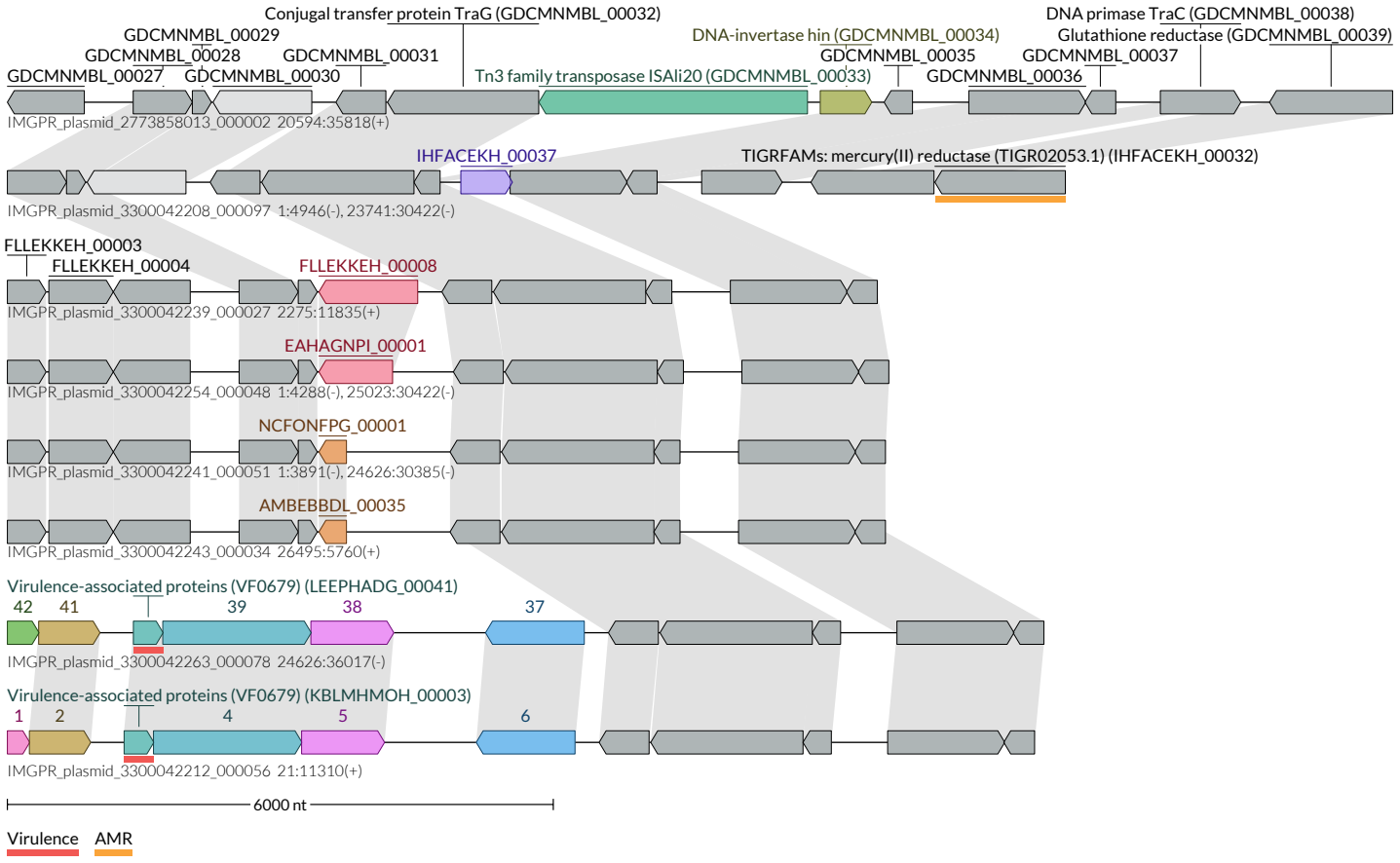
Conjugal transfer protein TraG (GDCMNMBL_00032)
GDCMNMBL_00029
GDCMNMBL_00028
GDCMNMBL_00031
GDCMNMBL_00027
GDCMNMBL_00030
DNA-invertase hin (GDCMNMBL_00034)
DNA primase TraC (GDCMNMBL_00038)
Glutathione reductase (GDCMNMBL_00039)
GDCMNMBL_00035
GDCMNMBL_00037
Tn3 family transposase ISAli20 (GDCMNMBL_00033)
GDCMNMBL_00036
IMGPR_plasmid_2773858013_000002 20594:35818(+)
IHFACEKH_00037
TIGRFAMs: mercury(II) reductase (TIGR02053.1) (IHFACEKH_00032)
IMGPR_plasmid_3300042208_000097 1:4946(-), 23741:30422(-)
FLLEKKEH_00003
FLLEKKEH_00004
FLLEKKEH_00008
IMGPR_plasmid_3300042239_000027 2275:11835(+)
EAHAGNPI_00001
IMGPR_plasmid_3300042254_000048 1:4288(-), 25023:30422(-)
NCFONFPG_00001
IMGPR_plasmid_3300042241_000051 1:3891(-), 24626:30385(-)
AMBEBBDL_00035
IMGPR_plasmid_3300042243_000034 26495:5760(+)
Virulence-associated proteins (VF0679) (LEEPHADG_00041)
42
41
39
38
37
IMGPR_plasmid_3300042263_000078 24626:36017(-)
Virulence-associated proteins (VF0679) (KBLMHMOH_00003)
1
2
4
5
6
IMGPR_plasmid_3300042212_000056 21:11310(+)
6000 nt
Virulence
AMR