```
logo
NTDB id 287812 CDG57 RS13720 WP 004958893.1  MWNKINFRSELSSHWWQDSIFVGAVILALLLHLAVLAVQFAMPTPSDTSTKEIAVAIRPSTDKVENADFLAQADQKGSGE   80
NTDB id 1078 ABD1 RS15030 WP 000914774.1     MLKKRSLSIEPFYYWWQDRVFIAAIILAILLHIFVLLIHFAMPAPSEQSTKEIAISIRPTEDVIKHADFLAQADQQGSGA   80
consensus                                    !  !     !*  *!!!! *!**!*!!!*!!!* !! **!!!! !!* !!!!!!* !!!* ! * *!!!!!!!!! !!!

logo
NTDB id 287812 CDG57 RS13720 WP 004958893.1  FREAHRMSSDMPSPMQADATTGETEFESLEKIQQKRELKFEEKVLMTVLSWQKQAEQSERKKALEDLQSQFQAKAAMVAS  160
NTDB id 1078 ABD1 RS15030 WP 000914774.1     FREAHRMSSNSPAPMPSDASTGEAQLETLEKVQQQRELKFEEKVLMTVLSWQKQAEESQRKKALEQLQSQFQAKAAMVAS  160
consensus                                    !!!!!!!!!  ! !!* !!*!!!   !*!!!*!! !!!!!!!!!!!!!!!!!!!!!! ! !!!!!! !!!!!!!!!!!!!!

logo
NTDB id 287812 CDG57 RS13720 WP 004958893.1  LEAQYLQRQQNFSRKQRIKTVDGIQAKQDASAAYLDKFRQKVELYGNRYYPDAAKQQRLSGEVRLMVILNSQGGIRAIRL  240
NTDB id 1078 ABD1 RS15030 WP 000914774.1     LEAQYLQRQQDFSRQQKIKTVDGIQAKKDASAAYLDKFREKVELYGNRYYPEEAKQQQLKGEVRLMVILNAQGGIRAIRL  240
consensus                                    !!!!!!!!!! !!! !*!!!!!!!!!! !!!!!!!!!!! !!!!!!!!!!!* !!!! ! !!!!!!!!!! !!!!!!!!!

logo
NTDB id 287812 CDG57 RS13720 WP 004958893.1  IESSGHSILDEAAKASVRRGAPFGAFDSTMKKDISELRIIRTWRFDPVDAEFEVH   295
NTDB id 1078 ABD1 RS15030 WP 000914774.1     LESSGHPVLDEAAKASVRRGAPFGRFDANM.KDISELRIIRTWRFDPAEAEFEVH   294
consensus                                    *!!!!!**!!!!!!!!!!!!!!!! !!   !*!!!!!!!!!!!!!!!! *!!!!!!
```

X non conserved

X similar

X ≥ 50% conserved