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gi |154800437|ref|NP_001946.3|MDVLLMIFSLLFVACGAPETAVLGAELSAVGENGGKPTTPSPWRLRRSKRCSSSLMDKCEVYFCHLDIIWVNTPEHVVPVYGLGSP-RSKRALENLLPTKATDRENRCQASQKDKKCWNFCOAGKELRAEDIMEKDWNNHKKGKDCS 149
gi |114605505|ref|XP_518241.2|MDVLPMIFSLLFVACGAPETAVLGAELSVAGEDGGEKPTTPSPWRLRRSKRCSSSLMDKCEVYFCHLDIIWVNTPEHVVPVYGLGSP-RSKRALENLLPTKATDRENRCQANQKDKKCWNFCOAGKELRAEDIMEKDWNNHKKGKDCS 149
gi |50950175|ref|NP_001002956.1|MDYFPMIFSLLFVACGAPETAVLGAELSTGPDSTRGKPPAPWRPRRSKRCSSSLMDKCEVYFCHLDIIWVNTPEHVVPVYGLGSPSRSKRSLKDLFPTKATDHRKRCQASQKDKKCWNFCOAGKELRAEDIMEKDWNNHKKGKDCS 150
gi |30794314|ref|NP_851353.1|MDYFPMIFALLFVAFQGAPEAAVLGTELSAGAEDGGEKPAPAPWRPRRSKRCSSSLMDKCEVYFCHLDIIWVNTPEHVVPVYGLGSPSRSKRSLKDLFPTKATVHRKRCQASQKDKKCWNFCOAGKELRAEDIMEKAWNNQKRGKDCS 150
gi |6753720|ref|NP_034234.1|MDYFVIFSLLFVTFQGAPEAVLGAELSTGAENGVSPPTPTWRPRRSKRCSSSLMDKCEVYFCHLDIIWVNTPEHVVPVYGLGSPSRSKRSLKDLLEPKATDQAVRCQAHQKDKKCWNFCOAGKELRAEDIMEKSLKDSKKGKPCP 150
gi |6978791|ref|NP_036680.1|MDYFVIFSLLFVAFQGAPEAVLGAELSPRAEKEVSPPTPTWRPRRSKRCSSSLMDKCEVYFCHLDIIWVNTPEHVVPVYGLGSPSRSKRSLKDLLEPKATDQGNRCQAHQKDKKCWNFCOAGKELRAEDIMEKGVKDFKKGKPCP 150
gi |118086398|ref|XP_418943.2|MDCSRLFLPLLVALCP-ALLPAAPGAEVNAAS-----PSPAAASH-RRARRCSSSLLEDECEVYFCHLDIIWVNTPEKIVPYGLGGPSRKRSLKDIPEMLAGASSRCRCASQKDKKCLNFCOTGKDLWAGSTAEKTSRHRNKAGGC- 142
gi |18858581|ref|NP_571594.1|-MHLRIIFPVLITMLISGFFDFGAPASLG-----PGTAPARHSRNKRCSCASFLDKCEVYFCHLDIIWVNTPEKIVSYGLGNAPRKRKRSVTEFVVLES-----RCRCADSQDKTCSFCCQADSALQFKAAADRRAIR---AAQGH 130
1.....10.....20.....30.....40.....50.....60.....70.....80.....90.....100.....110.....120.....130.....140.....150

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gi |154800437|ref|NP_001946.3|KLGKKCIYQQLVGRKIRRSSEHLRQTRSETMNRNSVKSFSFHDPKLKGKPSRERYVTHNRAHW----- 212
gi |114605505|ref|XP_518241.2|KLGKKCIYQQLVGRKIRRSSEHLRQTRSETMNRNSVKSFSFHDPKLKGKPSRERYATHNRAHW----- 212
gi |50950175|ref|NP_001002956.1|ELGEKSKQQLVAGRKIRR-----LDAISNSIKTAFRVAKLKAELYRENKVTNHRTH----- 202
gi |30794314|ref|NP_851353.1|KLGEKCLHQQLVAGRKTRR-----LEAISNSIKTAFRVAKLKAELYRDKKVIYNRAH----- 202
gi |6753720|ref|NP_034234.1|KLGKKCIYQQLVEGRKLR-----LEAISNSIKAFRVAKLKAELYRDKLTHNRAH----- 202
gi |6978791|ref|NP_036680.1|KLGKKCIYQQLVEGRKLR-----LEAISNSIKTAFRVAKLKAELYRDKLTHNRAH----- 202
gi |118086398|ref|XP_418943.2|-IGPKMNVQFVDSRKMKR-----LEAVGNSIKAFSLAKLKAELQGRKCLKHNRASKRSIWKSLKTF 205
gi |18858581|ref|NP_571594.1|CAGKQCKHRLAETQTKIRR-----LRTTKQTDVLSGKKKIQLLLEKWRMRNRHRSQAWISENS----- 188
.....160.....170.....180.....190.....200.....210.....220.....

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