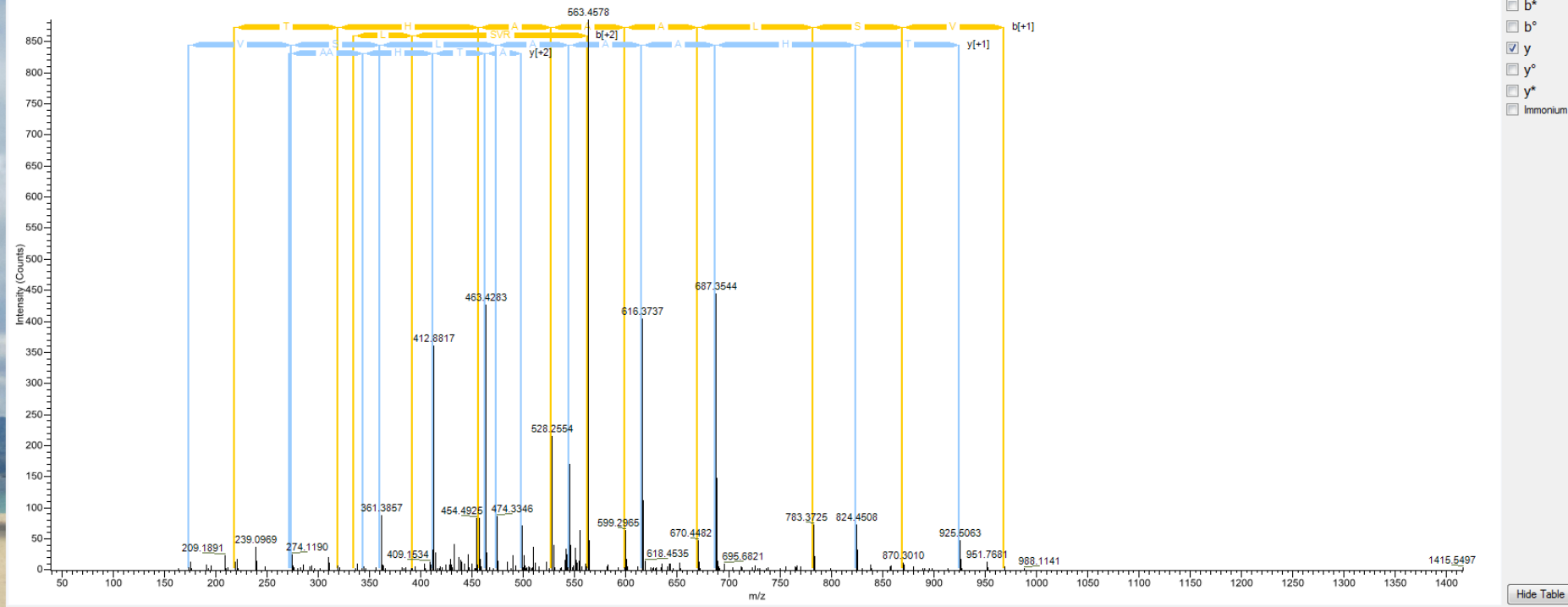


Peptide Consensus View

Sequence: FATHAALSVR  
 Charge: +2, Monoisotopic m/z: 572.67944 Da (+15.93 mmu/+27.82 ppm), MH+: 1144.35161 Da, RT: 39.34 min.

Extracted from: E:\XCalibur\data\CELL\cell224\_bd2.RAW #6237 RT: 39.34  
 ITMS, CID, z=+2, Mono m/z=572.67944 Da, MH+=1144.35161 Da, Match Tol.=1 Da



	(NH3+)-	A>	T>	H>	A>	A>	A>	<L	L>	<S	S>	V>	R-(COOH)
b (+1)	148.07570	219.11282	320.16050	457.21941	528.25653	599.29365	670.33077	783.41484	870.44687	969.51529	1125.61641		
b (+2)	74.54149	110.06005	160.58389	229.11334	264.63190	300.15046	335.66902	392.21106	435.72707	485.26128	563.31184		
b*(+1)	131.04915	202.08627	303.13395	440.19286	511.22998	582.26710	653.30422	766.38829	853.42032	952.48874	1108.58986		
b*(+2)	66.02821	101.54677	152.07061	220.60007	256.11863	291.63719	327.15575	383.69778	427.21380	476.74801	554.79857		
b'(+1)	130.06513	201.10225	302.14993	439.20884	510.24596	581.28308	652.32020	765.40427	852.43630	951.50472	1107.60584		
b'(+2)	65.53620	101.05476	151.57860	220.10806	255.62662	291.14518	326.66374	383.20577	426.72179	476.25600	554.30656		

