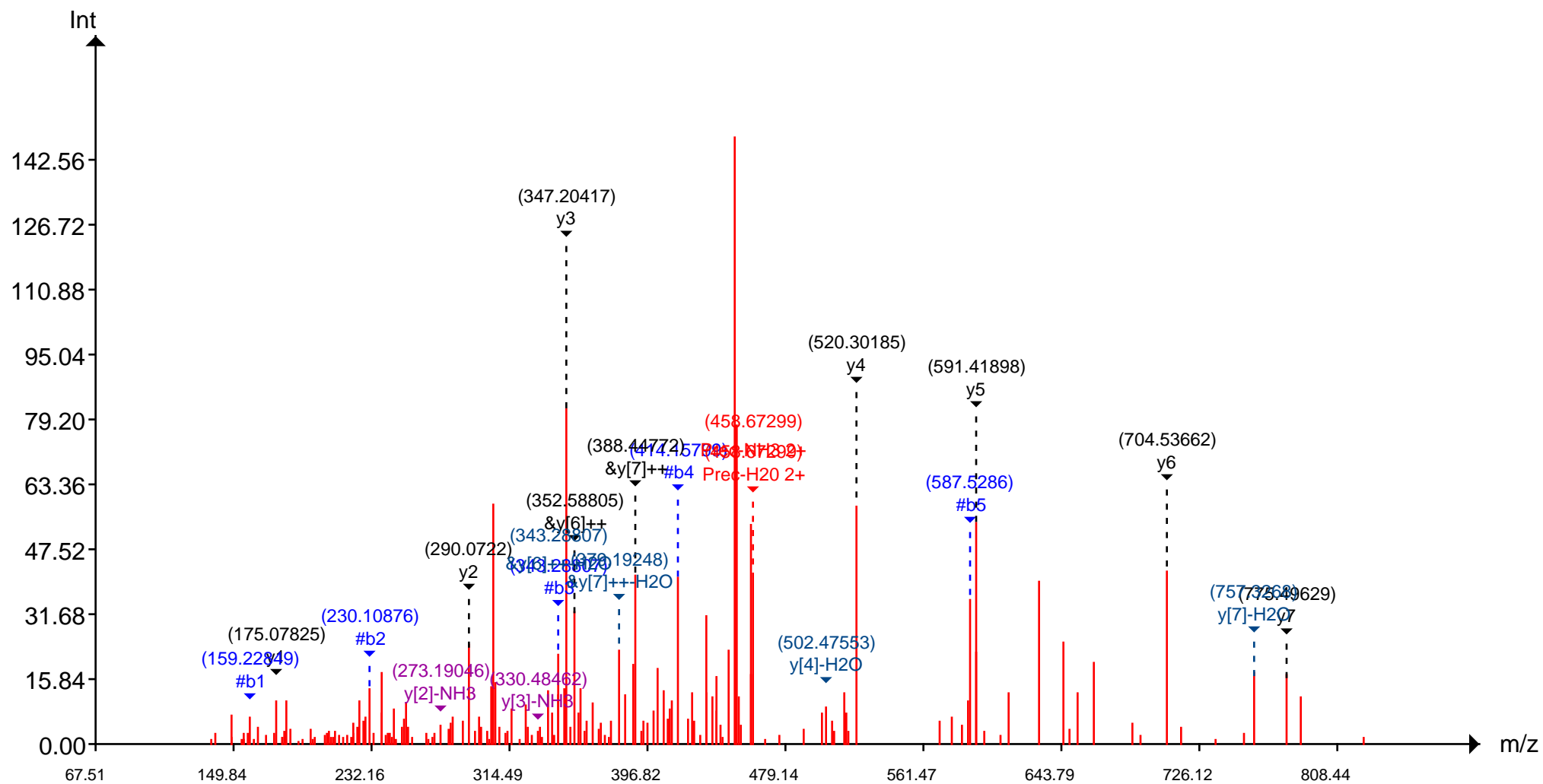
Info

Modified Sequence	AcD3-IVQAVSSSK<AcD3K*>ER<C13>-COOH	Ion score	48.16
Protein	P43275	Confidence level	99.00%
Precursor m/z	650.37829	Identity Threshold	46.23
Charge	2+	Mass error	0.001214



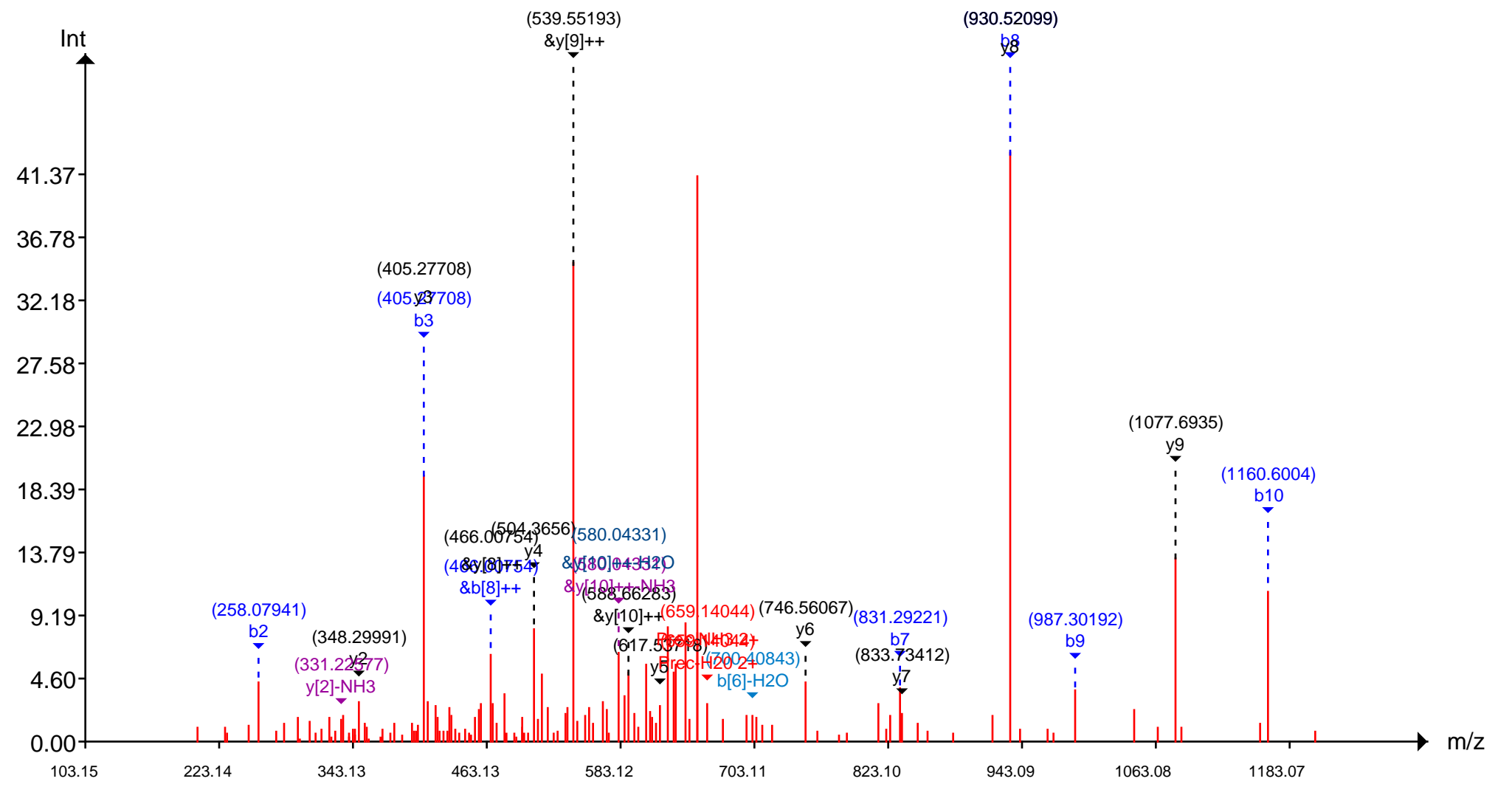
Info

Modified Sequence	AcD3-NIC<Cmm*>VGESGDR<C13>-COOH	Ion score	67.43
Protein	Q9CXW4	Confidence level	99.00%
Precursor m/z	579.27322	Identity Threshold	45.46
Charge	2+	Mass error	1.34E-4

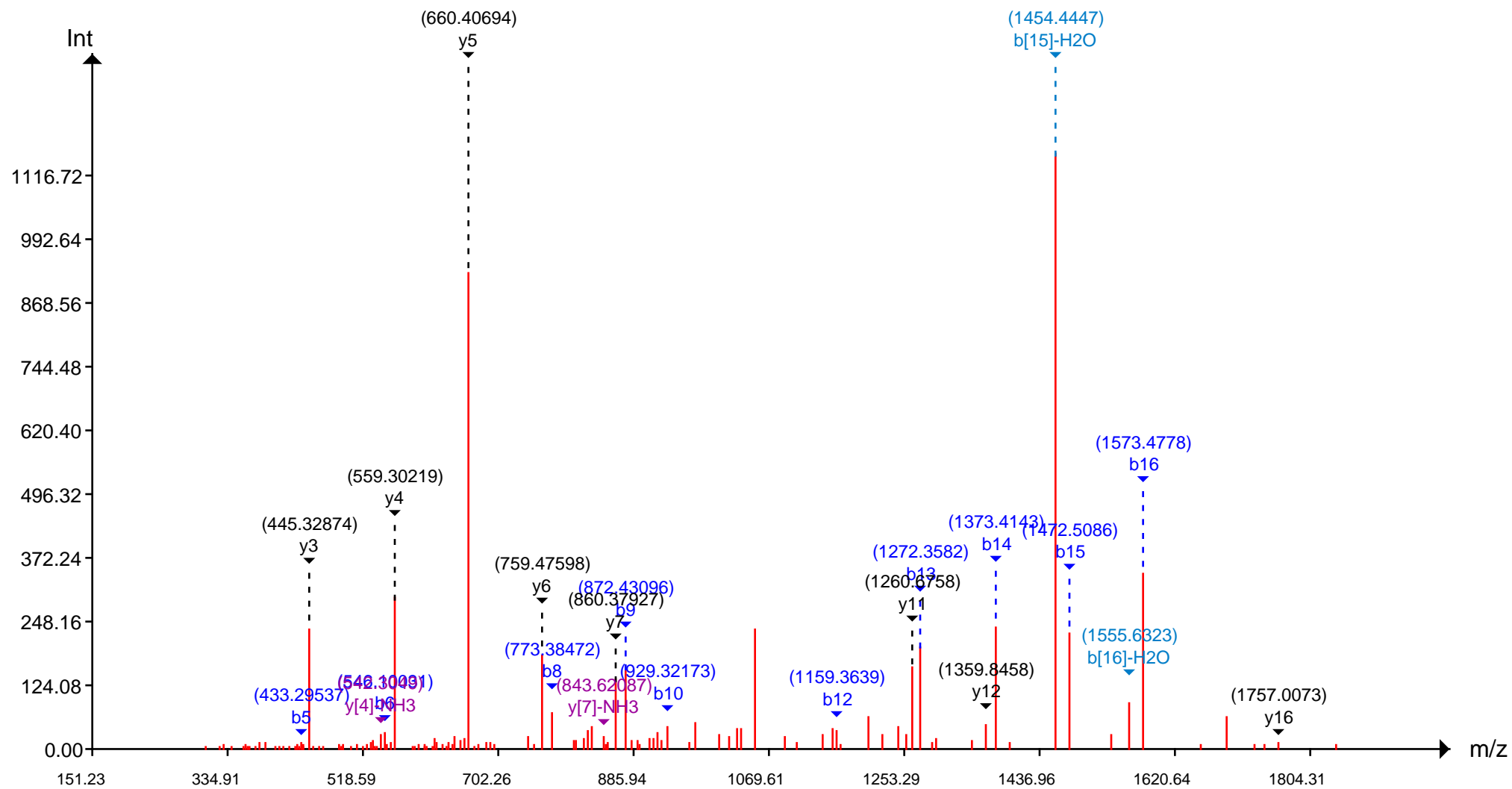


Info

Modified Sequence	AcD3-LALAK<AcD3K*>GDR-COOH	Ion score	51.1
Protein	P10107	Confidence level	99.00%
Precursor m/z	467.28533	Identity Threshold	44.87
Charge	2+	Mass error	-6.0E-5

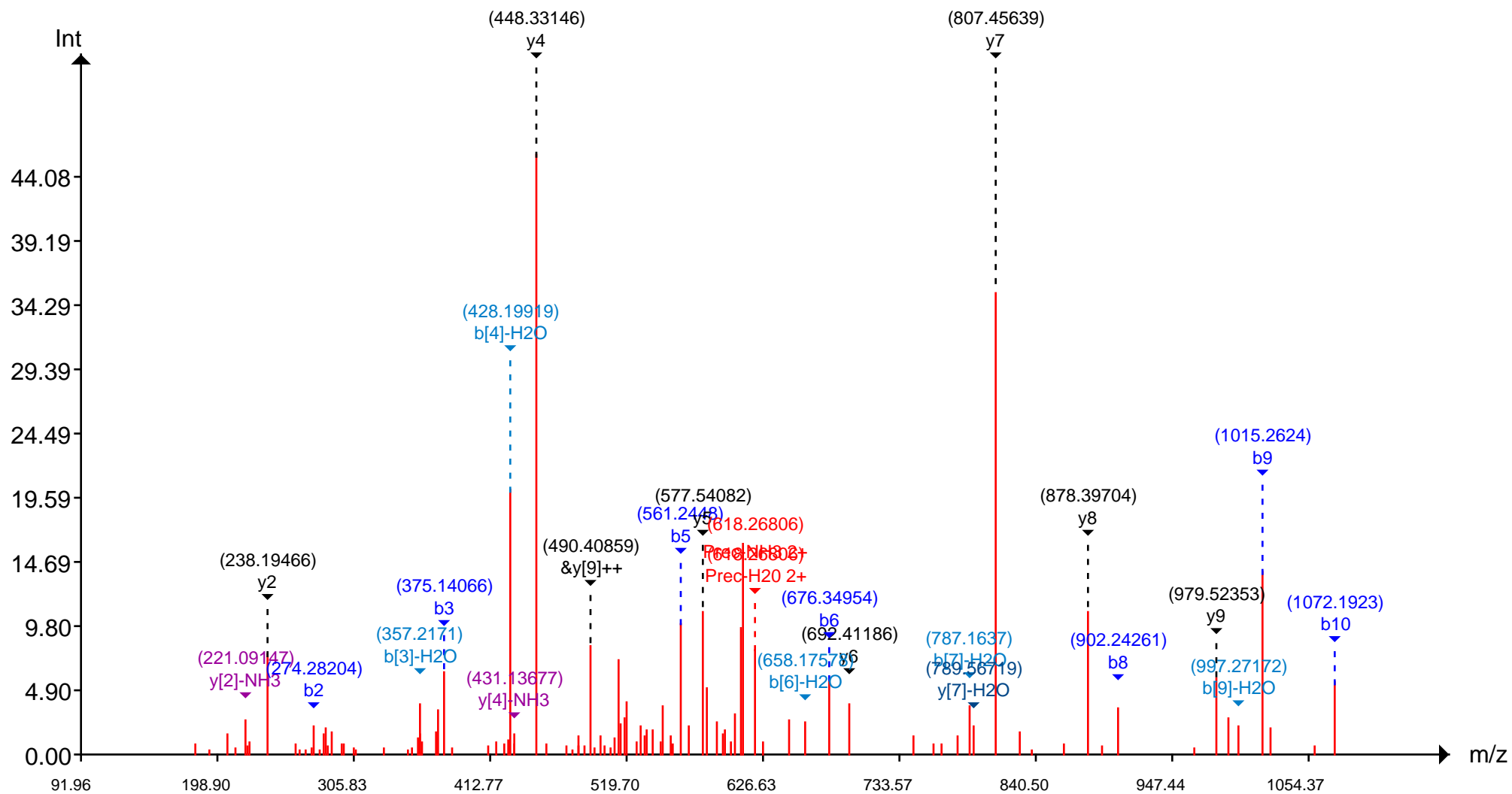


Info			
Modified Sequence	AcD3-LVFPSEIVGK<AcD3K*>R-COOH	Ion score	54.11
Protein	P62082	Confidence level	99.00%
Precursor m/z	667.9016	Identity Threshold	44.10
Charge	2+	Mass error	0.001005



Info

Modified Sequence	AcD3-TASAGIQVVGDDTLVTNPK<AcD3K*>R-COOH	Ion score	66.63
Protein	P17182	Confidence level	99.00%
Precursor m/z	1066.578	Identity Threshold	47.45
Charge	2+	Mass error	0.002257



Info

Modified Sequence AcD3-DITADDEPLGR<C13>-COOH

Protein P17742

Precursor m/z 626.8127

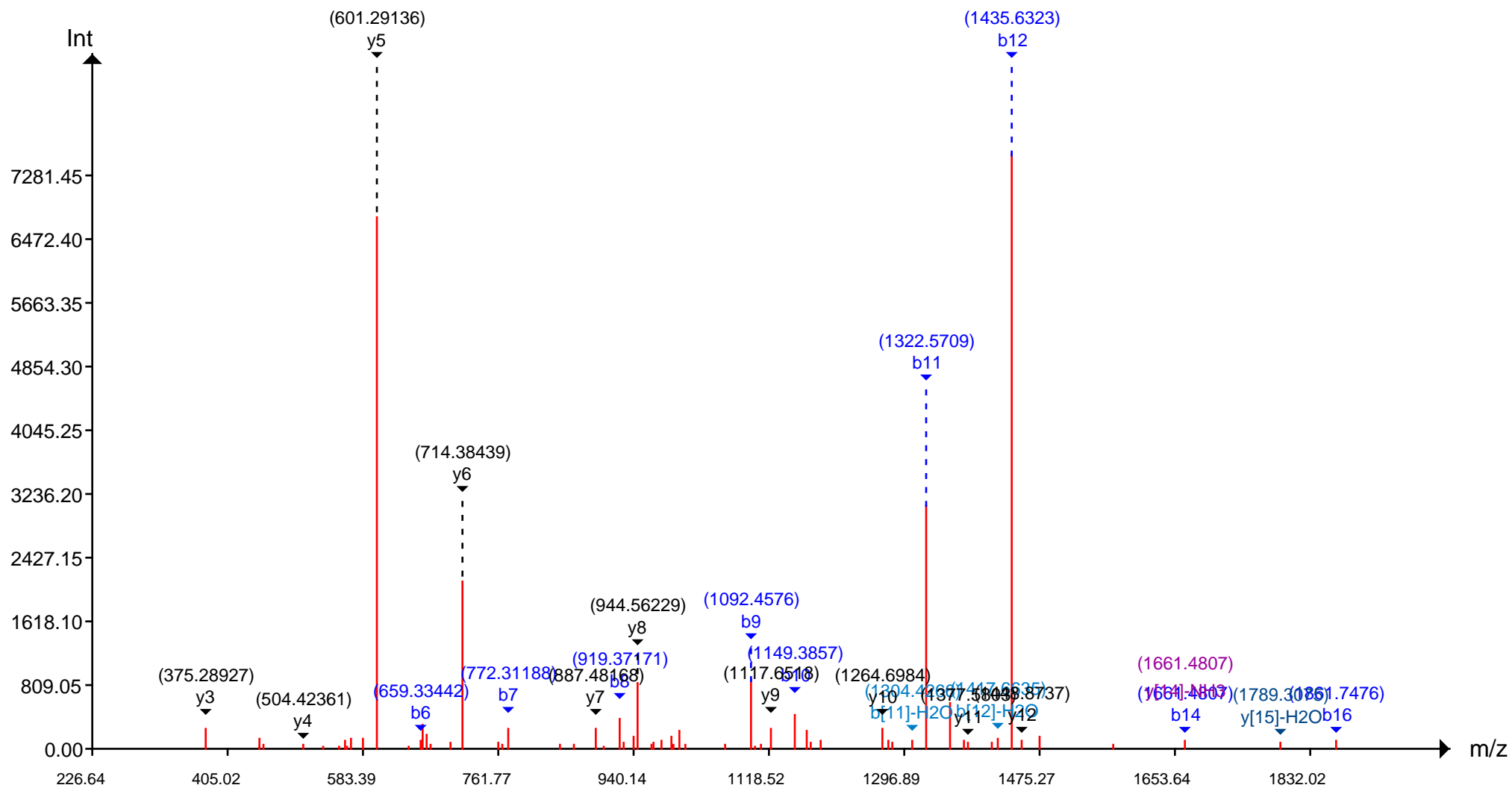
Charge 2+

Ion score 70.12

Confidence level 99.00%

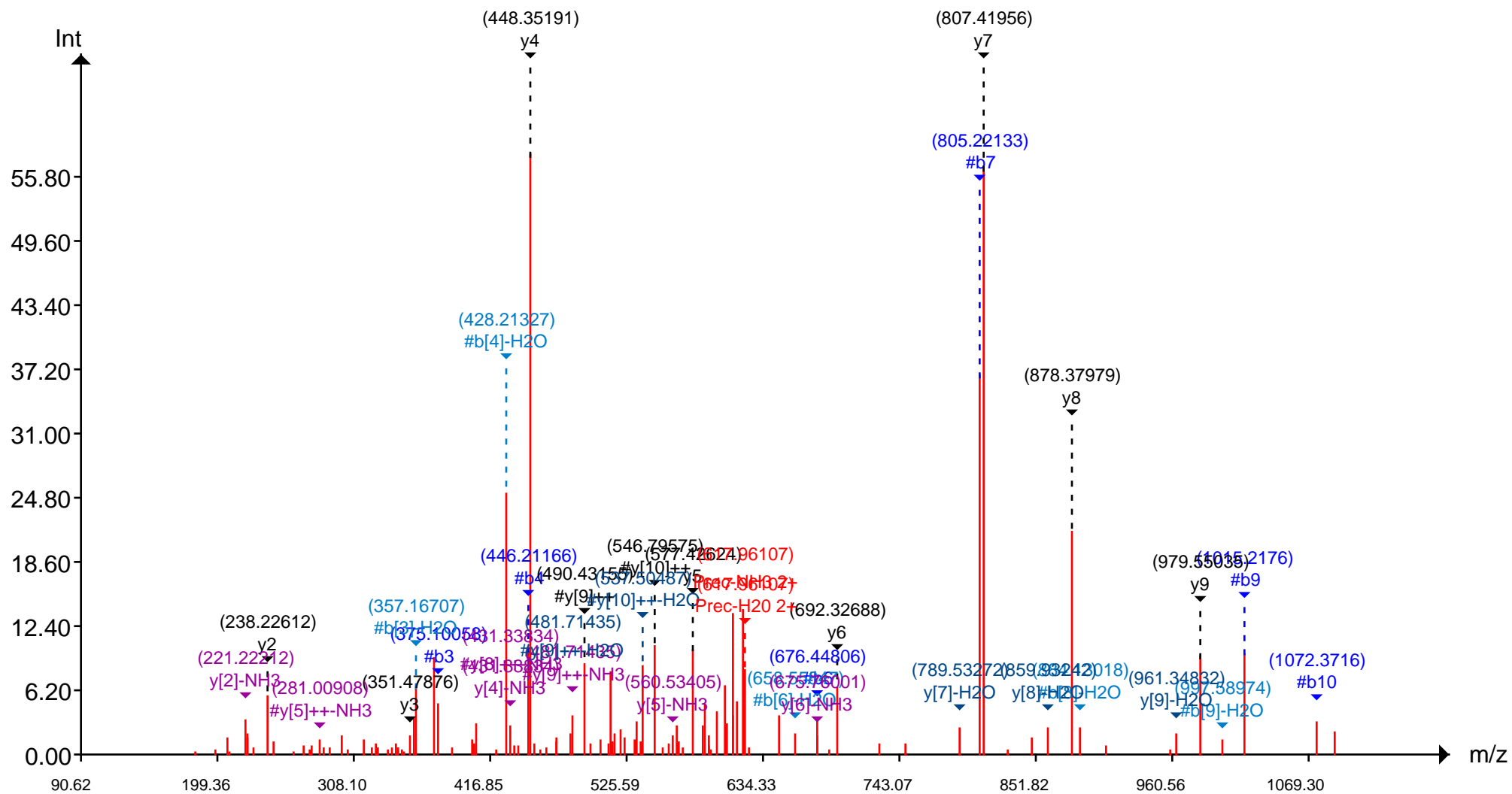
Identity Threshold 47.47

Mass error -9.23E-4



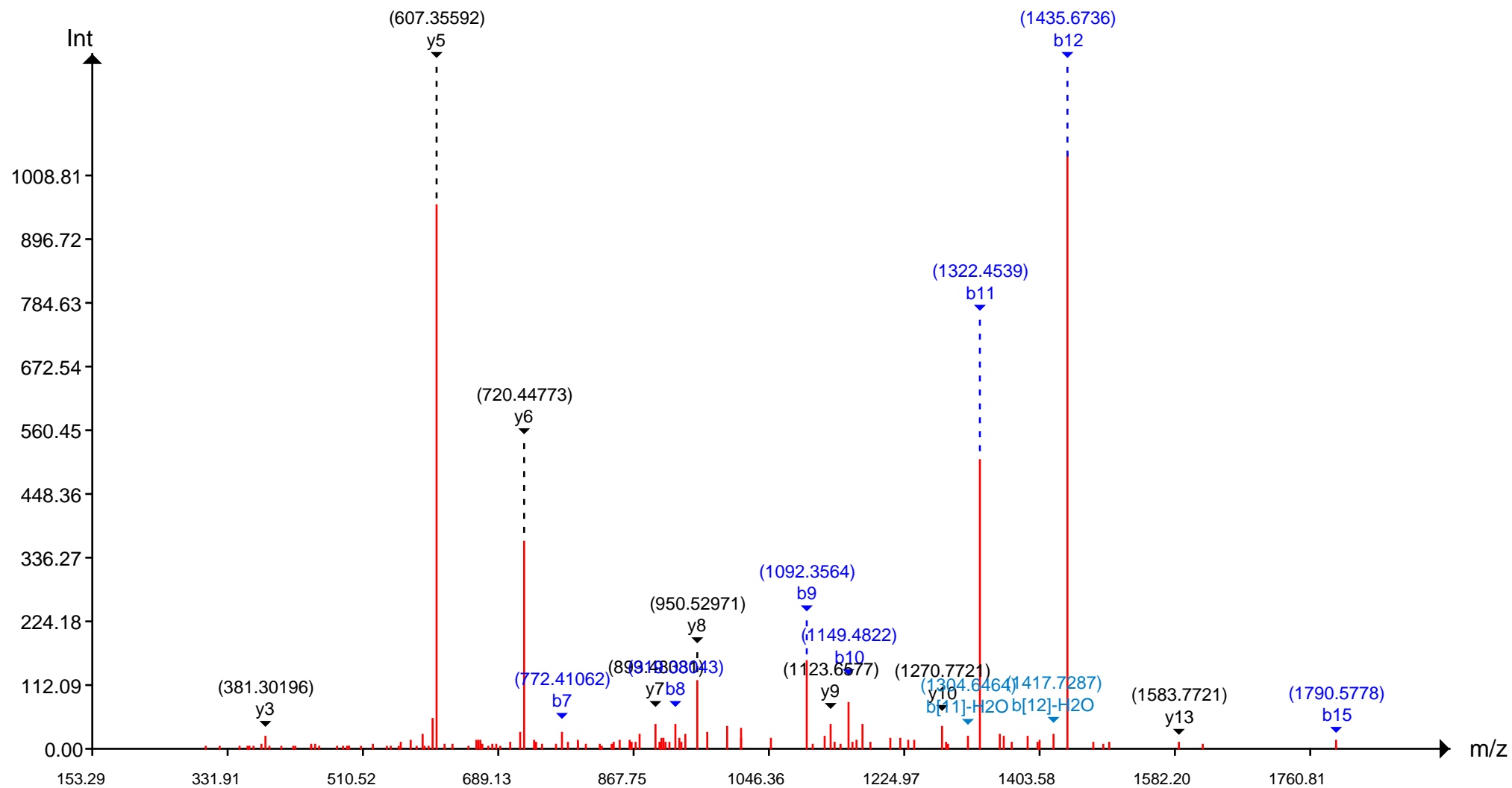
Info

Modified Sequence	AcD3-LAQTEALM<Mox*>K<AcD3K*>GK<AcD3K*>LPPEAR-COOH	Ion score	50.22
Protein	P06745	Confidence level	99.00%
Precursor m/z	1018.557	Identity Threshold	47.30
Charge	2+	Mass error	0.002509



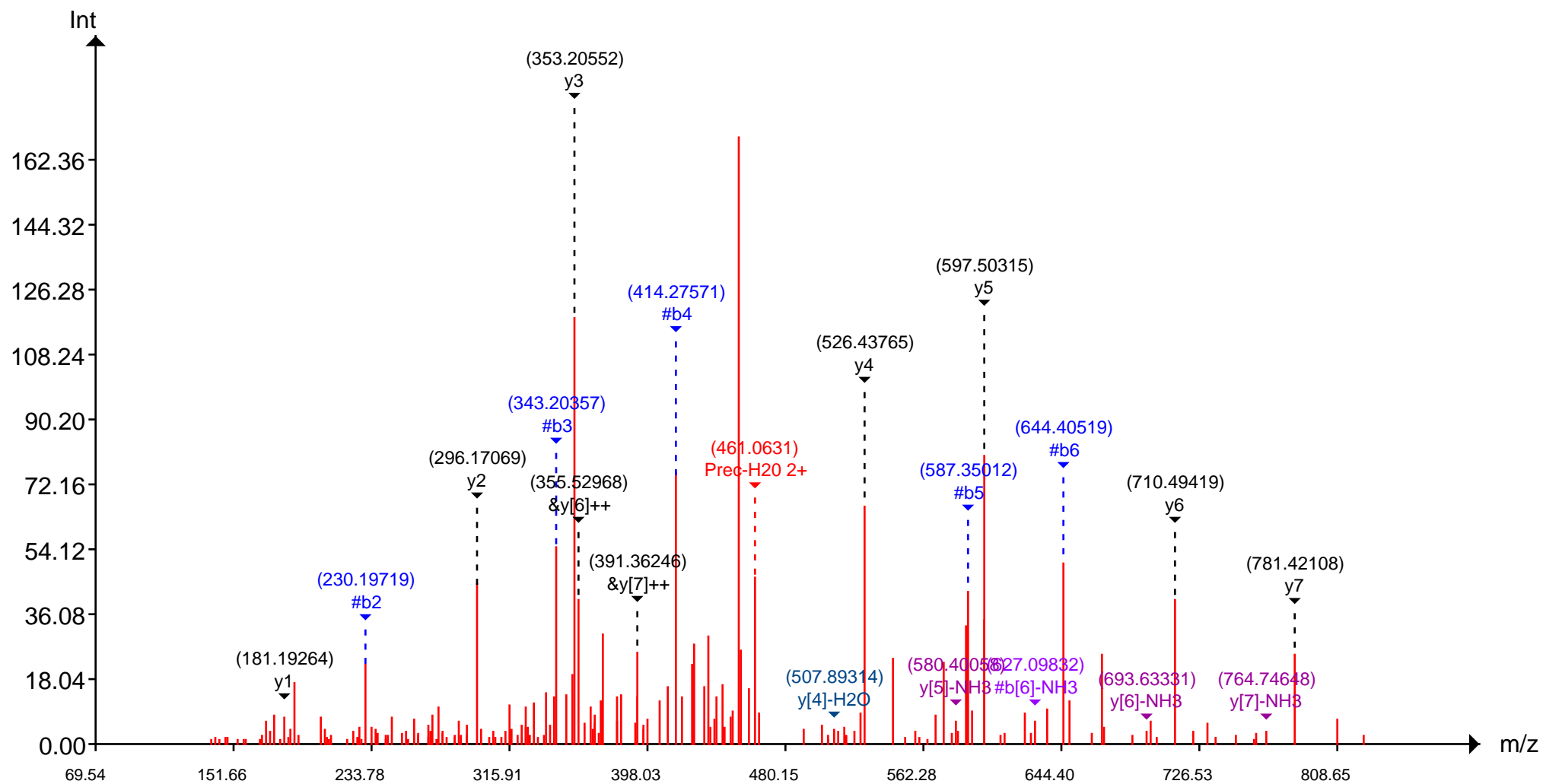
Info

Modified Sequence	AcD3-DITADDEPLGR<C13>-COOH	Ion score	51.94
Protein	P17742	Confidence level	99.00%
Precursor m/z	626.81366	Identity Threshold	47.50
Charge	2+	Mass error	9.97E-4



Info

Modified Sequence	AcD3-LAQTEALM<Mox*>K<AcD3K*>GK<AcD3K*>LPPEAR<C13>-COOH	Ion score	54.8
Protein	P06745	Confidence level	99.00%
Precursor m/z	1021.5673	Identity Threshold	46.73
Charge	2+	Mass error	0.002967



Info

Modified Sequence AcD3-LALAK<AcD3K*>GDR<C13>-COOH

Ion score 48.71

Protein P10107

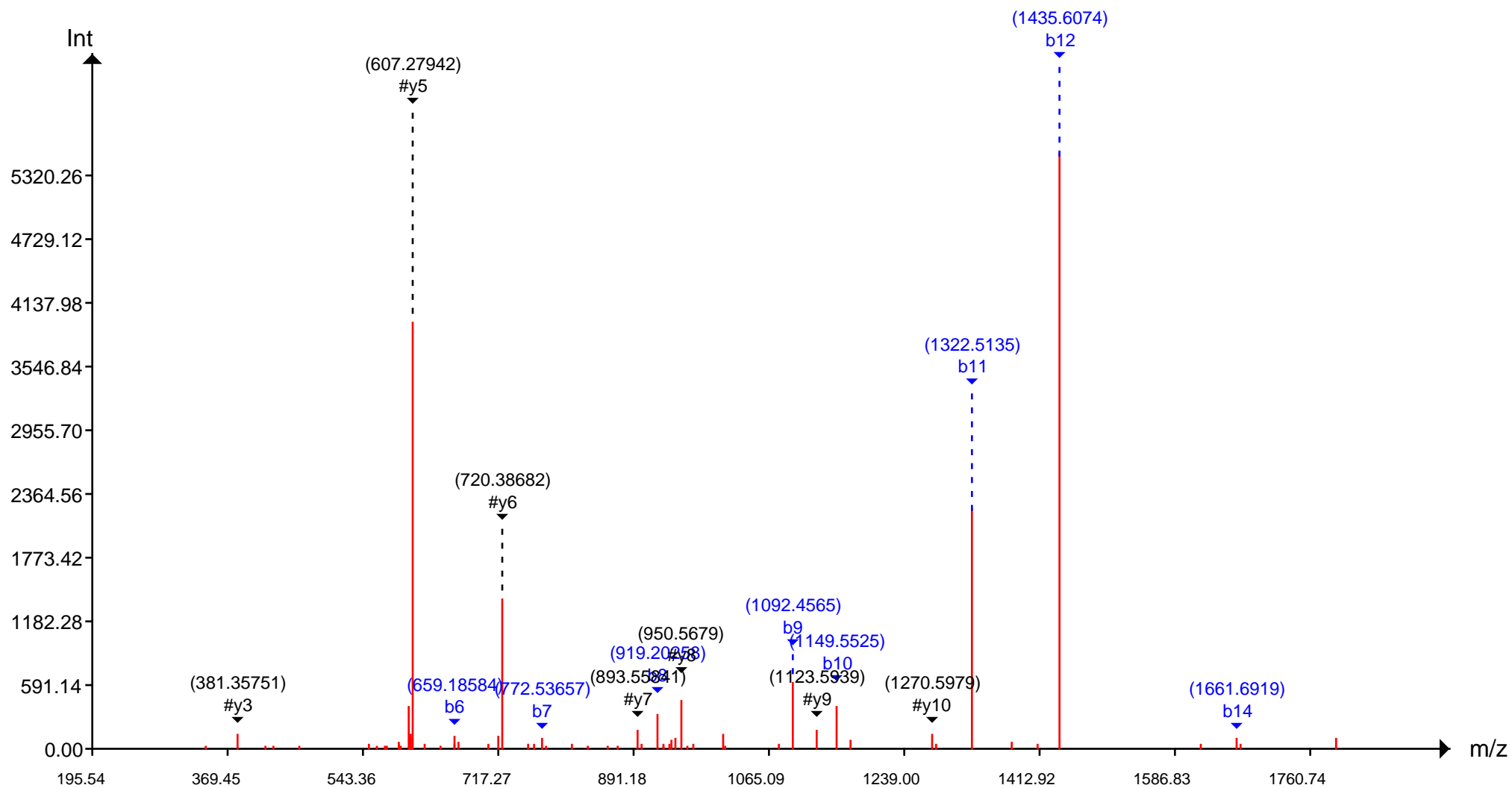
Confidence level 99.00%

Precursor m/z 470.2955

Identity Threshold 44.11

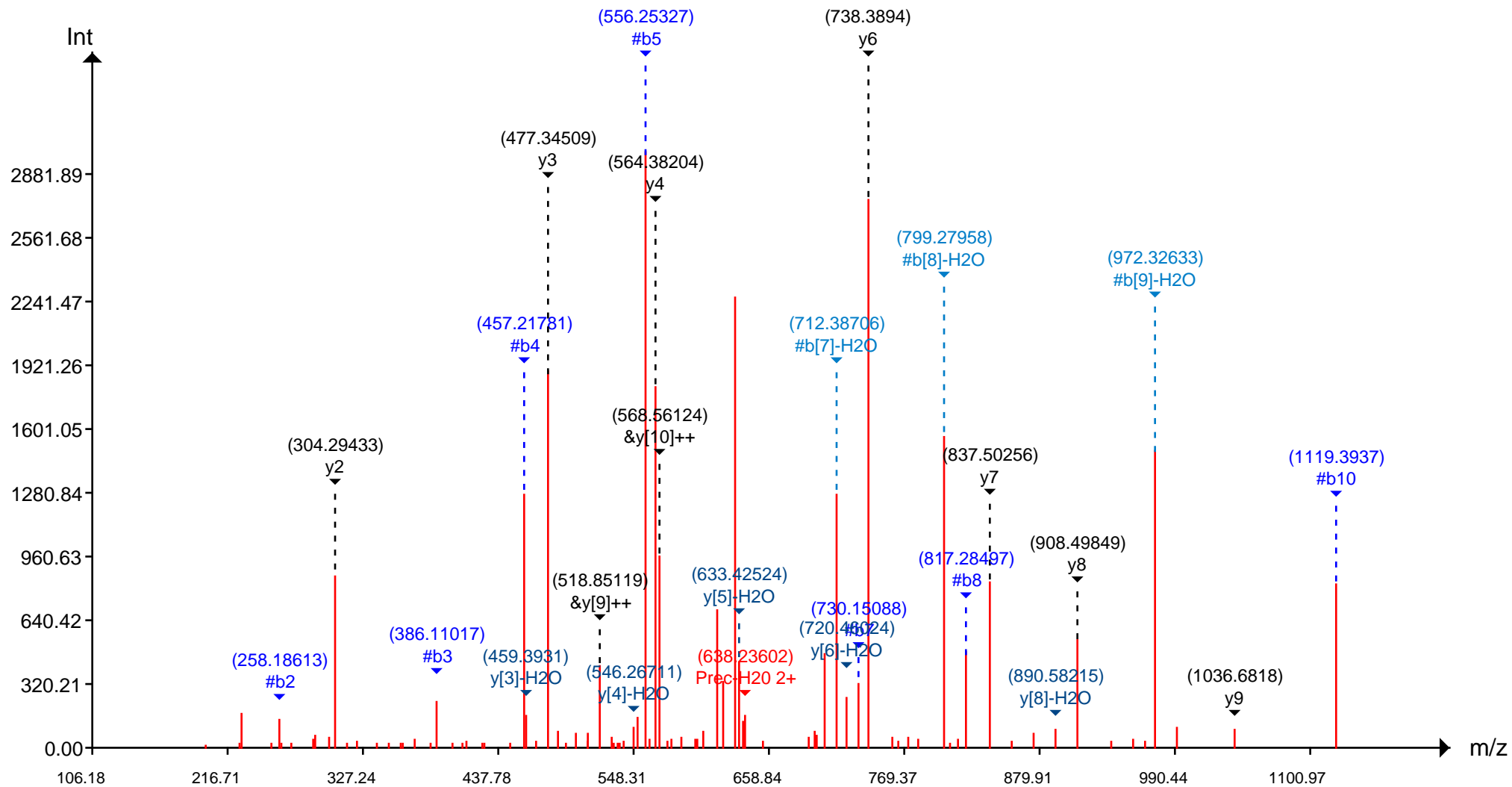
Charge 2+

Mass error 1.39E-4



Info

Modified Sequence	AcD3-LAQTEALM<Mox*>K<AcD3K*>GK<AcD3K*>LPPEAR<C13>-COOH	Ion score	54.35
Protein	P06745	Confidence level	99.00%
Precursor m/z	1021.5666	Identity Threshold	46.74
Charge	2+	Mass error	0.001567



Info

Modified Sequence AcD3-IVQAVSSSK<AcD3K*>ER-COOH

Ion score 66.13

Protein P43275

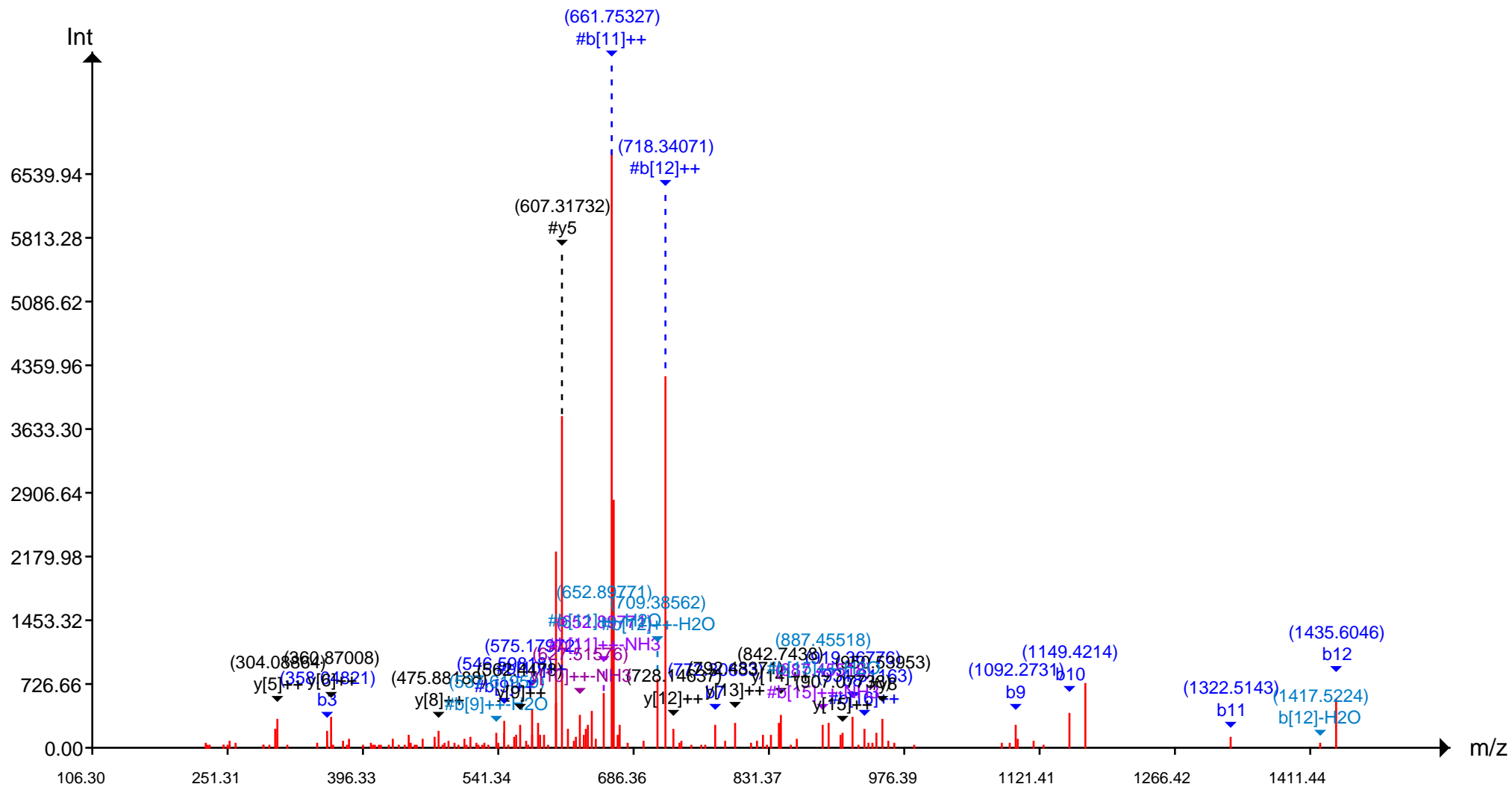
Confidence level 99.00%

Precursor m/z 647.36898

Identity Threshold 47.02

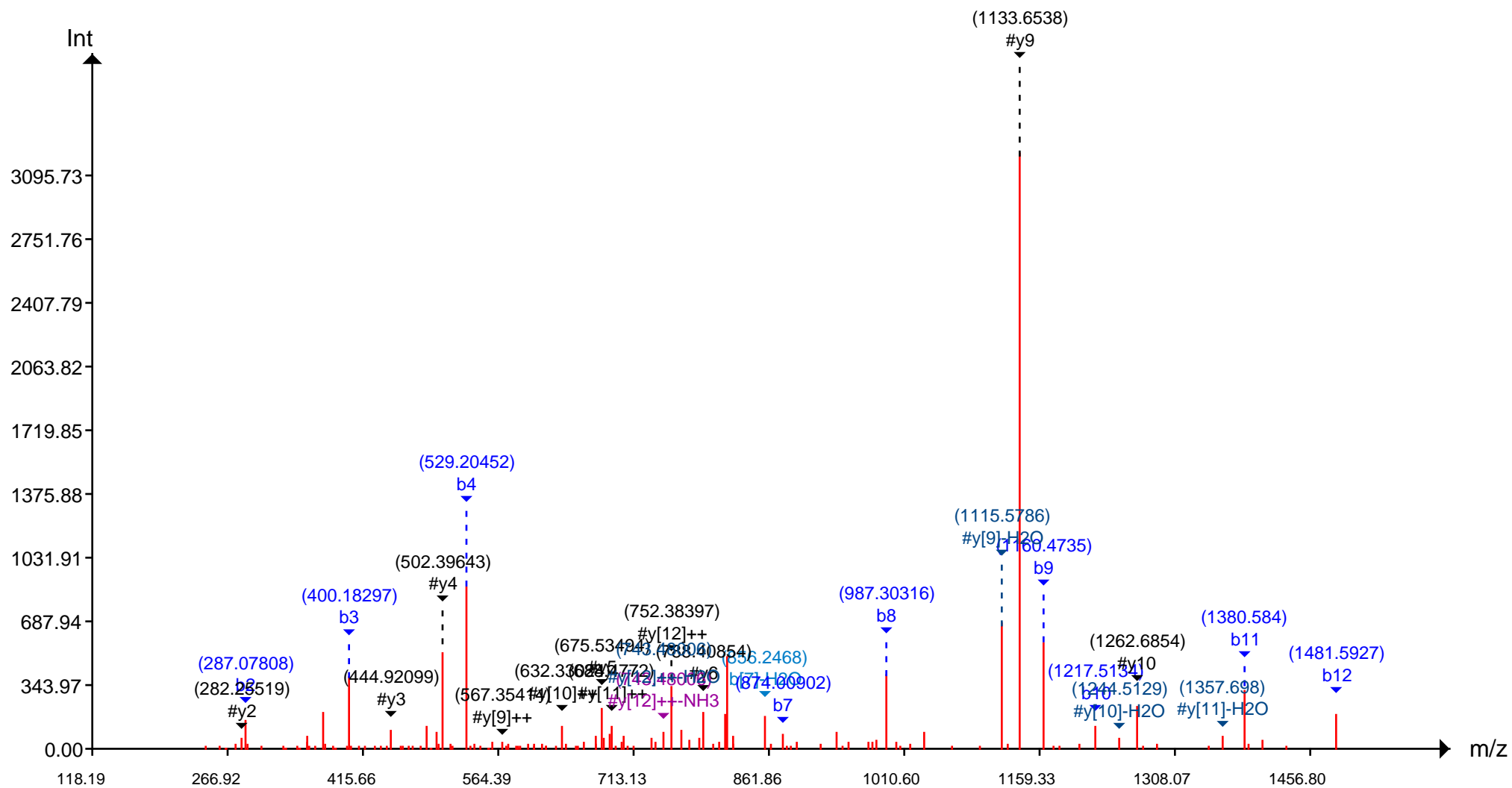
Charge 2+

Mass error 0.002735



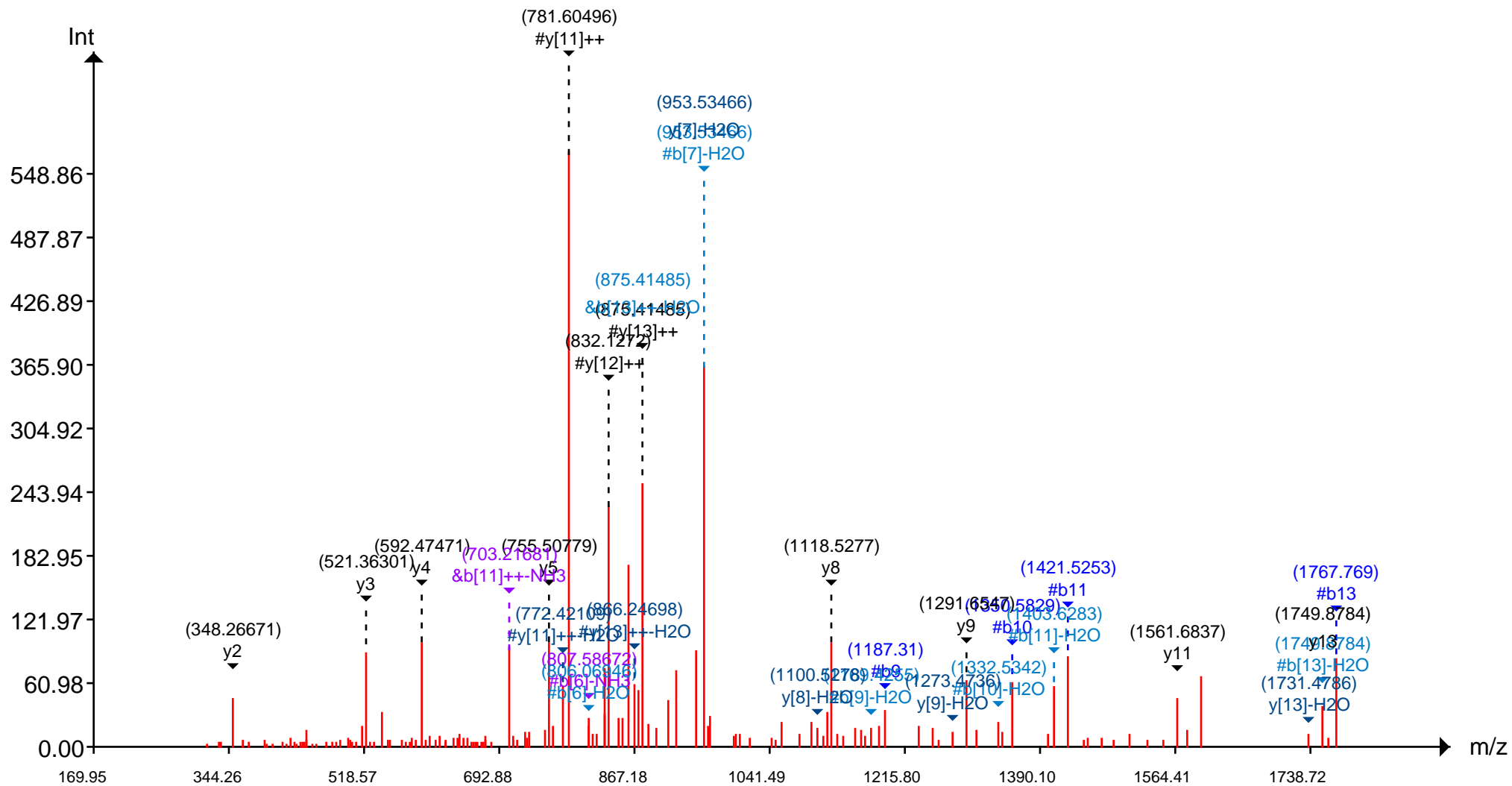
Info

Modified Sequence	AcD3-LAQTEALM<Mox*>K<AcD3K*>GK<AcD3K*>LPEEAR<C13>-COOH	Ion score	48.74
Protein	P06745	Confidence level	99.00%
Precursor m/z	681.37971	Identity Threshold	46.79
Charge	3+	Mass error	2.21E-4



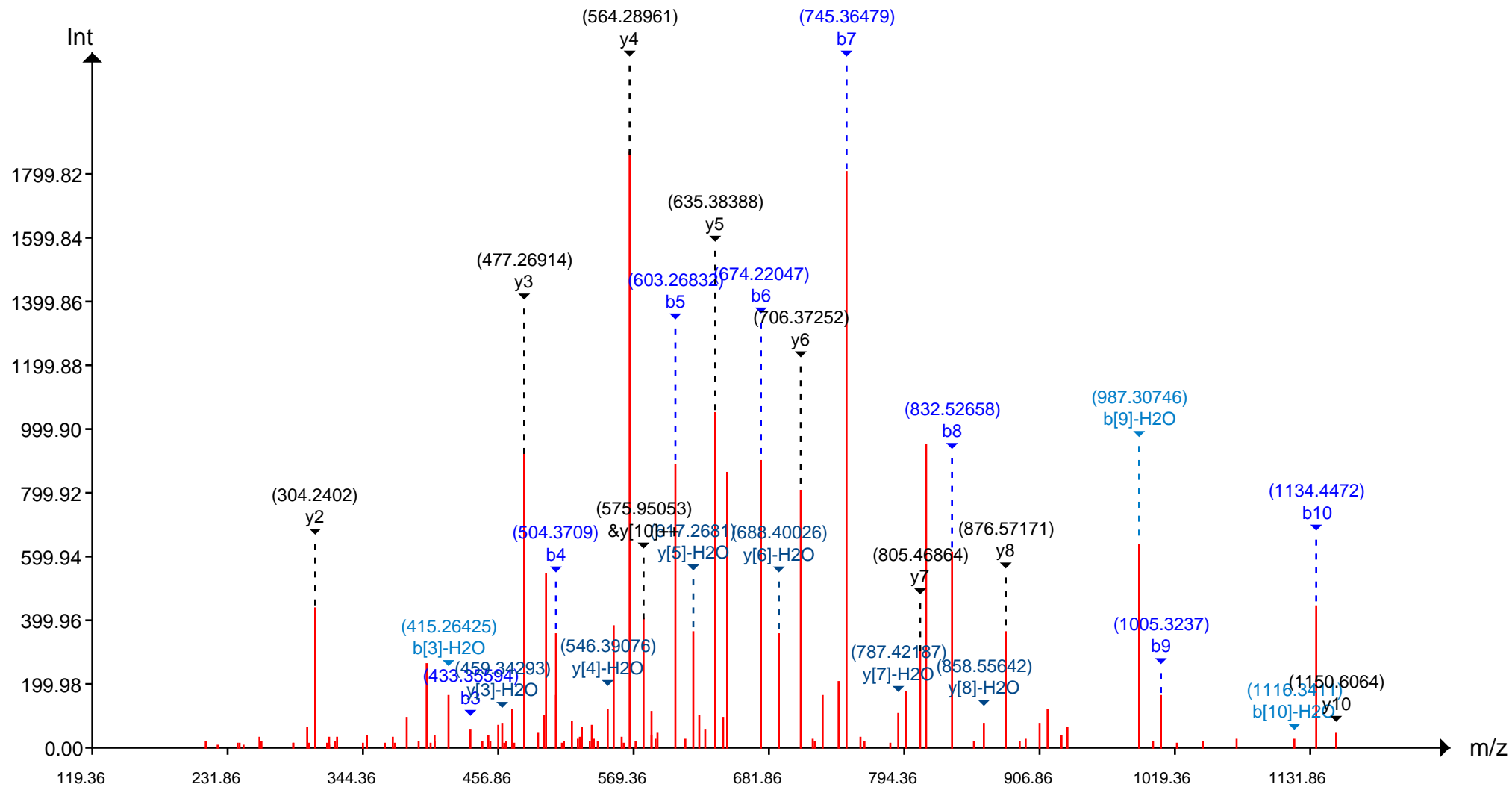
Info

Modified Sequence	AcD3-IQLEPTFIK<AcD3K*>GYTR<C13>-COOH	Ion score	71.22
Protein	Q60864	Confidence level	99.00%
Precursor m/z	831.47902	Identity Threshold	45.59
Charge	2+	Mass error	0.003241



Info

Modified Sequence	AcD3-FSTPK<AcD3K*>K<AcD3K*>M<Mox*>TDYAK<AcD3K*>K<AcD3K*>R-COOH	Ion score	50.07
Protein	Q9CX56	Confidence level	99.00%
Precursor m/z	971.53354	Identity Threshold	47.20
Charge	2+	Mass error	0.002678



Info

Modified Sequence AcD3-ITK<AcD3K*>AVAASK<AcD3K*>ER-COOH

Ion score 49.72

Protein P15864, P43277, P43274

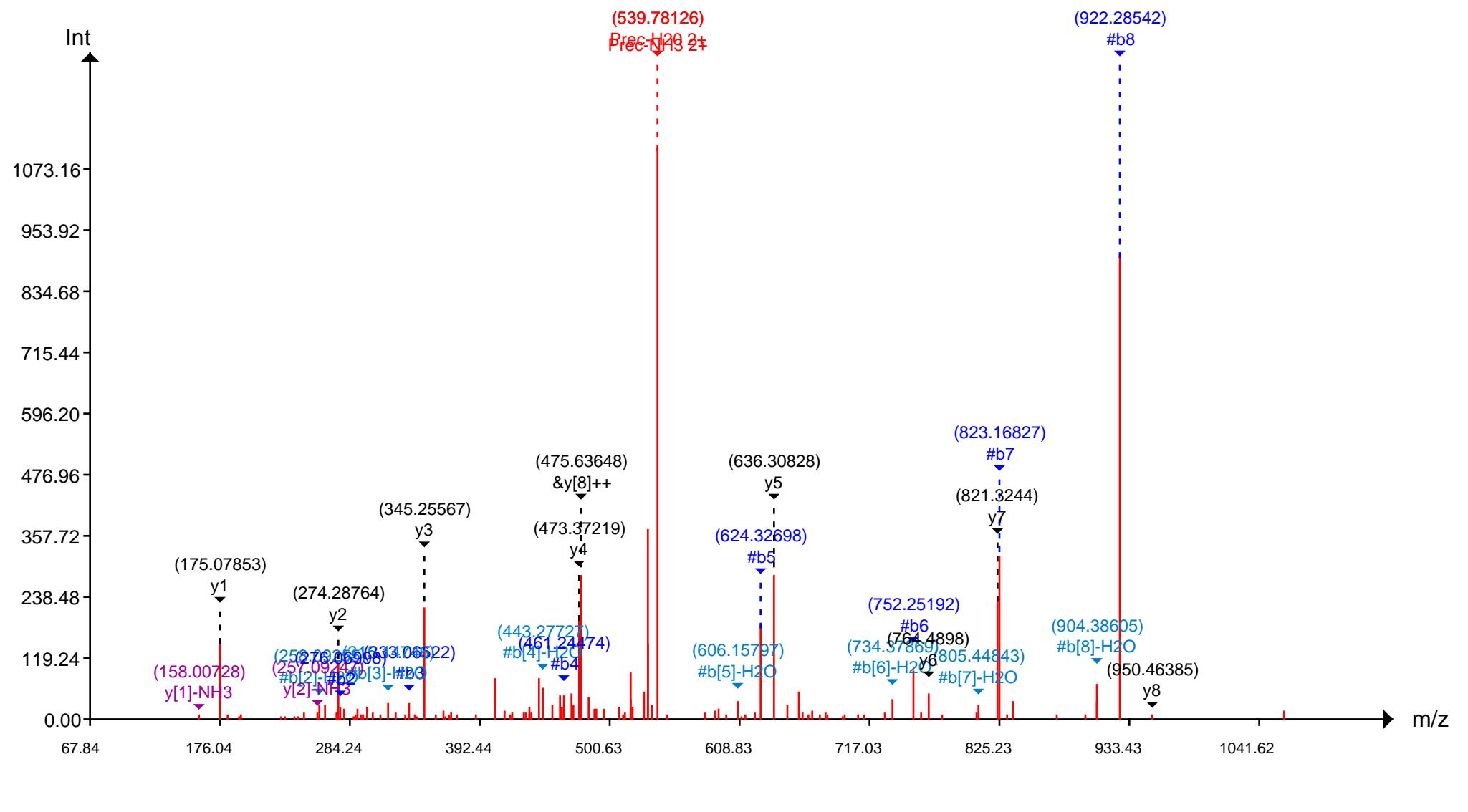
Confidence level 99.00%

Precursor m/z 654.89575

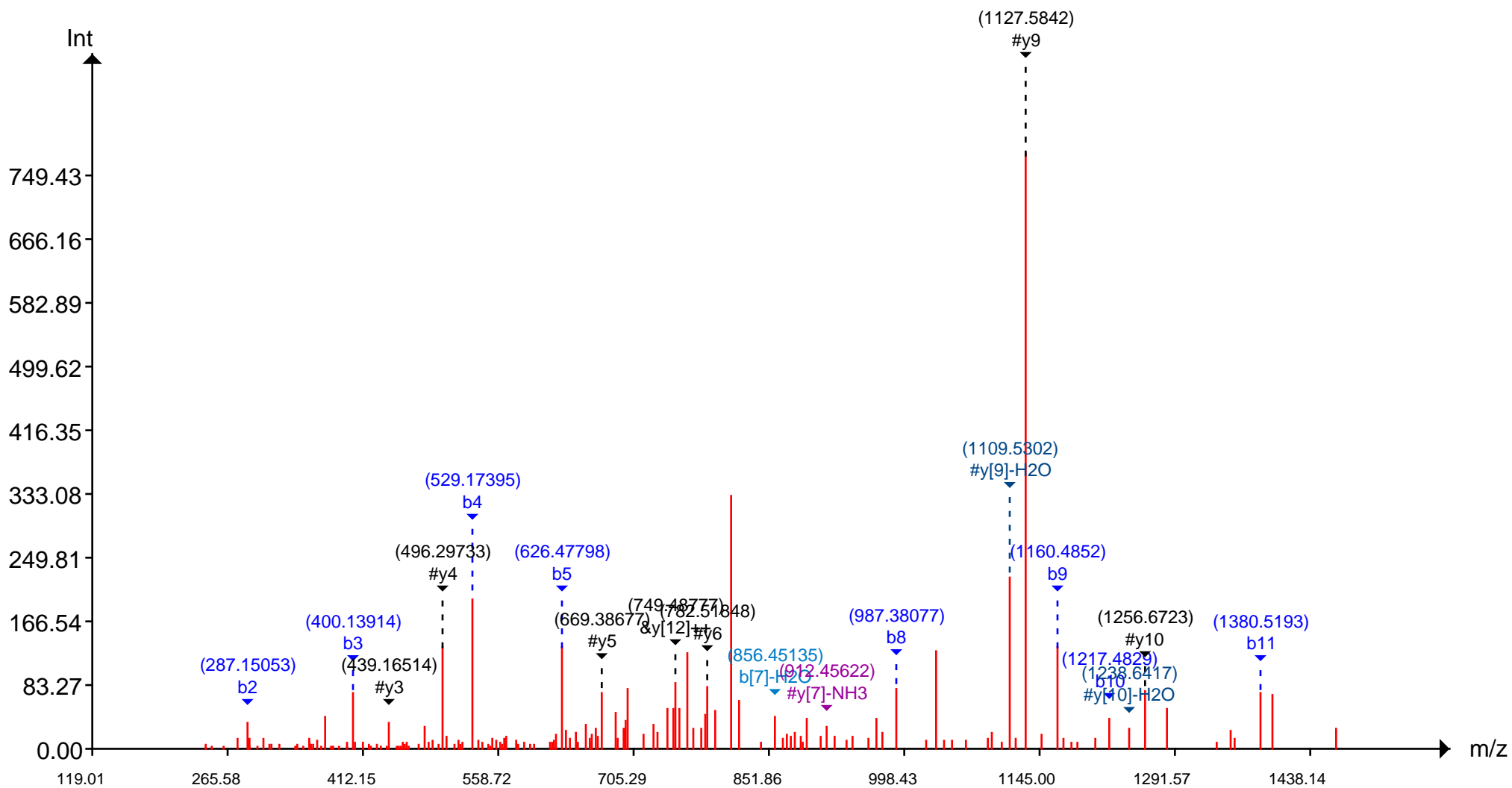
Identity Threshold 44.08

Charge 2+

Mass error 0.001069



Info			
Modified Sequence	AcD3-TEGQYQAVR-COOH	Ion score	58.72
Protein	Q9EPB4	Confidence level	99.00%
Precursor m/z	548.77672	Identity Threshold	47.13
Charge	2+	Mass error	8.5E-5



