```
logo
NTDB id 374 SMU RS06885 WP 002262930.1        ......MKKILIVDDEKPISDIIKFNLAKEGYDTITAFDGREALSKYEEENPDLIILDLMLPELDGLEVAKEVRK...NS   71
NTDB id 1417 SGL RS04200 WP 014407071.1       ......MNAVLLVEDSSSQREMISGILKDHGWQVTIACDGVEALEKLQNFSPDLVVLDIVMPRMNGYEVCRRIKSDPKTK   74
NTDB id 72497 YC6258 RS06195 WP 044616249.1   .MDNFENLKVMVIDDSKTIRRTAETLLKKVGCEVFTATDGFDALAKIADTKPNIIFVDIMMPRLDGYQTCALIKNNSSFK   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 1417 SGL RS04200 WP 014407071.1       NVPVIMCSSKGEEFDRFWGMRQGADAYIAKPFQPMELVGTIKQLLRN................................  121
NTDB id 72497 YC6258 RS06195 WP 044616249.1   NTPVIMLSSKDGLFDKAKGRIVGSDQYLTKPFGKEELLGAIRQHIKD.................................  126
NTDB id 1047 HON27 RS03110 WP 000389061.1     NIPVIMLSSKDGLFDQAKGRVVGSDEYLTKPFSKDELLNAIRNHVSS.................................  127
consensus                                     **!*!!*!*!** !!***!***! !*!**!!!***!!*********

logo
NTDB id 374 SMU RS06885 WP 002262930.1        KKRGTEVELTHREFELLHHLATHTGQVMTREHLLETVWGYDYFGDVRTVDVTVRRLREKIEDTPSRPEYILTRRGVGYYM  231
NTDB id 1417 SGL RS04200 WP 014407071.1       ......................................................................................  121
NTDB id 72497 YC6258 RS06195 WP 044616249.1   ......................................................................................  126
NTDB id 1047 HON27 RS03110 WP 000389061.1     ......................................................................................  127
consensus

logo
NTDB id 374 SMU RS06885 WP 002262930.1        KSYD   235
NTDB id 1417 SGL RS04200 WP 014407071.1       ....   121
NTDB id 72497 YC6258 RS06195 WP 044616249.1   ....   126
NTDB id 1047 HON27 RS03110 WP 000389061.1     ....   127
consensus
```

X non conserved

X similar

X ≥ 50% conserved