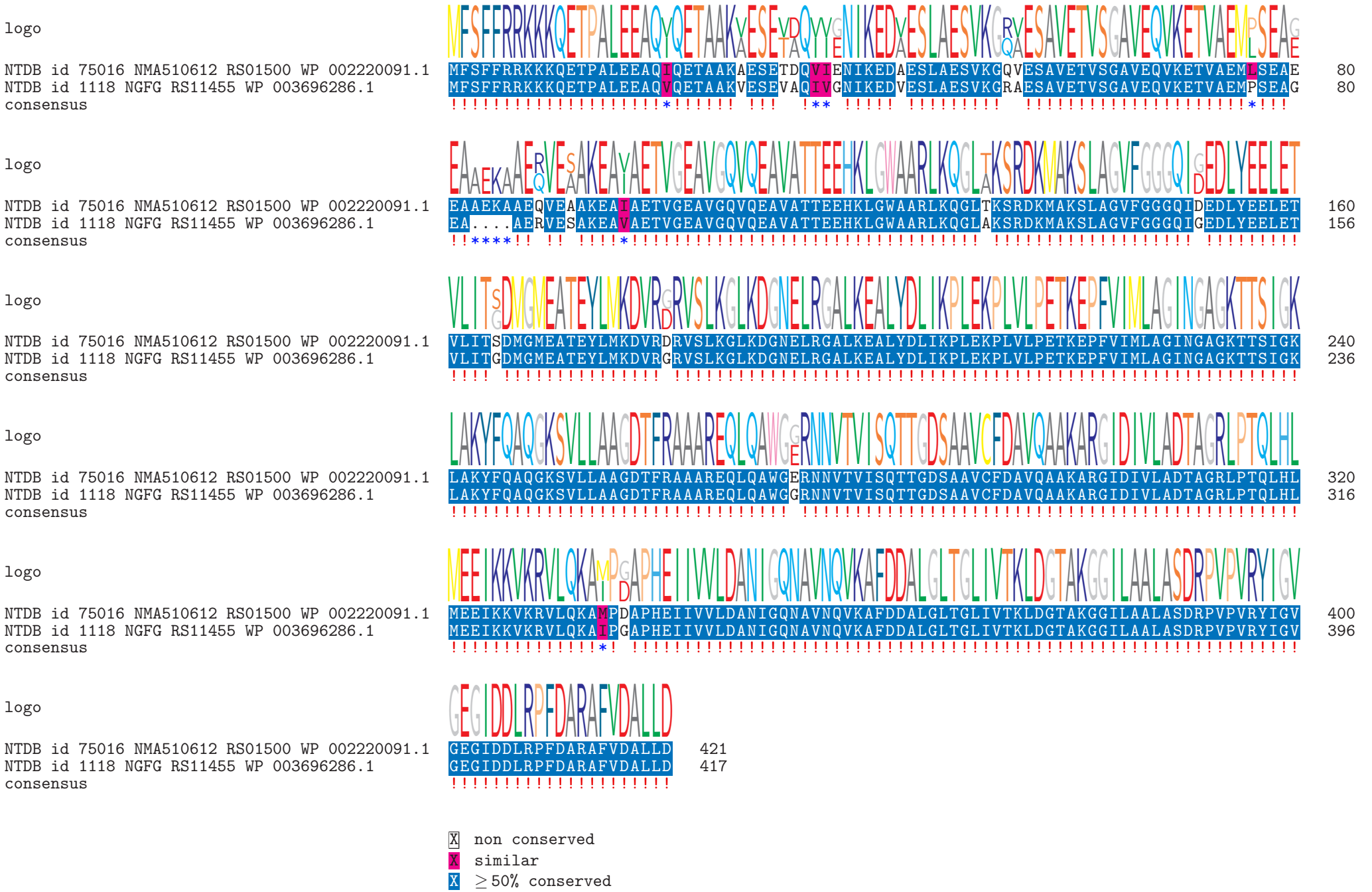
logo
NTDB id 75016 NMA510612 RS01500 WP 002220091.1 MFSFFRRKKKQETPALEEAQIQETAAKAESETDQVIENIKEDAESLAESVKGQVESAVETVSGAVEQVKETVAEMLSEAE 80
NTDB id 1118 NGFG RS11455 WP 003696286.1 MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG 80
consensus
logo
NTDB id 75016 NMA510612 RS01500 WP 002220091.1 EAAEKAAEQVEAAKEAIAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLTKSRDKMAKSLAGVFGGGQIDEDLYEELET 160
NTDB id 1118 NGFG RS11455 WP 003696286.1 EA....AERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELET 156
consensus
logo
NTDB id 75016 NMA510612 RS01500 WP 002220091.1 VLITSDMGMEATEYLMKDVRDRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGK 240
NTDB id 1118 NGFG RS11455 WP 003696286.1 VLITGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGK 236
consensus
logo
NTDB id 75016 NMA510612 RS01500 WP 002220091.1 LAKYFQAQGKSVLLAAGDTFRAAAREQLQAWGERNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHL 320
NTDB id 1118 NGFG RS11455 WP 003696286.1 LAKYFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHL 316
consensus
logo
NTDB id 75016 NMA510612 RS01500 WP 002220091.1 MEEIKKVKRVLQKAMPDAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGV 400
NTDB id 1118 NGFG RS11455 WP 003696286.1 MEEIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGV 396
consensus
logo
NTDB id 75016 NMA510612 RS01500 WP 002220091.1 GEGIDDLRPFDARAFVDALLD 421
NTDB id 1118 NGFG RS11455 WP 003696286.1 GEGIDDLRPFDARAFVDALLD 417
consensus
X non conserved
X similar
X ≥ 50% conserved