```
logo

NTDB id 213377 BVG18 RS02540 WP 216986319.1  ........MPLVRTWWKDPVFSSAVLAAGILHLIVLTLEFGMPQDRDTSTKEIAVSLRQTDEKIEHADFLAQTDQQGSGQ   72
NTDB id 1078 ABD1 RS15030 WP 000914774.1     MLKKRSLSIEPFYYWWQDRVFIAAIILAILLHIFVLLIHFAMPAPSEQSTKEIAISIRPTEDVIKHADFLAQADQQGSGA   80
consensus                                    ***********   !! !*!!  !** ! *!!* !! **!*!! * * !!!!!!*!*!*!** ! !!!!!!! !!!!!!

logo

NTDB id 213377 BVG18 RS02540 WP 216986319.1  FREAHRMSSDMPSPLMEQ.HAGEELQESLDMLQQQQELSFEEKVLMTTLSWQKQAEENQRKKMQEQLQSQFQAKAAMVAS   151
NTDB id 1078 ABD1 RS15030 WP 000914774.1     FREAHRMSSNSPAPMPSDASTGEAQLETLEKVQQQRELKFEEKVLMTVLSWQKQAEESQRKKALEQLQSQFQAKAAMVAS   160
consensus                                    !!!!!!!!!  ! !**  ** !!   !*!* *!!! !! !!!!!!!! !!!!!!!!! !!!!  !!!!!!!!!!!!!!!!

logo

NTDB id 213377 BVG18 RS02540 WP 216986319.1  LEAQYLQRQQNFSRQQKIKTVDGIQSKQDVSAAYLEKFREKVEFYGNRYYPEEAKQQNLAGEVRLMVILNQNGGIRAIRL   231
NTDB id 1078 ABD1 RS15030 WP 000914774.1     LEAQYLQRQQDFSRQQKIKTVDGIQAKKDASAAYLDKFREKVELYGNRYYPEEAKQQQLKGEVRLMVILNAQGGIRAIRL   240
consensus                                    !!!!!!!!!! !!!!!!!!!!!!!! ! ! !!!!!*!!!!!!! !!!!!!!! !!!!*! !!!!!!!!!! *!!!!!!!!

logo

NTDB id 213377 BVG18 RS02540 WP 216986319.1  IDSSGHAMLDEAAKASVRKAAPFGAFDSKMKEISELRVIRTWRFDPAEAEFEVR   285
NTDB id 1078 ABD1 RS15030 WP 000914774.1     LESSGHPVLDEAAKASVRRGAPFGRFDANMKDISELRIIRTWRFDPAEAEFEVH   294
consensus                                    **!!!!**!!!!!!!!!!**!!!! !!   !!*!!!!!*!!!!!!!!!!!!!!!*
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X non conserved

X similar

X ≥ 50% conserved