

gi | 4758226 | ref | NP_004082.1 | MLQGPRALASAAGTTPKVVPAMSPTLWPSGLSSPOLCPAT--ATYYTPLYPTAPPAAPCTCLDAIPHGPEGQVVRCLPAGRLPAKRLDLEIGRPVVVEFPFPGKGCIRVDGLPSPKTPKSPGKTRYDTSGLLTKKFIYLLSES 148
gi | 114554629 | ref | XP_524538.2 | MLQGPRALASAAGTTPKVVPAMSPTLWPSGLSSPOLCPAT--ATYYTPLYPTAPPAAPCTCLDAIPHGPEGQVVRCLPAGRLPAKRLDLEIGRPVVVEFPFPGKGCIRVDGLPSPKTPKSPGKTRYDTSGLLTKKFIYLLSES 148
gi | 76611569 | ref | XP_874289.1 | MLL-----KLCLELLSLP-----KAKRLDLEIGRPVVVEFPFPGKGCIRVDGLPSPKTPKSPGKTRYDTSGLLTKKFIYLLSES 78
gi | 29244208 | ref | NP_808401.1 | MLRAPRILAPAIAPTKSLPALNPTELWPSGLSSPOLCPATATYYTSLYTDIVPSSVALGTCCLDAIPHGPEGQIVRCAPAGRLPAKRLDLEIGRPVVVEFPFPGKGCIRVDGLPSPKTPKSPGKTRYDTSGLLTKKFIYLLSES 150



gi | 4758226 | ref | NP_004082.1 | EDGVLDLNWAAEVLVQKRRYDIINVLEGIQLIRKKAKNNIQWVGRGMFEDPTRPGKQQLGQELKELMNEQALDQLIQSCSLSPFKHLTEDKANKRLAYVTYQDIRAVGNFKKQTVIAVKAPPQTRLEVPRTEDELQIYLKSTQGPI 298
gi | 114554629 | ref | XP_524538.2 | EDGVLDLNWAAEVLVQKRRYDIINVLEGIQLIRKKAKNNIQWVGRGMFEDPTRPGKQQLGQELKELMNEQALDQLIQSCSLSPFKHLTEDKANKRLAYVTYQDIRAVGNFKKQTVIAVKAPPQTRLEVPRTEDELQIYLKSTQGPI 298
gi | 76611569 | ref | XP_874289.1 | EDGVLDLNWAAEVLVQKRRYDIINVLEGIQLIRKKAKNNIQWVGRGLFEDPTRPGKQQLGQELKELMNEQALDQLIHSCSLNFKHLTEDKANKRLAYVTYQDIRAVGNFKKQTVIAVKAPPQTRLEVPRSEENLQIHLKSTQGP 228
gi | 29244208 | ref | NP_808401.1 | EDGVLDLNWAAEVLVQKRRYDIINVLEGIQLIRKKAKNNIQWVGRELFEDPTRPSRQQLGQELKELMNAEQLDQLIQSCSLSPFKHLTEDNANKRLAYVTYQDIRAVGNFKKQTVIAVKAPPQTRLEVPRAEENLQIYLKSTQGP 300



gi | 4758226 | ref | NP_004082.1 | EVVLCPEEVQEPDPSPEEPLPSTSLCPSDAPSSSTDPSSIMEPTASSV---PAPAPTPQQAPPPPSLVPLEATDLSLELPHPLLQOTEDQFLSPTLACSSPLISFSPPLDQDDVWGLGEGEGISDLFDSYDLGDLLIN 437
gi | 114554629 | ref | XP_524538.2 | EVVLCPEEVQEPDPSPEEPLPSTSLCPSDAPSSSTDPSSIMEPTASSV---PAPAPTPQQAPPPPSLVPLEATDLSLELPHPLLQOTEDQFLSPTLACSSPLISFSPPLDQDDVWGLGEGEGISDLFDSYDLGDLLIN 437
gi | 76611569 | ref | XP_874289.1 | EVVLCPEEVQEPHSPAKEPLPSTALSPPDSTQLNSDPIGITEPTASSE---PALISPVQVPPPPPPPLVPLEATENMLELPHPLLQOTEDQFLSPTLPCSSPLISFSPPLDQDDVWGLDGGEGISDLFDYDLGDLLIN 367
gi | 29244208 | ref | NP_808401.1 | EVVLCPEEGQEPDSPAALPSTALSPIPCAPGCGTDSGIAETIEPSSVLIPIPPPPPPPLPAPSLVPLEATDNMLELPHPLLQOTEDQFLSPTLAANSPLISFSPPLDQDDVWGLMDEGEGISDLFDSYDLGDLLIN 443

