```
logo
NTDB id 80168 L6483 RS25960 WP 236787299.1  ..................MVAGKQLLLEELPSDLRRELSD.LKKEGEVICVQGVIKKASKYICQRCGNIEQRLFASFLCKRCSKVC   67
NTDB id 108 BSU 35470 NP 391427.1           MNVPVEKNSSFSKELQQTLRSRHLLRTELS..FSDEMIEWHIKNGYITAENSISINKRRYRCNRCGQTDQRYFSFYHS..SGKNK   81
consensus                                   **************** *  **!!  !!***   !* *** ! ! * * * *     *! !*!!!* *!! !  ****** ! *

logo
NTDB id 80168 L6483 RS25960 WP 236787299.1  TYCRKCITMGRVSECAVLVRGIHERKGERELHSLQWKGSLSLGQELAAQGIIEAIKQKESFFIWAVCGAGKTEMLFYGIESALQK  152
NTDB id 108 BSU 35470 NP 391427.1           LYCRSCVMMGRVSEEVPLYSWKEENESNWKSIKLTWDGKLSSGQQKAANVLIEAISKKEELLIWAVCGAGKTEMLFPGIESALNQ  166
consensus                                    !!! !* !!!!!!* *!    *!       * ! ! ! !! !!  !!* *!!!!  !!   !!!!!!!!!!!!!!*!!! !!*

logo
NTDB id 80168 L6483 RS25960 WP 236787299.1  GERVCIATPRTDVVLELVPRLQEVFPSINVAALYGGSVDREKDAALVVATTHQLLRYYRAFHVMIVDEIDAFPYHADQMLQYAVQ  237
NTDB id 108 BSU 35470 NP 391427.1           GLRVCIATPRTDVVLELAPRLKAAFGGADISALYGGSDDKGRLSPLMISTTHQLLRYKDAIDVMIIDEVDAFPYSADQTLQFAVQ  251
consensus                                   ! !!!!!!!!!!!!!!! !!!   !*   * !!!!!! !* *  *!** !!!!!!!!  ! *!!!*!!*!!!!!*!!! !!*!!!

logo
NTDB id 80168 L6483 RS25960 WP 236787299.1  QAMKERAARIYLTATPDEKWKRNFRTGKQKGIIVSGRYHRHPLPVPLFSWCGNWKKSLHHKKIPRVLLQWLKMYVNKKYPIFLFV  322
NTDB id 108 BSU 35470 NP 391427.1           KARKKNSTLVYLSATPPKELKRKALNGQLHSVRIPARHHRKPLPEPRFVWCGNWKKKLNRNKIPPAVKRWIEFHVKEGRPVFLFV  336
consensus                                    ! !     *!!*!!!*   !!    !  * * ***!*!!*!!! ! ! !!!!!!! !** !!!* *  !*  *!    !*!!!!

logo
NTDB id 80168 L6483 RS25960 WP 236787299.1  PHVRYIEEISLLLKELDHRIDGVHAEDPMRKEKVAAFRKGDIPLLVTTTILERGVTVKNLQVAVLGAEEEIFSESALVQIAGRAG  407
NTDB id 108 BSU 35470 NP 391427.1           PSVSILEKAAACFKGVHCRTASVHAEDKHRKEKVQQFRDGQLDLLITTTILERGVTVPKVQTGVLGAESSIFTESALVQIAGRTG  421
consensus                                   !*!  *!    * ! ** !   !!!!!**!!!!!  !! ! **!!*!!!!!!!!!!!* *! *!!!!!  !!*!!!!!!!!!! !

logo
NTDB id 80168 L6483 RS25960 WP 236787299.1  RSFEEPYGEVVYFHYGKTESMVRAKRHIQSMNKSAKEQGLID   449
NTDB id 108 BSU 35470 NP 391427.1           RHKEYADGDVIYFHFGKTKSMLDARKHIKEMNELAAKVECTD   463
consensus                                   !* ! * !*!*!!!*!!! !!* !**!!  !!  !    * !
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X non conserved

X similar

X ≥50% conserved