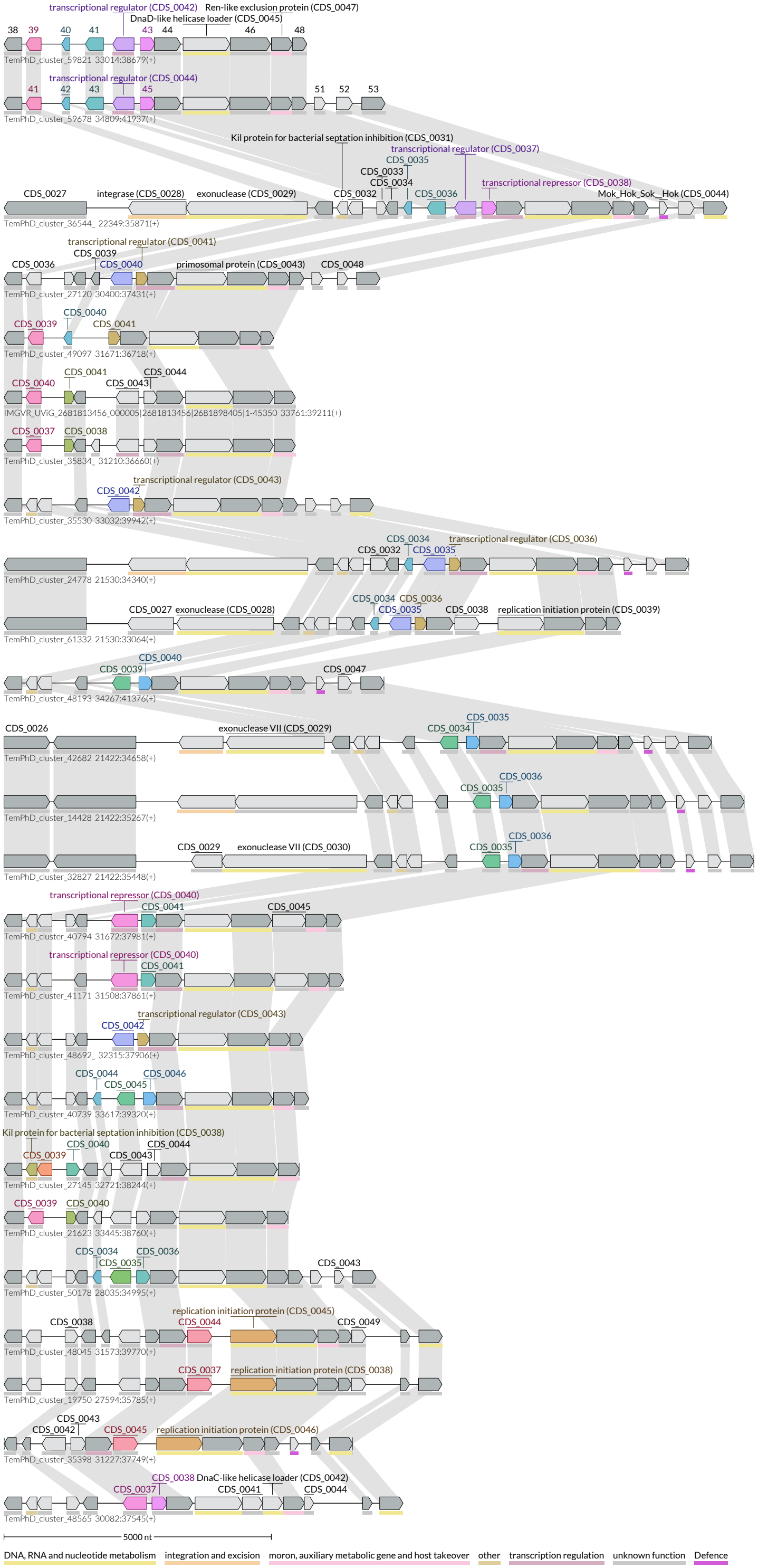
transcriptional regulator (CDS_0042)
Ren-like exclusion protein (CDS_0047)
DnaD-like helicase loader (CDS_0045)
38 39 40 41 43 44 46 48
TemPhD_cluster_59821 33014:38679(+)
transcriptional regulator (CDS_0044)
41 42 43 45 51 52 53
TemPhD_cluster_59678 34809:41937(+)
Kil protein for bacterial septation inhibition (CDS_0031)
transcriptional regulator (CDS_0037)
CDS_0035
CDS_0033
CDS_0034
transcriptional repressor (CDS_0038)
CDS_0027
integrase (CDS_0028)
exonuclease (CDS_0029)
CDS_0032
CDS_0036
Mok_Hok_Sok__Hok (CDS_0044)
TemPhD_cluster_36544_ 22349:35871(+)
transcriptional regulator (CDS_0041)
CDS_0039
CDS_0036
CDS_0040
primosomal protein (CDS_0043)
CDS_0048
TemPhD_cluster_27120 30400:37431(+)
CDS_0040
CDS_0039
CDS_0041
TemPhD_cluster_49097 31671:36718(+)
CDS_0041
CDS_0044
CDS_0040
CDS_0043
IMGVR_UViG_2681813456_000005|2681813456|2681898405|1-45350 33761:39211(+)
CDS_0037
CDS_0038
TemPhD_cluster_35834_ 31210:36660(+)
transcriptional regulator (CDS_0043)
CDS_0042
TemPhD_cluster_35530 33032:39942(+)
CDS_0034
transcriptional regulator (CDS_0036)
CDS_0032
CDS_0035
TemPhD_cluster_24778 21530:34340(+)
CDS_0034
CDS_0036
CDS_0027
exonuclease (CDS_0028)
CDS_0035
CDS_0038
replication initiation protein (CDS_0039)
TemPhD_cluster_61332 21530:33064(+)
CDS_0040
CDS_0039
CDS_0047
TemPhD_cluster_48193 34267:41376(+)
CDS_0035
CDS_0026
