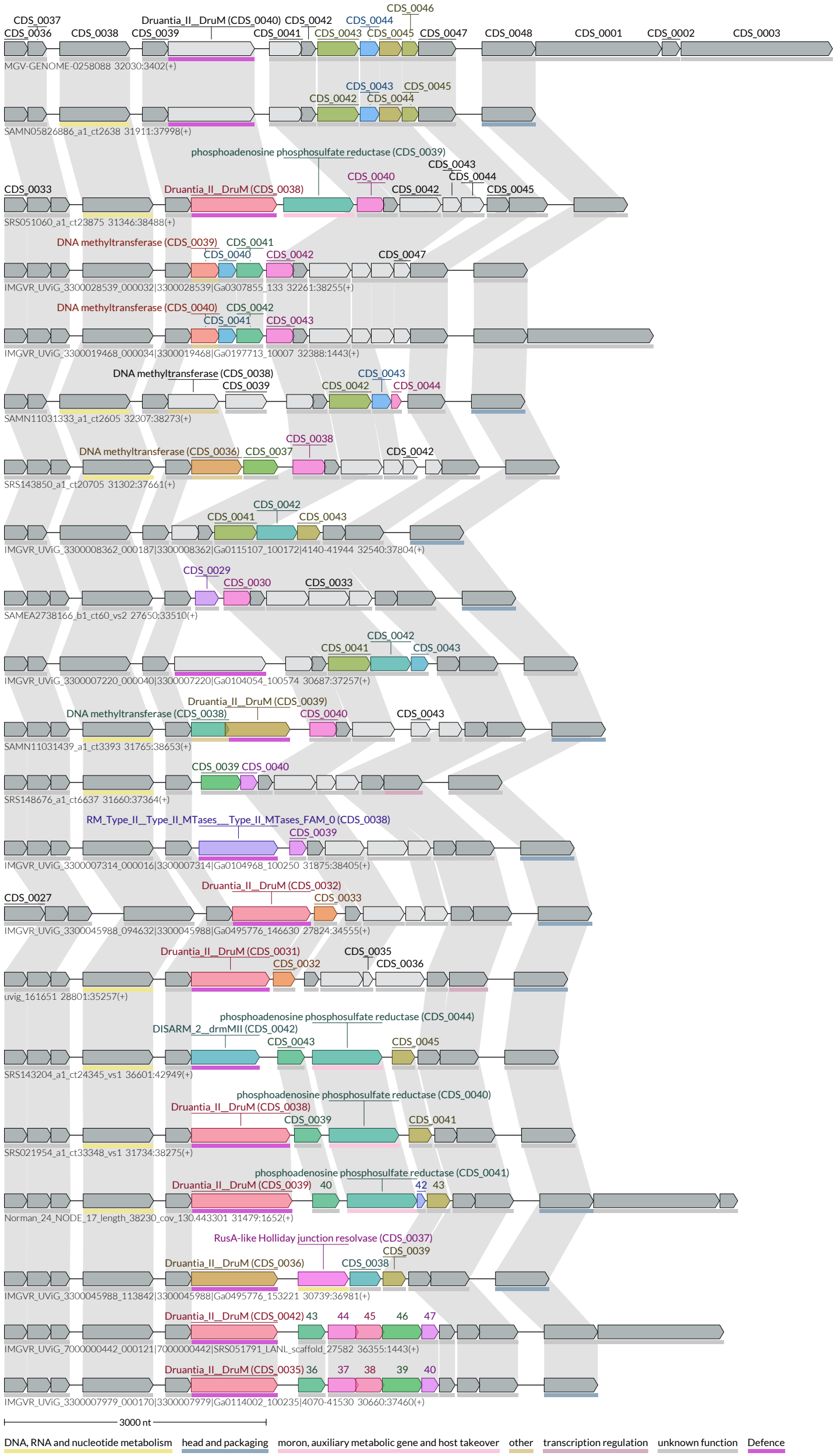
CDS_0037
CDS_0036
CDS_0038
CDS_0039
Druantia_II__DruM (CDS_0040)
CDS_0041
CDS_0042
CDS_0043
CDS_0044
CDS_0045
CDS_0046
CDS_0047
CDS_0048
CDS_0001
CDS_0002
CDS_0003
MGV-GENOME-0258088 32030:3402(+)
SAMN05826886_a1_ct2638 31911:37998(+)
phosphoadenosine phosphosulfate reductase (CDS_0039)
CDS_0033
Druantia_II__DruM (CDS_0038)
CDS_0040
SRS051060_a1_ct23875 31346:38488(+)
DNA methyltransferase (CDS_0039)
IMGVR_UViG_3300028539_000032|3300028539|Ga0307855_133 32261:38255(+)
DNA methyltransferase (CDS_0040)
IMGVR_UViG_3300019468_000034|3300019468|Ga0197713_10007 32388:1443(+)
DNA methyltransferase (CDS_0038)
SAMN11031333_a1_ct2605 32307:38273(+)
DNA methyltransferase (CDS_0036)
SRS143850_a1_ct20705 31302:37661(+)
IMGVR_UViG_3300008362_000187|3300008362|Ga0115107_100172|4140-41944 32540:37804(+)
SAMEA2738166_b1_ct60_vs2 27650:33510(+)
IMGVR_UViG_3300007220_000040|3300007220|Ga0104054_100574 30687:37257(+)
Druantia_II__DruM (CDS_0039)
SAMN11031439_a1_ct3393 31765:38653(+)
SRS148676_a1_ct6637 31660:37364(+)
RM_Type_II__Type_II_MTases___Type_II_MTases_FAM_0 (CDS_0038)
IMGVR_UViG_3300007314_000016|3300007314|Ga0104968_100250 31875:38405(+)
Druantia_II__DruM (CDS_0032)
CDS_0027
IMGVR_UViG_3300045988_094632|3300045988|Ga0495776_146630 27824:34555(+)
Druantia_II__DruM (CDS_0031)
CDS_0035
uvig_161651 28801:35257(+)
phosphoadenosine phosphosulfate reductase (CDS_0044)
DISARM_2__drmMII (CDS_0042)
SRS143204_a1_ct24345_vs1 36601:42949(+)
phosphoadenosine phosphosulfate reductase (CDS_0040)
SRS021954_a1_ct33348_vs1 31734:38275(+)
phosphoadenosine phosphosulfate reductase (CDS_0041)
Druantia_II__DruM (CDS_0039)
Norman_24_NODE_17_length_38230_cov_130.443301 31479:1652(+)
RusA-like Holliday junction resolvase (CDS_0037)
Druantia_II__DruM (CDS_0036)
IMGVR_UViG_3300045988_113842|3300045988|Ga0495776_153221 30739:36981(+)
Druantia_II__DruM (CDS_0042)
IMGVR_UViG_7000000442_000121|7000000442|SRS051791_LANL_scaffold_27582 36355:1443(+)
Druantia_II__DruM (CDS_0035)
IMGVR_UViG_3300007979_000170|3300007979|Ga0114002_100235|4070-41530 30660:37460(+)
3000 nt
DNA, RNA and nucleotide metabolism
head and packaging
moron, auxiliary metabolic gene and host takeover
other
transcription regulation
unknown function
Defence