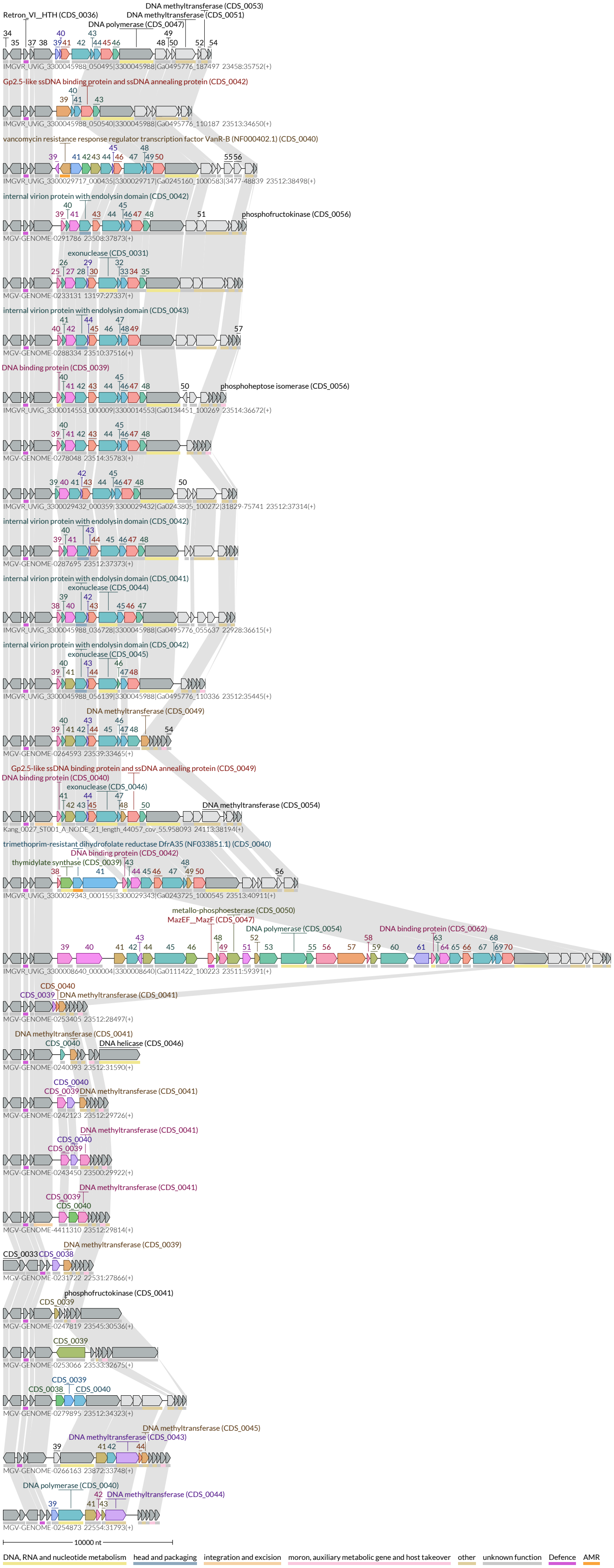
DNA methyltransferase (CDS_0053)
Retron_VI__HTH (CDS_0036)
DNA methyltransferase (CDS_0051)
DNA polymerase (CDS_0047)
IMGVR_UViG_3300045988_050495|3300045988|Ga0495776_187497 23458:35752(+)
Gp2.5-like ssDNA binding protein and ssDNA annealing protein (CDS_0042)
IMGVR_UViG_3300045988_050540|3300045988|Ga0495776_110187 23513:34650(+)
vancomycin resistance response regulator transcription factor VanR-B (NF000402.1) (CDS_0040)
IMGVR_UViG_3300029717_000435|3300029717|Ga0245160_1000583|3477-48839 23512:38498(+)
internal virion protein with endolysin domain (CDS_0042)
phosphofructokinase (CDS_0056)
MGV-GENOME-0291786 23508:37873(+)
exonuclease (CDS_0031)
MGV-GENOME-0233131 13197:27337(+)
internal virion protein with endolysin domain (CDS_0043)
MGV-GENOME-0288334 23510:37516(+)
DNA binding protein (CDS_0039)
phosphoheptose isomerase (CDS_0056)
IMGVR_UViG_3300014553_000009|3300014553|Ga0134451_100269 23514:36672(+)
MGV-GENOME-0278048 23514:35783(+)
IMGVR_UViG_3300029432_000359|3300029432|Ga0243805_100272|31829-75741 23512:37314(+)
internal virion protein with endolysin domain (CDS_0042)
MGV-GENOME-0287695 23512:37373(+)
internal virion protein with endolysin domain (CDS_0041)
exonuclease (CDS_0044)
IMGVR_UViG_3300045988_036728|3300045988|Ga0495776_055637 22928:36615(+)
internal virion protein with endolysin domain (CDS_0042)
exonuclease (CDS_0045)
IMGVR_UViG_3300045988_056139|3300045988|Ga0495776_110336 23512:35445(+)
DNA methyltransferase (CDS_0049)
MGV-GENOME-0264593 23539:33465(+)
Gp2.5-like ssDNA binding protein and ssDNA annealing protein (CDS_0049)
DNA binding protein (CDS_0040)
exonuclease (CDS_0046)
DNA methyltransferase (CDS_0054)
Kang_0027_ST001_A_NODE_21_length_44057_cov_55.958093 24113:38194(+)
trimethoprim-resistant dihydrofolate reductase DfrA35 (NF033851.1) (CDS_0040)
DNA binding protein (CDS_0042)
thymidylate synthase (CDS_0039)
IMGVR_UViG_3300029343_000155|3300029343|Ga0243725_1000545 23513:40911(+)
metallo-phosphoesterase (CDS_0050)
MazEF__MazF (CDS_0047)
DNA polymerase (CDS_0054)
DNA binding protein (CDS_0062)
IMGVR_UViG_3300008640_000004|3300008640|Ga0111422_100223 23511:59391(+)
CDS_0040
CDS_0039
DNA methyltransferase (CDS_0041)
MGV-GENOME-0253405 23512:28497(+)
DNA methyltransferase (CDS_0041)
CDS_0040
DNA helicase (CDS_0046)
MGV-GENOME-0240093 23512:31590(+)
CDS_0040
CDS_0039
DNA methyltransferase (CDS_0041)
MGV-GENOME-0242123 23512:29726(+)
DNA methyltransferase (CDS_0041)
CDS_0040
CDS_0039
MGV-GENOME-0243450 23500:29922(+)
DNA methyltransferase (CDS_0041)
CDS_0039
CDS_0040
MGV-GENOME-4411310 23512:29814(+)
DNA methyltransferase (CDS_0039)
CDS_0033
CDS_0038
MGV-GENOME-0231722 22531:27866(+)
phosphofructokinase (CDS_0041)
CDS_0039
MGV-GENOME-0247819 23545:30536(+)
CDS_0039
MGV-GENOME-0253066 23533:32675(+)
CDS_0039
CDS_0038
CDS_0040
MGV-GENOME-0279895 23512:34323(+)
DNA methyltransferase (CDS_0045)
DNA methyltransferase (CDS_0043)
MGV-GENOME-0266163 23872:33748(+)
DNA polymerase (CDS_0040)
DNA methyltransferase (CDS_0044)
MGV-GENOME-0254873 22554:31793(+)
10000 nt
DNA, RNA and nucleotide metabolism
head and packaging
integration and excision
moron, auxiliary metabolic gene and host takeover
other
unknown function
Defence
AMR