

MKTIIIALSYILCLVFAQKLP GNDNSTATLCLGHHA V P NGTIVKTITNDQIEVTNATELVQSSSTGEICDSPHQILDGKNCT

LIDALLGDPQCDDFQNKKWDLFVERSKAYSNCYPYDVPDYASLRSLVAS SGTLEFNNESFNWTGVTQNGTSSACIRRSKNS

FFSRLNWLTHLNFKYPALNVTMPNNEQFDKLYIWGVLHPGTDKDKQIFLYAQASGRITVSTKRSQQIVSPNIGSRPRVRNIP

SRI SIYWTIVKPGDILLINSTGNLIAAPRGYFKIRSGKSSIMRSDAPIGKCNSECITPNGSIPNDKPFQNVNRITYGACPRY

VKQNTLKLATGMRNVPEKQTRGIFGATAGFIENGWEGMVDGWYGFRRHQNSEGRGQAADLKSTQAAIDQINGKLNRLIGKTN

HKCDNACIGSIRNGTYDHDVYRDEALNNRFQIKGVLELKS GYKDWILWISFAISCFLLCVALLGFI MWACQKGNIRCNICI

HKCDNACIGSIRNGTYDHDVYRDEALNNRFQIKGVLELKS GYKDWILWISFAISCFLLCVALLGFI MWACQKGNIRCNICI