```
logo
NTDB id 374 SMU RS06885 WP 002262930.1          ......MKKILIVDDEKPISDIIKFNLAKEGYDTITAFDGREALSKYEEENPDLIILDLMLPELDGLEVAKEVRK...NS   71
NTDB id 1027315 AB4P94 RS08415 WP 395610168.1   MEQHSNALRVMVIDDSKTIRRTAETLLKNMGCDVITAIDGFDALARIVDHHPHIIFVDIMMPRLDGYQTCALVKNNPAFK   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 1027315 AB4P94 RS08415 WP 395610168.1   SIPVIMLSSKDGLFDKAKGRIVGVDQFLTKPFSKEELLGAIKAYVPAFAAVEQAH........................   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 1027315 AB4P94 RS08415 WP 395610168.1   ..................................................................................   135
NTDB id 1047 H0N27 RS03110 WP 000389061.1       ..................................................................................   127
consensus

logo
NTDB id 374 SMU RS06885 WP 002262930.1          KSYD   235
NTDB id 1027315 AB4P94 RS08415 WP 395610168.1   ....   135
NTDB id 1047 H0N27 RS03110 WP 000389061.1       ....   127
consensus
```

X non conserved
X similar
X ≥ 50% conserved