```
logo
NTDB id 1017 ACIAD RS02540 WP 011182074.1     MSKWTINKTLGCAAIVGAFILSGCQTVPT.TKDPEKGVKVRTQLAAEYIKSGDLDSAKRALDQALKVNPKDSSANMMMGV   79
NTDB id 181513 AsACE RS02615 WP 076753067.1   MRN.LTSKIAFISTVCVSIALSGCQTAEVGKKDPKKATQVRTQLAVEYLKTGDLDSAKRALDEALKINSRDAQANMMMGV   79
NTDB id 1051 ABD1 RS02465 WP 004738086.1      MST.PKLKIMLCMGVAVALLASGCQTSQTVKKDPEKAVKVRTQLAAEYIRSGDLDSAKRSLDQALSVDSRDATANMMMGI   79
consensus                                     !*     !** * ** ******!!!!!  * *!!!*!***!!!!!!*!!***!!!!!!!!*!!*!!*****!**!!!!!!*

logo
NTDB id 1017 ACIAD RS02540 WP 011182074.1     LLQQEGSRPNLEKADAYFERAVQLDPENAQAHNNYGTYLYQMKRYHDAIQQLTLAGSSLGYDQRYRALENIGRIYLHLGG   159
NTDB id 181513 AsACE RS02615 WP 076753067.1   LLQQEGSRLNLDRAETYFKRAISAEPKNAQARNNYGTYLYQLERYNDAIEQFAIAGATLGYDQRYLALENMGRIYLKQGD   159
NTDB id 1051 ABD1 RS02465 WP 004738086.1      LLQQEGSKPNLEKAEHYFKRAISSEPDNAQARNNYGTYLYQMERYNDAIEQFRIAGATLGYDQRYQALENLGRIYLKLGD   159
consensus                                     !!!!!!!**!!**!* !!*!!** *!*!!!!*!!!!!!!!!**!!*!!!*!* *!!**!!!!!!! !!!!*!!!!!**!*

logo
NTDB id 1017 ACIAD RS02540 WP 011182074.1     VANAEKMFSQALQANRDSSVSMLELAEIYYLQQRIPAATGMYEQYVRTVGQKNQGARALWIGIRVARANADRTGVQVLVN   239
NTDB id 181513 AsACE RS02615 WP 076753067.1   LVNAEKSFKQALQANRDSYISMLELAEIFYLNQQFPAASNMYQQFVRSVGQKNQGARALWIGIRTARAEGDQMGMQVLVN   239
NTDB id 1051 ABD1 RS02465 WP 004738086.1      IASAEKTFKQALLANRDSYISMLELAEIFYLQQQIPAATQMYEQYVRTVGQKNQGARALWIGLRVARANADKMGMQVLVN   239
consensus                                     ***!!!*!*!!!*!!!!!**!!!!!!!!!*!!*!**!!!**!!*!*!!*!!!!!!!!!!!!!!!!*!*!!!**!**!*!!!!!

logo
NTDB id 1017 ACIAD RS02540 WP 011182074.1     QLRIFYPDSSEYQRYLKLQYSTEAVWK   266
NTDB id 181513 AsACE RS02615 WP 076753067.1   QLRALFPESPEYQRYLQLQYTTEAVWK   266
NTDB id 1051 ABD1 RS02465 WP 004738086.1      QLRALFPESPEYQRYLQLQYSTEAVWK   266
consensus                                     !!!***!*!*!!!!!!*!!!*!!!!!!
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X non conserved

X similar

X ≥ 50% conserved