Info

Modified Sequence AcD3-IQLEPTFIK<AcD3K*>GYTR-COOH

Ion score 49.33

Protein Q60864

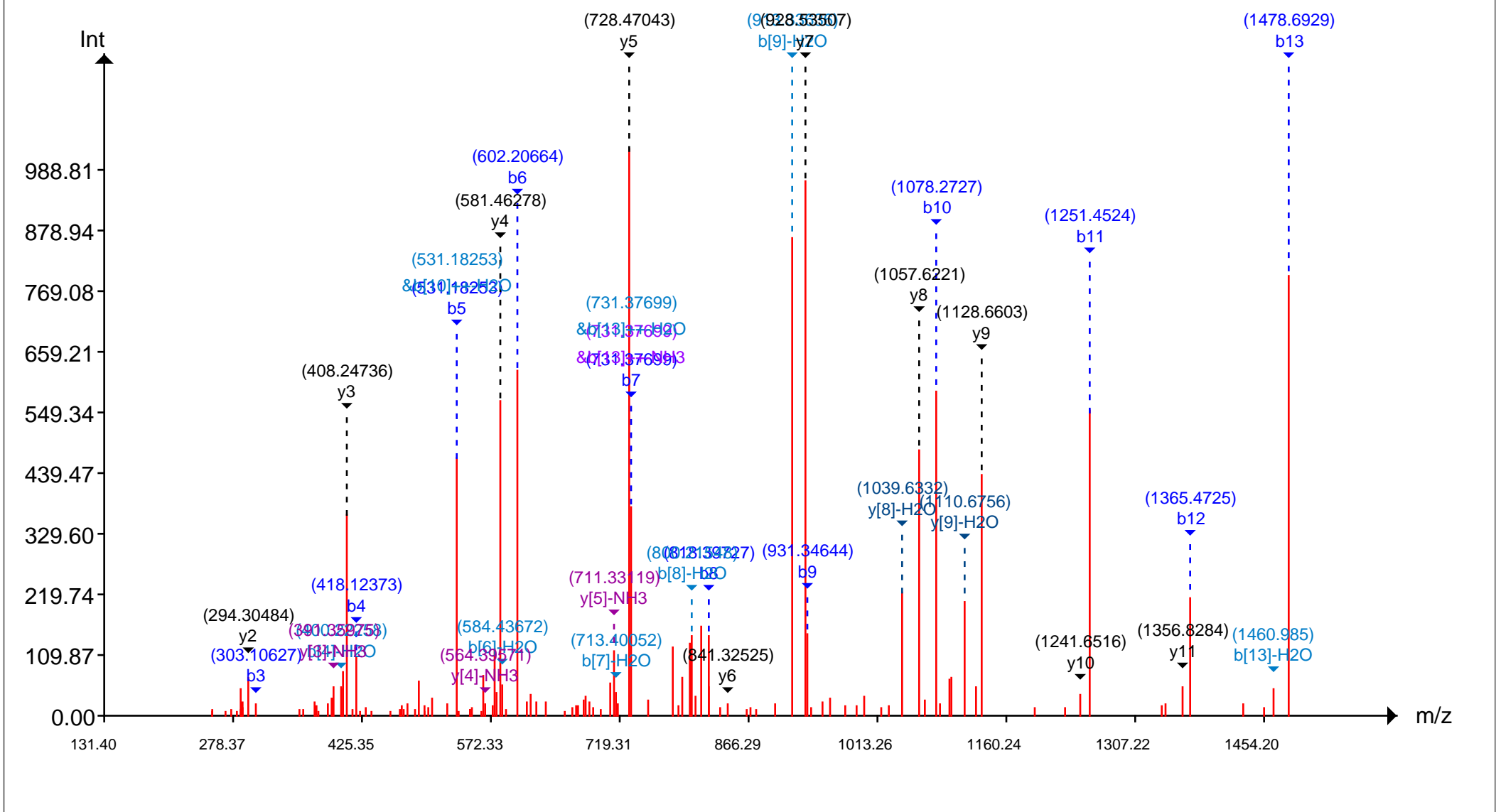
Confidence level 99.00%

Precursor m/z 828.46879

Identity Threshold 46.59

Charge 2+

Mass error 0.002923



Info

Modified Sequence AcD3-SVADLAESIM<Mox*>K<AcD3K*>NLR<C13>-COOH

Ion score 82.04

Protein P06151

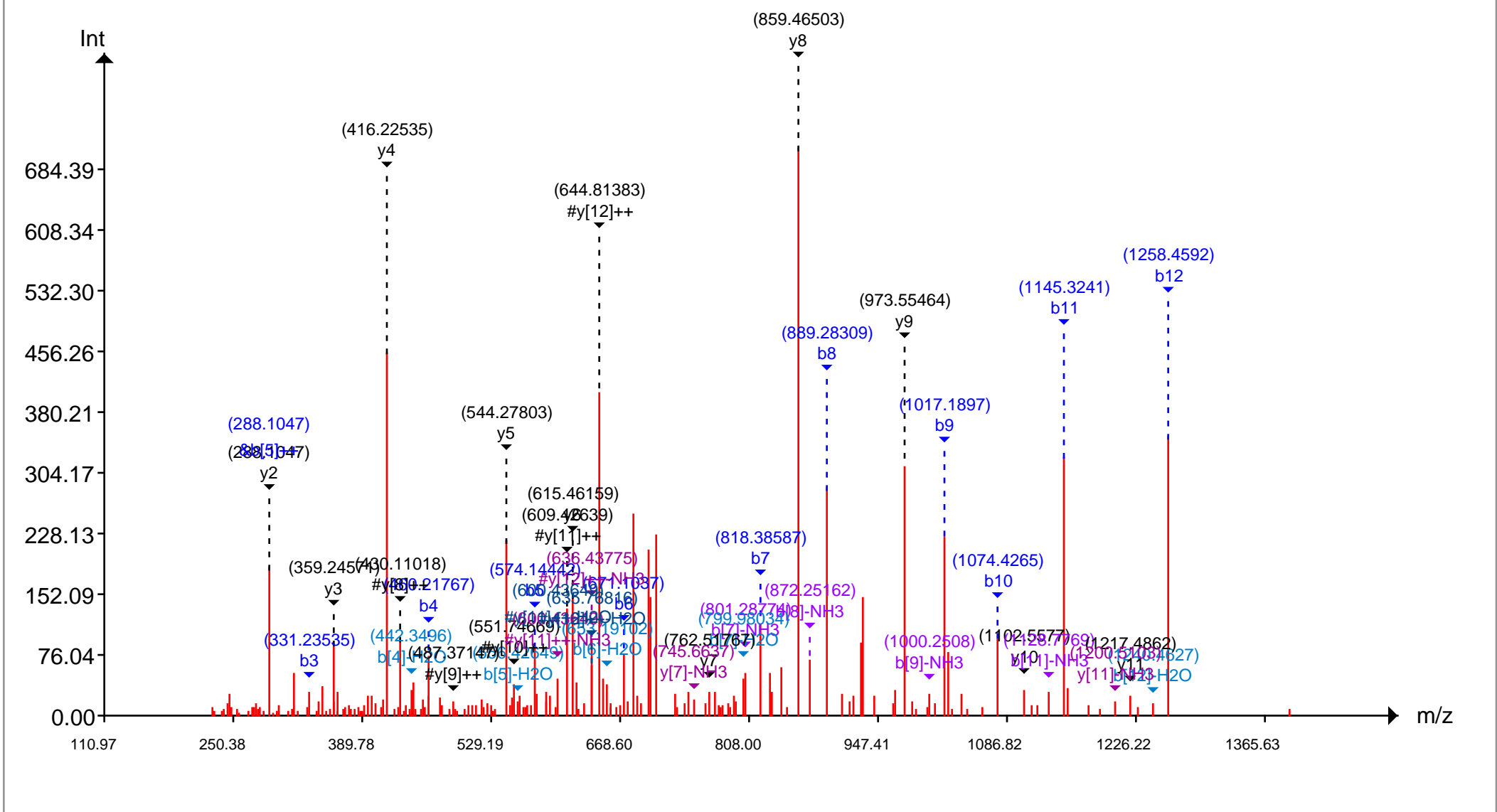
Confidence level 99.00%

Precursor m/z 829.95423

Identity Threshold 47.69

Charge 2+

Mass error 0.00162



Info

Modified Sequence AcD3-VADENPFAQGALR-COOH

Ion score 72.63

Protein P13020

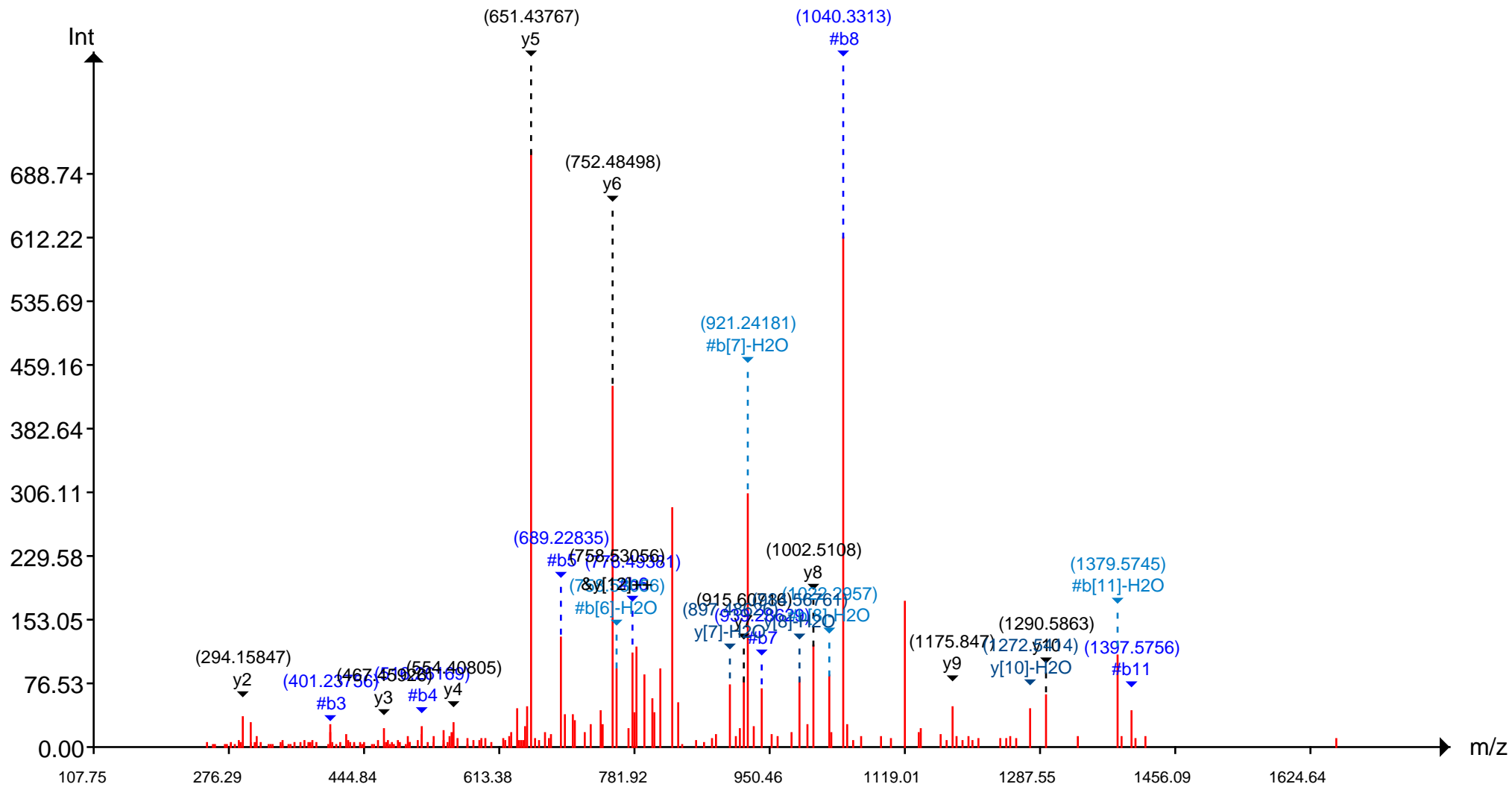
Confidence level 99.00%

Precursor m/z 716.86736

Identity Threshold 47.91

Charge 2+

Mass error 0.001616



Info

Modified Sequence AcD3-EIIDK<AcD3K*>SYTPSK<AcD3K*>IR<C13>-COOH

Ion score 58.01

Protein O70492

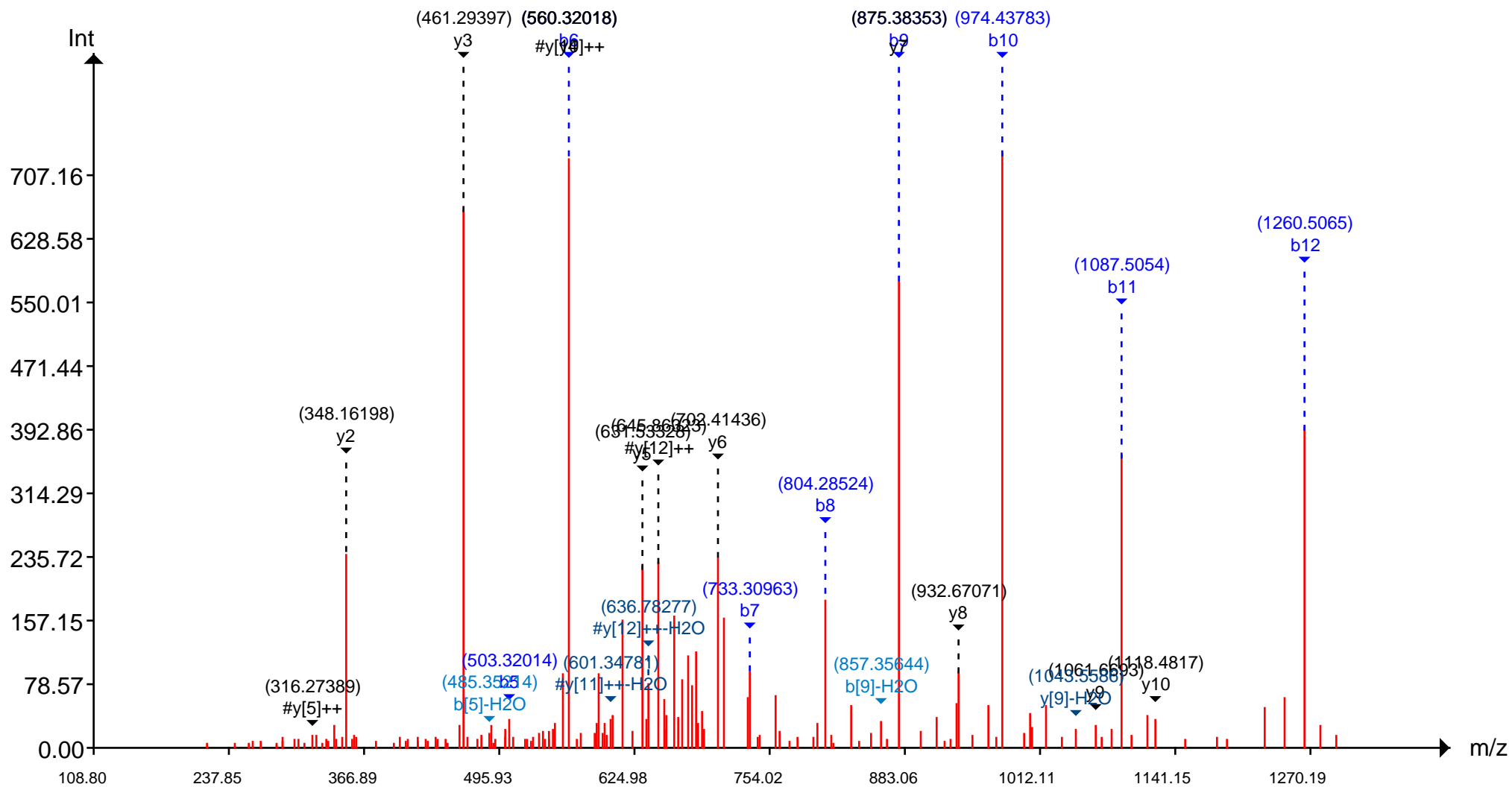
Confidence level 99.00%

Precursor m/z 845.98758

Identity Threshold 45.48

Charge 2+

Mass error 0.001151



Info

Modified Sequence AcD3-VATGEGK<AcD3K*>AAVLK<AcD3K*>R-COOH

Ion score 69.07

Protein Q9CQ60

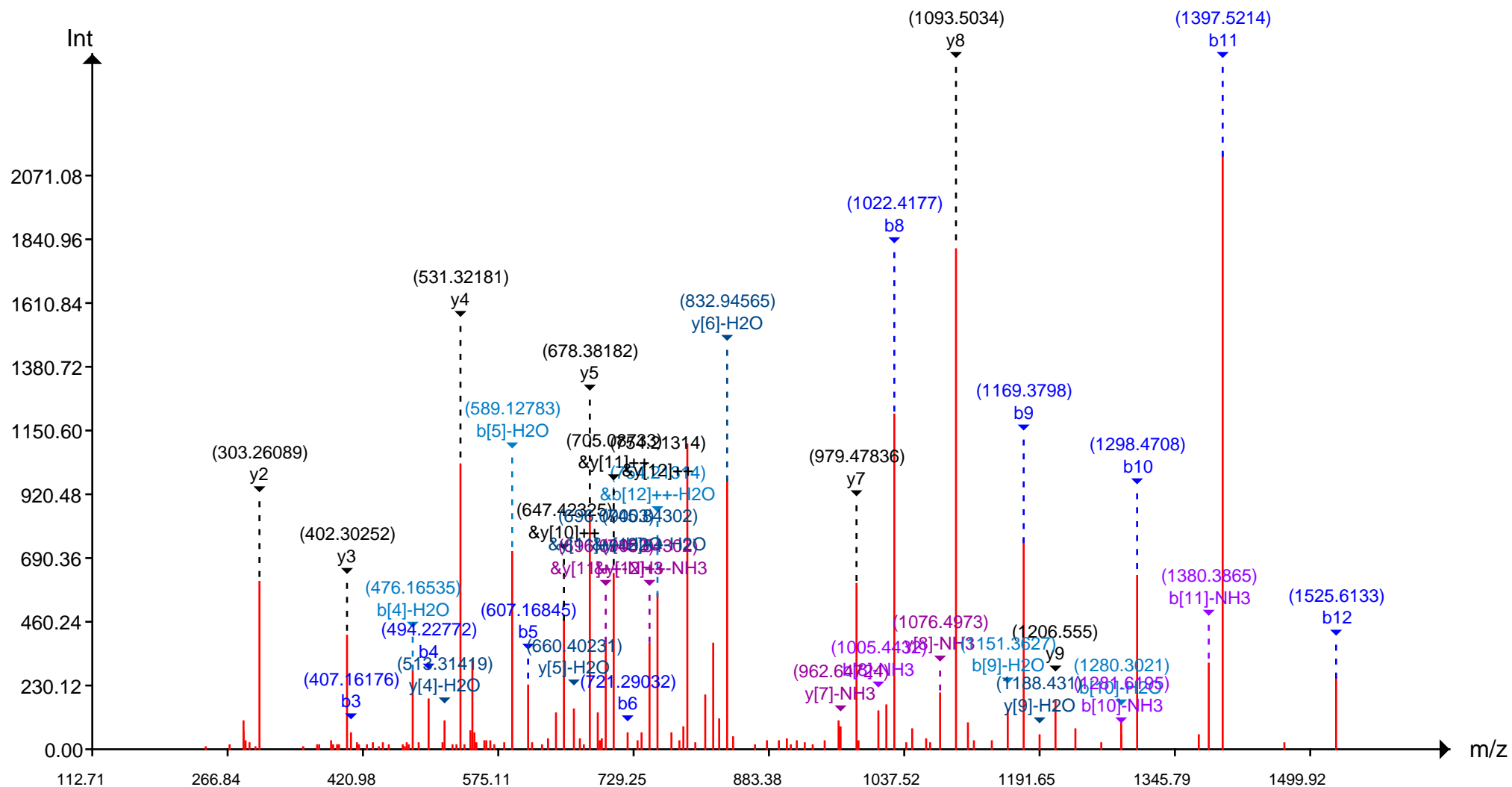
Confidence level 99.00%

Precursor m/z 717.93532

Identity Threshold 43.04

Charge 2+

Mass error 8.79E-4



Info

Modified Sequence AcD3-FVDSL NQK<AcD3K*>M<Mox*>EVQR-COOH

Ion score 54.65

Protein P16675

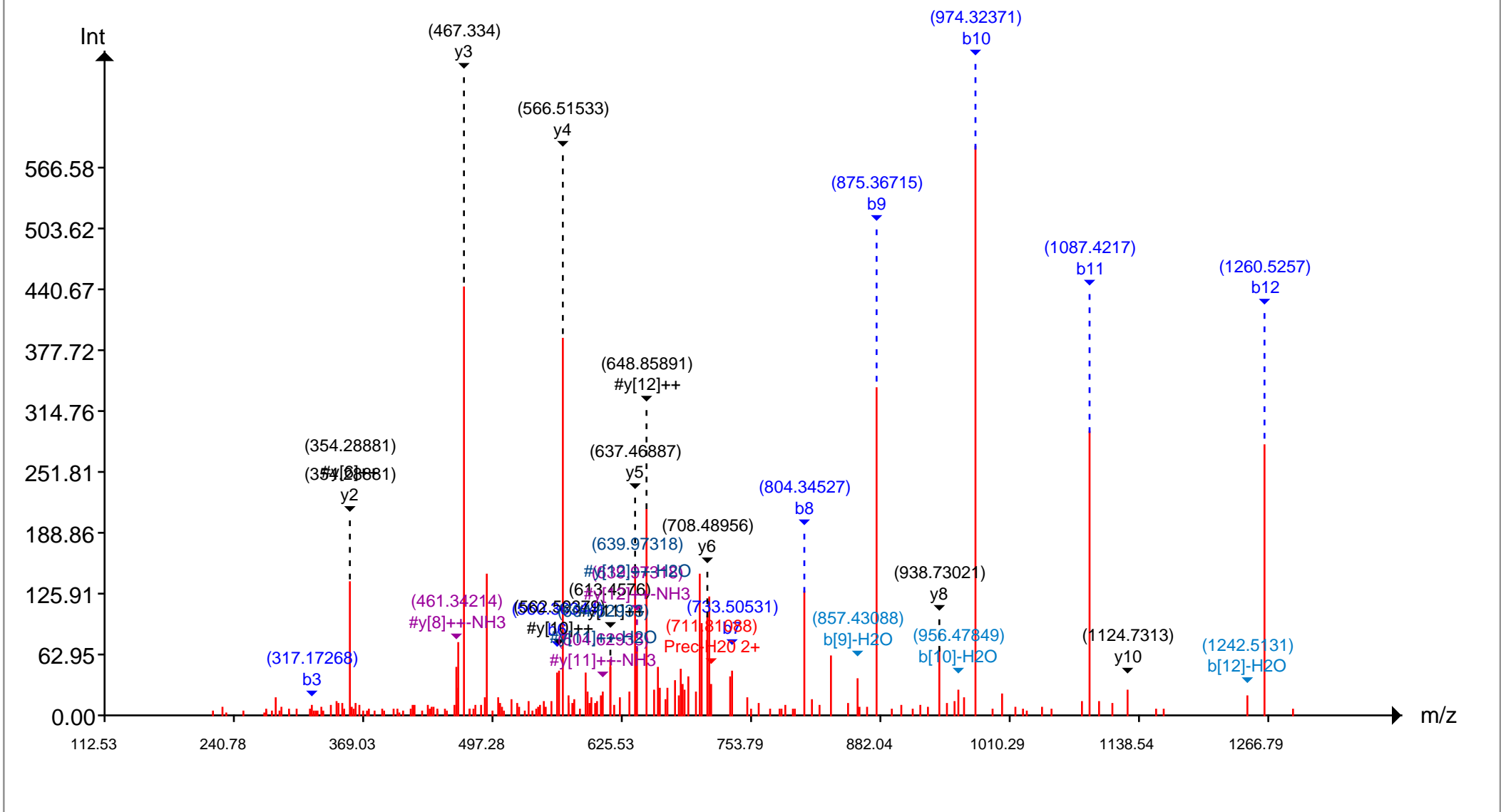
Confidence level 99.00%

Precursor m/z 850.4336

Identity Threshold 47.88

Charge 2+

Mass error 8.72E-4



Info

Modified Sequence AcD3-VATGEGK<AcD3K*>AAVLK<AcD3K*>R<C13>-COOH

Ion score 55.27

Protein Q9CQ60

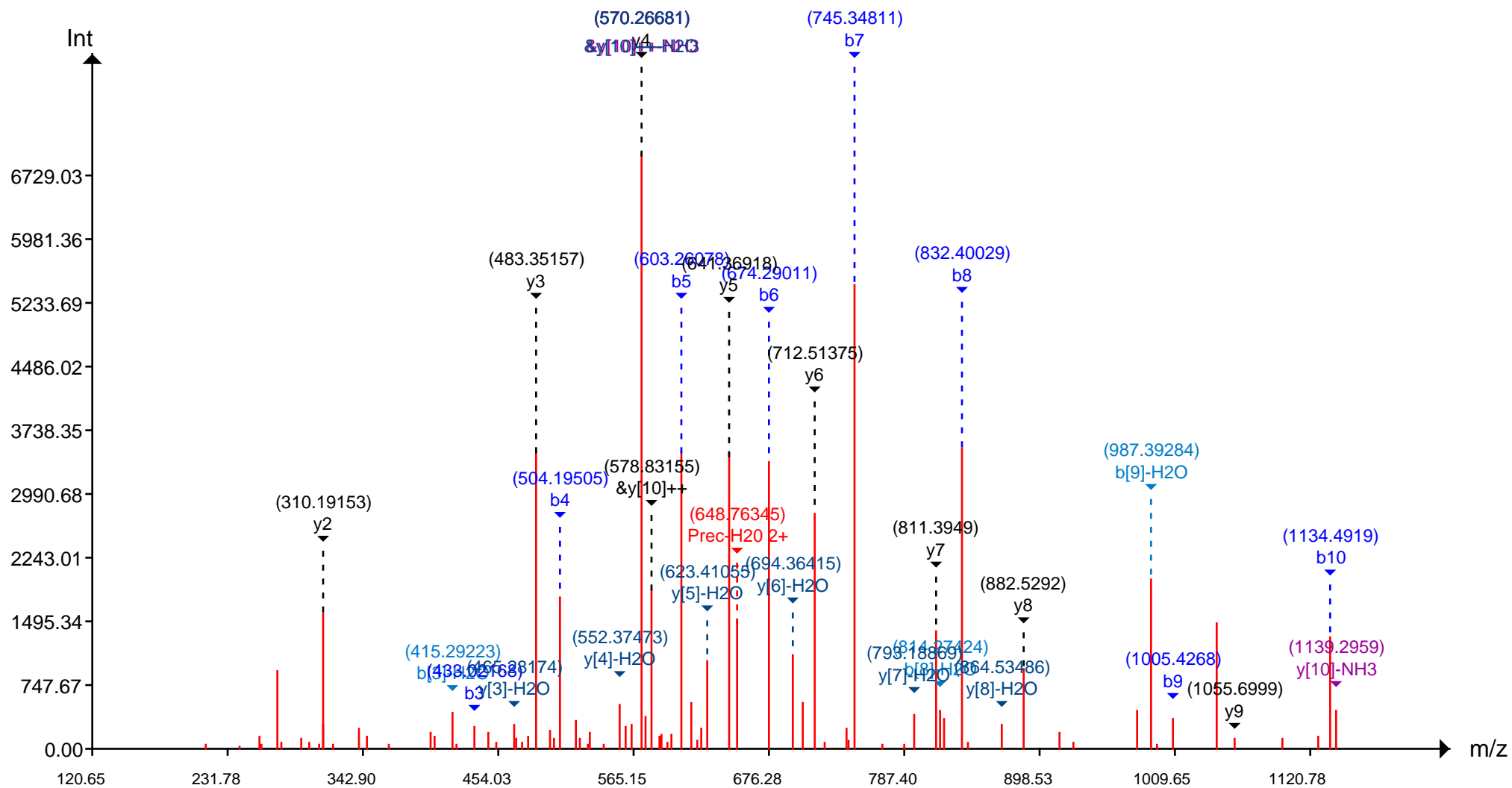
Confidence level 99.00%

Precursor m/z 720.9454

Identity Threshold 41.96

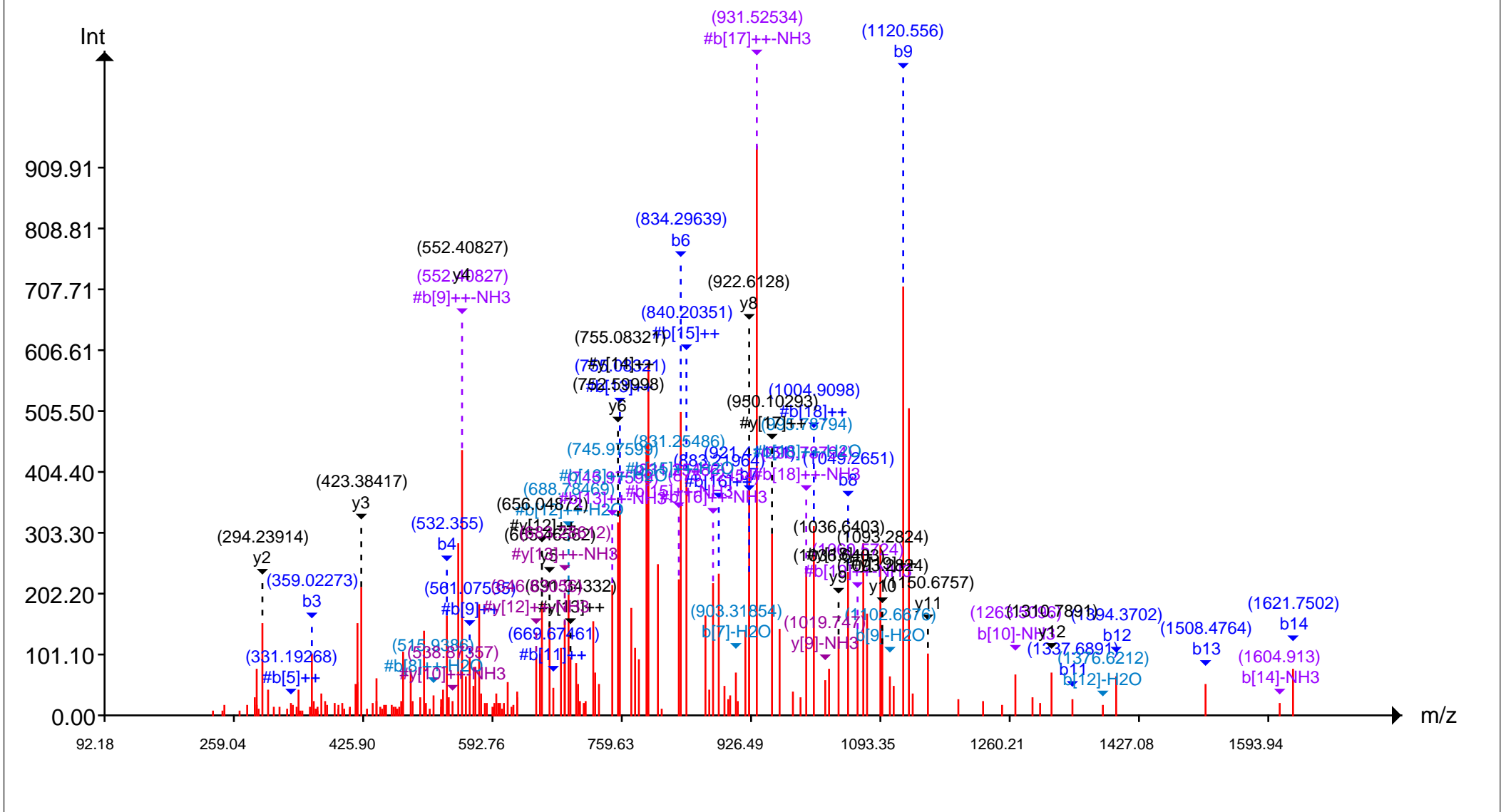
Charge 2+

Mass error 8.97E-4



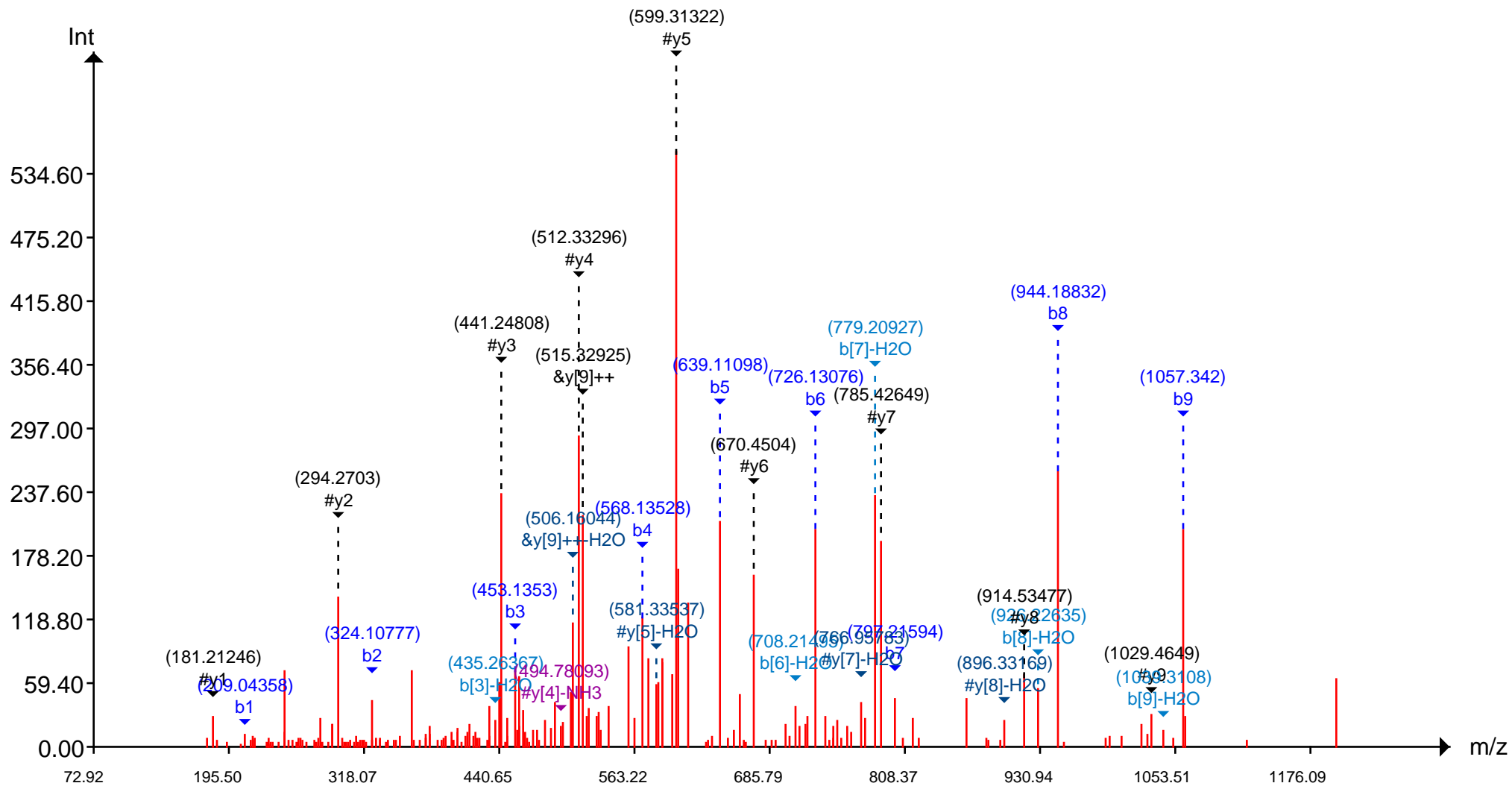
Info

Modified Sequence	AcD3-ITK<AcD3K*>AVAASK<AcD3K*>ER<C13>-COOH	Ion score	63.43
Protein	P15864, P43277, P43274	Confidence level	99.00%
Precursor m/z	657.90526	Identity Threshold	42.82
Charge	2+	Mass error	-5.3E-5



Info

Modified Sequence	AcD3-AELK<AcD3K*>EK<AcD3K*>SQAC<Cmm*>GGNLGSIEELR<C13>-COOH	Ion score	63.92
Protein	Q99KH8	Confidence level	99.00%
Precursor m/z	810.7584	Identity Threshold	47.89
Charge	3+	Mass error	0.002029



Info

Modified Sequence AcD3-YDEDASAM<Mox*>LR<C13>-COOH

Ion score 68.34

Protein Q99L47

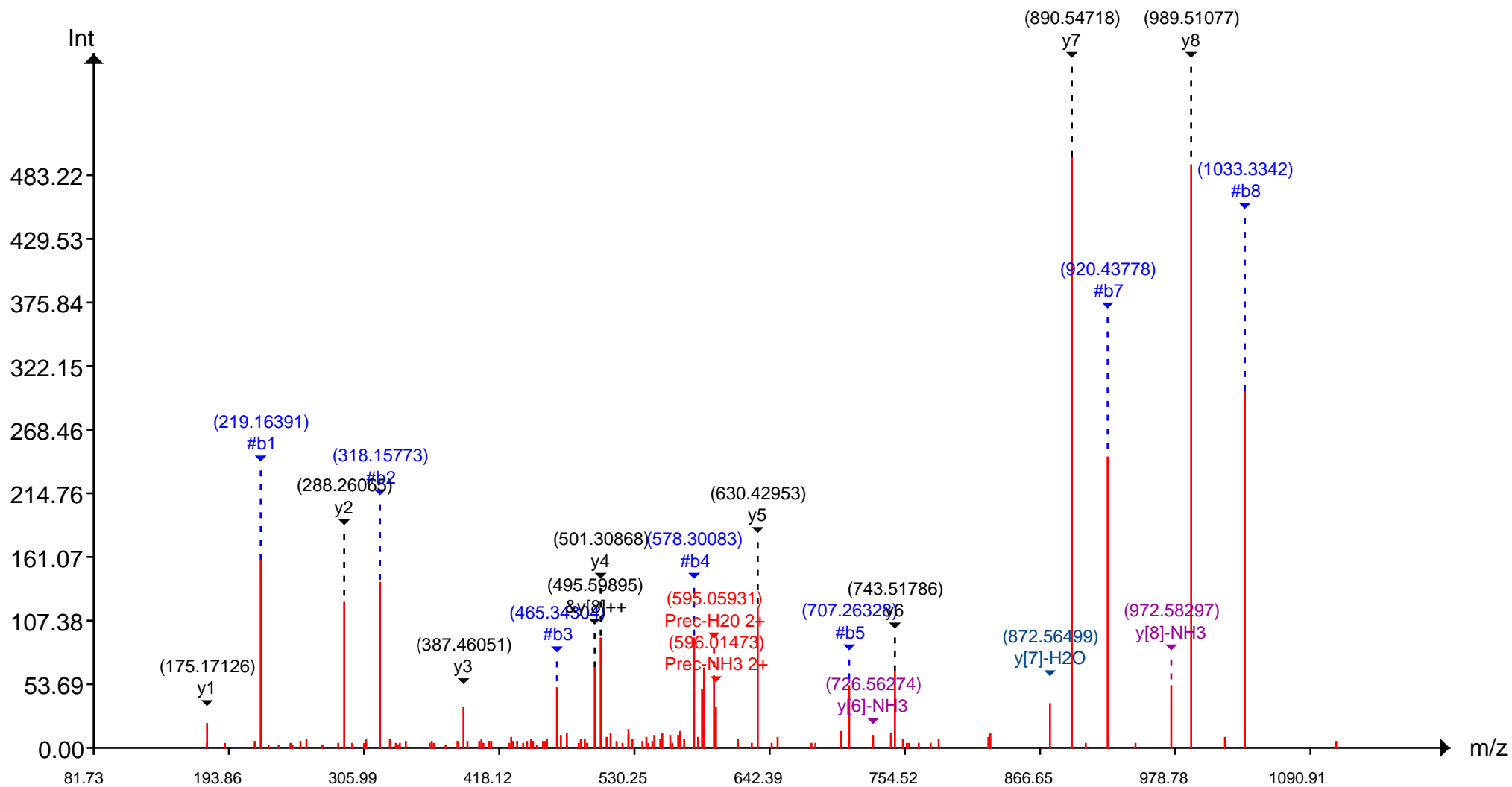
Confidence level 99.00%

Precursor m/z 619.28105

Identity Threshold 44.46

Charge 2+

Mass error 8.41E-4



Info

Modified Sequence AcD3-K<AcD3K*>VFLENVIR-COOH

Ion score 78.16

Protein P62806

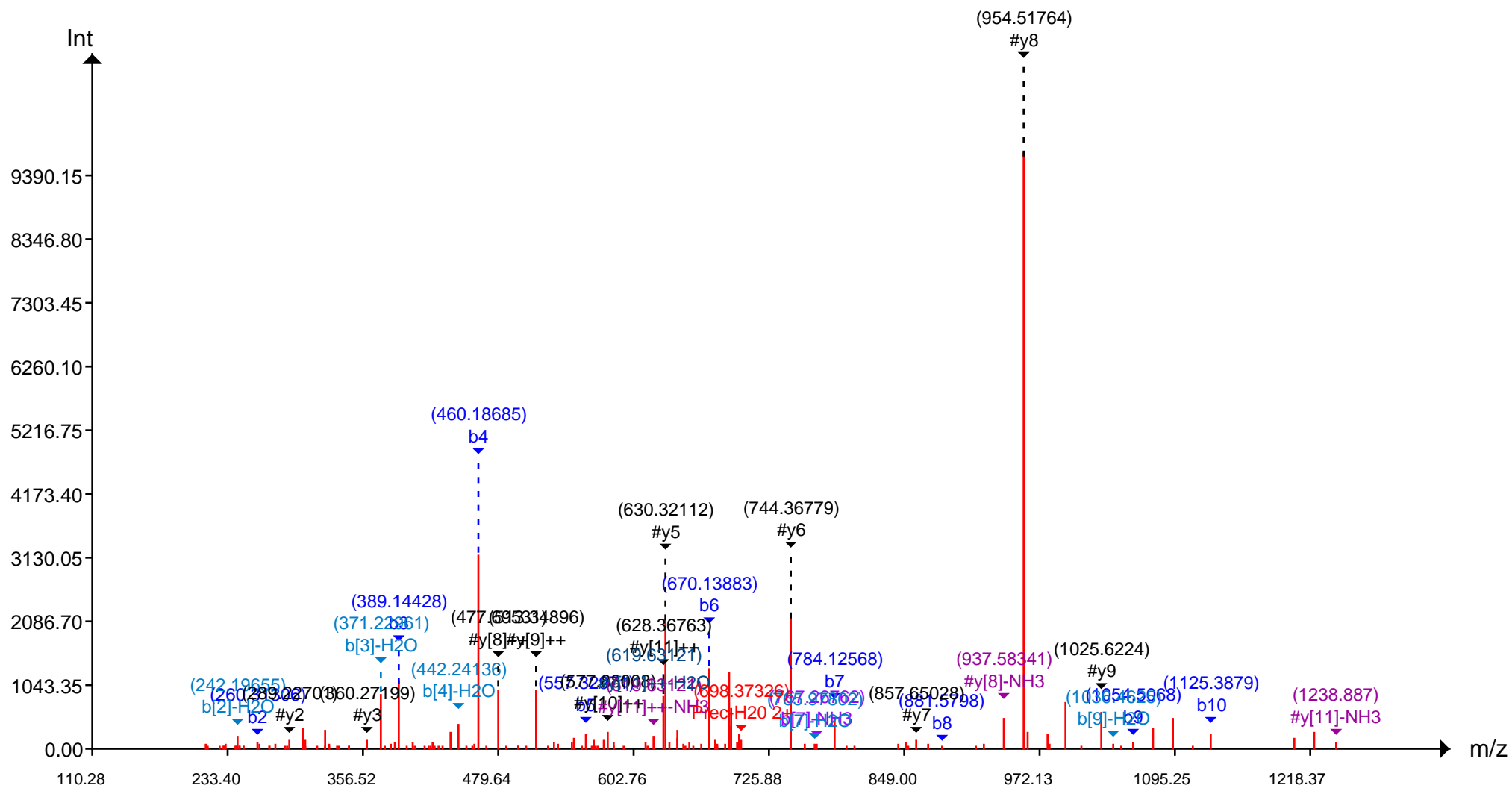
Confidence level 99.00%

Precursor m/z 604.36969

Identity Threshold 43.65

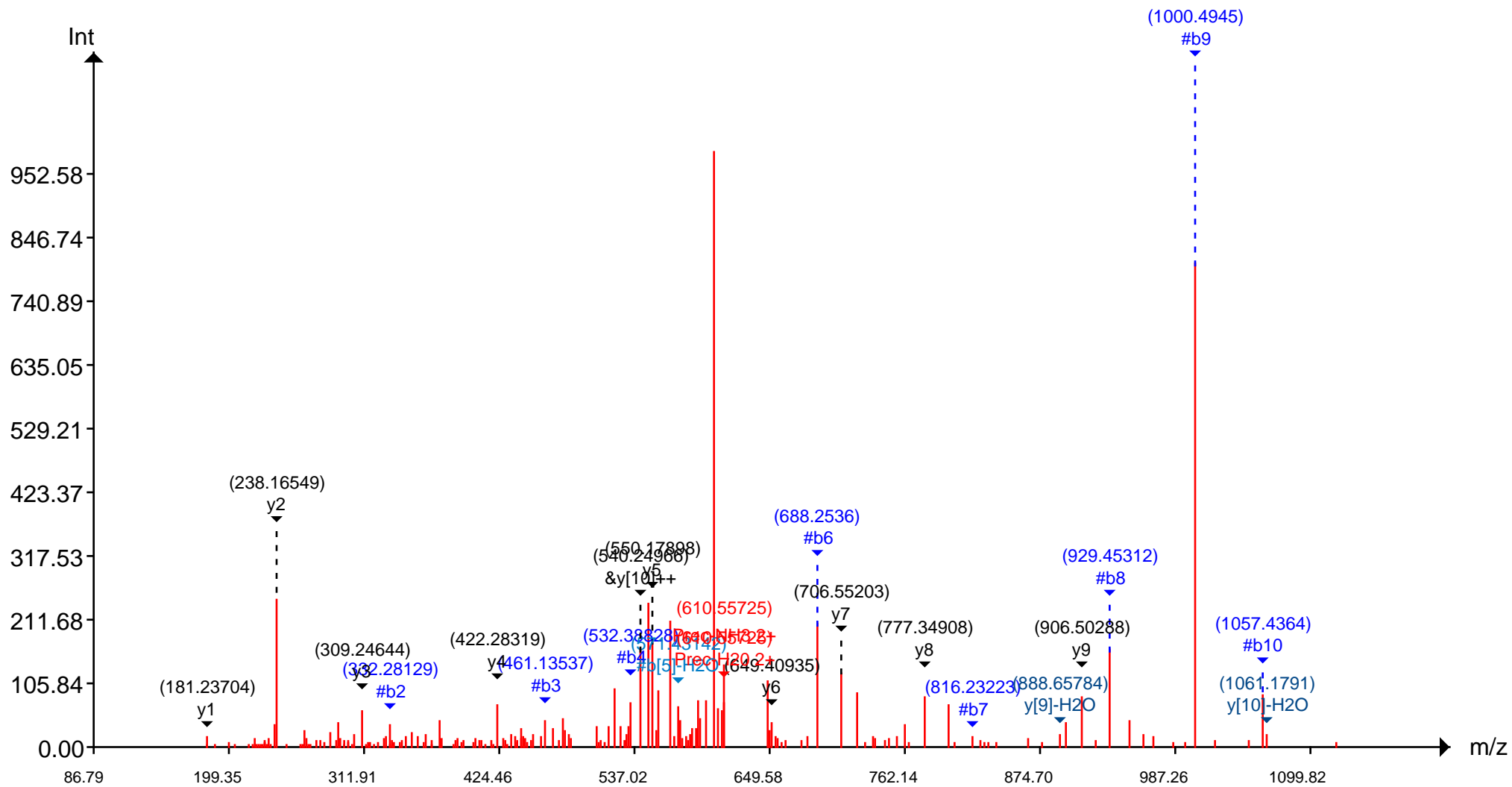
Charge 2+

Mass error 5.24E-4



Info

Modified Sequence	AcD3-LTEAPLNPK<AcD3K*>ANR-COOH	Ion score	58.82
Protein	P62737, P60710, P68033, P63260, P63268, P68134	Confidence level	99.00%
Precursor m/z	707.40359	Identity Threshold	46.76
Charge	2+	Mass error	0.00323



Info

Modified Sequence AcD3-IK<AcD3K*>EAGVQIAGR<C13>-COOH

Ion score 46.55

Protein Q922D8

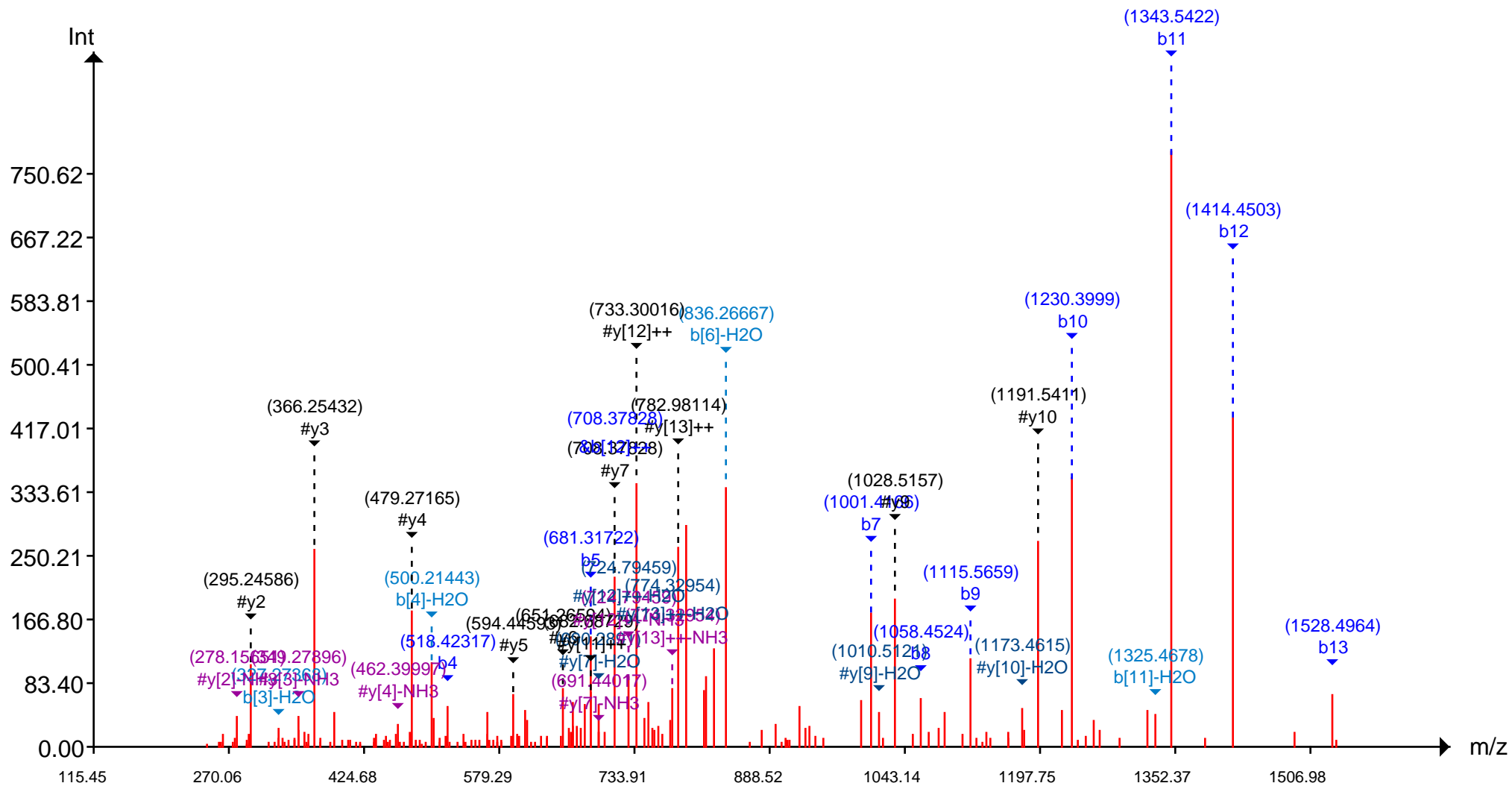
Confidence level 99.00%

Precursor m/z 619.37819

Identity Threshold 43.50

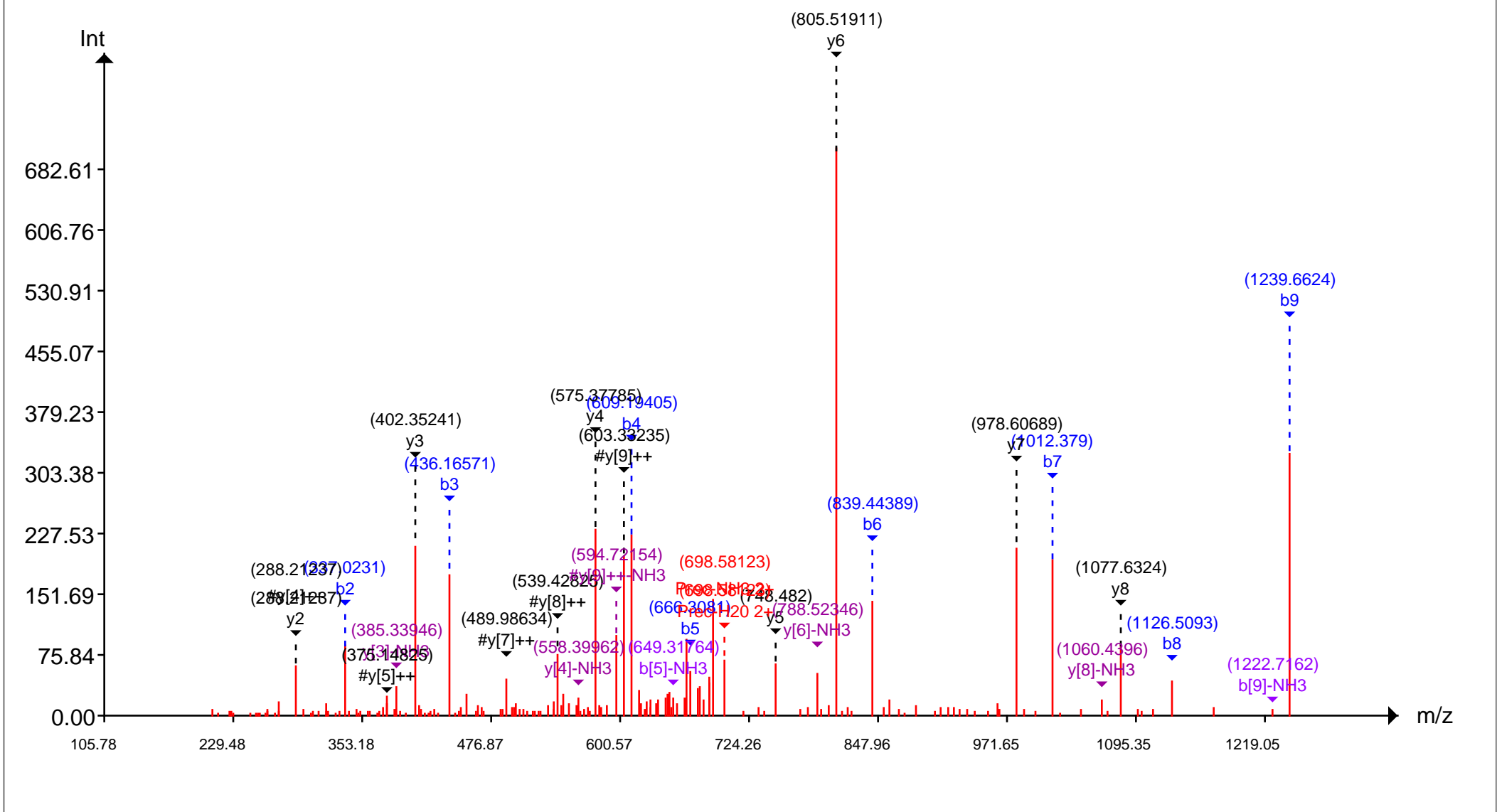
Charge 2+

Mass error 0.001411



Info

Modified Sequence	AcD3-VVTK<AcD3K*>YK<AcD3K*>M<Mox*>GGDIANR<C13>-COOH	Ion score	62.12
Protein	P50580	Confidence level	99.00%
Precursor m/z	854.9722	Identity Threshold	47.44
Charge	2+	Mass error	0.002709



Info

Modified Sequence AcD3-YQVK<AcD3K*>GK<AcD3K*>K<AcD3K*>NIR-COOH

Ion score 61.79

Protein P24452

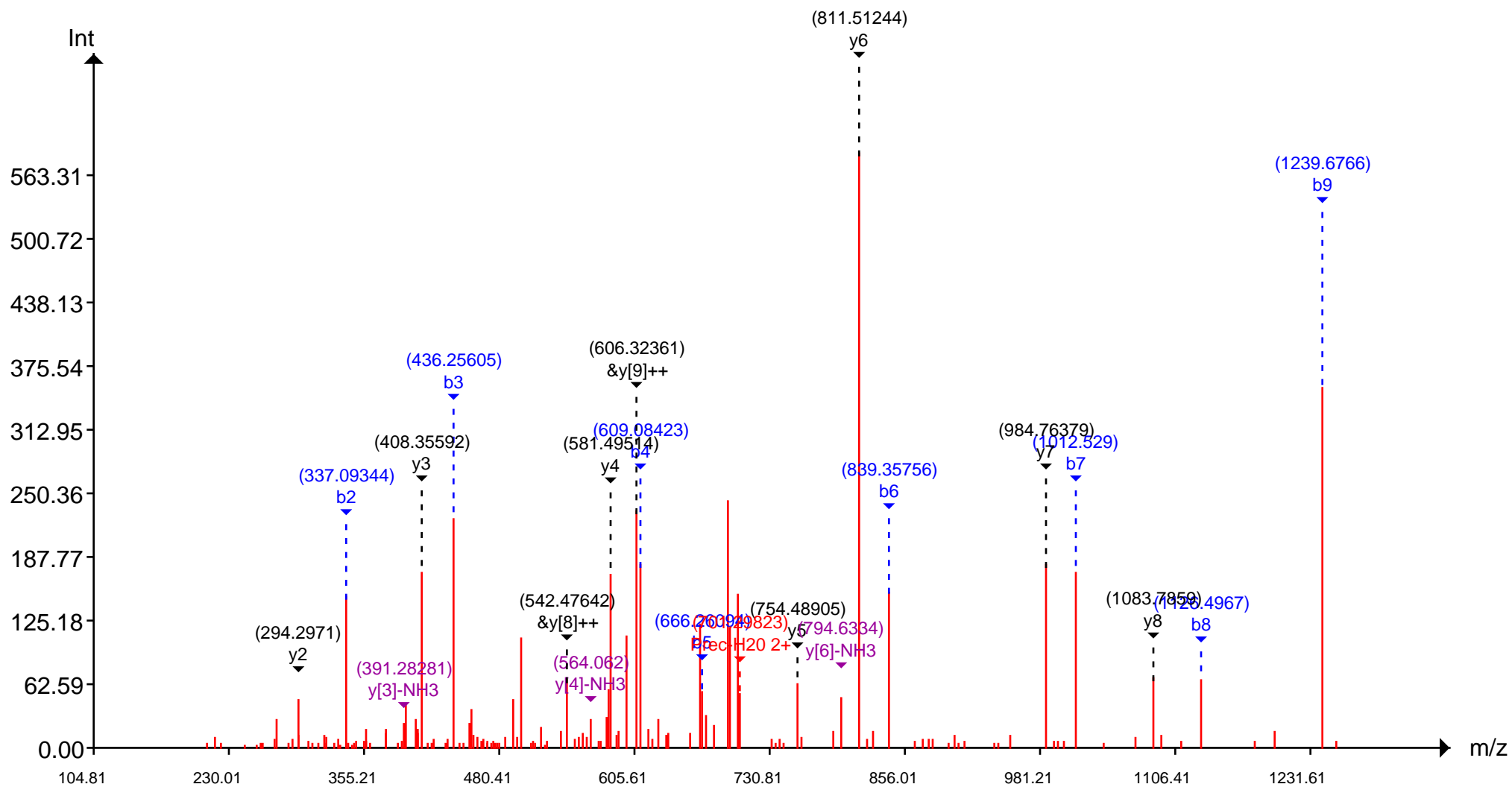
Confidence level 99.00%

Precursor m/z 707.43405

Identity Threshold 42.44

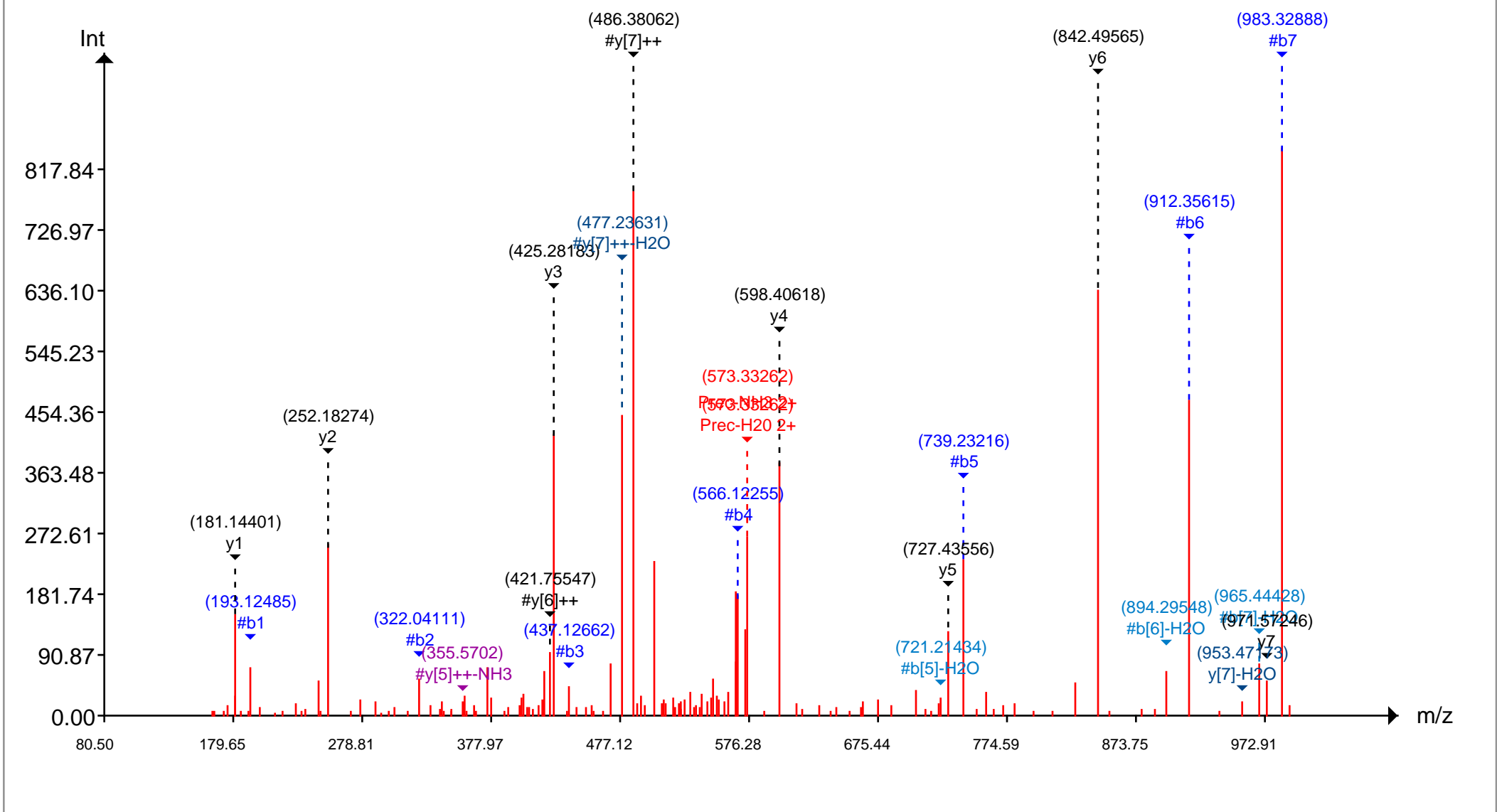
Charge 2+

Mass error 6.43E-4



Info

Modified Sequence	AcD3-YQVK<AcD3K*>GK<AcD3K*>K<AcD3K*>NIR<C13>-COOH	Ion score	50.03
Protein	P24452	Confidence level	99.00%
Precursor m/z	710.44437	Identity Threshold	40.64
Charge	2+	Mass error	0.001141



Info

Modified Sequence AcD3-FEDEK<AcD3K*>K<AcD3K*>AR<C13>-COOH

Ion score 55.96

Protein P50580

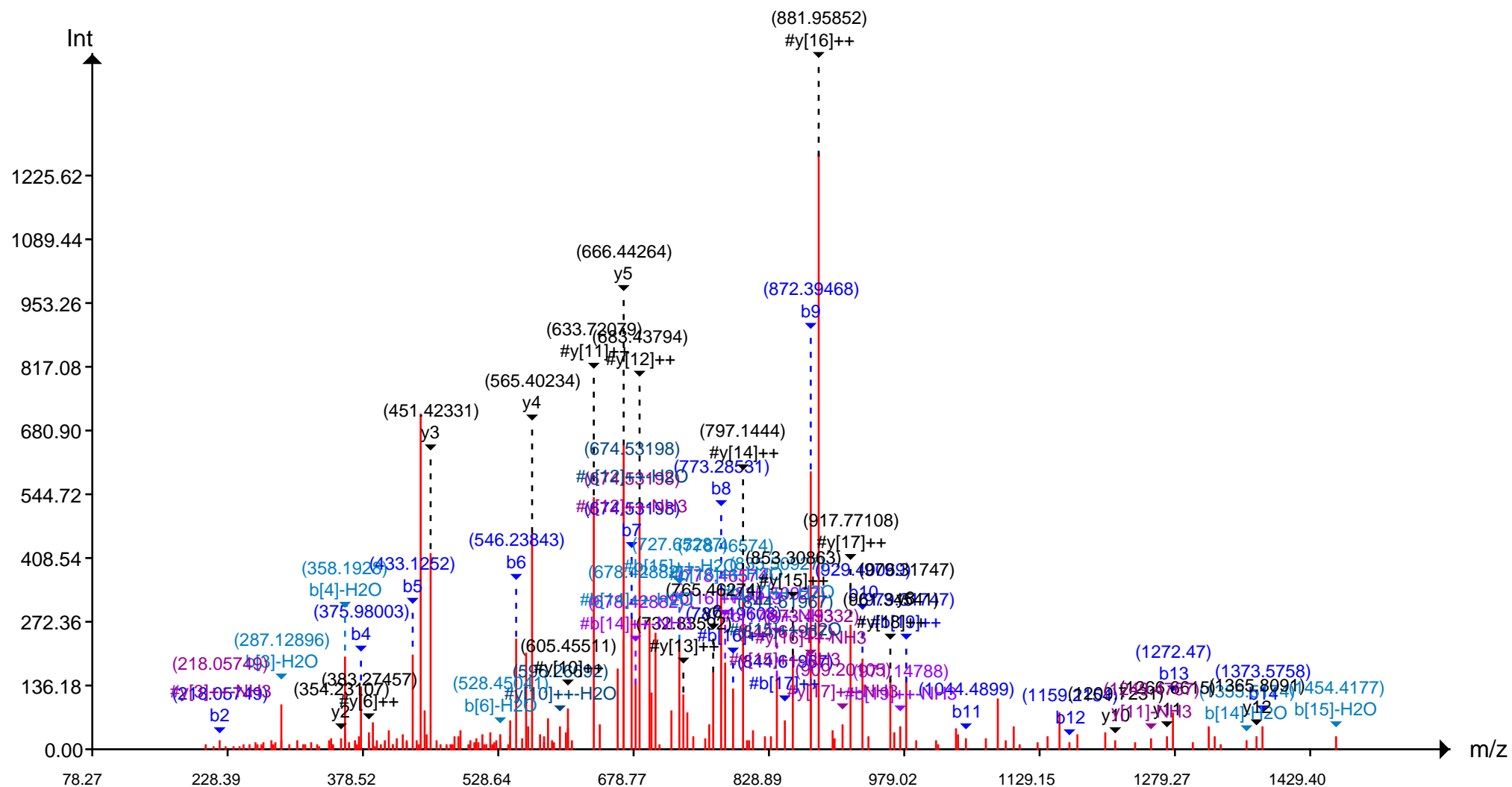
Confidence level 99.00%

Precursor m/z 582.3211

Identity Threshold 47.87

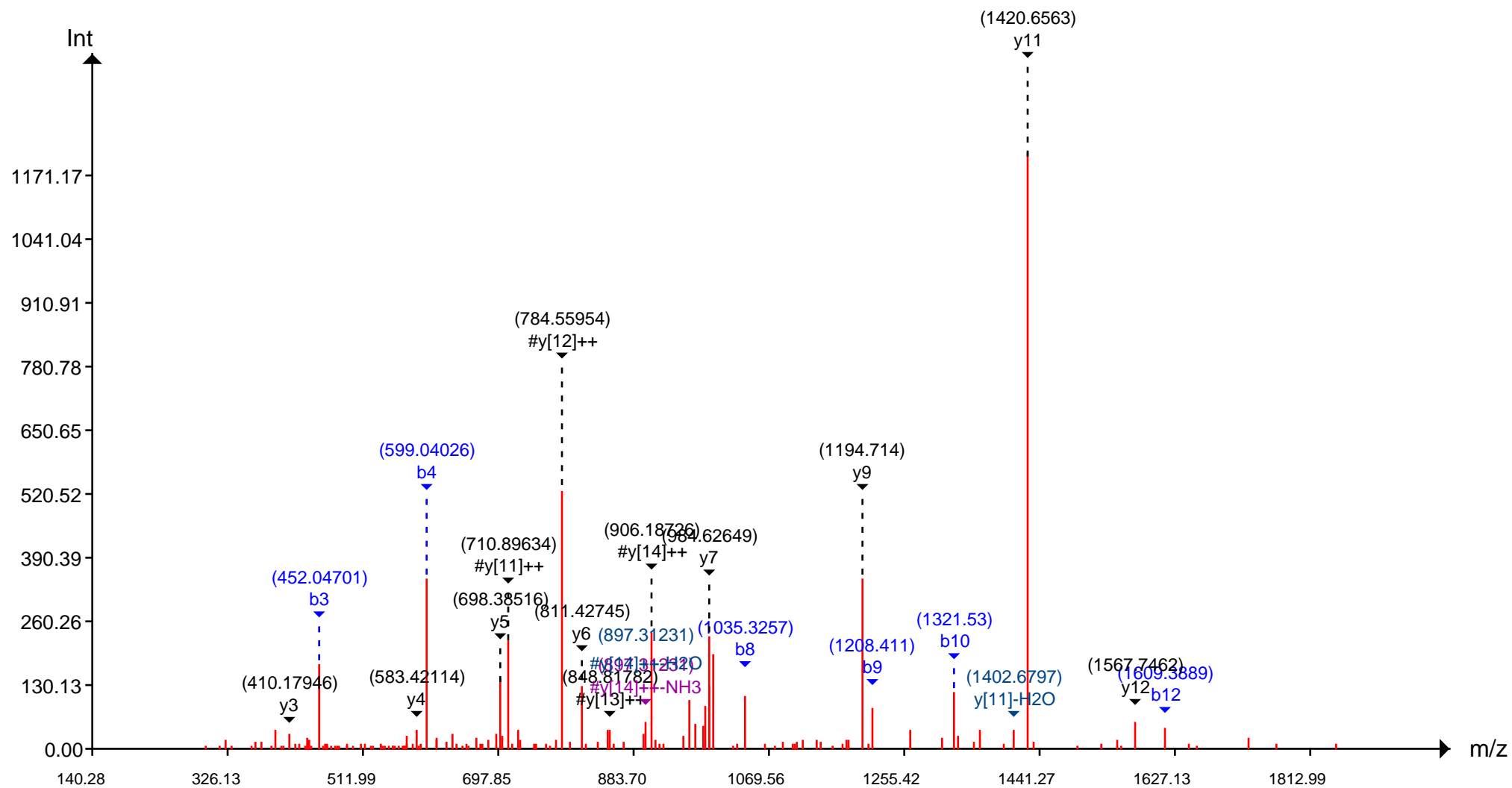
Charge 2+

Mass error 8.5E-5



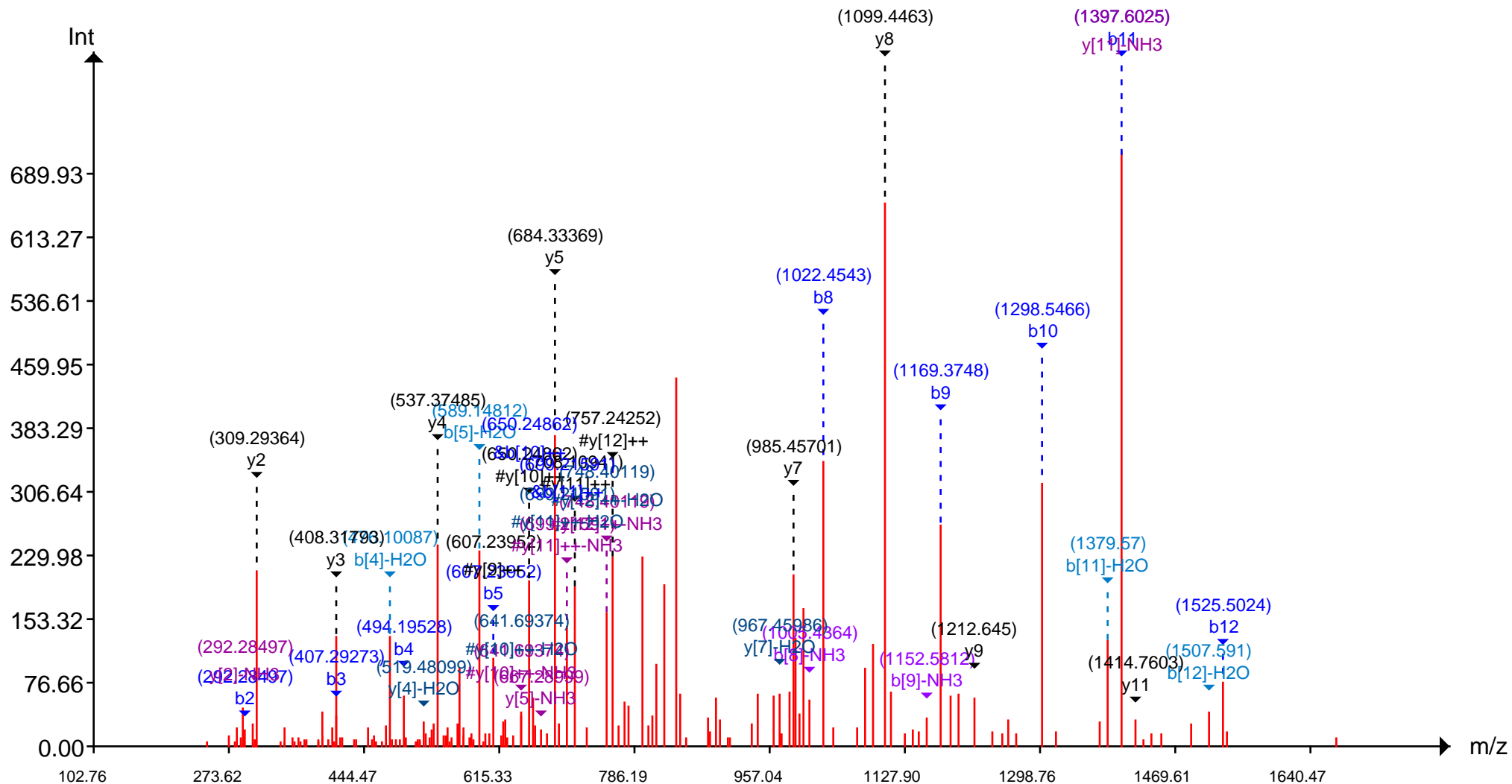
Info

Modified Sequence	AcD3-TASAGIQVVGDDLTVTNPK<AcD3K*>R<C13>-COOH	Ion score	71.32
Protein	P17182	Confidence level	99.00%
Precursor m/z	713.39419	Identity Threshold	46.96
Charge	3+	Mass error	0.00141



Info

Modified Sequence	AcD3-YNEFPEPIK<AcD3K*>LDK<AcD3K*>NDR<C13>-COOH	Ion score	50.24
Protein	P51150	Confidence level	99.00%
Precursor m/z	1010.0283	Identity Threshold	48.07
Charge	2+	Mass error	0.001826



Info

Modified Sequence AcD3-FVDSL NQK<AcD3K*>M<Mox*>EVQR<C13>-COOH

Ion score 49.41

Protein P16675

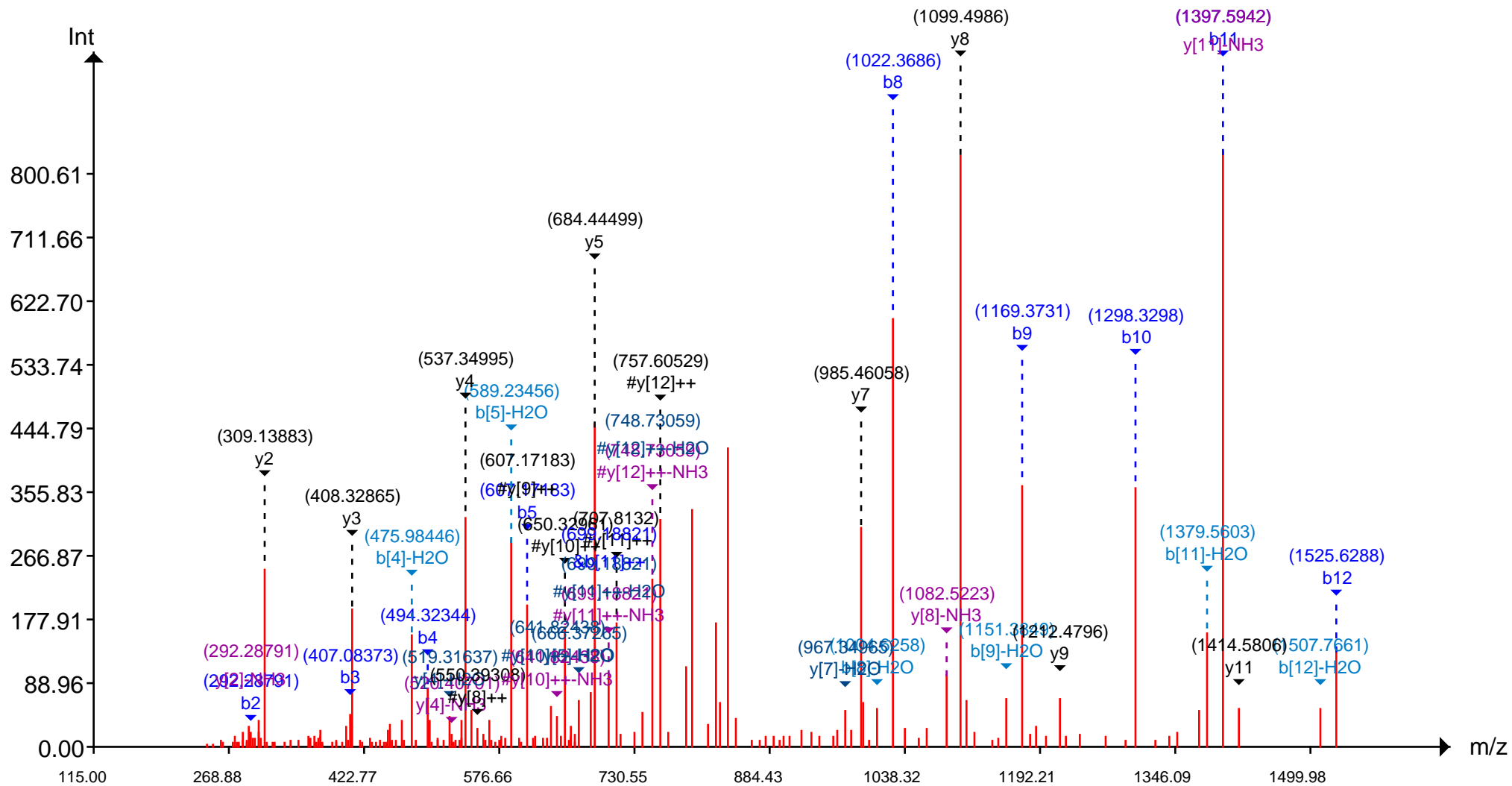
Confidence level 99.00%

Precursor m/z 853.44445

Identity Threshold 48.09

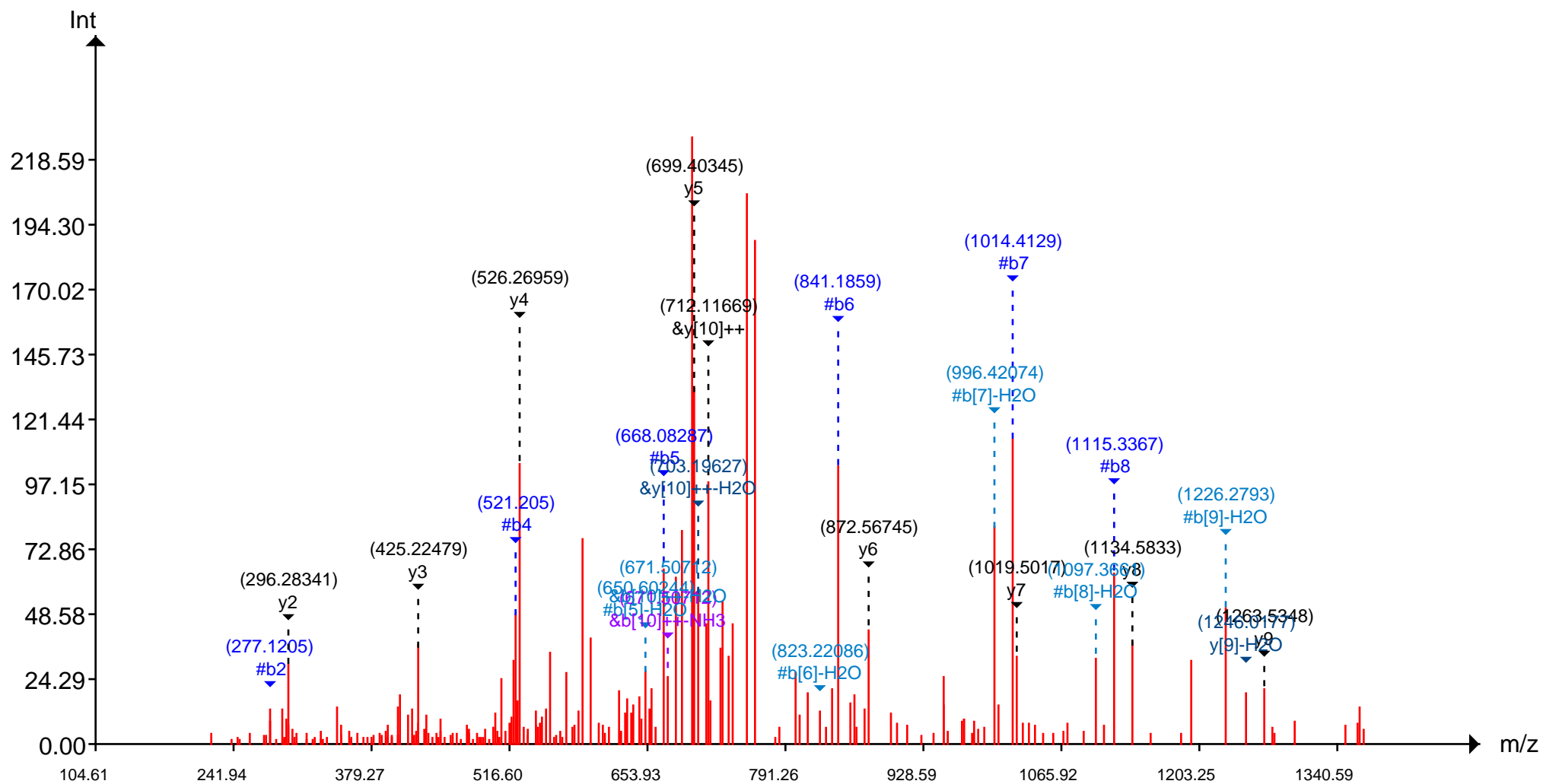
Charge 2+

Mass error 0.00243



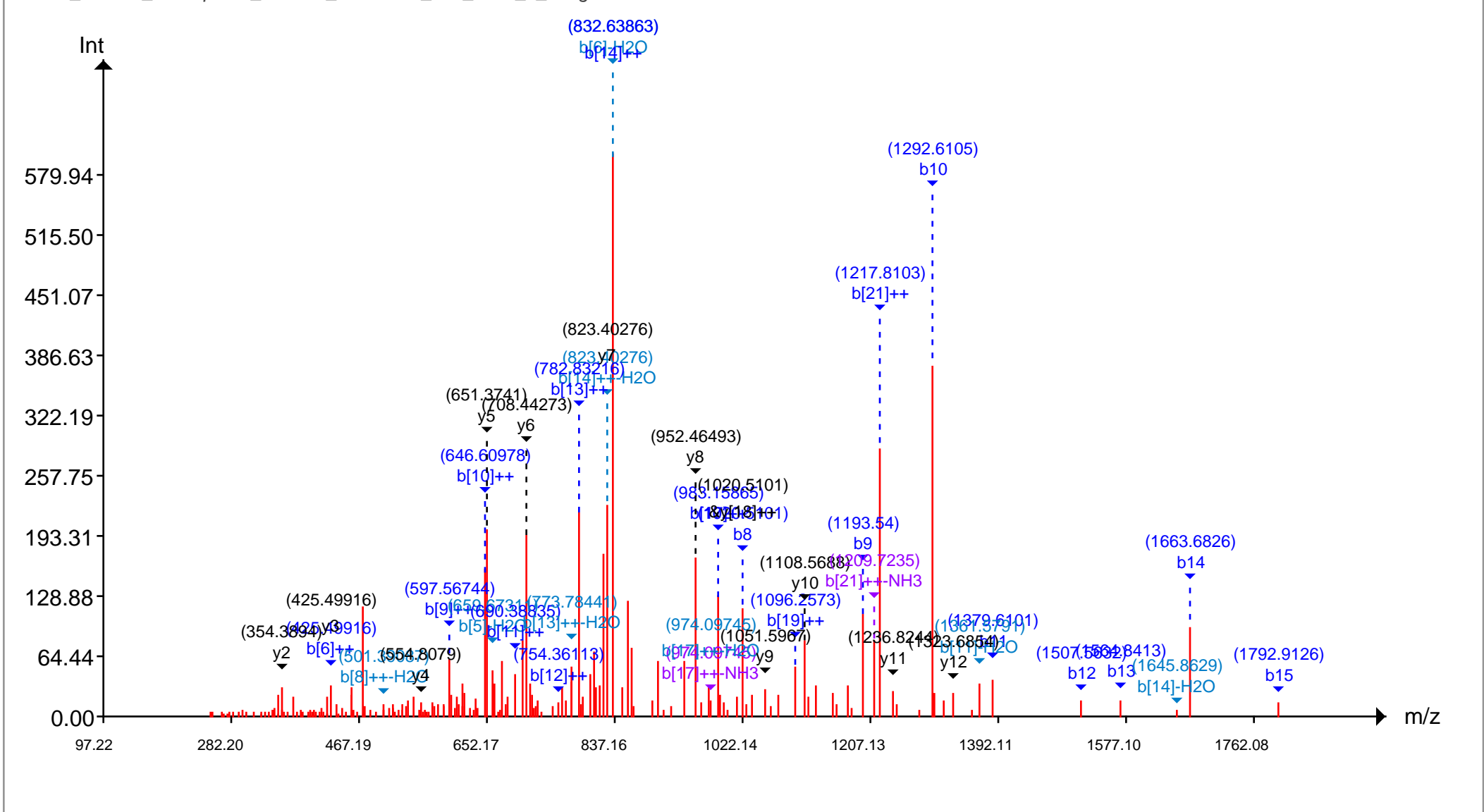
Info

Modified Sequence	AcD3-FVDSL NQK<AcD3K*>M<Mox*>EVQR<C13>-COOH	Ion score	60.64
Protein	P16675	Confidence level	99.00%
Precursor m/z	853.44429	Identity Threshold	48.06
Charge	2+	Mass error	0.00211



