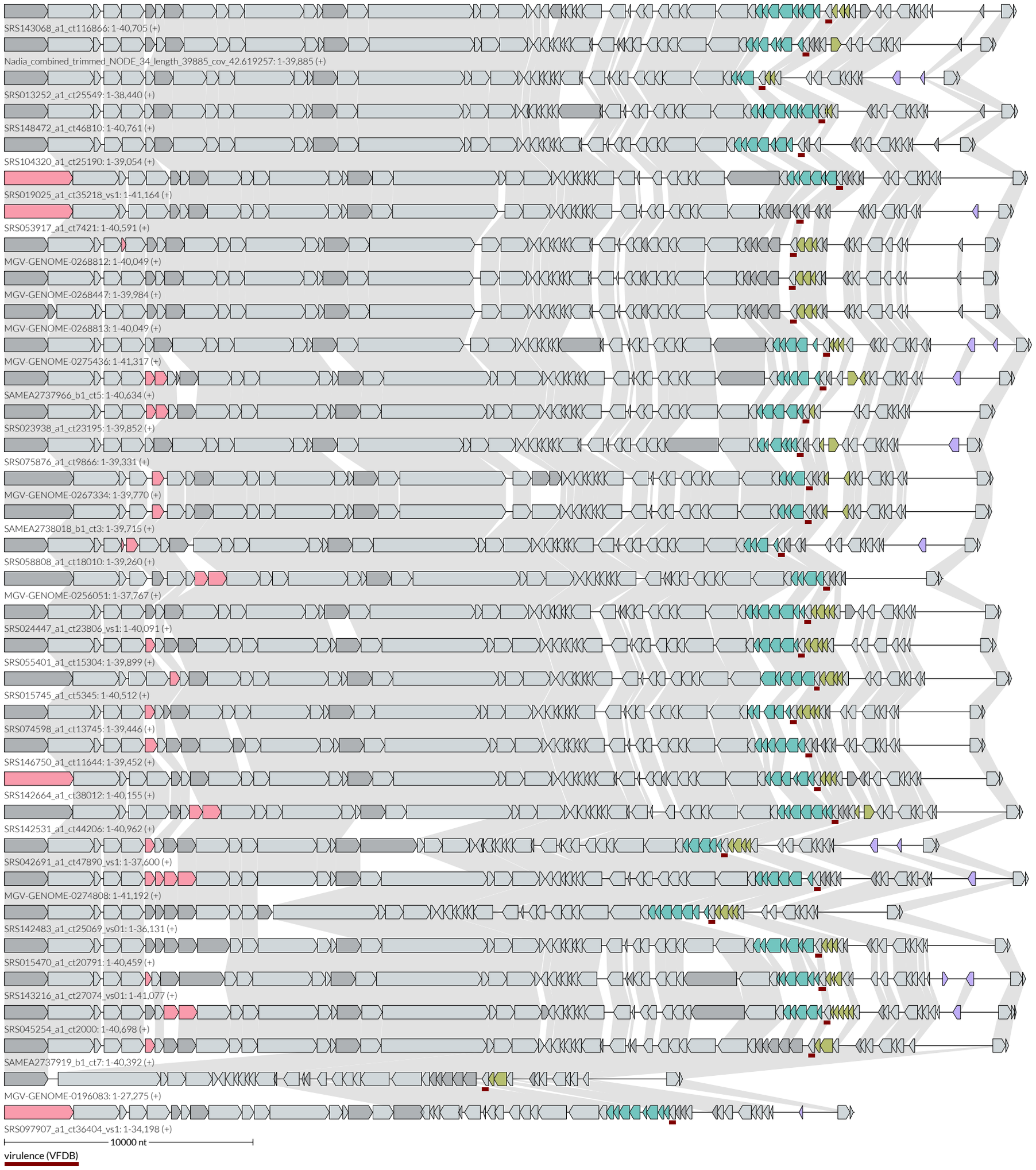
SRS143068_a1_ct116866: 1-40,705 (+)
Nadia_combined_trimmed_NODE_34_length_39885_cov_42.619257: 1-39,885 (+)
SRS013252_a1_ct25549: 1-38,440 (+)
SRS148472_a1_ct46810: 1-40,761 (+)
SRS104320_a1_ct25190: 1-39,054 (+)
SRS019025_a1_ct35218_vs1: 1-41,164 (+)
SRS053917_a1_ct7421: 1-40,591 (+)
MGV-GENOME-0268812: 1-40,049 (+)
MGV-GENOME-0268447: 1-39,984 (+)
MGV-GENOME-0268813: 1-40,049 (+)
MGV-GENOME-0275436: 1-41,317 (+)
SAMEA2737966_b1_ct5: 1-40,634 (+)
SRS023938_a1_ct23195: 1-39,852 (+)
SRS075876_a1_ct9866: 1-39,331 (+)
MGV-GENOME-0267334: 1-39,770 (+)
SAMEA2738018_b1_ct3: 1-39,715 (+)
SRS058808_a1_ct18010: 1-39,260 (+)
MGV-GENOME-0256051: 1-37,767 (+)
SRS024447_a1_ct23806_vs1: 1-40,091 (+)
SRS055401_a1_ct15304: 1-39,899 (+)
SRS015745_a1_ct5345: 1-40,512 (+)
SRS074598_a1_ct13745: 1-39,446 (+)
SRS146750_a1_ct11644: 1-39,452 (+)
SRS142664_a1_ct38012: 1-40,155 (+)
SRS142531_a1_ct44206: 1-40,962 (+)
SRS042691_a1_ct47890_vs1: 1-37,600 (+)
MGV-GENOME-0274808: 1-41,192 (+)
SRS142483_a1_ct25069_vs01: 1-36,131 (+)
SRS015470_a1_ct20791: 1-40,459 (+)
SRS143216_a1_ct27074_vs01: 1-41,077 (+)
SRS045254_a1_ct2000: 1-40,698 (+)
SAMEA2737919_b1_ct7: 1-40,392 (+)
MGV-GENOME-0196083: 1-27,275 (+)
SRS097907_a1_ct36404_vs1: 1-34,198 (+)
10000 nt
virulence (VFDB)