```
logo
NTDB id 65513 ANG RS03030 WP 003035495.1  MENKKSSLFSQVEPYIKGFQFPLLVAFVGVILSSVITVYGPVKLKEITNLITEGMRSRIDLKAISNIALFLAVLYVVGAVLNYMQ   85
NTDB id 370 SMU RS04260 WP 002263294.1    MENKKKSLLSQMAPYLKGYKALFGLAVIFTIVSSTITVIGPDRLKEMTDTMTKGLAGKIDLDKIGEIALTLALLYFAGALVSYTA   85
consensus                                 !!!!! !! !!* !!*!!*  *  *! *  !*!! !!! !! *!!!*!  *! !*  *!!!  !  !!! !!*!!  !!** !

logo
NTDB id 65513 ANG RS03030 WP 003035495.1  SFIISSVVQHFSKRLRTAIAEKINKLPLRYFDSHSQGDTLSRVTNDVDTVGQSLNQSLGTVISSSLLLIAVLFMMFYNNVVLSFV  170
NTDB id 370 SMU RS04260 WP 002263294.1    SFIVSTLIQKFSQRLRNAIADKINKVPLKYFDSHSQGDTLSRVTNDVDLMTQSFNQSLVSMVAAIILLIGSIFMMIKTNGALAAT  170
consensus                                 !!!*!***!*!! !!! !!!*!!!!*!!*!!!!!!!!!!!!!!!!!!! * !! !!!! ***   *!!!* *!!!   !  !

logo
NTDB id 65513 ANG RS03030 WP 003035495.1  TIGSVLIGFVFVALIMGKSQRFFRTQQENLASVNGYVEEMYSGHNVIVSYNAAEESKATFQTLNTNLYNSMWKSQFISGIMMPLM  255
NTDB id 370 SMU RS04260 WP 002263294.1    AILSVFAGFVLSTVIMAKSQPLFKKQQANLADVSGYVEEVYSGHNVVSSYNAIQQSKKQFENLNDQLFASMWKSQFFSGIMMPLM  255
consensus                                  ! !!  !!!   *!!*!!!* !* !! !!! ! !!!!!*!!!!!!* !!!!   !!  !  !! *!* !!!!!!! !!!!!!!!

logo
NTDB id 65513 ANG RS03030 WP 003035495.1  MFVGNFGYVMVVLVGASMALNGDVTMGTIVAFMVYVRIFSQPLSQIAQGITTLQQASAAMGRVFEFLAEPEMENDEHKAQQLTTL  340
NTDB id 370 SMU RS04260 WP 002263294.1    QFIGNFGYVMVCIVGATMAINGDITMGTIVAFMTYVRIFTQPIAQIAQGITQLQSANAAMGRVFEFLDEEEIEDENHKVKQLEKV  340
consensus                                  !*!!!!!!!!**!!!*!!*!!!*!!!!!!!!! !!!!!*!!* !!!!!!! !! ! !!!!!!!!!! !*!*! * !!  !!  *

logo
NTDB id 65513 ANG RS03030 WP 003035495.1  KGDVTFDKVFFGYNPDRTIIHDFSANAKAGQKIAIVGPTGAGKTTIVNLLMKFYEIDKGRITIDGVDTKQMKRSEVHDAFSMVLQ  425
NTDB id 370 SMU RS04260 WP 002263294.1    EGNVNFDNVFFGYSPDKTIIHDFSAHAKAGQKIAIVGPTGAGKTTIVNLLMRFYEVDRGMISIDGVNIHDMTRKEVHDAFAMVLQ  425
consensus                                  ! ! !! !!!!! !!*!!!!!!!!!*!!!!!!!!!!!!!!!!!!
```

X non conserved
X similar
X $\geq$ 50% conserved