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 112540 K663 RS00740 WP 062112879.1   .MAGSVNKVILVGNLGADPEVRSFQNGGKVCNIRIATSESWKDRNSGERKERTEWHSVAIFSEGLAGVAERYLRKGSKVY   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....NEMKMLGGRNENSGGA....PYDEGY..GQS.......QEAYQRPAQQSRQ   138
NTDB id 1103 NMB RS07590 WP 002212976.1      LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGA....PYEEGY..GQS.......QEAYQRPAQQSRQ   138
NTDB id 112540 K663 RS00740 WP 062112879.1   VEGKLQTRKWQDQSGNDRYTTEVVLQGPSAVLTMLDGAGGGGGGQGGGRSQGGGSDWGASSGGDYDDFGGSS.....GG.   153
NTDB id 1166 A1552VC RS00795 WP 000168289.1  IEGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGM.....PAQGGM...NVPAQ....QGSWGQPQQPAK.   146
NTDB id 1390 A4U84 RS00055 WP 021115958.1    VEGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDF.....GGNQGSGWNQAPAQTNYNQGGYSDNYAQNN.   147
consensus                                    *!!****!!*!** !**!*******      ** !!***  ****      *   !   *** *      ****  * ** *
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NTDB id 1131 NGFG RS05740 WP 003695064.1     PAPDAPSHP.QEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1      PASDAPSHP.QEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 112540 K663 RS00740 WP 062112879.1   ...GFGGGSSGGRPSGGARGGAGGPNFDNDLDDEVPF   187
NTDB id 1166 A1552VC RS00795 WP 000168289.1  ...QHQ..P.MQQSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 1390 A4U84 RS00055 WP 021115958.1    ...NFNGGN.ATRPQPAQKPAAQAEPPMDNFDDDIPF   180
consensus                                          *    ***  *    *  *****!!**!!
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X non conserved

X similar

X ≥ 50% conserved