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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 970526 V1467 RS02245 WP 416307175.1 MASRGINKVILVGNLGQDPEVRYTPNGKAVANLTLATSESWKDQS.GQIQEKTEWHRVSMFGKLAEIAGEYLKKGSQVYI   79
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                     **!!!!**!*!!*!!**!**!*!*!!*****!!!!*!*!*  !** !*!!!!********!!*!!*!!*!!* !!*

logo

NTDB id 1103 NMB RS07590 WP 002212976.1      EGRIQSRKYQGKDGIERTAYDIVAN....EMKMLGGRNENSGGAPYEEGYGQSQEA......YQRPAQQS.RQP....AS   141
NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGKDGIERTAYDIVAN....EMKMLGGRNENSGGAPYDEGYGQSQEA......YQRPAQQS.RQP....AP   141
NTDB id 1390 A4U84 RS00055 WP 021115958.1    EGKLKTRKWQDQNGQDRYTTEIQGD....VLQMLDSRSSGGDFG.GNQGSGWNQAPAQTNYNQGGYSDNYAQNNNFNGGN   153
NTDB id 970526 V1467 RS02245 WP 416307175.1 EGKLQTRKWQDQQGQDRYTTEIVLDPFNGVMQMLGGGSGRSQSE....G.GGYQQNQQRQQPQQNPQAQYQQYPNQGGGQ   154
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMP....AQGGMNVPAQ.QGSWGQP.....QQP....AK   146
consensus                                   !!*****!!*!** !**!*******     ***!!***  *** *     * !  *   * *  ** *  *  ***    *

logo

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    ATRPQPAQKPAA..QAEPPM....DNFDDDIPF   180
NTDB id 970526 V1467 RS02245 WP 416307175.1 AQQSRPQNPPAQ..QNTPPMAEPDFDFDDDIPF   185
NTDB id 1166 A1552VC RS00795 WP 000168289.1 QHQPMQQSAPQQ..YSQPQYNEPPMDFDDDIPF   177
consensus                                     * ** *!**   * ** * * ***!!!!!!
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X non conserved

X similar

X ≥ 50% conserved