

gi | 4507233 | ref | NP\_003736.1 | MVAHNQVAADNAVSTAAEPRRRPEPSSSSSSS-----PAAPARPRPCPA---VPAPAPGDTHFRTRSHADYRRIIRASALLDACGFYWGPLSVHGAHERLRAEVPVGTFLVRDSRQRNCFVALSVKMASGPTSIRVHFOAGRFHLD 150  
gi | 114660995 | ref | XP\_001141793.1 | MVAHNQVAADNAVSTAAEPRRRPEPSSSSSSS-----SAAPARPRPCPA---VPAPAPGDTHFRTRSHADYRRIIRASALLDACGFYWGPLSVHGAHERLRAEVPVGTFLVRDSRQRNCFVALSVKMASGPTSIRVHFOAGRFHLD 150  
gi | 6753424 | ref | NP\_034026.1 | MVARNOVAADNAISPAEPRRRPEPSSSSSSS-----PAAPVRPRPCPA---VPAPAPGDTHFRTRSHSDYRRIIRTSALLDACGFYWGPLSVHGAHERLRAEVPVGTFLVRDSRQRNCFVALSVKMASGPTSIRVHFOAGRFHLD 150  
gi | 22024396 | ref | NP\_665886.1 | MVARNOVAADNAISPAEPRRRPEPSSSSSSS-----PAAPRRPRPCPV---VPAPAPGDTHFRTRSHSDYRRIIRTSALLDACGFYWGPLSVHGAHERLRAEVPVGTFLVRDSRQRNCFVALSVKMASGPTSIRVHFOAGRFHLD 150  
gi | 76652907 | ref | XP\_869409.1 | MVAHNQVAADNAISPAEPRRRPEPSSSSSSSSSSSSSSS-----SAAPARLRPCAAAAAPAPAPGDTHFRTRSHAEYRRIIRASALLDACGFYWGPLSVHGAHERLRAEVPVGTFLVRDSRQRNCFVALSVKMASGPTSIRVHFOAGRFHLD 150  
gi | 73959153 | ref | XP\_547126.2 | -----MRRP-----CAPPAAPTAPA-----PAAAPGDTHFRTRSOAEYRRIIRASALLEACGFYWGPLNVHGAHERLRAEVPVGTFLVRDSRQRNCFVALSVKMASGPTSIRVHFOAGRFHLD 150  
gi | 50755896 | ref | XP\_414929.1 | MVAHSKVSADNAV--AADPRCLLDPPARDRQ-----ARGYRGTGRPG-----AAQAPSSTHFRTRSOADEFSSIRASALLDACGFYWGPLTVSAAHEKLSKSEPEGTFLLRDSTQKNCFVALSVKMASGPTSIRVHFOAGRFHLD 150  
gi | 51011015 | ref | NP\_001003467.1 | MVAHSSVEGNQGI---EDRQVSTEASSDVFQ-----SQRSRPRQS-----TVAFQTHFRKPFNYARDFKIIENITSMLEESGFYWGPMTEVEEAHLKLLKESVGTFLLRDSRQSDVFFTLVYKAQNGFVIRINFKNSKFSIT 150



gi | 4507233 | ref | NP\_003736.1 | GRSRESFDCLFELLEHYVAAPRRMLGAPLRORRVRPQELCRORIVAIVGRENLARIPLPVLRDYLSSFFPFOI 223  
gi | 114660995 | ref | XP\_001141793.1 | GRSRESFDCLFELLEHYVAAPRRMLGAPLRORRVRPQELCRORIVAIVGRENLARIPLPVLRDYLSSFFPFOI 223  
gi | 6753424 | ref | NP\_034026.1 | GRSRESFDCLFELLEHYVAAPRRMLGAPLRORRVRPQELCRORIVAIVGRENLARIPLPVLRDYLSSFFPFOI 223  
gi | 22024396 | ref | NP\_665886.1 | GRSRESFDCLFELLEHYVAAPRRMLGAPLRORRVRPQELCRORIVAIVGRENLARIPLPVLRDYLSSFFPFOI 223  
gi | 76652907 | ref | XP\_869409.1 | GRSRESFDCLFELLEHYVAAPRRMLGAPLRORRVRPQELCRORIVAIVGRENLARIPLPVLRDYLSSFFPFOI 223  
gi | 73959153 | ref | XP\_547126.2 | GRSRESFDCLFELLEHYVAAPRRMLGAPLRORRVRPQELCRORIVAIVGRENLARIPLPVLRDYLSSFFPFOI 223  
gi | 50755896 | ref | XP\_414929.1 | GSKETFDFCLFKLLEHYLSSPRKVLVPLKRVVQPLQELCRKSVKIFGRENLNQIPLPVLRDYLKSEFFPFOI 223  
gi | 51011015 | ref | NP\_001003467.1 | GSKESFDSLFLKLEHYISSPKGLIRPCRKPEVQSLQQLCRQIMERCDDGKIDICIPVQPIILKDFLHAFPHPL 223

