

logo

NTDB id 981452 NSU09 RS05735 WP 113713002.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



80
80

logo

NTDB id 981452 NSU09 RS05735 WP 113713002.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



160
160

logo

NTDB id 981452 NSU09 RS05735 WP 113713002.1
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consensus



240
240

logo

NTDB id 981452 NSU09 RS05735 WP 113713002.1
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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 981452 NSU09 RS05735 WP 113713002.1
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consensus

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

720
720

logo

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VAQSFTASQLRLWTREYDISDVVWSTYNVLMSEQDRLQSKKLFSSLFFRNEVKQLERSVSRQLYGERIQGSVSRMETFNA
VAQSFTASQLRLWTREYDISDVVWSTYNVLMSEQDRLQSKKLFSSLFFRNEVKQLERSVSRQLYGERIQGSVSRMETFNA
VAQSFTASQLRLWTREYDISDVVWSTYNVLMSEQDRLQSKKLFSSLFFRNEVKQLERSVSRQLYGERIQGSVSRMETFNA
!!

800
800

logo

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CPFSHFASHGLHLKERQFFKLEAPDIGQLFHSSLKLISDRLREQKLDWRDLTKEQCELFSDAVERLAPKLQKEILLSSN
CPFSHFASHGLHLKERQFFKLEAPDIGQLFHSSLKLISDRLREQKLDWRDLTKEQCELFSDAVERLAPKLQKEILLSSN
CPFSHFASHGLHLKERQFFKLEAPDIGQLFHSSLKLISDRLREQKLDWRDLTKEQCELFSDAVERLAPKLQKEILLSSN
!!

880
880

logo

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RHYVKEKLQKIVTRVSGILSEHAKASGFVPIGLELFGGKGPLPPLTFQLKNGCTMELVGRIDRVDKAESSKGLLLRIV
RHYVKEKLQKIVTRVSGILSEHAKASGFVPIGLELFGGKGPLPPLTFQLKNGCTMELVGRIDRVDKAESSKGLLLRIV
RHYVKEKLQKIVTRVSGILSEHAKASGFVPIGLELFGGKGPLPPLTFQLKNGCTMELVGRIDRVDKAESSKGLLLRIV
!!

960
960

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DYKSSDKGLDLAEVYGLALQMLTYLDSLITHSADWLGMRATPAGVLYFHIHDPMIQSNLPLGLDEIEQEIEFKKFKMKGL
DYKSSDKGLDLAEVYGLALQMLTYLDSLITHSADWLGMRATPAGVLYFHIHDPMIQSNLPLGLDEIEQEIEFKKFKMKGL
DYKSSDKGLDLAEVYGLALQMLTYLDSLITHSADWLGMRATPAGVLYFHIHDPMIQSNLPLGLDEIEQEIEFKKFKMKGL
!!

1040
1040

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LLGDQEWRLMDTTLQEGRSNIINAGLKKDGSLSRSDSAAVG⁵EKEFDLLTKHVRRTFQEAGEQITDGRVSI⁵EPYKMKNKTP
LLGDQEVVRLMDTTLQEGRSNIINAGLKKDGSLSRSDSAAVG⁵EKEFDLLTKHVRRTFQEAGEQITDGRVSI⁵EPYKMKNKTP
LLGDQEVVRLMDTTLQEGRSNIINAGLKKDGSLSRSDSAAVG⁵EKEFDLLTKHVRRTFQEAGEQITDGRVSI⁵EPYKMKNKTP
!!

1120
1120

