

logo

NTDB id 398476 GIJ09 RS01785 WP 003690743.1
NTDB id 1129 NGFG RS01435 WP 003687618.1
consensus

MSENKQNEVLTGYEQLKRRNRRRLVTASSLVAASCILLAAALSSDPADSNPAPQAGETGATESQTANTAQTTPALKSAAEN
MSENKQNEVLTGYEQLKRRNRRRLVTASSLVAASCILLAAALSSDPADSNPAPQAGETGATESQTANTAQTTPALKSAAEN
MSENKQNEVLTGYEQLKRRNRRRLVTASSLVAASCILLAAALSSDPADSNPAPQAGETGATESQTANTAQTTPALKSAAEN
!!

80
80

logo

NTDB id 398476 GIJ09 RS01785 WP 003690743.1
NTDB id 1129 NGFG RS01435 WP 003687618.1
consensus

GETAADKPQDLAGEDKPSAADSEISEPENVGAPLVLINDRLEDSNIKGLEESEKLQQAETAKTEPKQAKQRAAEKVSATA
GETAADKPQDLAGEDKPSAADSEISEPENVGAPLVLINDRLEDSNIKGLEESEKLQQAETAKTEPKQAKQRAAEKVSATA
GETAADKPQDLAGEDKPSAADSEISEPENVGAPLVLINDRLEDSNIKGLEESEKLQQAETAKTEPKQAKQRAAEKVSATA
!!

160
160

logo

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consensus

DSTDTVAVEKPKR₁AEPKPQKAERTAEAKPKAKETKTAEKVADKPKTAAEKTTPDTAKSDSAVKEAKKADKAEGKKTAEK
DSTDTVAVEKPKR₁AEPKPQKAERTAEAKPKAKETKTAEKVADKPKTAAEKTTPDTAKSDSAVKEAKKADKAEGKKTAEK
DSTDTVAVEKPKR₁AEPKPQKAERTAEAKPKAKETKTAEKVADKPKTAAEKTTPDTAKSDSAVKEAKKADKAEGKKTAEK
!!

240
240

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consensus

DRSDGKKHETAQKTDKADKTKTAEKEKSGKAGKAAIQAGYAEKERALSQRKMKAAAGIDSTITEIMTDNGKVYRVKSSN
DRSDGKKHETAQKTDKADKTKTAEKEKSGKAGKAAIQAGYAEKERALSQRKMKAAAGIDSTITEIMTDNGKVYRVKSSN
DRSDGKKHETAQKTDKADKTKTAEKEKSGKAGKAAIQAGYAEKERALSQRKMKAAAGIDSTITEIMTDNGKVYRVKSSN
!!

320
320

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NTDB id 398476 GIJ09 RS01785 WP 003690743.1
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consensus

YKNARDAERDLNKLRVHGIAGQVTNE
YKNARDAERDLNKLRVHGIAGQVTNE
YKNARDAERDLNKLRVHGIAGQVTNE
!!

346
346

non conserved
 similar
 ≥ 50% conserved