

[illegible][illegible]

QLVTSGSESTGNLFMVGDVKQSIYRFLAEPGLFLSKYKRFTQSGGTGRKIDLNKNFRSRHEYLDS TNFLT FKQIMGKYGEYDY
 KFTVKDSESEGNLFMVGDVKQSIYRFLAEPGLFLGKYKRFTQEGLGGMKIDLAKNFRSRHEVL AGTNFLT FKQIMGEEVGEIDY
 QLVTSGPEETGNLFMVGDVKQSIYRFLAEPGLFLSKYKRFTESGEGTGRKIDLNKNFRSRADILDS TNFLT FKQLMGKIGEVYDY
 * * * * *

[illegible]

594
589

logo

NTDB id 73644 FNQ92 RS00075 WP 000970458.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



679
674

logo

NTDB id 73644 FNQ92 RS00075 WP 000970458.1
NTDB id 119 BSU 10630 NP 388944.2
consensus

764
757

logo

NTDB id 73644 FNQ92 RS00075 WP 000970458.1
NTDB id 119 BSU 10630 NP 388944.2
consensus

849
842

logo

NTDB id 73644 FNQ92 RS00075 WP 000970458.1
NTDB id 119 BSU 10630 NP 388944.2
consensus

934
927

logo

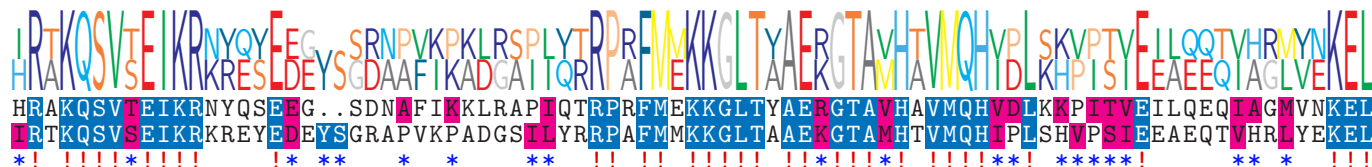
NTDB id 73644 FNQ92 RS00075 WP 000970458.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



1017
1009

logo

NTDB id 73644 FNQ92 RS00075 WP 000970458.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



1100
1094

logo

NTDB id 73644 FNQ92 RS00075 WP 000970458.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



1185
1178

logo

NTDB id 73644 FNQ92 RS00075 WP 000970458.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



 non conserved
 similar
 ≥ 50% conserved