```
logo
NTDB id 1108006 ACN19N RS01865 WP 269229511.1  MFKFISRSTRQLFDYLTPCSLCDIGIKRHFGVCQSCWEQLPWLKQSIERNQQQIFVTCHYQYPIDRIIQQFKYEQKLHHQ   80
NTDB id 1042 H0N27 RS01930 WP 000472273.1      MFKFLN..PQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHYQ   78
NTDB id 1071 ABD1 RS15375 WP 000472269.1       MFKFLN..PQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHYQ   78
consensus                                      !!!!**  ***!**!*!!*!!*!!*******!**!!!!!!!!!!*!*!!*!***!*!!!*!!!*!!!!!!!!!!!!!!*!

logo
NTDB id 1108006 ACN19N RS01865 WP 269229511.1  RLLEGLLQQLKLPKVHAIVPMPISTDRLVERGFNQTLLLAQALSKSLNVPIWQPVQRLAQHSQKGLSRIERLENIEQQFV  160
NTDB id 1042 H0N27 RS01930 WP 000472273.1      TLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLSKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQFV  158
NTDB id 1071 ABD1 RS15375 WP 000472269.1       TLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLGKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQFV  158
consensus                                      *!!***!!!!!*!!!*!!!!!!!!**!!*!!!!!!*!!!!**!*!*!**!*!!!!!!!**!!!!!!!!*!!*!!!!!!!!

logo
NTDB id 1108006 ACN19N RS01865 WP 269229511.1  AAPPNHLRYRKVLIIDDVVTTGSSITALSYVLQQLGCQQIYSICLAAAVNHQASTLDFADTMESPHVE  228
NTDB id 1042 H0N27 RS01930 WP 000472273.1      ALTQEKRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY...............  211
NTDB id 1071 ABD1 RS15375 WP 000472269.1       ALTQENRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY...............  211
consensus                                      !**** *!!!*!!!!!!!*!!!!!!*!!!**!*!!!!**!***!!!!!*****
```

X non conserved

X similar

X $\geq$ 50% conserved