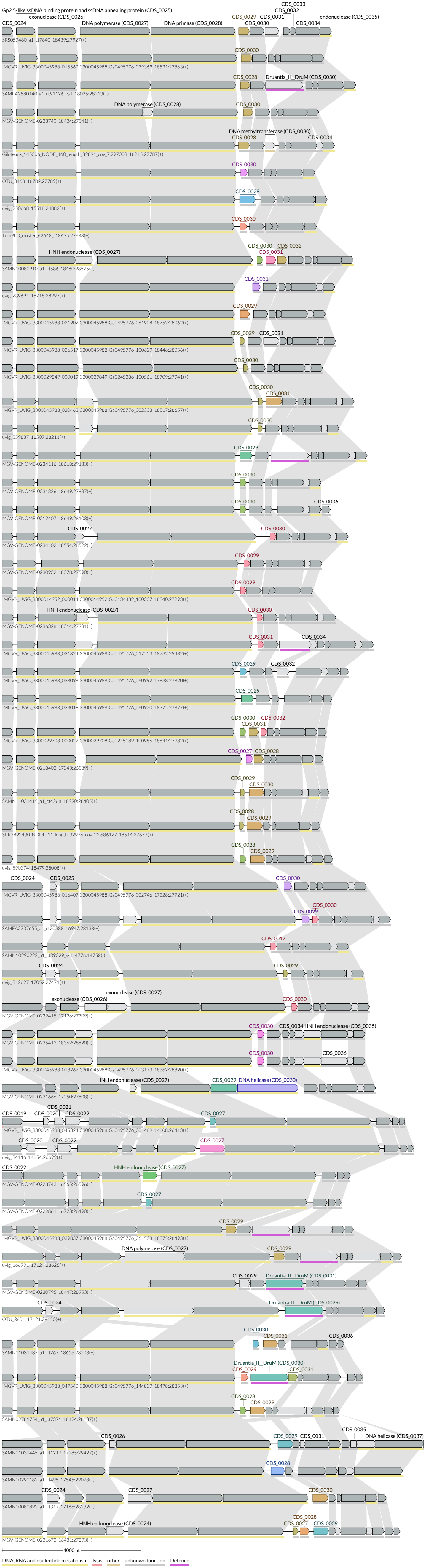
Gp2.5-like ssDNA binding protein and ssDNA annealing protein (CDS_0025)
exonuclease (CDS_0026)
CDS_0024
DNA polymerase (CDS_0027)
DNA primase (CDS_0028)
CDS_0029
CDS_0031
CDS_0033
CDS_0032
CDS_0030
CDS_0034
endonuclease (CDS_0035)
SRS057480_a1_ct7840 18439:27927(+)
CDS_0030
IMGVR_UVIG_3300045988_015560|3300045988|Ga0495776_079369 18591:27863(+)
CDS_0028
Druantia_II__DruM (CDS_0030)
SAMEA2580140_a1_ct91126_vs1 18025:28213(+)
DNA polymerase (CDS_0028)
CDS_0030
MGV-GENOME-0223740 18424:27541(+)
DNA methyltransferase (CDS_0030)
CDS_0028
CDS_0034
Giloteaux_145306_NODE_460_length_32891_cov_7.297003 18215:27787(+)
CDS_0030
OTU_3468 18782:27789(+)
CDS_0028
uvig_250668 15518:24882(+)
CDS_0030
TemPhD_cluster_62648_ 18635:27688(+)
CDS_0030
CDS_0032
HNH endonuclease (CDS_0027)
CDS_0031
SAMN10080910_a1_ct586 18460:28575(+)
CDS_0031
uvig_239694 18718:28297(+)
CDS_0029
IMGVR_UVIG_3300045988_021902|3300045988|Ga0495776_061908 18752:28062(+)
CDS_0029
CDS_0031
IMGVR_UVIG_3300045988_026517|3300045988|Ga0495776_100629 18446:28056(+)
CDS_0030
IMGVR_UVIG_3300029849_000019|3300029849|Ga0245286_100561 18709:27941(+)
CDS_0030
CDS_0031
IMGVR_UVIG_3300045988_020463|3300045988|Ga0495776_002303 18517:28657(+)
CDS_0030
uvig_559837 18507:28211(+)
CDS_0029
MGV-GENOME-0234116 18618:29133(+)
CDS_0030
MGV-GENOME-0231326 18649:27837(+)
CDS_0030
CDS_0036
MGV-GENOME-0212407 18649:28103(+)
CDS_0027
CDS_0030
MGV-GENOME-0234102 18554:28522(+)
CDS_0029
MGV-GENOME-0230932 18378:27590(+)
CDS_0029
IMGVR_UVIG_3300014952_000014|3300014952|Ga0134432_100337 18340:27293(+)
HNH endonuclease (CDS_0027)
CDS_0030
MGV-GENOME-0236328 18314:27931(+)
CDS_0031
CDS_0034
IMGVR_UVIG_3300045988_021824|3300045988|Ga0495776_017553 18732:29432(+)
CDS_0029
CDS_0032
IMGVR_UVIG_3300045988_028098|3300045988|Ga0495776_060992 17838:27820(+)
CDS_0029
IMGVR_UVIG_3300045988_023019|3300045988|Ga0495776_060920 18375:27877(+)
CDS_0030
CDS_0032
CDS_0031
IMGVR_UVIG_3300029708_000027|3300029708|Ga0245189_100986 18641:27982(+)
