```
logo
NTDB id 162 SP RS10370 WP 000738627.1        MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKNE.DASLRKLQADGRI   84
NTDB id 530 SMSK321 RS09230 WP 000738643.1  MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLLLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKNE.DASLSKLQADGRI   84
NTDB id 98 BSU 24710 NP 390351.1             ..........MNEKGFTLVEMLIVLFIISILLLITIPNVTKHNQTIQKKGCEGLQNMVKAQMTAFELDHEGQTPSLADLQSEGYV   75
NTDB id 111315 RZ44 RS03820 WP 231856248.1  MRKTMNRKFKQKEEGFTLIEMLLVLLVISVLIILIIPNIAAQSRNVQDTGCEAQVRMVQSQIEAYTLNEGGRPSSIQDLVPE.YL   84
NTDB id 32 MW RS07995 WP 000472256.1         .....MFKFLKKTQAFTLIEMLLVLLIISLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNPSSIEDLIADGFI   80
NTDB id 16 SA RS07765 WP 000472256.1         .....MFKFLKKTQAFTLIEMLLVLLIISLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNPSSIEDLIADGFI   80
consensus                                          *     *  *!!!*!!!*!!**!!*!*** *!!* **    **  !* * ***! *!**** !  *    !* *! *****

logo
NTDB id 162 SP RS10370 WP 000738627.1        TEEQAKAYKEYN..DKNGGANRKVND   108
NTDB id 530 SMSK321 RS09230 WP 000738643.1  TEEQAKAYNEYY..AKNGGANRKVND   108
NTDB id 98 BSU 24710 NP 390351.1             KKD.AVCPNGKRIIITGGEVKVEH..    98
NTDB id 111315 RZ44 RS03820 WP 231856248.1  TAEQVSCQNGESITITNGEAARE...   107
NTDB id 32 MW RS07995 WP 000472256.1         KEAQKTCKSGETITISNGEAVAN...   103
NTDB id 16 SA RS07765 WP 000472256.1         KEAQKTCKSGETITISNGEAVAN...   103
consensus                                     ***  *  *  * ***!**
```

X non conserved

X similar

X ≥ 50% conserved