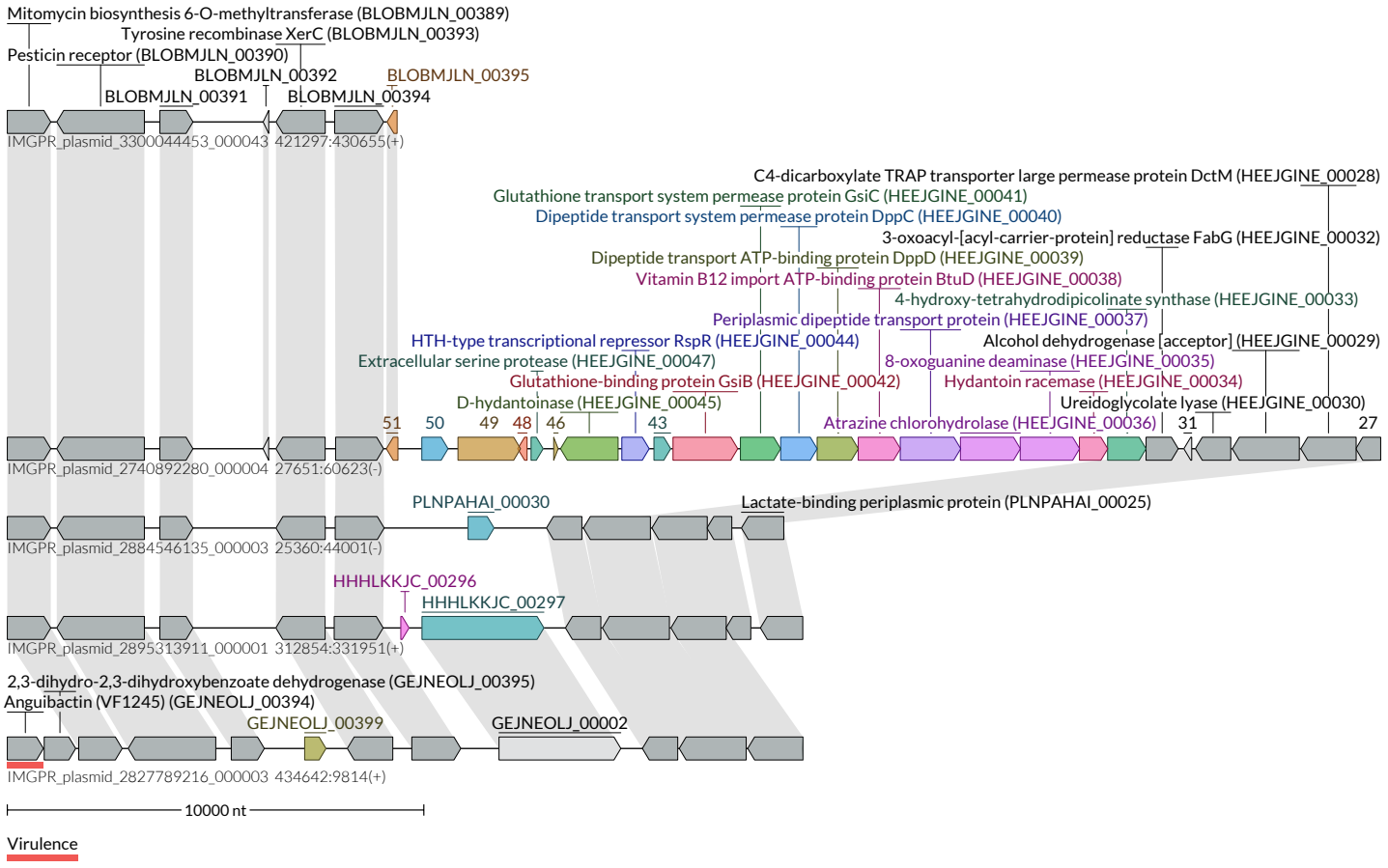
Mitomycin biosynthesis 6-O-methyltransferase (BLOBMJLN_00389)
Tyrosine recombinase XerC (BLOBMJLN_00393)
Pesticin receptor (BLOBMJLN_00390)
BLOBMJLN_00392
BLOBMJLN_00395
BLOBMJLN_00391
BLOBMJLN_00394
IMGPR_plasmid_3300044453_000043 421297:430655(+)
C4-dicarboxylate TRAP transporter large permease protein DctM (HEEJGINE_00028)
Glutathione transport system permease protein GsiC (HEEJGINE_00041)
Dipeptide transport system permease protein DppC (HEEJGINE_00040)
3-oxoacyl-[acyl-carrier-protein] reductase FabG (HEEJGINE_00032)
Dipeptide transport ATP-binding protein DppD (HEEJGINE_00039)
Vitamin B12 import ATP-binding protein BtuD (HEEJGINE_00038)
4-hydroxy-tetrahydrodipicolinate synthase (HEEJGINE_00033)
Periplasmic dipeptide transport protein (HEEJGINE_00037)
HTH-type transcriptional repressor RspR (HEEJGINE_00044)
Alcohol dehydrogenase [acceptor] (HEEJGINE_00029)
Extracellular serine protease (HEEJGINE_00047)
8-oxoguanine deaminase (HEEJGINE_00035)
Glutathione-binding protein GsiB (HEEJGINE_00042)
Hydantoin racemase (HEEJGINE_00034)
D-hydantoinase (HEEJGINE_00045)
Ureidoglycolate lyase (HEEJGINE_00030)
51
50
49
48
46
43
Atrazine chlorohydrolase (HEEJGINE_00036)
31
27
IMGPR_plasmid_2740892280_000004 27651:60623(-)
PLNPAHAI_00030
Lactate-binding periplasmic protein (PLNPAHAI_00025)
IMGPR_plasmid_2884546135_000003 25360:44001(-)
HHHLKKJC_00296
HHHLKKJC_00297
IMGPR_plasmid_2895313911_000001 312854:331951(+)
2,3-dihydro-2,3-dihydroxybenzoate dehydrogenase (GEJNEOLJ_00395)
Anguibactin (VF1245) (GEJNEOLJ_00394)
GEJNEOLJ_00399
GEJNEOLJ_00002
IMGPR_plasmid_2827789216_000003 434642:9814(+)
10000 nt
Virulence