```
logo
NTDB id 156587 AN159 RS01740 WP 002246121.1  MRKQNTLTGIPTSDGQRGFALFIVLMVMIVVAFLVVTAAQSYNTEQRISANESDRKLALSLAEAALREGELQVLDLEYDT   80
NTDB id 1139 NGFG RS02435 WP 003687918.1     MRKQNTLTGIPTSDGQRGSALFIVLMVMIVVAFLVVTAAQSYNTEQRISANESDRKLALSLAEAALREGEFQVLDLEYTA   80
consensus                                    !!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo
NTDB id 156587 AN159 RS01740 WP 002246121.1  DSKVTFSENCAKGLCTAVNVRTNNTGNEEAFDNIVVQGNPTVEAVKRSCLAK....STGLCIDNKGMEYKKGTRSVSKPP  156
NTDB id 1139 NGFG RS02435 WP 003687918.1     DSKVTFSENCEKGLCTAVNVRTNNNGNEEVFGNIVVQGTPTVEAVKRSCPAKSGKNSTGLCIDNQGVEYEKGTGNVSKMP  160
consensus                                    !!!!!!!!!!! !!!!!!!!!!!!!! !!!!! ! !!!!!! !!!!!!!!!!!*!!****!!!!!!!!! !*!! !!!  !!!*!

logo
NTDB id 156587 AN159 RS01740 WP 002246121.1  RYIIEYLGVKNGENVYRVTAKAWGKNANTVVVLQSYVSNNDE.   198
NTDB id 1139 NGFG RS02435 WP 003687918.1     RYIIEYLGEKNNQNIYRVTAKAWGKNANTVVVLQSYVGNNDEQ   203
consensus                                    !!!!!!!! !!  !*!!!!!!!!!!!!!!!!!!!!!! !!!!*
```

X non conserved

X similar

X $\geq$ 50% conserved