

Supplementary material

Table S1 Peptide sequence data for the 43 different expression proteins identified by MS

Spot	Peptide sequences
49	VLVISGER, KFVLPENADMEK, DGVLTVTVDKLPPEPK, ELPGAYAFVVDMPGLGSGDIK
57	NLVDHLQER, TEFDVVIEVPSSAR
66	NLVDHLQER, TEFDVVIEVPSSAR, VLELGDAIAGLTLEEAR
76	EHNSSPGYYDGR, KFETLSYLPPLSTEALLK, QVQCVSFIAPRPPGCEESGK, QVQCVSFIAPRPPGCEESGKA
84	CHFVAIDIFNGK, DDLKLPTDDALLTQIK
97	LEDKPVGQYLEK
109	SLNENDVVEFEVITGDDGR, GFGFISPEDGSEDLFVHQSAIK
112	GVGPGDFYYHGLATAGNTNLLIK
113	EYLTFLSGFR, DAAEYVYDVPEGWK, DAAEYVYDVPEGWKER, DADGQVYYEYEVAGAGAHSLISVTCAR, LAPVGAVLDNLALSDVGLQDQISSADDV TSAER
123	VTFLLDDAQVK
127	ESEIYHCR, YPGGAFDPLGFSK, FKESEIYHCR
138	LLIDKLLK
142	WSPELAAACEVWK
145	LEDKPVGQYLEK

148 GLPEDVIAGWR,
NWSVPDALAHVK,
TIFSMDSFVNTR
LVDLANKPDWLF TINPEGK
149 GGGGGGGGCFSCGESGHFSR,
DCPQGGGGGGGYGGGGYGGGGGGGR,
SLNENDTVEFEVITGDDGR,
GFGFISPEDGSEDLFVHQSAIK
167 AVCTPEFYLFK,
YFPYLYDESEQEVAK
218 AEQYLADSGLPYTIIR,
ADVAEVCIQALLFEEAK,
KAEQYLADSGLPYTIIR
220 SEVSSLIAELASAAAER
242 SEVSSLIAELASAAAER
261 GLSFDEGMEER,
SEVSSLIAELASAAAER
306 YCTFPHR,
QFSDVAAER,
GLPMEELWK,
VMLRPACPGSGVTAGGAVR
316 IQGIGAGFVPR,
VDIFIGIGTGGTISGAGR,
SVLVEATSGNTGIGLAFIAASR
345 VPFVCGCR,
SIAAPYDLVMQTK,
GGVIMDVVNAEQAR,
IAEEAGACAVMALER
347 SYIEFAER,
IFISELLVDELSAPSQEIR
366 VIGFDNLR,
EYPDAYVR,
EHNSSPGYYDGR

369 TFSYYHPESR,
 IFLEDGHQIGCAQSYAK,
 NHFPFFDMAYQGFASGDPER
 375 LFGVTTLDVVR,
 KLVGVTTLDVVR,
 GFVGGDDQLGEALEGADLVIIPAGVPR,
 LNPLVSSLSLYDIAATPGVAADVSHINTR
 394 FYWAPTR,
 IVDTFPGQSIDFFGALR,
 VPIIVTGNDFSTLYAPLIR
 428 ENVADAQATFLAR,
 VGAEVIAEYTVAAALR,
 YGGAAAAGDAAASESLHVAGYKY
 432 DRYPGVLVDVVASAR,
 SRGDDDCLLPLELWAEIAK
 441 DTDILAAFR,
 DNGLLLHIHR,
 FVFCAEAIYK,
 LTYYTPEYETK,
 TFQPPHGIQVER
 444 FYWAPTR,
 GLAYDISDDQQDITR,
 IVDTFPGQSIDFFGALR,
 VPIVVTGNDFSTLYAPLIR
 447 AHGGVSVFGGVGER,
 IVGNEHYETAQR,
 FVQAGSEVSALLGR,
 LPYIYNALVVQSR
 455(JMC) AHGGVSVFGGVGER,
 IVGNEHYETAQR,
 FVQAGSEVSALLGR,
 VGLTALTMAEYFR,
 LPYIYNALVVKSR,

455(LYC) VALVYQMNEPPGAR
IAQIPVSEAYLGR,
ASSVAQVVTNFQER
571 HYPVLFER,
FWPTTCR,
APPMQFTGK,
EYAAFPAAWLR
640 GLAYDISDDQQDITR,
IVDTFPGQSIDFFGALR,
VPIVVTGNDSTLYAPLIR
653 GYFVLNDILR,
EFAQSFFLAPQEK,
NLPLHATPSQLEEEFKR
667 FVFCAEALYK,
TFKGPPHGIQVER
682 EAAWGLAR,
EAAYYQQGAR,
IVDILVEQGIVPGIK,
TWGGRPENVAQAALLR,
GLVPLVGSNDESWCQGLDGLASR,
YAAISQDNGLVPIVEPEIMLDGEHGIER
689 EAAWGLAR,
AAQEALLR,
TWGGRPENVK,
EAAYYQQGAR,
GLVPLTGSNDESWCQGLDGLASR
692 QFGSLAEYTAVEEK,
GPFSFPQVVEAFSYLETGR
