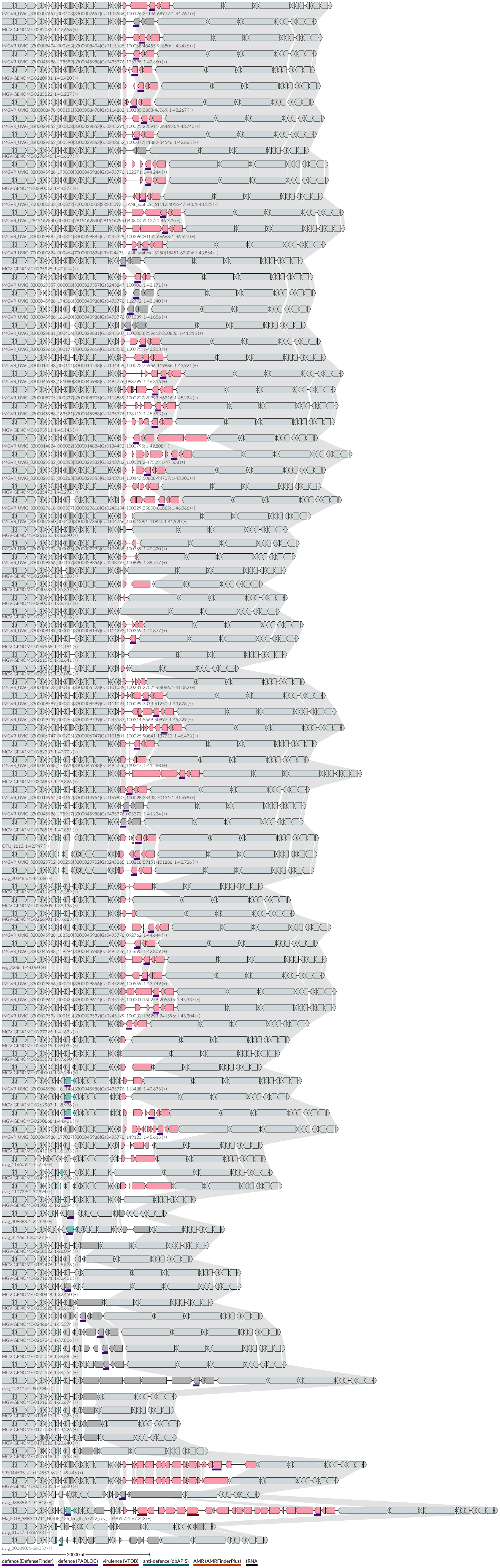
IMGVR_UViG_3300007657_000403|3300007657|Ga0105536_100126|24146-68912: 1-44,767 (+)
MGV-GENOME-0282085: 1-42,658 (+)
IMGVR_UViG_3300008404_000263|3300008404|Ga0115185_100068|48455-91880: 1-43,426 (+)
IMGVR_UViG_3300045988_178599|3300045988|Ga0495776_135298: 1-42,660 (+)
MGV-GENOME-0280931: 1-42,420 (+)
MGV-GENOME-0285322: 1-43,237 (+)
IMGVR_UViG_3300008478_000151|3300008478|Ga0114882_100230|3803-46069: 1-42,267 (+)
IMGVR_UViG_3300029852_000308|3300029852|Ga0245291_100025|220911-264650: 1-43,740 (+)
IMGVR_UViG_3300029362_000590|3300029362|Ga0243852_1000077|11882-54546: 1-42,665 (+)
MGV-GENOME-0276945: 1-41,619 (+)
IMGVR_UViG_3300045988_179800|3300045988|Ga0495776_132271: 1-44,244 (+)
MGV-GENOME-0290012: 1-44,277 (+)
IMGVR_UViG_7000000333_000372|7000000333|SRS050925_LANL_scaffold_61112|4016-47540: 1-43,525 (+)
IMGVR_UViG_2915162840_000005|2915162840|2915162961|43803-90127: 1-46,325 (+)
IMGVR_UViG_3300029885_000314|3300029885|Ga0245329_100296|20142-66668: 1-46,527 (+)
IMGVR_UViG_7000000624_000464|7000000624|SRS024435_LANL_scaffold_5250|18451-62304: 1-43,854 (+)
MGV-GENOME-0295925: 1-40,814 (+)
IMGVR_UViG_3300029357_000008|3300029357|Ga0243847_100906: 1-42,171 (+)
IMGVR_UViG_3300045988_174568|3300045988|Ga0495776_112772: 1-42,540 (+)
IMGVR_UViG_3300045988_161430|3300045988|Ga0495776_071039: 1-41,856 (+)
IMGVR_UViG_3300029881_000406|3300029881|Ga0245307_1000003|259612-300826: 1-41,215 (+)
IMGVR_UViG_3300029616_000177|3300029616|Ga0245133_100775: 1-43,203 (+)
IMGVR_UViG_3300014548_000111|3300014548|Ga0134459_100022|75966-119886: 1-43,921 (+)
IMGVR_UViG_3300045988_181080|3300045988|Ga0495776_098799: 1-46,316 (+)
IMGVR_UViG_3300008705_000337|3300008705|Ga0113869_1000227|20993-66216: 1-45,224 (+)
IMGVR_UViG_3300045988_169521|3300045988|Ga0495776_138113: 1-45,593 (+)
MGV-GENOME-0293911: 1-45,145 (+)
IMGVR_UViG_3300014824_000022|3300014824|Ga0134492_1000795: 1-42,808 (+)
IMGVR_UViG_3300029332_000353|3300029332|Ga0243782_100521|2-47509: 1-47,508 (+)
IMGVR_UViG_3300029355_000263|3300029355|Ga0243784_100143|50808-94707: 1-43,900 (+)
MGV-GENOME-0285473: 1-43,272 (+)
IMGVR_UViG_3300029618_000307|3300029618|Ga0245134_100229|35800-81865: 1-46,066 (+)
IMGVR_UViG_3300007360_000400|3300007360|Ga0104766_1000129|1-41920: 1-41,920 (+)
MGV-GENOME-0261350: 1-38,690 (+)
IMGVR_UViG_3300007793_000025|3300007793|Ga0105888_100759: 1-40,320 (+)
IMGVR_UViG_3300029356_000137|3300029356|Ga0243797_100899: 1-39,777 (+)
MGV-GENOME-0260443: 1-38,528 (+)
MGV-GENOME-0240783: 1-35,337 (+)
MGV-GENOME-0290087: 1-38,737 (+)
MGV-GENOME-0273150: 1-37,616 (+)
IMGVR_UViG_3300008149_000014|3300008149|Ga0114293_100369: 1-40,877 (+)
MGV-GENOME-0269568: 1-40,191 (+)
