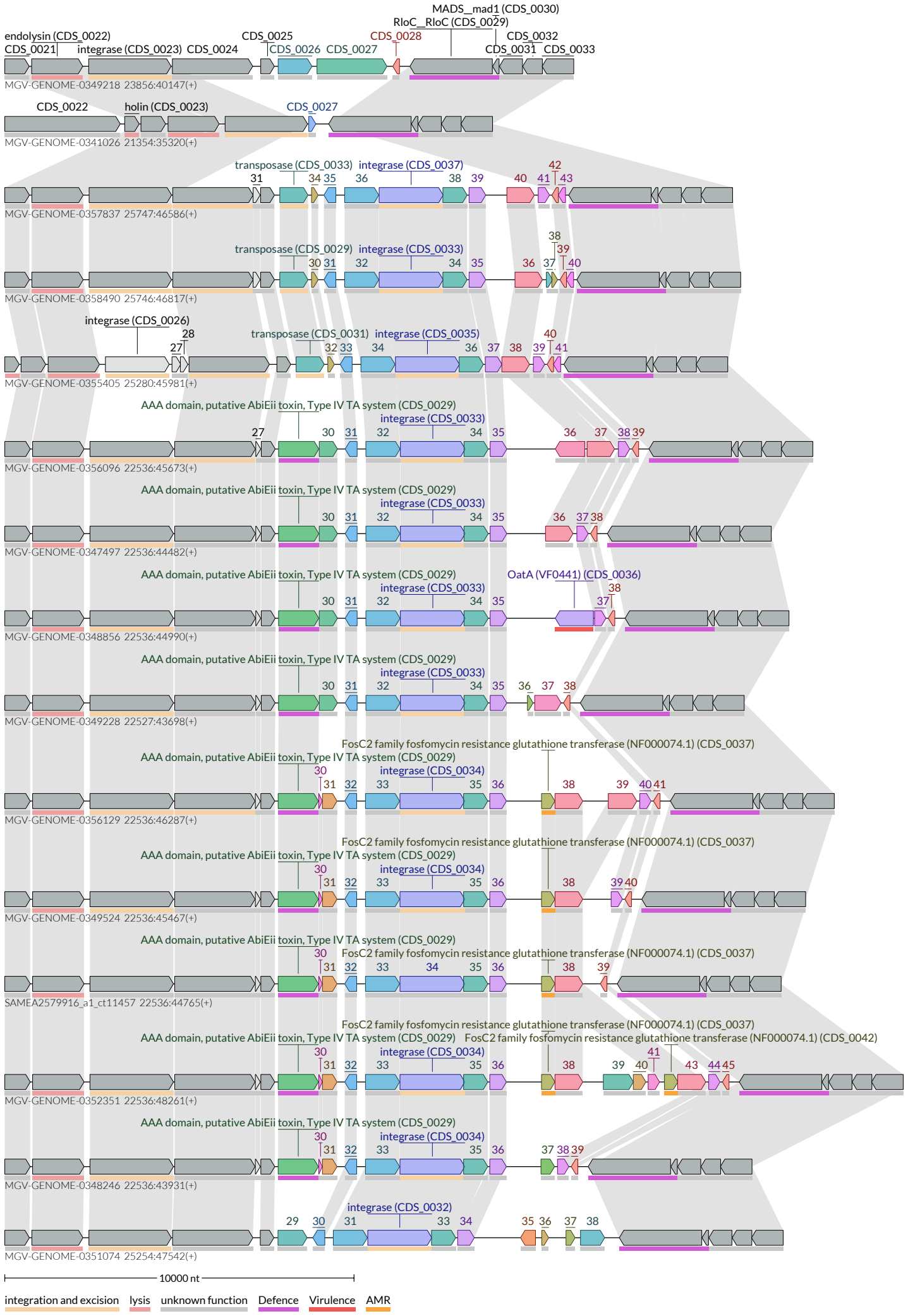
MADS_mad1 (CDS_0030)
RloC_RloC (CDS_0029)
endolysin (CDS_0022)
CDS_0021
integrase (CDS_0023)
CDS_0024
CDS_0025
CDS_0026
CDS_0027
CDS_0028
CDS_0032
CDS_0031
CDS_0033
MGV-GENOME-0349218 23856:40147(+)
CDS_0022
holin (CDS_0023)
CDS_0027
MGV-GENOME-0341026 21354:35320(+)
transposase (CDS_0033)
integrase (CDS_0037)
MGV-GENOME-0357837 25747:46586(+)
transposase (CDS_0029)
integrase (CDS_0033)
MGV-GENOME-0358490 25746:46817(+)
integrase (CDS_0026)
transposase (CDS_0031)
integrase (CDS_0035)
MGV-GENOME-0355405 25280:45981(+)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0029)
integrase (CDS_0033)
MGV-GENOME-0356096 22536:45673(+)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0029)
integrase (CDS_0033)
MGV-GENOME-0347497 22536:44482(+)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0029)
OatA (VF0441) (CDS_0036)
integrase (CDS_0033)
MGV-GENOME-0348856 22536:44990(+)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0029)
integrase (CDS_0033)
MGV-GENOME-0349228 22527:43698(+)
FosC2 family fosfomycin resistance glutathione transferase (NF000074.1) (CDS_0037)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0029)
integrase (CDS_0034)
MGV-GENOME-0356129 22536:46287(+)
FosC2 family fosfomycin resistance glutathione transferase (NF000074.1) (CDS_0037)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0029)
integrase (CDS_0034)
MGV-GENOME-0349524 22536:45467(+)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0029)
FosC2 family fosfomycin resistance glutathione transferase (NF000074.1) (CDS_0037)
SAMEA2579916_a1_ct11457 22536:44765(+)
FosC2 family fosfomycin resistance glutathione transferase (NF000074.1) (CDS_0037)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0029)
FosC2 family fosfomycin resistance glutathione transferase (NF000074.1) (CDS_0042)
integrase (CDS_0034)
MGV-GENOME-0352351 22536:48261(+)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0029)
integrase (CDS_0034)
MGV-GENOME-0348246 22536:43931(+)
integrase (CDS_0032)
MGV-GENOME-0351074 25254:47542(+)
10000 nt
integration and excision
lysis
unknown function
Defence
Virulence
AMR