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Peptide number	Experimental mass	Predicted mass	Position of peptide	MC*	Peptide sequence
1	1458.74	1459.7362	70-80	0	VIHNELELYCR
2	894.3085	894.4468	118 - 124	0	WYTAPTR
3	844.2945	844.4206	125-132	1	GPAANCRR
4	1402.479	1401.7597	205-216	1	TASYLLIHDGRR
5	892.2887	891.5047	238 - 244	1	SWIRTTK
6	1566.675	1565.8645	242-255	1	TTKVTGDHPLVIER
7	1235.465	1235.6742	245-255	0	VTGDHPLVIER
8	1940.871	1940.9422	291 - 309	0	SIFGPGGTPSLFPTSCSTK
9	2031.072	2032.9289	323 - 339	0	LMLFGATLDDQAFCCSR

\* Number of modifications