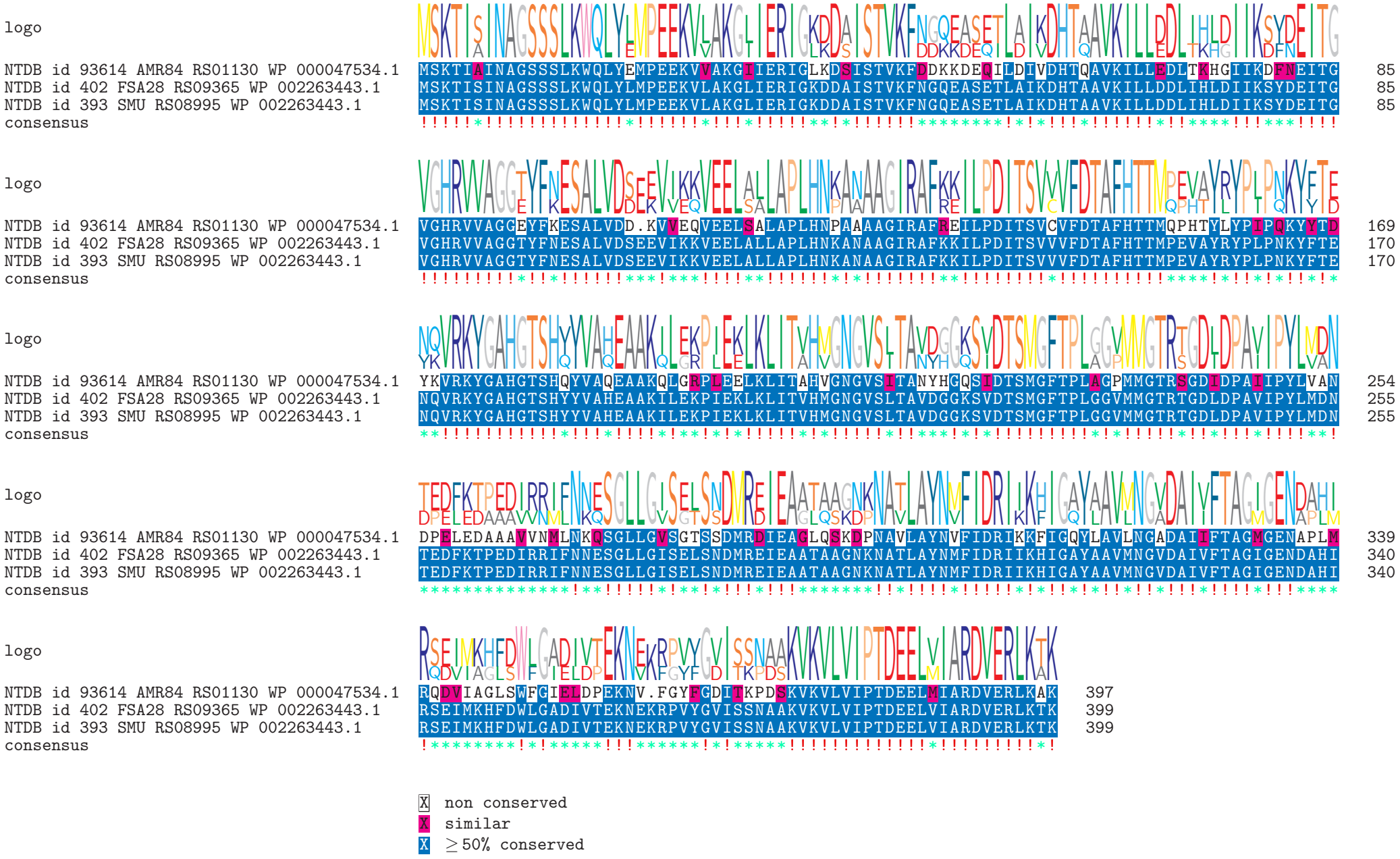

logo
NTDB id 93614 AMR84 RS01130 WP 000047534.1 MSKTIAINAGSSSLKWQLYEMPEEKVVAKGIIERIGLKDSISTVKFDDKKDEQILDIVDHTQAVKILLEDLTKHGIIKDFNEITG 85
NTDB id 402 FSA28 RS09365 WP 002263443.1 MSKTISINAGSSSLKWQLYLMPEEKVLAKGLIERIGKDDAISTVKFNGQEASETLAIKDHTAAVKILLDDLIHLDIIKSYDEITG 85
NTDB id 393 SMU RS08995 WP 002263443.1 MSKTISINAGSSSLKWQLYLMPEEKVLAKGLIERIGKDDAISTVKFNGQEASETLAIKDHTAAVKILLDDLIHLDIIKSYDEITG 85
consensus
logo
NTDB id 93614 AMR84 RS01130 WP 000047534.1 VGHRVVAGGEYFKESALVDD.KVVEQVEELSALAPLHNPAAAAGIRAFREILPDITSVCVFDTAFHTTMQPHTYLYPIPQKYYTD 169
NTDB id 402 FSA28 RS09365 WP 002263443.1 VGHRVVAGGTYFNESALVDSEEVIKKVEELALLAPLHNKANAAGIRAFKKILPDITSVVVFDTAFHTTMPEVAYRYPLPNKYFTE 170
NTDB id 393 SMU RS08995 WP 002263443.1 VGHRVVAGGTYFNESALVDSEEVIKKVEELALLAPLHNKANAAGIRAFKKILPDITSVVVFDTAFHTTMPEVAYRYPLPNKYFTE 170
consensus
logo
NTDB id 93614 AMR84 RS01130 WP 000047534.1 YKVRKYGAHGTSHQYVAQEAAKQLGRPLEELKLITAHVGNGVSITANYHGQQSIDTSMGFTPLAGPMMGTRSGDIDPAIIPYLVAN 254
