```
logo

NTDB id 374 SMU RS06885 WP 002262930.1        ......MKKILIVDDEKPISDIIKFNLAKEGYDTITAFDGREALSKYEEENPDLIILDLMLPELDGLEVAKEVR...KNS    71
NTDB id 669544 MRY16 RS11915 WP 004640629.1  MEDKFQNLKVMVIDDSKTIRRTAETLLQREGCEVITAVDGFEALSKIAEANPDIVFVDIMMPRLDGYQTCALIKNSLNYQ    80
NTDB id 1047 HON27 RS03110 WP 000389061.1    MEDAFQNLKVMVIDDSKTIRRTAETLLQREGCEVITAVDGFEALSKIAEANPDIVFVDIMMPRLDGYQTCALIKNSQNYQ    80
NTDB id 1417 SGL RS04200 WP 014407071.1      ......MNAVLLVEDSSSQREMISGILKDHGWQVTIACDGVEALEKLQNFSPDLVVLDIVMPRMNGYEVCRRIKSDPKTK    74
consensus                                    *** ** *******!******* ***!***!*****!*!!*!!!*!*****!!****!***!***!*  *******  **

logo

NTDB id 374 SMU RS06885 WP 002262930.1        HVPIIMLSAKDSEFDKVIGLEIGADDYVTKPFSNRELLARVKAHLRRTENIESAVAEENASGIPEIIIGDLQILPDAFVA   151
NTDB id 669544 MRY16 RS11915 WP 004640629.1  NIPVIMLSSKDGLFDQAKGRVVGSDEYLTKPFSKDELLSAIRNHVSA...............................   127
NTDB id 1047 HON27 RS03110 WP 000389061.1    NIPVIMLSSKDGLFDQAKGRVVGSDEYLTKPFSKDELLNAIRNHVSS...............................   127
NTDB id 1417 SGL RS04200 WP 014407071.1      NVPVIMCSSKGEEFDRFWGMRQGADAYIAKPFQPMELVGTIKQLLRN................................   121
consensus                                    **!*!!*!*!** !!***!***! !*!**!!!***!!*******

logo

NTDB id 374 SMU RS06885 WP 002262930.1        KKRGTEVELTHREFELLHHLATHTGQVMTREHLLETVWGYDYFGDVRTVDVTVRRLREKIEDTPSRPEYILTRRGVGYYM   231
NTDB id 669544 MRY16 RS11915 WP 004640629.1  ......................................................................................   127
NTDB id 1047 HON27 RS03110 WP 000389061.1    ......................................................................................   127
NTDB id 1417 SGL RS04200 WP 014407071.1      ......................................................................................   121
consensus

logo

NTDB id 374 SMU RS06885 WP 002262930.1        KSYD   235
NTDB id 669544 MRY16 RS11915 WP 004640629.1  ....   127
NTDB id 1047 HON27 RS03110 WP 000389061.1    ....   127
NTDB id 1417 SGL RS04200 WP 014407071.1      ....   121
consensus
```

X non conserved

X similar

X ≥ 50% conserved