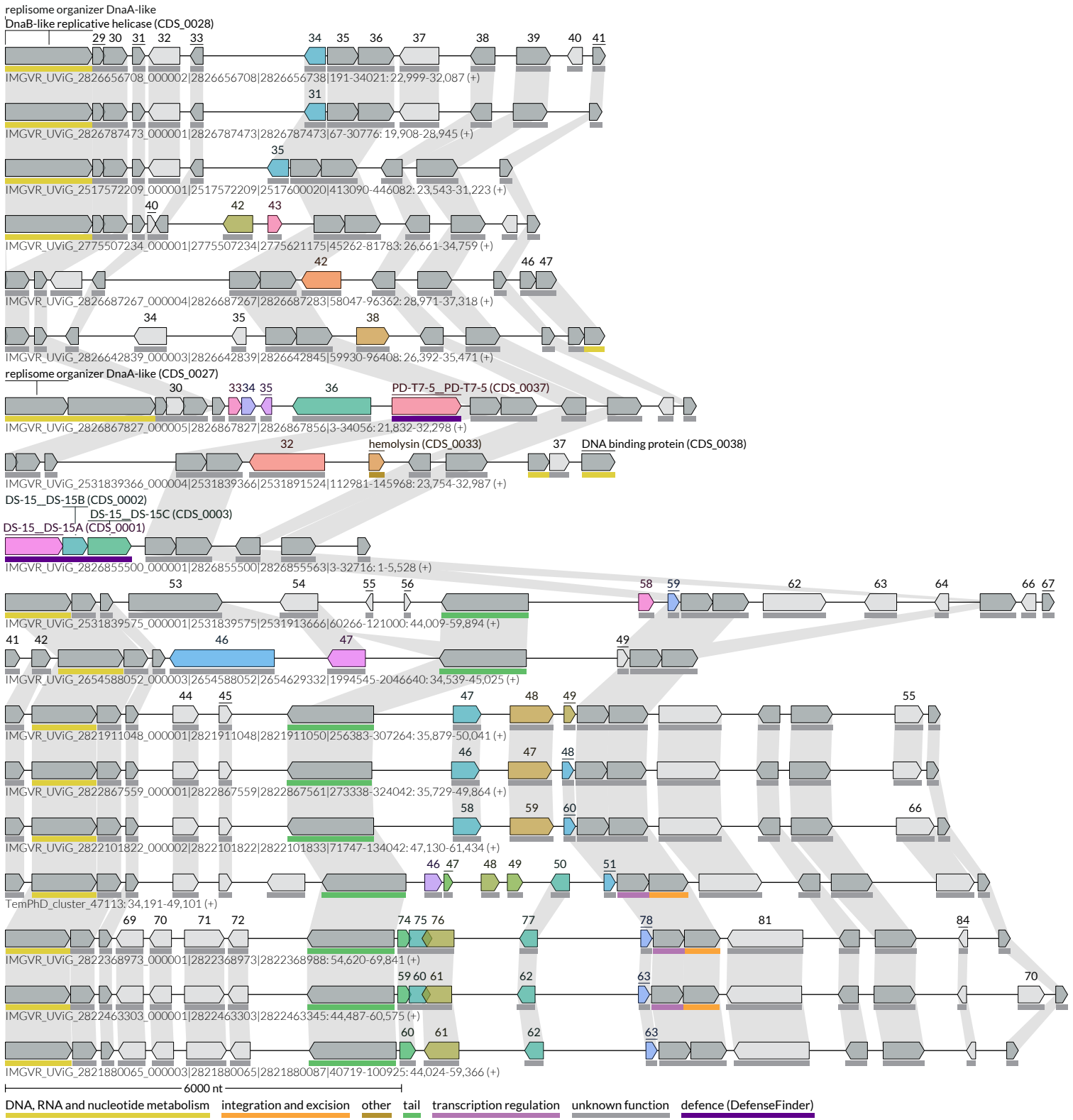
replisome organizer DnaA-like
DnaB-like replicative helicase (CDS_0028)
IMGVR_UViG_2826656708_000002|2826656708|2826656738|191-34021: 22,999-32,087 (+)
IMGVR_UViG_2826787473_000001|2826787473|2826787473|67-30776: 19,908-28,945 (+)
IMGVR_UViG_2517572209_000001|2517572209|2517600020|413090-446082: 23,543-31,223 (+)
IMGVR_UViG_2775507234_000001|2775507234|2775621175|45262-81783: 26,661-34,759 (+)
IMGVR_UViG_2826687267_000004|2826687267|2826687283|58047-96362: 28,971-37,318 (+)
IMGVR_UViG_2826642839_000003|2826642839|2826642845|59930-96408: 26,392-35,471 (+)
replisome organizer DnaA-like (CDS_0027)
PD-T7-5__PD-T7-5 (CDS_0037)
IMGVR_UViG_2826867827_000005|2826867827|2826867856|3-34056: 21,832-32,298 (+)
hemolysin (CDS_0033)
DNA binding protein (CDS_0038)
IMGVR_UViG_2531839366_000004|2531839366|2531891524|112981-145968: 23,754-32,987 (+)
DS-15__DS-15B (CDS_0002)
DS-15__DS-15C (CDS_0003)
DS-15__DS-15A (CDS_0001)
IMGVR_UViG_2826855500_000001|2826855500|2826855563|3-32716: 1-5,528 (+)
IMGVR_UViG_2531839575_000001|2531839575|2531913666|60266-121000: 44,009-59,894 (+)
IMGVR_UViG_2654588052_000003|2654588052|2654629332|1994545-2046640: 34,539-45,025 (+)
IMGVR_UViG_2821911048_000001|2821911048|2821911050|256383-307264: 35,879-50,041 (+)
IMGVR_UViG_2822867559_000001|2822867559|2822867561|273338-324042: 35,729-49,864 (+)
IMGVR_UViG_2822101822_000002|2822101822|2822101833|71747-134042: 47,130-61,434 (+)
TemPhD_cluster_47113: 34,191-49,101 (+)
IMGVR_UViG_2822368973_000001|2822368973|2822368988: 54,620-69,841 (+)
IMGVR_UViG_2822463303_000001|2822463303|2822463345: 44,487-60,575 (+)
IMGVR_UViG_2821880065_000003|2821880065|2821880087|40719-100925: 44,024-59,366 (+)
6000 nt
DNA, RNA and nucleotide metabolism
integration and excision
other
tail
transcription regulation
unknown function
defence (DefenseFinder)