```
logo

NTDB id 1002641 ABEF85 RS13030 WP 347453547.1  MPMFKLIQQGKHWLGKLQPCQLCHHDYQAIHSVCQDCWQQLPWACQTVQRQEIPVQIACHYAYPMDRLIQLFKYEQKLYL   80
NTDB id 1042 HON27 RS01930 WP 000472273.1      ..MFKFLNP.QYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHY   77
NTDB id 1071 ABD1 RS15375 WP 000472269.1       ..MFKFLNP.QYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHY   77
consensus                                        !!!**** ******!*!!*!!********!*!**!!*!!!!**!!*!!****!**!!!!!!!**!*!!*!!!!!!!**

logo

NTDB id 1002641 ABEF85 RS13030 WP 347453547.1  ENLLAGTLLSLNLPKVSAVVPMPISTEKLVQRGYNQSLLLAKHVARHLNVPVWQPITRLKQHSQKGLCRLERIEDIDTQF  160
NTDB id 1042 HON27 RS01930 WP 000472273.1      QTLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLSKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQF  157
NTDB id 1071 ABD1 RS15375 WP 000472269.1       QTLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLGKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQF  157
consensus                                      **!!!**!**!**!!!*!*!!!!!*!**!**!!*!!!!!!!***********!**!!!!!**!!**!!!!!!*!!!!*!*!**!!

logo

NTDB id 1002641 ABEF85 RS13030 WP 347453547.1  QMNTSSKLRYRKVLIIDDVITTGSSIRALSQKLEELGCQQINAACVAGAKL...  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 $\geq$ 50% conserved