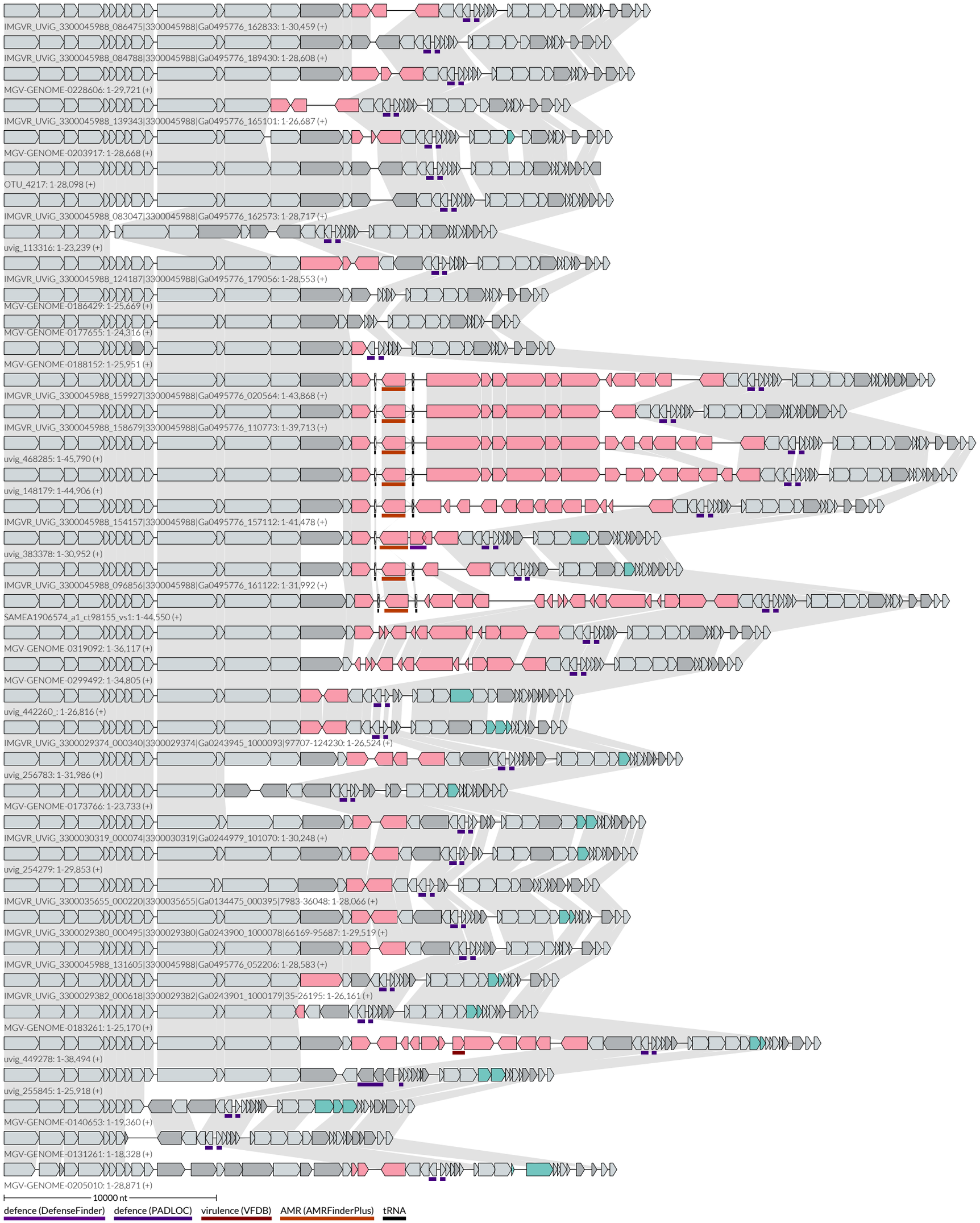
IMGVR_UViG_3300045988_086475|3300045988|Ga0495776_162833: 1-30,459 (+)
IMGVR_UViG_3300045988_084788|3300045988|Ga0495776_189430: 1-28,608 (+)
MGV-GENOME-0228606: 1-29,721 (+)
IMGVR_UViG_3300045988_139343|3300045988|Ga0495776_165101: 1-26,687 (+)
MGV-GENOME-0203917: 1-28,668 (+)
OTU_4217: 1-28,098 (+)
IMGVR_UViG_3300045988_083047|3300045988|Ga0495776_162573: 1-28,717 (+)
uvig_113316: 1-23,239 (+)
IMGVR_UViG_3300045988_124187|3300045988|Ga0495776_179056: 1-28,553 (+)
MGV-GENOME-0186429: 1-25,669 (+)
MGV-GENOME-0177655: 1-24,316 (+)
MGV-GENOME-0188152: 1-25,951 (+)
IMGVR_UViG_3300045988_159927|3300045988|Ga0495776_020564: 1-43,868 (+)
IMGVR_UViG_3300045988_158679|3300045988|Ga0495776_110773: 1-39,713 (+)
uvig_468285: 1-45,790 (+)
uvig_148179: 1-44,906 (+)
IMGVR_UViG_3300045988_154157|3300045988|Ga0495776_157112: 1-41,478 (+)
uvig_383378: 1-30,952 (+)
IMGVR_UViG_3300045988_096856|3300045988|Ga0495776_161122: 1-31,992 (+)
SAMEA1906574_a1_ct98155_vs1: 1-44,550 (+)
MGV-GENOME-0319092: 1-36,117 (+)
MGV-GENOME-0299492: 1-34,805 (+)
uvig_442260_: 1-26,816 (+)
IMGVR_UViG_3300029374_000340|3300029374|Ga0243945_1000093|97707-124230: 1-26,524 (+)
uvig_256783: 1-31,986 (+)
MGV-GENOME-0173766: 1-23,733 (+)
IMGVR_UViG_3300030319_000074|3300030319|Ga0244979_101070: 1-30,248 (+)
uvig_254279: 1-29,853 (+)
IMGVR_UViG_3300035655_000220|3300035655|Ga0134475_000395|7983-36048: 1-28,066 (+)
IMGVR_UViG_3300029380_000495|3300029380|Ga0243900_1000078|66169-95687: 1-29,519 (+)
IMGVR_UViG_3300045988_131605|3300045988|Ga0495776_052206: 1-28,583 (+)
IMGVR_UViG_3300029382_000618|3300029382|Ga0243901_1000179|35-26195: 1-26,161 (+)
MGV-GENOME-0183261: 1-25,170 (+)
uvig_449278: 1-38,494 (+)
uvig_255845: 1-25,918 (+)
MGV-GENOME-0140653: 1-19,360 (+)
MGV-GENOME-0131261: 1-18,328 (+)
MGV-GENOME-0205010: 1-28,871 (+)
10000 nt
defence (DefenseFinder)
defence (PADLOC)
virulence (VFDB)
AMR (AMRFinderPlus)
tRNA