

logo

NTDB id 328373 EGH17 RS02025 WP 003690743.1
NTDB id 1129 NGFG RS01435 WP 003687618.1
consensus

MSENKQNEVLTGYEQLKRRNRRRLVTASSLVAASCILLAAALSSDPADSNPAPQAGETGATESQTANTAQTTPALKSAAEN
MSENKQNEVLTGYEQLKRRNRRRLVTASSLVAASCILLAAALSSDPADSNPAPQAGETGATESQTANTAQTTPALKSAAEN
MSENKQNEVLTGYEQLKRRNRRRLVTASSLVAASCILLAAALSSDPADSNPAPQAGETGATESQTANTAQTTPALKSAAEN
!!

80
80

logo

NTDB id 328373 EGH17 RS02025 WP 003690743.1
NTDB id 1129 NGFG RS01435 WP 003687618.1
consensus

GETAADKPQDLAGEDKPSAADSEISEPENVGAPLVLINDRLEDSNIKGLEESEKLLQAETAETKTEPKQAKQRAAEKVSATA
GETAADKPQDLAGEDKPSAADSEISEPENVGAPLVLINDRLEDSNIKGLEESEKLLQAETAETKTEPKQAKQRAAEKVSATA
GETAADKPQDLAGEDKPSAADSEISEPENVGAPLVLINDRLEDSNIKGLEESEKLLQAETAETKTEPKQAKQRAAEKVSATA
!!

160
160

logo

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NTDB id 1129 NGFG RS01435 WP 003687618.1
consensus

DSTDTVAVEKPKR[†]SAEPKPQKAERTAEAKPKAKETKTAEKVADKPKTAAEKTTPDTAKSDSAVKEAKKADKAEGKKTAEK
DSTDTVAVEKPKR[†]SAEPKPQKAERTAEAKPKAKETKTAEKVADKPKTAAEKTTPDTAKSDSAVKEAKKADKAEGKKTAEK
DSTDTVAVEKPKR[†]SAEPKPQKAERTAEAKPKAKETKTAEKVADKPKTAAEKTTPDTAKSDSAVKEAKKADKAEGKKTAEK
!!

240
240

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consensus

DRSDGKKHETAQKTDKADKTKTAEKEKSGKAGKAAIQAGYAEKERALSQRKMKAAAGIDSTITEIMTDNGKVYRVKSSN
DRSDGKKHETAQKTDKADKTKTAEKEKSGKAGKAAIQAGYAEKERALSQRKMKAAAGIDSTITEIMTDNGKVYRVKSSN
DRSDGKKHETAQKTDKADKTKTAEKEKSGKAGKAAIQAGYAEKERALSQRKMKAAAGIDSTITEIMTDNGKVYRVKSSN
!!

320
320

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consensus

YKNARDAERDLNKLRVHGIAGQVTNE
YKNARDAERDLNKLRVHGIAGQVTNE
YKNARDAERDLNKLRVHGIAGQVTNE
!!

346
346

non conserved
 similar
 ≥ 50% conserved