```
logo
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 1390 A4U84 RS00055 WP 021115958.1  ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
NTDB id 63900 X636 RS02875 WP 023873012.1  ..MASVNKVILVGNLGADPEVRYMPSGDAVANIRLATTDRYKDKQSGEFKEMTEWHRVSFFGRLAEIVNEYLKKGSSVYI   78
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                     *!*!***! !* !!**!**!*!*!!** **!***** ** *!** * **** ** ****!!*****!*!!* ***

logo
NTDB id 113 BSU 40900 NP 391970.1          DGRLQTRNYENQQGQRVFVTEVQAE....SVQFLEPKNGGGSG..SGGYNEGNSG........GGQYFGG.GQNDNPFGG  138
NTDB id 1103 NMB RS07590 WP 002212976.1    EGRIQSRKYQGKDGIERTAYDIVAN....EMKMLGGRNENSGG..........APYEEGYGQSQEAYQRPAQQSRQPASD  142
NTDB id 1131 NGFG RS05740 WP 003695064.1   EGRIQSRKYQGKDGIERTAYDIVAN....EMKMLGGRNENSGG..........APYDEGYGQSQEAYQRPAQQSRQPAPD  142
NTDB id 1390 A4U84 RS00055 WP 021115958.1  EGKLKTRKWQDQNGQDRYTTEIQGD....VLQMLDSRSSGGDFGGNQGSGWNQAPAQTNYN..QGGYSDNYAQNNNFNGG  152
NTDB id 63900 X636 RS02875 WP 023873012.1  EGRIRTRKYQAQDGTDRYSTEIVAD....QMQMLGGRSGGGEGGGGGGYSRGGGD.EGGGG..GGGYSRGGGG...GGGG  148
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGGMNV..................PAQQ...GSWG  139
consensus                                  *!****!*** * ! *** *****      ***!***   * *    *     *            **   * *  *   *

logo
NTDB id 113 BSU 40900 NP 391970.1          NQNNQRRN.....QGNSFNDDPFANDGKPIDISDDDLPF   172
NTDB id 1103 NMB RS07590 WP 002212976.1    A......PSHPQEAPAAPRRQ.PVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1   A......PSHPQEAPAAPRRQ.PVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1  NATRP...........QPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 63900 X636 RS02875 WP 023873012.1  RQQAPAKP..................SGGFDDMDDDIPF   169
NTDB id 1166 A1552VC RS00795 WP 000168289.1 QPQQPAKQHQPMQ.QSAPQQYSQPQYNEPPMDFDDDIPF   177
consensus                                                *          **** !!!*!!
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X non conserved

X similar

X ≥ 50% conserved