```
logo

NTDB id 1006852 ABLB98 RS12410 WP 348882346.1  MFKLIQTSAKQIFDYLTPCSLCDIGIRKQYGVCQSCWNQLPWLKQSIQRNQQQIFVVCHYQYPIDRIIQQFKYEQKLHHQ   80
NTDB id 1042 HON27 RS01930 WP 000472273.1      MFKFL..NPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHYQ   78
NTDB id 1071 ABD1 RS15375 WP 000472269.1       MFKFL..NPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHYQ   78
consensus                                      !!!**  *****!**!*!!*!!*!!*!**!**!**!!*!!!!!!!*!!!!!*!***!*!!!*!!!*!!!!!!!!!!!!!!*!

logo

NTDB id 1006852 ABLB98 RS12410 WP 348882346.1  RLLEGLLQQLKLPKVHAIVPMPISTDRLVERGFNQTLLLAQALSKSLNIPIWQPVQRLDQHSQKGLSRIERLENIEQQFV  160
NTDB id 1042 HON27 RS01930 WP 000472273.1      TLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLSKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQFV  158
NTDB id 1071 ABD1 RS15375 WP 000472269.1       TLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLGKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQFV  158
consensus                                      *!!***!!!!!*!!!*!!!!!!!!**!!*!!!!!!*!!!!**!*!*!*!!*!!!!!!!**!!!!!!!!*!!*!!!!!!!!

logo

NTDB id 1006852 ABLB98 RS12410 WP 348882346.1  AAPPNHLRYRKVLIVDDVVTTGSSITALSQVLQQLGCQQVYSICLAAPIRNNASSLNFADTMESPRVE  228
NTDB id 1042 HON27 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