```
logo
NTDB id 207842 BSR56 RS14805 WP 005085668.1  MKKVLNGIQSFHALGLKKQLIAAAICAGLGMSV.NLAHAASPNVSPPSVKVGAPHVYVVKKGDTLWDISGKFLSKPWRWP   79
NTDB id 1053 ABD1 RS00870 WP 000755276.1     MKKVFNGMVNFHALGIKKHLLALALCTGVAIGTIAEVHATSPNHNPPSLKSNAPNVYVVKRGDTLWDISGHFLNKPWRWP   80
consensus                                    !!!! !!*  !!!!!*!!*!*! !*! !*** *   !! !!!* !!!*!  !!*!!!!!*!!!!!!!!!*!! !!!!!!

logo
NTDB id 207842 BSR56 RS14805 WP 005085668.1  EIWASNKHVKNPHWIYPGDRLLLCSLNGKPLVGKDEGDGCAGIIRRYTGNTSTLRPQVRVEALNNSIPVIPLAHIQQWLE  159
NTDB id 1053 ABD1 RS00870 WP 000755276.1     EIWASNQHVKNPHWIYPGDRLLLCSLDGRPLVGKDEGDGCVGIIRRYTGQTTHLQPQVRVEALNNSVPVIPLEHIKQWLE  160
consensus                                    !!!!!! !!!!!!!!!!!!!!!!!!! !*!!!!!!!!!!! !!!!!!!!*!**! !!!!!!!!!!!*!!!!! !! !!!!

logo
NTDB id 207842 BSR56 RS14805 WP 005085668.1  RTSVLPADAIAHTPYILGTADQRVLAGKGQRVYVRGNGIENGQRYGVFREGEPYTILDEHGKKHILGVELTQVASGVAIS  239
NTDB id 1053 ABD1 RS00870 WP 000755276.1     NSTILPADSITNTPYIVGTADQRVLAGKGQTIYARGQGLINGQRYAVYREGEPYYFTDNKGKKHSLGIELLQVASGVAVS  240
consensus                                     ***!!!! ! *!!!!*!!!!!!!!!!!!! *! !!*!* !!!!!*!*!!!!!!   ! *!!!! !!*!! !!!!!!!*!

logo
NTDB id 207842 BSR56 RS14805 WP 005085668.1  TEKDITTVELTDSYNGEVRRGDRVMPEQDAMLPTLFYPTDANQVTDGGKIIRVMGSIGTAAKNGVVTVDRGTADGIEIGQ  319
NTDB id 1053 ABD1 RS00870 WP 000755276.1     SEKDITTLELTDSYNAEVRRGDRVMPEEQATLPTLFYPVDAKQVTDGGKIIRVMGSIGRAAKNSVVTLDRGTTQGIQVGQ  320
consensus                                    *!!!!!!*!!!!!!!*!!!!!!!!!!!  ! !!!!!!! !! !!!!!!!!!!!!!!!! !!!! !!!*!!!!  !! *!!

logo
NTDB id 207842 BSR56 RS14805 WP 005085668.1  VFNAYQDGETVKDPKTKENVKLPGQYVGSVMIFKSFDRISYAYVLESELPIKMGSSIKPPRLDN  383
NTDB id 1053 ABD1 RS00870 WP 000755276.1     VFDITQQGESIRDPKTKEVIQLPGQQIGSLMVFRTFDQLSYAYVLESDLPIKVGSSIQSPQFND  384
consensus                                    !!   ! !!***!!!!!! * !!!! *!!*!*!**!! *!!!!!!!!*!!!!*!!!! *!
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X non conserved

X similar

X $\geq$ 50% conserved