```
logo

NTDB id 1043282 ACEN3N RS18740 WP 004825113.1  MIKKLFANTQHTGYWWQDALFVGAVSIAVIVHGLFLSVQFGMPQENNTATKEIAVSIRPSEDKVKDADFLAQADQQGSGT   80
NTDB id 1078 ABD1 RS15030 WP 000914774.1       MLKKRSLSIEPFYYWWQDRVFIAAIILAILLHIFVLLIHFAMPAPSEQSTKEIAISIRPTEDVIKHADFLAQADQQGSGA   80
consensus                                      !*!!          !!!!! *!**!* *!***!   ! **!*!! *    !!!!!*!!!!*!! *!*!!!!!!!!!!!!!!

logo

NTDB id 1043282 ACEN3N RS18740 WP 004825113.1  FRKEHRMSSDMPAQA.EDKSAGDQQLEALEQIQQKRELKFEEKVLMTVLSWQKQAEQNDRKKAMEELQSQMQAKAAMVAS  159
NTDB id 1078 ABD1 RS15030 WP 000914774.1       FREAHRMSSNSPAPMPSDASTGEAQLETLEKVQQQRELKFEEKVLMTVLSWQKQAEESQRKKALEQLQSQFQAKAAMVAS  160
consensus                                      !!  !!!!!  !!* * ! ! !* !!! !! *!! !!!!!!!!!!!!!!!!!!!!!!   !!!!*! !!!!! !!!!!!!!!

logo

NTDB id 1043282 ACEN3N RS18740 WP 004825113.1  LEAQYLQKQQNFSKQQKIKTLDGIQAKQDASAGYLDKFREKVEMYGNRYYPDLAKQQRLSGEVRLMVILNNEGGIRAIRL  239
NTDB id 1078 ABD1 RS15030 WP 000914774.1       LEAQYLQRQQDFSRQQKIKTVDGIQAKKDASAAYLDKFREKVELYGNRYYPEEAKQQQLKGEVRLMVILNAQGGIRAIRL  240
consensus                                      !!!!!!!*!! !!*!!!!!!*!!!!!!! !!!!*!!!!!!!!!!*!!!!!!!* !!!! ! !!!!!!!!!!  !!!!!!!!

logo

NTDB id 1043282 ACEN3N RS18740 WP 004825113.1  IESSGHAILDEAAKNSVRKGAPFGRFDAKMKDISELRIVRTWRFDPAESEFEVH  293
NTDB id 1078 ABD1 RS15030 WP 000914774.1       LESSGHPVLDEAAKASVRRGAPFGRFDANMKDISELRIIRTWRFDPAEAEFEVH  294
consensus                                      *!!!!!**!!!!!! !!!*!!!!!!!!!! !!!!!!!!!*!!!!!!!!!! !!!!!
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X non conserved

X similar

X ≥ 50% conserved