

gi|4506885|ref|NP_002999.1| MKSITILFVLSLLILEKQAAVMGQKGGSKGQLPSGSSQFPHGQKGOHYFGQKDDQHTKSKGSFSIQHTYHVDINDHDWTRKSOQYDLNALHKATKSKOHLGGSOQLLNKQEGRDHDKSKGHFHMIVIHKGGQAHGTONPSODQGNSP 150
gi|57114137|ref|NP_001009138.1| MKSITILFVLSLVLILEKQAAVMGQKDGSKGQLPSGSSQFPHGQKGOHYFGQKDDQHTKSKGSFSIQHTYHVDINDHDQTRKSOQYDLNALHKVTKSKOHLGGSOQLLNKQEGRDHDKSEGHFHMIVIHKGGQAHCGTONPSODQGNSP 150
1.....10.....20.....30.....40.....50.....60.....70.....80.....90.....100.....110.....120.....130.....140.....150



gi|4506885|ref|NP_002999.1| SGKGLSSQCSNTEKRLWVHGLSKEQASASGAQKGRTOGGSSQSSYVLQTEELVNVKQORETKNSHONKGHYONVVDVREEHSSKLOSLHPAHODRLOHGPKDIFTTQDELLVYNKNOHOTKNLSODQEHGRKAHKISYPSRTEERQLHH 300
gi|57114137|ref|NP_001009138.1| SGKGLSSQYSNTEKRLWVHGLSKEQASASGAQKGRTOGGSSQSSYVLQTEELVNVKQOLETKNSHONKGHYONVVDVREEHSGKLOSLHPAHODRLOHGPKDIFTTQDELLVYNKNOHOTKNLNDQEHGQKAHKISYQSSRTEERQLNH 300
.....160.....170.....180.....190.....200.....210.....220.....230.....240.....250.....260.....270.....280.....290.....300



gi|4506885|ref|NP_002999.1| GEKSVQKDVSKGSIISIQTEEKIHGKSONQVTIHSQDQEHGHKENKISYQSSSTEERHLNCGEKIQKGVSKGSIISIQTEEQIHGKSONVRIIPSAQAEYGHKENKISYQSSSTEERLNSGEKDVQKGVSKGSIISIQTEEKIHGKSONQV 450
gi|57114137|ref|NP_001009138.1| GEKSVQKDVSKGSIISIQTEEKIHGKSONQVTIHSQDQEHGHKENKMSYQSSSTEERHLNCGEKIQKGVSKGSIISIQTEEQIHGKSONXVRIIPSAQAEYGRKENKIS----- 407
.....310.....320.....330.....340.....350.....360.....370.....380.....390.....400.....410.....420.....430.....440.....450



gi|4506885|ref|NP_002999.1| TIPSQDQEHGHKENKMSYQSSSTEERLNYGKSTQKDVSSSISFQIEKLVGKSGIQTPNPNQDQWSGONAKGKSGQASADSKODLLSHEOKGRYKQESSESHNIVIIEHEVAQDDHLTCQYNEDRNPISL 582
gi|57114137|ref|NP_001009138.1|460.....470.....480.....490.....500.....510.....520.....530.....540.....550.....560.....570.....580.. 407

