

logo

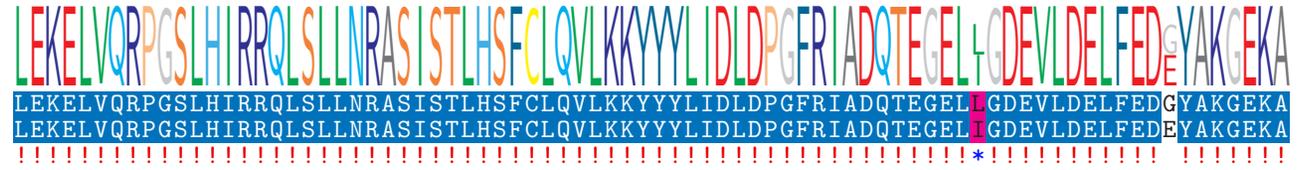
NTDB id 311292 D0808 RS08375 WP 153255881.1  
NTDB id 119 BSU 10630 NP 388944.2  
consensus



80  
80

logo

NTDB id 311292 D0808 RS08375 WP 153255881.1  
NTDB id 119 BSU 10630 NP 388944.2  
consensus



160  
160

logo

NTDB id 311292 D0808 RS08375 WP 153255881.1  
NTDB id 119 BSU 10630 NP 388944.2  
consensus



240  
240

logo

NTDB id 311292 D0808 RS08375 WP 153255881.1  
NTDB id 119 BSU 10630 NP 388944.2  
consensus



320  
320

logo

NTDB id 311292 D0808 RS08375 WP 153255881.1  
NTDB id 119 BSU 10630 NP 388944.2  
consensus



400  
400

logo

NTDB id 311292 D0808 RS08375 WP 153255881.1  
NTDB id 119 BSU 10630 NP 388944.2  
consensus



480  
480

logo

NTDB id 311292 D0808 RS08375 WP 153255881.1  
NTDB id 119 BSU 10630 NP 388944.2  
consensus



560  
560

logo

NTDB id 311292 D0808 RS08375 WP 153255881.1  
NTDB id 119 BSU 10630 NP 388944.2  
consensus

KLISSPFKVYDGKKKTHRNIQYRDIVILLRSMPWAPQIMEELRAQGIPVYANLTSYGYFEAVEVAVALSVLKVIDNPHYQDI  
KLISSPFKVYDGKKKTHRNIQYRDIVILLRSMPWAPQIMEELRAQGIPVYANLTSYGYFEAVEVAVALSVLKVIDNPHYQDI  
!!

640  
640

logo

NTDB id 311292 D0808 RS08375 WP 153255881.1  
NTDB id 119 BSU 10630 NP 388944.2  
consensus

PLASVLRSPVIGADENELSLIRLENKKAPYYEAMKDYLAAAGDR<sup>s</sup>DELYQKLNFTFYGHLQKWRAFSKNHSVSELIWEVYRD  
PLASVLRSPVIGADENELSLIRLENKKAPYYEAMKDYLAAAGDRNDELYQKLNFTFYGHLQKWRAFSKNHSVSELIWEVYRD  
PLASVLRSPVIGADENELSLIRLENKKAPYYEAMKDYLAAAGDRSDELYQKLNFTFYGHLQKWRAFSKNHSVSELIWEVYRD  
!!

720  
720

logo

NTDB id 311292 D0808 RS08375 WP 153255881.1  
NTDB id 119 BSU 10630 NP 388944.2  
consensus

TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVRLMTIHSSKGLE  
TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVRLMTIHSSKGLE  
TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVRLMTIHSSKGLE  
!!

800  
800

logo

NTDB id 311292 D0808 RS08375 WP 153255881.1  
NTDB id 119 BSU 10630 NP 388944.2  
consensus

FPVVF<sub>A</sub>AGLGRNFMMDLNKSYLLDKELGFGTKYIHPQLRISYPTLPLIAMKKMRRELLSEELRVLYVALTRAKEKLFEL  
FPVVF<sub>A</sub>AGLGRNFMMDLNKSYLLDKELGFGTKYIHPQLRISYPTLPLIAMKKMRRELLSEELRVLYVALTRAKEKLFEL  
FPVVFV<sub>A</sub>AGLGRNFMMDLNKSYLLDKELGFGTKYIHPQLRISYPTLPLIAMKKMRRELLSEELRVLYVALTRAKEKLFEL  
!!

880  
880

logo

NTDB id 311292 D0808 RS08375 WP 153255881.1  
NTDB id 119 BSU 10630 NP 388944.2  
consensus

IGSCKDHQKQLAKWQASASQTDWLLPEFDYQARTYLDFIGPALARHRL<sup>E</sup>DLAGVPAHADISGHPARFAVQMIHSYDLL  
IGSCKDHQKQLAKWQASASQTDWLLPEFDYQARTYLDFIGPALARHRL<sup>E</sup>DLAGVPAHADISGHPARFAVQMIHSYDLL  
IGSCKDHQKQLAKWQASASQTDWLLPEFDYQARTYLDFIGPALARHRL<sup>G</sup>DLAGVPAHADISGHPARFAVQMIHSYDLL  
!!

960  
960

logo

NTDB id 311292 D0808 RS08375 WP 153255881.1  
NTDB id 119 BSU 10630 NP 388944.2  
consensus

DDDLEERMEEKSERLEAIRRGEVPGSFAFDEKAREQLSWTYPH<sup>H</sup>EVTQIRTKQSVSEIKRKREYEDEYSGRAPVKPADG  
DDDLEERMEEKSERLEAIRRGEVPGSFAFDEKAREQLSWTYPH<sup>H</sup>EVTQIRTKQSVSEIKRKREYEDEYSGRAPVKPADG  
DDDLEERMEEKSERLEAIRRGEVPGSFAFDEKAREQLSWTYPH<sup>Q</sup>EVTQIRTKQSVSEIKRKREYEDEYSGRAPVKPADG  
!!

1040  
1040

logo

NTDB id 311292 D0808 RS08375 WP 153255881.1  
NTDB id 119 BSU 10630 NP 388944.2  
consensus

SILYRRP<sub>A</sub>FMMKKGLTAAEKGTAMHTVMQHIPLSHVPSIEEAEQTVHRLYEKELLTEEQKDAIDIEEIVQFFHTEIGGQL  
SILYRRP<sub>A</sub>FMMKKGLTAAEKGTAMHTVMQHIPLSHVPSIEEAEQTVHRLYEKELLTEEQKDAIDIEEIVQFFHTEIGGQL  
SILYRRP<sub>A</sub>FMMKKGLTAAEKGTAMHTVMQHIPLSHVPSIEEAEQTVHRLYEKELLTEEQKDAIDIEEIVQFFHTEIGGQL  
!!

1120  
1120

logo

NTDB id 311292 D0808 RS08375 WP 153255881.1  
NTDB id 119 BSU 10630 NP 388944.2  
consensus



1200  
1200

logo

NTDB id 311292 D0808 RS08375 WP 153255881.1  
NTDB id 119 BSU 10630 NP 388944.2  
consensus



1232  
1232

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