```
logo

NTDB id 1107162 ACLKMZ RS02625 WP 411691453.1  ........MPLVKEWWKDPVFTTAVLLAALLHLAFLTMQFTMPQEGEYNTKEIAVSLRPATDEVEHADFLAQADQQGSGT   72
NTDB id 1078 ABD1 RS15030 WP 000914774.1       MLKKRSLSIEPFYYWWQDRVFIAAIILAILLHIFVLLIHFAMPAPSEQSTKEIAISIRPTEDVIKHADFLAQADQQGSGA   80
consensus                                      ***********   !! !*!!  !**!! !!!*  ! **! !! * !  !!!!!*!*!!  ! * !!!!!!!!!!!!!!

logo

NTDB id 1107162 ACLKMZ RS02625 WP 411691453.1  FRESHRMSSDMPAPMIEQ.KAGEQKNEALEQLQQKKQLSFEEKVLMTTLSWQKQAEENSRKKAVEELNSQFQAKAAMVAS   151
NTDB id 1078 ABD1 RS15030 WP 000914774.1       FREAHRMSSNSPAPMPSDASTGEAQLETLEKVQQQRELKFEEKVLMTVLSWQKQAEESQRKKALEQLQSQFQAKAAMVAS   160
consensus                                      !!! !!!!!  !!!!*  *  !!   ! !! *!! * ! !!!!!!!!! !!!!!!!!!!  !!!!*! !*!!!!!!!!!!!!!

logo

NTDB id 1107162 ACLKMZ RS02625 WP 411691453.1  IEAQYLQRQQNFSRQQKIKTIDGIQAKQDVSAAYMEKFRQKVEFYGNRYYPEAAKQQQLSGEVRLMVILNTNGGIRAIRL   231
NTDB id 1078 ABD1 RS15030 WP 000914774.1       LEAQYLQRQQDFSRQQKIKTVDGIQAKKDASAAYLDKFREKVELYGNRYYPEEAKQQQLKGEVRLMVILNAQGGIRAIRL   240
consensus                                      *!!!!!!!!! !!!!!!!!!*!!!!!! ! !!!!**!!! !!! !!!!!!!! !!!!!! !!!!!!!!!! *!!!!!!!!

logo

NTDB id 1107162 ACLKMZ RS02625 WP 411691453.1  IESSGSTLLDDAAKTSVRKAAPFGQFDAKMKDISELRIIRTWRFDPAEAEFEVR   285
NTDB id 1078 ABD1 RS15030 WP 000914774.1       LESSGHPVLDEAAKASVRRGAPFGRFDANMKDISELRIIRTWRFDPAEAEFEVH   294
consensus                                      *!!!!***!!*!!! !!!**!!!! !!! !!!!!!!!!!!!!!!!!!!!!!!!*
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X non conserved

X similar

X ≥ 50% conserved