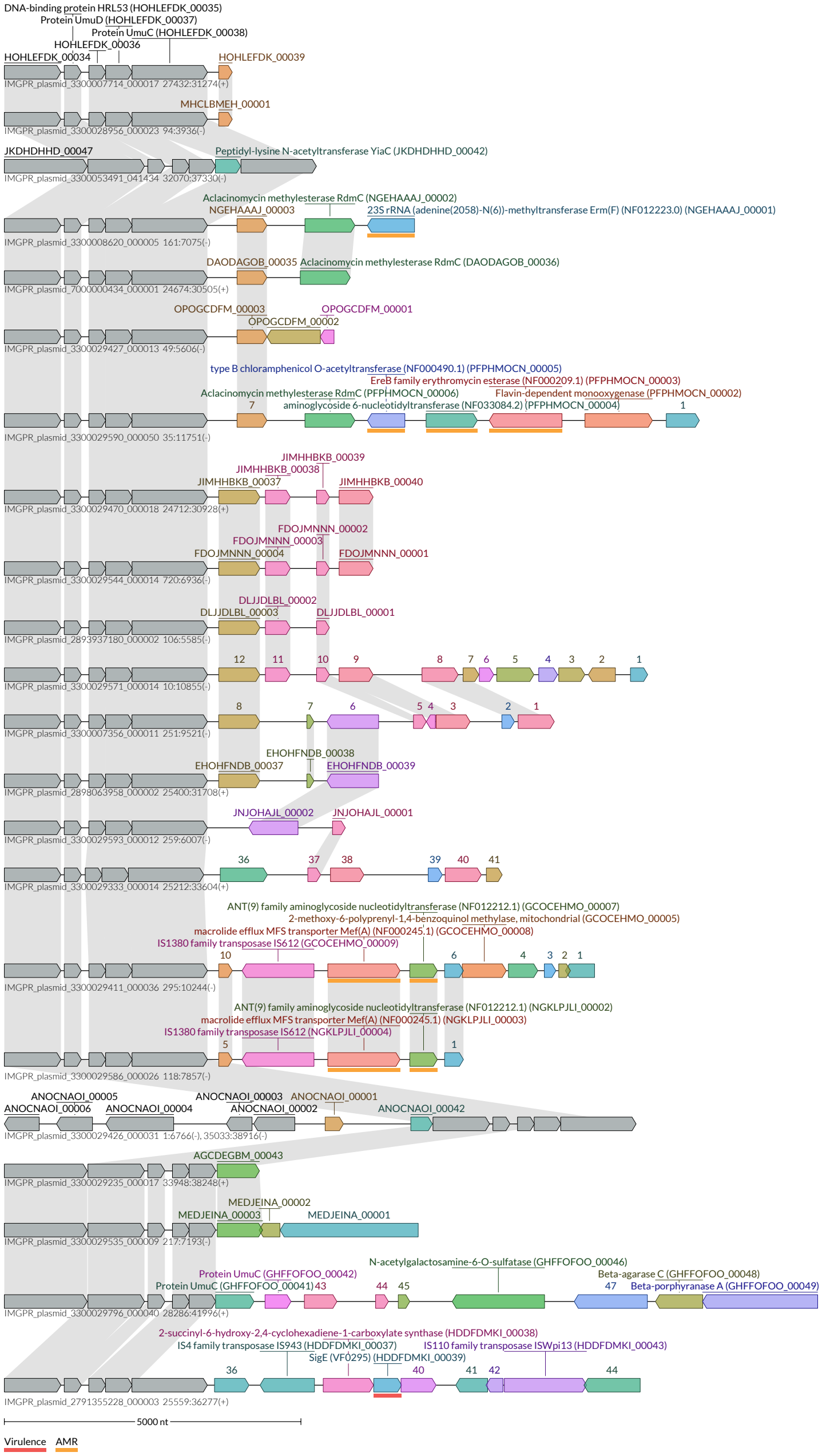
DNA-binding protein HRL53 (HOHLEFDK_00035)
Protein UmuD (HOHLEFDK_00037)
Protein UmuC (HOHLEFDK_00038)
HOHLEFDK_00036
HOHLEFDK_00034
HOHLEFDK_00039
IMGPR_plasmid_3300007714_000017 27432:31274(+)
MHCLBMEH_00001
IMGPR_plasmid_3300028956_000023 94:3936(-)
JKDHDHHD_00047
Peptidyl-lysine N-acetyltransferase YiaC (JKDHDHHD_00042)
IMGPR_plasmid_3300053491_041434 32070:37330(-)
Aclacinomycin methylesterase RdmC (NGEHAAAJ_00002)
NGEHAAAJ_00003
23S rRNA (adenine(2058)-N(6))-methyltransferase Erm(F) (NF012223.0) (NGEHAAAJ_00001)
IMGPR_plasmid_3300008620_000005 161:7075(-)
DAODAGOB_00035
Aclacinomycin methylesterase RdmC (DAODAGOB_00036)
IMGPR_plasmid_7000000434_000001 24674:30505(+)
OPOGCDFM_00003
OPOGCDFM_00001
OPOGCDFM_00002
IMGPR_plasmid_3300029427_000013 49:5606(-)
type B chloramphenicol O-acetyltransferase (NF000490.1) (PFPHMOCN_00005)
EreB family erythromycin esterase (NF000209.1) (PFPHMOCN_00003)
Aclacinomycin methylesterase RdmC (PFPHMOCN_00006)
Flavin-dependent monooxygenase (PFPHMOCN_00002)
7
aminoglycoside 6-nucleotidyltransferase (NF033084.2) (PFPHMOCN_00004)
1
IMGPR_plasmid_3300029590_000050 35:11751(-)
JIMHHBKB_00039
JIMHHBKB_00038
JIMHHBKB_00037
JIMHHBKB_00040
IMGPR_plasmid_3300029470_000018 24712:30928(+)
FDOJMNNN_00002
FDOJMNNN_00003
FDOJMNNN_00004
FDOJMNNN_00001
IMGPR_plasmid_3300029544_000014 720:6936(-)
DLJJDLBL_00002
DLJJDLBL_00003
DLJJDLBL_00001
IMGPR_plasmid_2893937180_000002 106:5585(-)
12 11 10 9 8 7 6 5 4 3 2 1
IMGPR_plasmid_3300029571_000014 10:10855(-)
8 7 6 5 4 3 2 1
IMGPR_plasmid_3300007356_000011 251:9521(-)
EHOHFNDB_00038
EHOHFNDB_00037
