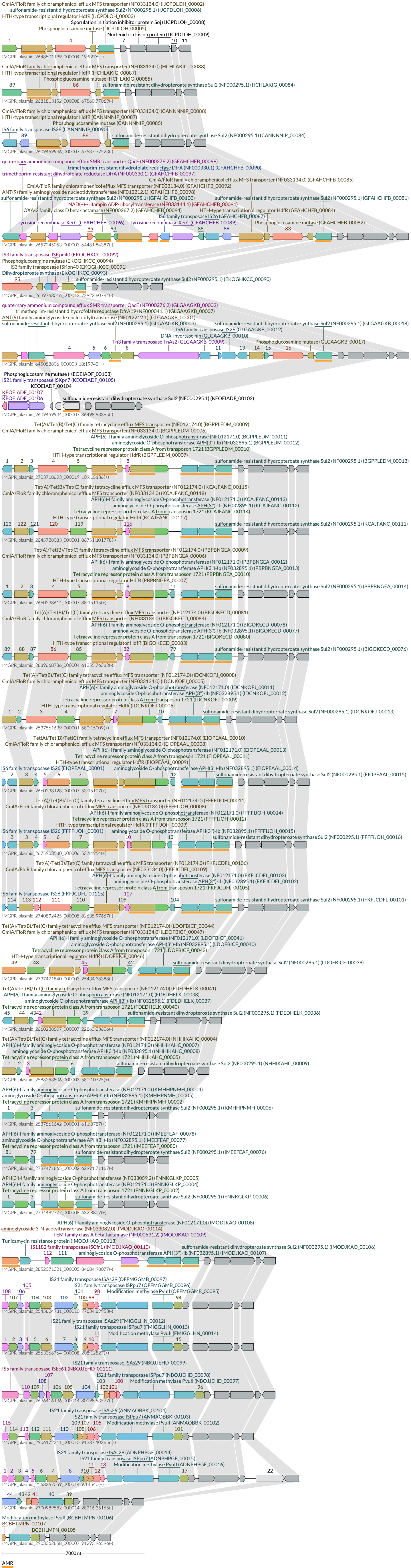
CmlA/FloR family chloramphenicol efflux MFS transporter (NF033134.0) (IJCPDLOH_00002)
sulfonamide-resistant dihydropteroate synthase Sul2 (NF000295.1) (IJCPDLOH_00006)
HTH-type transcriptional regulator HdfR (IJCPDLOH_00003)
Sporulation initiation inhibitor protein Soj (IJCPDLOH_00008)
Phosphoglucosamine mutase (IJCPDLOH_00005)
Nucleoid occlusion protein (IJCPDLOH_00009)
IMGPR_plasmid_2648501199_000004 19:9276(+)
CmlA/FloR family chloramphenicol efflux MFS transporter (NF033134.0) (HCHLAKIG_00088)
HTH-type transcriptional regulator HdfR (HCHLAKIG_00087)
Phosphoglucosamine mutase (HCHLAKIG_00085)
sulfonamide-resistant dihydropteroate synthase Sul2 (NF000295.1) (HCHLAKIG_00084)
IMGPR_plasmid_2681813152_000008 67560:77069(-)
CmlA/FloR family chloramphenicol efflux MFS transporter (NF033134.0) (CANNNNIP_00088)
HTH-type transcriptional regulator HdfR (CANNNNIP_00087)
Phosphoglucosamine mutase (CANNNNIP_00085)
IS6 family transposase IS26 (CANNNNIP_00090)
sulfonamide-resistant dihydropteroate synthase Sul2 (NF000295.1) (CANNNNIP_00084)
IMGPR_plasmid_2609459946_000007 67537:77523(-)
quaternary ammonium compound efflux SMR transporter QacE (NF000276.2) (GFAHCHFB_00099)
trimethoprim-resistant dihydrofolate reductase DfrA (NF000330.1) (GFAHCHFB_00090)
trimethoprim-resistant dihydrofolate reductase DfrA (NF000330.1) (GFAHCHFB_00097)
CmlA/FloR family chloramphenicol efflux MFS transporter (NF033134.0) (GFAHCHFB_00085)
CmlA/FloR family chloramphenicol efflux MFS transporter (NF033134.0) (GFAHCHFB_00092)
ANT(9) family aminoglycoside nucleotidyltransferase (NF012212.1) (GFAHCHFB_00098)
sulfonamide-resistant dihydropteroate synthase Sul2 (NF000295.1) (GFAHCHFB_00100)
sulfonamide-resistant dihydropteroate synthase Sul2 (NF000295.1) (GFAHCHFB_00081)
NAD(+)--rifampin ADP-ribosyltransferase (NF033144.1) (GFAHCHFB_00091)
OXA-2 family class D beta-lactamase (NF000267.2) (GFAHCHFB_00094)
HTH-type transcriptional regulator HdfR (GFAHCHFB_00084)
IS6 family transposase IS26 (GFAHCHFB_00087)
Tyrosine recombinase XerC (GFAHCHFB_00096)
Tyrosine recombinase XerC (GFAHCHFB_00089)
Phosphoglucosamine mutase (GFAHCHFB_00082)
IMGPR_plasmid_2657245053_000003 64481:84387(-)
IS3 family transposase ISKpn40 (EKOGHKCC_00092)
Phosphoglucosamine mutase (EKOGHKCC_00094)
IS3 family transposase ISKpn40 (EKOGHKCC_00091)
Dihydropteroate synthase (EKOGHKCC_00093)
sulfonamide-resistant dihydropteroate synthase Sul2 (NF000295.1) (EKOGHKCC_00090)
IMGPR_plasmid_2639763016_000013 72923:80784(-)
