

RYTSDRSDDLQRMVLQVITYSRSHPNPEKMLESL VHLVDVSGKSTIELVYYSVYL EDYKMVLNTAKQKLRRATELTMLPSGPA
RYTSDRSDDDLQRMITALYTESRAHPNPEKWLDKLVEAYDVEGK.TIEDLVYASYLLEDVKFQLETAEQHIRKATELAMLPDGPA
RYTTDRHDLDLQFFLVKKQVYEYSRSHPNPEAWLESFVHLVDVSEKSAIEELPFYQYVKEDIAMVNLNGAKEKLLRLALELTKAPGGPA
* * * * * * * * * * * * * *

QLVTSGSESTGNLFMVGDVKQSIYRFRLAEPGLFLSKYKRFTQSGGTGRKIDLNKNFRSRHEYLDSNFTFKQLMGSKYEYDY
 KFTVTKDSESEGNLFMVGDVKQSIYRFRLAEPGLFLGKYKRFTQEGLGGGMKIDLAKNFRSRHEVLAGTNFTFKQLMGEEVGEIDY
 QLVTSGPEETGNLFMVGDVKQSIYRFRLAEPGLFLSKYKRFTESGEGTGRKIDLNKNFRSRADILDSNFTLFKQLMGKIGEVYDY
 * * * * *

[illegible]

594
589

logo

NTDB id 121691 DJ48 RS15815 WP 000970460.1
NTDB id 119 BSU 10630 NP 388944.2
consensus

679
674

logo

NTDB id 121691 DJ48 RS15815 WP 000970460.1
NTDB id 119 BSU 10630 NP 388944.2
consensus

764
757

logo

NTDB id 121691 DJ48 RS15815 WP 000970460.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



849
842

logo

NTDB id 121691 DJ48 RS15815 WP 000970460.1
NTDB id 119 BSU 10630 NP 388944.2
consensus

934
927

logo

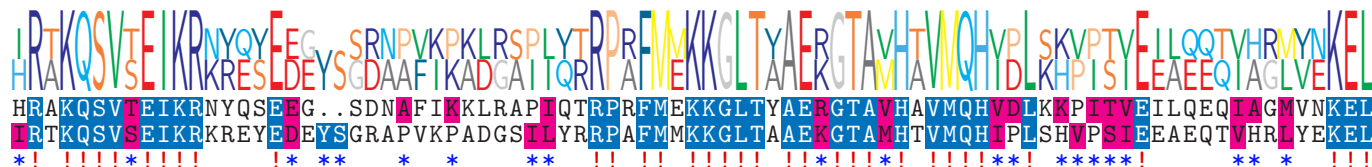
NTDB id 121691 DJ48 RS15815 WP 000970460.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



1017
1009

logo

NTDB id 121691 DJ48 RS15815 WP 000970460.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



1100
1094

logo

NTDB id 121691 DJ48 RS15815 WP 000970460.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



1185
1178

logo

NTDB id 121691 DJ48 RS15815 WP 000970460.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



non conserved
 similar
 ≥ 50% conserved