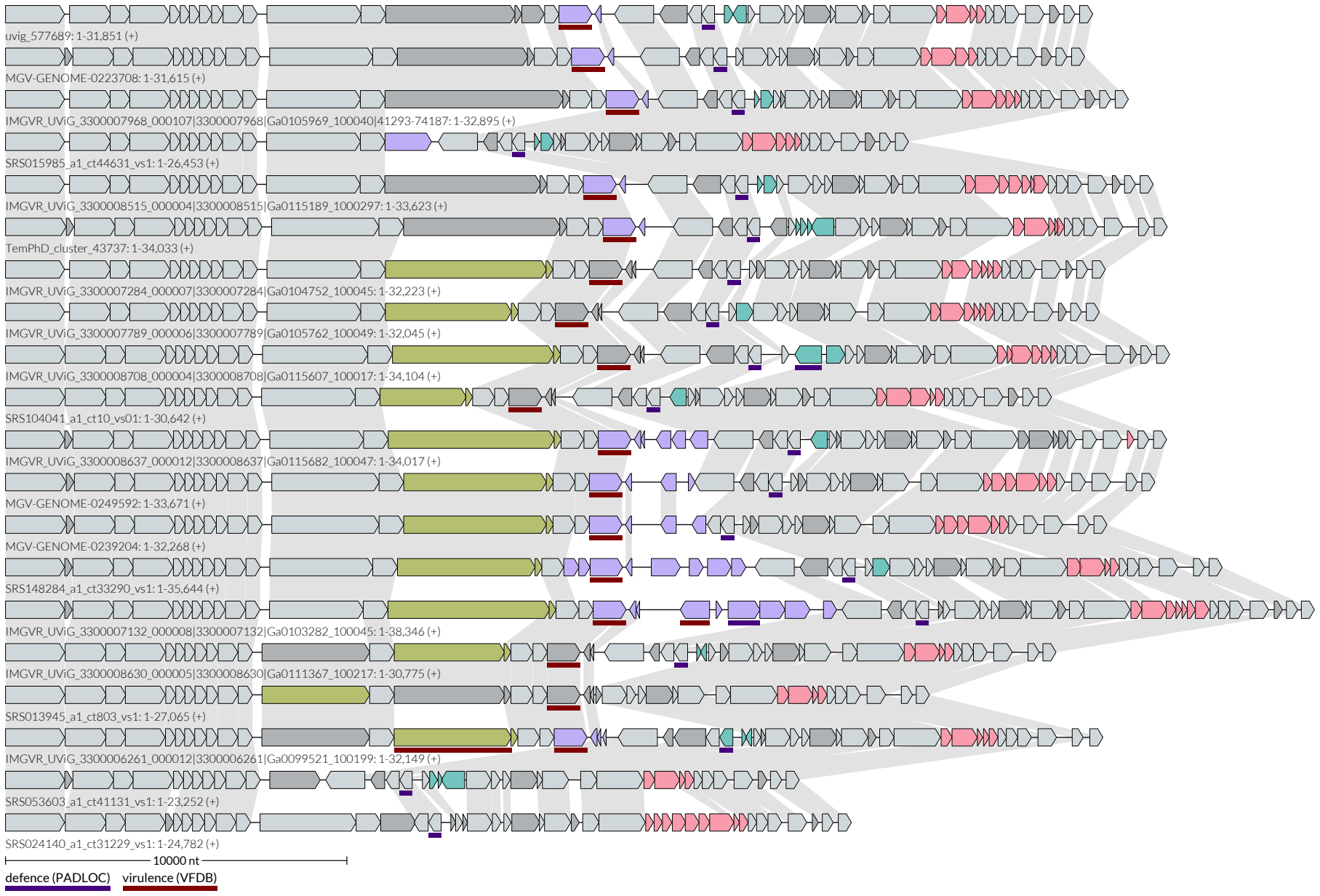
uvig_577689: 1-31,851 (+)
MGV-GENOME-0223708: 1-31,615 (+)
IMGVR_UViG_3300007968_000107|3300007968|Ga0105969_100040|41293-74187: 1-32,895 (+)
SRS015985_a1_ct44631_vs1: 1-26,453 (+)
IMGVR_UViG_3300008515_000004|3300008515|Ga0115189_1000297: 1-33,623 (+)
TemPhD_cluster_43737: 1-34,033 (+)
IMGVR_UViG_3300007284_000007|3300007284|Ga0104752_100045: 1-32,223 (+)
IMGVR_UViG_3300007789_000006|3300007789|Ga0105762_100049: 1-32,045 (+)
IMGVR_UViG_3300008708_000004|3300008708|Ga0115607_100017: 1-34,104 (+)
SRS104041_a1_ct10_vs01: 1-30,642 (+)
IMGVR_UViG_3300008637_000012|3300008637|Ga0115682_100047: 1-34,017 (+)
MGV-GENOME-0249592: 1-33,671 (+)
MGV-GENOME-0239204: 1-32,268 (+)
SRS148284_a1_ct33290_vs1: 1-35,644 (+)
IMGVR_UViG_3300007132_000008|3300007132|Ga0103282_100045: 1-38,346 (+)
IMGVR_UViG_3300008630_000005|3300008630|Ga0111367_100217: 1-30,775 (+)
SRS013945_a1_ct803_vs1: 1-27,065 (+)
IMGVR_UViG_3300006261_000012|3300006261|Ga0099521_100199: 1-32,149 (+)
SRS053603_a1_ct41131_vs1: 1-23,252 (+)
SRS024140_a1_ct31229_vs1: 1-24,782 (+)
10000 nt
defence (PADLOC)
virulence (VFDB)