```
logo
NTDB id 101156 ACKTTY RS06435 WP 308703019.1  MLQELSITNFAIIEQLDIAFENGMTVLTGETGAGKSIIIDAVGLLAGGRGSQEFIRTGADKAVLQGMFILPADGVTAHML   80
NTDB id 352 BSU 24240 NP 390304.2             MLAELSIKNFAIIEELTVSFERGLTVLTGETGAGKSIIIDAISLLVGGRGSSEFVRYGEAKAELEGLFLLESGHPVLGVC   80
consensus                                     !! !!!! !!!!!! ! * !! !*!!!!!!!!!!!!!!!!* !! !!!!! !!*! !  !! ! !*!*!*  **  ***

logo
NTDB id 101156 ACKTTY RS06435 WP 308703019.1  DEAGIDHGDNTVILQREIAKSGRNTCRINGMLVNTTTLKRIGETIVDIHGQNEHQELMQPDKHLGLLDEFAAHKIHKLKA  160
NTDB id 352 BSU 24240 NP 390304.2             AEQGIDVSDEMIVMRRDISTSGKSVCRVNGKLVTIASLREIGRLLLDIHGQHDNQLLMEDENHLQLLDKFAGAEVESALK  160
consensus                                      ! !!!* !  *** !*!  !!*  !!*!! !!   *!* !!  **!!!!!***! !! ** !! !!! !!** **

logo
NTDB id 101156 ACKTTY RS06435 WP 308703019.1  QYTQEYEAYQQLNQELRQKNANEKEWAQRLDMLNFQVEEISAAQVKVGEEADLTAERDRLDNYQTINQALQQSYNLLVAE  240
NTDB id 352 BSU 24240 NP 390304.2             TYQEGYQRYVKLLKKLKQLSESEQEMAHRLDLIQFQLEEIESAKLELNEDEQLQEERQQISNFEKIYESLQNAYNALRSE  240
consensus                                      !   !  !  !   !*!    !  ! !*!!!***!!*!!!  ! * * !*  !  !!  * !*  !   !!* !! !  !

logo
NTDB id 101156 ACKTTY RS06435 WP 308703019.1  EETPGAVDVVGEAMNALEPIANLDSAFNDITVNVKNAFYALQDATGQISNQLDLQEFDEGRLDEIEQRLDVLSQLKRKYG  320
NTDB id 352 BSU 24240 NP 390304.2             ...QGGLDWVGMASAQLEDISDINEPLKKMSESVSNSYYLLEDATFQMRNMLDELEFDPERLNYIETRLNEIKQLKRKYG  317
consensus                                     ****!**! !! !   !!*!  *  *   **  ! ! *! ! !!! !* ! !!  !!!* !!  !! !!  * !!!!!!!

logo
NTDB id 101156 ACKTTY RS06435 WP 308703019.1  DSEQQILDYYQKISTELAKMTDSEENSEGLSQRVADKKAALLKTGQALSDKRRTCAKSLQRQIHQELKDLYMDKAIFEVR  400
NTDB id 352 BSU 24240 NP 390304.2             ATVEDILEYASKIEEEIDQIENRDSHLQSLKKELDSVGKDVAVEAANVSQIRKTWAKKLADEIHRELKSLYMEKSTFDTE  397
consensus                                      *   !!*!  !!  !*  *   * *   !   *     *   *  *!  !*!*!! !   !! !!! !!!*!  !*

logo
NTDB id 101156 ACKTTY RS06435 WP 308703019.1  FKPTK..............GQIYRSGLDSVEFYIQTNPGERMRPLAKIASGGELSRMMLALKTIFSESQGITSIIFDEVD  466
NTDB id 352 BSU 24240 NP 390304.2             FKVRTASRNEEAPLVNGQPVQLTEQGIDLVKFLISTNTGEPLKSLSKVASGGELSRVMLAIKSIFSSQQDVTSIIFDEVD  477
consensus                                     !!*  ************** !*   !*! ! ! ! !!*!!****! !*!!!!!!!!*!!!*!*!!!   ! *!!!!!!!!!

logo
NTDB id 101156 ACKTTY RS06435 WP 308703019.1  TGVSGRVAQAIAEKISGIAKFSQVLCITHLPQVAAMADHHYFIAKKITGRRTKTSVTKLAQPDRVQELARMLAGTKVTKL  546
NTDB id 352 BSU 24240 NP 390304.2             TGVSGRVAQAIAEKIHKVSIGSQVLCITHLPQVAAMADTHLYIAKELKDGRTTTRVKPLSKQEKVAEIGRMIAGVEVTDL  557
consensus                                     !!!!!!!!!!!!!!!* *   !!!!!!!!!!!!!!!!!!*! *!!! *   !! ! ! *!  ***! !**!!*!!  !! !
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logo
                                                  TLRHAKELLRQAGQVKVT
                                                  SKEHAEELLKLADEEKTEG

NTDB id 101156 ACKTTY RS06435 WP 308703019.1      SLEHAEELLRLAGEEKVE.   564
NTDB id 352 BSU 24240 NP 390304.2                 TKRHAKELLKQADQVKTTG   576
consensus                                         *  !! !!!* !   !  *
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X non conserved

X similar

X $\geq$ 50% conserved