```
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 1166 A1552VC RS00795 WP 000168289.1    MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
NTDB id 1051080 ACEWX4 RS13720 WP 045795510.1 ..MRGVNKVILVGTLGRDPETKTFPNGGSLTQFSIATSDAWTDKTTGERKEQTEWHRIVLHNRLGEIAQQYLRKGSKVYI   78
NTDB id 1390 A4U84 RS00055 WP 021115958.1      ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
consensus                                          **!!!!**! !!*!!*****!*!*** ****!!!**!*!* *!***!*!!!!**********!*!**!!*!!* !!*

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQSQE................AYQRPAQQS   136
NTDB id 1131 NGFG RS05740 WP 003695064.1       EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQSQE................AYQRPAQQS   136
NTDB id 1166 A1552VC RS00795 WP 000168289.1    EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQG..GMNVPA............QQGSWGQPQQP.   144
NTDB id 1051080 ACEWX4 RS13720 WP 045795510.1 EGSLRTRQWTDQNGQERYTTEIRG....EQMQMLDSGRQQGE.QGDN...GFNQPRFNNNQGGGYSNNNQGGYAPQAQGG   150
NTDB id 1390 A4U84 RS00055 WP 021115958.1      EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGD.FGGNQGSGWNQAP.......AQTNYNQGGYSDNYAQ.   145
consensus                                      !!*****!***** !**!******       **!!***  ***  * * * ! **             *****  ****

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        RQPASDAPSHPQEAPAAPRRQPVP.AAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1       RQPAPDAPSHPQEAPAAPRRQPVP.AAAPVEDIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1    ......AKQHQPMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 1051080 ACEWX4 RS13720 WP 045795510.1 FNN.NNAGGGYGNQGGYQQPKPAPAATPAPADLDDDLPF   188
NTDB id 1390 A4U84 RS00055 WP 021115958.1      ..N.NNFNGGNA.TRPQPAQKPAAQAEPPMDNFDDDIPF   180
consensus                                       *    *  *     ***   * * *  *****!!!*!!
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X non conserved

X similar

X $\geq$ 50% conserved