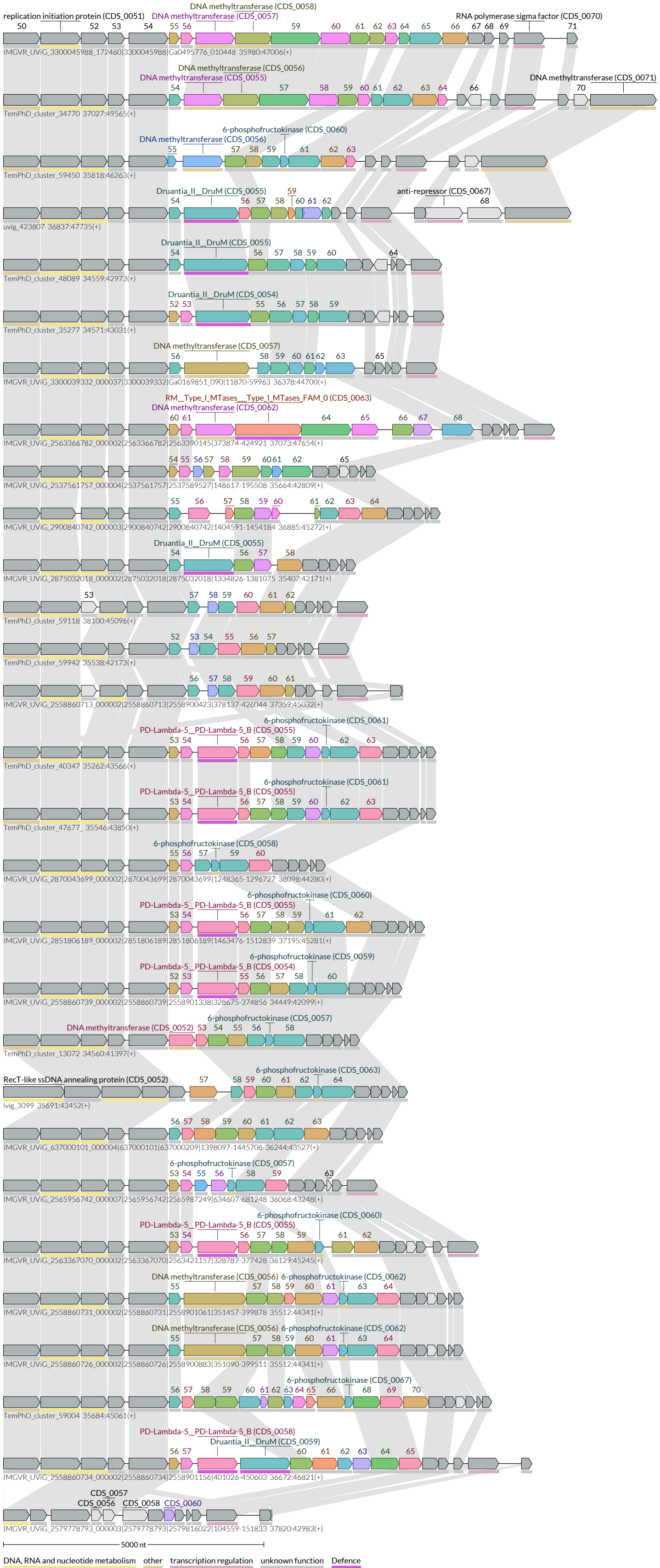
replication initiation protein (CDS_0051)
DNA methyltransferase (CDS_0058)
DNA methyltransferase (CDS_0057)
RNA polymerase sigma factor (CDS_0070)
IMGVR_UViG_3300045988_172460|3300045988|Ga0495776_010448 35980:47006(+)
DNA methyltransferase (CDS_0056)
DNA methyltransferase (CDS_0055)
DNA methyltransferase (CDS_0071)
TemPhD_cluster_34770 37027:49565(+)
6-phosphofructokinase (CDS_0060)
DNA methyltransferase (CDS_0056)
TemPhD_cluster_59450 35818:46263(+)
Druantia_II__DruM (CDS_0055)
anti-repressor (CDS_0067)
uvig_423807 36837:47735(+)
Druantia_II__DruM (CDS_0055)
TemPhD_cluster_48089 34559:42973(+)
Druantia_II__DruM (CDS_0054)
TemPhD_cluster_35277 34571:43031(+)
DNA methyltransferase (CDS_0057)
IMGVR_UViG_3300039332_000037|3300039332|Ga0169851_090|11870-59963 36378:44700(+)
RM__Type_I_MTases__Type_I_MTases_FAM_0 (CDS_0063)
DNA methyltransferase (CDS_0062)
IMGVR_UViG_2563366782_000002|2563366782|2563390145|373874-424921 37073:47654(+)
IMGVR_UViG_2537561757_000004|2537561757|2537589527|148617-195508 35664:42809(+)
IMGVR_UViG_2900840742_000003|2900840742|2900840742|1404591-1454184 36885:45272(+)
Druantia_II__DruM (CDS_0055)
IMGVR_UViG_2875032018_000002|2875032018|2875032018|1334826-1381075 35407:42171(+)
TemPhD_cluster_59118 38100:45096(+)
TemPhD_cluster_59942 35538:42173(+)
IMGVR_UViG_2558860713_000002|2558860713|2558900423|378137-426044 37359:45032(+)
6-phosphofructokinase (CDS_0061)
PD-Lambda-5__PD-Lambda-5_B (CDS_0055)
TemPhD_cluster_40347 35262:43566(+)
6-phosphofructokinase (CDS_0061)
PD-Lambda-5__PD-Lambda-5_B (CDS_0055)
TemPhD_cluster_47677_ 35546:43850(+)
6-phosphofructokinase (CDS_0058)
