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 gi | 5729836 | ref | NP_006696.1 | MTLLEVRGQDTVP ESTARMOGAGKALHELLLSAQRGGCLTAGVYESAKVLNVDPDNTVFCVLAAGEEDEGDIALQIHFTLIQAFCCENDIDIVRVGDVORLAAIVGAGEEAGAPGDLHCILIS----- 150
 gi | 114625449 | ref | XP_001141972.1 | MTLLEVRGQDTVP ESTARMOGAGKALHELLLSAQRGGCLTAGVYESAKVLNVDPDNTVFCVLAAGEEDEGDIALQIHFTLIQAFCCENDIDIVRVGDVORLAAIVGAGEEAGAPGDLHCILIS----- 150
 gi | 6753938 | ref | NP_035947.1 | MTLLEVRGQDTVP ESTARMOGAGKALHELLLSAQRGGCLTAGVYESAKVLNVDPDNTVFCVLAAGEEDEGDIALQIHFTLIQAFCCENDIDIVRVGDVORLAAIVGAGEEAGAPGDLHCILIS----- 150
 gi | 109504685 | ref | XP_237999.4 | MTLLEVRGQDTVP ESTARMOGAGKALHELLLSAQRGGCLTAGVYESAKVLNVDPDNTVFCVLAAGEEDEGDIALQIHFTLIQAFCCENDIDIVRVGDVORLAAIVGADDEGGAPGDLHCILIS----- 150
 gi | 114052512 | ref | NP_001039366.1 | MTLLEVRGQDTVP ESTARMOGAGKALHELLLSAQRGGCLTAGVYESAKVLNVDPDNTVFCVLAAGEEDEGDIALQIHFTLIQAFCCENDIDIVRVGDVORLAAIVGTGDESGAPVDLHCILIS----- 150
 gi | 73947066 | ref | XP_541314.2 | MTLLEIRGQDTVP ESTARMOGAGKALHELLLSAQRGGCLTAGVYESAKVLNVDPDNTVFCVLAAGEEDEGDIALQIHFTLIQAFCCENDIDIVRVGDVORLAAIVGADDEGGAPGDLHCILISVAVSPORRPPRAAPRORPARLTSAP----- 150
 gi | 118129613 | ref | XP_001232966.1 | -----XGCLPAGVYESAKLMNVDPDNTVFCVLAAGEEDEGDIALQIHFTLIQAFCCNDIIVRVVAGMRRLLAAVGEFAPGAEPRDLHCLLV----- 150
 gi | 45387791 | ref | NP_991254.1 | MTFEELSGDNTATGDRMGTAGTALAEELLVSAKQDCLTVGVYESAQMNVDPDSVAFCVLATDEEYDYDIALQIHFTLIQAFCDNDINIVRVNDIERLADIVGCDQD-EEPKDAHCILV----- 150
 1.....10.....20.....30.....40.....50.....60.....70.....80.....90.....100.....110.....120.....130.....140.....150



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 gi | 5729836 | ref | NP_006696.1 | ---NPNEDAWKDP ALEKLSLFCEESRSVNDWVPSITLPE- 190
 gi | 114625449 | ref | XP_001141972.1 | ---NPNEDAWKDP ALEKLSLFCEESRSVNDWVPSITLPE- 190
 gi | 6753938 | ref | NP_035947.1 | ---NPNETWKPDP ALEKLSLFCEESRSFNDWVPSITLPE- 190
 gi | 109504685 | ref | XP_237999.4 | ---NPNETWKPDP ALEKLSLFCEESRSFNDWVPSITLPE- 190
 gi | 114052512 | ref | NP_001039366.1 | ---NPNEDAWKDP ALEKLSLFCEESRSVNDWVPSITLPE- 190
 gi | 73947066 | ref | XP_541314.2 | SPQNPNEDAWKDP ALEKLSVFCESRSVNDWVPSITLPE- 190
 gi | 118129613 | ref | XP_001232966.1 | -----VSAG----- 190
 gi | 45387791 | ref | NP_991254.1 | ---RPNDDSWKDSALDKLGLFCESRSNVYEWVPSITLPER 190
160.....170.....180.....190

