

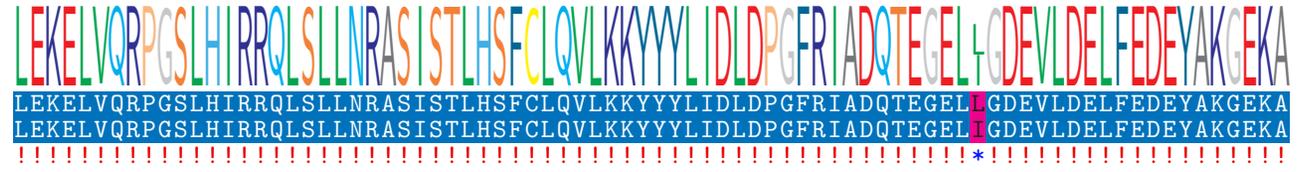
logo

NTDB id 340076 ETA18 RS05840 WP 128992613.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



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KLISSPFKVDGKKKTHRNIQYRDIVILLRSMWPAPQIMEELRAQGIPVYANLTSYGFEEAVEVAVALSVLKVIDNPYQDI
KLISSPFKVDGKKKTHRNIQYRDIVILLRSMWPAPQIMEELRAQGIPVYANLTSYGFEEAVEVAVALSVLKVIDNPYQDI
!!

640
640

logo

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consensus

PLASVLRSPVIGADENELSLIRLENKKAPYYEAMKDYLAAAGDR^sDELYQKLNFTFYGHLQKWRAFSKNHSVSELIWEVYRD
PLASVLRSPVIGADENELSLIRLENKKAPYYEAMKDYLAAAGDRNDELYQKLNFTFYGHLQKWRAFSKNHSVSELIWEVYRD
PLASVLRSPVIGADENELSLIRLENKKAPYYEAMKDYLAAAGDRSDELYQKLNFTFYGHLQKWRAFSKNHSVSELIWEVYRD
!!

720
720

logo

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consensus

TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVRLMTIHSSKGLE
TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVRLMTIHSSKGLE
TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVRLMTIHSSKGLE
!!

800
800

logo

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FPVVF_AAGLGRNFNMDLNKSYLLDKELGFGTKYIHPQLRISYPTLPLIAMKKMRRELLSEELRVLYVALTRAKEKLFEL
FPVVF_AAGLGRNFNMDLNKSYLLDKELGFGTKYIHPQLRISYPTLPLIAMKKMRRELLSEELRVLYVALTRAKEKLFEL
FPVVFV_AAGLGRNFNMDLNKSYLLDKELGFGTKYIHPQLRISYPTLPLIAMKKMRRELLSEELRVLYVALTRAKEKLFEL
!!

880
880

logo

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consensus

IGSCKDHQKQLAKWQASASQTDWLLPEFDYQARTYLDFIGPALARHRL^EDLAGVPAHADIS_GHPARFAVQMIHSYDLL
IGSCKDHQKQLAKWQASASQTDWLLPEFDYQARTYLDFIGPALARHRL^EDLAGVPAHADIS_GHPARFAVQMIHSYDLL
IGSCKDHQKQLAKWQASASQTDWLLPEFDYQARTYLDFIGPALARHRL^EDLAGVPAHADIS_GHPARFAVQMIHSYDLL
!!

960
960

logo

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DDDLEERMEEKSERLEAIRRGEVPVGSFAFDEKAREQLSWTYPH^HEVTQIRTKQSVSEIKRKREYEDEYSGRAPVKPADG
DDDLEERMEEKSERLEAIRRGEVPVGSFAFDEKAREQLSWTYPH^HEVTQIRTKQSVSEIKRKREYEDEYSGRAPVKPADG
DDDLEERMEEKSERLEAIRRGEVPVGSFAFDEKAREQLSWTYPH^HEVTQIRTKQSVSEIKRKREYEDEYSGRAPVKPADG
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1040
1040

logo

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consensus

SIL_FRRPAFMMKKGLTAAEKGTAMHTVMQHIPLSHVPSIEEAEQTVHRLYEKELLTEEQKDAIDIEEIVQFFHTEIGGQL
SIL_FRRPAFMMKKGLTAAEKGTAMHTVMQHIPLSHVPSIEEAEQTVHRLYEKELLTEEQKDAIDIEEIVQFFHTEIGGQL
SIL_FRRPAFMMKKGLTAAEKGTAMHTVMQHIPLSHVPSIEEAEQTVHRLYEKELLTEEQKDAIDIEEIVQFFHTEIGGQL
!!

1120
1120

