```
logo
NTDB id 1103 NMB RS07590 WP 002212976.1        ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1131 NGFG RS05740 WP 003695064.1       ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1390 A4U84 RS00055 WP 021115958.1      ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
NTDB id 1008744 ABNP46 RS01870 WP 349920742.1 MASRGINKVILIGNLGQDPEVRYMPSGGAVTNITIATSETWRDKQTGEQKERTEWHRVVFMGKLAEVAGEYLKKGSQVYV   80
NTDB id 1166 A1552VC RS00795 WP 000168289.1   MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                         **!!!!**!*!!*!!**!*!!*!*!! ****!!!!*!*!* *!***!*!!!!********!!*!!*!!*!!* !!*

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQ..SQEAYQRPAQQSRQ...PASDAP...   144
NTDB id 1131 NGFG RS05740 WP 003695064.1       EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQ..SQEAYQRPAQQSRQ...PAPDAP...   144
NTDB id 1390 A4U84 RS00055 WP 021115958.1      EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGD.FGGNQG.......SGWNQAPAQTNYNQGGYSDNYAQN   146
NTDB id 1008744 ABNP46 RS01870 WP 349920742.1 EGKLQTRKWQDQSGQERYTTEVLVDSFTGVMQMLGGRPQGGQ......GMGGQQSQGNWGQPQQTPQQNMQRPAAAPQQN   154
NTDB id 1166 A1552VC RS00795 WP 000168289.1   EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQGGMNVPAQQGSWGQPQQPAKQHQPMQQSAP...   156
consensus                                     !!****!!*!** !**!******      ***!!**!  ***  *   ! *   ** ***** *****      ***

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        ........SHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1       ........SHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1      NNFNGGNATRP..Q..PAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 1008744 ABNP46 RS01870 WP 349920742.1 MQQQQGGYGRPQQQ..PQSAPPVYNEPPM.DFDDDIPF   189
NTDB id 1166 A1552VC RS00795 WP 000168289.1   ................QQYSQPQYNEPPM.DFDDDIPF   177
consensus                                             * **        **** ***!****!!!!!!
```

X non conserved

X similar

X $\geq$ 50% conserved