```
logo
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 1140134 DQO23 RS30260 WP 026382027.1  ..MASVNKVILVGNLGRDPEVRYSPDGAAICNMSIATTSTWKDKASGERREETEWHRVVMYNRLAEIAGEYLKKGRSVYI   78
NTDB id 1166 A1552VC RS00795 WP 000168289.1  MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                        **!!!!**!*!!*!!**!**!*!*!******!!***!*!* *!***!*!!!!********!!*!!*!!*!!  !!*

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1      EGRIQSRKYQGK.DGIERTAYDIVAN....EMKMLGGRNENSGGAPYDEGYGQSQ.........EAYQ.RPAQ.QSRQPA  140
NTDB id 1103 NMB RS07590 WP 002212976.1       EGRIQSRKYQGK.DGIERTAYDIVAN....EMKMLGGRNENSGGAPYEEGYGQSQ.........EAYQ.RPAQ.QSRQPA  140
NTDB id 1390 A4U84 RS00055 WP 021115958.1     EGKLKTRKWQDQ.NGQDRYTTEIQGD....VLQMLDSRSSGGDFGG.NQGSGWNQAPAQTNYNQGGYSDNYAQNNNFNGG  152
NTDB id 1140134 DQO23 RS30260 WP 026382027.1  EGRLKTRKWQDKDTGADRYSTEIVAD....QMQMLGGREDGGGGGG.SYGGGG.......GY.....DDAPAR.QPQQRA  140
NTDB id 1166 A1552VC RS00795 WP 000168289.1  EGQLQTRKWQDQ.SGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPA.QGGMNV...PAQQGS.....WGQPQQ.PAKQHQ  149
consensus                                    !!****!!*!**  ! *!********     **!!**! *******   ! *  *          **   *** * ** *

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1      PDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1       SDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1     NATRPQPAQKP.AAQA.....EPPMDNFDDDIPF   180
NTDB id 1140134 DQO23 RS30260 WP 026382027.1  PAQRPAPQQRP.APQAAPAGGGANLADMDDDIPF   173
NTDB id 1166 A1552VC RS00795 WP 000168289.1  PMQQSAPQQYS.QPQY.....NEPPMDFDDDIPF   177
consensus                                    *   * !** * ***  *  ********!!!!!!
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X non conserved

X similar

X ≥ 50% conserved