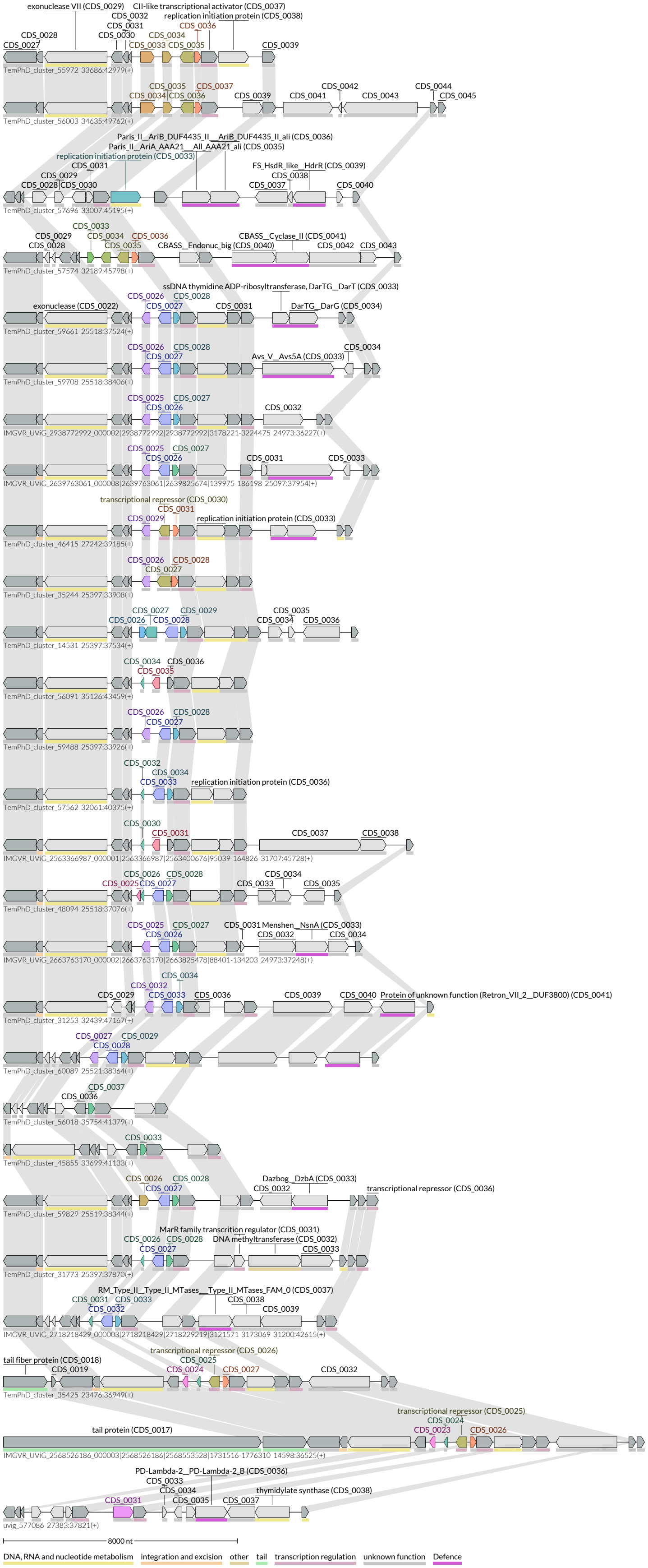
exonuclease VII (CDS_0029)
CII-like transcriptional activator (CDS_0037)
CDS_0032
replication initiation protein (CDS_0038)
CDS_0031
CDS_0036
CDS_0028
CDS_0030
CDS_0034
CDS_0027
CDS_0033
CDS_0035
CDS_0039
TemPhD_cluster_55972 33686:42979(+)
CDS_0035
CDS_0037
CDS_0042
CDS_0044
CDS_0034
CDS_0036
CDS_0039
CDS_0041
CDS_0043
CDS_0045
TemPhD_cluster_56003 34635:49762(+)
Paris_II__AriB_DUF4435_II__AriB_DUF4435_II_ali (CDS_0036)
Paris_II__AriA_AAA21__AII_AAA21_ali (CDS_0035)
replication initiation protein (CDS_0033)
CDS_0031
FS_HsdR_like__HdrR (CDS_0039)
CDS_0029
CDS_0038
CDS_0028
CDS_0030
CDS_0037
CDS_0040
TemPhD_cluster_57696 33007:45195(+)
CDS_0033
CDS_0029
CDS_0034
CDS_0036
CBASS__Cyclase_II (CDS_0041)
CDS_0028
CDS_0035
CBASS__Endonuc_big (CDS_0040)
CDS_0042
CDS_0043
TemPhD_cluster_57574 32189:45798(+)
ssDNA thimidine ADP-ribosyltransferase, DarTG__DarT (CDS_0033)
CDS_0026
CDS_0028
exonuclease (CDS_0022)
CDS_0027
CDS_0031
DarTG__DarG (CDS_0034)
TemPhD_cluster_59661 25518:37524(+)
CDS_0026
CDS_0028
CDS_0034
CDS_0027
Avs_V__Avs5A (CDS_0033)
TemPhD_cluster_59708 25518:38406(+)
CDS_0025
CDS_0027
CDS_0026
CDS_0032
IMGVR_UViG_2938772992_000002|2938772992|2938772992|3178221-3224475 24973:36227(+)
CDS_0025
CDS_0027
CDS_0026
CDS_0031
CDS_0033
IMGVR_UViG_2639763061_000008|2639763061|2639825674|139975-186198 25097:37954(+)
transcriptional repressor (CDS_0030)
CDS_0031
CDS_0029
replication initiation protein (CDS_0033)
TemPhD_cluster_46415 27242:39185(+)
CDS_0026
CDS_0028
CDS_0027
TemPhD_cluster_35244 25397:33908(+)
CDS_0027
CDS_0029
CDS_0035
CDS_0026
CDS_0028
CDS_0034
CDS_0036
TemPhD_cluster_14531 25397:37534(+)
CDS_0034
CDS_0036
CDS_0035
TemPhD_cluster_56091 35126:43459(+)
CDS_0026
CDS_0028
CDS_0027
TemPhD_cluster_59488 25397:33926(+)
CDS_0032
CDS_0034
CDS_0033
