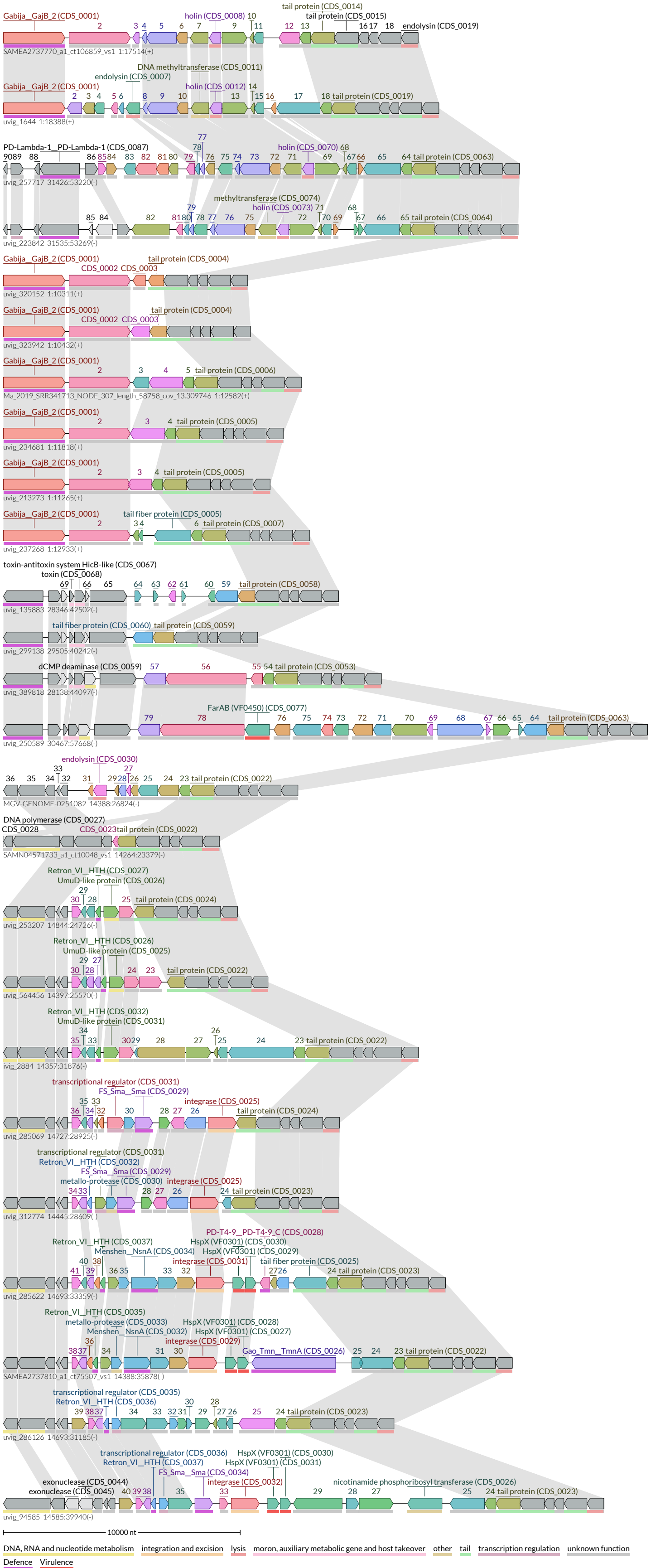
Gabija__GajB_2 (CDS_0001)
holin (CDS_0008)
tail protein (CDS_0014)
tail protein (CDS_0015)
endolysin (CDS_0019)
SAMEA2737770_a1_ct106859_vs1 1:17514(+)
DNA methyltransferase (CDS_0011)
endolysin (CDS_0007)
Gabija__GajB_2 (CDS_0001)
holin (CDS_0012)
18 tail protein (CDS_0019)
uvig_1644 1:18388(+)
PD-Lambda-1__PD-Lambda-1 (CDS_0087)
holin (CDS_0070)
64 tail protein (CDS_0063)
uvig_257717 31426:53220(-)
methyltransferase (CDS_0074)
holin (CDS_0073)
65 tail protein (CDS_0064)
uvig_223842 31535:53269(-)
Gabija__GajB_2 (CDS_0001)
tail protein (CDS_0004)
CDS_0002 CDS_0003
uvig_320152 1:10311(+)
Gabija__GajB_2 (CDS_0001)
tail protein (CDS_0004)
CDS_0002 CDS_0003
uvig_323942 1:10432(+)
Gabija__GajB_2 (CDS_0001)
5 tail protein (CDS_0006)
Ma_2019_SRR341713_NODE_307_length_58758_cov_13.309746 1:12582(+)
Gabija__GajB_2 (CDS_0001)
4 tail protein (CDS_0005)
uvig_234681 1:11818(+)
Gabija__GajB_2 (CDS_0001)
4 tail protein (CDS_0005)
uvig_213273 1:11265(+)
Gabija__GajB_2 (CDS_0001)
tail fiber protein (CDS_0005)
6 tail protein (CDS_0007)
uvig_237268 1:12933(+)
toxin-antitoxin system HicB-like (CDS_0067)
toxin (CDS_0068)
tail protein (CDS_0058)
uvig_135883 28346:42502(-)
tail fiber protein (CDS_0060)
tail protein (CDS_0059)
uvig_299138 29505:40242(-)
dCMP deaminase (CDS_0059)
55 54 tail protein (CDS_0053)
uvig_389818 28138:44097(-)
FarAB (VF0450) (CDS_0077)
tail protein (CDS_0063)
uvig_250589 30467:57668(-)
endolysin (CDS_0030)
23 tail protein (CDS_0022)
MGV-GENOME-0251082 14388:26824(-)
DNA polymerase (CDS_0027)
CDS_0028
CDS_0023tail protein (CDS_0022)
SAMN04571733_a1_ct10048_vs1 14264:23379(-)
Retron_VI__HTH (CDS_0027)
UmuD-like protein (CDS_0026)
