```
logo
NTDB id 374 SMU RS06885 WP 002262930.1          ...........MKKILIVDDEKPISDIIKFNLAKEGYDTITAFDGREALSKYEEENPDLIILDLMLPELDGLEVAKEVRK   69
NTDB id 1009308 ABNK63 RS04150 WP 274378300.1  MNLNVSANDLAGLKVMVIDDSKTIRRTAETLLKKEGCDVLTAVDGFEALAKISDQKPAIIFVDIMMPRLDGYQTCALIKN   80
NTDB id 1047 HON27 RS03110 WP 000389061.1       .....MEDAFQNLKVMVIDDSKTIRRTAETLLQREGCEVITAVDGFEALSKIAEANPDIVFVDIMMPRLDGYQTCALIKN   75
consensus                                                 *!****!!*!*!*******!  *!!****!!*!!*!!!*!* * *!*****!*!*!*!!!*********

logo
NTDB id 374 SMU RS06885 WP 002262930.1          ...NSHVPIIMLSAKDSEFDKVIGLEIGADDYVTKPFSNRELLARVKAHLRRTENIESAVAEENASGIPEIIIGDLQILP   146
NTDB id 1009308 ABNK63 RS04150 WP 274378300.1  NPQFRATPVIMLSSKDGLFDKARGRIVGAEQYLTKPFTRDELLGAIHRHVSTVSSH.......................   136
NTDB id 1047 HON27 RS03110 WP 000389061.1       SQNYQNIPVIMLSSKDGLFDQAKGRVVGSDEYLTKPFSKDELLNAIRNHVSS...........................   127
consensus                                         **  *!*!!!!*!!**!!***!***!***!*!!!!**!!!**** !***

logo
NTDB id 374 SMU RS06885 WP 002262930.1          DAFVAKKRGTEVELTHREFELLHHLATHTGQVMTREHLLETVWGYDYFGDVRTVDVTVRRLREKIEDTPSRPEYILTRRG   226
NTDB id 1009308 ABNK63 RS04150 WP 274378300.1  ......................................................................................   136
NTDB id 1047 HON27 RS03110 WP 000389061.1       ......................................................................................   127
consensus

logo
NTDB id 374 SMU RS06885 WP 002262930.1          VGYYMKSYD   235
NTDB id 1009308 ABNK63 RS04150 WP 274378300.1  .........   136
NTDB id 1047 HON27 RS03110 WP 000389061.1       .........   127
consensus
```

X non conserved
X similar
X ≥ 50% conserved