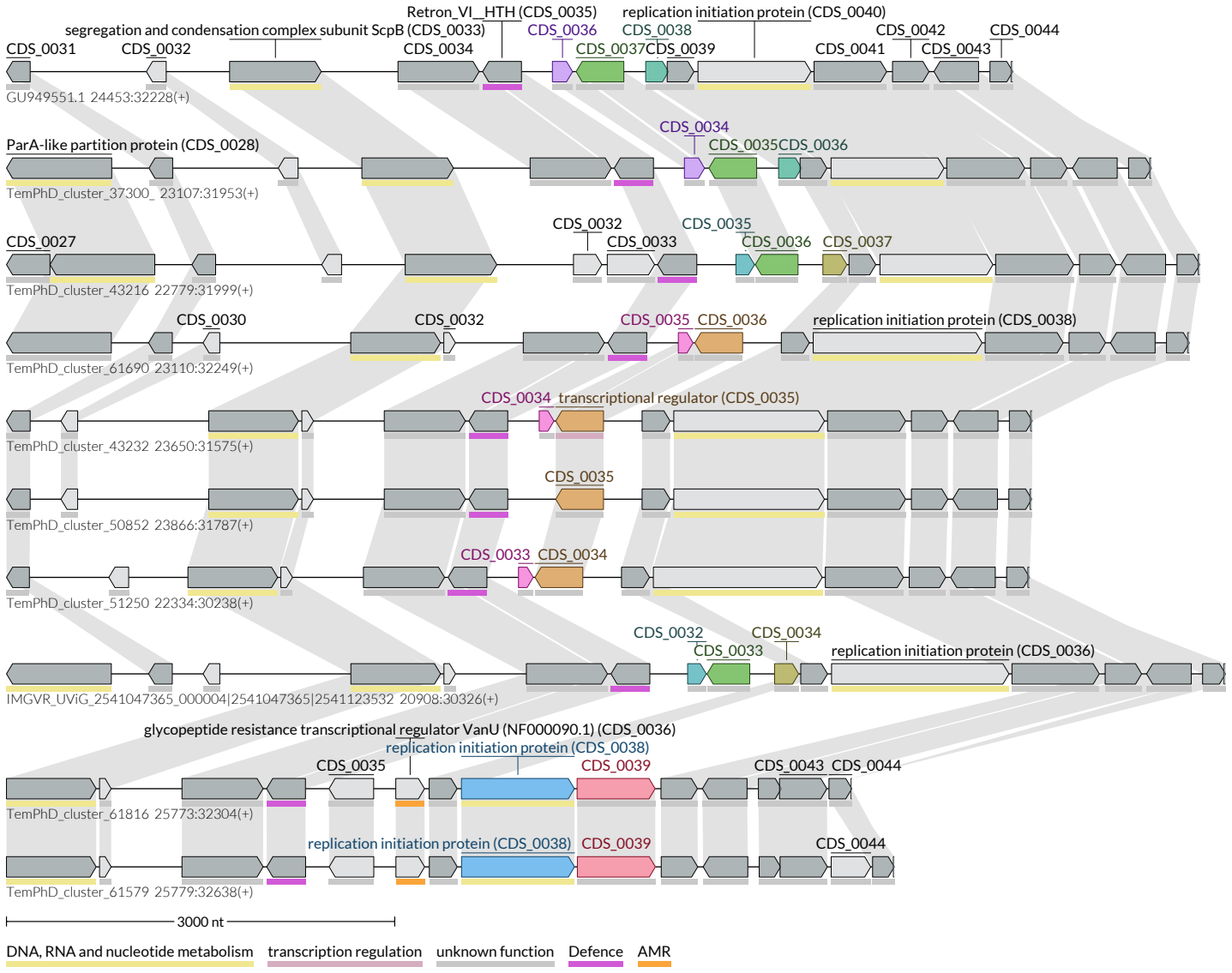
Retron_VI__HTH (CDS_0035)
replication initiation protein (CDS_0040)
segregation and condensation complex subunit ScpB (CDS_0033)
CDS_0036
CDS_0038
CDS_0042
CDS_0044
CDS_0031
CDS_0032
CDS_0034
CDS_0037
CDS_0039
CDS_0041
CDS_0043
GU949551.1 24453:32228(+)
CDS_0034
ParA-like partition protein (CDS_0028)
CDS_0035
CDS_0036
TemPhD_cluster_37300_ 23107:31953(+)
CDS_0032
CDS_0035
CDS_0027
CDS_0033
CDS_0036
CDS_0037
TemPhD_cluster_43216 22779:31999(+)
CDS_0030
CDS_0032
CDS_0035
CDS_0036
replication initiation protein (CDS_0038)
TemPhD_cluster_61690 23110:32249(+)
CDS_0034
transcriptional regulator (CDS_0035)
TemPhD_cluster_43232 23650:31575(+)
CDS_0035
TemPhD_cluster_50852 23866:31787(+)
CDS_0033
CDS_0034
TemPhD_cluster_51250 22334:30238(+)
CDS_0032
CDS_0034
CDS_0033
replication initiation protein (CDS_0036)
IMGVR_UViG_2541047365_000004|2541047365|2541123532 20908:30326(+)
glycopeptide resistance transcriptional regulator VanU (NF000090.1) (CDS_0036)
replication initiation protein (CDS_0038)
CDS_0035
CDS_0039
CDS_0043
CDS_0044
TemPhD_cluster_61816 25773:32304(+)
replication initiation protein (CDS_0038)
CDS_0039
CDS_0044
TemPhD_cluster_61579 25779:32638(+)
3000 nt
DNA, RNA and nucleotide metabolism
transcription regulation
unknown function
Defence
AMR