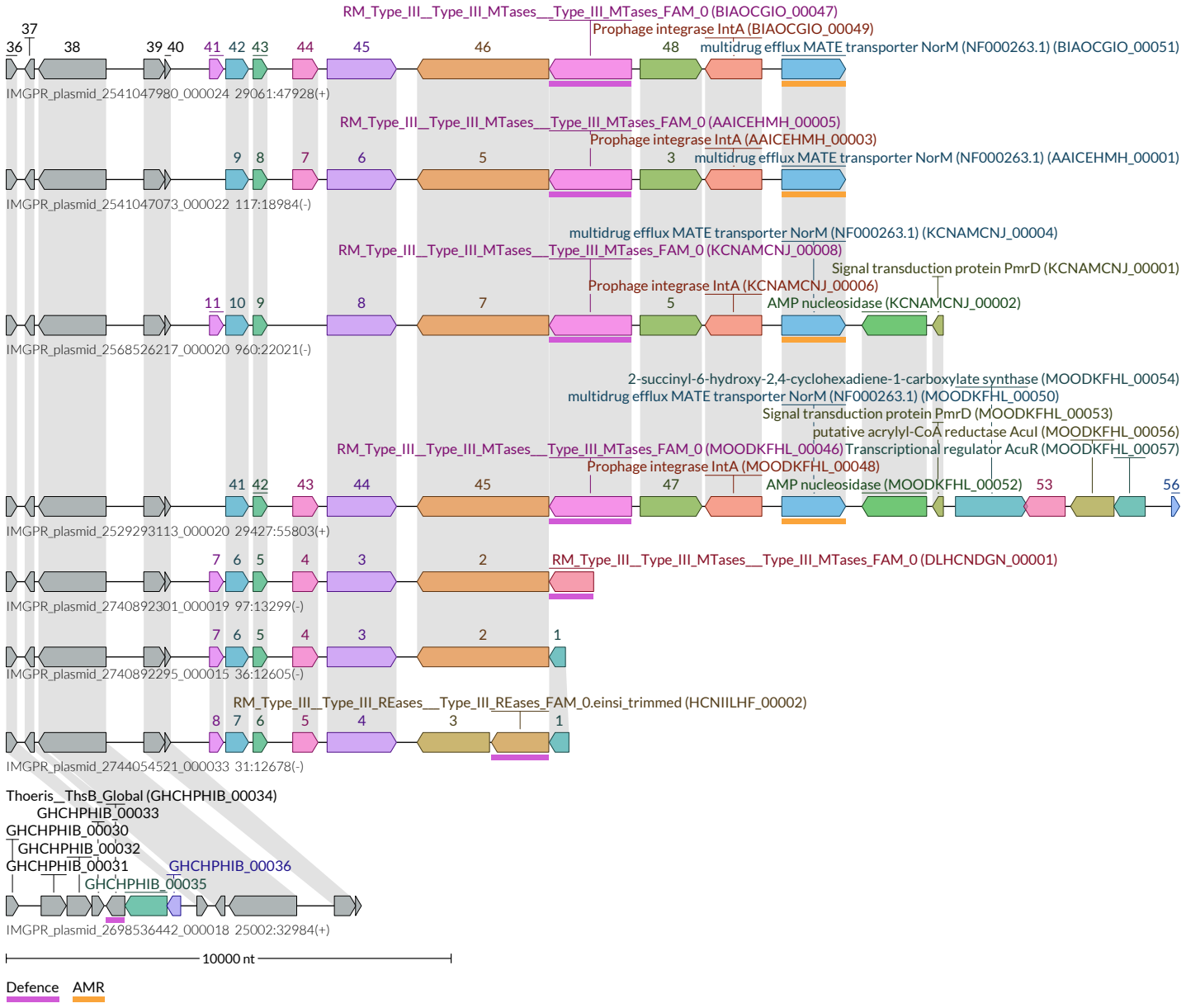
RM_Type_III__Type_III_MTases___Type_III_MTases_FAM_0 (BIAOCGIO_00047)
Prophage integrase IntA (BIAOCGIO_00049)
multidrug efflux MATE transporter NorM (NF000263.1) (BIAOCGIO_00051)
IMGPR_plasmid_2541047980_000024 29061:47928(+)
RM_Type_III__Type_III_MTases___Type_III_MTases_FAM_0 (AAICEHMH_00005)
Prophage integrase IntA (AAICEHMH_00003)
multidrug efflux MATE transporter NorM (NF000263.1) (AAICEHMH_00001)
IMGPR_plasmid_2541047073_000022 117:18984(-)
multidrug efflux MATE transporter NorM (NF000263.1) (KCNAMCNJ_00004)
RM_Type_III__Type_III_MTases___Type_III_MTases_FAM_0 (KCNAMCNJ_00008)
Signal transduction protein PmrD (KCNAMCNJ_00001)
Prophage integrase IntA (KCNAMCNJ_00006)
AMP nucleosidase (KCNAMCNJ_00002)
IMGPR_plasmid_2568526217_000020 960:22021(-)
2-succinyl-6-hydroxy-2,4-cyclohexadiene-1-carboxylate synthase (MOODKFHL_00054)
multidrug efflux MATE transporter NorM (NF000263.1) (MOODKFHL_00050)
Signal transduction protein PmrD (MOODKFHL_00053)
putative acrylyl-CoA reductase AcuI (MOODKFHL_00056)
RM_Type_III__Type_III_MTases___Type_III_MTases_FAM_0 (MOODKFHL_00046)
Transcriptional regulator AcuR (MOODKFHL_00057)
Prophage integrase IntA (MOODKFHL_00048)
AMP nucleosidase (MOODKFHL_00052)
IMGPR_plasmid_2529293113_000020 29427:55803(+)
RM_Type_III__Type_III_MTases___Type_III_MTases_FAM_0 (DLHCNDGN_00001)
IMGPR_plasmid_2740892301_000019 97:13299(-)
IMGPR_plasmid_2740892295_000015 36:12605(-)
RM_Type_III__Type_III_REases___Type_III_REases_FAM_0.einsi_trimmed (HCNIILHF_00002)
IMGPR_plasmid_2744054521_000033 31:12678(-)
Thoeris__ThsB_Global (GHCHPHIB_00034)
GHCHPHIB_00033
GHCHPHIB_00030
GHCHPHIB_00032
GHCHPHIB_00031
GHCHPHIB_00036
GHCHPHIB_00035
IMGPR_plasmid_2698536442_000018 25002:32984(+)
10000 nt
Defence
AMR