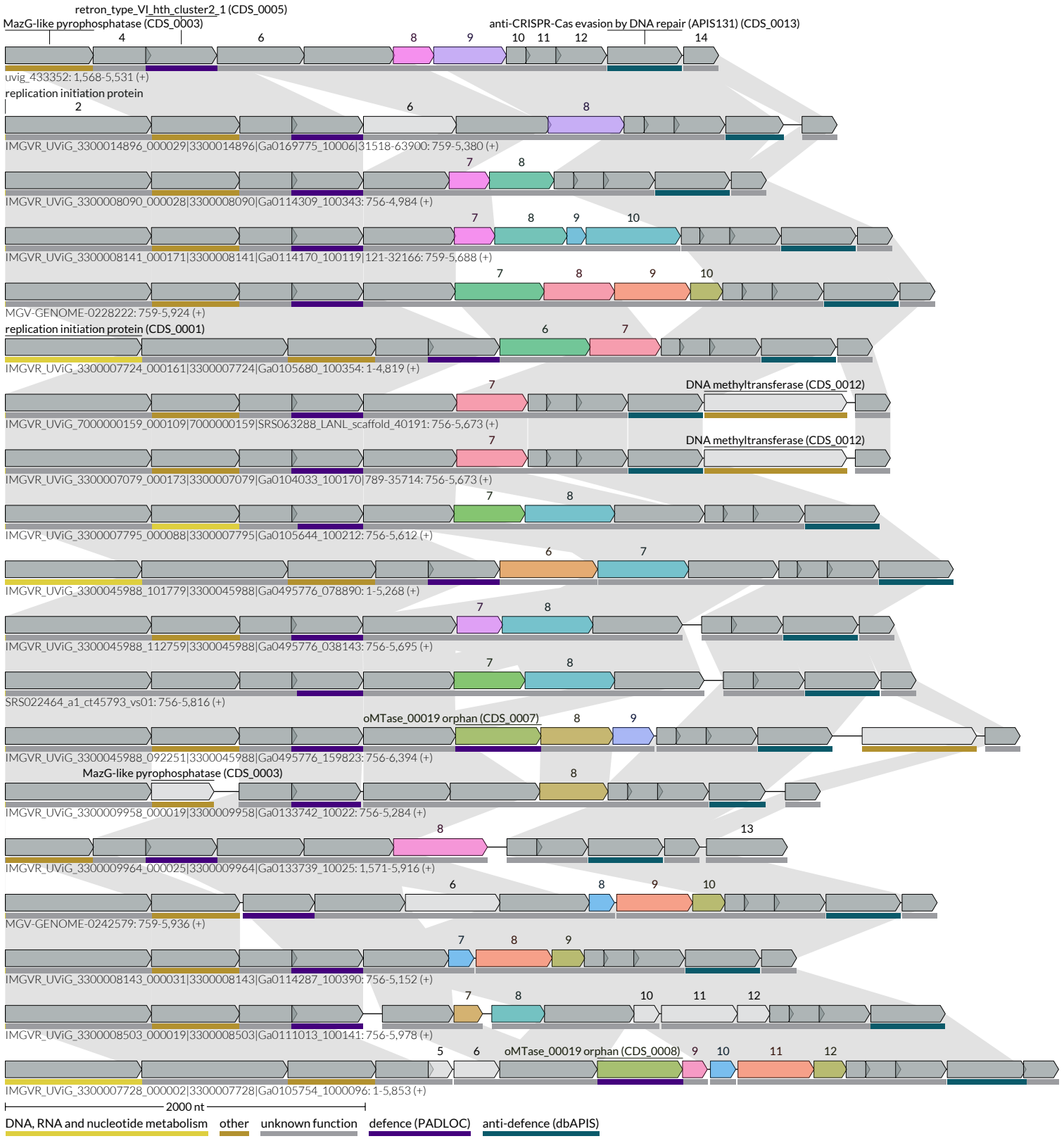
retron_type_VI_hth_cluster2_1 (CDS_0005)
MazG-like pyrophosphatase (CDS_0003)
anti-CRISPR-Cas evasion by DNA repair (APIS131) (CDS_0013)
uvig_433352: 1,568-5,531 (+)
replication initiation protein
IMGVR_UViG_3300014896_000029|3300014896|Ga0169775_10006|31518-63900: 759-5,380 (+)
IMGVR_UViG_3300008090_000028|3300008090|Ga0114309_100343: 756-4,984 (+)
IMGVR_UViG_3300008141_000171|3300008141|Ga0114170_100119|121-32166: 759-5,688 (+)
MGV-GENOME-0228222: 759-5,924 (+)
replication initiation protein (CDS_0001)
IMGVR_UViG_3300007724_000161|3300007724|Ga0105680_100354: 1-4,819 (+)
DNA methyltransferase (CDS_0012)
IMGVR_UViG_7000000159_000109|7000000159|SRS063288_LANL_scaffold_40191: 756-5,673 (+)
IMGVR_UViG_3300007079_000173|3300007079|Ga0104033_100170|789-35714: 756-5,673 (+)
IMGVR_UViG_3300007795_000088|3300007795|Ga0105644_100212: 756-5,612 (+)
IMGVR_UViG_3300045988_101779|3300045988|Ga0495776_078890: 1-5,268 (+)
IMGVR_UViG_3300045988_112759|3300045988|Ga0495776_038143: 756-5,695 (+)
SRS022464_a1_ct45793_vs01: 756-5,816 (+)
oMTase_00019 orphan (CDS_0007)
IMGVR_UViG_3300045988_092251|3300045988|Ga0495776_159823: 756-6,394 (+)
MazG-like pyrophosphatase (CDS_0003)
IMGVR_UViG_3300009958_000019|3300009958|Ga0133742_10022: 756-5,284 (+)
IMGVR_UViG_3300009964_000025|3300009964|Ga0133739_10025: 1,571-5,916 (+)
MGV-GENOME-0242579: 759-5,936 (+)
IMGVR_UViG_3300008143_000031|3300008143|Ga0114287_100390: 756-5,152 (+)
IMGVR_UViG_3300008503_000019|3300008503|Ga0111013_100141: 756-5,978 (+)
oMTase_00019 orphan (CDS_0008)
IMGVR_UViG_3300007728_000002|3300007728|Ga0105754_1000096: 1-5,853 (+)
2000 nt
DNA, RNA and nucleotide metabolism
other
unknown function
defence (PADLOC)
anti-defence (dbAPIS)