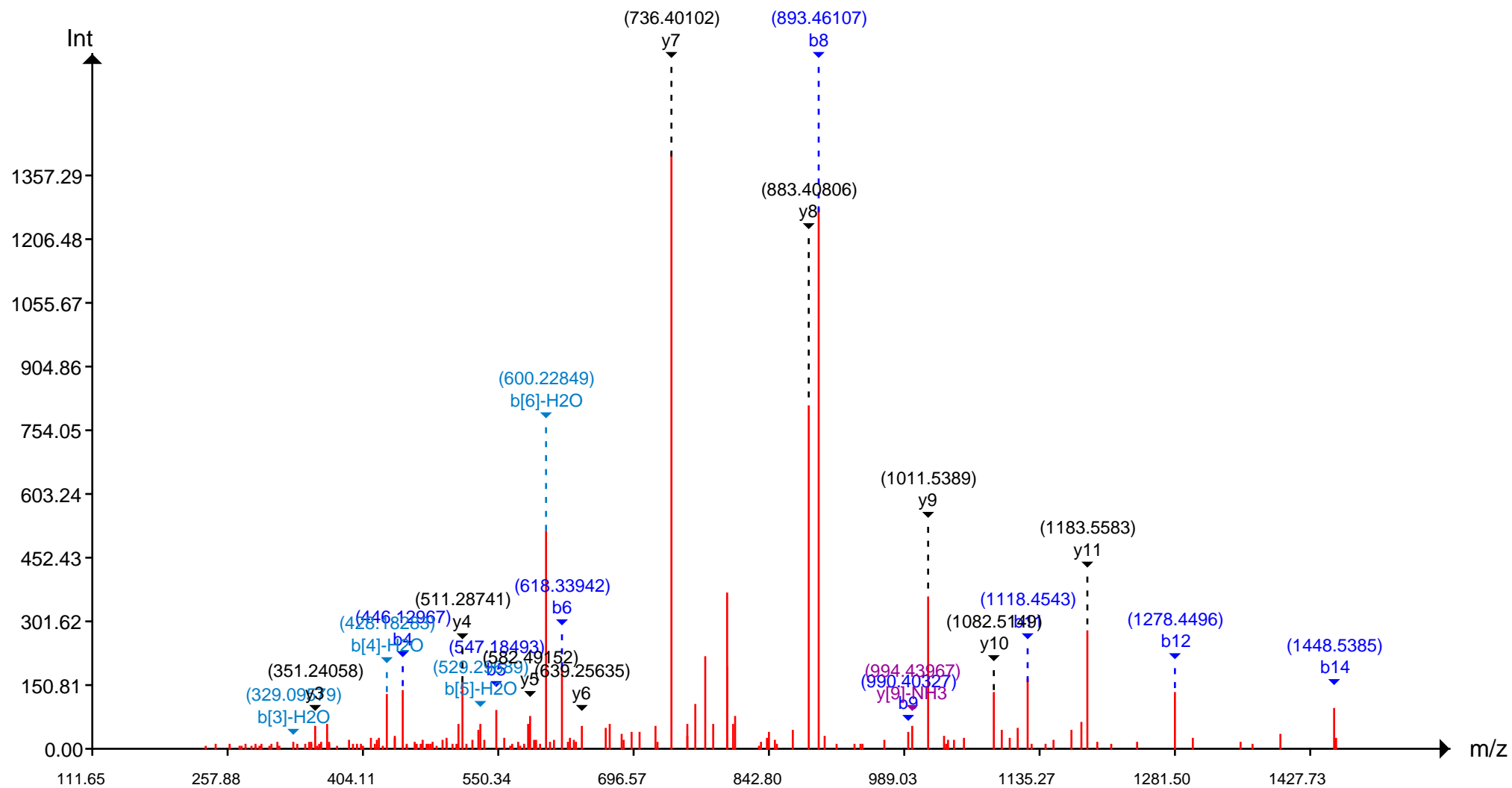
Info

Modified Sequence	AcD3-AC<Cmm*>EDFK<AcD3K*>K<AcD3K*>TEDR<C13>-COOH	Ion score	55.16
Protein	Q9CQE5	Confidence level	99.00%
Precursor m/z	770.37413	Identity Threshold	46.90
Charge	2+	Mass error	8.59E-4



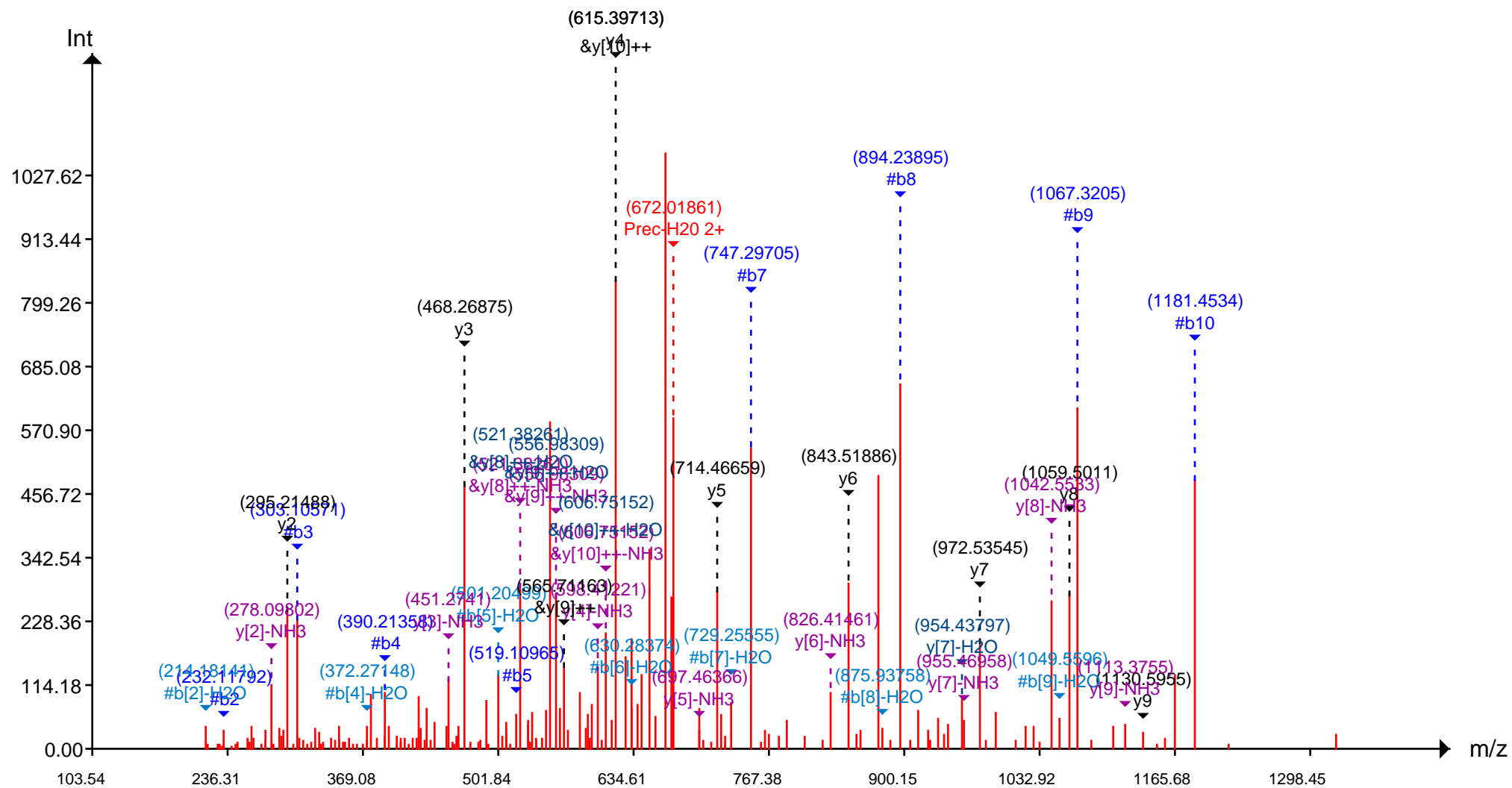
Info

Modified Sequence	AcD3-VEEK<AcD3K*>TK<AcD3K*>VAK<AcD3K*>VSQGVEDGPEAK<AcD3K*>R<C13>-COOH	Ion score	65.22
Protein	Q9R1T2	Confidence level	99.00%
Precursor	872.48787	Identity Threshold	45.56
Charge	3+	Mass error	0.00396



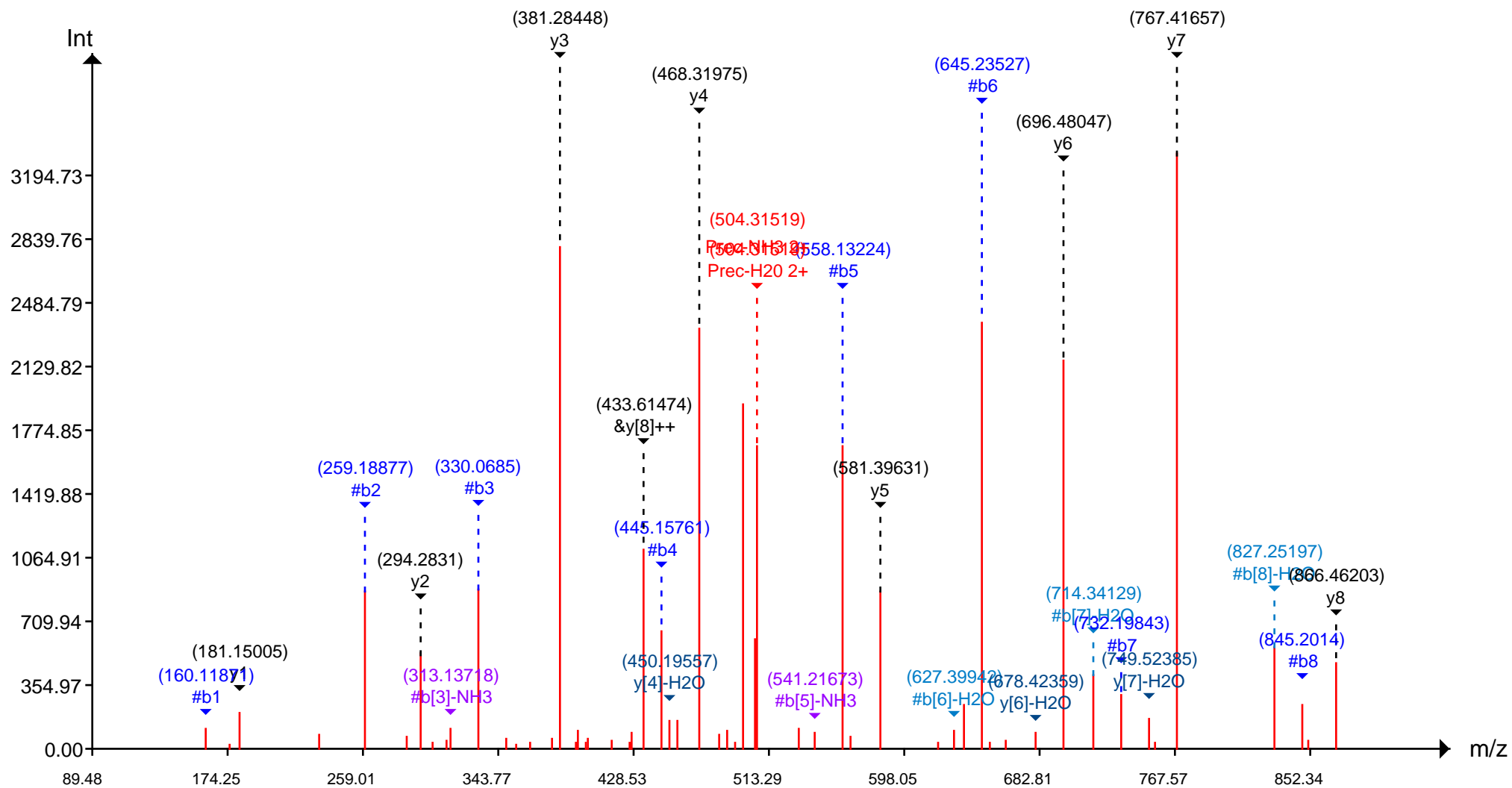
Info

Modified Sequence	AcD3-LSTVTAQFPGAC<Cmm*>GLR<C13>-COOH	Ion score	54.32
Protein	Q921F2	Confidence level	99.00%
Precursor m/z	814.93485	Identity Threshold	48.11
Charge	2+	Mass error	0.002441



Info

Modified Sequence	AcD3-SVASEEVM<Mox*>K<AcD3K*>NR<C13>-COOH	Ion score	59.95
Protein	Q9JIH2	Confidence level	99.00%
Precursor m/z	681.35046	Identity Threshold	47.99
Charge	2+	Mass error	-6.89E-4



Info

Modified Sequence AcD3-NVADLSSLR<C13>-COOH

Ion score 88.29

Protein P97372

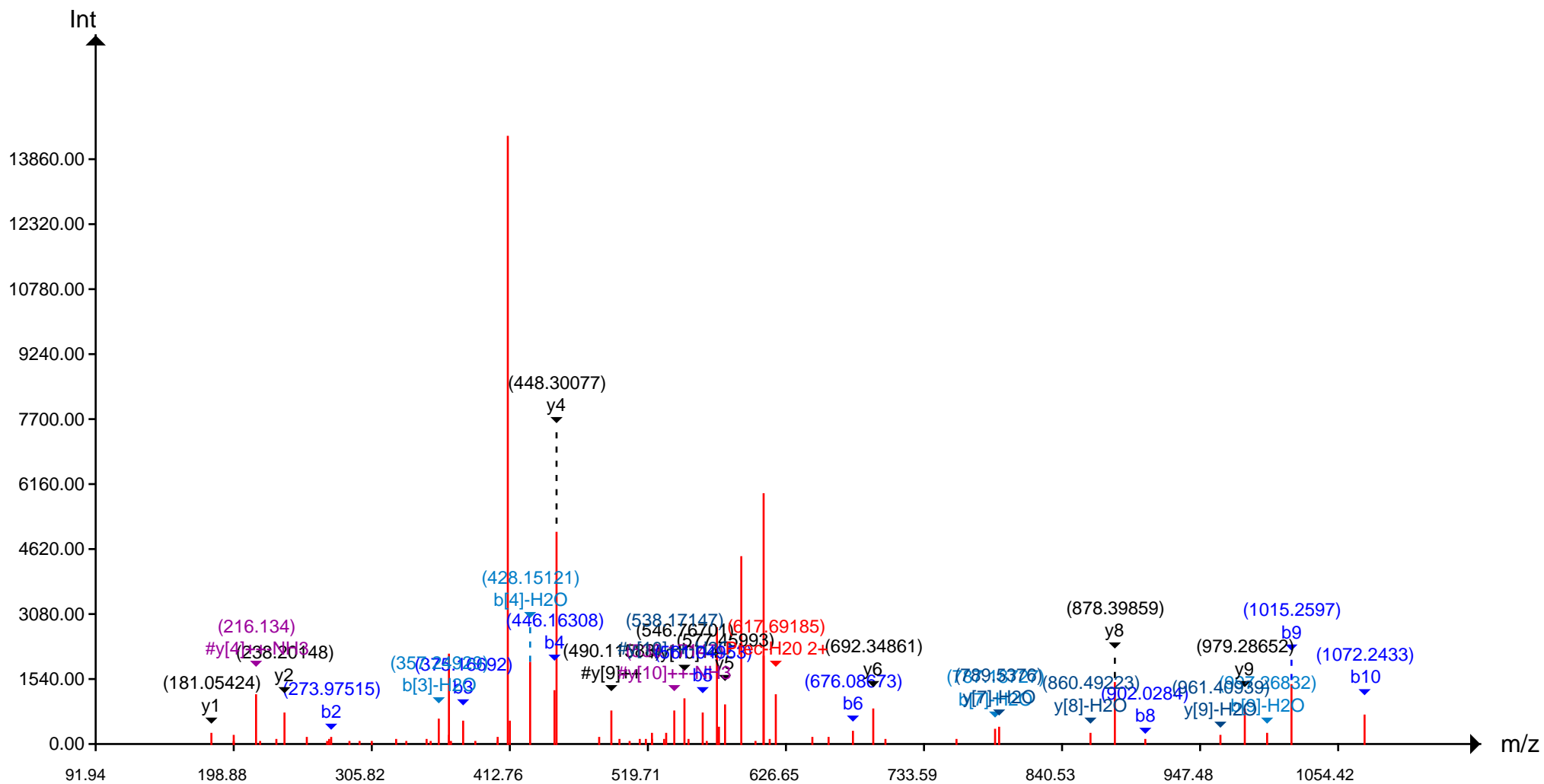
Confidence level 99.00%

Precursor m/z 513.29156

Identity Threshold 46.94

Charge 2+

Mass error -2.03E-4



Info

Modified Sequence AcD3-DITADDEPLGR<C13>-COOH

Ion score 51.58

Protein P17742

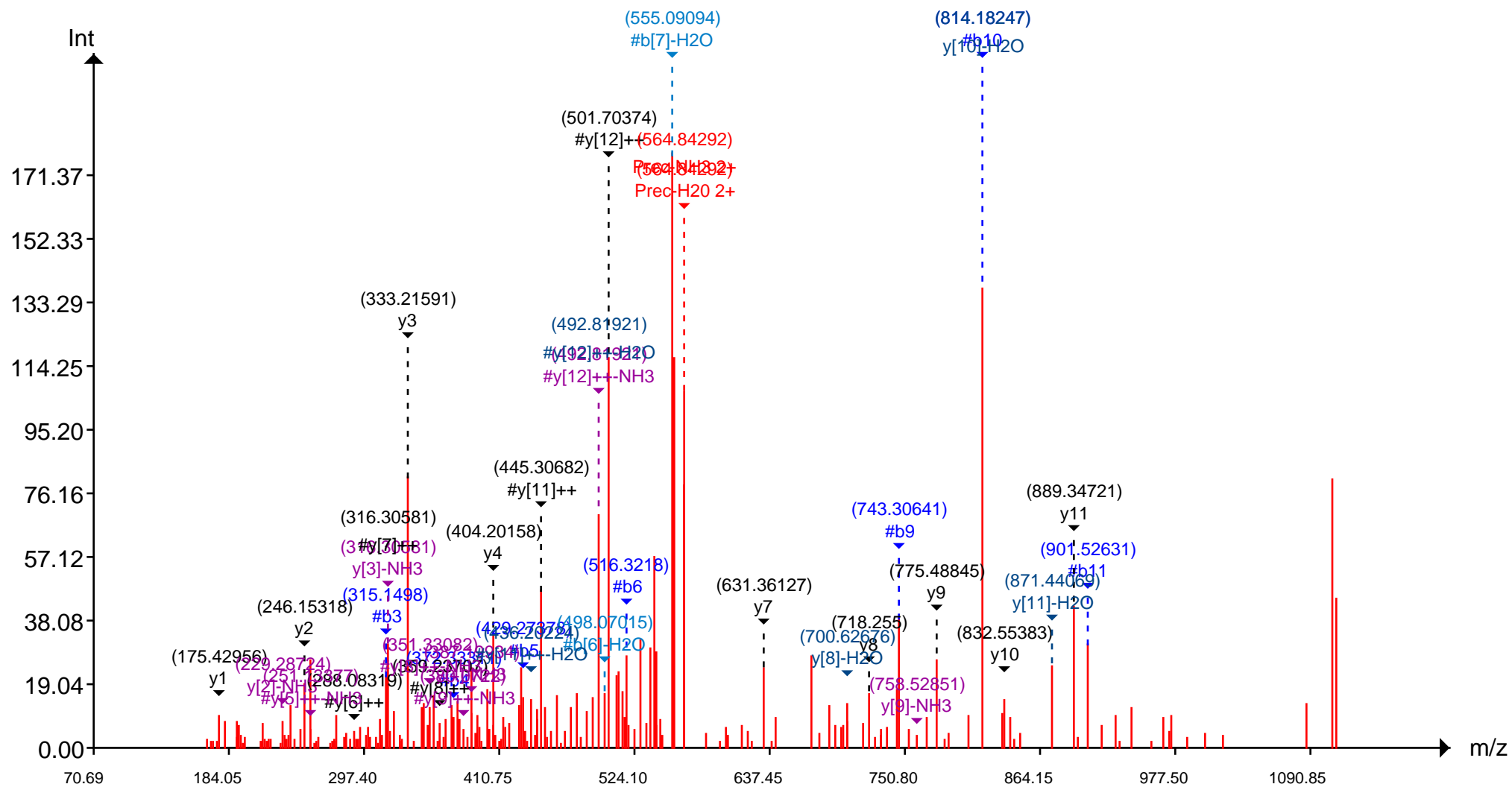
Confidence level 99.00%

Precursor m/z 626.81228

Identity Threshold 47.40

Charge 2+

Mass error -0.001763



Info

Modified Sequence AcD3-VIGGGSGGLASAR-COOH

Ion score 58.69

Protein P47791

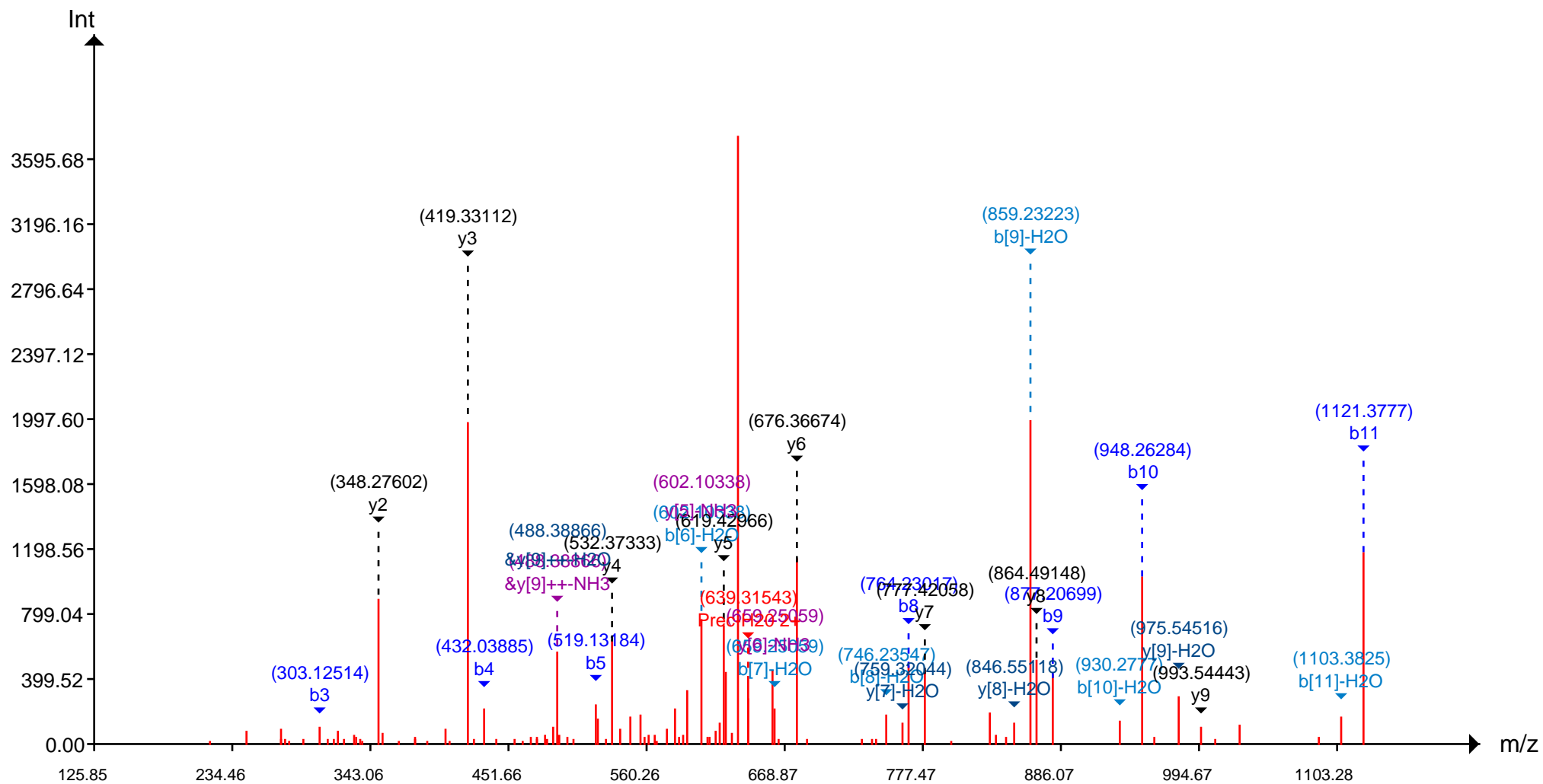
Confidence level 99.00%

Precursor m/z 573.81772

Identity Threshold 47.95

Charge 2+

Mass error -0.002327



Info

Modified Sequence AcD3-AADESTGSIK<AcD3K*>R-COOH

Ion score 62.69

Protein P05064

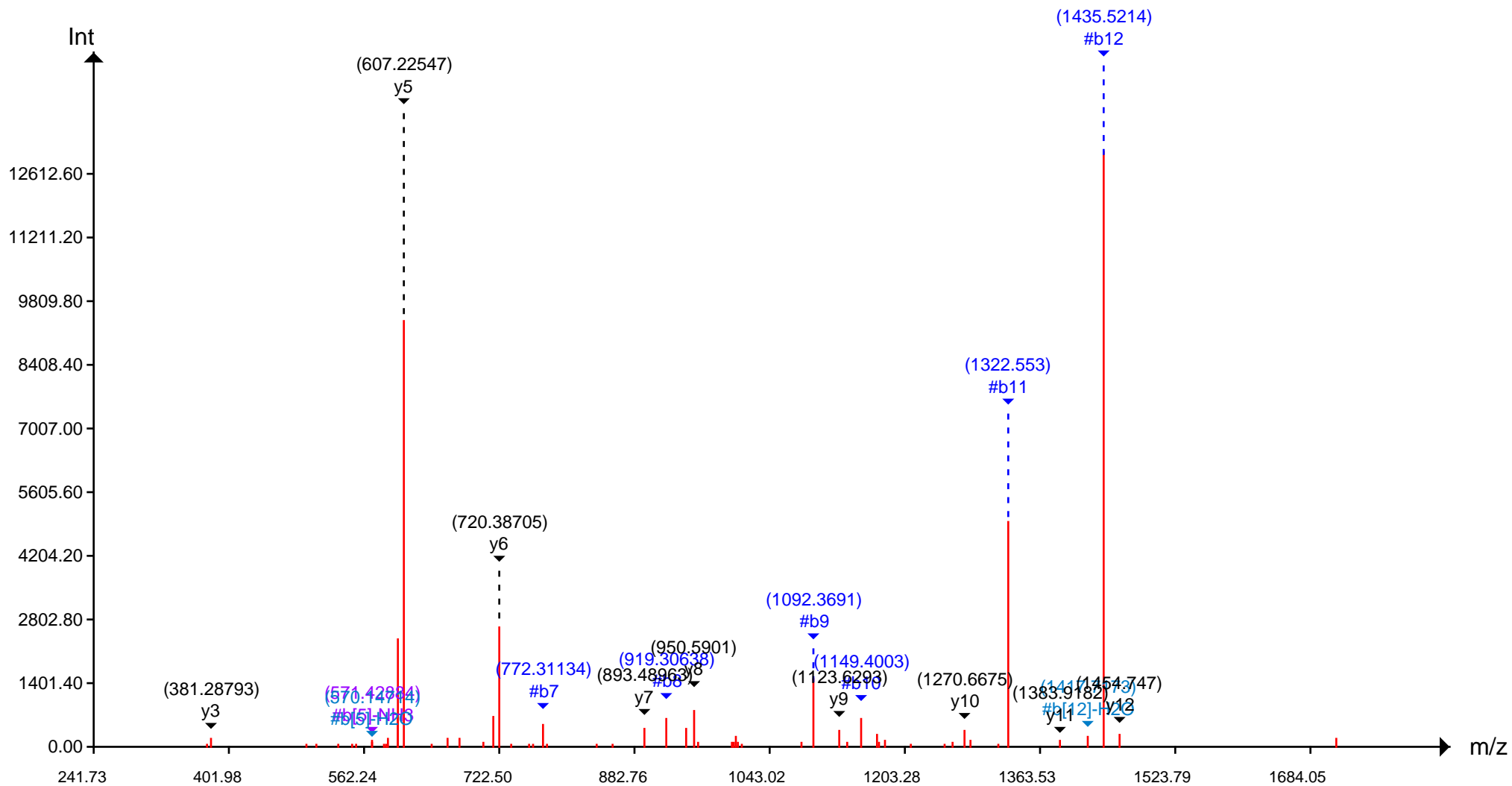
Confidence level 99.00%

Precursor m/z 648.34008

Identity Threshold 47.94

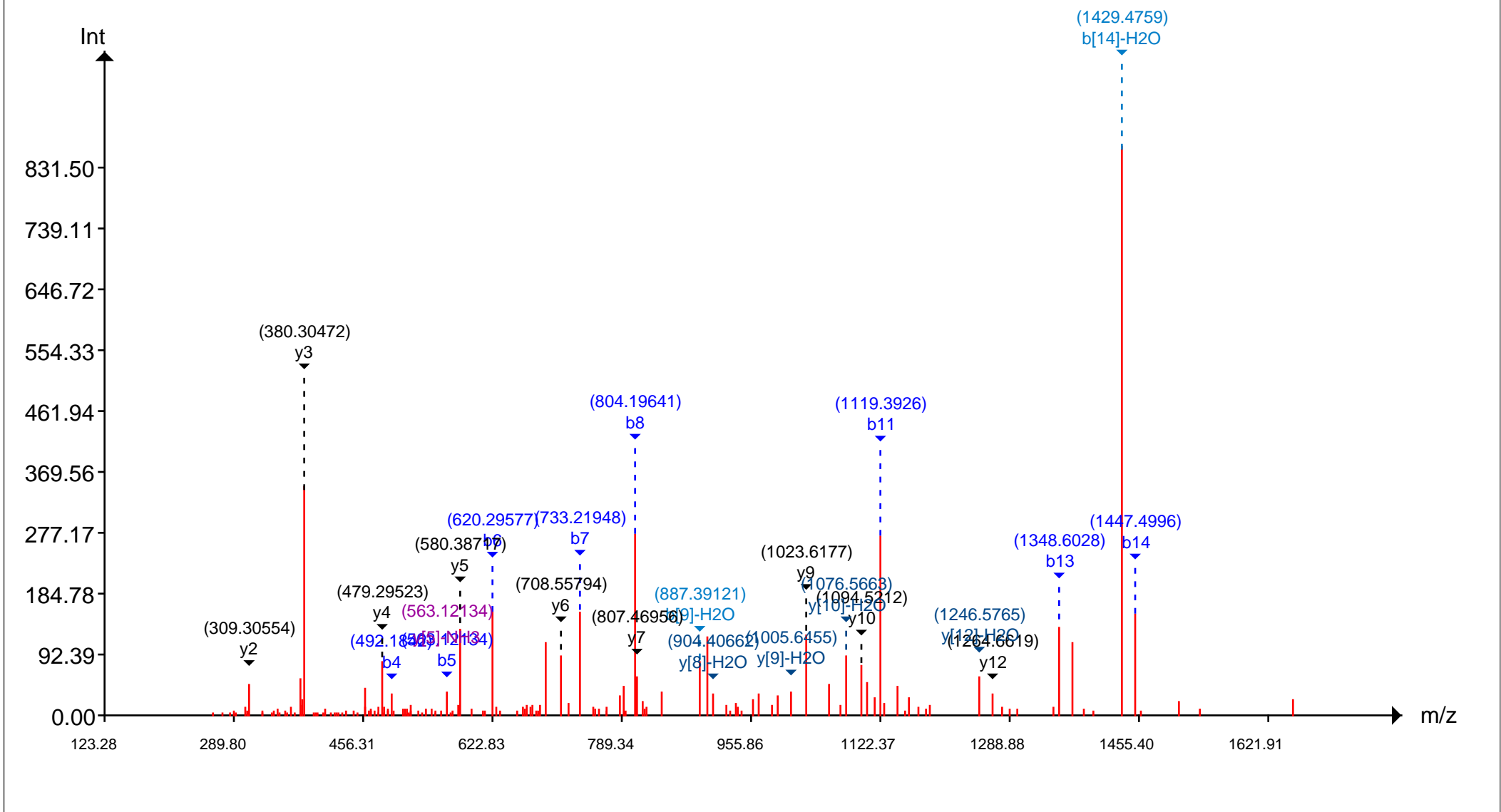
Charge 2+

Mass error 0.002064



Info

Modified Sequence	AcD3-LAQTEALM<Mox*>K<AcD3K*>GK<AcD3K*>LPPEAR<C13>-COOH	Ion score	47.39
Protein	P06745	Confidence level	99.00%
Precursor m/z	1021.567	Identity Threshold	46.70
Charge	2+	Mass error	0.002367



Info

Modified Sequence AcD3-YIGLAGLATDVQTVAR<C13>-COOH

Ion score 67.26

Protein Q9R1P1

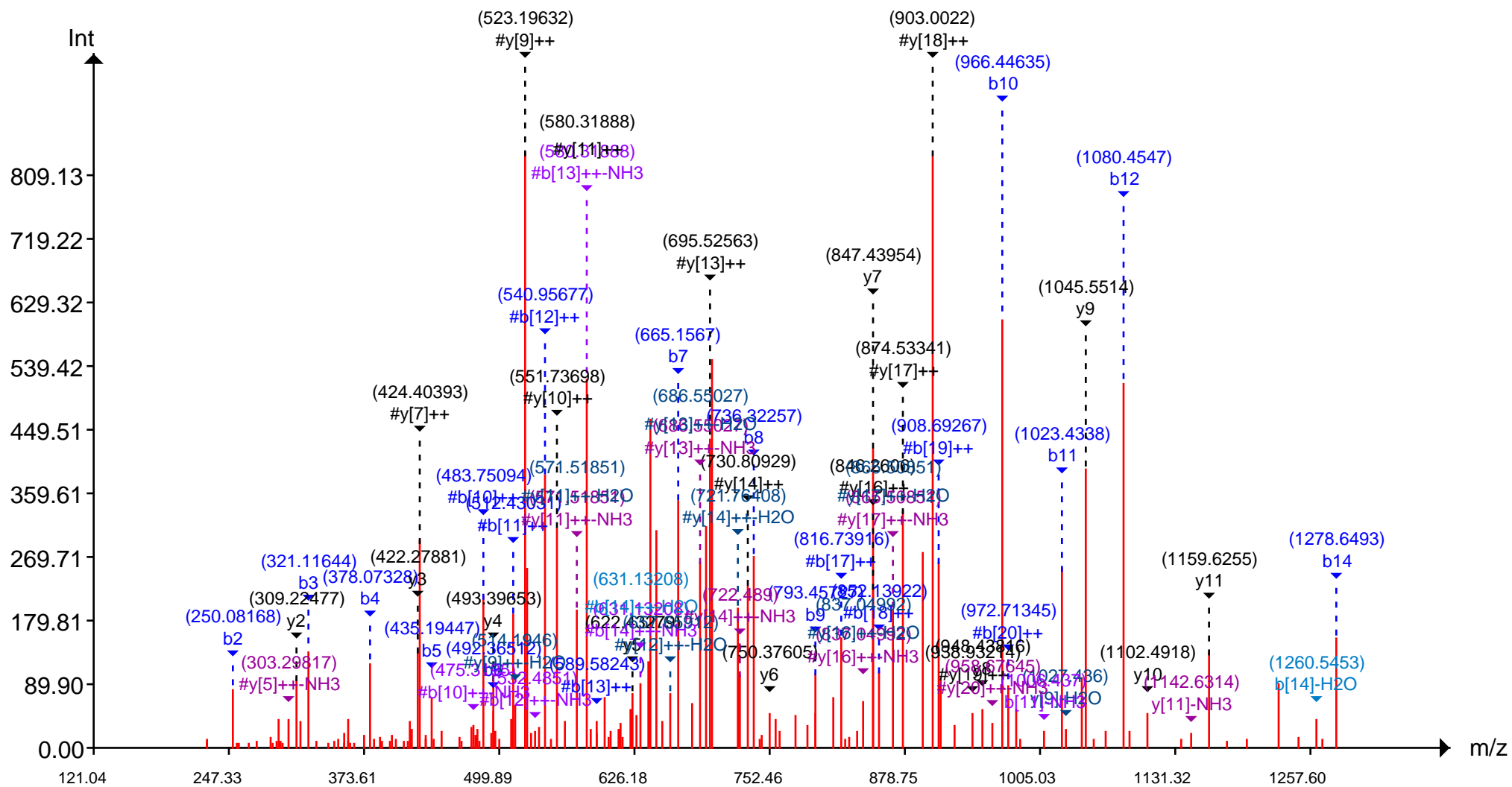
Confidence level 99.00%

Precursor m/z 914.01249

Identity Threshold 46.65

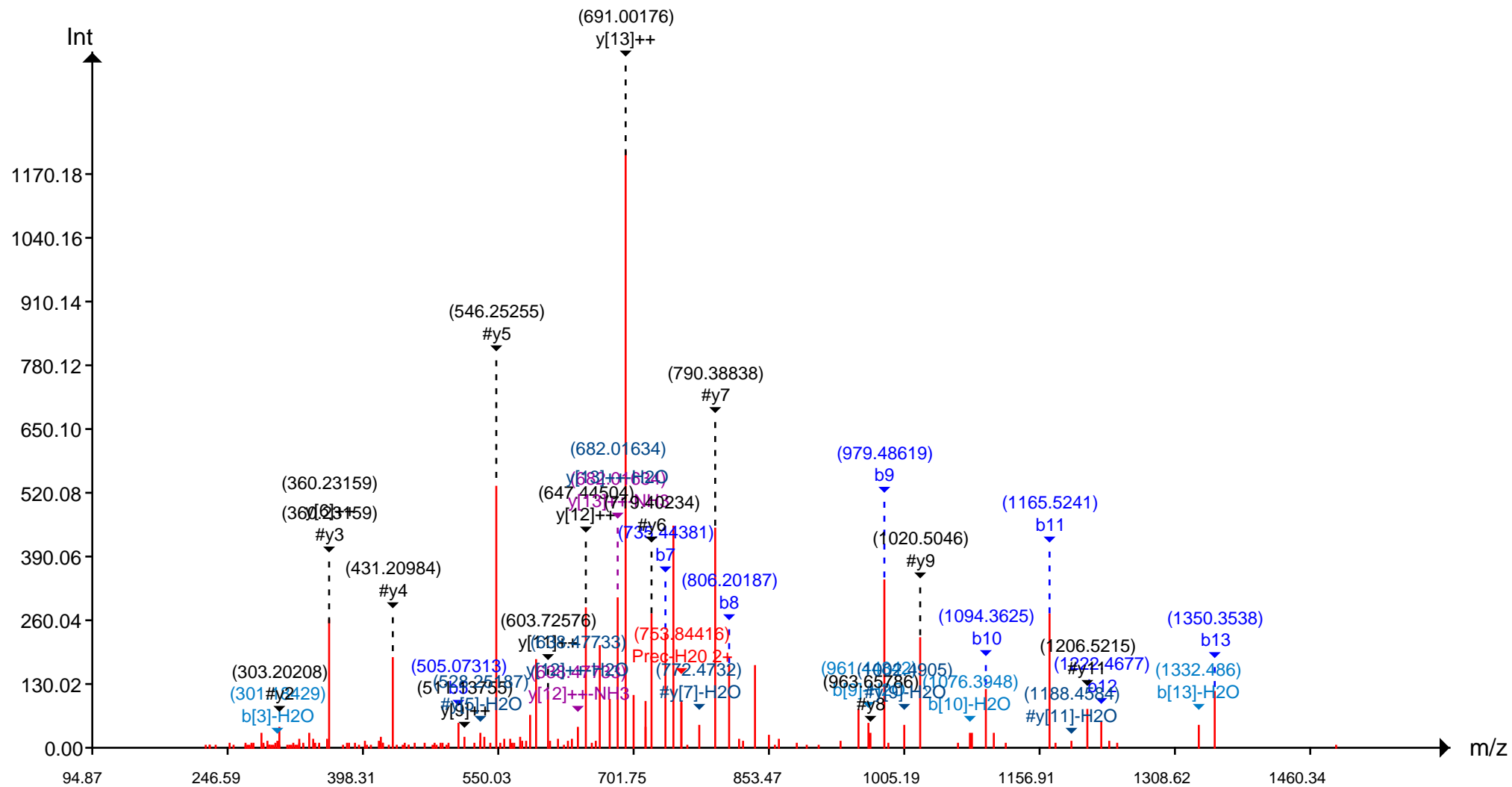
Charge 2+

Mass error 0.00315



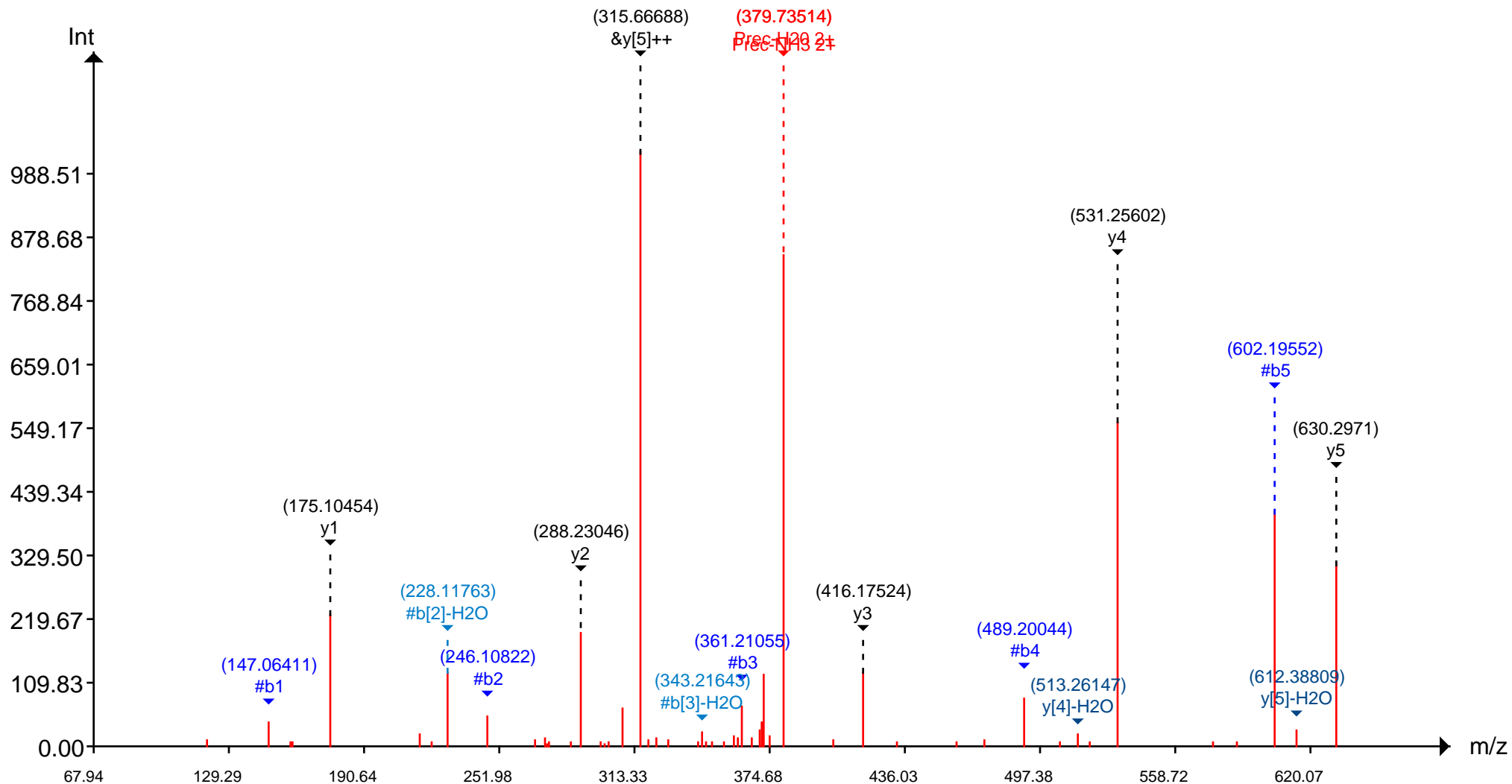
Info

Modified Sequence	AcD3-FGAGGGK<AcD3K*>AGK<AcD3K*>GGPTPQEAIQR<C13>-COOH	Ion score	91.0
Protein	Q9D8B3	Confidence level	99.00%
Precursor m/z	709.05333	Identity Threshold	47.36
Charge	3+	Mass error	0.001199



Info

Modified Sequence	AcD3-VSSGEGK<AcD3K*>AK<AcD3K*>DAGQR-COOH	Ion score	56.44
Protein	P50580	Confidence level	99.00%
Precursor m/z	762.90241	Identity Threshold	48.06
Charge	2+	Mass error	0.001282



Info

Modified Sequence AcD3-TVDQIR-COOH

Protein P58252

Precursor m/z 388.72103

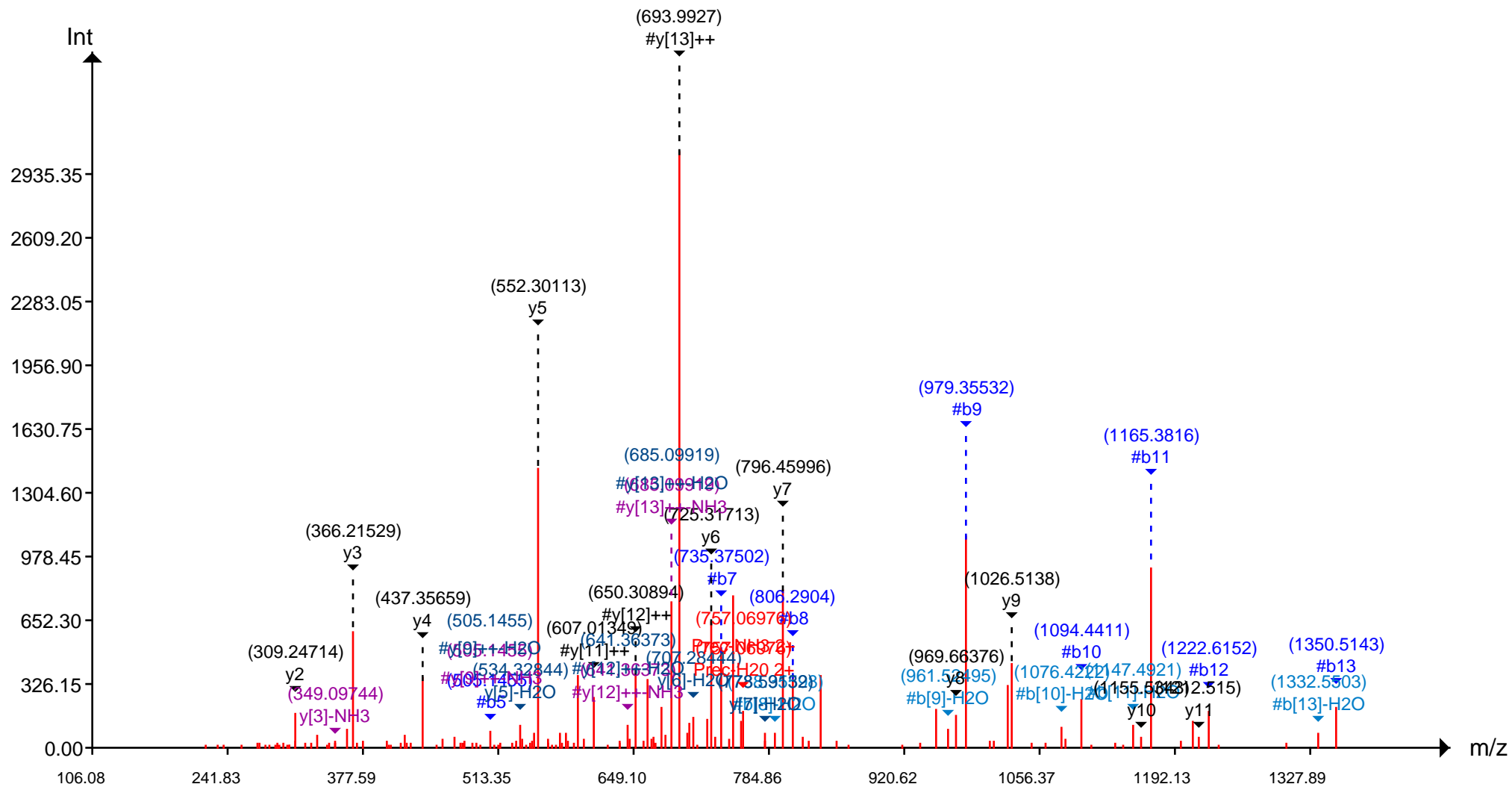
Charge 2+

Ion score 49.04

Confidence level 99.00%

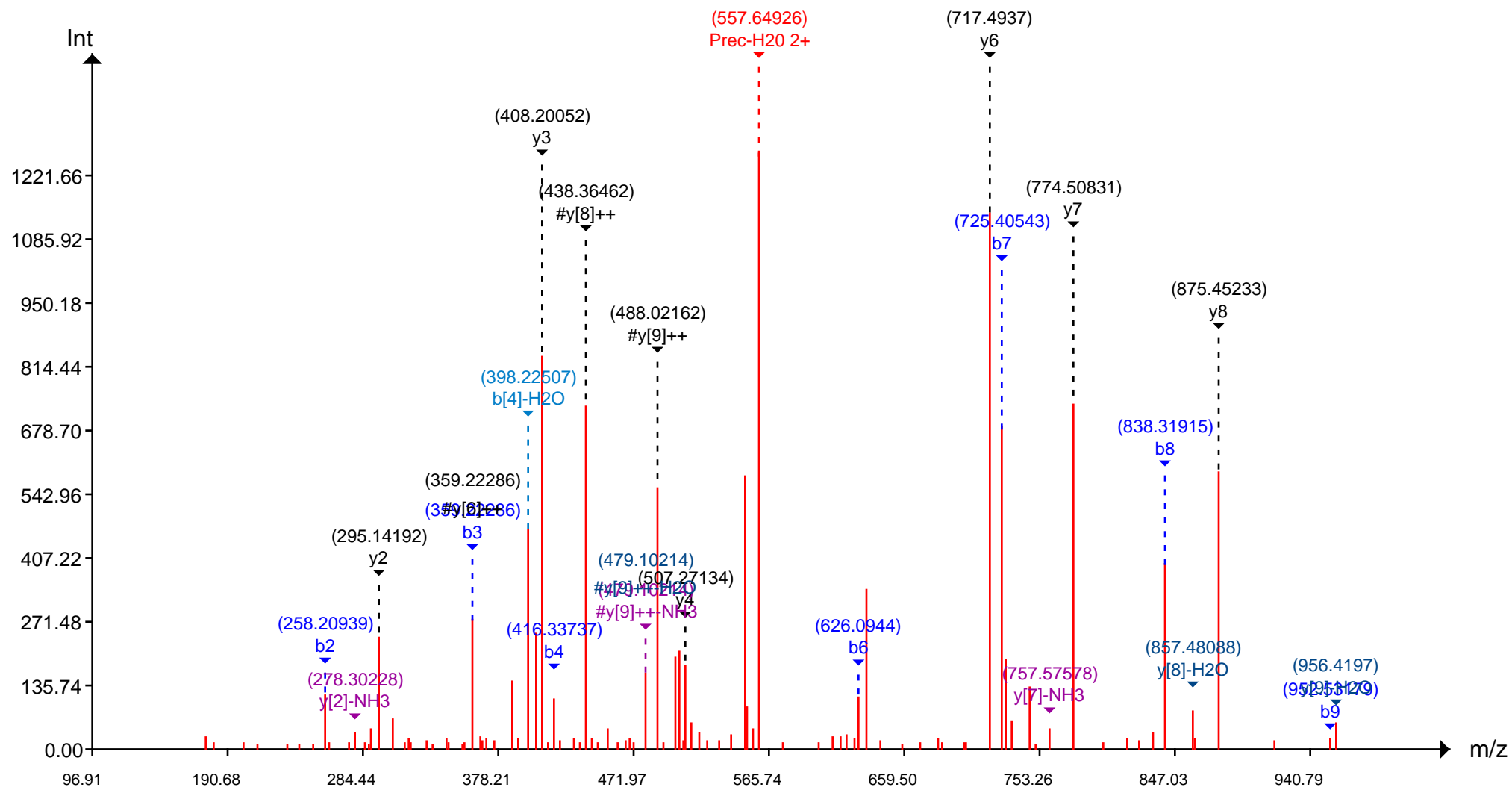
Identity Threshold 47.88

Mass error 7.65E-4



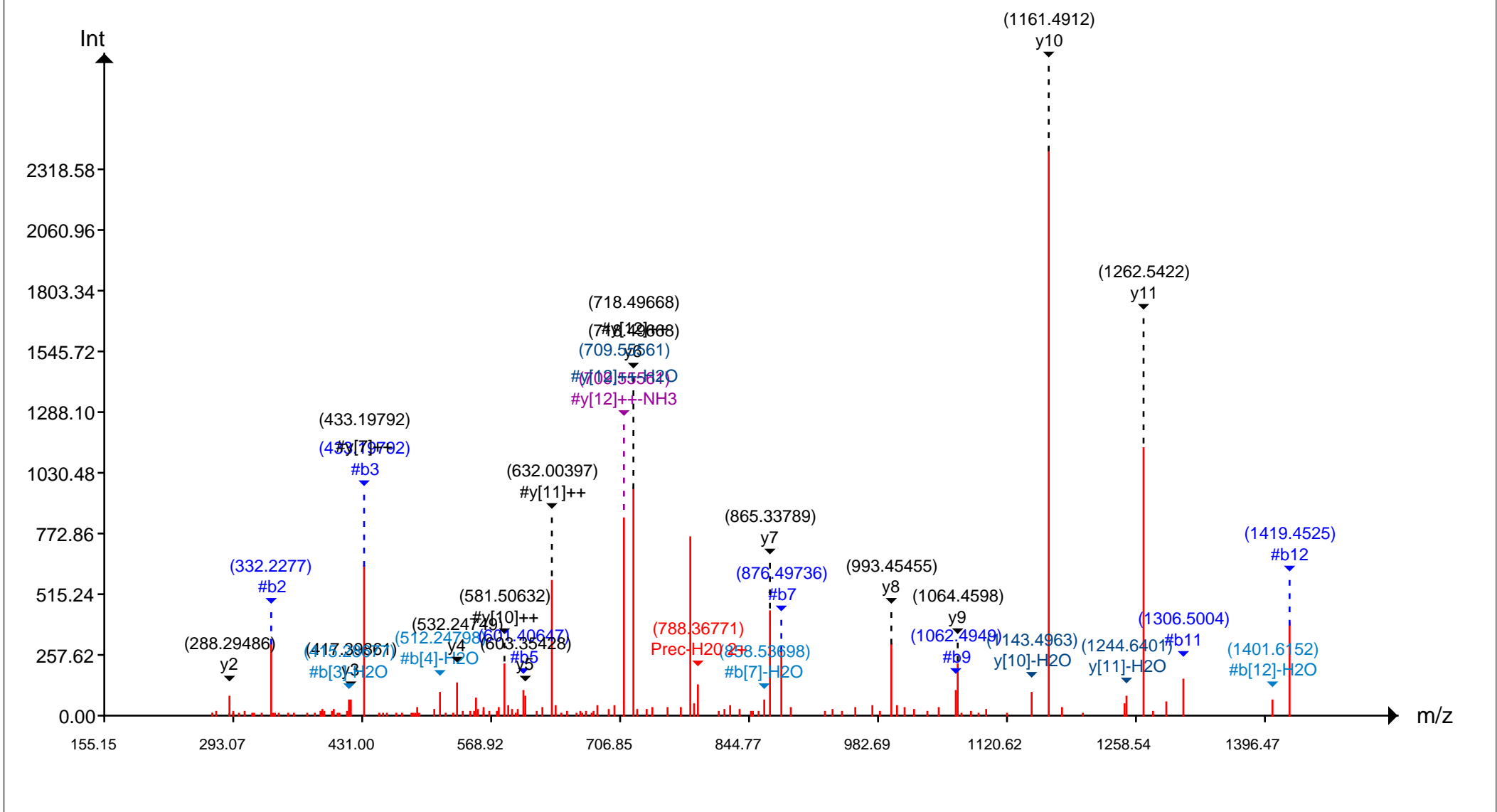
Info

Modified Sequence	AcD3-VSSGEGK<AcD3K*>AK<AcD3K*>DAGQR<C13>-COOH	Ion score	61.2
Protein	P50580	Confidence level	99.00%
Precursor m/z	765.91251	Identity Threshold	48.06
Charge	2+	Mass error	0.00134



Info

Modified Sequence	AcD3-LVTGPLVINR<C13>-COOH	Ion score	43.29
Protein	P47911	Confidence level	99.00%
Precursor m/z	566.86492	Identity Threshold	40.27
Charge	2+	Mass error	2.31E-4



Info

Modified Sequence AcD3-LK<AcD3K*>TPAQFDADELK-COOH

Ion score 87.7

Protein P10107

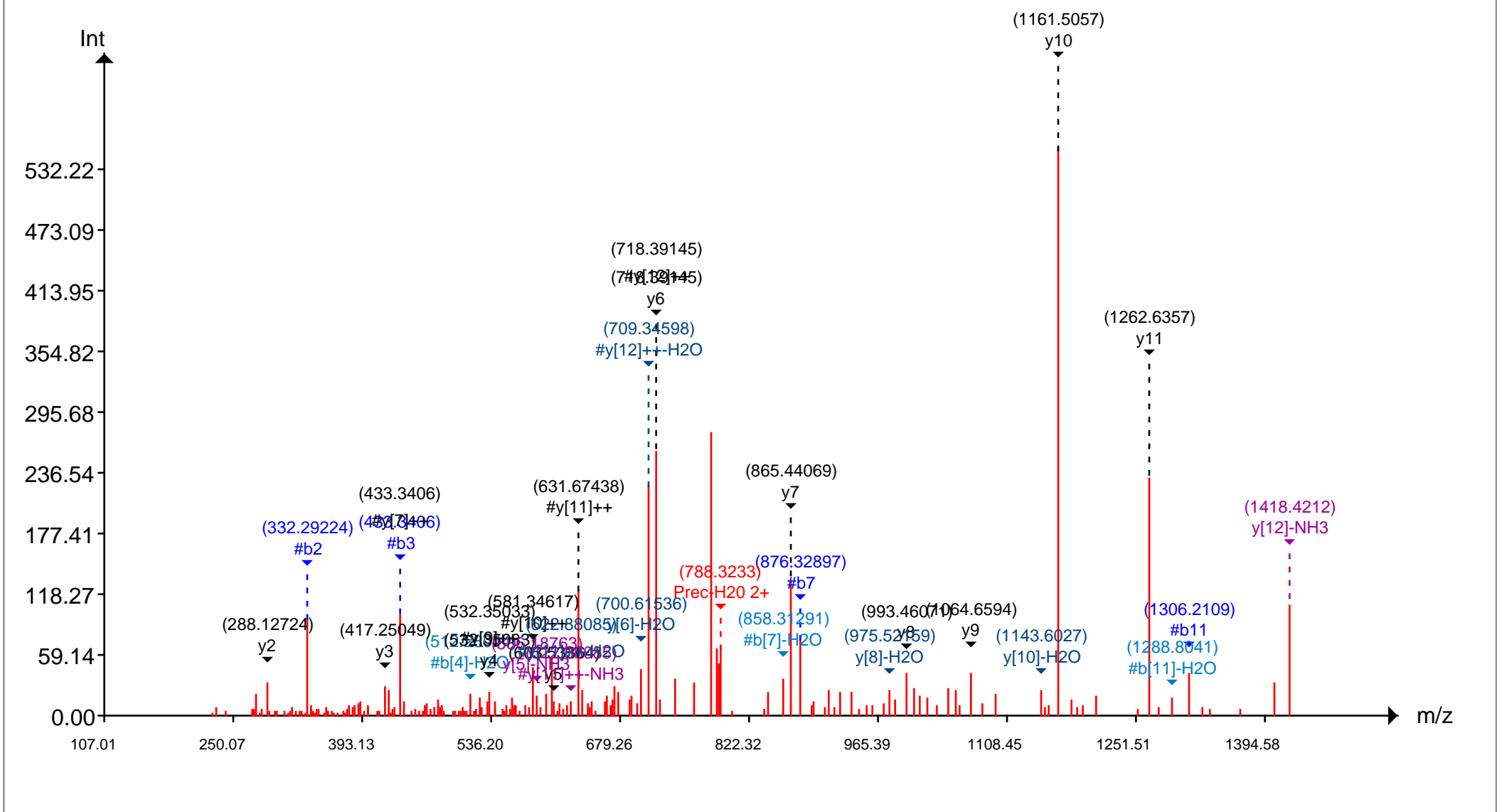
Confidence level 99.00%

Precursor m/z 797.4239

Identity Threshold 48.09

Charge 2+

Mass error 0.001568



Info

Modified Sequence AcD3-LK<AcD3K*>TPAQFDADELK-COOH

Ion score 63.68

Protein P10107

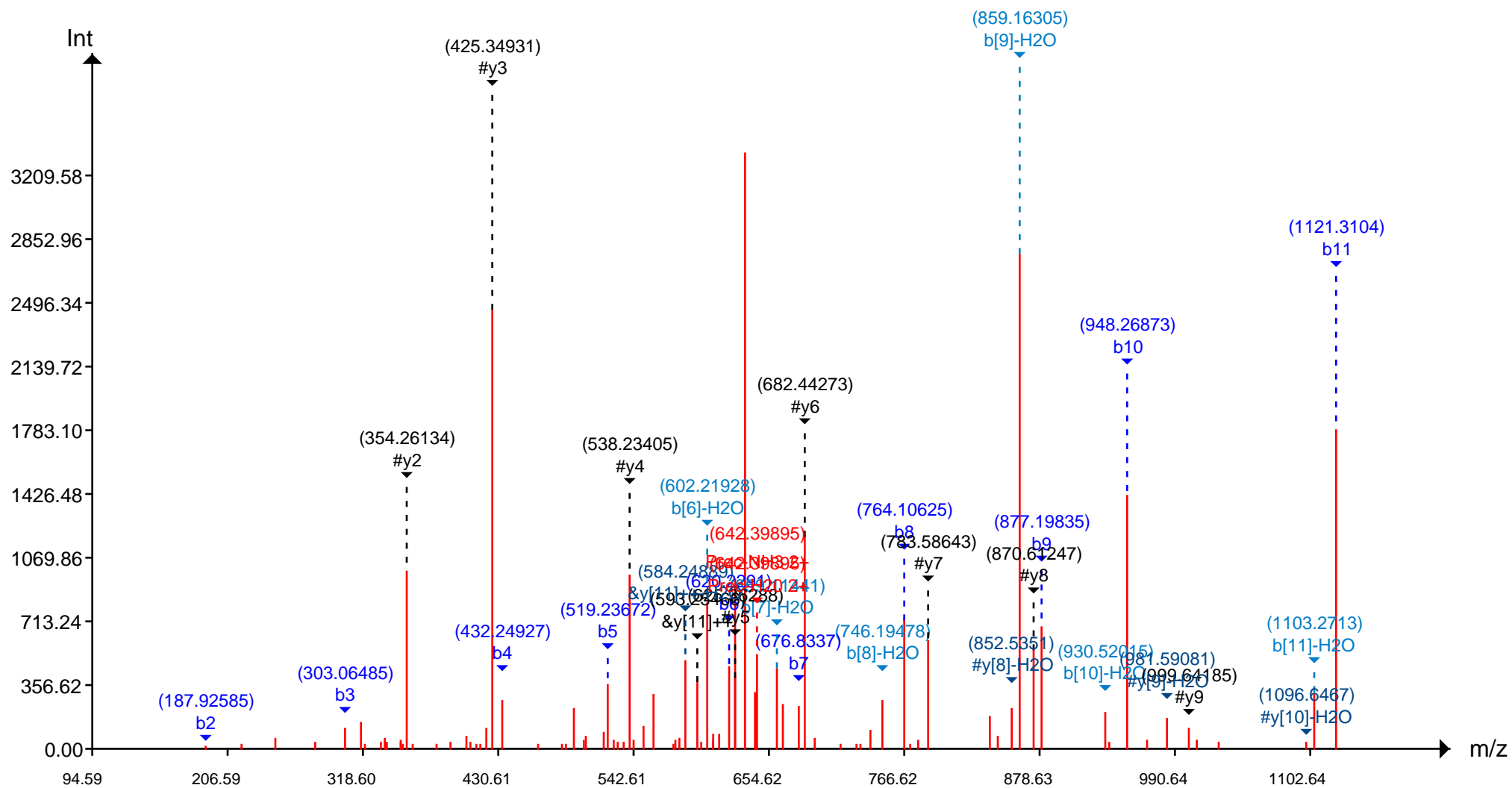
Confidence level 99.00%

Precursor m/z 797.42377

Identity Threshold 48.11

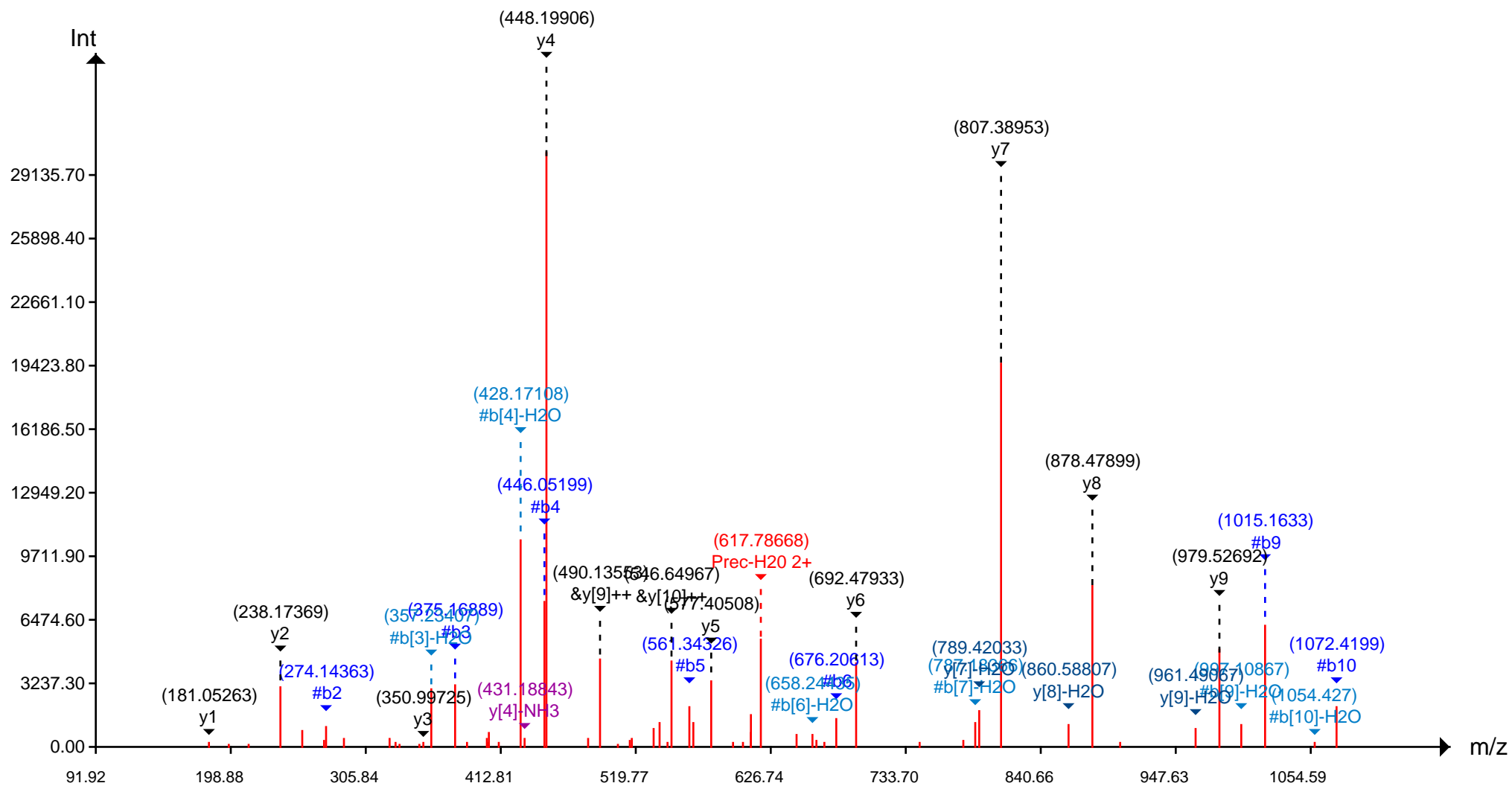
Charge 2+

Mass error 0.001308



Info

Modified Sequence	AcD3-AADESTGSIK<AcD3K*>R<C13>-COOH	Ion score	59.55
Protein	P05064	Confidence level	99.00%
Precursor m/z	651.35002	Identity Threshold	47.89
Charge	2+	Mass error	0.001803



