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logo

NTDB id 274602 FORC77 RS10545 WP 011149911.1  MQQITTTKQRYLLMVEDTASVAALYRSYLMPLGIDINIVGTGRDAIDSLKHRIPDLILLDLRLPDMTGMDVLHAVKASHP   80
NTDB id 1157 A1552VC RS03895 WP 001888250.1  .............MVEDTASVAALYRSYLTPLDIDINIVGTGRDAIESIGRREPDLILLDLRLPDMTGMDVLYAVKEKSP   67
consensus                                    *************!!!!!!!!!!!!!!!! !! !!!!!!!!!!!!!*!* *! !!!!!!!!!!!!!!!!!!!!*!!!  *!

logo

NTDB id 274602 FORC77 RS10545 WP 011149911.1  DVPIIFMTAHGSIDTAVEAMRHGSQDFLIKPCEADRLRVTVNNAIRKASKLKNDADSAGSQNYQGFIGSSQKMQQVYRTI  160
NTDB id 1157 A1552VC RS03895 WP 001888250.1  DVPIVFMTAHGSIDTAVEAMRHGAQDFLIKPCEADRLRVTVNNAIRKASKLKNDVDN.KNQNYQGFIGSSQTMQAVYRTI  146
consensus                                    !!!!*!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!! ! *  !!!!!!!!!!! !! !!!!!

logo

NTDB id 274602 FORC77 RS10545 WP 011149911.1  DSAASSKASIFITGESGTGKEVCAEAIHAASRRGDKPFIAINCAAIPKDLIESELFGHVKGAFTGAATDRQGAAELADGG  240
NTDB id 1157 A1552VC RS03895 WP 001888250.1  DSAASSKASIFITGESGTGKEVCAEAIHAASKRGDKPFIAINCAAIPKDLIESELFGHVKGAFTGAATERQGAAEAADGG  226
consensus                                    !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!! !!!!

logo

NTDB id 274602 FORC77 RS10545 WP 011149911.1  TLFLDELCEMDLDLQTKLLRFIQTGTFQKVGSSKMKSVDVRFVCATNRDPWKEVQEGRFREDLYYRLYVIPLHLPPLRER  320
NTDB id 1157 A1552VC RS03895 WP 001888250.1  TLFLDELCEMDLDLQTKLLRFIQTGTFQKVGSSKMKSVDVRFVCATNRDPWKEVQEGRFREDLYYRLYVIPLHLPPLRAR  306
consensus                                    !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !

logo

NTDB id 274602 FORC77 RS10545 WP 011149911.1  GEDVIEIAYSLLGYMSHEEGKNFVRFSQPVIDRFNEYEWPGNVRQLQNVLRNVVVLNNGKEITMEMLPPPLNQPFERKES  400
NTDB id 1157 A1552VC RS03895 WP 001888250.1  GDDVIEIAYSLLGFMSKEEGKDFVRLSAEVVERFRQYEWPGNVRQLQNVLRNVVVLNEGREITLDMLPPPLNQMSAPINR  386
consensus                                    !*!!!!!!!!!!!*!!*!!!! !!! ! *!**!! !!!!!!!!!!!!!!!!!!!!! !*!!!**!!!!!!!!!*  *

logo

NTDB id 274602 FORC77 RS10545 WP 011149911.1  VQP.DLSELISVRDICPLWLTEKLAIEQAIKACDGNIPRAAGYLDVSPSTIYRKLQAWNEKEEKQKA..  466
NTDB id 1157 A1552VC RS03895 WP 001888250.1  ALPLAHENKVSVHEIFPLWMTEKQAIEQAIEACDGNIPRAATYLDVSPSTIYRKLQTWNEKVKEKEKER  455
consensus                                      !* *   *!!**!*!!!*!!! !!!!!! !!!!!!!!!! !!!!!!!!!!!!!! !!!!       **
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X non conserved

X similar

X ≥ 50% conserved