```
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 1017170 AB0T70 RS00960 WP 002068963.1  ..MRGVNKVILVGTLGRDPETKTFPNGGSLTQFSIATSEVWTDKNTGERKEQTEWHRIVLHNRLGEIAQQFLRKGSKVYI   78
NTDB id 1390 A4U84 RS00055 WP 021115958.1      ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
consensus                                          **!!!!**! !!*!!*****!*!*** ****!!!!*!*!***!***!*!!!!**********!*!***!*!!* !!*

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQSQEAYQRPAQQSRQ..............   138
NTDB id 1131 NGFG RS05740 WP 003695064.1       EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQSQEAYQRPAQQSRQ..............   138
NTDB id 1166 A1552VC RS00795 WP 000168289.1    EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQGGM.......NVPAQQG.........SWGQPQQ   143
NTDB id 1017170 AB0T70 RS00960 WP 002068963.1  EGSLRTRQWTDQNGQERYTTEIRG....DQMQMLDARQQGEQ..GFAGGNDFNQPRFNAPQQGGGYQNNNQGGGYGQNNG   152
NTDB id 1390 A4U84 RS00055 WP 021115958.1      EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGD.FGGNQGSGWNQAPAQTNYNQGGYSD.....NYAQN..   146
consensus                                      !!*****!***** !**!*******       **!!**!* * *  **  ! *  *  ** *****          ***

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        .........PASDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1       .........PAPDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1    PA........KQHQP...MQQSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 1017170 AB0T70 RS00960 WP 002068963.1  GYGGQGGFGNGSNSPQGGGFAPKAPQQP....ASAPADLDDDLPF   193
NTDB id 1390 A4U84 RS00055 WP 021115958.1      .....NNFNGGN......ATRPQPAQKPAAQAEPPMDNFDDDIPF   180
consensus                                                *   *      *******     **  *****!!!*!!
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X non conserved

X similar

X $\geq$ 50% conserved