```
logo
NTDB id 314739 CDG62 RS05310 WP 087528037.1  .MKKPVSKLYLALTLGVSVWYLVGCQTVNT.KGDPSKAVQVRTQLAAEYIKSNDYDAAKRTLDQALDIDSRDASANMMMG   78
NTDB id 1051 ABD1 RS02465 WP 004738086.1     MS.TPKLKIMLCM.GVAVALLASGCQTSQTVKKDPEKAVKVRTQLAAEYIRSGDLDSAKRSLDQALSVDSRDATANMMMG   78
NTDB id 1017 ACIAD RS02540 WP 011182074.1    MSKWTINKTLGCA.AIVGAFILSGCQTVPT.TKDPEKGVKVRTQLAAEYIKSGDLDSAKRALDQALKVNPKDSSANMMMG   78
consensus                                    *** ** !***** *** *****!!!!**! **!!*!*!*!!!!!!!!!!*!*!*!!!*!!!!! ****!**!!!!!!

logo
NTDB id 314739 CDG62 RS05310 WP 087528037.1  VLLQREGSPQNVEKAERYFKHAIAAEPKNAQARNNYGTYLYQIGRYNDAIDQLQVAGSTLGYDQRYRALENLGRAYIQVG  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 314739 CDG62 RS05310 WP 087528037.1  RVTEAEAAFKQALQVNSGAYLAMIEMAEISYLRQQNAEATQYYEQFVQAVGEKNLDARALWIGIRIARANHDTMGSQVLV  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 314739 CDG62 RS05310 WP 087528037.1  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