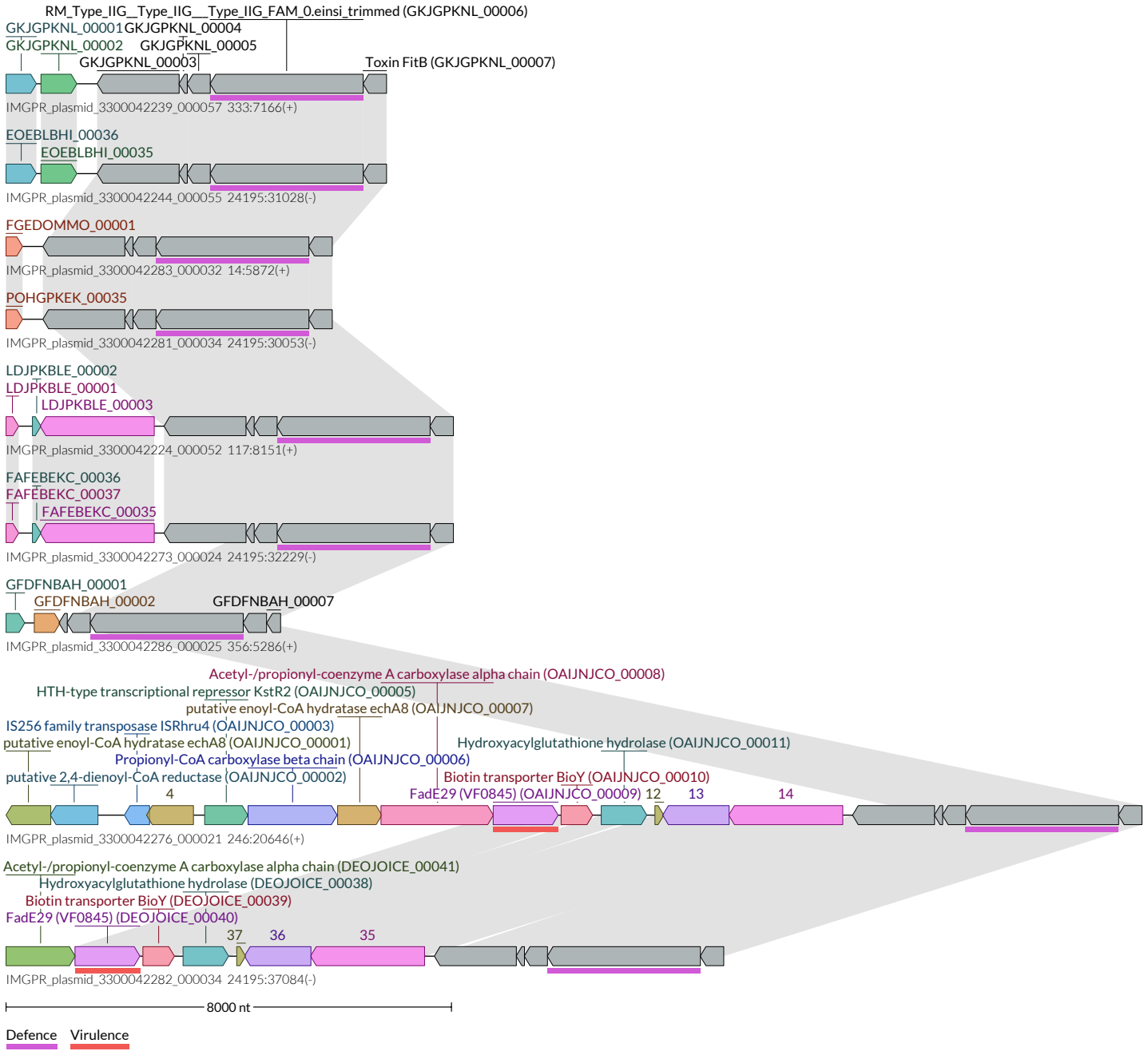
RM_Type_IIG__Type_IIG__Type_IIG_FAM_0.einsi_trimmed (GKJGPKNL_00006)
GKJGPKNL_00001
GKJGPKNL_00004
GKJGPKNL_00002
GKJGPKNL_00005
GKJGPKNL_00003
Toxin FitB (GKJGPKNL_00007)
IMGPR_plasmid_3300042239_000057 333:7166(+)
EOEBLBHI_00036
EOEBLBHI_00035
IMGPR_plasmid_3300042244_000055 24195:31028(-)
FGEDOMMO_00001
IMGPR_plasmid_3300042283_000032 14:5872(+)
POHGPKEK_00035
IMGPR_plasmid_3300042281_000034 24195:30053(-)
LDJPKBLE_00002
LDJPKBLE_00001
LDJPKBLE_00003
IMGPR_plasmid_3300042224_000052 117:8151(+)
FAFEBEKC_00036
FAFEBEKC_00037
FAFEBEKC_00035
IMGPR_plasmid_3300042273_000024 24195:32229(-)
GFDFNBAH_00001
GFDFNBAH_00002
GFDFNBAH_00007
IMGPR_plasmid_3300042286_000025 356:5286(+)
Acetyl-/propionyl-coenzyme A carboxylase alpha chain (OAIJNJCO_00008)
HTH-type transcriptional repressor KstR2 (OAIJNJCO_00005)
putative enoyl-CoA hydratase echA8 (OAIJNJCO_00007)
IS256 family transposase ISRhru4 (OAIJNJCO_00003)
putative enoyl-CoA hydratase echA8 (OAIJNJCO_00001)
Hydroxyacylglutathione hydrolase (OAIJNJCO_00011)
Propionyl-CoA carboxylase beta chain (OAIJNJCO_00006)
putative 2,4-dienoyl-CoA reductase (OAIJNJCO_00002)
Biotin transporter BioY (OAIJNJCO_00010)
4
FadE29 (VF0845) (OAIJNJCO_00009)
12
13
14
IMGPR_plasmid_3300042276_000021 246:20646(+)
Acetyl-/propionyl-coenzyme A carboxylase alpha chain (DEOJOICE_00041)
Hydroxyacylglutathione hydrolase (DEOJOICE_00038)
Biotin transporter BioY (DEOJOICE_00039)
FadE29 (VF0845) (DEOJOICE_00040)
37
36
35
IMGPR_plasmid_3300042282_000034 24195:37084(-)
8000 nt
Defence
Virulence