```
logo
NTDB id 1107001 ACLKM3 RS01600 WP 236162857.1  MIQPINLAQRGIQWLRQLQPCQLCHSDTQWQHSVCQTCWQELPWRSQAVQRQEMQIQVAFDYHYPINRIIQQFKYEQQLQ   80
NTDB id 1042 HON27 RS01930 WP 000472273.1      MFKF....LNPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLH   76
NTDB id 1071 ABD1 RS15375 WP 000472269.1       MFKF....LNPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLH   76
consensus                                      !***    *********!*!!*!!***!****!*!**!!**!!!**!**!!******!!**!*!!!!!!!!!!!!!!!*!*

logo
NTDB id 1107001 ACLKM3 RS01600 WP 236162857.1  YQLLLVGALKQLSYPKIQAIVPMPISSTRLTERGYNQSLLLAKGLARYLDVPIWQPVTRLQKHAQKGLSRLERLEDIQQQ  160
NTDB id 1042 HON27 RS01930 WP 000472273.1      YQTLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLSKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQ  156
NTDB id 1071 ABD1 RS15375 WP 000472269.1       YQTLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLGKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQ  156
consensus                                      !!*!!***!*!!**!!*!!!!!!!!!**!!!!!!*!!!!!!!**!*****!**!*!!!!!*!!**!*!!!!!!!!!*!*!*!!

logo
NTDB id 1107001 ACLKM3 RS01600 WP 236162857.1  FQIQAAPKIRYRRVLIVDDVVTTGSSIHALKQQLLKLGCQKVEAVCLASAE....  211
NTDB id 1042 HON27 RS01930 WP 000472273.1      FVALTQEKRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY  211
NTDB id 1071 ABD1 RS15375 WP 000472269.1       FVALTQENRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY  211
consensus                                      !********!!!!!!!*!!!*!!!!!!!!!!*!*!**!!!****!*!!!*!*****
```

X non conserved

X similar

X $\geq$ 50% conserved