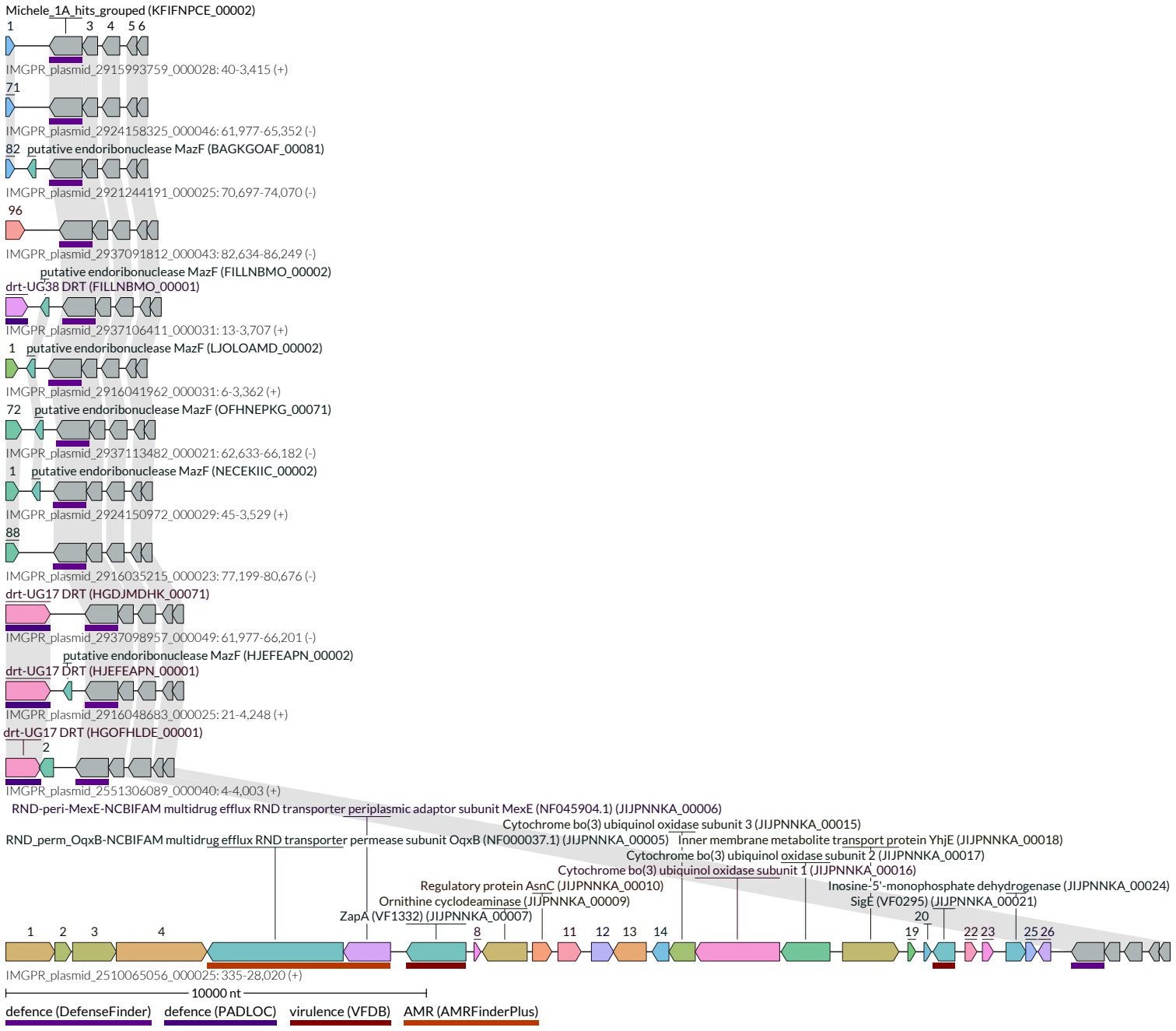
Michele_1A_hits_grouped (KFIFNPCE_00002)
1
3
4
5 6
IMGPR_plasmid_2915993759_000028: 40-3,415 (+)
71
IMGPR_plasmid_2924158325_000046: 61,977-65,352 (-)
82
putative endoribonuclease MazF (BAGKGOAF_00081)
IMGPR_plasmid_2921244191_000025: 70,697-74,070 (-)
96
IMGPR_plasmid_2937091812_000043: 82,634-86,249 (-)
putative endoribonuclease MazF (FILLNBMO_00002)
drt-UG38 DRT (FILLNBMO_00001)
IMGPR_plasmid_2937106411_000031: 13-3,707 (+)
1
putative endoribonuclease MazF (LJOLOAMD_00002)
IMGPR_plasmid_2916041962_000031: 6-3,362 (+)
72
putative endoribonuclease MazF (OFHNEPKG_00071)
IMGPR_plasmid_2937113482_000021: 62,633-66,182 (-)
1
putative endoribonuclease MazF (NECEKIIC_00002)
IMGPR_plasmid_2924150972_000029: 45-3,529 (+)
88
IMGPR_plasmid_2916035215_000023: 77,199-80,676 (-)
drt-UG17 DRT (HGDJMDHK_00071)
IMGPR_plasmid_2937098957_000049: 61,977-66,201 (-)
putative endoribonuclease MazF (HJEFEAPN_00002)
drt-UG17 DRT (HJEFEAPN_00001)
IMGPR_plasmid_2916048683_000025: 21-4,248 (+)
drt-UG17 DRT (HGOFHLDE_00001)
2
IMGPR_plasmid_2551306089_000040: 4-4,003 (+)
RND-peri-MexE-NCBIFAM multidrug efflux RND transporter periplasmic adaptor subunit MexE (NF045904.1) (JIJPNNKA_00006)
Cytochrome bo(3) ubiquinol oxidase subunit 3 (JIJPNNKA_00015)
RND_perm_OqxB-NCBIFAM multidrug efflux RND transporter permease subunit OqxB (NF000037.1) (JIJPNNKA_00005)
Inner membrane metabolite transport protein YhjE (JIJPNNKA_00018)
Cytochrome bo(3) ubiquinol oxidase subunit 2 (JIJPNNKA_00017)
Cytochrome bo(3) ubiquinol oxidase subunit 1 (JIJPNNKA_00016)
Regulatory protein AsnC (JIJPNNKA_00010)
Inosine-5'-monophosphate dehydrogenase (JIJPNNKA_00024)
Ornithine cyclodeaminase (JIJPNNKA_00009)
SigE (VF0295) (JIJPNNKA_00021)
ZapA (VF1332) (JIJPNNKA_00007)
1
2
3
4
8
11
12
13
14
20
19
22 23
25 26
IMGPR_plasmid_2510065056_000025: 335-28,020 (+)
10000 nt
defence (DefenseFinder)
defence (PADLOC)
virulence (VFDB)
AMR (AMRFinderPlus)