logo

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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 1086321 KI216 RS15175 WP 005006829.1 ..MRGVNKVILVGTLGRDPETKTFPNGGSLTQFSIATSDSWTDKSTGERKEQTEWHRIVLHNRLGEIAQQYLRKGSKVYI   78
NTDB id 1390 A4U84 RS00055 WP 021115958.1    ..MAGVNKVITVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
consensus                                       **!!!!**! !!*!!*****!*!*** ****!!!**!*!* *!***!*!!!!**********!*!**!!*!!* !!*
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NTDB id 1103 NMB RS07590 WP 002212976.1      EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQSQE.........AYQRPAQQSR.....   137
NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQSQE.........AYQRPAQQSR.....   137
NTDB id 1166 A1552VC RS00795 WP 000168289.1  EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQGG..MNV.P.....AQQGSWGQPQQPA......   145
NTDB id 1086321 KI216 RS15175 WP 005006829.1 EGSLRTRQWTDQNGQERYTTEIRG....EQMQMLDSSRPQGE.QG....QGFEQPRFNNNNGQQGGYNNNQSGGYGNQGG   149
NTDB id 1390 A4U84 RS00055 WP 021115958.1    EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGD.FGGNQGSGWNQAPAQTN.YNQGGYSDNYAQN......   146
consensus                                    !!*****!***** !**!******       **!!***  ***  *   * *  *        ****** * **
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NTDB id 1103 NMB RS07590 WP 002212976.1      ............QPASDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1     ............QPAPDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1  ............KQHQPMQQ.....SAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 1086321 KI216 RS15175 WP 005006829.1 NSQSGYGNQGGNNQSNGYANNNPSGF.APKAPAAPA.AAPADLDDDLPF   196
NTDB id 1390 A4U84 RS00055 WP 021115958.1    ..............NNFNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
consensus                                              * *       *******   ** ******!!!*!!
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X non conserved
X similar
X ≥ 50% conserved