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**** .:** ***** : *.**.*.*.*.*****:***:*****:*.:.: * * :**:* ***** **.*: :*.:.:**: **.:**:* ** *:* *****:*. * :* *.* ***** .:** **:* ** *
gi |50978922|ref|NP_001003182.1| -MAPFPALASGILLLLWVTAAPSRACTCAPPHPOTALCNSQIVIRAKFVGTAEVNOITDLNRRYEIKMTKMFKGFSAALGNASDIRFVDTPALESVCGYLHRSQNRSEEFVLVAGNLRDGHLOINICSFVAPWSSLTAORRGFTKTYAAGCEG 150
gi |27806161|ref|NP_776896.1| -MAPFPAPMASGILLLLWLTAPSRACTCVPPHPOTALCNSDVVIRAKFVGTAEVNETALYORYEIKMTKMFKGFSAALRDAPDIRFIYTPAMESVCGYFHRSONRSEEFLIAGQLSNGLHITICSFVAPWNSMSAORRGFTKTYAAGCEE 150
gi |4507509|ref|NP_003245.1| -MAPFPALASGILLLLWLIAPSRACTCVPPHPOTALCNSDLVIRAKFVGTPEVNOITLYORYEIKMTKMYKGFQALGDAADIRFVYTPAMESVCGYFHRSHNRSEEFLIAGKLDGGLHITICSFVAPWNSLSLAORRGFTKTYTVGCEE 150
gi |113205063|ref|NP_001037849.1| MMAPFASLASGILLLLSLIASKACSCAPPHPOTALCNSDLVIRAKFMGSPETNETLYORYEIKMTKMLKGFQAVGNAADIRYAYTPVMSLGGYVHKSQNRSEEFLLITGRLRNGNLHISACSFVLPWRITLSPAQRAPSKTYSAGCGV 150
gi |16758666|ref|NP_446271.1| -MAPFASLASGILLLLSLIASKACSCAPTHPOTALCNSDLVIRAKFMGSPETNETLYORYEIKMTKMLKGFDAVGNATGFRFAYTPAMESLCGYVHKSQNRSEEFLLIAGRLRNGNLHITACSFVLPWHNLSPAQRAPSKTYSAGCGV 150
1.....10.....20.....30.....40.....50.....60.....70.....80.....90.....100.....110.....120.....130.....140.....150

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gi |50978922|ref|NP_001003182.1| CTVFTCSIPCKLQSDTHCLWTDHFLTGSDDKGFQSRHLACLPREPGICTWOSLRPRMA----- 218
gi |27806161|ref|NP_776896.1| CTVFPSSIPCKLQSDTHCLWTDOLLTGSDDKGFQSRHLACLPREPGICTWOSLRAQMA----- 218
gi |4507509|ref|NP_003245.1| CTVFPCLIPCKLQSGTHCLWTDOLLQSGSEKGFQSRHLACLPREPGICTWOSLRSQIA----- 218
gi |113205063|ref|NP_001037849.1| CTVFPCLIPCKLESSTHCLWTDQVLVGSSEDFQSRHFACLPRNPGLCTWRSLGAR----- 218
gi |16758666|ref|NP_446271.1| CTVFPCLIPCKLESSTHCLWTDQILMGSEKGYQSDHFACLPRNPDLCTWQYLGVSMTSLPLAKAEA 218
.....160.....170.....180.....190.....200.....210.....

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