```
logo
NTDB id 1131 NGFG RS05740 WP 003695064.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1103 NMB RS07590 WP 002212976.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
NTDB id 390185 FSC10 RS14105 WP 071320869.1 ..MRGVNKVILVGTLGRDPETKTFPNGGSLTQFSIATSDSWTDKNTGERKEQTEWHRIVLHNRLGEIAQQYLRKGSKVYI   78
NTDB id 1390 A4U84 RS00055 WP 021115958.1   ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
consensus                                      **!!!!**! !!*!!*****!*!*** ****!!!**!*!***!***!*!!!!**********!*!**!!*!!* !!*

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGY................GQS...QEAYQRPA   133
NTDB id 1103 NMB RS07590 WP 002212976.1     EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGY................GQS...QEAYQRPA   133
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQG.GMPAQGGMNVPA.......QQGSWG..........QPQ   142
NTDB id 390185 FSC10 RS14105 WP 071320869.1 EGSLRTRQWTDQSGQERYTTEIRG....EQMQMLDSGRPQG.EQSDNGGF....AQPRFNNNQGGYGNNNAGQQGYNAPQ   149
NTDB id 1390 A4U84 RS00055 WP 021115958.1   EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGG.DFGGNQGSGWNQAPAQTNYNQGGYSDNYAQN.......   146
consensus                                   !!*****!***** !**!*******      **!!***  ** * * * !*           *** ***    * *** *

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1     QQSRQPAPDAPSH...PQEAPAAPRRQP.VPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1     QQSRQPASDAPSH...PQEAPAAPRRQP.VPAAAPVEDIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1 QPAKQHQP..........MQQSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 390185 FSC10 RS14105 WP 071320869.1 QGGFNNAPAGGNGYGNNNPSGFAPKSAPA.PVAAPAADLDDDLPF   193
NTDB id 1390 A4U84 RS00055 WP 021115958.1   ...........NNFNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
consensus                                   *  ** ** *        *    *!*  *  ****!****!!!*!!
```

X non conserved

X similar

X ≥ 50% conserved