

logo

NTDB id 116968 A412 RS15210 WP 014930816.1
NTDB id 11 SA RS00430 WP 001186602.1
consensus

MQQLKTKRVGIYVRVSTEMQSTEGYSIDGQINQIKEYCDFHHEVVDIYADRGISGKSMNRPELQRI LKDAKEGYIDCVMVYKTN
.....
MQQLKTKRVGIYVRVSTEMQSTEGYSIDGQINQIKEYCDFHHEVVDIYADRGISGKSMNRPELQRI LKDAKEGYIDCVMVYKTN

0
85

logo

NTDB id 116968 A412 RS15210 WP 014930816.1
NTDB id 11 SA RS00430 WP 001186602.1
consensus

RLARNTS DLLKIVEDLHKQNV E F F S L S E R M E V N T S S G K L M L Q I L A S F S E F E R N N I V E N V F M G Q T R R A Q E G Y Y Q G N L P L G Y D K I P N
.....
RLARNTS DLLKIVEDLHKQNV E F F S L S E R M E V N T S S G K L M L Q I L A S F S E F E R N N I V E N V F M G Q T R R A Q E G Y Y Q G N L P L G Y D K I P N

0
170

logo

NTDB id 116968 A412 RS15210 WP 014930816.1
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consensus

SKHELMINQHEANIVKYIFESYAKGHGYRKIANALNHKGYVTKKGKPFSSITYILANPFYIGKIQFAKYKDWSEKRRKGLNDK
.....
SKHELMINQHEANIVKYIFESYAKGHGYRKIANALNHKGYVTKKGKPFSSITYILANPFYIGKIQFAKYKDWSEKRRKGLNDK

0
255

logo

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consensus

PVIAEGKHSPINQDLWDKVMRKKQVSQKPVHKGKTNLLTGIIHCPQCGAPMAASNTTNTLKDGTKKRIRYYSCSNFRNKGSK
.....
PVI AEGKHSPINQDLWDKVMRKKQVSQKPVHKGKTNLLTGIIHCPQCGAPMAASNTTNTLKDGTKKRIRYYSCSNFRNKGSK

32
340

logo

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consensus

VCHANSYRADVIEQYVMKRLLETVQSPQVQRVWTSV
.....
VCHANSYRADVIEQYVMKRLLETVQSPQVQRVWTSV

69
425

logo

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consensus

SIQKYEKQLNDITNQINQLKNQQNEDKPLFDAKEISKLLQHFHDIKHIEKSRLKALYLSVIDRIDIKKDGNHKKQFYVTLKLNN
.....
SIQKYEKQLNDITNQINQLKNQQNEDKPLFDAKEISKLLQHFHDIKHIEKSRLKALYLSVIDRIDIKKDGNHKKQFYVTLKLNN

69
510

logo

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consensus

EIIKQLFNNKQLDEVHLSSTSLFLPQTLYLTI
.....
EIIKQLFNNKQLDEVHLSSTSLFLPQTLYLTI

69
542

non conserved
 similar
 $\geq 50\%$ conserved