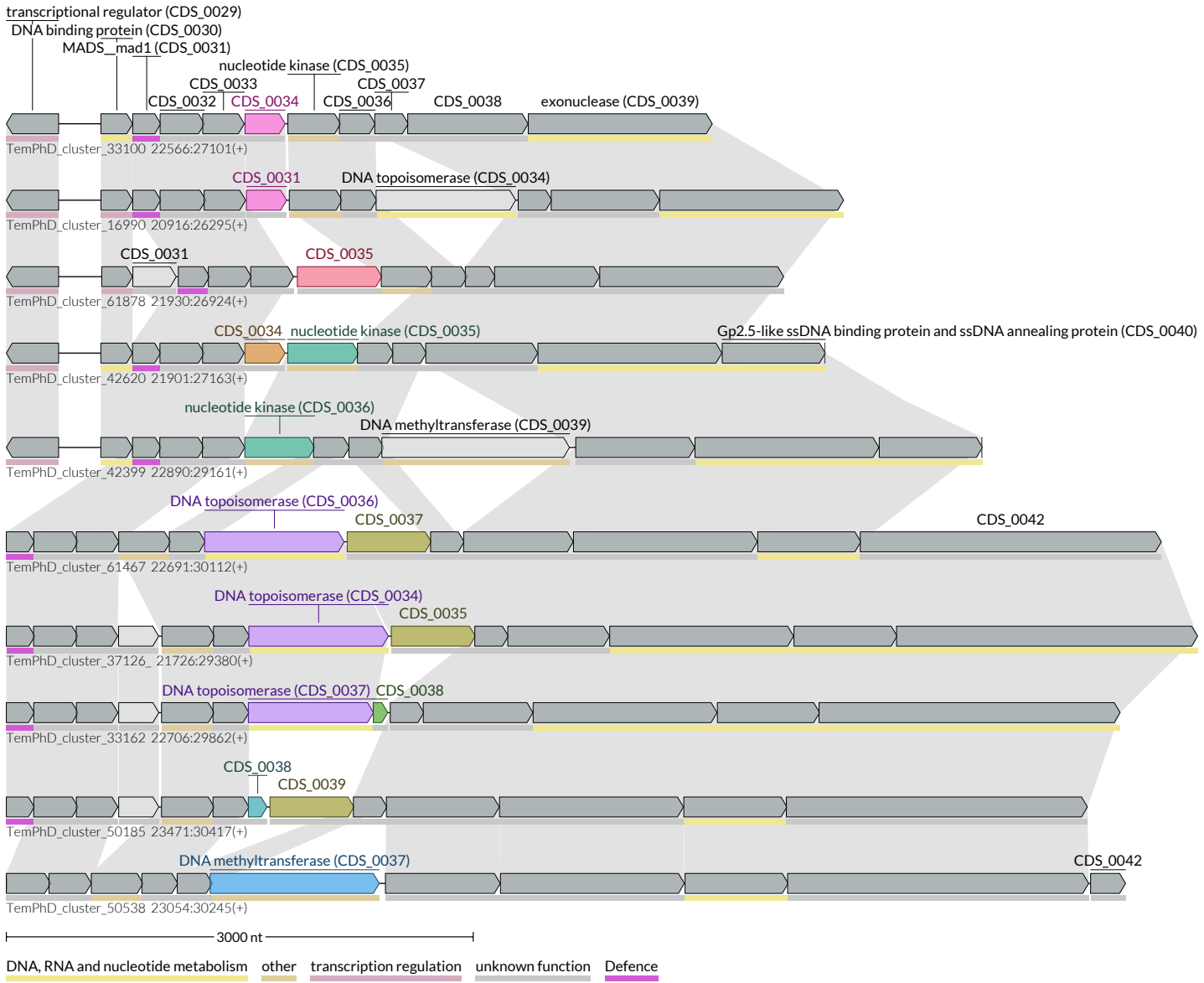
transcriptional regulator (CDS_0029)
DNA binding protein (CDS_0030)
MADS_mad1 (CDS_0031)
nucleotide kinase (CDS_0035)
CDS_0033
CDS_0037
CDS_0032
CDS_0034
CDS_0036
CDS_0038
exonuclease (CDS_0039)
TemPhD_cluster_33100 22566:27101(+)
CDS_0031
DNA topoisomerase (CDS_0034)
TemPhD_cluster_16990 20916:26295(+)
CDS_0031
CDS_0035
TemPhD_cluster_61878 21930:26924(+)
CDS_0034
nucleotide kinase (CDS_0035)
Gp2.5-like ssDNA binding protein and ssDNA annealing protein (CDS_0040)
TemPhD_cluster_42620 21901:27163(+)
nucleotide kinase (CDS_0036)
DNA methyltransferase (CDS_0039)
TemPhD_cluster_42399 22890:29161(+)
DNA topoisomerase (CDS_0036)
CDS_0037
CDS_0042
TemPhD_cluster_61467 22691:30112(+)
DNA topoisomerase (CDS_0034)
CDS_0035
TemPhD_cluster_37126_ 21726:29380(+)
DNA topoisomerase (CDS_0037)
CDS_0038
TemPhD_cluster_33162 22706:29862(+)
CDS_0038
CDS_0039
TemPhD_cluster_50185 23471:30417(+)
DNA methyltransferase (CDS_0037)
CDS_0042
TemPhD_cluster_50538 23054:30245(+)
3000 nt
DNA, RNA and nucleotide metabolism
other
transcription regulation
unknown function
Defence