

Sequence	Prob	Mascot Ion score	Spectral count
SFLEFAEDVIQVPR	100%	69.34	75
MEELVVEVR	100%	70.36	52
VLVASSVVAGESQKPELK	100%	73.15	36
LIQEIVDK	99%	36.68	27
LQIDEQLR	99%	48.9	27
DINESDEVEVYSR	100%	84.36	19
VGPNAPPEEK	100%	59.24	19
FPPVGYNK	99%	32.39	18
EDLMGLAIGTHGANIQQAR	100%	68.42	17
DVHEDSITVAFENNWQPDR	100%	90.12	15
LDVPEDLR	99%	33.81	14
TTDGSLQIR	100%	70.09	13
SYVTDDGQGMGR	100%	58.75	12

Prob= probability, indicative of the peptide identification confidence; Mascot Ion score= Mascot algorithm metric for evaluating the identity of the peptide/protein; Spectral count= based on total MS/MS data.