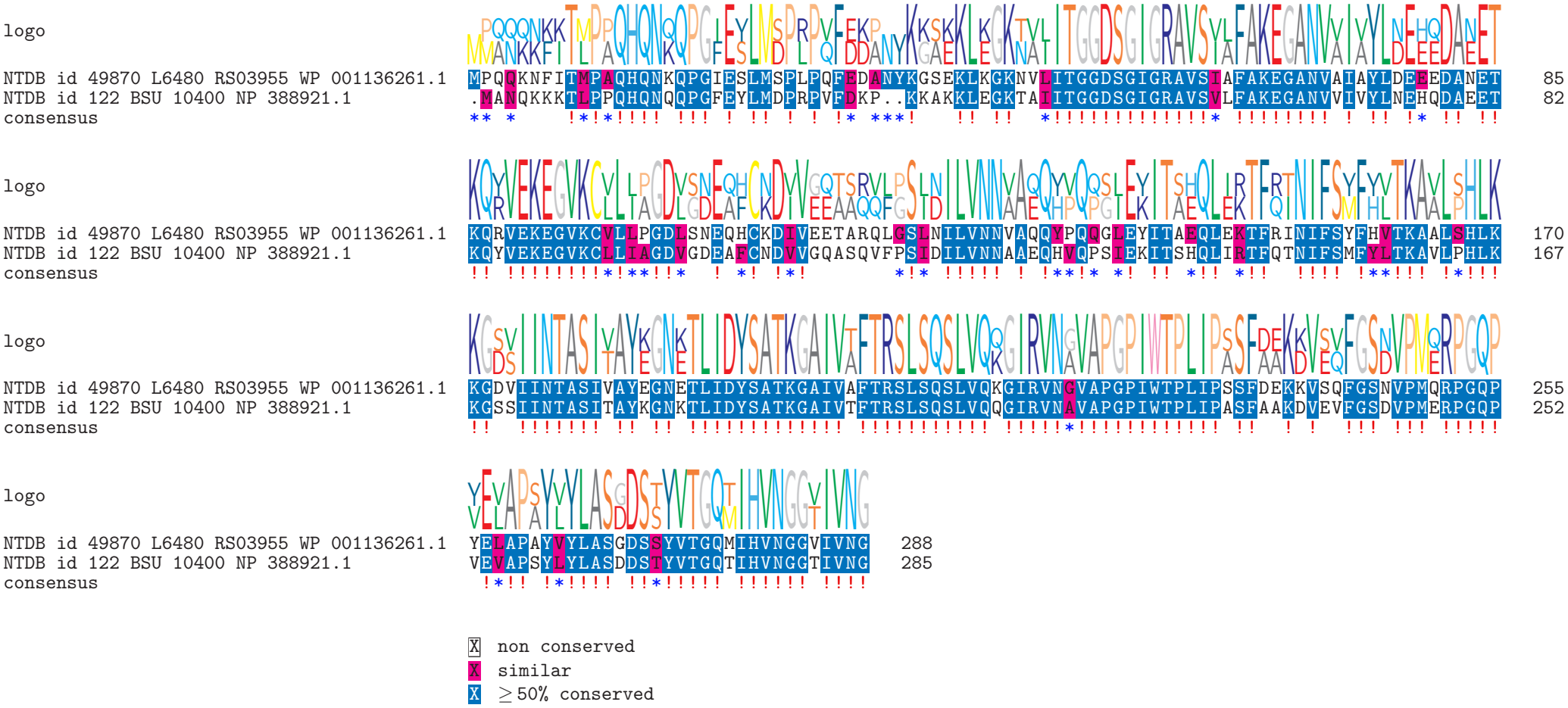

logo
NTDB id 49870 L6480 RS03955 WP 001136261.1 MPQQKNFITMPAQHQNKQPGIESLMSPLPQFEDANYKGSEKLKGKNVLITGGDSGIGRAVSIAFAKEGANVAIAYLDEEEDANET 85
NTDB id 122 BSU 10400 NP 388921.1 .MANQKKKTLPPQHQNQQPGFEYLMDPRPVFDKP..KKAKKLEGKTAIITGGDSGIGRAVSVLFAKEGANVVIVYLNEHQDAEET 82
consensus
logo
NTDB id 49870 L6480 RS03955 WP 001136261.1 KQRVEKEGVKCVLLPGDLSNEQHCKDIVEETARQLGSLNILVNNVAQQYPQQGLEYITAEQLEKTFRINIFSYFHVTKAALSHLK 170
NTDB id 122 BSU 10400 NP 388921.1 KQYVEKEGVKCLLIAGDVGDEAFCNDVVGQASQVFPSIDILVNNAAEQHVQPSIEKITSHQLIRTFQTNIFSMFYLTKAVLPHLK 167
consensus
logo
NTDB id 49870 L6480 RS03955 WP 001136261.1 KGDVIINTASIVAYEGNETLIDYSATKGAIVAFTRSLSQSLVQKGIRVNGVAPGPIWTPLIPSSFDEKKVSQFGSNVPMQRPGQP 255
NTDB id 122 BSU 10400 NP 388921.1 KGSSIINTASITAYKGNKTLIDYSATKGAIVTFTRSLSQSLVQQGIRVNAVAPGPIWTPLIPASFAAKDVEVFGSDVPMERPGQP 252
consensus
logo
NTDB id 49870 L6480 RS03955 WP 001136261.1 YELAPAYVYLASGDSSYVTGQMIHVNGGVIVNG 288
NTDB id 122 BSU 10400 NP 388921.1 VEVAPSYLYLASDDSTYVTGQTIHVNGGTIVNG 285
consensus
X non conserved
X similar
X ≥ 50% conserved