MGV-GENOME-0261075: 1-38,641 (+)
MGV-GENOME-0223053: 1-32,074 (+)
IMGVR_UViG_3300006523_000165|3300006523|Ga0101039_100231|27025-68086: 1-41,062 (+)
IMGVR_UViG_3300008599_000313|3300008599|Ga0111091_100099|7373-51250: 1-43,878 (+)
IMGVR_UViG_3300029739_000265|3300029739|Ga0245187_100314|5669-50997: 1-45,329 (+)
IMGVR_UViG_3300006747_000205|3300006747|Ga0101801_100025|90841-137313: 1-46,473 (+)
MGV-GENOME-0282337: 1-42,703 (+)
IMGVR_UViG_3300045988_174970|3300045988|Ga0495776_110347: 1-41,588 (+)
MGV-GENOME-0306857: 1-48,826 (+)
IMGVR_UViG_3300014954_000152|3300014954|Ga0169837_100098|28433-70131: 1-41,699 (+)
IMGVR_UViG_3300045988_175917|3300045988|Ga0495776_025372: 1-41,234 (+)
MGV-GENOME-0298515: 1-40,851 (+)
OTU_1613: 1-42,947 (+)
IMGVR_UViG_3300029702_000258|3300029702|Ga0245265_100218|59151-101886: 1-42,736 (+)
uvig_203485: 1-42,338 (+)
MGV-GENOME-0241120: 1-35,389 (+)
MGV-GENOME-0263909: 1-39,138 (+)
MGV-GENOME-0266901: 1-39,683 (+)
IMGVR_UViG_3300045988_182560|3300045988|Ga0495776_092762: 1-44,644 (+)
IMGVR_UViG_3300045988_159294|3300045988|Ga0495776_135690: 1-42,809 (+)
ivig_3286: 1-44,060 (+)
IMGVR_UViG_3300029856_000210|3300029856|Ga0245296_100569: 1-43,749 (+)
IMGVR_UViG_3300029614_000325|3300029614|Ga0245155_100001|160279-205615: 1-45,337 (+)
IMGVR_UViG_3300029592_000163|3300029592|Ga0245129_100032|196293-241596: 1-45,304 (+)
MGV-GENOME-0277226: 1-41,673 (+)
MGV-GENOME-0263319: 1-39,035 (+)
MGV-GENOME-0255591: 1-37,690 (+)
MGV-GENOME-0240101: 1-35,240 (+)
IMGVR_UViG_3300045988_181146|3300045988|Ga0495776_113438: 1-40,675 (+)
MGV-GENOME-0262987: 1-38,976 (+)
MGV-GENOME-0290668: 1-44,431 (+)
IMGVR_UViG_3300045988_177027|3300045988|Ga0495776_149123: 1-45,615 (+)
MGV-GENOME-0241019: 1-35,375 (+)
uvig_116609: 1-35,778 (+)
MGV-GENOME-0249713: 1-36,696 (+)
uvig_110729: 1-37,994 (+)
MGV-GENOME-0190250: 1-26,294 (+)
uvig_409388: 1-22,328 (+)
uvig_45166: 1-30,227 (+)
MGV-GENOME-0208522: 1-28,094 (+)
MGV-GENOME-0192476: 1-25,876 (+)
MGV-GENOME-0271891: 1-32,401 (+)
MGV-GENOME-0240444: 1-32,450 (+)
MGV-GENOME-0203628: 1-28,617 (+)
MGV-GENOME-0246843: 1-35,374 (+)
MGV-GENOME-0267343: 1-37,806 (+)
MGV-GENOME-0275048: 1-38,385 (+)
MGV-GENOME-0275576: 1-38,554 (+)
uvig_122104: 1-50,794 (+)
MGV-GENOME-0191655: 1-23,674 (+)
MGV-GENOME-0170913: 1-23,325 (+)
MGV-GENOME-0177033: 1-24,226 (+)
MGV-GENOME-0191226: 1-23,640 (+)
MGV-GENOME-0209418: 1-27,953 (+)
SRS044535_a1_ct14552_vs2: 1-49,446 (+)
MGV-GENOME-0317335: 1-45,630 (+)
uvig_389899: 1-34,942 (+)
Ma_2019_SRR341711_NODE_126_length_67222_cov_5.318907: 1-67,222 (+)
uvig_81017: 1-28,993 (+)
uvig_200810: 1-36,057 (+)
20000 nt
defence (DefenseFinder)
defence (PADLOC)
virulence (VFDB)
anti-defence (dbAPIS)
AMR (AMRFinderPlus)
tRNA