
logo
NTDB id 75894 EG63 RS01665 WP 038417325.1 MKSNFQYSTLENIPKAFDILKDPPKKLYCVGDTKLLEAPLKVAIIGTRRPTPYSKQHTITLARELAKNGAVIVSGGALGVDIIAQ 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 75894 EG63 RS01665 WP 038417325.1 ENALPTTIMFSPCSLDFVYPTNNYKVIQEIAQNGLILSEYEKDFMPIKGSFLARNRLVIALSDAVIIPQADLKSGSMSSARLAQK 170
NTDB id 1210 C694 RS01710 WP 001862983.1 ENALPKTIMLSPCSLDFIYPTNNHKVIQEIAQNGLILSEYEKDFMPIKGSFLARNRLVIALSDVVIIPQADLKSGSMSSARLAQK 170
NTDB id 1239 Cj0634 YP 002344063.1 MAAMPNTIGIFANGLDQIYPRTNEKIIKQIYENALALSEYEDDYLPKNYDFLLRNRLVIALSKAVVVAQADIKSGSMQSAKLALE 166
consensus
logo
NTDB id 75894 EG63 RS01665 WP 038417325.1 YQKPLFVLPQRLNESDGTNELLEKGQAQGIFNIQNFINTLLKDYHLKEMPEMKDEFLEYCAKNPSYEEAYLKFGDKLLEYELLGK 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 75894 EG63 RS01665 WP 038417325.1 IKRINHIVVLA. 266
NTDB id 1210 C694 RS01710 WP 001862983.1 IKRINHIVVLA. 266
NTDB id 1239 Cj0634 YP 002344063.1 ISIEGLFIRVLA 257
consensus
X non conserved
X similar
X ≥ 50% conserved