```
logo

NTDB id 395916 GFH30 RS11050 WP 171501032.1  ........MPLMKSWWKDPIFTGAVIIAVALHIMLLALQFGMPQEQDAMSKEIAVTLRQADKEPEHADFLAQTNQQGSGE   72
NTDB id 1078 ABD1 RS15030 WP 000914774.1     MLKKRSLSIEPFYYWWQDRVFIAAIILAILLHIFVLLIHFAMPAPSEQSTKEIAISIRPTEDVIKHADFLAQADQQGSGA   80
consensus                                    ***********   !! !**! *!*!*!* !!! *! **!*!! * *  *!!!!***!* *  * !!!!!!!  !!!!!

logo

NTDB id 395916 GFH30 RS11050 WP 171501032.1  FREAHRMSSDMPAPMLEQ.KAGESQQESLDMLEQKQQLSFEEKILMTTLSWQKQAEDAKRKKIQEQLNSQFQAKAAMVAS   151
NTDB id 1078 ABD1 RS15030 WP 000914774.1     FREAHRMSSNSPAPMPSDASTGEAQLETLEKVQQQRELKFEEKVLMTVLSWQKQAEESQRKKALEQLQSQFQAKAAMVAS   160
consensus                                    !!!!!!!!!  !!!!*  *  !! ! !*!* * !   ! !!!!*!!! !!!!!!!!*  !!!  !!!*!!!!!!!!!!!!!

logo

NTDB id 395916 GFH30 RS11050 WP 171501032.1  LEAQYLQRQKNFSRQQKIKTVDGIQAKQDVSAAYLEKFRQKVEFYGNRFYPDAARAQKLAGEVRLMVILNQNGGIRAIRL   231
NTDB id 1078 ABD1 RS15030 WP 000914774.1     LEAQYLQRQQDFSRQQKIKTVDGIQAKKDASAAYLDKFREKVELYGNRYYPEEAKQQQLKGEVRLMVILNAQGGIRAIRL   240
consensus                                    !!!!!!!!!  !!!!!!!!!!!!!!!! ! !!!!!*!!! !!! !!!!*!!* !* ! ! !!!!!!!!!! *!!!!!!!!

logo

NTDB id 395916 GFH30 RS11050 WP 171501032.1  IDSSGHGILDEAAKASVRKSAPFGAFDAKMSDISELRIIRTWRFDPADAEFEVR   285
NTDB id 1078 ABD1 RS15030 WP 000914774.1     LESSGHPVLDEAAKASVRRGAPFGRFDANMKDISELRIIRTWRFDPAEAEFEVH   294
consensus                                    **!!!!**!!!!!!!!!!* !!!! !!! ! !!!!!!!!!!!!!!!!!*!!!!!*
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X non conserved

X similar

X ≥ 50% conserved