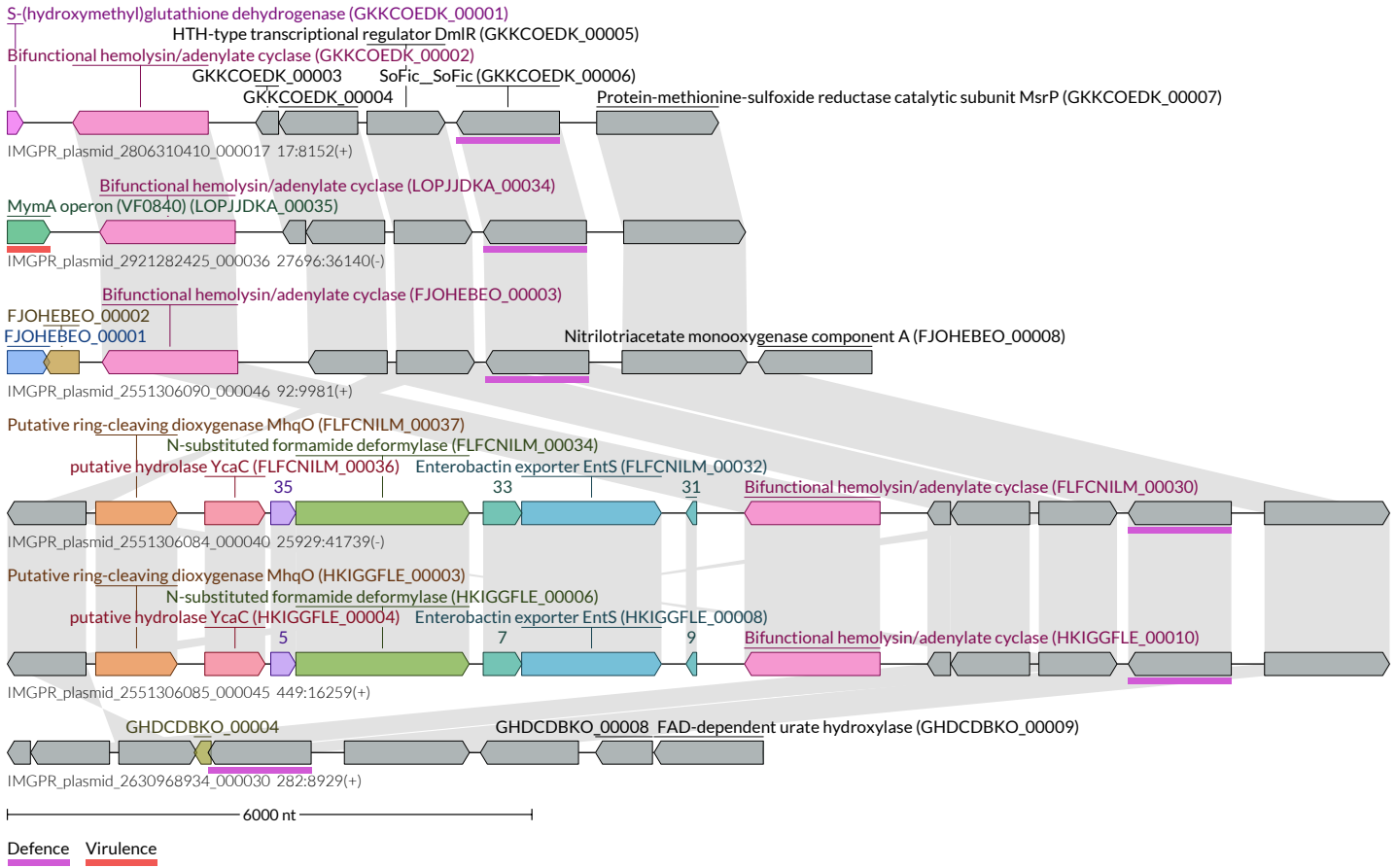
S-(hydroxymethyl)glutathione dehydrogenase (GKKCOEDK_00001)
HTH-type transcriptional regulator DmlR (GKKCOEDK_00005)
Bifunctional hemolysin/adenylate cyclase (GKKCOEDK_00002)
GKKCOEDK_00003
SoFic_SoFic (GKKCOEDK_00006)
GKKCOEDK_00004
Protein-methionine-sulfoxide reductase catalytic subunit MsrP (GKKCOEDK_00007)
IMGPR_plasmid_2806310410_000017 17:8152(+)
Bifunctional hemolysin/adenylate cyclase (LOPJJDKA_00034)
MymA operon (VF0840) (LOPJJDKA_00035)
IMGPR_plasmid_2921282425_000036 27696:36140(-)
Bifunctional hemolysin/adenylate cyclase (FJOHEBEO_00003)
FJOHEBEO_00002
FJOHEBEO_00001
Nitrilotriacetate monooxygenase component A (FJOHEBEO_00008)
IMGPR_plasmid_2551306090_000046 92:9981(+)
Putative ring-cleaving dioxygenase MhqO (FLFCNILM_00037)
N-substituted formamide deformylase (FLFCNILM_00034)
putative hydrolase YcaC (FLFCNILM_00036)
Enterobactin exporter EntS (FLFCNILM_00032)
35
33
31
Bifunctional hemolysin/adenylate cyclase (FLFCNILM_00030)
IMGPR_plasmid_2551306084_000040 25929:41739(-)
Putative ring-cleaving dioxygenase MhqO (HKIGGFLE_00003)
N-substituted formamide deformylase (HKIGGFLE_00006)
putative hydrolase YcaC (HKIGGFLE_00004)
Enterobactin exporter EntS (HKIGGFLE_00008)
5
7
9
Bifunctional hemolysin/adenylate cyclase (HKIGGFLE_00010)
IMGPR_plasmid_2551306085_000045 449:16259(+)
GHDCDBKO_00004
GHDCDBKO_00008
FAD-dependent urate hydroxylase (GHDCDBKO_00009)
IMGPR_plasmid_2630968934_000030 282:8929(+)
6000 nt
Defence
Virulence