

logo

NTDB id 92959 AK066 RS14730 WP 057079727.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



85
85

logo

NTDB id 92959 AK066 RS14730 WP 057079727.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



170
170

logo

NTDB id 92959 AK066 RS14730 WP 057079727.1
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consensus



255
255

logo

NTDB id 92959 AK066 RS14730 WP 057079727.1
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consensus



340
340

logo

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consensus



425
425

logo

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510
510

logo

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595
595

logo

NTDB id 92959 AK066 RS14730 WP 057079727.1
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SRMYQTSCTFVIGVNDGVLPARPSSENSVLEDDREWLRATIGVELSSTGRERLLDEQFLIYQALSSPSHYLYVSYSPASDTEGRSLL
SRMYQAKCTFVIGVNDGVLPARPSDESVLSEDDREWLRAGAELAEETGKERLLDEQFLIYQALSSPSHYLYVSYSPASDTEGRSLL
SRMYGTSCTFVLGANDGVLPARPDENGVLSDDDREWLRATIGVELSSSGRERLLDEHFLIYMAFSSPSDRLYVSYPIADAEGRKTL

680
680

logo

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765
765

logo

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850
850

logo

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LTKEQCVTYSYHAEERLAPRLQKEILLSSNRHYVYKEKLQKIVTRVSGILSEHAKVSGFVPIVGLGFGGQGPLPPFTFQLKNGC
 LTKEQCELFSDYDAVERLAPRLQKEILLSSNRHYVYKEKLQKIVTRVSGILSEHAKVSGFVPIVGLGFGGQGPLPPFTFQLKNGC
 * * * * *

935
935

logo

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1020
1020

logo

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PLGLDE|EQE|FKKFKMKGLLL|EDVEVRLMD|TLQSEGRSQV|NAGLKKDGSFRSDSAVWS|EKHFH|LTKHVRRTFE|EAGE|RLTS
 PLAEDAIEQEILKKFKMKGLLLEDVEAVKLMDDQTLSEGRSQVINAGLKKDGSFRSDSAVLS|EDHFH|LTKHVRRTFE|EAGER|ITS
 PLGLDEIEQEIFKKFKMKGLLLGDQEVRLMD|TLQEGRSNNI|NAGLKKDGSFRSDSAVGEK|EFDLLTKHVRRTFE|EAGE|QITD
 * * * * *

1105
1105

logo

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1169
1166

 non conserved
 similar
 $\geq 50\%$ conserved