| NP_068576.1 XP_011543854.1 NP_001358344.1 XP_011543851.1 XP_011543853.1 | 1 10 MSSSSWLLLSLVAVT MSSSSWLLLSLVAVT MSSSSWLLLSLVAVT MSSSSWLLLSLVAVT MSSSSWLLLSLVAVT | AAQSTIEEQ AAQSTIEEQ AAQSTIEEQ | AKTFLDKFNHE AKTFLDKFNHE AKTFLDKFNHE | AEDLFYQSS: AEDLFYQSS: AEDLFYQSS: | LASWNYNTNI' LASWNYNTNI' LASWNYNTNI' | FEENVQ FEENVQ |
|---|--|--|---|--|---|----------------------------|
| NP_068576.1 XP_011543854.1 NP_001358344.1 XP_011543851.1 XP_011543853.1 | 70 NMNNAGDKWSAFLKE NMNNAGDKWSAFLKE NMNNAGDKWSAFLKE NMNNAGDKWSAFLKE NMNNAGDKWSAFLKE | QSTLAQMYP QSTLAQMYP QSTLAQMYP | LQEIQNLTVKI LQEIQNLTVKI | QLQALQQNG QLQALQQNG QLQALQQNG | SSVLSEDKSKI SSVLSEDKSKI SSVLSEDKSKI | RLNTIL RLNTIL RLNTIL |
| NP_068576.1 XP_011543854.1 NP_001358344.1 XP_011543851.1 XP_011543853.1 | 130 NTMSTIYSTGKVCNP NTMSTIYSTGKVCNP NTMSTIYSTGKVCNP NTMSTIYSTGKVCNP NTMSTIYSTGKVCNP | DNPQECLLL DNPQECLLL DNPQECLLL | EPGLNEIMANS EPGLNEIMANS EPGLNEIMANS | LDYNERLWAI LDYNERLWAI LDYNERLWAI | WESWRSEVGK WESWRSEVGK WESWRSEVGK | QLRPLY QLRPLY QLRPLY |
| NP_068576.1 XP_011543854.1 NP_001358344.1 XP_011543851.1 XP_011543853.1 | 190 EEYVVLKNEMARANH EEYVVLKNEMARANH EEYVVLKNEMARANH EEYVVLKNEMARANH | YEDYGDYWR YEDYGDYWR YEDYGDYWR | GDYEVNGVDGY GDYEVNGVDGY GDYEVNGVDGY | DYSRGQLIE DYSRGQLIE DYSRGQLIE | DVEHTFEEIKI DVEHTFEEIKI DVEHTFEEIKI | PLYEHL PLYEHL |
| NP_068576.1 XP_011543854.1 NP_001358344.1 XP_011543851.1 XP_011543853.1 | 250 HAYVRAKLMNAYPSY HAYVRAKLMNAYPSY HAYVRAKLMNAYPSY HAYVRAKLMNAYPSY HAYVRAKLMNAYPSY | ISPIGCLPA ISPIGCLPA ISPIGCLPA | HLLGDMWGRFW HLLGDMWGRFW HLLGDMWGRFW | TNLYSLTVP: TNLYSLTVP: TNLYSLTVP: | FGÖKPNIDVTI FGOKPNIDVTI FGOKPNIDVTI | DAMVDQ DAMVDQ DAMVDQ |
| NP_068576.1 XP_011543854.1 NP_001358344.1 XP_011543851.1 XP_011543853.1 | 310 AWDAQRIFKEAEKFF AWDAQRIFKEAEKFF AWDAQRIFKEAEKFF AWDAQRIFKEAEKFF AWDAQRIFKEAEKFF | VSVGLPNMT VSVGLPNMT VSVGLPNMT | QGFWENSMLTD QGFWENSMLTD QGFWENSMLTD | PGNVQKAVC PGNVQKAVC PGNVQKAVC | HPTAWDLGKGI HPTAWDLGKGI HPTAWDLGKGI | DFRILM DFRILM DFRILM |
| NP_068576.1 XP_011543854.1 NP_001358344.1 XP_011543851.1 XP_011543853.1 | 370 CTKVTMDDFLTAHHE CTKVTMDDFLTAHHE CTKVTMDDFLTAHHE CTKVTMDDFLTAHHE CTKVTMDDFLTAHHE | MGHIQYDMA MGHIQYDMA MGHIQYDMA | YAAQPFLLRNG YAAQPFLLRNG YAAQPFLLRNG | ANEGFHEAV ANEGFHEAV ANEGFHEAV | GEIMSLSAATI GEIMSLSAATI GEIMSLSAATI | PKHLKS PKHLKS PKHLKS |
