

logo

NTDB id 328031 EGH15 RS01645 WP 003690743.1  
NTDB id 1129 NGFG RS01435 WP 003687618.1  
consensus

MSENKQNEVLTGYEQLKRRNRRRLVTASSLVAASCILLAAALSSDPADSNPAPQAGETGATESQTANTAQTTPALKSAAEN  
MSENKQNEVLTGYEQLKRRNRRRLVTASSLVAASCILLAAALSSDPADSNPAPQAGETGATESQTANTAQTTPALKSAAEN  
MSENKQNEVLTGYEQLKRRNRRRLVTASSLVAASCILLAAALSSDPADSNPAPQAGETGATESQTANTAQTTPALKSAAEN  
!!

80  
80

logo

NTDB id 328031 EGH15 RS01645 WP 003690743.1  
NTDB id 1129 NGFG RS01435 WP 003687618.1  
consensus

GETAADKPQDLAGEDKPSAADSEISEPENVGAPLVLINDRLEDSNIKGLEESEKLLQAETAETKTEPKQAKQRAAEKVSATA  
GETAADKPQDLAGEDKPSAADSEISEPENVGAPLVLINDRLEDSNIKGLEESEKLLQAETAETKTEPKQAKQRAAEKVSATA  
GETAADKPQDLAGEDKPSAADSEISEPENVGAPLVLINDRLEDSNIKGLEESEKLLQAETAETKTEPKQAKQRAAEKVSATA  
!!

160  
160

logo

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consensus

DSTDTVAVEKPKR<sup>†</sup>AEPKPQKAERTAEAKPKAKETKTAEKVADKPKTAAEKTTPDTAKSDSAVKEAKKADKAEGKKTAEK  
DSTDTVAVEKPKR<sup>†</sup>AEPKPQKAERTAEAKPKAKETKTAEKVADKPKTAAEKTTPDTAKSDSAVKEAKKADKAEGKKTAEK  
DSTDTVAVEKPKR<sup>†</sup>AEPKPQKAERTAEAKPKAKETKTAEKVADKPKTAAEKTTPDTAKSDSAVKEAKKADKAEGKKTAEK  
!!

240  
240

logo

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consensus

DRSDGKKHETAQKTDKADKTKTAEKEKSGKAGKAAIQAGYAEKERALSQRKMKAAAGIDSTITEIMTDNGKVYRVKSSN  
DRSDGKKHETAQKTDKADKTKTAEKEKSGKAGKAAIQAGYAEKERALSQRKMKAAAGIDSTITEIMTDNGKVYRVKSSN  
DRSDGKKHETAQKTDKADKTKTAEKEKSGKAGKAAIQAGYAEKERALSQRKMKAAAGIDSTITEIMTDNGKVYRVKSSN  
!!

320  
320

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NTDB id 328031 EGH15 RS01645 WP 003690743.1  
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consensus

YKNARDAERDLNKLRVHGIAGQVTNE  
YKNARDAERDLNKLRVHGIAGQVTNE  
YKNARDAERDLNKLRVHGIAGQVTNE  
!!

346  
346

non conserved  
 similar  
 ≥ 50% conserved