Info

Modified Sequence AcD3-DITADDEPLGR<C13>-COOH

Ion score 69.74

Protein P17742

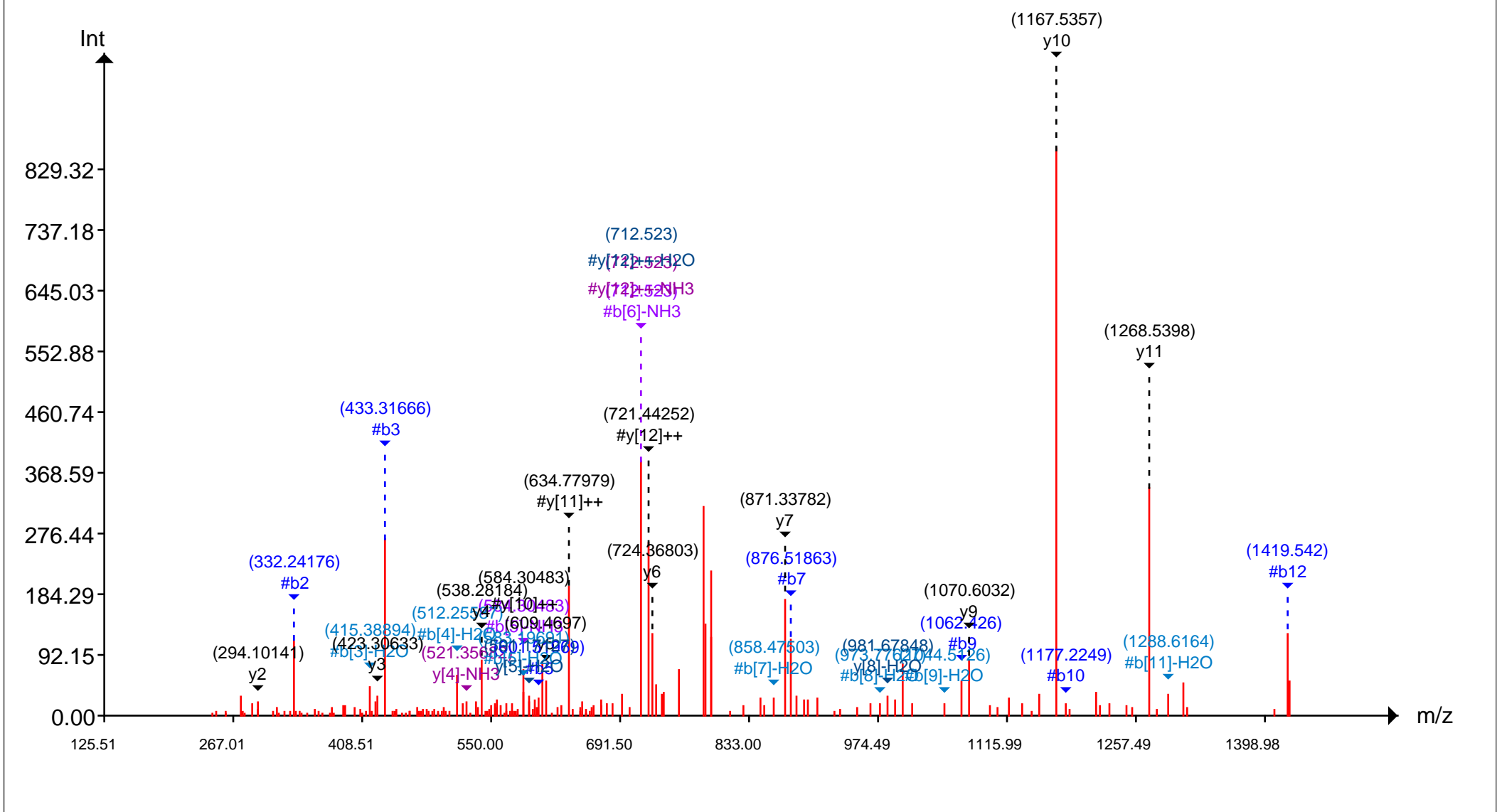
Confidence level 99.00%

Precursor m/z 626.8136

Identity Threshold 47.49

Charge 2+

Mass error 8.77E-4



Info

Modified Sequence AcD3-LK<AcD3K*>TPAQFDADELRL<C13>-COOH

Ion score 52.31

Protein P10107

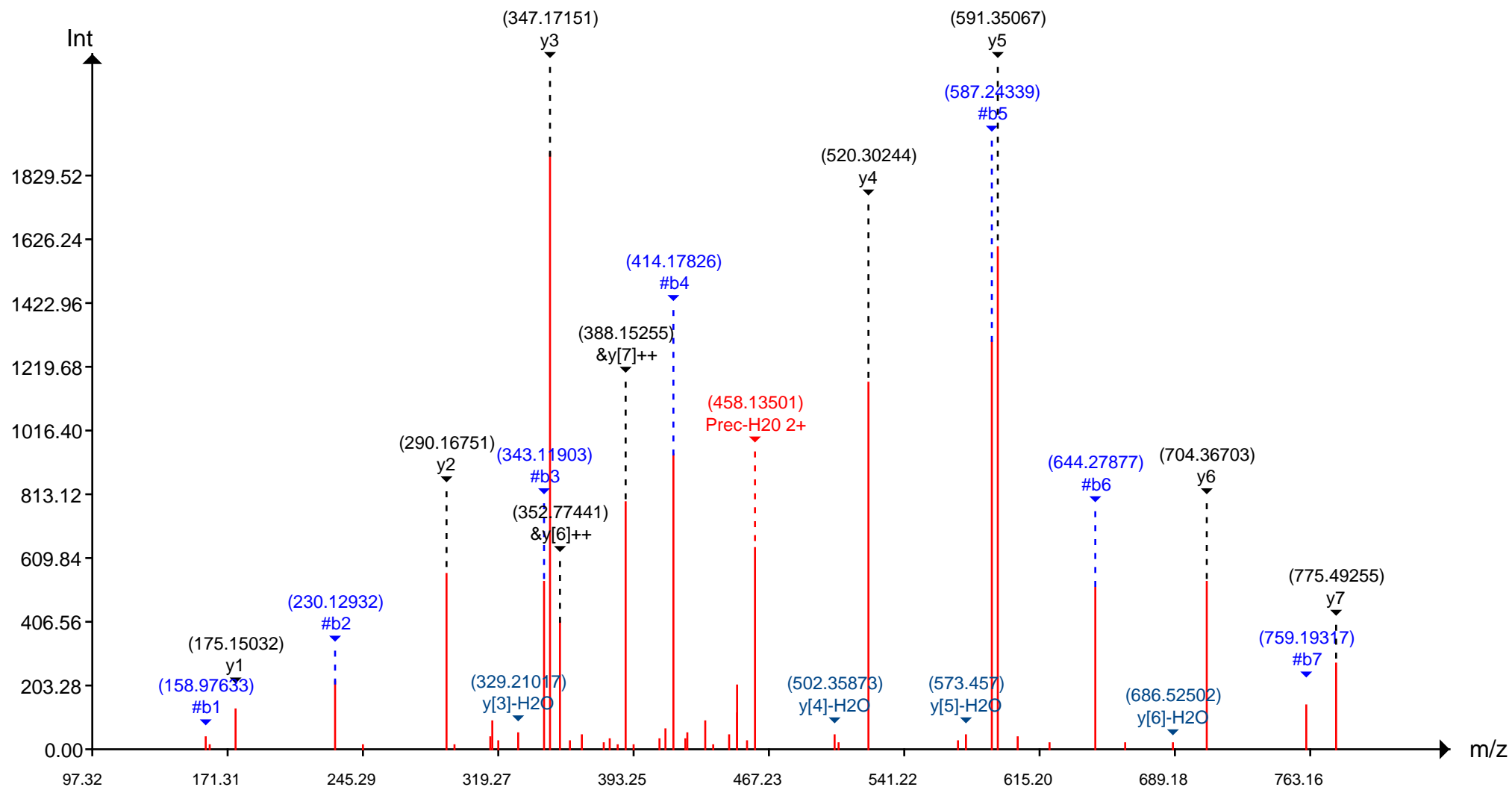
Confidence level 99.00%

Precursor m/z 800.43418

Identity Threshold 47.86

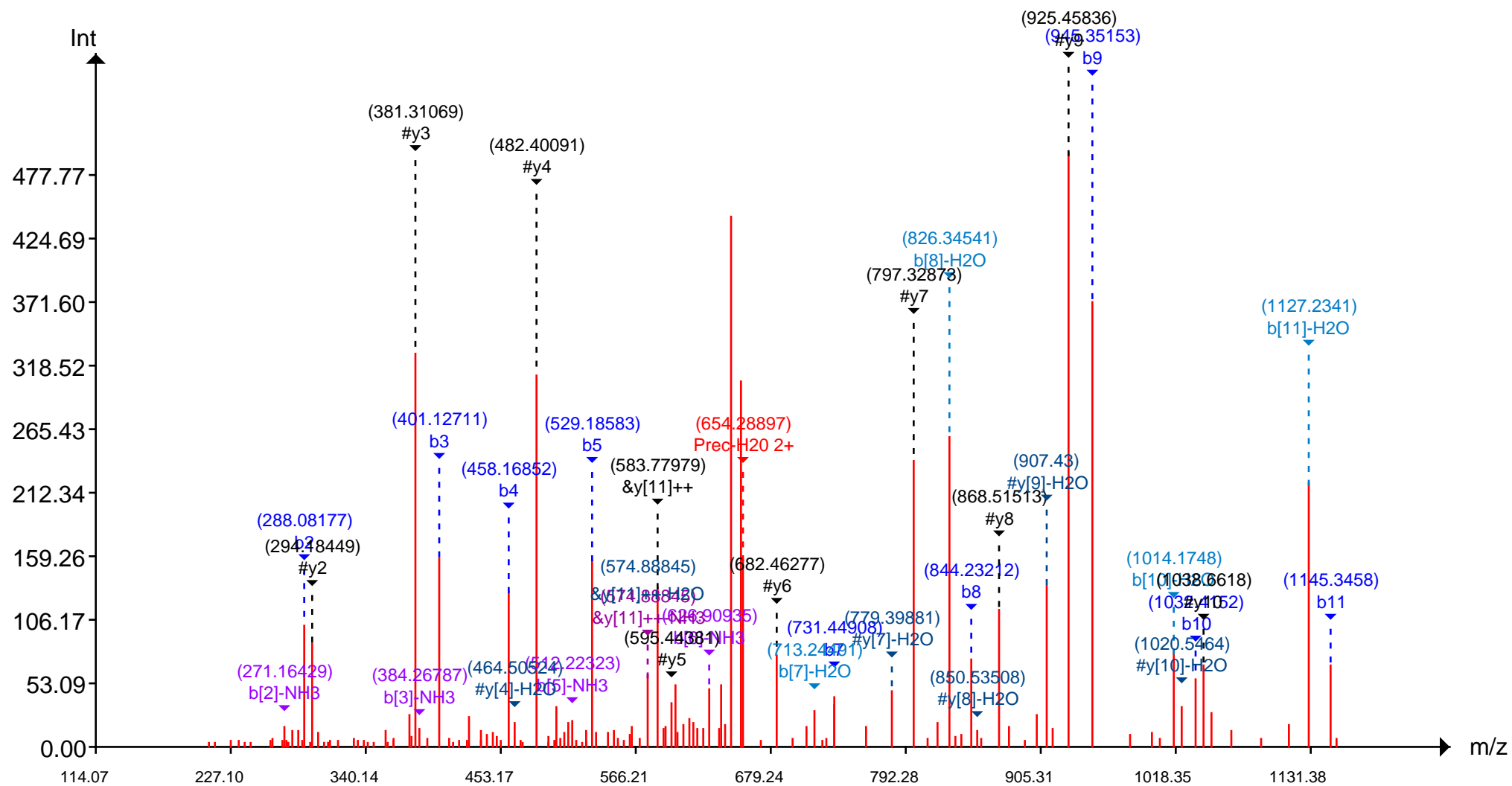
Charge 2+

Mass error 0.001986



Info

Modified Sequence	AcD3-LALAK<AcD3K*>GDR-COOH	Ion score	76.28
Protein	P10107	Confidence level	99.00%
Precursor m/z	467.28535	Identity Threshold	44.85
Charge	2+	Mass error	-2.0E-5



Info

Modified Sequence AcD3-NQLGADSLTSLR<C13>-COOH

Ion score 71.26

Protein Q64152

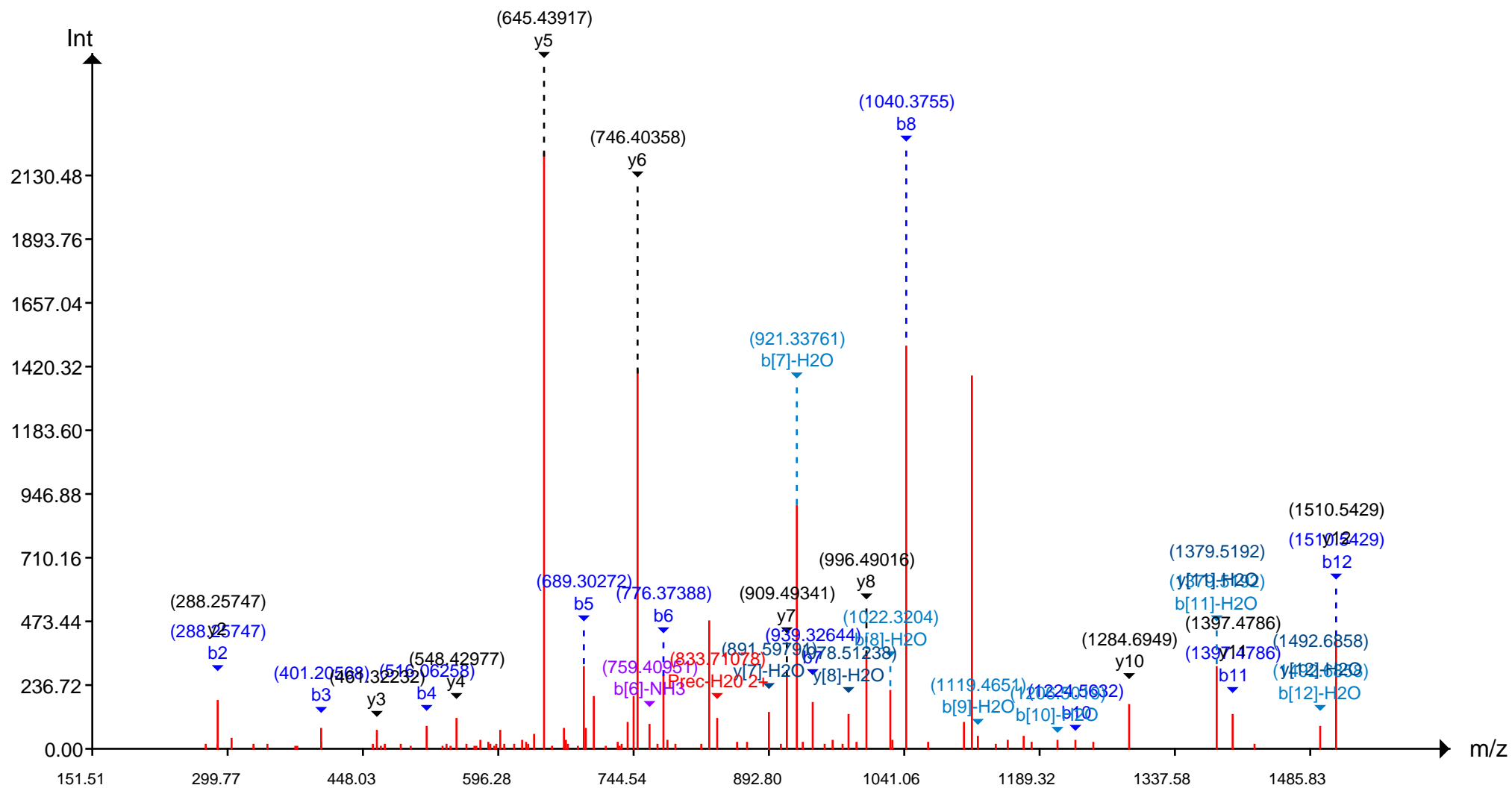
Confidence level 99.00%

Precursor m/z 663.36355

Identity Threshold 47.70

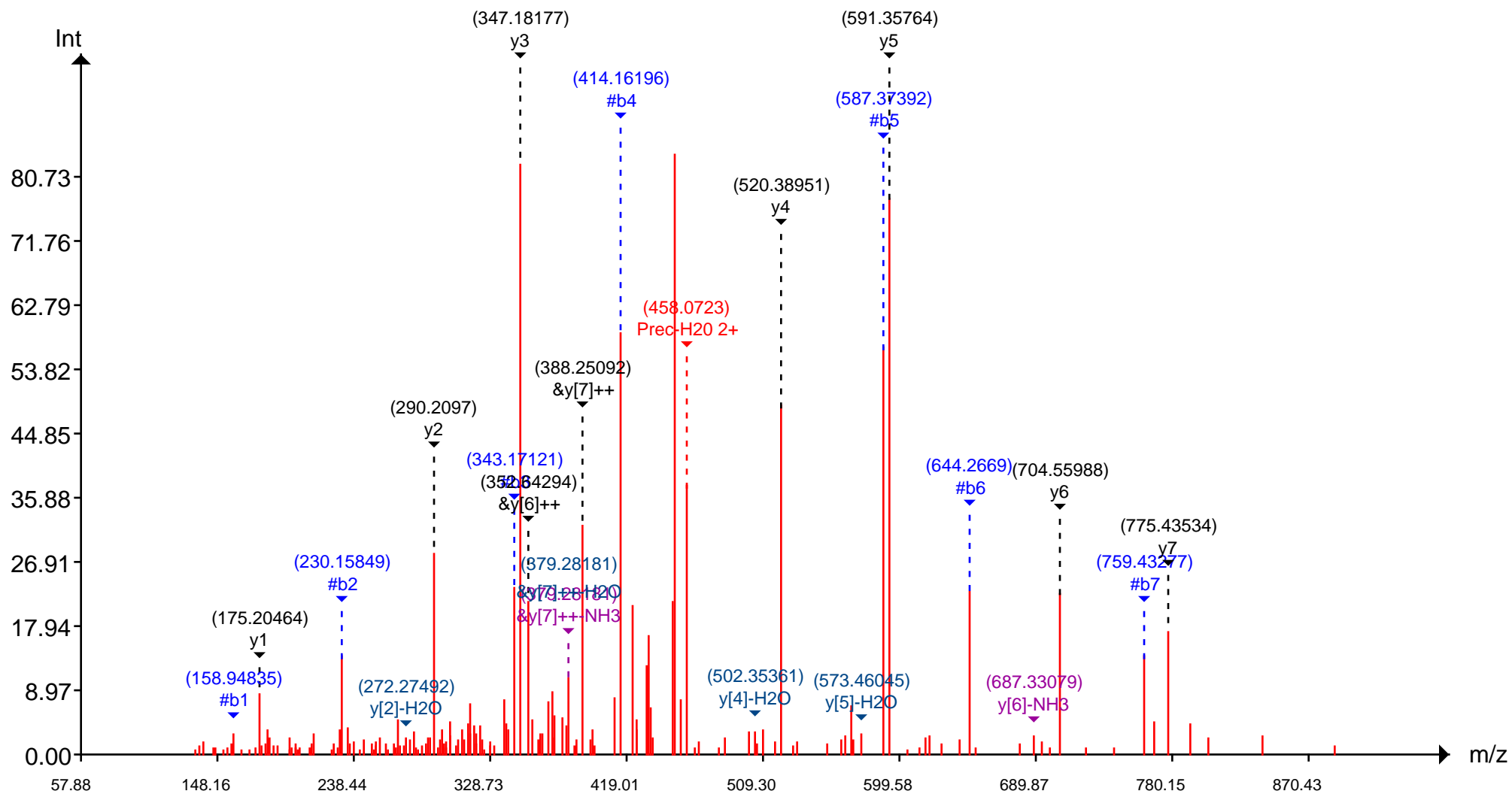
Charge 2+

Mass error 4.05E-4



Info

Modified Sequence	AcD3-EIIDK<AcD3K*>SYTPSK<AcD3K*>IR-COOH	Ion score	94.49
Protein	O70492	Confidence level	99.00%
Precursor m/z	842.97836	Identity Threshold	46.34
Charge	2+	Mass error	0.002852



Info

Modified Sequence AcD3-LALAK<AcD3K*>GDR-COOH

Ion score 67.76

Protein P10107

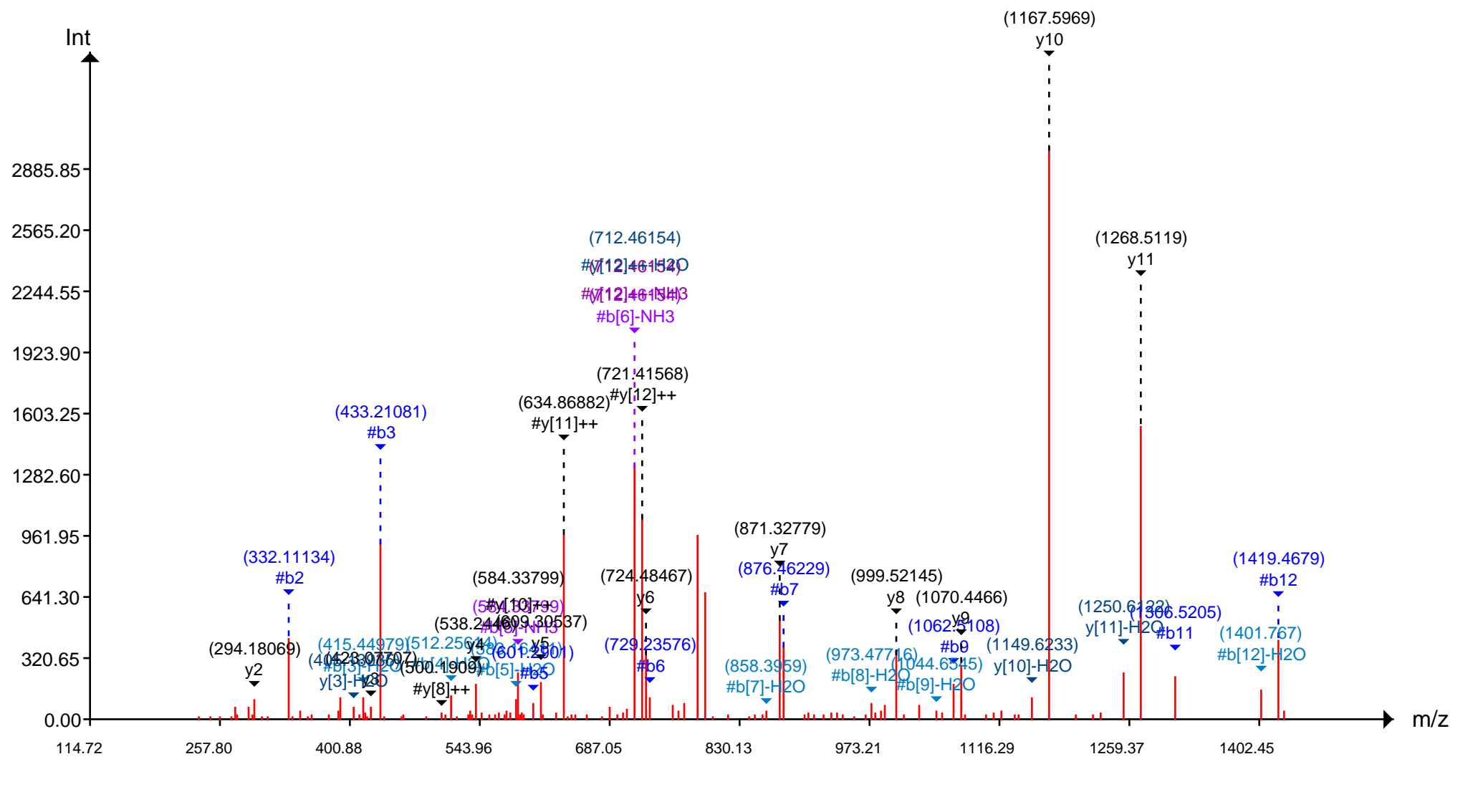
Confidence level 99.00%

Precursor m/z 467.28526

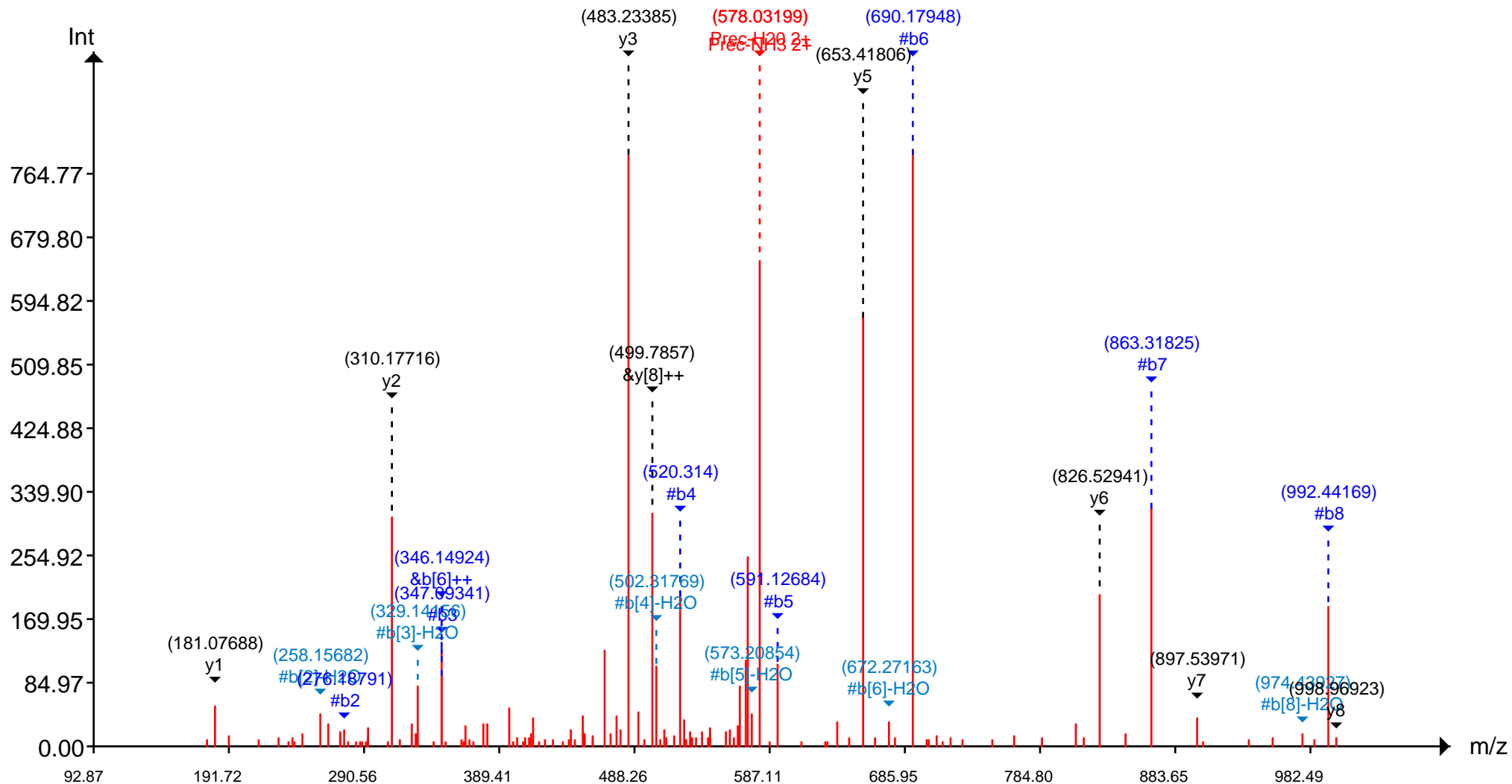
Identity Threshold 45.01

Charge 2+

Mass error -2.0E-4



Info			
Modified Sequence	AcD3-LK<AcD3K*>TPAQFDAELR<C13>-COOH	Ion score	72.66
Protein	P10107	Confidence level	99.00%
Precursor m/z	800.43389	Identity Threshold	47.90
Charge	2+	Mass error	0.001406



Info

Modified Sequence AcD3-ETAK<AcD3K*>AVK<AcD3K*>ER<C13>-COOH

Ion score 58.19

Protein Q99JY9

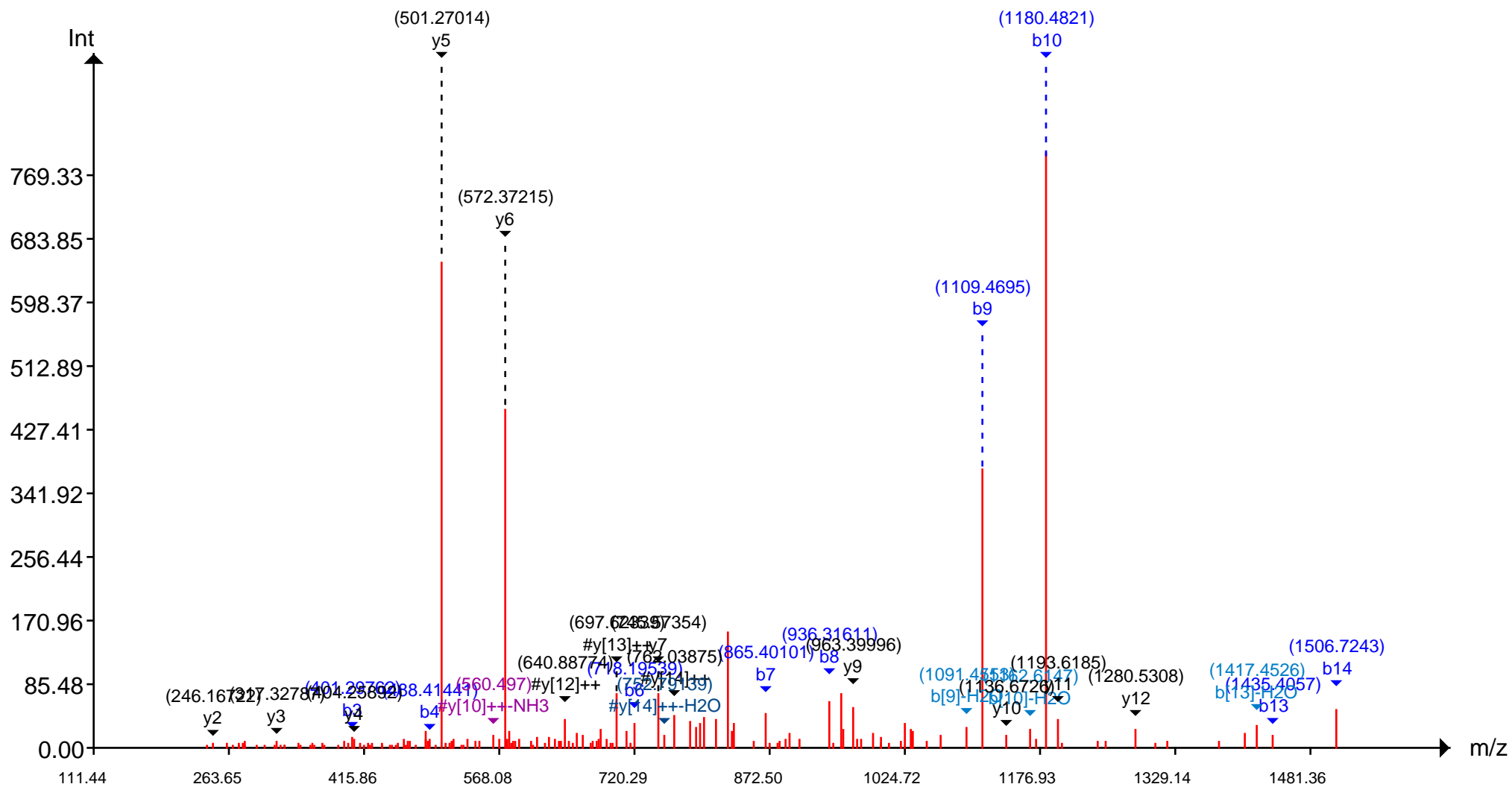
Confidence level 99.00%

Precursor m/z 586.85053

Identity Threshold 45.36

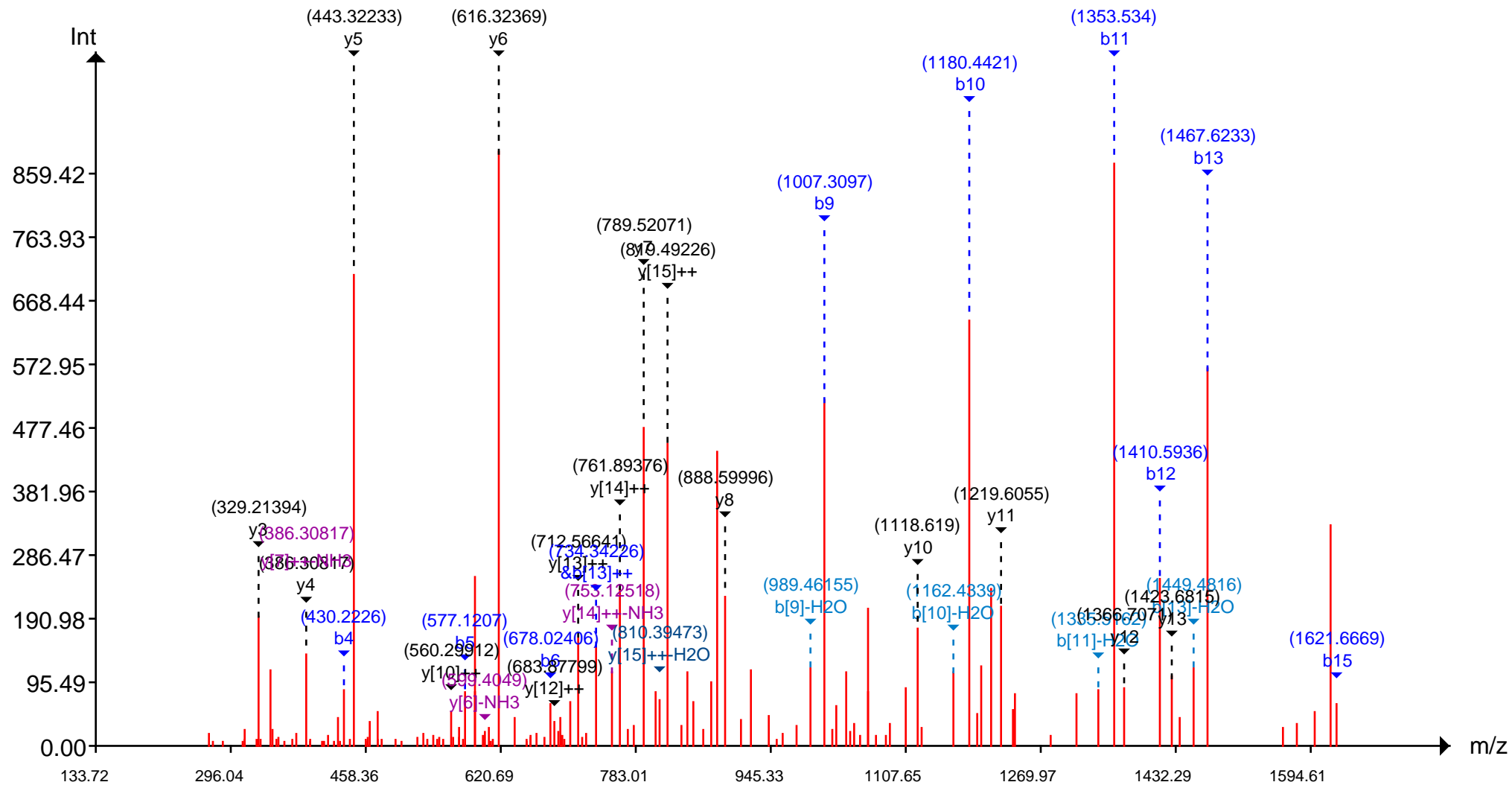
Charge 2+

Mass error 0.001098



Info

Modified Sequence	AcD3-IEISGK<AcD3K*>M<Mox*>AK<AcD3K*>APSAAR-COOH	Ion score	83.83
Protein	Q8BJF9	Confidence level	99.00%
Precursor m/z	840.96988	Identity Threshold	46.90
Charge	2+	Mass error	0.002601



Info

Modified Sequence AcD3-IDVGFTDDVK<AcD3K*>K<AcD3K*>GGPGR-COOH

Ion score 57.35

Protein Q9CX00

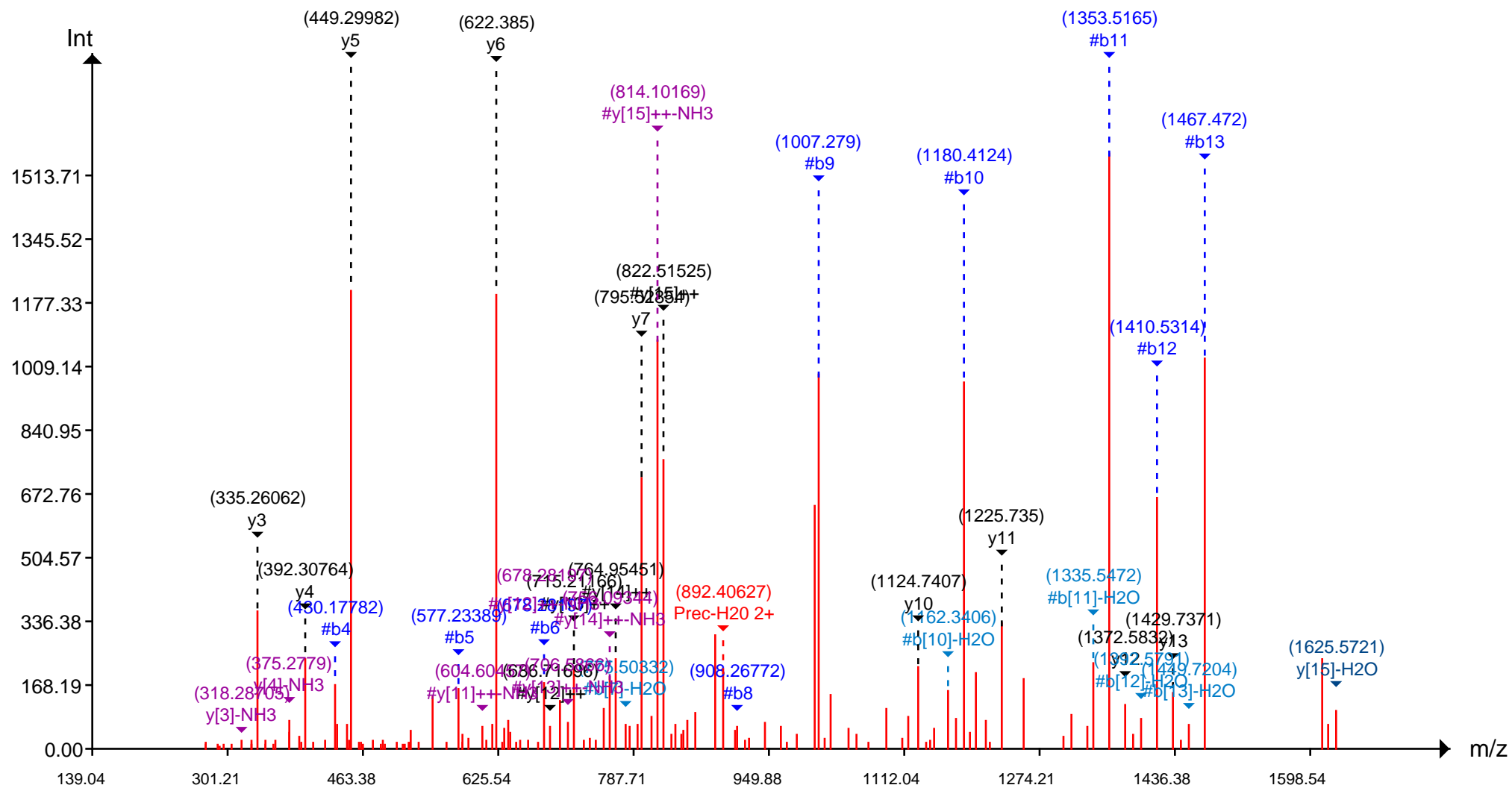
Confidence level 99.00%

Precursor m/z 898.48158

Identity Threshold 47.82

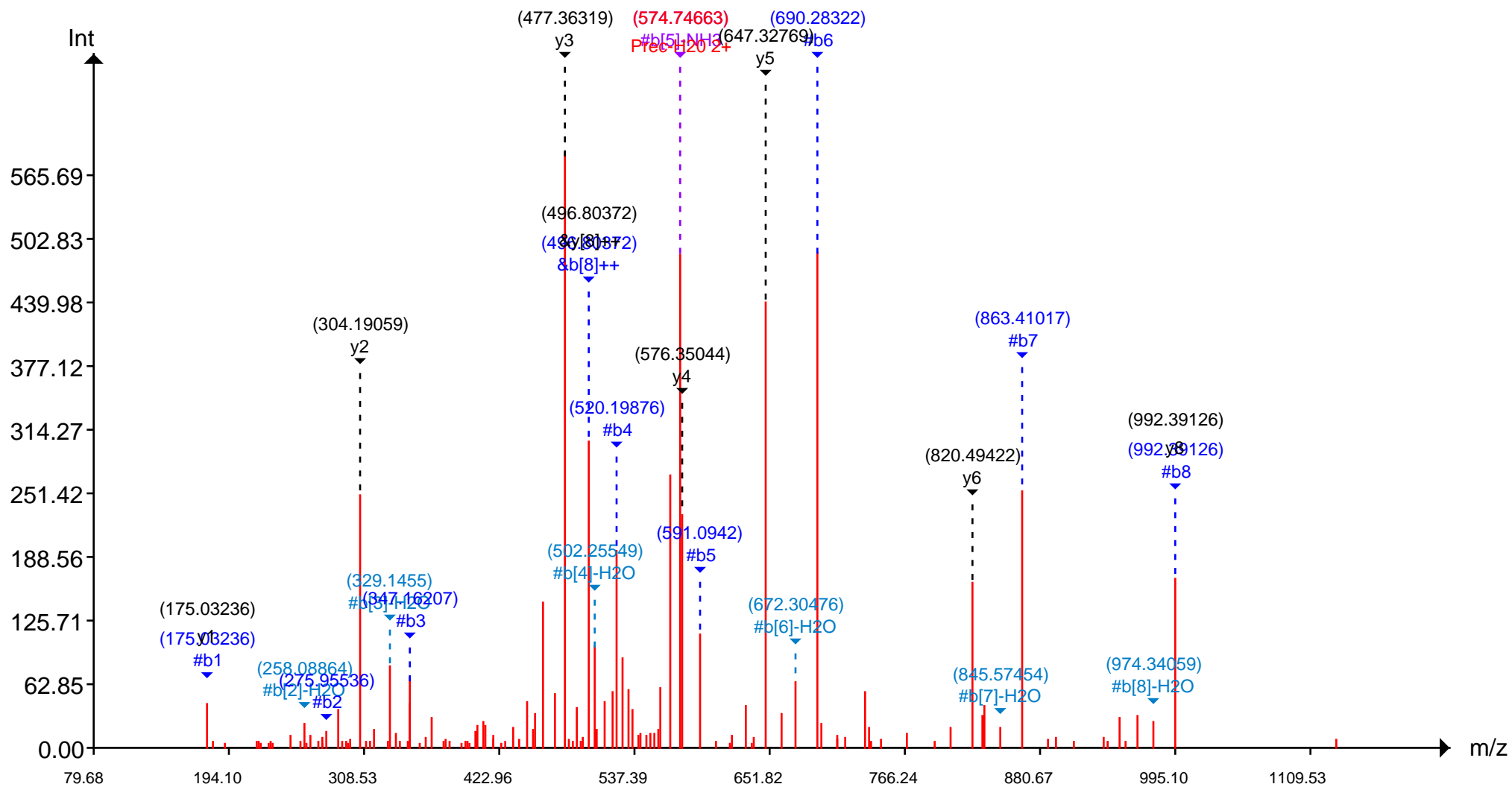
Charge 2+

Mass error 0.002349



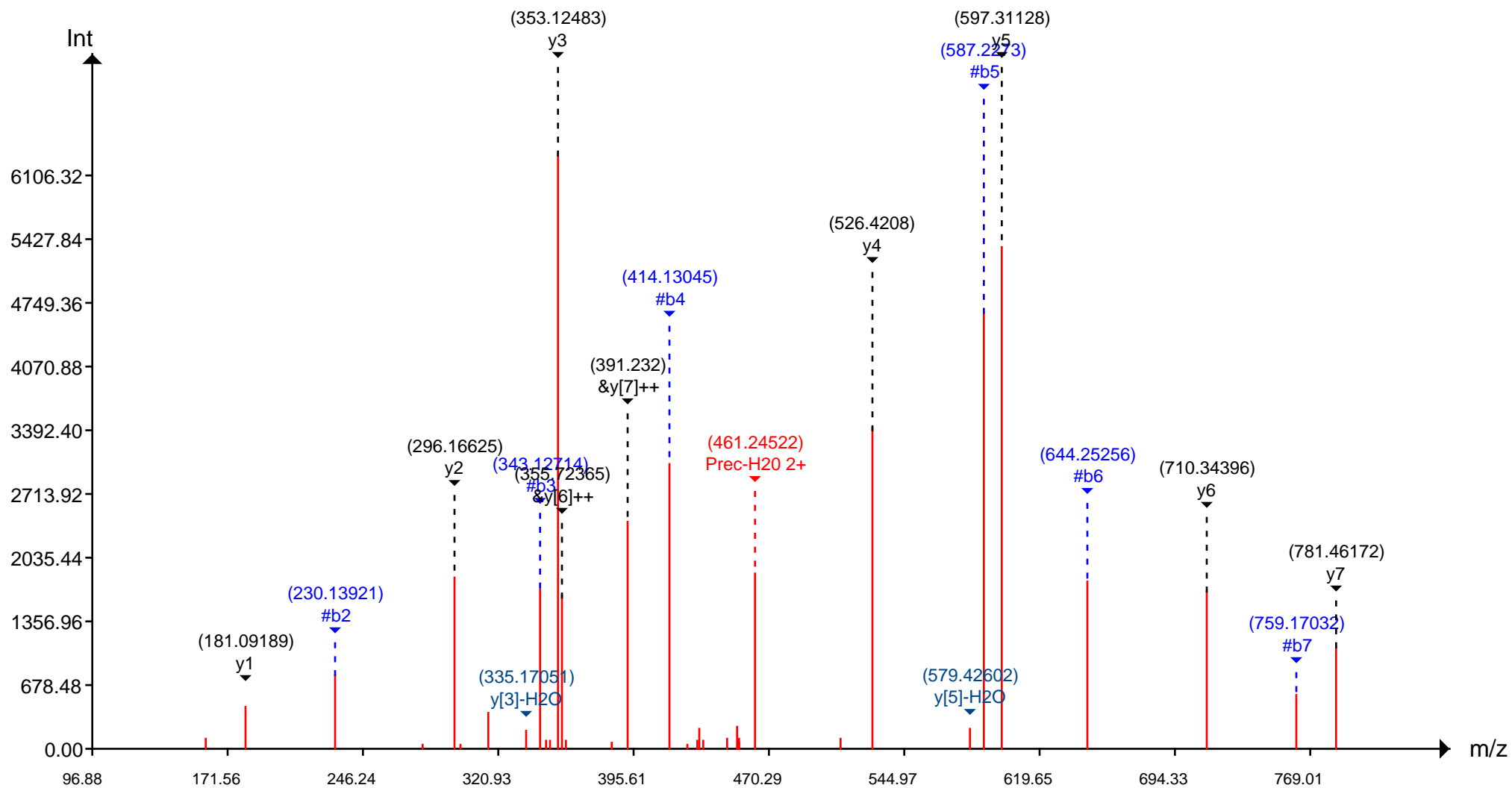
Info

Modified Sequence	AcD3-IDVGFTDDVK<AcD3K*>K<AcD3K*>GGPGR<C13>-COOH	Ion score	59.61
Protein	Q9CX00	Confidence level	99.00%
Precursor m/z	901.4912	Identity Threshold	47.61
Charge	2+	Mass error	0.001448



Info

Modified Sequence	AcD3-ETAK<AcD3K*>AVK<AcD3K*>ER-COOH	Ion score	57.43
Protein	Q99JY9	Confidence level	99.00%
Precursor m/z	583.8401	Identity Threshold	46.18
Charge	2+	Mass error	3.8E-4



Info

Modified Sequence AcD3-LALAK<AcD3K*>GDR<C13>-COOH

Ion score 72.12

Protein P10107

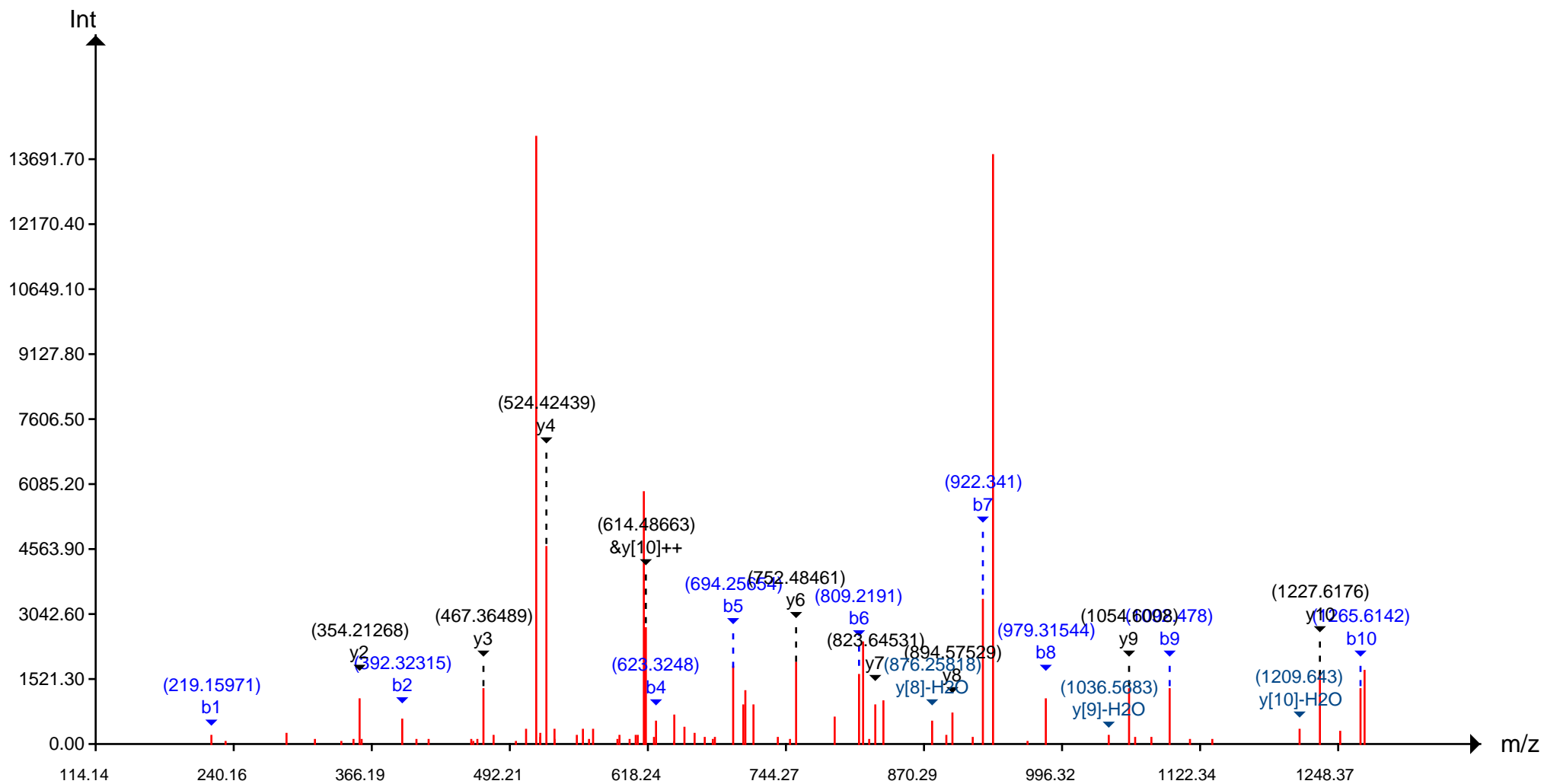
Confidence level 99.00%

Precursor m/z 470.29514

Identity Threshold 43.99

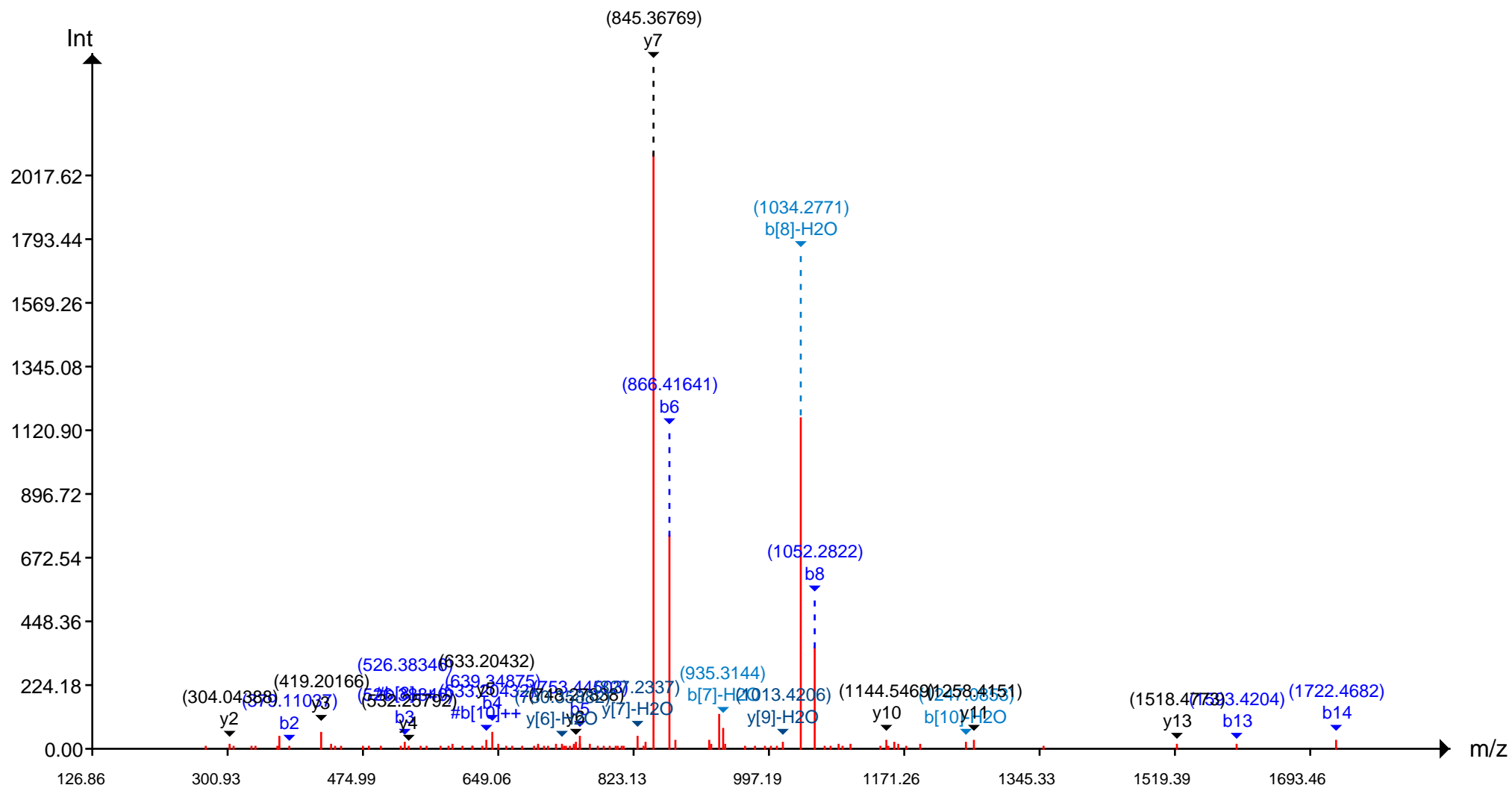
Charge 2+

Mass error -5.81E-4



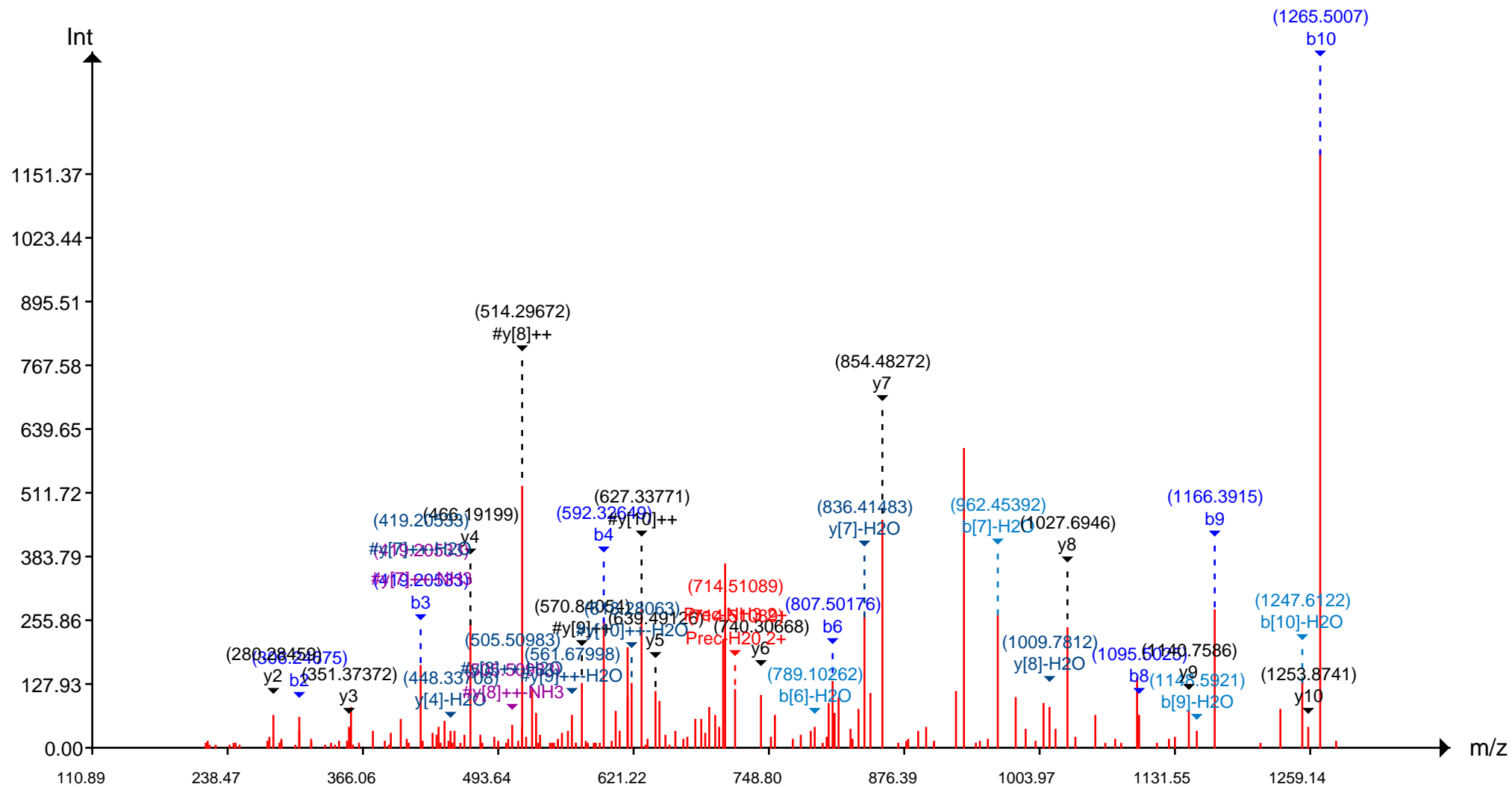
Info

Modified Sequence	AcD3-K<AcD3K*>K<AcD3K*>C<Cmm*>AADLGLK<AcD3K*>R<C13>-COOH	Ion score	57.39
Protein	P70349	Confidence level	99.00%
Precursor m/z	723.4345	Identity Threshold	43.91
Charge	2+	Mass error	-0.001234



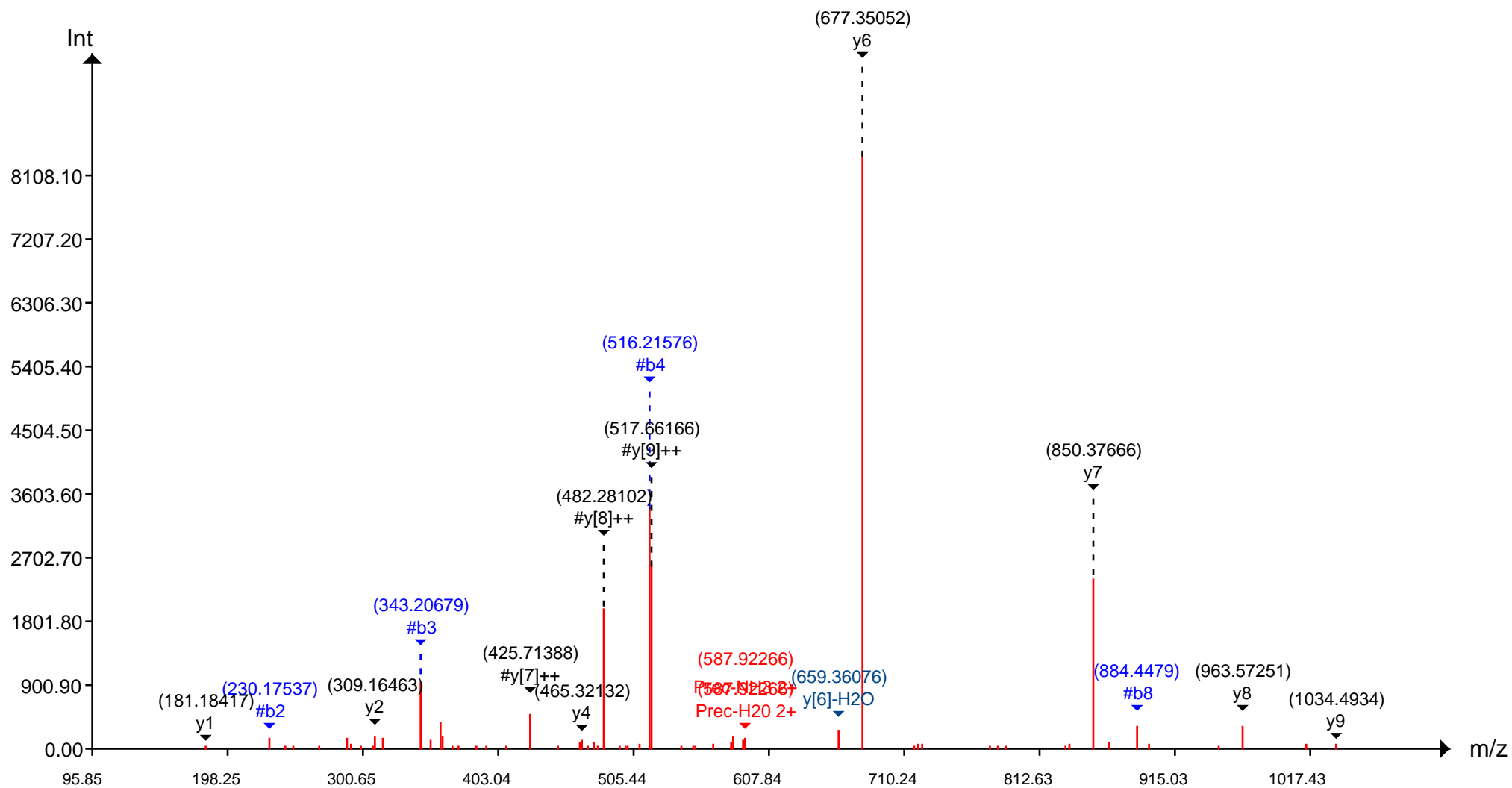
Info

Modified Sequence	AcD3-C<Cmm*>K<AcD3K*>M<Mox*>INLSVPDTIDER-COOH	Ion score	72.18
Protein	Q61233, Q99K51	Confidence level	99.00%
Precursor m/z	948.97058	Identity Threshold	47.50
Charge	2+	Mass error	0.002658



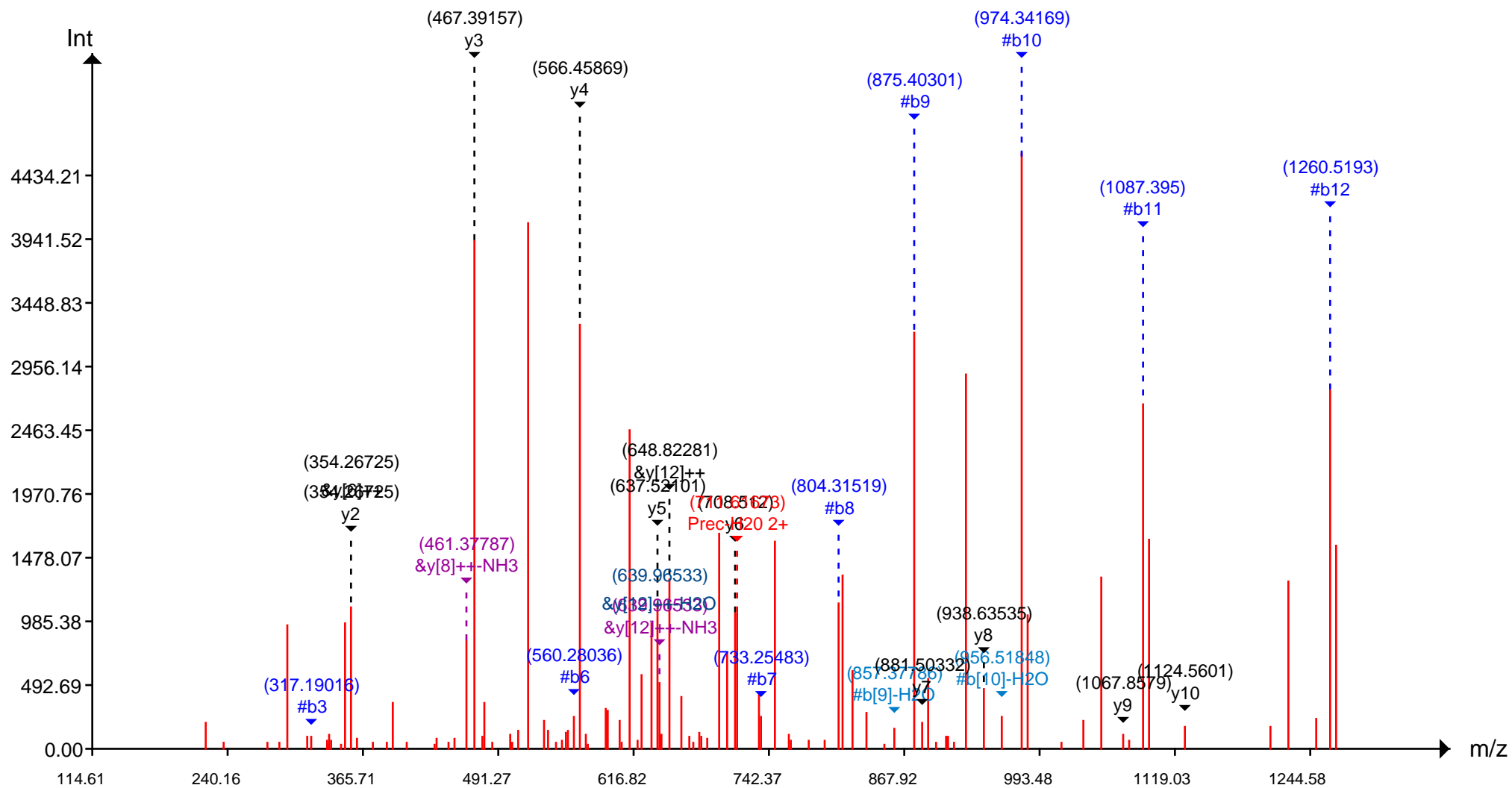
Info

Modified Sequence	AcD3-M<Mox*>ILK<AcD3K*>NTK<AcD3K*>DAVR<C13>-COOH	Ion score	52.58
Protein	Q3TXS7	Confidence level	99.00%
Precursor m/z	723.42539	Identity Threshold	45.23
Charge	2+	Mass error	-2.89E-4



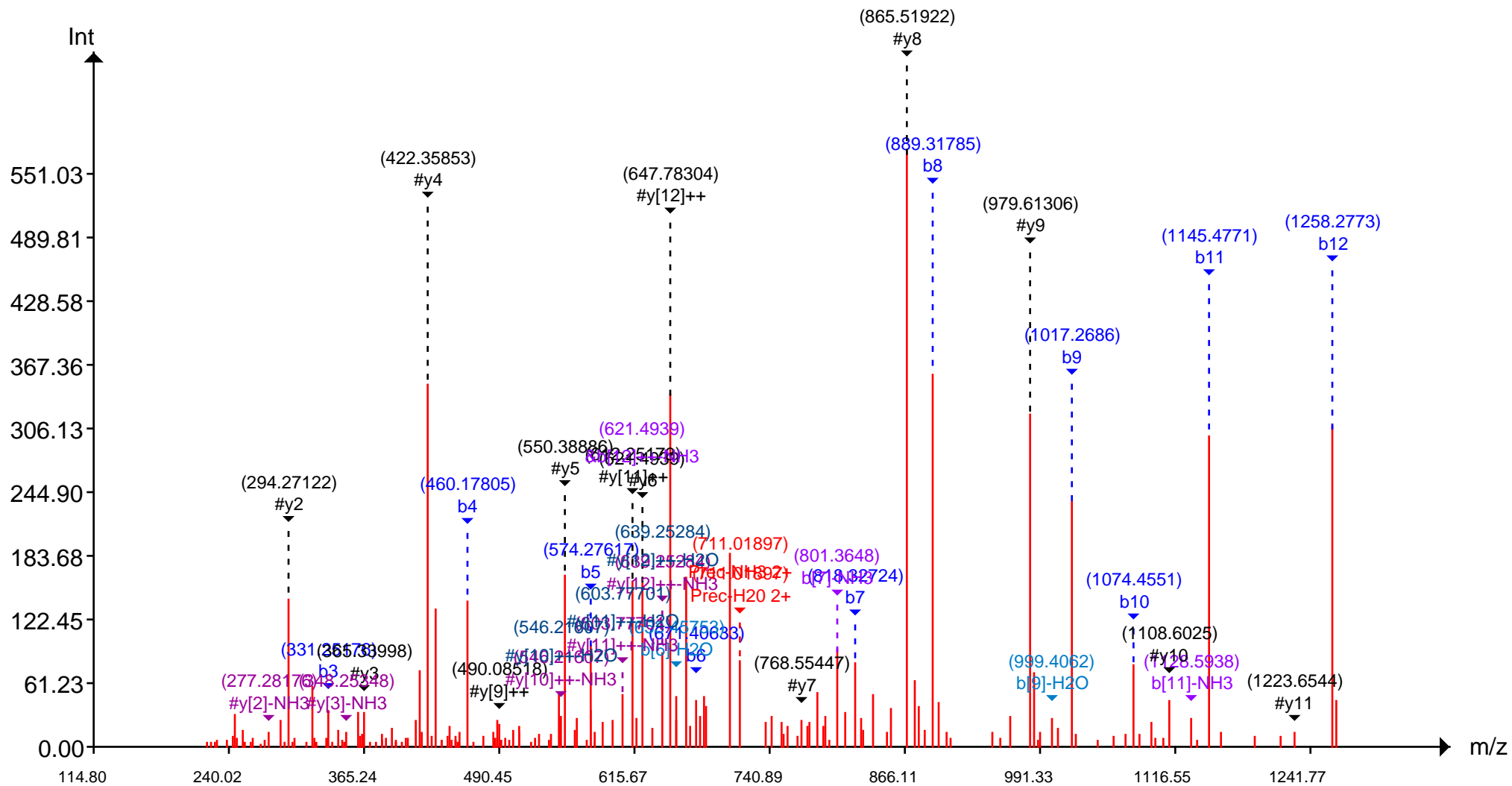
Info

Modified Sequence	AcD3-IAIK<AcD3K*>PDGVQR<C13>-COOH	Ion score	47.15
Protein	P15532, Q01768	Confidence level	99.00%
Precursor m/z	596.86754	Identity Threshold	43.11
Charge	2+	Mass error	0.001564



