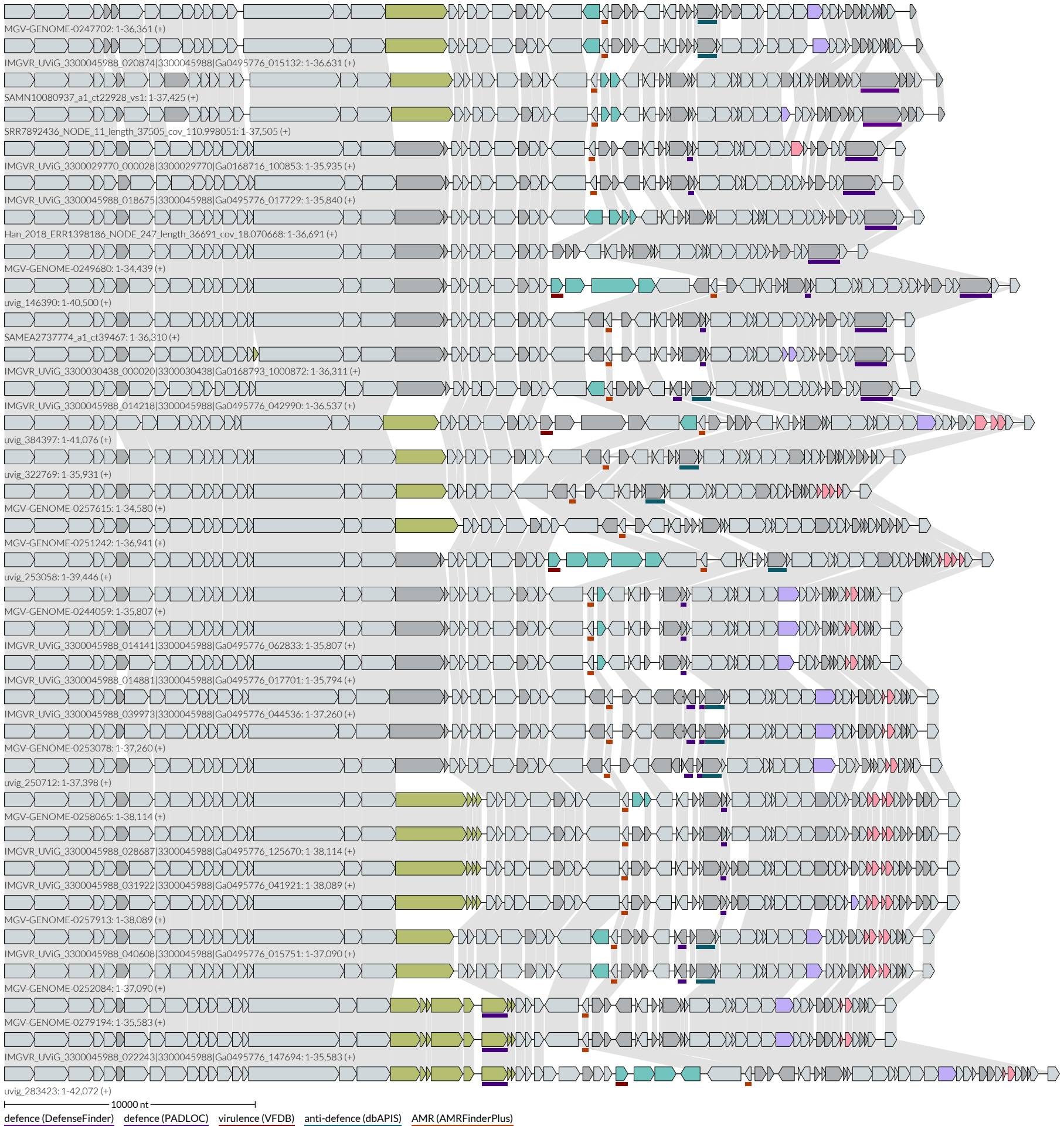
MGV-GENOME-0247702: 1-36,361 (+)
IMGVR_UViG_3300045988_020874|3300045988|Ga0495776_015132: 1-36,631 (+)
SAMN10080937_a1_ct22928_vs1: 1-37,425 (+)
SRR7892436_NODE_11_length_37505_cov_110.998051: 1-37,505 (+)
IMGVR_UViG_3300029770_000028|3300029770|Ga0168716_100853: 1-35,935 (+)
IMGVR_UViG_3300045988_018675|3300045988|Ga0495776_017729: 1-35,840 (+)
Han_2018_ERR1398186_NODE_247_length_36691_cov_18.070668: 1-36,691 (+)
MGV-GENOME-0249680: 1-34,439 (+)
uvig_146390: 1-40,500 (+)
SAMEA2737774_a1_ct39467: 1-36,310 (+)
IMGVR_UViG_3300030438_000020|3300030438|Ga0168793_1000872: 1-36,311 (+)
IMGVR_UViG_3300045988_014218|3300045988|Ga0495776_042990: 1-36,537 (+)
uvig_384397: 1-41,076 (+)
uvig_322769: 1-35,931 (+)
MGV-GENOME-0257615: 1-34,580 (+)
MGV-GENOME-0251242: 1-36,941 (+)
uvig_253058: 1-39,446 (+)
MGV-GENOME-0244059: 1-35,807 (+)
IMGVR_UViG_3300045988_014141|3300045988|Ga0495776_062833: 1-35,807 (+)
IMGVR_UViG_3300045988_014881|3300045988|Ga0495776_017701: 1-35,794 (+)
IMGVR_UViG_3300045988_039973|3300045988|Ga0495776_044536: 1-37,260 (+)
MGV-GENOME-0253078: 1-37,260 (+)
uvig_250712: 1-37,398 (+)
MGV-GENOME-0258065: 1-38,114 (+)
IMGVR_UViG_3300045988_028687|3300045988|Ga0495776_125670: 1-38,114 (+)
IMGVR_UViG_3300045988_031922|3300045988|Ga0495776_041921: 1-38,089 (+)
MGV-GENOME-0257913: 1-38,089 (+)
IMGVR_UViG_3300045988_040608|3300045988|Ga0495776_015751: 1-37,090 (+)
MGV-GENOME-0252084: 1-37,090 (+)
MGV-GENOME-0279194: 1-35,583 (+)
IMGVR_UViG_3300045988_022243|3300045988|Ga0495776_147694: 1-35,583 (+)
uvig_283423: 1-42,072 (+)
10000 nt
defence (DefenseFinder)
defence (PADLOC)
virulence (VFDB)
anti-defence (dbAPIS)
AMR (AMRFinderPlus)