quaternary ammonium compound efflux SMR transporter QacE (NF000276.2) (GLGAAGKB_00002)
trimethoprim-resistant dihydrofolate reductase DfrA19 (NF000041.1) (GLGAAGKB_00007)
ANT(9) family aminoglycoside nucleotidyltransferase (NF012212.1) (GLGAAGKB_00001)
sulfonamide-resistant dihydropteroate synthase Sul2 (NF000295.1) (GLGAAGKB_00003)
sulfonamide-resistant dihydropteroate synthase Sul2 (NF000295.1) (GLGAAGKB_00018)
IS6 family transposase IS26 (GLGAAGKB_00012)
DNA-invertase hin (GLGAAGKB_00010)
Tn3 family transposase TnAs2 (GLGAAGKB_00009)
Phosphoglucosamine mutase (GLGAAGKB_00017)
IMGPR_plasmid_645058806_000003 18:19943(+)
Phosphoglucosamine mutase (KEOEIADF_00103)
IS21 family transposase ISKpn7 (KEOEIADF_00105)
KEOEIADF_00104
KEOEIADF_00107
KEOEIADF_00106
sulfonamide-resistant dihydropteroate synthase Sul2 (NF000295.1) (KEOEIADF_00102)
IMGPR_plasmid_2609459934_000007 86498:93365(-)
Tet(A)/Tet(B)/Tet(C) family tetracycline efflux MFS transporter (NF012174.0) (BGPPLEDM_00009)
CmlA/FloR family chloramphenicol efflux MFS transporter (NF033134.0) (BGPPLEDM_00006)
APH(6)-I family aminoglycoside O-phosphotransferase (NF012171.0) (BGPPLEDM_00011)
aminoglycoside O-phosphotransferase APH(3'')-Ib (NF032895.1) (BGPPLEDM_00012)
Tetracycline repressor protein class A from transposon 1721 (BGPPLEDM_00010)
HTH-type transcriptional regulator HdfR (BGPPLEDM_00007)
sulfonamide-resistant dihydropteroate synthase Sul2 (NF000295.1) (BGPPLEDM_00013)
IMGPR_plasmid_2703718693_000019 109:15136(+)
Tet(A)/Tet(B)/Tet(C) family tetracycline efflux MFS transporter (NF012174.0) (KCAJFANC_00115)
CmlA/FloR family chloramphenicol efflux MFS transporter (NF033134.0) (KCAJFANC_00118)
APH(6)-I family aminoglycoside O-phosphotransferase (NF012171.0) (KCAJFANC_00113)
aminoglycoside O-phosphotransferase APH(3'')-Ib (NF032895.1) (KCAJFANC_00112)
Tetracycline repressor protein class A from transposon 1721 (KCAJFANC_00114)
HTH-type transcriptional regulator HdfR (KCAJFANC_00117)
sulfonamide-resistant dihydropteroate synthase Sul2 (NF000295.1) (KCAJFANC_00111)
IMGPR_plasmid_2645728083_000001 86751:101778(-)
Tet(A)/Tet(B)/Tet(C) family tetracycline efflux MFS transporter (NF012174.0) (PBPBNGEA_00009)
CmlA/FloR family chloramphenicol efflux MFS transporter (NF033134.0) (PBPBNGEA_00006)
APH(6)-I family aminoglycoside O-phosphotransferase (NF012171.0) (PBPBNGEA_00012)
aminoglycoside O-phosphotransferase APH(3'')-Ib (NF032895.1) (PBPBNGEA_00013)
Tetracycline repressor protein class A from transposon 1721 (PBPBNGEA_00010)
