```
logo
NTDB id 238190 CFH90 RS03055 WP 126035299.1  .MKKPVSKLYLALTLGVSAWYLVGCQTVNT.KGDPSKAVQVRTQLAAEYIKSNDYDAAKRTLDQALDIDSRDASANMMMG   78
NTDB id 1051 ABD1 RS02465 WP 004738086.1     MS.TPKLKIML.CMGVAVALLASGCQTSQTVKKDPEKAVKVRTQLAAEYIRSGDLDSAKRSLDQALSVDSRDATANMMMG   78
NTDB id 1017 ACIAD RS02540 WP 011182074.1    MSKWTINKTLG.CAAIVGAFILSGCQTVPT.TKDPEKGVKVRTQLAAEYIKSGDLDSAKRALDQALKVNPKDSSANMMMG   78
consensus                                    *** ** !*** * *** !****!!!!**! **!!*!*!*!!!!!!!!!!*!*!*!!!*!!!!! ****!**!!!!!!

logo
NTDB id 238190 CFH90 RS03055 WP 126035299.1  VLLQREGSPQNVEKAERYFKHAIAAEPKNAQARNNYGTYLYQIGRYNDAIDQLQVAGSTLGYDQRYRALENLGRAYLQVG  158
NTDB id 1051 ABD1 RS02465 WP 004738086.1     ILLQQEGSKPNLEKAEHYFKRAISSEPDNAQARNNYGTYLYQMERYNDAIEQFRIAGATLGYDQRYQALENLGRIYLKLG  158
NTDB id 1017 ACIAD RS02540 WP 011182074.1    VLLQQEGSRPNLEKADAYFERAVQLDPENAQAHNNYGTYLYQMKRYHDAIQQLTLAGSSLGYDQRYRALENIGRIYLHLG  158
consensus                                    *!!!*!!!**!*!!!* !!**!*  *!*!!!!*!!!!!!!!!* !!*!!!*!* *!!**!!!!!!!*!!!!*!!*!! *!

logo
NTDB id 238190 CFH90 RS03055 WP 126035299.1  RVAEAEAAFKQALQVNSGAYLAMIEMAEISYLRQQNAEATQYYEQFVRAVGEKNLDARALWIGIRIARASHDTMGSQVLV  238
NTDB id 1051 ABD1 RS02465 WP 004738086.1     DIASAEKTFKQALLANRDSYISMLELAEIFYLQQQIPAATQMYEQYVRTVGQKNQGARALWIGLRVARANADKMGMQVLV  238
NTDB id 1017 ACIAD RS02540 WP 011182074.1    GVANAEKMFSQALQANRDSSVSMLELAEIYYLQQRIPAATGMYEQYVRTVGQKNQGARALWIGIRVARANADRTGVQVLV  238
consensus                                     *! !!* !*!!!**!****!*!*!!!*!!*!****!!**!!!*!!*!!*!!**!!!!!!!*!*!!!**!**!*!!!!

logo
NTDB id 238190 CFH90 RS03055 WP 126035299.1  NQLRALYPDSQEYKRYLQLQYTTEAVWK   266
NTDB id 1051 ABD1 RS02465 WP 004738086.1     NQLRALFPESPEYQRYLQLQYSTEAVWK   266
NTDB id 1017 ACIAD RS02540 WP 011182074.1    NQLRIFYPDSSEYQRYLKLQYSTEAVWK   266
consensus                                    !!!!***!*! !!*!!!*!!!*!!!!!!
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X non conserved

X similar

X ≥ 50% conserved