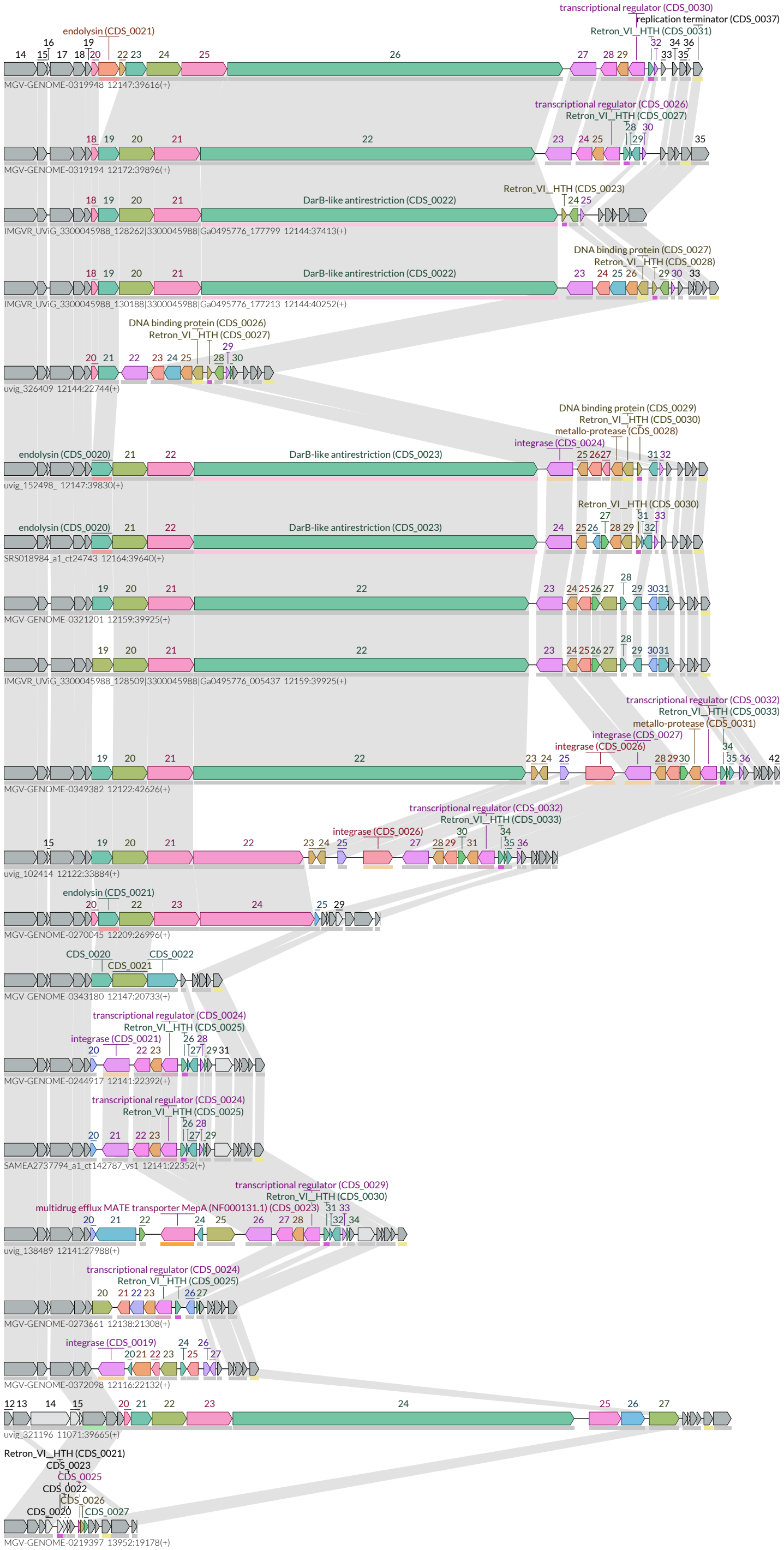
transcriptional regulator (CDS_0030)
replication terminator (CDS_0037)
endolysin (CDS_0021)
Retron_VI_HTH (CDS_0031)
MGV-GENOME-0319948 12147:39616(+)
transcriptional regulator (CDS_0026)
Retron_VI_HTH (CDS_0027)
MGV-GENOME-0319194 12172:39896(+)
Retron_VI_HTH (CDS_0023)
DarB-like antirestriction (CDS_0022)
IMGVR_UViG_3300045988_128262|3300045988|Ga0495776_177799 12144:37413(+)
DNA binding protein (CDS_0027)
Retron_VI_HTH (CDS_0028)
DarB-like antirestriction (CDS_0022)
IMGVR_UViG_3300045988_130188|3300045988|Ga0495776_177213 12144:40252(+)
DNA binding protein (CDS_0026)
Retron_VI_HTH (CDS_0027)
uvig_326409 12144:22744(+)
DNA binding protein (CDS_0029)
Retron_VI_HTH (CDS_0030)
metallo-protease (CDS_0028)
integrase (CDS_0024)
endolysin (CDS_0020)
DarB-like antirestriction (CDS_0023)
uvig_152498_ 12147:39830(+)
Retron_VI_HTH (CDS_0030)
endolysin (CDS_0020)
DarB-like antirestriction (CDS_0023)
SRS018984_a1_ct24743 12164:39640(+)
MGV-GENOME-0321201 12159:39925(+)
IMGVR_UViG_3300045988_128509|3300045988|Ga0495776_005437 12159:39925(+)
transcriptional regulator (CDS_0032)
Retron_VI_HTH (CDS_0033)
metallo-protease (CDS_0031)
integrase (CDS_0027)
integrase (CDS_0026)
MGV-GENOME-0349382 12122:42626(+)
transcriptional regulator (CDS_0032)
Retron_VI_HTH (CDS_0033)
integrase (CDS_0026)
uvig_102414 12122:33884(+)
endolysin (CDS_0021)
MGV-GENOME-0270045 12209:26996(+)
CDS_0020
CDS_0022
CDS_0021
MGV-GENOME-0343180 12147:20733(+)
transcriptional regulator (CDS_0024)
Retron_VI_HTH (CDS_0025)
integrase (CDS_0021)
MGV-GENOME-0244917 12141:22392(+)
transcriptional regulator (CDS_0024)
Retron_VI_HTH (CDS_0025)
SAMEA2737794_a1_ct142787_vs1 12141:22352(+)
transcriptional regulator (CDS_0029)
Retron_VI_HTH (CDS_0030)
multidrug efflux MATE transporter MepA (NF000131.1) (CDS_0023)
uvig_138489 12141:27988(+)
transcriptional regulator (CDS_0024)
Retron_VI_HTH (CDS_0025)
MGV-GENOME-0273661 12138:21308(+)
integrase (CDS_0019)
MGV-GENOME-0372098 12116:22132(+)
uvig_321196 11071:39665(+)
Retron_VI_HTH (CDS_0021)
CDS_0023
CDS_0025
CDS_0022
CDS_0026
CDS_0020
CDS_0027
MGV-GENOME-0219397 13952:19178(+)
10000 nt
DNA, RNA and nucleotide metabolism
integration and excision
lysis
moron, auxiliary metabolic gene and host takeover
other
tail
transcription regulation
unknown function
Defence
AMR