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logo

NTDB id 1017 ACIAD RS02540 WP 011182074.1       MSKWTINKTLGCAAIVGAFILSGCQTV....PTTKDPEKGVKVRTQLAAEYIKSGDLDSAKRALDQALKVNPKDSSANMM   76
NTDB id 1043280 ACEN3N RS18515 WP 373350541.1  ..MYQKAVVILAIGL.SALVVQGCQTTNSDIIIKKDPEKAVKVRTQLAAEYIRSGDLDAAKRALDQALETDSRDSAANMM   77
NTDB id 1051 ABD1 RS02465 WP 004738086.1       .MSTPKLKIMLCMGVAVALLASGCQTS...QTVKKDPEKAVKVRTQLAAEYIRSGDLDSAKRSLDQALSVDSRDATANMM   76
consensus                                         * *  *******  !****!!!!*     **!!!!!*!!!!!!!!!!!!*!!!!!*!!!*!!!!! ****!**!!!!

logo

NTDB id 1017 ACIAD RS02540 WP 011182074.1       MGVLLQQEGSRPNLEKADAYFERAVQLDPENAQAHNNYGTYLYQMKRYHDAIQQLTLAGSSLGYDQRYRALENIGRIYLH   156
NTDB id 1043280 ACEN3N RS18515 WP 373350541.1  MGVLLQQEGSKISMEKADGYFKRAISSDSKNAQARNNYGTYLFVSERYNDAIEQLSIAGASLGYDQRYKALENLGRVYLK   157
NTDB id 1051 ABD1 RS02465 WP 004738086.1       MGILLQQEGSKPNLEKAEHYFKRAISSEPDNAQARNNYGTYLYQMERYNDAIEQFRIAGATLGYDQRYQALENLGRIYLK   156
consensus                                      !!*!!!!!!!****!!!*!!*!!*****!!!!*!!!!!!!****!!*!!!*!***!!**!!!!!!!*!!!!*!!*!!*

logo

NTDB id 1017 ACIAD RS02540 WP 011182074.1       LGGVANAEKMFSQALQANRDSSVSMLELAEIYYLQQRIPAATGMYEQYVRTVGQKNQGARALWIGIRVARANADRTGVQV   236
NTDB id 1043280 ACEN3N RS18515 WP 373350541.1  VGDTANAEKAFKNALQVNRDSIISLVELSEIFYLQQKNAQASQLYEQYVRLVGQDKQGARALWIGLRIARANNDQMGMQV   237
NTDB id 1051 ABD1 RS02465 WP 004738086.1       LGDIASAEKTFKQALLANRDSYISMLELAEIFYLQQQIPAATQMYEQYVRTVGQKNQGARALWIGLRVARANADKMGMQV   236
consensus                                      *!**!*!!! !**!!**!!!! *!**!!*!!*!!!!*****!***!!!!!!*!!!**!!!!!!!!!*!*!!!!*!**!*!!

logo

NTDB id 1017 ACIAD RS02540 WP 011182074.1       LVNQLRIFYPDSSEYQRYLKLQYSTEAVWK   266
NTDB id 1043280 ACEN3N RS18515 WP 373350541.1  LVNQLRALFPDSKEYQRYLQLQYSTEAVWK   267
NTDB id 1051 ABD1 RS02465 WP 004738086.1       LVNQLRALFPESPEYQRYLQLQYSTEAVWK   266
consensus                                      !!!!!!***!*! !!!!!!*!!!!!!!!!!!
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X non conserved

X similar

X ≥ 50% conserved