```
logo
NTDB id 301318 DPR03 RS01790 WP 071319847.1  MPMSKLIQQGKHWLDKLQPCQLCHNDYQTIYSVCQDCWNQLPWAHQTIQRQEMQVQIACHYDYPMDRLIQLFKYEQKLYL   80
NTDB id 1042 HON27 RS01930 WP 000472273.1    ..MFKFL.NPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHY   77
NTDB id 1071 ABD1 RS15375 WP 000472269.1     ..MFKFL.NPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHY   77
consensus                                      !*!** ********!*!!*!!*******!!*!**!!*!!!!**!!!!!****!**!!!!*!!**!*!!*!!!!!!!**

logo
NTDB id 301318 DPR03 RS01790 WP 071319847.1  ENLLAGALISLKLPKVSAVVPMPISTDKLVQRGYNQSLLLAKQVAKHLQVPVWQPVSRLKQHSQKGLSRLERIEDIEAQF  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 301318 DPR03 RS01790 WP 071319847.1  QISTSSKLRYRKVLIIDDVVTTGSSIRALNQKLQELGCQQIYTACVAGAKL...  211
NTDB id 1042 HON27 RS01930 WP 000472273.1    VALTQEKRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY  211
NTDB id 1071 ABD1 RS15375 WP 000472269.1     VALTQENRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY  211
consensus                                    ***!****!!!*!!!!!!!*!!!!!!*!!*!*!**!!!**!***!*!*!*****
```

X non conserved

X similar

X $\geq$ 50% conserved