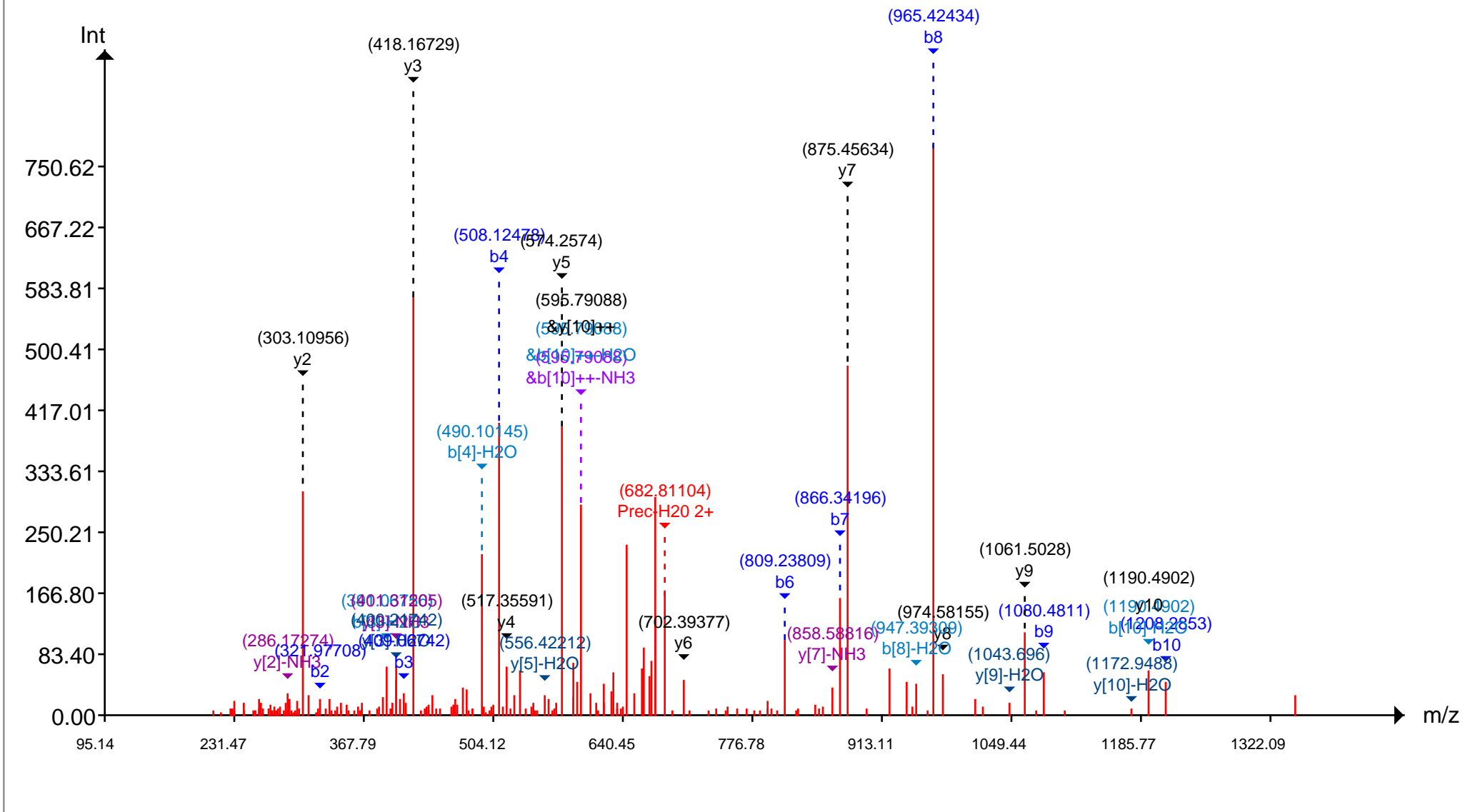
Info

Modified Sequence	AcD3-VATGEGK<AcD3K*>AAVLK<AcD3K*>R<C13>-COOH	Ion score	43.2
Protein	Q9CQ60	Confidence level	99.00%
Precursor m/z	720.94708	Identity Threshold	41.37
Charge	2+	Mass error	0.004257



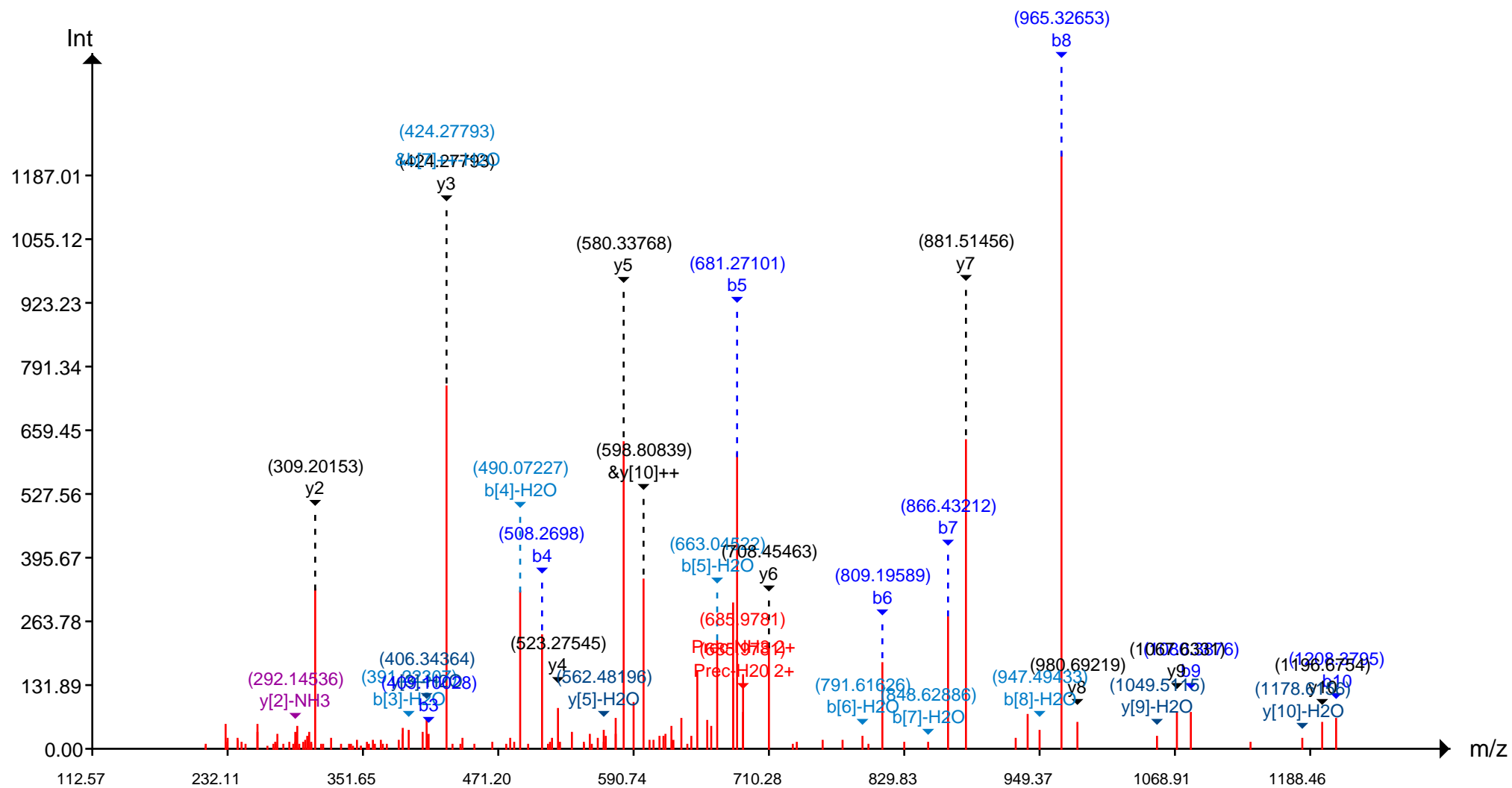
Info

Modified Sequence	AcD3-VADENPFAQGALR<C13>-COOH	Ion score	73.16
Protein	P13020	Confidence level	99.00%
Precursor m/z	719.87748	Identity Threshold	48.02
Charge	2+	Mass error	0.001715



Info

Modified Sequence	AcD3-M<Mox*>ESVK<AcD3K*>QGVDQR-COOH	Ion score	55.39
Protein	Q9DBG5	Confidence level	99.00%
Precursor m/z	691.84663	Identity Threshold	47.56
Charge	2+	Mass error	8.67E-4



Info

Modified Sequence AcD3-M<Mox*>ESVK<AcD3K*>QGVDQR<C13>-COOH

Ion score 70.6

Protein Q9DBG5

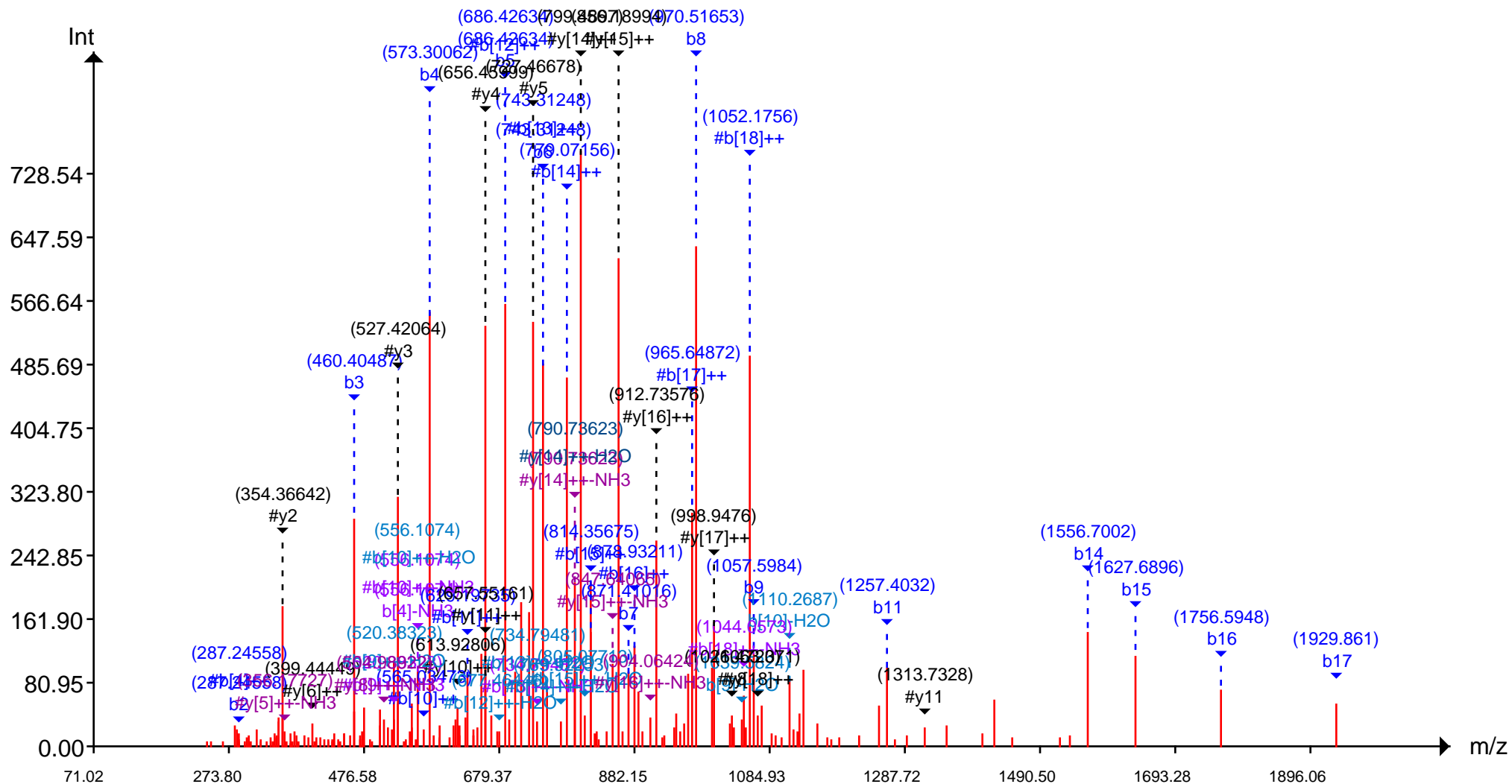
Confidence level 99.00%

Precursor m/z 694.85663

Identity Threshold 47.88

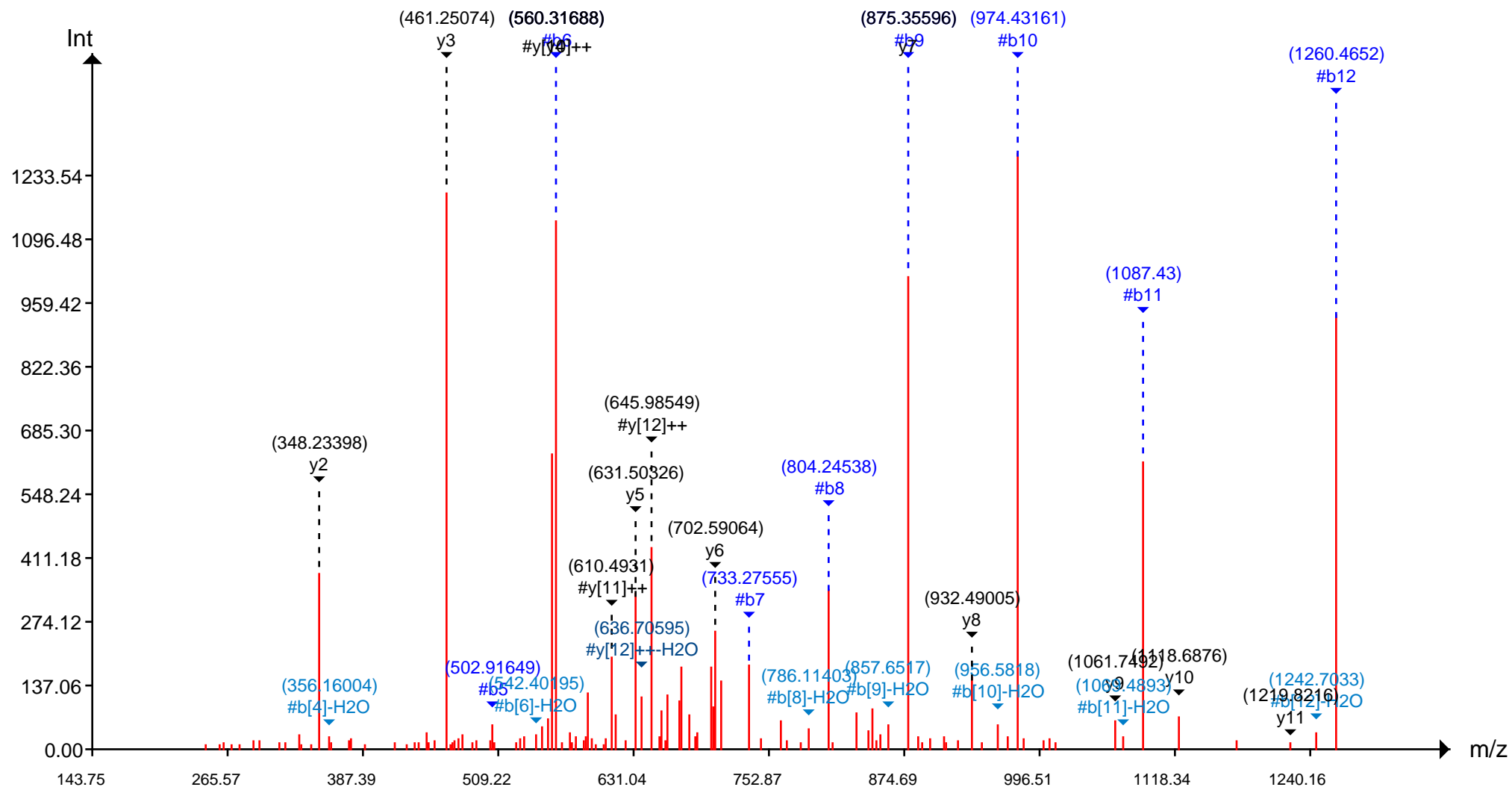
Charge 2+

Mass error 7.25E-4



Info

Modified Sequence	AcD3-IQK<AcD3K*>LLGQVSAEDLAAEK<AcD3K*>K<AcD3K*>R<C13>-COOH	Ion score	78.15
Protein	Q9CY64	Confidence level	99.00%
Precursor m/z	761.78557	Identity Threshold	41.58
Charge	3+	Mass error	0.001844



Info

Modified Sequence AcD3-VATGEGK<AcD3K*>AAVLK<AcD3K*>R-COOH

Ion score 82.59

Protein Q9CQ60

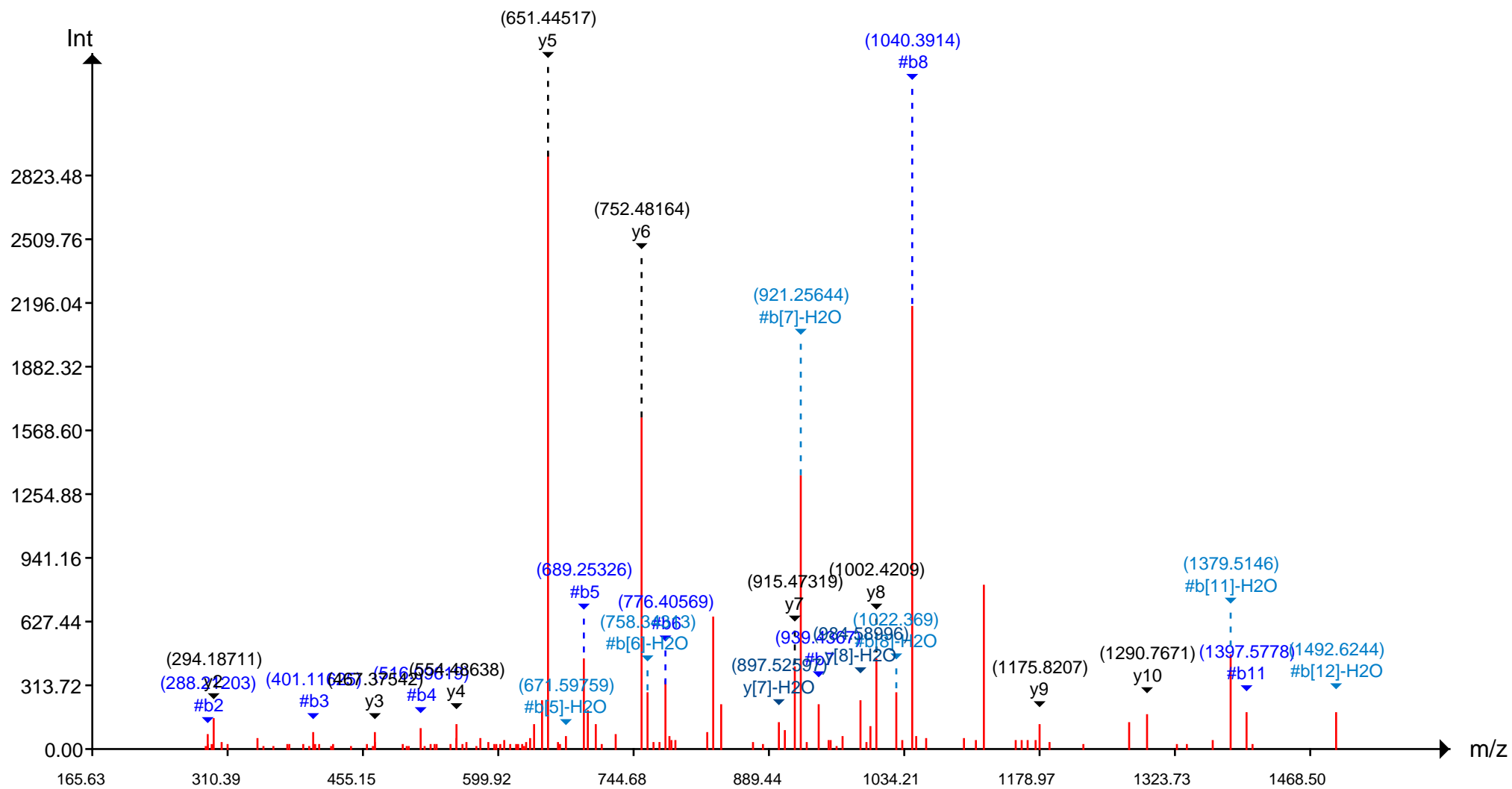
Confidence level 99.00%

Precursor m/z 717.93524

Identity Threshold 43.06

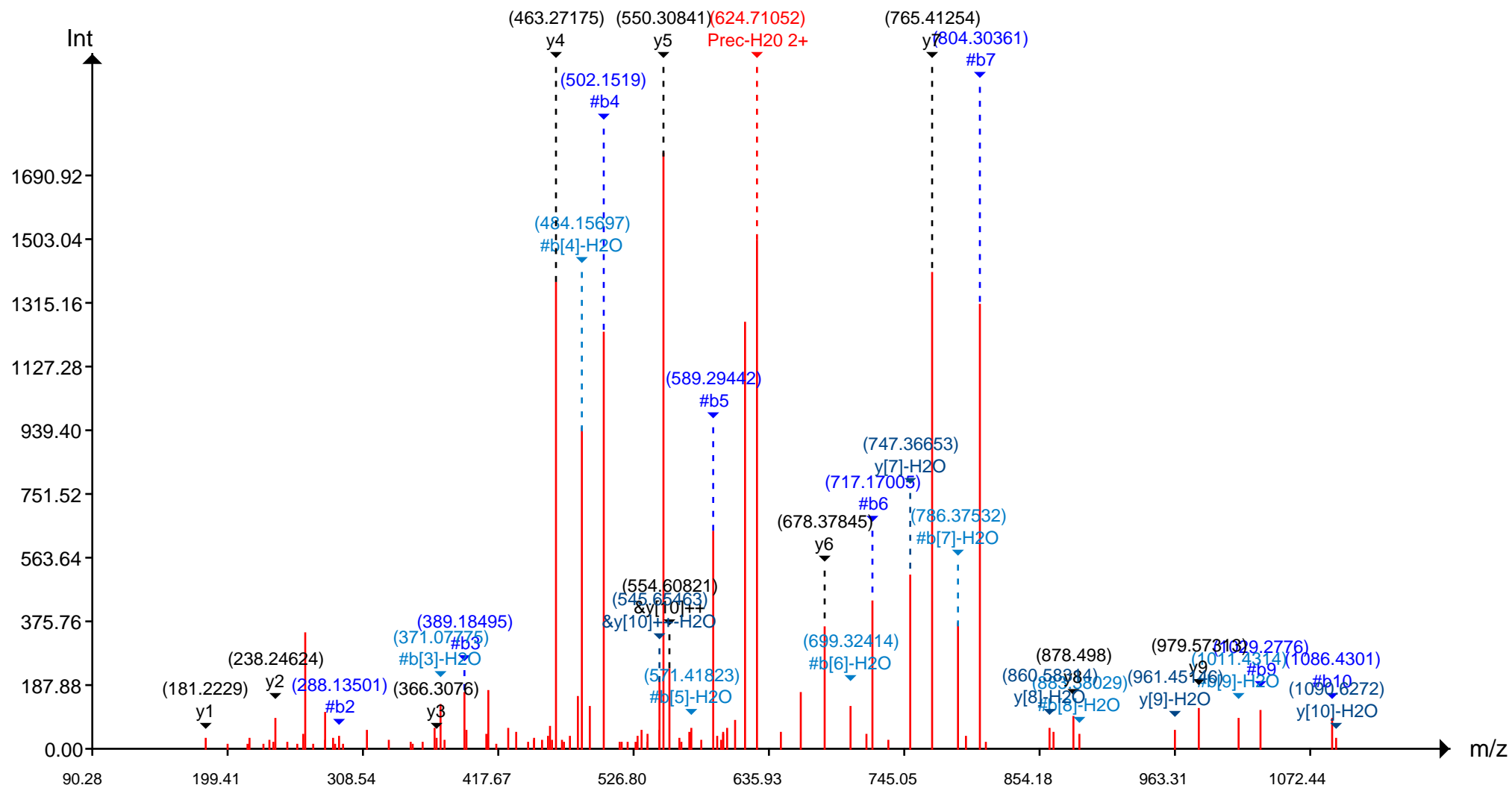
Charge 2+

Mass error 7.19E-4



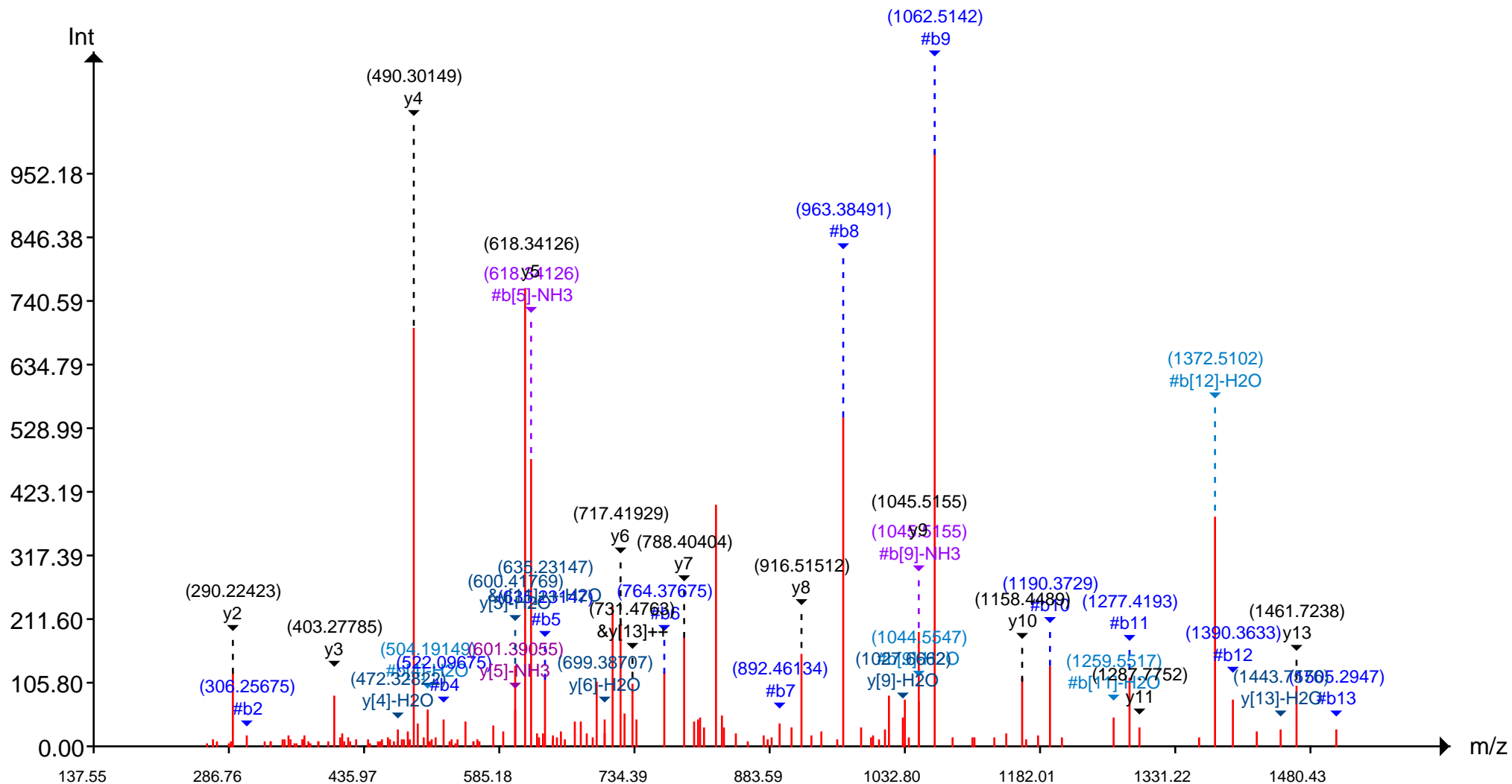
Info

Modified Sequence	AcD3-EIIDK<AcD3K*>SYTPSK<AcD3K*>IR<C13>-COOH	Ion score	72.48
Protein	O70492	Confidence level	99.00%
Precursor m/z	845.98808	Identity Threshold	45.51
Charge	2+	Mass error	0.002151



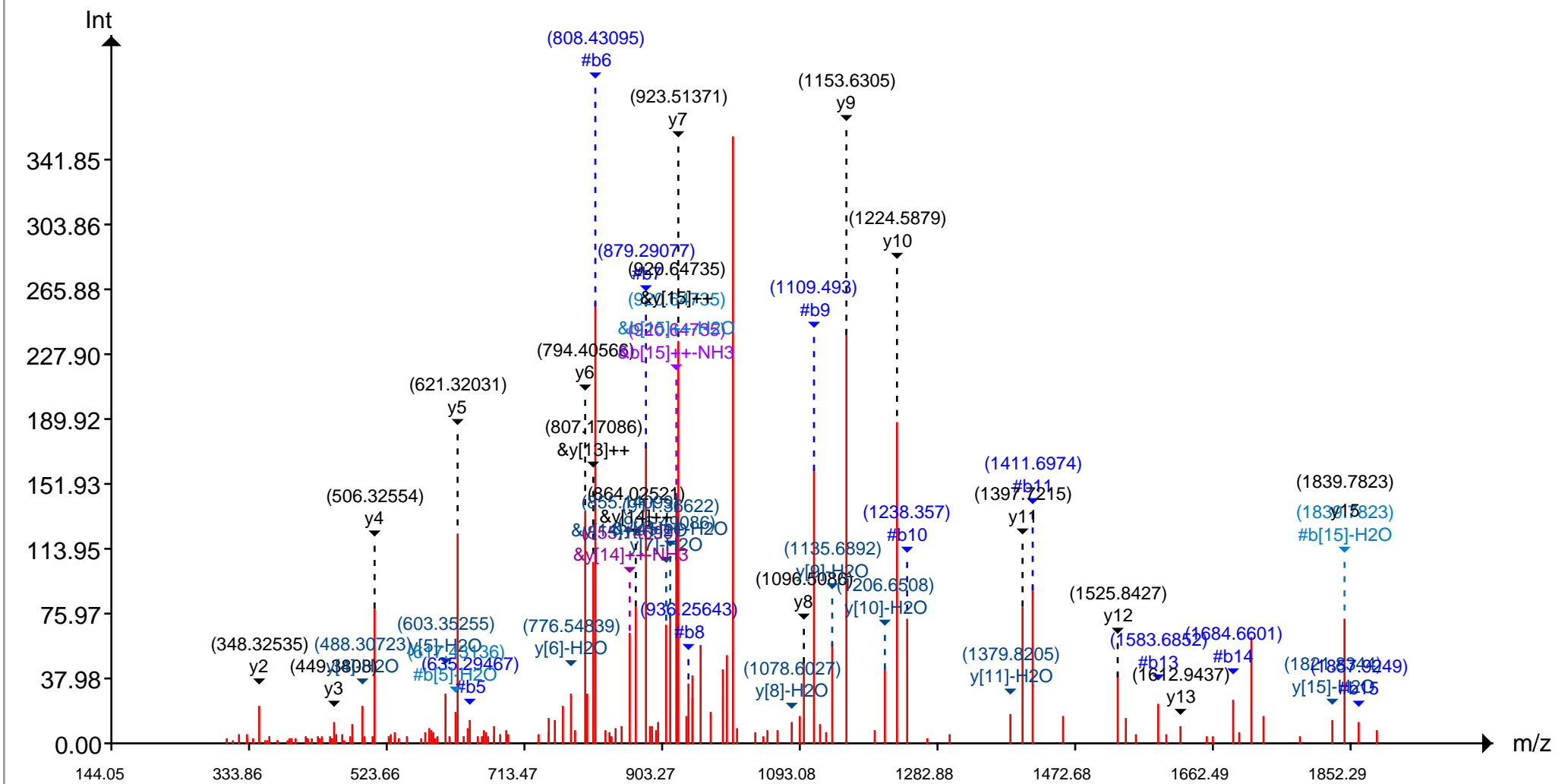
Info

Modified Sequence	AcD3-LETLSQSPQGR<C13>-COOH	Ion score	64.48
Protein	P60335	Confidence level	99.00%
Precursor m/z	633.84508	Identity Threshold	47.81
Charge	2+	Mass error	5.74E-4



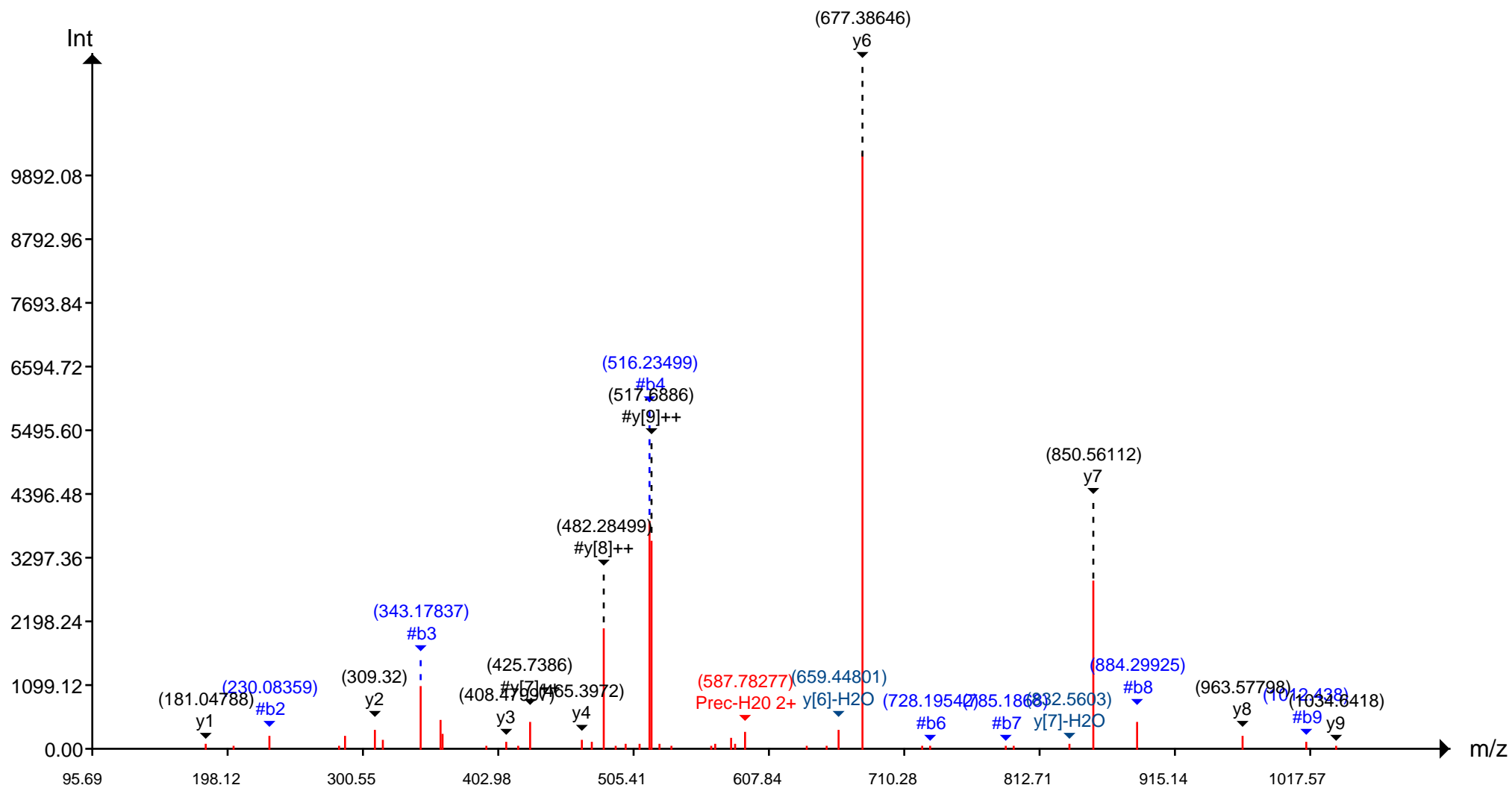
Info

Modified Sequence	AcD3-K<AcD3K*>SSEIEQAVQSLDR-COOH	Ion score	107.07
Protein	Q9D898	Confidence level	99.00%
Precursor m/z	840.44067	Identity Threshold	48.12
Charge	2+	Mass error	0.002362



Info

Modified Sequence	AcD3-FLNSQ<AcD3K*>AGK<AcD3K*>EK<AcD3K*>DGTK<AcD3K*>R-COOH	Ion score	117.61
Protein	P23198	Confidence level	99.00%
Precursor m/z	1016.5693	Identity Threshold	46.30
Charge	2+	Mass error	0.002283



Info

Modified Sequence AcD3-IAIK<AcD3K*>PDGVQR<C13>-COOH

Ion score 76.37

Protein P15532, Q01768

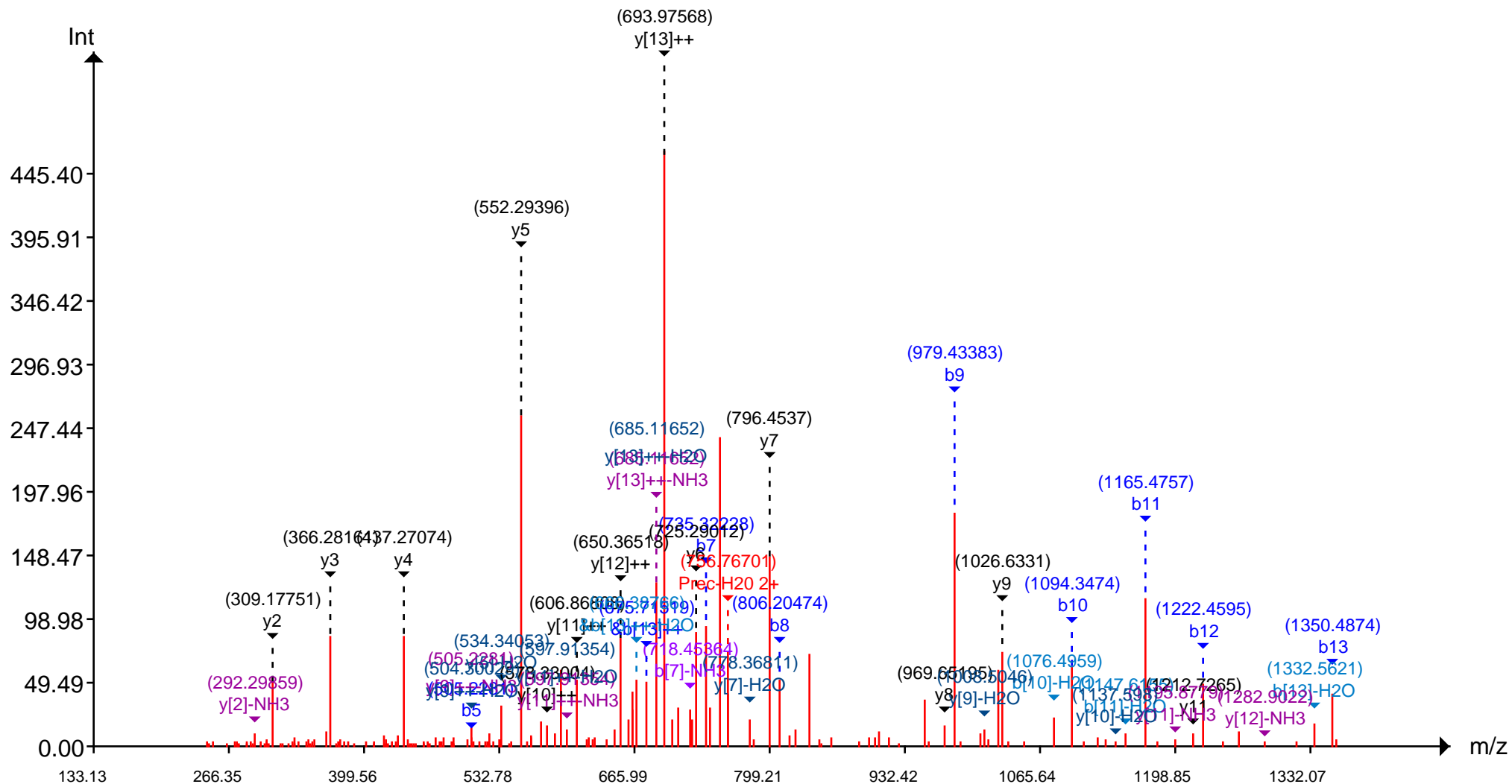
Confidence level 99.00%

Precursor m/z 596.86709

Identity Threshold 42.91

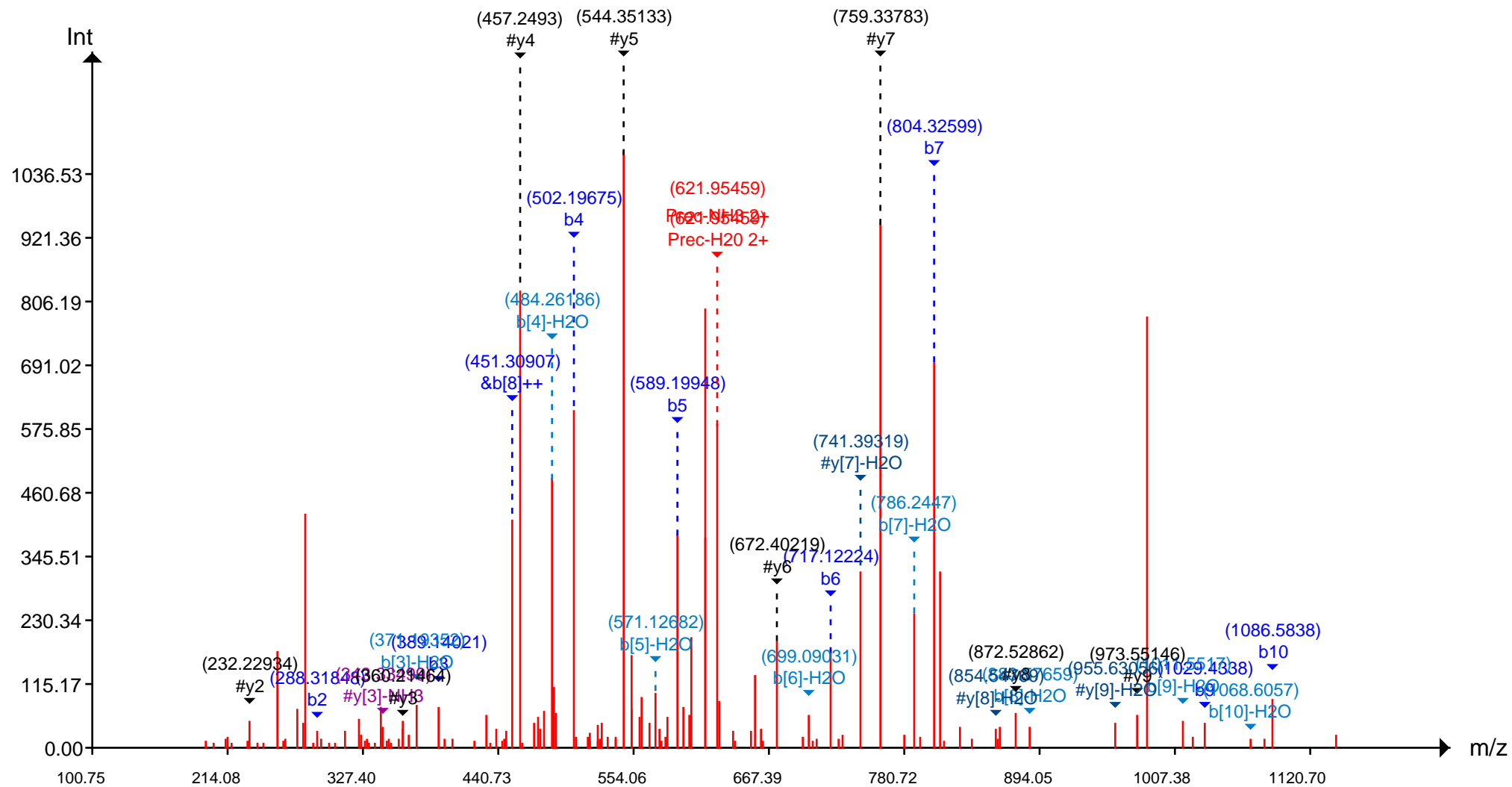
Charge 2+

Mass error 6.64E-4



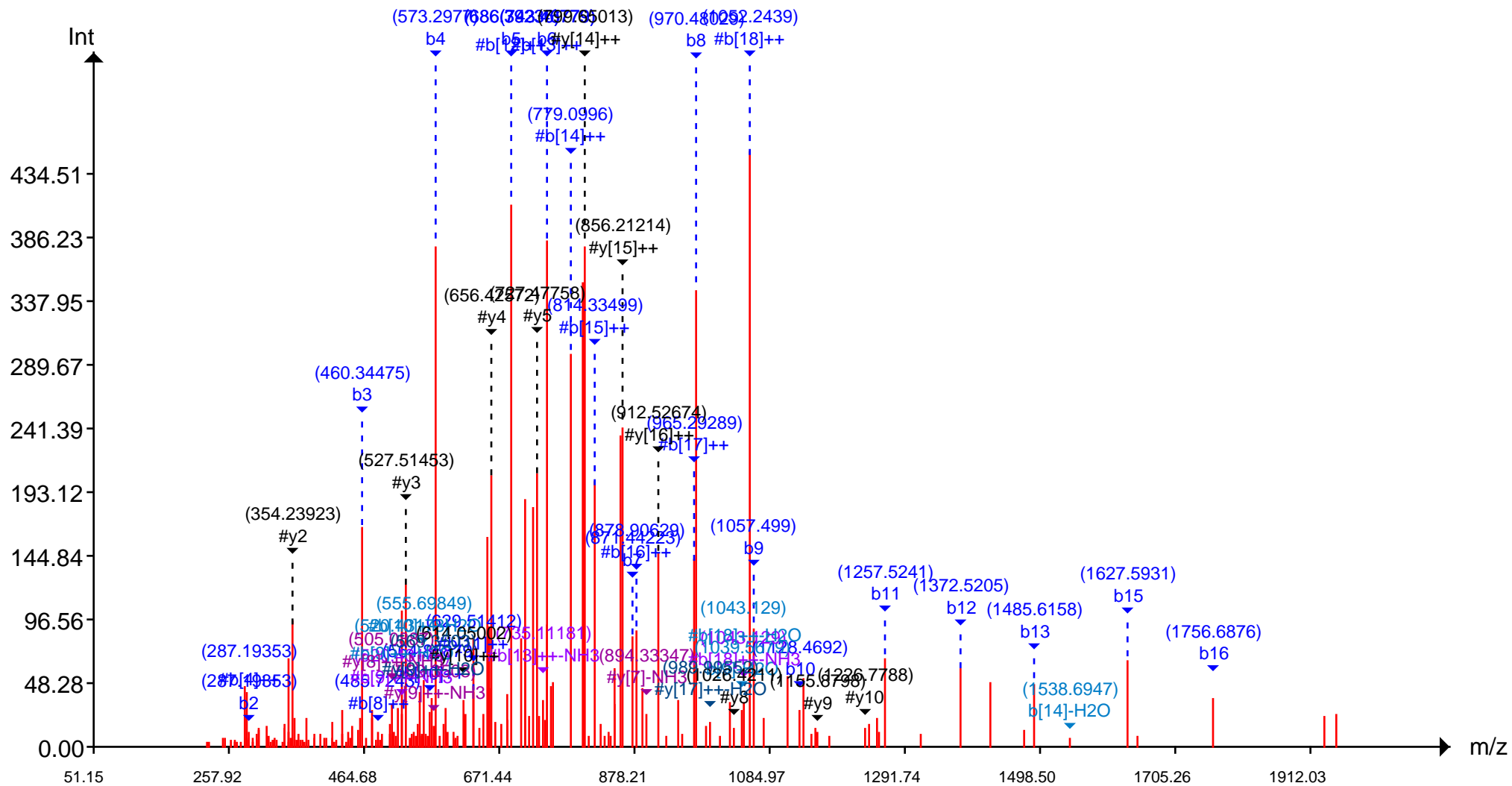
Info

Modified Sequence	AcD3-VSSGEGK<AcD3K*>AK<AcD3K*>DAGQR<C13>-COOH	Ion score	54.93
Protein	P50580	Confidence level	99.00%
Precursor m/z	765.91202	Identity Threshold	48.04
Charge	2+	Mass error	3.6E-4



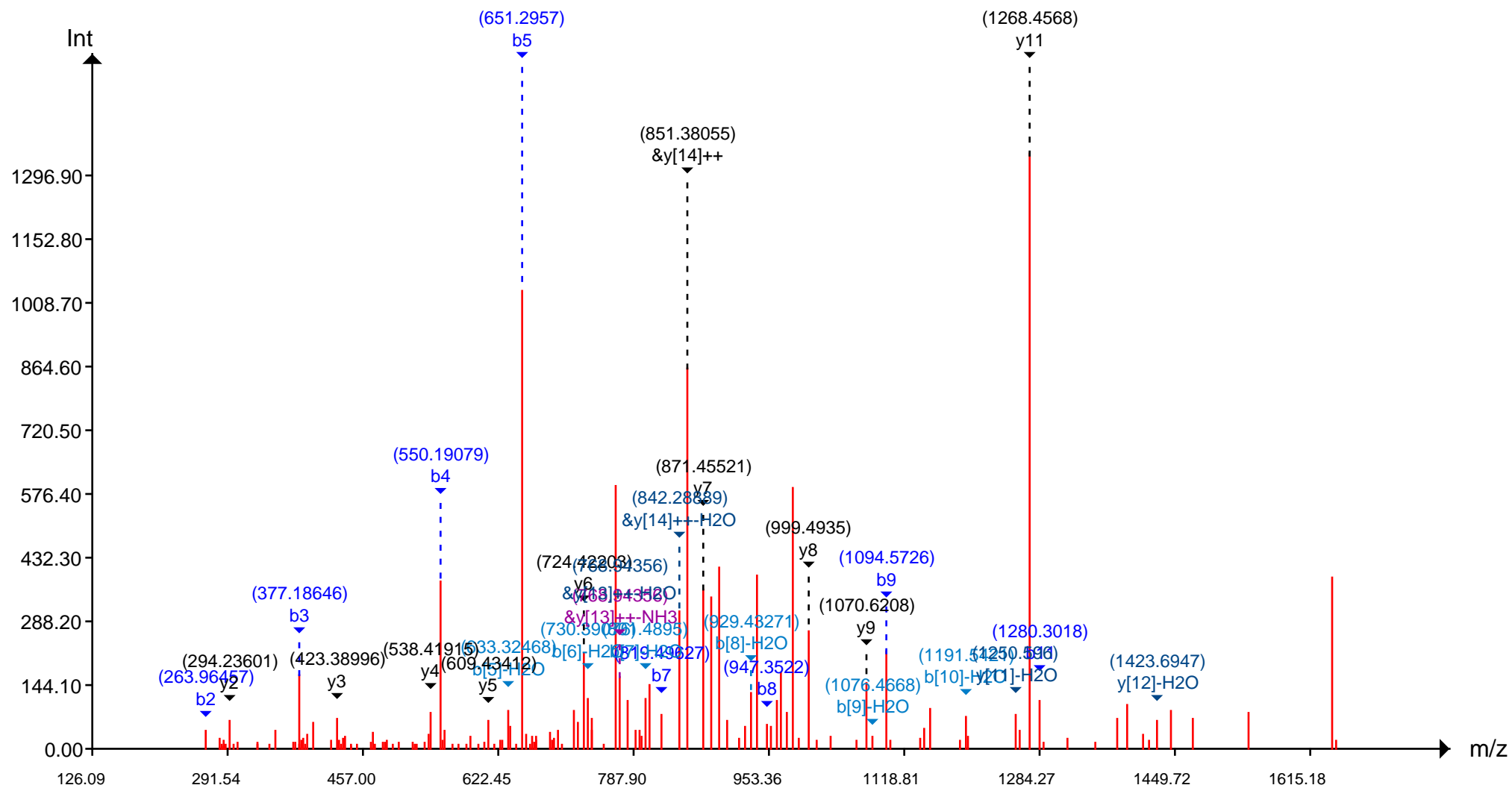
Info

Modified Sequence	AcD3-LETLSQSPQGR-COOH	Ion score	48.56
Protein	P60335	Confidence level	99.00%
Precursor m/z	630.83531	Identity Threshold	48.05
Charge	2+	Mass error	0.001176



Info

Modified Sequence	AcD3-IQK<AcD3K*>LLGQVSAEDLAAEK<AcD3K*>K<AcD3K*>R<C13>-COOH	Ion score	85.82
Protein	Q9CY64	Confidence level	99.00%
Precursor m/z	761.78572	Identity Threshold	41.59
Charge	3+	Mass error	0.002294



Info

Modified Sequence AcD3-AM<Mox*>LK<AcD3K*>TPAQFDADELRL<C13>-COOH

Ion score 54.83

Protein P10107

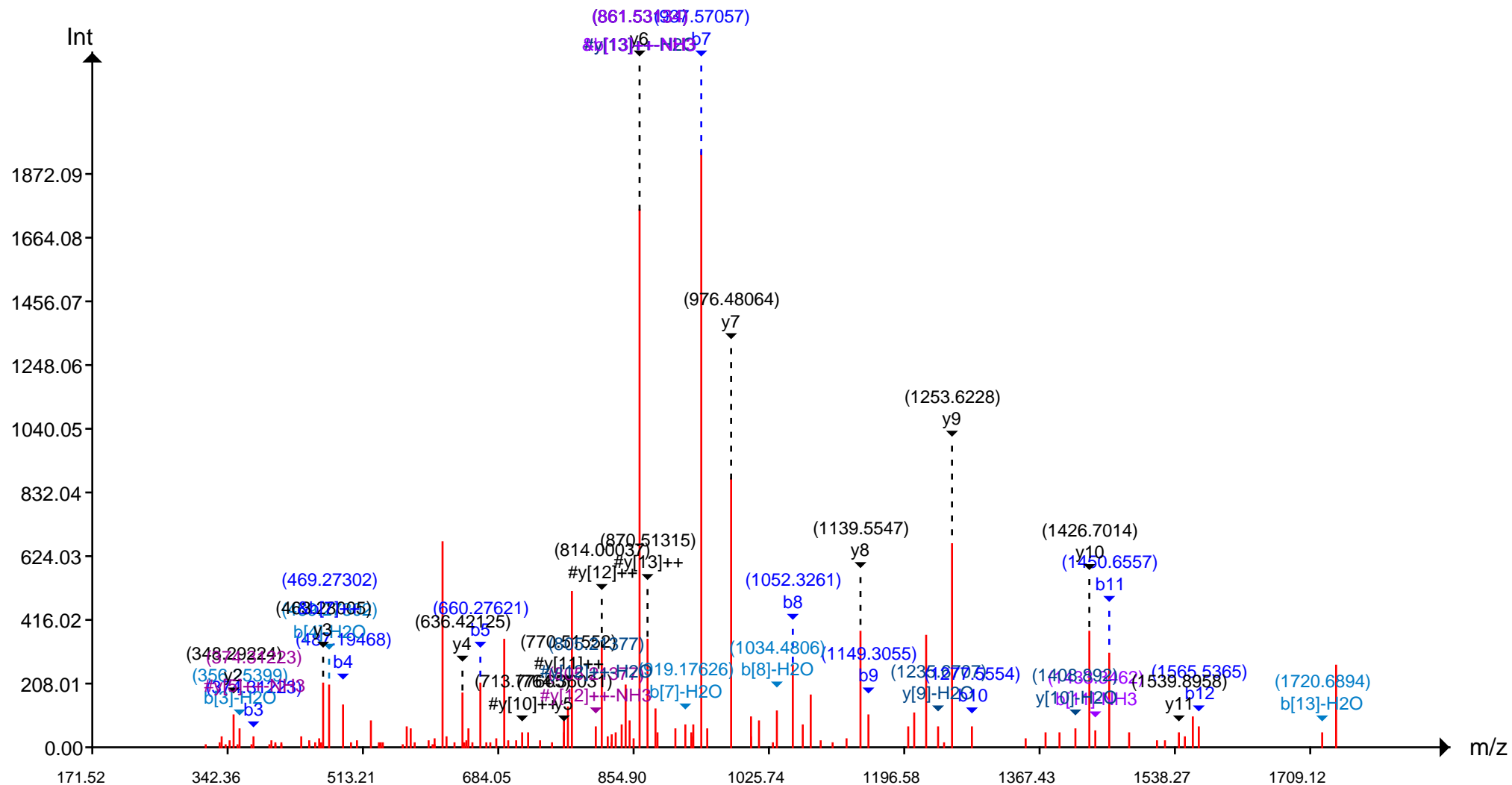
Confidence level 99.00%

Precursor m/z 909.46965

Identity Threshold 48.10

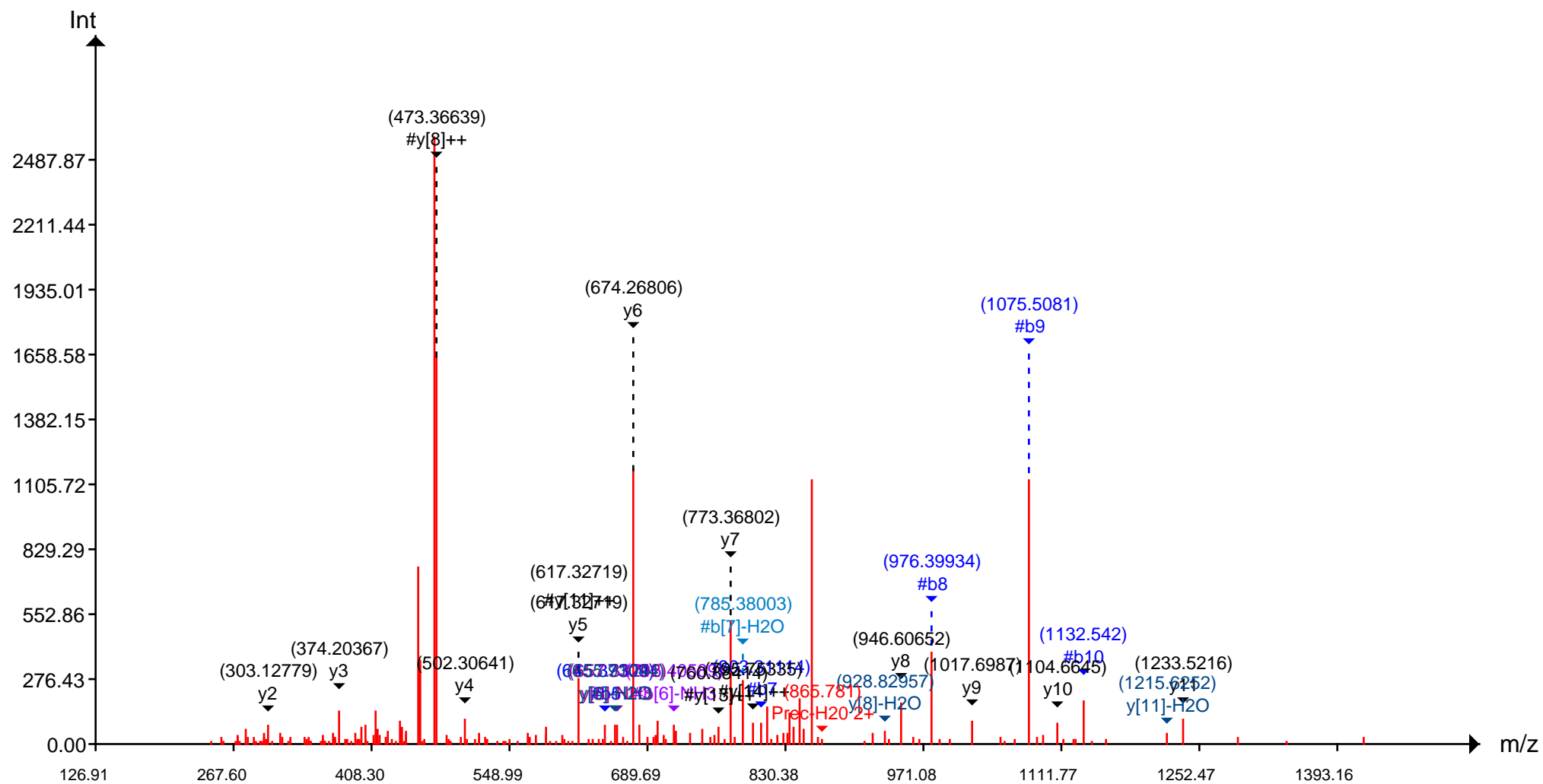
Charge 2+

Mass error 4.16E-4



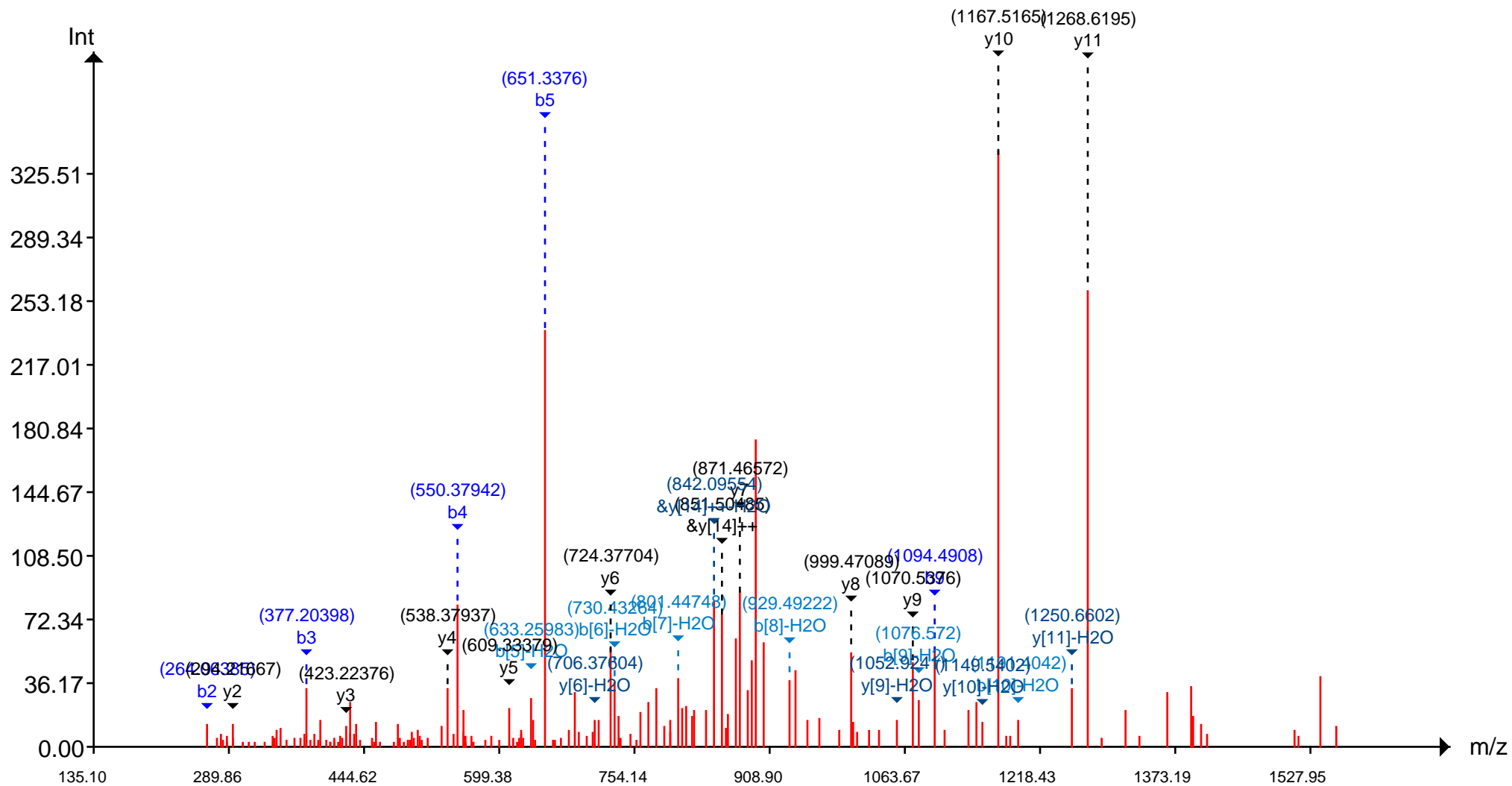
Info

Modified Sequence	AcD3-QISLK<AcD3K*>NYDPQK<AcD3K*>DK<AcD3K*>R-COOH	Ion score	59.72
Protein	P53026	Confidence level	99.00%
Precursor m/z	957.03027	Identity Threshold	46.88
Charge	2+	Mass error	0.001646



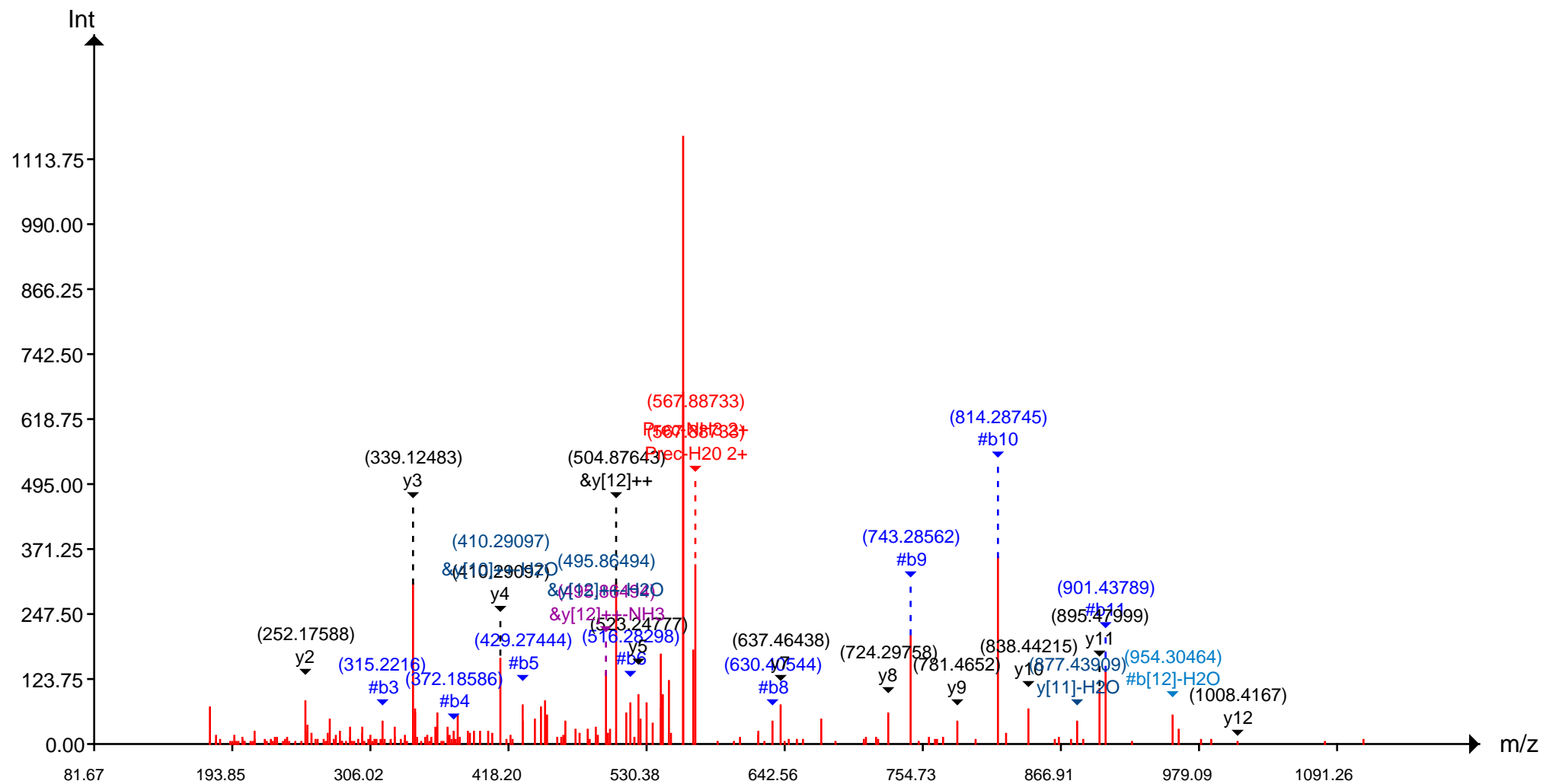
Info

Modified Sequence	AcD3-IAIK<AcD3K*>ESAK<AcD3K*>VGDQAQR-COOH	Ion score	53.52
Protein	Q99JY9	Confidence level	99.00%
Precursor m/z	874.99721	Identity Threshold	46.22
Charge	2+	Mass error	0.002039



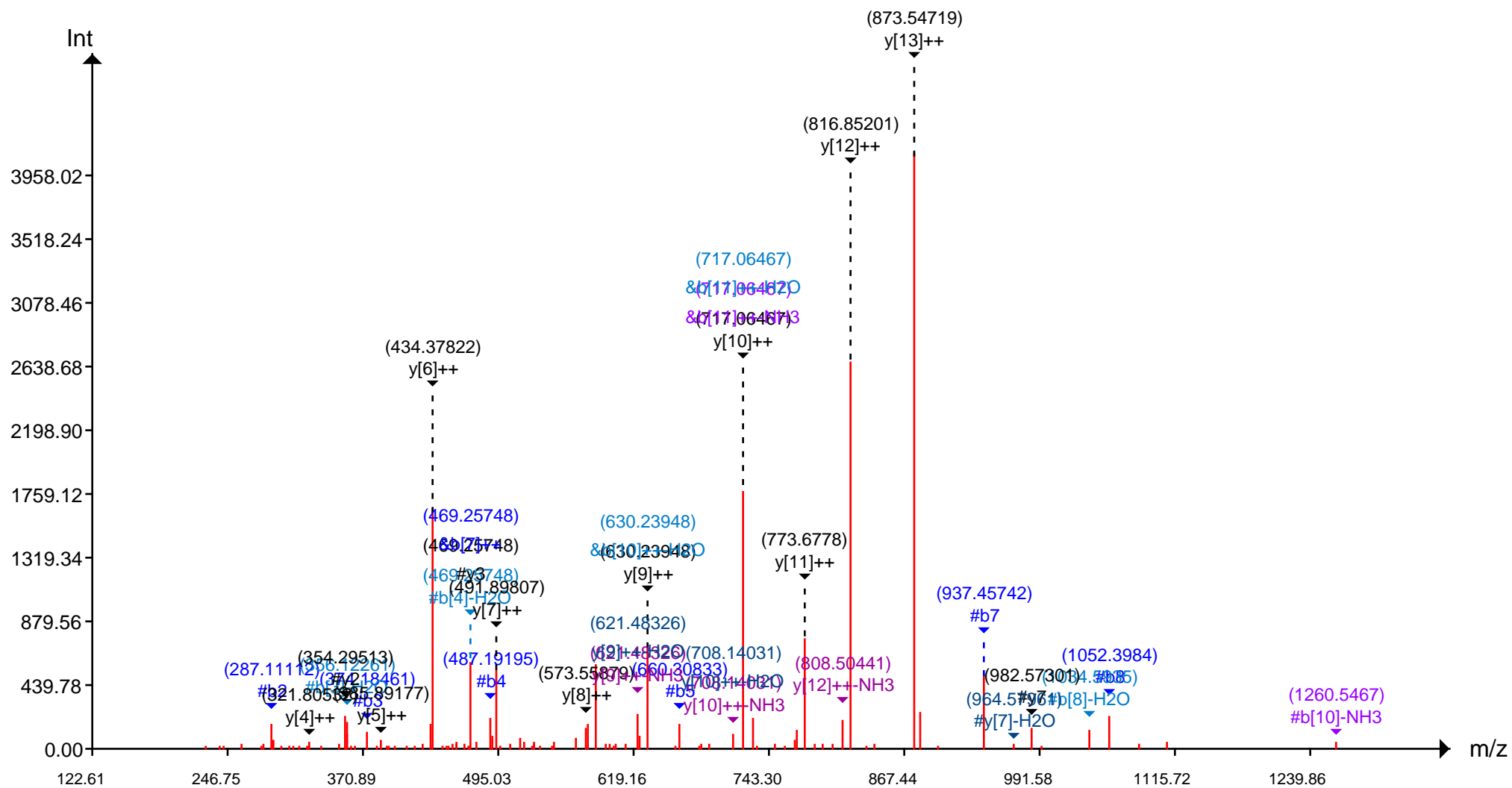
Info

Modified Sequence	AcD3-AM<Mox*>LK<AcD3K*>TPAQFDAELR<C13>-COOH	Ion score	57.4
Protein	P10107	Confidence level	99.00%
Precursor m/z	909.47056	Identity Threshold	48.09
Charge	2+	Mass error	0.002236



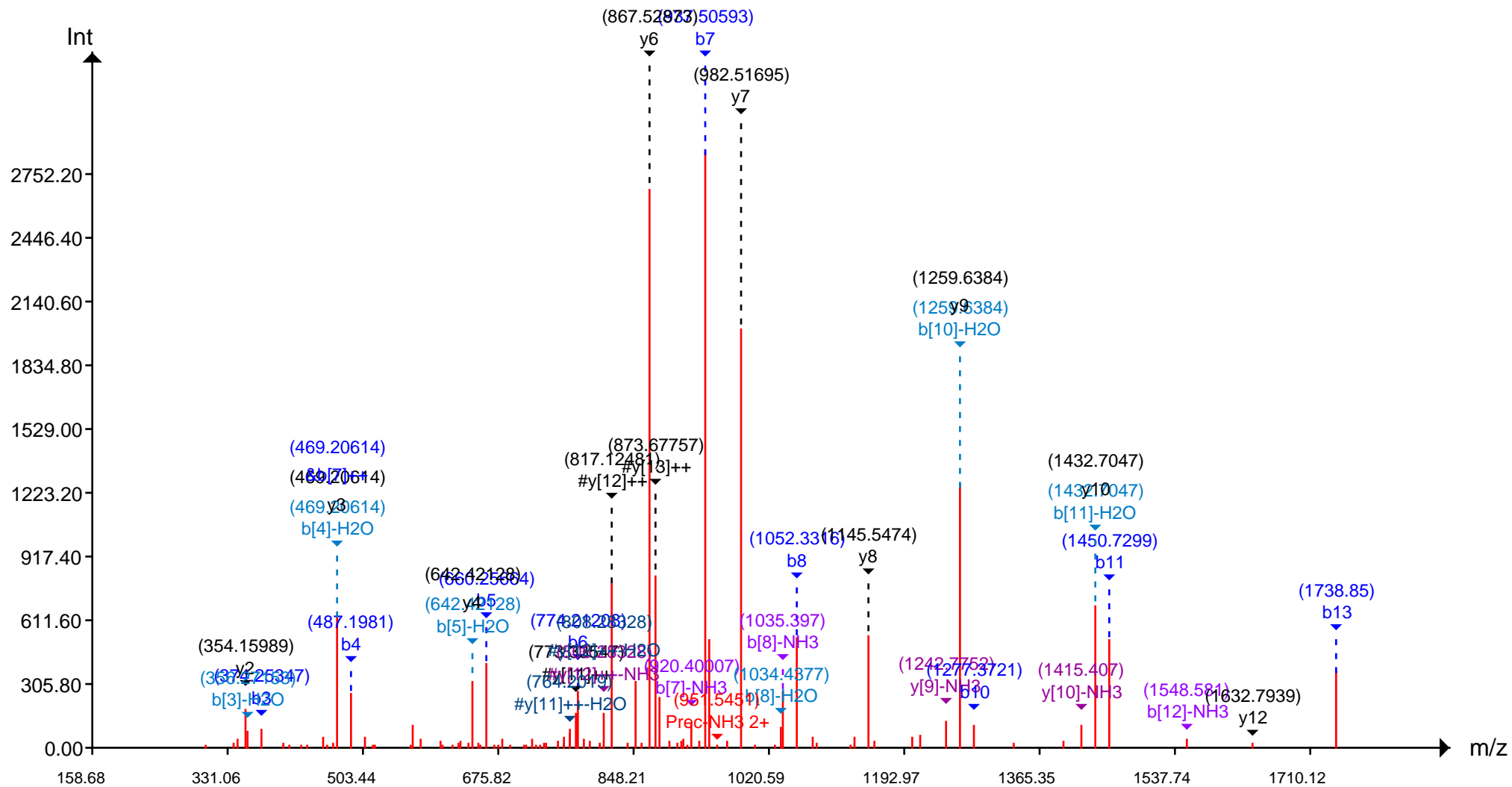
Info

Modified Sequence	AcD3-VIGGGSGGLASAR<C13>-COOH	Ion score	60.73
Protein	P47791	Confidence level	99.00%
Precursor m/z	576.82884	Identity Threshold	47.38
Charge	2+	Mass error	-2.28E-4



