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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 300564 LFZ12 RS03820 WP 058111616.1 MASKGVNKVILVGNLGQDPEVRYMPNGGAVVNLSLATSDTWTDKQTGDKKERTEWHRVVLYGKLAEIASEYLRKGSQVYI   80
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                  **!!!!**!*!!*!!**!*!!*!*!! ****!!!**!*!* *!***!*!!!!********!!*!**!!*!!* !!*
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NTDB id 1131 NGFG RS05740 WP 003695064.1   EGRIQSRKYQGKDGIERTAYDIV....ANEMKMLGGRNENSGGAPYDEGY..GQSQEAYQRPAQQSR.........QPAPD   142
NTDB id 1103 NMB RS07590 WP 002212976.1    EGRIQSRKYQGKDGIERTAYDIV....ANEMKMLGGRNENSGGAPYEEGY..GQSQEAYQRPAQQSR.........QPASD   142
NTDB id 1390 A4U84 RS00055 WP 021115958.1  EGKLKTRKWQDQNGQDRYTTEIQ....GDVLQMLDSRSSGGDF.GG.......NQGSGWNQAPAQTNYNQGGYSDNYAQN   146
NTDB id 300564 LFZ12 RS03820 WP 058111616.1 EGALRTRKWTDQSGVEKYTTEVVVS.QSGTMHMLGGRNSAGS..........GQQQGGWGQPQQPAA..............   136
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGM.PAQGGMNVPAQQGSWGQPQQPAK..............   146
consensus                                  !!*****!!**** !********** * * !!**!* *** * * **** ***** **** * *
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NTDB id 1131 NGFG RS05740 WP 003695064.1   APSHP..QEAPAAPRRQPVPAAAPVEDIDDDIPF.................   174
NTDB id 1103 NMB RS07590 WP 002212976.1    APSHP..QEAPAAPRRQPVPAAAPVEDIDDDIPF.................   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1  NNFNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF.................   180
NTDB id 300564 LFZ12 RS03820 WP 058111616.1 .PSHSG...TP....PQQHPANEPPMDFDDDIPFAPFGHSVARHALYVLS   178
NTDB id 1166 A1552VC RS00795 WP 000168289.1 .QHQPMQQSAPQ..QYSQPQYNEPPMDFDDDIPF.................   177
consensus                                   **** * *!    **  **  !****!!!!!!
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X non conserved
X similar
X ≥ 50% conserved