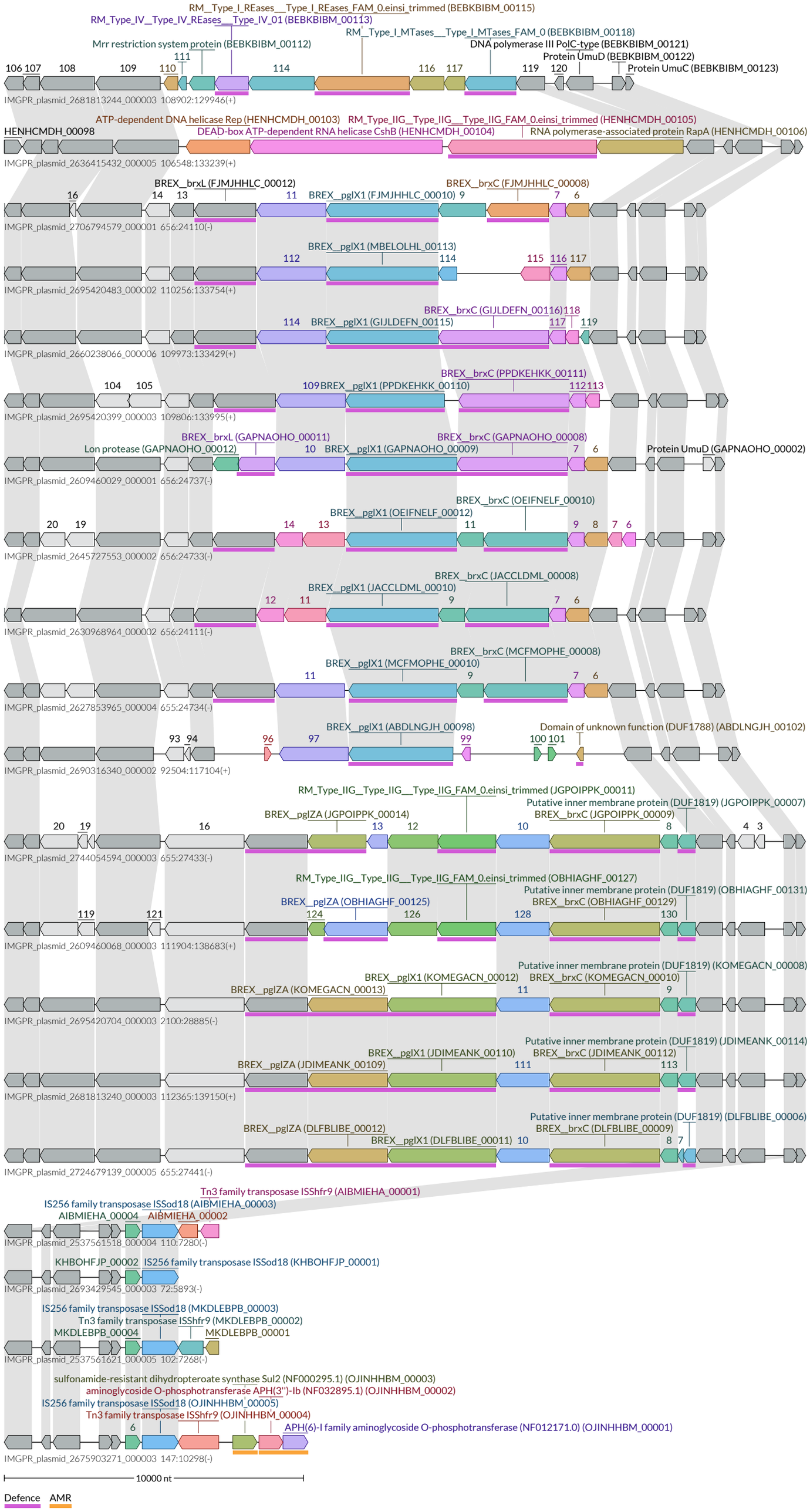
RM_Type_I_REases__Type_I_REases_FAM_0.einsi_trimmed (BEBKBIBM_00115)
RM_Type_IV__Type_IV_REases__Type_IV_01 (BEBKBIBM_00113)
RM_Type_I_MTases__Type_I_MTases_FAM_0 (BEBKBIBM_00118)
Mrr restriction system protein (BEBKBIBM_00112)
DNA polymerase III PolC-type (BEBKBIBM_00121)
Protein UmuD (BEBKBIBM_00122)
Protein UmuC (BEBKBIBM_00123)
IMGPR_plasmid_2681813244_000003 108902:129946(+)
ATP-dependent DNA helicase Rep (HENHCMDH_00103)
RM_Type_IIG__Type_IIG__Type_IIG_FAM_0.einsi_trimmed (HENHCMDH_00105)
HENHCMDH_00098
DEAD-box ATP-dependent RNA helicase CshB (HENHCMDH_00104)
RNA polymerase-associated protein RapA (HENHCMDH_00106)
IMGPR_plasmid_2636415432_000005 106548:133239(+)
BREX__brxL (FJMJHHLC_00012)
BREX__pglX1 (FJMJHHLC_00010)
BREX__brxC (FJMJHHLC_00008)
IMGPR_plasmid_2706794579_000001 656:24110(-)
BREX__pglX1 (MBELOLHL_00113)
IMGPR_plasmid_2695420483_000002 110256:133754(+)
BREX__brxC (GIJLDEFN_00116)
BREX__pglX1 (GIJLDEFN_00115)
IMGPR_plasmid_2660238066_000006 109973:133429(+)
BREX__brxC (PPDKEHKK_00111)
BREX__pglX1 (PPDKEHKK_00110)
IMGPR_plasmid_2695420399_000003 109806:133995(+)
BREX__brxL (GAPNAOHO_00011)
Lon protease (GAPNAOHO_00012)
BREX__brxC (GAPNAOHO_00008)
BREX__pglX1 (GAPNAOHO_00009)
Protein UmuD (GAPNAOHO_00002)
IMGPR_plasmid_2609460029_000001 656:24737(-)
BREX__brxC (OEIFNELF_00010)
BREX__pglX1 (OEIFNELF_00012)
IMGPR_plasmid_2645727553_000002 656:24733(-)
BREX__brxC (JACCLDML_00008)
BREX__pglX1 (JACCLDML_00010)
IMGPR_plasmid_2630968964_000002 656:24111(-)
BREX__brxC (MCFMOPHE_00008)
BREX__pglX1 (MCFMOPHE_00010)
IMGPR_plasmid_2627853965_000004 655:24734(-)
BREX__pglX1 (ABDLNGJH_00098)
Domain of unknown function (DUF1788) (ABDLNGJH_00102)
IMGPR_plasmid_2690316340_000002 92504:117104(+)
RM_Type_IIG__Type_IIG__Type_IIG_FAM_0.einsi_trimmed (JGPOIPPK_00011)
