

NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 180631 A2I68 RS06350 WP 235318543.1
NTDB id 30 MW RS08005 WP 000697220.1
NTDB id 14 SA RS07775 WP 000697220.1
consensus

80
0
0
0

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[illegible]

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K E N D F K H L M N K K Q G L I L F T G P T G S G K S T L M Y Q M V S Y A N K A L N L N V I S I E D P V E M Q I P G I V Q I N V N D K A G I N Y V N S F K A I L
G P K H L L N A L K Q P . . . V L I T L S A L N L I Q M D K H I L T A . . . I E D L I T Q . R S I Q L D F S Q L L R T F L
Q Q T T L L N A L K Q P Q G L I L V T G P T G S G K S I T L Y S A L N Y L N Q . M D K H I L T A E D P V E I E I D G L I Q T Q V N R S I Q L D F S Q L L R T F L 318
G P K H F Y H L M N K K Q G L I L F T G P T G V L 141
K F N D F K H L M N K K Q G L I L F S G P T G S G K S T L M Y Q M V S Y A N K A L N L N V I S I E D P V E M Q I P G I V Q I N V N D K A G I N Y V N S F K A I L 196
K F N D F K H L M N K K Q G L I L F S G P T G S G K S T L M Y Q M V S Y A N K A L N L N V I S I E D P V E M Q I P G I V Q I N V N D K A G I N Y V N S F K A I L 196
***** 196

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RCDPDVILIGEIRDKDVAKCVIQASLSGHLVLTTTLHATDCKGAILRLLEMGISVQELIQAATNLIINQRLVTTIKKQRQL
RQDPDIIMLGEIRDDEESAKMALRASQTGHVLSTLHTNDALSAYERLLQLGIQEYEIQNSLLLVIAQRLVRKLCKKCRGV 398

RCDPDVILIGEIRDKDVAKCVIQASLSGHLVLTTTLHATDCKGAILRLLEMGISVQELIQAATNLIINQRLVTTIKKQRQL 141
RCDPDVILIGEIRDKDVAKCVIQASLSGHLVLTTTLHATDCKGAILRLLEMGISVQELIQAATNLIINQRLVTTIKKQRQL 275
***** 275

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consensus

 non conserved
 similar
 $\geq 50\%$ conserved