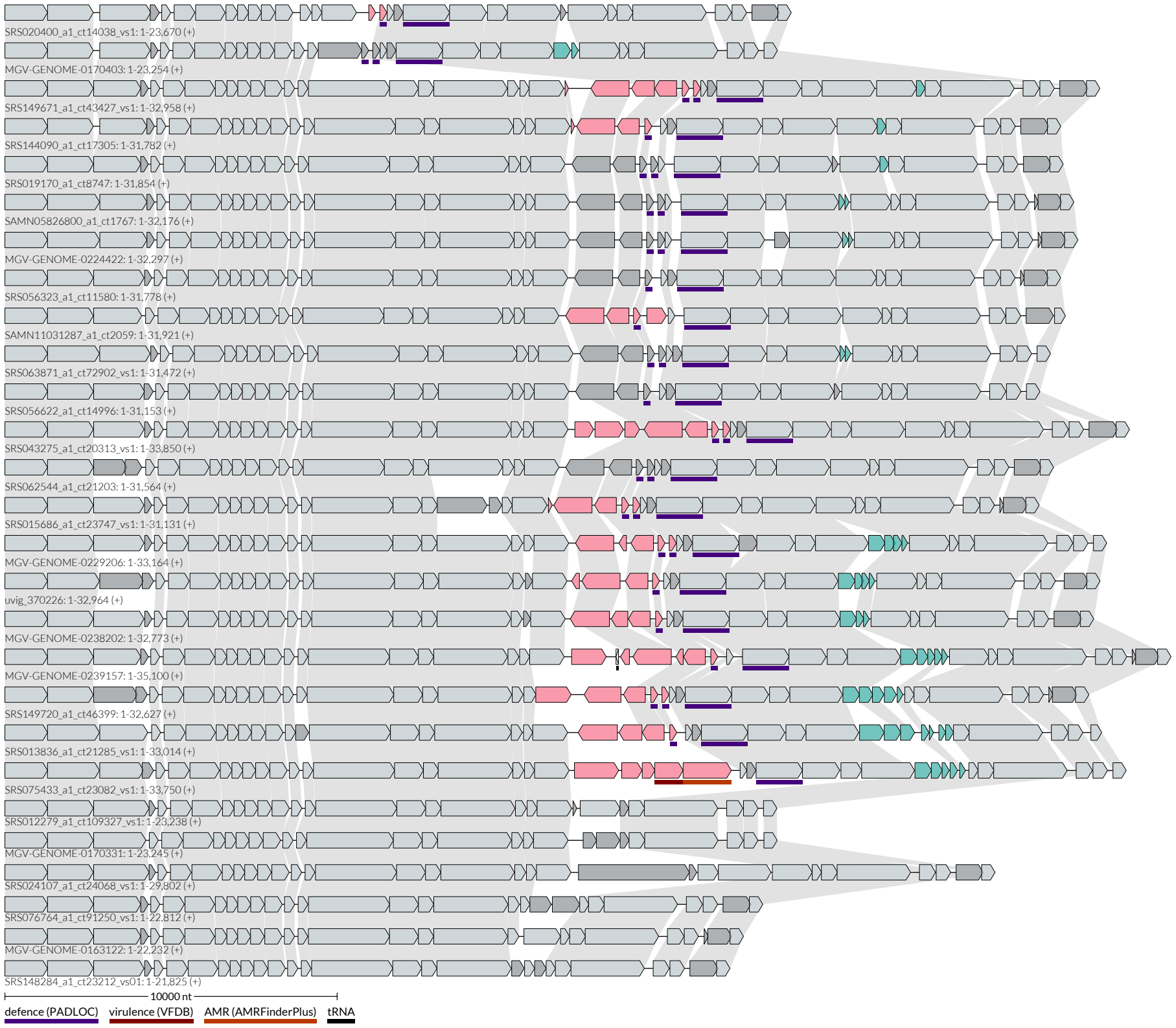
SRS020400_a1_ct14038_vs1: 1-23,670 (+)
MGV-GENOME-0170403: 1-23,254 (+)
SRS149671_a1_ct43427_vs1: 1-32,958 (+)
SRS144090_a1_ct17305: 1-31,782 (+)
SRS019170_a1_ct8747: 1-31,854 (+)
SAMN05826800_a1_ct1767: 1-32,176 (+)
MGV-GENOME-0224422: 1-32,297 (+)
SRS056323_a1_ct11580: 1-31,778 (+)
SAMN11031287_a1_ct2059: 1-31,921 (+)
SRS063871_a1_ct72902_vs1: 1-31,472 (+)
SRS056622_a1_ct14996: 1-31,153 (+)
SRS043275_a1_ct20313_vs1: 1-33,850 (+)
SRS062544_a1_ct21203: 1-31,564 (+)
SRS015686_a1_ct23747_vs1: 1-31,131 (+)
MGV-GENOME-0229206: 1-33,164 (+)
uvig_370226: 1-32,964 (+)
MGV-GENOME-0238202: 1-32,773 (+)
MGV-GENOME-0239157: 1-35,100 (+)
SRS149720_a1_ct46399: 1-32,627 (+)
SRS013836_a1_ct21285_vs1: 1-33,014 (+)
SRS075433_a1_ct23082_vs1: 1-33,750 (+)
SRS012279_a1_ct109327_vs1: 1-23,238 (+)
MGV-GENOME-0170331: 1-23,245 (+)
SRS024107_a1_ct24068_vs1: 1-29,802 (+)
SRS076764_a1_ct91250_vs1: 1-22,812 (+)
MGV-GENOME-0163122: 1-22,232 (+)
SRS148284_a1_ct23212_vs01: 1-21,825 (+)
10000 nt
defence (PADLOC)
virulence (VFDB)
AMR (AMRFinderPlus)
tRNA