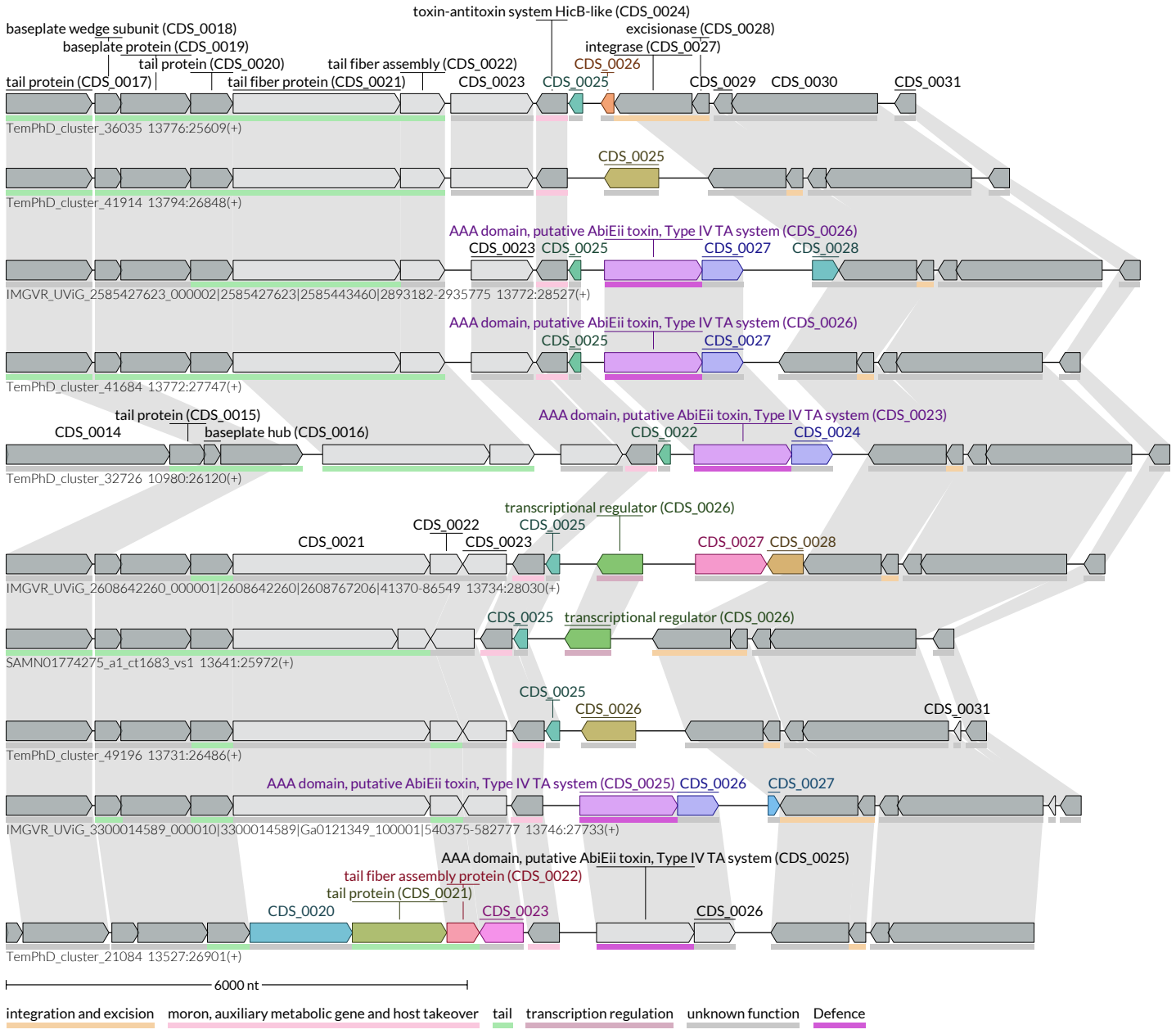
toxin-antitoxin system HicB-like (CDS_0024)
baseplate wedge subunit (CDS_0018)
excisionase (CDS_0028)
baseplate protein (CDS_0019)
integrase (CDS_0027)
tail protein (CDS_0020)
tail fiber assembly (CDS_0022)
CDS_0026
tail protein (CDS_0017)
tail fiber protein (CDS_0021)
CDS_0023
CDS_0025
CDS_0029
CDS_0030
CDS_0031
TemPhD_cluster_36035 13776:25609(+)
CDS_0025
TemPhD_cluster_41914 13794:26848(+)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0026)
CDS_0023
CDS_0025
CDS_0027
CDS_0028
IMGVR_UViG_2585427623_000002|2585427623|2585443460|2893182-2935775 13772:28527(+)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0026)
CDS_0025
CDS_0027
TemPhD_cluster_41684 13772:27747(+)
tail protein (CDS_0015)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0023)
CDS_0014
baseplate hub (CDS_0016)
CDS_0022
CDS_0024
TemPhD_cluster_32726 10980:26120(+)
transcriptional regulator (CDS_0026)
CDS_0022
CDS_0025
CDS_0021
CDS_0023
CDS_0027
CDS_0028
IMGVR_UViG_2608642260_000001|2608642260|2608767206|41370-86549 13734:28030(+)
CDS_0025
transcriptional regulator (CDS_0026)
SAMN01774275_a1_ct1683_vs1 13641:25972(+)
CDS_0025
CDS_0026
CDS_0031
TemPhD_cluster_49196 13731:26486(+)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0025)
CDS_0026
CDS_0027
IMGVR_UViG_3300014589_000010|3300014589|Ga0121349_100001|540375-582777 13746:27733(+)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0025)
tail fiber assembly protein (CDS_0022)
tail protein (CDS_0021)
CDS_0020
CDS_0023
CDS_0026
TemPhD_cluster_21084 13527:26901(+)
6000 nt
integration and excision
moron, auxiliary metabolic gene and host takeover
tail
transcription regulation
unknown function
Defence