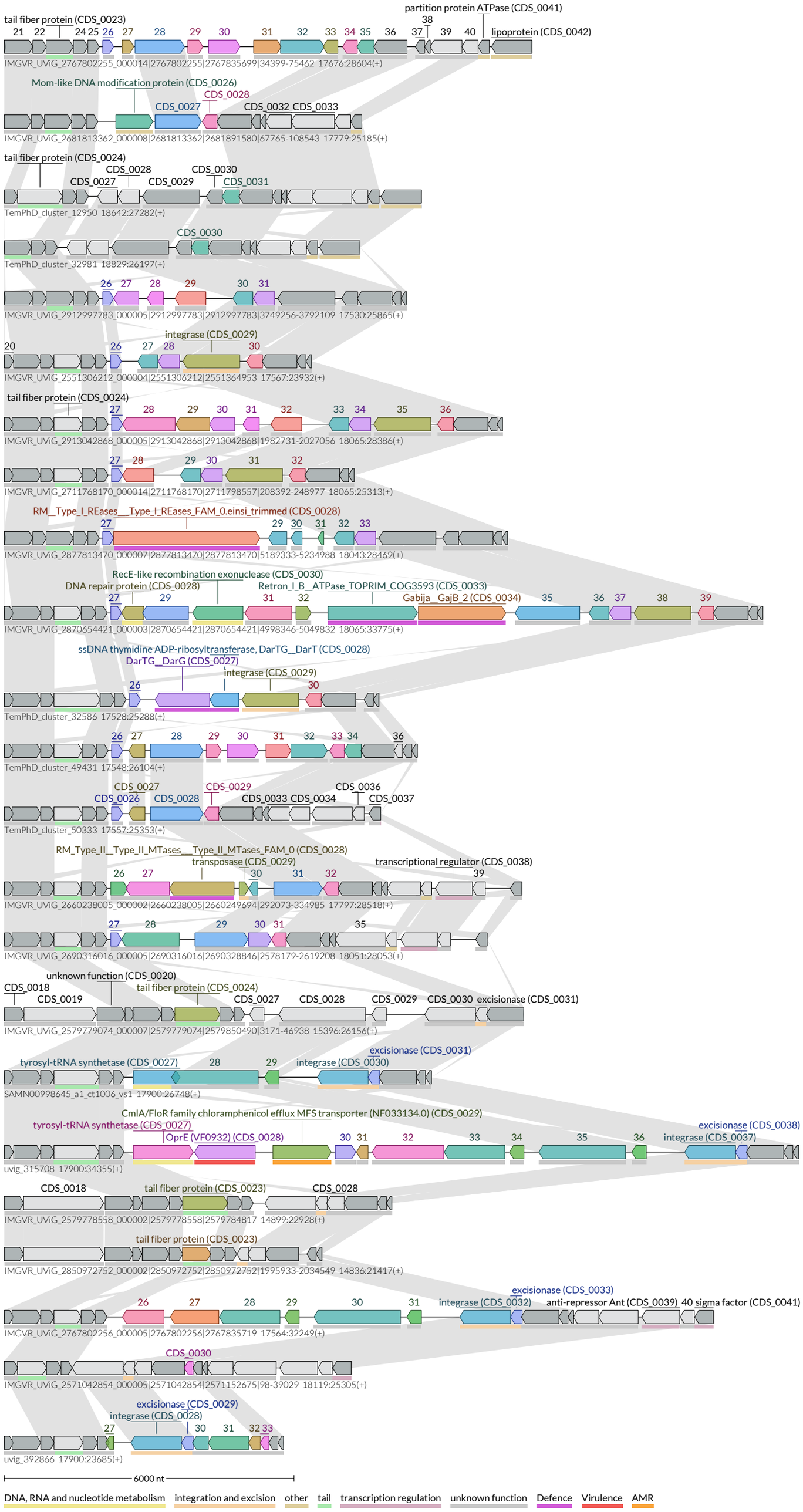
tail fiber protein (CDS_0023)
partition protein ATPase (CDS_0041)
lipoprotein (CDS_0042)
IMGVR_UViG_2767802255_000014|2767802255|2767835699|34399-75462 17676:28604(+)
Mom-like DNA modification protein (CDS_0026)
CDS_0027
CDS_0028
CDS_0032 CDS_0033
IMGVR_UViG_2681813362_000008|2681813362|2681891580|67765-108543 17779:25185(+)
tail fiber protein (CDS_0024)
CDS_0028
CDS_0030
CDS_0027
CDS_0029
CDS_0031
TemPhD_cluster_12950 18642:27282(+)
CDS_0030
TemPhD_cluster_32981 18829:26197(+)
IMGVR_UViG_2912997783_000005|2912997783|2912997783|3749256-3792109 17530:25865(+)
integrase (CDS_0029)
IMGVR_UViG_2551306212_000004|2551306212|2551364953 17567:23932(+)
tail fiber protein (CDS_0024)
IMGVR_UViG_2913042868_000005|2913042868|2913042868|1982731-2027056 18065:28386(+)
IMGVR_UViG_2711768170_000014|2711768170|2711798557|208392-248977 18065:25313(+)
RM__Type_I_REases___Type_I_REases_FAM_0.einsi_trimmed (CDS_0028)
IMGVR_UViG_2877813470_000007|2877813470|2877813470|5189333-5234988 18043:28469(+)
RecE-like recombination exonuclease (CDS_0030)
DNA repair protein (CDS_0028)
Retron_I_B__ATPase_TOPRIM_COG3593 (CDS_0033)
Gabija__GajB_2 (CDS_0034)
IMGVR_UViG_2870654421_000003|2870654421|2870654421|4998346-5049832 18065:33775(+)
ssDNA thymidine ADP-ribosyltransferase, DarTG__DarT (CDS_0028)
DarTG__DarG (CDS_0027)
integrase (CDS_0029)
TemPhD_cluster_32586 17528:25288(+)
TemPhD_cluster_49431 17548:26104(+)
CDS_0027
CDS_0029
CDS_0036
CDS_0026
CDS_0028
CDS_0033 CDS_0034
CDS_0037
TemPhD_cluster_50333 17557:25353(+)
