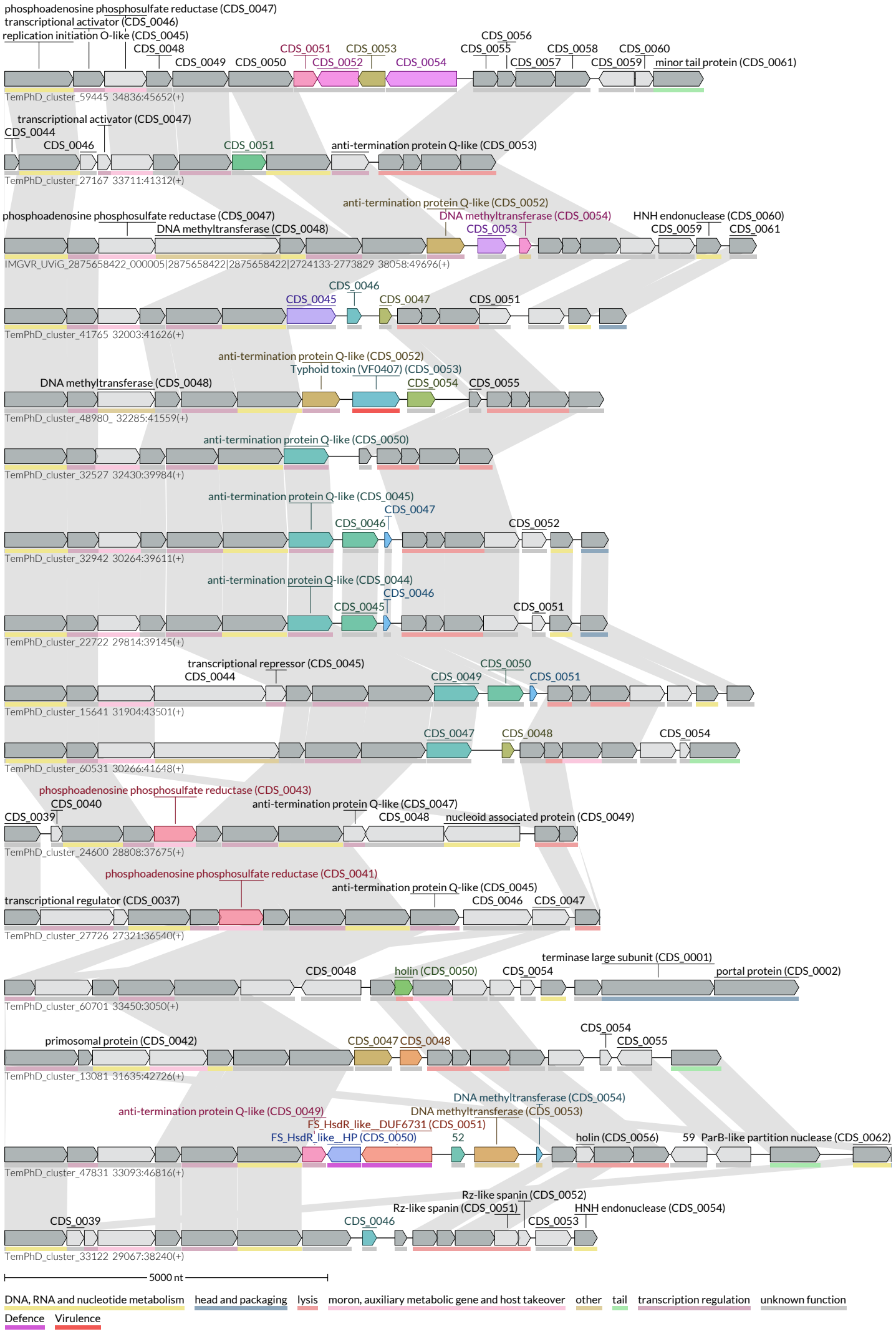
phosphoadenosine phosphosulfate reductase (CDS_0047)
transcriptional activator (CDS_0046)
replication initiation O-like (CDS_0045)
CDS_0048
CDS_0049
CDS_0050
CDS_0051
CDS_0052
CDS_0053
CDS_0054
CDS_0056
CDS_0055
CDS_0057
CDS_0058
CDS_0059
CDS_0060
minor tail protein (CDS_0061)
TemPhD_cluster_59445 34836:45652(+)
transcriptional activator (CDS_0047)
CDS_0044
CDS_0046
CDS_0051
anti-termination protein Q-like (CDS_0053)
TemPhD_cluster_27167 33711:41312(+)
anti-termination protein Q-like (CDS_0052)
phosphoadenosine phosphosulfate reductase (CDS_0047)
DNA methyltransferase (CDS_0048)
DNA methyltransferase (CDS_0054)
CDS_0053
HNH endonuclease (CDS_0060)
CDS_0059
CDS_0061
IMGVR_UViG_2875658422_000005|2875658422|2875658422|2724133-2773829 38058:49696(+)
CDS_0046
CDS_0045
CDS_0047
CDS_0051
TemPhD_cluster_41765 32003:41626(+)
anti-termination protein Q-like (CDS_0052)
Typhoid toxin (VF0407) (CDS_0053)
DNA methyltransferase (CDS_0048)
CDS_0054
CDS_0055
TemPhD_cluster_48980_ 32285:41559(+)
anti-termination protein Q-like (CDS_0050)
TemPhD_cluster_32527 32430:39984(+)
anti-termination protein Q-like (CDS_0045)
CDS_0047
CDS_0046
CDS_0052
TemPhD_cluster_32942 30264:39611(+)
anti-termination protein Q-like (CDS_0044)
CDS_0046
CDS_0045
CDS_0051
TemPhD_cluster_22722 29814:39145(+)
transcriptional repressor (CDS_0045)
