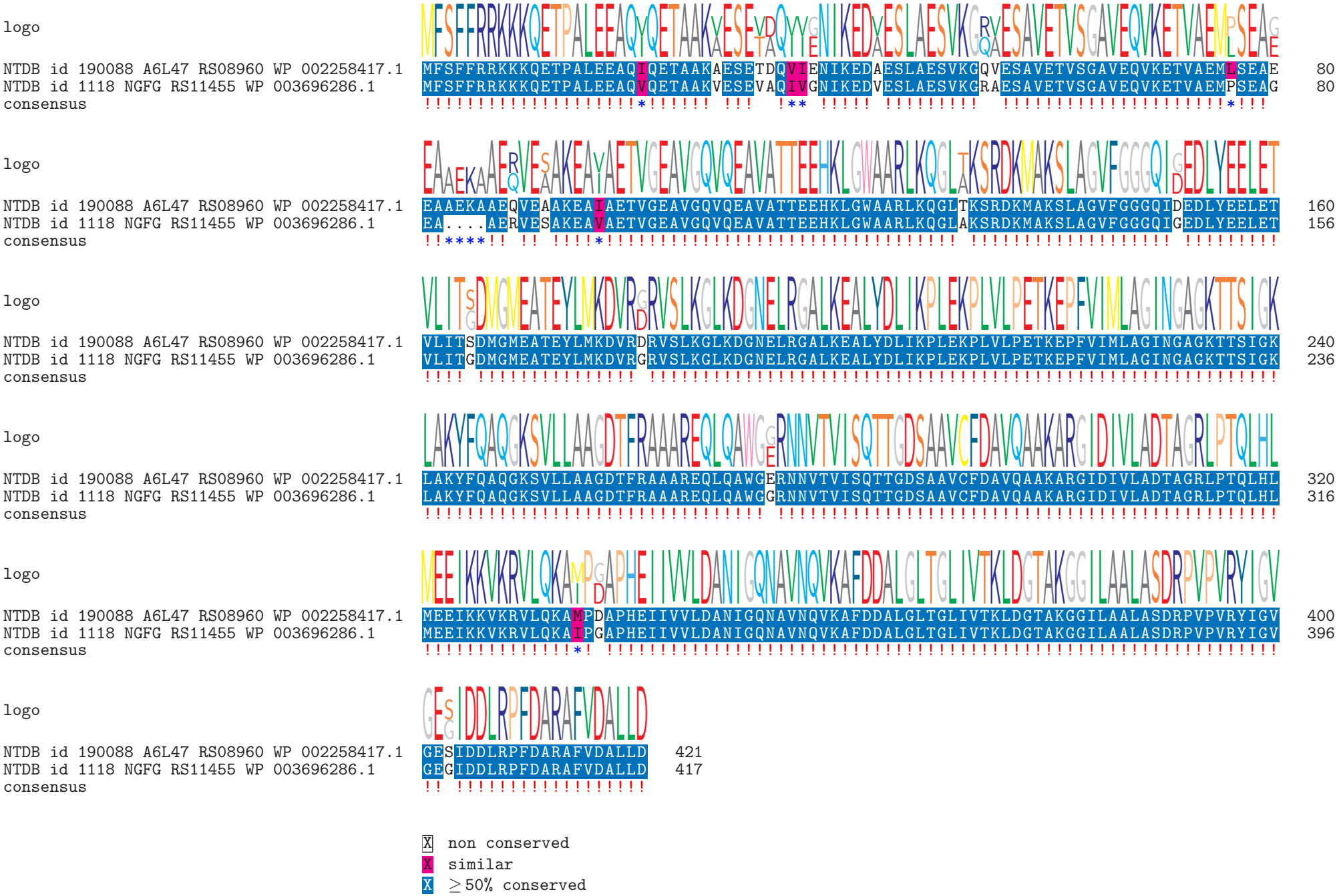

logo
NTDB id 190088 A6L47 RS08960 WP 002258417.1 MFSFFRRKKKQETPALEEAQIQETAAKAESETDQVIENIKEDAESLAESVKGQVESAVETVSGAVEQVKETVAEMLSEAE 80
NTDB id 1118 NGFG RS11455 WP 003696286.1 MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG 80
consensus
NTDB id 190088 A6L47 RS08960 WP 002258417.1 EAAEKAAEQVEAAKEAIAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLTKSRDKMAKSLAGVFGGGQIDEDLYEELET 160
NTDB id 1118 NGFG RS11455 WP 003696286.1 EA....AERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELET 156
consensus
NTDB id 190088 A6L47 RS08960 WP 002258417.1 VLITSDMGMEATEYLMKDVRDRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGK 240
NTDB id 1118 NGFG RS11455 WP 003696286.1 VLITGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGK 236
consensus
NTDB id 190088 A6L47 RS08960 WP 002258417.1 LAKYFQAQGKSVLLAAGDTFRAAAREQLQAWGERNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHL 320
NTDB id 1118 NGFG RS11455 WP 003696286.1 LAKYFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHL 316
consensus
NTDB id 190088 A6L47 RS08960 WP 002258417.1 MEEIKKVKRVLQKAMPDAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGV 400
NTDB id 1118 NGFG RS11455 WP 003696286.1 MEEIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGV 396
consensus
NTDB id 190088 A6L47 RS08960 WP 002258417.1 GESIDDLRPFDARAFVDALLD 421
NTDB id 1118 NGFG RS11455 WP 003696286.1 GEGIDDLRPFDARAFVDALLD 417
consensus
X non conserved
X similar
X ≥ 50% conserved