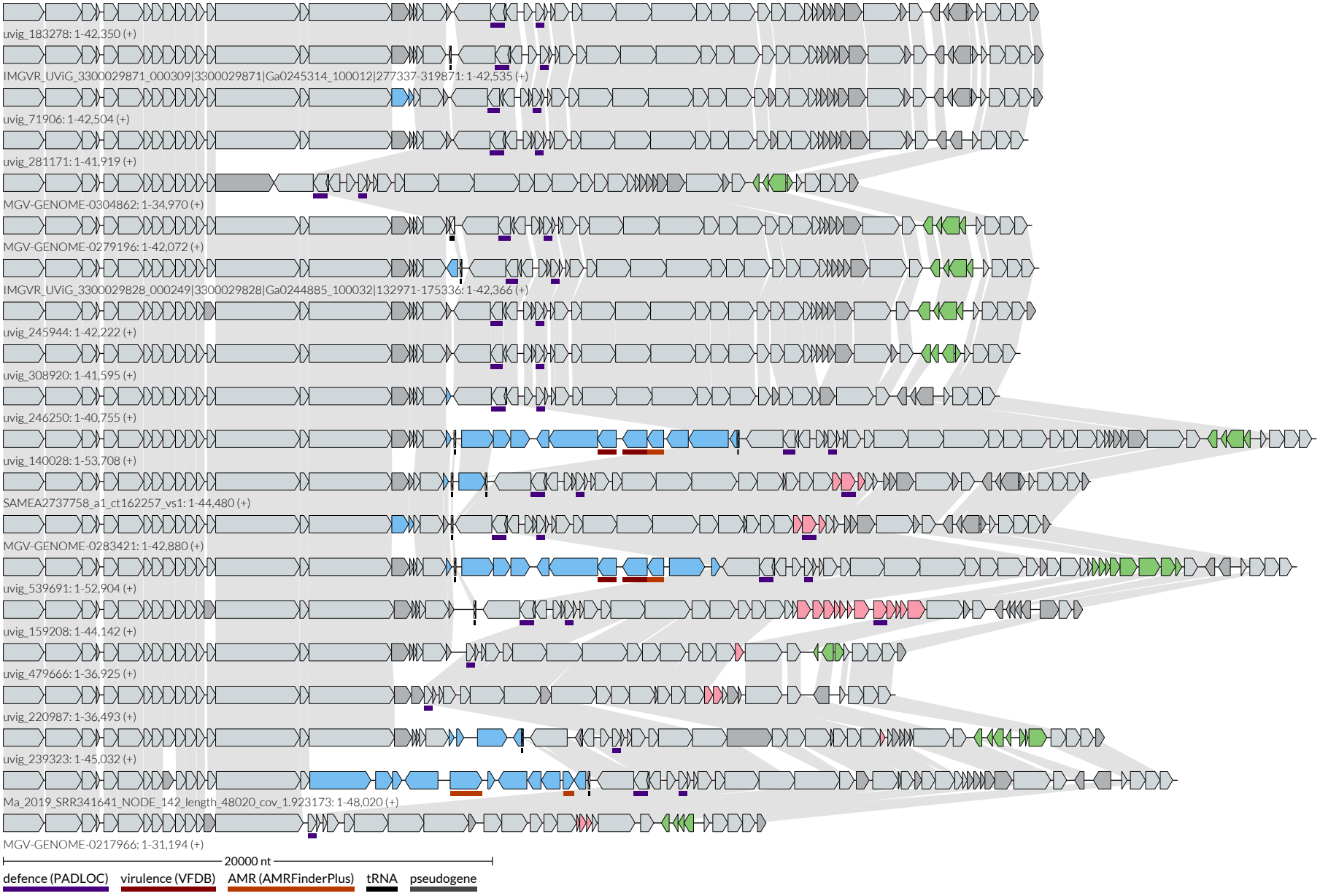
uvig_183278: 1-42,350 (+)
IMGVR_UViG_3300029871_000309|3300029871|Ga0245314_100012|277337-319871: 1-42,535 (+)
uvig_71906: 1-42,504 (+)
uvig_281171: 1-41,919 (+)
MGV-GENOME-0304862: 1-34,970 (+)
MGV-GENOME-0279196: 1-42,072 (+)
IMGVR_UViG_3300029828_000249|3300029828|Ga0244885_100032|132971-175336: 1-42,366 (+)
uvig_245944: 1-42,222 (+)
uvig_308920: 1-41,595 (+)
uvig_246250: 1-40,755 (+)
uvig_140028: 1-53,708 (+)
SAMEA2737758_a1_ct162257_vs1: 1-44,480 (+)
MGV-GENOME-0283421: 1-42,880 (+)
uvig_539691: 1-52,904 (+)
uvig_159208: 1-44,142 (+)
uvig_479666: 1-36,925 (+)
uvig_220987: 1-36,493 (+)
uvig_239323: 1-45,032 (+)
Ma_2019_SRR341641_NODE_142_length_48020_cov_1.923173: 1-48,020 (+)
MGV-GENOME-0217966: 1-31,194 (+)
20000 nt
defence (PADLOC)
virulence (VFDB)
AMR (AMRFinderPlus)
tRNA
pseudogene