

RYTSDRSDLDLQRMVLQVYTYSRSHPNPEKWLDSLVEAYDVGK.TIEDLVYASYLLEDVKFQLETAEQHIRKATELAMLPDGPA
RYTSDRSDLDLQFLVKQVYEYSRSHPNPEAWLSFSVHLVDVSEKSAIEELPFYQYVKEDIAMVLNGAKEKLRLALELTAKAPGGPA
* * * * * * * * * * * * * * *

[illegible]

QLVTSGSESTGNLFMVGDVKQSIYRFRLAEPGLFLSKYKRFTQSGGTGRKIDLNKNFRSRHEYLDS TNFLT FKQLMGSKYEYDY
 KFTVKDSESEGNLFMVGDVKQSIYRFRLAEPGLFLGKYKRFTQEGLGGGMKIDLAKNFRSRHEVL AGTNFLT FKQLMGEEYGEIDY
 QLVTSGP EETGNLFMVGDVKQSIYRFRLAEP LFLSKYKRFTESGEGTGRKIDLNKNFRSRADILDS TNFLT FKQLMGKIGEV DY
 * * * * *

[illegible]594
589

logo

NTDB id 129698 BF26 RS09265 WP 000970458.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



679
674

logo

NTDB id 129698 BF26 RS09265 WP 000970458.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



764
757

logo

NTDB id 129698 BF26 RS09265 WP 000970458.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



849
842

logo

NTDB id 129698 BF26 RS09265 WP 000970458.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



934
927

logo

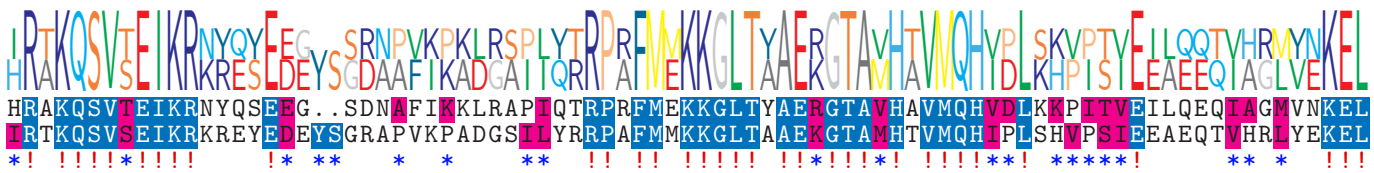
NTDB id 129698 BF26 RS09265 WP 000970458.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



1017
1009

logo

NTDB id 129698 BF26 RS09265 WP 000970458.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



1100
1094

logo

NTDB id 129698 BF26 RS09265 WP 000970458.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



1185
1178

logo

NTDB id 129698 BF26 RS09265 WP 000970458.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



 non conserved
 similar
 ≥ 50% conserved