```
logo
NTDB id 1131 NGFG RS05740 WP 003695064.1        ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL    76
NTDB id 1103 NMB RS07590 WP 002212976.1         ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL    76
NTDB id 1166 A1552VC RS00795 WP 000168289.1     MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI    80
NTDB id 1076369 ACJEHK RS02200 WP 011214630.1  .MARGINKVILVGNVGADPDVRYLPNGNAVTTLSVATSETWKDKTTGEKQDRTEWHRVVCFNRLGEIAGEYIRKGSKLYV    79
NTDB id 1390 A4U84 RS00055 WP 021115958.1       ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV    78
consensus                                          **!!!!**!**! !!**!**!*!*!! ****!!!!*!*!* *!** **!!!!***********!*!!*!**!!* *!*

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1        EGRIQSRKYQGKDGIERTAYDIVAN....EMKMLGGRNENSGGAPYDEGYGQSQE.........AYQRPAQQSRQPAPDA   143
NTDB id 1103 NMB RS07590 WP 002212976.1         EGRIQSRKYQGKDGIERTAYDIVAN....EMKMLGGRNENSGGAPYEEGYGQSQE.........AYQRPAQQSRQPASDA   143
NTDB id 1166 A1552VC RS00795 WP 000168289.1     EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQGG...MNVPAQ....QGSWGQP....QQPA..K   146
NTDB id 1076369 ACJEHK RS02200 WP 011214630.1  EGSLRTRKWQDQQGQDRYTTEIVAS....DIQMLDSKGSSTT.........................NY..DDMPSFQG   127
NTDB id 1390 A4U84 RS00055 WP 021115958.1       EGKLKTRKWQDQNGQDRYTTEIQGD....VLQMLDSRSSGGD.FGGNQGSGWNQAPAQTNYNQGGYSDNY..AQNNNFNG   151
consensus                                       !!*****!!*!** !**!*******    ***!!***  ***  *   * *  *          **  *     ***  *

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1        PSHPQEAPAAPRRQPV.PAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1         PSHPQEAPAAPRRQPV.PAAAPVEDIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1     QHQPMQ.QSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 1076369 ACJEHK RS02200 WP 011214630.1  TSTPQQASTKNQVTPTSTAQDAFDELDDDIPF   159
NTDB id 1390 A4U84 RS00055 WP 021115958.1       GNAT...RPQPAQKPAAQAEPPMDNFDDDIPF   180
consensus                                        * ** *  **   *   *  *****!!!!!!
```

X non conserved

X similar

X $\geq$ 50% conserved