```
logo
NTDB id 374 SMU RS06885 WP 002262930.1      ......MKKILIVDDEKPISDIIKFNLAKEGYDTITAFDGREALSKYEEENPDLIILDLMLPELDGLEVAKEVR...KNSHVPII   76
NTDB id 79460 SKTS RS01200 WP 173068798.1   ..........MVIDDSNTIRRSAEIFLLKSGCEVILAEDGFDAMAKITDYQPDIIFVDIMMPRLDGYQTTMLIKKNPRFKHTPVI   75
NTDB id 1047 HON27 RS03110 WP 000389061.1   MEDAFQNLKVMVIDDSKTIRRTAETLLQREGCEVITAVDGFEALSKIAEANPDIVFVDIMMPRLDGYQTCALIKNSQNYQNIPVI   85
consensus                                          *****!!***!*****  ! **!***!*! !!**!**!* * *!!****!*!*!*!!!***  ***    ** **!*!

logo
NTDB id 374 SMU RS06885 WP 002262930.1      MLSAKDSEFDKVIGLEIGADDYVTKPFSNRELLARVKAHLRRTENIESAVAEENASGIPEIIIGDLQILPDAFVAKKRGTEVELT   161
NTDB id 79460 SKTS RS01200 WP 173068798.1   MLSSKDGLFDRARGRMVGSDEYLTKPFTKDSLLEAVRKHTTARSGA.....................................   121
NTDB id 1047 HON27 RS03110 WP 000389061.1   MLSSKDGLFDQAKGRVVGSDEYLTKPFSKDELLNAIRNHVSS.........................................   127
consensus                                   !!!*!!**!!***!***!*!*!*!!!!****!! *** !**

logo
NTDB id 374 SMU RS06885 WP 002262930.1      HREFELLHHLATHTGQVMTREHLLETVWGYDYFGDVRTVDVTVRRLREKIEDTPSRPEYILTRRGVGYYMKSYD   235
NTDB id 79460 SKTS RS01200 WP 173068798.1   ......................................................................   121
NTDB id 1047 HON27 RS03110 WP 000389061.1   ......................................................................   127
consensus
```

X non conserved

X similar

X $\geq$ 50% conserved