logo

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NTDB id 113 BSU 40900 NP 391970.1           ....MLNRVVLVGRLTKDPELRYTPNGAAVATFTLAVNRTFTNQ.SGER..EADFINCVTWRRQAENVANFLKKGSLAGV    73
NTDB id 1103 NMB RS07590 WP 002212976.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL    76
NTDB id 1131 NGFG RS05740 WP 003695064.1    ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL    76
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI    80
NTDB id 136200 RO07 RS02995 WP 039407932.1  ..MASVNKVILVGNLGADPEVRYMPSGDAVANIRLATTDRYKDKQSGEFKEMTEWHRVSFFGRLAEIVNEYLKKGSSVYI    78
NTDB id 1390 A4U84 RS00055 WP 021115958.1   ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV    78
consensus                                        *!*!***! !* !!**!**!*!*!!** **!***** ** *!** * **** ** ****!!*****!*!!* ***
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NTDB id 113 BSU 40900 NP 391970.1           DGRLQTRNYENQQGQRVFVTEVQA....ESVQFLEPKNGGGSG...SGGYNEGNSGGGQYFGGGQNDNPFGGNQNNQ...   143
NTDB id 1103 NMB RS07590 WP 002212976.1     EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQS.........QEAYQRPAQQSRQPASDA   143
NTDB id 1131 NGFG RS05740 WP 003695064.1    EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQS.........QEAYQRPAQQSRQPAPDA   143
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQGGMNVP.......AQQGSWGQPQQPAKQ....H   148
NTDB id 136200 RO07 RS02995 WP 039407932.1  EGRIRTRKYQAQDGTDRYSTEIVA....DQMQMLGGRSGGGE.GGGGGGYS..RGGGD.EGGGGGYSR....GGG....G   142
NTDB id 1390 A4U84 RS00055 WP 021115958.1   EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGD.FGGNQGSGWNQAPAQTNYNQGGYSDNYAQNNN....F   149
consensus                                   *!****!*** * ! *** *****      ***!***   *       !*            ****  *     *
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logo

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NTDB id 113 BSU 40900 NP 391970.1           ..RRNQGNSFNDDPFANDGKPIDISDDDLPF    172
NTDB id 1103 NMB RS07590 WP 002212976.1     PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF    174
NTDB id 1131 NGFG RS05740 WP 003695064.1    PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF    174
NTDB id 1166 A1552VC RS00795 WP 000168289.1 ..QPMQQSAPQQYSQPQYNEPPMDFDDDIPF    177
NTDB id 136200 RO07 RS02995 WP 039407932.1  GGGGGGR....QQAPAKQSGGFDDMDDDIPF    169
NTDB id 1390 A4U84 RS00055 WP 021115958.1   NGGNATRPQPAQKPAAQAEPPMDNFDDDIPF    180
consensus                                                       **** !!!*!!
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X non conserved

X similar

X ≥ 50% conserved