tail protein (CDS_0024)
uvig_253207 14844:24726(-)
Retron_VI__HTH (CDS_0026)
UmuD-like protein (CDS_0025)
tail protein (CDS_0022)
uvig_564456 14397:25570(-)
Retron_VI__HTH (CDS_0032)
UmuD-like protein (CDS_0031)
23 tail protein (CDS_0022)
ivig_2884 14357:31876(-)
transcriptional regulator (CDS_0031)
FS_Sma__Sma (CDS_0029)
integrase (CDS_0025)
tail protein (CDS_0024)
uvig_285069 14727:28925(-)
transcriptional regulator (CDS_0031)
Retron_VI__HTH (CDS_0032)
FS_Sma__Sma (CDS_0029)
metallo-protease (CDS_0030)
integrase (CDS_0025)
24 tail protein (CDS_0023)
uvig_312774 14445:28609(-)
PD-T4-9__PD-T4-9_C (CDS_0028)
Retron_VI__HTH (CDS_0037)
HspX (VF0301) (CDS_0030)
Menshen__NsnA (CDS_0034)
HspX (VF0301) (CDS_0029)
integrase (CDS_0031)
tail fiber protein (CDS_0025)
24 tail protein (CDS_0023)
uvig_285622 14693:33359(-)
Retron_VI__HTH (CDS_0035)
metallo-protease (CDS_0033)
HspX (VF0301) (CDS_0028)
Menshen__NsnA (CDS_0032)
HspX (VF0301) (CDS_0027)
integrase (CDS_0029)
Gao_Tmn__TmnA (CDS_0026)
23 tail protein (CDS_0022)
SAMEA2737810_a1_ct75507_vs1 14388:35878(-)
transcriptional regulator (CDS_0035)
Retron_VI__HTH (CDS_0036)
24 tail protein (CDS_0023)
uvig_286126 14693:31185(-)
transcriptional regulator (CDS_0036)
HspX (VF0301) (CDS_0030)
Retron_VI__HTH (CDS_0037)
HspX (VF0301) (CDS_0031)
FS_Sma__Sma (CDS_0034)
exonuclease (CDS_0044)
integrase (CDS_0032)
nicotinamide phosphoribosyl transferase (CDS_0026)
exonuclease (CDS_0045)
24 tail protein (CDS_0023)
uvig_94585 14585:39940(-)
10000 nt
DNA, RNA and nucleotide metabolism
integration and excision
lysis
moron, auxiliary metabolic gene and host takeover
other
tail
transcription regulation
unknown function
Defence
Virulence