

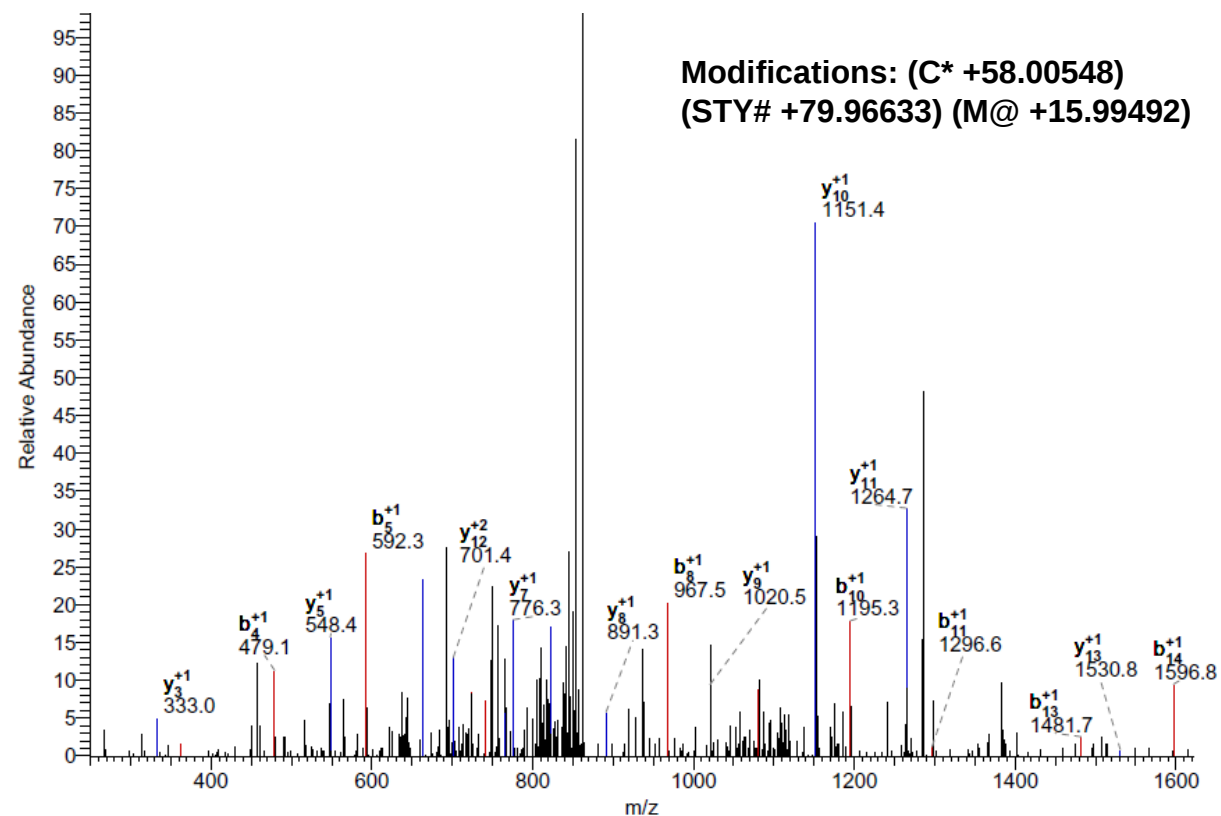
a) MTCMSQLER NIETIINTFH QYSVKLGHPD TLNQGEFKEL VRKDLQNFLK
 KENKNEK**VIE** **HIMEDLDTNA** **DK**QLSFEEFI MLMARLTWAS HEKMHEGDEG
 PGHHHKPGLG EGTP

Protein Coverage:

Sequence
VIEHIMEDLDTNADK

	MH+	% Mass	AA	% AA
Totals:	1742.83	13.17	58 - 72	13.16
	1742.83	13.17	15	13.16

b)



c)

AA	B ions	Y ions
V	100.08	
I	213.16	1643.76
E	342.20	1530.67
H	479.26	1401.63
I	592.35	1264.57
M	723.39	1151.49
E	852.43	1020.45
D	967.46	891.41
L	1080.54	776.38
D	1195.57	663.29
T	1296.61	548.27
N	1410.66	447.22
A	1481.69	333.18
D	1596.72	262.14
K		147.11