```
logo

NTDB id 48511 H7R46 RS02040 WP 005067250.1  MFNFLNFQPLIQLFSPCSLCELGIREKYSLCKDCWEQLPWLKQTIQRNNQSVLVACNYAYPVNRIIQQFKYEQKLHYQILLGEIL   85
NTDB id 1042 H0N27 RS01930 WP 000472273.1  MFKFLNPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHYQTLLAEVL   85
NTDB id 1071 ABD1 RS15375 WP 000472269.1  MFKFLNPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHYQTLLAEVL   85
consensus                                  !!*!!!*!*!*!!*!!!!*!!!*!*!!!!!!!!!*!!!!!!!!!!!!!!!!*!!!!!!!*!!!!*!!!!!!!!!!!!!!!!!*!!*!*!

logo

NTDB id 48511 H7R46 RS02040 WP 005067250.1  KQLKFPKVQAIVPMPISNQRLIERGFNQSLVLANILSKHLKIPVWQPIQRLNEHSQKGLSRLERFENIEQQFLPHHQEKRRYRRV   170
NTDB id 1042 H0N27 RS01930 WP 000472273.1  QQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLSKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQFVALTQEKRRYRRV   170
NTDB id 1071 ABD1 RS15375 WP 000472269.1  QQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLGKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQFVALTQENRRYRRV   170
consensus                                  *!!!!!!!!!!!!!!!!!*!!!*!!!!!!!!*!!!*!*!*!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!****!!*!!!!!!

logo

NTDB id 48511 H7R46 RS02040 WP 005067250.1  LIIDDVITTGSSVHALSQALKQLGCTSIHTACLAATPKS..   209
NTDB id 1042 H0N27 RS01930 WP 000472273.1  LIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY   211
NTDB id 1071 ABD1 RS15375 WP 000472269.1  LIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY   211
consensus                                  !!!!!!!!!!!!!!!!!!!!
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