logo

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NTDB id 1131 NGFG RS05740 WP 003695064.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1103 NMB RS07590 WP 002212976.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1390 A4U84 RS00055 WP 021115958.1    ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
NTDB id 70069 ALSL RS11495 WP 126539301.1    .MARGVNKVILIGNLGADPEVRYTGSGTAITSLRIATSEQWTDKQSGERQERTEWHRVKLFGKLAEIAGEYLKKGRQVYI   79
NTDB id 1166 A1552VC RS00795 WP 000168289.1  MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                        **!!!!**!*!! !!**!****!*!* ****!!!!*!*!* *!** !*!!!!*********!!*!!*!!*!!  !!*
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NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGKDGIERTAYDIVANE....MKMLGGRNENSGGAPYDE.GYGQSQ..EAYQ.............RPAQQSRQ   138
NTDB id 1103 NMB RS07590 WP 002212976.1     EGRIQSRKYQGKDGIERTAYDIVANE....MKMLGGRNENSGGAPYEE.GYGQSQ..EAYQ.............RPAQQSRQ   138
NTDB id 1390 A4U84 RS00055 WP 021115958.1    EGKLKTRKWQDQNGQDRYTTEIQGDV....LQMLDSRSSGGDFGGNQGSGWNQAPAQTNYNQGGYSDNYAQNNNFNGGNA   154
NTDB id 70069 ALSL RS11495 WP 126539301.1    EGSLRTDKYTDKDGIERYSTDIVASD....MQMLGGMGGEGGGQGGQG.GFR.ERAPA....GPR....G..PAARGGEE   143
NTDB id 1166 A1552VC RS00795 WP 000168289.1  EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQG.GMPAQG.GMN.VPAQQ....GSW....G..QPQQPAKQ   147
consensus                                    !!*****!*****!**!********      **!!***  ****** ** !* *  *   ** *       *   *** ***
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logo

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NTDB id 1131 NGFG RS05740 WP 003695064.1     PAPDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1     PASDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1    TRPQ....PAQKP......AAQAEPPMDNFDDDIPF   180
NTDB id 70069 ALSL RS11495 WP 126539301.1    RAPR....P.SAPA.........PESTGFDDDDIPF   165
NTDB id 1166 A1552VC RS00795 WP 000168289.1  HQPM....QQSAPQ..QYSQPQYNEPPMDFDDDIPF   177
consensus                                     **      **  *!*         *   **** !!!!!!
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X non conserved

X similar

X ≥ 50% conserved