```
logo
NTDB id 22942 BC RS25890 WP 000225360.1  ..................MVAGKQLLLEELSSDLQRKLNDLKKKGEIVCVQGVKKKNSKYMCQRCGNVDRRLFASFLCKRCSKVCT   68
NTDB id 108 BSU 35470 NP 391427.1        MNVPVEKNSSFSKELQQTLRSRHLLRTELSFS.DEMIEWHIKNGYITAENSISINKRRYRCNRCGQTDQRYFSFYHS..SGKNKL   82
consensus                                ***************** *  **!!  !!!  *   *  * ! ! ! * * *     *! !*!!!* ! ! !  ****** ! *

logo
NTDB id 22942 BC RS25890 WP 000225360.1  YCRKCITMGRVSECAVLVRGIAERKREKNSNLLQWNGTLSTGQNLAAQGVIEAIKQKESFFIWAVCGAGKTEMLFYGINEALQKG  153
NTDB id 108 BSU 35470 NP 391427.1        YCRSCVMMGRVSEEVPLYSWKEENESNWKSIKLTWDGKLSSGQQKAANVLIEAISKKEELLIWAVCGAGKTEMLFPGIESALNQG  167
consensus                                !!! !* !!!!!!* *!     !      !  ! ! ! !!*!!* !!* *!!!!  !!   !!!!!!!!!!!!!!*!!  !!* !

logo
NTDB id 22942 BC RS25890 WP 000225360.1  ERVCIATPRTDVVLELAPRLQKVFPYIKVAALYGGSVDKEKDAVLVVATTHQLLRYYRAFHVIVVDEIDAFPYCADQMLQYAVKQ  238
NTDB id 108 BSU 35470 NP 391427.1        LRVCIATPRTDVVLELAPRLKAAFQGADISALYGGSDDKGRLSPLMISTTHQLLRYKDAIDVMIIDEVDAFPYSADQTLQFAVQK  252
consensus                                 !!!!!!!!!!!!!!!!!!!   !*   * !!!!!! !! *  *!** !!!!!!!!  ! *!***!!*!!!!!*!!! !!*!!

logo
NTDB id 22942 BC RS25890 WP 000225360.1  AMKEKAARIYLTATPDETWKRKFRKGEQKGVIVSGRYHRHPLPVPLFCWCGNWKKSLNRERIPRVLLQWLKMYLNKKYPVFLFVP  323
NTDB id 108 BSU 35470 NP 391427.1        ARKKNSTLVYLSATPPKELKRKALNGQLHSVRIPARHHRKPLPEPRFVWCGNWKKKLNRNKIPPAVKRWIEFHVKEGRPVFLFVP  337
consensus                                ! !     *!!*!!!*   !!!    !  * ! ***!*!!*!!! ! !*!!!!!!! !!! *!!* *  !*  **    !!!!!!!

logo
NTDB id 22942 BC RS25890 WP 000225360.1  HVRYIEEISSLLKSLHNKVEGVHAEDPMRKEKVAAFRKGEIPLLVTTTILERGVTVINLQVAVLGAEEEIFSESALVQIAGRAGR  408
NTDB id 108 BSU 35470 NP 391427.1        SVSILEKAAACFKGVHCRTASVHAEDKHRKEKVQQFRDGQLDLLITTTILERGVTVPKVQTGVLGAESSIFTESALVQIAGRTGR  422
consensus                                *!  *!    * ! *!**   !!!!!**!!!!!  !! ! **!!*!!!!!!!!!!!* *! *!!!!!  !!*!!!!!!!!!! !!

logo
NTDB id 22942 BC RS25890 WP 000225360.1  SFEAPYGEVIYFHYGKTEAMVRAKKHIQGMNKNAKEQGLID   449
NTDB id 108 BSU 35470 NP 391427.1        HKEYADGDVIYFHFGKTKSMLDARKHIKEMNELAAKVECTD   463
consensus                                * ! * !*!!!!!*!!!  !* !*!!!  !!  !    * !
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X non conserved

X similar

X ≥50% conserved