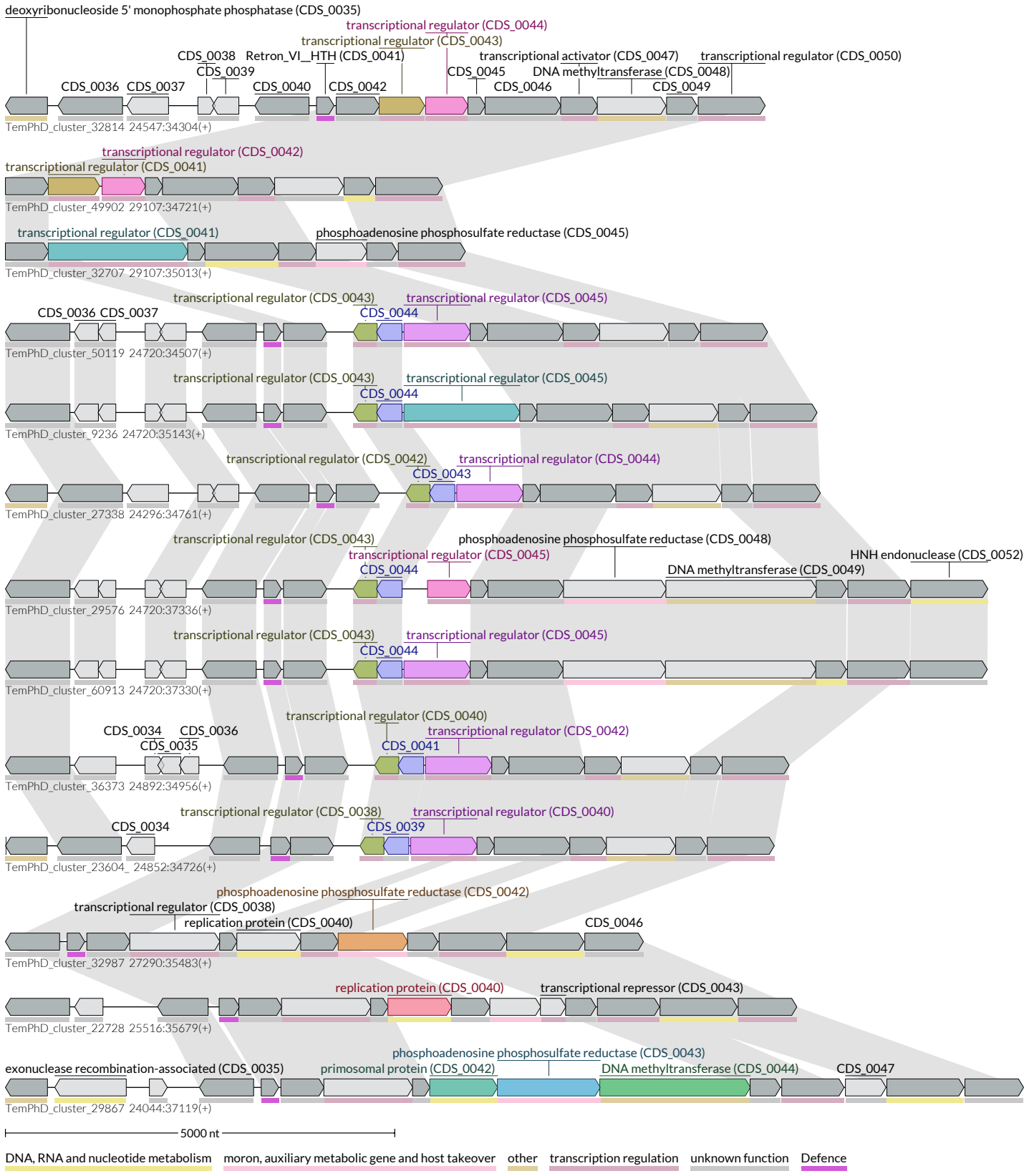
deoxyribonucleoside 5' monophosphate phosphatase (CDS_0035)
transcriptional regulator (CDS_0044)
transcriptional regulator (CDS_0043)
CDS_0038
Retron_VI__HTH (CDS_0041)
transcriptional activator (CDS_0047)
transcriptional regulator (CDS_0050)
CDS_0039
CDS_0045
DNA methyltransferase (CDS_0048)
CDS_0036
CDS_0037
CDS_0040
CDS_0042
CDS_0046
CDS_0049
TemPhD_cluster_32814 24547:34304(+)
transcriptional regulator (CDS_0042)
transcriptional regulator (CDS_0041)
TemPhD_cluster_49902 29107:34721(+)
transcriptional regulator (CDS_0041)
phosphoadenosine phosphosulfate reductase (CDS_0045)
TemPhD_cluster_32707 29107:35013(+)
transcriptional regulator (CDS_0043)
transcriptional regulator (CDS_0045)
CDS_0036 CDS_0037
CDS_0044
TemPhD_cluster_50119 24720:34507(+)
transcriptional regulator (CDS_0043)
transcriptional regulator (CDS_0045)
CDS_0044
TemPhD_cluster_9236 24720:35143(+)
transcriptional regulator (CDS_0042)
transcriptional regulator (CDS_0044)
CDS_0043
TemPhD_cluster_27338 24296:34761(+)
transcriptional regulator (CDS_0043)
phosphoadenosine phosphosulfate reductase (CDS_0048)
transcriptional regulator (CDS_0045)
HNH endonuclease (CDS_0052)
CDS_0044
DNA methyltransferase (CDS_0049)
TemPhD_cluster_29576 24720:37336(+)
transcriptional regulator (CDS_0043)
transcriptional regulator (CDS_0045)
CDS_0044
TemPhD_cluster_60913 24720:37330(+)
transcriptional regulator (CDS_0040)
CDS_0034
CDS_0036
transcriptional regulator (CDS_0042)
CDS_0035
CDS_0041
TemPhD_cluster_36373 24892:34956(+)
transcriptional regulator (CDS_0038)
transcriptional regulator (CDS_0040)
CDS_0034
CDS_0039
TemPhD_cluster_23604_ 24852:34726(+)
phosphoadenosine phosphosulfate reductase (CDS_0042)
transcriptional regulator (CDS_0038)
replication protein (CDS_0040)
CDS_0046
TemPhD_cluster_32987 27290:35483(+)
replication protein (CDS_0040)
transcriptional repressor (CDS_0043)
TemPhD_cluster_22728 25516:35679(+)
phosphoadenosine phosphosulfate reductase (CDS_0043)
exonuclease recombination-associated (CDS_0035)
primosomal protein (CDS_0042)
DNA methyltransferase (CDS_0044)
CDS_0047
TemPhD_cluster_29867 24044:37119(+)
5000 nt
DNA, RNA and nucleotide metabolism
moron, auxiliary metabolic gene and host takeover
other
transcription regulation
unknown function
Defence