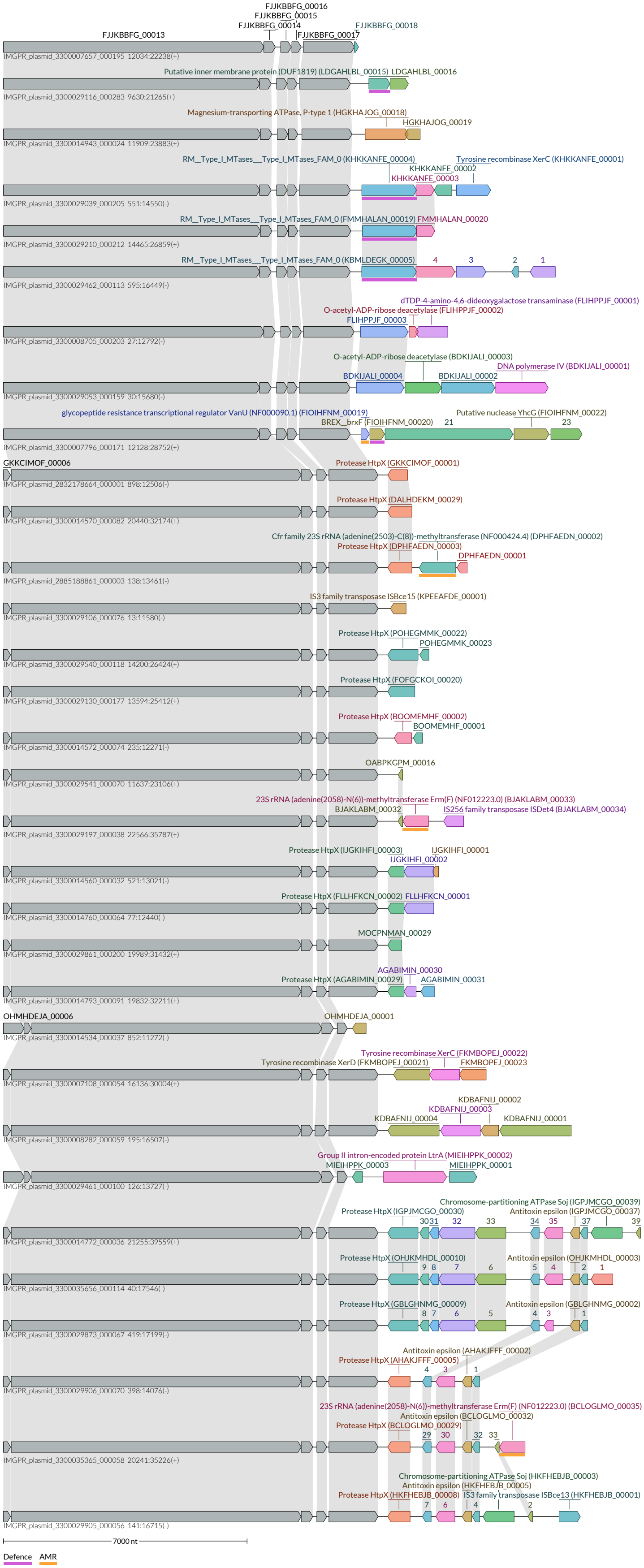
FJJKBBFG_00016
FJJKBBFG_00015
FJJKBBFG_00014
FJJKBBFG_00018
FJJKBBFG_00013
FJJKBBFG_00017
IMGPR_plasmid_3300007657_000195 12034:22238(+)
Putative inner membrane protein (DUF1819) (LDGAHLBL_00015) LDGAHLBL_00016
IMGPR_plasmid_3300029116_000283 9630:21265(+)
Magnesium-transporting ATPase, P-type 1 (HGKHAJOG_00018)
HGKHAJOG_00019
IMGPR_plasmid_3300014943_000024 11909:23883(+)
RM__Type_I_MTases__Type_I_MTases_FAM_0 (KHKKANFE_00004)
Tyrosine recombinase XerC (KHKKANFE_00001)
KHKKANFE_00002
KHKKANFE_00003
IMGPR_plasmid_3300029039_000205 551:14550(-)
RM__Type_I_MTases__Type_I_MTases_FAM_0 (FMMHALAN_00019) FMMHALAN_00020
IMGPR_plasmid_3300029210_000212 14465:26859(+)
RM__Type_I_MTases__Type_I_MTases_FAM_0 (KBMLDEGK_00005)
4
3
2
1
IMGPR_plasmid_3300029462_000113 595:16449(-)
dTDP-4-amino-4,6-dideoxygalactose transaminase (FLIHPPJF_00001)
O-acetyl-ADP-ribose deacetylase (FLIHPPJF_00002)
FLIHPPJF_00003
IMGPR_plasmid_3300008705_000203 27:12792(-)
O-acetyl-ADP-ribose deacetylase (BDKIJALI_00003)
DNA polymerase IV (BDKIJALI_00001)
BDKIJALI_00004
BDKIJALI_00002
IMGPR_plasmid_3300029053_000159 30:15680(-)
glycopeptide resistance transcriptional regulator VanU (NF000090.1) (FIOIHFNM_00019)
Putative nuclease YhcG (FIOIHFNM_00022)
BREX__brxF (FIOIHFNM_00020)
21
23
IMGPR_plasmid_3300007796_000171 12128:28752(+)
GKKCIMOF_00006
Protease HtpX (GKKCIMOF_00001)
IMGPR_plasmid_2832178664_000001 898:12506(-)
Protease HtpX (DALHDEKM_00029)
IMGPR_plasmid_3300014570_000082 20440:32174(+)
Cfr family 23S rRNA (adenine(2503)-C(8))-methyltransferase (NF000424.4) (DPHFAEDN_00002)
Protease HtpX (DPHFAEDN_00003)
DPHFAEDN_00001
IMGPR_plasmid_2885188861_000003 138:13461(-)
IS3 family transposase ISBce15 (KPEEAFDE_00001)
IMGPR_plasmid_3300029106_000076 13:11580(-)
Protease HtpX (POHEGMMK_00022)
POHEGMMK_00023
IMGPR_plasmid_3300029540_000118 14200:26424(+)
Protease HtpX (FOFGCKOI_00020)
IMGPR_plasmid_3300029130_000177 13594:25412(+)
Protease HtpX (BOOMEMHF_00002)
BOOMEMHF_00001
IMGPR_plasmid_3300014572_000074 235:12271(-)
OABPKGPM_00016
