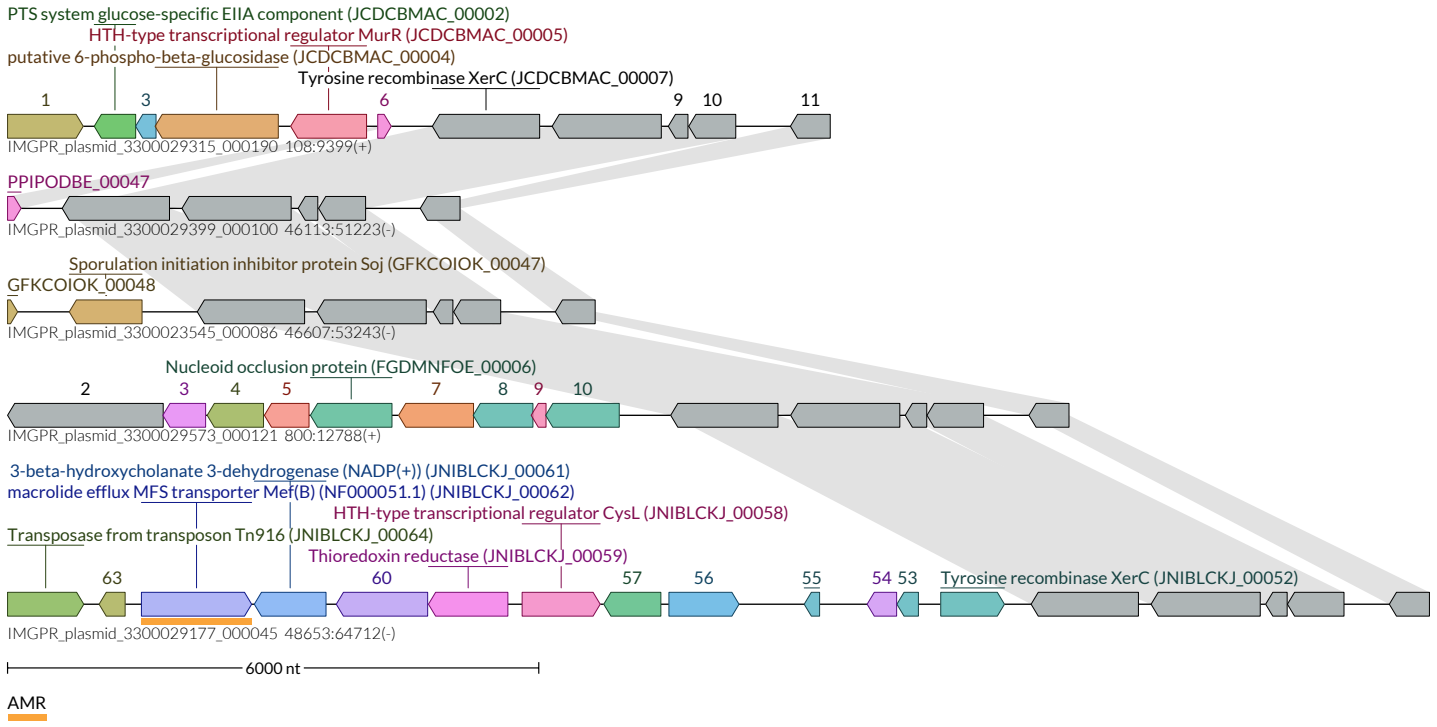
PTS system glucose-specific EIIA component (JCDCBMAC_00002)
HTH-type transcriptional regulator MurR (JCDCBMAC_00005)
putative 6-phospho-beta-glucosidase (JCDCBMAC_00004)
Tyrosine recombinase XerC (JCDCBMAC_00007)
1
3
6
9
10
11
IMGPR_plasmid_3300029315_000190 108:9399(+)
PPIPODBE_00047
IMGPR_plasmid_3300029399_000100 46113:51223(-)
Sporulation initiation inhibitor protein Soj (GFKCOIOK_00047)
GFKCOIOK_00048
IMGPR_plasmid_3300023545_000086 46607:53243(-)
Nucleoid occlusion protein (FGDMNFOE_00006)
2
3
4
5
7
8
9
10
IMGPR_plasmid_3300029573_000121 800:12788(+)
3-beta-hydroxycholanate 3-dehydrogenase (NADP(+)) (JNIBLCKJ_00061)
macrolide efflux MFS transporter Mef(B) (NF000051.1) (JNIBLCKJ_00062)
HTH-type transcriptional regulator CysL (JNIBLCKJ_00058)
Transposase from transposon Tn916 (JNIBLCKJ_00064)
Thioredoxin reductase (JNIBLCKJ_00059)
63
60
57
56
55
54
53
Tyrosine recombinase XerC (JNIBLCKJ_00052)
IMGPR_plasmid_3300029177_000045 48653:64712(-)
6000 nt
AMR