

1 MKTKNNDLILDKNNFNHDAYRRMRVSHSAEALEGQELVDYVNGRQNLRTATSNNKFAKYT  
61 KEEKKQLMGVLNVRNPIKARRNMSPTRYMDIQLPESFDSRKQWGS MCRSVTEVRD QSSCG  
121 SCWSFGAVEAMSDRICIHSNGSIQVDLSAKDLLSCCKACGFGCNGGEPFEAWSFWHTSGI  
181 VTGSNYTTKKGCQPYTFPECEHHSTKAHFKPCPKELYPTPKCEKTCQEGYDKEYNADKYY  
241 GQEPYAVDEDQEAIKKELYLNPLEVSFEVYDDFLTYSGGVYVHQGRIDGGHAVKLVGW  
301 GTENEVPYWIVANSWNTDFGEDGYFRILRGSNECGIEFGVVGGLPDLKRSPHRVHYHHHH  
361 SSASSYDIF