```
logo
NTDB id 174317 BaDB11 RS09770 WP 059260886.1  .MIHIEQPAFYTRELRSCLEQRHLLKSELPFRESVVDWHIQEGLIKTEEGIKKTKKGFICLRCGQHERSFFARYPCYRCS   79
NTDB id 108 BSU 35470 NP 391427.1             MNVPVEKNSSFSKELQQTLRSRHLLRTELSFSDEMIEWHIKNGYITAENSISINKRRYRCNRCGQTDQRYFSFYHS..SG   78
consensus                                     * * *! *  ***!!  *!  !!!!**!!*! * ***!!!  ! !  !  !   !* * ! !!!!**  *!  ! ****

logo
NTDB id 174317 BaDB11 RS09770 WP 059260886.1  KCCVYCRSCVMMGRVSDCTPLLTWHMHVSHKWAPVHTEWKGALSAGQEKAAKSIINAIRRKEELLVWAVCGSGKTELLFQ   159
NTDB id 108 BSU 35470 NP 391427.1             KNKLYCRSCVMMGRVSEEVPLYSWKEENESNWKSIKLTWDGKLSSGQQKAANVLIEAISKKEELLIWAVCGAGKTEMLFP   158
consensus                                     !***!!!!!!!!!!!!** !! *!* *  * ! ***  ! ! !! !! !!!  *! !! *!!!!!*!!!!! !!!!*!!*

logo
NTDB id 174317 BaDB11 RS09770 WP 059260886.1  GIEFALNNGLTVCIATPRTDVVLELAPRFRQAFPGVEIAALYGGSSDVGNLSPLIISTTHQLLRYKEAFDVIIIDEVDAF   239
NTDB id 108 BSU 35470 NP 391427.1             GIESALNQGLRVCIATPRTDVVLELAPRLKAAFQGADISALYGGSDDKGRLSPLMISTTHQLLRYKDAIDVMIIDEVDAF   238
consensus                                     !!!  !!!*!! !!!!!!!!!!!!!!!!! * !!*! *! !!!!!! ! ! !!!!*!!!!!!!!!!!*! !!*!!!!!!!!

logo
NTDB id 174317 BaDB11 RS09770 WP 059260886.1  PYSIDNTLQYAVKKSAKRQSAHIYLTATPSADMKKRAESGKLDTVRIPARFHRSPLPEPTLLWCGNWKKSLKRRKVPFRL   319
NTDB id 108 BSU 35470 NP 391427.1             PYSADQTLQFAVQKARKKNSTLVYLSATPPKELKRKALNGQLHSVRIPARHHRKPLPEPRFVWCGNWKKKLNRNKIPPAV   318
consensus                                     !!!  !*!!!*!! !  !**! **!!*!!!* **!**!  ! !**!!!!!!*!! !!!!!  *!!!!!!! ! ! !*!* *

logo
NTDB id 174317 BaDB11 RS09770 WP 059260886.1  KKWLFKHQELQQPVFLFVPSVPVLKSVVSVLKKETFRAEGVYADDPDRNEKVNRFRESKLEVLVTTTILERGVTVKKAQV   399
NTDB id 108 BSU 35470 NP 391427.1             KRWIEFHVKEGRPVFLFVPSVSILEKAAACFKGVHCRTASVHAEDKHRKEKVQQFRDGQLDLLITTTILERGVTVPKVQT   398
consensus                                     !*!*  !      !!!!!!!!!**!     * !  **!   !*!*!**! !!!* !!*  !**!*!!!!!!!!!!!*! !

logo
NTDB id 174317 BaDB11 RS09770 WP 059260886.1  GVLGAESAVFTESALVQMAGRAGRHPEHTDGDVCFFHYGRTKSMNEARRHIQNMNKMSKKEMLID   464
NTDB id 108 BSU 35470 NP 391427.1             GVLGAESSIFTESALVQIAGRTGRHKEYADGDVIYFHFGKTKSMLDARKHIKEMNELAAKVECTD   463
consensus                                     !!!!!!! *!!!!!!!!!*!!! !!!*!* !!!!**!!*!*!!!! *!!*!!  !! *  !  * !
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X non conserved

X similar

X ≥50% conserved