

logo

NTDB id 73411 FNP71 RS05950 WP 003244988.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



85
85

logo

NTDB id 73411 FNP71 RS05950 WP 003244988.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



170
170

logo

NTDB id 73411 FNP71 RS05950 WP 003244988.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



255
255

logo

NTDB id 73411 FNP71 RS05950 WP 003244988.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



340
340

logo

NTDB id 73411 FNP71 RS05950 WP 003244988.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



425
425

logo

NTDB id 73411 FNP71 RS05950 WP 003244988.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



510
510

logo

NTDB id 73411 FNP71 RS05950 WP 003244988.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



595
595

logo

NTDB id 73411 FNP71 RS05950 WP 003244988.1
NTDB id 120 BSU 10620 NP 388943.2
consensus

SRMYGTSCTFVLGANDGVLPARPDENGVLSDDDREWLKTI GVELSSGGRERLLDEHFLIYMAFSSPSDRLYVSYPIADAEGKTL

680
680

logo

NTDB id 73411 FNP71 RS05950 WP 003244988.1
NTDB id 120 BSU 10620 NP 388943.2
consensus

PSMIVKRLEELFPHHKERLLTNEPEQVSDEEQLMYVVKNSVAQSFTASQLRLWTREYDISDVVWSTYNVLMSEQDRLQSKKLFSS
PSMIVKRLEELFPHHKERLLTNEPEQVSDEEQLMYVVKNSVAQSFTASQLRLWTREYDISDVVWSTYNVLMSEQDRLQSKKLFSS
PSMIVKRLEELFPHHKERLLTNEPEQVSDEEQLMYVVKNSVAQSFTASQLRLWTREYDISDVVWSTYNVLMSEQDRLQSKKLFSS

765
765

logo

NTDB id 73411 FNP71 RS05950 WP 003244988.1
NTDB id 120 BSU 10620 NP 388943.2
consensus

LFFRNEVKQLERSVSRQLYGERIQGSVSRMETFNACPFSHFASHGLHLKERQFFKLEAPDIGQLFHSSKLISDRLREQKLDWRD
LFFRNEVKQLERSVSRQLYGERIQGSVSRMETFNACPFSHFASHGLHLKERQFFKLEAPDIGQLFHSSKLISDRLREQKLDWRD
LFFRNEVKQLERSVSRQLYGERIQGSVSRMETFNACPFSHFASHGLHLKERQFFKLEAPDIGQLFHSSKLISDRLREQKLDWRD

850
850

logo

NTDB id 73411 FNP71 RS05950 WP 003244988.1
NTDB id 120 BSU 10620 NP 388943.2
consensus

LTKEQCELFSDAVERLAPKLQKEILLSSNRHYVKEKLQKIVTRVSGILSEHAKASGFVPIGLELFGGKGPLPPLTFQLKNGC
 LTKEQCELFSDAVERLAPKLQKEILLSSNRHYVKEKLQKIVTRVSGILSEHAKASGFVPIGLELFGGKGGLPPLTFQLKNGC
 LTKEQCELFSDAVERLAPKLQKEILLSSNRHYVKEKLQKIVTRVSGILSEHAKASGFVPIGLELFGGKGPLPPLTFQLKNGC

935
935

logo

NTDB id 73411 FNP71 RS05950 WP 003244988.1
NTDB id 120 BSU 10620 NP 388943.2
consensus

TMELVGRIDRVDKAESSKGILLRIVDYKSSDKGLDLAEVYGLALQMLTYLDLSITHSADWLGMRA TPAGVLYFHIHDPMIQSNL
 TMELVGRIDRVDKAESSKGILLRIVDYKSSDKGLDLAEVYGLALQMLTYLDLSITHSADWLGMRA TPAGVLYFHIHDPMIQSNL
 TMELVGRIDRVDKAESSKGILLRIVDYKSSDKGLDLAEVYGLALQMLTYLDLSITHSADWLGMRA TPAGVLYFHIHDPMIQSNL

1020
1020

logo

NTDB id 73411 FNP71 RS05950 WP 003244988.1
NTDB id 120 BSU 10620 NP 388943.2
consensus

PLGLDEIEQEIFKKFKMKGLLLGDQEWRLMDTTLQEGRSNIINAGLKKDGSLSRSDSAAVGEKEFDLLTKHVRRTFQEAGEQITD
PLGLDEIEQEIFKKFKMKGLLLGDQEVVRLMDTTLQEGRSNIINAGLKKDGSLSRSDSAAVGEKEFDLLTKHVRRTFQEAGEQITD
PLGLDEIEQEIFKKFKMKGLLLGDQEVVRLMDTTLQEGRSNIINAGLKKDGSLSRSDSAAVGEKEFDLLTKHVRRTFQEAGEQITD

1105
1105

logo

NTDB id 73411 FNP71 RS05950 WP 003244988.1
NTDB id 120 BSU 10620 NP 388943.2
consensus

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

1166
1166

 non conserved
 similar
 $\geq 50\%$ conserved