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logo

NTDB id 1239 Cj0634 YP 002344063.1      MASEI...LPDEFLELFKDLKKPPKKLHYKGNLSLLK.QDKIAIIGSRRMSVYTKNCVFSLASMLKNAHLCVVSGGALGV   76
NTDB id 79095 HPYLSS1 RS01680 WP 077231809.1 MKSHFQYSALENIPKAFDILKDPPKKLYCVGDTKLLEAPLKVAIIGTRRPTPYSKQHTITLARELAKNGAVIVSGGALGV   80
NTDB id 1210 C694 RS01710 WP 001862983.1 MKSHFQYSTLENIPKAFDILKDPPKKLYCVGDTKLLDTPLKVAIIGTRRPTPYSKQHTITLARELAKNGAVIVSGGALGV   80
consensus                               !*!***** ********!**!!*!!!!!***!***!!* **!*!!!!*!!***!*!*****!!**!********!!!!!!!!

logo

NTDB id 1239 Cj0634 YP 002344063.1      DITASMAAMPNTIGIFANGLDQIYPRTNEKIIKQIYENALALSEYEDDYLPKNYDFLLRNRLVIALSKAVVVAQADIKSG   156
NTDB id 79095 HPYLSS1 RS01680 WP 077231809.1 DIIAQENALPKTIMLSPCSLDLIYPTNNHKVIQEIAQNGLILSEYEKDFMPVKGSFLARNRLVIALSDVVIIPQADLKSG   160
NTDB id 1210 C694 RS01710 WP 001862983.1 DIIAQENALPKTIMLSPCSLDFIYPTNNHKVIQEIAQNGLILSEYEKDFMPIKGSFLARNRLVIALSDVVIIPQADLKSG   160
consensus                               !!*!***!*!*!!******!! !!!**!*!*!**!**!*!*!!!!!*!**!****!!*!!!!!!!!!!**!***!!!*!!!

logo

NTDB id 1239 Cj0634 YP 002344063.1      SMQSAKLALELNKPLYVLPQRLGESTATNLLLKENKAKLICDFKEFVSEF......ASIDMNQDEFLEFCKKGVSVDEAL   230
NTDB id 79095 HPYLSS1 RS01680 WP 077231809.1 SMSSARLAQKYQKPLFVLPQRLNESDGTNELLEKGQAQGIFNIQNFINTLLKDYHLKEMPEMKDEFLEYCAKNPSYEEAY   240
NTDB id 1210 C694 RS01710 WP 001862983.1 SMSSARLAQKYQKPLFVLPQRLNESDGTNELLEKGQAQGIFNIQNFINTLLKDYHLKEMPEMEDEFLEYCAKNPSYEEAY   240
consensus                               !!*!!*!!****!!!*!!!!!!*!!**!!*!!****!**!*****!**************** !!!!!*!*!**!**!!*

logo

NTDB id 1239 Cj0634 YP 002344063.1      KIYGQKVYEYELEGKISIEGLFIRVLA   257
NTDB id 79095 HPYLSS1 RS01680 WP 077231809.1 LKFGDKLLEYELLGKIKRINHLVVLA.   266
NTDB id 1210 C694 RS01710 WP 001862983.1 LKFGDKLLEYELLGKIKRINHIVVLA.   266
consensus                               ***!*!**!!!!*!!!*****!****
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X non conserved

X similar

X ≥ 50% conserved