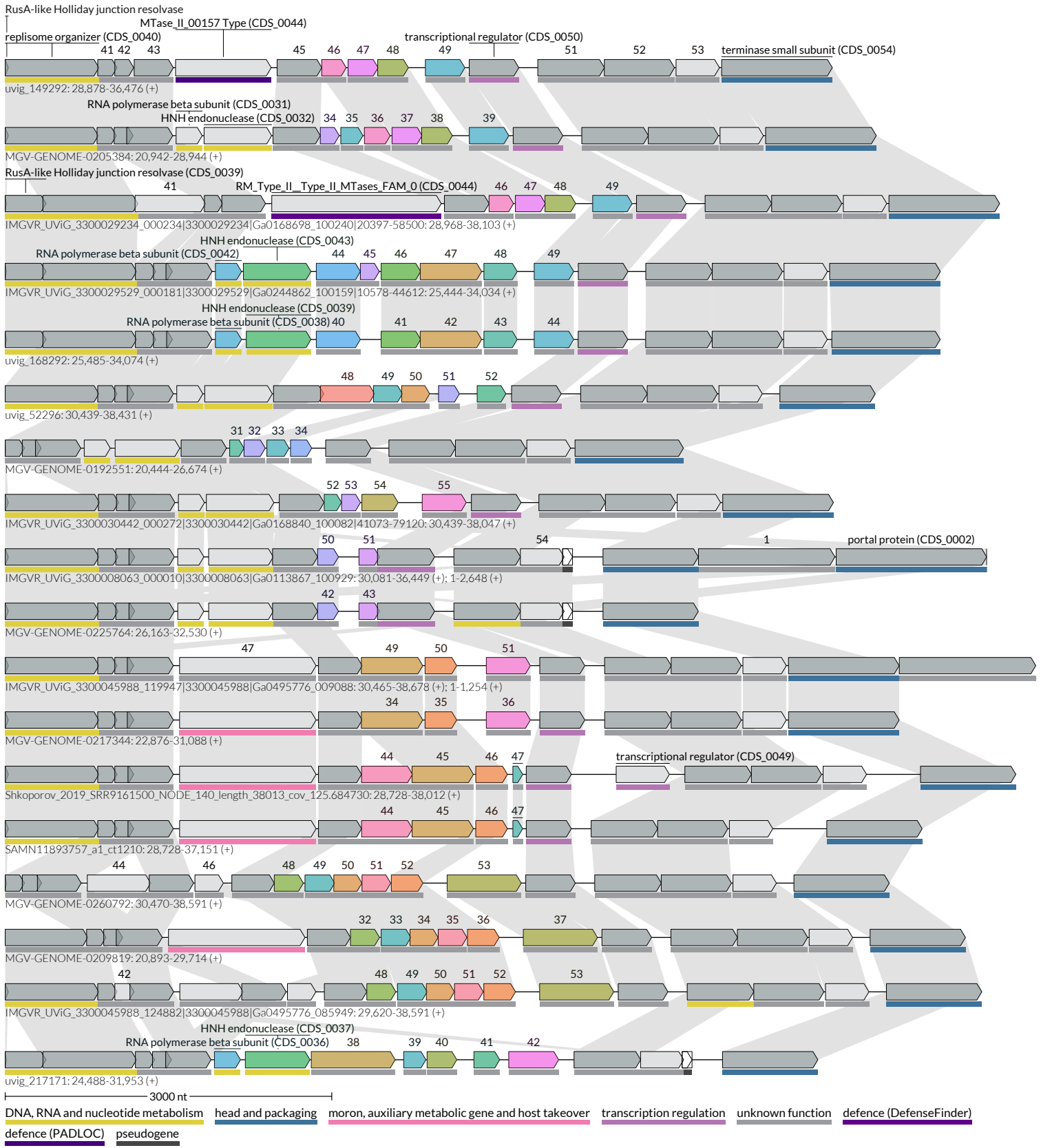
RusA-like Holliday junction resolvase
MTase_II_00157 Type (CDS_0044)
replisome organizer (CDS_0040)
transcriptional regulator (CDS_0050)
terminase small subunit (CDS_0054)
uvig_149292: 28,878-36,476 (+)
RNA polymerase beta subunit (CDS_0031)
HNH endonuclease (CDS_0032)
MGV-GENOME-0205384: 20,942-28,944 (+)
RusA-like Holliday junction resolvase (CDS_0039)
RM_Type_II__Type_II_MTases_FAM_0 (CDS_0044)
IMGVR_UVIG_3300029234_000234|3300029234|Ga0168698_100240|20397-58500: 28,968-38,103 (+)
HNH endonuclease (CDS_0043)
RNA polymerase beta subunit (CDS_0042)
IMGVR_UVIG_3300029529_000181|3300029529|Ga0244862_100159|10578-44612: 25,444-34,034 (+)
HNH endonuclease (CDS_0039)
RNA polymerase beta subunit (CDS_0038)
uvig_168292: 25,485-34,074 (+)
uvig_52296: 30,439-38,431 (+)
MGV-GENOME-0192551: 20,444-26,674 (+)
IMGVR_UVIG_3300030442_000272|3300030442|Ga0168840_100082|41073-79120: 30,439-38,047 (+)
portal protein (CDS_0002)
IMGVR_UVIG_3300008063_000010|3300008063|Ga0113867_100929: 30,081-36,449 (+); 1-2,648 (+)
MGV-GENOME-0225764: 26,163-32,530 (+)
IMGVR_UVIG_3300045988_119947|3300045988|Ga0495776_009088: 30,465-38,678 (+); 1-1,254 (+)
MGV-GENOME-0217344: 22,876-31,088 (+)
transcriptional regulator (CDS_0049)
Shkoporov_2019_SRR9161500_NODE_140_length_38013_cov_125.684730: 28,728-38,012 (+)
SAMN11893757_a1_ct1210: 28,728-37,151 (+)
MGV-GENOME-0260792: 30,470-38,591 (+)
MGV-GENOME-0209819: 20,893-29,714 (+)
IMGVR_UVIG_3300045988_124882|3300045988|Ga0495776_085949: 29,620-38,591 (+)
HNH endonuclease (CDS_0037)
RNA polymerase beta subunit (CDS_0036)
uvig_217171: 24,488-31,953 (+)
3000 nt
DNA, RNA and nucleotide metabolism
head and packaging
moron, auxiliary metabolic gene and host takeover
transcription regulation
unknown function
defence (DefenseFinder)
defence (PADLOC)
pseudogene