

1	MKALRLSASA	LFCLLLINGL	GAAPPGRPEA	QPPPLSSEHK	EPVAGDAVPG	PKDGSAPEVR	60
61	GARNSEPQDE	GELFQGVDP	ALAAVLLQAL	DRPASPPAPS	GSQQGP EEEA	AEALLTETVR	120
121	SQTHSLPAPE	SPEPAAPRP	QTPENGPEAS	DPSEELEALA	SLLQELRDFS	PSSAKRQQET	180
181	AAAE TETRTH	TLTRVNLESP	GPERVWRASW	GEFQARVPER	APLPPPAPSQ	FQARMPSGP	240
241	LPETHKFGEG	VSSPKTHLGE	ALAPLSKAYQ	GVAAPFPKAR	RPESALLGGS	EAGERLLQQG	300
301	LAQVEAGRRQ	AEATRQAAAQ	EERLADLASD	LLLOYLLQGG	ARQRGLGGRG	LQEAAEERES	360
361	AREEEAEQE	<u>RRGGEERVGE</u>	<u>EDEEAAEAEA</u>	<u>EAE EAERARQ</u>	<u>NALLFAEED</u>	<u>GEAGAED</u>KRS	420
421	QEETPGHRRK	EAEGTEEGGE	EEDDEEMDPQ	TIDSLIELST	KLHLPADDVV	SIIEEVEEKR	480
481	KRKKNAPPEP	VPPPRAAPAP	THVRSPPPPP	PAPAPARDEL	PDWNEVLPPW	DREEDEVYPP	540
541	GPYHPFPNYI	RPRTLQPPSA	LRRRHYPHAL	PPSRHYPGRE	AQARRAQEEA	EAEERRLQEQ	600
601	EELENYIEHV	LLRRP					615