

logo

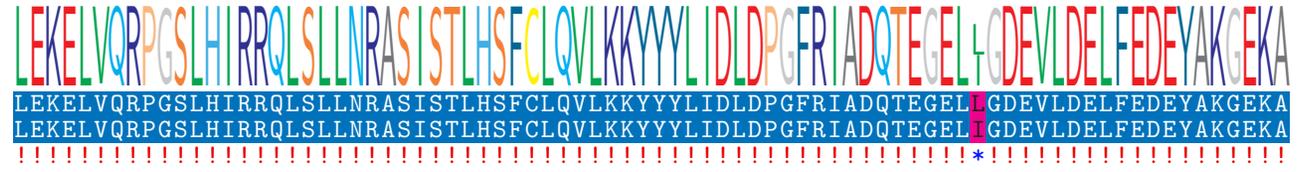
NTDB id 393969 GCU76 RS05755 WP 152425414.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



80
80

logo

NTDB id 393969 GCU76 RS05755 WP 152425414.1
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consensus



160
160

logo

NTDB id 393969 GCU76 RS05755 WP 152425414.1
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consensus



240
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NTDB id 393969 GCU76 RS05755 WP 152425414.1
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consensus



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NTDB id 393969 GCU76 RS05755 WP 152425414.1
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consensus



400
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NTDB id 393969 GCU76 RS05755 WP 152425414.1
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consensus



480
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560
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logo

NTDB id 393969 GCU76 RS05755 WP 152425414.1
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consensus

KLISSPFKVDGKKKTHRNQYRDIVILLRSMWPAPQIMEELRAQGIPVYANLTSYGFEEAVEVAVALSVLKVIDNPYQDI
KLISSPFKVDGKKKTHRNQYRDIVILLRSMWPAPQIMEELRAQGIPVYANLTSYGFEEAVEVAVALSVLKVIDNPYQDI
!!

640
640

logo

NTDB id 393969 GCU76 RS05755 WP 152425414.1
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consensus

PLASVLRSPVIGADENELSLIRLENKKAPYYEAMKDYLAAAGDRSDELYQKLNFTFYGHLQKWRAFSKNHSVSELIWEVYRD
PLASVLRSPVIGADENELSLIRLENKKAPYYEAMKDYLAAAGDRNDELYQKLNFTFYGHLQKWRAFSKNHSVSELIWEVYRD
PLASVLRSPVIGADENELSLIRLENKKAPYYEAMKDYLAAAGDRSDELYQKLNFTFYGHLQKWRAFSKNHSVSELIWEVYRD
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720
720

logo

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TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVRLMTIHSSKGLE
TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVRLMTIHSSKGLE
TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVRLMTIHSSKGLE
!!

800
800

logo

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FPVVF_AAGLGRNFMMDLNKSYLLDKELGFGTKYIHPQLRISYPTLPLIAMKKMRRELLSEELRVLYVALTRAKEKLFEL
FPVVF_AAGLGRNFMMDLNKSYLLDKELGFGTKYIHPQLRISYPTLPLIAMKKMRRELLSEELRVLYVALTRAKEKLFEL
FPVVF_AAGLGRNFMMDLNKSYLLDKELGFGTKYIHPQLRISYPTLPLIAMKKMRRELLSEELRVLYVALTRAKEKLFEL
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880
880

logo

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IGSCKDHQKQLAKWQASASQTDWLLPEFDYQARTYLDFIGPALARHRL_EDLAGVPAHADISGHPARFAVQMIHSYDLL
IGSCKDHQKQLAKWQASASQTDWLLPEFDYQARTYLDFIGPALARHRL_EDLAGVPAHADISGHPARFAVQMIHSYDLL
IGSCKDHQKQLAKWQASASQTDWLLPEFDYQARTYLDFIGPALARHRL_EDLAGVPAHADISGHPARFAVQMIHSYDLL
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960
960

logo

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DDDLEERMEEKSERLEAIRRGEVPGSF_AFDEKAREQLSWTYPH_HEVTQIRTKQSVSEIKRKREYEDEYSGRAPVKPADG
DDDLEERMEEKSERLEAIRRGEVPGSFTFDEKAREQLSWTYPH_HEVTQIRTKQSVSEIKRKREYEDEYSGRAPVKPADG
DDDLEERMEEKSERLEAIRRGEVPGSFAFDEKAREQLSWTYPH_HEVTQIRTKQSVSEIKRKREYEDEYSGRAPVKPADG
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1040
1040

logo

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SILYRRP_AFMMKKGLTAAEKGTAMHTVMQHIPLSHVPSIEEAEQTVHRLYE_RELLTEEQKDAIDIEEIVQFFHTEIGGQL
SILYRRP_AFMMKKGLTAAEKGTAMHTVMQHIPLSHVPSIEEAEQTVHRLYE_RELLTEEQKDAIDIEEIVQFFHTEIGGQL
SILYRRP_AFMMKKGLTAAEKGTAMHTVMQHIPLSHVPSIEEAEQTVHRLYE_RELLTEEQKDAIDIEEIVQFFHTEIGGQL
!!

1120
1120

