

logo

NTDB id 1046874 AB2Q58 RS01525 WP 014930816.1  
NTDB id 11 SA RS00430 WP 001186602.1  
consensus

MQQLKTKRVGIYVRVSTEMQSTEGYSIDGQINQIKEYCDFHHFEVKDIYADRGISGKSMNRPELQRI LKDAKEGYIDCVM  
.....  
MQQLKTKRVGIYVRVSTEMQSTEGYSIDGQINQIKEYCDFHHFEVKDIYADRGISGKSMNRPELQRI LKDAKEGYIDCVM  
\*\*\*\*\*

0  
80

logo

NTDB id 1046874 AB2Q58 RS01525 WP 014930816.1  
NTDB id 11 SA RS00430 WP 001186602.1  
consensus

VYKTNRLARNTS DLLKIVEDLHKQNV EFFSLSERMEVNTSSGKMLMLQILASFSEFERNNIVENVFMGQTRRAQEGYYQGN  
.....  
VYKTNRLARNTS DLLKIVEDLHKQNV EFFSLSERMEVNTSSGKMLMLQILASFSEFERNNIVENVFMGQTRRAQEGYYQGN  
\*\*\*\*\*

0  
160

logo

NTDB id 1046874 AB2Q58 RS01525 WP 014930816.1  
NTDB id 11 SA RS00430 WP 001186602.1  
consensus

LPLGYDKIPNSKHELMINQHEANIVKYIFESYAKGHGYRKI ANALNHKGYVTKKGKPFSSITYILANPFYIGKIQFAK  
.....  
LPLGYDKIPNSKHELMINQHEANIVKYIFESYAKGHGYRKI ANALNHKGYVTKKGKPFSSITYILANPFYIGKIQFAK  
\*\*\*\*\*

0  
240

logo

NTDB id 1046874 AB2Q58 RS01525 WP 014930816.1  
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consensus

YKDWSEKRRKGLNDKPVIAEGKHSPINQDLWDKVMRKKQVSQKPQVHGKGTNLLTGIHCPQCGAPMAASNTTNTLKD  
.....  
YKDWSEKRRKGLNDKPVIAEGKHSPINQDLWDKVMRKKQVSQKPQVHGKGTNLLTGIHCPQCGAPMAASNTTNTLKD  
\*\*\*\*\*!!!!!!

12  
320

logo

NTDB id 1046874 AB2Q58 RS01525 WP 014930816.1  
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consensus

GTKKRIRYYSCFQFRNKGASVCHANSIRADQAEQFVAERTKETVQHPQTIKEVNSSST.....  
.....  
GTKKRIRYYSCFQFRNKGASVCHANSIRADQAEQFVAERTKETVQHPQTIKEVNSSST.....  
GTKKRIRYYSCSIFRNKGSKVCANSVRADVIDEYVMKQILEIVKSDKVIQRVVIHVNQENQVDGAALHHDIA YKQQQYD  
!!!!!!!!!!!! \*!!!!!! !!\*!!!!\*!!!! !\*! \*!!\*\* \*!!\*\* \*!!!!!!

69  
400

logo

NTDB id 1046874 AB2Q58 RS01525 WP 014930816.1  
NTDB id 11 SA RS00430 WP 001186602.1  
consensus

EVQIKLNNLIKTI EDNPDLT SVIRPSIQKYEKQLNDITNQINQLKNQQNEDKPLFDAKEISKLLQHFHDIKHIEKSRLK  
.....  
EVQIKLNNLIKTI EDNPDLT SVIRPSIQKYEKQLNDITNQINQLKNQQNEDKPLFDAKEISKLLQHFHDIKHIEKSRLK  
\*\*\*\*\*

69  
480

logo

NTDB id 1046874 AB2Q58 RS01525 WP 014930816.1  
NTDB id 11 SA RS00430 WP 001186602.1  
consensus

ALYLSVIDRIDIKKDG NHKKQFYVTLKLNNEI IKQLFNNKQLDEVHLSTSSFLPQTLTYLTI  
.....  
ALYLSVIDRIDIKKDG NHKKQFYVTLKLNNEI IKQLFNNKQLDEVHLSTSSFLPQTLTYLTI  
\*\*\*\*\*

69  
542

 non conserved  
 similar  
  $\geq 50\%$  conserved