replication initiation protein (CDS_0036)
TemPhD_cluster_57562 32061:40375(+)
CDS_0030
CDS_0031
CDS_0037
CDS_0038
IMGVR_UViG_2563366987_000001|2563366987|2563400676|95039-164826 31707:45728(+)
CDS_0026
CDS_0028
CDS_0034
CDS_0025
CDS_0027
CDS_0033
CDS_0035
TemPhD_cluster_48094 25518:37076(+)
CDS_0025
CDS_0027
CDS_0031
Menshen__NsnA (CDS_0033)
CDS_0026
CDS_0032
CDS_0034
IMGVR_UViG_2663763170_000002|2663763170|2663825478|88401-134203 24973:37248(+)
CDS_0034
CDS_0032
CDS_0029
CDS_0033
CDS_0036
CDS_0039
CDS_0040
Protein of unknown function (Retron_VII_2__DUF3800) (CDS_0041)
TemPhD_cluster_31253 32439:47167(+)
CDS_0027
CDS_0029
CDS_0028
TemPhD_cluster_60089 25521:38364(+)
CDS_0037
CDS_0036
TemPhD_cluster_56018 35754:41379(+)
CDS_0033
TemPhD_cluster_45855 33699:41133(+)
CDS_0026
CDS_0028
Dazbog__DzbA (CDS_0033)
CDS_0027
CDS_0032
transcriptional repressor (CDS_0036)
TemPhD_cluster_59829 25519:38344(+)
MarR family transcrition regulator (CDS_0031)
CDS_0026
CDS_0028
DNA methyltransferase (CDS_0032)
CDS_0027
CDS_0033
TemPhD_cluster_31773 25397:37870(+)
RM_Type_II__Type_II_MTases__Type_II_MTases_FAM_0 (CDS_0037)
CDS_0031
CDS_0033
CDS_0038
CDS_0032
CDS_0039
IMGVR_UViG_2718218429_000003|2718218429|2718229219|3121571-3173069 31200:42615(+)
transcriptional repressor (CDS_0026)
tail fiber protein (CDS_0018)
CDS_0025
CDS_0019
CDS_0024
CDS_0027
CDS_0032
TemPhD_cluster_35425 23476:36949(+)
transcriptional repressor (CDS_0025)
CDS_0024
tail protein (CDS_0017)
CDS_0023
CDS_0026
IMGVR_UViG_2568526186_000003|2568526186|2568553528|1731516-1776310 14598:36525(+)
PD-Lambda-2__PD-Lambda-2_B (CDS_0036)
CDS_0033
CDS_0034
thymidylate synthase (CDS_0038)
CDS_0031
CDS_0035
CDS_0037
uvig_577086 27383:37821(+)
8000 nt
DNA, RNA and nucleotide metabolism
integration and excision
other
tail
transcription regulation
unknown function
Defence