

gi | 31560568 | ref | NP\_034884.2 | MSSILPFPPVVKRLLGWKKSAGGSGGAGGG-EQNGQEEKWCEKAVKSLVKLKKLGRLEDELEKAITTQNCNTKCVIIPSTCSEIWLSTANTVDQWDTGLYFSFSEQTRSLDGRLOVSHRKGKLPVVIYCRLWRWPDLSHSHLKAIEHC 149  
gi | 9506873 | ref | NP\_062064.1 | MSSILPFPPVVKRLLGWKKSAGGSGGAGGG-EQNGQEEKWCEKAVKSLVKLKKLGRLEDELEKAITTQNCNTKCVIIPSTCSEIWLSTANTVDQWDTGLYFSFSEQTRSLDGRLOVSHRKGKLPVVIYCRLWRWPDLSHSHLKAIEHC 149  
gi | 114051079 | ref | NP\_001039683.1 | MSSILPFPPVVKRLLGWKKSAGGSGGAGGG-EQNGQEEKWCEKAVKSLVKLKKLGRLEDELEKAITTQNCNTKCVIIPSTCSEIWLSTANTVDQWDTGLYFSFSEQTRSLDGRLOVSHRKGKLPVVIYCRLWRWPDLSHSHLKAIEHC 149  
gi | 114673061 | ref | XP\_001149364.1 | MSSILPFPPVVKRLLGWKKSAGGSGGAGGG-EQNGQEEKWCEKAVKSLVKLKKLGRLEDELEKAITTQNCNTKCVIIPSTCSEIWLSTANTVDQWDTGLYFSFSEQTRSLDGRLOVSHRKGKLPVVIYCRLWRWPDLSHSHLKAIEHC 149  
gi | 73961195 | ref | XP\_852799.1 | MSSILPFPPVVKRLLGWKKSAGGSGGAGGG-EQNGQEEKWCEKAVKSLVKLKKLGRLEDELEKAITTQNCNTKCVIIPSTCSEIWLSTANTVDQWDTGLYFSFSEQTRSLDGRLOVSHRKGKLPVVIYCRLWRWPDLSHSHLKAIEHC 149  
gi | 5174511 | ref | NP\_005892.1 | MSSILPFPPVVKRLLGWKKSAGGSGGAGGG-EQNGQEEKWCEKAVKSLVKLKKLGRLEDELEKAITTQNCNTKCVIIPSTCSEIWLSTANTVDQWDTGLYFSFSEQTRSLDGRLOVSHRKGKLPVVIYCRLWRWPDLSHSHLKAIEHC 149  
gi | 46048929 | ref | NP\_989892.1 | MSSILPFPPVVKRLLGWKKSAGGSGGAGGG-EQNGQEEKWCEKAVKSLVKLKKLGRLEDELEKAITTQNCNTKCVIIPSTCSEIWLSTANTVDQWDTGLYFSFSEQTRSLDGRLOVSHRKGKLPVVIYCRLWRWPDLSHSHLKAIEHC 149  
gi | 40254710 | ref | NP\_571441.2 | MSSILPFPPVVKRLLGWKKSAGGSGGAGGG-EQNGQEEKWCEKAVKSLVKLKKLGRLEDELEKAITTQNCNTKCVIIPSTCSEIWLSTANTVDQWDTGLYFSFSEQTRSLDGRLOVSHRKGKLPVVIYCRLWRWPDLSHSHLKAIEHC 149



gi | 31560568 | ref | NP\_034884.2 | EYAFNLKKDEVCVNPYHYQRVETPVLPPVLPVPRHTEIILTELPPLDDYTHSIPENTNFPAGIEPQSNYIPEPPPGYISEDGETSDQQLNQSMDTGPSAELSPITLSPVNHSLDLPVITYSEPAFWCSIAAYELNQRVGETFHASQPSLTV 299  
gi | 9506873 | ref | NP\_062064.1 | EYAFNLKKDEVCVNPYHYQRVETPVLPPVLPVPRHTEIILTELPPLDDYTHSIPENTNFPAGIEPQSNYIPEPPPGYISEDGETSDQQLNQSMDTGPSAELSPITLSPVNHSLDLPVITYSEPAFWCSIAAYELNQRVGETFHASQPSLTV 299  
gi | 114051079 | ref | NP\_001039683.1 | EYAFNLKKDEVCVNPYHYQRVETPVLPPVLPVPRHTEIILTELPPLDDYTHSIPENTNFPAGIEPQSNYIPEPPPGYISEDGETSDQQLNQSMDTGPSAELSPITLSPVNHSLDLPVITYSEPAFWCSIAAYELNQRVGETFHASQPSLTV 299  
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gi | 46048929 | ref | NP\_989892.1 | EYAFNLKKDEVCVNPYHYQRVETPVLPPVLPVPRHTEIILTELPPLDDYTHSIPENTNFPAGIEPQSNYIPEPPPGYISEDGETSDQQLNQSMDTGPSAELSPITLSPVNHSLDLPVITYSEPAFWCSIAAYELNQRVGETFHASQPSLTV 299  
gi | 40254710 | ref | NP\_571441.2 | EYAFNLKKDEVCVNPYHYQRVETPVLPPVLPVPRHTEIILTELPPLDDYTHSIPENTNFPAGIEPQSNYIPEPPPGYISEDGETSDQQLNQSMDTGPSAELSPITLSPVNHSLDLPVITYSEPAFWCSIAAYELNQRVGETFHASQPSLTV 300



gi | 31560568 | ref | NP\_034884.2 | DGFIDPSNSERFCLGLLSNVNRNATVEMTRRHIGRGRVRLYYIGGEVFAECLSDSAIFVQSPNCNORYGWHPATVCKIPPGCNLKIIFNNOEFAALLAQSVNGGFEAVYQLTRMCTIRMSFVKGWGAEYRROTVTSTPCWIELHLNGPLOWL 449  
gi | 9506873 | ref | NP\_062064.1 | DGFIDPSNSERFCLGLLSNVNRNATVEMTRRHIGRGRVRLYYIGGEVFAECLSDSAIFVQSPNCNORYGWHPATVCKIPPGCNLKIIFNNOEFAALLAQSVNGGFEAVYQLTRMCTIRMSFVKGWGAEYRROTVTSTPCWIELHLNGPLOWL 449  
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gi | 114673061 | ref | XP\_001149364.1 | DGFIDPSNSERFCLGLLSNVNRNATVEMTRRHIGRGRVRLYYIGGEVFAECLSDSAIFVQSPNCNORYGWHPATVCKIPPGCNLKIIFNNOEFAALLAQSVNGGFEAVYQLTRMCTIRMSFVKGWGAEYRROTVTSTPCWIELHLNGPLOWL 449  
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gi | 31560568 | ref | NP\_034884.2 | DKVLTQMGSPSVRCSSMS 467  
gi | 9506873 | ref | NP\_062064.1 | DKVLTQMGSPSVRCSSMS 467  
gi | 114051079 | ref | NP\_001039683.1 | DKVLTQMGSPSVRCSSMS 467  
gi | 114673061 | ref | XP\_001149364.1 | DKVLTQMGSPSVRCSSMS 467  
gi | 73961195 | ref | XP\_852799.1 | DKVLTQMGSPSVRCSSMS 467  
gi | 5174511 | ref | NP\_005892.1 | DKVLTQMGSPSVRCSSMS 467  
gi | 46048929 | ref | NP\_989892.1 | DKVLTQMGSPSVRCSSMS 467  
gi | 40254710 | ref | NP\_571441.2 | DKVLTQMGSPSVRCSSMS 468

