```
logo
NTDB id 313232 Ahae2126ch RS15960 WP 005092484.1  MKKVLNGIQSFHALGLKKQLIAAAICAGLGMSV.NLAHAASPNVSPPSVKVGAPHVYVVKKGDTLWDISGKFLSKPWRWP   79
NTDB id 1053 ABD1 RS00870 WP 000755276.1          MKKVFNGMVNFHALGIKKHLLALALCTGVAIGTIAEVHATSPNHNPPSLKSNAPNVYVVKRGDTLWDISGHFLNKPWRWP   80
consensus                                         !!!! !!*  !!!!!*!!*!*! !*! !***  *   !! !!!* !!!*!  !!*!!!!!*!!!!!!!!!*!! !!!!!!

logo
NTDB id 313232 Ahae2126ch RS15960 WP 005092484.1  EIWASNKHVKNPHWIYPGDRLLLCSLNGKPLVGKDEGDGCAGIIRRYTGNTSTLRPQVRVEALNNSIPVIPLAHIQQWLE  159
NTDB id 1053 ABD1 RS00870 WP 000755276.1          EIWASNQHVKNPHWIYPGDRLLLCSLDGRPLVGKDEGDGCVGIIRRYTGQTTHLQPQVRVEALNNSVPVIPLEHIKQWLE  160
consensus                                         !!!!!! !!!!!!!!!!!!!!!!!!! !*!!!!!!!!!!! !!!!!!!!*!**! !!!!!!!!!!!!*!!!!! !! !!!!

logo
NTDB id 313232 Ahae2126ch RS15960 WP 005092484.1  RTSVLPADAIAHTPYILGTADQRVLAGKGQRVYVRGNGIENGQRYGVFREGEPYTILDEHGKKQVLGVELTQVASGVAIS  239
NTDB id 1053 ABD1 RS00870 WP 000755276.1          NSTILPADSITNTPYIVGTADQRVLAGKGQTIYARGQGLINGQRYAVYREGEPYYFTDNKGKKHSLGIELLQVASGVAVS  240
consensus                                          ***!!!! ! *!!!!*!!!!!!!!!!!!! *! !!*!* !!!!!*!*!!!!!!   ! *!!!* !!*!! !!!!!!!*!

logo
NTDB id 313232 Ahae2126ch RS15960 WP 005092484.1  TEKDITTVELTDSYNGEVRRGDRVMPEQDAMLPTLFYPTDANQVTDGGKIIRVMGSIGTAAKNGVVTVDRGTADGIEIGQ  319
NTDB id 1053 ABD1 RS00870 WP 000755276.1          SEKDITTLELTDSYNAEVRRGDRVMPEEQATLPTLFYPVDAKQVTDGGKIIRVMGSIGRAAKNSVVTLDRGTTQGIQVGQ  320
consensus                                         *!!!!!!*!!!!!!!*!!!!!!!!!!!!  ! !!!!!!! !! !!!!!!!!!!!!!!!! !!!! !!!*!!!!  !! *!!

logo
NTDB id 313232 Ahae2126ch RS15960 WP 005092484.1  VFNAYQDGETVKDPKTKENVKLPGQYVGSVMIFKSFDRISYAYVLESELPIKMGSSIKPPRLDN  383
NTDB id 1053 ABD1 RS00870 WP 000755276.1          VFDITQQGESIRDPKTKEVIQLPGQQIGSLMVFRTFDQLSYAYVLESDLPIKVGSSIQSPQFND  384
consensus                                         !!   ! !!***!!!!!! * !!!! *!!*!*!**!! *!!!!!!!!*!!!!*!!!! *!
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X non conserved

X similar

X ≥ 50% conserved