

logo

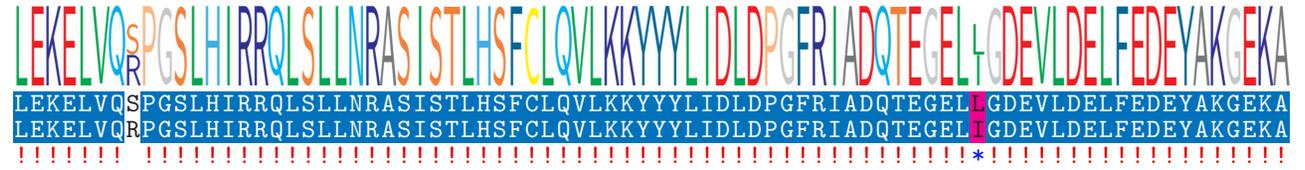
NTDB id 234779 CD084 RS04845 WP 040083514.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



80
80

logo

NTDB id 234779 CD084 RS04845 WP 040083514.1
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consensus



160
160

logo

NTDB id 234779 CD084 RS04845 WP 040083514.1
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consensus



240
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NTDB id 234779 CD084 RS04845 WP 040083514.1
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consensus



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320

logo

NTDB id 234779 CD084 RS04845 WP 040083514.1
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consensus



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logo

NTDB id 234779 CD084 RS04845 WP 040083514.1
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consensus



480
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560
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logo

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consensus

KLISSPFKVDGKTKTHRNIQYRDIVILLRSMWPAPQIMEELRAQGIPVYANLTSGYFEAVEVAVALSVLKVIDNPHYQDI
KLISSPFKVDGKTKTHRNIQYRDIVILLRSMWPAPQIMEELRAQGIPVYANLTSGYFEAVEVAVALSVLKVIDNPHYQDI
!!

640
640

logo

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consensus

PLASVLRSPVIGADENELSLIRLENKKAPYYEAMKDYLAAAGDRNDELYQKLHTFYGHLQKWRVAFSKNHSVSELIWEVYRD
PLASVLRSPVIGADENELSLIRLENKKAPYYEAMKDYLAAAGDRNDELYQKLHTFYGHLQKWRVAFSKNHSVSELIWEVYRD
PLASVLRSPVIGADENELSLIRLENKKAPYYEAMKDYLAAAGDRSDELYQKLHTFYGHLQKWRVAFSKNHSVSELIWEVYRD
!!

720
720

logo

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consensus

TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVRLMTIHSSKGLE
TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVRLMTIHSSKGLE
TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVRLMTIHSSKGLE
!!

800
800

logo

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FPVVFAGLGRNFMMDLNKSYLLDKELGFGTKYIHPQLRISYPTLPLIAMKKMRRELLSEELRVLYVALTRAKEKLFEL
FPVVFAGLGRNFMMDLNKSYLLDKELGFGTKYIHPQLRISYPTLPLIAMKKMRRELLSEELRVLYVALTRAKEKLFEL
FPVVFAGLGRNFMMDLNKSYLLDKELGFGTKYIHPQLRISYPTLPLIAMKKMRRELLSEELRVLYVALTRAKEKLFEL
!!

880
880

logo

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IGSCKNHQKQLAKWQASASQTDWLLPEFDYQARTYLDFIGSALARHRDLEALAGNGTPVHADISGHPARFAVQMIHSYD
IGSCKNHQKQLAKWQASASQTDWLLPEFDYQARTYLDFIGSALARHRDLEALAGNGTPVHADISGHPARFAVQMIHSYD
IGSCKDHQKQLAKWQASASQTDWLLPEFDYQARTYLDFIGSALARHRDLEALAGNGTPVHADISGHPARFAVQMIHSYD
!!

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958

logo

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LLDDDYEERMEEKSERLEAIRRGEVPGSFADFDEKAREQLSWTYPHQEVTVIRTKQSVSEIKRKREYEDEYSGRAPVKPA
LLDDDYEERMEEKSERLEAIRRGEVPGSFADFDEKAREQLSWTYPHQEVTVIRTKQSVSEIKRKREYEDEYSGRAPVKPA
LLDDDYEERMEEKSERLEAIRRGEVPGSFADFDEKAREQLSWTYPHQEVTVIRTKQSVSEIKRKREYEDEYSGRAPVKPA
!!

1040
1038

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DGSILYRRPAFMMKKGLTAAEKGTAMHTVMQHIPLSHVPSIEEAEQTVHRLYEKELLTEEQKDAIDIEEIVQFFHTEIGG
DGSILYRRPAFMMKKGLTAAEKGTAMHTVMQHIPLSHVPSIEEAEQTVHRLYEKELLTEEQKDAIDIEEIVQFFHTEIGG
DGSILYRRPAFMMKKGLTAAEKGTAMHTVMQHIPLSHVPSIEEAEQTVHRLYEKELLTEEQKDAIDIEEIVQFFHTEIGG
!!

1120
1118

