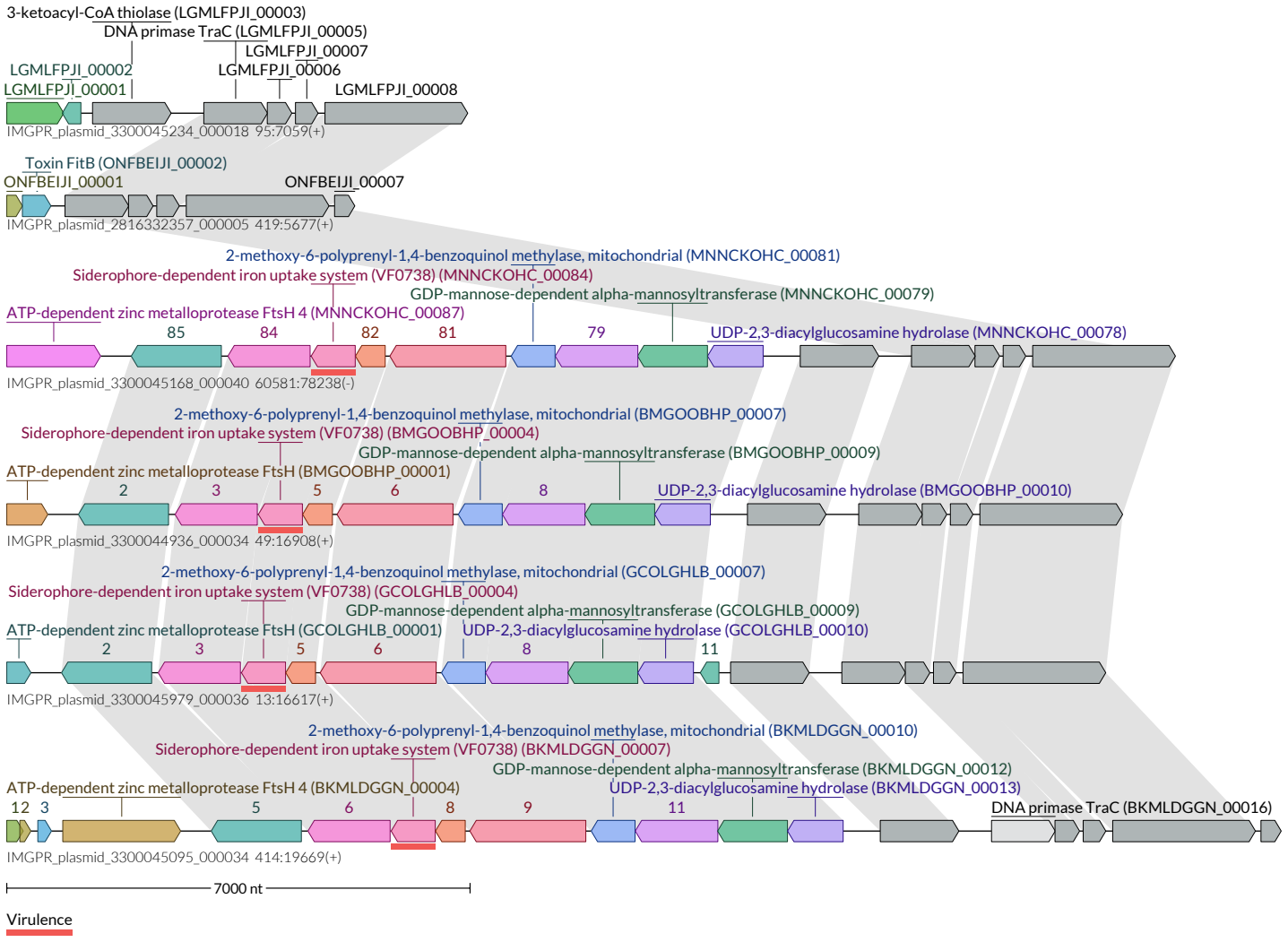
3-ketoacyl-CoA thiolase (LGMLFPJI_00003)
DNA primase TraC (LGMLFPJI_00005)
LGMLFPJI_00007
LGMLFPJI_00002
LGMLFPJI_00006
LGMLFPJI_00001
LGMLFPJI_00008
IMGPR_plasmid_3300045234_000018 95:7059(+)
Toxin FitB (ONFBEIJI_00002)
ONFBEIJI_00001
ONFBEIJI_00007
IMGPR_plasmid_2816332357_000005 419:5677(+)
2-methoxy-6-polyprenyl-1,4-benzoquinol methylase, mitochondrial (MNNCKOHC_00081)
Siderophore-dependent iron uptake system (VF0738) (MNNCKOHC_00084)
GDP-mannose-dependent alpha-mannosyltransferase (MNNCKOHC_00079)
ATP-dependent zinc metalloprotease FtsH 4 (MNNCKOHC_00087)
85
84
82
81
79
UDP-2,3-diacylglucosamine hydrolase (MNNCKOHC_00078)
IMGPR_plasmid_3300045168_000040 60581:78238(-)
2-methoxy-6-polyprenyl-1,4-benzoquinol methylase, mitochondrial (BMGOOBHP_00007)
Siderophore-dependent iron uptake system (VF0738) (BMGOOBHP_00004)
GDP-mannose-dependent alpha-mannosyltransferase (BMGOOBHP_00009)
ATP-dependent zinc metalloprotease FtsH (BMGOOBHP_00001)
2
3
5
6
8
UDP-2,3-diacylglucosamine hydrolase (BMGOOBHP_00010)
IMGPR_plasmid_3300044936_000034 49:16908(+)
2-methoxy-6-polyprenyl-1,4-benzoquinol methylase, mitochondrial (GCOLGHLB_00007)
Siderophore-dependent iron uptake system (VF0738) (GCOLGHLB_00004)
GDP-mannose-dependent alpha-mannosyltransferase (GCOLGHLB_00009)
ATP-dependent zinc metalloprotease FtsH (GCOLGHLB_00001)
UDP-2,3-diacylglucosamine hydrolase (GCOLGHLB_00010)
2
3
5
6
8
11
IMGPR_plasmid_3300045979_000036 13:16617(+)
2-methoxy-6-polyprenyl-1,4-benzoquinol methylase, mitochondrial (BKMLDGGN_00010)
Siderophore-dependent iron uptake system (VF0738) (BKMLDGGN_00007)
GDP-mannose-dependent alpha-mannosyltransferase (BKMLDGGN_00012)
ATP-dependent zinc metalloprotease FtsH 4 (BKMLDGGN_00004)
UDP-2,3-diacylglucosamine hydrolase (BKMLDGGN_00013)
12
3
5
6
8
9
11
DNA primase TraC (BKMLDGGN_00016)
IMGPR_plasmid_3300045095_000034 414:19669(+)
7000 nt
Virulence