```
logo                                

NTDB id 374 SMU RS06885 WP 002262930.1      .....MKKILIVDDEKPISDIIKFNLAKEGYDTITAFDGREALSKYEEENPDLIILDLMLPELDGLEVAKEVR...KNSHVPII   76
NTDB id 50208 OCUAC RS03295 WP 000389061.1  MEDAFQNLKVMVIDDSKTIRRTAETLLQREGCEVITAVDGFEALSKIAEANPDIVFVDIMMPRLDGYQTCALIKNSQNYQNIPVI   85
NTDB id 1047 HON27 RS03110 WP 000389061.1   MEDAFQNLKVMVIDDSKTIRRTAETLLQREGCEVITAVDGFEALSKIAEANPDIVFVDIMMPRLDGYQTCALIKNSQNYQNIPVI   85
NTDB id 1417 SGL RS04200 WP 014407071.1     .....MNAVLLVEDSSSQREMISGILKDHGWQVTIACDGVEALEKLQNFSPDLVVLDIVMPRMNGYEVCRRIKSDPKTKNVPVI   79
consensus                                   ****** *******!******* ***!***!*****!*!!*!!!*!*****!!****!***!***!*  ******** ****!*!
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logo                                

NTDB id 374 SMU RS06885 WP 002262930.1      MLSAKDSEFDKVIGLEIGADDYVTKPFSNRELLARVKAHLRRTENIESAVAEENASGIPEIIIGDLQILPDAFVAKKRGTEVELT   161
NTDB id 50208 OCUAC RS03295 WP 000389061.1  MLSSKDGLFDQAKGRVVGSDEYLTKPFSKDELLNAIRNHVSS.....................................................   127
NTDB id 1047 HON27 RS03110 WP 000389061.1   MLSSKDGLFDQAKGRVVGSDEYLTKPFSKDELLNAIRNHVSS.....................................................   127
NTDB id 1417 SGL RS04200 WP 014407071.1     MCSSKGEEFDRFWGMRQGADAYIAKPFQPMELVGTIKQLLRN.....................................................   121
consensus                                   !*!*!** !!***!***! !*!**!!!***!!********* *
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logo                                

NTDB id 374 SMU RS06885 WP 002262930.1      HREFELLHHLATHTGQVMTREHLLETVWGYDYFGDVRTVDVTVRRLREKIEDTPSRPEYILTRRGVGYYMKSYD   235
NTDB id 50208 OCUAC RS03295 WP 000389061.1  ..............................................................................   127
NTDB id 1047 HON27 RS03110 WP 000389061.1   ..............................................................................   127
NTDB id 1417 SGL RS04200 WP 014407071.1     ..............................................................................   121
consensus
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X non conserved
X similar
X ≥ 50% conserved