```
logo

NTDB id 98 BSU 24710 NP 390351.1              ..........MNEKGFTLVEMLIVLFIISILLLITIPNVTKHNQTIQKKGCEGLQNMVKAQMTAFELDHEGQTPSLADLQ   70
NTDB id 530 SMSK321 RS09230 WP 000738643.1    MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLLLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKN.EDASLSKLQ   79
NTDB id 431 SGO RS09410 WP 012130940.1        ....MNKLKKLRVKAFTLVEMLVVLLIISVLMLLFVPNLTKQKEAVSDTGNAAVVKVVESQAELYELKNTGDQATLSKLV   76
NTDB id 396 FSA28 RS09395 WP 002263437.1      ....MKKIKRVSVRAFTLIEMLVVLLIISILLLLFVPNLTKQKDKVKDTGGSAVVKVVESQAELYELNKT.DSATLSKLV   75
NTDB id 387 SMU RS09025 WP 002263437.1        ....MKKIKRVSVRAFTLIEMLVVLLIISILLLLFVPNLTKQKDKVKDTGGSAVVKVVESQAELYELNKT.DSATLSKLV   75
NTDB id 280889 C7J89 RS08170 WP 103295107.1   ....MNKIKH..FKAFTLIEMLLVLLIISLLLVLIIPNISKQSKHIQTTGCKAQLKLIDSQIQAYTLKFGQPPNTIDNLV   74
NTDB id 32 MW RS07995 WP 000472256.1          ...MFKFLKK..TQAFTLIEMLLVLLIISLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNPSSIEDLI   75
NTDB id 16 SA RS07765 WP 000472256.1          ...MFKFLKK..TQAFTLIEMLLVLLIISLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNPSSIEDLI   75
consensus                                         *  ***   **!!!*!!!*!!*!!!*!*** *!!**!*   ** *!  ********! * * !       **  !*

logo

NTDB id 98 BSU 24710 NP 390351.1              SEGYVKKDA.VCPNGKRIIITGGEVKVEH   98
NTDB id 530 SMSK321 RS09230 WP 000738643.1    ADGRITEEQAKAYNEYYAKNGGANRKVND  108
NTDB id 431 SGO RS09410 WP 012130940.1        AACNISQKQADSYKAYYGKNNSETQAVAN  105
NTDB id 396 FSA28 RS09395 WP 002263437.1      ASGDITQEQANSYKDYYAKHPSKTRHVAN  104
NTDB id 387 SMU RS09025 WP 002263437.1        ASGDITQEQANSYKDYYAKHPSKTRHVAN  104
NTDB id 280889 C7J89 RS08170 WP 103295107.1   SEGFIKDEQKSCKNGQTIA.INNGEAVAS  102
NTDB id 32 MW RS07995 WP 000472256.1          ADGFIKEAQKTCKSGETIT.ISNGEAVAN  103
NTDB id 16 SA RS07765 WP 000472256.1          ADGFIKEAQKTCKSGETIT.ISNGEAVAN  103
consensus                                     **! *  **      *       *    !**
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X non conserved

X similar

X $\geq$ 50% conserved