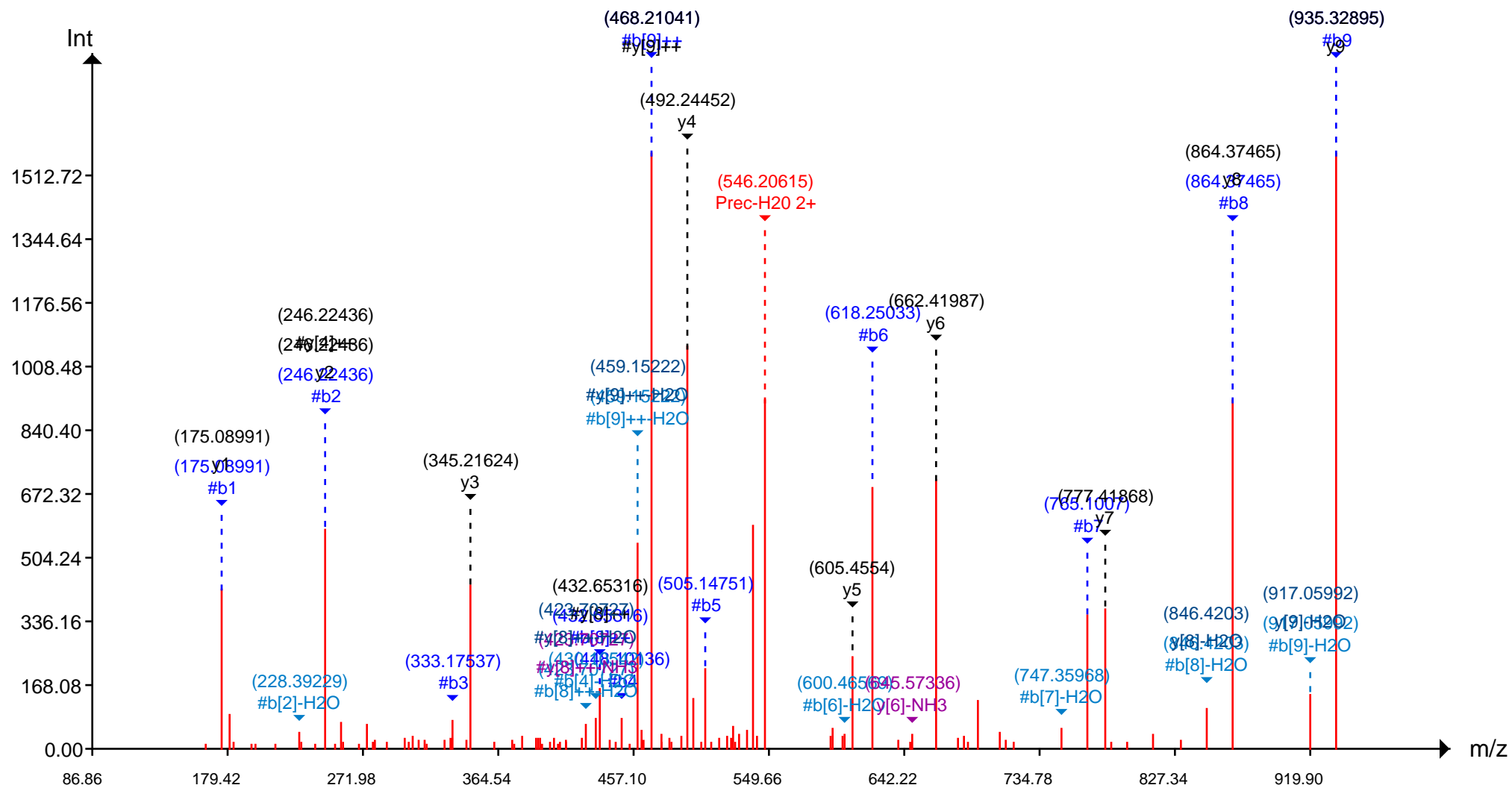
Info

Modified Sequence	AcD3-QISLK<AcD3K*>NYDPQK<AcD3K*>DK<AcD3K*>R<C13>-COOH	Ion score	62.11
Protein	P53026	Confidence level	99.00%
Precursor m/z	640.36203	Identity Threshold	46.36
Charge	3+	Mass error	-2.22E-4



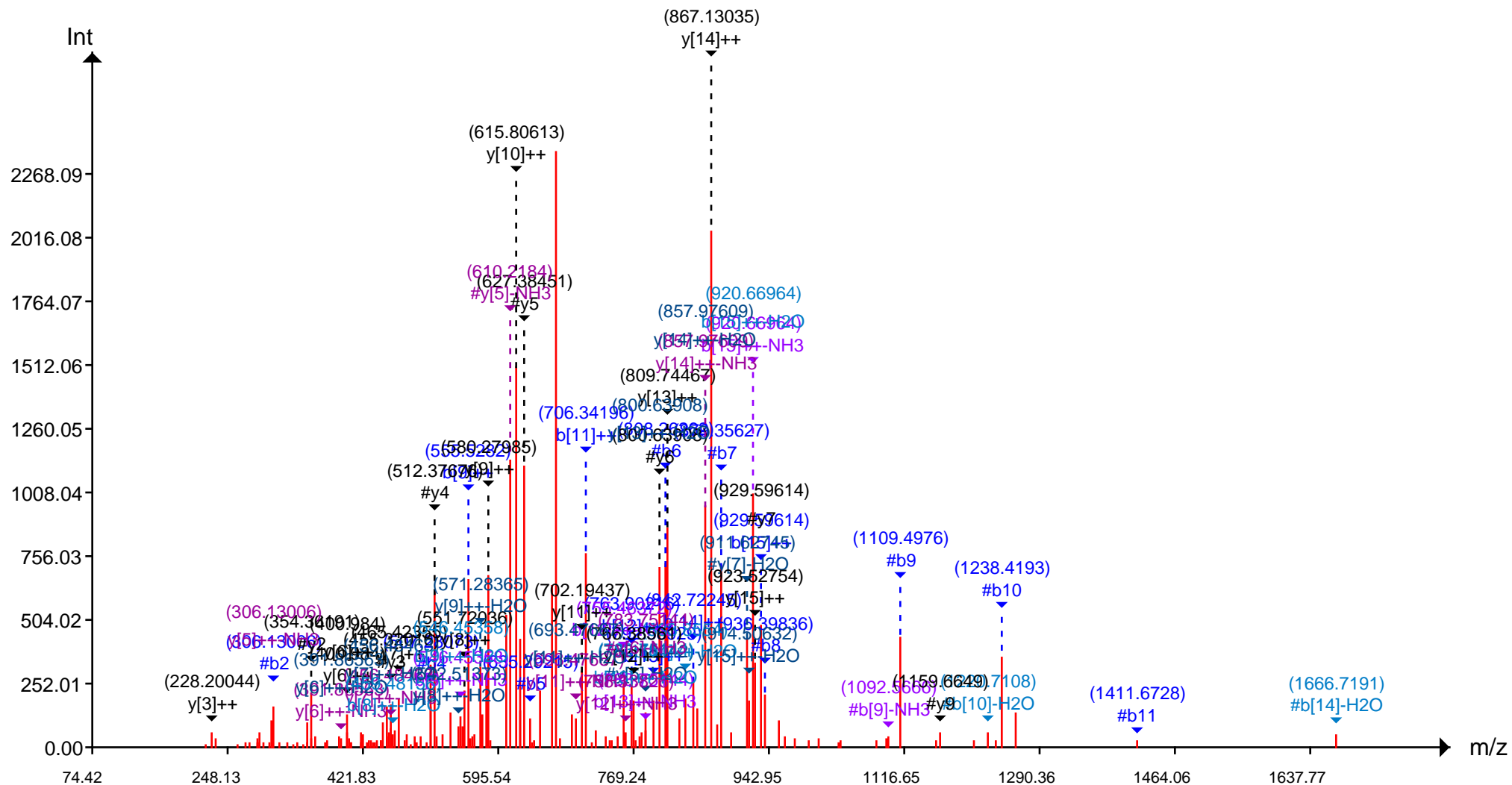
Info

Modified Sequence	AcD3-QISLK<AcD3K*>NYDPQK<AcD3K*>DK<AcD3K*>R<C13>-COOH	Ion score	84.96
Protein	P53026	Confidence level	99.00%
Precursor m/z	960.04011	Identity Threshold	46.32
Charge	2+	Mass error	0.001184



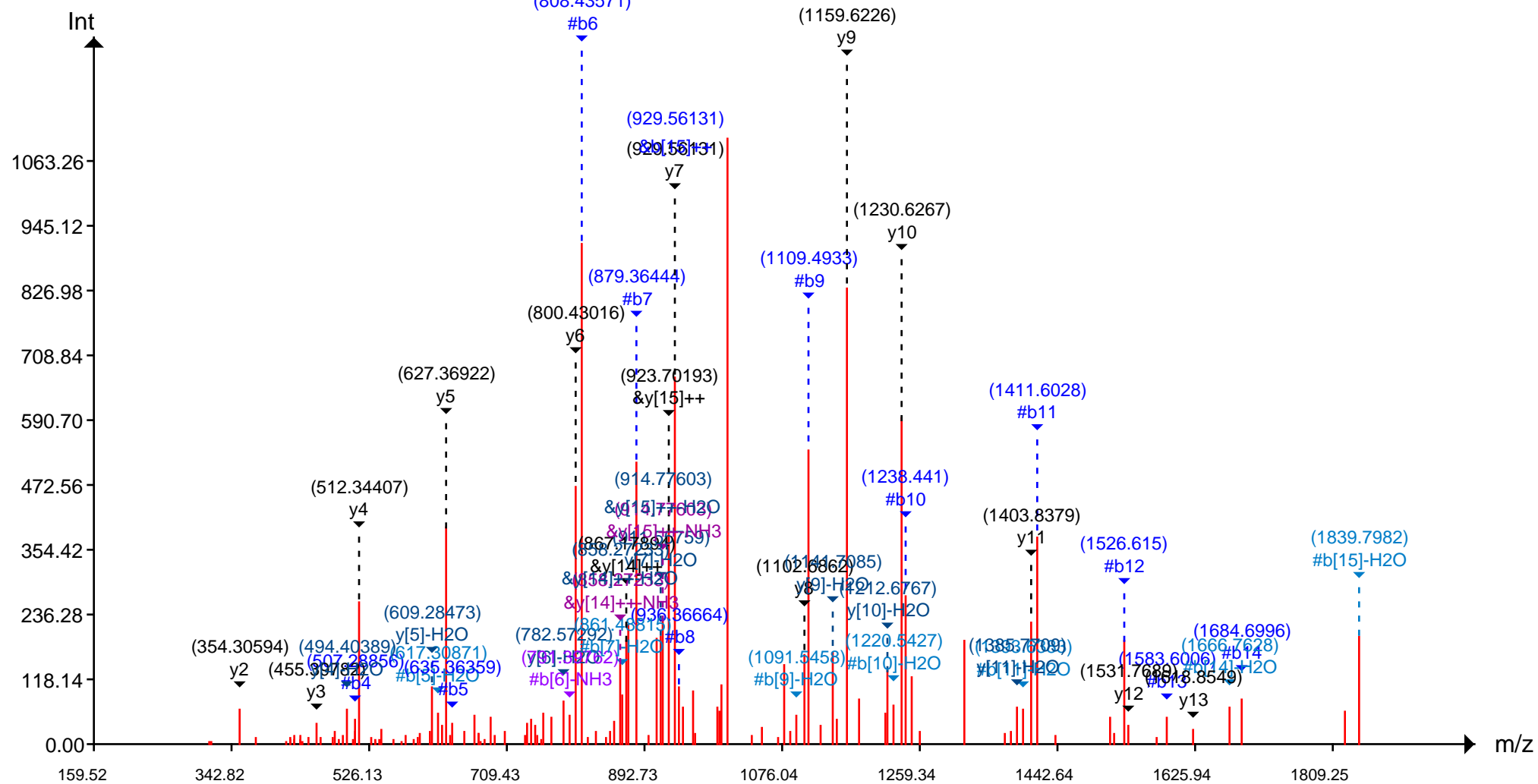
Info

Modified Sequence	AcD3-EASDGIM<Mox*>VAR-COOH	Ion score	93.33
Protein	P52480	Confidence level	99.00%
Precursor m/z	555.27135	Identity Threshold	46.64
Charge	2+	Mass error	0.001964



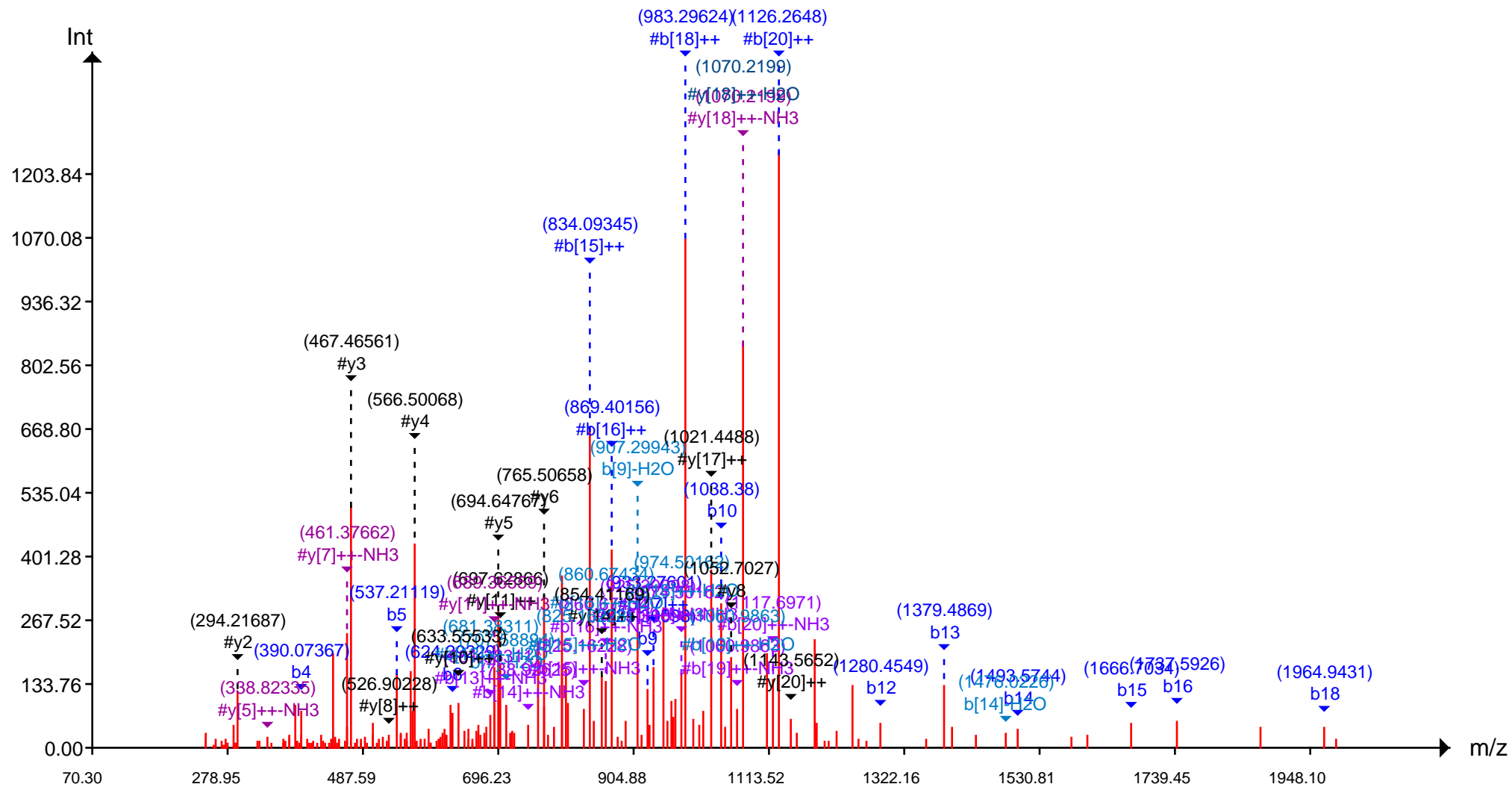
Info

Modified Sequence	AcD3-FLNSQ<AcD3K*>AGK<AcD3K*>EK<AcD3K*>DGTK<AcD3K*>R<C13>-COOH	Ion score	63.81
Protein	P23198	Confidence level	99.00%
Precursor m/z	680.05437	Identity Threshold	45.55
Charge	3+	Mass error	-6.25E-4



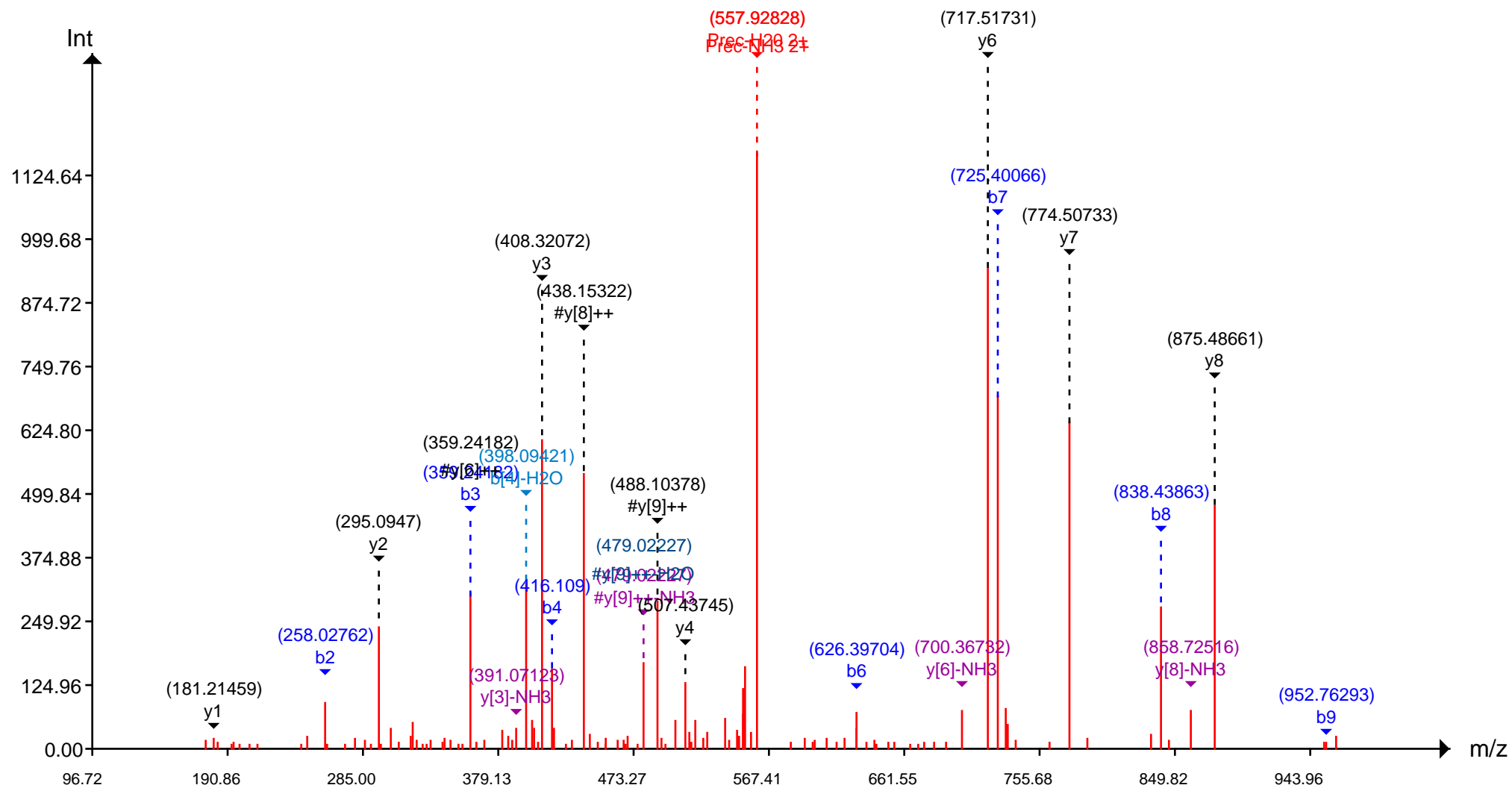
Info

Modified Sequence	AcD3-FLNSQK<AcD3K*>AGK<AcD3K*>EK<AcD3K*>DGTK<AcD3K*>R<C13>-COOH	Ion score	92.12
Protein	P23198	Confidence level	99.00%
Precursor m/z	1019.5795	Identity Threshold	45.45
Charge	2+	Mass error	0.002541



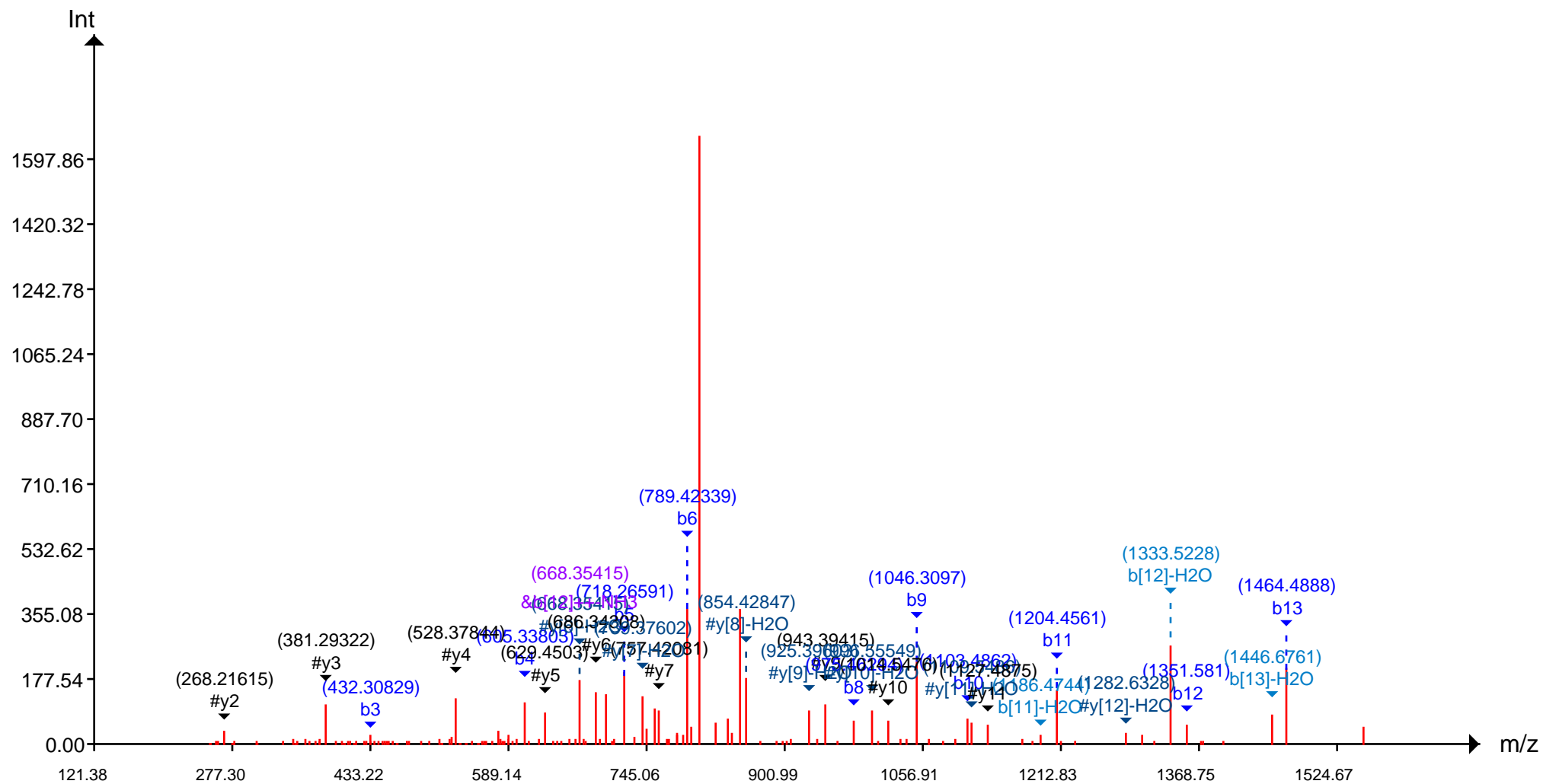
Info

Modified Sequence	AcD3-TAGDFSTTVIQNVNK<AcD3K*>AQVK<AcD3K*>IR<C13>-COOH	Ion score	49.4
Protein	P57746	Confidence level	99.00%
Precursor m/z	811.45983	Identity Threshold	45.14
Charge	3+	Mass error	1.00523



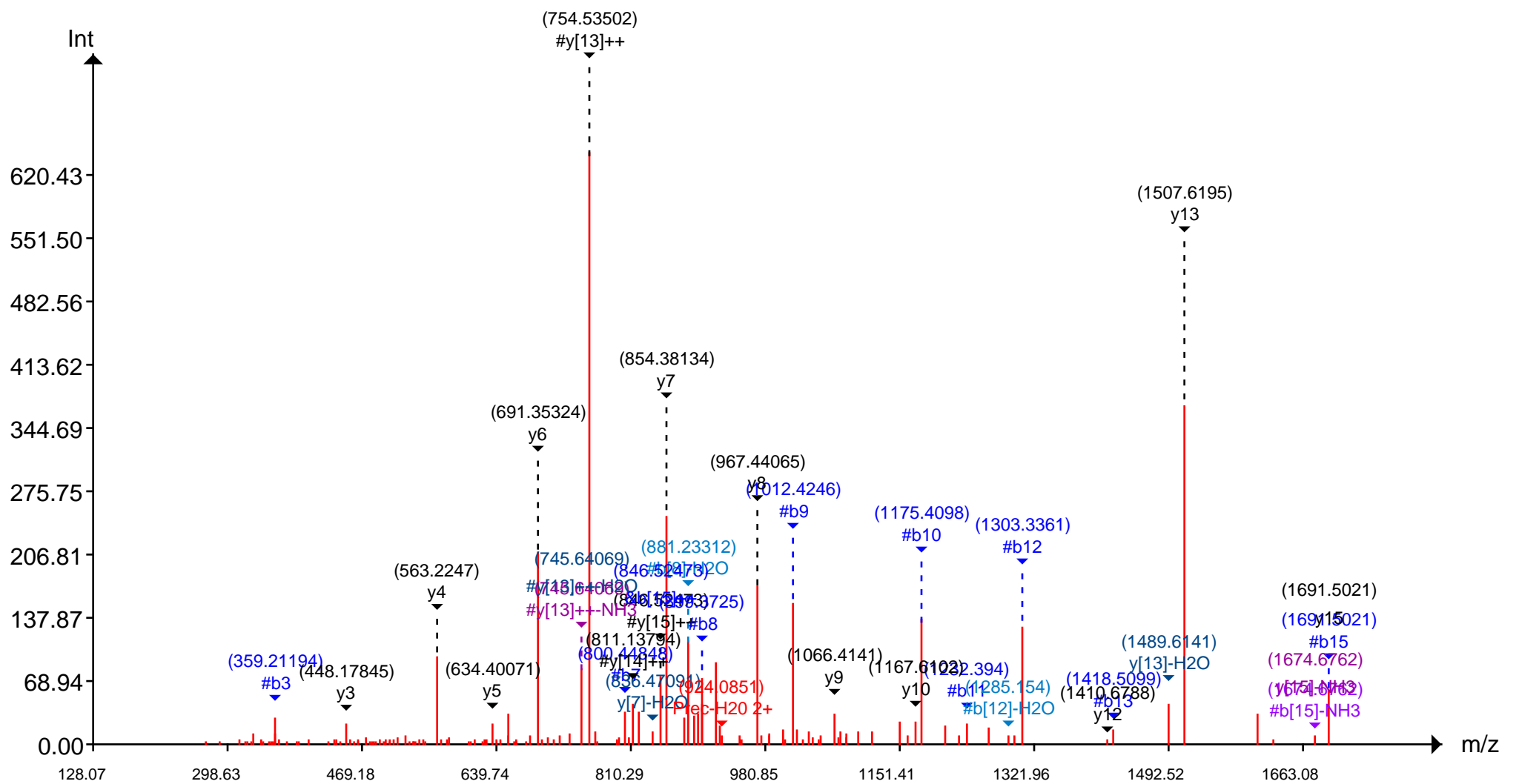
Info

Modified Sequence	AcD3-LVTGPLVINR<C13>-COOH	Ion score	48.97
Protein	P47911	Confidence level	99.00%
Precursor m/z	566.86478	Identity Threshold	40.30
Charge	2+	Mass error	-4.9E-5

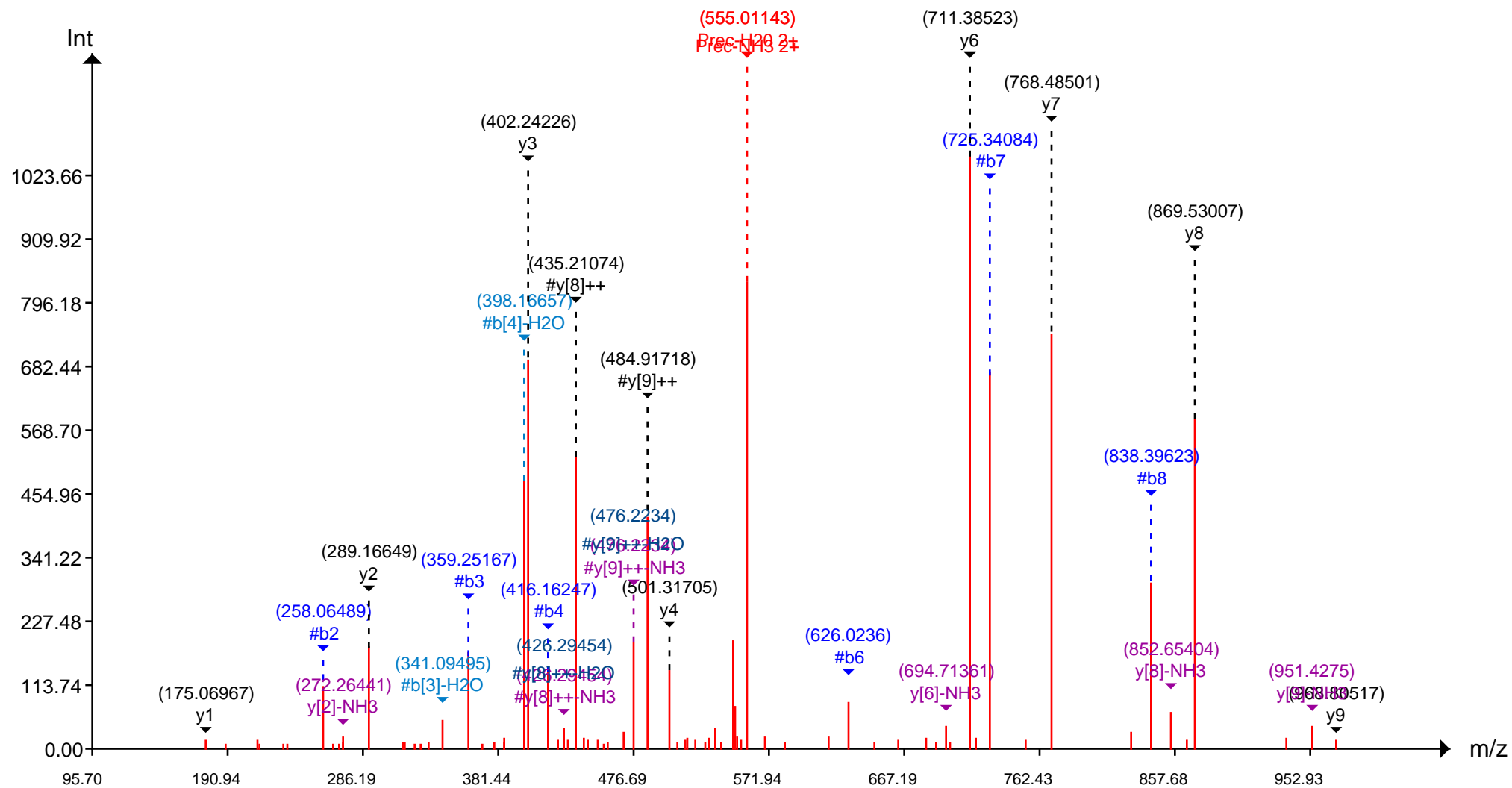


Info

Modified Sequence	AcD3-NVK<AcD3K*>K<AcD3K*>LAADAGTFLSR<C13>-COOH	Ion score	76.35
Protein	Q9JK48	Confidence level	99.00%
Precursor m/z	866.50706	Identity Threshold	44.54
Charge	2+	Mass error	0.002315

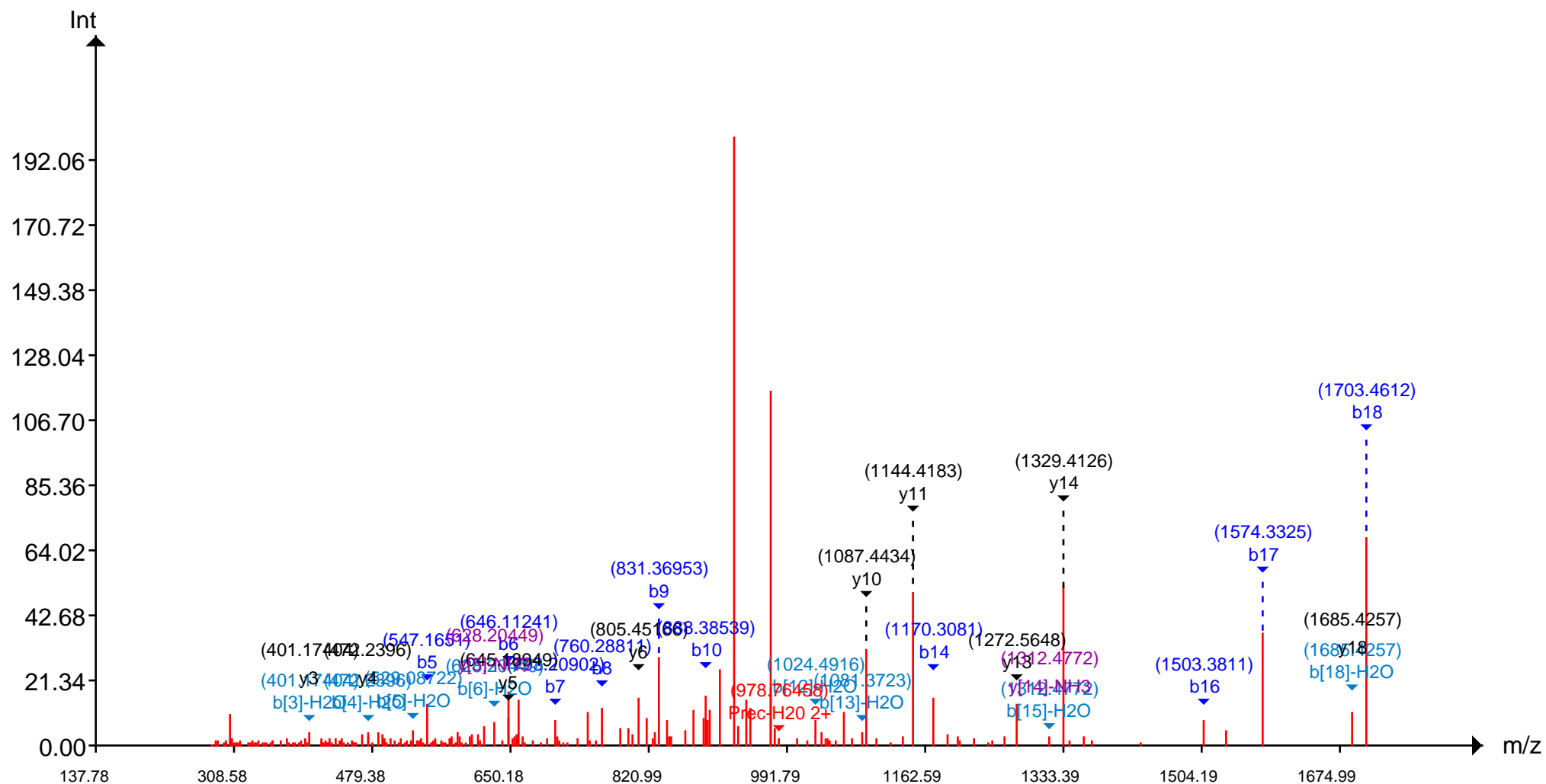


Info			
Modified Sequence	AcD3-EAIPENTVIYGADC<Cmm*>LR-COOH	Ion score	93.24
Protein	Q9WUB4	Confidence level	99.00%
Precursor m/z	933.46268	Identity Threshold	47.50
Charge	2+	Mass error	0.003993



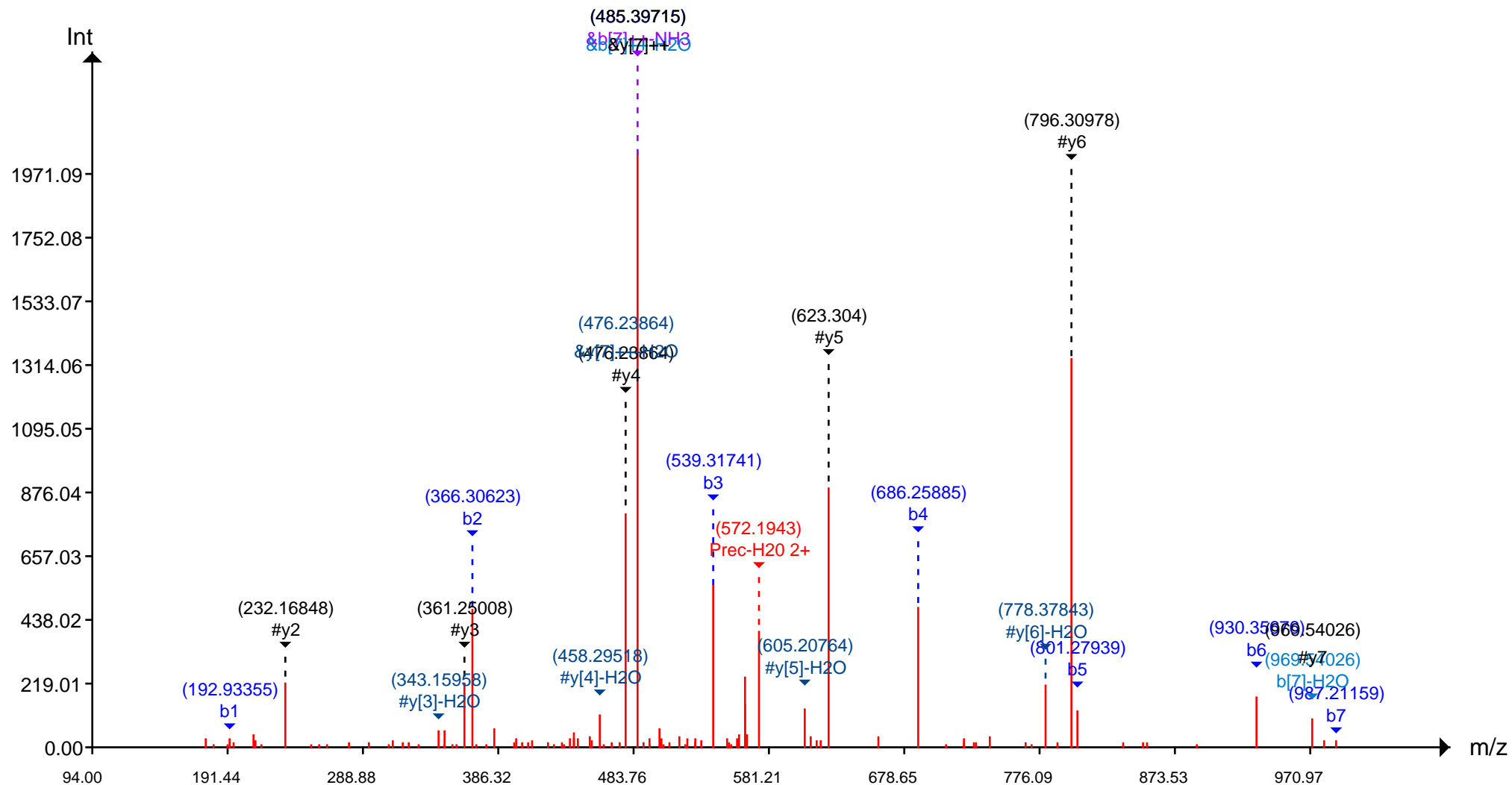
Info

Modified Sequence	AcD3-LVTGPLVINR-COOH	Ion score	58.66
Protein	P47911	Confidence level	99.00%
Precursor m/z	563.8548	Identity Threshold	41.60
Charge	2+	Mass error	1.32E-4



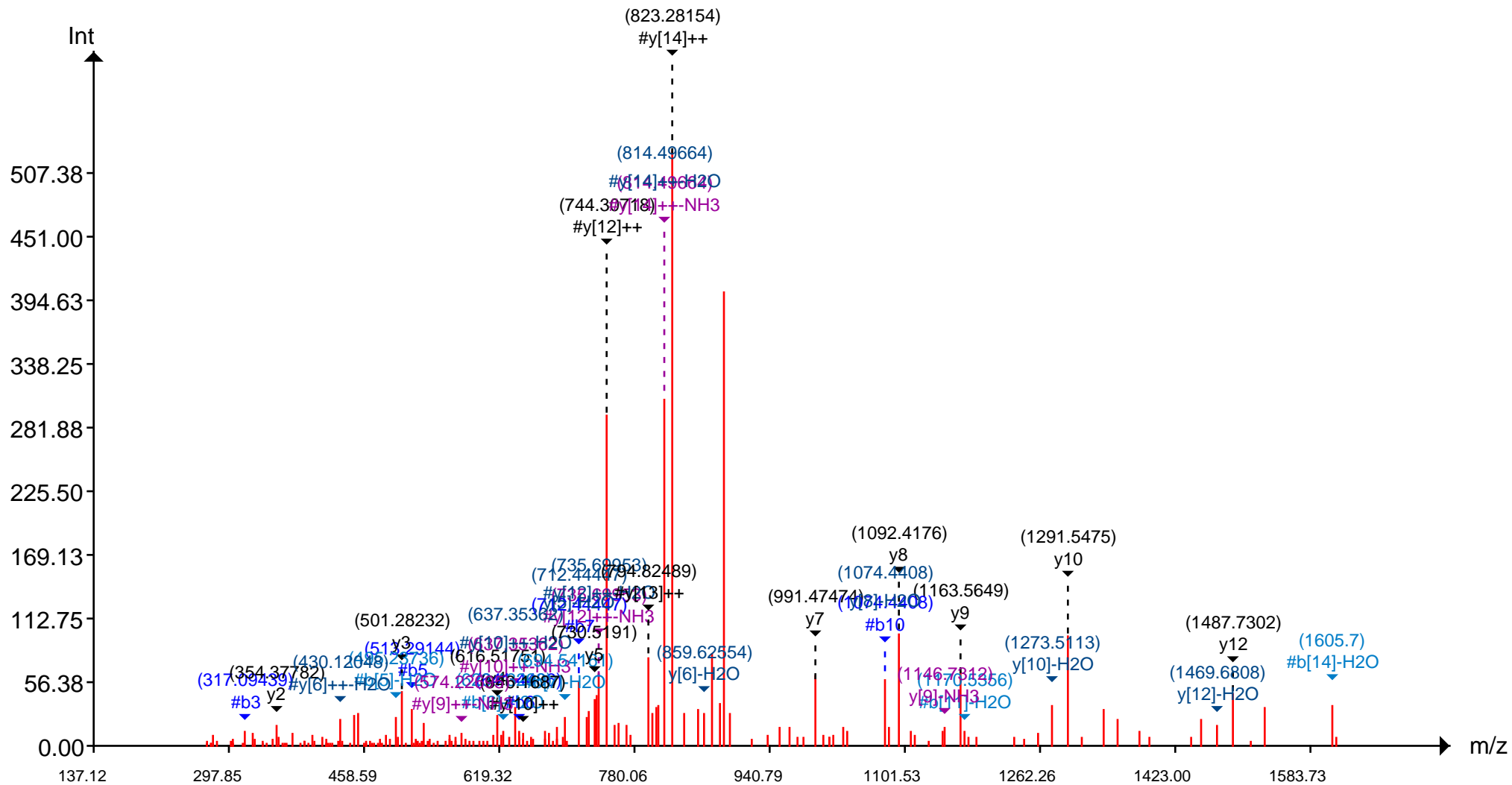
Info

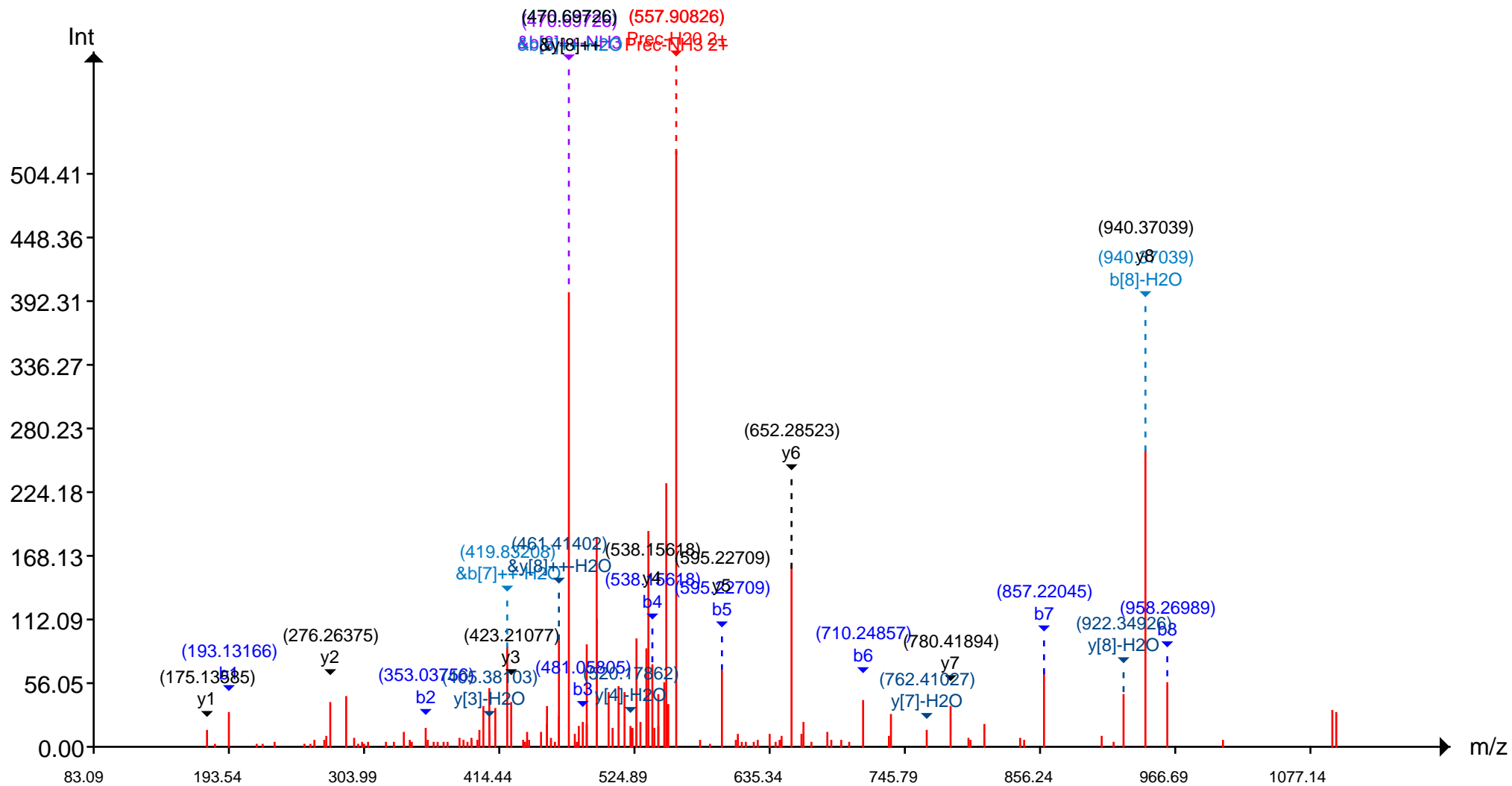
Modified Sequence	AcD3-EDEAGVGGAGPGGAC<Cmm*>K<AcD3K*>AEPR-COOH	Ion score	60.52
Protein	Q00P19	Confidence level	99.00%
Precursor m/z	987.95948	Identity Threshold	45.27
Charge	2+	Mass error	0.00243



Info

Modified Sequence	AcD3-FK<AcD3K*>K<AcD3K*>FDEGR-COOH	Ion score	52.03
Protein	P09103	Confidence level	99.00%
Precursor m/z	581.31601	Identity Threshold	47.73
Charge	2+	Mass error	-1.47E-4





Info

Modified Sequence AcD3-M<Mox*>C<Cmm*>QGGDFTR-COOH

Ion score 47.91

Protein P17742

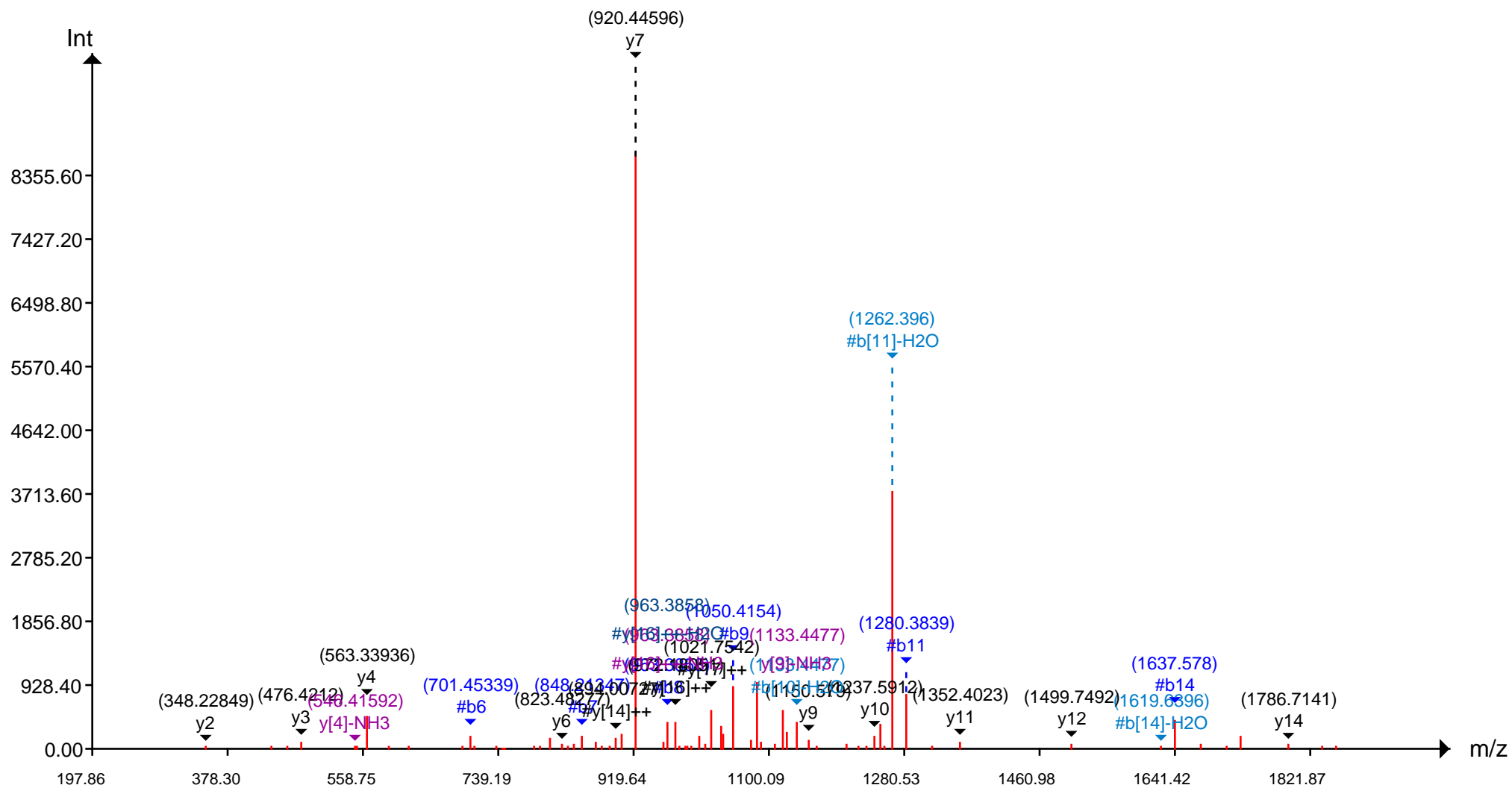
Confidence level 99.00%

Precursor m/z 566.73294

Identity Threshold 39.19

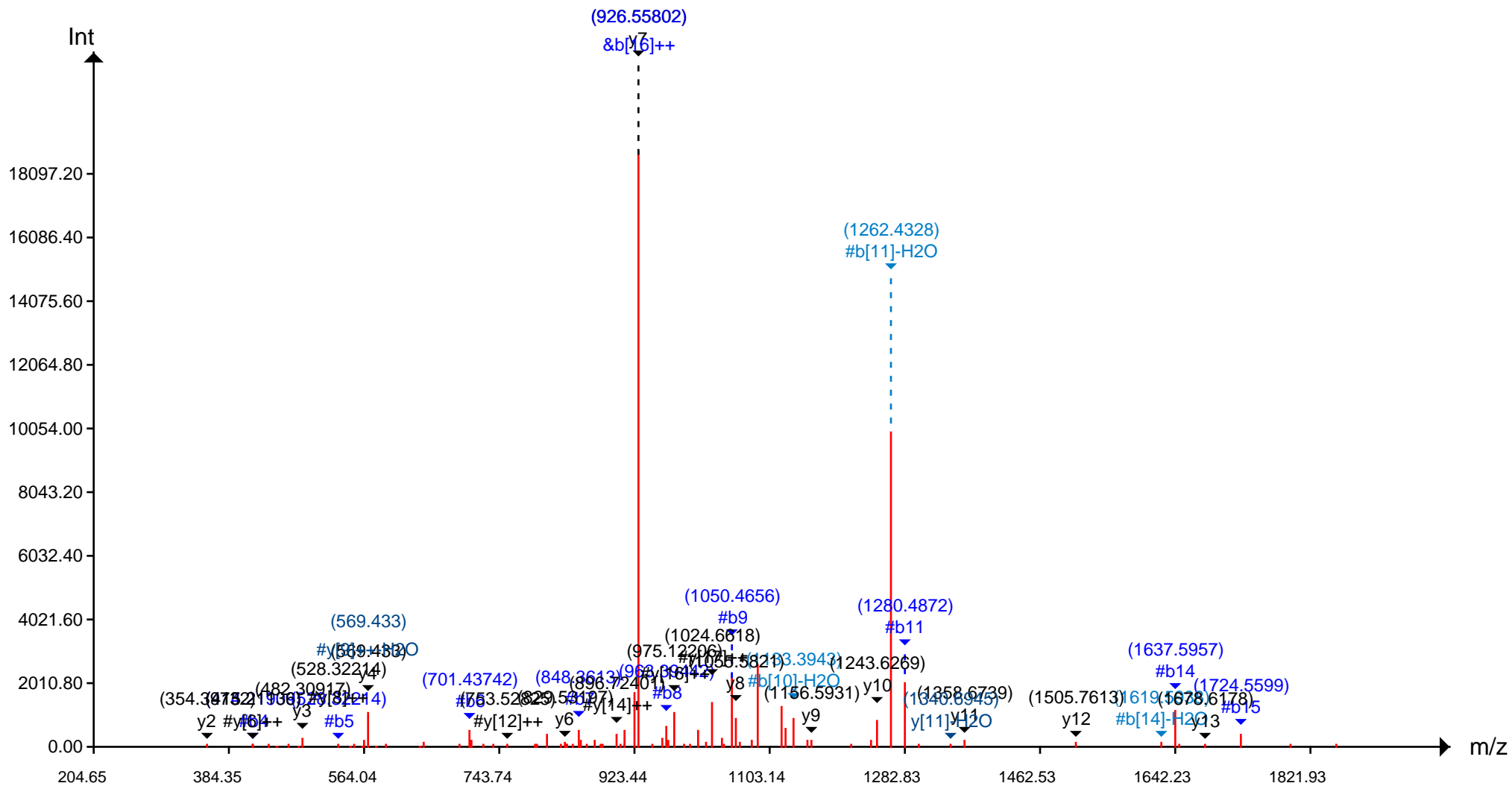
Charge 2+

Mass error -3.47E-4



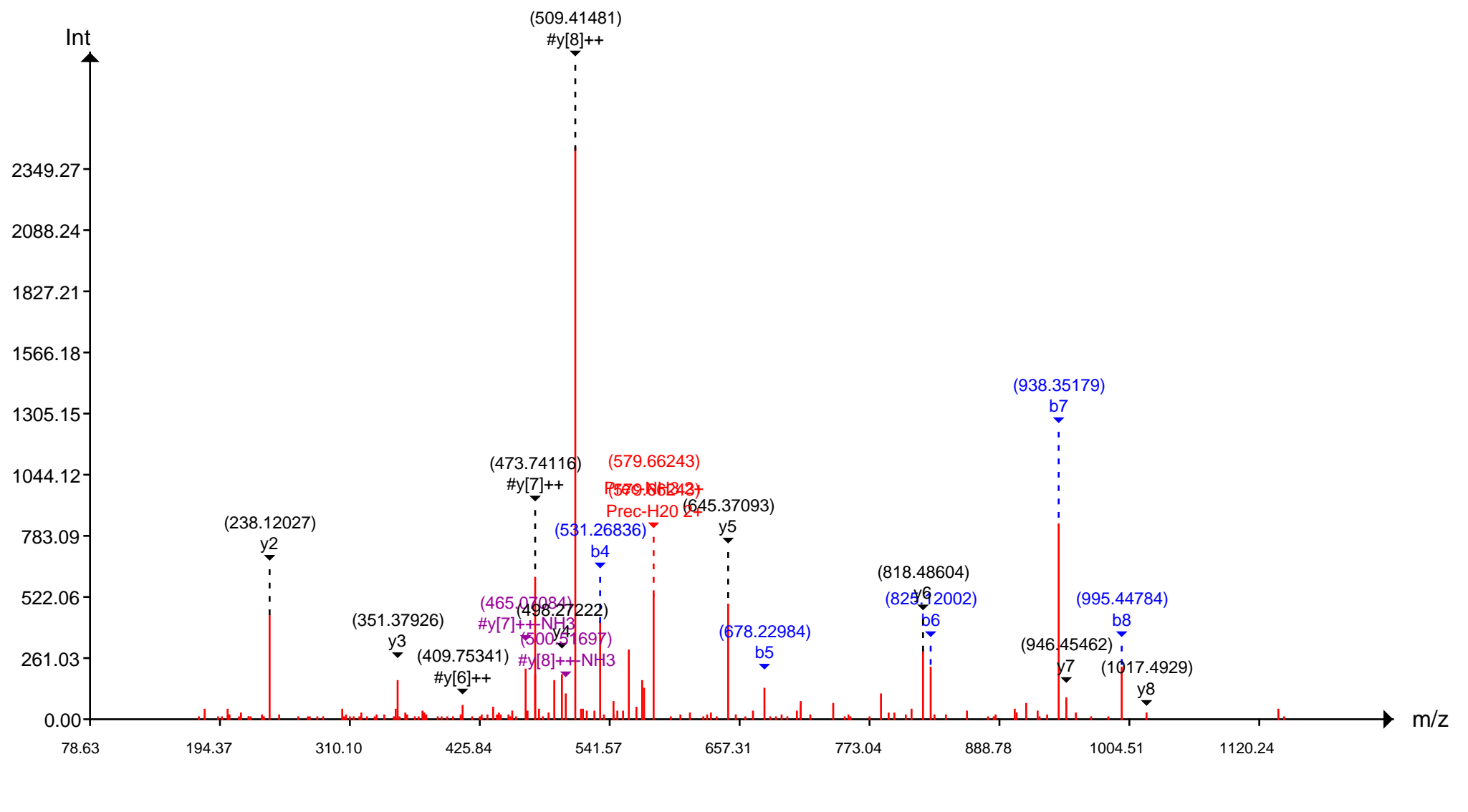
Info

Modified Sequence	AcD3-IVGVNK<AcD3K*>M<Mox*>DSTEPPYSQK<AcD3K*>R-COOH	Ion score	64.56
Protein	P10126	Confidence level	99.00%
Precursor m/z	1100.5681	Identity Threshold	47.95
Charge	2+	Mass error	0.002477

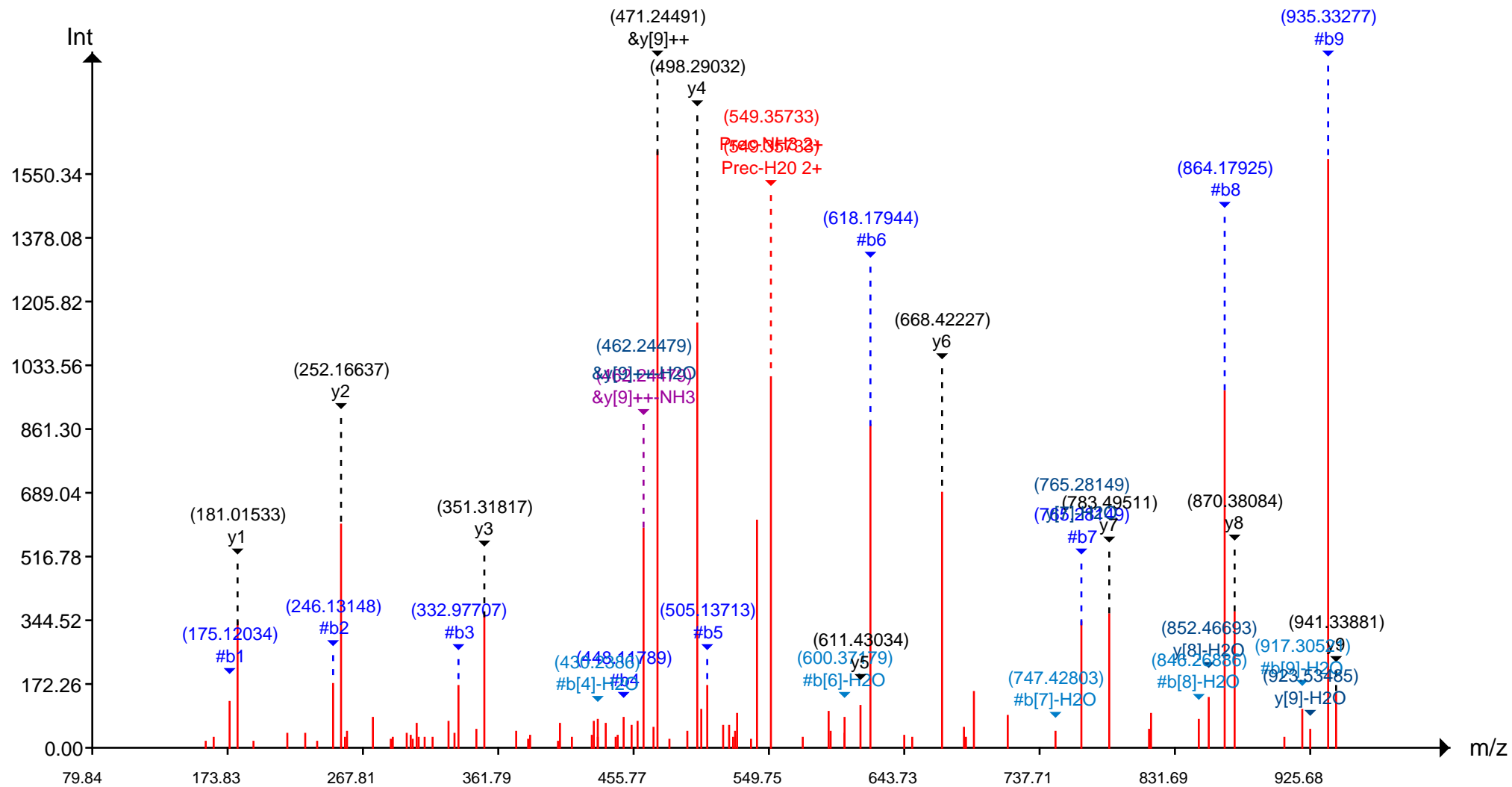


Info

Modified Sequence	AcD3-IVGVNK<AcD3K*>M<Mox*>DSTEPPYSQK<AcD3K*>R<C13>-COOH	Ion score	70.55
Protein	P10126	Confidence level	99.00%
Precursor m/z	1103.578	Identity Threshold	47.98
Charge	2+	Mass error	0.002135

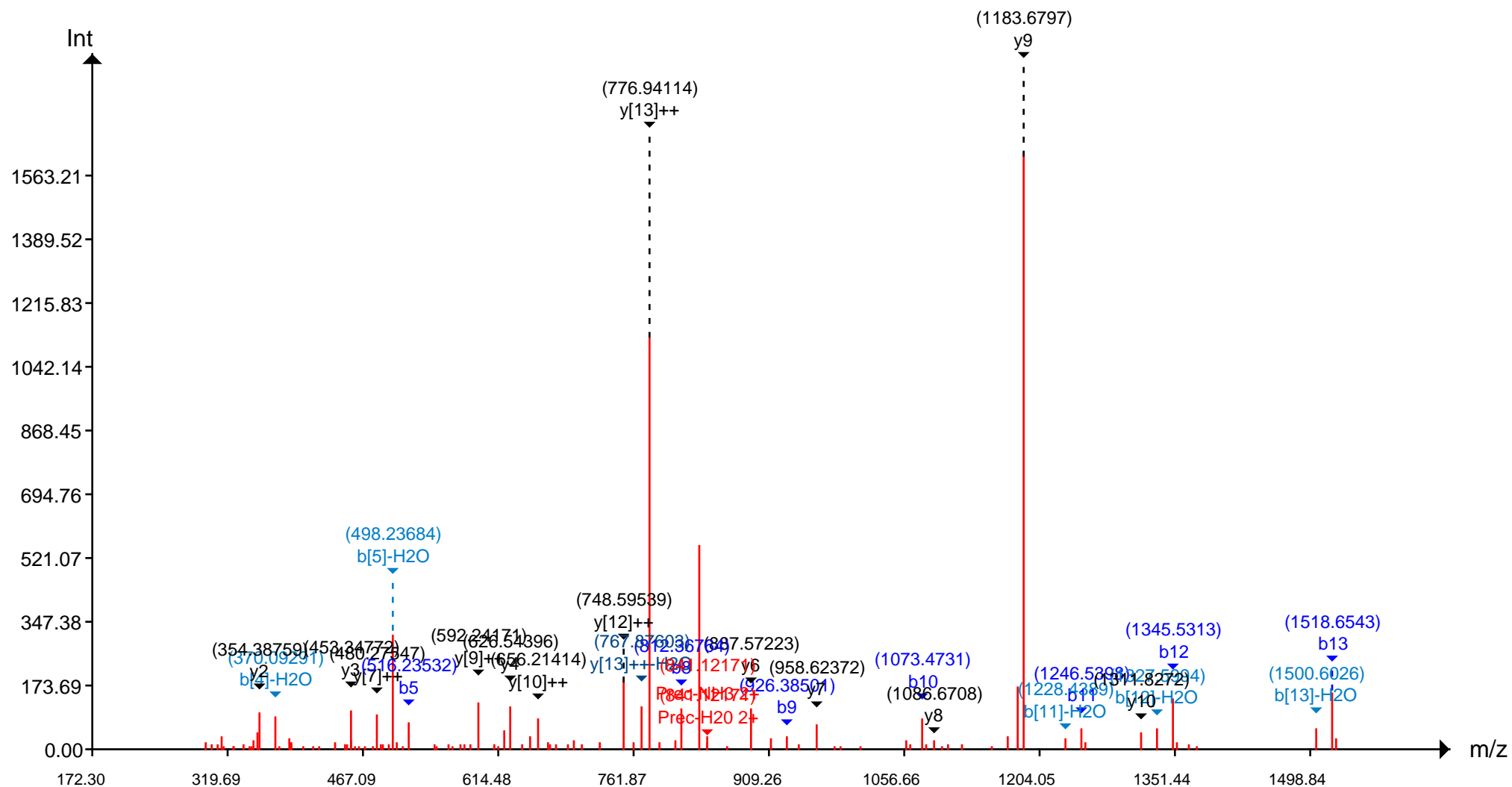


Info			
Modified Sequence	AcD3-LAQK<AcD3K*>M<Mox*>M<Mox*>IGR<C13>-COOH	Ion score	51.37
Protein	P52480, P53657	Confidence level	99.00%
Precursor m/z	588.32818	Identity Threshold	47.61
Charge	2+	Mass error	1.76E-4



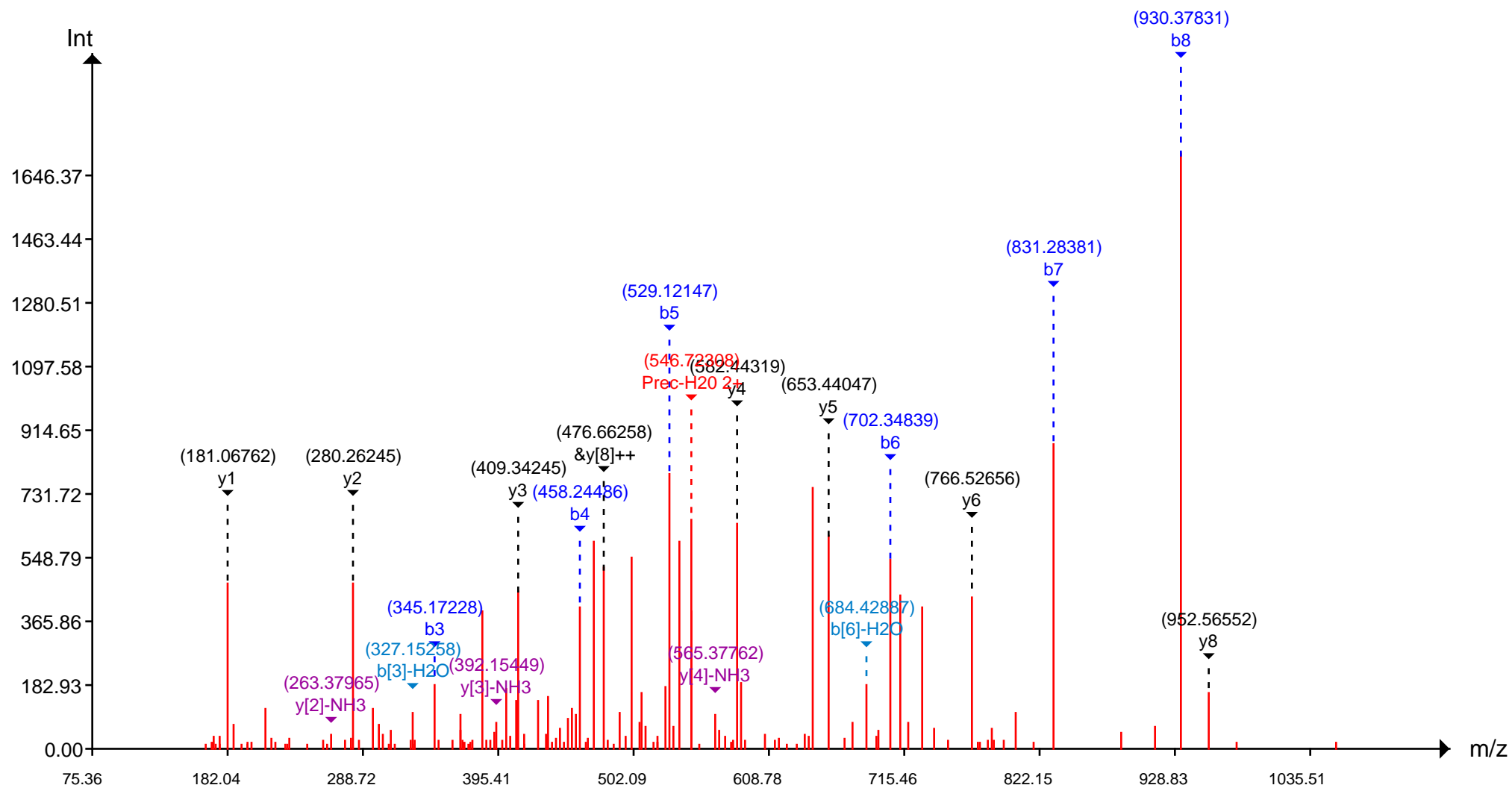
Info

Modified Sequence	AcD3-EASDGIM<Mox*>VAR<C13>-COOH	Ion score	78.26
Protein	P52480	Confidence level	99.00%
Precursor m/z	558.2813	Identity Threshold	47.14
Charge	2+	Mass error	0.001722



Info

Modified Sequence	AcD3-TGPSQPQANFK<AcD3K*>VK<AcD3K*>R<C13>-COOH	Ion score	78.35
Protein	Q9JHS9	Confidence level	99.00%
Precursor m/z	849.98346	Identity Threshold	46.25
Charge	2+	Mass error	0.001715



