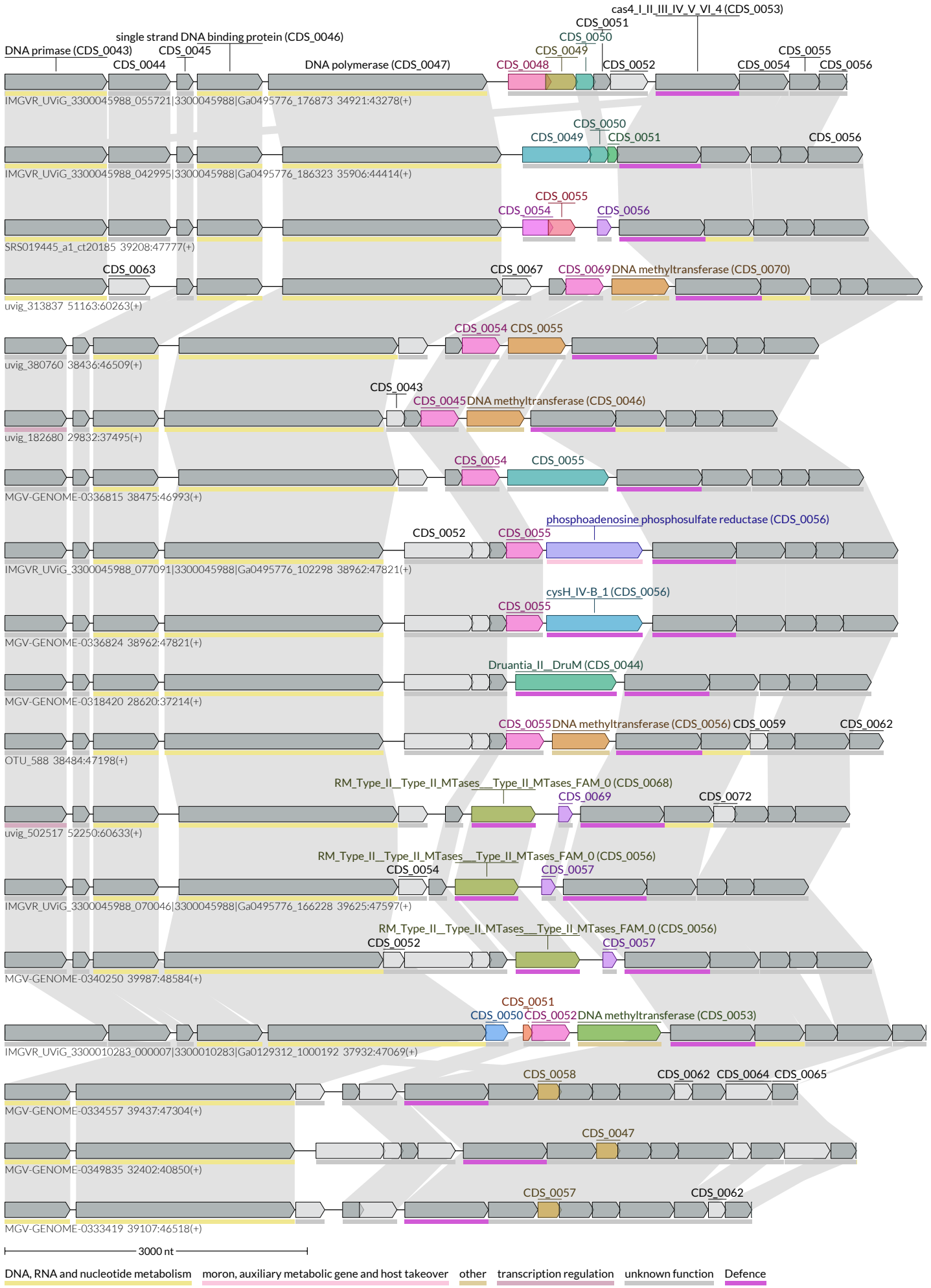
cas4_I_II_III_IV_V_VI_4 (CDS_0053)
CDS_0051
CDS_0050
CDS_0049
CDS_0048
CDS_0052
CDS_0054
CDS_0056
CDS_0055
single strand DNA binding protein (CDS_0046)
DNA primase (CDS_0043)
CDS_0045
CDS_0044
DNA polymerase (CDS_0047)
IMGVR_UViG_3300045988_055721|3300045988|Ga0495776_176873 34921:43278(+)
IMGVR_UViG_3300045988_042995|3300045988|Ga0495776_186323 35906:44414(+)
SRS019445_a1_ct20185 39208:47777(+)
CDS_0063
CDS_0067
CDS_0069
DNA methyltransferase (CDS_0070)
uvig_313837 51163:60263(+)
uvig_380760 38436:46509(+)
CDS_0043
DNA methyltransferase (CDS_0046)
uvig_182680 29832:37495(+)
MGV-GENOME-0336815 38475:46993(+)
phosphoadenosine phosphosulfate reductase (CDS_0056)
IMGVR_UViG_3300045988_077091|3300045988|Ga0495776_102298 38962:47821(+)
cysH_IV-B_1 (CDS_0056)
MGV-GENOME-0336824 38962:47821(+)
Druantia_II__DruM (CDS_0044)
MGV-GENOME-0318420 28620:37214(+)
DNA methyltransferase (CDS_0056)
CDS_0059
CDS_0062
OTU_588 38484:47198(+)
RM_Type_II__Type_II_MTases___Type_II_MTases_FAM_0 (CDS_0068)
CDS_0072
uvig_502517 52250:60633(+)
RM_Type_II__Type_II_MTases___Type_II_MTases_FAM_0 (CDS_0056)
CDS_0057
IMGVR_UViG_3300045988_070046|3300045988|Ga0495776_166228 39625:47597(+)
MGV-GENOME-0340250 39987:48584(+)
DNA methyltransferase (CDS_0053)
IMGVR_UViG_3300010283_000007|3300010283|Ga0129312_1000192 37932:47069(+)
CDS_0058
CDS_0062
CDS_0064
CDS_0065
MGV-GENOME-0334557 39437:47304(+)
CDS_0047
MGV-GENOME-0349835 32402:40850(+)
MGV-GENOME-0333419 39107:46518(+)
3000 nt
DNA, RNA and nucleotide metabolism
moron, auxiliary metabolic gene and host takeover
other
transcription regulation
unknown function
Defence