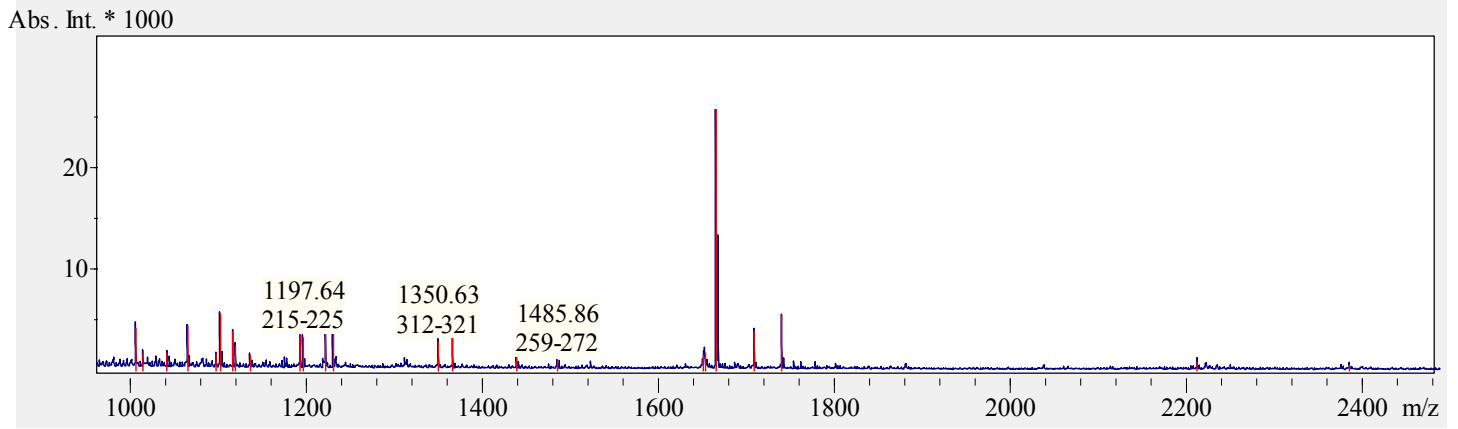


A



B

Protein View | Match Errors | MS/MS Fragments | MSMS Analysis

Protein: Peak threshold: 0.0

Intensity coverage: 3.3% (7374 cnts) Sequence coverage MS: 9.4% Sequence coverage MS/MS: 0.0% pI: 8.6 MW (kDa): 43.3

10	20	30	40	50	60	70	80	90	100	110	120
DLNCKPDYSY	AIAKSDRVGP	LGAEDLTTIW	KDYSQEMTLE	DIMVIANCKD	GKFFVYLPRCT	RETRYLATLH	SRALPTNVVF	EKLFEGLRQG	DTVEMEDNFE	FGLCPCDAKP	IVRGRFNITL
130	140	150	160	170	180	190	200	210	220	230	240
LNGSAFQMYC	PIGWIGSVSC	MLANRDILKT	AVVRTYRRSK	PFYRQGCIT	QKTLGEDLYN	CDLGGNNTCV	TGDQLQYTG	SIESCWCWGF	KFQKSEGLPH	YPIGKCRLN	ETGYRLVDGT
250	260	270	280	290	300	310	320	330	340	350	360
SCNREGVAIV	PIGQVKCKIG	DTIVQVIALD	TKLGPMPCKP	HEIISSEGPV	EKTACTFNVT	KILKNKYFEP	RDSYFQQYML	RGEYQYWFDL	EVIDHHRDYF	AESILVVVVA	LLGGRYVLWL
370	380										
LVTYMLVLEQ	KALG										