EHOHFNDB_00039
IMGPR_plasmid_2898063958_000002 25400:31708(+)
JNJOHAJL_00002
JNJOHAJL_00001
IMGPR_plasmid_3300029593_000012 259:6007(-)
36 37 38 39 40 41
IMGPR_plasmid_3300029333_000014 25212:33604(+)
ANT(9) family aminoglycoside nucleotidyltransferase (NF012212.1) (GCOCEHMO_00007)
2-methoxy-6-polyprenyl-1,4-benzoquinol methylase, mitochondrial (GCOCEHMO_00005)
macrolide efflux MFS transporter Mef(A) (NF000245.1) (GCOCEHMO_00008)
IS1380 family transposase IS612 (GCOCEHMO_00009)
10 6 4 3 2 1
IMGPR_plasmid_3300029411_000036 295:10244(-)
ANT(9) family aminoglycoside nucleotidyltransferase (NF012212.1) (NGKLPJLI_00002)
macrolide efflux MFS transporter Mef(A) (NF000245.1) (NGKLPJLI_00003)
IS1380 family transposase IS612 (NGKLPJLI_00004)
5 1
IMGPR_plasmid_3300029586_000026 118:7857(-)
ANOCNAOI_00005
ANOCNAOI_00006
ANOCNAOI_00004
ANOCNAOI_00003
ANOCNAOI_00001
ANOCNAOI_00002
ANOCNAOI_00042
IMGPR_plasmid_3300029426_000031 1:6766(-), 35033:38916(-)
AGCDEGBM_00043
IMGPR_plasmid_3300029235_000017 33948:38248(+)
MEDJEINA_00002
MEDJEINA_00003
MEDJEINA_00001
IMGPR_plasmid_3300029535_000009 217:7193(-)
N-acetylgalactosamine-6-O-sulfatase (GHFFOFOO_00046)
Protein UmuC (GHFFOFOO_00042)
Beta-agarase C (GHFFOFOO_00048)
Protein UmuC (GHFFOFOO_00041) 43
44 45
47
Beta-porphyranase A (GHFFOFOO_00049)
IMGPR_plasmid_3300029796_000040 28286:41996(+)
2-succinyl-6-hydroxy-2,4-cyclohexadiene-1-carboxylate synthase (HDDFDMKI_00038)
IS4 family transposase IS943 (HDDFDMKI_00037)
IS110 family transposase ISWpi13 (HDDFDMKI_00043)
SigE (VF0295) (HDDFDMKI_00039)
36 40 41 42 44
IMGPR_plasmid_2791355228_000003 25559:36277(+)
5000 nt
Virulence
AMR