```
logo
NTDB id 269244 C3F22 RS15645 WP 104426934.1  MKKVLKGMPFFSALGLKNQMLALAVCVGVSMGTMYSAEAAPARNVNPPALKASAPNVYVVKKGDTLWDISKRFLKNPVRW   80
NTDB id 1053 ABD1 RS00870 WP 000755276.1     MKKVFNGMVNFHALGIKKHLLALALCTGVAIGTIAEVHA.TSPNHNPPSLKSNAPNVYVVKRGDTLWDISGHFLNKPWRW   79
consensus                                    !!!!  !!* !*!!!*! **!!!!*! !! *!!*   *!** *!*!!! !!  !!!!!!!!*!!!!!!!! *!!  ! !!

logo
NTDB id 269244 C3F22 RS15645 WP 104426934.1  PEIWASNKHVKNPHWIFPGDRLLMCDYQGRPIIGKDEGDGCEGIISRYIGST.SLQPQVRVESLNNTIPVIPLAHIQQWL   159
NTDB id 1053 ABD1 RS00870 WP 000755276.1     PEIWASNQHVKNPHWIYPGDRLLLCSLDGRPLVGKDEGDGCVGIIRRYTGQTTHLQPQVRVEALNNSVPVIPLEHIKQWL   159
consensus                                    !!!!!!! !!!!!!!!*!!!!!!*!   !!!**!!!!!!!! !!! !! ! !**!!!!!!!! !!!**!!!!! !! !!!

logo
NTDB id 269244 C3F22 RS15645 WP 104426934.1  ERYTIVAADSVQGTPYILGTADQRVLAGKGQKVYARGNGLVVGQRYAVYREADPYVFTDANGKKYTAALELQQVASGIAV   239
NTDB id 1053 ABD1 RS00870 WP 000755276.1     ENSTILPADSITNTPYIVGTADQRVLAGKGQTIYARGQGLINGQRYAVYREGEPYYFTDNKGKKHSLGIELLQVASGVAV   239
consensus                                    !  !!**!!!*  !!!!*!!!!!!!!!!!!! *!!!!*!!* !!!!!!!!!**!! !!!  !!!** **!! !!!!!*!!

logo
NTDB id 269244 C3F22 RS15645 WP 104426934.1  RGEGDVTTLELTDSYNAEVRRGDRVLPEYDPMLPTLFYPVNAENTVAGGQIVRVLGSIGTAARHSVVTIDRGTAHGVQTG   319
NTDB id 1053 ABD1 RS00870 WP 000755276.1     SSEKDITTLELTDSYNAEVRRGDRVMPEEQATLPTLFYPVDAKQVTDGGKIIRVMGSIGRAAKNSVVTLDRGTTQGIQVG   319
consensus                                      ! !*!!!!!!!!!!!!!!!!!!!*!!  * !!!!!!!! ! *   !! !*!!*!!!! !!**!!!!*!!!! *!*! !

logo
NTDB id 269244 C3F22 RS15645 WP 104426934.1  HVFSVNQKGEVVTDPKTKERVQLPGERIGNVMVFKTFDQLSYAYVLESELPIKVGANIQAPLLDE   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