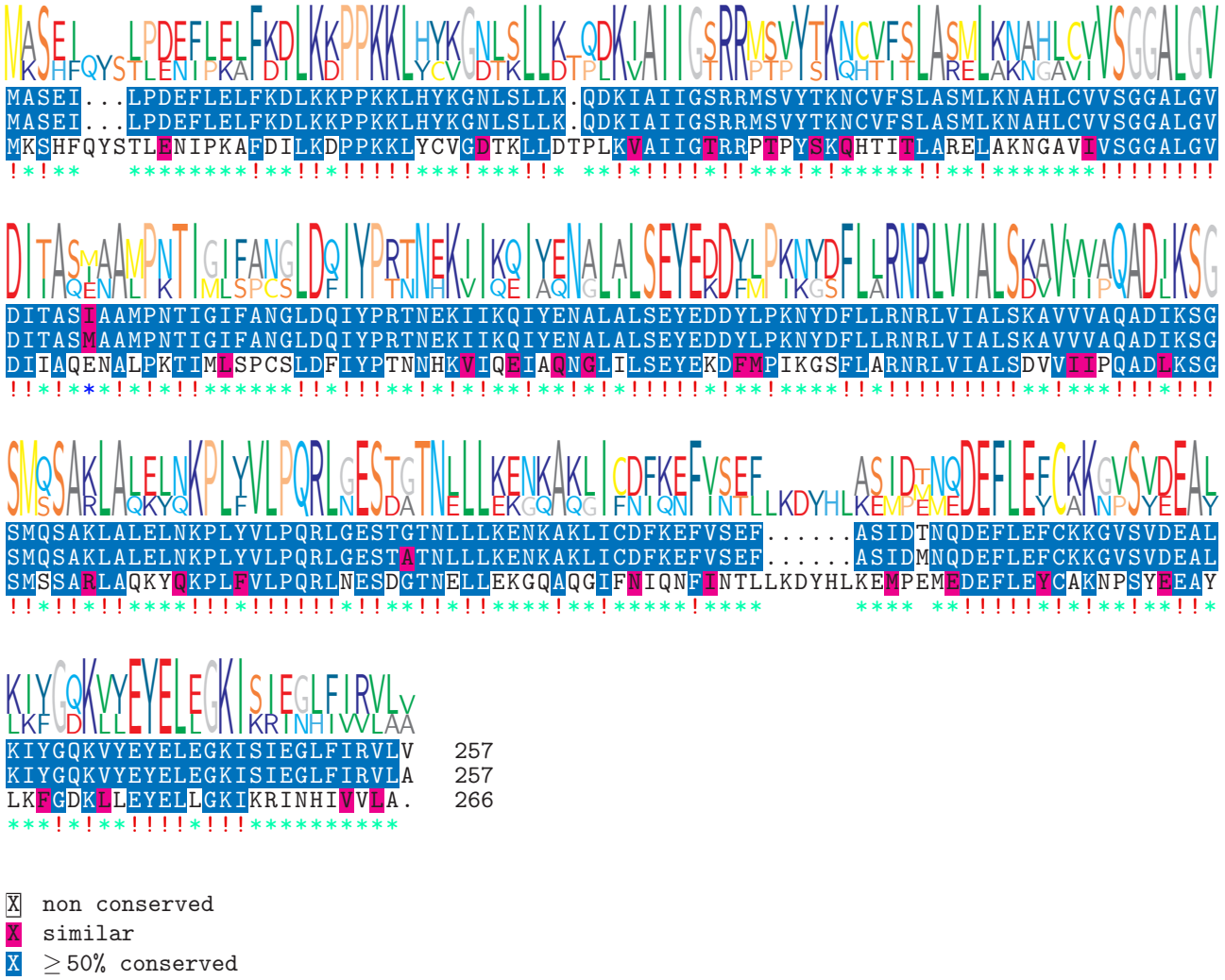
logo
NTDB id 105327 R8922 RS03130 WP 002857196.1 MASEI...LPDEFLELFKDLKKPPKKLHYKGNLSLLK.QDKIAIIGSRRMSVYTKNCVFSLASMLKNAHLCVVSGGALGV 76
NTDB id 1239 Cj0634 YP 002344063.1 MASEI...LPDEFLELFKDLKKPPKKLHYKGNLSLLK.QDKIAIIGSRRMSVYTKNCVFSLASMLKNAHLCVVSGGALGV 76
NTDB id 1210 C694 RS01710 WP 001862983.1 MKSHFQYSTLENIPKAFDILKDPPKKLYCVGDTKLLDTPLKVAIIGTRRPTPYSKQHTITLARELAKNGAVIVSGGALGV 80
consensus
logo
NTDB id 105327 R8922 RS03130 WP 002857196.1 DITASIAAMPNTIGIFANGLDQIYPRTNEKIIKQIYENALALSEYEDDYLPKNYDFLLRNRLVIALSKAVVVAQADIKSG 156
NTDB id 1239 Cj0634 YP 002344063.1 DITASMAAMPNTIGIFANGLDQIYPRTNEKIIKQIYENALALSEYEDDYLPKNYDFLLRNRLVIALSKAVVVAQADIKSG 156
NTDB id 1210 C694 RS01710 WP 001862983.1 DIIAQENALPKTIMLSPCSLDFIYPTNNHKVIQEIAQNGLILSEYEKDFMPIKGSFLARNRLVIALSDVVIIPQADLKSG 160
consensus
logo
NTDB id 105327 R8922 RS03130 WP 002857196.1 SMQSAKLALELNKPLYVLPQRLGESTGTNLLLKENKAKLICDFKEFVSEF.....ASIDTNQDEFLEFCKKGVSVDEAL 230
NTDB id 1239 Cj0634 YP 002344063.1 SMQSAKLALELNKPLYVLPQRLGESTATNLLLKENKAKLICDFKEFVSEF.....ASIDMNQDEFLEFCKKGVSVDEAL 230
NTDB id 1210 C694 RS01710 WP 001862983.1 SMSSARLAQKYQKPLFVLPQRLNESDGTNELLEKGQAQGIFNIQNFINTLLKDYHLKEMPEMEDEFLEYCAKNPSYEEAY 240
consensus
logo
NTDB id 105327 R8922 RS03130 WP 002857196.1 KIYGQKVYEYELEGKISIEGLFIRVLV 257
NTDB id 1239 Cj0634 YP 002344063.1 KIYGQKVYEYELEGKISIEGLFIRVLA 257
NTDB id 1210 C694 RS01710 WP 001862983.1 LKFGDKLLEYELLGKIKRINHIVVLA. 266
consensus
X non conserved
X similar
X ≥ 50% conserved