```
logo
NTDB id 54229 C695 RS01710 WP 001862983.1  MKSHFQYSTLENIPKAFDILKDPPKKLYCVGDTKLLDTPLKVAIIGTRRPTPYSKQHTITLARELAKNGAVIVSGGALGVDIIAQ   85
NTDB id 1210 C694 RS01710 WP 001862983.1   MKSHFQYSTLENIPKAFDILKDPPKKLYCVGDTKLLDTPLKVAIIGTRRPTPYSKQHTITLARELAKNGAVIVSGGALGVDIIAQ   85
NTDB id 1239 Cj0634 YP 002344063.1         MASEI...LPDEFLELFKDLKKPPKKLHYKGNLSLLK.QDKIAIIGSRRMSVYTKNCVFSLASMLKNAHLCVVSGGALGVDITAS   81
consensus                                  !*!*************!**!!*!!!!!***!***!!****!*!!!!*!!***!*!*****!!**!*******!!!!!!!!!!*!*

logo
NTDB id 54229 C695 RS01710 WP 001862983.1  ENALPKTIMLSPCSLDFIYPTNNHKVIQEIAQNGLILSEYEKDFMPIKGSFLARNRLVIALSDVVIIPQADLKSGSMSSARLAQK  170
NTDB id 1210 C694 RS01710 WP 001862983.1   ENALPKTIMLSPCSLDFIYPTNNHKVIQEIAQNGLILSEYEKDFMPIKGSFLARNRLVIALSDVVIIPQADLKSGSMSSARLAQK  170
NTDB id 1239 Cj0634 YP 002344063.1         MAAMPNTIGIFANGLDQIYPRTNEKIIKQIYENALALSEYEDDYLPKNYDFLLRNRLVIALSKAVVVAQADIKSGSMQSAKLALE  166
consensus                                  **!*!*!!******!!*!!!**!*!*!**!**!*!*!!!!!*!**!****!!*!!!!!!!!!!**!***!!!*!!!!!*!!*!!**

logo
NTDB id 54229 C695 RS01710 WP 001862983.1  YQKPLFVLPQRLNESDGTNELLEKGQAQGIFNIQNFINTLLKDYHLKEMPEMEDEFLEYCAKNPSYEEAYLKFGDKLLEYELLGK  255
NTDB id 1210 C694 RS01710 WP 001862983.1   YQKPLFVLPQRLNESDGTNELLEKGQAQGIFNIQNFINTLLKDYHLKEMPEMEDEFLEYCAKNPSYEEAYLKFGDKLLEYELLGK  255
NTDB id 1239 Cj0634 YP 002344063.1         LNKPLYVLPQRLGESTATNLLLKENKAKLICDFKEFVSEF......ASIDMNQDEFLEFCKKGVSVDEALKIYGQKVYEYELEGK  245
consensus                                  **!!!*!!!!!!*!!**!!*!!****!**!*****!*****************!!!!!*!*!**!**!!****!*!**!!!!*!!

logo
NTDB id 54229 C695 RS01710 WP 001862983.1  IKRINHIVVLA.   266
NTDB id 1210 C694 RS01710 WP 001862983.1   IKRINHIVVLA.   266
NTDB id 1239 Cj0634 YP 002344063.1         ISIEGLFIRVLA   257
consensus                                  !**********
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X non conserved

X similar

X ≥ 50% conserved