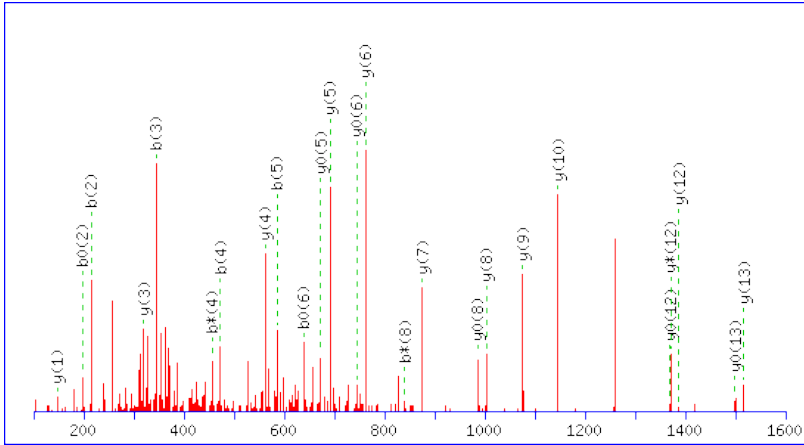


MS/MS Fragmentation of **SQEQLAAELAEpY<sup>424</sup>TAK**  
 Found in EZRI\_HUMAN, **P15311**|EZRI\_HUMAN Ezrin  
 Homo sapiens (Human)  
 Match from (866.393092,2+)



Monoisotopic mass of neutral peptide Mr(calc): 1730.78

Variable modifications:

Y12 : Phospho (Y)

Ions Score: 75 Expect: 4.1e-06

Matches (Bold Red): 25/162 fragment ions using 46 most intense peaks

#	b	b <sup>++</sup>	b <sup>*</sup>	b <sup>+++</sup>	b <sup>0</sup>	b <sup>0++</sup>	Seq.	y	y <sup>++</sup>	y <sup>*</sup>	y <sup>+++</sup>	y <sup>0</sup>	y <sup>0++</sup>	#
1	88.04	44.52			70.03	35.52	S							15
2	<b>216.10</b>	108.55	199.07	100.04	<b>198.09</b>	99.55	Q	1644.75	822.88	1627.73	814.37	1626.74	813.87	14
3	<b>345.14</b>	173.07	328.11	164.56	327.13	164.07	E	<b>1516.69</b>	758.85	1499.67	750.34	<b>1498.68</b>	749.84	13
4	<b>473.20</b>	237.10	<b>456.17</b>	228.59	455.19	228.10	Q	<b>1387.65</b>	694.33	<b>1370.62</b>	685.82	<b>1369.64</b>	685.32	12
5	<b>586.28</b>	293.65	569.26	285.13	568.27	284.64	L	1259.59	630.30	1242.57	621.79	1241.58	621.29	11
6	657.32	329.16	640.29	320.65	<b>639.31</b>	320.16	A	<b>1146.51</b>	573.76	1129.48	565.24	1128.50	564.75	10
7	728.36	364.68	711.33	356.17	710.35	355.68	A	<b>1075.47</b>	538.24	1058.44	529.73	1057.46	529.23	9
8	857.40	429.20	<b>840.37</b>	420.69	839.39	420.20	E	<b>1004.43</b>	502.72	987.41	494.21	<b>986.42</b>	493.72	8
9	970.48	485.75	953.46	477.23	952.47	476.74	L	<b>875.39</b>	438.20	858.36	429.69	857.38	429.19	7
10	1041.52	521.26	1024.49	512.75	1023.51	512.26	A	<b>762.31</b>	381.66	745.28	373.14	<b>744.30</b>	372.65	6
11	1170.56	585.79	1153.54	577.27	1152.55	576.78	E	<b>691.27</b>	346.14	674.24	337.63	<b>673.26</b>	337.13	5
12	1413.59	707.30	1396.57	698.79	1395.58	698.30	Y	<b>562.23</b>	281.62	545.20	273.10	544.22	272.61	4
13	1514.64	757.82	1497.61	749.31	1496.63	748.82	T	<b>319.20</b>	160.10	302.17	151.59	301.19	151.10	3
14	1585.68	793.34	1568.65	784.83	1567.67	784.34	A	218.15	109.58	201.12	101.07			2
15							K	<b>147.11</b>	74.06	130.09	65.55			1