logo

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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 1166 A1552VC RS00795 WP 000168289.1 .MSGSVNKVILVGNLGADPEVRSLPNGNPVVNLRIATSERWRDGGGGEPREKTEWHRVVIFSEGLAKVAQNYLKKGAKVY   79
NTDB id 42680 KKY RS07055 WP 014130627.1    MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYG.KLAEVAGEYLRKGSQVY   79
consensus                                       **!!!!**!*!! !!**!**!*!**! ****!!!!*!*!* !***!*!!!!*****  **!**!**!!*!!* !!
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NTDB id 1103 NMB RS07590 WP 002212976.1     LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQSQE......AYQRPAQQSRQPASDAPS   145
NTDB id 1131 NGFG RS05740 WP 003695064.1    LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQSQE......AYQRPAQQSRQPAPDAPS   145
NTDB id 1390 A4U84 RS00055 WP 021115958.1   VEGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGG.DFGGNQGSGWNQAPAQTNYNQGGYSDNYAQNN.NFNG   151
NTDB id 42680 KKY RS07055 WP 014130627.1    IEGQLQTRKWEK.DGQDQYTTEIVLQGFNSTLTMLDGRGEGG.GDRDSGGYG...GGSRQVENRGRP.............   141
NTDB id 1166 A1552VC RS00795 WP 000168289.1 IEGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQG.GMPAQGGMN...VPAQQG.SWGQPQQPAKQH......   148
consensus                                   *!!*****!!**  *!**********       * !!**! *** * *   !**  *        **** ** **
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NTDB id 1103 NMB RS07590 WP 002212976.1     HPQEAPAAPRRQP.VPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1    HPQEAPAAPRRQP.VPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1   GNATRP.QPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 42680 KKY RS07055 WP 014130627.1    ..........SSAPA..FEPGGMDDDIPF   158
NTDB id 1166 A1552VC RS00795 WP 000168289.1 QPMQQS.APQQYSQPQYNEPPMDFDDDIPF   177
consensus                                    *   * **   *   *  *****!!!!!!
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X non conserved
X similar
X ≥ 50% conserved