



logo

NTDB id 283292 C7M28 RS17585 WP 160216283.1  
NTDB id 119 BSU 10630 NP 388944.2  
consensus

KLISSPFKVYDGGKKKTHRNIQYRDIVILLRSMPWAPQIMEELRAQGIPVYANLTSGYFEAVEVAVALSVLKVIDNPHYQDI  
KLISSPFKVYDGGKKKTHRNIQYRDIVILLRSMPWAPQIMEELRAQGIPVYANLTSGYFEAVEVAVALSVLKVIDNPHYQDI  
!!

640  
640

logo

NTDB id 283292 C7M28 RS17585 WP 160216283.1  
NTDB id 119 BSU 10630 NP 388944.2  
consensus

PLASVLRSPVIGADENELSLIRLENKKAPYYEAMKDYLEAGDRSDELYQKLNFTFYGHLQKWRAFASKNHSVSELIWEVYRD  
PLASVLRSPVIGADENELSLIRLENKKAPYYEAMKDYLEAGDRNDELYQKLNFTFYGHLQKWRAFASKNHSVSELIWEVYRD  
PLASVLRSPVIGADENELSLIRLENKKAPYYEAMKDYLEAGDRSDELYQKLNFTFYGHLQKWRAFASKNHSVSELIWEVYRD  
!!

720  
720

logo

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consensus

TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVRLMTIHSSKGLE  
TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVRLMTIHSSKGLE  
TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVRLMTIHSSKGLE  
!!

800  
800

logo

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consensus

FPVVF<sub>A</sub>AGLGRNFNMDLNKSYLLDKELGFGTKYIHPQLRISYPTLPLIAMKKMRRELLSEELRVLYVALTRAKEKLFEL  
FPVVF<sub>A</sub>AGLGRNFNMDLNKSYLLDKELGFGTKYIHPQLRISYPTLPLIAMKKMRRELLSEELRVLYVALTRAKEKLFEL  
FPVVF<sub>A</sub>AGLGRNFNMDLNKSYLLDKELGFGTKYIHPQLRISYPTLPLIAMKKMRRELLSEELRVLYVALTRAKEKLFEL  
!!

880  
880

logo

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IGSCKDHQKQLAKWQASASQTDWLLPEFDYQARTYLDFIGPALARHRLG<sub>E</sub>DLAGVPAHADIS<sub>G</sub>HPARFAVQMIHSYDLL  
IGSCKDHQKQLAKWQASASQTDWLLPEFDYQARTYLDFIGPALARHRLG<sub>E</sub>DLAGVPAHADIS<sub>G</sub>HPARFAVQMIHSYDLL  
IGSCKDHQKQLAKWQASASQTDWLLPEFDYQARTYLDFIGPALARHRLG<sub>E</sub>DLAGVPAHADIS<sub>G</sub>HPARFAVQMIHSYDLL  
!!

960  
960

logo

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DDDLEERMEEKSERLEAIRRGEVPVPGSF<sub>A</sub>FDEKAREQLSWTYPH<sub>H</sub>EVTQIRTKQSVSEIKRKREYEDEYSGRAPVKPADG  
DDDLEERMEEKSERLEAIRRGEVPVPGSF<sub>A</sub>FDEKAREQLSWTYPH<sub>H</sub>EVTQIRTKQSVSEIKRKREYEDEYSGRAPVKPADG  
DDDLEERMEEKSERLEAIRRGEVPVPGSF<sub>A</sub>FDEKAREQLSWTYPH<sub>H</sub>EVTQIRTKQSVSEIKRKREYEDEYSGRAPVKPADG  
!!

1040  
1040

logo

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consensus

SIL<sub>F</sub>RRPAFMMKKGLTAAEKGTAMHTVMQHIPLSHVPSIEEAEQTVHRLYEKELLTEEQKDAIDIEEIVQFFHTEIGGQL  
SIL<sub>F</sub>RRPAFMMKKGLTAAEKGTAMHTVMQHIPLSHVPSIEEAEQTVHRLYEKELLTEEQKDAIDIEEIVQFFHTEIGGQL  
SIL<sub>F</sub>RRPAFMMKKGLTAAEKGTAMHTVMQHIPLSHVPSIEEAEQTVHRLYEKELLTEEQKDAIDIEEIVQFFHTEIGGQL  
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1120  
1120

