```
logo
NTDB id 1101203 ACNANF RS24505 WP 033696100.1  ..................MLAGKQLLLEEISSDLQRELND.LKKKGEVICVQGVKKKNSKYMCQRCGNVDWRLFASFLCKR   62
NTDB id 108 BSU 35470 NP 391427.1              MNVPVEKNSSFSKELQQTLRSRHLLRTELSF..SDEMIEWHIKNGYITAENSISINKRRYRCNRCGQTDQRYFSFYHS..   76
consensus                                      ***************** !  **!!  !*! **  !* *** ! ! * * * *     *! !*!!!* ! ! !  *****

logo
NTDB id 1101203 ACNANF RS24505 WP 033696100.1  CSKVCTYCRKCITMGRVSECAVLVRGIAERKREKNSNLLQWNGTLSTGQNLAVQGVIEAIKQKESFFIWAVCGAGKTEML   142
NTDB id 108 BSU 35470 NP 391427.1              SGKNKLYCRSCVMMGRVSEEVPLYSWKEENESNWKSIKLTWDGKLSSGQQKAANVLIEAISKKEELLIWAVCGAGKTEML   156
consensus                                      * ! * !!! !* !!!!!!* *!     !       !  ! ! ! !!*!!* ! * *!!!!  !!   !!!!!!!!!!!!!

logo
NTDB id 1101203 ACNANF RS24505 WP 033696100.1  FYGINEALQKGERVCIATPRTDVVLELAPRLQEVFPYIKVAALYGGSVDKEKDAVLVVATTHQLLRYYRAFHVIVVDEID   222
NTDB id 108 BSU 35470 NP 391427.1              FPGIESALNQGLRVCIATPRTDVVLELAPRLKAAFQGADISALYGGSDDKGRLSPLMISTTHQLLRYKDAIDVMIIDEVD   236
consensus                                      !*!!  !!* ! !!!!!!!!!!!!!!!!!!!!   !*   * !!!!!! !! *  *!** !!!!!!!!  ! *!***!!*!

logo
NTDB id 1101203 ACNANF RS24505 WP 033696100.1  AFPYCADQMLQYAVKQAMKEKAARIYLTATPDETWKRKFRKGEQKGVIVSGRYHRHPLPVPLFCWCGNWKKSLNRERIPR   302
NTDB id 108 BSU 35470 NP 391427.1              AFPYSADQTLQFAVQKARKKNSTLVYLSATPPKELKRKALNGQLHSVRIPARHHRKPLPEPRFVWCGNWKKKLNRNKIPP   316
consensus                                      !!!!*!!! !!*!!  ! !     *!!*!!!*   !!!   !  * ! ***!*!!*!!! ! !*!!!!!!! !!! *!!*

logo
NTDB id 1101203 ACNANF RS24505 WP 033696100.1  VLLQWLKMYLNKKYPVFLFVPHVRYIEEISSLLKSLHNKVEGVHAEDPMRKEKVAAFRKGEIPLLVTTTILERGVTVKNL   382
NTDB id 108 BSU 35470 NP 391427.1              AVKRWIEFHVKEGRPVFLFVPSVSILEKAAACFKGVHCRTASVHAEDKHRKEKVQQFRDGQLDLLITTTILERGVTVPKV   396
consensus                                       *  !*  **    !!!!!!!*!  *!    * ! *!**   !!!!!**!!!!!  !! ! **!!*!!!!!!!!!!!* *

logo
NTDB id 1101203 ACNANF RS24505 WP 033696100.1  QVAVLGAEEEIFSESALVQIAGRAGRSFEAPYGEVIYFHYGKTEAMVRAKKHIQGMNKNAKEQGLID   449
NTDB id 108 BSU 35470 NP 391427.1              QTGVLGAESSIFTESALVQIAGRTGRHKEYADGDVIYFHFGKTKSMLDARKHIKEMNELAAKVECTD   463
consensus                                      ! *!!!!!  !!*!!!!!!!!!! !!* ! * !*!!!!!*!!!  !* !*!!!  !!  !    * !
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X non conserved

X similar

X ≥ 50% conserved