```
logo
NTDB id 345136 C9E88 RS01075 WP 106985327.1  MKKVLKGMPLFSALGFKKQMLALAICAGVGLGAISTVEAAPARNVNPPSLKASAPHVYVVKKGDTLWDISKKFLKNPVRW   80
NTDB id 1053 ABD1 RS00870 WP 000755276.1     MKKVFNGMVNFHALGIKKHLLALALCTGVAIGTIAEVHAT.SPNHNPPSLKSNAPNVYVVKRGDTLWDISGHFLNKPWRW   79
consensus                                    !!!!  !!* !*!!! !!**!!!!*! !!**! !  !*! * *!*!!!!!!  !!*!!!!!*!!!!!!!! *!!  ! !!

logo
NTDB id 345136 C9E88 RS01075 WP 106985327.1  PEIWASNKHVKNPHWIFPGDRLLMCTYNGKPIIGKDEGDGCEGIIRRYTGGT.KLQPQVRVESLSNAIPVIPLEHIKQWL  159
NTDB id 1053 ABD1 RS00870 WP 000755276.1     PEIWASNQHVKNPHWIYPGDRLLLCSLDGRPLVGKDEGDGCVGIIRRYTGQTTHLQPQVRVEALNNSVPVIPLEHIKQWL  159
consensus                                    !!!!!!! !!!!!!!!*!!!!!!*!*  !*!**!!!!!!!! !!!!!!!! !**!!!!!!!! ! ! *!!!!!!!!!!!!

logo
NTDB id 345136 C9E88 RS01075 WP 106985327.1  ERSSIINPESLEQTPYIVGAADHRVLAAAGQTVYARGHGLEVGQRYAIYREGQPYVLTDNQGKKYTAGLEINQVASGIAV  239
NTDB id 1053 ABD1 RS00870 WP 000755276.1     ENSTILPADSITNTPYIVGTADQRVLAGKGQTIYARGQGLINGQRYAVYREGEPYYFTDNKGKKHSLGIELLQVASGVAV  239
consensus                                    ! !*!****!* *!!!!!! !!*!!!!* !!!*!!!!*!!  !!!!!*!!!! !!  !!! !!!** !*!* !!!!!*!!

logo
NTDB id 345136 C9E88 RS01075 WP 106985327.1  RGENDITTLELTDTYTGEVRRGDRVLPEYDPMLPTVFYPTHANNVTSGGSIVRVLGSIGTAAIHSVVTIDRGSSQGVQAG  319
NTDB id 1053 ABD1 RS00870 WP 000755276.1     SSEKDITTLELTDSYNAEVRRGDRVMPEEQATLPTLFYPVDAKQVTDGGKIIRVMGSIGRAAKNSVVTLDRGTTQGIQVG  319
consensus                                      ! !!!!!!!!!*! *!!!!!!!!*!!  * !!!*!!! *! *!! !! !*!!*!!!! !! *!!!!*!!!**!!*! !

logo
NTDB id 345136 C9E88 RS01075 WP 106985327.1  YVFSVEDKGDIVNDPKTRERIQLPDQRVGQIMVFKTFDNLSYAYVLESELPIKVGASIQPPLMDD  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