```
logo
NTDB id 374 SMU RS06885 WP 002262930.1      ......MKKILIVDDEKPISDIIKFNLAKEGYDTITAFDGREALSKYEEENPDLIILDLMLPELDGLEVAKEVRK...NSHVPII   76
NTDB id 31110 CJA RS00390 WP 007639219.1    MEVTYESLKVMVIDDSKTIRRTAETLLKKAGCMVITATDGFDALAKIADTRPDIIFVDIMMPRLDGYQTCALIKNNSEFKTTPVI   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 31110 CJA RS00390 WP 007639219.1    MLSSKDGLFDKAKGRIVGSDQYLTKPFSKSELLGAIEAHVKHLKAS.......................................  131
NTDB id 1047 HON27 RS03110 WP 000389061.1   MLSSKDGLFDQAKGRVVGSDEYLTKPFSKDELLNAIRNHVSS.............................................  127
consensus                                   !!!*!!**!!***!***!*!*!*!!!!!* !!!*****!**

logo
NTDB id 374 SMU RS06885 WP 002262930.1      HREFELLHHLATHTGQVMTREHLLETVWGYDYFGDVRTVDVTVRRLREKIEDTPSRPEYILTRRGVGYYMKSYD  235
NTDB id 31110 CJA RS00390 WP 007639219.1    ......................................................................  131
NTDB id 1047 HON27 RS03110 WP 000389061.1   ......................................................................  127
consensus
```

X non conserved

X similar

X ≥ 50% conserved