```
logo
NTDB id 374 SMU RS06885 WP 002262930.1      ...........MKKILIVDDEKPISDIIKFNLAKEGYDTITAFDGREALSKYEEENPDLIILDLMLPELDGLEVAKEVRK...NS    71
NTDB id 70056 ALSL RS07185 WP 269433057.1   MNLNSNANGLAGLKVMVIDDSKTIRRTAETLLRKEGCEVLTAVDGFEALAKIAEHKPAIIFVDIMMPRLDGYQTCALIKNNPQFR    85
NTDB id 1047 HON27 RS03110 WP 000389061.1   .....MEDAFQNLKVMVIDDSKTIRRTAETLLQREGCEVITAVDGFEALSKIAEANPDIVFVDIMMPRLDGYQTCALIKNSQNYQ    80
consensus                                            *    *!****!!*!*!*******! *!!****!!*!!*!!!*!**! *!*****!*!*!*!!!*********  **

logo
NTDB id 374 SMU RS06885 WP 002262930.1      HVPIIMLSAKDSEFDKVIGLEIGADDYVTKPFSNRELLARVKAHLRRTENIESAVAEENASGIPEIIIGDLQILPDAFVAKKRGT   156
NTDB id 70056 ALSL RS07185 WP 269433057.1   GTPVIMLSSKDGLFDKARGRIVGAEQYLTKPFTRDELLGAIQRHVGQPA.................................   134
NTDB id 1047 HON27 RS03110 WP 000389061.1   NIPVIMLSSKDGLFDQAKGRVVGSDEYLTKPFSKDELLNAIRNHVSS.....................................   127
consensus                                    *!*!!!!*!!**!!***!***!***!*!!!!***!!!**** !*

logo
NTDB id 374 SMU RS06885 WP 002262930.1      EVELTHREFELLHHLATHTGQVMTREHLLETVWGYDYFGDVRTVDVTVRRLREKIEDTPSRPEYILTRRGVGYYMKSYD   235
NTDB id 70056 ALSL RS07185 WP 269433057.1   ......................................................................................   134
NTDB id 1047 HON27 RS03110 WP 000389061.1   ......................................................................................   127
consensus
```

X non conserved

X similar

X $\geq$ 50% conserved