```
logo
NTDB id 395917 GFH30 RS11385 WP 153372705.1  MPMSNLIKQAKTWLQQLQPCQLCGSDYQHIHSVCEDCWSQLPWAKTTVQRQEMQVHIACDYTYPMDRMIQLFKYEQKLHL   80
NTDB id 1042 HON27 RS01930 WP 000472273.1    ..MFK.FLNPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHY   77
NTDB id 1071 ABD1 RS15375 WP 000472269.1     ..MFK.FLNPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHY   77
consensus                                      !** ********!*!*!!*!!********!*!**!!*!!!!*!*!*!!****!**!!*!*!!**!*!!*!!!!!!!!*

logo
NTDB id 395917 GFH30 RS11385 WP 153372705.1  QNVLAGTLLSLRLPKVSAIVPMPISTERLVERGYNQSLVLAKHVAKHLNVPIWQPITRRSQHSQKGLSRLERMDDIEAQF  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 395917 GFH30 RS11385 WP 153372705.1  EISTQSKLCYRRVLIIDDVVTTGSSIHALKLKLEALGCREIYAACIAGAKI...  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