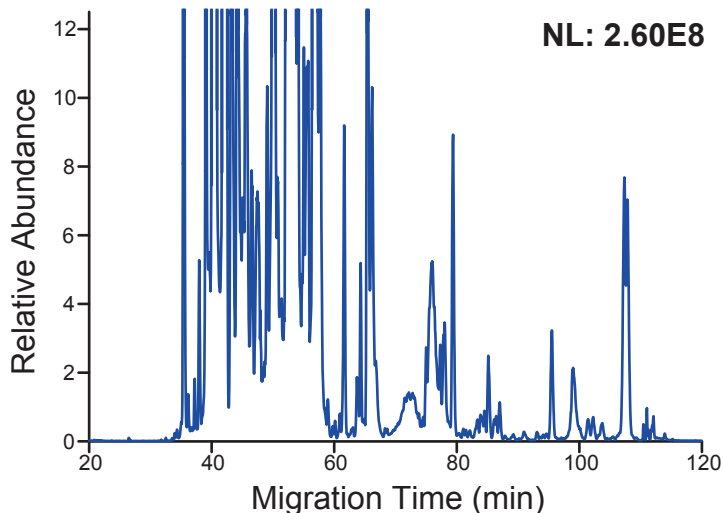




Protein-Spectrum-Match #827 for Spectrum #1119

PrSM ID:	827	Scan(s):	2427	Precursor charge:	9
Precursor m/z:	1030.5272	Precursor mass:	9265.6797	Proteome mass:	9265.6800
# matched peaks:	79	# matched fragment ions:	60	# unexpected modifications:	0
E value:	2.11e-48	P value:	2.11e-48	Q-value (Special FDR):	0

↑
General proteoform information

1 M T M S L L E L V F E L K L L E A T K I V Q Q A L I D T I T L L L Q M E L I L E 30
 31 E L L K L E L K L N L N S L L S L Q L E L V Q N A Q H Q R E E L E L R E N N H L 60
 61 K E Q Q I N L G W L Q E L R L Q A L L L G R M E E T V 81

↑
Protein sequence, observed fragmentation pattern and modifications