```
logo
NTDB id 241224 BaGK RS06590 WP 094231807.1  MFSAVTSEGKTVYLMKERYAATEASELRKTKFFCPVCREQLDVKLGSLKAPHFAHKQNKACTIEFEPESAYHLEGKKQLYSWLKE   85
NTDB id 123 BSU 11530 NP 389035.1           ..............MFHLLGAQQNQKLKRRRFFCPVCGGELAVKLGLQKAPHFAHKQNKSCAIDIEPESAYHLEGKRQLYVWLKT   71
consensus                                   **************! *  *!     !** *!!!!!!   ! !!!!  !!!!!!!!!!! ! !* !!!!!!!!!!!*!!! !!!

logo
NTDB id 241224 BaGK RS06590 WP 094231807.1  KGCSPVLEPYVKSIQQRPDIMANVQGDMLAFEFQCANLAPDLLYKRTQGLKKAGAEPLWIIGGNRLKRTSRHFFQLSAFHWQFMK  170
NTDB id 123 BSU 11530 NP 389035.1           QRASPILEPYIRTINQRPDVMARIKEHMLAVEYQCATIAPDVFQKRTEGFKQEGIIPQWIMGYSRLKRTASSFYQLSTFHWQFIN  156
consensus                                    *!!*!!!!***!*!!!!*!! *  *!!! !*!!! *!!!*  !!! ! !  !  ! !!*!  !!!!!  *!*!!! !!!!!*

logo
NTDB id 241224 BaGK RS06590 WP 094231807.1  ENPQQFLLFYCPEARSFYKLHHIIPFYANYACASLDAIPLTKANVKDMWFSASAPSFQPAAWQNAITRFRHKPPRFLSKEANHLR  255
NTDB id 123 BSU 11530 NP 389035.1           ASPYRELICYCPERRSFLRLSHIIPFYTNHSYSSVQTIPIHRAGAGDLFFTEPKPSIQYSGWTKAIHRFRHKPHRFNSKETNRLR  241
consensus                                     !   !**!!!! !!! *!*!!!!!! !* * !*  !!***!   !**!* * !! !* *!  !!*!!!!!! !! !!! !*!!

logo
NTDB id 241224 BaGK RS06590 WP 094231807.1  LLFYEKRQTPLPFLPPEVFAPIPLGAVFLSPVFIWQGHLYLYMTDLAEKRMSIRFSAVFRFCKMQIQKKRIHLRYEYSEQLLREA  340
NTDB id 123 BSU 11530 NP 389035.1           LLFYEKRQTPFSFLPTEVFVPVRKGAVFKSPVFVWQGFLYLFMTDLGGKRAPIRFSAVLQQCKLHIHNKNIALRSECSEECLSEA  326
consensus                                   !!!!!!!!!! *!!!*!!! !** !!!! !!!!*!!!*!!!*!!!!* !! *!!!!!!   !!**!* ! !*!! !*!! *! !!

logo
NTDB id 241224 BaGK RS06590 WP 094231807.1  LKQYINFLCGKEFLIEGEKEVYMVKNPVHDCRTLEGVMERDQTCFGE   387
NTDB id 123 BSU 11530 NP 389035.1           VKQYIDFLCKKGFLRETQKEVYVLNQPAGGIHSMQDLIERDRSCFIE   373
consensus                                   *!!!! !!! ! !! !  !!!!** *! * ****  **!!! *!! !
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X non conserved

X similar

X $\geq$ 50% conserved