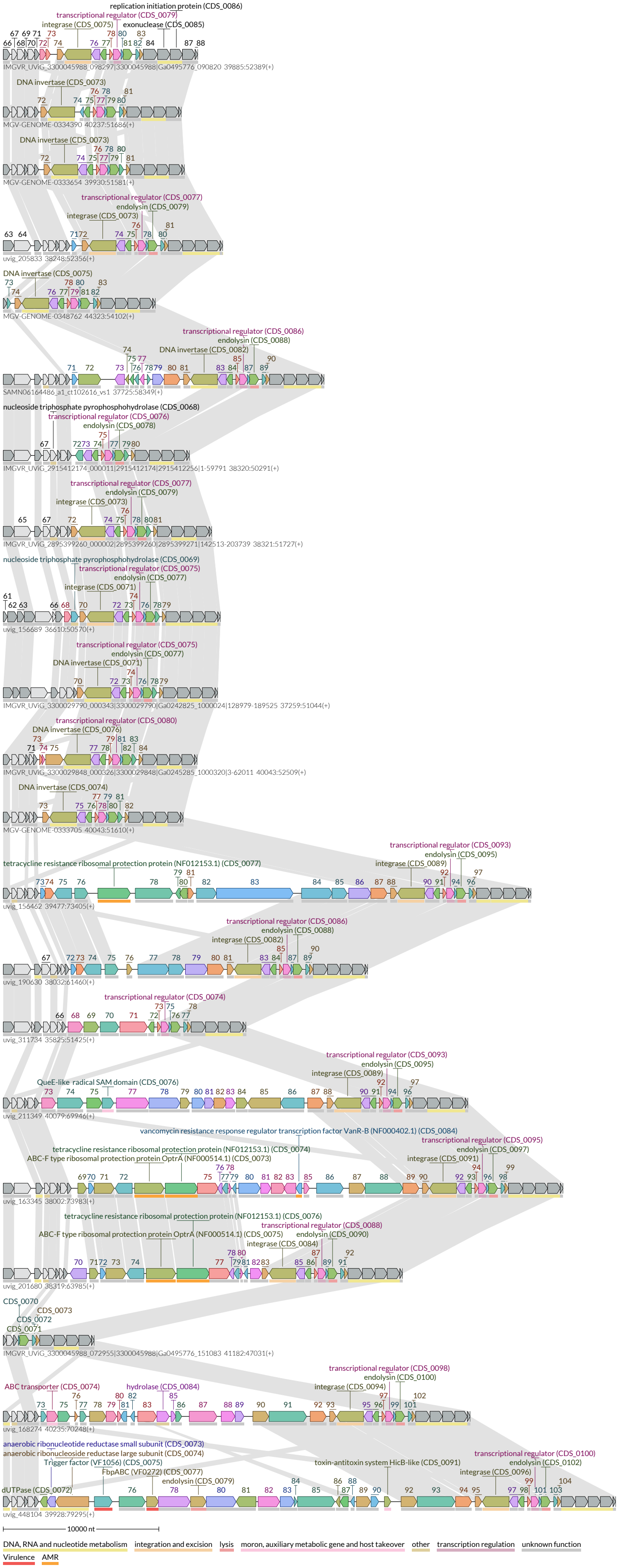
replication initiation protein (CDS_0086)
transcriptional regulator (CDS_0079)
integrase (CDS_0075)
exonuclease (CDS_0085)
IMGVR_UViG_3300045988_098297|3300045988|Ga0495776_090820 39885:52389(+)
DNA invertase (CDS_0073)
MGV-GENOME-0334390 40237:51686(+)
DNA invertase (CDS_0073)
MGV-GENOME-0333654 39930:51581(+)
transcriptional regulator (CDS_0077)
endolysin (CDS_0079)
integrase (CDS_0073)
uvig_205833 38248:52356(+)
DNA invertase (CDS_0075)
MGV-GENOME-0348762 44323:54102(+)
transcriptional regulator (CDS_0086)
endolysin (CDS_0088)
DNA invertase (CDS_0082)
SAMN06164486_a1_ct102616_vs1 37725:58349(+)
nucleoside triphosphate pyrophosphohydrolase (CDS_0068)
transcriptional regulator (CDS_0076)
endolysin (CDS_0078)
IMGVR_UViG_2915412174_000011|2915412174|2915412256|1-59791 38320:50291(+)
transcriptional regulator (CDS_0077)
endolysin (CDS_0079)
integrase (CDS_0073)
IMGVR_UViG_2895399260_000002|2895399260|2895399271|142513-203739 38321:51727(+)
nucleoside triphosphate pyrophosphohydrolase (CDS_0069)
transcriptional regulator (CDS_0075)
endolysin (CDS_0077)
integrase (CDS_0071)
uvig_156689 36610:50570(+)
transcriptional regulator (CDS_0075)
endolysin (CDS_0077)
DNA invertase (CDS_0071)
IMGVR_UViG_3300029790_000343|3300029790|Ga0242825_1000024|128979-189525 37259:51044(+)
transcriptional regulator (CDS_0080)
DNA invertase (CDS_0076)
IMGVR_UViG_3300029848_000326|3300029848|Ga0245285_1000320|3-62011 40043:52509(+)
DNA invertase (CDS_0074)
MGV-GENOME-0333705 40043:51610(+)
transcriptional regulator (CDS_0093)
endolysin (CDS_0095)
integrase (CDS_0089)
tetracycline resistance ribosomal protection protein (NF012153.1) (CDS_0077)
uvig_156462 39477:73405(+)
