```
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 1054176 BCV36 RS00115 WP 010434324.1  MASRGVNKVILVGNLGNDPEIRYMPNGGAVANITIATSESWRDKATGEQREKTEWHRVALFGKLAEVAGEYLRKGSQVYI   80
NTDB id 1166 A1552VC RS00795 WP 000168289.1   MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                         **!!!!**!*!!*!!**!*!!*!*!!*****!!!!*!*!* *!***!*!!!!********!!*!!*!!*!!* !!*

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEG.........YGQSQEAYQRPAQQS.......   136
NTDB id 1131 NGFG RS05740 WP 003695064.1       EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEG.........YGQSQEAYQRPAQQS.......   136
NTDB id 1390 A4U84 RS00055 WP 021115958.1      EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGG.DFGG.......NQGSGWNQAPAQTNYNQGGYSDNYAQN   146
NTDB id 1054176 BCV36 RS00115 WP 010434324.1  EGQLQTRKWQDQSGQDRYTTEVVVQGFNGVMQMLGGRAQ.G.GAPAQGGMGN.QQQGGWGQPQQPQQ.QQQQY.......   149
NTDB id 1166 A1552VC RS00795 WP 000168289.1   EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQG.GMPAQGGMNVPAQQGSWGQPQQPAK.QHQPM.......   151
consensus                                     !!****!!*!** !**!*******     ***!!**!  ** ****  *     *   **! *   * * **

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        ...RQPASDAPSHPQEAPAAPRRQ.PVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1       ...RQPAPDAPSHPQEAPAAPRRQ.PVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1      NNFNGGNATRP........QPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 1054176 BCV36 RS00115 WP 010434324.1  ...SAPAQQQPKA....PQQAPQQAQPQYNEPPMDFDDDIPF   184
NTDB id 1166 A1552VC RS00795 WP 000168289.1   ...QQ.............SAPQQYSQPQYNEPPMDFDDDIPF   177
consensus                                         ***   *       * ** **   **  !****!!!!!!
```

X non conserved

X similar

X $\geq$ 50% conserved