RM_Type_II__Type_II_MTases___Type_II_MTases_FAM_0 (CDS_0028)
transposase (CDS_0029)
transcriptional regulator (CDS_0038)
IMGVR_UViG_2660238005_000002|2660238005|2660249694|292073-334985 17797:28518(+)
IMGVR_UViG_2690316016_000005|2690316016|2690328846|2578179-2619208 18051:28053(+)
unknown function (CDS_0020)
CDS_0018
CDS_0019
tail fiber protein (CDS_0024)
CDS_0027
CDS_0028
CDS_0029
CDS_0030
excisionase (CDS_0031)
IMGVR_UViG_2579779074_000007|2579779074|2579850490|3171-46938 15396:26156(+)
excisionase (CDS_0031)
tyrosyl-tRNA synthetase (CDS_0027)
integrase (CDS_0030)
SAMN00998645_a1_ct1006_vs1 17900:26748(+)
CmlA/FloR family chloramphenicol efflux MFS transporter (NF033134.0) (CDS_0029)
tyrosyl-tRNA synthetase (CDS_0027)
OprE (VF0932) (CDS_0028)
excisionase (CDS_0038)
integrase (CDS_0037)
uvig_315708 17900:34355(+)
CDS_0018
tail fiber protein (CDS_0023)
CDS_0028
IMGVR_UViG_2579778558_000002|2579778558|2579784817 14899:22928(+)
tail fiber protein (CDS_0023)
IMGVR_UViG_2850972752_000002|2850972752|2850972752|1995933-2034549 14836:21417(+)
excisionase (CDS_0033)
integrase (CDS_0032)
anti-repressor Ant (CDS_0039)
sigma factor (CDS_0041)
IMGVR_UViG_2767802256_000005|2767802256|2767835719 17564:32249(+)
CDS_0030
IMGVR_UViG_2571042854_000005|2571042854|2571152675|98-39029 18119:25305(+)
excisionase (CDS_0029)
integrase (CDS_0028)
uvig_392866 17900:23685(+)
6000 nt
DNA, RNA and nucleotide metabolism
integration and excision
other
tail
transcription regulation
unknown function
Defence
Virulence
AMR