```
logo
NTDB id 313065 AhaeAN43 RS15785 WP 005092484.1  MKKVLNGIQSFHALGLKKQLIAAAICAGLGMSV.NLAHAASPNVSPPSVKVGAPHVYVVKKGDTLWDISGKFLSKPWRWP   79
NTDB id 1053 ABD1 RS00870 WP 000755276.1        MKKVFNGMVNFHALGIKKHLLALALCTGVAIGTIAEVHATSPNHNPPSLKSNAPNVYVVKRGDTLWDISGHFLNKPWRWP   80
consensus                                       !!!! !!*  !!!!!*!!*!*! !*! !***  *   !! !!!* !!!*!  !!*!!!!!*!!!!!!!!!*!! !!!!!!

logo
NTDB id 313065 AhaeAN43 RS15785 WP 005092484.1  EIWASNKHVKNPHWIYPGDRLLLCSLNGKPLVGKDEGDGCAGIIRRYTGNTSTLRPQVRVEALNNSIPVIPLAHIQQWLE  159
NTDB id 1053 ABD1 RS00870 WP 000755276.1        EIWASNQHVKNPHWIYPGDRLLLCSLDGRPLVGKDEGDGCVGIIRRYTGQTTHLQPQVRVEALNNSVPVIPLEHIKQWLE  160
consensus                                       !!!!!! !!!!!!!!!!!!!!!!!!! !*!!!!!!!!!!!! !!!!!!!!*!**! !!!!!!!!!!!!*!!!!! !! !!!!

logo
NTDB id 313065 AhaeAN43 RS15785 WP 005092484.1  RTSVLPADAIAHTPYILGTADQRVLAGKGQRVYVRGNGIENGQRYGVFREGEPYTILDEHGKKQVLGVELTQVASGVAIS  239
NTDB id 1053 ABD1 RS00870 WP 000755276.1        NSTILPADSITNTPYIVGTADQRVLAGKGQTIYARGQGLINGQRYAVYREGEPYYFTDNKGKKHSLGIELLQVASGVAVS  240
consensus                                        ***!!!! ! *!!!!*!!!!!!!!!!!!!! *! !!*!* !!!!!*!*!!!!!!   ! *!!!* !!*!! !!!!!!!*!

logo
NTDB id 313065 AhaeAN43 RS15785 WP 005092484.1  TEKDITTVELTDSYNGEVRRGDRVMPEQDAMLPTLFYPTDANQVTDGGKIIRVMGSIGTAAKNGVVTVDRGTADGIEIGQ  319
NTDB id 1053 ABD1 RS00870 WP 000755276.1        SEKDITTLELTDSYNAEVRRGDRVMPEEQATLPTLFYPVDAKQVTDGGKIIRVMGSIGRAAKNSVVTLDRGTTQGIQVGQ  320
consensus                                       *!!!!!!*!!!!!!!*!!!!!!!!!!!!  ! !!!!!!! !! !!!!!!!!!!!!!!!! !!!! !!!*!!!!  !! *!!

logo
NTDB id 313065 AhaeAN43 RS15785 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