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 66855 PBDP RS26620 WP 024778338.1       .MARGVNKVILVGTCGQDPEVRYLPNGNAVTNLSLATSEQWTDKQTGQKVEKTEWHRVSMFGKVAEIAGEYLRKGSQVYI   79
NTDB id 1166 A1552VC RS00795 WP 000168289.1     MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                           **!!!!**! *!*!!**!**!*!*!! ****!!!!*!*!* *!***!*!!!!********!!*!!*!!*!!* !!*
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NTDB id 1103 NMB RS07590 WP 002212976.1         EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEE..GYGQSQEAYQRPAQQSRQPAS.........   141
NTDB id 1131 NGFG RS05740 WP 003695064.1        EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDE..GYGQSQEAYQRPAQQSRQPAP.........   141
NTDB id 1390 A4U84 RS00055 WP 021115958.1       EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGD.FGGNQGSG.......WNQAPAQTNYNQGGYSDNYAQN   146
NTDB id 66855 PBDP RS26620 WP 024778338.1       EGKLQTREWEK.DGIKRYTTEIVV.DMQGTMQLLGGRPQGDQ...QQGGNNYQQSAP.RQQAPRPQSAP..........   142
NTDB id 1166 A1552VC RS00795 WP 000168289.1     EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQGGMNVPAQQGSWGQPQQPAKQH...........   148
consensus                                       !!****!***  *!**!*******      ***!**!  * *  * * * ** ***  **** ******
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NTDB id 1103 NMB RS07590 WP 002212976.1         .DAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1        .DAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1       NNFNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 66855 PBDP RS26620 WP 024778338.1       .....QPQRERPAPQQAAPQPAPDFDSFDDDIPF   171
NTDB id 1166 A1552VC RS00795 WP 000168289.1     .....QPMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
consensus                                             **  ******   *** *****!!!!!!
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X non conserved
X similar
X ≥50% conserved