

logo

NTDB id 398203 GII78 RS05730 WP 113713002.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



80
80

logo

NTDB id 398203 GII78 RS05730 WP 113713002.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



160
160

logo

NTDB id 398203 GII78 RS05730 WP 113713002.1
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consensus



240
240

logo

NTDB id 398203 GII78 RS05730 WP 113713002.1
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consensus



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

SLDLFQQMIEAGAESLTFSLIPPALDQVFVGNMDSLRYGTSCTFVLGANDGVLPARPDENGVLSDDDREWLKTI GVELS
SLDLFQQMIEAGAESLTFSLIPPALDQVFVGNMDSLRYGTSCTFVLGANDGVLPARPDENGVLSDDDREWLKTI GVELS
!!

640
640

logo

NTDB id 398203 GII78 RS05730 WP 113713002.1
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consensus

SGGRERLL⁵EHFLIYMAFSSPSDRLVSYPIADAEGKTL⁵PSMIVKRLEELFPHHKERLLTNEPEQVSDEEQLMYV⁵VNKS
SGGRERLL⁵EHFLIYMAFSSPSDRLVSYPIADAEGKTL⁵PSMIVKRLEELFPHHKERLLTNEPEQVSDEEQLMYV⁵VNKS
SGGRERLL⁵EHFLIYMAFSSPSDRLVSYPIADAEGKTL⁵PSMIVKRLEELFPHHKERLLTNEPEQVSDEEQLMYV⁵VNKS
!!!!!!!!!!!!!!!!*!!

720
720

logo

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consensus

VAQSFTASQLRLWTREYDISDVVWSTYNVLMSEQDRLQSKKLFSSLFFRNEVKQLERSVSRQLYGERIQGSVSRMETFNA
VAQSFTASQLRLWTREYDISDVVWSTYNVLMSEQDRLQSKKLFSSLFFRNEVKQLERSVSRQLYGERIQGSVSRMETFNA
VAQSFTASQLRLWTREYDISDVVWSTYNVLMSEQDRLQSKKLFSSLFFRNEVKQLERSVSRQLYGERIQGSVSRMETFNA
!!

800
800

logo

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CPFSHFASHGLHLKERQFFKLEAPDIGQLFHSSLKLISDRLREQKLDWRDLTKEQCELFSDAVERLAPKLQKEILLSSN
CPFSHFASHGLHLKERQFFKLEAPDIGQLFHSSLKLISDRLREQKLDWRDLTKEQCELFSDAVERLAPKLQKEILLSSN
CPFSHFASHGLHLKERQFFKLEAPDIGQLFHSSLKLISDRLREQKLDWRDLTKEQCELFSDAVERLAPKLQKEILLSSN
!!

880
880

logo

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RHYVKEK⁵LQKIVTRVSGILSEHAKASGFVPIGLELFGGKGPLPPLTFQLKNGCTMELVGRIDRVDKAESSKGLLLRIV
RHYVKEK⁵LQKIVTRVSGILSEHAKASGFVPIGLELFGGKGPLPPLTFQLKNGCTMELVGRIDRVDKAESSKGLLLRIV
RHYVKEK⁵LQKIVTRVSGILSEHAKASGFVPIGLELFGGKGPLPPLTFQLKNGCTMELVGRIDRVDKAESSKGLLLRIV
!!

960
960

logo

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DYKSSDKGLDLAEVYGLALQMLTYL⁵DL⁵SITHSADWLGMRATPAGVLYFHIHDPMIQSNLPLGLDEIEQEIFKKFKMKGL
DYKSSDKGLDLAEVYGLALQMLTYL⁵DL⁵SITHSADWLGMRATPAGVLYFHIHDPMIQSNLPLGLDEIEQEIFKKFKMKGL
DYKSSDKGLDLAEVYGLALQMLTYL⁵DL⁵SITHSADWLGMRATPAGVLYFHIHDPMIQSNLPLGLDEIEQEIFKKFKMKGL
!!

1040
1040

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LLGDQEWRLMDTTLQEGRSNIINAGLKKDGSLSRSDSAAVG⁵KEFDLLTKHVRRTFQEAGEQITDGRVSI⁵EPYKMK⁵NKTP
LLGDQEWRLMDTTLQEGRSNIINAGLKKDGSLSRSDSAAVG⁵KEFDLLTKHVRRTFQEAGEQITDGRVSI⁵EPYKMK⁵NKTP
LLGDQEWRLMDTTLQEGRSNIINAGLKKDGSLSRSDSAAVG⁵KEFDLLTKHVRRTFQEAGEQITDGRVSI⁵EPYKMK⁵NKTP
!!

1120
1120

