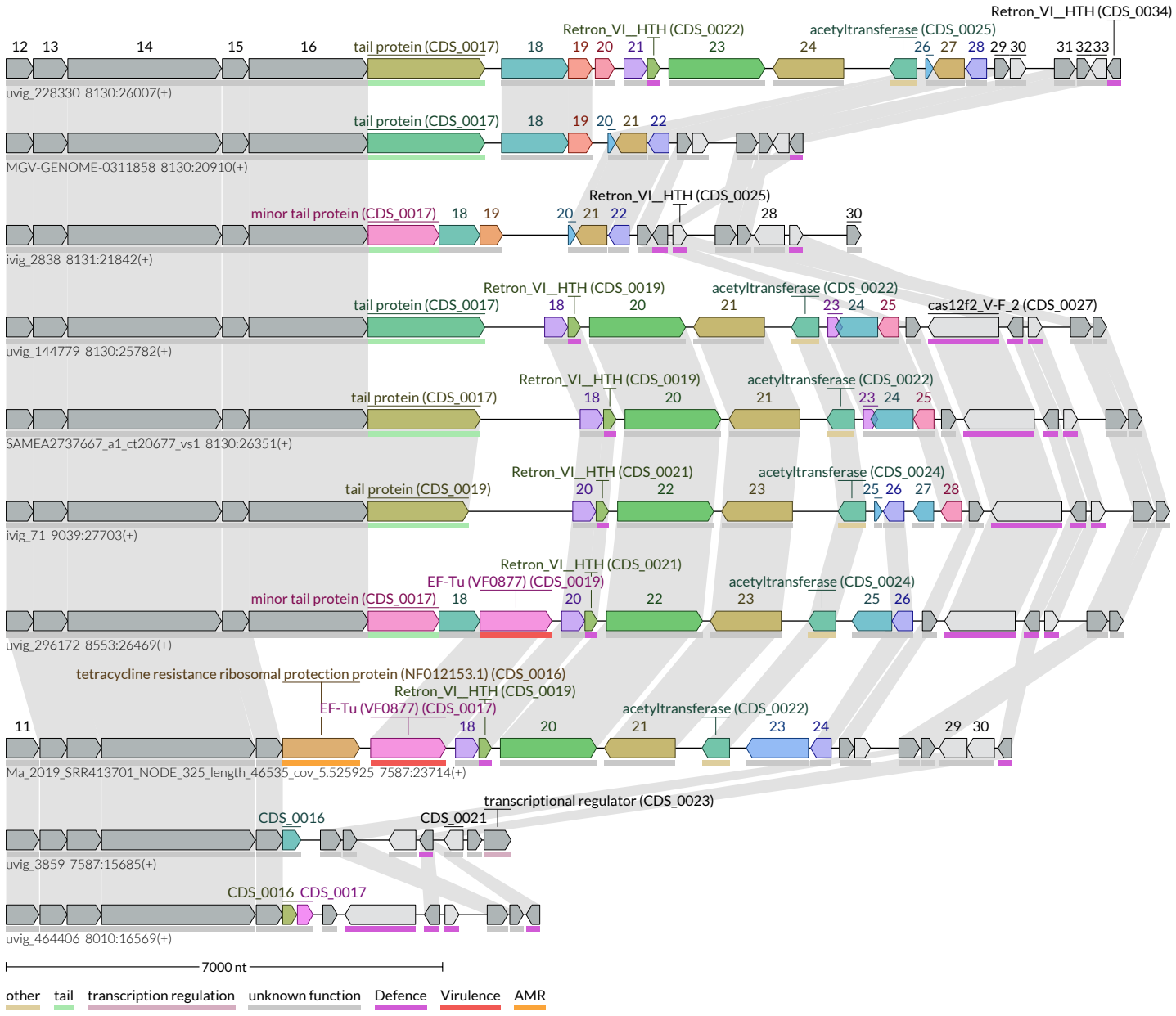
Retron_VI__HTH (CDS_0034)
Retron_VI__HTH (CDS_0022)
acetyltransferase (CDS_0025)
12 13 14 15 16 tail protein (CDS_0017) 18 19 20 21 23 24 26 27 28 29 30 31 32 33
uvig_228330 8130:26007(+)
tail protein (CDS_0017) 18 19 20 21 22
MGV-GENOME-0311858 8130:20910(+)
Retron_VI__HTH (CDS_0025)
minor tail protein (CDS_0017) 18 19 20 21 22 28 30
ivig_2838 8131:21842(+)
Retron_VI__HTH (CDS_0019)
acetyltransferase (CDS_0022)
tail protein (CDS_0017) 18 20 21 23 24 25 cas12f2_V-F_2 (CDS_0027)
uvig_144779 8130:25782(+)
Retron_VI__HTH (CDS_0019)
acetyltransferase (CDS_0022)
tail protein (CDS_0017) 18 20 21 23 24 25
SAMEA2737667_a1_ct20677_vs1 8130:26351(+)
Retron_VI__HTH (CDS_0021)
acetyltransferase (CDS_0024)
tail protein (CDS_0019) 20 22 23 25 26 27 28
ivig_71 9039:27703(+)
Retron_VI__HTH (CDS_0021)
EF-Tu (VF0877) (CDS_0019)
acetyltransferase (CDS_0024)
minor tail protein (CDS_0017) 18 20 22 23 25 26
uvig_296172 8553:26469(+)
tetracycline resistance ribosomal protection protein (NF012153.1) (CDS_0016)
Retron_VI__HTH (CDS_0019)
EF-Tu (VF0877) (CDS_0017)
acetyltransferase (CDS_0022)
11 18 20 21 23 24 29 30
Ma_2019_SRR413701_NODE_325_length_46535_cov_5.525925 7587:23714(+)
transcriptional regulator (CDS_0023)
CDS_0016 CDS_0021
uvig_3859 7587:15685(+)
CDS_0016 CDS_0017
uvig_464406 8010:16569(+)
7000 nt
other tail transcription regulation unknown function Defence Virulence AMR