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logo

NTDB id 359 SMU RS01430 WP 002263524.1  MDPKFLQSAEFYRRRYHNFATLLIVPLVCLIIFLVIFLCFAKKEITVISTGEVAPTKVVDVIQSYSDSSIIKNNLDNNAAVEKGD   85
NTDB id 404 SMU RS08660 WP 002263739.1  MNPNLFKSAEFYRRRYHNFATVLIIPLVCLLVFLFIFLLFAKKEVTVLSDGEISPIKVIDVIQSKSDYPISENNLTDNHAVKKRD   85
consensus                               ! !    !!!!!!!!!!!!!!*!!*!!!!!**!! !!!*!!!!!*!!*! !!* ! !!*!!!!! !! *!  !!!  !*!! ! !

logo

NTDB id 359 SMU RS01430 WP 002263524.1  VLIEYSENASPNRQTEQKNIIKERQKREEKEK...KKHQK...........SKKKKSKS...........K..KASKDKK....   139
NTDB id 404 SMU RS08660 WP 002263739.1  VLIQYVNPAANTRQGGNNQNQNQSQNKQNKNQQQNNKNQKNGNQGQKNQNQNQNKKDQKNNNQSQQNKQGQKKTNQSQNQQNKNQ   170
consensus                               !!! !  *! * !!  * *     ! *  !  *** !*!!***********   !!  ! ***********!**  !    ****

logo

NTDB id 359 SMU RS01430 WP 002263524.1  ..KKSKDKESS....SDDENETKKVSIFASEDGIIHTNPKYDGANIIPKQTEIAQIYPDIQKTRKVLITYYASSDDVVSMKKGQT   218
NTDB id 404 SMU RS08660 WP 002263739.1  NKNQSQNKQNGNQNQNNWNQNNQNDKVTASQDGVIHTNPKYEGENLIPRGTEIAQLYPDINKTRQVLITYYVSSKDVTRLKQGQK   255
consensus                               **  !  !   ****    *      * !! !!*!!!!!!!*! !*!!* !!!!!*!!!!*!!! !!!!!! !! !!  *! !!

logo

NTDB id 359 SMU RS01430 WP 002263524.1  ARLSLEKKGNDKVVIEGKINNVASSATTTKKGNLFKVTAKVKVSKKNSKLIKYGMTGKTVTVIDKKTYFDYFKDKLLHKMDN   300
NTDB id 404 SMU RS08660 WP 002263739.1  THLALTKVGNDQISIRGKINSIASAATTTKKGNVFKVTAKAKVSKKDSQTIKYGLQGKTITVVEKKTFFDYYKDKLLNKNQK   337
consensus                                *! ! ! !!! * ! !!!! *!! !!!!!!!!*!!!!!! !!!!! !  !!!!* !!!*!!**!!!*!!!*!!!!!*!
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X non conserved

X similar

X $\geq$50% conserved