Info

Modified Sequence AcD3-IGELAK<AcD3K*>EVR<C13>-COOH

Ion score 49.09

Protein P97352

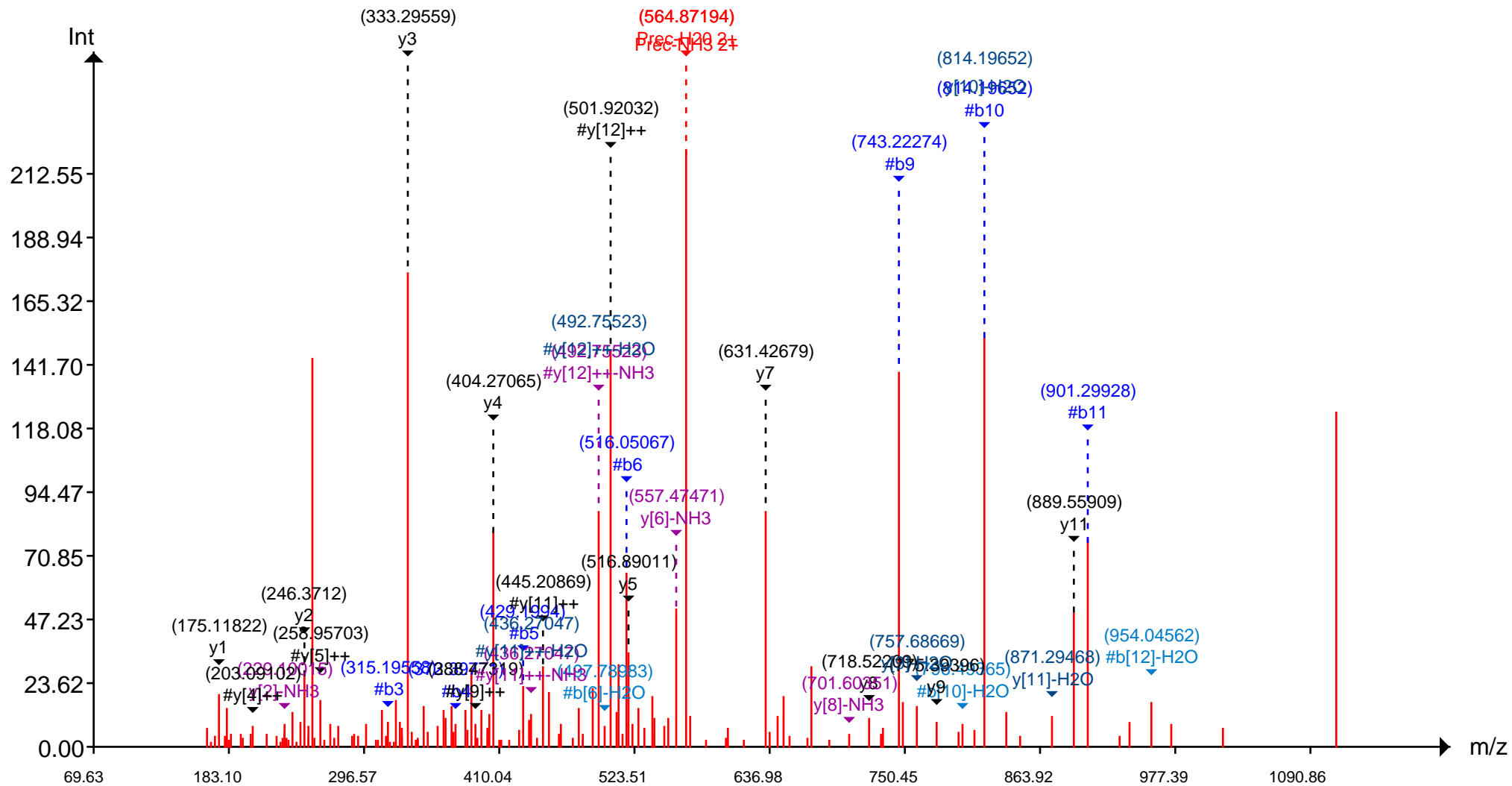
Confidence level 99.00%

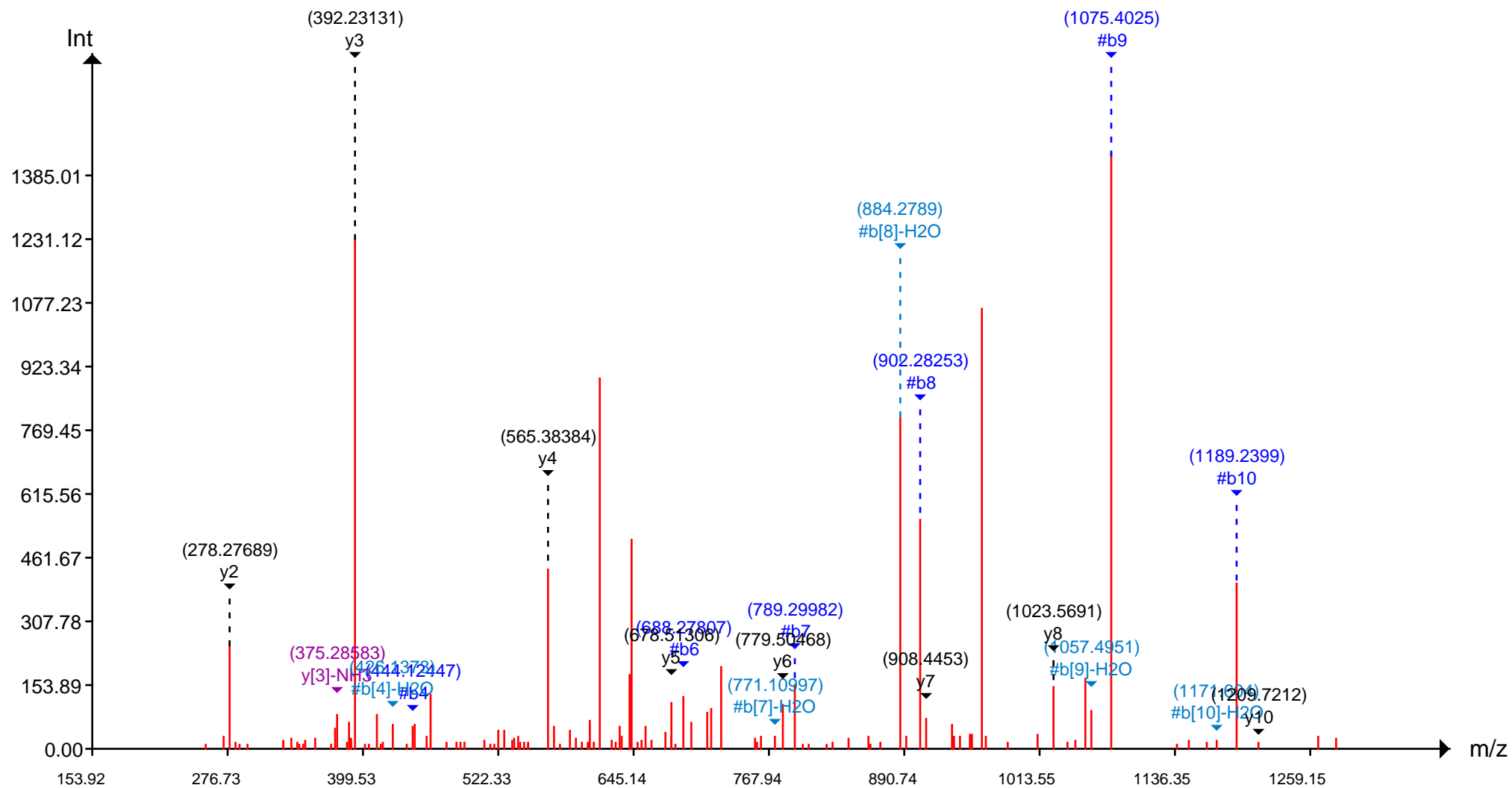
Precursor m/z 555.83992

Identity Threshold 44.15

Charge 2+

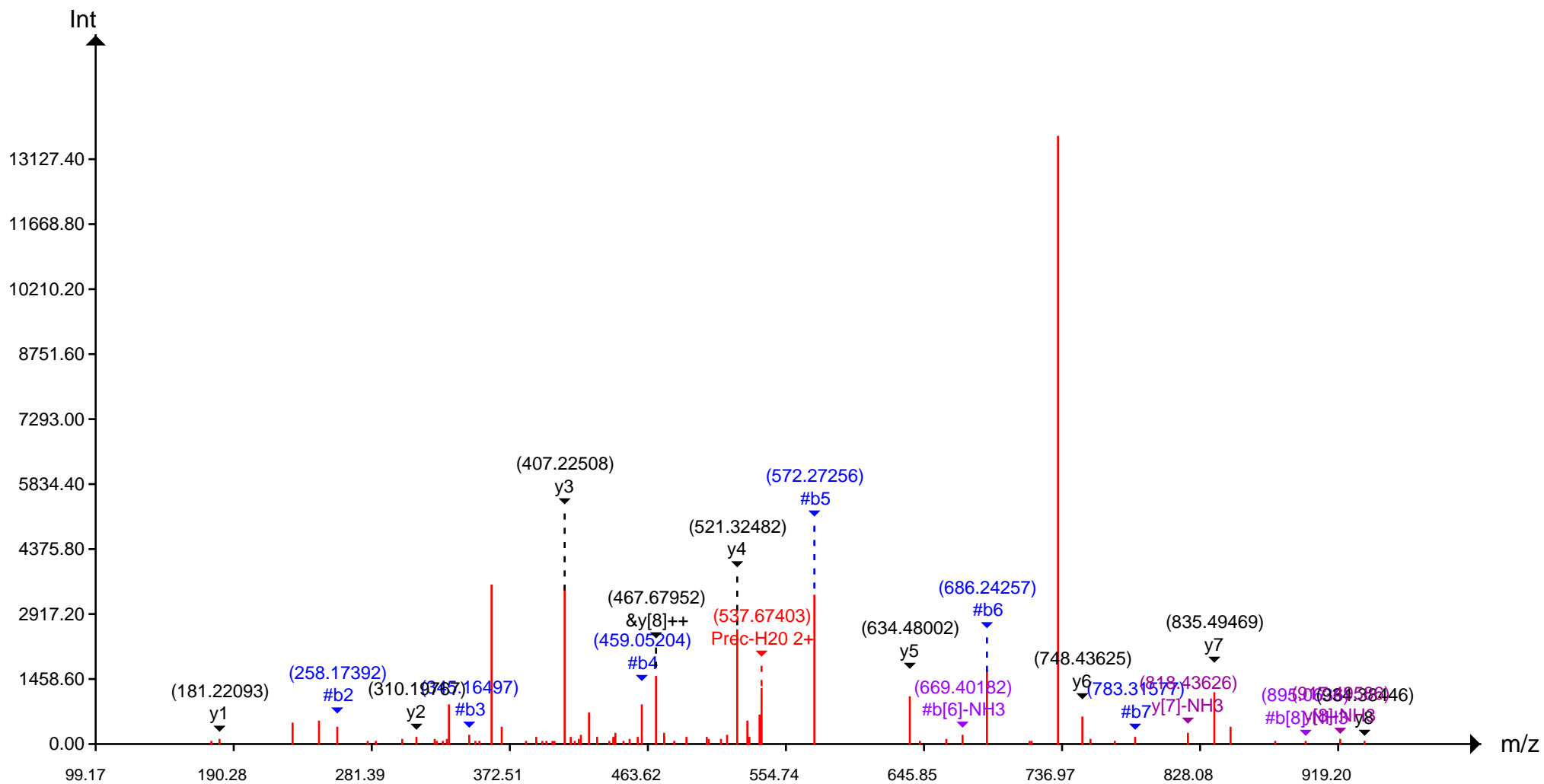
Mass error -5.6E-4





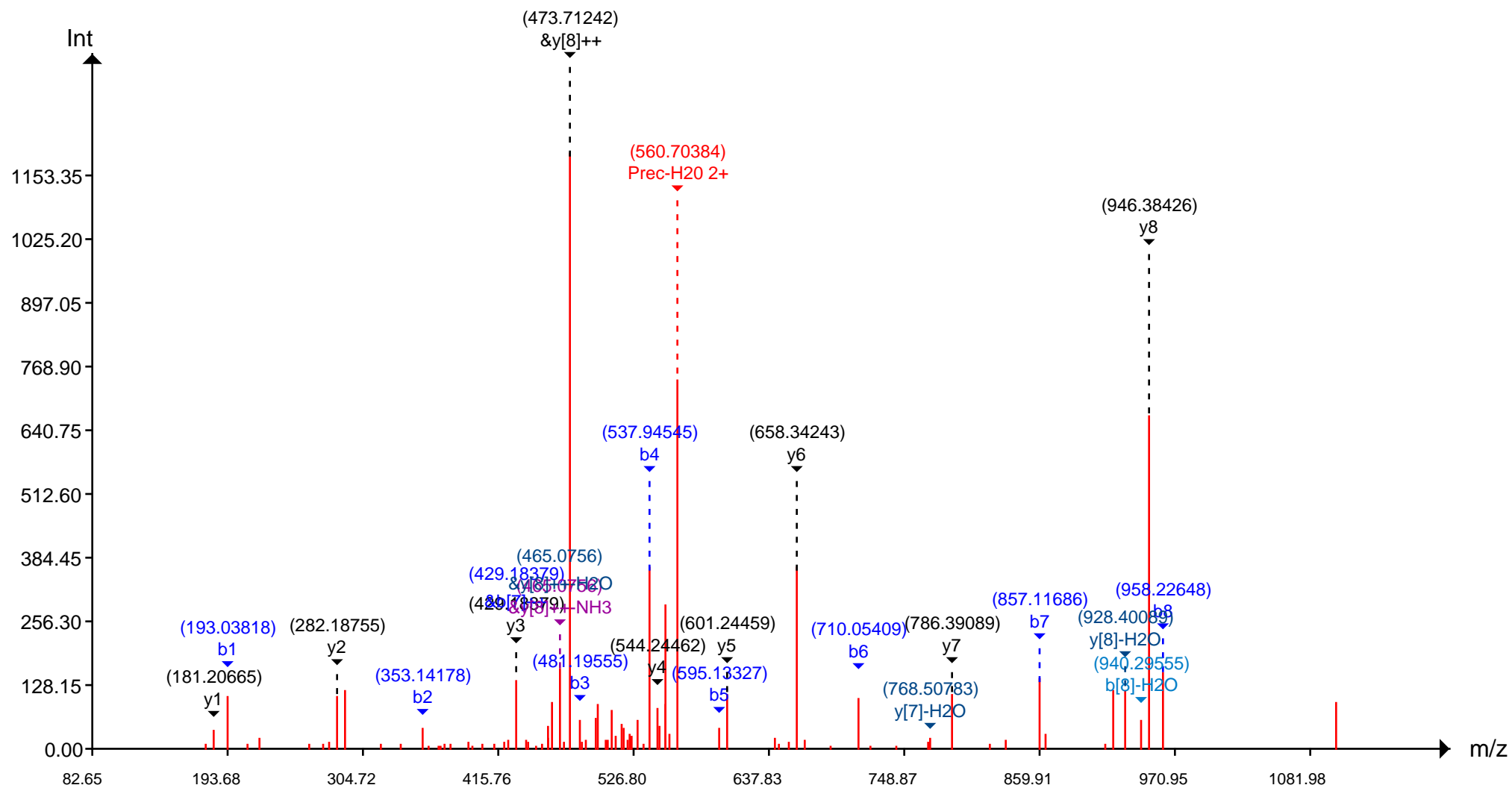
Info

Modified Sequence	AcD3-IVSVDETIK<AcD3K*>NPR<C13>-COOH	Ion score	48.76
Protein	P80313	Confidence level	99.00%
Precursor m/z	733.92568	Identity Threshold	45.78
Charge	2+	Mass error	0.001359



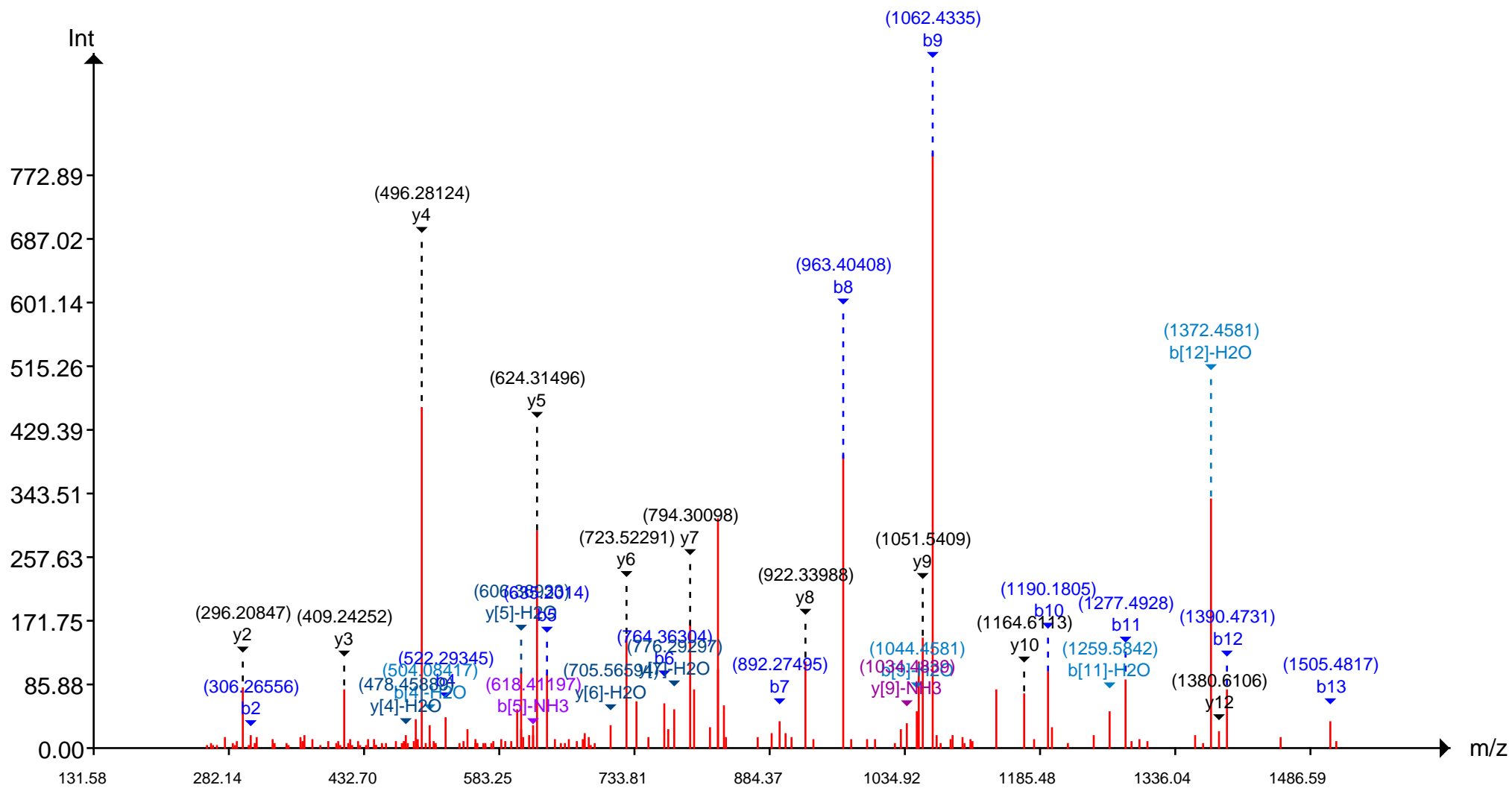
Info

Modified Sequence	AcD3-LVSNLNPER<C13>-COOH	Ion score	50.07
Protein	P17225	Confidence level	99.00%
Precursor m/z	546.81263	Identity Threshold	46.96
Charge	2+	Mass error	-2.54E-4



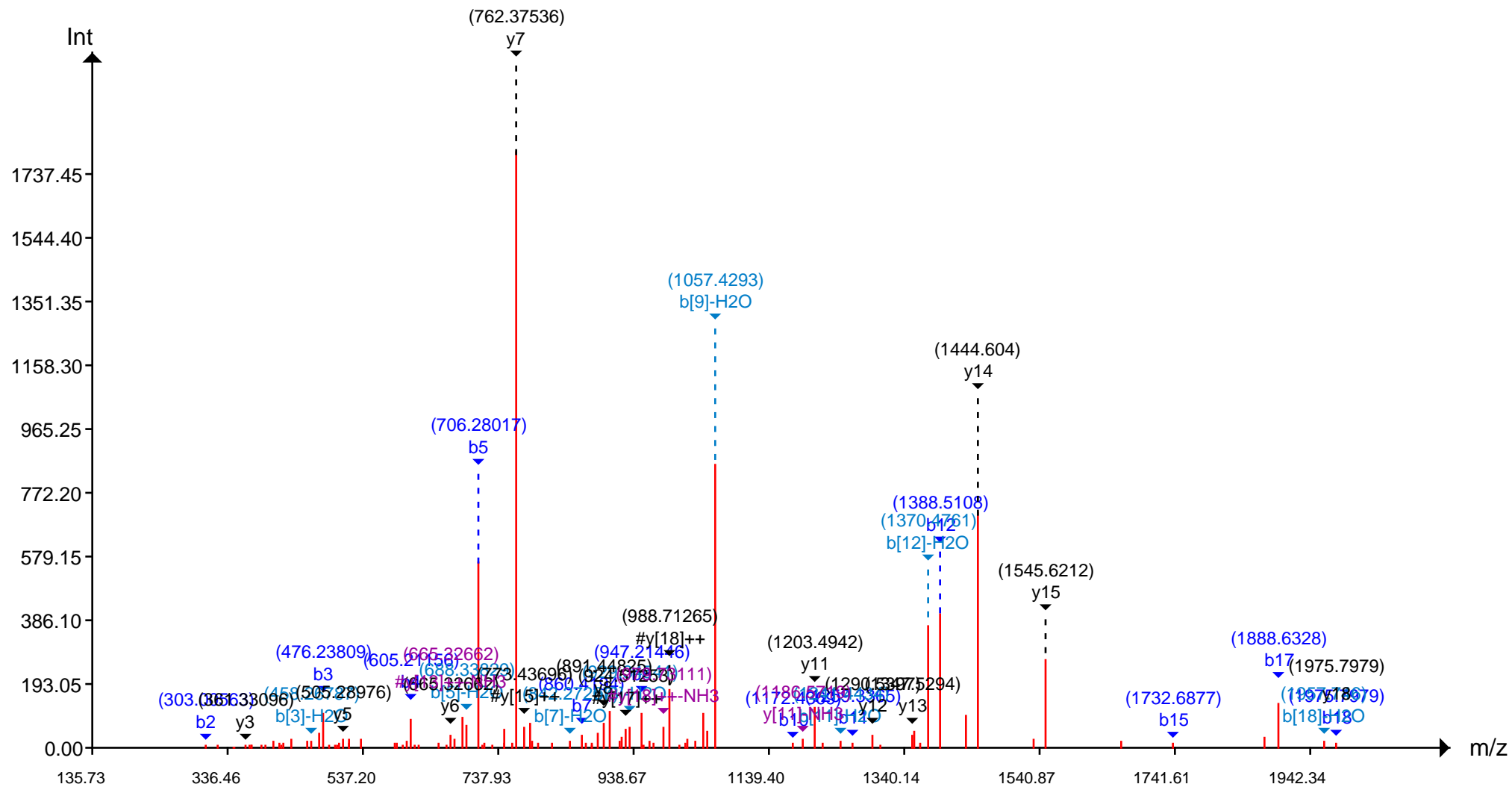
Info

Modified Sequence	AcD3-M<Mox*>C<Cmm*>QGGDFTR<C13>-COOH	Ion score	58.06
Protein	P17742	Confidence level	99.00%
Precursor m/z	569.74278	Identity Threshold	41.37
Charge	2+	Mass error	-8.09E-4



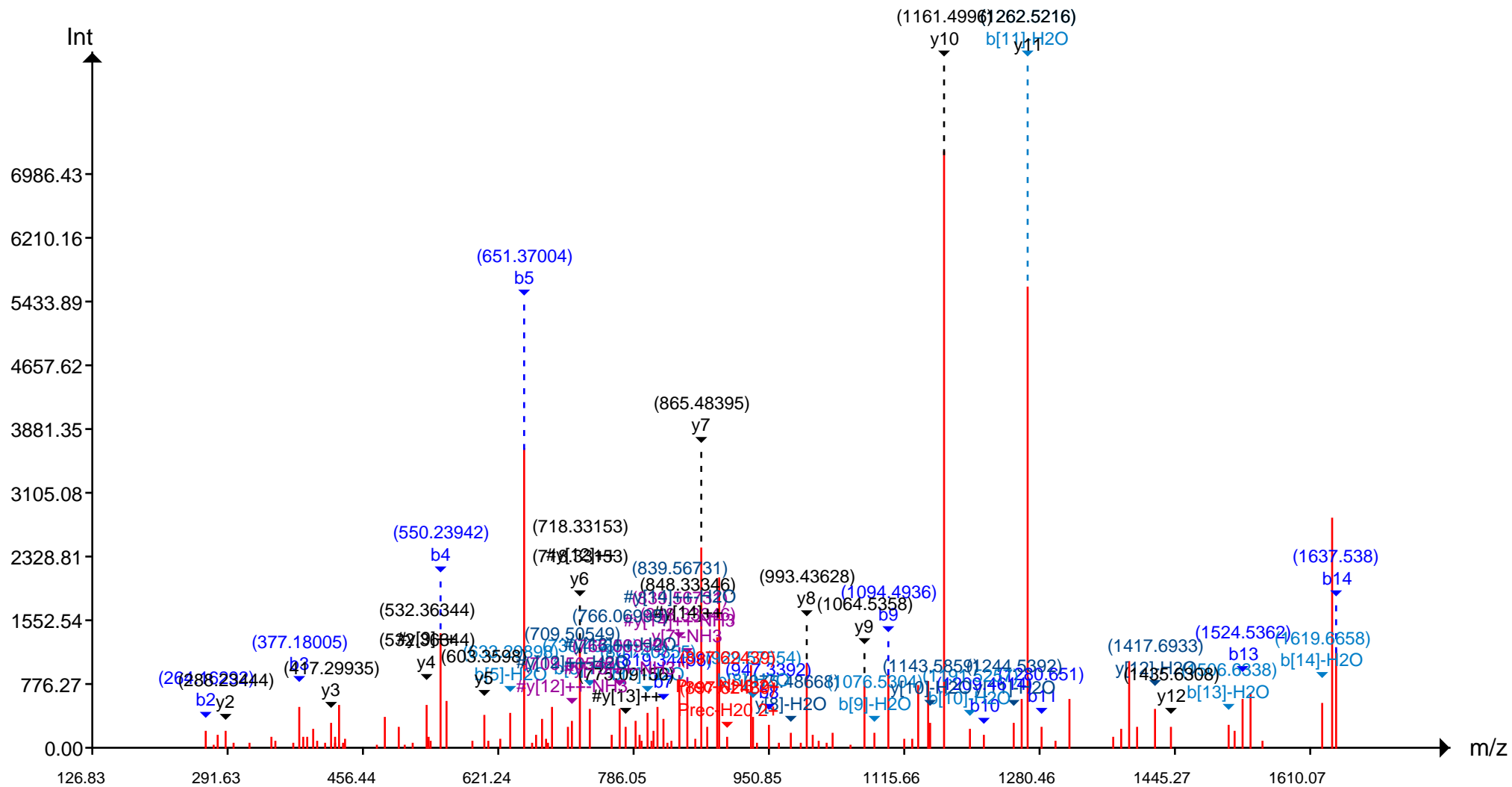
Info

Modified Sequence	AcD3-K<AcD3K*>SSEIEQAVQSLDR<C13>-COOH	Ion score	90.76
Protein	Q9D898	Confidence level	99.00%
Precursor m/z	843.45042	Identity Threshold	47.85
Charge	2+	Mass error	0.001721



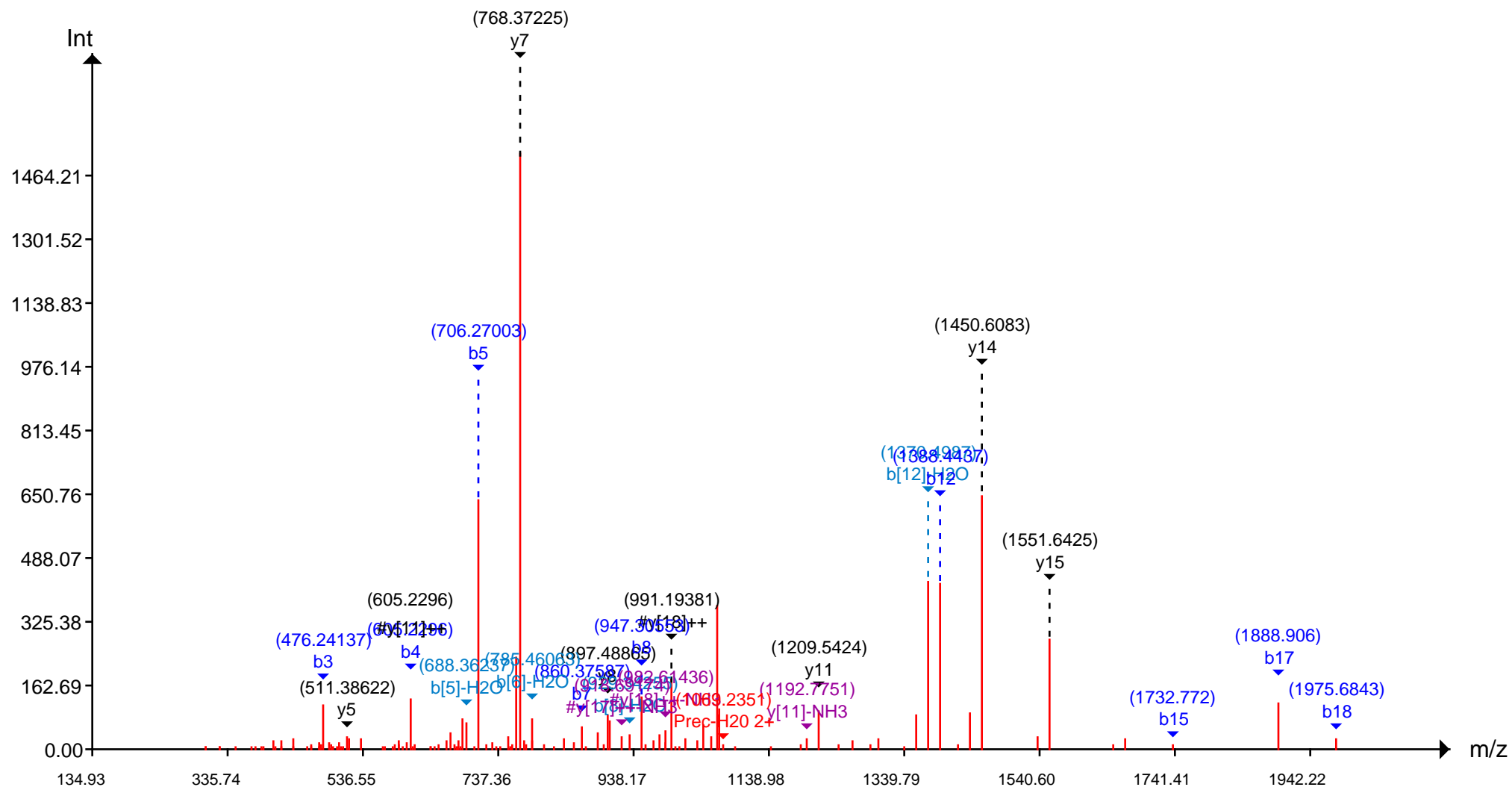
Info

Modified Sequence	AcD3-EQK<AcD3K*>ETPGSQPSEPC<Cmm*>SGVSR-COOH	Ion score	90.55
Protein	P27546	Confidence level	99.00%
Precursor m/z	1075.502	Identity Threshold	45.42
Charge	2+	Mass error	0.003013



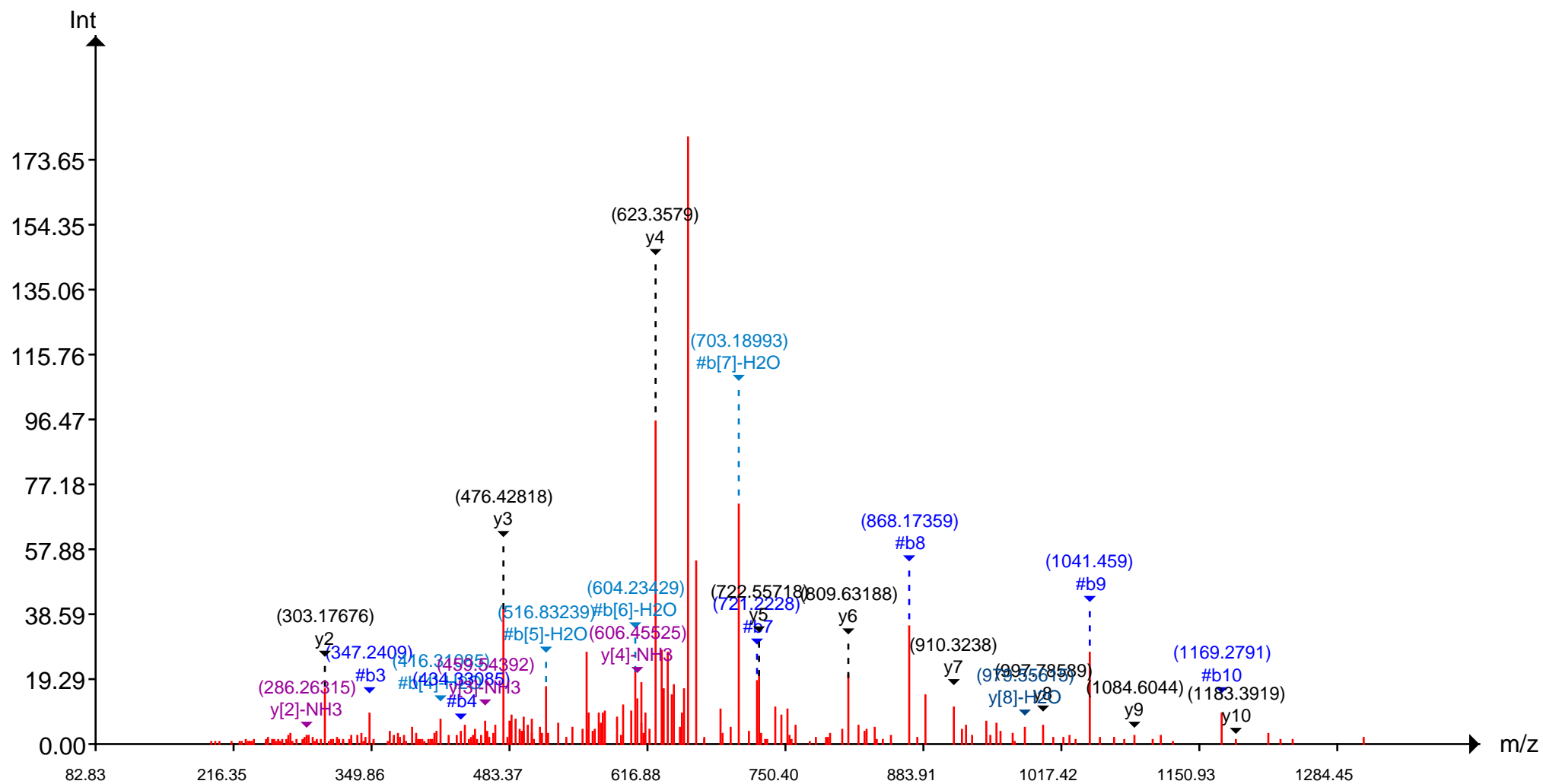
Info

Modified Sequence	AcD3-AM<Mox*>LK<AcD3K*>TPAQFDAELR-COOH	Ion score	76.0
Protein	P10107	Confidence level	99.00%
Precursor m/z	906.46075	Identity Threshold	47.94
Charge	2+	Mass error	0.002758



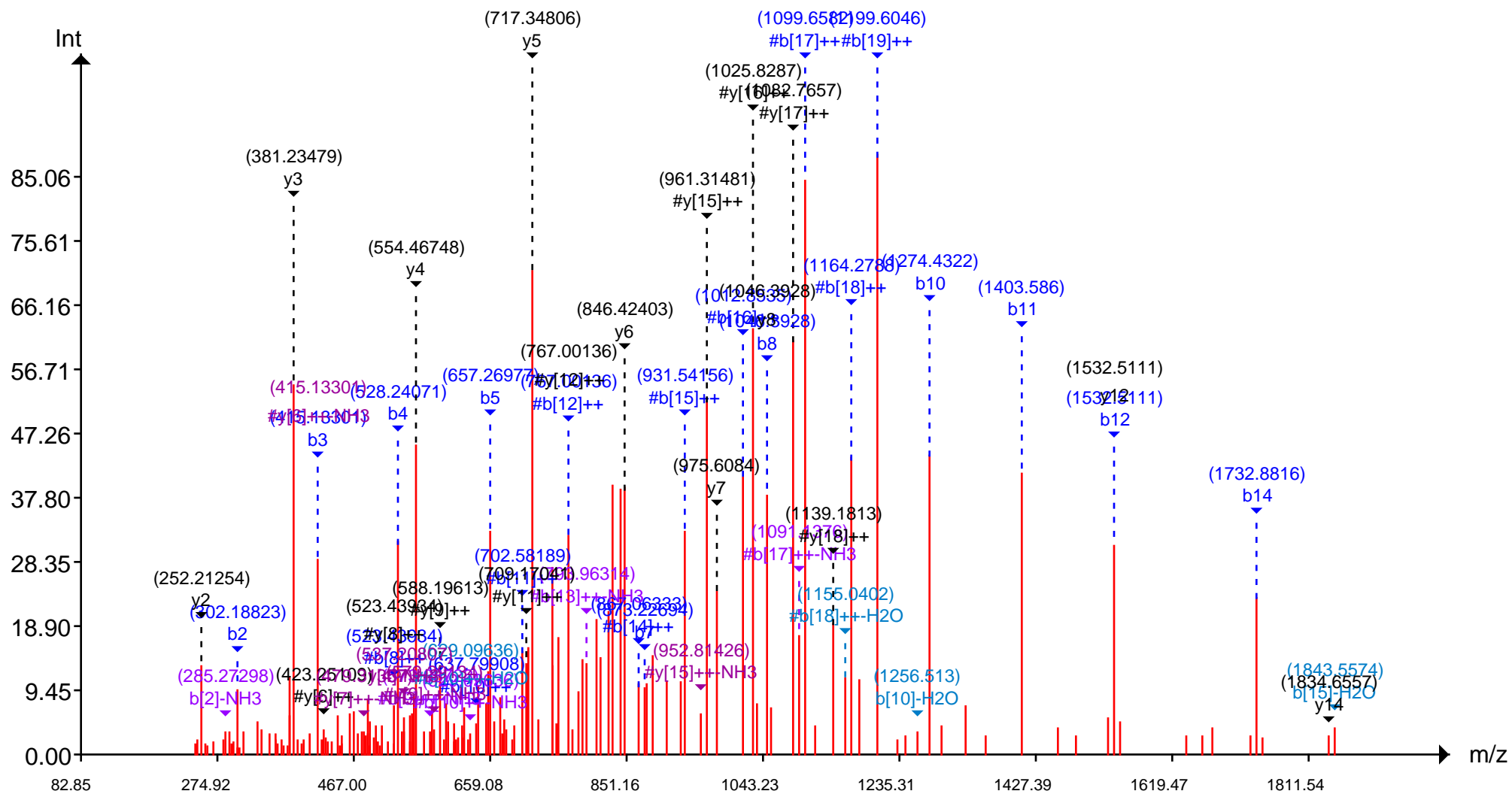
Info

Modified Sequence	AcD3-EQK<AcD3K*>ETPGSQPSEPC<Cmm*>SGVSR<C13>-COOH	Ion score	65.18
Protein	P27546	Confidence level	99.00%
Precursor m/z	1078.5115	Identity Threshold	46.07
Charge	2+	Mass error	0.001871



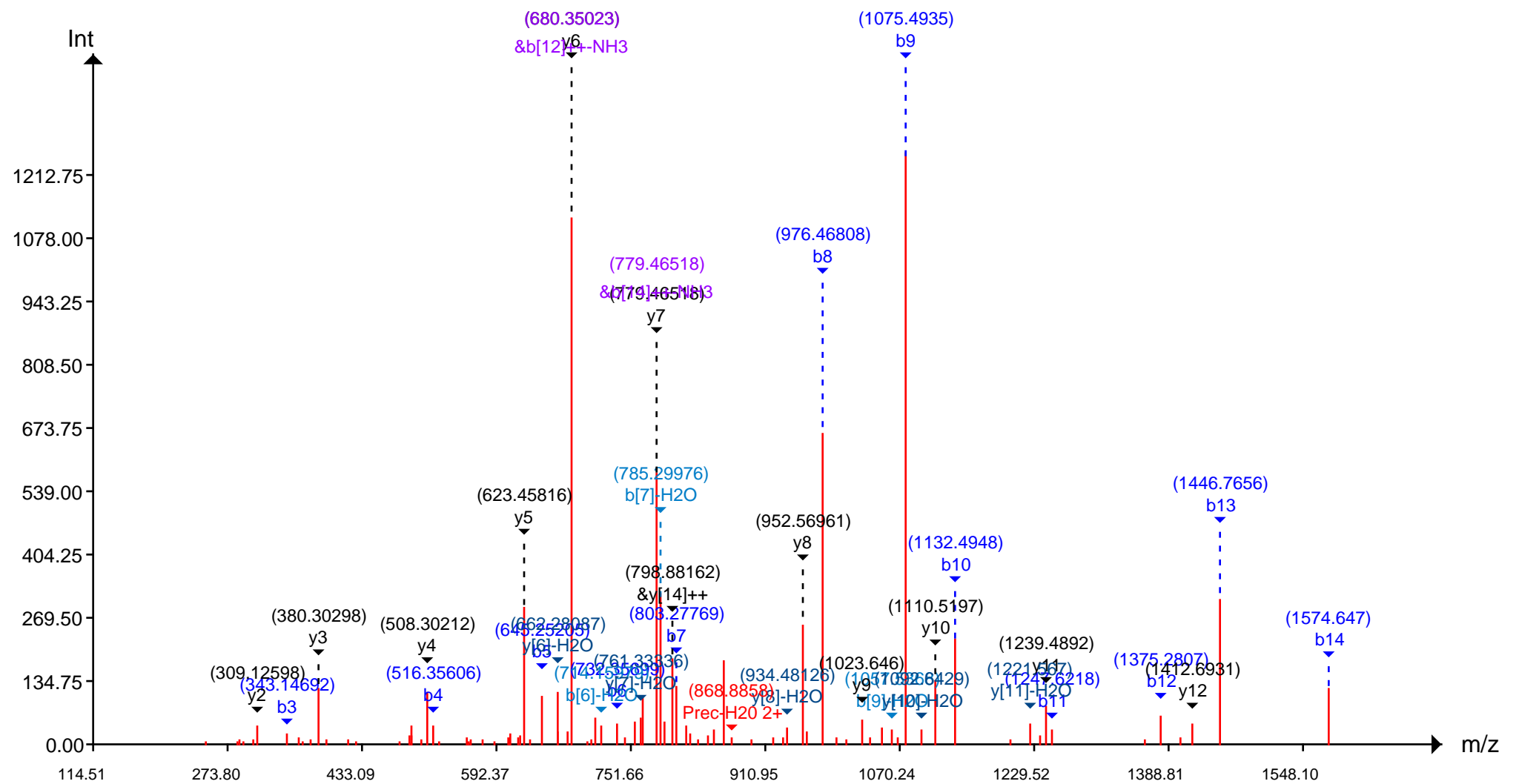
Info

Modified Sequence	AcD3-DVSSTSVFK<AcD3K*>QR-COOH	Ion score	48.41
Protein	Q8R1F1	Confidence level	99.00%
Precursor m/z	672.35818	Identity Threshold	48.00
Charge	2+	Mass error	0.001842



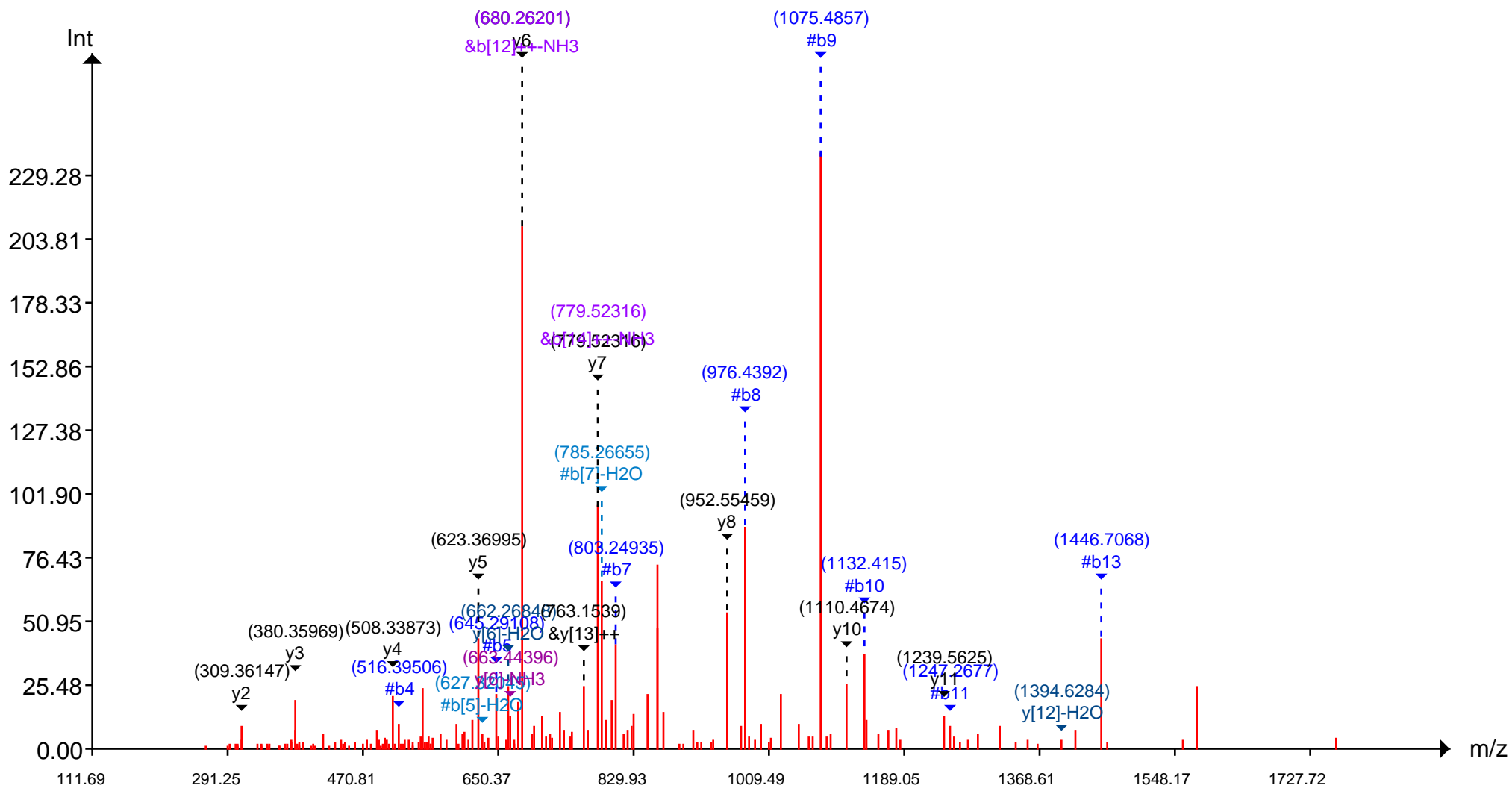
Info

Modified Sequence	AcD3-QQILESEK<AcD3K*>DLEEAEYK<AcD3K*>EAR<C13>-COOH	Ion score	65.8
Protein	P48428	Confidence level	99.00%
Precursor m/z	860.09976	Identity Threshold	47.56
Charge	3+	Mass error	0.003572



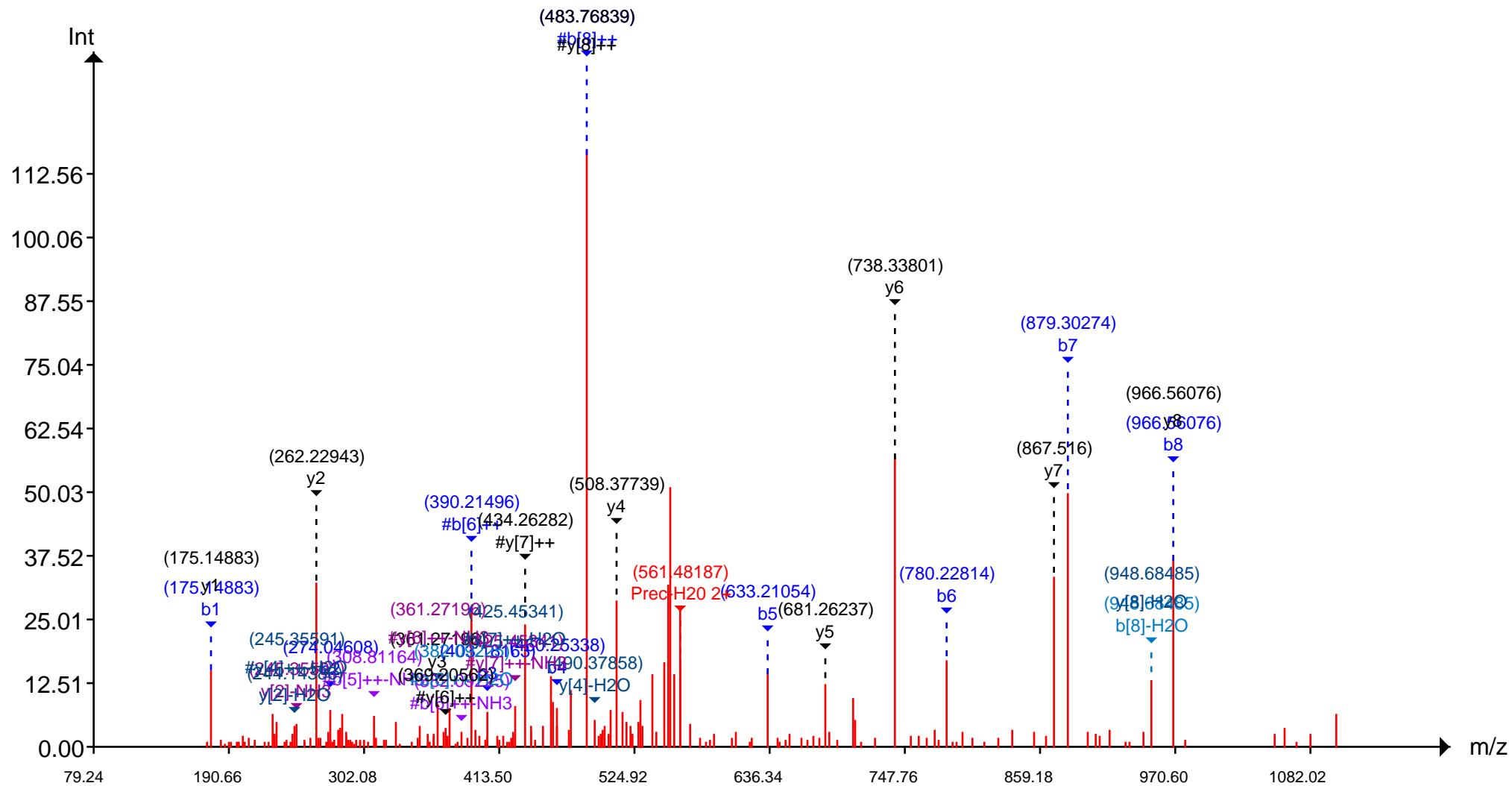
Info

Modified Sequence	AcD3-IAIK<AcD3K*>ESAK<AcD3K*>VGDQAQR<C13>-COOH	Ion score	112.34
Protein	Q99JY9	Confidence level	99.00%
Precursor m/z	878.00739	Identity Threshold	45.18
Charge	2+	Mass error	0.002257



Info

Modified Sequence	AcD3-IAIK<AcD3K*>ESAK<AcD3K*>VGDQAQR<C13>-COOH	Ion score	53.57
Protein	Q99JY9	Confidence level	99.00%
Precursor m/z	878.0073	Identity Threshold	45.21
Charge	2+	Mass error	0.002077



Info

Modified Sequence AcD3-EVEGK<AcD3K*>M<Mox*>VSR-COOH

Ion score 62.9

Protein P63028

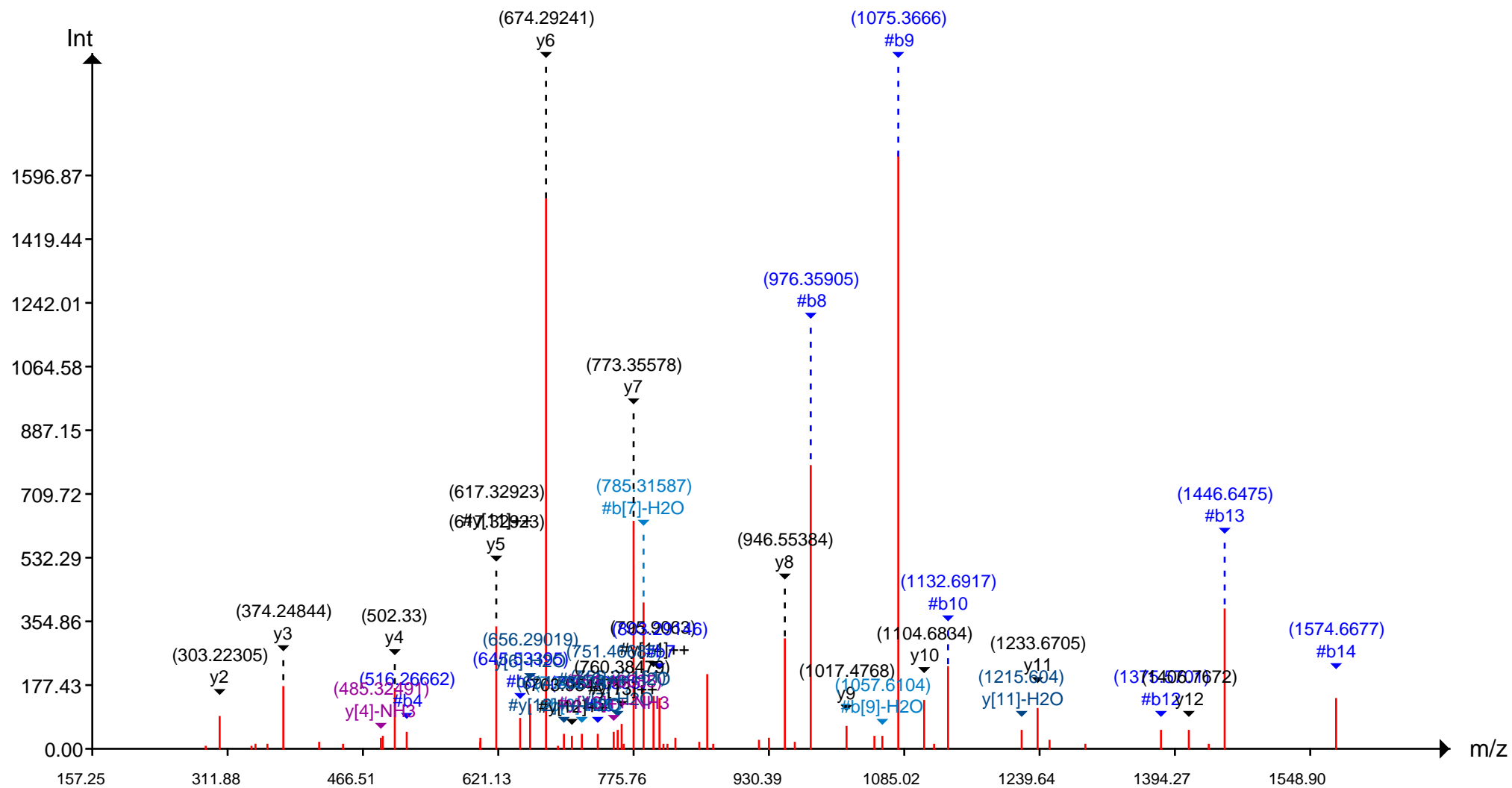
Confidence level 99.00%

Precursor m/z 570.79563

Identity Threshold 47.46

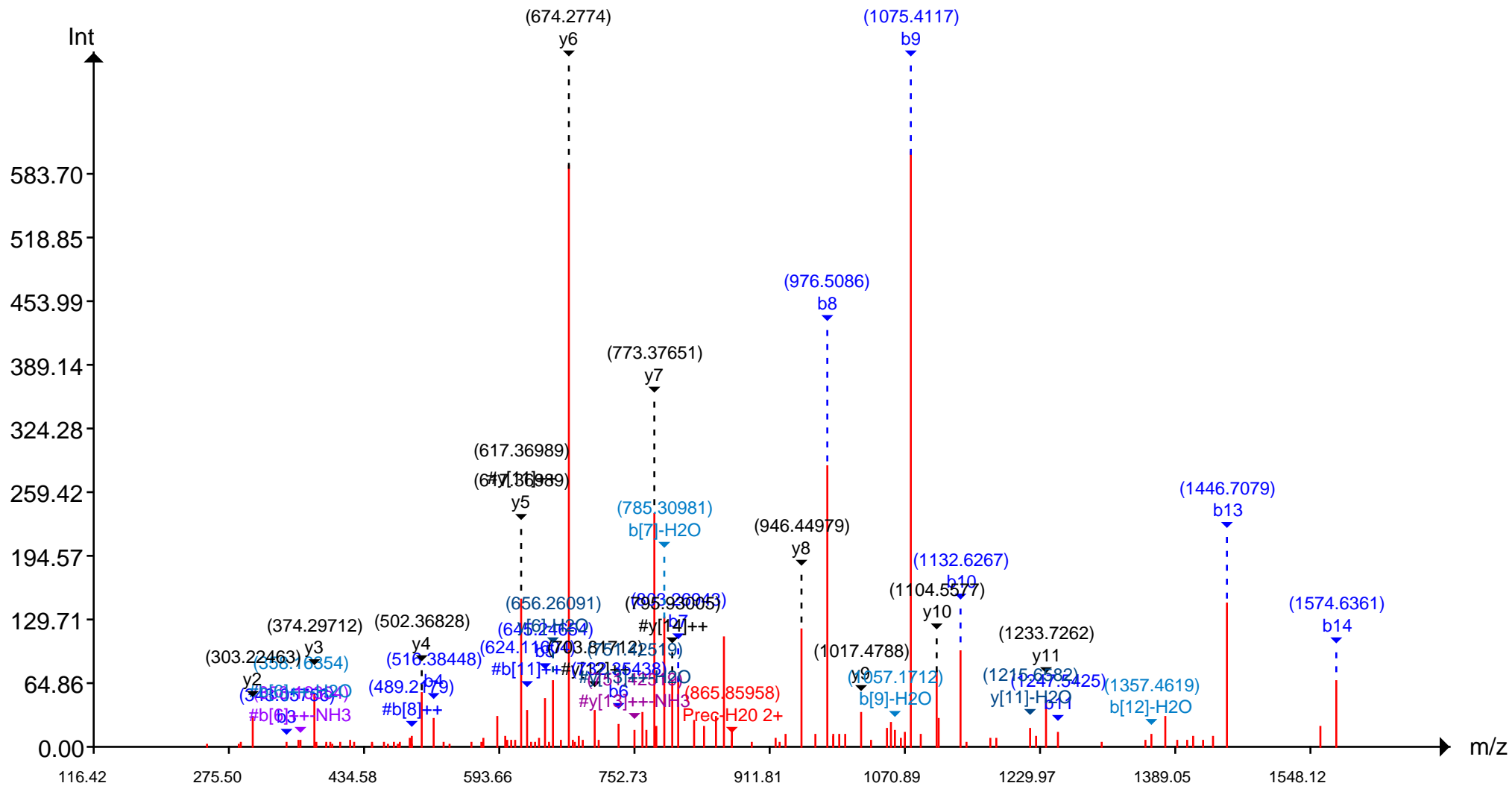
Charge 2+

Mass error 3.84E-4



Info

Modified Sequence	AcD3-IAIK<AcD3K*>ESAK<AcD3K*>VGDQAQR-COOH	Ion score	112.97
Protein	Q99JY9	Confidence level	99.00%
Precursor m/z	874.99762	Identity Threshold	46.08
Charge	2+	Mass error	0.002859



Info

Modified Sequence AcD3-IAIK<AcD3K*>ESAK<AcD3K*>VGDQAQR-COOH

Ion score 92.34

Protein Q99JY9

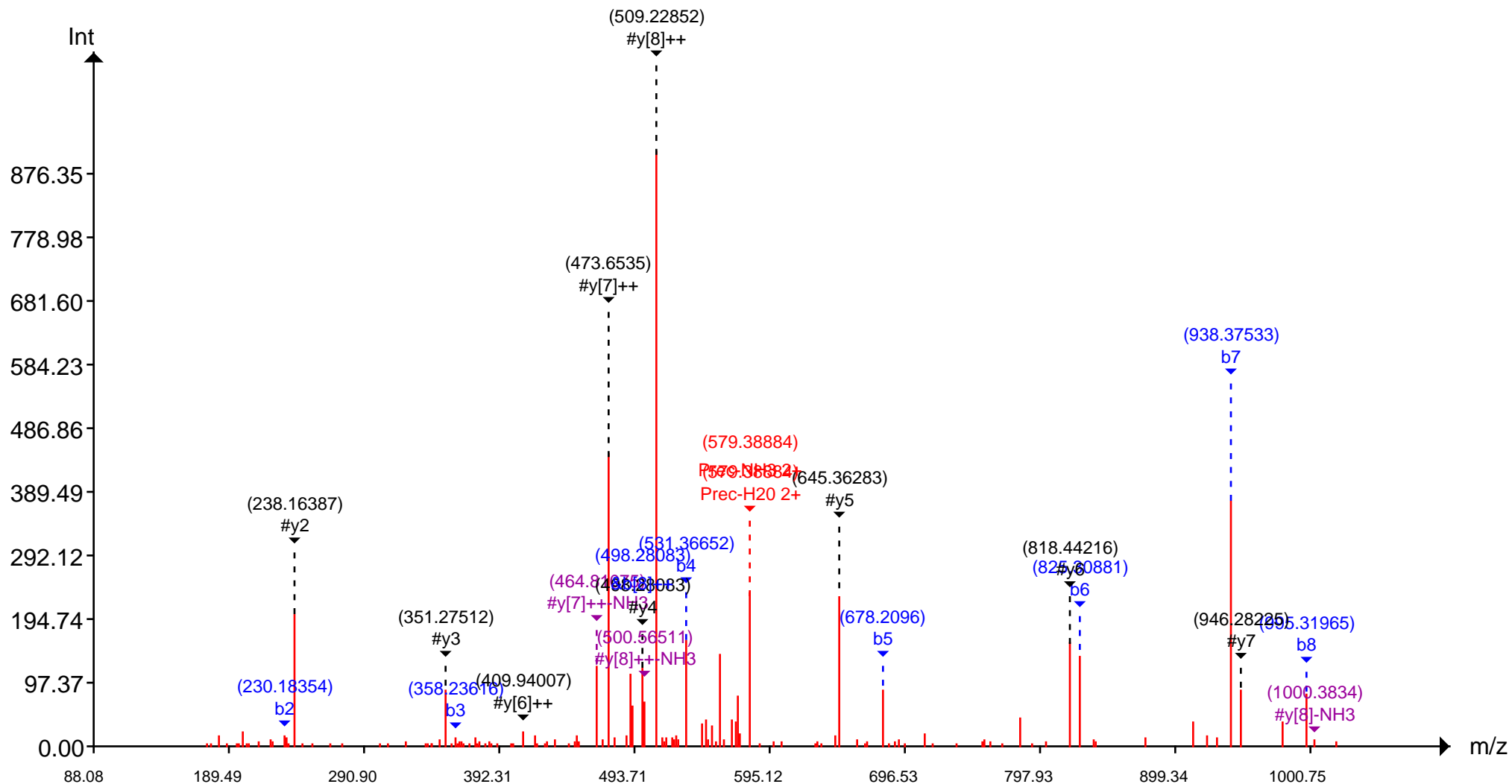
Confidence level 99.00%

Precursor m/z 874.99731

Identity Threshold 46.19

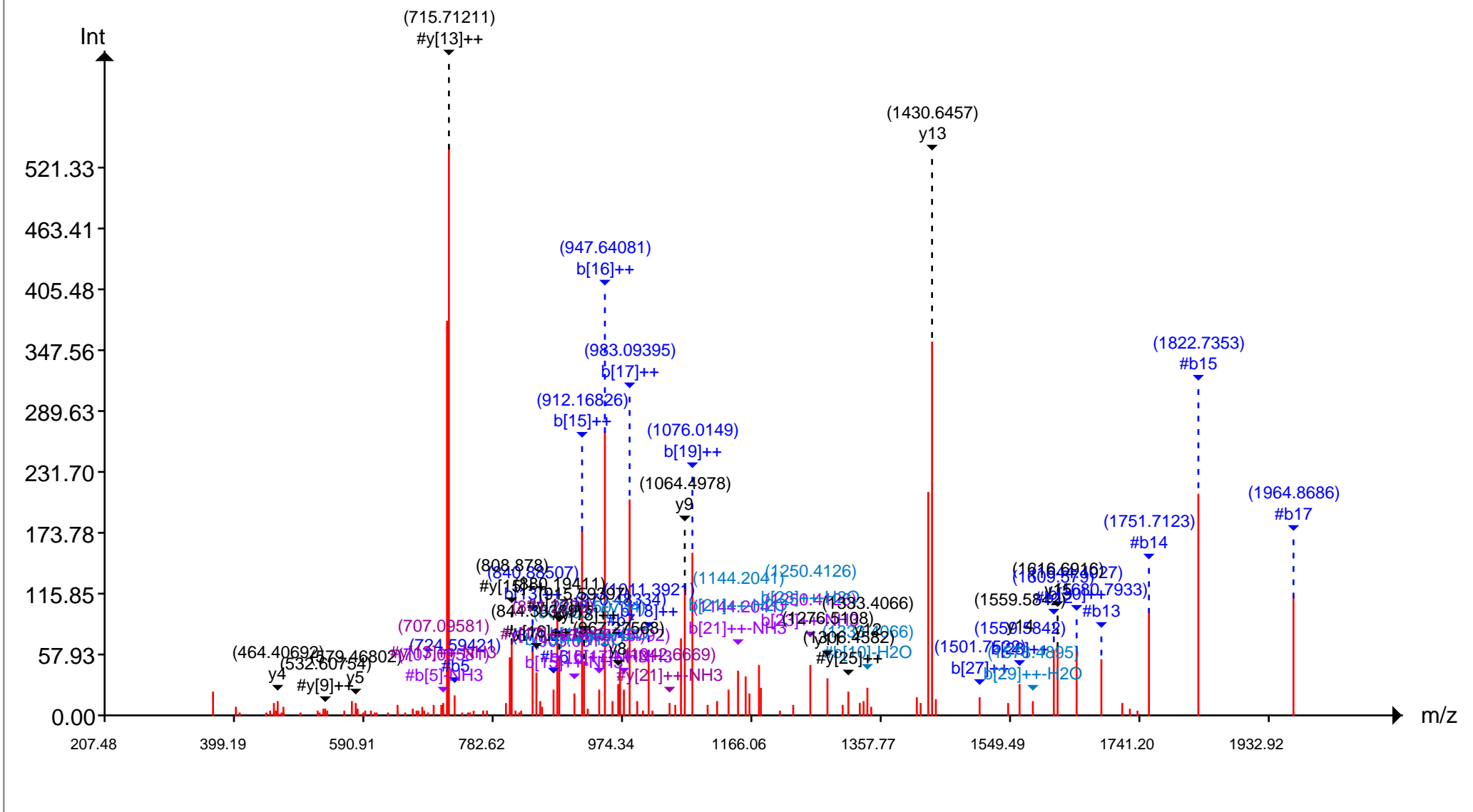
Charge 2+

Mass error 0.002239



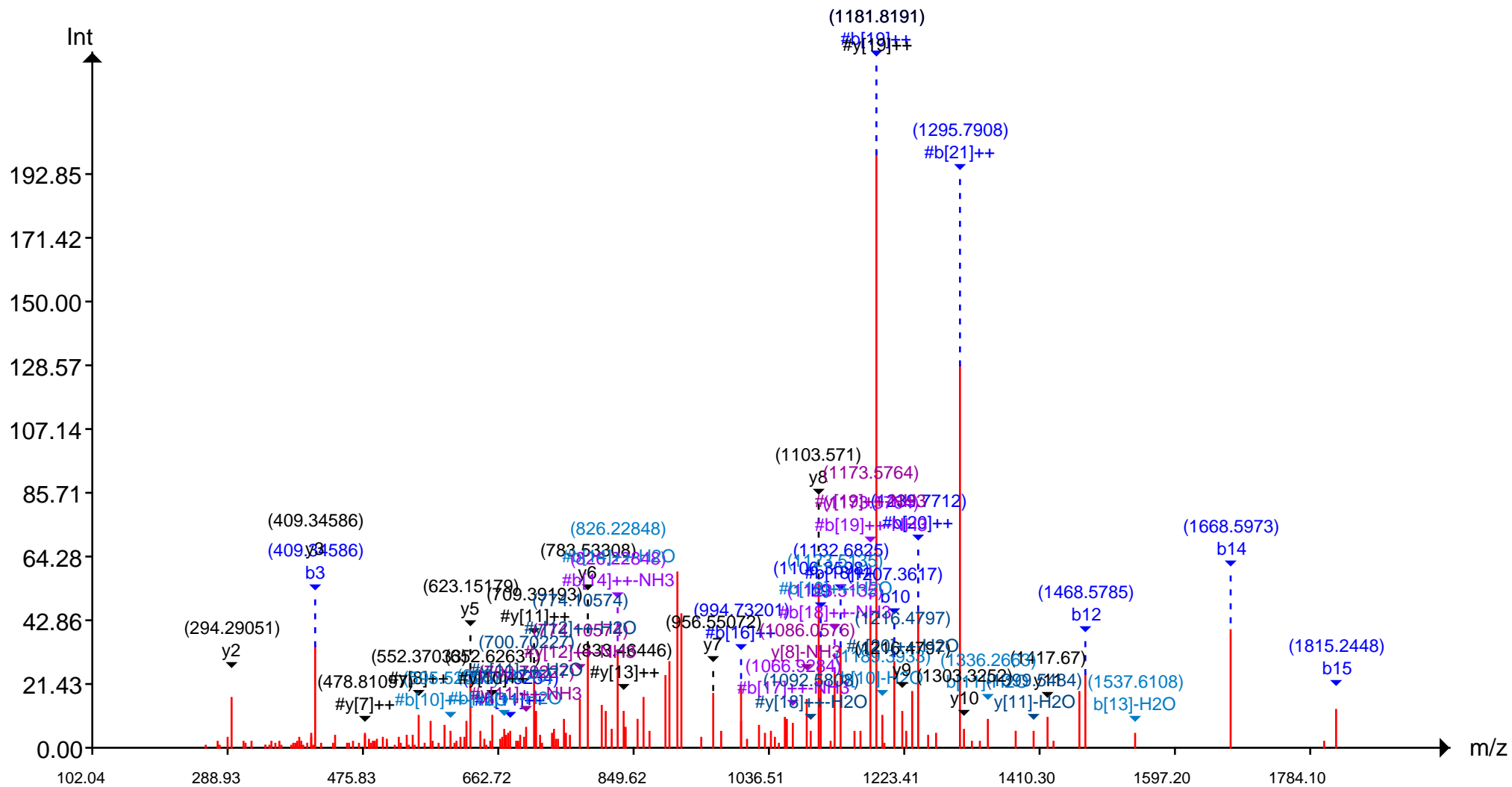
Info

Modified Sequence	AcD3-LAQK<AcD3K*>M<Mox*>M<Mox*>IGR<C13>-COOH	Ion score	50.14
Protein	P52480, P53657	Confidence level	99.00%
Precursor m/z	588.32796	Identity Threshold	47.60
Charge	2+	Mass error	-2.64E-4



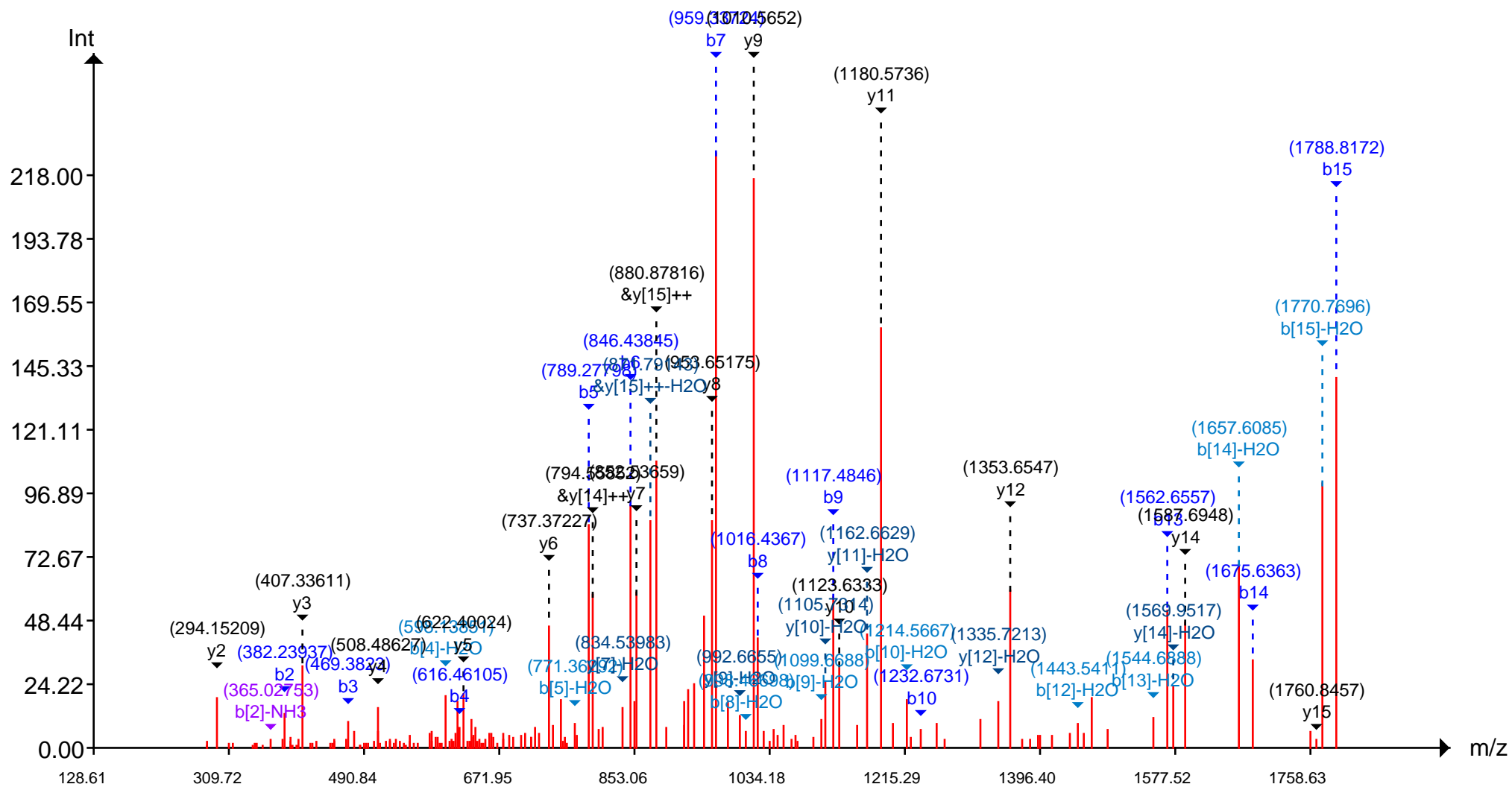
Info

Modified Sequence	AcD3-FDENK<AcD3K*>FVDEHEEAAAAAGEPGPDPC<Cmm*>EVDGLLR<C13>-COOH	Ion score	62.88
Protein	Q9D898	Confidence level	99.00%
Precursor m/z	1194.2189	Identity Threshold	43.26
Charge	3+	Mass error	0.003463