IMGPR_plasmid_3300029541_000070 11637:23106(+)
23S rRNA (adenine(2058)-N(6))-methyltransferase Erm(F) (NF012223.0) (BJAKLABM_00033)
BJAKLABM_00032
IS256 family transposase ISDet4 (BJAKLABM_00034)
IMGPR_plasmid_3300029197_000038 22566:35787(+)
Protease HtpX (IJGKIHFI_00003)
IJGKIHFI_00001
IJGKIHFI_00002
IMGPR_plasmid_3300014560_000032 521:13021(-)
Protease HtpX (FLLHFKCN_00002) FLLHFKCN_00001
IMGPR_plasmid_3300014760_000064 77:12440(-)
MOCPNMAN_00029
IMGPR_plasmid_3300029861_000200 19989:31432(+)
AGABIMIN_00030
Protease HtpX (AGABIMIN_00029)
AGABIMIN_00031
IMGPR_plasmid_3300014793_000091 19832:32211(+)
OHMHDEJA_00006
OHMHDEJA_00001
IMGPR_plasmid_3300014534_000037 852:11272(-)
Tyrosine recombinase XerC (FKMBOPEJ_00022)
Tyrosine recombinase XerD (FKMBOPEJ_00021)
FKMBOPEJ_00023
IMGPR_plasmid_3300007108_000054 16136:30004(+)
KDBAFNIJ_00002
KDBAFNIJ_00003
KDBAFNIJ_00004
KDBAFNIJ_00001
IMGPR_plasmid_3300008282_000059 195:16507(-)
Group II intron-encoded protein LtrA (MIEIHPPK_00002)
MIEIHPPK_00003
MIEIHPPK_00001
IMGPR_plasmid_3300029461_000100 126:13727(-)
Chromosome-partitioning ATPase Soj (IGPJMCGO_00039)
Protease HtpX (IGPJMCGO_00030)
Antitoxin epsilon (IGPJMCGO_00037)
30 31 32 33 34 35 37 39
IMGPR_plasmid_3300014772_000036 21255:39559(+)
Protease HtpX (OHJKMHDL_00010)
Antitoxin epsilon (OHJKMHDL_00003)
9 8 7 6 5 4 2 1
IMGPR_plasmid_3300035656_000114 40:17546(-)
Protease HtpX (GBLGHNMG_00009)
Antitoxin epsilon (GBLGHNMG_00002)
8 7 6 5 4 3 1
IMGPR_plasmid_3300029873_000067 419:17199(-)
Antitoxin epsilon (AHAKJFFF_00002)
Protease HtpX (AHAKJFFF_00005)
4 3 1
IMGPR_plasmid_3300029906_000070 398:14076(-)
23S rRNA (adenine(2058)-N(6))-methyltransferase Erm(F) (NF012223.0) (BCLOGLMO_00035)
Antitoxin epsilon (BCLOGLMO_00032)
Protease HtpX (BCLOGLMO_00029)
29 30 32 33
IMGPR_plasmid_3300035365_000058 20241:35226(+)
Chromosome-partitioning ATPase Soj (HKFHEBJB_00003)
Antitoxin epsilon (HKFHEBJB_00005)
Protease HtpX (HKFHEBJB_00008)
IS3 family transposase ISBce13 (HKFHEBJB_00001)
7 6 4 2
IMGPR_plasmid_3300029905_000056 141:16715(-)
7000 nt
Defence
AMR