```
logo
NTDB id 270905 C3Y93 RS07860 WP 104852593.1  MTQPINLLQCGIQWLRQLQPCQLCHSDTQWQHSVCQTCWQQLPWRSQAVQRQEIQIQVAFDYHYPINRIIQQFKYEQQLH   80
NTDB id 1042 H0N27 RS01930 WP 000472273.1    ....MFKFLNPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLH   76
NTDB id 1071 ABD1 RS15375 WP 000472269.1     ....MFKFLNPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLH   76
consensus                                     *************!*!!*!!***!****!*!**!!*!!!!**!**!!******!!**!*!!!!!!!!!!!!!!*!!

logo
NTDB id 270905 C3Y93 RS07860 WP 104852593.1  YQLLLVGALKQLSYPKIQAIVPMPISTARLTERGYNQSLLLAKGLARHLNVPIWQPVTRLQQHAQKGLSRLERLEDIQQQ  160
NTDB id 1042 H0N27 RS01930 WP 000472273.1    YQTLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLSKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQ  156
NTDB id 1071 ABD1 RS15375 WP 000472269.1     YQTLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLGKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQ  156
consensus                                    !!*!!***!*!!**!!*!!!!!!!!!**!!!!!!*!!!!!!!**!***!**!*!!!!*!!**!*!!!!!!!!!*!*!*!!

logo
NTDB id 270905 C3Y93 RS07860 WP 104852593.1  FQIQNAPKIRFRRVLIVDDVVTTGSSIHALKQQLLKLGCQKVEAVCLASAE....  211
NTDB id 1042 H0N27 RS01930 WP 000472273.1    FVALTQEKRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY  211
NTDB id 1071 ABD1 RS15375 WP 000472269.1     FVALTQENRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY  211
consensus                                    !********!*!!!!!*!!!*!!!!!!!!!!*!*!**!!!****!*!!!*!*****
```

X non conserved

X similar

X ≥ 50% conserved