

logo

NTDB id 185118 AS012 RS01420 WP 003687618.1
NTDB id 1129 NGFG RS01435 WP 003687618.1
consensus

MSENKQNEVLTGYEQLKRRNRRRLVTASSLVAASCILLAAALSSDPADSNPAPQAGETGATESQTANTAQTTPALKSAAEN
MSENKQNEVLTGYEQLKRRNRRRLVTASSLVAASCILLAAALSSDPADSNPAPQAGETGATESQTANTAQTTPALKSAAEN
MSENKQNEVLTGYEQLKRRNRRRLVTASSLVAASCILLAAALSSDPADSNPAPQAGETGATESQTANTAQTTPALKSAAEN
!!

80
80

logo

NTDB id 185118 AS012 RS01420 WP 003687618.1
NTDB id 1129 NGFG RS01435 WP 003687618.1
consensus

GETAADKPQDLAGEDKPSAADSEISEPENVGAPLVLINDRLEDSNIKGLEESEKLLQAETAETKTEPKQAKQRAAEKVSATA
GETAADKPQDLAGEDKPSAADSEISEPENVGAPLVLINDRLEDSNIKGLEESEKLLQAETAETKTEPKQAKQRAAEKVSATA
GETAADKPQDLAGEDKPSAADSEISEPENVGAPLVLINDRLEDSNIKGLEESEKLLQAETAETKTEPKQAKQRAAEKVSATA
!!

160
160

logo

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consensus

DSTDTVAVEKPKRSAEPKPKQAERTAEAKPKAKETKTAEKVADKPKTAAEKTTPDTAKSDSAVKEAKKADKAEKGGKTAEK
DSTDTVAVEKPKRSAEPKPKQAERTAEAKPKAKETKTAEKVADKPKTAAEKTTPDTAKSDSAVKEAKKADKAEKGGKTAEK
DSTDTVAVEKPKRSAEPKPKQAERTAEAKPKAKETKTAEKVADKPKTAAEKTTPDTAKSDSAVKEAKKADKAEKGGKTAEK
!!

240
240

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consensus

DRSDGKKHETAQKTDKADKTKTAEKEKSGKAGKAAIQAGYAEKERALSQRKMKAAAGIDSTITEIMTDNGKVYRVKSSN
DRSDGKKHETAQKTDKADKTKTAEKEKSGKAGKAAIQAGYAEKERALSQRKMKAAAGIDSTITEIMTDNGKVYRVKSSN
DRSDGKKHETAQKTDKADKTKTAEKEKSGKAGKAAIQAGYAEKERALSQRKMKAAAGIDSTITEIMTDNGKVYRVKSSN
!!

320
320

logo

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consensus

YKNARDAERDLNKLRVHGIAGQVTNE
YKNARDAERDLNKLRVHGIAGQVTNE
YKNARDAERDLNKLRVHGIAGQVTNE
!!

346
346

⊠ non conserved
⊠ similar
⊠ ≥ 50% conserved