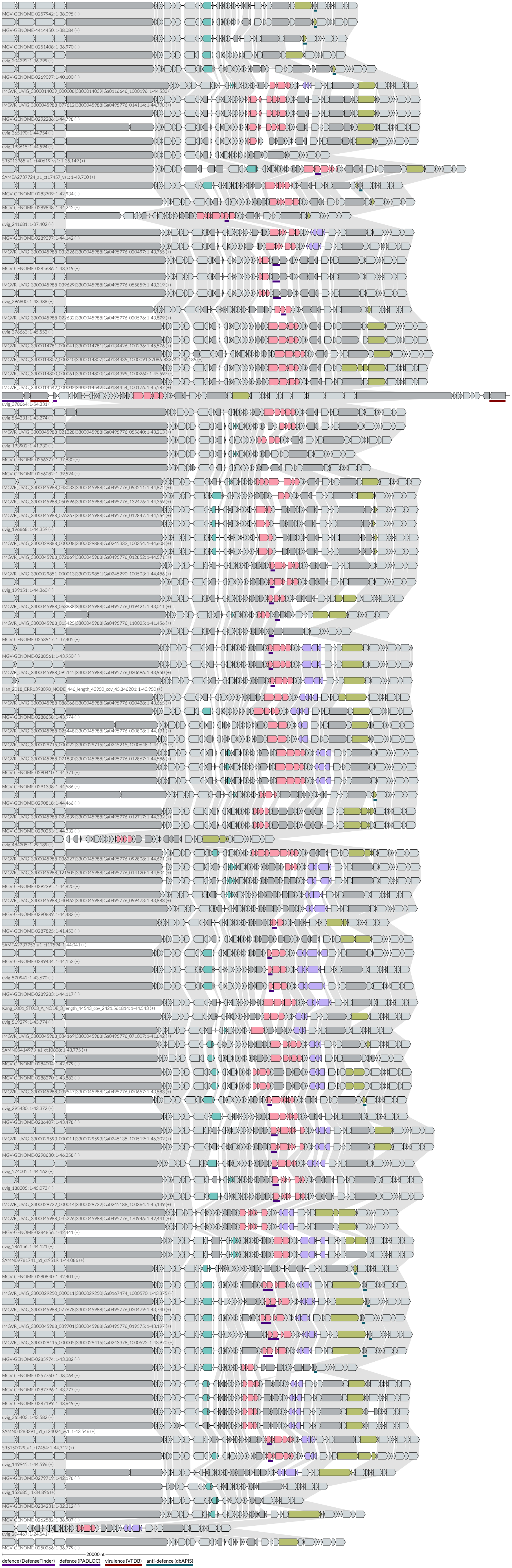
MGV-GENOME-0257942: 1-38,095 (+)
MGV-GENOME-4414450: 1-38,084 (+)
MGV-GENOME-0251408: 1-36,970 (+)
uvig_204292: 1-36,799 (+)
MGV-GENOME-0269097: 1-40,100 (+)
IMGVR_UViG_3300014039_000008|3300014039|Ga0116646_1000196: 1-44,533 (+)
IMGVR_UViG_3300045988_077612|3300045988|Ga0495776_014114: 1-44,798 (+)
MGV-GENOME-0292286: 1-44,798 (+)
uvig_365190: 1-44,754 (+)
uvig_193615: 1-44,594 (+)
SRS013965_a1_ct40619_vs1: 1-35,149 (+)
SAMEA2737724_a1_ct17457_vs1: 1-49,700 (+)
MGV-GENOME-0283709: 1-42,934 (+)
MGV-GENOME-0289848: 1-44,242 (+)
uvig_241681: 1-37,402 (+)
MGV-GENOME-0289397: 1-44,142 (+)
IMGVR_UViG_3300045988_033226|3300045988|Ga0495776_020497: 1-43,755 (+)
MGV-GENOME-0285686: 1-43,319 (+)
IMGVR_UViG_3300045988_039629|3300045988|Ga0495776_055859: 1-43,319 (+)
uvig_296800: 1-43,388 (+)
IMGVR_UViG_3300045988_022632|3300045988|Ga0495776_020576: 1-43,879 (+)
uvig_376663: 1-45,552 (+)
IMGVR_UViG_3300014781_000041|3300014781|Ga0134426_100236: 1-45,576 (+)
IMGVR_UViG_3300014807_000240|3300014807|Ga0134439_1000091|37086-83274: 1-46,189 (+)
IMGVR_UViG_3300014800_000061|3300014800|Ga0134399_1000260: 1-45,597 (+)
IMGVR_UViG_3300014542_000002|3300014542|Ga0134454_100176: 1-45,587 (+)
uvig_378664: 1-54,331 (+)
uvig_554331: 1-43,274 (+)
IMGVR_UViG_3300045988_021328|3300045988|Ga0495776_055640: 1-43,213 (+)
uvig_193902: 1-41,730 (+)
MGV-GENOME-0256377: 1-37,830 (+)
MGV-GENOME-0266082: 1-39,524 (+)
IMGVR_UViG_3300045988_043033|3300045988|Ga0495776_093211: 1-44,872 (+)
IMGVR_UViG_3300045988_050596|3300045988|Ga0495776_132476: 1-44,359 (+)
IMGVR_UViG_3300045988_076267|3300045988|Ga0495776_012847: 1-44,564 (+)
uvig_196868: 1-44,359 (+)
IMGVR_UViG_3300029888_000008|3300029888|Ga0245333_100354: 1-44,608 (+)
IMGVR_UViG_3300045988_072869|3300045988|Ga0495776_012852: 1-44,571 (+)
IMGVR_UViG_3300029851_000013|3300029851|Ga0245290_100503: 1-44,486 (+)
uvig_199151: 1-44,360 (+)
IMGVR_UViG_3300045988_063888|3300045988|Ga0495776_019421: 1-43,011 (+)
IMGVR_UViG_3300045988_015425|3300045988|Ga0495776_110025: 1-41,456 (+)
MGV-GENOME-0253917: 1-37,405 (+)
MGV-GENOME-0288561: 1-43,950 (+)
IMGVR_UViG_3300045988_095145|3300045988|Ga0495776_020696: 1-43,950 (+)
Han_2018_ERR1398098_NODE_446_length_43950_cov_45.846201: 1-43,950 (+)
IMGVR_UViG_3300045988_088066|3300045988|Ga0495776_020428: 1-43,665 (+)
MGV-GENOME-0288658: 1-43,974 (+)
IMGVR_UViG_3300045988_025448|3300045988|Ga0495776_020808: 1-44,131 (+)
