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gi 281485608 |ref| NP_003217.3 |-----MKRVLSCVPEPTVVMARALCMLGL-VLALLSSSSAEYVGLSANQCAVPAKDRVDCGYPHVTPKECNRGCCFDSSRIPGVWCFKPLQEAECTF 131
gi 114684421 |ref| XP_531572.2 |MQERTGAATARRRESLPQANNPEQLCKQRCINEASWTMKRVLSCVLEPTVVMARALCMLGL-VLALLSSSSAEYVGLSANQCAVPAKDRVDCGYPQVTPKECNRGCCFDSSRIPGVWCFKPLQEAECTF 131
gi 50950221 |ref| NP_001002990.1 |-----MEARVLWLLVV-VLVLGSSSLAVAYQGLATNLCEVPPKDRVDCGYPEITSEQCVNRGCCFDSSIHGVWCFKPLQDTECRF 131
gi 194663768 |ref| XP_596070.4 |-----MFSEQSGFQSQPVIMEARTFWLLVAVLALGSSSSTGGYVGLSANQCAVPAKDRVDCGYPEVTPEQCNRGCCFDSSIHGVWCFKPLQEAECTF 131
gi 6755773 |ref| NP_035705.1 |-----METRALWMLLVVLVAGSSGIAADYVGLSPSQCMV PANVRVDCGYPSVTSEQCNRGCCFDSSIPNVWCFKPLQETECTF 131
gi 6981644 |ref| NP_037174.1 |-----METRAFWITLLLVLVAGSSCKAQEFVGLSPSQCMAPTINVRVDCNYPTVTSEQCNRGCCFDSSIPNVWCFKPLQETECTF 131
1.....10.....20.....30.....40.....50.....60.....70.....80.....90.....100.....110.....120.....130.

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