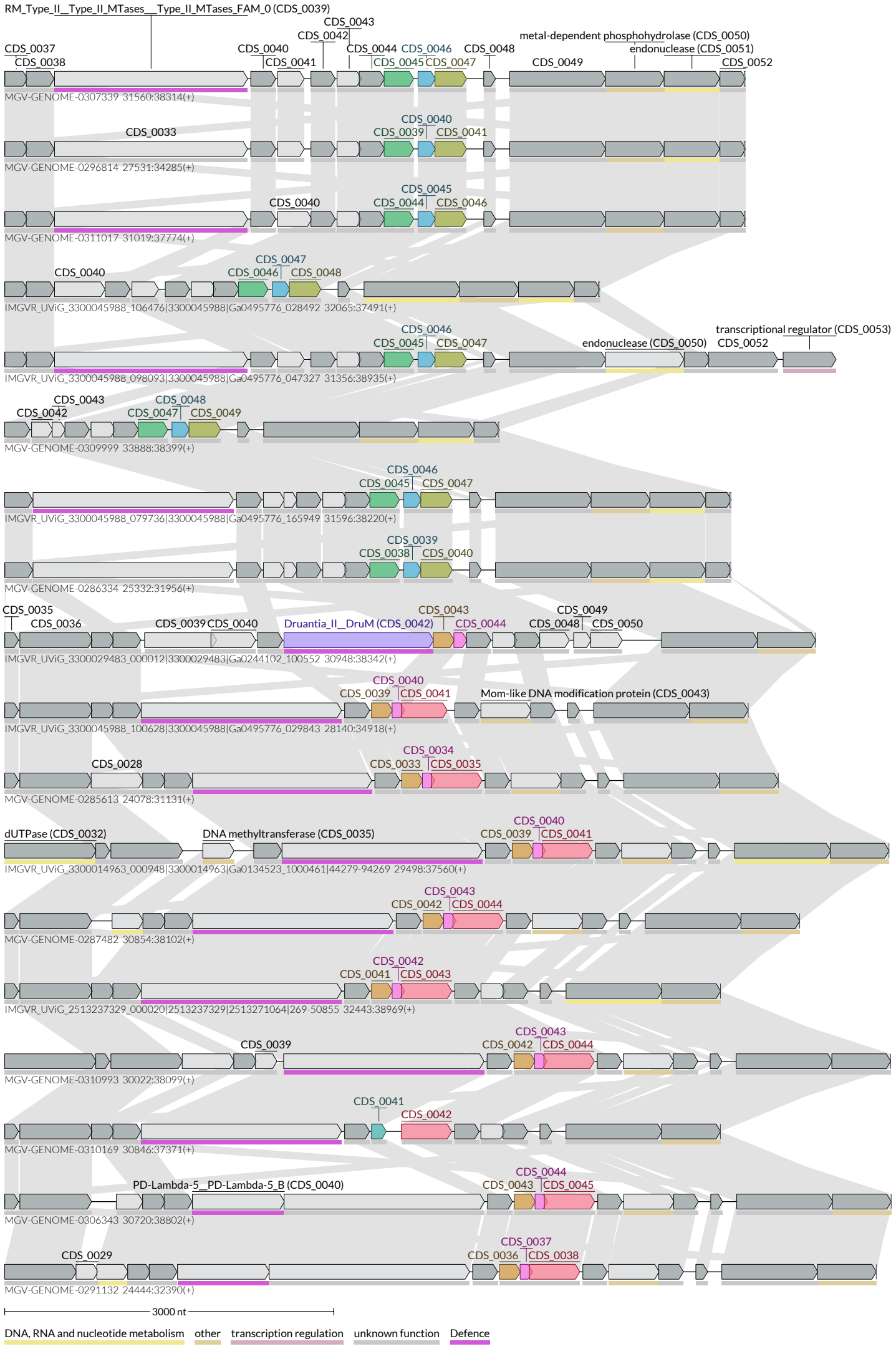
RM_Type_II__Type_II_MTases__Type_II_MTases_FAM_0 (CDS_0039)
CDS_0037
CDS_0038
CDS_0040
CDS_0041
CDS_0042
CDS_0043
CDS_0044
CDS_0045
CDS_0046
CDS_0047
CDS_0048
metal-dependent phosphohydrolase (CDS_0050)
endonuclease (CDS_0051)
CDS_0049
CDS_0052
MGV-GENOME-0307339 31560:38314(+)
CDS_0033
CDS_0039
MGV-GENOME-0296814 27531:34285(+)
MGV-GENOME-0311017 31019:37774(+)
IMGVR_UViG_3300045988_106476|3300045988|Ga0495776_028492 32065:37491(+)
transcriptional regulator (CDS_0053)
endonuclease (CDS_0050)
IMGVR_UViG_3300045988_098093|3300045988|Ga0495776_047327 31356:38935(+)
MGV-GENOME-0309999 33888:38399(+)
IMGVR_UViG_3300045988_079736|3300045988|Ga0495776_165949 31596:38220(+)
CDS_0038
MGV-GENOME-0286334 25332:31956(+)
CDS_0035
CDS_0036
CDS_0039
Druantia_II__DruM (CDS_0042)
IMGVR_UViG_3300029483_000012|3300029483|Ga0244102_100552 30948:38342(+)
Mom-like DNA modification protein (CDS_0043)
IMGVR_UViG_3300045988_100628|3300045988|Ga0495776_029843 28140:34918(+)
CDS_0028
CDS_0033
MGV-GENOME-0285613 24078:31131(+)
dUTPase (CDS_0032)
DNA methyltransferase (CDS_0035)
IMGVR_UViG_3300014963_000948|3300014963|Ga0134523_1000461|44279-94269 29498:37560(+)
MGV-GENOME-0287482 30854:38102(+)
IMGVR_UViG_2513237329_000020|2513237329|2513271064|269-50855 32443:38969(+)
MGV-GENOME-0310993 30022:38099(+)
MGV-GENOME-0310169 30846:37371(+)
PD-Lambda-5__PD-Lambda-5_B (CDS_0040)
MGV-GENOME-0306343 30720:38802(+)
CDS_0029
CDS_0036
MGV-GENOME-0291132 24444:32390(+)
3000 nt
DNA, RNA and nucleotide metabolism
other
transcription regulation
unknown function
Defence