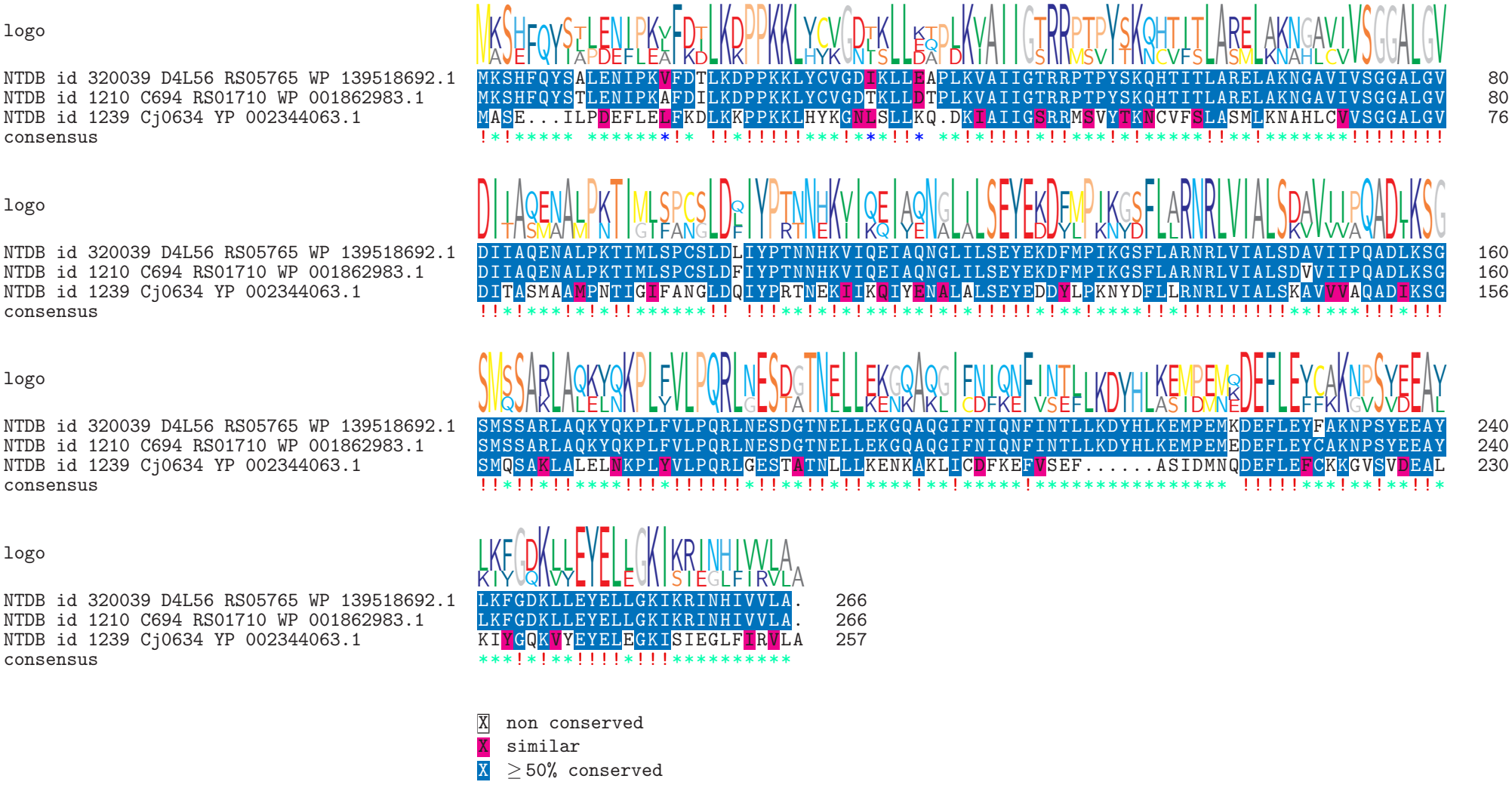

logo
NTDB id 320039 D4L56 RS05765 WP 139518692.1 MKSHFQYSALENIPKVFDTLKDPPKKLYCVGDIKLLEAPLKVAIIGTRRPTPYSKQHTITLARELAKNGAVIVSGGALGV 80
NTDB id 1210 C694 RS01710 WP 001862983.1 MKSHFQYSTLENIPKAFDILKDPPKKLYCVGDTKLLDTPLKVAIIGTRRPTPYSKQHTITLARELAKNGAVIVSGGALGV 80
NTDB id 1239 Cj0634 YP 002344063.1 MASE...ILPDEFLELFKDLKKPPKKLHYKGNLSLLKQ.DKIAIIGSRRMSVYTKNCVFSLASMLKNAHLCVVSGGALGV 76
consensus
logo
NTDB id 320039 D4L56 RS05765 WP 139518692.1 DIIAQENALPKTIMLSPCSLDLIYPTNNHKVIQEIAQNGLILSEYEKDFMPIKGSFLARNRLVIALSDAVIIPQADLKSG 160
NTDB id 1210 C694 RS01710 WP 001862983.1 DIIAQENALPKTIMLSPCSLDFIYPTNNHKVIQEIAQNGLILSEYEKDFMPIKGSFLARNRLVIALSDVVIIPQADLKSG 160
NTDB id 1239 Cj0634 YP 002344063.1 DITASMAAMPNTIGIFANGLDQIYPRTNEKIIKQIYENALALSEYEDDYLPKNYDFLLRNRLVIALSKAVVVAQADIKSG 156
consensus
logo
NTDB id 320039 D4L56 RS05765 WP 139518692.1 SMSSARLAQKYQKPLFVLPQRLNESDGTNELLEKGQAQGIFNIQNFINTLLKDYHLKEMPEMKDEFLEYFAKNPSYEEAY 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 320039 D4L56 RS05765 WP 139518692.1 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