```
logo
NTDB id 23414 ACIAD RS14675 WP 011182736.1  MLMSNLLATTLQWLKKECVPCPLCELSPQQKQQILCKDCWQELPWFKDYVTRHELNILAACEYRYPIDQIVQQFKYHDRLHYQSL   85
NTDB id 1042 HON27 RS01930 WP 000472273.1  ..MF..KFLNPQYLFQLLSPCLLCEIGTREK.YSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHYQTL   80
NTDB id 1071 ABD1 RS15375 WP 000472269.1  ..MF..KFLNPQYLFQLLSPCLLCEIGTREK.YSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHYQTL   80
consensus                                   !*  *****!*!*****!!*!!!*****! **!!!*!!**!!!*!****!*****!*!!*!*!!!**!*!!!!!***!!!!*!

logo
NTDB id 23414 ACIAD RS14675 WP 011182736.1  LSSLLLTLPLPRVQAIVPMPISEKRLIERGYNQSLLLAQSIAKHLNVPVWQPVERLAQHSQKGLSRLERISQIDEQFIVRQPVKE   170
NTDB id 1042 HON27 RS01930 WP 000472273.1  LAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLSKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQFVALTQEKR   165
NTDB id 1071 ABD1 RS15375 WP 000472269.1  LAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLGKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQFVALTQENR   165
consensus                                  !***!**!**!*!!!!!!!!!!**!!*!!!*!!!!!!!***!*!*!**!!!!!!*!!**!!!!!!!!!!!***!**!!********

logo
NTDB id 23414 ACIAD RS14675 WP 011182736.1  RYRRVMIVDDVVTTGSSIHALTHVLQNLGCVHIYPVCIAAVFD...   213
NTDB id 1042 HON27 RS01930 WP 000472273.1  RYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY   211
NTDB id 1071 ABD1 RS15375 WP 000472269.1  RYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY   211
consensus                                  !!!!!*!*!!!!*!!!!!!!!!***!**!!!**!***!*!!******
```

X non conserved

X similar

X ≥ 50% conserved