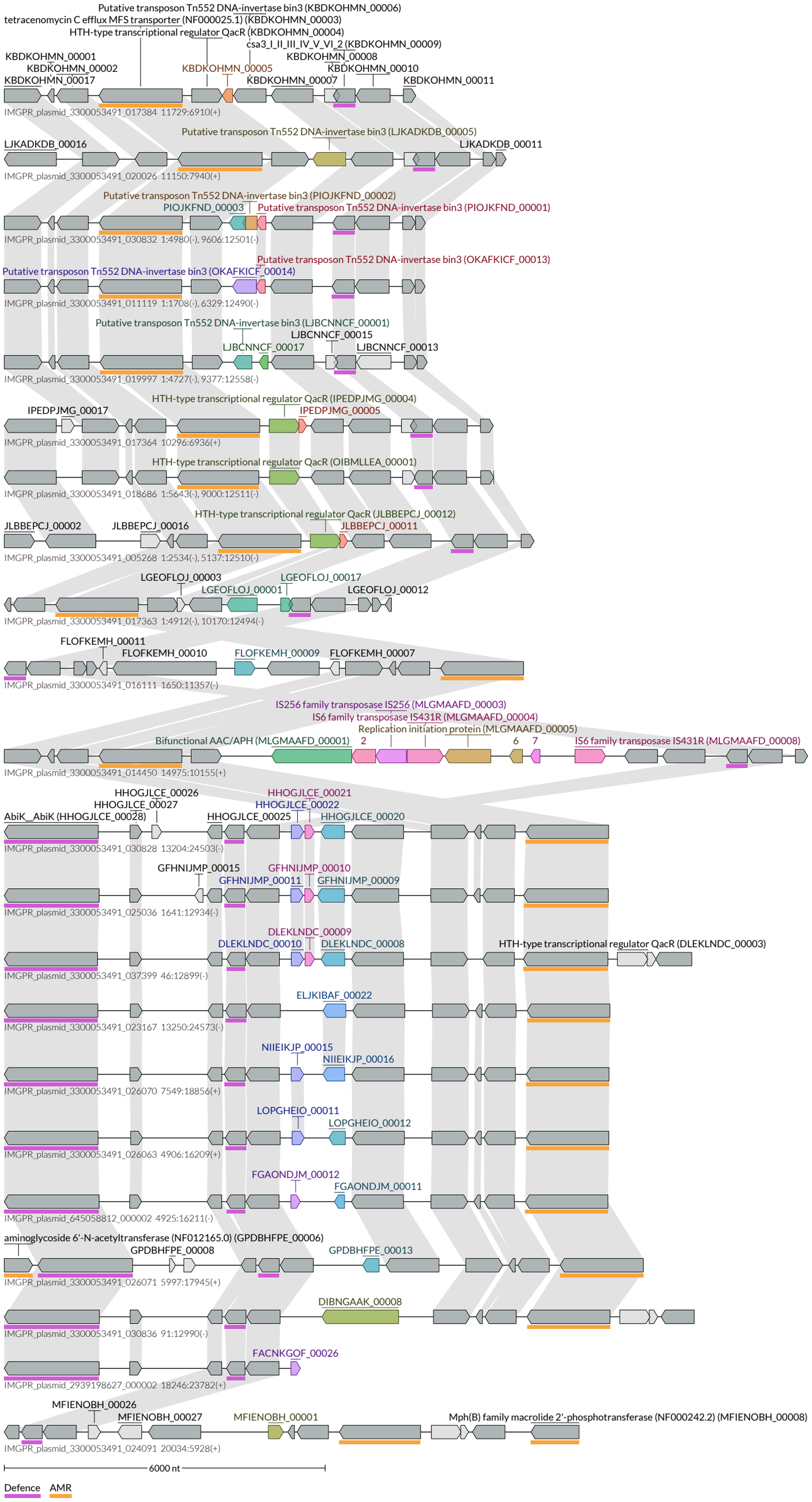
Putative transposon Tn552 DNA-invertase bin3 (KBDKOHMN_00006)
tetracenomycin C efflux MFS transporter (NF000025.1) (KBDKOHMN_00003)
HTH-type transcriptional regulator QacR (KBDKOHMN_00004)
csa3_I_II_III_IV_V_VI_2 (KBDKOHMN_00009)
KBDKOHMN_00001
KBDKOHMN_00008
KBDKOHMN_00002
KBDKOHMN_00005
KBDKOHMN_00010
KBDKOHMN_00017
KBDKOHMN_00007
KBDKOHMN_00011
IMGPR_plasmid_3300053491_017384 11729:6910(+)
Putative transposon Tn552 DNA-invertase bin3 (LJKADKDB_00005)
LJKADKDB_00016
LJKADKDB_00011
IMGPR_plasmid_3300053491_020026 11150:7940(+)
Putative transposon Tn552 DNA-invertase bin3 (PIOJKFND_00002)
PIOJKFND_00003
Putative transposon Tn552 DNA-invertase bin3 (PIOJKFND_00001)
IMGPR_plasmid_3300053491_030832 1:4980(-), 9606:12501(-)
Putative transposon Tn552 DNA-invertase bin3 (OKAFKICF_00013)
Putative transposon Tn552 DNA-invertase bin3 (OKAFKICF_00014)
IMGPR_plasmid_3300053491_011119 1:1708(-), 6329:12490(-)
Putative transposon Tn552 DNA-invertase bin3 (LJBCNNCF_00001)
LJBCNNCF_00015
LJBCNNCF_00017
LJBCNNCF_00013
IMGPR_plasmid_3300053491_019997 1:4727(-), 9377:12558(-)
HTH-type transcriptional regulator QacR (IPEDPJMG_00004)
IPEDPJMG_00017
IPEDPJMG_00005
IMGPR_plasmid_3300053491_017364 10296:6936(+)
HTH-type transcriptional regulator QacR (OIBMLLEA_00001)
IMGPR_plasmid_3300053491_018686 1:5643(-), 9000:12511(-)
HTH-type transcriptional regulator QacR (JLBBEPCJ_00012)
JLBBEPCJ_00002
JLBBEPCJ_00016
JLBBEPCJ_00011
IMGPR_plasmid_3300053491_005268 1:2534(-), 5137:12510(-)
LGEOFLOJ_00003
LGEOFLOJ_00017
LGEOFLOJ_00001
LGEOFLOJ_00012
