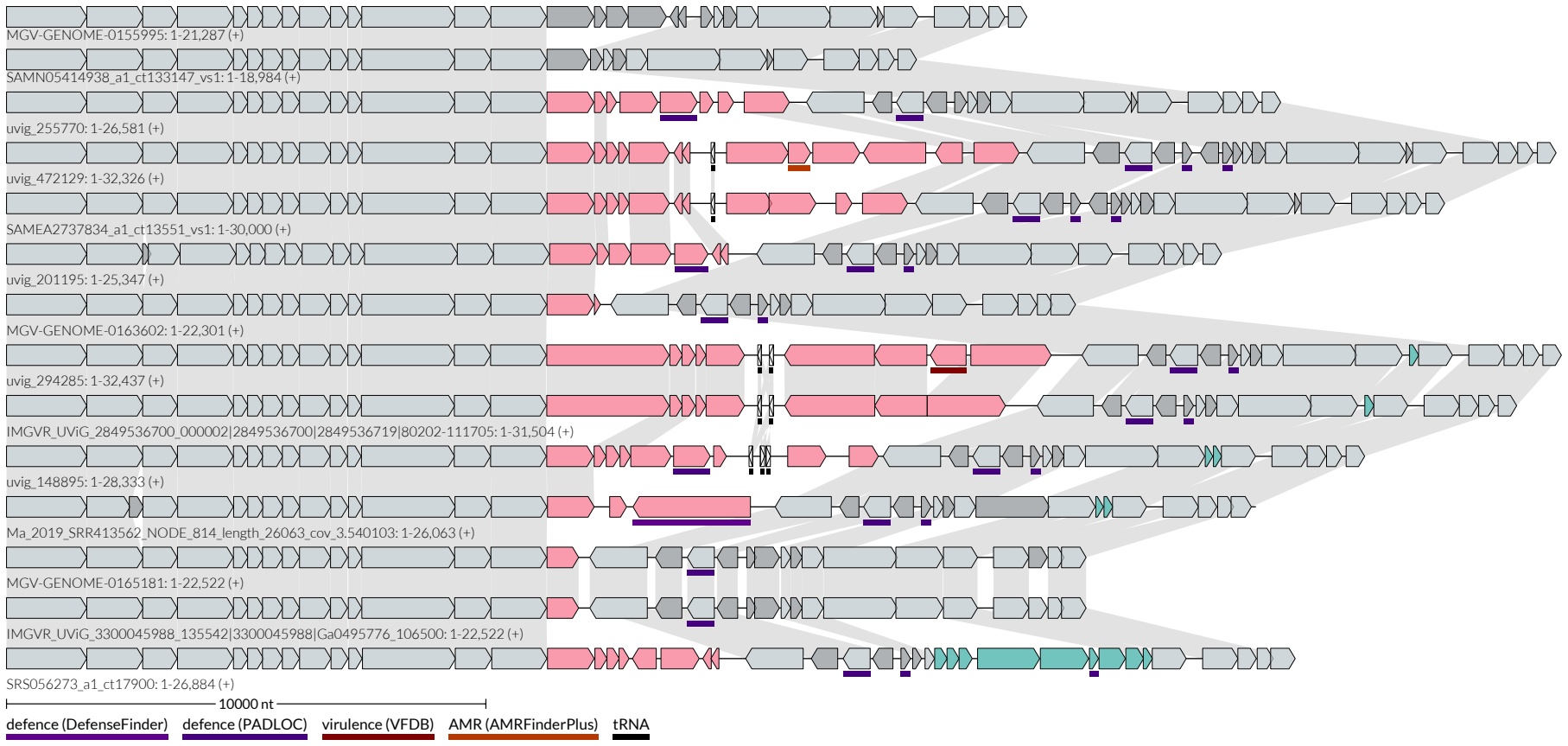
MGV-GENOME-0155995: 1-21,287 (+)
SAMN05414938_a1_ct133147_vs1: 1-18,984 (+)
uvig_255770: 1-26,581 (+)
uvig_472129: 1-32,326 (+)
SAMEA2737834_a1_ct13551_vs1: 1-30,000 (+)
uvig_201195: 1-25,347 (+)
MGV-GENOME-0163602: 1-22,301 (+)
uvig_294285: 1-32,437 (+)
IMGVR_UViG_2849536700_000002|2849536700|2849536719|80202-111705: 1-31,504 (+)
uvig_148895: 1-28,333 (+)
Ma_2019_SRR413562_NODE_814_length_26063_cov_3.540103: 1-26,063 (+)
MGV-GENOME-0165181: 1-22,522 (+)
IMGVR_UViG_3300045988_135542|3300045988|Ga0495776_106500: 1-22,522 (+)
SRS056273_a1_ct17900: 1-26,884 (+)
10000 nt
defence (DefenseFinder)
defence (PADLOC)
virulence (VFDB)
AMR (AMRFinderPlus)
tRNA