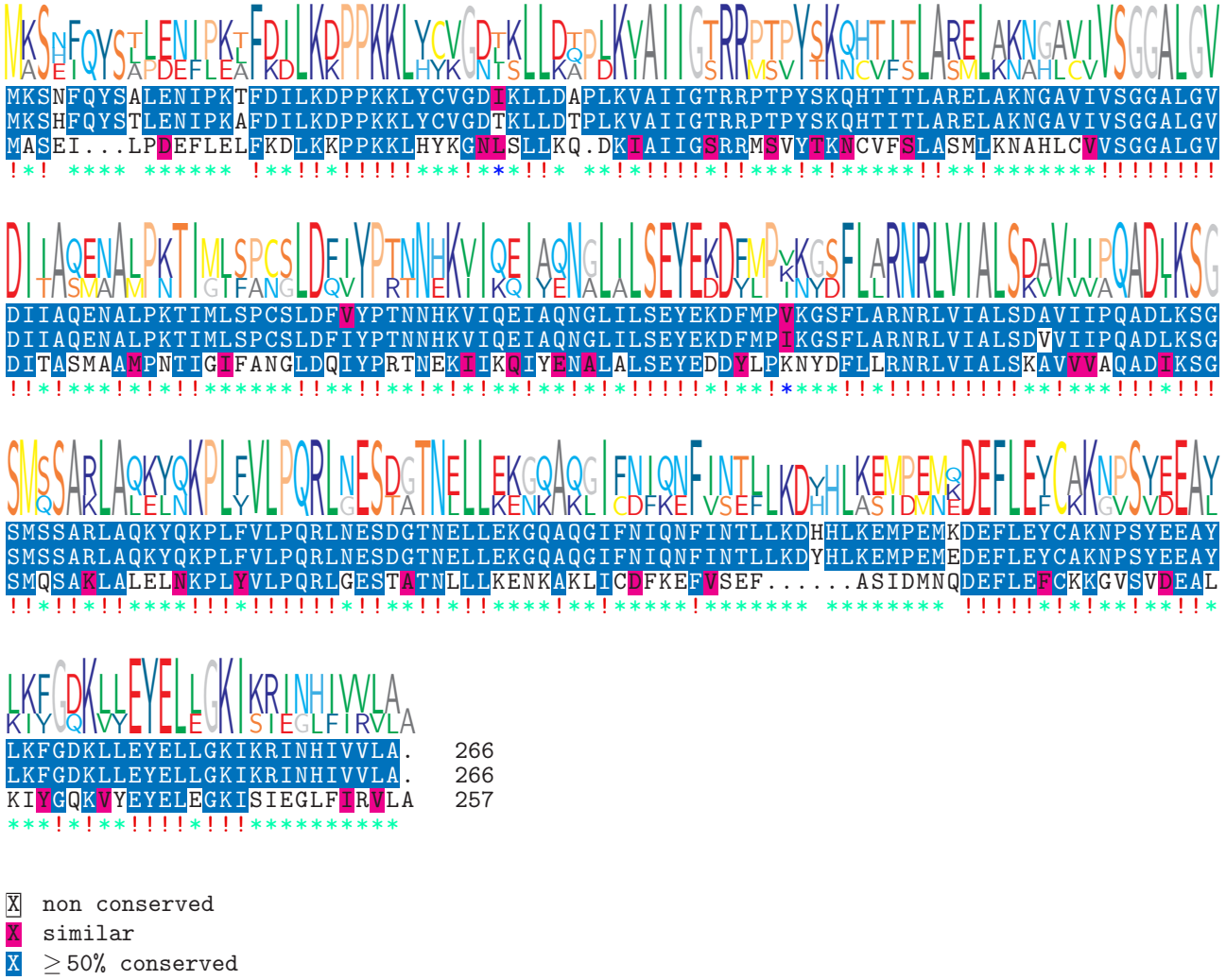
logo
NTDB id 667156 MPG25 RS05470 WP 245018519.1 MKSNFQYSALENIPKTFDILKDPPKKLYCVGDIKLLDAPLKVAIIGTRRPTPYSKQHTITLARELAKNGAVIVSGGALGV 80
NTDB id 1210 C694 RS01710 WP 001862983.1 MKSHFQYSTLENIPKAFDILKDPPKKLYCVGDTKLLDTPLKVAIIGTRRPTPYSKQHTITLARELAKNGAVIVSGGALGV 80
NTDB id 1239 Cj0634 YP 002344063.1 MASEI...LPDEFLELFKDLKKPPKKLHYKGNLSLLKQ.DKIAIIGSRRMSVYTKNCVFSLASMLKNAHLCVVSGGALGV 76
consensus
logo
NTDB id 667156 MPG25 RS05470 WP 245018519.1 DIIAQENALPKTIMLSPCSLDFVYPTNNHKVIQEIAQNGLILSEYEKDFMPVKGSFLARNRLVIALSDAVIIPQADLKSG 160
NTDB id 1210 C694 RS01710 WP 001862983.1 DIIAQENALPKTIMLSPCSLDFIYPTNNHKVIQEIAQNGLILSEYEKDFMPIKGSFLARNRLVIALSDVVIIPQADLKSG 160
NTDB id 1239 Cj0634 YP 002344063.1 DITASMAAMPNTIGIFANGLDQIYPRTNEKIIKQIYENALALSEYEDDYLPKNYDFLLRNRLVIALSKAVVVAQADIKSG 156
consensus
logo
NTDB id 667156 MPG25 RS05470 WP 245018519.1 SMSSARLAQKYQKPLFVLPQRLNESDGTNELLEKGQAQGIFNIQNFINTLLKDHHLKEMPEMKDEFLEYCAKNPSYEEAY 240
NTDB id 1210 C694 RS01710 WP 001862983.1 SMSSARLAQKYQKPLFVLPQRLNESDGTNELLEKGQAQGIFNIQNFINTLLKDYHLKEMPEMEDEFLEYCAKNPSYEEAY 240
NTDB id 1239 Cj0634 YP 002344063.1 SMQSAKLALELNKPLYVLPQRLGESTATNLLLKENKAKLICDFKEFVSEF.....ASIDMNQDEFLEFCKKGVSVDEAL 230
consensus
logo
NTDB id 667156 MPG25 RS05470 WP 245018519.1 LKFGDKLLEYELLGKIKRINHIVVLA. 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