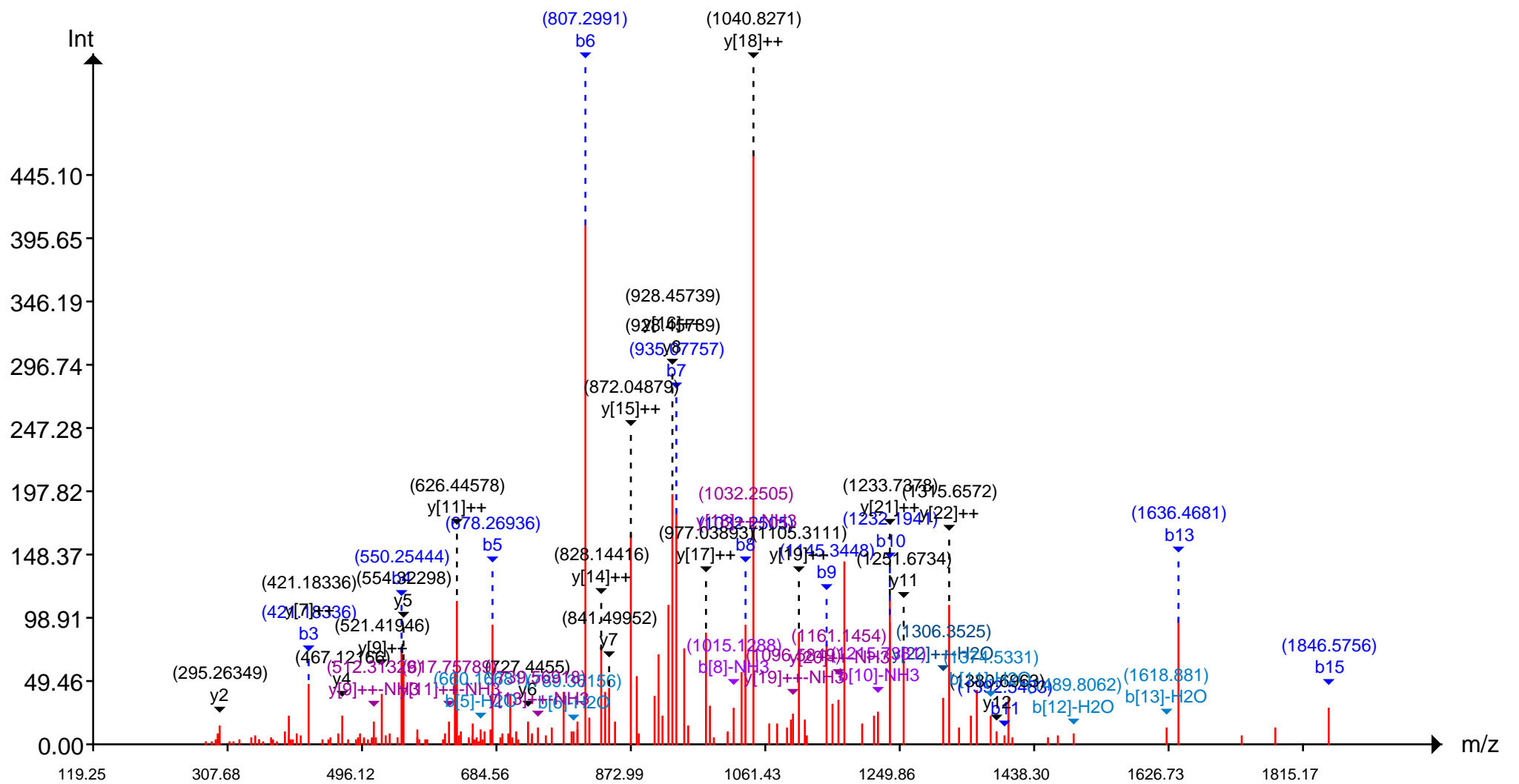
Info

Modified Sequ...	AcD3-M<Mox*>ESC<Cmm*>GIHETTFNSIM<Mox*>K<AcD3K*>C<Cmm*>DVIDIR<C13>-COOH	Ion score	49.23
Pro...	P60710, P63260	Confidence level	99.00%
Precursor ...	924.41751	Identity Threshold	42.58
Ch...	3+	Mass error	0.00221

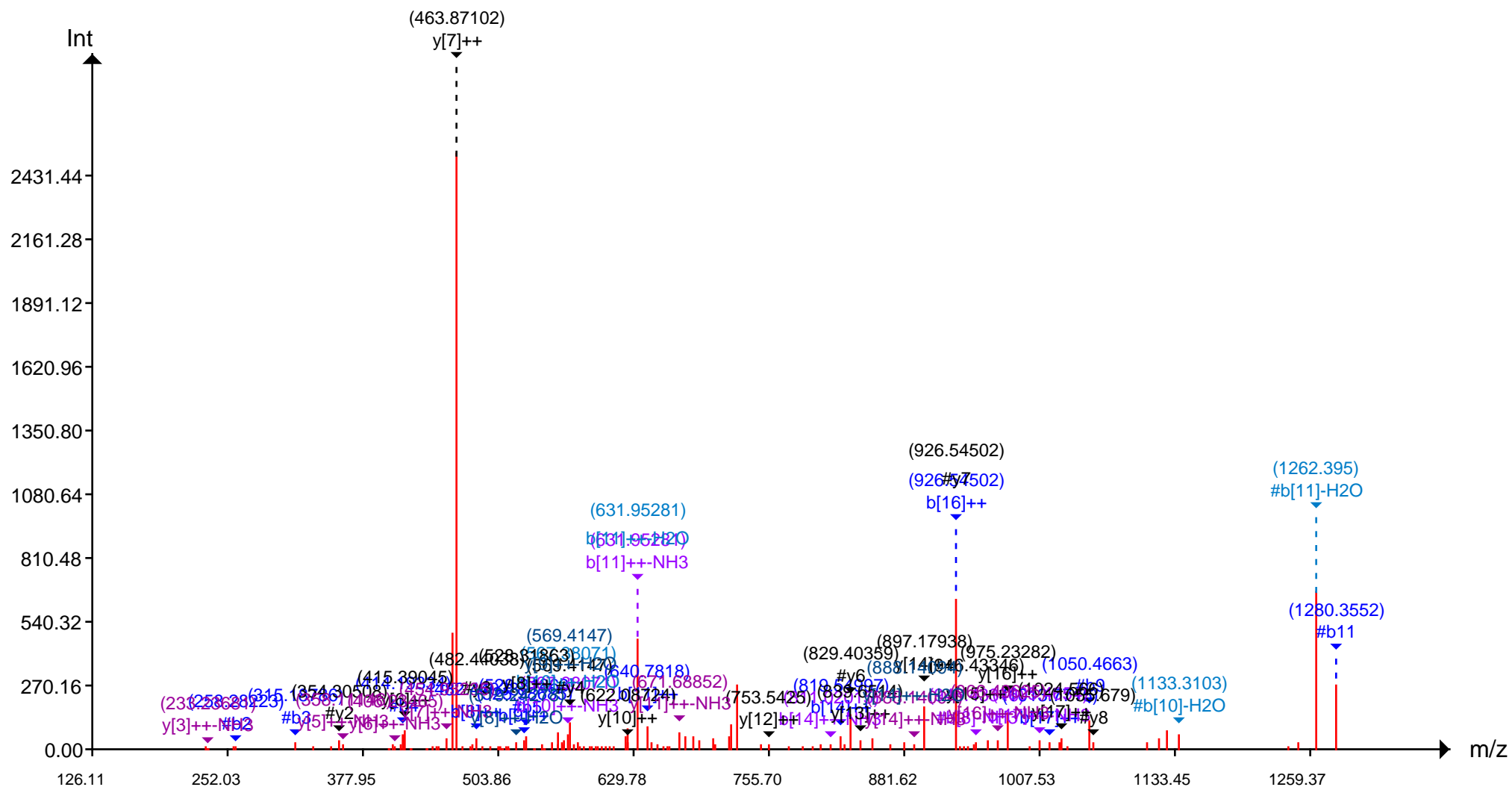


Info

Modified Sequence	AcD3-YK<AcD3K*>SM<Mox*>K<AcD3K*>GLGTDDNTLIR<C13>-COOH	Ion score	105.72
Protein	P97429	Confidence level	99.00%
Precursor m/z	985.02264	Identity Threshold	47.82
Charge	2+	Mass error	0.002759

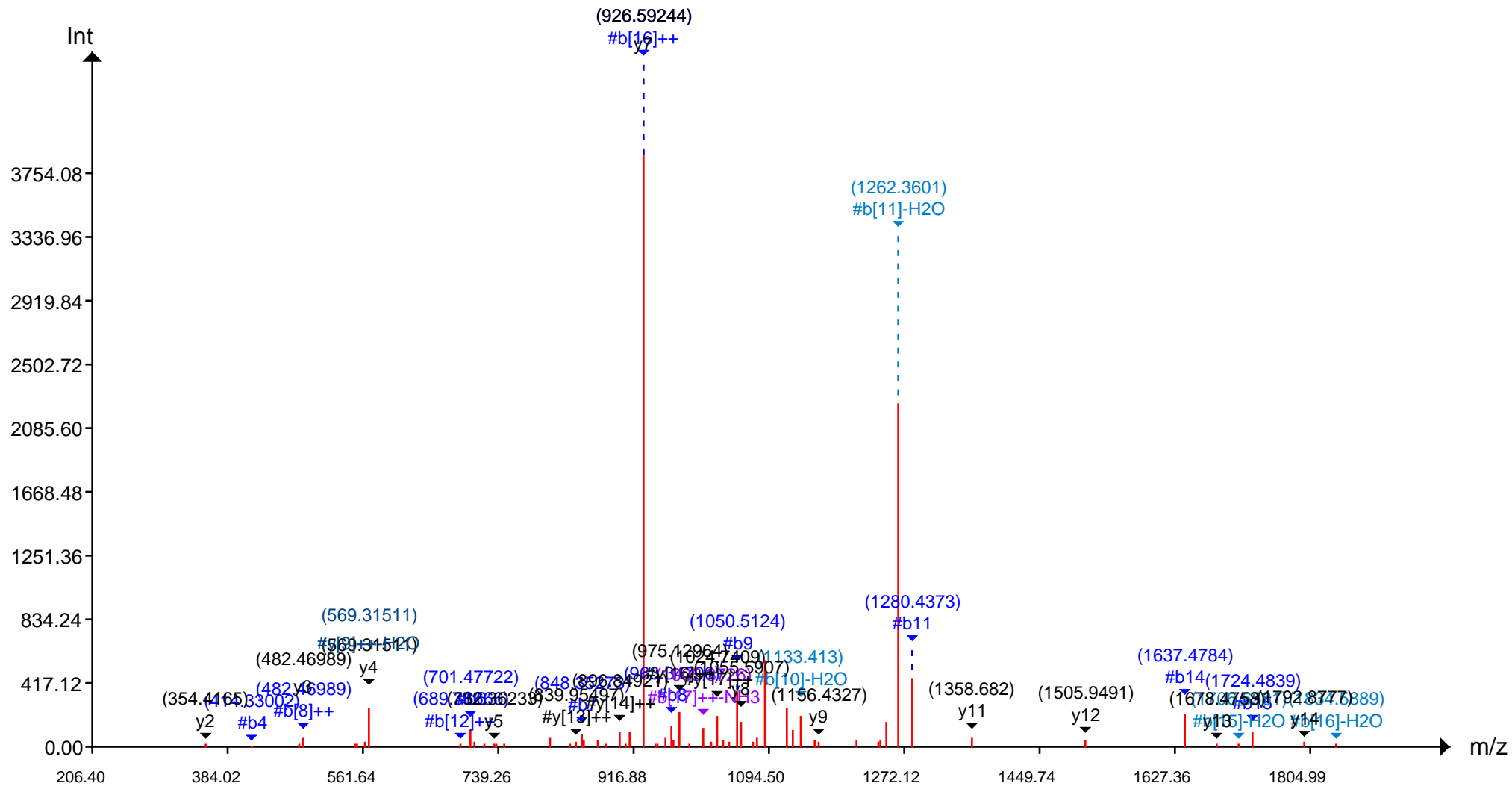


Info			
Modified Sequence	AcD3-LVYEQEQPLSC<Cmm*>DEPILSNK<AcD3K*>SGDNR<C13>-COOH	Ion score	92.08
Protein	P70296	Confidence level	99.00%
Precursor m/z	963.13925	Identity Threshold	46.92
Charge	3+	Mass error	0.003985



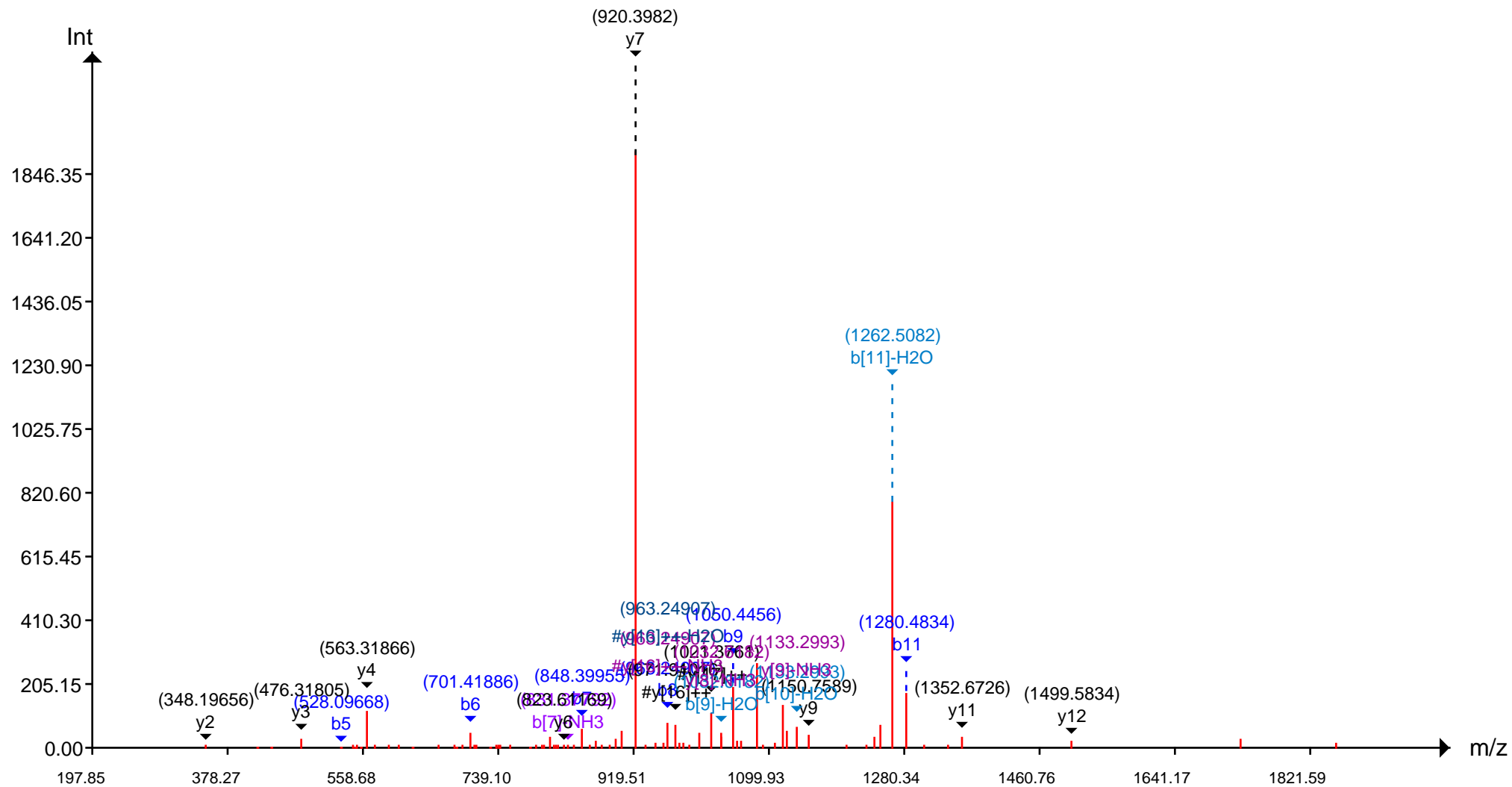
Info

Modified Sequence	AcD3-IVGVNK<AcD3K*>M<Mox*>DSTEPPYSQK<AcD3K*>R<C13>-COOH	Ion score	54.11
Protein	P10126	Confidence level	99.00%
Precursor m/z	736.05443	Identity Threshold	47.98
Charge	3+	Mass error	0.002149



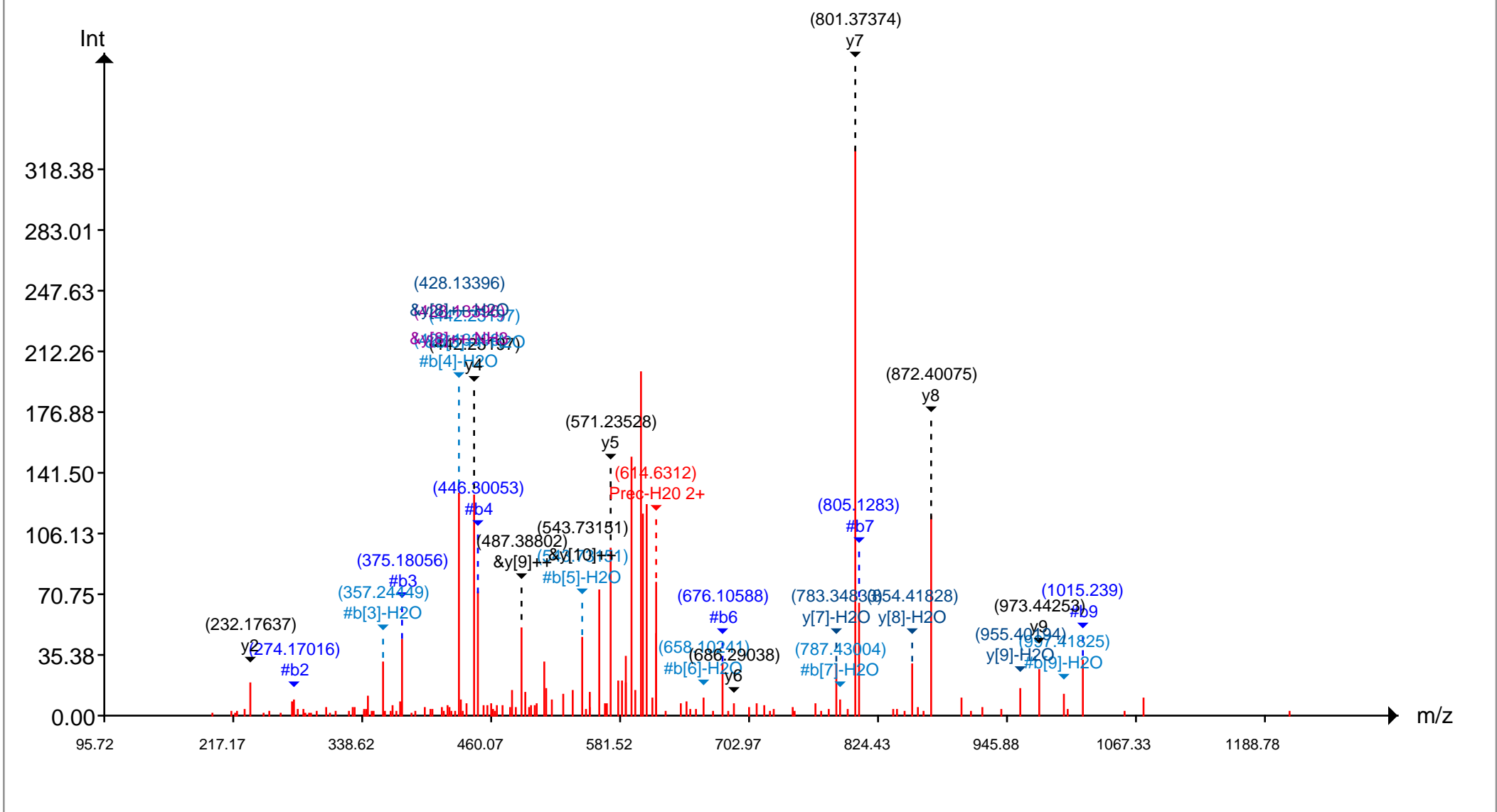
Info

Modified Sequence	AcD3-IVGVNK<AcD3K*>M<Mox*>DSTEPPYSQK<AcD3K*>R<C13>-COOH	Ion score	72.39
Protein	P10126	Confidence level	99.00%
Precursor m/z	1103.578	Identity Threshold	47.98
Charge	2+	Mass error	0.002135



Info

Modified Sequence	AcD3-IVGVNK<AcD3K*>M<Mox*>DSTEPPEYSQK<AcD3K*>R-COOH	Ion score	48.89
Protein	P10126	Confidence level	99.00%
Precursor m/z	1100.5683	Identity Threshold	47.93
Charge	2+	Mass error	0.002877



Info

Modified Sequence AcD3-DITADDEPLGR-COOH

Protein P17742

Precursor m/z 623.80361

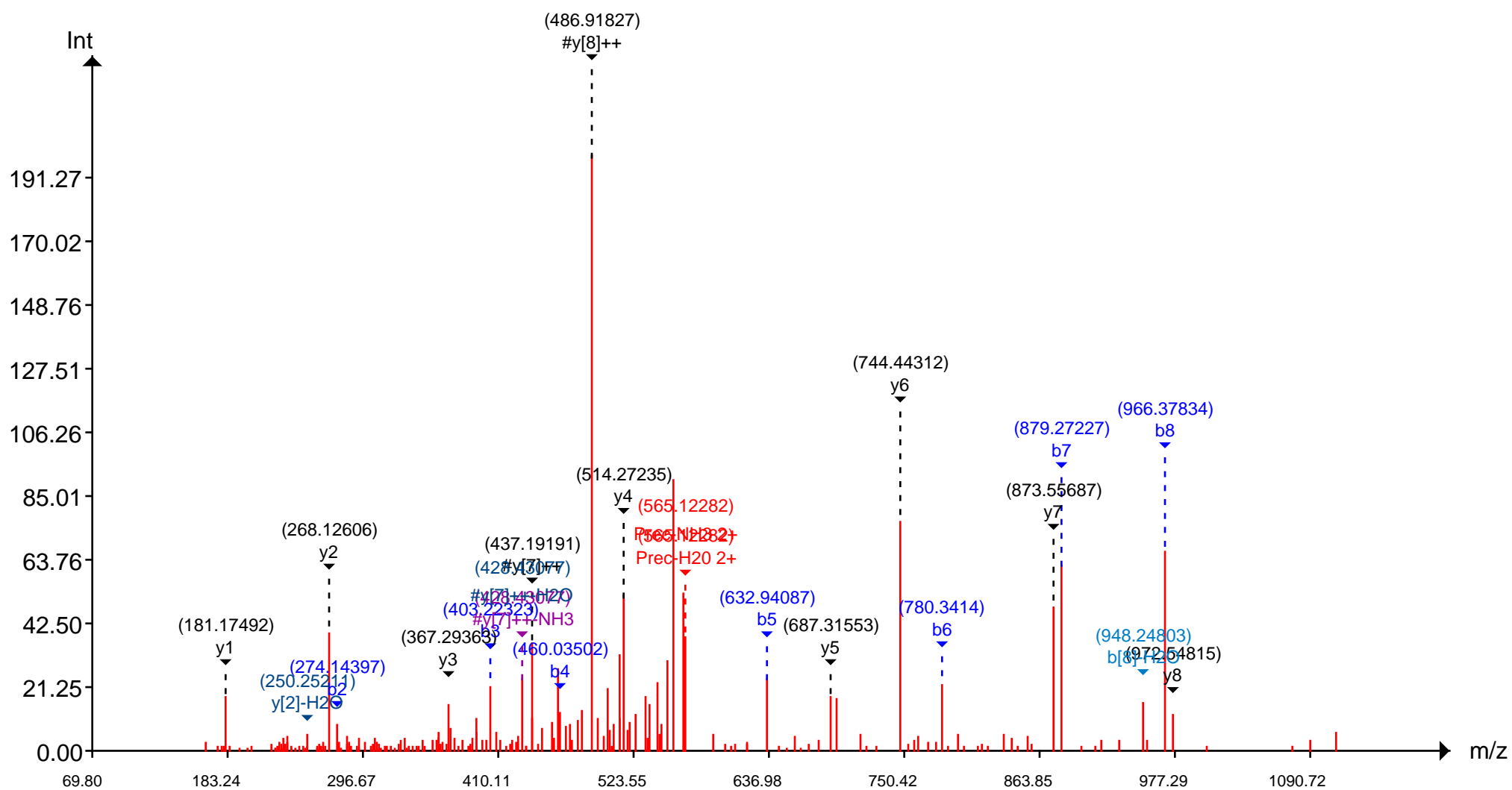
Charge 2+

Ion score 48.75

Confidence level 99.00%

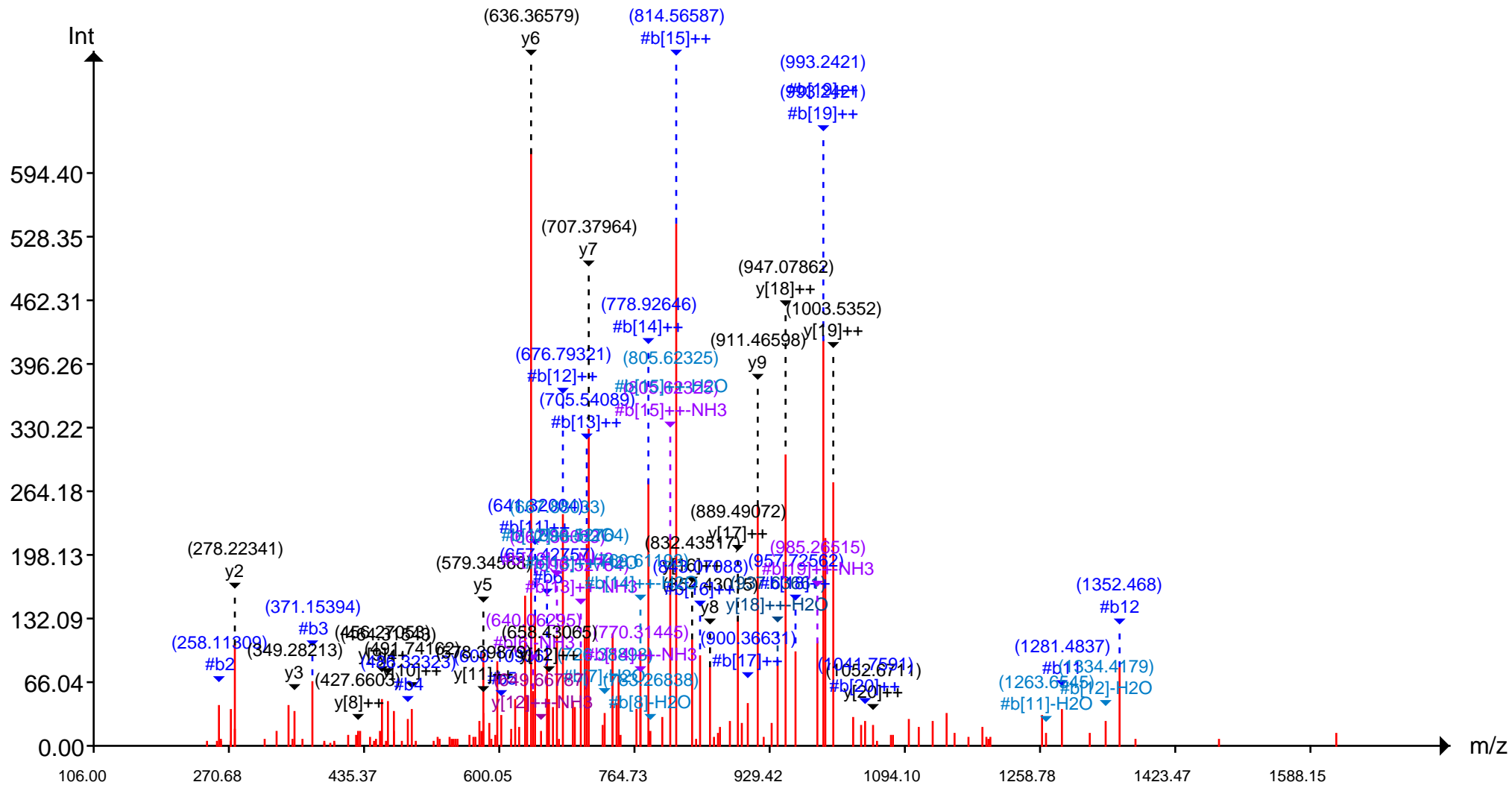
Identity Threshold 46.88

Mass error 0.001039



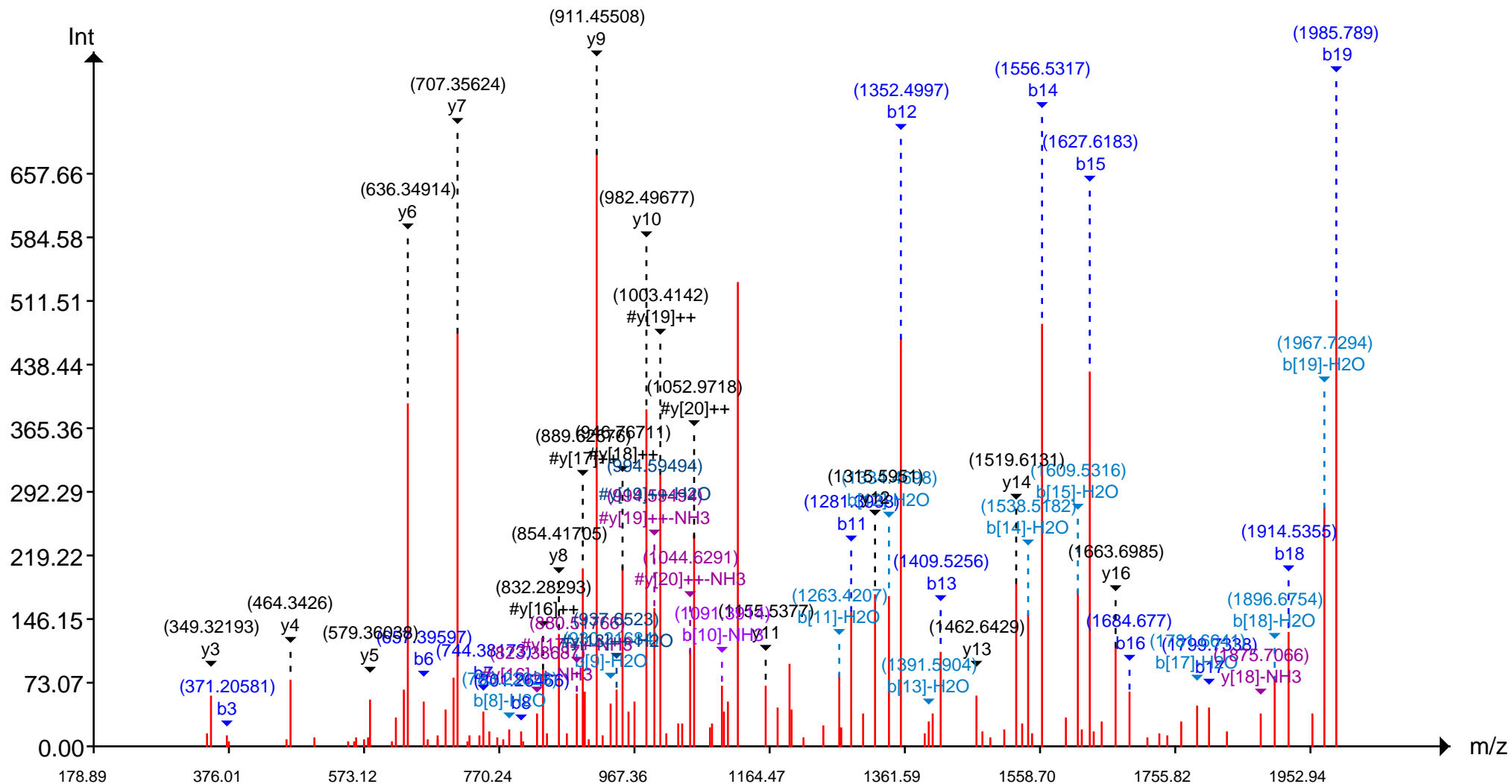
Info

Modified Sequence	AcD3-EVEGK<AcD3K*>M<Mox*>VSR<C13>-COOH	Ion score	52.99
Protein	P63028	Confidence level	99.00%
Precursor m/z	573.80566	Identity Threshold	47.94
Charge	2+	Mass error	3.02E-4



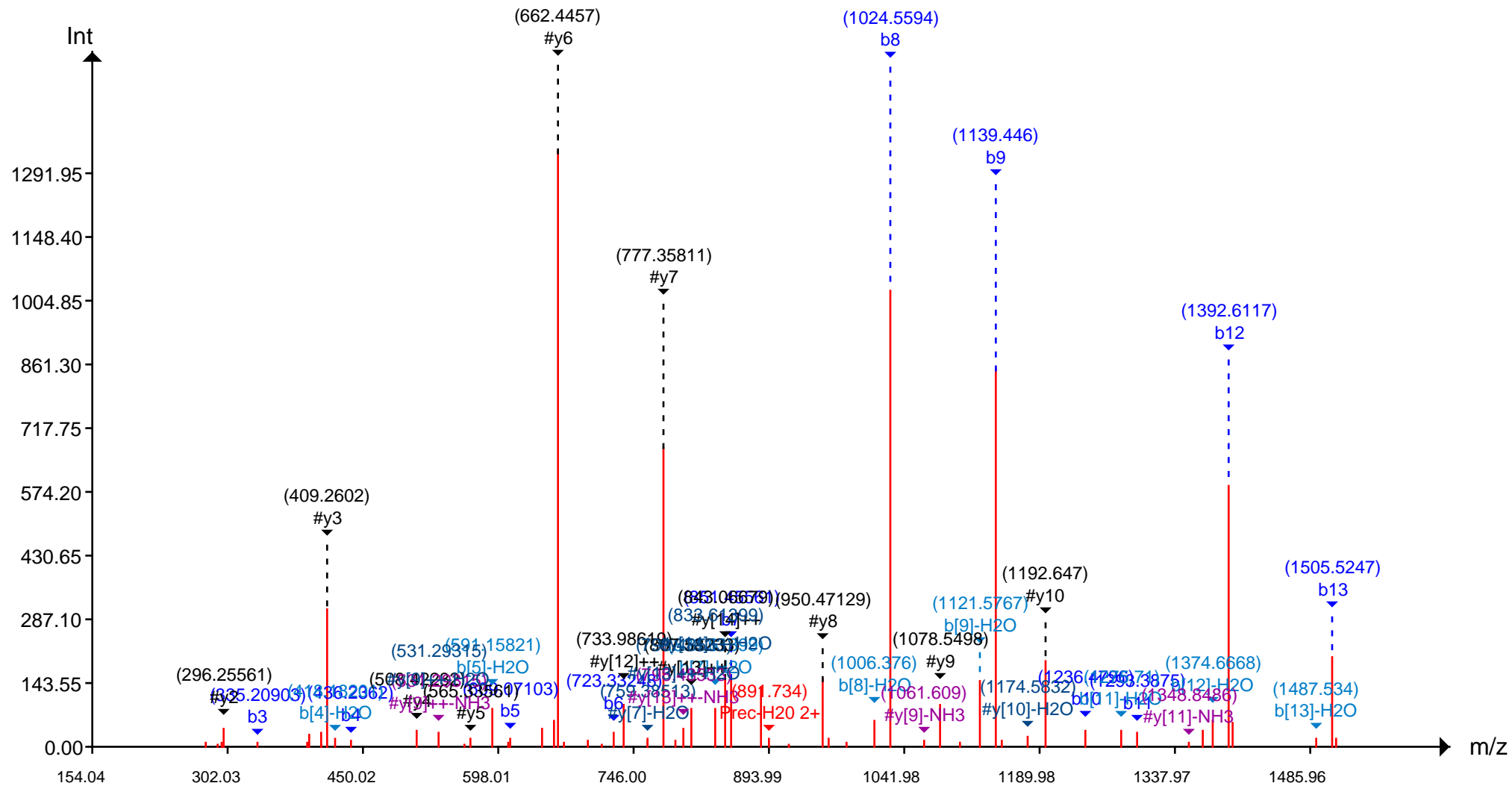
Info

Modified Sequence	AcD3-LVIDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH	Ion score	65.12
Protein	P63260	Confidence level	99.00%
Precursor m/z	755.02815	Identity Threshold	46.30
Charge	3+	Mass error	2.9E-4



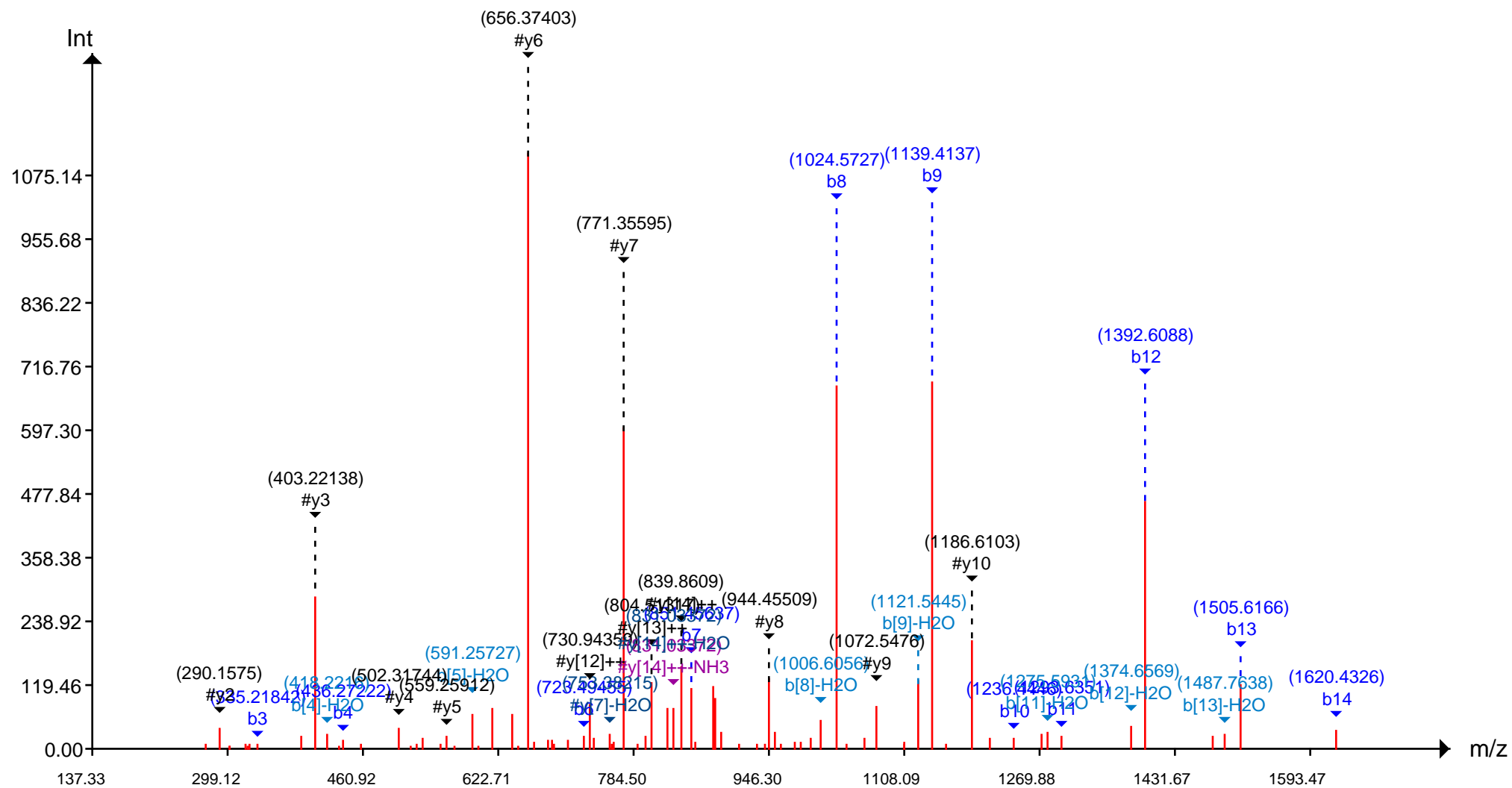
Info

Modified Sequence	AcD3-LVIDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH	Ion score	118.94
Protein	P63260	Confidence level	99.00%
Precursor m/z	1132.0392	Identity Threshold	46.37
Charge	2+	Mass error	0.001516



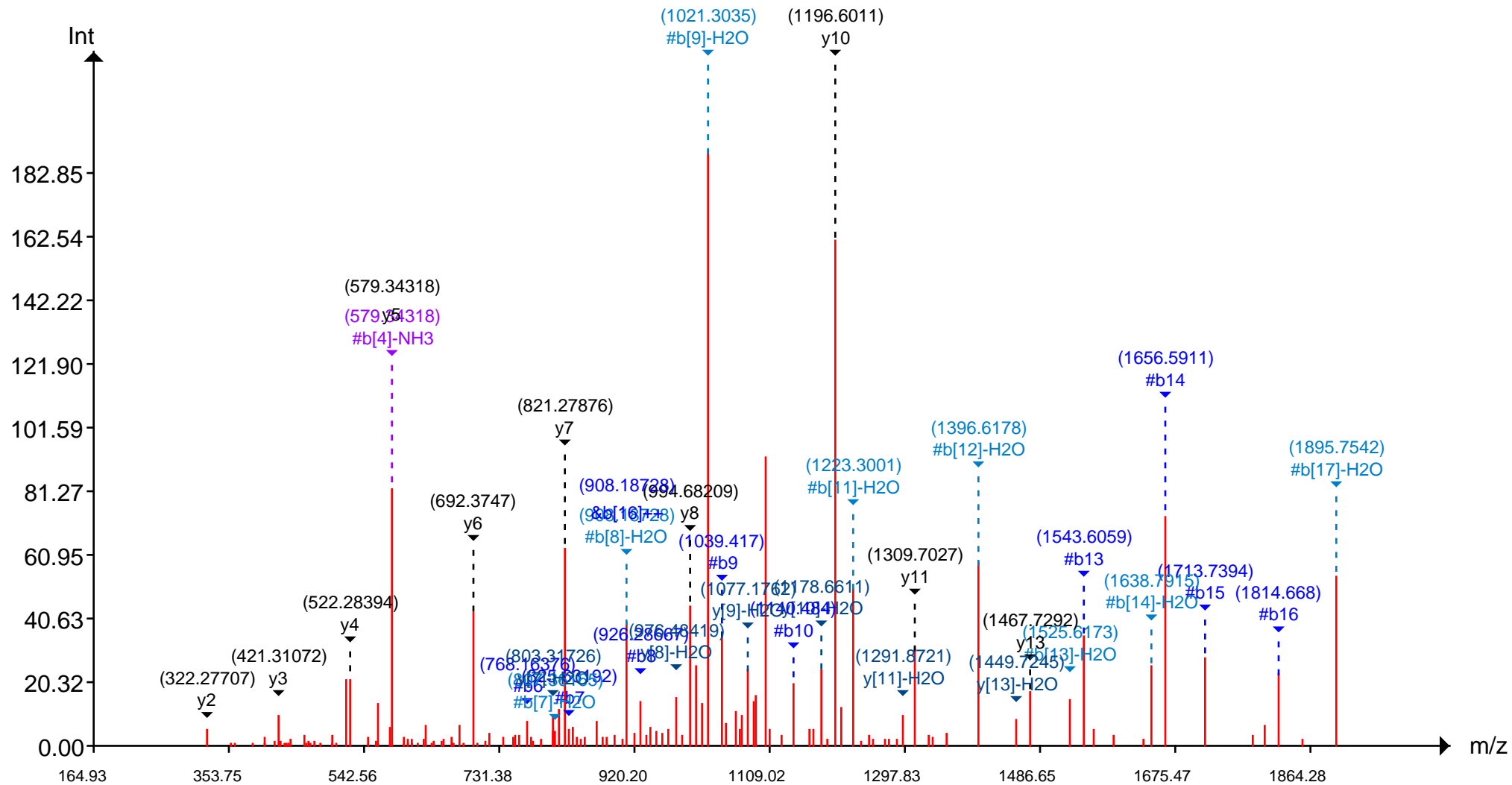
Info

Modified Sequence	AcD3-AAFTK<AcD3K*>NQK<AcD3K*>DPGVLDK<C13>-COOH	Ion score	83.77
Protein	P50543	Confidence level	99.00%
Precursor m/z	900.99923	Identity Threshold	47.12
Charge	2+	Mass error	0.001562



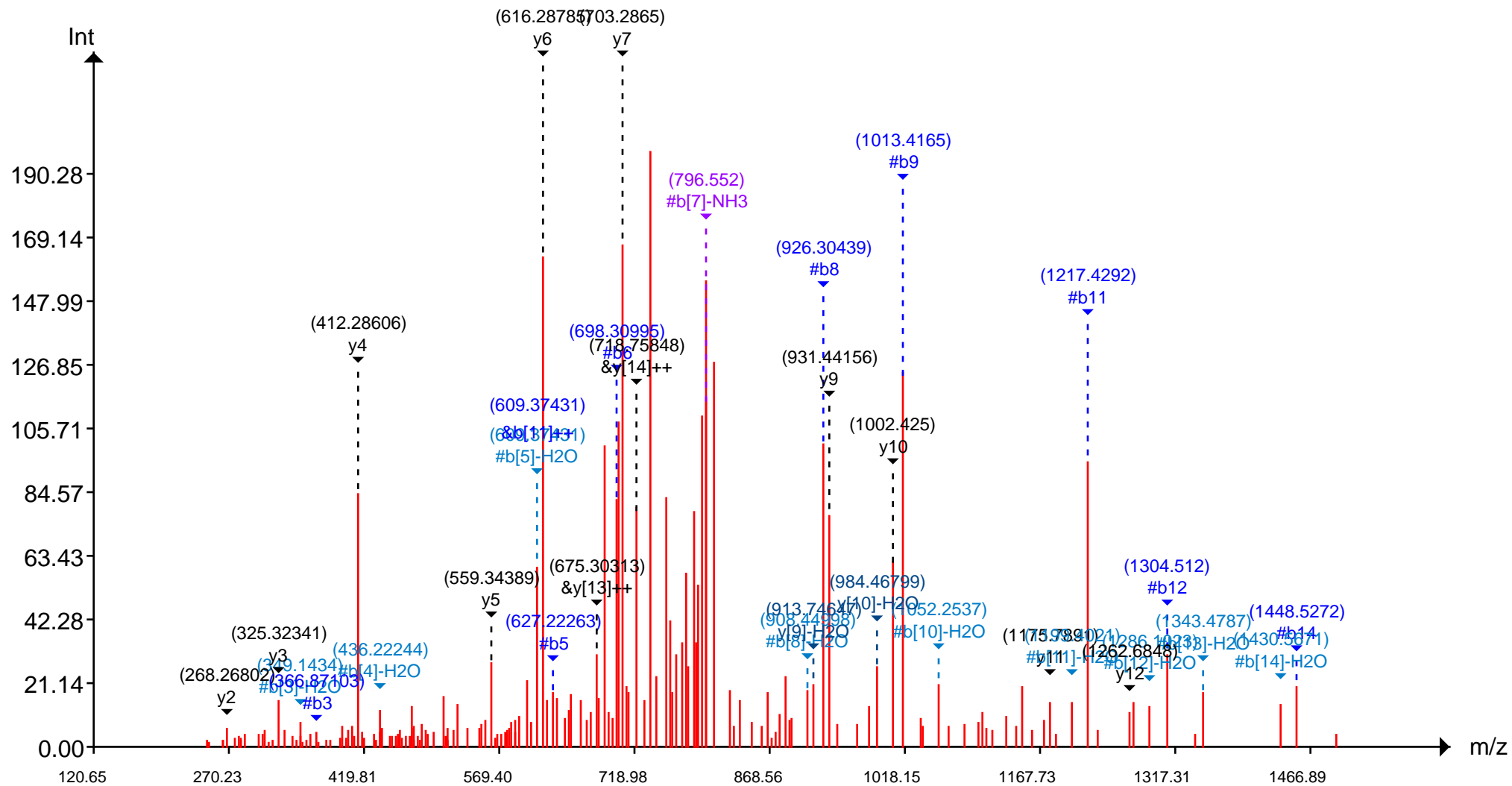
Info

Modified Sequence	AcD3-AAFTK<AcD3K*>NQK<AcD3K*>DPGVLDLDR-COOH	Ion score	84.19
Protein	P50543	Confidence level	99.00%
Precursor m/z	897.98939	Identity Threshold	47.70
Charge	2+	Mass error	0.002024



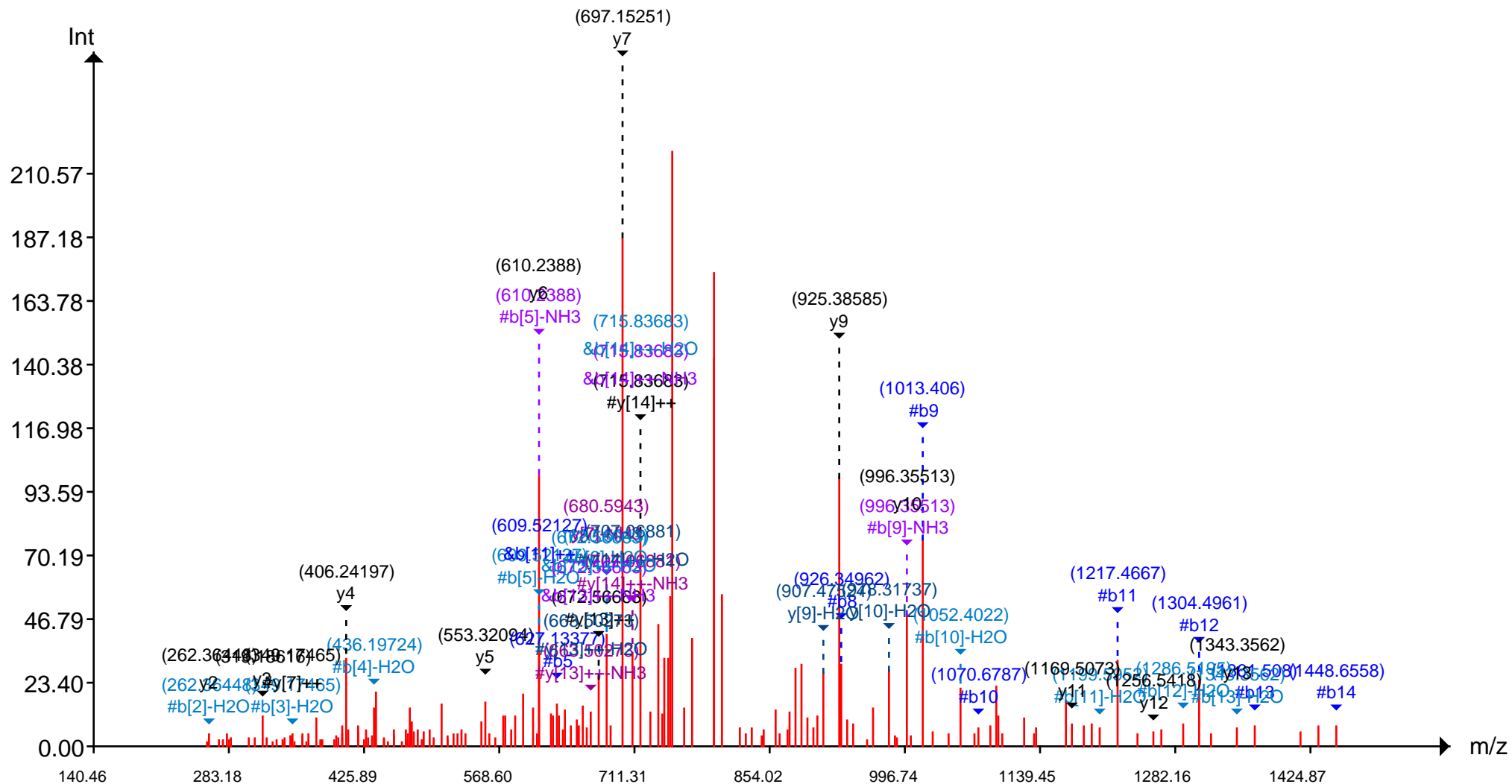
Info

Modified Sequence	AcD3-FDK<AcD3K*>DGDGTITTK<AcD3K*>ELGTVM<MoX*>R-COOH	Ion score	73.04
Protein	P62204	Confidence level	99.00%
Precursor m/z	1118.0628	Identity Threshold	47.78
Charge	2+	Mass error	0.002344



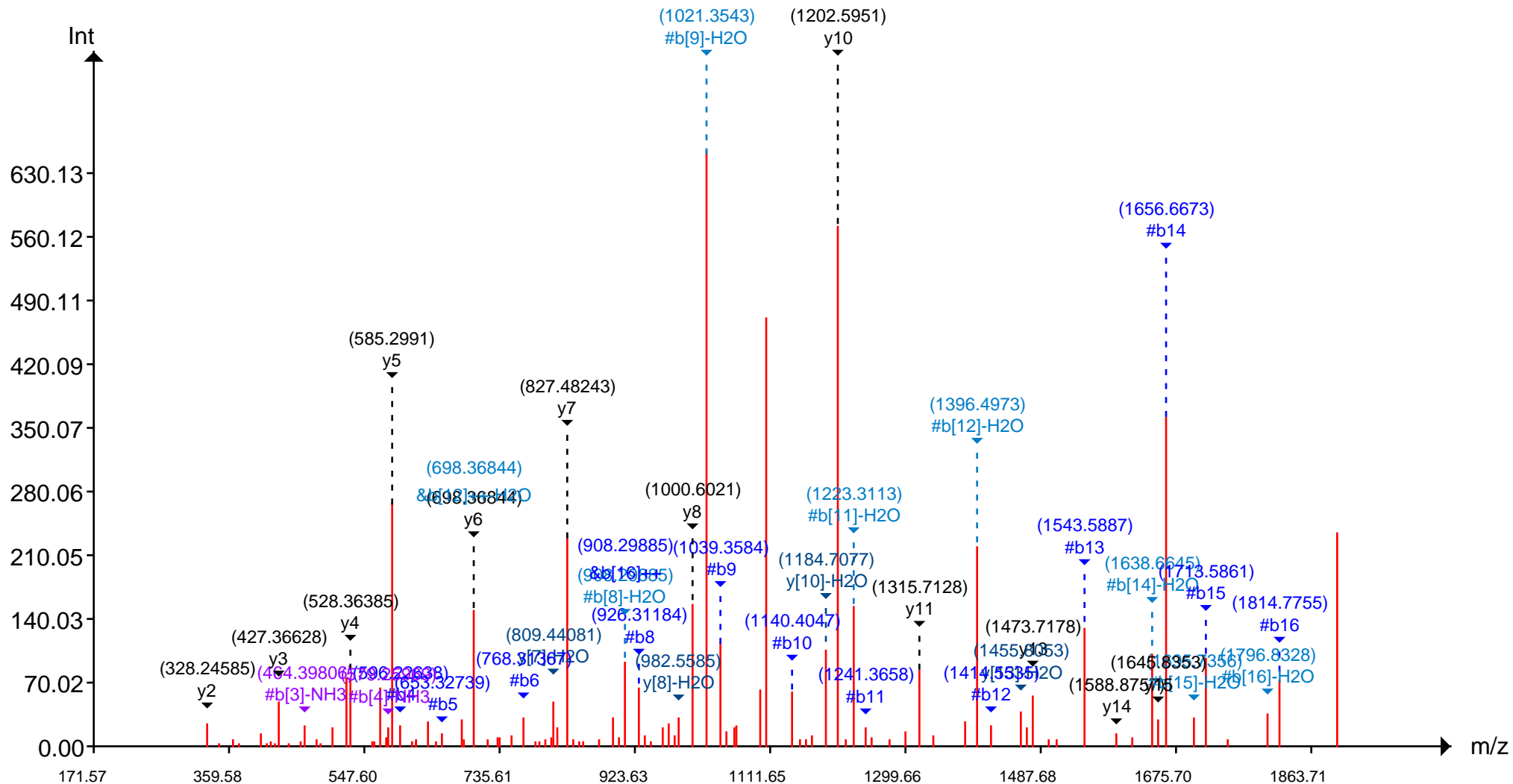
Info

Modified Sequence	AcD3-FSSSK<AcD3K*>ADLSGM<Mox*>SGSR<C13>-COOH	Ion score	59.69
Protein	Q9D154, Q5SV42	Confidence level	99.00%
Precursor m/z	814.89419	Identity Threshold	46.74
Charge	2+	Mass error	0.00123



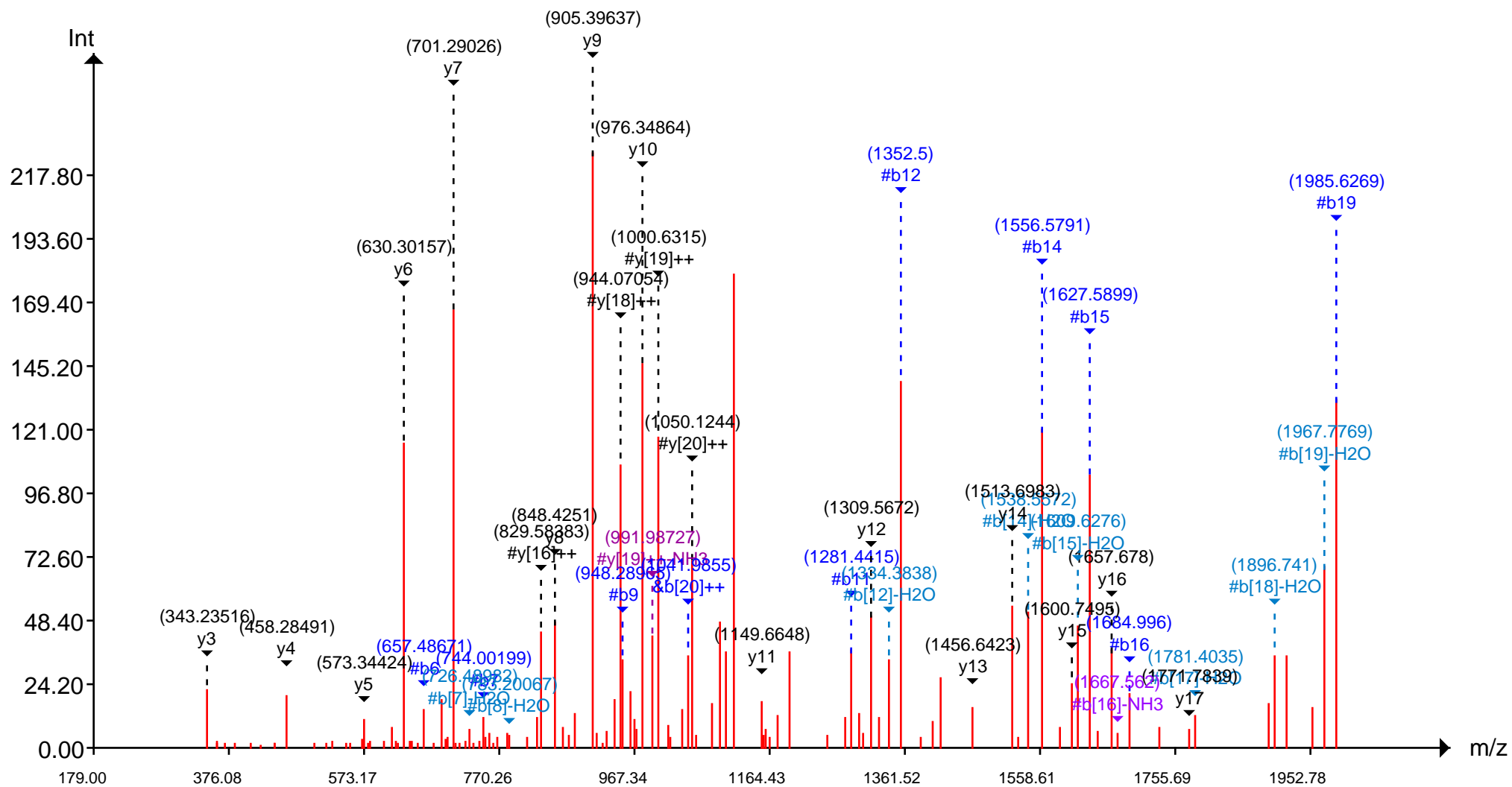
Info

Modified Sequence	AcD3-FSSSK<AcD3K*>ADLSGM<Mox*>SGSR-COOH	Ion score	62.0
Protein	Q9D154, Q5SV42	Confidence level	99.00%
Precursor m/z	811.88599	Identity Threshold	46.28
Charge	2+	Mass error	0.004971



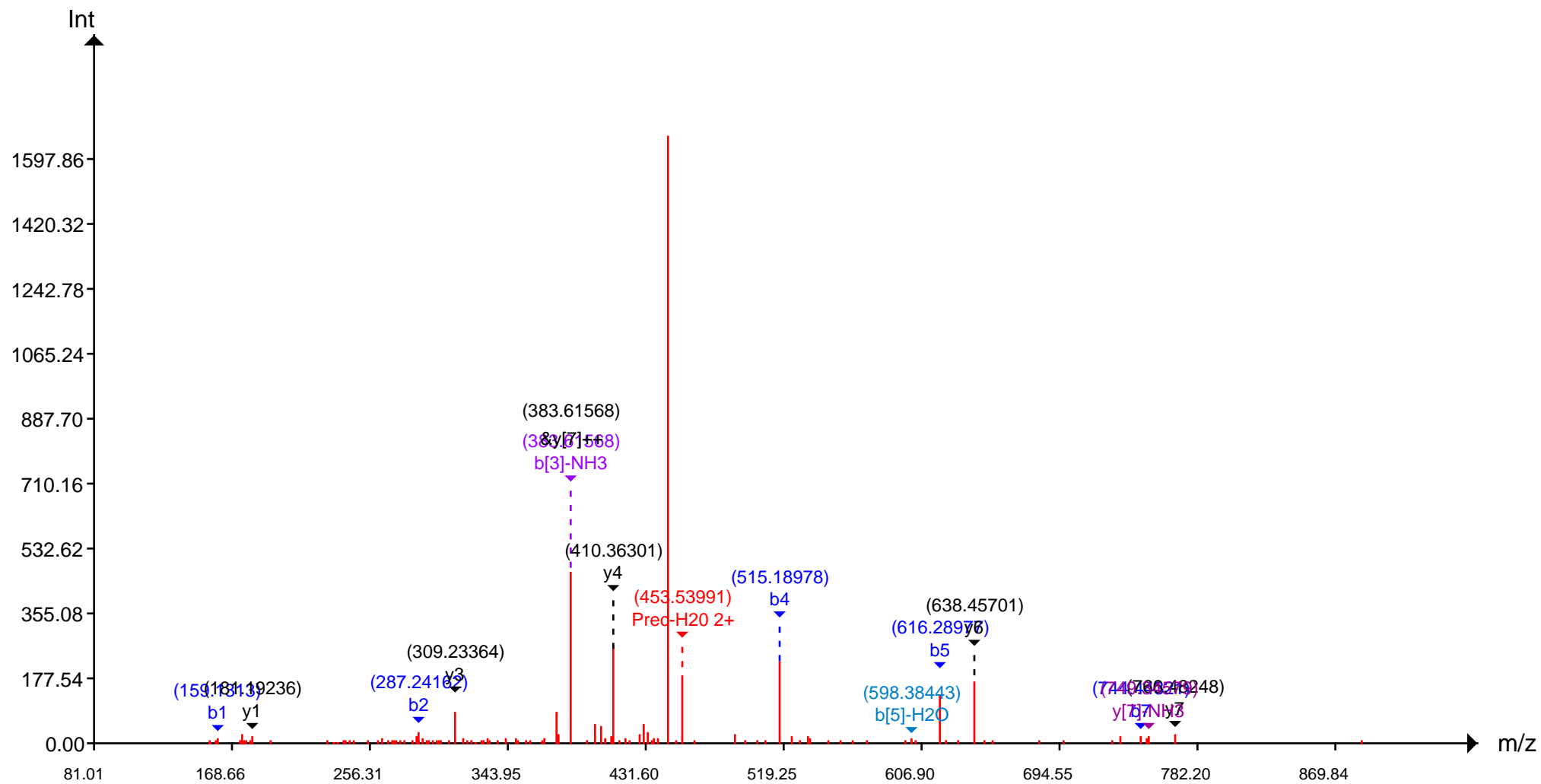
Info

Modified Sequence	AcD3-FDK<AcD3K*>DGDGTITTK<AcD3K*>ELGTVM<Mox*>R<C13>-COOH	Ion score	99.72
Protein	P62204	Confidence level	99.00%
Precursor m/z	1121.073	Identity Threshold	47.94
Charge	2+	Mass error	0.002603



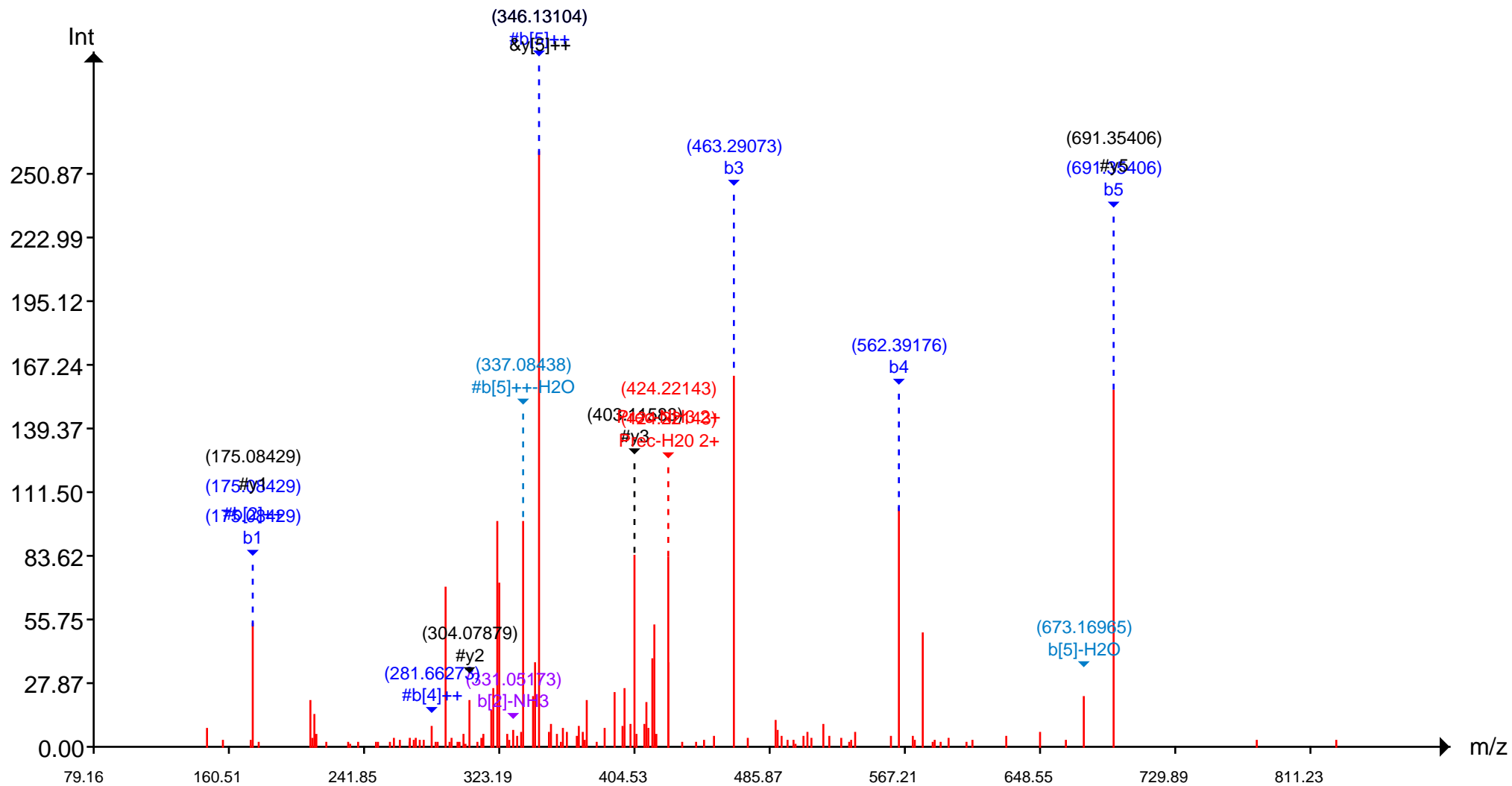
Info

Modified Sequence	AcD3-LVIDNGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR-COOH	Ion score	114.22
Protein	P63260	Confidence level	99.00%
Precursor m/z	1129.0304	Identity Threshold	45.62
Charge	2+	Mass error	0.004057



Info

Modified Sequence	AcD3-LQLDTAGR<C13>-COOH	Ion score	37.74
Protein	Q8C7B8	Confidence level	99.00%
Precursor m/z	462.76765	Identity Threshold	32.56
Charge	2+	Mass error	-3.55E-4



Info

Modified Sequence AcD3-EK<AcD3K*>DVER-COOH

Protein O35326

Precursor m/z 433.23045

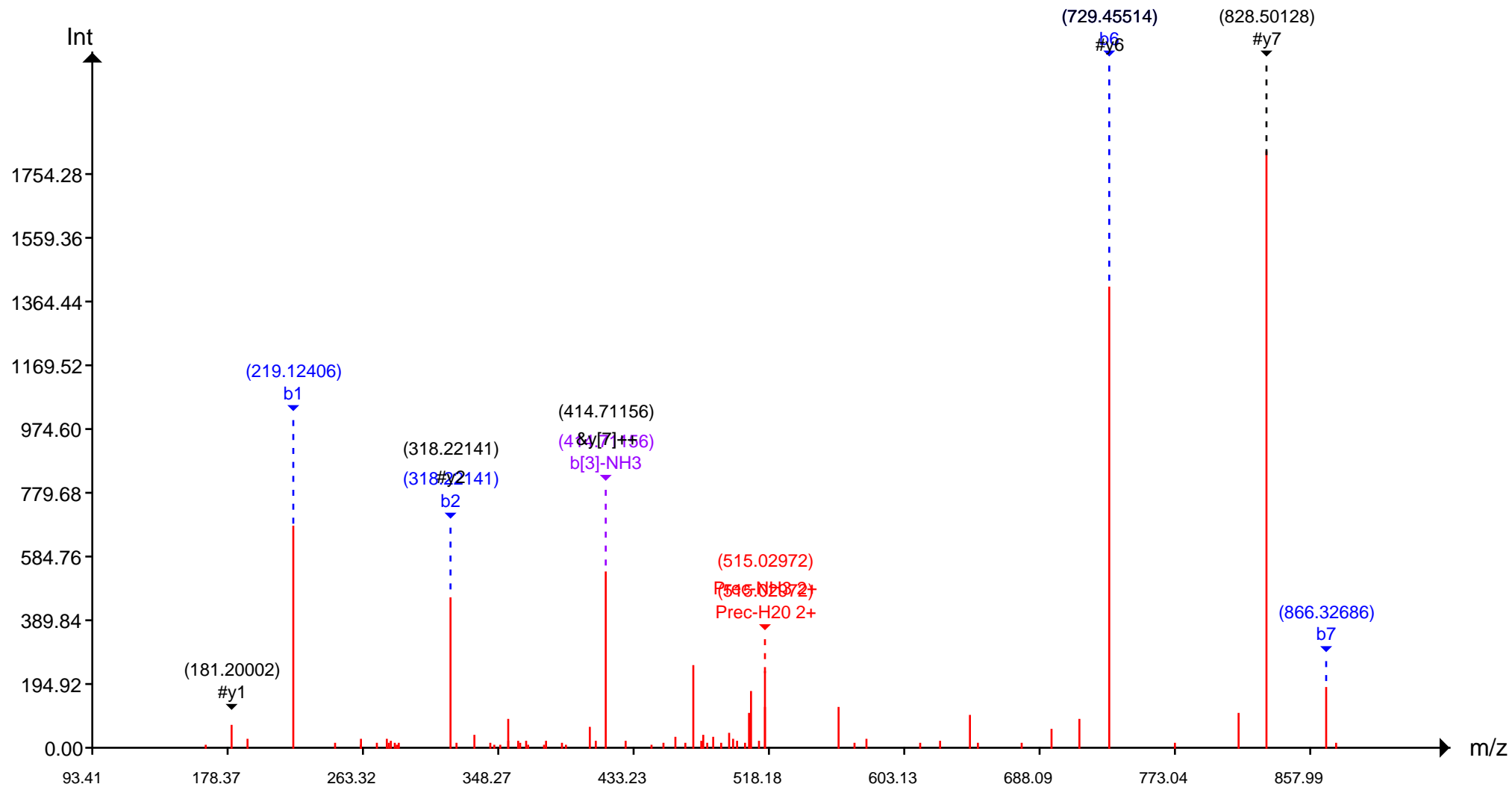
Charge 2+

Ion score 34.91

Confidence level 99.00%

Identity Threshold 33.69

Mass error 3.95E-4



Info

Modified Sequence AcD3-K<AcD3K*>VLQAVHR<C13>-COOH

Ion score 31.37

Protein O70622

Confidence level 99.00%

Precursor m/z 523.83756

Identity Threshold 29.12

Charge 2+

Mass error -4.89E-4