

logo

NTDB id 123686 BSUB RS05885 WP 003244988.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



85
85

logo

NTDB id 123686 BSUB RS05885 WP 003244988.1
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consensus



170
170

logo

NTDB id 123686 BSUB RS05885 WP 003244988.1
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consensus



255
255

logo

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340
340

logo

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425
425

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510
510

logo

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595
595

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SRMYGTSCTFVLGANDGVLPARPDENGVLSDDDREWLKTI GVELSSGGRERLLDEHFLIYMAFSSPSDRLYVSYPIADAEGKTLI
SRMYGTSCTFVLGANDGVLPARPDENGVLSDDDREWLKTI GVELSSGGRERLLDEHFLIYMAFSSPSDRLYVSYPIADAEGKTLI
SRMYGTSCTFVLGANDGVLPARPDENGVLSDDDREWLKTI GVELSSGGRERLLDEHFLIYMAFSSPSDRLYVSYPIADAEGKTLI

680
680

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PSMIVKRLEELFPHHKERLLTNEPEQVSDEEQLMYVVKNSVAQSFTASQLRLWTREYDISDVVWSTYNVLMSEQDRLQSKKLFSS
PSMIVKRLEELFPHHKERLLTNEPEQVSDEEQLMYVVKNSVAQSFTASQLRLWTREYDISDVVWSTYNVLMSEQDRLQSKKLFSS
PSMIVKRLEELFPHHKERLLTNEPEQVSDEEQLMYVVKNSVAQSFTASQLRLWTREYDISDVVWSTYNVLMSEQDRLQSKKLFSS

765
765

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LFFRNEVKQLERSVSRQLYGERIQGSVSRMETFNACPFSHFASHGLHLKERQFFKLEAPDIGQLFHSSSLKLISDRLEQKLDWRDLFFRNEVKQLERSVSRQLYGERIQGSVSRMETFNACPFSHFASHGLHLKERQFFKLEAPDIGQLFHSSSLKLISDRLEQKLDWRDLFFRNEVKQLERSVSRQLYGERIQGSVSRMETFNACPFSHFASHGLHLKERQFFKLEAPDIGQLFHSSSLKLISDRLEQKLDWRDL

850
850

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LTKEQCELFSDAVERLAPKLQKEILLSSNRHYVKEKLQKIVTRVSGILSEHAKASGFVPIGLELGFGGKGPLPPLTFQLKNGC
LTKEQCELFSDAVERLAPKLQKEILLSSNRHYVKEKLQKIVTRVSGILSEHAKASGFVPIGLELGFGGKGPLPPLTFQLKNGC
LTKEQCELFSDAVERLAPKLQKEILLSSNRHYVKEKLQKIVTRVSGILSEHAKASGFVPIGLELGFGGKGPLPPLTFQLKNGC

935
935

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TMELVGRIDRVDKAESSKGLLLRIVDYKSSDKGLDLAEVYYGLALQMLTYLDLSITHSADWLGMRA TPAGVLYFHIHDPMIQSNL
TMELVGRIDRVDKAESSKGLLLRIVDYKSSDKGLDLAEVYYGLALQMLTYLDLSITHSADWLGMRA TPAGVLYFHIHDPMIQSNL
TMELVGRIDRVDKAESSKGLLLRIVDYKSSDKGLDLAEVYYGLALQMLTYLDLSITHSADWLGMRA TPAGVLYFHIHDPMIQSNL

1020
1020

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PLGLDEIEQEIFKKFKMKGLLLGDQEWRLMDTTLQEGRSNIINAGLKKDGSLSRSDSAAVGEKEFDLLTKHVRRTFQEAGEQITD
PLGLDEIEQEIFKKFKMKGLLLGDQEVVRLMDTTLQEGRSNIINAGLKKDGSLSRSDSAAVGEKEFDLLTKHVRRTFQEAGEQITD
PLGLDEIEQEIFKKFKMKGLLLGDQEVVRLMDTTLQEGRSNIINAGLKKDGSLSRSDSAAVGEKEFDLLTKHVRRTFQEAGEQITD

1105
1105

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GRVSI EPYKMKNKTPCTYCAFKSVCQFDESLEENEYRPLKAEKDKTILEWIKKEADGNEHS 1166
GRVSI EPYKMKNKTPCTYCAFKSVCQFDESLEENEYRPLKAEKDKTILEWIKKEADGNEHS 1166

1166
1166

 non conserved
 similar
 $\geq 50\%$ conserved