```
logo

NTDB id 1107091 ACLKM1 RS11080 WP 005094077.1  ........MPLVRTWWKDPVFSSAVLTAGILHLIVLTLEFGMPQDRETSTKEIAVSLRQTDEKIEHADFLAQTDQQGSGK   72
NTDB id 1078 ABD1 RS15030 WP 000914774.1       MLKKRSLSIEPFYYWWQDRVFIAAIILAILLHIFVLLIHFAMPAPSEQSTKEIAISIRPTEDVIKHADFLAQADQQGSGA   80
consensus                                      ***********   !! !*!!  !** ! *!!* !! **!*!! * ! !!!!!!*!*!*!** ! !!!!!!! !!!!!!

logo

NTDB id 1107091 ACLKM1 RS11080 WP 005094077.1  FREAHRMSSDMPSPLLEQ.KAGEEQQESPDLLQQKQELSFEEKVLMTTLSWQKQAEENQRKKMQEQLQSQFQAKAAMVAS  151
NTDB id 1078 ABD1 RS15030 WP 000914774.1       FREAHRMSSNSPAPMPSDASTGEAQLETLEKVQQQRELKFEEKVLMTVLSWQKQAEESQRKKALEQLQSQFQAKAAMVAS  160
consensus                                      !!!!!!!!!  ! !**  *  !! ! !*** *!!  !! !!!!!!!! !!!!!!!!! !!!!  !!!!!!!!!!!!!!!!

logo

NTDB id 1107091 ACLKM1 RS11080 WP 005094077.1  LEAQYLQRQQNFSRQQKIKTVDGIQSKQDVSAAYLEKFRQKVEFYGNQYYPEAAKQQNLAGEVRLMVILNQNGGIRAIRL  231
NTDB id 1078 ABD1 RS15030 WP 000914774.1       LEAQYLQRQQDFSRQQKIKTVDGIQAKKDASAAYLDKFREKVELYGNRYYPEEAKQQQLKGEVRLMVILNAQGGIRAIRL  240
consensus                                      !!!!!!!!!! !!!!!!!!!!!!!! ! ! !!!!!*!!! !!! !!! !!!! !!!!*! !!!!!!!!!! *!!!!!!!!

logo

NTDB id 1107091 ACLKM1 RS11080 WP 005094077.1  IDSSGHAMLDEAAKSSVRKAAPFGAFDAKMKEISELRIIRTWRFDPAEAEFEVR  285
NTDB id 1078 ABD1 RS15030 WP 000914774.1       LESSGHPVLDEAAKASVRRGAPFGRFDANMKDISELRIIRTWRFDPAEAEFEVH  294
consensus                                      **!!!!**!!!!!! !!!**!!!! !!! !!*!!!!!!!!!!!!!!!!!!!!!*
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X non conserved

X similar

X ≥ 50% conserved