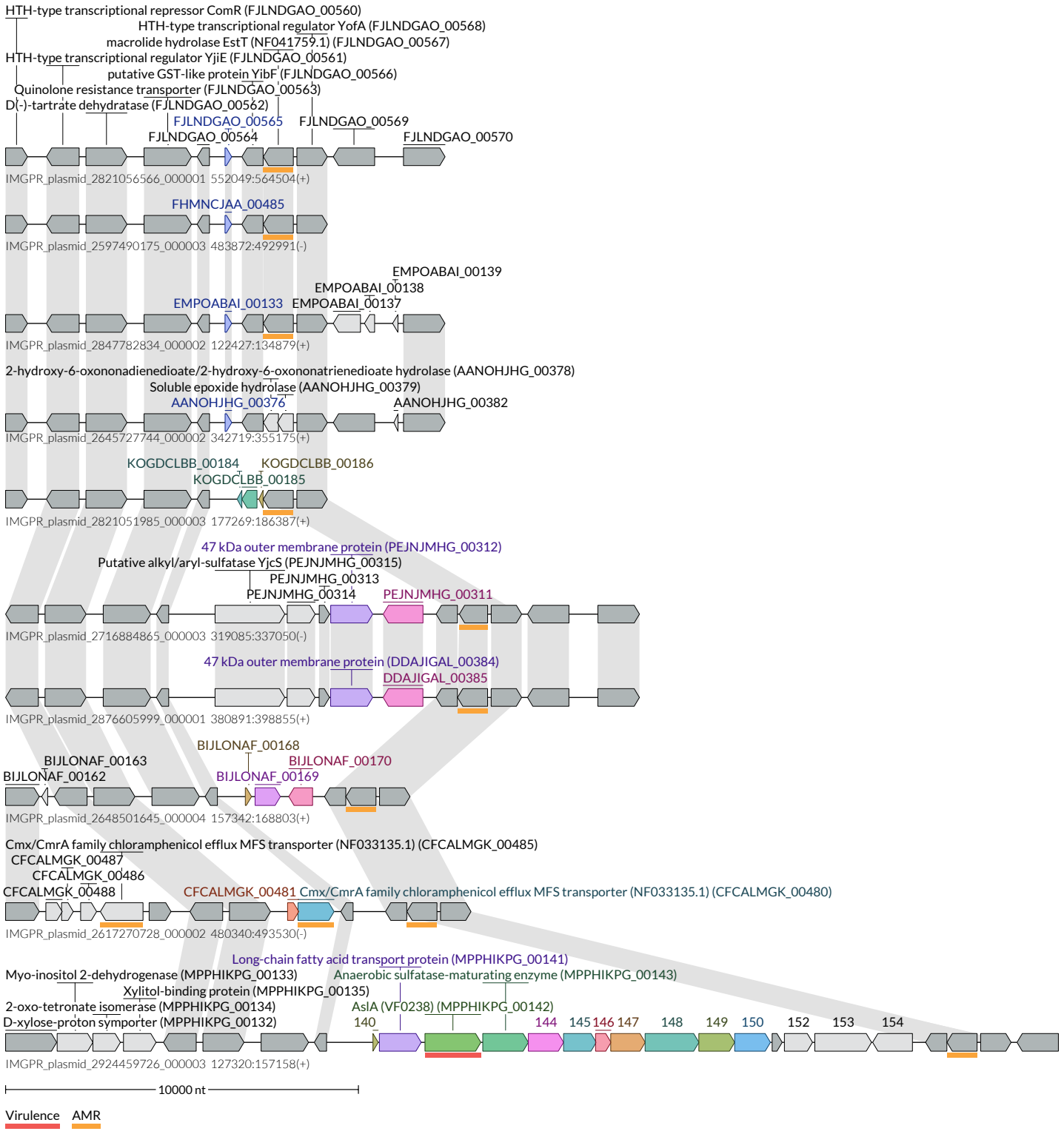
HTH-type transcriptional repressor ComR (FJLNDGAO_00560)
HTH-type transcriptional regulator YofA (FJLNDGAO_00568)
macrolide hydrolase EstT (NF041759.1) (FJLNDGAO_00567)
HTH-type transcriptional regulator YjiE (FJLNDGAO_00561)
putative GST-like protein YibF (FJLNDGAO_00566)
Quinolone resistance transporter (FJLNDGAO_00563)
D(-)-tartrate dehydratase (FJLNDGAO_00562)
FJLNDGAO_00565
FJLNDGAO_00569
FJLNDGAO_00564
FJLNDGAO_00570
IMGPR_plasmid_2821056566_000001 552049:564504(+)
FHMNCJAA_00485
IMGPR_plasmid_2597490175_000003 483872:492991(-)
EMPOABAI_00139
EMPOABAI_00138
EMPOABAI_00133
EMPOABAI_00137
IMGPR_plasmid_2847782834_000002 122427:134879(+)
2-hydroxy-6-oxononadienedioate/2-hydroxy-6-oxononatrienedioate hydrolase (AANOHJHG_00378)
Soluble epoxide hydrolase (AANOHJHG_00379)
AANOHJHG_00376
AANOHJHG_00382
IMGPR_plasmid_2645727744_000002 342719:355175(+)
KOGDCLBB_00184
KOGDCLBB_00186
KOGDCLBB_00185
IMGPR_plasmid_2821051985_000003 177269:186387(+)
47 kDa outer membrane protein (PEJNJMHG_00312)
Putative alkyl/aryl-sulfatase YjcS (PEJNJMHG_00315)
PEJNJMHG_00313
PEJNJMHG_00314
PEJNJMHG_00311
IMGPR_plasmid_2716884865_000003 319085:337050(-)
47 kDa outer membrane protein (DDAJIGAL_00384)
DDAJIGAL_00385
IMGPR_plasmid_2876605999_000001 380891:398855(+)
BIJLONAF_00168
BIJLONAF_00163
BIJLONAF_00170
BIJLONAF_00162
BIJLONAF_00169
IMGPR_plasmid_2648501645_000004 157342:168803(+)
Cmx/CmrA family chloramphenicol efflux MFS transporter (NF033135.1) (CFCALMGK_00485)
CFCALMGK_00487
CFCALMGK_00486
CFCALMGK_00488
CFCALMGK_00481
Cmx/CmrA family chloramphenicol efflux MFS transporter (NF033135.1) (CFCALMGK_00480)
IMGPR_plasmid_2617270728_000002 480340:493530(-)
Long-chain fatty acid transport protein (MPPHIKPG_00141)
Myo-inositol 2-dehydrogenase (MPPHIKPG_00133)
Anaerobic sulfatase-maturating enzyme (MPPHIKPG_00143)
Xylitol-binding protein (MPPHIKPG_00135)
2-oxo-tetronate isomerase (MPPHIKPG_00134)
AslA (VF0238) (MPPHIKPG_00142)
D-xylose-proton symporter (MPPHIKPG_00132)
140
144
145
146
147
148
149
150
152
153
154
IMGPR_plasmid_2924459726_000003 127320:157158(+)
10000 nt
Virulence
AMR