

logo

NTDB id 71660 D5070 RS00225 WP 001186608.1
NTDB id 11 SA RS00430 WP 001186602.1
consensus

85
85

logo

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consensus

RLARNTSDLKIVEDLHKQNEFFSLSERMEVNTSSGKMLQLASFSEFERNIVENVFMGQTRRAQEGYYQGNLPLGYDKIP
 RLARNTSDLLKIVEDLHKQNEFFSLSERMEVNTSSGKMLQLILASFSEFERNIVENVFMGQTRRAQEGYYQGNLPLGYDKIP
 RLARNTSDLLKIVEDLHKQNEFFSLSERMEVNTSSGKMLQLILASFSEFERNIVENVFMGQTRRAQEGYYQGNLPLGYDKIP

170
170

logo

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consensus

SKHELMINQHEANIVKYIFE_CYAKGHGYRKIANALNHKGVTKKGKPFSSITYILANPFYIGKIQFAKYKDISEKRRKGLNDK
SKHELMINQHEANIVKYIFE_CYAKGHGYRKIANALNHKGVTKKGKPFSSITYILANPFYIGKIQFAKYKDISEKRRKGLNDK
SKHELMINQHEANIVKYIFE_CYAKGHGYRKIANALNHKGVTKKGKPFSSITYILANPFYIGKIQFAKYKDISEKRRKGLNDK

255
255

logo

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P V I A E G K H S P I I N Q D L W D K V Q M R K K Q V S Q K P Q V H G K G T N L L T G I I H C P Q C G A P M A A S N T T N T L K D G T K K R I R Y Y S C S N F R N K G S K
 P V I A E G K H S P I I N Q D L W D K V Q M R K K Q V S Q K P Q V H G K G T N L L T G I I H C P Q C G A P M A A S N T T N T L K D G T K K R I R Y Y S C S N F R N K G S K
 P V I A E G K H S P I I N Q D L W D K V Q M R K K Q V S Q K P Q V H G K G T N L L T G I I H C P Q C G A P M A A S N T T N T L K D G T K K R I R Y Y S C S N F R N K G S K

340
340

logo

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VCSANSVRADVIEDYVMKQILEIVKSDKVIQRVTVHVNQENQVDGAALHHDIA YKQQQYDEVQIKLNNL IKTIEDNPDLTSVIRP
VCSANSVRADVIEDYVMKQILEIVKSDKVIQRVTVHVNQENQVDGAALHHDIA YKQQQYDEVQIKLNNL IKTIEDNPDLTSVIRP
VCSANSVRADVIEDYVMKQILEIVKSDKVIQRVTVHVNQENQVDGAALHHDIA YKQQQYDEVQIKLNNL IKTIEDNPDLTSVIRP

425
425

logo

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SIQKYEKQLNDITNQINQLKNQQNEDK¹FLDAKEISKLLQHFHDIKHIEKSRLKALYLSVIDRIDIKKDGNHKKQFYVTLKLN

510
510

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E I I K Q L F N N K Q L D E V H L S T S S L F L P Q T I Y L T I
 E I I K Q L F N N K Q L D E V H L S T S S L F L P Q T I Y L T I
 E I I K Q L F N N K Q L D E V H L S T S S L F L P Q T I Y L T I

542
542

 non conserved
 similar
 $\geq 50\%$ conserved