logo

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NTDB id 1131 NGFG RS05740 WP 003695064.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRK.LAEIAGQYLKKGGLVY   75
NTDB id 1103 NMB RS07590 WP 002212976.1      ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRK.LAEIAGQYLKKGGLVY   75
NTDB id 1166 A1552VC RS00795 WP 000168289.1  MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGK.LAEVAGEYLRKGSQVY   79
NTDB id 77783 KU6B RS13955 WP 074620477.1    .MAGSVNKVILVGNLGRDPETRTFQNGGKVCNLRIATSENWKDRNTGERRERTEWHSVAIFSEPLARIAEQYLRKGSKVY   79
NTDB id 1390 A4U84 RS00055 WP 021115958.1    ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRR.QAEVAGQYLRKGSKVY   77
consensus                                        **!!!!**!*!!*!!**!****!**!******!!!!*!*!***!***!*!!!!  ****** *!**!**!!*!!* !!
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NTDB id 1131 NGFG RS05740 WP 003695064.1     LEGRIQSRKYQGKDGIERTAYDIVAN....EMKMLGGRNENSGGAPYDE....GYGQ.......SQEAYQRPAQQSRQPA   140
NTDB id 1103 NMB RS07590 WP 002212976.1      LEGRIQSRKYQGKDGIERTAYDIVAN....EMKMLGGRNENSGGAPYEE....GYGQ.......SQEAYQRPAQQSRQPA   140
NTDB id 1166 A1552VC RS00795 WP 000168289.1  IEGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMP.AQGGMN.........VPAQQGSWGQPQQPAKQH.   148
NTDB id 77783 KU6B RS13955 WP 074620477.1    IEGQLETRKWQDQNGQDRYSTEVVLRPYRGELTLLDSRSEGGGGG.FGGGSGGGYGGGGGGY.DDRGGYDDPGY.GGGS.   155
NTDB id 1390 A4U84 RS00055 WP 021115958.1    VEGKLKTRKWQDQNGQDRYTTEIQGD....VLQMLDSRSS..GGD.FGGNQGSGWNQAPAQTNYNQGGYSDNYAQNNNF.   149
consensus                                    *!!****!!*!** !**!*******     ** *!**! ** !** * *    ****         ****  **** **
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NTDB id 1131 NGFG RS05740 WP 003695064.1     PDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1      SDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1  .....QPMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 77783 KU6B RS13955 WP 074620477.1    ...GGGSGGGSG.......PSRAPARDIDDEIPF   179
NTDB id 1390 A4U84 RS00055 WP 021115958.1    ...NGGNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
consensus                                        *  ***   *   ** *!****!!*!!!
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X non conserved
X similar
X ≥ 50% conserved