

logo

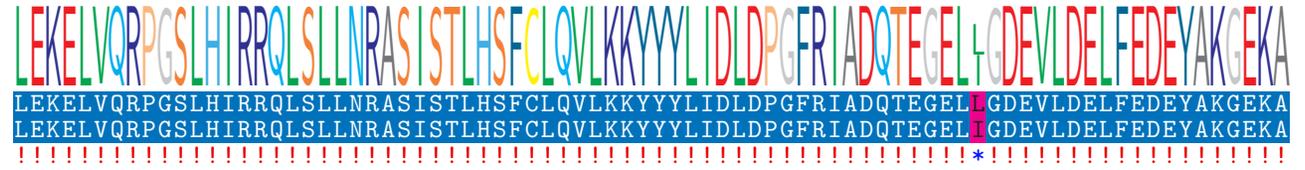
NTDB id 339763 ES967 RS05795 WP 128992275.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



80
80

logo

NTDB id 339763 ES967 RS05795 WP 128992275.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



160
160

logo

NTDB id 339763 ES967 RS05795 WP 128992275.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



240
240

logo

NTDB id 339763 ES967 RS05795 WP 128992275.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



320
320

logo

NTDB id 339763 ES967 RS05795 WP 128992275.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



400
400

logo

NTDB id 339763 ES967 RS05795 WP 128992275.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



480
480

logo

NTDB id 339763 ES967 RS05795 WP 128992275.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



560
560

logo

NTDB id 339763 ES967 RS05795 WP 128992275.1
NTDB id 119 BSU 10630 NP 388944.2
consensus

KLISSPFKVDGKKKTHRNQYRDIVILLRSMWPAPQIMEELRAQGIPVYANLTSYGFEEAVEVAVALSVLKVIDNPYQDI
KLISSPFKVDGKKKTHRNQYRDIVILLRSMWPAPQIMEELRAQGIPVYANLTSYGFEEAVEVAVALSVLKVIDNPYQDI
!!

640
640

logo

NTDB id 339763 ES967 RS05795 WP 128992275.1
NTDB id 119 BSU 10630 NP 388944.2
consensus

PLASVLRSPVIGADENELSLIRLENKKAPYYEAMKDYLAAAGDRSDELYQKLNFTFYGHLQKWRAFSKNHSVSELIWEVYRD
PLASVLRSPVIGADENELSLIRLENKKAPYYEAMKDYLAAAGDRNDELYQKLNFTFYGHLQKWRAFSKNHSVSELIWEVYRD
PLASVLRSPVIGADENELSLIRLENKKAPYYEAMKDYLAAAGDRSDELYQKLNFTFYGHLQKWRAFSKNHSVSELIWEVYRD
!!

720
720

logo

NTDB id 339763 ES967 RS05795 WP 128992275.1
NTDB id 119 BSU 10630 NP 388944.2
consensus

TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVRLMTIHSSKGLE
TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVRLMTIHSSKGLE
TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVRLMTIHSSKGLE
!!

800
800

logo

NTDB id 339763 ES967 RS05795 WP 128992275.1
NTDB id 119 BSU 10630 NP 388944.2
consensus

FPVVF_AAGLGRNFNMDLNKSYLLDKELGFGTKYIHPQLRISYPTLPLIAMKKMRRELLSEELRVLYVALTRAKEKLFEL
FPVVF_AAGLGRNFNMDLNKSYLLDKELGFGTKYIHPQLRISYPTLPLIAMKKMRRELLSEELRVLYVALTRAKEKLFEL
FPVVF_AAGLGRNFNMDLNKSYLLDKELGFGTKYIHPQLRISYPTLPLIAMKKMRRELLSEELRVLYVALTRAKEKLFEL
!!

880
880

logo

NTDB id 339763 ES967 RS05795 WP 128992275.1
NTDB id 119 BSU 10630 NP 388944.2
consensus

IGSCKDHQKQLAKWQASASQTDWLLPEFDYQARTYLDFIGPALARHRL_EDLAGVPAHADIS_GHPARFAVQMIHSYDLL
IGSCKDHQKQLAKWQASASQTDWLLPEFDYQARTYLDFIGPALARHRL_EDLAGVPAHADIS_GHPARFAVQMIHSYDLL
IGSCKDHQKQLAKWQASASQTDWLLPEFDYQARTYLDFIGPALARHRL_EDLAGVPAHADIS_GHPARFAVQMIHSYDLL
!!

960
960

logo

NTDB id 339763 ES967 RS05795 WP 128992275.1
NTDB id 119 BSU 10630 NP 388944.2
consensus

DDDLEERMEEKSERLEAIRRGEVPVGSFAFDEKAREQLSWTYPH_HEVTQIRTKQSVSEIKRKREYEDEYSGRAPVKPADG
DDDLEERMEEKSERLEAIRRGEVPVGSFAFDEKAREQLSWTYPH_HEVTQIRTKQSVSEIKRKREYEDEYSGRAPVKPADG
DDDLEERMEEKSERLEAIRRGEVPVGSFAFDEKAREQLSWTYPH_HEVTQIRTKQSVSEIKRKREYEDEYSGRAPVKPADG
!!

1040
1040

logo

NTDB id 339763 ES967 RS05795 WP 128992275.1
NTDB id 119 BSU 10630 NP 388944.2
consensus

SIL_FRRPAFMMKKGLTAAEKGTAMHTVMQHIPLSHVPSIEEAEQTVHRLYEKELLTEEQKDAIDIEEIVQFFHTEIGGQL
SIL_FRRPAFMMKKGLTAAEKGTAMHTVMQHIPLSHVPSIEEAEQTVHRLYEKELLTEEQKDAIDIEEIVQFFHTEIGGQL
SIL_FRRPAFMMKKGLTAAEKGTAMHTVMQHIPLSHVPSIEEAEQTVHRLYEKELLTEEQKDAIDIEEIVQFFHTEIGGQL
!!

1120
1120

logo

NTDB id 339763 ES967 RS05795 WP 128992275.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



1200
1200

logo

NTDB id 339763 ES967 RS05795 WP 128992275.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



1232
1232

- X non conserved
- X similar
- X ≥ 50% conserved