```
logo
NTDB id 1035742 ABKQ09 RS06145 WP 328184600.1  MFSAVTSEGKTVYLMKERYAATEASELRKTKFFCPVCREQLDVKLGSLKAPHFAHKQNKACTIEFEPESAYHLEGKKQLY   80
NTDB id 123 BSU 11530 NP 389035.1              ..............MFHLLGAQQNQKLKRRRFFCPVCGGELAVKLGLQKAPHFAHKQNKSCAIDIEPESAYHLEGKRQLY   66
consensus                                      **************! *  *!     !** *!!!!!!   ! !!!!  !!!!!!!!!!! ! !* !!!!!!!!!!!*!!!

logo
NTDB id 1035742 ABKQ09 RS06145 WP 328184600.1  SWLKAKGCSPILEPYVKSIQQRPDIMANVQGDMLAFEEFQCANLAPDLLYKRTQGLKQAGAEPLWIIGGNRLKRTSRHFFQ  160
NTDB id 123 BSU 11530 NP 389035.1              VWLKTQRASPILEPYIRTINQRPDVMARIKEHMLAVEYQCATIAPDVFQKRTEGFKQEGIIPQWIMGYSRLKRTASSFYQ  146
consensus                                       !!!   *!!!!!!***!*!!!!*!! *  *!!! !*!!! *!!!*  !!! ! !! !  ! !!*!  !!!!!  *!*!

logo
NTDB id 1035742 ABKQ09 RS06145 WP 328184600.1  LSAFHWQFMKENPRQFLLFYCPEARSFYKLHHITPFYANYACASLEAIPLTKANVKDMWFSASAPSFQPAAWQSAITRFR  240
NTDB id 123 BSU 11530 NP 389035.1              LSTFHWQFINASPYRELICYCPERRSFLRLSHIIPFYTNHSYSSVQTIPIHRAGAGDLFFTEPKPSIQYSGWTKAIHRFR  226
consensus                                       !! !!!!!*   !   !**!!!! !!! *!*!! !!! !* * !*  !!***!   !**!* * !! !* *!  !!*!!!

logo
NTDB id 1035742 ABKQ09 RS06145 WP 328184600.1  HKPPRFLSKEANHLRLLFYEKRQTPLPFLPPEVFAPIPLGAVFLSPVFIWQGHLYLYMTDLAEKRMSIRFSAVFRFCKMQ  320
NTDB id 123 BSU 11530 NP 389035.1              HKPHRFNSKETNRLRLLFYEKRQTPFSFLPTEVFVPVRKGAVFKSPVFVWQGFLYLFMTDLGGKRAPIRFSAVLQQCKLH  306
consensus                                       !!! !! !!! !*!!!!!!!!!!!! *!!!*!!! !** !!!! !!!!*!!!*!!!*!!!!* !! *!!!!!!   !!**

logo
NTDB id 1035742 ABKQ09 RS06145 WP 328184600.1  IRKKRIHLRYENSEHLLREALKQYINFLCRKEFLIEGEKEVYMVKYPVHDCRTLEGVMERDQTCFGE  387
NTDB id 123 BSU 11530 NP 389035.1              IHNKNIALRSECSEECLSEAVKQYIDFLCKKGFLRETQKEVYVLNQPAGGIHSMQDLIERDRSCFIE  373
consensus                                       !* ! !*!! !*!!**! !!*!!!! !!!*! !! !  !!!!**  ! * ****  **!!! *!! !
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X non conserved

X similar

X $\geq$ 50% conserved