

logo

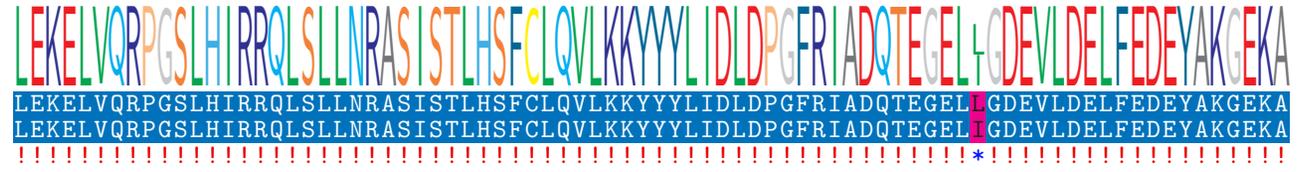
NTDB id 283406 C7M30 RS04960 WP 160244063.1  
NTDB id 119 BSU 10630 NP 388944.2  
consensus



80  
80

logo

NTDB id 283406 C7M30 RS04960 WP 160244063.1  
NTDB id 119 BSU 10630 NP 388944.2  
consensus



160  
160

logo

NTDB id 283406 C7M30 RS04960 WP 160244063.1  
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consensus



240  
240

logo

NTDB id 283406 C7M30 RS04960 WP 160244063.1  
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consensus



320  
320

logo

NTDB id 283406 C7M30 RS04960 WP 160244063.1  
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consensus



400  
400

logo

NTDB id 283406 C7M30 RS04960 WP 160244063.1  
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consensus



480  
480

logo

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consensus



560  
560

logo

NTDB id 283406 C7M30 RS04960 WP 160244063.1  
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consensus

KLISSPFKVYDGKKKTHRN IQYRDIVILLR SMPWAPQIMEELRAQGI PVYANL TSGYFEAVEVAVALSVLKVIDNPYQDI  
KLISSPFKVYDGKKKTHRN IQYRDIVILLR SMPWAPQIMEELRAQGI PVYANL TSGYFEAVEVAVALSVLKVIDNPYQDI  
!!

640  
640

logo

NTDB id 283406 C7M30 RS04960 WP 160244063.1  
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consensus

PLASVLRSPVIGADENELSLIRLENKKAPYYEAMKDYL AAGDR<sup>s</sup>DELYQKLNTFYGHLQKWRAFSKNHSVSELIWEVYRD  
PLASVLRSPVIGADENELSLIRLENKKAPYYEAMKDYL AAGDR<sup>s</sup>DELYQKLNTFYGHLQKWRAFSKNHSVSELIWEVYRD  
PLASVLRSPVIGADENELSLIRLENKKAPYYEAMKDYL AAGDR<sup>s</sup>DELYQKLNTFYGHLQKWRAFSKNHSVSELIWEVYRD  
!!

720  
720

logo

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TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVRLMTIHSSKGLE  
TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVRLMTIHSSKGLE  
TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVRLMTIHSSKGLE  
!!

800  
800

logo

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consensus

FPVVF<sup>A</sup>AGLGRNFNMMDLNKS YLLDKELGFGTKYIHPQLRISYPTLPLIAMKKMRRELLSEELRVLYVALTRAKEKLFEL  
FPVVF<sup>A</sup>AGLGRNFNMMDLNKS YLLDKELGFGTKYIHPQLRISYPTLPLIAMKKMRRELLSEELRVLYVALTRAKEKLFEL  
FPVVF<sup>A</sup>AGLGRNFNMMDLNKS YLLDKELGFGTKYIHPQLRISYPTLPLIAMKKMRRELLSEELRVLYVALTRAKEKLFEL  
!!

880  
880

logo

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consensus

IGSCKDHQKQLAKWQASASQTDWLLPEFD RYQARTYLDFIGPALARHRL<sup>E</sup>DLAGVPAH<sup>s</sup>DISGHPARFAVQMIHSDLL  
IGSCKDHQKQLAKWQASASQTDWLLPEFD RYQARTYLDFIGPALARHRL<sup>E</sup>DLAGVPAH<sup>s</sup>DISGHPARFAVQMIHSDLL  
IGSCKDHQKQLAKWQASASQTDWLLPEFD RYQARTYLDFIGPALARHRL<sup>E</sup>DLAGVPAH<sup>s</sup>DISGHPARFAVQMIHSDLL  
!!

960  
960

logo

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DDDLEERMEEKSERLEAIRRGE PVPGSF AFDEKAREQLSWTYPH<sup>Q</sup>EVTQIRTKQSVSEIKRKREYEDEYSGRAPVKPADG  
DDDLEERMEEKSERLEAIRRGE PVPGSF AFDEKAREQLSWTYPH<sup>Q</sup>EVTQIRTKQSVSEIKRKREYEDEYSGRAPVKPADG  
DDDLEERMEEKSERLEAIRRGE PVPGSF AFDEKAREQLSWTYPH<sup>Q</sup>EVTQIRTKQSVSEIKRKREYEDEYSGRAPVKPADG  
!!

1040  
1040

logo

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consensus

SILYRRP AFMMKKGLTAAEKGTAMHTVMQHIPL<sup>S</sup>HVPSIEEAEQTVHRLYEKELLTEEQKDAIDIEEIVQFFHTEIGGQL  
SILYRRP AFMMKKGLTAAEKGTAMHTVMQHIPL<sup>S</sup>HVPSIEEAEQTVHRLYEKELLTEEQKDAIDIEEIVQFFHTEIGGQL  
SILYRRP AFMMKKGLTAAEKGTAMHTVMQHIPL<sup>S</sup>HVPSIEEAEQTVHRLYEKELLTEEQKDAIDIEEIVQFFHTEIGGQL  
!!

1120  
1120

logo

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consensus



1200  
1200

logo

NTDB id 283406 C7M30 RS04960 WP 160244063.1  
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consensus



1232  
1232

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