```
logo

NTDB id 1021138 ABZP31 RS13165 WP 367156824.1  .MPMFNPSSLLRPISRLLPCPLCSSSLQMQHGVCHDCWEQLPWYRQTVERQELNIHVAYGYQYPIDRIIQQFKYEQQLNY   79
NTDB id 1042 HON27 RS01930 WP 000472273.1      MFKFLNPQYL...FQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHY   77
NTDB id 1071 ABD1 RS15375 WP 000472269.1       MFKFLNPQYL...FQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHY   77
consensus                                      *****!!**!    ***!*!!*!!**********!**!!!!!!!**!!**!******!!**!*!!!*!!!!!!!!!!!*!*!

logo

NTDB id 1021138 ABZP31 RS13165 WP 367156824.1  QQLLTASILTLRLPKVQAIVPMPISTERLIERGYNQSLVIAKILARELNVPVWQPVIRLAQHSQKGLSRLERLNQIEMQF   159
NTDB id 1042 HON27 RS01930 WP 000472273.1      QTLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLSKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQF   157
NTDB id 1071 ABD1 RS15375 WP 000472269.1       QTLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLGKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQF   157
consensus                                      !*!!******!**!!!!!!!!!!!!!**!!*!!!*!!!!**!**!***!**!!!!!!*!!**!!!!!!!!!!!!***!!*!!

logo

NTDB id 1021138 ABZP31 RS13165 WP 367156824.1  QPILTERRRYRRVLMLDDVVTTGSSLSALAQALTQLGCQHIFAACIAAAETA..   211
NTDB id 1042 HON27 RS01930 WP 000472273.1      VALTQEKRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY   211
NTDB id 1071 ABD1 RS15375 WP 000472269.1       VALTQENRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY   211
consensus                                      *****!*!!!!!!!!**!!!*!!!!!**!!*!!!*!!!!**!*!*!*!!!*****
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X non conserved

X similar

X ≥ 50% conserved