

logo

NTDB id 420221 GYN04 RS01345 WP 003687618.1  
NTDB id 1129 NGFG RS01435 WP 003687618.1  
consensus

MSENKQNEVLTGYEQLKRRNRRRLVTASSLVAASCILLAAALSSDPADSNPAPQAGETGATESQTANTAQTTPALKSAAEN  
MSENKQNEVLTGYEQLKRRNRRRLVTASSLVAASCILLAAALSSDPADSNPAPQAGETGATESQTANTAQTTPALKSAAEN  
MSENKQNEVLTGYEQLKRRNRRRLVTASSLVAASCILLAAALSSDPADSNPAPQAGETGATESQTANTAQTTPALKSAAEN  
!!

80  
80

logo

NTDB id 420221 GYN04 RS01345 WP 003687618.1  
NTDB id 1129 NGFG RS01435 WP 003687618.1  
consensus

GETAADKPQDLAGEDKPSAADSEISEPENVGAPLVLINDRLEDSNIKGLEESEKLQQAETAKTEPKQAKQRAAEKVSATA  
GETAADKPQDLAGEDKPSAADSEISEPENVGAPLVLINDRLEDSNIKGLEESEKLQQAETAKTEPKQAKQRAAEKVSATA  
GETAADKPQDLAGEDKPSAADSEISEPENVGAPLVLINDRLEDSNIKGLEESEKLQQAETAKTEPKQAKQRAAEKVSATA  
!!

160  
160

logo

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consensus

DSTDTVAVEKPKRSAEPKPKQAERTAEAKPKAKETKTAEKVADKPKTAAEKTTPDTAKSDSAVKEAKKADKAEKGGKTAEK  
DSTDTVAVEKPKRSAEPKPKQAERTAEAKPKAKETKTAEKVADKPKTAAEKTTPDTAKSDSAVKEAKKADKAEKGGKTAEK  
DSTDTVAVEKPKRSAEPKPKQAERTAEAKPKAKETKTAEKVADKPKTAAEKTTPDTAKSDSAVKEAKKADKAEKGGKTAEK  
!!

240  
240

logo

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consensus

DRSDGKKHETAQKTDKADKTKTAEKEKSGKAGKAAIQAGYAEKERALSQRKMKAAAGIDSTITEIMTDNGKVYRVKSSN  
DRSDGKKHETAQKTDKADKTKTAEKEKSGKAGKAAIQAGYAEKERALSQRKMKAAAGIDSTITEIMTDNGKVYRVKSSN  
DRSDGKKHETAQKTDKADKTKTAEKEKSGKAGKAAIQAGYAEKERALSQRKMKAAAGIDSTITEIMTDNGKVYRVKSSN  
!!

320  
320

logo

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consensus

YKNARDAERDLNKL RVHGIAGQVTNE  
YKNARDAERDLNKL RVHGIAGQVTNE  
YKNARDAERDLNKL RVHGIAGQVTNE  
!!

346  
346

⊠ non conserved  
⊠ similar  
⊠ ≥ 50% conserved