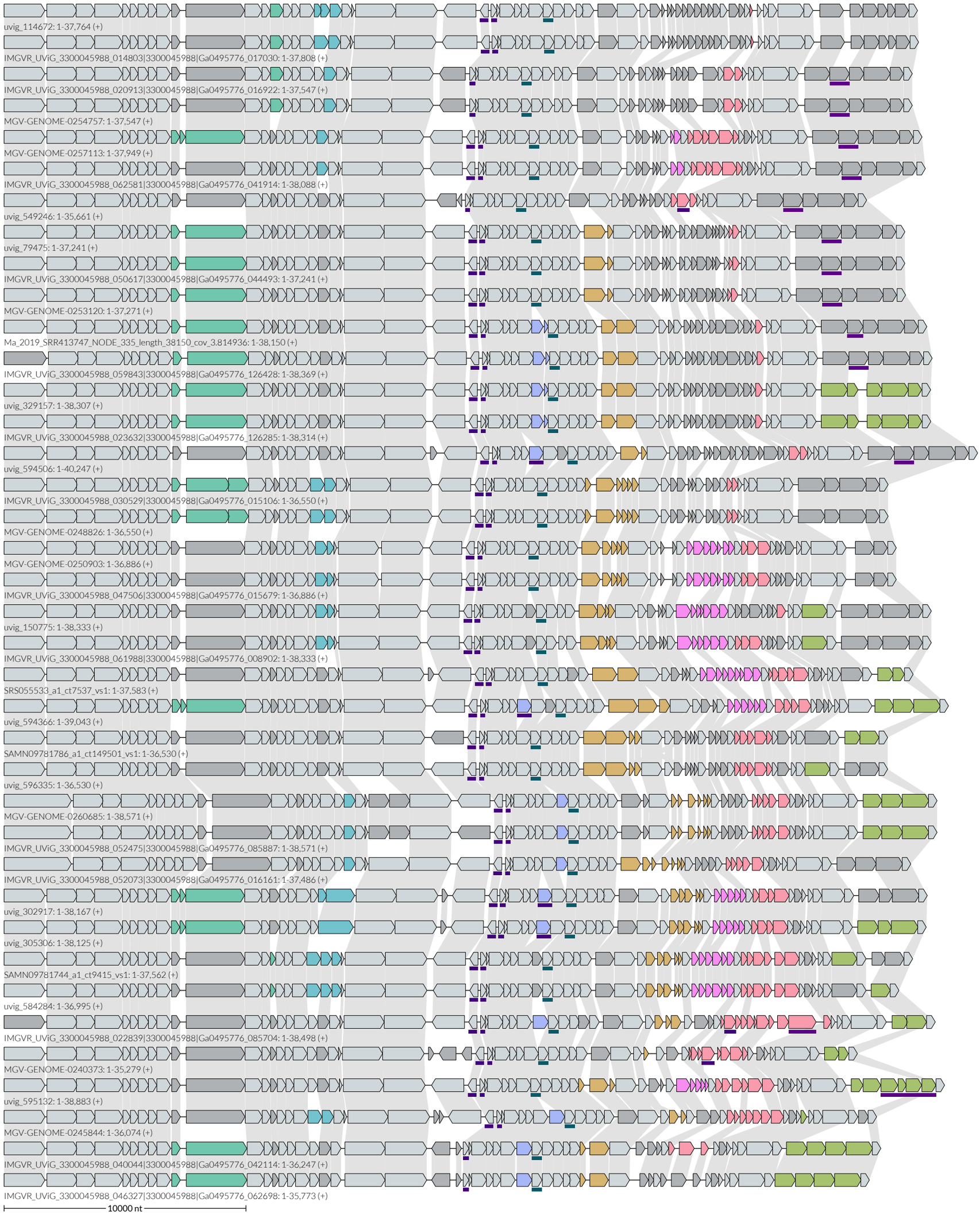
uvig_114672: 1-37,764 (+)
IMGVR_UViG_3300045988_014803|3300045988|Ga0495776_017030: 1-37,808 (+)
IMGVR_UViG_3300045988_020913|3300045988|Ga0495776_016922: 1-37,547 (+)
MGV-GENOME-0254757: 1-37,547 (+)
MGV-GENOME-0257113: 1-37,949 (+)
IMGVR_UViG_3300045988_062581|3300045988|Ga0495776_041914: 1-38,088 (+)
uvig_549246: 1-35,661 (+)
uvig_79475: 1-37,241 (+)
IMGVR_UViG_3300045988_050617|3300045988|Ga0495776_044493: 1-37,241 (+)
MGV-GENOME-0253120: 1-37,271 (+)
Ma_2019_SRR413747_NODE_335_length_38150_cov_3.814936: 1-38,150 (+)
IMGVR_UViG_3300045988_059843|3300045988|Ga0495776_126428: 1-38,369 (+)
uvig_329157: 1-38,307 (+)
IMGVR_UViG_3300045988_023632|3300045988|Ga0495776_126285: 1-38,314 (+)
uvig_594506: 1-40,247 (+)
IMGVR_UViG_3300045988_030529|3300045988|Ga0495776_015106: 1-36,550 (+)
MGV-GENOME-0248826: 1-36,550 (+)
MGV-GENOME-0250903: 1-36,886 (+)
IMGVR_UViG_3300045988_047506|3300045988|Ga0495776_015679: 1-36,886 (+)
uvig_150775: 1-38,333 (+)
IMGVR_UViG_3300045988_061988|3300045988|Ga0495776_008902: 1-38,333 (+)
SRS055533_a1_ct7537_vs1: 1-37,583 (+)
uvig_594366: 1-39,043 (+)
SAMN09781786_a1_ct149501_vs1: 1-36,530 (+)
uvig_596335: 1-36,530 (+)
MGV-GENOME-0260685: 1-38,571 (+)
IMGVR_UViG_3300045988_052475|3300045988|Ga0495776_085887: 1-38,571 (+)
IMGVR_UViG_3300045988_052073|3300045988|Ga0495776_016161: 1-37,486 (+)
uvig_302917: 1-38,167 (+)
uvig_305306: 1-38,125 (+)
SAMN09781744_a1_ct9415_vs1: 1-37,562 (+)
uvig_584284: 1-36,995 (+)
IMGVR_UViG_3300045988_022839|3300045988|Ga0495776_085704: 1-38,498 (+)
MGV-GENOME-0240373: 1-35,279 (+)
uvig_595132: 1-38,883 (+)
MGV-GENOME-0245844: 1-36,074 (+)
IMGVR_UViG_3300045988_040044|3300045988|Ga0495776_042114: 1-36,247 (+)
IMGVR_UViG_3300045988_046327|3300045988|Ga0495776_062698: 1-35,773 (+)
10000 nt
defence (DefenseFinder)
defence (PADLOC)
anti-defence (dbAPIS)