```
logo
NTDB id 374 SMU RS06885 WP 002262930.1        ......MKKILIVDDEKPISDIIKFNLAKEGYDTITAFDGREALSKYEEENPDLIILDLMLPELDGLEVAKEVRK...NS   71
NTDB id 1027243 AB4P90 RS27670 WP 014720321.1 MEQHSNALKVMVIDDSKTIRRTAETLLKNMGCEVITAIDGFDALARIVDHHPHIIFVDIMMPRLDGYQTCALVKNNPAFK   80
NTDB id 1047 H0N27 RS03110 WP 000389061.1     MEDAFQNLKVMVIDDSKTIRRTAETLLQREGCEVITAVDGFEALSKIAEANPDIVFVDIMMPRLDGYQTCALIKNSQNYQ   80
consensus                                     **    * *!****!!*!*!*******! **!***!!!*!!**!!*** * *!*****!*!*!*!!!*********    *

logo
NTDB id 374 SMU RS06885 WP 002262930.1        HVPIIMLSAKDSEFDKVIGLEIGADDYVTKPFSNRELLARVKAHLRRTENIESAVAEENASGIPEIIIGDLQILPDAFVA  151
NTDB id 1027243 AB4P90 RS27670 WP 014720321.1 SIPVIMLSSKDGLFDKAKGRIVGVDQFLTKPFSKEELLGAIKAYVPGFAAVEQAH........................  135
NTDB id 1047 H0N27 RS03110 WP 000389061.1     NIPVIMLSSKDGLFDQAKGRVVGSDEYLTKPFSKDELLNAIRNHVSS...............................  127
consensus                                      *!*!!!!*!!**!!***!***! !***!!!!!**!!!*******     ** *

logo
NTDB id 374 SMU RS06885 WP 002262930.1        KKRGTEVELTHREFELLHHLATHTGQVMTREHLLETVWGYDYFGDVRTVDVTVRRLREKIEDTPSRPEYILTRRGVGYYM  231
NTDB id 1027243 AB4P90 RS27670 WP 014720321.1 ................................................................................  135
NTDB id 1047 H0N27 RS03110 WP 000389061.1     ................................................................................  127
consensus

logo
NTDB id 374 SMU RS06885 WP 002262930.1        KSYD  235
NTDB id 1027243 AB4P90 RS27670 WP 014720321.1 ....  135
NTDB id 1047 H0N27 RS03110 WP 000389061.1     ....  127
consensus
```

X non conserved
X similar
X ≥ 50% conserved