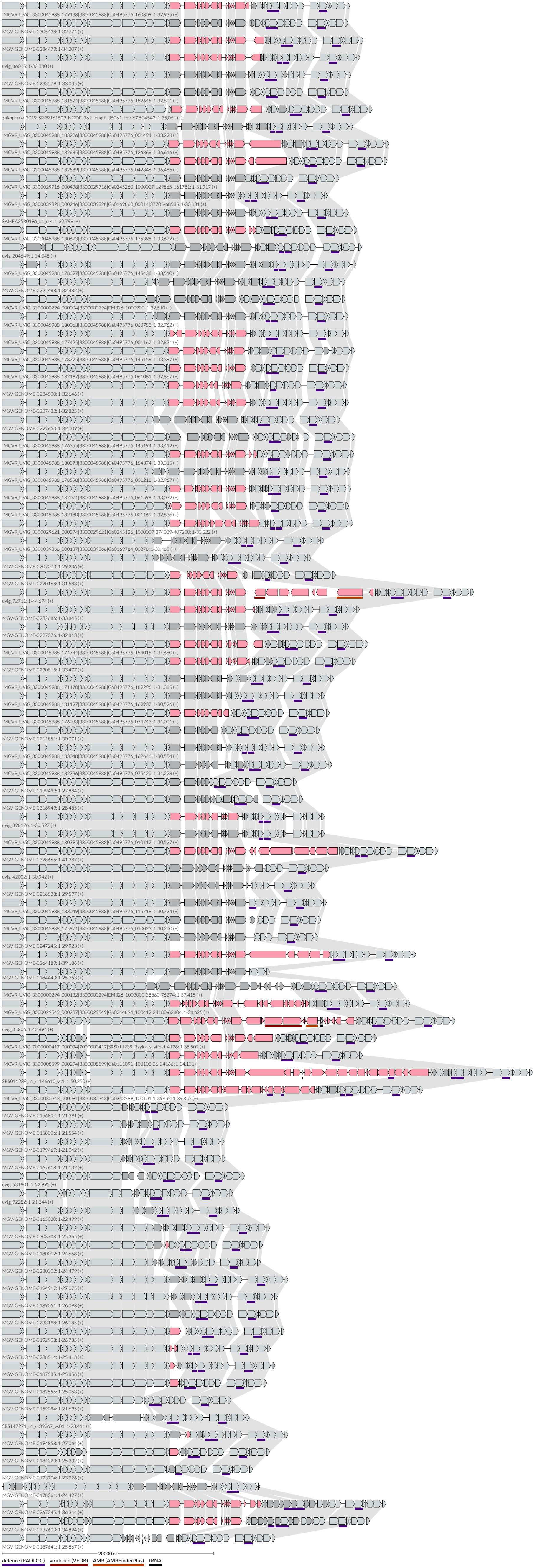
IMGVR_UViG_3300045988_179138|3300045988|Ga0495776_160809: 1-32,935 (+)
MGV-GENOME-0305438: 1-32,774 (+)
MGV-GENOME-0234479: 1-34,207 (+)
uvig_86015: 1-33,880 (+)
MGV-GENOME-0233579: 1-33,035 (+)
IMGVR_UViG_3300045988_181574|3300045988|Ga0495776_182645: 1-32,801 (+)
Shkoporov_2019_SRR9161509_NODE_362_length_35061_cov_67.504542: 1-35,061 (+)
IMGVR_UViG_3300045988_183226|3300045988|Ga0495776_001494: 1-33,228 (+)
IMGVR_UViG_3300045988_182685|3300045988|Ga0495776_126868: 1-36,616 (+)
IMGVR_UViG_3300045988_182589|3300045988|Ga0495776_042846: 1-36,485 (+)
IMGVR_UViG_3300029716_000498|3300029716|Ga0245260_1000027|129865-161781: 1-31,917 (+)
IMGVR_UViG_3300039328_000246|3300039328|Ga0169860_00014|37705-68535: 1-30,831 (+)
SAMEA2580196_b1_ct4: 1-32,798 (+)
IMGVR_UViG_3300045988_180673|3300045988|Ga0495776_175398: 1-33,622 (+)
uvig_204649: 1-34,048 (+)
IMGVR_UViG_3300045988_178697|3300045988|Ga0495776_145436: 1-33,510 (+)
MGV-GENOME-0225488: 1-32,482 (+)
IMGVR_UViG_3300000294_000004|3300000294|EM326_1000900: 1-32,510 (+)
IMGVR_UViG_3300045988_180063|3300045988|Ga0495776_060758: 1-32,762 (+)
IMGVR_UViG_3300045988_177425|3300045988|Ga0495776_001167: 1-32,831 (+)
IMGVR_UViG_3300045988_178225|3300045988|Ga0495776_145159: 1-33,397 (+)
IMGVR_UViG_3300045988_182197|3300045988|Ga0495776_061081: 1-32,867 (+)
MGV-GENOME-0234500: 1-32,646 (+)
MGV-GENOME-0227432: 1-32,825 (+)
MGV-GENOME-0222653: 1-32,009 (+)
IMGVR_UViG_3300045988_176355|3300045988|Ga0495776_145194: 1-33,412 (+)
IMGVR_UViG_3300045988_180373|3300045988|Ga0495776_154374: 1-33,315 (+)
IMGVR_UViG_3300045988_178598|3300045988|Ga0495776_001218: 1-32,967 (+)
IMGVR_UViG_3300045988_182071|3300045988|Ga0495776_061598: 1-33,032 (+)
IMGVR_UViG_3300045988_182180|3300045988|Ga0495776_001169: 1-32,836 (+)
IMGVR_UViG_3300029621_000374|3300029621|Ga0245126_1000007|374029-407250: 1-33,222 (+)
IMGVR_UViG_3300039366_000137|3300039366|Ga0169784_00278: 1-30,465 (+)
MGV-GENOME-0207073: 1-29,236 (+)
MGV-GENOME-0220168: 1-31,583 (+)
uvig_72711: 1-44,674 (+)
MGV-GENOME-0232686: 1-33,845 (+)
MGV-GENOME-0227376: 1-32,813 (+)
IMGVR_UViG_3300045988_174744|3300045988|Ga0495776_154015: 1-34,660 (+)
MGV-GENOME-0230818: 1-33,477 (+)
IMGVR_UViG_3300045988_171170|3300045988|Ga0495776_189296: 1-31,385 (+)