NTDB id 402 FSA28 RS09365 WP 002263443.1 NQVRKYGAHGTSHYYVAHEAAKILEKPIEKLKLITVHMGNGVSLTAVDGGKSVDTSMGFTPLGGVMMGTRTGDLDPAVIPYLMDN 255
NTDB id 393 SMU RS08995 WP 002263443.1 NQVRKYGAHGTSHYYVAHEAAKILEKPIEKLKLITVHMGNGVSLTAVDGGKSVDTSMGFTPLGGVMMGTRTGDLDPAVIPYLMDN 255
consensus
logo
NTDB id 93614 AMR84 RS01130 WP 000047534.1 DPELEDAAAVVNMLNKQSGLLGVSGTSSDMRDIEAGLQSKDPNAVLAYNVFIDRIKKFIGQYLAVLNGADAIIFTAGMGENAPLM 339
NTDB id 402 FSA28 RS09365 WP 002263443.1 TEDFKTPEDIRRIFNNESGLLGISELSNDMREIEAATAAGNKNATLAYNMFIDRIIKHIGAYAAVMNGVDAIVFTAGIGENDAHI 340
NTDB id 393 SMU RS08995 WP 002263443.1 TEDFKTPEDIRRIFNNESGLLGISELSNDMREIEAATAAGNKNATLAYNMFIDRIIKHIGAYAAVMNGVDAIVFTAGIGENDAHI 340
consensus
logo
NTDB id 93614 AMR84 RS01130 WP 000047534.1 RQDVIAGLSWFGIELDPEKNV.FGYFGDITKPDSKVKVLVIPTDEELMIARDVERLKAK 397
NTDB id 402 FSA28 RS09365 WP 002263443.1 RSEIMKHFDWLGADIVTEKNEKRPVYGVISSNAAKVKVLVIPTDEELVIARDVERLKTK 399
NTDB id 393 SMU RS08995 WP 002263443.1 RSEIMKHFDWLGADIVTEKNEKRPVYGVISSNAAKVKVLVIPTDEELVIARDVERLKTK 399
consensus
X non conserved
X similar
X ≥ 50% conserved