```
logo
NTDB id 23958 STR RS02265 WP 011226912.1  MENKPKTSSYKRLKPYIKGFQVPFVLAIVGAIISAVITVIGPDKLKEITNLITDGITPTKMGVIPGIDLDKVGSIALTLAVLYAI   85
NTDB id 370 SMU RS04260 WP 002263294.1    MEN.KKKSLLSQMAPYLKGYKALFGLAVIFTIVSSTITVIGPDRLKEMTDTMTKGL.......AGKIDLDKIGEIALTLALLYFA   77
consensus                                 !!!**! !    * !!*!!*  *! !!**  !*!  !!!!!!!*!!!*!  *! !******** * !!!!!*! !!!!!!*!!

logo
NTDB id 23958 STR RS02265 WP 011226912.1  SAIVGYLQSFTVVTITQRFSQRFRTAIQKKINTVPLNYFDSHSQGDILSRVTNDVDLLGQSLSQGLGTLITSSILLVAAIIMMFY  170
NTDB id 370 SMU RS04260 WP 002263294.1    GALVSYTASFIVSTLIQKFSQRLRNAIADKINKVPLKYFDSHSQGDTLSRVTNDVDLMTQSFNQSLVSMVAAIILLIGSIFMMIK  162
consensus                                  !*! !  !! ! !* !*!!!! ! !!  !!! !!! !!!!!!!!! !!!!!!!!!!* !!  ! ! ***   !!!** ! !!

logo
NTDB id 23958 STR RS02265 WP 011226912.1  SNVTMAFTAIGSVLIGFVLVAFIMGFSQPLFKRQQENLANINGYIEEIYTGQAVVTSYNASKESSQDFKGLNDKLYNSMWQSQFI  255
NTDB id 370 SMU RS04260 WP 002263294.1    TNGALAATAILSVFAGFVLSTVIMAKSQPLFKKQQANLADVSGYVEEVYSGHNVVSSYNAIQQSKKQFENLNDQLFASMWKSQFF  247
consensus                                 *!  *! !!! !!  !!!!   !!* !!!!!!*!! !!! * !!*!!*!*!* !!*!!!!   !   !  !!! !* !!! !!!

logo
NTDB id 23958 STR RS02265 WP 011226912.1  SGIMMPLMIFIGNFGYVMVCLVGAVKVIDGSLTIGDVVAFMTYVRIFSQPLSQIAQGITQLQSATAAIGSIFEFLEEKDMDDESV  340
NTDB id 370 SMU RS04260 WP 002263294.1    SGIMMPLMQFIGNFGYVMVCIVGATMAINGDITMGTIVAFMTYVRIFTQPIAQIAQGITQLQSANAAMGRVFEFLDEEEIEDENH  332
consensus                                 !!!!!!!! !!!!!!!!!!!*!!!   ! ! *!*! *!!!!!!!!!!*!!* !!!!!!!!!!!! !!*! *!!!!*! ***!! *

logo
NTDB id 23958 STR RS02265 WP 011226912.1  KTAELTDTKGQVVFDHVSFGYTPEKTIIHDFSALAKPGQKIAIVGPTGAGKTTIVNLLMKFYNIDKGRITIDGVDTNDMKRETVH  425
NTDB id 370 SMU RS04260 WP 002263294.1    KVKQLEKVEGNVNFDNVFFGYSPDKTIIHDFSAHAKAGQKIAIVGPTGAGKTTIVNLLMRFYEVDRGMISIDGVNIHDMTRKEVH  417
consensus                                 !   !    !*! !!*! !!!*!*!!!!!!!!!*!!*!!!!!!!!!!
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X non conserved
X similar
X $\geq$ 50% conserved