logo

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NTDB id 1103 NMB RS07590 WP 002212976.1      ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1131 NGFG RS05740 WP 003695064.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1390 A4U84 RS00055 WP 021115958.1    ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
NTDB id 301433 CYM74 RS21325 WP 000168277.1  MASRGVNKVILLGRVGQDPEVRYSPSGTAFANLTIATSEQWRDKNTGEQKELTEWHRVAVSGKLAEVVGRYVKKGDQIYF   80
NTDB id 1166 A1552VC RS00795 WP 000168289.1  MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                        **!!!!**!**!*!!**!**!*!*!*******!!!!*!*!***!***!*!!!!*********!!**!*!**!!  *!*
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NTDB id 1103 NMB RS07590 WP 002212976.1      EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQSQ.........EAYQRPAQQSRQPASDA   143
NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQSQ.........EAYQRPAQQSRQPAPDA   143
NTDB id 1390 A4U84 RS00055 WP 021115958.1    EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDF.GGNQGSGWNQAPAQTNYNQGGYSDNYAQ....NNNF   149
NTDB id 301433 CYM74 RS21325 WP 000168277.1  EGMLRTRKWKDQSGQDRYTTEVH.VGINGVMQMLGGIGDSKQQ.AAS.....................RQSQ.........   129
NTDB id 1166 A1552VC RS00795 WP 000168289.1  EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGM.PAQ...GGMNVPAQ....QGSWGQPQQP....AKQH   148
consensus                                    !!*****!!**** !**!*******     ***!!*** ** *  **  * *  *          **  * **    *
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NTDB id 1103 NMB RS07590 WP 002212976.1      PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF.............   174
NTDB id 1131 NGFG RS05740 WP 003695064.1     PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF.............   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1    NGGNATRPQPAQKPAAQAEPPMDNFDDDIPF.............   180
NTDB id 301433 CYM74 RS21325 WP 000168277.1  .K.....PQQQSSPAQHNEPP.MDFDDDIPFAPVTLPFPRHAIHAI   168
NTDB id 1166 A1552VC RS00795 WP 000168289.1  .QPMQQSAPQQYSQPQYNEPP.MDFDDDIPF.............   177
consensus                                        *  *    ***  ***!****!!!!!!
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X non conserved
X similar
X ≥ 50% conserved