

logo
NTDB id 246774 Bateq7PJ16 RS06145 WP 159376757.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



80
80

logo
NTDB id 246774 Bateq7PJ16 RS06145 WP 159376757.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



160
160

logo
NTDB id 246774 Bateq7PJ16 RS06145 WP 159376757.1
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consensus



240
240

logo
NTDB id 246774 Bateq7PJ16 RS06145 WP 159376757.1
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consensus



320
320

logo
NTDB id 246774 Bateq7PJ16 RS06145 WP 159376757.1
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consensus



400
400

logo
NTDB id 246774 Bateq7PJ16 RS06145 WP 159376757.1
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consensus



480
480

logo
NTDB id 246774 Bateq7PJ16 RS06145 WP 159376757.1
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consensus



560
560

logo

NTDB id 246774 Bateq7PJ16 RS06145 WP 159376757.1
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consensus

KLISSPFKVVYDGKKKTHRN IQYRDIVILLRSMPWAPQIMEELRAQGIPVYANLTSGYFEAVEVAVALSVLKVIDNPYQDI
KLISSPFKVVYDGKKKTHRN IQYRDIVILLRSMPWAPQIMEELRAQGIPVYANLTSGYFEAVEVAVALSVLKVIDNPYQDI
!!

640
640

logo

NTDB id 246774 Bateq7PJ16 RS06145 WP 159376757.1
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PLASVLRSPVIGADENELSLIRLENKKAPYYEAMKDYLAAAGDRSDELYQKLNIFYGHLQKWRAFASKNHVSSELIWEVYRD
PLASVLRSPVIGADENELSLIRLENKKAPYYEAMKDYLAAAGDRSDELYQKLNIFYGHLQKWRAFASKNHVSSELIWEVYRD
PLASVLRSPVIGADENELSLIRLENKKAPYYEAMKDYLAAAGDRSDELYQKLNIFYGHLQKWRAFASKNHVSSELIWEVYRD
!!

720
720

logo

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TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVRLMTIHSSKGLE
TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVRLMTIHSSKGLE
TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVRLMTIHSSKGLE
!!

800
800

logo

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FPWF_AAGLGRNFNMDLNKSYLLDKELGFGTKYIHPQLRISYPTLPLIAMKKMRRELLSEELRVLYVALTRAKEKLFEL
FPVVF_AAGLGRNFNMDLNKSYLLDKELGFGTKYIHPQLRISYPTLPLIAMKKMRRELLSEELRVLYVALTRAKEKLFEL
FPVVFV_AAGLGRNFNMDLNKSYLLDKELGFGTKYIHPQLRISYPTLPLIAMKKMRRELLSEELRVLYVALTRAKEKLFEL
!!

880
880

logo

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IGSCKDHQKQLAKWQASASQTDWLLPEFDYQAR_RTYLDFIGPALARHRDLGDLAGVPAHADISGHPARFAVQMIHSYDLL
IGSCKDHQKQLAKWQASASQTDWLLPEFDYQAR_KTYLDFIGPALARHRDLGDLAGVPAHADISGHPARFAVQMIHSYDLL
IGSCKDHQKQLAKWQASASQTDWLLPEFDYQAR_RTYLDFIGPALARHRDLGDLAGVPAHADISGHPARFAVQMIHSYDLL
!!

960
960

logo

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DDDLEERMEEKSERLEAIRRGEVPGSFAFDEKAREQLSWTYPH_HEVTQIRTKQSVSEIKRKREYEDEYSGRAPVKPADG
DDDLEERMEEKSERLEAIRRGEVPGSFAFDEKAREQLSWTYPH_HEVTQIRTKQSVSEIKRKREYEDEYSGRAPVKPADG
DDDLEERMEEKSERLEAIRRGEVPGSFAFDEKAREQLSWTYPH_QEVTQIRTKQSVSEIKRKREYEDEYSGRAPVKPADG
!!

1040
1040

logo

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SILYRRP_AFMMKKGLTAAEKGTAMHTVMQHIPLSHVPSIEEAEQTVHRLYEKELL_SEEQKDAIDIEEIVQFFHTEIGGQL
SILYRRP_AFMMKKGLTAAEKGTAMHTVMQHIPLSHVPSIEEAEQTVHRLYEKELL_SEEQKDAIDIEEIVQFFHTEIGGQL
SILYRRP_AFMMKKGLTAAEKGTAMHTVMQHIPLSHVPSIEEAEQTVHRLYEKELL_TEEQKDAIDIEEIVQFFHTEIGGQL
!!

1120
1120

logo

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consensus



1200
1200

logo

NTDB id 246774 Bateq7PJ16 RS06145 WP 159376757.1
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consensus



1232
1232

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