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logo
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NTDB id 71510 HPOKI898 RS01875 WP 025275690.1  MKSNFQYSTLENIPKAFDILKDPPKKLYCVGDIKLLRAPLKVAIIGTRRPTPYSKQHTITLARELAKNGAVIVSGGALGV  80
NTDB id 1210 C694 RS01710 WP 001862983.1       MKSHFQYSTLENIPKAFDILKDPPKKLYCVGDTKLLDTPLKVAIIGTRRPTPYSKQHTITLARELAKNGAVIVSGGALGV  80
NTDB id 1239 Cj0634 YP 002344063.1             MASEI...LPDEFLELFKDLKKPPKKLHYKGNLSLLKQ.DKIAIIGSRRMSVYTKNCVFSLASMLKNAHLCVVSGGALGV  76
consensus                                      !*!  ************!**!!*!!!!!***!***!!* **!*!!!!*!!***!*!*****!!**!*******!!!!!!!!
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logo
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NTDB id 71510 HPOKI898 RS01875 WP 025275690.1  DIIAQENALPKTIMLSPCSLDFIYPTNNHKVIQEIAQNGLILSEYEKDFMPIKGSFLARNRLVIALSDAVIIPQADLKSG  160
NTDB id 1210 C694 RS01710 WP 001862983.1       DIIAQENALPKTIMLSPCSLDFIYPTNNHKVIQEIAQNGLILSEYEKDFMPIKGSFLARNRLVIALSDVVIIPQADLKSG  160
NTDB id 1239 Cj0634 YP 002344063.1             DITASMAAMPNTIGIFANGLDQIYPRTNEKIIKQIYENALALSEYEDDYLPKNYDFLLRNRLVIALSKAVVVAQADIKSG  156
consensus                                      !!*!***!*!*!!******!!*!!!**!*!*!**!**!*!*!!!!!*!**!****!!*!!!!!!!!!**!***!!!*!!!
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logo
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NTDB id 71510 HPOKI898 RS01875 WP 025275690.1  SMSSARLAQKYQKPLFVLPQRLNESDGTNELLEKGQAQGIFNIQNFINTLLKDYHLKEMPEMEDEFLEYCAKNPSYEEAY  240
NTDB id 1210 C694 RS01710 WP 001862983.1       SMSSARLAQKYQKPLFVLPQRLNESDGTNELLEKGQAQGIFNIQNFINTLLKDYHLKEMPEMEDEFLEYCAKNPSYEEAY  240
NTDB id 1239 Cj0634 YP 002344063.1             SMQSAKLALELNKPLYVLPQRLGESTATNLLLKENKAKLICDFKEFVSEF......ASIDMNQDEFLEFCKKGVSVDEAL  230
consensus                                      !!*!!*!!****!!!*!!!!!!*!!**!!*!!****!**!*****!*******************!!!!!*!*!**!**!!*
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logo
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NTDB id 71510 HPOKI898 RS01875 WP 025275690.1  LKFGDKLLEYELLGKIKRINHLVVLA.  266
NTDB id 1210 C694 RS01710 WP 001862983.1       LKFGDKLLEYELLGKIKRINHIVVLA.  266
NTDB id 1239 Cj0634 YP 002344063.1             KIYGQKVYEYELEGKISIEGLFIRVLA  257
consensus                                      ***!*!**!!!!*!!!*****!****
```

X non conserved
X similar
X ≥ 50% conserved