transcriptional regulator (CDS_0086)
endolysin (CDS_0088)
integrase (CDS_0082)
uvig_190630 38032:61460(+)
transcriptional regulator (CDS_0074)
uvig_311734 35825:51425(+)
transcriptional regulator (CDS_0093)
endolysin (CDS_0095)
integrase (CDS_0089)
QueE-like radical SAM domain (CDS_0076)
uvig_211349 40079:69946(+)
vancomycin resistance response regulator transcription factor VanR-B (NF000402.1) (CDS_0084)
transcriptional regulator (CDS_0095)
endolysin (CDS_0097)
tetracycline resistance ribosomal protection protein (NF012153.1) (CDS_0074)
ABC-F type ribosomal protection protein OptrA (NF000514.1) (CDS_0073)
integrase (CDS_0091)
uvig_163345 38002:73983(+)
tetracycline resistance ribosomal protection protein (NF012153.1) (CDS_0076)
transcriptional regulator (CDS_0088)
ABC-F type ribosomal protection protein OptrA (NF000514.1) (CDS_0075)
endolysin (CDS_0090)
integrase (CDS_0084)
uvig_201680 38319:63985(+)
CDS_0070
CDS_0073
CDS_0072
CDS_0071
IMGVR_UViG_3300045988_072955|3300045988|Ga0495776_151083 41182:47031(+)
transcriptional regulator (CDS_0098)
endolysin (CDS_0100)
integrase (CDS_0094)
ABC transporter (CDS_0074)
hydrolase (CDS_0084)
uvig_168274 40235:70248(+)
anaerobic ribonucleotide reductase small subunit (CDS_0073)
anaerobic ribonucleoside reductase large subunit (CDS_0074)
Trigger factor (VF1056) (CDS_0075)
FbpABC (VF0272) (CDS_0077)
endolysin (CDS_0079)
dUTPase (CDS_0072)
toxin-antitoxin system HicB-like (CDS_0091)
transcriptional regulator (CDS_0100)
endolysin (CDS_0102)
integrase (CDS_0096)
uvig_448104 39928:79295(+)
10000 nt
DNA, RNA and nucleotide metabolism
integration and excision
lysis
moron, auxiliary metabolic gene and host takeover
other
transcription regulation
unknown function
Virulence
AMR