exonuclease VII (CDS_0029)
CDS_0034
TemPhD_cluster_42682 21422:34658(+)
CDS_0036
CDS_0035
TemPhD_cluster_14428 21422:35267(+)
CDS_0036
CDS_0029
exonuclease VII (CDS_0030)
CDS_0035
TemPhD_cluster_32827 21422:35448(+)
transcriptional repressor (CDS_0040)
CDS_0041
CDS_0045
TemPhD_cluster_40794 31672:37981(+)
transcriptional repressor (CDS_0040)
CDS_0041
TemPhD_cluster_41171 31508:37861(+)
transcriptional regulator (CDS_0043)
CDS_0042
TemPhD_cluster_48692_ 32315:37906(+)
CDS_0044
CDS_0046
CDS_0045
TemPhD_cluster_40739 33617:39320(+)
Kil protein for bacterial septation inhibition (CDS_0038)
CDS_0040
CDS_0044
CDS_0039
CDS_0043
TemPhD_cluster_27145 32721:38244(+)
CDS_0039
CDS_0040
TemPhD_cluster_21623 33445:38760(+)
CDS_0034
CDS_0036
CDS_0035
CDS_0043
TemPhD_cluster_50178 28035:34995(+)
replication initiation protein (CDS_0045)
CDS_0038
CDS_0044
CDS_0049
TemPhD_cluster_48045 31573:39770(+)
CDS_0037
replication initiation protein (CDS_0038)
TemPhD_cluster_19750 27594:35785(+)
CDS_0043
CDS_0042
CDS_0045
replication initiation protein (CDS_0046)
TemPhD_cluster_35398 31227:37749(+)
CDS_0038
DnaC-like helicase loader (CDS_0042)
CDS_0037
CDS_0041
CDS_0044
TemPhD_cluster_48565 30082:37545(+)
5000 nt
DNA, RNA and nucleotide metabolism
integration and excision
moron, auxiliary metabolic gene and host takeover
other
transcription regulation
unknown function
Defence