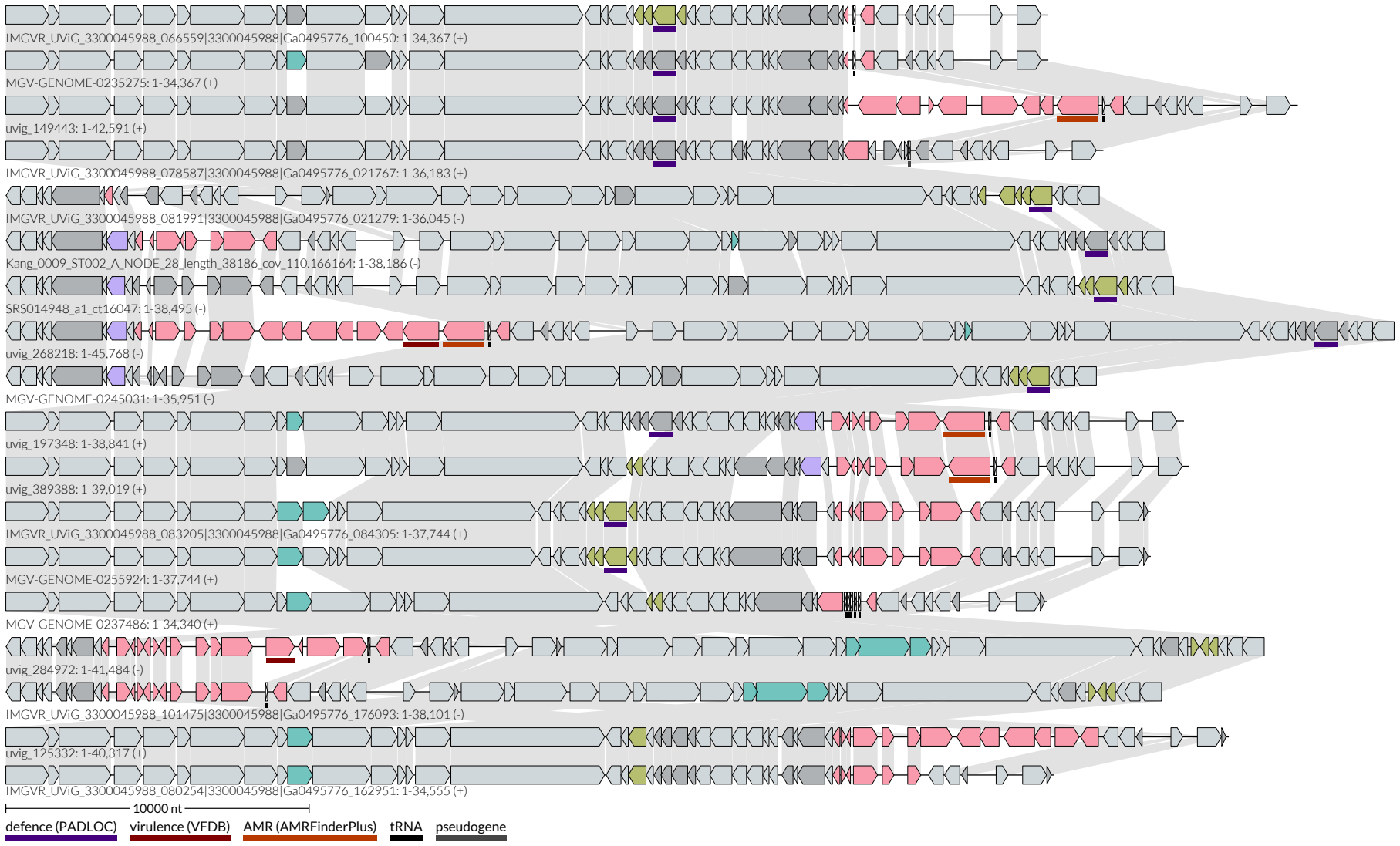
IMGVR_UViG_3300045988_066559|3300045988|Ga0495776_100450: 1-34,367 (+)
MGV-GENOME-0235275: 1-34,367 (+)
uvig_149443: 1-42,591 (+)
IMGVR_UViG_3300045988_078587|3300045988|Ga0495776_021767: 1-36,183 (+)
IMGVR_UViG_3300045988_081991|3300045988|Ga0495776_021279: 1-36,045 (-)
Kang_0009_ST002_A_NODE_28_length_38186_cov_110.166164: 1-38,186 (-)
SRS014948_a1_ct16047: 1-38,495 (-)
uvig_268218: 1-45,768 (-)
MGV-GENOME-0245031: 1-35,951 (-)
uvig_197348: 1-38,841 (+)
uvig_389388: 1-39,019 (+)
IMGVR_UViG_3300045988_083205|3300045988|Ga0495776_084305: 1-37,744 (+)
MGV-GENOME-0255924: 1-37,744 (+)
MGV-GENOME-0237486: 1-34,340 (+)
uvig_284972: 1-41,484 (-)
IMGVR_UViG_3300045988_101475|3300045988|Ga0495776_176093: 1-38,101 (-)
uvig_125332: 1-40,317 (+)
IMGVR_UViG_3300045988_080254|3300045988|Ga0495776_162951: 1-34,555 (+)
10000 nt
defence (PADLOC)
virulence (VFDB)
AMR (AMRFinderPlus)
tRNA
pseudogene