```
logo
NTDB id 1103 NMB RS07590 WP 002212976.1      ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1131 NGFG RS05740 WP 003695064.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1166 A1552VC RS00795 WP 000168289.1  MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
NTDB id 310666 BEN71 RS01555 WP 068972970.1 ..MRGVNKVILVGTLGRDPETKTFPNGGSLTQFSIATSESWTDKNTGERKEQTEWHRIVLHNRLGEIAQQYLRKGSKVYI   78
NTDB id 1390 A4U84 RS00055 WP 021115958.1   ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
consensus                                       **!!!!**! !!*!!*****!*!*** ****!!!!*!*!***!***!*!!!!**********!*!**!!*!!* !!*

logo

NTDB id 1103 NMB RS07590 WP 002212976.1      EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQSQEAYQRPAQ..................Q   135
NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQSQEAYQRPAQ..................Q   135
NTDB id 1166 A1552VC RS00795 WP 000168289.1  EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQ.GGMPAQGGM..NVPA.......QQG...........SWGQ   140
NTDB id 310666 BEN71 RS01555 WP 068972970.1 EGSLRTRQWTDQNGQERYTTEIRG....EQMQMLDSNRQQ.GGEGGENG.GYNQPRFNNNNQGNQGGYGNQGGNQGGYGN   152
NTDB id 1390 A4U84 RS00055 WP 021115958.1   EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSG.GDFGGNQGSGWNQAPAQ..TNYNQGGYSD......NYAQ   145
consensus                                   !!*****!***** !**!*******       **!!***  * !* *** ! * ** ***   * ***          ***

logo

NTDB id 1103 NMB RS07590 WP 002212976.1      SRQPASDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1     SRQPAPDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1  PQQPAK..QHQPMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 310666 BEN71 RS01555 WP 068972970.1  NNQGGYGNAGNNVNQGFQSPKPAPAPAAAPADLDDDLPF   191
NTDB id 1390 A4U84 RS00055 WP 021115958.1    NNN....FNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
consensus                                     ****       *   **   *    ********!!!*!!
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X non conserved

X similar

X $\geq$ 50% conserved