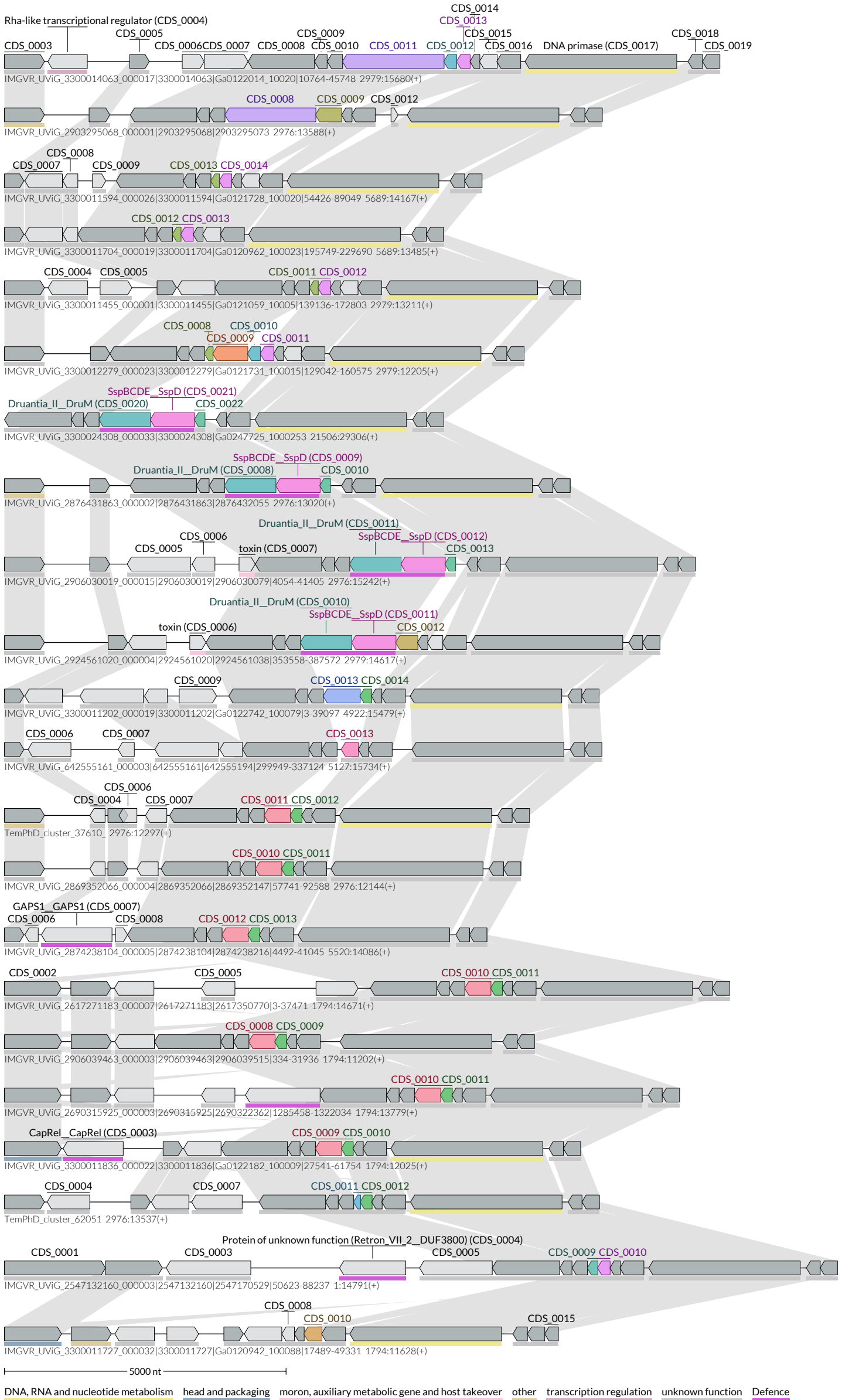
Rha-like transcriptional regulator (CDS_0004)
CDS_0003
CDS_0005
CDS_0006
CDS_0007
CDS_0008
CDS_0009
CDS_0010
CDS_0011
CDS_0012
CDS_0013
CDS_0014
CDS_0015
CDS_0016
DNA primase (CDS_0017)
CDS_0018
CDS_0019
IMGVR_UViG_3300014063_000017|3300014063|Ga0122014_10020|10764-45748 2979:15680(+)
CDS_0008
CDS_0009
CDS_0012
IMGVR_UViG_2903295068_000001|2903295068|2903295073 2976:13588(+)
CDS_0008
CDS_0007
CDS_0009
CDS_0013
CDS_0014
IMGVR_UViG_3300011594_000026|3300011594|Ga0121728_100020|54426-89049 5689:14167(+)
CDS_0012
CDS_0013
IMGVR_UViG_3300011704_000019|3300011704|Ga0120962_100023|195749-229690 5689:13485(+)
CDS_0004
CDS_0005
CDS_0011
CDS_0012
IMGVR_UViG_3300011455_000001|3300011455|Ga0121059_10005|139136-172803 2979:13211(+)
CDS_0008
CDS_0010
CDS_0009
CDS_0011
IMGVR_UViG_3300012279_000023|3300012279|Ga0121731_100015|129042-160575 2979:12205(+)
SspBCDE__SspD (CDS_0021)
Druantia_II__DruM (CDS_0020)
CDS_0022
IMGVR_UViG_3300024308_000033|3300024308|Ga0247725_1000253 21506:29306(+)
SspBCDE__SspD (CDS_0009)
Druantia_II__DruM (CDS_0008)
CDS_0010
IMGVR_UViG_2876431863_000002|2876431863|2876432055 2976:13020(+)
Druantia_II__DruM (CDS_0011)
CDS_0006
SspBCDE__SspD (CDS_0012)
CDS_0005
toxin (CDS_0007)
CDS_0013
IMGVR_UViG_2906030019_000015|2906030019|2906030079|4054-41405 2976:15242(+)
Druantia_II__DruM (CDS_0010)
SspBCDE__SspD (CDS_0011)
toxin (CDS_0006)
CDS_0012
IMGVR_UViG_2924561020_000004|2924561020|2924561038|353558-387572 2979:14617(+)
