

logo

NTDB id 1026772 AB3470 RS13250 WP 029726732.1
NTDB id 352 BSU 24240 NP 390304.2
consensus

MLAELSIKNFAIIEELTVSFERGLTVLTGETGAGKSIIDAISLLVGGGRGSSEFVRYGEAKAELEGLFLLLESHPVLGVC
MLAELSIKNFAIIEELTVSFERGLTVLTGETGAGKSIIDAISLLVGGGRGSSEFVRYGEAKAELEGLFLLLESHPVLGVC
MLAELSIKNFAIIEELTVSFERGLTVLTGETGAGKSIIDAISLLVGGGRGSSEFVRYGEAKAELEGLFLLLESHPVLGVC
!!

80
80

logo

NTDB id 1026772 AB3470 RS13250 WP 029726732.1
NTDB id 352 BSU 24240 NP 390304.2
consensus

AEQGI DVSD EMI V MRRDI STSGKSVCRVNGKLVTIASLREI GRLLLDIHGQHDNQLLMEDENHLQLLDKFAGAEVESALK
AEQGIDVSD EMI V MRRDISTSGKSVCRVNGKLVTIASLREI GRLLLDIHGQHDNQLLMEDENHLQLLDKFAGAEVESALK
AEQGIDVSD EMI V MRRDISTSGKSVCRVNGKLVTIASLREI GRLLLDIHGQHDNQLLMEDENHLQLLDKFAGAEVESALK
!!

160
160

logo

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consensus

TYQEGYQRYM KLLK LKQLSESEQEMAHRLDLIQFQLEEI ESAKLELNEDQLQEERQQISNFEKIYESLQAYNALRSE
TYQEGYQRYM KLLK LKQLSESEQEMAHRLDLIQFQLEEI ESAKLELNEDQLQEERQQISNFEKIYESLQAYNALRSE
TYQEGYQRYM KLLK LKQLSESEQEMAHRLDLIQFQLEEI ESAKLELNEDQLQEERQQISNFEKIYESLQAYNALRSE
!!

240
240

logo

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consensus

QGGLDIVGMSAQL EDI SDINEPLK KMSSES VSN SYLLEDATFQMRNMLDELEFDPERLNYIETRLNEIKQLKRKYGATV
QGGLDIVGMSAQL EDI SDINEPLK KMSSES VSN SYLLEDATFQMRNMLDELEFDPERLNYIETRLNEIKQLKRKYGATV
QGGLDIVGMSAQL EDI SDINEPLK KMSSES VSN SYLLEDATFQMRNMLDELEFDPERLNYIETRLNEIKQLKRKYGATV
!!

320
320

logo

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consensus

EDILEYASKIEEEEIDQIENRDSHLQSLKKELDVSGKDVAVEAANVSQIRKTWAKKLAD EIHRELKSLYMEKSTFDTEFKV
EDILEYASKIEEEEIDQIENRDSHLQSLKKELDVSGKDVAVEAANVSQIRKTWAKKLAD EIHRELKSLYMEKSTFDTEFKV
EDILEYASKIEEEEIDQIENRDSHLQSLKKELDVSGKDVAVEAANVSQIRKTWAKKLAD EIHRELKSLYMEKSTFDTEFKV
!!

400
400

logo

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consensus

RTASRNEEAPLVNGQPVLTEQGI DLVKFLISTNTGEPLKSLSKVASGGELSRVMLA IKSIFSSQQDVTSIIFDEVDTGV
RTASRNEEAPLVNGQPVLTEQGI DLVKFLISTNTGEPLKSLSKVASGGELSRVMLA IKSIFSSQQDVTSIIFDEVDTGV
RTASRNEEAPLVNGQPVLTEQGI DLVKFLISTNTGEPLKSLSKVASGGELSRVMLA IKSIFSSQQDVTSIIFDEVDTGV
!!

480
480

logo

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consensus

SGRVAQAIAEKIHKVSI GSVL CITHLPQVAAMADTHLYIAKELKDGRITTTTRVKPLSKQEKVAEIGRMIAGVEVTDLTKR
SGRVAQAIAEKIHKVSI GSVL CITHLPQVAAMADTHLYIAKELKDGRITTTTRVKPLSKQEKVAEIGRMIAGVEVTDLTKR
SGRVAQAIAEKIHKVSI GSVL CITHLPQVAAMADTHLYIAKELKDGRITTTTRVKPLSKQEKVAEIGRMIAGVEVTDLTKR
!!

560
560

logo

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consensus

HAKE~~LL~~KQADQVKTG
HAKE~~LL~~KQADQVKTG
HAKE~~LL~~KQADQVKTG
!!!!!!!!!!!!!!!!!!!!

576
576

non conserved
 similar
 $\geq 50\%$ conserved