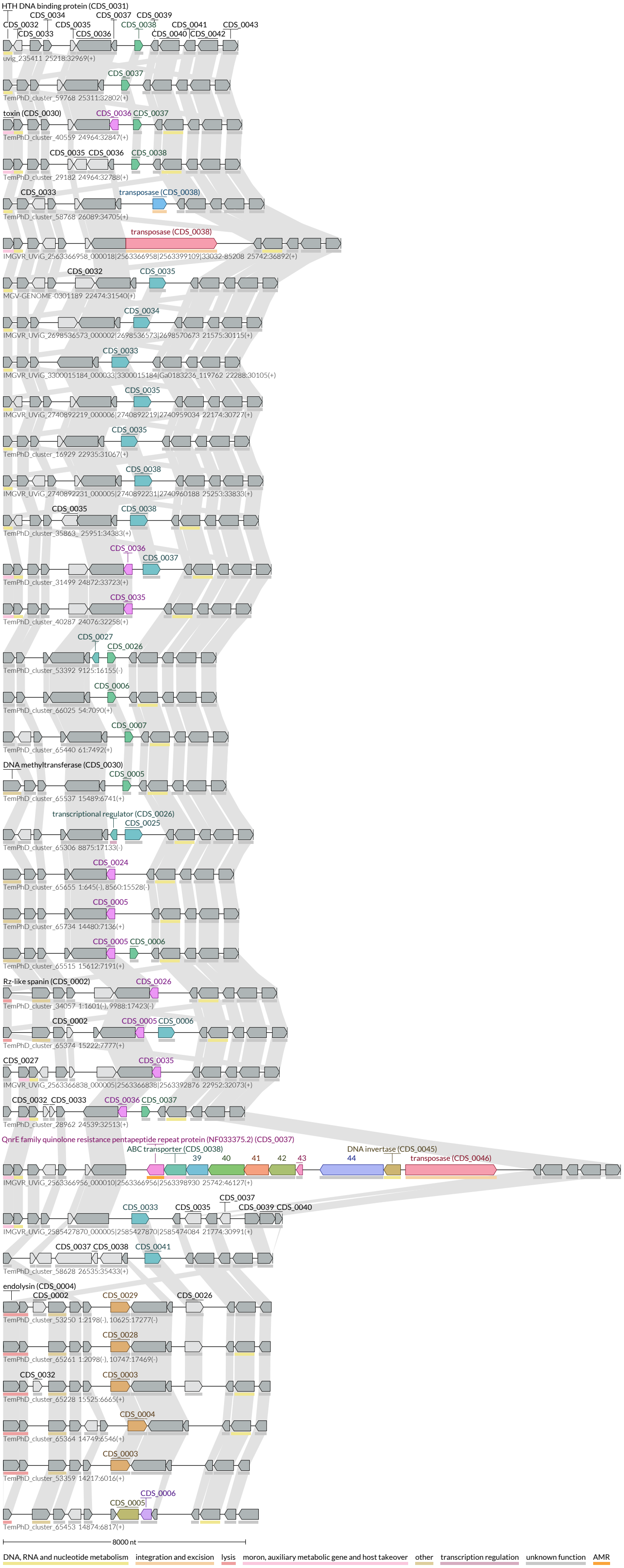
HTH DNA binding protein (CDS_0031)
CDS_0034
CDS_0037
CDS_0039
CDS_0032
CDS_0035
CDS_0038
CDS_0041
CDS_0043
CDS_0033
CDS_0036
CDS_0040
CDS_0042
uvig_235411 25218:32969(+)
CDS_0037
TemPhD_cluster_59768 25311:32802(+)
toxin (CDS_0030)
CDS_0036
CDS_0037
TemPhD_cluster_40559 24964:32847(+)
CDS_0035
CDS_0036
CDS_0038
TemPhD_cluster_29182 24964:32788(+)
CDS_0033
transposase (CDS_0038)
TemPhD_cluster_58768 26089:34705(+)
transposase (CDS_0038)
IMGVR_UViG_2563366958_000018|2563366958|2563399109|33032-85208 25742:36892(+)
CDS_0032
CDS_0035
MGV-GENOME-0301189 22474:31540(+)
CDS_0034
IMGVR_UViG_2698536573_000002|2698536573|2698570673 21575:30115(+)
CDS_0033
IMGVR_UViG_3300015184_000033|3300015184|Ga0183236_119762 22288:30105(+)
CDS_0035
IMGVR_UViG_2740892219_000006|2740892219|2740959034 22174:30727(+)
CDS_0035
TemPhD_cluster_16929 22935:31067(+)
CDS_0038
IMGVR_UViG_2740892231_000005|2740892231|2740960188 25253:33833(+)
CDS_0035
CDS_0038
TemPhD_cluster_35863_ 25951:34383(+)
CDS_0036
CDS_0037
TemPhD_cluster_31499 24872:33723(+)
CDS_0035
TemPhD_cluster_40287 24076:32258(+)
CDS_0027
CDS_0026
TemPhD_cluster_53392 9125:16155(-)
CDS_0006
TemPhD_cluster_66025 54:7090(+)
CDS_0007
TemPhD_cluster_65440 61:7492(+)
DNA methyltransferase (CDS_0030)
CDS_0005
TemPhD_cluster_65537 15489:6741(+)
transcriptional regulator (CDS_0026)
CDS_0025
TemPhD_cluster_65306 8875:17133(-)
CDS_0024
TemPhD_cluster_65655 1:645(-), 8560:15528(-)
CDS_0005
TemPhD_cluster_65734 14480:7136(+)
CDS_0005
CDS_0006
TemPhD_cluster_65515 15612:7191(+)
Rz-like spanin (CDS_0002)
CDS_0026
TemPhD_cluster_34057 1:1601(-), 9988:17423(-)
CDS_0002
CDS_0005
CDS_0006
TemPhD_cluster_65374 15222:7777(+)
CDS_0027
CDS_0035
IMGVR_UViG_2563366838_000005|2563366838|2563392876 22952:32073(+)
CDS_0032
CDS_0033
CDS_0036
CDS_0037
TemPhD_cluster_28962 24539:32513(+)
QnrE family quinolone resistance pentapeptide repeat protein (NF033375.2) (CDS_0037)
ABC transporter (CDS_0038)
39
40
41
42
43
DNA invertase (CDS_0045)
44
transposase (CDS_0046)
IMGVR_UViG_2563366956_000010|2563366956|2563398930 25742:46127(+)
CDS_0037
CDS_0033
CDS_0035
CDS_0039
CDS_0040
IMGVR_UViG_2585427870_000005|2585427870|2585474084 21774:30991(+)
CDS_0037
CDS_0038
CDS_0041
TemPhD_cluster_58628 26535:35433(+)
endolysin (CDS_0004)
CDS_0002
CDS_0029
CDS_0026
TemPhD_cluster_53250 1:2198(-), 10625:17277(-)
CDS_0028
TemPhD_cluster_65261 1:2098(-), 10747:17469(-)
CDS_0032
CDS_0003
TemPhD_cluster_65228 15525:6665(+)
CDS_0004
TemPhD_cluster_65364 14749:6546(+)
CDS_0003
TemPhD_cluster_53359 14217:6016(+)
CDS_0006
CDS_0005
TemPhD_cluster_65453 14874:6817(+)
8000 nt
DNA, RNA and nucleotide metabolism
integration and excision
lysis
moron, auxiliary metabolic gene and host takeover
other
transcription regulation
unknown function
AMR