logo
NTDB id 1210 C694 RS01710 WP 001862983.1 MKSHFQYSTLENIPKAFDILKDPPKKLYCVGDTKLLDTPLKVAIIGTRRPTPYSKQHTITLARELAKNGAVIVSGGALGVDIIAQ 85
NTDB id 117216 YSQ RS05990 WP 002815152.1 ....MQSELPSAYLELFKELKNLPKKLYYKGNLKLLEQ.RKIAIIGSRRMSTYTKNCVFELSNLLKNSKISVVSGGALGVDISAS 80
NTDB id 1239 Cj0634 YP 002344063.1 ...MASEILPDEFLELFKDLKKPPKKLHYKGNLSLLKQ.DKIAIIGSRRMSVYTKNCVFSLASMLKNAHLCVVSGGALGVDITAS 81
consensus
logo
NTDB id 1210 C694 RS01710 WP 001862983.1 ENALPKTIMLSPCSLDFIYPTNNHKVIQEIAQNGLILSEYEKDFMPIKGSFLARNRLVIALSDVVIIPQADLKSGSMSSARLAQK 170
NTDB id 117216 YSQ RS05990 WP 002815152.1 MAAMPSTIGVFANGLDEIYPRSNEKIIKQIYENALALSENESYYKPKPYDFLLRNRLIVALSEAVVVAQADLQSGSMQSARLALE 165
NTDB id 1239 Cj0634 YP 002344063.1 MAAMPNTIGIFANGLDQIYPRTNEKIIKQIYENALALSEYEDDYLPKNYDFLLRNRLVIALSKAVVVAQADIKSGSMQSAKLALE 166
consensus
logo
NTDB id 1210 C694 RS01710 WP 001862983.1 YQKPLFVLPQRLNESDGTNELLEKGQAQGIFNIQNFINTLLKDYHLKEMPEMEDEFLEYCAKNPSYEEAYLKFGDKLLEYELLGK 255
NTDB id 117216 YSQ RS05990 WP 002815152.1 LNKPLYVLPQRKNESDGTNLLLQEKNANLLLDFKEFASCF......GVIEEEKDEFLDFCRQGVSVDEATQIYGEKVYEYELEGK 244
NTDB id 1239 Cj0634 YP 002344063.1 LNKPLYVLPQRLGESTATNLLLKENKAKLICDFKEFVSEF......ASIDMNQDEFLEFCKKGVSVDEALKIYGQKVYEYELEGK 245
consensus
logo
NTDB id 1210 C694 RS01710 WP 001862983.1 IKRINHIVVLA. 266
NTDB id 117216 YSQ RS05990 WP 002815152.1 ISIEGLFIRVLV 256
NTDB id 1239 Cj0634 YP 002344063.1 ISIEGLFIRVLA 257
consensus
X non conserved
X similar
X ≥ 50% conserved