IMGVR_UViG_3300029715_000022|3300029715|Ga0245215_1000648: 1-44,175 (+)
IMGVR_UViG_3300045988_071830|3300045988|Ga0495776_012867: 1-44,586 (+)
MGV-GENOME-0290410: 1-44,371 (+)
MGV-GENOME-0291338: 1-44,586 (+)
MGV-GENOME-0290818: 1-44,466 (+)
IMGVR_UViG_3300045988_022639|3300045988|Ga0495776_012717: 1-44,332 (+)
MGV-GENOME-0290253: 1-44,332 (+)
uvig_484205: 1-29,189 (+)
IMGVR_UViG_3300045988_036227|3300045988|Ga0495776_092808: 1-44,671 (+)
IMGVR_UViG_3300045988_121505|3300045988|Ga0495776_014120: 1-44,804 (+)
MGV-GENOME-0292395: 1-44,820 (+)
IMGVR_UViG_3300045988_040462|3300045988|Ga0495776_099473: 1-43,883 (+)
MGV-GENOME-0290889: 1-44,482 (+)
MGV-GENOME-0287825: 1-41,453 (+)
SAMEA2737753_a1_ct17594: 1-44,041 (+)
MGV-GENOME-0289434: 1-44,152 (+)
uvig_570942: 1-43,670 (+)
MGV-GENOME-0289283: 1-44,117 (+)
Kang_0001_ST003_A_NODE_3_length_44543_cov_2421.561814: 1-44,543 (+)
uvig_519279: 1-43,774 (+)
IMGVR_UViG_3300045988_034169|3300045988|Ga0495776_071007: 1-41,842 (+)
SAMN05414973_a1_ct10808: 1-43,775 (+)
MGV-GENOME-0284004: 1-42,979 (+)
MGV-GENOME-0288270: 1-43,883 (+)
IMGVR_UViG_3300045988_039547|3300045988|Ga0495776_020657: 1-43,883 (+)
uvig_295430: 1-43,372 (+)
MGV-GENOME-0286407: 1-43,478 (+)
IMGVR_UViG_3300029593_000011|3300029593|Ga0245135_100519: 1-46,302 (+)
MGV-GENOME-0298630: 1-46,258 (+)
uvig_574005: 1-44,162 (+)
uvig_188305: 1-45,073 (+)
IMGVR_UViG_3300029722_000014|3300029722|Ga0245188_100364: 1-45,139 (+)
IMGVR_UViG_3300045988_045226|3300045988|Ga0495776_170946: 1-42,441 (+)
MGV-GENOME-0284856: 1-42,441 (+)
uvig_586156: 1-44,121 (+)
SAMN09781741_a1_ct9519: 1-44,086 (+)
MGV-GENOME-0280840: 1-42,401 (+)
IMGVR_UViG_3300029250_000011|3300029250|Ga0167474_1000570: 1-43,375 (+)
IMGVR_UViG_3300045988_077678|3300045988|Ga0495776_020479: 1-43,740 (+)
IMGVR_UViG_3300045988_039701|3300045988|Ga0495776_019575: 1-43,197 (+)
IMGVR_UViG_3300029415_000005|3300029415|Ga0243378_1000522: 1-43,970 (+)
MGV-GENOME-0285974: 1-43,382 (+)
MGV-GENOME-0257760: 1-38,064 (+)
MGV-GENOME-0287796: 1-43,777 (+)
MGV-GENOME-0287199: 1-43,649 (+)
uvig_365403: 1-43,582 (+)
SAMN03283291_a1_ct24024_vs1: 1-43,546 (+)
SRS150029_a1_ct7454: 1-44,712 (+)
uvig_149945: 1-44,596 (+)
MGV-GENOME-0279719: 1-42,178 (+)
uvig_152685_: 1-34,896 (+)
MGV-GENOME-0234231: 1-32,312 (+)
MGV-GENOME-0262582: 1-38,907 (+)
uvig_204467: 1-24,541 (+)
MGV-GENOME-0250266: 1-36,779 (+)
20000 nt
defence (DefenseFinder)
defence (PADLOC)
virulence (VFDB)
anti-defence (dbAPIS)