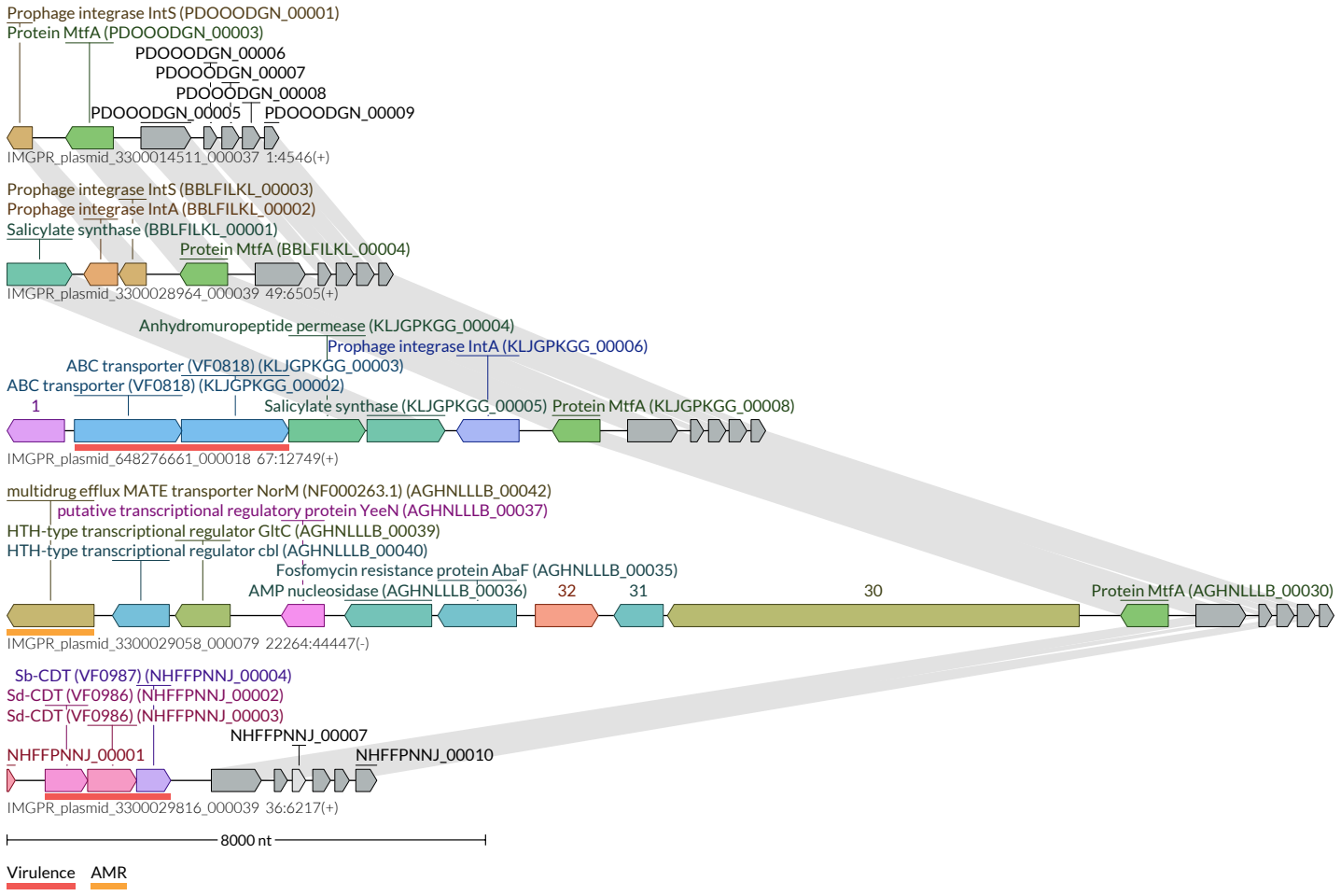
Prophage integrase IntS (PDOOODGN_00001)
Protein MtfA (PDOOODGN_00003)
PDOOODGN_00006
PDOOODGN_00007
PDOOODGN_00008
PDOOODGN_00005
PDOOODGN_00009
IMGPR_plasmid_3300014511_000037 1:4546(+)
Prophage integrase IntS (BBLFILKL_00003)
Prophage integrase IntA (BBLFILKL_00002)
Salicylate synthase (BBLFILKL_00001)
Protein MtfA (BBLFILKL_00004)
IMGPR_plasmid_3300028964_000039 49:6505(+)
Anhydromuropeptide permease (KLJGPKGG_00004)
Prophage integrase IntA (KLJGPKGG_00006)
ABC transporter (VF0818) (KLJGPKGG_00003)
ABC transporter (VF0818) (KLJGPKGG_00002)
1
Salicylate synthase (KLJGPKGG_00005)
Protein MtfA (KLJGPKGG_00008)
IMGPR_plasmid_648276661_000018 67:12749(+)
multidrug efflux MATE transporter NorM (NF000263.1) (AGHNLLLB_00042)
putative transcriptional regulatory protein YeeN (AGHNLLLB_00037)
HTH-type transcriptional regulator GltC (AGHNLLLB_00039)
HTH-type transcriptional regulator cbl (AGHNLLLB_00040)
Fosfomycin resistance protein AbaF (AGHNLLLB_00035)
AMP nucleosidase (AGHNLLLB_00036)
32
31
30
Protein MtfA (AGHNLLLB_00030)
IMGPR_plasmid_3300029058_000079 22264:44447(-)
Sb-CDT (VF0987) (NHFFPNNJ_00004)
Sd-CDT (VF0986) (NHFFPNNJ_00002)
Sd-CDT (VF0986) (NHFFPNNJ_00003)
NHFFPNNJ_00007
NHFFPNNJ_00001
NHFFPNNJ_00010
IMGPR_plasmid_3300029816_000039 36:6217(+)
8000 nt
Virulence
AMR