logo

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NTDB id 1103 NMB RS07590 WP 002212976.1      ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL    76
NTDB id 1131 NGFG RS05740 WP 003695064.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL    76
NTDB id 1390 A4U84 RS00055 WP 021115958.1    ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV    78
NTDB id 213670 LM70 RS18020 WP 102030570.1   MASRGVNKVIILGRVGQDPEVRYSPSGTAFANLTVATSERWRDKQTGEQKEQTEWHRVAVVGKLAEVVGQYVKKGDQVYF    80
NTDB id 1166 A1552VC RS00795 WP 000168289.1  MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI    80
consensus                                        **!!!!**!**!*!!**!**!*!*!******!!!!*!*!* *!***!*!!!!********!!**!*!**!!  !!*
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NTDB id 1103 NMB RS07590 WP 002212976.1      EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQSQE.........AYQRPAQQSRQPASDA   143
NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQSQE.........AYQRPAQQSRQPAPDA   143
NTDB id 1390 A4U84 RS00055 WP 021115958.1    EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGD.FGGNQGSGWNQAPAQTNYNQGGYSDNYAQNN....NF   149
NTDB id 213670 LM70 RS18020 WP 102030570.1   EGMLRTRKWQDQTGQDRYTTEIN.VGINGVMQMLGGTGDSKQ.QAADR..................QS...........   128
NTDB id 1166 A1552VC RS00795 WP 000168289.1  EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQGGM..N.VPAQ....QGSWGQPQQPAK....QH   148
consensus                                    !!*****!!*!** !**!*******    ***!!*** ** *  *** * *  *          **  * ** *
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logo

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NTDB id 1103 NMB RS07590 WP 002212976.1      PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF...............   174
NTDB id 1131 NGFG RS05740 WP 003695064.1     PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF...............   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1    NGGNATRPQPAQKPAAQAEPPMDNFDDDIPF...............   180
NTDB id 213670 LM70 RS18020 WP 102030570.1   .....QKPQQQPSPTQHNEPP.MDFDDDIPFAPVTLPFPRHAIHAI   168
NTDB id 1166 A1552VC RS00795 WP 000168289.1  .QPMQQSAPQQYSQPQYNEPP.MDFDDDIPF...............   177
consensus                                        *  *    ***  ***!****!!!!!!
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X non conserved

X similar

X ≥ 50% conserved