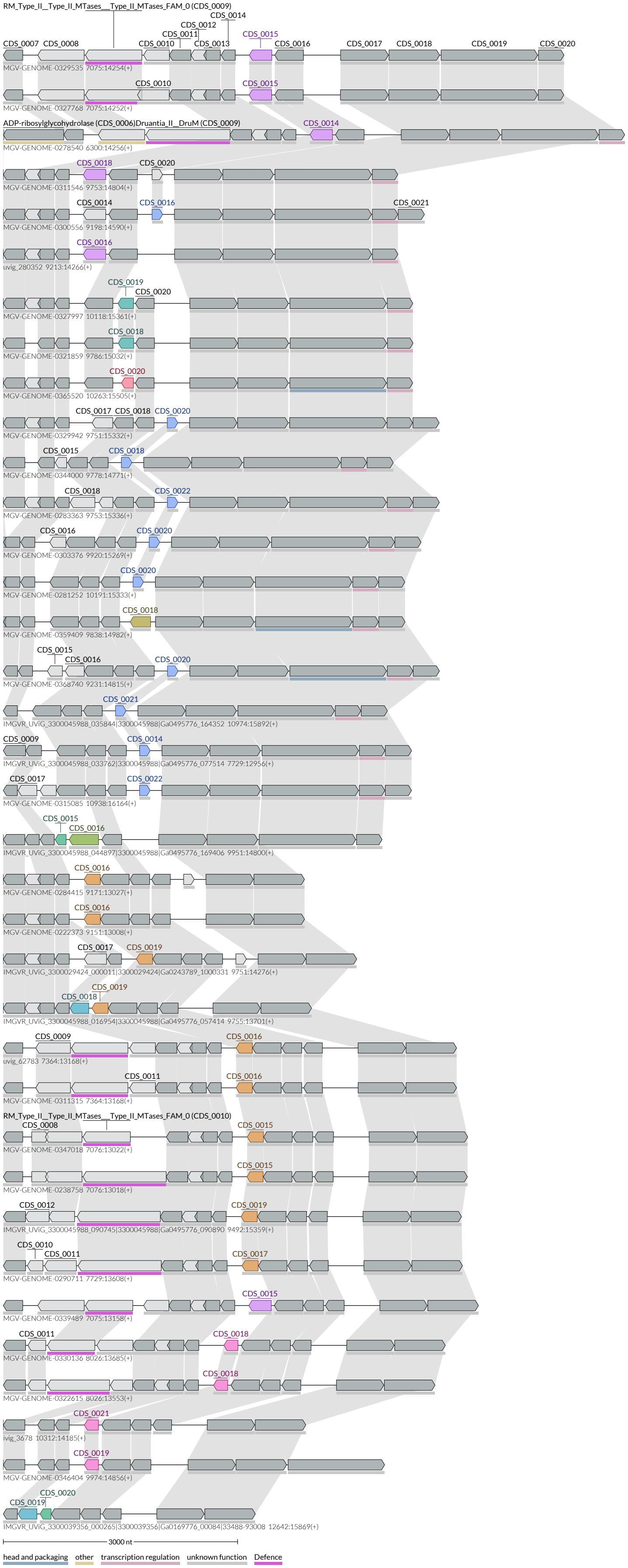
RM_Type_II__Type_II_MTases___Type_II_MTases_FAM_0 (CDS_0009)
CDS_0014
CDS_0012
CDS_0011
CDS_0015
CDS_0007 CDS_0008 CDS_0010 CDS_0013 CDS_0016 CDS_0017 CDS_0018 CDS_0019 CDS_0020
MGV-GENOME-0329535 7075:14254(+)
CDS_0010 CDS_0015
MGV-GENOME-0327768 7075:14252(+)
ADP-ribosylglycohydrolase (CDS_0006)Druantia_II__DruM (CDS_0009)
CDS_0014
MGV-GENOME-0278540 6300:14256(+)
CDS_0018 CDS_0020
MGV-GENOME-0311546 9753:14804(+)
CDS_0014 CDS_0016 CDS_0021
MGV-GENOME-0300556 9198:14590(+)
CDS_0016
uvig_280352 9213:14266(+)
CDS_0019 CDS_0020
MGV-GENOME-0327997 10118:15361(+)
CDS_0018
MGV-GENOME-0321859 9786:15032(+)
CDS_0020
MGV-GENOME-0365520 10263:15505(+)
CDS_0017 CDS_0018 CDS_0020
MGV-GENOME-0329942 9751:15332(+)
CDS_0015 CDS_0018
MGV-GENOME-0344000 9778:14771(+)
CDS_0018 CDS_0022
MGV-GENOME-0283363 9753:15336(+)
CDS_0016 CDS_0020
MGV-GENOME-0303376 9920:15269(+)
CDS_0020
MGV-GENOME-0281252 10191:15333(+)
CDS_0018
MGV-GENOME-0359409 9838:14982(+)
CDS_0015 CDS_0016 CDS_0020
MGV-GENOME-0368740 9231:14815(+)
CDS_0021
IMGVR_UViG_3300045988_035844|3300045988|Ga0495776_164352 10974:15892(+)
CDS_0009 CDS_0014
IMGVR_UViG_3300045988_033762|3300045988|Ga0495776_077514 7729:12956(+)
CDS_0017 CDS_0022
MGV-GENOME-0315085 10938:16164(+)
CDS_0015 CDS_0016
IMGVR_UViG_3300045988_044897|3300045988|Ga0495776_169406 9951:14800(+)
CDS_0016
MGV-GENOME-0284415 9171:13027(+)
CDS_0016
MGV-GENOME-0222373 9151:13008(+)
CDS_0017 CDS_0019
IMGVR_UViG_3300029424_000011|3300029424|Ga0243789_1000331 9751:14276(+)
CDS_0019 CDS_0018
IMGVR_UViG_3300045988_016954|3300045988|Ga0495776_057414 9755:13701(+)
CDS_0009 CDS_0016
uvig_62783 7364:13168(+)
CDS_0011 CDS_0016
MGV-GENOME-0311315 7364:13168(+)
RM_Type_II__Type_II_MTases___Type_II_MTases_FAM_0 (CDS_0010)
CDS_0008 CDS_0015
MGV-GENOME-0347018 7076:13022(+)
CDS_0015
MGV-GENOME-0238758 7076:13018(+)
CDS_0012 CDS_0019
IMGVR_UViG_3300045988_090745|3300045988|Ga0495776_090890 9492:15359(+)
CDS_0010 CDS_0011 CDS_0017
MGV-GENOME-0290711 7729:13608(+)
CDS_0015
MGV-GENOME-0339489 7075:13158(+)
CDS_0011 CDS_0018
MGV-GENOME-0330136 8026:13685(+)
CDS_0018
MGV-GENOME-0322615 8026:13553(+)
CDS_0021
ivig_3678 10312:14185(+)
CDS_0019
MGV-GENOME-0346404 9974:14856(+)
CDS_0020 CDS_0019
IMGVR_UViG_3300039356_000265|3300039356|Ga0169776_00084|33488-93008 12642:15869(+)
3000 nt
head and packaging
other
transcription regulation
unknown function
Defence