```
logo
NTDB id 269177 C3F34 RS05810 WP 044739597.1  MKKVLKGMPSFSALGFKQQLLALTVGLAISVGSIAVVEAAPARNVNPPSLKAGAPQVYVVKKGDTLWDISGKFLDKPWRW   80
NTDB id 1053 ABD1 RS00870 WP 000755276.1     MKKVFNGMVNFHALGIKKHLLALALCTGVAIGTIAEVHAT.SPNHNPPSLKSNAPNVYVVKRGDTLWDISGHFLNKPWRW   79
consensus                                    !!!!  !!* !*!!! ! *!!!! ** ** *!*!! !*! * *!*!!!!!!  !!*!!!!!*!!!!!!!!!*!! !!!!!

logo
NTDB id 269177 C3F34 RS05810 WP 044739597.1  PEIWASNRHVKNPHWIFPGDKLLLCSYQGKPLIGKDEGDGCDGIIRRYAGG.TKLQPQVRVESLNDAIPVIPLDNIKQWL  159
NTDB id 1053 ABD1 RS00870 WP 000755276.1     PEIWASNQHVKNPHWIYPGDRLLLCSLDGRPLVGKDEGDGCVGIIRRYTGQTTHLQPQVRVEALNNSVPVIPLEHIKQWL  159
consensus                                    !!!!!!! !!!!!!!!*!!!*!!!!!  !*!!*!!!!!!!! !!!!!! ! *!*!!!!!!!! !!  *!!!!!**!!!!!

logo
NTDB id 269177 C3F34 RS05810 WP 044739597.1  EHSTIVAPESVENTPYVLGTADQRVLAAKGQTIYARGNGLEVGQQYAVYREGDPYTLSNEKGKKYNAGLELNQVATGIAV  239
NTDB id 1053 ABD1 RS00870 WP 000755276.1     ENSTILPADSITNTPYIVGTADQRVLAGKGQTIYARGQGLINGQRYAVYREGEPYYFTDNKGKKHSLGIELLQVASGVAV  239
consensus                                    !*!!!****!* !!!!**!!!!!!!!!*!!!!!!!!!*!! !! !!!!!!!*!! * !!!!* !*!! !!!*!*!!

logo
NTDB id 269177 C3F34 RS05810 WP 044739597.1  ASENDITTLELTDSFNAEVRRGDRVLPVYDPMLPTLFYPTMAEDVTAGGQIVRVMGSIGTAAQHSVVTIDRGSFEGVQVG  319
NTDB id 1053 ABD1 RS00870 WP 000755276.1     SSEKDITTLELTDSYNAEVRRGDRVMPEEQATLPTLFYPVDAKQVTDGGKIIRVMGSIGRAAKNSVVTLDRGTTQGIQVG  319
consensus                                     !! !!!!!!!!!!*!!!!!!!!!!*!   * !!!!!!!  !  !! !! !*!!!!!!! !! *!!!!*!!!*  !*!!!

logo
NTDB id 269177 C3F34 RS05810 WP 044739597.1  HIFSINQKGEQVRDPKTKEMVQLPGQKIGNLMVFRTFDHFSYGYVLDSSLPIKIGAYIQPPLMDD   384
NTDB id 1053 ABD1 RS00870 WP 000755276.1     QVFDITQQGESIRDPKTKEVIQLPGQQIGSLMVFRTFDQLSYAYVLESDLPIKVGSSIQSPQFND   384
consensus                                    **! ! ! !! *!!!!!!!**!!!!! !! !!!!!!!!* !!*!!!*! !!!!*!  !!*!   !
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X non conserved

X similar

X $\geq$ 50% conserved