

logo

NTDB id 967939 MKY10 RS05690 WP 044052572.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



80
80

logo

NTDB id 967939 MKY10 RS05690 WP 044052572.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



160
160

logo

NTDB id 967939 MKY10 RS05690 WP 044052572.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



240
240

logo

NTDB id 967939 MKY10 RS05690 WP 044052572.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



320
320

logo

NTDB id 967939 MKY10 RS05690 WP 044052572.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



400
400

logo

NTDB id 967939 MKY10 RS05690 WP 044052572.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



480
480

logo

NTDB id 967939 MKY10 RS05690 WP 044052572.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



560
560

logo

NTDB id 967939 MKY10 RS05690 WP 044052572.1
NTDB id 120 BSU 10620 NP 388943.2
consensus

SLDLFQQMIEAGAESLTFSLIPPALDQVFVGNMDSLRYMGTSCTFVLGANDGVLPARPDENGVLSDDDREWLKTI GVELS
SLDLFQQMIEAGAESLTFSLIPPALDQVFVGNMDSLRYMGTSCTFVLGANDGVLPARPDENGVLSDDDREWLKTI GVELS
!!

640
640

logo

NTDB id 967939 MKY10 RS05690 WP 044052572.1
NTDB id 120 BSU 10620 NP 388943.2
consensus

SGGRERLLDEHFLIYMAFSSPSDRLVSYPIADAEGKTLPSMIVKRLGELFPHHKERLLTNEPEQVSDEEQLMYVVKNS
SGGRERLLDEHFLIYMAFSSPSDRLVSYPIADAEGKTLPSMIVKRLGELFPHHKERLLTNEPEQVSDEEQLMYVVKNS
!!

720
720

logo

NTDB id 967939 MKY10 RS05690 WP 044052572.1
NTDB id 120 BSU 10620 NP 388943.2
consensus

VAQSFTASQLRLWTREYDISDVVWSTYNVLMSEQDRLQSKKLFSSLFFRNEVKQLERSVSRQLYGERIQGSVSRMETFNA
VAQSFTASQLRLWTREYDISDVVWSTYNVLMSEQDRLQSKKLFSSLFFRNEVKQLERSVSRQLYGERIQGSVSRMETFNA
VAQSFTASQLRLWTREYDISDVVWSTYNVLMSEQDRLQSKKLFSSLFFRNEVKQLERSVSRQLYGERIQGSVSRMETFNA
!!

800
800

logo

NTDB id 967939 MKY10 RS05690 WP 044052572.1
NTDB id 120 BSU 10620 NP 388943.2
consensus

CPFSHFASHGLHLKERQFFKLEAPDIGQLFHSSLKLISDRLREQKLDWRDLTKEQCELFSDAVERLAPKLQKEILLSSN
CPFSHFASHGLHLKERQFFKLEAPDIGQLFHSSLKLISDRLREQKLDWRDLTKEQCELFSDAVERLAPKLQKEILLSSN
CPFSHFASHGLHLKERQFFKLEAPDIGQLFHSSLKLISDRLREQKLDWRDLTKEQCELFSDAVERLAPKLQKEILLSSN
!!

880
880

logo

NTDB id 967939 MKY10 RS05690 WP 044052572.1
NTDB id 120 BSU 10620 NP 388943.2
consensus

RHYVKEKLVKIVTRVSGILSEHAKASGFVPIGLELFGGKGPLPPLTFQLKNGCTMELVGRIDRVDKAESSKGLLLRIV
RHYVKEKLVKIVTRVSGILSEHAKASGFVPIGLELFGGKGPLPPLTFQLKNGCTMELVGRIDRVDKAESSKGLLLRIV
RHYVKEKLVKIVTRVSGILSEHAKASGFVPIGLELFGGKGPLPPLTFQLKNGCTMELVGRIDRVDKAESSKGLLLRIV
!!

960
960

logo

NTDB id 967939 MKY10 RS05690 WP 044052572.1
NTDB id 120 BSU 10620 NP 388943.2
consensus

DYKSSDKGLDLAEVYGLALQMLTYLDSLITHSADWLGMRATPAGVLYFHIHDPMIQSNLPLGLDEIEQEIEFKKFKMKGL
DYKSSDKGLDLAEVYGLALQMLTYLDSLITHSADWLGMRATPAGVLYFHIHDPMIQSNLPLGLDEIEQEIEFKKFKMKGL
DYKSSDKGLDLAEVYGLALQMLTYLDSLITHSADWLGMRATPAGVLYFHIHDPMIQSNLPLGLDEIEQEIEFKKFKMKGL
!!

1040
1040

logo

NTDB id 967939 MKY10 RS05690 WP 044052572.1
NTDB id 120 BSU 10620 NP 388943.2
consensus

LLGDQEWRLMDTTLQEGRSNIINAGLKKDGSLSRSDSAAVGEKEFDLLTKHVRRTFQEAGEQITDGRVSI EPYKMKNKTP
LLGDQEWRLMDTTLQEGRSNIINAGLKKDGSLSRSDSAAVGEKEFDLLTKHVRRTFQEAGEQITDGRVSI EPYKMKNKTP
LLGDQEWRLMDTTLQEGRSNIINAGLKKDGSLSRSDSAAVGEKEFDLLTKHVRRTFQEAGEQITDGRVSI EPYKMKNKTP
!!

1120
1120

