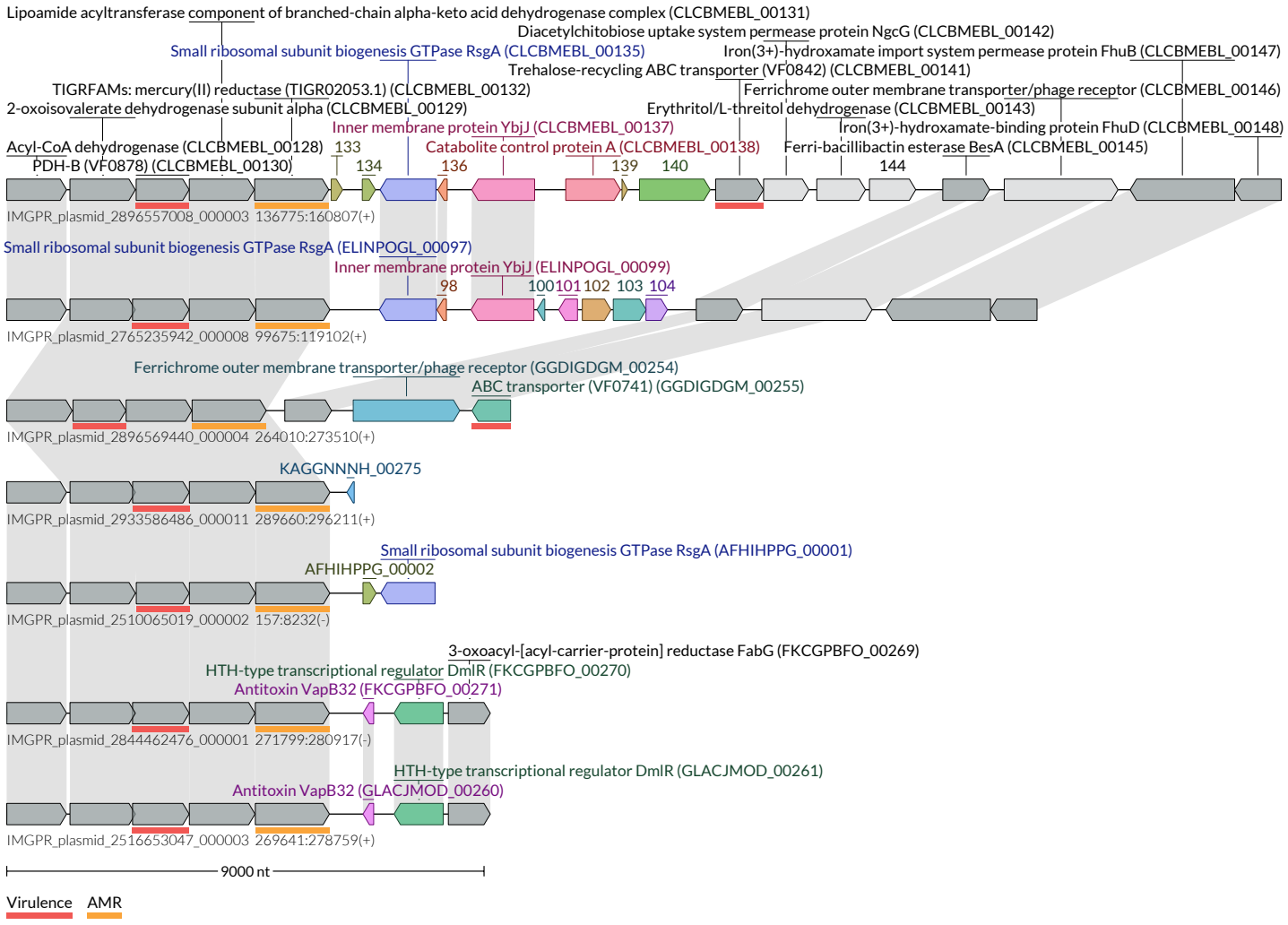
Lipoamide acyltransferase component of branched-chain alpha-keto acid dehydrogenase complex (CLCBMEBL_00131)
Diacetylchitobiose uptake system permease protein NgcG (CLCBMEBL_00142)
Small ribosomal subunit biogenesis GTPase RsgA (CLCBMEBL_00135)
Iron(3+)-hydroxamate import system permease protein FhuB (CLCBMEBL_00147)
Trehalose-recycling ABC transporter (VF0842) (CLCBMEBL_00141)
TIGRFAMs: mercury(II) reductase (TIGR02053.1) (CLCBMEBL_00132)
Ferrichrome outer membrane transporter/phage receptor (CLCBMEBL_00146)
2-oxoisovalerate dehydrogenase subunit alpha (CLCBMEBL_00129)
Erythritol/L-threitol dehydrogenase (CLCBMEBL_00143)
Inner membrane protein YbjJ (CLCBMEBL_00137)
Iron(3+)-hydroxamate-binding protein FhuD (CLCBMEBL_00148)
Acyl-CoA dehydrogenase (CLCBMEBL_00128)
133
Catabolite control protein A (CLCBMEBL_00138)
Ferri-bacillibactin esterase BesA (CLCBMEBL_00145)
PDH-B (VF0878) (CLCBMEBL_00130)
134
136
139
140
144
IMGPR_plasmid_2896557008_000003 136775:160807(+)
Small ribosomal subunit biogenesis GTPase RsgA (ELINPOGL_00097)
Inner membrane protein YbjJ (ELINPOGL_00099)
98
100
101
102
103
104
IMGPR_plasmid_2765235942_000008 99675:119102(+)
Ferrichrome outer membrane transporter/phage receptor (GGDIGDGM_00254)
ABC transporter (VF0741) (GGDIGDGM_00255)
IMGPR_plasmid_2896569440_000004 264010:273510(+)
KAGGNNNH_00275
IMGPR_plasmid_2933586486_000011 289660:296211(+)
Small ribosomal subunit biogenesis GTPase RsgA (AFHIHPPG_00001)
AFHIHPPG_00002
IMGPR_plasmid_2510065019_000002 157:8232(-)
3-oxoacyl-[acyl-carrier-protein] reductase FabG (FKCGPBFO_00269)
HTH-type transcriptional regulator DmlR (FKCGPBFO_00270)
Antitoxin VapB32 (FKCGPBFO_00271)
IMGPR_plasmid_2844462476_000001 271799:280917(-)
HTH-type transcriptional regulator DmlR (GLACJMOD_00261)
Antitoxin VapB32 (GLACJMOD_00260)
IMGPR_plasmid_2516653047_000003 269641:278759(+)
9000 nt
Virulence
AMR