

logo

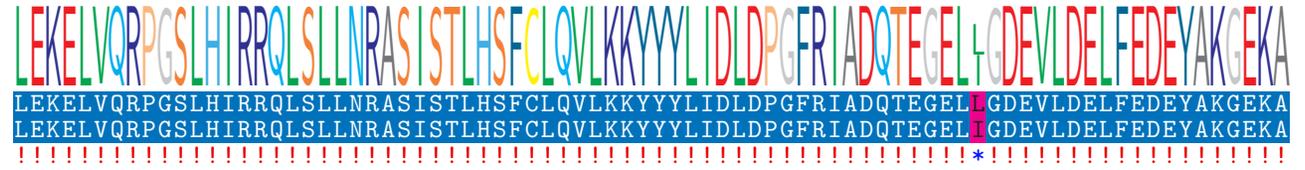
NTDB id 339601 ES965 RS05770 WP 124059687.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



80
80

logo

NTDB id 339601 ES965 RS05770 WP 124059687.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



160
160

logo

NTDB id 339601 ES965 RS05770 WP 124059687.1
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consensus



240
240

logo

NTDB id 339601 ES965 RS05770 WP 124059687.1
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consensus



320
320

logo

NTDB id 339601 ES965 RS05770 WP 124059687.1
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consensus



400
400

logo

NTDB id 339601 ES965 RS05770 WP 124059687.1
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consensus



480
480

logo

NTDB id 339601 ES965 RS05770 WP 124059687.1
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consensus



560
560

logo

NTDB id 339601 ES965 RS05770 WP 124059687.1
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consensus

KLISSPFKVDGKKKTHRNQYRDIVILLRSMWPAPQIMEELRAQGIPVYANLTSYGFEEAVEVAVALSVLKVIDNPYQDI
KLISSPFKVDGKKKTHRNQYRDIVILLRSMWPAPQIMEELRAQGIPVYANLTSYGFEEAVEVAVALSVLKVIDNPYQDI
!!

640
640

logo

NTDB id 339601 ES965 RS05770 WP 124059687.1
NTDB id 119 BSU 10630 NP 388944.2
consensus

PLASVLRSPVIGADENELSLIRLENKKAPYYEAMKDYLAAAGDRSDELYQKLNFTFYGHLQKWRAFSKNHSVSELIWEVYRD
PLASVLRSPVIGADENELSLIRLENKKAPYYEAMKDYLAAAGDRNDELYQKLNFTFYGHLQKWRAFSKNHSVSELIWEVYRD
PLASVLRSPVIGADENELSLIRLENKKAPYYEAMKDYLAAAGDRSDELYQKLNFTFYGHLQKWRAFSKNHSVSELIWEVYRD
!!

720
720

logo

NTDB id 339601 ES965 RS05770 WP 124059687.1
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consensus

TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVRLMTIHSSKGLE
TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVRLMTIHSSKGLE
TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVRLMTIHSSKGLE
!!

800
800

logo

NTDB id 339601 ES965 RS05770 WP 124059687.1
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consensus

FPVVF_AAGLGRNFNMDLNKSYLLDKELGFGTKYIHPQLRISYPTLPLIAMKKMRRELLSEELRVLYVALTRAKEKLFEL
FPVVF_AAGLGRNFNMDLNKSYLLDKELGFGTKYIHPQLRISYPTLPLIAMKKMRRELLSEELRVLYVALTRAKEKLFEL
FPVVF_VAGLGRNFNMDLNKSYLLDKELGFGTKYIHPQLRISYPTLPLIAMKKMRRELLSEELRVLYVALTRAKEKLFEL
!!

880
880

logo

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consensus

IGSCKDHQKQLAKWQASASQTDWLLPEFDRYQARTYLDFIGPALARHRL_EDLAGVPAHADIS_GHPARFAVQMIHSYDLL
IGSCKDHQKQLAKWQASASQTDWLLPEFDRYQARTYLDFIGPALARHRL_EDLAGVPAHADIS_DHPARFAVQMIHSYDLL
IGSCKDHQKQLAKWQASASQTDWLLPEFDRYQARTYLDFIGPALARHRL_GDLAGVPAHADIS_GHPARFAVQMIHSYDLL
!!

960
960

logo

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DDDLEERMEEKSERLEAIRRGEVPVPGSF_AFDEKAREQLSWTYPH_HEVTQIRTKQSVSEIKRKREYEDEYSGRAPVKPADG
DDDLEERMEEKSERLEAIRRGEVPVPGSF_AFDEKAREQLSWTYPH_HEVTQIRTKQSVSEIKRKREYEDEYSGRAPVKPADG
DDDLEERMEEKSERLEAIRRGEVPVPGSF_AFDEKAREQLSWTYPH_QEVTQIRTKQSVSEIKRKREYEDEYSGRAPVKPADG
!!

1040
1040

logo

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consensus

SILYRRP_AFMMKKGLTAAEKGTAMHTVMQHIPLSHVPSIEEAEQTVHRLYEKELLTEEQKDAIDIEEIVQFFHTEIGGQL
SILYRRP_AFMMKKGLTAAEKGTAMHTVMQHIPLSHVPSIEEAEQTVHRLYEKELLTEEQKDAIDIEEIVQFFHTEIGGQL
SILYRRP_AFMMKKGLTAAEKGTAMHTVMQHIPLSHVPSIEEAEQTVHRLYEKELLTEEQKDAIDIEEIVQFFHTEIGGQL
!!

1120
1120

logo

NTDB id 339601 ES965 RS05770 WP 124059687.1
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consensus



1200
1200

logo

NTDB id 339601 ES965 RS05770 WP 124059687.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



1232
1232

- X non conserved
- X similar
- X \geq 50% conserved