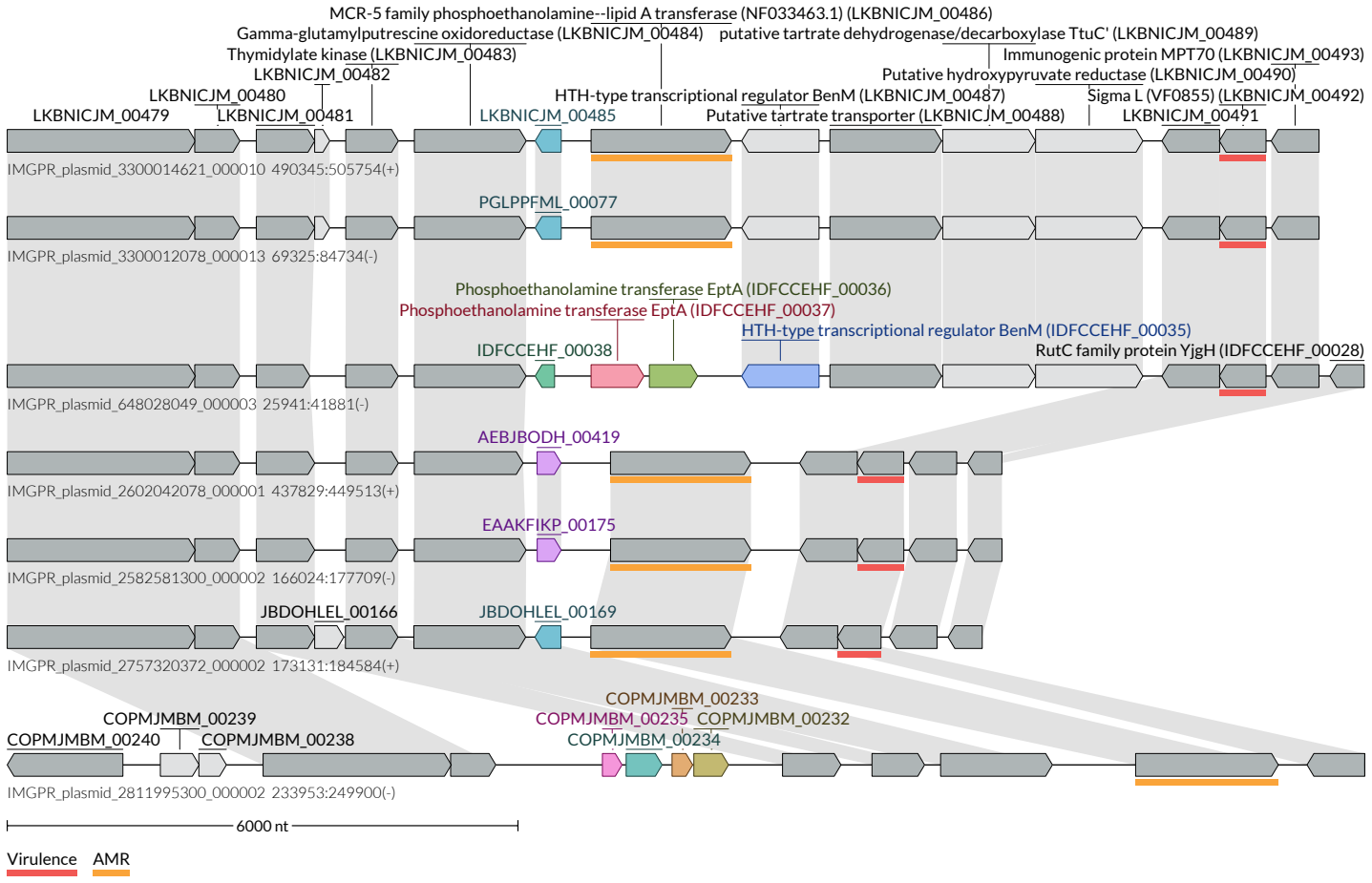
MCR-5 family phosphoethanolamine--lipid A transferase (NF033463.1) (LKBNICJM_00486)
Gamma-glutamylputrescine oxidoreductase (LKBNICJM_00484)
putative tartrate dehydrogenase/decarboxylase TtuC' (LKBNICJM_00489)
Thymidylate kinase (LKBNICJM_00483)
Immunogenic protein MPT70 (LKBNICJM_00493)
LKBNICJM_00482
Putative hydroxypyruvate reductase (LKBNICJM_00490)
LKBNICJM_00480
HTH-type transcriptional regulator BenM (LKBNICJM_00487)
Sigma L (VF0855) (LKBNICJM_00492)
LKBNICJM_00479
LKBNICJM_00481
LKBNICJM_00485
Putative tartrate transporter (LKBNICJM_00488)
LKBNICJM_00491
IMGPR_plasmid_3300014621_000010 490345:505754(+)
PGLPPFML_00077
IMGPR_plasmid_3300012078_000013 69325:84734(-)
Phosphoethanolamine transferase EptA (IDFCCEHF_00036)
Phosphoethanolamine transferase EptA (IDFCCEHF_00037)
HTH-type transcriptional regulator BenM (IDFCCEHF_00035)
IDFCCEHF_00038
RutC family protein YjgH (IDFCCEHF_00028)
IMGPR_plasmid_648028049_000003 25941:41881(-)
AEBJBODH_00419
IMGPR_plasmid_2602042078_000001 437829:449513(+)
EAAKFIKP_00175
IMGPR_plasmid_2582581300_000002 166024:177709(-)
JBDOHLEL_00166
JBDOHLEL_00169
IMGPR_plasmid_2757320372_000002 173131:184584(+)
COPMJMBM_00233
COPMJMBM_00239
COPMJMBM_00235
COPMJMBM_00232
COPMJMBM_00240
COPMJMBM_00238
COPMJMBM_00234
IMGPR_plasmid_2811995300_000002 233953:249900(-)
6000 nt
Virulence
AMR