



logo

NTDB id 364996 FC605 RS05765 WP 080529152.1  
NTDB id 120 BSU 10620 NP 388943.2  
consensus

SLDLFQQMIEAGAESLTFSLIPPALDQVFGNMDLSRMYGTSCTFVLGANDGVLPARPDENGVLSDDDREWLKTI GVELS  
SLDLFQQMIEAGAESLTFSLIPPALDQVFGNMDLSRMYGTSCTFVLGANDGVLPARPDENGVLSDDDREWLKTI GVELS  
!!

640  
640

logo

NTDB id 364996 FC605 RS05765 WP 080529152.1  
NTDB id 120 BSU 10620 NP 388943.2  
consensus

SGGRERLLDEHFLIYMAFSSPSDRLVSYPIADAEGKTL LPSMIVKRLEELFPHHMERLLTNEPEQVSDEEQLMYVVKNS  
SGGRERLLDEHFLIYMAFSSPSDRLVSYPIADAEGKTL LPSMIVKRLEELFPHHMERLLTNEPEQVSDEEQLMYVVKNS  
!!

720  
720

logo

NTDB id 364996 FC605 RS05765 WP 080529152.1  
NTDB id 120 BSU 10620 NP 388943.2  
consensus

VAQSFTASQLRLWTREYDISDVVWSTYNVLMSE<sup>Q</sup>DRLQSKKLFSSLFFRNEVKQLERSVSRQLYGERIQGSVSRMETFNA  
VAQSFTASQLRLWTREYDISDVVWSTYNVLMSE<sup>Q</sup>DRLQSKKLFSSLFFRNEVKQLERSVSRQLYGERIQGSVSRMETFNA  
!!

800  
800

logo

NTDB id 364996 FC605 RS05765 WP 080529152.1  
NTDB id 120 BSU 10620 NP 388943.2  
consensus

CPFSHFASHGLHLKERQFFKLEAPDIGQLFHSSLKLISDRLREQKLDWRDLTKEQCELFSDAVERLAPKLQKEILLSSN  
CPFSHFASHGLHLKERQFFKLEAPDIGQLFHSSLKLISDRLREQKLDWRDLTKEQCELFSDAVERLAPKLQKEILLSSN  
!!

880  
880

logo

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NTDB id 120 BSU 10620 NP 388943.2  
consensus

RHYVVEKLEKIVTRVSGILSEHAKASGFVPIGLELFGGKGPLPPLTFQLKNGCTMELVGRIDRVDKAESSKGLLLRIV  
RHYVVEKLEKIVTRVSGILSEHAKASGFVPIGLELFGGKGPLPPLTFQLKNGCTMELVGRIDRVDKAESSKGLLLRIV  
!!

960  
960

logo

NTDB id 364996 FC605 RS05765 WP 080529152.1  
NTDB id 120 BSU 10620 NP 388943.2  
consensus

DYKSSDKGLDLAEVYGLALQMLTYLDSLITHSADWLGMRATPAGVLYFHIHDPMIQSNLPLGLDEIEQEIEFKKFKMKGL  
DYKSSDKGLDLAEVYGLALQMLTYLDSLITHSADWLGMRATPAGVLYFHIHDPMIQSNLPLGLDEIEQEIEFKKFKMKGL  
!!

1040  
1040

logo

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NTDB id 120 BSU 10620 NP 388943.2  
consensus

LLGDQEVRLMDTTLQEGRSNIINAGLKKDGSLSRSDSAAVGEKEFDLLTKHVRRTFQEAGEQITDGRVSI EPYKMKNKTP  
LLGDQEVRLMDTTLQEGRSNIINAGLKKDGSLSRSDSAAVGEKEFDLLTKHVRRTFQEAGEQITDGRVSI EPYKMKNKTP  
!!

1120  
1120

