```
logo
                                                  MNEKGFTLVEMLIVLFIISILLLITIPNVTKHNQTIQKKGCEGLQNMIKAQMTAYELDHEGQTPSLADLQSEGY
NTDB id 292741 DKG76 RS12965 WP 003236957.1      ......MNEKGFTLVEMLIVLFIISILLLITIPNVTKHNQTIQKKGCEGLQNMIKAQMTAYELDHEGQTPSLADLQSEGY   74
NTDB id 98 BSU 24710 NP 390351.1                 ......MNEKGFTLVEMLIVLFIISILLLITIPNVTKHNQTIQKKGCEGLQNMVKAQMTAFELDHEGQTPSLADLQSEGY   74
NTDB id 460 GPW51 RS00790 WP 011921720.1         MKKLVEMKVKGFTLVEMLVVLGIISLLLLLFVPNLSQQKDAIQKKGDAAVVKVVESQMELYELEHDEEA.TVADLQAKGY   79
NTDB id 32 MW RS07995 WP 000472256.1             MFK.FLKKTQAFTLIEMLLVLLIISLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNPSSIEDLIADGF   79
NTDB id 16 SA RS07765 WP 000472256.1             MFK.FLKKTQAFTLIEMLLVLLIISLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNPSSIEDLIADGF   79
consensus                                        * *   ** **!!!*!!!*!! !!!*!!** *!!****   !!**!* ****** *!*****!*!* *  ***!!***!*

logo
                                                  VKKD.AVCPNGKHIIISGGEVKVEH
NTDB id 292741 DKG76 RS12965 WP 003236957.1      VKKD.AVCPNGKHIIISGGEVKVEH   98
NTDB id 98 BSU 24710 NP 390351.1                 VKKD.AVCPNGKRIIITGGEVKVEH   98
NTDB id 460 GPW51 RS00790 WP 011921720.1         ITEKQAKQYATAKK..........    93
NTDB id 32 MW RS07995 WP 000472256.1             IKEAQKTCKSGETITISNGEAVAN.  103
NTDB id 16 SA RS07765 WP 000472256.1             IKEAQKTCKSGETITISNGEAVAN.  103
consensus                                        *** ** *  *  * ** **
```

X non conserved

X similar

X ≥ 50% conserved