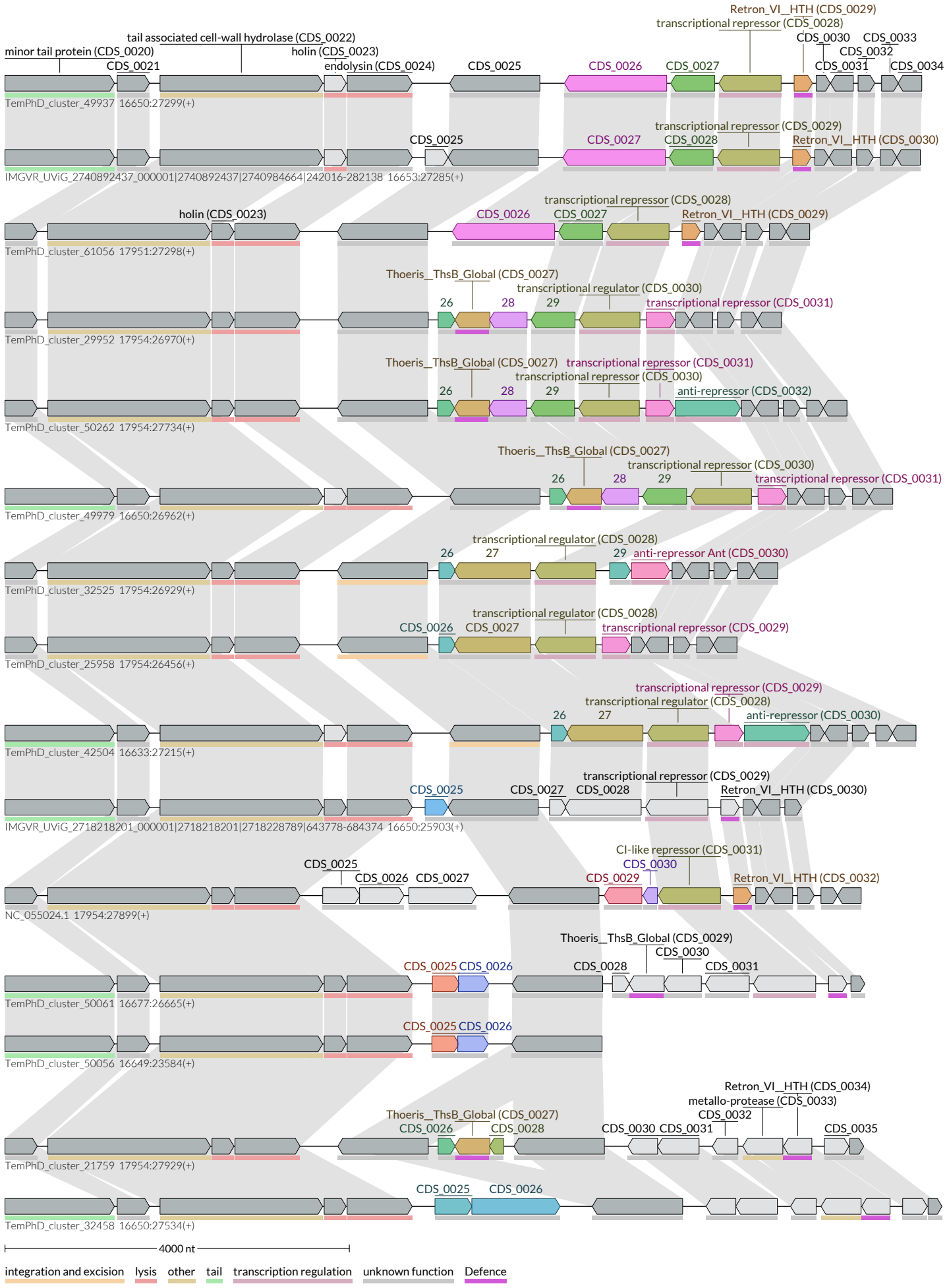
Retron_VI__HTH (CDS_0029)
transcriptional repressor (CDS_0028)
tail associated cell-wall hydrolase (CDS_0022)
minor tail protein (CDS_0020)
holin (CDS_0023)
CDS_0030
CDS_0033
CDS_0032
CDS_0021
endolysin (CDS_0024)
CDS_0025
CDS_0026
CDS_0027
CDS_0031
CDS_0034
TemPhD_cluster_49937 16650:27299(+)
transcriptional repressor (CDS_0029)
CDS_0025
CDS_0027
CDS_0028
Retron_VI__HTH (CDS_0030)
IMGVR_UViG_2740892437_000001|2740892437|2740984664|242016-282138 16653:27285(+)
transcriptional repressor (CDS_0028)
holin (CDS_0023)
CDS_0026
CDS_0027
Retron_VI__HTH (CDS_0029)
TemPhD_cluster_61056 17951:27298(+)
Thoeris__ThsB_Global (CDS_0027)
transcriptional regulator (CDS_0030)
26
28
29
transcriptional repressor (CDS_0031)
TemPhD_cluster_29952 17954:26970(+)
Thoeris__ThsB_Global (CDS_0027)
transcriptional repressor (CDS_0031)
transcriptional repressor (CDS_0030)
26
28
29
anti-repressor (CDS_0032)
TemPhD_cluster_50262 17954:27734(+)
Thoeris__ThsB_Global (CDS_0027)
transcriptional repressor (CDS_0030)
26
28
29
transcriptional repressor (CDS_0031)
TemPhD_cluster_49979 16650:26962(+)
transcriptional regulator (CDS_0028)
26
27
29
anti-repressor Ant (CDS_0030)
TemPhD_cluster_32525 17954:26929(+)
transcriptional regulator (CDS_0028)
CDS_0026
CDS_0027
transcriptional repressor (CDS_0029)
TemPhD_cluster_25958 17954:26456(+)
transcriptional repressor (CDS_0029)
transcriptional regulator (CDS_0028)
26
27
anti-repressor (CDS_0030)
TemPhD_cluster_42504 16633:27215(+)
transcriptional repressor (CDS_0029)
CDS_0025
CDS_0027
CDS_0028
Retron_VI__HTH (CDS_0030)
IMGVR_UViG_2718218201_000001|2718218201|2718228789|643778-684374 16650:25903(+)
CI-like repressor (CDS_0031)
CDS_0025
CDS_0030
CDS_0026
CDS_0027
CDS_0029
Retron_VI__HTH (CDS_0032)
NC_055024.1 17954:27899(+)
Thoeris__ThsB_Global (CDS_0029)
CDS_0030
CDS_0025
CDS_0026
CDS_0028
CDS_0031
TemPhD_cluster_50061 16677:26665(+)
CDS_0025
CDS_0026
TemPhD_cluster_50056 16649:23584(+)
Retron_VI__HTH (CDS_0034)
metallo-protease (CDS_0033)
Thoeris__ThsB_Global (CDS_0027)
CDS_0032
CDS_0026
CDS_0028
CDS_0030
CDS_0031
CDS_0035
TemPhD_cluster_21759 17954:27929(+)
CDS_0025
CDS_0026
TemPhD_cluster_32458 16650:27534(+)
4000 nt
integration and excision
lysis
other
tail
transcription regulation
unknown function
Defence