CDS_0009
CDS_0013
CDS_0014
IMGVR_UViG_3300011202_000019|3300011202|Ga0122742_100079|3-39097 4922:15479(+)
CDS_0006
CDS_0007
CDS_0013
IMGVR_UViG_642555161_000003|642555161|642555194|299949-337124 5127:15734(+)
CDS_0006
CDS_0004
CDS_0007
CDS_0011
CDS_0012
TemPhD_cluster_37610_ 2976:12297(+)
CDS_0010
CDS_0011
IMGVR_UViG_2869352066_000004|2869352066|2869352147|57741-92588 2976:12144(+)
GAPS1__GAPS1 (CDS_0007)
CDS_0006
CDS_0008
CDS_0012
CDS_0013
IMGVR_UViG_2874238104_000005|2874238104|2874238216|4492-41045 5520:14086(+)
CDS_0002
CDS_0005
CDS_0010
CDS_0011
IMGVR_UViG_2617271183_000007|2617271183|2617350770|3-37471 1794:14671(+)
CDS_0008
CDS_0009
IMGVR_UViG_2906039463_000003|2906039463|2906039515|334-31936 1794:11202(+)
CDS_0010
CDS_0011
IMGVR_UViG_2690315925_000003|2690315925|2690322362|1285458-1322034 1794:13779(+)
CapRel__CapRel (CDS_0003)
CDS_0009
CDS_0010
IMGVR_UViG_3300011836_000022|3300011836|Ga0122182_100009|27541-61754 1794:12025(+)
CDS_0004
CDS_0007
CDS_0011
CDS_0012
TemPhD_cluster_62051 2976:13537(+)
Protein of unknown function (Retron_VII_2__DUF3800) (CDS_0004)
CDS_0001
CDS_0003
CDS_0005
CDS_0009
CDS_0010
IMGVR_UViG_2547132160_000003|2547132160|2547170529|50623-88237 1:14791(+)
CDS_0008
CDS_0010
CDS_0015
IMGVR_UViG_3300011727_000032|3300011727|Ga0120942_100088|17489-49331 1794:11628(+)
5000 nt
DNA, RNA and nucleotide metabolism
head and packaging
moron, auxiliary metabolic gene and host takeover
other
transcription regulation
unknown function
Defence