CDS_0027
CDS_0028
MGV-GENOME-0218403 17343:26589(+)
CDS_0029
CDS_0030
SAMN11031415_a1_ct4268 18990:28405(+)
CDS_0028
CDS_0029
SRR7892430_NODE_11_length_32976_cov_22.686127 18514:27677(+)
CDS_0028
CDS_0029
uvig_590374 18479:28008(+)
CDS_0024
CDS_0025
CDS_0030
IMGVR_UVIG_3300045988_016407|3300045988|Ga0495776_002746 17228:27721(+)
CDS_0030
CDS_0029
SAMEA2737655_a1_ct20388 16947:28138(+)
CDS_0017
SAMN10290222_a1_ct39229_vs1 4776:14758(-)
CDS_0024
CDS_0029
uvig_312627 17052:27471(+)
exonuclease (CDS_0027)
exonuclease (CDS_0026)
CDS_0030
MGV-GENOME-0232415 17126:27709(+)
CDS_0030
CDS_0034 HNH endonuclease (CDS_0035)
MGV-GENOME-0235412 18362:28820(+)
CDS_0030
CDS_0036
IMGVR_UVIG_3300045988_018262|3300045988|Ga0495776_003173 18362:28820(+)
HNH endonuclease (CDS_0027)
CDS_0029
DNA helicase (CDS_0030)
MGV-GENOME-0231666 17050:27808(+)
CDS_0021
CDS_0019
CDS_0020
CDS_0022
CDS_0027
IMGVR_UVIG_3300045988_045324|3300045988|Ga0495776_001489 14808:26413(+)
CDS_0020
CDS_0022
CDS_0027
uvig_34116 14854:26699(+)
CDS_0022
HNH endonuclease (CDS_0027)
MGV-GENOME-0228743 16565:26596(+)
CDS_0027
MGV-GENOME-0229861 16723:26490(+)
CDS_0029
IMGVR_UVIG_3300045988_039837|3300045988|Ga0495776_061533 18375:28493(+)
DNA polymerase (CDS_0027)
CDS_0029
uvig_166791 17124:28625(+)
CDS_0029
Druantia_II__DruM (CDS_0031)
MGV-GENOME-0230795 18447:28953(+)
CDS_0024
Druantia_II__DruM (CDS_0029)
OTU_3601 17121:28150(+)
CDS_0030
CDS_0031
CDS_0036
SAMN11031437_a1_ct267 18656:28503(+)
Druantia_II__DruM (CDS_0030)
CDS_0029
CDS_0031
IMGVR_UVIG_3300045988_047540|3300045988|Ga0495776_144837 18478:28853(+)
CDS_0028
CDS_0029
SAMN09781754_a1_ct7371 18424:28137(+)
CDS_0026
CDS_0029
CDS_0031
CDS_0035
DNA helicase (CDS_0037)
SAMN11031445_a1_ct1217 17285:29427(+)
CDS_0028
SAMN10290182_a1_ct495 17545:29078(+)
CDS_0024
CDS_0027
CDS_0030
SAMN10080892_a1_ct317 17166:28232(+)
CDS_0028
HNH endonuclease (CDS_0024)
CDS_0027
CDS_0029
MGV-GENOME-0221672 16431:27893(+)
4000 nt
DNA, RNA and nucleotide metabolism
lysis
other
unknown function
Defence