```
logo
NTDB id 1123722 EL276 RS03390 WP 107879254.1  MRKSY.ILRHNLKAAQQGFSLFIVLIVMLVIAFLVVSATQSYNTEQRISTNDADRKFALSLAEAALQVGEAEILNF...K   76
NTDB id 1139 NGFG RS02435 WP 003687918.1      MRKQNTLTGIPTSDGQRGSALFIVLMVMIVVAFLVVTAAQSYNTEQRISANESDRKLALSLAEAALREGEFQVLDLEYTA   80
consensus                                     !!!  **  **   *! !  !!!!!*!!*!*!!!!!*! !!!!!!!!!! !* !!! !!!!!!!!!  !!  *!  ***

logo
NTDB id 1123722 EL276 RS03390 WP 107879254.1  DSQVVFSTDCSNGLCAAAGSEVSTTPRRV......VQAGGDTPAW.....KRKKECGGSLCIDTNGKIYNVGSSRGRRNA  145
NTDB id 1139 NGFG RS02435 WP 003687918.1      DSKVTFSENCEKGLCTAVNVRTNNNGNEEVFGNIVVQGTPTVEAVKRSCPAKSGKNSTGLCIDNQGVEYEKGTGNVSKMP  160
consensus                                     !! ! !!  !  !!! !        *   ******!!* *  *! ***** *   *   !!!! *!  !  !!*    * *

logo
NTDB id 1123722 EL276 RS03390 WP 107879254.1  RYIVEYVGTQPDGSPIYRVTSKAWGKNNNTVVMLQSYISVE...   186
NTDB id 1139 NGFG RS02435 WP 003687918.1      RYIIEYLGEKN.NQNIYRVTAKAWGKNANTVVVLQSYVGNNDEQ   203
consensus                                     !!!*!!*!  **  *!!!!! !!!!!! !!!!*!!!!*   ***
```

X non conserved

X similar

X $\geq$ 50% conserved