HTH-type transcriptional regulator HdfR (PBPBNGEA_00007)
sulfonamide-resistant dihydropteroate synthase Sul2 (NF000295.1) (PBPBNGEA_00014)
IMGPR_plasmid_2660238614_000007 88:15115(+)
Tet(A)/Tet(B)/Tet(C) family tetracycline efflux MFS transporter (NF012174.0) (BIGOKECD_00081)
CmlA/FloR family chloramphenicol efflux MFS transporter (NF033134.0) (BIGOKECD_00084)
APH(6)-I family aminoglycoside O-phosphotransferase (NF012171.0) (BIGOKECD_00078)
aminoglycoside O-phosphotransferase APH(3'')-Ib (NF032895.1) (BIGOKECD_00077)
Tetracycline repressor protein class A from transposon 1721 (BIGOKECD_00080)
HTH-type transcriptional regulator HdfR (BIGOKECD_00083)
sulfonamide-resistant dihydropteroate synthase Sul2 (NF000295.1) (BIGOKECD_00076)
IMGPR_plasmid_2889668736_000004 61355:76382(-)
Tet(A)/Tet(B)/Tet(C) family tetracycline efflux MFS transporter (NF012174.0) (IDCNKOFJ_00008)
CmlA/FloR family chloramphenicol efflux MFS transporter (NF033134.0) (IDCNKOFJ_00005)
APH(6)-I family aminoglycoside O-phosphotransferase (NF012171.0) (IDCNKOFJ_00011)
aminoglycoside O-phosphotransferase APH(3'')-Ib (NF032895.1) (IDCNKOFJ_00012)
Tetracycline repressor protein class A from transposon 1721 (IDCNKOFJ_00009)
HTH-type transcriptional regulator HdfR (IDCNKOFJ_00006)
sulfonamide-resistant dihydropteroate synthase Sul2 (NF000295.1) (IDCNKOFJ_00013)
IMGPR_plasmid_2537561639_000001 580:15009(+)
Tet(A)/Tet(B)/Tet(C) family tetracycline efflux MFS transporter (NF012174.0) (EIOPEAAL_00010)
CmlA/FloR family chloramphenicol efflux MFS transporter (NF033134.0) (EIOPEAAL_00008)
APH(6)-I family aminoglycoside O-phosphotransferase (NF012171.0) (EIOPEAAL_00013)
Tetracycline repressor protein class A from transposon 1721 (EIOPEAAL_00011)
HTH-type transcriptional regulator HdfR (EIOPEAAL_00009)
IS6 family transposase IS26 (EIOPEAAL_00001)
aminoglycoside O-phosphotransferase APH(3'')-Ib (NF032895.1) (EIOPEAAL_00014)
sulfonamide-resistant dihydropteroate synthase Sul2 (NF000295.1) (EIOPEAAL_00015)
IMGPR_plasmid_2660238128_000007 53:15107(+)
Tet(A)/Tet(B)/Tet(C) family tetracycline efflux MFS transporter (NF012174.0) (FFFFIJOH_00011)
CmlA/FloR family chloramphenicol efflux MFS transporter (NF033134.0) (FFFFIJOH_00008)
APH(6)-I family aminoglycoside O-phosphotransferase (NF012171.0) (FFFFIJOH_00014)
Tetracycline repressor protein class A from transposon 1721 (FFFFIJOH_00012)
HTH-type transcriptional regulator HdfR (FFFFIJOH_00009)
IS6 family transposase IS26 (FFFFIJOH_00001)
aminoglycoside O-phosphotransferase APH(3'')-Ib (NF032895.1) (FFFFIJOH_00015)
