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 55559 MESAU RS19230 WP 015317710.1   .MAGSVNKVILVGNLGADPEIRRLNSGEPVVNIRIATSESWRDKNSGERKEKTEWHNVVIFNEGIAKVAEQYLKKGMKVY   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....NEMKMLGGRNENSGGAPY....DEGYGQS..Q.....EAYQRPAQQSRQPA   140
NTDB id 1103 NMB RS07590 WP 002212976.1      LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPY....EEGYGQS..Q.....EAYQRPAQQSRQPA   140
NTDB id 55559 MESAU RS19230 WP 015317710.1   VEGQLQTRKWQDQTGADKYTTEVVLQKFRGELQMLDARGQGEGGQVGGSSRGSDFGQSSPNEGFNRGGGNAP.......   151
NTDB id 1166 A1552VC RS00795 WP 000168289.1  IEGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQ...GGM...NVPAQ....QGSWGQPQQPAKQ..   147
NTDB id 1390 A4U84 RS00055 WP 021115958.1    VEGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGGN...QGSGWNQAPAQTNYNQGGYSDNYAQNNN..   148
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 55559 MESAU RS19230 WP 015317710.1   .........RG.......GGGGGSSRELDDEIPF   169
NTDB id 1166 A1552VC RS00795 WP 000168289.1  ..HQ..PMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 1390 A4U84 RS00055 WP 021115958.1    ..FNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
consensus                                          *   **   *    *********!!*!!!
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X non conserved
X similar
X ≥ 50% conserved