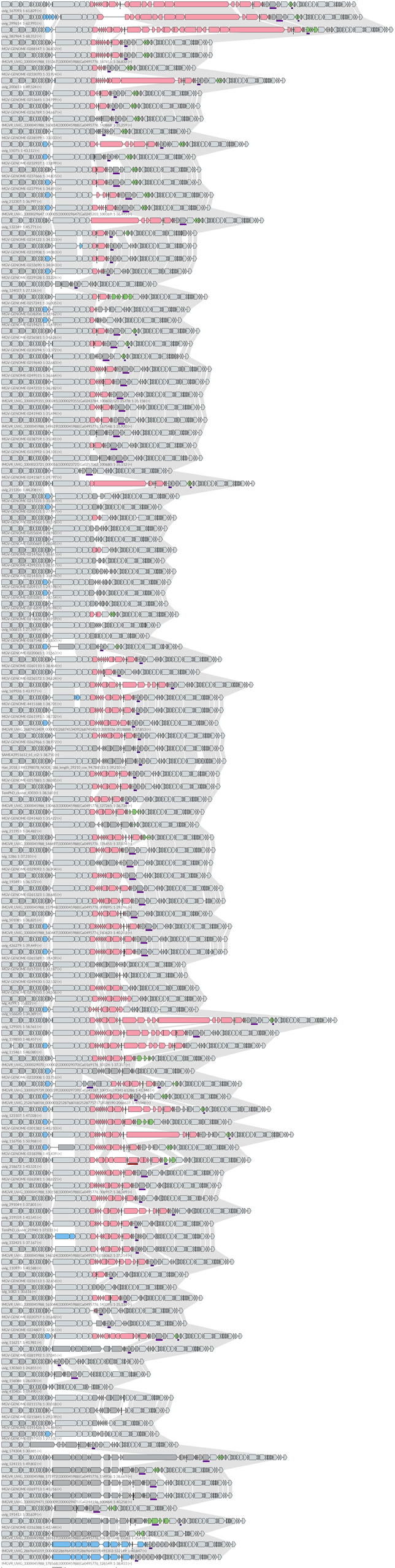
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scaffold_27960:1-62,995 (+)
scaffold_382964:1-68,312 (+)
MGV-GENOME-0268147:1-36,812 (+)
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MGV-GENOME-0233070:1-33,914 (+)
scaffold_200653:1-49,524 (+)
MGV-GENOME-0253665:1-34,999 (+)
MGV-GENOME-0236789:1-34,671 (+)
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MGV-GENOME-0228099:1-33,833 (+)
scaffold_55075:1-43,113 (+)
MGV-GENOME-0229597:1-33,899 (+)
MGV-GENOME-0237666:1-34,815 (+)
MGV-GENOME-0237954:1-34,881 (+)
scaffold_212307:1-36,997 (+)
IMGVR_UViG_3300029647_...:1-36,495 (+)
scaffold_13234:1-45,771 (+)
MGV-GENOME-0234122:1-34,113 (+)
MGV-GENOME-0233908:1-34,083 (+)
MGV-GENOME-0233690:1-34,041 (+)
MGV-GENOME-0229528:1-33,258 (+)
scaffold_124027:1-27,126 (+)
MGV-GENOME-0257241:1-36,805 (+)
MGV-GENOME-0258656:1-32,420 (+)
MGV-GENOME-0236581:1-34,626 (+)
MGV-GENOME-0230294:1-31,372 (+)
MGV-GENOME-0239640:1-32,483 (+)
MGV-GENOME-0249515:1-34,644 (+)
MGV-GENOME-0247233:1-34,782 (+)
IMGVR_UViG_3300029645:...:1-35,158 (+)
MGV-GENOME-0241940:1-31,494 (+)
IMGVR_UViG_3300049988:149:19:3300049988|Ga0495779_...:1-35,955 (+)
MGV-GENOME-0238759:1-31,411 (+)
MGV-GENOME-0233992:1-34,101 (+)
IMGVR_UViG_3300027271_...:1-35,122 (+)
MGV-GENOME-0241587:1-29,797 (+)
scaffold_211206:1-44,208 (+)
MGV-GENOME-0237227:1-31,679 (+)
MGV-GENOME-0250110:1-27,797 (+)
MGV-GENOME-0252552:1-30,381 (+)
MGV-GENOME-0260644:1-28,833 (+)
MGV-GENOME-0250366:1-30,383 (+)
MGV-GENOME-0241961:1-30,155 (+)
MGV-GENOME-0239551:1-28,170 (+)
MGV-GENOME-0241810:1-29,944 (+)
MGV-GENOME-0249177:1-31,780 (+)
MGV-GENOME-0250885:1-31,541 (+)
MGV-GENOME-0236587:1-29,539 (+)
scaffold_208815:1-29,755 (+)
MGV-GENOME-0238548:1-28,850 (+)
MGV-GENOME-0230063:1-31,563 (+)
MGV-GENOME-0260110:1-38,464 (+)
MGV-GENOME-0230372:1-34,824 (+)
scaffold_169926:1-43,917 (+)
MGV-GENOME-0445188:1-38,701 (+)
MGV-GENOME-0261591:1-38,732 (+)
IMGVR_...:3687453409:3687454540:3:3830106_3959808:... (+)
MGV-GENOME-0262966:1-38,972 (+)
SAMEA3951612.31_c0:1-38,758 (+)
Han_2018_1.fill39078_NODE_186_length_39210_cov_94.784820:1-39,210 (+)
MGV-GENOME-0257885:1-38,845 (+)