IMGPR_plasmid_3300053491_017363 1:4912(-), 10170:12494(-)
FLOFKEMH_00011
FLOFKEMH_00010
FLOFKEMH_00009
FLOFKEMH_00007
IMGPR_plasmid_3300053491_016111 1650:11357(-)
IS256 family transposase IS256 (MLGMAAFD_00003)
IS6 family transposase IS431R (MLGMAAFD_00004)
Replication initiation protein (MLGMAAFD_00005)
Bifunctional AAC/APH (MLGMAAFD_00001)
2
6
7
IS6 family transposase IS431R (MLGMAAFD_00008)
IMGPR_plasmid_3300053491_014450 14975:10155(+)
HHOGJLCE_00026
HHOGJLCE_00021
HHOGJLCE_00027
HHOGJLCE_00022
AbiK__AbiK (HHOGJLCE_00028)
HHOGJLCE_00025
HHOGJLCE_00020
IMGPR_plasmid_3300053491_030828 13204:24503(-)
GFHNIJMP_00015
GFHNIJMP_00010
GFHNIJMP_00011
GFHNIJMP_00009
IMGPR_plasmid_3300053491_025036 1641:12934(-)
DLEKLNDC_00009
DLEKLNDC_00010
DLEKLNDC_00008
HTH-type transcriptional regulator QacR (DLEKLNDC_00003)
IMGPR_plasmid_3300053491_037399 46:12899(-)
ELJKIBAF_00022
IMGPR_plasmid_3300053491_023167 13250:24573(-)
NIIEIKJP_00015
NIIEIKJP_00016
IMGPR_plasmid_3300053491_026070 7549:18856(+)
LOPGHEIO_00011
LOPGHEIO_00012
IMGPR_plasmid_3300053491_026063 4906:16209(+)
FGAONDJM_00012
FGAONDJM_00011
IMGPR_plasmid_645058812_000002 4925:16211(-)
aminoglycoside 6'-N-acetyltransferase (NF012165.0) (GPDBHFPE_00006)
GPDBHFPE_00008
GPDBHFPE_00013
IMGPR_plasmid_3300053491_026071 5997:17945(+)
DIBNGAAK_00008
IMGPR_plasmid_3300053491_030836 91:12990(-)
FACNKGOF_00026
IMGPR_plasmid_2939198627_000002 18246:23782(+)
MFIENOBH_00026
MFIENOBH_00027
MFIENOBH_00001
Mph(B) family macrolide 2'-phosphotransferase (NF000242.2) (MFIENOBH_00008)
IMGPR_plasmid_3300053491_024091 20034:5928(+)
6000 nt
Defence
AMR