Putative inner membrane protein (DUF1819) (JGPOIPPK_00007)
BREX__pglZA (JGPOIPPK_00014)
BREX__brxC (JGPOIPPK_00009)
IMGPR_plasmid_2744054594_000003 655:27433(-)
RM_Type_IIG__Type_IIG__Type_IIG_FAM_0.einsi_trimmed (OBHIAGHF_00127)
Putative inner membrane protein (DUF1819) (OBHIAGHF_00131)
BREX__pglZA (OBHIAGHF_00125)
BREX__brxC (OBHIAGHF_00129)
IMGPR_plasmid_2609460068_000003 111904:138683(+)
Putative inner membrane protein (DUF1819) (KOMEGACN_00008)
BREX__pglX1 (KOMEGACN_00012)
BREX__brxC (KOMEGACN_00010)
BREX__pglZA (KOMEGACN_00013)
IMGPR_plasmid_2695420704_000003 2100:28885(-)
Putative inner membrane protein (DUF1819) (JDIMEANK_00114)
BREX__pglX1 (JDIMEANK_00110)
BREX__brxC (JDIMEANK_00112)
BREX__pglZA (JDIMEANK_00109)
IMGPR_plasmid_2681813240_000003 112365:139150(+)
Putative inner membrane protein (DUF1819) (DLFBLIBE_00006)
BREX__pglZA (DLFBLIBE_00012)
BREX__brxC (DLFBLIBE_00009)
BREX__pglX1 (DLFBLIBE_00011)
IMGPR_plasmid_2724679139_000005 655:27441(-)
Tn3 family transposase ISShfr9 (AIBMIEHA_00001)
IS256 family transposase ISSod18 (AIBMIEHA_00003)
AIBMIEHA_00004
AIBMIEHA_00002
IMGPR_plasmid_2537561518_000004 110:7280(-)
KHBOHFJP_00002
IS256 family transposase ISSod18 (KHBOHFJP_00001)
IMGPR_plasmid_2693429545_000003 72:5893(-)
IS256 family transposase ISSod18 (MKDLEBPB_00003)
Tn3 family transposase ISShfr9 (MKDLEBPB_00002)
MKDLEBPB_00004
MKDLEBPB_00001
IMGPR_plasmid_2537561621_000005 102:7268(-)
sulfonamide-resistant dihydropteroate synthase Sul2 (NF000295.1) (OJINHHBM_00003)
aminoglycoside O-phosphotransferase APH(3'')-Ib (NF032895.1) (OJINHHBM_00002)
IS256 family transposase ISSod18 (OJINHHBM_00005)
Tn3 family transposase ISShfr9 (OJINHHBM_00004)
APH(6)-I family aminoglycoside O-phosphotransferase (NF012171.0) (OJINHHBM_00001)
IMGPR_plasmid_2675903271_000003 147:10298(-)
10000 nt
Defence
AMR