

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 Proteoform 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 K L L E A K I V Q Q L A I D T L I T L L L Q M E L I L E 30
31 E L L K L E K L N L N S L L S Q L E L V L 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 I V 81



Protein sequence, observed fragmentation pattern and modifications