

logo

NTDB id 81292 KI220 RS10560 WP 212791793.1
NTDB id 119 BSU 10630 NP 388944.2
consensus

MEISKP^{AD}SGSTNDDQKNAIVSSGRDILVAAAAGSGKTAVLVERMIRKITDQENHPVDVDRLLVFTFTNASAAEMKHRIEAELEKEL
 MNIPKPADSTWDDQWN AIVSSGRDILVAAAAGSGKTAVLVERMIRKITDQENHPVDVDRLLVFTFTNASAAEMKHRIEAELEKEL

85
85

logo

NTDB id 81292 KI220 RS10560 WP 212791793.1
NTDB id 119 BSU 10630 NP 388944.2
consensus

VQRPGSLHRRQLALLNRASTLHSFCLQVIRKYYYLIDDPGFRIADQTGELTGDEVLDLFEDEYKKQEPKAFFELVDRYTT
AENPGSLHRRRLAALLNKASISTLHSFCLQVIRKYYYLLIDVPDGFRIADQTGELLGDEVLDLFEDEYKKQDPKAFFELVDRYTT
VQRPGSLHIRRQLSLLNRASISTLHSFCLQVLRKYYYLLIDDPGFRIADQTGELTGDEVLDLFEDEYAKGEKKAFFELVDRYTT

* * * * *

170
170

logo

NTDB id 81292 KI220 RS10560 WP 212791793.1
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consensus

DRHDLDLQHFLVKRWVEYFSRSHPNPEAGWLHTFVHLVDVSSKTKYAELEPFYSYVKEDIALVLENGMRQKLTRLALDLTKQSGGPAPRAE
DRHDLDLQHFLVKRWVEYFSRSHPNPEAGWLHTFVHLVDVSSKTKYAELEPFYSYVKEDIALVLENGMRQKLTRLALDLTKQSGGPAPRAE
DRHDLDLQLFLVKRWVEYFSRSHPNPEAWLESFVHLYDVSEKSAIELEPFYQYVKEDIAMVLNGAEKLLRLALELTKAPGGPAPRAD

255
255

logo

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340
340

logo

NTDB id 81292 KI220 RS10560 WP 212791793.1
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consensus

425
425

logo

NTDB id 81292 KI220 RS10560 WP 212791793.1
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consensus

509
510

logo

NTDB id 81292 KI220 RS10560 WP 212791793.1
NTDB id 119 BSU 10630 NP 388944.2
consensus

[illegible]593
595

logo

NTDB id 81292 KI220 RS10560 WP 212791793.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



non conserved
 similar
 ≥ 50% conserved