TemPhO_cluster_43010:1-38,160 (+)
IMGVR_UViG_3300049988:...:1-38,663 (+)
MGV-GENOME-0241460:1-35,422 (+)
scaffold_211951:1-34,480 (+)
IMGVR_UViG_3300045988_146:97:3300045988|Ga0495779_...:1-37,074 (+)
scaffold_1286:1-37,210 (+)
MGV-GENOME-0229092:1-36,934 (+)
scaffold_193489:1-36,572 (+)
MGV-GENOME-0261523:1-38,645 (+)
IMGVR_UViG_3300049988:...:1-39,076 (+)
scaffold_50108:1-36,825 (+)
IMGVR_UViG_3300049988:...:1-38,620 (+)
scaffold_426279:1-39,449 (+)
MGV-GENOME-0261589:1-39,439 (+)
MGV-GENOME-0257355:1-37,170 (+)
MGV-GENOME-0249430:1-32,132 (+)
MGV-GENOME-0230228:1-36,501 (+)
scaffold_47997:1-37,227 (+)
scaffold_55220:1-34,369 (+)
scaffold_129505:1-38,561 (+)
scaffold_159850:1-38,457 (+)
scaffold_115463:1-46,880 (+)
IMGVR_UViG_3300029070_...:1-37,157 (+)
MGV-GENOME-0232008:1-33,716 (+)
IMGVR_UViG_3300029739_...:1-35,193 (+)
IMGVR_UViG_2528768058_...:1-37,048 (+)
scaffold_123101:1-47,028 (+)
MGV-GENOME-0301382:1-40,310 (+)
scaffold_114706:1-50,968 (+)
MGV-GENOME-0318398:1-41,439 (+)
scaffold_218672:1-43,131 (+)
MGV-GENOME-0262081:1-38,622 (+)
IMGVR_UViG_3300049988:...:1-38,599 (+)
scaffold_291049:1-37,801 (+)
scaffold_319509:1-43,145 (+)
TemPhO_cluster_21940:1-37,031 (+)
scaffold_102421:1-37,167 (+)
IMGVR_UViG_3300045988_146:124:3300045988|Ga0495779_...:1-37,114 (+)
scaffold_119705:1-40,188 (+)
MGV-GENOME-0226113:1-32,614 (+)
IMGVR_UViG_3300049988:...:1-37,154 (+)
MGV-GENOME-0220757:1-31,642 (+)
MGV-GENOME-0249807:1-31,316 (+)
scaffold_116217:1-41,981 (+)
MGV-GENOME-0281992:1-37,041 (+)
scaffold_130360:1-24,855 (+)
scaffold_154088:1,063,930 (+)
scaffold_410300:1-39,490 (+)
MGV-GENOME-0211576:1-30,918 (+)
MGV-GENOME-0215845:1-29,339 (+)
MGV-GENOME-0197660:1-31,626 (+)
MGV-GENOME-0197559:1-27,537 (+)
scaffold_174304:1-30,001 (+)
scaffold_124115:1-49,852 (+)
IMGVR_UViG_3300049988:...:1-36,665 (+)
MGV-GENOME-0269713:1-40,156 (+)
IMGVR_UViG_3300029947_...:1-40,258 (+)
scaffold_191452:1-30,639 (+)
MGV-GENOME-0315388:1-42,644 (+)
IMGVR_UViG_3300049988:...:1-36,746 (+)
IMGVR_UViG_3300045988_178568:3300045988|Ga0495776_...:1-38,411 (+)
20000 nt
defence (DefenseFinder)
defence (PADLOC)
virulence (VFDB)
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