CDS_0044
CDS_0050
CDS_0049
CDS_0051
TemPhD_cluster_15641 31904:43501(+)
CDS_0047
CDS_0048
CDS_0054
TemPhD_cluster_60531 30266:41648(+)
phosphoadenosine phosphosulfate reductase (CDS_0043)
CDS_0040
CDS_0039
anti-termination protein Q-like (CDS_0047)
CDS_0048
nucleoid associated protein (CDS_0049)
TemPhD_cluster_24600 28808:37675(+)
phosphoadenosine phosphosulfate reductase (CDS_0041)
anti-termination protein Q-like (CDS_0045)
transcriptional regulator (CDS_0037)
CDS_0046
CDS_0047
TemPhD_cluster_27726 27321:36540(+)
terminase large subunit (CDS_0001)
CDS_0048
holin (CDS_0050)
CDS_0054
portal protein (CDS_0002)
TemPhD_cluster_60701 33450:3050(+)
CDS_0054
primosomal protein (CDS_0042)
CDS_0047
CDS_0048
CDS_0055
TemPhD_cluster_13081 31635:42726(+)
DNA methyltransferase (CDS_0054)
anti-termination protein Q-like (CDS_0049)
DNA methyltransferase (CDS_0053)
FS_HsdR_like__DUF6731 (CDS_0051)
FS_HsdR_like__HP (CDS_0050)
52
holin (CDS_0056)
59
ParB-like partition nuclease (CDS_0062)
TemPhD_cluster_47831 33093:46816(+)
Rz-like spanin (CDS_0052)
Rz-like spanin (CDS_0051)
HNH endonuclease (CDS_0054)
CDS_0039
CDS_0046
CDS_0053
TemPhD_cluster_33122 29067:38240(+)
5000 nt
DNA, RNA and nucleotide metabolism
head and packaging
lysis
moron, auxiliary metabolic gene and host takeover
other
tail
transcription regulation
unknown function
Defence
Virulence