

logo

NTDB id 240726 CF389 RS09070 WP 003155147.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



80
80

logo

NTDB id 240726 CF389 RS09070 WP 003155147.1
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consensus



160
160

logo

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consensus



240
240

logo

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320
320

logo

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400
400

logo

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480
480

logo

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560
560

logo

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consensus

KLYSSPFKVDGKTKTHRN IQYRDIVILLRSPWPAPQMEELRNQGIPVYANLTSGYFEAVEVAAALSVLKVIDNPYQDI
 KLVSPPFKVYDGGTKTHRN IQYRDIVILLRSPWPAPQMEELRNQGIPVYANLTSGYFEAVEVAAALSVLKVIDNPYQDI
 KLISPPFKVYDGGTKTHRN IQYRDIVILLRSPWPAPQMEELRNQGIPVYANLTSGYFEAVEVAAALSVLKVIDNPYQDI

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640
640

logo

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720
720

logo

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T K Y M D Y A G G M P G G K Q R Q A N L R V L Y D R A R S Y E A T A F R G L F R F L R F I E R M Q E R G D D L G T A R A L S E Q E D W R L M T I H S S K G L E
 T G Y F D Y A G G M P G G K Q R Q A N L R V L Y D R A R S Y E A T A F R G L F R F L R F I E R M Q E R G D D L G T A R A L S E Q E D V V R L M T I H S S K G L E
 T K Y M D Y V G G M P G G K Q R Q A N L R V L Y D R A R Q Y E S T A F R G L F R F L R F I E R M Q E R G D D L G T A R A L S E Q E D V V R L M T I H S S K G L E

800
800

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FPWF^TAGLGR^SFVMDLNKSYLLDKELGFGTKYIHP^QELRISYPTLP^VAMKKKMRRELLSEELRVLYVALTRAKEKLF^I
 FPVVF^TAGLGR^SFNMMDLNKSYLLDKELGFGTKYIHP^QELRISYPTLP^VAMKKKMRRELLSEELRVLYVALTRAKEKLF^I
 FPVVF^TAGLGR^NFNMMDLNKSYLLDKELGFGTKYIHP^QELRISYPTLP^VAMKKKMRRELLSEELRVLYVALTRAKEKLF^I
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880
880

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960
958

logo

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consensus

1040
1038

logo

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GGT L YRR P G F M K K G L T A E K G T A M H T V M Q H I P L T H V P T I E E A E R T V R R L Y E K E L L T E E Q Q E A I D E E V R F F H T E I G
D D S T A F M K K G L T A E K G T A M H T V M Q H I P L T H V P T I E E A E R T V R R L Y E K E L L T E E Q Q E A I D E E V R F F H T E I G

G D T L Y R R P C F M M K K G L T A A E K G T A M H T V M Q H I P L T H V P T A E E A B R T V R M L Y E K E L L T E E Q Q E A I D I E E I V R F F C T E I G K
D G S I L Y R R P A F M M K K G L T A A E K G T A M H T V M Q H I P L S H V P S T E E A B Q T V H R L Y E K E L L T E E Q K D A I D I E E I V Q F F H T E I G

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1120
1118

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-  non conserved
-  similar
-  $\geq 50\%$ conserved