

logo

NTDB id 122329 EQ80 RS00200 WP 001186608.1
NTDB id 11 SA RS00430 WP 001186602.1
consensus

85
85

logo

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consensus

RLARNTSDLKIVEDLHKQNEFFSLSERMEVNTSSGKMLQLASFSEFERNIVENVFMGQTRRAQEGYYQGNLPLGYDKIP
 RLARNTSDLLKIVEDLHKQNEFFSLSERMEVNTSSGKMLQLILASFSEFERNNIVENVFMGQTRRAQEGYYQGNLPLGYDKIP
 RLARNTSDLLKIVEDLHKQNEFFSLSERMEVNTSSGKMLQLILASFSEFERNNIVENVFMGQTRRAQEGYYQGNLPLGYDKIP

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170

logo

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consensus

SKHELMINQHEANIVKYIFE_CYAKGHGYRKIANALNHKGVTKKGKPFSSITYILANPFYIGKIQFAKYKDISEKRRKGLNDK
SKHELMINQHEANIVKYIFE_CYAKGHGYRKIANALNHKGVTKKGKPFSSITYILANPFYIGKIQFAKYKDISEKRRKGLNDK
SKHELMINQHEANIVKYIFE_CYAKGHGYRKIANALNHKGVTKKGKPFSSITYILANPFYIGKIQFAKYKDISEKRRKGLNDK

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255

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PVIAEGKHSPINQDLWDKQMRKKQVSQKPQVHGKGTNLLTGIHCPQCGAPMAASNTTNTLKDGTKKRIRYYSCSNFRNKGSK
PVIAEGKHSPINQDLWDKQMRKKQVSQKPQVHGKGTNLLTGIHCPQCGAPMAASNTTNTLKDGTKKRIRYYSCSNFRNKGSK
PVIAEGKHSPINQDLWDKQMRKKQVSQKPQVHGKGTNLLTGIHCPQCGAPMAASNTTNTLKDGTKKRIRYYSCSNFRNKGSK

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340

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VCSANSVRADVIEDYVMKQILEIVKSDKVIQRVTVHVNQENQVDGAALHHDIA YKQQQYDEVQIKLNNL IKTI EDNPD L TSVIRP
VCSANSVRADVIEDYVMKQILEIVKSDKVIQRVTVHVNQENQVDGAALHHDIA YKQQQYDEVQIKLNNL IKTI EDNPD L TSVIRP
VCSANSVRADVIEDYVMKQILEIVKSDKVIQRVTVHVNQENQVDGAALHHDIA YKQQQYDEVQIKLNNL IKTI EDNPD L TSVIRP

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425

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SIQKYEKQLNDITNQINQLKNQQNEDK¹FLDAKEISKLLQHFHDIKHIEKSRLKALYLSVIDRIDIKKDGNHKKQFYVTLKLN

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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 L 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 L 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 L Y L T I

542
542

 non conserved
 similar
 $\geq 50\%$ conserved