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logo

NTDB id 1103 NMB RS07590 WP 002212976.1      ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY    75
NTDB id 1131 NGFG RS05740 WP 003695064.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY    75
NTDB id 1390 A4U84 RS00055 WP 021115958.1    ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFR.RQAEVAGQYLRKGSKVY    77
NTDB id 200542 BJP26 RS02910 WP 017980474.1 .MAGSVNKVILVGNLGRDPESRSFQNGGKVVNLRIATSESWKDRNSGERKEKTEWHSVAIFNEGLANVAERFLRKGSKVY    79
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYG.KLAEVAGEYLRKGSQVY    79
consensus                                       **!!!!**!*!!*!!**!****!**! ****!!!!*!*!***!***!*!!!! ***** **!**!***!*!!* !!

logo

NTDB id 1103 NMB RS07590 WP 002212976.1      LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYE........EGYGQSQ.......EAYQRPAQQS   136
NTDB id 1131 NGFG RS05740 WP 003695064.1     LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYD........EGYGQSQ.......EAYQRPAQQS   136
NTDB id 1390 A4U84 RS00055 WP 021115958.1    VEGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGGN.......QGSGWNQAPAQTNYNQGGYSDNYAQN   146
NTDB id 200542 BJP26 RS02910 WP 017980474.1 IEGQLQTRKWQDQNGQDKYSTEIVLQGFNSVLTMLDGPNGGQGGG...GGGRSGGGNDWGGGDDFGGGSSGSYGGGSRGG   156
NTDB id 1166 A1552VC RS00795 WP 000168289.1 IEGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGGMNVPAQQGSWGQPQQPAKQHQ.........   149
consensus                                   *!!****!!*!** !*********       ** !!**** * ***            **** *         **      *

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NTDB id 1103 NMB RS07590 WP 002212976.1      RQPASDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1     RQPAPDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1    NNFNGGNATRPQPAQ....KPAAQAEPPMDNFDDDIPF   180
NTDB id 200542 BJP26 RS02910 WP 017980474.1  S...SGGASGGGRGGFGGGQSGGPGGGFADDLDDDVPF   191
NTDB id 1166 A1552VC RS00795 WP 000168289.1  ......PMQQSAPQQYSQPQY....NEPPMDFDDDIPF   177
consensus                                     *    * * ** *  *  *   *********!!!*!!
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X non conserved

X similar

X ≥ 50% conserved