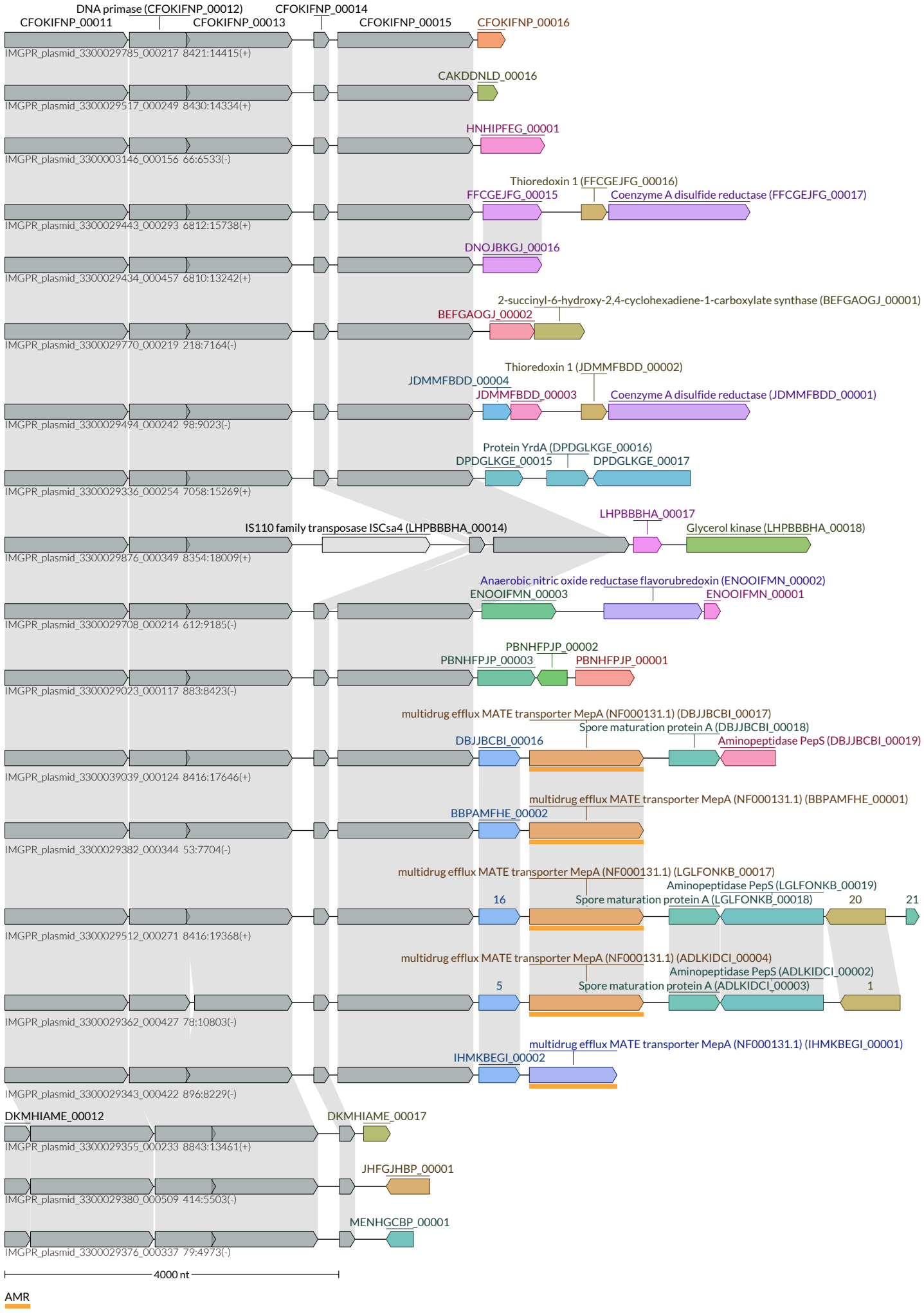
DNA primase (CFOKIFNP_00012)
CFOKIFNP_00014
CFOKIFNP_00011
CFOKIFNP_00013
CFOKIFNP_00015
CFOKIFNP_00016
IMGPR_plasmid_3300029785_000217 8421:14415(+)
CAKDDNLD_00016
IMGPR_plasmid_3300029517_000249 8430:14334(+)
HNHIPFEG_00001
IMGPR_plasmid_3300003146_000156 66:6533(-)
Thioredoxin 1 (FFCGEJFG_00016)
FFCGEJFG_00015
Coenzyme A disulfide reductase (FFCGEJFG_00017)
IMGPR_plasmid_3300029443_000293 6812:15738(+)
DNOJBKGJ_00016
IMGPR_plasmid_3300029434_000457 6810:13242(+)
2-succinyl-6-hydroxy-2,4-cyclohexadiene-1-carboxylate synthase (BEFGAOGJ_00001)
BEFGAOGJ_00002
IMGPR_plasmid_3300029770_000219 218:7164(-)
Thioredoxin 1 (JDMMFBDD_00002)
JDMMFBDD_00004
JDMMFBDD_00003
Coenzyme A disulfide reductase (JDMMFBDD_00001)
IMGPR_plasmid_3300029494_000242 98:9023(-)
Protein YrdA (DPDGLKGE_00016)
DPDGLKGE_00015
DPDGLKGE_00017
IMGPR_plasmid_3300029336_000254 7058:15269(+)
LHPBBBHA_00017
IS110 family transposase ISCsa4 (LHPBBBHA_00014)
Glycerol kinase (LHPBBBHA_00018)
IMGPR_plasmid_3300029876_000349 8354:18009(+)
Anaerobic nitric oxide reductase flavorubredoxin (ENOOIFMN_00002)
ENOOIFMN_00003
ENOOIFMN_00001
IMGPR_plasmid_3300029708_000214 612:9185(-)
PBNHFPJP_00002
PBNHFPJP_00003
PBNHFPJP_00001
IMGPR_plasmid_3300029023_000117 883:8423(-)
multidrug efflux MATE transporter MepA (NF000131.1) (DBJJBCBI_00017)
Spore maturation protein A (DBJJBCBI_00018)
DBJJBCBI_00016
Aminopeptidase PepS (DBJJBCBI_00019)
IMGPR_plasmid_3300039039_000124 8416:17646(+)
multidrug efflux MATE transporter MepA (NF000131.1) (BBPAMFHE_00001)
BBPAMFHE_00002
IMGPR_plasmid_3300029382_000344 53:7704(-)
multidrug efflux MATE transporter MepA (NF000131.1) (LGLFONKB_00017)
Aminopeptidase PepS (LGLFONKB_00019)
16
Spore maturation protein A (LGLFONKB_00018)
20
21
IMGPR_plasmid_3300029512_000271 8416:19368(+)
multidrug efflux MATE transporter MepA (NF000131.1) (ADLKIDCI_00004)
Aminopeptidase PepS (ADLKIDCI_00002)
5
Spore maturation protein A (ADLKIDCI_00003)
1
IMGPR_plasmid_3300029362_000427 78:10803(-)
multidrug efflux MATE transporter MepA (NF000131.1) (IHMKBEGI_00001)
IHMKBEGI_00002
IMGPR_plasmid_3300029343_000422 896:8229(-)
DKMHIAME_00012
DKMHIAME_00017
IMGPR_plasmid_3300029355_000233 8843:13461(+)
JHFGJHBP_00001
IMGPR_plasmid_3300029380_000509 414:5503(-)
MENHGCBP_00001
IMGPR_plasmid_3300029376_000337 79:4973(-)
4000 nt
AMR