```
logo
NTDB id 1131 NGFG RS05740 WP 003695064.1        ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRK.LAEIAGQYLKKGGLVY    75
NTDB id 1103 NMB RS07590 WP 002212976.1         ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRK.LAEIAGQYLKKGGLVY    75
NTDB id 1390 A4U84 RS00055 WP 021115958.1       ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRR.QAEVAGQYLRKGSKVY    77
NTDB id 1012087 ABMK85 RS19895 WP 011750041.1  .MAGSVNKVILVGNLGQDPEVRSFQNGGKVANLRIATSEQWKDRNTGERRERTEWHTVVIYSEPLVRVAEQFLKKGSKVY    79
NTDB id 1166 A1552VC RS00795 WP 000168289.1    MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGK.LAEVAGEYLRKGSQVY    79
consensus                                          **!!!!**!*!!*!!**!****!**!*****!!!!*!*!***!***!*!!!! ****** ****!***!*!!* !!

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1        LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQSQ.........EAYQRPAQQSRQPAPD   142
NTDB id 1103 NMB RS07590 WP 002212976.1         LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQSQ.........EAYQRPAQQSRQPASD   142
NTDB id 1390 A4U84 RS00055 WP 021115958.1       VEGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFG.GNQGSGWNQAPAQTNYNQGGYSDNYAQNNNFNGG   152
NTDB id 1012087 ABMK85 RS19895 WP 011750041.1  VEGQLETRKWQDQSGNDRYSTEVALRPFRSELHMLDSRGGAGGGG....G..RDE.....GYGGGSYGGGSSGGSS....   144
NTDB id 1166 A1552VC RS00795 WP 000168289.1    IEGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMP.AQGG..MNV.....PAQQGSWGQPQQPAKQHQPM   151
consensus                                      *!!****!!*!** !**!******     ** !!**!  *****    ! *  *         ***  * ** **

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1        APSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1         APSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1       NATRPQ.PAQKP...AAQAEPPMDNFDDDIPF   180
NTDB id 1012087 ABMK85 RS19895 WP 011750041.1  ...SGGGYSSGG...GGGQSQSRPDFDDDIPF   170
NTDB id 1166 A1552VC RS00795 WP 000168289.1    QQSAPQQYS......QPQYNEPPMDFDDDIPF   177
consensus                                        * **     **       *  *****!!!!!!
```

X non conserved

X similar

X $\geq$ 50% conserved