logo

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NTDB id 1131 NGFG RS05740 WP 003695064.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY   75
NTDB id 1103 NMB RS07590 WP 002212976.1      ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY   75
NTDB id 148820 AM2010 RS08870 WP 047806755.1 .MAGSLNKVMLIGNLGADPEVREFQNGGAVCNLRIATSEQWKDRNTGERQERTEWHSVAIFSEGLINVAKNYLRKGSKVF   79
NTDB id 1166 A1552VC RS00795 WP 000168289.1  MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYG.KLAEVAGEYLRKGSQVY   79
NTDB id 1390 A4U84 RS00055 WP 021115958.1    ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFR.RQAEVAGQYLRKGSKVY   77
consensus                                        **!!!***!*!! !!**!****!*!!******!!!!*!*!***!** !*!!!! ***** *****!**!!*!!* !*
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NTDB id 1131 NGFG RS05740 WP 003695064.1     LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGG.........APYDEGY.......GQSQEAYQRPAQQS   136
NTDB id 1103 NMB RS07590 WP 002212976.1      LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGG.........APYEEGY.......GQSQEAYQRPAQQS   136
NTDB id 148820 AM2010 RS08870 WP 047806755.1 VEGQLQTRKWQDQNGQDRYSTEIVLRGFTGTLTMLDGPPGAGGGGGGQRSGGGNWDRGSSGGGSGGGASGGGWNQGG...   156
NTDB id 1166 A1552VC RS00795 WP 000168289.1  IEGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGGMPAQG...GM...NVPAQ........QGSWGQPQQPA   145
NTDB id 1390 A4U84 RS00055 WP 021115958.1    VEGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGGNQ...GSGWNQAPAQTN....YNQGGYSDNYAQN   146
consensus                                    *!!****!!*!** !**!*******       * !!***  **** *     ** ** *         * ****** ****
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NTDB id 1131 NGFG RS05740 WP 003695064.1     RQPAPDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1      RQPASDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 148820 AM2010 RS08870 WP 047806755.1 .........GAGSSGGSVGGGAASGGSNYDDLDDDIPF   185
NTDB id 1166 A1552VC RS00795 WP 000168289.1  KQHQ......PMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 1390 A4U84 RS00055 WP 021115958.1    NNFN....GGNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
consensus                                    **         *   **   *    ** *****!!!!!!
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X non conserved
X similar
X ≥ 50% conserved