FATHAALSVR

Tolerance: [ ] Da Charge State: All

- b
- b\*
- b°
- y
- y°
- y\*
- Immorium

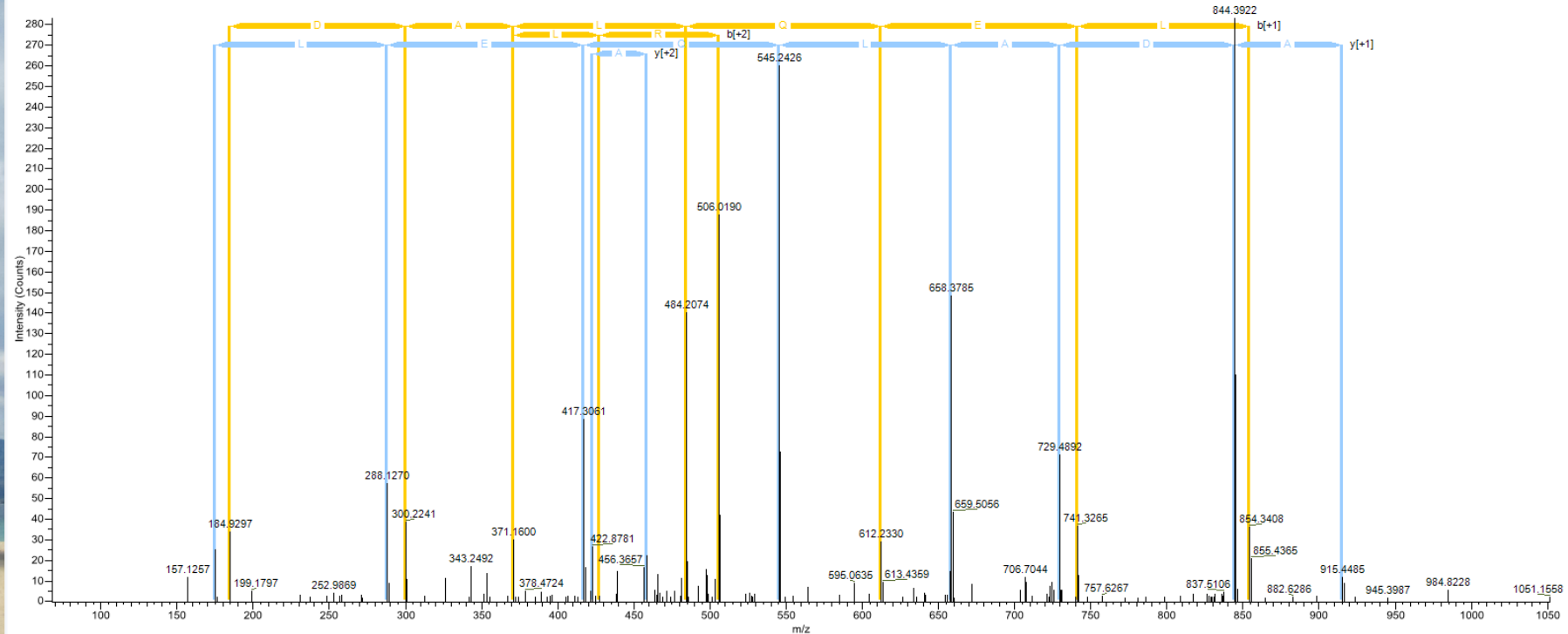
Hide Table

Value Type:  
 Theo. Mass [Da]

Peptide Consensus View

Sequence: LADALQELR  
 Charge: +2, Monoisotopic m/z: 515.34229 Da (+247.64 mmu/+480.54 ppm), MH+: 1029.67729 Da, RT: 47.14 min.

Extracted from: E:\XCalibur\data\CELL\cell224\_bd2\_RAW #7511 RT: 47.14  
 ITMS, CID, z=+2, Mono m/z=515.34229 Da, MH+=1029.67729 Da, Match Tol.=1 Da



- b
- b\*
- b°
- y
- y°
- y\*
- Immonium

Hide Table

	(NH3+)-	A>	D>	A>	<L	L>	Q>	E>	<L	L>	R-(COOH)
▶ b (+1)	114.09135	185.12847	300.15542	371.19254	484.27661	612.33519	741.37779	854.46186	1010.56298		
b (+2)	57.54931	93.06787	150.58135	186.09991	242.64194	306.67123	371.19253	427.73457	505.78513		
b*(+1)	97.06480	168.10192	283.12887	354.16599	467.25006	595.30864	724.35124	837.43531	993.53643		
b*(+2)	49.03604	84.55460	142.06807	177.58663	234.12867	298.15796	362.67926	419.22129	497.27185		
b*(+1)	96.08078	167.11790	282.14485	353.18197	466.26604	594.32462	723.36722	836.45129	992.55241		
b*(-2)	48.54403	84.06259	141.57606	177.09462	233.63666	297.66595	362.18725	418.72928	496.77984		

Value Type:  
 Theo. Mass [Da]

