

gi|10835256|ref|NP_037390.2| ** *****:.*:****:*:*..*****:** .*:*****.*** ** ** *****:*:* * :** : * :*. ***** **:* * :****.*:*: *****
 gi|114662166|ref|XP_001158687.1| MGRARDAILDALLENLTAEELKKFKLKLLEVPVLRGYGRIPRGALLSMDALDLTDKLVSFYLETYGAELTANVLRDMGLOEMAGQLAAATHGSGAEPAGIQAPPQSAAKPG-LHFIDQHRRAALIARVTNVEWLLDALYGVLTDEQYQAV 150
 gi|73958280|ref|XP_848608.1| MGRARDAILDALENLTAEELKKFKLKLLEVPVLRGYGRIPRGALLSMDALDLTDKLVSFYLETYGAELTANVLRDMGLOEMAGQLAAATHGSGAEPAGIQAPPQSAAKPG-LHFIDQHRRAALIARVTNVEWLLDALYGVLTDEQYQAV 150
 gi|27807409|ref|NP_777155.1| MGRARDAILDALENLTAEELKKFKLKLLEVPVLRGYGRIPRGALLSMDALDLTDKLVSFYLETYGAELTANVLRDMGLOEMAGQLAAATHGSGAEPAGIQAPPQSAAKPG-LHFIDQHRRAALIARVTNVEWLLDALYGVLTDEQYQAV 150
 gi|31560222|ref|NP_075747.2| MGRARDAILDALENLTAEELKKFKLKLLEVPVLRGYGRIPRGALLSMDALDLTDKLVSFYLETYGAELTANVLRDMGLOEMAGQLAAATHGSGAEPAGIQAPPQSAAKPG-LHFIDQHRRAALIARVTNVEWLLDALYGVLTDEQYQAV 150
 gi|27229294|ref|NP_758825.1| MGRARDAILDALENLTAEELKKFKLKLLEVPVLRGYGRIPRGALLSMDALDLTDKLVSFYLETYGAELTANVLRDMGLOEMAGQLAAATHGSGAEPAGIQAPPQSAAKPG-LHFIDQHRRAALIARVTNVEWLLDALYGVLTDEQYQAV 150
 1.....10.....20.....30.....40.....50.....60.....70.....80.....90.....100.....110.....120.....130.....140.....150



gi|10835256|ref|NP_037390.2| *** ..**:*:* *:* ** * : * ** : :.* ** * : *
 gi|114662166|ref|XP_001158687.1| RAEPINPSKMRKLFSPAWNNTCKDLLLQALRESQSVLVEDLERS 196
 gi|73958280|ref|XP_848608.1| RAEPINPSKMRKLFSPAWNNTCKDLLLQALRESQSVLVEDLERS 196
 gi|27807409|ref|NP_777155.1| RAEPINPSKMRKLFSPAWNNTCKDLLLQALRESQSVLVEDLERS 196
 gi|31560222|ref|NP_075747.2| RAEPINPSKMRKLFSPAWNNTCKDLLLQALRESQSVLVEDLERS 196
 gi|27229294|ref|NP_758825.1| RAEPINPSKMRKLFSPAWNNTCKDLLLQALRESQSVLVEDLERS 196
160.....170.....180.....190.....

