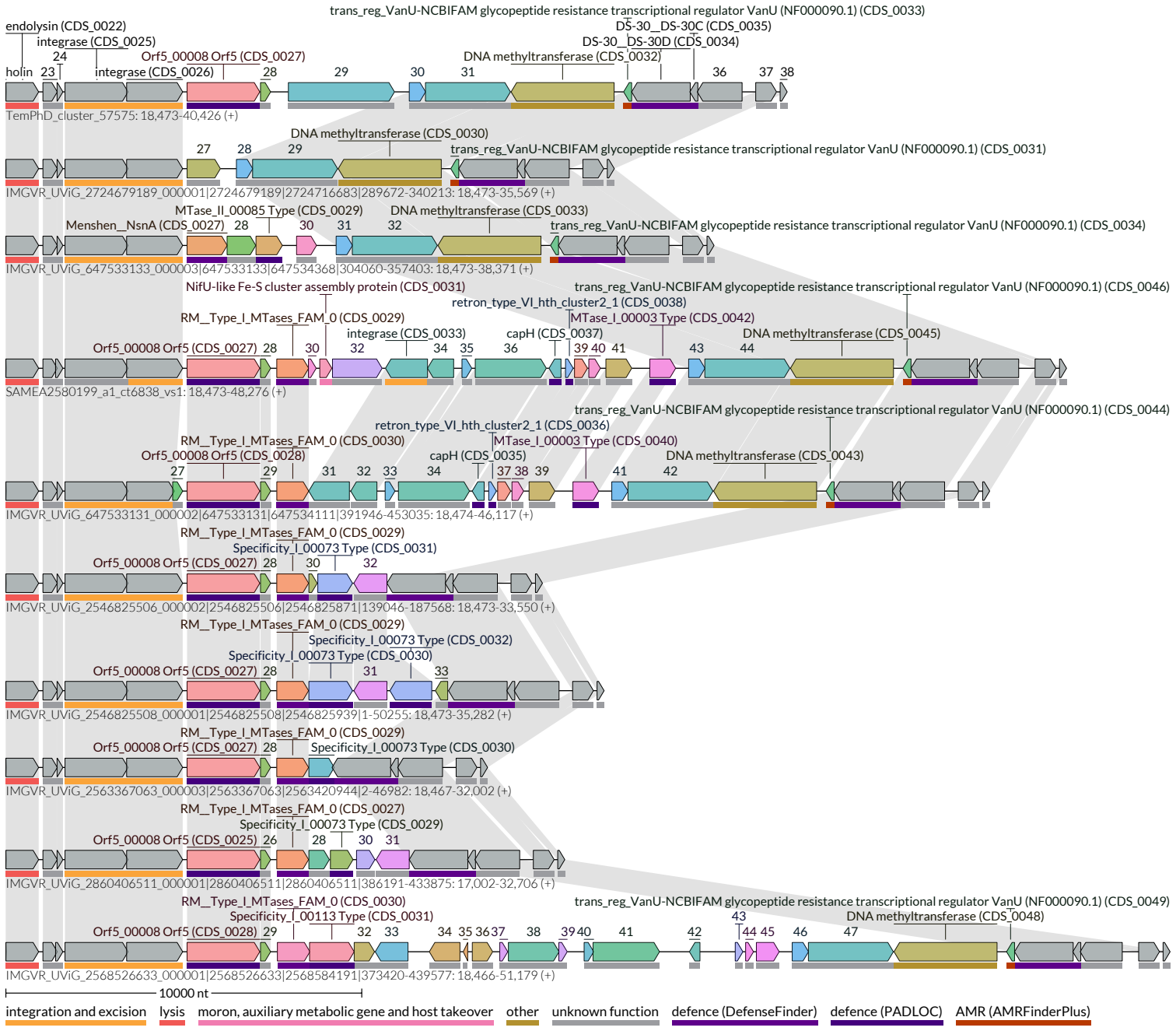
trans_reg_VanU-NCBIFAM glycopeptide resistance transcriptional regulator VanU (NF000090.1) (CDS_0033)
endolysin (CDS_0022)
integrase (CDS_0025)
DS-30__DS-30C (CDS_0035)
DS-30__DS-30D (CDS_0034)
Orf5_00008 Orf5 (CDS_0027)
DNA methyltransferase (CDS_0032)
integrase (CDS_0026)
holin
TemPhD_cluster_57575: 18,473-40,426 (+)
DNA methyltransferase (CDS_0030)
trans_reg_VanU-NCBIFAM glycopeptide resistance transcriptional regulator VanU (NF000090.1) (CDS_0031)
IMGVR_UViG_2724679189_000001|2724679189|2724716683|289672-340213: 18,473-35,569 (+)
MTase_II_00085 Type (CDS_0029)
DNA methyltransferase (CDS_0033)
Menshen__NsnA (CDS_0027)
trans_reg_VanU-NCBIFAM glycopeptide resistance transcriptional regulator VanU (NF000090.1) (CDS_0034)
IMGVR_UViG_647533133_000003|647533133|647534368|304060-357403: 18,473-38,371 (+)
NifU-like Fe-S cluster assembly protein (CDS_0031)
trans_reg_VanU-NCBIFAM glycopeptide resistance transcriptional regulator VanU (NF000090.1) (CDS_0046)
retron_type_VI_hth_cluster2_1 (CDS_0038)
RM__Type_I_MTases_FAM_0 (CDS_0029)
MTase_I_00003 Type (CDS_0042)
integrase (CDS_0033)
capH (CDS_0037)
DNA methyltransferase (CDS_0045)
Orf5_00008 Orf5 (CDS_0027)
SAMEA2580199_a1_ct6838_vs1: 18,473-48,276 (+)
trans_reg_VanU-NCBIFAM glycopeptide resistance transcriptional regulator VanU (NF000090.1) (CDS_0044)
retron_type_VI_hth_cluster2_1 (CDS_0036)
RM__Type_I_MTases_FAM_0 (CDS_0030)
