Sequence: cds.comp144315_c1_seq1.prot (2500 aa) Features: 6 total MTVACIMVRLLVLTISLGLAACQRPKSIETVYSNGKEYLYDYESQILSGLPQGSRIYSGFKLKAQ 10 20 30 150 160 220 Pep_mark_1 290 340 350 360 420

KSEGGLQHSNYMYRSSYDKKTRLGHSYEQVLLHSAKERFPRAFAMQYTATLMGKTVDLFELGYKA 780 KGVDTIVDRMYGVLGDTSMTNIVRKLFDTKFMKTVSPMEQELLKMKEKLEVKSRFPEIPEGHVYM

800 810 820 blast hit 1 blast_hit_2

Pep_mark_2

KLFGQELIYFDLPSMLGPSLTSSPDGSDGSQPNDLKLTKILPELMNVVISLNRLLKTQKSAFFPL 850 860 890 880 blast hit 2

MSTMLPTETGLPLRFGIDKVAALKIDGDVKFEKEKDPNTSEKKFLLSSFDSLLKFRFGCVFEIRG
blast_hit_2
NMEVDFKYIRSTLSLQGRVSVEAPVTLAVNVDTTKSKLLLHLQDEKEFYPVKMVSRPSSEIRRMP 980 990 1000 1010 1020 1030 1040
SDADRYPIPPDVKEIIIIDGADVRLFSYDIGKDTLGRKLTVNGEMTVSRDDSCPLPPLNGRQVLY
IKYEPLTFTGLDIQMQRVREVSTKDPSISTVSKQTEPWWLPSWLSGTSTEEQSYSGPDRNKFEKN 1110 1120 1130 1140 1150 1160 1170
LDDPNKGFIVLDKTDLSLPYIKKKFERETELMKTGVKCGMVIRVSDHVPQPKHQIQLTTLYQSDS 1180 1190 1200 1210 1220 1230
SGSQQWYSIELMKAPKSDERMPWKMMTTVKLDLPETVVKPSFLLDNEYQKFIQDQIKVIVTENGK 1240 1250 1260 1270 1280 1290 1300
QTDSLLDLGSKFLTKAIKQLYKLQRRDSTDKIIEDVVSHVLERSVDKDSAQLRDSVQEQRKLLKE
TDKLKKQLSEPRKLKSQLELLKKILLKTKLVSETVSKNLDVIPWKQQKPEDSVMVLYVVSRQTQL 1370 1380 1390 1400 1410 1420 1430
KKKLRKIVEKLHPLLTTNEKLDLESKLKEQKQSLDDIRMKQNTLSLAKQSTEEQKSRIEMEPKER
SLQRQLLDIEQILISVSMERKNGKPFPSNKEKTYEIEKYLNKVIEKTKKVKDEMDERLLQKVKID 1500 1510 1520 1530 1540 1550 1560
KMLRVKLLEVLLHQKLTVLDIEDKLLYQDKKQKPLFLRSPYSSYNDLEYPLKKENIKLKDDLQRE
FGDLARRLAVSVDCEIETDSLVSFLSSEDQQQMDTLQKLKINQLRMLELLRTKSHPSETETDLPT 1630 1640 1650 1660 1670 1680 1690
LDSVKETVHLIKESHEKIKDIVETIKSEKMRGIKQMMNPVLIKSLKIISKQFDILNKVKQIIKHK
MIVQALKEDEPKLESYKLLKREVFDLKENTIKELKKIEHLTMKDDNKDMERSPTLTFPKEDNELP 1760 1770 1780 1790 1800 1810 1820
SRERTLRKPRSLPVSTLLKKALKVQTKQLQMKAEIEKLSMVLDVPQYRDTGMLTLRQIKQKQIVY
DVIGKAIDVQDELEELNKALQEKMEDTPEDKSVRDLIKVVDDRQDELKKTIKLFEDTVEHNSRMD 1890 1900 1910 1920 1930 1940 1950
NSLVVPLEKLQQLKDRIEQQQLKVQKTKERFDQLREAVDSDSKRTLPKYPKNLPSRSSMEMPNGR

