```
logo
NTDB id 374 SMU RS06885 WP 002262930.1          ......MKKILIVDDEKPISDIIKFNLAKEGYDTITAFDGREALSKYEEENPDLIILDLMLPELDGLEVAKEVRK...NS    71
NTDB id 1001619 ABFG87 RS01175 WP 136774096.1   .MSNLAGVKVMVIDDSNTIRRSAEIFLGQAGCEVILAEDGFDALAKISDHQPDLIFVDVMMPRLDGYQTCSLIKKNPRYK    79
NTDB id 1047 HON27 RS03110 WP 000389061.1       MEDAFQNLKVMVIDDSKTIRRTAETLLQREGCEVITAVDGFEALSKIAEANPDIVFVDIMMPRLDGYQTCALIKNSQNYQ    80
consensus                                              *!****!!***!*****  !***!***!*! !!**!!*!* * *!!****!*!*!*!!!**** ****   *

logo
NTDB id 374 SMU RS06885 WP 002262930.1          HVPIIMLSAKDSEFDKVIGLEIGADDYVTKPFSNRELLARVKAHLRRTENIESAVAEENASGIPEIIIGDLQILPDAFVA   151
NTDB id 1001619 ABFG87 RS01175 WP 136774096.1   STPVVMLSSKDGLFDRARGRMVGSDEYLTKPFTKDSLLAAVGAHVQSAA...............................   128
NTDB id 1047 HON27 RS03110 WP 000389061.1       NIPVIMLSSKDGLFDQAKGRVVGSDEYLTKPFSKDELLNAIRNHVSS.................................   127
consensus                                        *!**!!!*!!**!!***!***!*!*!*!!!!****!!*****!* *

logo
NTDB id 374 SMU RS06885 WP 002262930.1          KKRGTEVELTHREFELLHHLATHTGQVMTREHLLETVWGYDYFGDVRTVDVTVRRLREKIEDTPSRPEYILTRRGVGYYM   231
NTDB id 1001619 ABFG87 RS01175 WP 136774096.1   ................................................................................   128
NTDB id 1047 HON27 RS03110 WP 000389061.1       ................................................................................   127
consensus

logo
NTDB id 374 SMU RS06885 WP 002262930.1          KSYD   235
NTDB id 1001619 ABFG87 RS01175 WP 136774096.1   ....   128
NTDB id 1047 HON27 RS03110 WP 000389061.1       ....   127
consensus
```

X non conserved

X similar

X $\geq$ 50% conserved