| NP_068576.1 XP_011543854.1 NP_001358344.1 XP_011543851.1 XP_011543853.1 | 430 IGLLSPDFQEDNETE IGLLSPDFQEDNETE IGLLSPDFQEDNETE IGLLSPDFQEDNETE | INFLLKQAL INFLLKQAL INFLLKQAL | TIVGTLPFTYM TIVGTLPFTYM TIVGTLPFTYM | LEKWRWMVF LEKWRWMVF LEKWRWMVF | KGEIPKDQWMI KGEIPKDQWMI KGEIPKDQWMI | KKWWEM KKWWEM KKWWEM |
| NP_068576.1 XP_011543854.1 NP_001358344.1 XP_011543851.1 XP_011543853.1 | 490 KREIVGVVEPVPHDE KREIVGVVEPVPHDE KREIVGVVEPVPHDE KREIVGVVEPVPHDE | TYCDPASLF: TYCDPASLF: TYCDPASLF: | HVSNDYSFIRY HVSNDYSFIRY HVSNDYSFIRY | YTRTLYQFQI YTRTLYQFQI YTRTLYQFQI | FQEALCQAAKI FQEALCQAAKI FQEALCQAAKI | HEGPLH HEGPLH HEGPLH |

| | 5 5 O | 560 | 570 <u>.</u> | 580 | 590 | 60 Ó |
|---|--|--------------------------|---------------------------|---------------------------------------|-------------------------|---------------|
| NP_068576.1 | KCD I S N S T E A G Q K L F | NMLRLGKSEI | WTLALENVVG | AKNMNVRPLL | NYFEPLFTW | LKDQNK |
| XP_011543854.1 NP 001358344.1 | KCDISNSTEAGQKLL KCDISNSTEAGOKLF | NMI DI CECEI | NET AT ENVIS | AKNMNVRPLL | NVEEDIETW | TADONA |
| XP 011543851.1 | KCDISNSTEAGOKLE | | | | | |
| XP_011543853.1 | KCDISNSTEAGQKLF | NMLRLGKSEI | WTLALENVVG | AKNMNVRPLL | NYFEPLFTW | LKDQNK |
| NP_068576.1 XP 011543854.1 | 610 NSFVGWSTDWSPYAD | 620 QSIKVRISLE | 630 Ksalgdkayew | 640 NDNEMYLFRS | 650 SVAYAMRQY | 660 FLKVKN |
| NP_001358344.1 | NSFVGWSTDWSPYAD | QSIKVRISLE | KSALGDKAYEW | NDNEMYLFRS | SVAYAMRQY | FLKVKN |
| XP_011543851.1 | NSFVGWSTDWSPYAD | | | | | |
| XP_011543853.1 | NSFVGWSTDWSPYAD | <u> 680</u> | 690 | 700 | 710 | 720 |
| NP 068576.1 | OMILFGEEDVRVANL | · . | · | · · · · · · · · · · · · · · · · · · · | <u> </u> | • |
| XP_011543854.1 | EEDVRVANL | | TAPKNVSDII | | | |
| NP_001358344.1 XP 011543851.1 | QMILFGEEDVRVANL QMILFGEEDVRVANL | | | | | |
| XP_011543851.1 XP_011543853.1 | QMILF GEEDVRVANL | | | | | |
| | 730 | 740 | 750 | 760 | 770 | 780 |
| NP 068576.1 | SLEFLGIOPTLGPPN | | • | • | | • |
| XP_011543854.1 | SLEFLGIQPTLGPPN | QPPVSIWLIV | /FGVVMGVIV V | GIVILIFTGI | RDRKKKNKA | RSGENP |
| NP_001358344.1 XP 011543851.1 | SLEFLGIQPTLGPPN SLEFLGIQPTLGPPN | | | | | |
| XP_011543851.1 XP_011543853.1 | SLEFLGIQPTLGPPN | | | | | LGKSWL |
| NP_068576.1 XP_011543854.1 NP_001358344.1 XP_011543851.1 | 790 YASIDISKGENNPGF YASIDISKGENNPGF YASIDISKGENNPGF TAILKD | QNTDDVQTSE QNTDDVQTSE | 7 | | | |
| XP_011543853.1 | | | | | | |