MTase_I_00003 Type (CDS_0040)
Orf5_00008 Orf5 (CDS_0028)
capH (CDS_0035)
DNA methyltransferase (CDS_0043)
IMGVR_UViG_647533131_000002|647533131|647534111|391946-453035: 18,474-46,117 (+)
RM__Type_I_MTases_FAM_0 (CDS_0029)
Specificity_I_00073 Type (CDS_0031)
Orf5_00008 Orf5 (CDS_0027)
IMGVR_UViG_2546825506_000002|2546825506|2546825871|139046-187568: 18,473-33,550 (+)
RM__Type_I_MTases_FAM_0 (CDS_0029)
Specificity_I_00073 Type (CDS_0032)
Specificity_I_00073 Type (CDS_0030)
Orf5_00008 Orf5 (CDS_0027)
IMGVR_UViG_2546825508_000001|2546825508|2546825939|1-50255: 18,473-35,282 (+)
RM__Type_I_MTases_FAM_0 (CDS_0029)
Orf5_00008 Orf5 (CDS_0027)
Specificity_I_00073 Type (CDS_0030)
IMGVR_UViG_2563367063_000003|2563367063|2563420944|2-46982: 18,467-32,002 (+)
RM__Type_I_MTases_FAM_0 (CDS_0027)
Specificity_I_00073 Type (CDS_0029)
Orf5_00008 Orf5 (CDS_0025)
IMGVR_UViG_2860406511_000001|2860406511|2860406511|386191-433875: 17,002-32,706 (+)
RM__Type_I_MTases_FAM_0 (CDS_0030)
trans_reg_VanU-NCBIFAM glycopeptide resistance transcriptional regulator VanU (NF000090.1) (CDS_0049)
Specificity_I_00113 Type (CDS_0031)
DNA methyltransferase (CDS_0048)
Orf5_00008 Orf5 (CDS_0028)
IMGVR_UViG_2568526633_000001|2568526633|2568584191|373420-439577: 18,466-51,179 (+)
10000 nt
integration and excision
lysis
moron, auxiliary metabolic gene and host takeover
other
unknown function
defence (DefenseFinder)
defence (PADLOC)
AMR (AMRFinderPlus)