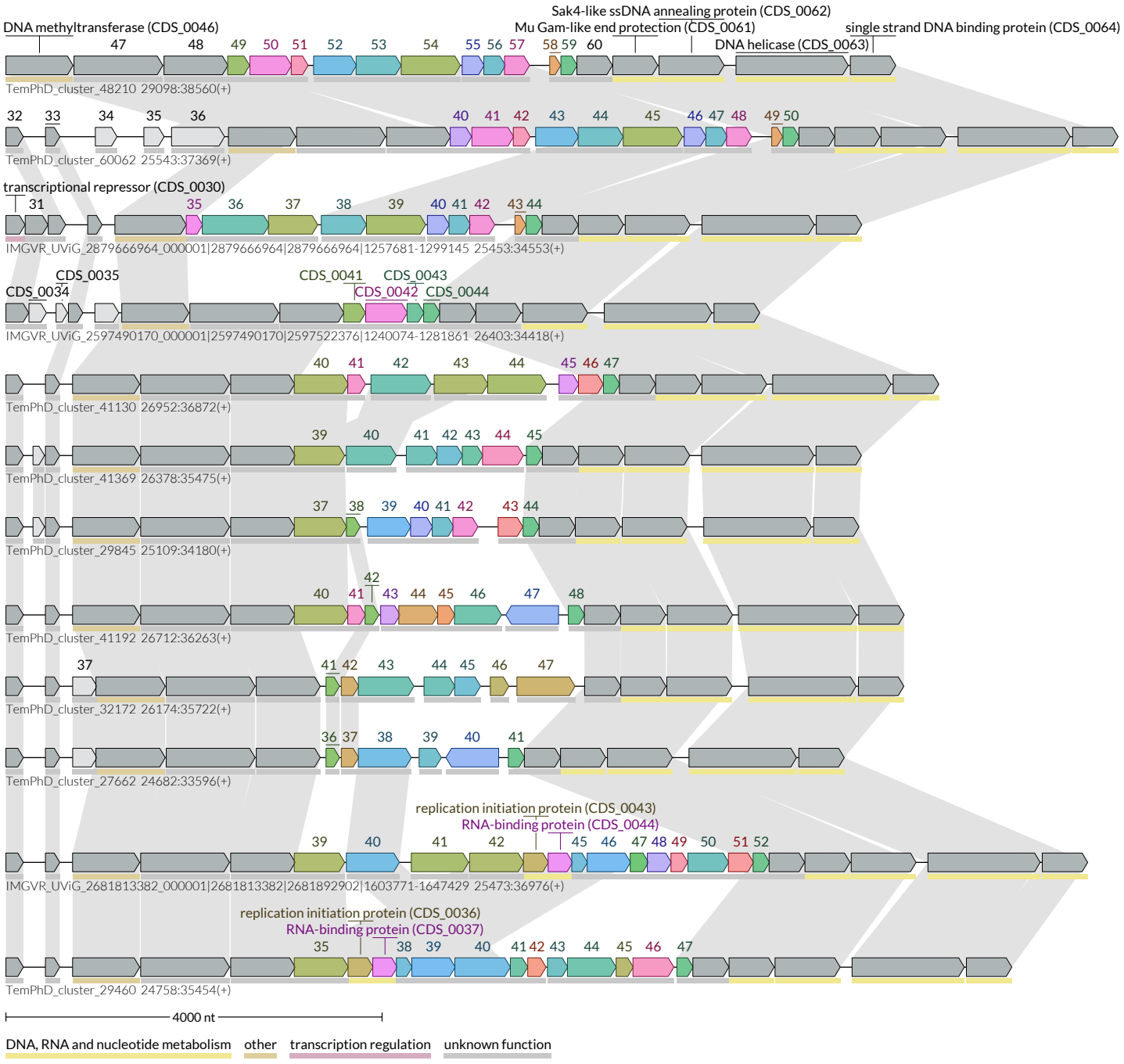
DNA methyltransferase (CDS_0046)
Sak4-like ssDNA annealing protein (CDS_0062)
Mu Gam-like end protection (CDS_0061)
single strand DNA binding protein (CDS_0064)
DNA helicase (CDS_0063)
TemPhD_cluster_48210 29098:38560(+)
TemPhD_cluster_60062 25543:37369(+)
transcriptional repressor (CDS_0030)
IMGVR_UViG_2879666964_000001|2879666964|2879666964|1257681-1299145 25453:34553(+)
CDS_0035
CDS_0034
CDS_0041
CDS_0043
CDS_0042
CDS_0044
IMGVR_UViG_2597490170_000001|2597490170|2597522376|1240074-1281861 26403:34418(+)
TemPhD_cluster_41130 26952:36872(+)
TemPhD_cluster_41369 26378:35475(+)
TemPhD_cluster_29845 25109:34180(+)
TemPhD_cluster_41192 26712:36263(+)
TemPhD_cluster_32172 26174:35722(+)
TemPhD_cluster_27662 24682:33596(+)
replication initiation protein (CDS_0043)
RNA-binding protein (CDS_0044)
IMGVR_UViG_2681813382_000001|2681813382|2681892902|1603771-1647429 25473:36976(+)
replication initiation protein (CDS_0036)
RNA-binding protein (CDS_0037)
TemPhD_cluster_29460 24758:35454(+)
4000 nt
DNA, RNA and nucleotide metabolism
other
transcription regulation
unknown function