

logo

NTDB id 558 61092..61127( ) (comS)  
NTDB id 119 BSU 10630 NP 388944.2  
consensus

MNI PKPADSTWTDQWNAIVSTGQDILVAAAAGSGKTAVLVERMIRKITAENPIDVDRLVWFTNASAAEMKHRIA EALEKELVQRPG  
.....  
MNIPKPADSTWTDQWNAIVSTGQDILVAAAAGSGKTAVLVERMIRKITAENPIDVDRLVWFTNASAAEMKHRIA EALEKELVQRPG  
\*\*\*\*\*

0  
90

logo

NTDB id 558 61092..61127( ) (comS)  
NTDB id 119 BSU 10630 NP 388944.2  
consensus

SLHIRRQLSLLNRASISTLHSFCLQVLKYYLIDLDPGFRIADQTEGELIGDEVLDLFEDEYAKGEKAFFELVDRYTTDRHDLDLQFL  
.....  
SLHIRRQLSLLNRASISTLHSFCLQVLKYYLIDLDPGFRIADQTEGELIGDEVLDLFEDEYAKGEKAFFELVDRYTTDRHDLDLQFL  
\*\*\*\*\*

0  
180

logo

NTDB id 558 61092..61127( ) (comS)  
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consensus

VKQVYEYSRSHPNPEAWLESFVHLYDVSEKSAIEELPFYQYVKEDIAMVLNGAKEKLLRALELTKAPGGPAPRADNFLDDLAQIDELIQH  
.....  
VKQVYEYSRSHPNPEAWLESFVHLYDVSEKSAIEELPFYQYVKEDIAMVLNGAKEKLLRALELTKAPGGPAPRADNFLDDLAQIDELIQH  
\*\*\*\*\*

0  
270

logo

NTDB id 558 61092..61127( ) (comS)  
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consensus

QDDFSELYKRVPAVSFKRAKAVKGDEFDPALLDEATDLRNGAKKLEKDKTDYFTRSPEQHLKSLAEMKPVIE TLVQLVISYGRFEAAK  
.....  
QDDFSELYKRVPAVSFKRAKAVKGDEFDPALLDEATDLRNGAKKLEKDKTDYFTRSPEQHLKSLAEMKPVIE TLVQLVISYGRFEAAK  
\*\*\*\*\*

0  
360

logo

NTDB id 558 61092..61127( ) (comS)  
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consensus

QEKSIIDFSLDLEHYCLAILTAEENDKGEREPSEAAARFYQEQFHEVLVDEYQDTNLVQESILQLVTSGPEETGNLFMVGDKQSIYRFLAE  
.....  
QEKSIIDFSLDLEHYCLAILTAEENDKGEREPSEAAARFYQEQFHEVLVDEYQDTNLVQESILQLVTSGPEETGNLFMVGDKQSIYRFLAE  
\*\*\*\*\*

0  
450

logo

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consensus

PLLFLSKYKRFTESGEGTGRKIDLNKNFRSRADILDSTNFLFKQLMGGKIGEVYDEQAELKLGAAYPDNDDETETELLLIDNAEDTDASE  
.....  
PLLFLSKYKRFTESGEGTGRKIDLNKNFRSRADILDSTNFLFKQLMGGKIGEVYDEQAELKLGAAYPDNDDETETELLLIDNAEDTDASE  
\*\*\*\*\*

0  
540

logo

NTDB id 558 61092..61127( ) (comS)  
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consensus

EAAELETVQFEAKAIAKEIRKLISPPFKVYDGGKKKTHRNIQYRDIIMVLLRALPWSL  
.....  
EAAELETVQFEAKAIAKEIRKLISPPFKVYDGGKKKTHRNIQYRDIIMVLLRALPWSL  
\*\*\*\*\*

12  
630

logo

NTDB id 558 61092..61127( ) (comS)  
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consensus

KVIDNPYQDIPLASVLRSPIVGADENELSLIRLENKKAPYYEAMKDYLEAGDRSDELYQKLNIFYGHLQKWRAF SKNHVSSELIWEVYRD  
KVIDNPYQDIPLASVLRSPIVGADENELSLIRLENKKAPYYEAMKDYLEAGDRSDELYQKLNIFYGHLQKWRAF SKNHVSSELIWEVYRD  
\*\*\*\*\*

12  
720

logo

NTDB id 558 61092..61127( ) (comS)  
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consensus

TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGGDLGTARALSEQEDVRLMTIHSSKGLEFPVVFVAGLG  
TKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGGDLGTARALSEQEDVRLMTIHSSKGLEFPVVFVAGLG  
\*\*\*\*\*

12  
810

logo

NTDB id 558 61092..61127( ) (comS)  
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consensus

RNFNMDLNKSYLLDKELGFGTKYIHPQLRISYPTLPLIAMKKKMRRELLSEELRVLYVALTRAKEKLFILGSCDKHQKQLAKWQASASQ  
RNFNMDLNKSYLLDKELGFGTKYIHPQLRISYPTLPLIAMKKKMRRELLSEELRVLYVALTRAKEKLFILGSCDKHQKQLAKWQASASQ  
\*\*\*\*\*

12  
900

logo

NTDB id 558 61092..61127( ) (comS)  
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consensus

TDWLLPEFDRYQARTYLDFIGPALARHRDLGDLAGVPAHADISGHPARFAVQMIHSYDLLDDDL EERMEEKSERLEAIRRGEVPVGSFAF  
TDWLLPEFDRYQARTYLDFIGPALARHRDLGDLAGVPAHADISGHPARFAVQMIHSYDLLDDDL EERMEEKSERLEAIRRGEVPVGSFAF  
\*\*\*\*\*

12  
990

logo

NTDB id 558 61092..61127( ) (comS)  
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consensus

DEKAREQLSWTYPHQEVTVIRTKQSVSEIKRKREYEDEYSGRAPVKPADGSI LYRRPAFMMKKGLTAAEKGTAMHTVMQHIPLSHVPSIE  
DEKAREQLSWTYPHQEVTVIRTKQSVSEIKRKREYEDEYSGRAPVKPADGSI LYRRPAFMMKKGLTAAEKGTAMHTVMQHIPLSHVPSIE  
\*\*\*\*\*

12  
1080

logo

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consensus

EAEQTVHRLYEKELLTEEQKDAIDIEEIVQFFHTEIGGQLIGAKWKDREIPFSLALPAKEIYPAHAHADEPLLVQGIIDCLYETEDGLYL  
EAEQTVHRLYEKELLTEEQKDAIDIEEIVQFFHTEIGGQLIGAKWKDREIPFSLALPAKEIYPAHAHADEPLLVQGIIDCLYETEDGLYL  
\*\*\*\*\*

12  
1170

logo

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consensus

LDYKSDRIEGKFQHGFE GAAPILKKRYETQIQLYTKAVEQIAKTKVKGCALYFFDGGHILTL  
LDYKSDRIEGKFQHGFE GAAPILKKRYETQIQLYTKAVEQIAKTKVKGCALYFFDGGHILTL  
\*\*\*\*\*

12  
1232

 non conserved  
 similar  
  $\geq 50\%$  conserved