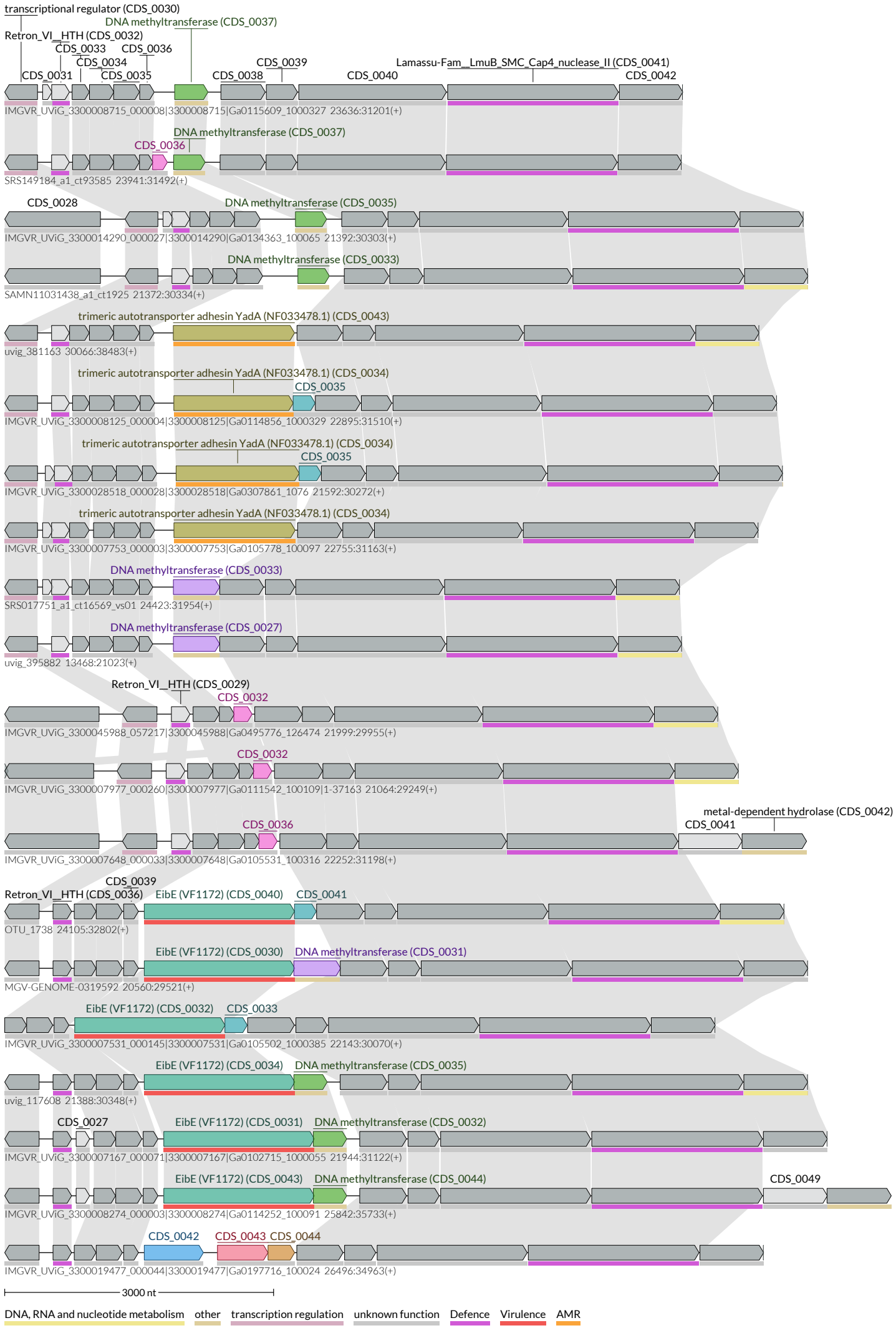
transcriptional regulator (CDS_0030)
DNA methyltransferase (CDS_0037)
Retron_VI__HTH (CDS_0032)
CDS_0033
CDS_0036
CDS_0034
CDS_0039
Lamassu-Fam__LmuB_SMC_Cap4_nuclease_II (CDS_0041)
CDS_0031
CDS_0035
CDS_0038
CDS_0040
CDS_0042
IMGVR_UViG_3300008715_000008|3300008715|Ga0115609_1000327 23636:31201(+)
DNA methyltransferase (CDS_0037)
CDS_0036
SRS149184_a1_ct93585 23941:31492(+)
CDS_0028
DNA methyltransferase (CDS_0035)
IMGVR_UViG_3300014290_000027|3300014290|Ga0134363_100065 21392:30303(+)
DNA methyltransferase (CDS_0033)
SAMN11031438_a1_ct1925 21372:30334(+)
trimeric autotransporter adhesin YadA (NF033478.1) (CDS_0043)
uvig_381163 30066:38483(+)
trimeric autotransporter adhesin YadA (NF033478.1) (CDS_0034)
CDS_0035
IMGVR_UViG_3300008125_000004|3300008125|Ga0114856_1000329 22895:31510(+)
trimeric autotransporter adhesin YadA (NF033478.1) (CDS_0034)
CDS_0035
IMGVR_UViG_3300028518_000028|3300028518|Ga0307861_1076 21592:30272(+)
trimeric autotransporter adhesin YadA (NF033478.1) (CDS_0034)
IMGVR_UViG_3300007753_000003|3300007753|Ga0105778_100097 22755:31163(+)
DNA methyltransferase (CDS_0033)
SRS017751_a1_ct16569_vs01 24423:31954(+)
DNA methyltransferase (CDS_0027)
uvig_395882 13468:21023(+)
Retron_VI__HTH (CDS_0029)
CDS_0032
IMGVR_UViG_3300045988_057217|3300045988|Ga0495776_126474 21999:29955(+)
CDS_0032
IMGVR_UViG_3300007977_000260|3300007977|Ga0111542_100109|1-37163 21064:29249(+)
metal-dependent hydrolase (CDS_0042)
CDS_0036
CDS_0041
IMGVR_UViG_3300007648_000033|3300007648|Ga0105531_100316 22252:31198(+)
CDS_0039
Retron_VI__HTH (CDS_0036)
EibE (VF1172) (CDS_0040)
CDS_0041
OTU_1738 24105:32802(+)
EibE (VF1172) (CDS_0030)
DNA methyltransferase (CDS_0031)
MGV-GENOME-0319592 20560:29521(+)
EibE (VF1172) (CDS_0032)
CDS_0033
IMGVR_UViG_3300007531_000145|3300007531|Ga0105502_1000385 22143:30070(+)
EibE (VF1172) (CDS_0034)
DNA methyltransferase (CDS_0035)
uvig_117608 21388:30348(+)
CDS_0027
EibE (VF1172) (CDS_0031)
DNA methyltransferase (CDS_0032)
IMGVR_UViG_3300007167_000071|3300007167|Ga0102715_1000055 21944:31122(+)
EibE (VF1172) (CDS_0043)
DNA methyltransferase (CDS_0044)
CDS_0049
IMGVR_UViG_3300008274_000003|3300008274|Ga0114252_100091 25842:35733(+)
CDS_0042
CDS_0043
CDS_0044
IMGVR_UViG_3300019477_000044|3300019477|Ga0197716_100024 26496:34963(+)
3000 nt
DNA, RNA and nucleotide metabolism
other
transcription regulation
unknown function
Defence
Virulence
AMR