IMGVR_UViG_3300045988_181197|3300045988|Ga0495776_169937: 1-30,526 (+)
IMGVR_UViG_3300045988_176033|3300045988|Ga0495776_074743: 1-31,001 (+)
MGV-GENOME-0211851: 1-30,071 (+)
IMGVR_UViG_3300045988_183048|3300045988|Ga0495776_162646: 1-30,554 (+)
IMGVR_UViG_3300045988_182736|3300045988|Ga0495776_075420: 1-31,228 (+)
MGV-GENOME-0199499: 1-27,884 (+)
MGV-GENOME-0316949: 1-28,485 (+)
uvig_398176: 1-30,527 (+)
IMGVR_UViG_3300045988_180395|3300045988|Ga0495776_010117: 1-30,527 (+)
MGV-GENOME-0328665: 1-41,287 (+)
uvig_42002: 1-30,942 (+)
MGV-GENOME-0216528: 1-29,597 (+)
IMGVR_UViG_3300045988_183049|3300045988|Ga0495776_115718: 1-30,724 (+)
IMGVR_UViG_3300045988_175871|3300045988|Ga0495776_010023: 1-30,200 (+)
MGV-GENOME-0247245: 1-29,923 (+)
MGV-GENOME-0264189: 1-39,186 (+)
MGV-GENOME-0184443: 1-25,353 (+)
IMGVR_UViG_3300000294_000132|3300000294|EM326_1003000|38860-76274: 1-37,415 (+)
IMGVR_UViG_3300029549_000237|3300029549|Ga0244894_100412|24180-62804: 1-38,625 (+)
uvig_35806: 1-42,894 (+)
IMGVR_UViG_7000000417_000094|7000000417|SRS011239_Baylor_scaffold_4178: 1-35,502 (+)
IMGVR_UViG_3300008599_000294|3300008599|Ga0111091_100108|36-34166: 1-34,131 (+)
SRS011239_a1_ct146610_vs1: 1-50,250 (+)
IMGVR_UViG_3300030343_000091|3300030343|Ga0243299_100101|1-39852: 1-39,852 (+)
MGV-GENOME-0156804: 1-21,391 (+)
MGV-GENOME-0158006: 1-21,554 (+)
MGV-GENOME-0179467: 1-21,042 (+)
MGV-GENOME-0167618: 1-21,132 (+)
uvig_531901: 1-22,995 (+)
uvig_92282: 1-21,844 (+)
MGV-GENOME-0165020: 1-22,499 (+)
MGV-GENOME-0303708: 1-25,365 (+)
MGV-GENOME-0180012: 1-24,668 (+)
MGV-GENOME-0230302: 1-24,479 (+)
MGV-GENOME-0194917: 1-27,075 (+)
MGV-GENOME-0189051: 1-26,093 (+)
MGV-GENOME-0233198: 1-26,185 (+)
MGV-GENOME-0192908: 1-26,735 (+)
MGV-GENOME-0238514: 1-25,413 (+)
MGV-GENOME-0187585: 1-25,856 (+)
MGV-GENOME-0182556: 1-25,063 (+)
MGV-GENOME-0159094: 1-21,695 (+)
SRS147271_a1_ct39267_vs01: 1-23,411 (+)
MGV-GENOME-0194858: 1-27,064 (+)
MGV-GENOME-0184323: 1-25,332 (+)
MGV-GENOME-0173704: 1-23,726 (+)
MGV-GENOME-0178361: 1-24,427 (+)
MGV-GENOME-0267245: 1-36,344 (+)
MGV-GENOME-0237603: 1-34,824 (+)
MGV-GENOME-0187641: 1-25,867 (+)
20000 nt
defence (PADLOC)
virulence (VFDB)
AMR (AMRFinderPlus)
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