logo

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NTDB id 1131 NGFG RS05740 WP 003695064.1        ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL    76
NTDB id 1103 NMB RS07590 WP 002212976.1         ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL    76
NTDB id 1390 A4U84 RS00055 WP 021115958.1       ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV    78
NTDB id 1066118 ACFWHB RS19590 WP 139744800.1   MASRGINKVILIGNLGQDPEVRYMPSGGAVTNITLATSDTWRDKQTGEQKERTEWHRVVFMGKLAEVAGEYLKKGSQVYV    80
NTDB id 1166 A1552VC RS00795 WP 000168289.1     MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI    80
consensus                                           **!!!!**!*!!*!!**!*!!*!*!! ****!!!**!*!* *!***!*!!!!********!!*!!*!!*!!* !!*
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NTDB id 1131 NGFG RS05740 WP 003695064.1        EGRIQSRKYQGKDGIERTAYDIVAN....EMKMLGGRNENSGGAPYDEGYG..QSQEAYQRPAQQSRQPAPDAP......   144
NTDB id 1103 NMB RS07590 WP 002212976.1         EGRIQSRKYQGKDGIERTAYDIVAN....EMKMLGGRNENSGGAPYEEGYG..QSQEAYQRPAQQSRQPASDAP......   144
NTDB id 1390 A4U84 RS00055 WP 021115958.1       EGKLKTRKWQDQNGQDRYTTEIQGD....VLQMLDSRSSGGDFGG.NQG.......SGWNQAPAQTNYNQGGYSDN.YAQ   145
NTDB id 1066118 ACFWHB RS19590 WP 139744800.1   EGKLQTRKWQDQGGQERYTTEVLVDSFTGVMQMLGGRPQGGAGQ....GMGGQQSQGNWGQPQQAAPQQAMQRPAAAPQQ   156
NTDB id 1166 A1552VC RS00795 WP 000168289.1     EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPA.QGGMNVPAQQGSWGQPQQPAKQHQPMQQ......   153
consensus                                       !!****!!*!** !**!********    ***!!**!  *****    ! *  *** ***** ***** *   *
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NTDB id 1131 NGFG RS05740 WP 003695064.1        .SHPQE.....APAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1         .SHPQE.....APAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1       NNNFNGGNATRPQPA.QKPAAQAEPPMDNFDDDIPF   180
NTDB id 1066118 ACFWHB RS19590 WP 139744800.1   NMQPQGGYSRPAQQPQSAPPVYNEPPM.DFDDDIPF   191
NTDB id 1166 A1552VC RS00795 WP 000168289.1     ...........SAPQQYSQPQYNEPPM.DFDDDIPF   177
consensus                                          **      *  *  **** ***!****!!!!!!
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X non conserved

X similar

X ≥ 50% conserved