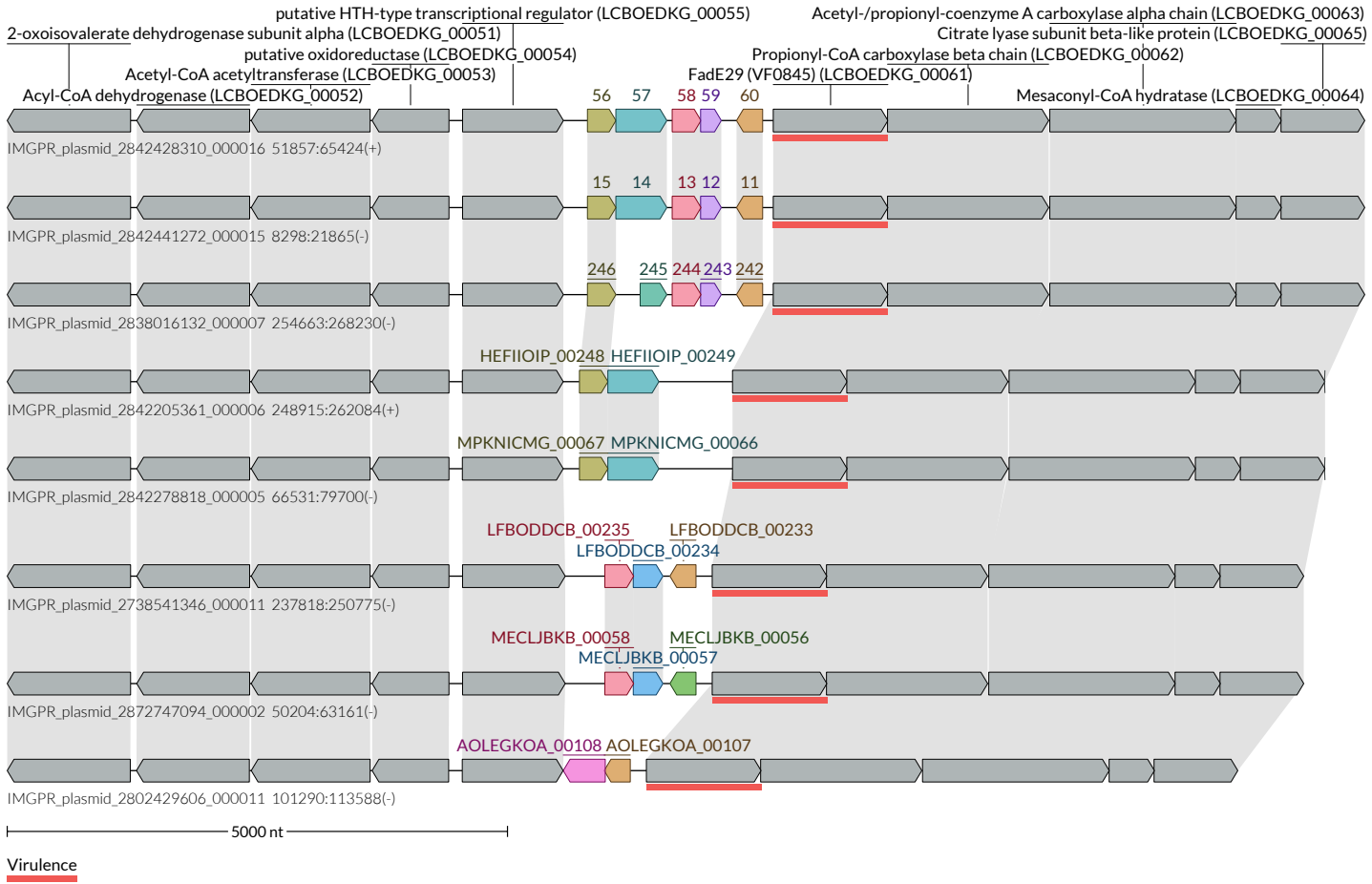
putative HTH-type transcriptional regulator (LCBOEDKG_00055)
Acetyl-/propionyl-coenzyme A carboxylase alpha chain (LCBOEDKG_00063)
2-oxoisovalerate dehydrogenase subunit alpha (LCBOEDKG_00051)
Citrate lyase subunit beta-like protein (LCBOEDKG_00065)
putative oxidoreductase (LCBOEDKG_00054)
Propionyl-CoA carboxylase beta chain (LCBOEDKG_00062)
Acetyl-CoA acetyltransferase (LCBOEDKG_00053)
FadE29 (VF0845) (LCBOEDKG_00061)
Acyl-CoA dehydrogenase (LCBOEDKG_00052)
Mesaconyl-CoA hydratase (LCBOEDKG_00064)
56 57 58 59 60
IMGPR_plasmid_2842428310_000016 51857:65424(+)
15 14 13 12 11
IMGPR_plasmid_2842441272_000015 8298:21865(-)
246 245 244 243 242
IMGPR_plasmid_2838016132_000007 254663:268230(-)
HEFIIOIP_00248 HEFIIOIP_00249
IMGPR_plasmid_2842205361_000006 248915:262084(+)
MPKNICMG_00067 MPKNICMG_00066
IMGPR_plasmid_2842278818_000005 66531:79700(-)
LFBODDCB_00235 LFBODDCB_00233
LFBODDCB_00234
IMGPR_plasmid_2738541346_000011 237818:250775(-)
MECLJBKB_00058 MECLJBKB_00056
MECLJBKB_00057
IMGPR_plasmid_2872747094_000002 50204:63161(-)
AOLEGKOA_00108 AOLEGKOA_00107
IMGPR_plasmid_2802429606_000011 101290:113588(-)
5000 nt
Virulence