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logo

NTDB id 1103 NMB RS07590 WP 002212976.1       ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1131 NGFG RS05740 WP 003695064.1      ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1390 A4U84 RS00055 WP 021115958.1     ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
NTDB id 284228 Tharo RS13650 WP 107221676.1  ..MASVNKVILIGNLGADPENRFAPSGDAICNLRIATAETWRDKNTGERREATEWHRVVFFGKLAEIAGQYLKKGSQVYI   78
NTDB id 1166 A1552VC RS00795 WP 000168289.1  MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                        **!!!!**!*!! !!**!**!*!*!*******!!*!*!*!***!***!*!!!!********!!*!!*!!*!!* !!*
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logo

NTDB id 1103 NMB RS07590 WP 002212976.1       EGRIQSRKYQGKDGIERTAYDIVANEM....KMLGGRNENSGGAPYEEGYGQ..SQEAYQRPAQQSRQ..........PA   140
NTDB id 1131 NGFG RS05740 WP 003695064.1      EGRIQSRKYQGKDGIERTAYDIVANEM....KMLGGRNENSGGAPYDEGYGQ..SQEAYQRPAQQSRQ..........PA   140
NTDB id 1390 A4U84 RS00055 WP 021115958.1     EGKLKTRKWQDQNGQDRYTTEIQGDVL....QMLDSRSSGG.DFGGNQG.......SGWNQAPAQTNYNQGGYSDNYAQN   146
NTDB id 284228 Tharo RS13650 WP 107221676.1  EGSLRTRKWQDQSGQERYTTEIRGDEM....KMLGR.REGG.DAPMRGG....AGEGGWDAPSPAA..............   134
NTDB id 1166 A1552VC RS00795 WP 000168289.1  EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQG.GMPAQGGMNVPAQQGSWGQPQQPA..............   145
consensus                                    !!*****!!*!** !**!**********     *!!*** *** ***   !       * *** * ***
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logo

NTDB id 1103 NMB RS07590 WP 002212976.1       SDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1      PDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1     NNFNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 284228 Tharo RS13650 WP 107221676.1  ARAPASAPQRPAAAP.SAAPQAGGFGDFDDDIPF   167
NTDB id 1166 A1552VC RS00795 WP 000168289.1  KQHQPMQQSAPQ.QY.SQPQYNEPPMDFDDDIPF   177
consensus                                      **    * *!**  *    *********!!!!!!
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X non conserved

X similar

X $\geq$ 50% conserved