```
logo
NTDB id 313838 CDG60 RS16095 WP 087512839.1  MLSFINQLLNKGVQSISPCLLCGIDHQQQHSLCNDCWEQLPWYKQHIQRHEHSILCAHHYDFPMNRIIQTYKYEQQLQYQ   80
NTDB id 1042 HON27 RS01930 WP 000472273.1    MFKFLNPQ..YLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHYQ   78
NTDB id 1071 ABD1 RS15375 WP 000472269.1     MFKFLNPQ..YLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHYQ   78
consensus                                    !**!*!**  ***!**!!!!!!*!******!!**!!!!!!!*!!*!!!***!*!*!*!!**!*!!!!!**!!!!*!*!!

logo
NTDB id 313838 CDG60 RS16095 WP 087512839.1  NLLARSLLNLRTPKVHAIVPMPISTERLKERGYNQMLIIAKIMARELKIPVWQPVIRAAQYSQKGLSRIERLENIEEQFQ  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 313838 CDG60 RS16095 WP 087512839.1  IIPAEKRKYKKVLIIDDVVTTGSSIHALSQALEKLGCQQIYTACIAAGGIKRQSQPGEVEINFKEND  227
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