```
logo

NTDB id 310693 BEN71 RS16285 WP 068973622.1  MIKKLYADTKHSDYWWQDPLFIGAVSVAVITHLLFLSIQFGMPQQNDAATKEIAISIRPSEDKVKDADFLAQEDQQGSGT   80
NTDB id 1078 ABD1 RS15030 WP 000914774.1     MLKKRSLSIEPFYYWWQDRVFIAAIILAILLHIFVLLIHFAMPAPSEQSTKEIAISIRPTEDVIKHADFLAQADQQGSGA   80
consensus                                    !*!!          !!!!!**!!*!* *!** !*  ! !*!*!! * *  !!!!!!!!!!*!! *!*!!!!!! !!!!!!

logo

NTDB id 310693 BEN71 RS16285 WP 068973622.1  FRKEHRMSSDMPAQA.EDQSAGDQQLQALEQMQQKRELNFEEKVLMTVLSWQKQAEQNDRKKAMEELQSQMQAKAAMVAS  159
NTDB id 1078 ABD1 RS15030 WP 000914774.1     FREAHRMSSNSPAPMPSDASTGEAQLETLEKVQQQRELKFEEKVLMTVLSWQKQAEESQRKKALEQLQSQFQAKAAMVAS  160
consensus                                    !!  !!!!!  !!* * ! ! !* !!  !! *!! !!! !!!!!!!!!!!!!!!!!!   !!!!*! !!!!! !!!!!!!!!

logo

NTDB id 310693 BEN71 RS16285 WP 068973622.1  LEAQYLQKQQNFSKQQKIKTLDGIQAKQDASAGYLDKFREKVEMYGNRYYPDLAKQQRLSGEVRLMVILNNEGGIRAIRL  239
NTDB id 1078 ABD1 RS15030 WP 000914774.1     LEAQYLQRQQDFSRQQKIKTVDGIQAKKDASAAYLDKFREKVELYGNRYYPEEAKQQQLKGEVRLMVILNAQGGIRAIRL  240
consensus                                    !!!!!!!*!! !!*!!!!!!*!!!!!! !!!!*!!!!!!!!!!*!!!!!!!* !!!! ! !!!!!!!!!!  !!!!!!!!

logo

NTDB id 310693 BEN71 RS16285 WP 068973622.1  VESSGHSILDEAAKNSVRKGAPFGRFDAKMKDISELRIIRTWRFDPAEAEFEVR   293
NTDB id 1078 ABD1 RS15030 WP 000914774.1     LESSGHPVLDEAAKASVRRGAPFGRFDANMKDISELRIIRTWRFDPAEAEFEVH   294
consensus                                    *!!!!!**!!!!!! !!!*!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!*
```

X non conserved

X similar

X $\geq$ 50% conserved