

logo

NTDB id 87100 BsBEST3145 RS05670 WP 003233100.1
NTDB id 119 BSU 10630 NP 388944.2
consensus

MNIPKPADSTWTDDQWNAIVSTGQDILVAAAAGSGKTAVLVERMIRKITAEEENPIDVDRLLVVTFTNASAAEMKHRIA
EA
MNIPKPADSTWTDDQWNAIVSTGQDILVAAAAGSGKTAVLVERMIRKITAEEENPIDVDRLLVVTFTFTNASAAEMKHRIA
EA
!!

80
80

logo

NTDB id 87100 BsBEST3145 RS05670 WP 003233100.1
NTDB id 119 BSU 10630 NP 388944.2
consensus

LEKELVQRPGLHRRQLSLLNRASISTLHSFCLQVLKYYLIDLDPGFRIADQTEGELIGDEVLDLFEDEYAKGEKA
LEKELVQRPGLHRRQLSLLNRASISTLHSFCLQVLKYYLIDLDPGFRIADQTEGELIGDEVLDLFEDEYAKGEKA
LEKELVQRPGLHRRQLSLLNRASISTLHSFCLQVLKYYLIDLDPGFRIADQTEGELIGDEVLDLFEDEYAKGEKA
!!

160
160

logo

NTDB id 87100 BsBEST3145 RS05670 WP 003233100.1
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consensus

FFELVDRYTDDRHDLDLQFLVKQVVEYSRSHNPPEAWLESFVHLYDVSEKSAIEELPFYQYVKEDIAMVLNGAKEKLLRA
FFELVDRYTDDRHDLDLQFLVKQVVEYSRSHNPPEAWLESFVHLYDVSEKSAIEELPFYQYVKEDIAMVLNGAKEKLLRA
FFELVDRYTDDRHDLDLQFLVKQVVEYSRSHNPPEAWLESFVHLYDVSEKSAIEELPFYQYVKEDIAMVLNGAKEKLLRA
!!

240
240

logo

NTDB id 87100 BsBEST3145 RS05670 WP 003233100.1
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consensus

LELTKAPGGPAPRADNFLDDLAQIDELIQHQDDFSELYKRVPVAVSFKRAKAVKGDEFDPALLDEATDLRNGAKKLLLEKLK
LELTKAPGGPAPRADNFLDDLAQIDELIQHQDDFSELYKRVPVAVSFKRAKAVKGDEFDPALLDEATDLRNGAKKLLLEKLK
LELTKAPGGPAPRADNFLDDLAQIDELIQHQDDFSELYKRVPVAVSFKRAKAVKGDEFDPALLDEATDLRNGAKKLLLEKLK
!!

320
320

logo

NTDB id 87100 BsBEST3145 RS05670 WP 003233100.1
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consensus

TDYFTRSPEQHLKSLAEMKPVETLVQLVISYGRFEAAKQEKSIIDFSDLEHYCLAILTAEENDKGEREPSEAAIFYQE
Q
TDYFTRSPEQHLKSLAEMKPVETLVQLVISYGRFEAAKQEKSIIDFSDLEHYCLAILTAEENDKGEREPSEAAIFYQE
Q
TDYFTRSPEQHLKSLAEMKPVETLVQLVISYGRFEAAKQEKSIIDFSDLEHYCLAILTAEENDKGEREPSEAAIFYQE
Q
!!

400
400

logo

NTDB id 87100 BsBEST3145 RS05670 WP 003233100.1
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consensus

FHEVLVDEYQDTNLVQESILQLVTSGPEETGNLFMVGDKQSIYRFRLAEPFLFLSKYKRFTESGEGTGRKIDLNKNFRS
FHEVLVDEYQDTNLVQESILQLVTSGPEETGNLFMVGDKQSIYRFRLAEPFLFLSKYKRFTESGEGTGRKIDLNKNFRS
FHEVLVDEYQDTNLVQESILQLVTSGPEETGNLFMVGDKQSIYRFRLAEPFLFLSKYKRFTESGEGTGRKIDLNKNFRS
!!

480
480

logo

NTDB id 87100 BsBEST3145 RS05670 WP 003233100.1
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consensus

RADILDSTNFLFKQLMGGKIGVDYDEQAELKLGAAYPDNDDETETELLLIDNAEDTDASEEAELETQFEAKIAKEIR
RADILDSTNFLFKQLMGGKIGVDYDEQAELKLGAAYPDNDDETETELLLIDNAEDTDASEEAELETQFEAKIAKEIR
RADILDSTNFLFKQLMGGKIGVDYDEQAELKLGAAYPDNDDETETELLLIDNAEDTDASEEAELETQFEAKIAKEIR
!!

560
560

NTDB id 87100 BsBEST3145 RS05670 WP 003233100.1
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consensus

640
640

NTDB id 87100 BsBEST3145 RS05670 WP 003233100.1
NTDB id 119 BSU 10630 NP 388944.2
consensus

720
720

NTDB id 87100 BsBEST3145 RS05670 WP 003233100.1
NTDB id 119 BSU 10630 NP 388944.2
consensus

800
800

NTDB id 87100 BsBEST3145 RS05670 WP 003233100.1
NTDB id 119 BSU 10630 NP 388944.2
consensus

880
880

NTDB id 87100 BsBEST3145 RS05670 WP 003233100.1
NTDB id 119 BSU 10630 NP 388944.2
consensus

960
960

NTDB id 87100 BsBEST3145 RS05670 WP 003233100.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



1040
1040

NTDB id 87100 BsBEST3145 RS05670 WP 003233100.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



1120
1120

logo

NTDB id 87100 BsBEST3145 RS05670 WP 003233100.1
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consensus



1200
1200

logo

NTDB id 87100 BsBEST3145 RS05670 WP 003233100.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



1232
1232

-  non conserved
-  similar
-  ≥ 50% conserved