

logo

NTDB id 266660 RL00 RS10785 WP 001186600.1
NTDB id 11 SA RS00430 WP 001186602.1
consensus

MQQLKTKRVGIYVRVSTEMQSTEGYSIDGQINQIKEYCDFHHFEVKDIYADRGISGKSMNRPELQRILKDAKGGYIDCVMVYKTN
MQQLKTKRVGIYVRVSTEMQSTEGYSIDGQINQIKEYCDFHHFEVKDIYADRGISGKSMNRPELQRILKDAKGGYIDCVMVYKTN
MQQLKTKRVGIYVRVSTEMQSTEGYSIDGQINQIKEYCDFHHFEVKDIYADRGISGKSMNRPELQRILKDAKGGYIDCVMVYKTN
!!

85
85

logo

NTDB id 266660 RL00 RS10785 WP 001186600.1
NTDB id 11 SA RS00430 WP 001186602.1
consensus

RLARNTSDLLKIVEDLHKQNVFFSLSERMEVNTSSGKMLMLQILASFSEFERNNIVENVMGQTRRAQEGYYQGNLPLGYDKIPN
RLARNTSDLLKIVEDLHKQNVFFSLSERMEVNTSSGKMLMLQILASFSEFERNNIVENVMGQTRRAQEGYYQGNLPLGYDKIPN
RLARNTSDLLKIVEDLHKQNVFFSLSERMEVNTSSGKMLMLQILASFSEFERNNIVENVMGQTRRAQEGYYQGNLPLGYDKIPN
!!

170
170

logo

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consensus

SKHELMINQHEANIVKYIFESYAKGHGYRKIANALNHKGYVTKKGKPFSSISITYILANPFYIGKIQFAKYKDWSEKRRKGLNDK
SKHELMINQHEANIVKYIFESYAKGHGYRKIANALNHKGYVTKKGKPFSSISITYILANPFYIGKIQFAKYKDWSEKRRKGLNDK
SKHELMINQHEANIVKYIFESYAKGHGYRKIANALNHKGYVTKKGKPFSSISITYILANPFYIGKIQFAKYKDWSEKRRKGLNDK
!!

255
255

logo

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consensus

PVIAEGKHSPINQDLWDKVMRKKQVSQKPQVHGKGTNLLTGTIHCPCGAPMAASNTTNTLKDGTKKRIRYYSCSNFRNKGSK
PVIAEGKHSPINQDLWDKVMRKKQVSQKPQVHGKGTNLLTGTIHCPCGAPMAASNTTNTLKDGTKKRIRYYSCSNFRNKGSK
PVIAEGKHSPINQDLWDKVMRKKQVSQKPQVHGKGTNLLTGTIHCPCGAPMAASNTTNTLKDGTKKRIRYYSCSNFRNKGSK
!!

340
340

logo

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consensus

VCSANSVRANVIEDYVMKQILEIVKSDKIERVVAHVNVQENQVDCASLHHDIA YKQQYDEVQIKLNNLIKTIEDNPDLTSVIRP
VCSANSVRANVIEDYVMKQILEIVKSDKIERVVAHVNVQENQVDCASLHHDIA YKQQYDEVQIKLNNLIKTIEDNPDLTSVIRP
VCSANSVRADVIEDYVMKQILEIVKSDKIQRVVTHVNVQENQVDCASLHHDIA YKQQYDEVQIKLNNLIKTIEDNPDLTSVIRP
!!

425
425

logo

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consensus

SIQKYEKQLNDITQINQLKNQQNEDKPLFDAKEISKLLQHFHDIKHIEKSRLKALYLSVIDRIDIKKDGNNHKKQFYVTLKLN
SIQKYEKQLNDITYQINQLKNQQNEDKPLFDAKEISKLLQHFHDIKHIEKSRLKALYLSVIDRIDIKKDGNNHKKQFYVTLKLN
SIQKYEKQLNDITNQINQLKNQQNEDKPLFDAKEISKLLQHFHDIKHIEKSRLKALYLSVIDRIDIKKDGNNHKKQFYVTLKLN
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510
510

logo

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consensus

EIIKQLFNNKQLDEVHLSTSSFLPQTLYLTI
EIIKQLFNNKQLDEVHLSTSSFLPQTLYLTI
EIIKQLFNNKQLDEVHLSTSSFLPQTLYLTI
!!

542
542

non conserved
 similar
 $\geq 50\%$ conserved