sulfonamide-resistant dihydropteroate synthase Sul2 (NF000295.1) (FFFFIJOH_00016)
IMGPR_plasmid_2675903580_000006 53:14954(+)
Tet(A)/Tet(B)/Tet(C) family tetracycline efflux MFS transporter (NF012174.0) (FKFJCDFL_00106)
CmlA/FloR family chloramphenicol efflux MFS transporter (NF033134.0) (FKFJCDFL_00109)
APH(6)-I family aminoglycoside O-phosphotransferase (NF012171.0) (FKFJCDFL_00103)
aminoglycoside O-phosphotransferase APH(3'')-Ib (NF032895.1) (FKFJCDFL_00102)
Tetracycline repressor protein class A from transposon 1721 (FKFJCDFL_00105)
IS6 family transposase IS26 (FKFJCDFL_00115)
sulfonamide-resistant dihydropteroate synthase Sul2 (NF000295.1) (FKFJCDFL_00101)
IMGPR_plasmid_2740892425_000005 82635:97667(-)
Tet(A)/Tet(B)/Tet(C) family tetracycline efflux MFS transporter (NF012174.0) (LDOFBICF_00044)
CmlA/FloR family chloramphenicol efflux MFS transporter (NF033134.0) (LDOFBICF_00047)
APH(6)-I family aminoglycoside O-phosphotransferase (NF012171.0) (LDOFBICF_00041)
aminoglycoside O-phosphotransferase APH(3'')-Ib (NF032895.1) (LDOFBICF_00040)
Tetracycline repressor protein class A from transposon 1721 (LDOFBICF_00043)
HTH-type transcriptional regulator HdfR (LDOFBICF_00046)
sulfonamide-resistant dihydropteroate synthase Sul2 (NF000295.1) (LDOFBICF_00039)
IMGPR_plasmid_2737471840_000003 25434:38388(-)
Tet(A)/Tet(B)/Tet(C) family tetracycline efflux MFS transporter (NF012174.0) (FDEDHELK_00041)
APH(6)-I family aminoglycoside O-phosphotransferase (NF012171.0) (FDEDHELK_00038)
aminoglycoside O-phosphotransferase APH(3'')-Ib (NF032895.1) (FDEDHELK_00037)
Tetracycline repressor protein class A from transposon 1721 (FDEDHELK_00040)
sulfonamide-resistant dihydropteroate synthase Sul2 (NF000295.1) (FDEDHELK_00036)
IMGPR_plasmid_2660238507_000007 22863:33606(-)
Tet(A)/Tet(B)/Tet(C) family tetracycline efflux MFS transporter (NF012174.0) (NHHIKAHC_00004)
APH(6)-I family aminoglycoside O-phosphotransferase (NF012171.0) (NHHIKAHC_00007)
aminoglycoside O-phosphotransferase APH(3'')-Ib (NF032895.1) (NHHIKAHC_00008)
Tetracycline repressor protein class A from transposon 1721 (NHHIKAHC_00005)
sulfonamide-resistant dihydropteroate synthase Sul2 (NF000295.1) (NHHIKAHC_00009)
IMGPR_plasmid_2588253806_000003 580:10725(+)
APH(6)-I family aminoglycoside O-phosphotransferase (NF012171.0) (KMHHPNMH_00004)
aminoglycoside O-phosphotransferase APH(3'')-Ib (NF032895.1) (KMHHPNMH_00005)
Tetracycline repressor protein class A from transposon 1721 (KMHHPNMH_00002)
