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gi |11386203|ref|NP_000380.1|-----MSEPAQDVRQNPCGSKACRRLFGPVDSEQLSRDCDALMAGCIQEARERWNFDFVTETPLEGDFAWERVRGLGLPKLYLPTGPRRGRDELGGRRPGTSPALLGG-- 104
gi |114607027|ref|XP_001157015.1|-----MSEPAQDVRQNPCGSKACRRLFGPVDSEQLSRDCDALMAGCIQEARERWNFDFVTETPLEGDFAWERVRGLGLPKLYLPTGPRRGRDELGGRRPGTSPALLGG-- 104
gi |73972722|ref|XP_532125.2|MSSTCTQAAPGGASSRAAAGDGLESGGGRGFAEPETLGGAAGAMSEPSRDALPIPHGSKACRRLFGPVDSEQLSRDCDALMAGCVOEARERWNFDFVTETPLEGDFAWERVRGLGLSKVSLPAGPRGRDDLGGGRPGTSPALLGG-- 148
gi |149643101|ref|NP_001092428.1|-----MSELSRDAHQIPRSSKACRRLFGPVDSEQLSRDCDALMAGCVOEARERWNFDFVTETPLEGDFAWERVRGLGLPKLYLPVGPR---DDLGGGRPGTSPALLGG-- 101
gi |6671726|ref|NP_031695.1|-----MSNP-GDVRPVPHRSKVCRCLFGPVDSEQLSRDCDALMAGCIQEARERWNFDFVTETPLEGNFVWERVRSLGLPKVYLSPGSRS-RDDLGGDKRPSTSSALLGG-- 102
gi |18158451|ref|NP_542960.1|-----MSDP-GDVRPVPHRSKVCRCLFGPVDSEQLSRDCDALMAGCIQEARERWNFDFATETPLEGNFVWERVRSPGLPKIYLSPGSRR-RDDLGGDKRPSTSSALLGGPG 104
1.....10.....20.....30.....40.....50.....60.....70.....80.....90.....100.....110.....120.....130.....140.....150

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gi |11386203|ref|NP_000380.1|TAEEDHVDLSLSCLVPRSGEQAEGSPGGPGDSQGRKRRQTSMDFYHSKRRLIFSKRKP 164
gi |114607027|ref|XP_001157015.1|TAEEDHVDLSLSCLVPRSGEQAEGSPGGPGDSQGRKRRQTSMDFYHSKRRLIFSKRKP 164
gi |73972722|ref|XP_532125.2|TAQEDHLDLSLTCLLPHSPERPEASPGVPGTSQGRKRRQTSMDFYHSKRRLIFSKRKP 208
gi |149643101|ref|NP_001092428.1|TSQEDHLDLSLSCLVTRSPERPEGTPGGPGPSQGRKRRQTSMDFYHSKRRLICSKRKP 161
gi |6671726|ref|NP_031695.1|PAPEDHVALSLSCLVS---ERPEDSPGGPGTSQGRKRRQTSLDFYHSKRRLVFCKRKP 159
gi |18158451|ref|NP_542960.1|PAPEDHVALSLSCLVSHAPERPEEDSPGGTTSQGRKRRQTSLDFYHSKRRLVFCKRKP 164
.....160.....170.....180.....190.....200.....210

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