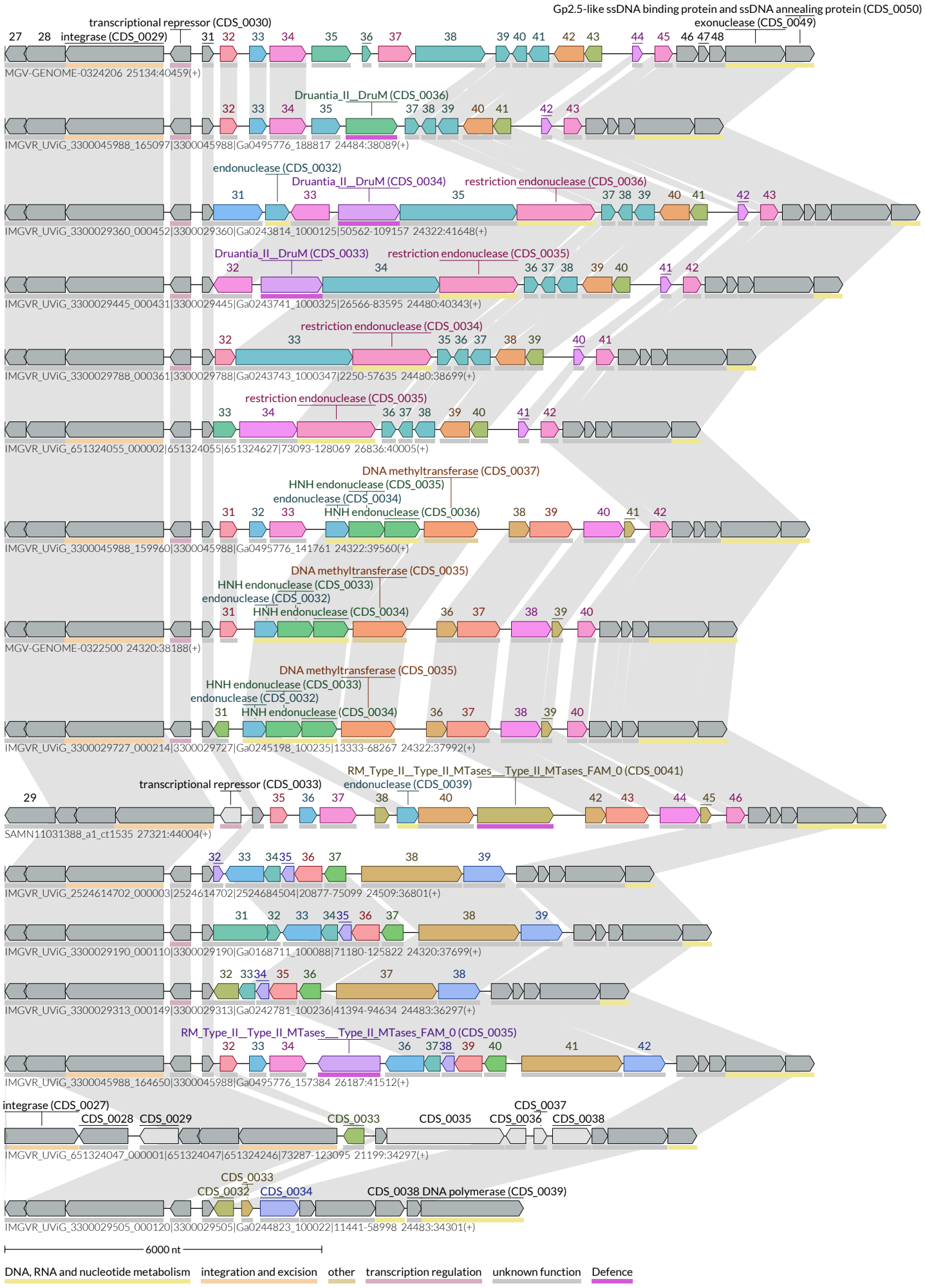
Gp2.5-like ssDNA binding protein and ssDNA annealing protein (CDS_0050)
transcriptional repressor (CDS_0030)
exonuclease (CDS_0049)
integrase (CDS_0029)
MGV-GENOME-0324206 25134:40459(+)
Druantia_II__DruM (CDS_0036)
IMGVR_UViG_3300045988_165097|3300045988|Ga0495776_188817 24484:38089(+)
endonuclease (CDS_0032)
Druantia_II__DruM (CDS_0034)
restriction endonuclease (CDS_0036)
IMGVR_UViG_3300029360_000452|3300029360|Ga0243814_1000125|50562-109157 24322:41648(+)
Druantia_II__DruM (CDS_0033)
restriction endonuclease (CDS_0035)
IMGVR_UViG_3300029445_000431|3300029445|Ga0243741_1000325|26566-83595 24480:40343(+)
restriction endonuclease (CDS_0034)
IMGVR_UViG_3300029788_000361|3300029788|Ga0243743_1000347|2250-57635 24480:38699(+)
restriction endonuclease (CDS_0035)
IMGVR_UViG_651324055_000002|651324055|651324627|73093-128069 26836:40005(+)
DNA methyltransferase (CDS_0037)
HNH endonuclease (CDS_0035)
endonuclease (CDS_0034)
HNH endonuclease (CDS_0036)
IMGVR_UViG_3300045988_159960|3300045988|Ga0495776_141761 24322:39560(+)
DNA methyltransferase (CDS_0035)
HNH endonuclease (CDS_0033)
endonuclease (CDS_0032)
HNH endonuclease (CDS_0034)
MGV-GENOME-0322500 24320:38188(+)
DNA methyltransferase (CDS_0035)
HNH endonuclease (CDS_0033)
endonuclease (CDS_0032)
HNH endonuclease (CDS_0034)
IMGVR_UViG_3300029727_000214|3300029727|Ga0245198_100235|13333-68267 24322:37992(+)
RM_Type_II__Type_II_MTases___Type_II_MTases_FAM_0 (CDS_0041)
transcriptional repressor (CDS_0033)
endonuclease (CDS_0039)
SAMN11031388_a1_ct1535 27321:44004(+)
IMGVR_UViG_2524614702_000003|2524614702|2524684504|20877-75099 24509:36801(+)
IMGVR_UViG_3300029190_000110|3300029190|Ga0168711_100088|71180-125822 24320:37699(+)
IMGVR_UViG_3300029313_000149|3300029313|Ga0242781_100236|41394-94634 24483:36297(+)
RM_Type_II__Type_II_MTases___Type_II_MTases_FAM_0 (CDS_0035)
IMGVR_UViG_3300045988_164650|3300045988|Ga0495776_157384 26187:41512(+)
integrase (CDS_0027)
CDS_0028
CDS_0029
CDS_0033
CDS_0035
CDS_0037
CDS_0036
CDS_0038
IMGVR_UViG_651324047_000001|651324047|651324246|73287-123095 21199:34297(+)
CDS_0033
CDS_0032
CDS_0034
CDS_0038 DNA polymerase (CDS_0039)
IMGVR_UViG_3300029505_000120|3300029505|Ga0244823_100022|11441-58998 24483:34301(+)
6000 nt
DNA, RNA and nucleotide metabolism
integration and excision
other
transcription regulation
unknown function
Defence