

logo

NTDB id 163544 AT706 RS12845 WP 015483058.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



80
80

logo

NTDB id 163544 AT706 RS12845 WP 015483058.1
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consensus



160
160

logo

NTDB id 163544 AT706 RS12845 WP 015483058.1
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240
240

logo

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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

SLDLFQQMIEAGAESLTFSLIPPALDQVFGNMDLSRMYGT_YCTFVLGANDGVLPARPDENGVLSDDDREWLKTI_YGVELS
SLDLFQQMIEAGAESLTFSLIPPALDQVFGNMDLSRMYGT_YCTFVLGANDGVLPARPDENGVLSDDDREWLKTI_YGVELS
!!

640
640

logo

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SGGRERLLDEHFLIYMAFSSPSDRLVSYPIADAEGKTL_LPSMIVKRLEELFPHHKERLLTNEPEQVSDEEQL_YMYV_NKS
SGGRERLLDEHFLIYMAFSSPSDRLVSYPIADAEGKTL_LPSMIVKRLEELFPHHKERLLTNEPEQVSDEEQL_YMYV_NKS
!!

720
720

logo

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VAQSFTASQLRLWTREYDISD_VVMSTYNVLMSE_PDRLQSKKLFSSLFFRNEVKQLERSV_SRQLYGERIQGSVSRMETFNA
VAQSFTASQLRLWTREYDISD_VVMSTYNVLMSE_PDRLQSKKLFSSLFFRNEVKQLERSV_SRQLYGERIQGSVSRMETFNA
VAQSFTASQLRLWTREYDISD_VVMSTYNVLMSE_PDRLQSKKLFSSLFFRNEVKQLERSV_SRQLYGERIQGSVSRMETFNA
!!

800
800

logo

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CPFSHFASHGLHLKERQFFKLEAPDIGQLFHSSLKLISDRLREQKLDWRDLTKEQCELF_SYDAVERLAPKLQKEILLSSN
CPFSHFASHGLHLKERQFFKLEAPDIGQLFHSSLKLISDRLREQKLDWRDLTKEQCELF_SYDAVERLAPKLQKEILLSSN
CPFSHFASHGLHLKERQFFKLEAPDIGQLFHSSLKLISDRLREQKLDWRDLTKEQCELF_SYDAVERLAPKLQKEILLSSN
!!

880
880

logo

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RHYVKEK_LQKIVTRVSGILSEHAKASGFVPIGLELFGGKGPLPPLTFQLKNGCTMELVGRIDRV_DKAESSKGLLLRIV
RHYVKEK_LQKIVTRVSGILSEHAKASGFVPIGLELFGGKGPLPPLTFQLKNGCTMELVGRIDRV_DKAESSKGLLLRIV
RHYVKEK_LQKIVTRVSGILSEHAKASGFVPIGLELFGGKGPLPPLTFQLKNGCTMELVGRIDRV_DKAESSKGLLLRIV
!!

960
960

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DYKSSDKGLDLAEVYGLALQMLTYL_DLSITHSADWLGM_RATPAGVLYFHIHDPMIQSNLPLGLDEIEQEIFKKFKMKGL
DYKSSDKGLDLAEVYGLALQMLTYL_DLSITHSADWLGM_RATPAGVLYFHIHDPMIQSNLPLGLDEIEQEIFKKFKMKGL
DYKSSDKGLDLAEVYGLALQMLTYL_DLSITHSADWLGM_RATPAGVLYFHIHDPMIQSNLPLGLDEIEQEIFKKFKMKGL
!!

1040
1040

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LLGDQEWRLMDTTLQEGRSNIINAGLKKDGS_LRS_DSAAVGEKEFDLLTKHVRRTFQEAGEQITDGRVSI_EPKMK_NKTP
LLGDQEWRLMDTTLQEGRSNIINAGLKKDGS_LRS_DSAAVGEKEFDLLTKHVRRTFQEAGEQITDGRVSI_EPKMK_NKTP
LLGDQEWRLMDTTLQEGRSNIINAGLKKDGS_LRS_DSAAVGEKEFDLLTKHVRRTFQEAGEQITDGRVSI_EPKMK_NKTP
!!

1120
1120

