```
logo
NTDB id 1210 C694 RS01710 WP 001862983.1          MKSHFQYSTLENIPKAFDILKDPPKKLYCVGDTKLLDTPLKVAIIGTRRPTPYSKQHTITLARELAKNGAVIVSGGALGV   80
NTDB id 115423 CFVI03293 RS05655 WP 002849743.1  ..MDFI...T..DILSLKKLKKPVNKLYYKGNLELLNFP.KVAIVGSRKCTSYTKNIVLSLASTLKNYGICVVSGAAIGT   72
NTDB id 1239 Cj0634 YP 002344063.1                MASEIL...PDEFLELFKDLKKPPKKLHYKGNLSLLKQD.KIAIIGSRRMSVYTKNCVFSLASMLKNAHLCVVSGGALGV   76
consensus                                         * ****    *  *  ** !!*!**!!***!** !!  * !*!!*!*!* * !*!* ***!!* !** ****!!!*!*!*

logo
NTDB id 1210 C694 RS01710 WP 001862983.1          DIIAQENALPKTIMLSPCSLDFIYPTNNHKVIQEIAQNGLILSEYEKDFMPIKGSFLARNRLVIALSDVVIIPQADLKSG  160
NTDB id 115423 CFVI03293 RS05655 WP 002849743.1  DIHAHEGAFPNTIAVFGNGLNNIYPAQNSKMIEKIYKDALALSEYEPNTFATPWSFLERNRIVVSLSDAVIIAEADIKSG  152
NTDB id 1239 Cj0634 YP 002344063.1                DITASMAAMPNTIGIFANGLDQIYPRTNEKIIKQIYENALALSEYEDDYLPKNYDFLLRNRLVIALSKAVVVAQADIKSG  156
consensus                                         !! ! **!*!*!!*******!**!!! *! !*!  !* **!*!!!!! ****  **!! !!!*!**!!**!****!!*!!!

logo
NTDB id 1210 C694 RS01710 WP 001862983.1          SMSSARLAQKYQKPLFVLPQRLNESDGTNELLEKGQAQGIFNIQNFINTLLKDYHLKEMPEMEDEFLEYCAKNPSYEEAY  240
NTDB id 115423 CFVI03293 RS05655 WP 002849743.1  SMSSARLALDMGIPLFVLPQRFNESNGTNDLLSKNRANLIADFDEFASKF....GSLNSDLSKDDVLEFIKINQNFDECY  228
NTDB id 1239 Cj0634 YP 002344063.1                SMQSAKLALELNKPLYVLPQRLGESTATNLLLKENKAKLICDFKEFVSEF....ASID..MNQDEFLEFCKKGVSVDEAL  230
consensus                                         !!*!!*!!*****!!*!!!!!**!! *!!*!! ***!**! ** *!** *    ****  *  !**!!***** ***!**

logo
NTDB id 1210 C694 RS01710 WP 001862983.1          LKFGDKLLEYELLGKIKRINHIVVLA.   266
NTDB id 115423 CFVI03293 RS05655 WP 002849743.1  ARFGDALYEYELEGKISIDRVYIRILS   255
NTDB id 1239 Cj0634 YP 002344063.1                KIYGQKVYEYELEGKISIEGLFIRVLA   257
consensus                                          **!****!!!!*!!!*** ******
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X non conserved

X similar

X ≥ 50% conserved