```
logo
NTDB id 1002843 ABEF90 RS02595 WP 347474256.1  ........MPLVKSWWKDPIFSGAVAIAAILHIMILLLQFGMPQERDTSSKEIAVSLRISNDKIEHADFLAQTDQQGSGQ   72
NTDB id 1078 ABD1 RS15030 WP 000914774.1       MLKKRSLSIEPFYYWWQDRVFIAAIILAILLHIFVLLIHFAMPAPSEQSTKEIAISIRPTEDVIKHADFLAQADQQGSGA   80
consensus                                      ***********   !! !**! *!* *! *!!! *!!**!*!! * * !*!!!!*!*!** ! ! !!!!!!! !!!!!!

logo
NTDB id 1002843 ABEF90 RS02595 WP 347474256.1  FREAHRMSSDAPMPIPADEAAVGEAQQQSEDMLRQKQELSFEEKVLMTALSWQKQAEENQRKKRQEELNSQFQVKAAMVA  152
NTDB id 1078 ABD1 RS15030 WP 000914774.1       FREAHRMSSNSPAPMPSDA.STGEAQLETLEKVQQQRELKFEEKVLMTVLSWQKQAEESQRKKALEQLQSQFQAKAAMVA  159
consensus                                      !!!!!!!!!!  ! !*! ! *  !!!!  * * * !  !! !!!!!!!! !!!!!!!!! !!!!  ! !*!!!! !!!!!!

logo
NTDB id 1002843 ABEF90 RS02595 WP 347474256.1  SLEAQYMQRQQNFSRQQKIKTVDGIQAKQDVSAAYLEKFRQKVEFYGNRFYPESARLQRLAGDVRLMVILNQNGGIRAIR  232
NTDB id 1078 ABD1 RS15030 WP 000914774.1       SLEAQYLQRQQDFSRQQKIKTVDGIQAKKDASAAYLDKFREKVELYGNRYYPEEAKQQQLKGEVRLMVILNAQGGIRAIR  239
consensus                                      !!!!!!*!!!! !!!!!!!!!!!!!!!! ! !!!!!*!!! !!! !!!!*!!! !* ! ! !*!!!!!!!! *!!!!!!!

logo
NTDB id 1002843 ABEF90 RS02595 WP 347474256.1  LLESSGHAVLDEAAKMSLRKAAPFGVFDAKMKDISELRIIRTWRFDPAEAEFEVR  287
NTDB id 1078 ABD1 RS15030 WP 000914774.1       LLESSGHPVLDEAAKASVRRGAPFGRFDANMKDISELRIIRTWRFDPAEAEFEVH  294
consensus                                      !!!!!!!*!!!!!!! !*!**!!!! !!! !!!!!!!!!!!!!!!!!!!!!!!!*
```

X non conserved

X similar

X $\geq$ 50% conserved