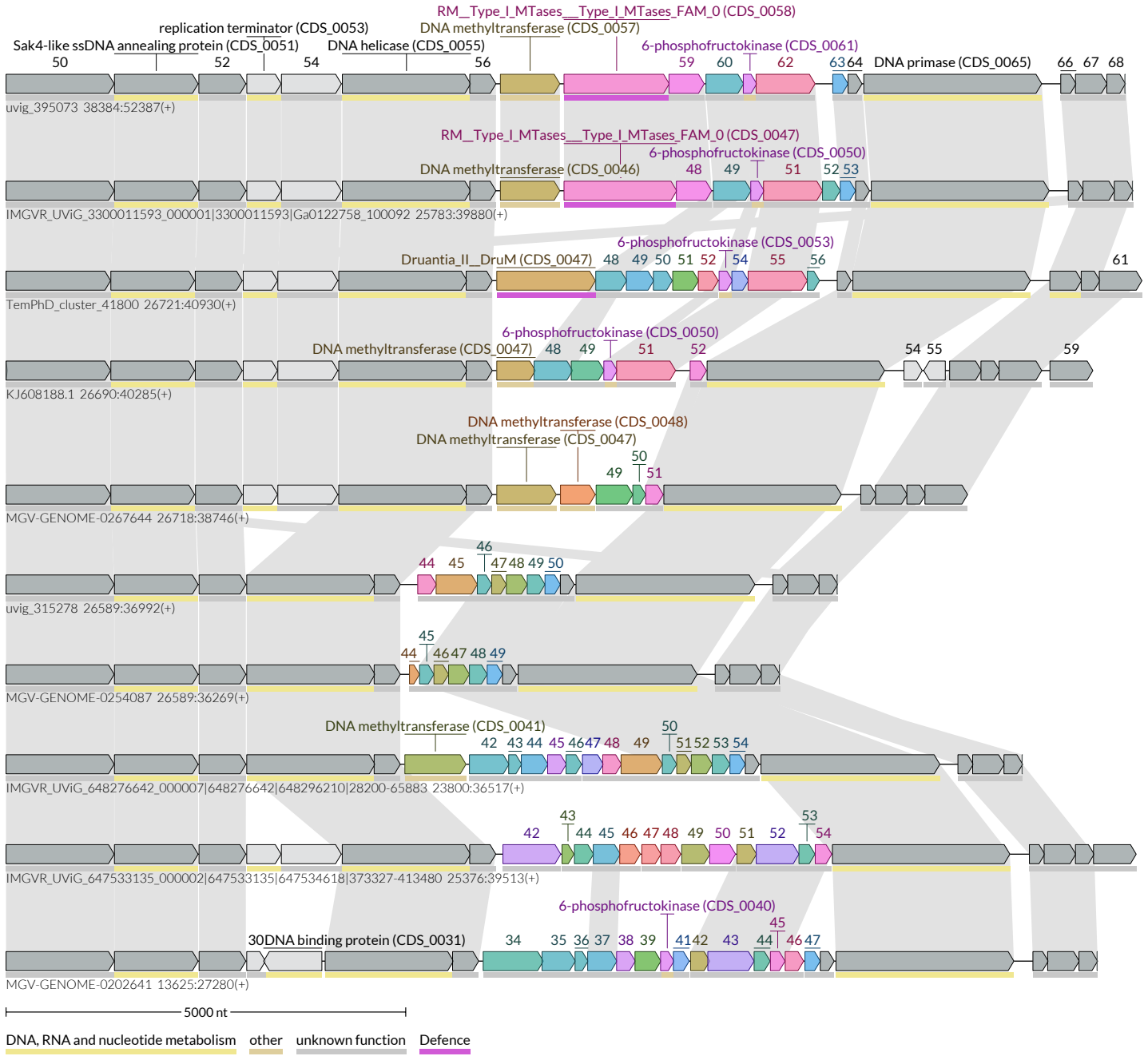
RM__Type_I_MTases__Type_I_MTases_FAM_0 (CDS_0058)
replication terminator (CDS_0053)
DNA methyltransferase (CDS_0057)
Sak4-like ssDNA annealing protein (CDS_0051)
DNA helicase (CDS_0055)
6-phosphofructokinase (CDS_0061)
50 52 54 56 59 60 62 63 64 DNA primase (CDS_0065) 66 67 68
uvig_395073 38384:52387(+)
RM__Type_I_MTases__Type_I_MTases_FAM_0 (CDS_0047)
6-phosphofructokinase (CDS_0050)
DNA methyltransferase (CDS_0046)
48 49 51 52 53
IMGVR_UViG_3300011593_000001|3300011593|Ga0122758_100092 25783:39880(+)
6-phosphofructokinase (CDS_0053)
Druantia_II__DruM (CDS_0047)
48 49 50 51 52 54 55 56 61
TemPhD_cluster_41800 26721:40930(+)
6-phosphofructokinase (CDS_0050)
DNA methyltransferase (CDS_0047)
48 49 51 52 54 55 59
KJ608188.1 26690:40285(+)
DNA methyltransferase (CDS_0048)
DNA methyltransferase (CDS_0047)
49 50 51
MGV-GENOME-0267644 26718:38746(+)
44 45 46 47 48 49 50
uvig_315278 26589:36992(+)
44 45 46 47 48 49
MGV-GENOME-0254087 26589:36269(+)
DNA methyltransferase (CDS_0041)
42 43 44 45 46 47 48 49 50 51 52 53 54
IMGVR_UViG_648276642_000007|648276642|648296210|28200-65883 23800:36517(+)
42 43 44 45 46 47 48 49 50 51 52 53 54
IMGVR_UViG_647533135_000002|647533135|647534618|373327-413480 25376:39513(+)
6-phosphofructokinase (CDS_0040)
30DNA binding protein (CDS_0031)
34 35 36 37 38 39 41 42 43 44 45 46 47
MGV-GENOME-0202641 13625:27280(+)
5000 nt
DNA, RNA and nucleotide metabolism
other
unknown function
Defence