```
logo
NTDB id 1210 C694 RS01710 WP 001862983.1        MKSHFQYSTLENIPKAFDILKDPPKKLYCVGDTKLLDTPLKVAIIGTRRPTPYSKQHTITLARELAKNGAVIVSGGALGV   80
NTDB id 1161386 AABJ57 RS09505 WP 002776372.1  ....MQSELPSAYLELFKELKNPPKKLYYKGNLKLLEQ.RKIAIIGSRRMSTYTKNCVFELSNLLKNSKISVVSGGALGV   75
NTDB id 1239 Cj0634 YP 002344063.1              ...MASEILPDEFLELFKDLKKPPKKLHYKGNLSLLKQ.DKIAIIGSRRMSVYTKNCVFSLASMLKNAHLCVVSGGALGV   76
consensus                                            *  *** ****!**!! !!!!!***!***!!**  !*!!!!*!!** !*!****!* *!**  * *!!!!!!!!

logo
NTDB id 1210 C694 RS01710 WP 001862983.1        DIIAQENALPKTIMLSPCSLDFIYPTNNHKVIQEIAQNGLILSEYEKDFMPIKGSFLARNRLVIALSDVVIIPQADLKSG  160
NTDB id 1161386 AABJ57 RS09505 WP 002776372.1  DISASMAAMPNTIGVFANGLDEIYPRSNEKIIKQIYENALALSENESYYKPKPYDFLLRNRLIVALSEAVVVAQADLQSG  155
NTDB id 1239 Cj0634 YP 002344063.1              DITASMAAMPNTIGIFANGLDQIYPRTNEKIIKQIYENALALSEYEDDYLPKNYDFLLRNRLVIALSKAVVVAQADIKSG  156
consensus                                       !!*!***!*!*!!******!! !!!**!*!*!**!**!*!!!!*! ***!* **!!*!!!!**!!!**!***!!!**!!

logo
NTDB id 1210 C694 RS01710 WP 001862983.1        SMSSARLAQKYQKPLFVLPQRLNESDGTNELLEKGQAQGIFNIQNFINTLLKDYHLKEMPEMEDEFLEYCAKNPSYEEAY  240
NTDB id 1161386 AABJ57 RS09505 WP 002776372.1  SMQSARLALELNKPLYVLPQRKNESDGTNLLLQEKNANLLLDFKEFASCF......GVIEEEKDEFLDFCRQGVSVDEAT  229
NTDB id 1239 Cj0634 YP 002344063.1              SMQSAKLALELNKPLYVLPQRLGESTATNLLLKENKAKLICDFKEFVSEF......ASIDMNQDEFLEFCKKGVSVDEAL  230
consensus                                       !!*!!*!!****!!!*!!!!!**!!**!!*!! * *!*** ****!** *       * ***  !!!!**!****!**!!

logo
NTDB id 1210 C694 RS01710 WP 001862983.1        LKFGDKLLEYELLGKIKRINHIVVLA.   266
NTDB id 1161386 AABJ57 RS09505 WP 002776372.1  QIYREKVYEYELEGKISIEGLFIRVLV   256
NTDB id 1239 Cj0634 YP 002344063.1              KIYGQKVYEYELEGKISIEGLFIRVLA   257
consensus                                        ***!**!!!!*!!!**********
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X non conserved

X similar

X ≥ 50% conserved