```
logo
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NTDB id 390162 FSC10 RS02135 WP 004896158.1  MPMSKLIQQGKHWLGKLQPCQLCHNDYQTIHSVCQDCWNQLPWAHQTIQRQEMQVQIACHYDYPMDRLIQLFKYEQKLYL  80
NTDB id 1042 HON27 RS01930 WP 000472273.1    ..MFKFL.NPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHY  77
NTDB id 1071 ABD1 RS15375 WP 000472269.1     ..MFKFL.NPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHY  77
consensus                                      !*!** ********!*!!*!!********!*!**!!*!!!!**!!!!!****!**!!!!*!!**!*!!*!!!!!!!**
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logo
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```
NTDB id 390162 FSC10 RS02135 WP 004896158.1  ENLLAGALISLKLPKVSAVVPMPISTDKLVQRGYNQSLLLAKQVAKHLQVPVWQPVSRLKQHSQKGLSRLERIEGIEAQF  160
NTDB id 1042 HON27 RS01930 WP 000472273.1    QTLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLSKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQF  157
NTDB id 1071 ABD1 RS15375 WP 000472269.1     QTLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLGKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQF  157
consensus                                    **!!!**!**!!*!!!*!*!!!!!!***!**!!*!!!!!!!***!*!**!!!!!!*!!**!!!!!!!!!!!*!*!!*!!
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logo
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NTDB id 390162 FSC10 RS02135 WP 004896158.1  QISTSSKLRYRKVLVIDDVVTTGSSIRALNQKLEELGCQQIYTACVAGAKL...  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 ≥ 50% conserved