IMGVR_UViG_2870043699_000002|2870043699|2870043699|1248365-1296727 38098:44280(+)
6-phosphofructokinase (CDS_0060)
PD-Lambda-5__PD-Lambda-5_B (CDS_0055)
IMGVR_UViG_2851806189_000002|2851806189|2851806189|1463476-1512839 37195:45281(+)
6-phosphofructokinase (CDS_0059)
PD-Lambda-5__PD-Lambda-5_B (CDS_0054)
IMGVR_UViG_2558860739_000002|2558860739|2558901338|328675-374856 34449:42099(+)
6-phosphofructokinase (CDS_0057)
DNA methyltransferase (CDS_0052)
TemPhD_cluster_13072 34560:41397(+)
6-phosphofructokinase (CDS_0063)
RecT-like ssDNA annealing protein (CDS_0052)
ivig_3099 35691:43452(+)
IMGVR_UViG_637000101_000004|637000101|637000209|1398097-1445706 36244:43527(+)
6-phosphofructokinase (CDS_0057)
IMGVR_UViG_2565956742_000007|2565956742|2565987249|634607-681248 36068:43248(+)
6-phosphofructokinase (CDS_0060)
PD-Lambda-5__PD-Lambda-5_B (CDS_0055)
IMGVR_UViG_2563367070_000002|2563367070|2563421157|328787-377428 36129:45245(+)
DNA methyltransferase (CDS_0056)
6-phosphofructokinase (CDS_0062)
IMGVR_UViG_2558860731_000002|2558860731|2558901061|351457-399878 35512:44341(+)
DNA methyltransferase (CDS_0056)
6-phosphofructokinase (CDS_0062)
IMGVR_UViG_2558860726_000002|2558860726|2558900883|351090-399511 35512:44341(+)
6-phosphofructokinase (CDS_0067)
TemPhD_cluster_59004 35684:45061(+)
PD-Lambda-5__PD-Lambda-5_B (CDS_0058)
Druantia_II__DruM (CDS_0059)
IMGVR_UViG_2558860734_000002|2558860734|2558901156|401026-450603 36672:46821(+)
CDS_0057
CDS_0056
CDS_0058
CDS_0060
IMGVR_UViG_2579778793_000003|2579778793|2579816022|104559-151833 37820:42983(+)
5000 nt
DNA, RNA and nucleotide metabolism
other
transcription regulation
unknown function
Defence