```
logo
NTDB id 374 SMU RS06885 WP 002262930.1      ......MKKILIVDDEKPISDIIKFNLAKEGYDTITAFDGREALSKYEEENPDLIILDLMLPELDGLEVAKEVRK...NSHVPII   76
NTDB id 93418 AK972 RS00880 WP 003213736.1  MEQHSNALKVMVIDDSKTIRRTAETLLKNVGCDVITAIDGFDALAKIVDHHPHIIFVDIMMPRLDGYQTCALVKNNPAFKSIPVI   85
NTDB id 1047 HON27 RS03110 WP 000389061.1   MEDAFQNLKVMVIDDSKTIRRTAETLLQREGCEVITAVDGFEALSKIAEANPDIVFVDIMMPRLDGYQTCALIKNSQNYQNIPVI   85
consensus                                   **    * *!****!!*!*!*******! **!***!!!*!!**!!*!* * *!*****!*!*!*!!!*********   *  *!*!

logo
NTDB id 374 SMU RS06885 WP 002262930.1      MLSAKDSEFDKVIGLEIGADDYVTKPFSNRELLARVKAHLRRTENIESAVAEENASGIPEIIIGDLQILPDAFVAKKRGTEVELT  161
NTDB id 93418 AK972 RS00880 WP 003213736.1  MLSSKDGLFDKAKGRIVGSDQFLTKPFSKEELLSAIKAYVPGFAAVEQAH.................................  135
NTDB id 1047 HON27 RS03110 WP 000389061.1   MLSSKDGLFDQAKGRVVGSDEYLTKPFSKDELLNAIRNHVSS.......................................  127
consensus                                   !!!*!!**!!***!***!*!***!!!!!**!!! ******     ** *

logo
NTDB id 374 SMU RS06885 WP 002262930.1      HREFELLHHLATHTGQVMTREHLLETVWGYDYFGDVRTVDVTVRRLREKIEDTPSRPEYILTRRGVGYYMKSYD  235
NTDB id 93418 AK972 RS00880 WP 003213736.1  ......................................................................  135
NTDB id 1047 HON27 RS03110 WP 000389061.1   ......................................................................  127
consensus
```

X non conserved

X similar

X ≥ 50% conserved