

logo

NTDB id 339860 ES968 RS12320 WP 129134317.1  
NTDB id 352 BSU 24240 NP 390304.2  
consensus

MLAELSIKNFAIIEELTVSFERGLTVLTGETGAGKSIIDAIISLLVGGRSSEFVRYGEAKAELEGLFLLLESGHPVLGVC  
MLAELSIKNFAIIEELTVSFERGLTVLTGETGAGKSIIDAIISLLVGGRSSEFVRYGEAKAELEGLFLLLESGHPVLGVC  
MLAELSIKNFAIIEELTVSFERGLTVLTGETGAGKSIIDAIISLLVGGRSSEFVRYGEAKAELEGLFLLLESGHPVLGVC  
!!

80  
80

logo

NTDB id 339860 ES968 RS12320 WP 129134317.1  
NTDB id 352 BSU 24240 NP 390304.2  
consensus

AEQGI DVSD EIMVRRD I STSGKSVCRVNGKLV TIASLREI GRLLLD IHGQHDNQLL MEDENHLQLL DKFAGAEVESALK  
AEQGIDVSD EIMVRRD I STSGKSVCRVNGKLV TIASLREI GRLLLD IHGQHDNQLL MEDENHLQLL DKFAGAEVESALK  
AEQGIDVSD EIMVRRD I STSGKSVCRVNGKLV TIASLREI GRLLLD IHGQHDNQLL MEDENHLQLL DKFAGAEVESALK  
!!

160  
160

logo

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consensus

TYQEGYQRYMKLLKLLKQLSESEQEMAHRLDLIQFQLEEIESAKLELNEDEQLQEERQQISNFEKIYESLQNAYNALRSE  
TYQEGYQRYMKLLKLLKQLSESEQEMAHRLDLIQFQLEEIESAKLELNEDEQLQEERQQISNFEKIYESLQNAYNALRSE  
TYQEGYQRYMKLLKLLKQLSESEQEMAHRLDLIQFQLEEIESAKLELNEDEQLQEERQQISNFEKIYESLQNAYNALRSE  
!!

240  
240

logo

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consensus

QGGLDWVGMASQALEDISDINEPLKKMSESVNSYLLLEDATFQMRNMLDELEFDPECLNYIETRLNEIKQLKRKYGATV  
QGGLDWVGMASQALEDISDINEPLKKMSESVNSYLLLEDATFQMRNMLDELEFDPECLNYIETRLNEIKQLKRKYGATV  
QGGLDWVGMASQALEDISDINEPLKKMSESVNSYLLLEDATFQMRNMLDELEFDPECLNYIETRLNEIKQLKRKYGATV  
!!

320  
320

logo

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consensus

EDILEYASKIEEEIDQIENRDSHLQSLKKELDVSGKDVAVEAANVSQIRKTWAKKLADEIHRELKSLYMEKSTFDTEFKV  
EDILEYASKIEEEIDQIENRDSHLQSLKKELDVSGKDVAVEAANVSQIRKTWAKKLADEIHRELKSLYMEKSTFDTEFKV  
EDILEYASKIEEEIDQIENRDSHLQSLKKELDVSGKDVAVEAANVSQIRKTWAKKLADEIHRELKSLYMEKSTFDTEFKV  
!!

400  
400

logo

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consensus

RTASRNEEAPLVNGQPVQLTEQGI DLVKFLISTNTGEPLKSLSKVASGGELSRVMLAIKSI FSSQQDVTSIIFDEVD TG  
RTASRNEEAPLVNGQPVQLTEQGI DLVKFLISTNTGEPLKSLSKVASGGELSRVMLAIKSI FSSQQDVTSIIFDEVD TG  
RTASRNEEAPLVNGQPVQLTEQGI DLVKFLISTNTGEPLKSLSKVASGGELSRVMLAIKSI FSSQQDVTSIIFDEVD TG  
!!

480  
480

logo

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consensus

SGRVAQAIAEKIHKVSI GSVQLCITHLPQVAAMADTHLYIAKELKDGRTTTRVKPLSKQEKVAEIGRMIAGVEVTDLTKR  
SGRVAQAIAEKIHKVSI GSVQLCITHLPQVAAMADTHLYIAKELKDGRTTTRVKPLSKQEKVAEIGRMIAGVEVTDLTKR  
SGRVAQAIAEKIHKVSI GSVQLCITHLPQVAAMADTHLYIAKELKDGRTTTRVKPLSKQEKVAEIGRMIAGVEVTDLTKR  
!!

560  
560

logo

NTDB id 339860 ES968 RS12320 WP 129134317.1  
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consensus

HAKELLKQADQVKTG  
HAKELLKQADQVKTG  
HAKELLKQADQVKTG  
!!!!!!!!!!!!!!!!

576  
576

non conserved  
 similar  
 ≥ 50% conserved