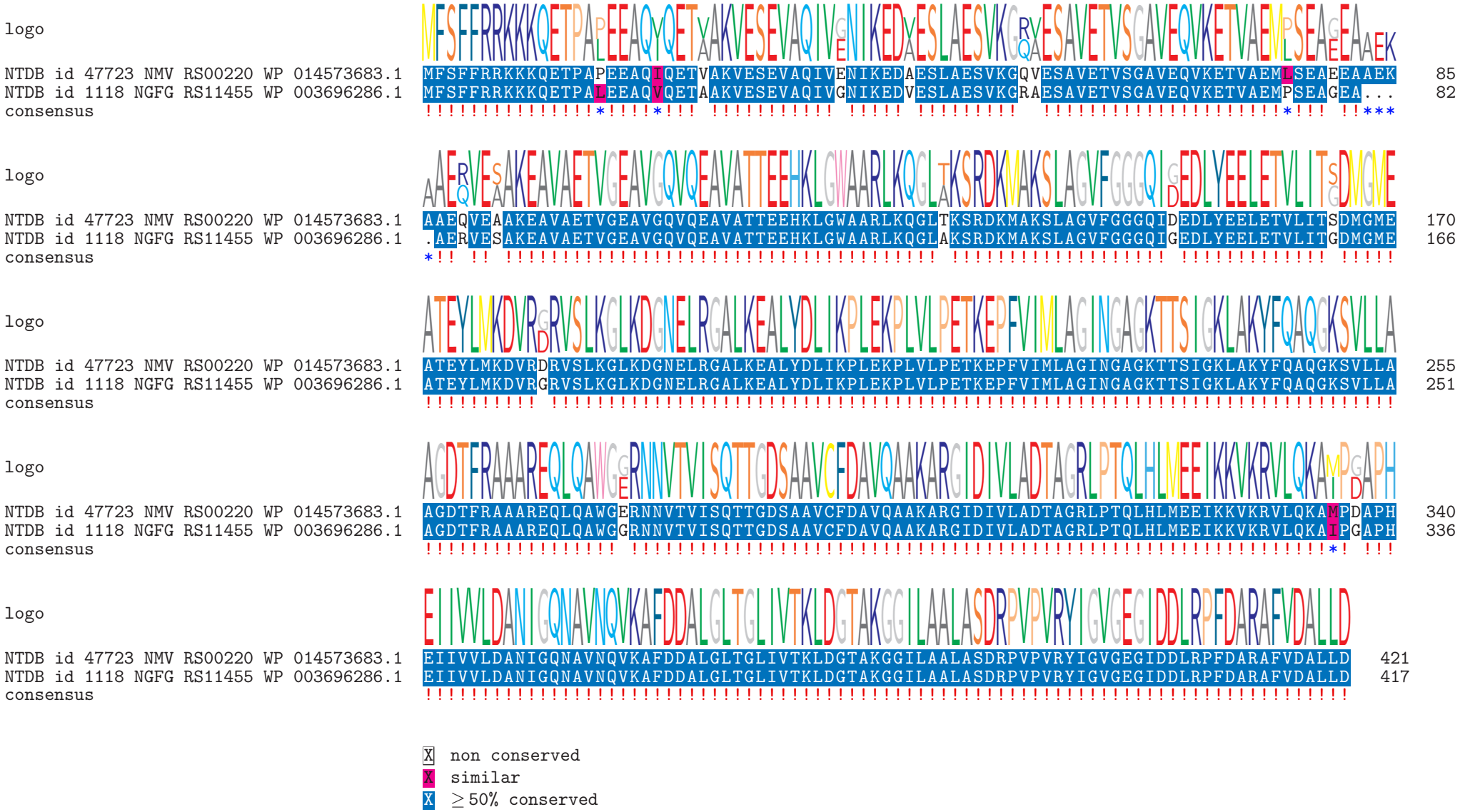
logo
NTDB id 47723 NMV RS00220 WP 014573683.1 MFSFFRRKKKQETPAPEEAQIQETVAKVESEVAQIVENIKEDAESLAESVKGQVESAVETVSGAVEQVKETVAEMLSEAEEAAEK 85
NTDB id 1118 NGFG RS11455 WP 003696286.1 MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAGEA... 82
consensus
logo
NTDB id 47723 NMV RS00220 WP 014573683.1 AAEQVEAAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLTKSRDKMAKSLAGVFGGGQIDEDLYEELETVLITSDMGME 170
NTDB id 1118 NGFG RS11455 WP 003696286.1 .AERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLITGDMGME 166
consensus
logo
NTDB id 47723 NMV RS00220 WP 014573683.1 ATEYLMKDVRDRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKYFQAQGKSVLLA 255
NTDB id 1118 NGFG RS11455 WP 003696286.1 ATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKYFQAQGKSVLLA 251
consensus
logo
NTDB id 47723 NMV RS00220 WP 014573683.1 AGDTFRAAAREQLQAWGERNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEIKKVKRVLQKAMPDAPH 340
NTDB id 1118 NGFG RS11455 WP 003696286.1 AGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEIKKVKRVLQKAIPGAPH 336
consensus
logo
NTDB id 47723 NMV RS00220 WP 014573683.1 EIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGIDDLRPFDARAFVDALLD 421
NTDB id 1118 NGFG RS11455 WP 003696286.1 EIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGIDDLRPFDARAFVDALLD 417
consensus
X non conserved
X similar
X ≥ 50% conserved