L [A] D [A] L [Q] E [L] R

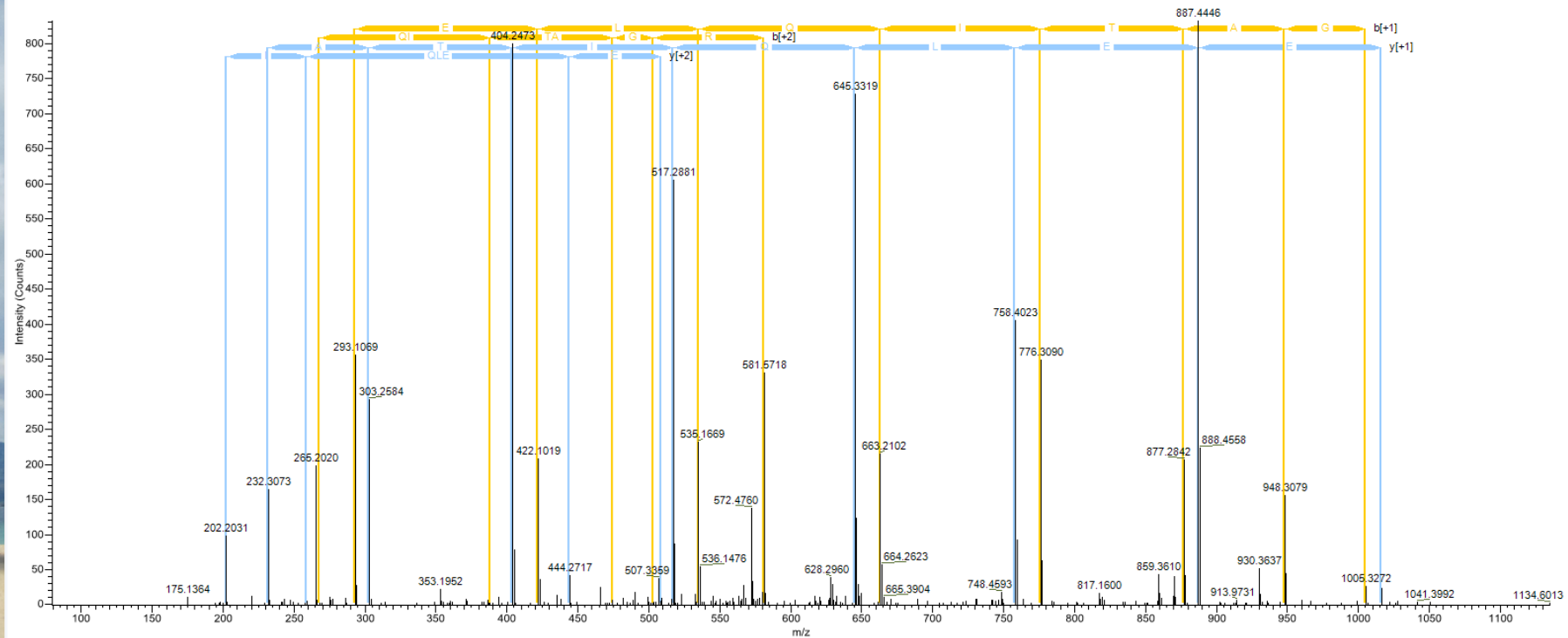
Tolerance: [ ] Da Charge State: All

z  
y  
b  
c

Peptide Consensus View

Sequence: YEELQITAGR  
 Charge: +2, Monoisotopic m/z: 590.48608 Da (-169.95 mmu/-287.81 ppm), MH+: 1179.96489 Da, RT: 44.94 min.

Extracted from: E:\XCalibur\data\CELL\cell224\_bd5\_RAW\_#7191 RT: 44.94  
 ITMS, CID, z=+2, Mono m/z=590.48608 Da, MH+=1179.96489 Da, Match Tol.=1 Da



- b
- b\*
- b<sup>o</sup>
- y
- y<sup>o</sup>
- y\*
- Immonium

Hide Table

	(NH3+)-	E>	E>	<L	L>	Q>	<I	I>	<T	T>	A>	G>	<R	R-(COOH)
▶ b (+1)	164.07060	293.11320	422.15580	535.23987	663.29845	776.38252	877.43020	948.46732	1005.48879	1161.58991				
b (+2)	82.53894	147.06024	211.58154	268.12357	332.15286	388.69490	439.21874	474.73730	503.24803	581.29859				
b*(+1)	147.04405	276.08665	405.12925	518.21332	646.27190	759.35597	860.40365	931.44077	988.46224	1144.56336				
b*(+2)	74.02566	138.54696	203.06826	259.61030	323.63959	380.18162	430.70546	466.22402	494.73476	572.78532				
b*(+1)	146.06003	275.10263	404.14523	517.22930	645.28788	758.37195	859.41963	930.45675	987.47822	1143.57934				
b*(+2)	73.53365	138.05495	202.57625	259.11829	323.14758	379.68961	430.21345	465.73201	494.24275	572.29331				

III

Value Type:  
 Theo. Mass [Da]

Y[E]E[L]Q[I]T[A]G[R]

z  
y  
b  
o

Tolerance: [ ] Da Charge State: All