sulfonamide-resistant dihydropteroate synthase Sul2 (NF000295.1) (KMHHPNMH_00006)
IMGPR_plasmid_2537561642_000001 611:8787(+)
APH(6)-I family aminoglycoside O-phosphotransferase (NF012171.0) (IMEEFEAF_00078)
aminoglycoside O-phosphotransferase APH(3'')-Ib (NF032895.1) (IMEEFEAF_00077)
Tetracycline repressor protein class A from transposon 1721 (IMEEFEAF_00080)
sulfonamide-resistant dihydropteroate synthase Sul2 (NF000295.1) (IMEEFEAF_00076)
IMGPR_plasmid_2737471865_000002 62991:71167(-)
APH(3')-I family aminoglycoside O-phosphotransferase (NF033059.2) (FNNKGLKP_00005)
APH(6)-I family aminoglycoside O-phosphotransferase (NF012171.0) (FNNKGLKP_00004)
Tetracycline repressor protein class A from transposon 1721 (FNNKGLKP_00002)
sulfonamide-resistant dihydropteroate synthase Sul2 (NF000295.1) (FNNKGLKP_00006)
IMGPR_plasmid_2734482777_000008 632:8807(+)
APH(6)-I family aminoglycoside O-phosphotransferase (NF012171.0) (IMODJKAO_00108)
aminoglycoside 3-N-acetyltransferase (NF033082.0) (IMODJKAO_00114)
TEM family class A beta-lactamase (NF000531.2) (IMODJKAO_00109)
Tunicamycin resistance protein (IMODJKAO_00113)
IS1182 family transposase ISCfr1 (IMODJKAO_00110)
sulfonamide-resistant dihydropteroate synthase Sul2 (NF000295.1) (IMODJKAO_00106)
aminoglycoside O-phosphotransferase APH(3'')-Ib (NF032895.1) (IMODJKAO_00107)
IMGPR_plasmid_2852071321_000005 84684:98077(-)
IS21 family transposase ISAs29 (OFFMGGMB_00097)
IS21 family transposase ISPpu7 (OFFMGGMB_00096)
Modification methylase PvuII (OFFMGGMB_00095)
IMGPR_plasmid_2545824781_000010 77634:89953(-)
IS21 family transposase ISAs29 (FMIGGLHN_00012)
IS21 family transposase ISPpu7 (FMIGGLHN_00013)
Modification methylase PvuII (FMIGGLHN_00014)
IMGPR_plasmid_2563366764_000008 208:12527(+)
IS21 family transposase ISAs29 (NBOJJEHD_00099)
IS5 family transposase ISEc61 (NBOJJEHD_00111)
IS21 family transposase ISPpu7 (NBOJJEHD_00098)
Modification methylase PvuII (NBOJJEHD_00097)
IMGPR_plasmid_2636416136_000014 80198:93577(-)
IS21 family transposase ISAs29 (ANMAOBBK_00104)
IS21 family transposase ISPpu7 (ANMAOBBK_00103)
Modification methylase PvuII (ANMAOBBK_00102)
IMGPR_plasmid_2906172311_000010 91337:103656(-)
IS21 family transposase ISAs29 (ADNPHPGE_00014)
IS21 family transposase ISPpu7 (ADNPHPGE_00015)
Modification methylase PvuII (ADNPHPGE_00016)
IMGPR_plasmid_2563367059_000014 9:14540(+)
IMGPR_plasmid_2700989582_000014 28216:35165(-)
Modification methylase PvuII (BCBHLMPN_00106)
BCBHLMPN_00107
BCBHLMPN_00105
IMGPR_plasmid_2903362858_000007 91293:96596(-)
7000 nt
AMR