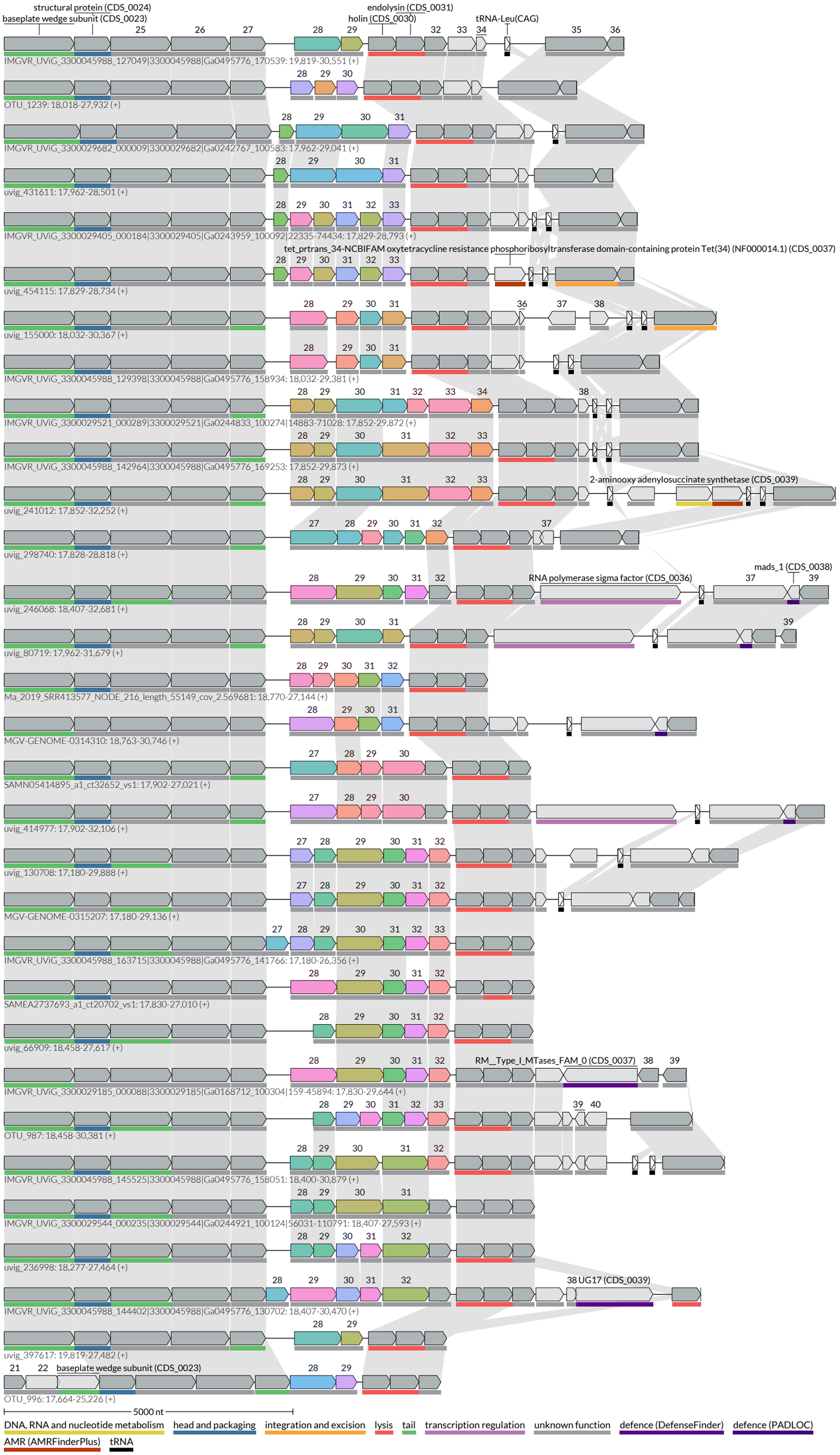
structural protein (CDS_0024)
baseplate wedge subunit (CDS_0023)
endolysin (CDS_0031)
holin (CDS_0030)
tRNA-Leu(CAG)
IMGVR_UViG_3300045988_127049|3300045988|Ga0495776_170539: 19,819-30,551 (+)
OTU_1239: 18,018-27,932 (+)
IMGVR_UViG_3300029682_000009|3300029682|Ga0242767_100583: 17,962-29,041 (+)
uvig_431611: 17,962-28,501 (+)
IMGVR_UViG_3300029405_000184|3300029405|Ga0243959_100092|22335-74434: 17,829-28,793 (+)
tet_prtrans_34-NCBIFAM oxytetracycline resistance phosphoribosyltransferase domain-containing protein Tet(34) (NF000014.1) (CDS_0037)
uvig_454115: 17,829-28,734 (+)
uvig_155000: 18,032-30,367 (+)
IMGVR_UViG_3300045988_129398|3300045988|Ga0495776_158934: 18,032-29,381 (+)
IMGVR_UViG_3300029521_000289|3300029521|Ga0244833_100274|14883-71028: 17,852-29,872 (+)
IMGVR_UViG_3300045988_142964|3300045988|Ga0495776_169253: 17,852-29,873 (+)
2-aminooxy adenylosuccinate synthetase (CDS_0039)
uvig_241012: 17,852-32,252 (+)
uvig_298740: 17,828-28,818 (+)
mads_1 (CDS_0038)
RNA polymerase sigma factor (CDS_0036)
uvig_246068: 18,407-32,681 (+)
uvig_80719: 17,962-31,679 (+)
Ma_2019_SRR413577_NODE_216_length_55149_cov_2.569681: 18,770-27,144 (+)
MGV-GENOME-0314310: 18,763-30,746 (+)
SAMN05414895_a1_ct32652_vs1: 17,902-27,021 (+)
uvig_414977: 17,902-32,106 (+)
uvig_130708: 17,180-29,888 (+)
MGV-GENOME-0315207: 17,180-29,136 (+)
IMGVR_UViG_3300045988_163715|3300045988|Ga0495776_141766: 17,180-26,356 (+)
SAMEA2737693_a1_ct20702_vs1: 17,830-27,010 (+)
uvig_66909: 18,458-27,617 (+)
RM__Type_I_MTases_FAM_0 (CDS_0037)
IMGVR_UViG_3300029185_000088|3300029185|Ga0168712_100304|159-45894: 17,830-29,644 (+)
OTU_987: 18,458-30,381 (+)
IMGVR_UViG_3300045988_145525|3300045988|Ga0495776_158051: 18,400-30,879 (+)
IMGVR_UViG_3300029544_000235|3300029544|Ga0244921_100124|56031-110791: 18,407-27,593 (+)
uvig_236998: 18,277-27,464 (+)
UG17 (CDS_0039)
IMGVR_UViG_3300045988_144402|3300045988|Ga0495776_130702: 18,407-30,470 (+)
uvig_397617: 19,819-27,482 (+)
baseplate wedge subunit (CDS_0023)
OTU_996: 17,664-25,226 (+)
5000 nt
DNA, RNA and nucleotide metabolism
head and packaging
integration and excision
lysis
tail
transcription regulation
unknown function
defence (DefenseFinder)
defence (PADLOC)
AMR (AMRFinderPlus)
tRNA