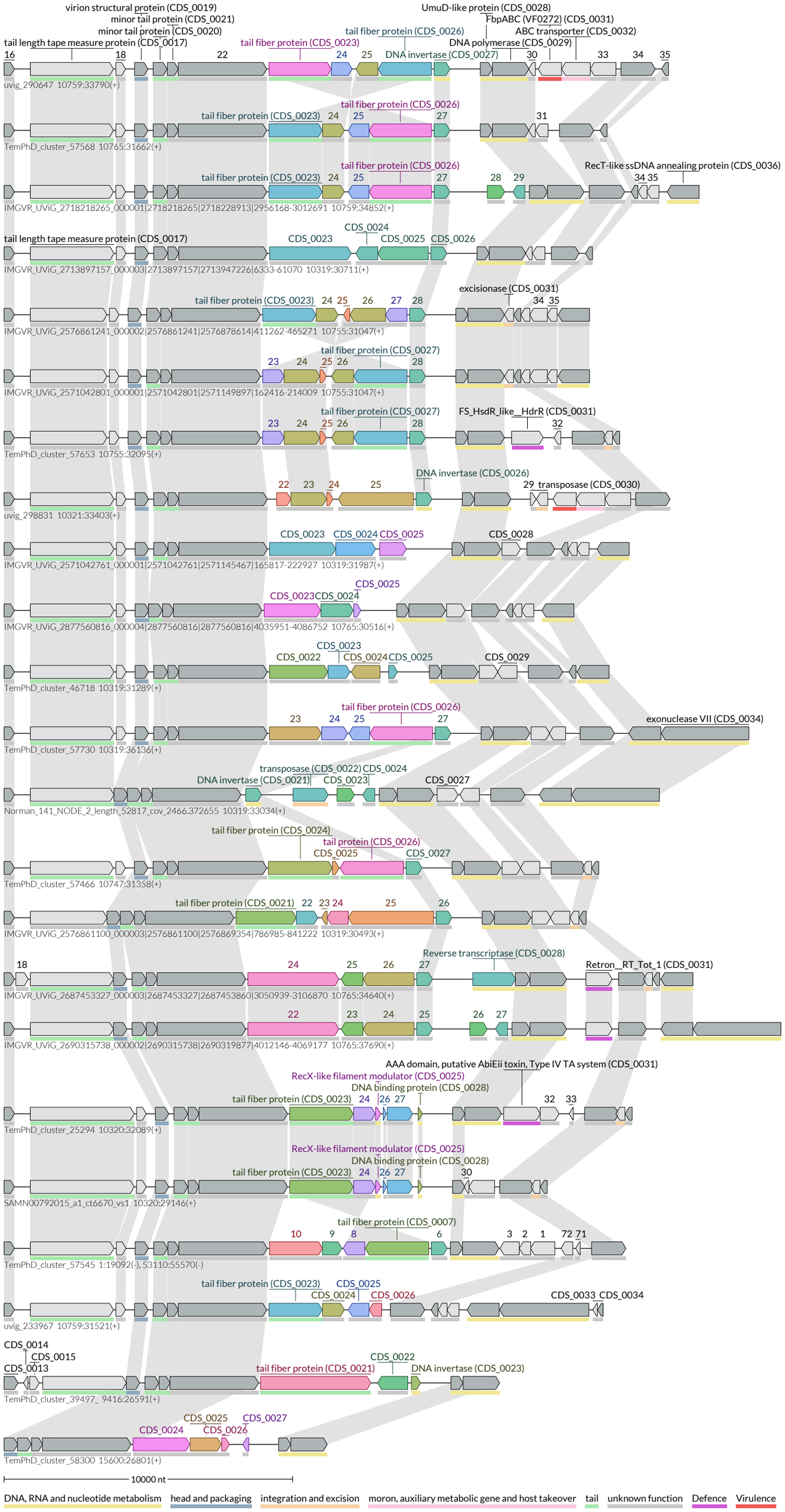
virion structural protein (CDS_0019)
minor tail protein (CDS_0021)
minor tail protein (CDS_0020)
tail length tape measure protein (CDS_0017)
tail fiber protein (CDS_0023)
tail fiber protein (CDS_0026)
DNA invertase (CDS_0027)
UmuD-like protein (CDS_0028)
FbpABC (VF0272) (CDS_0031)
ABC transporter (CDS_0032)
DNA polymerase (CDS_0029)
uvig_290647 10759:33790(+)
TemPhD_cluster_57568 10765:31662(+)
RecT-like ssDNA annealing protein (CDS_0036)
IMGVR_UViG_2718218265_000001|2718218265|2718228913|2956168-3012691 10759:34852(+)
CDS_0024
CDS_0023
CDS_0025
CDS_0026
IMGVR_UViG_2713897157_000003|2713897157|2713947226|6333-61070 10319:30711(+)
excisionase (CDS_0031)
IMGVR_UViG_2576861241_000002|2576861241|2576878614|411262-465271 10755:31047(+)
tail fiber protein (CDS_0027)
IMGVR_UViG_2571042801_000001|2571042801|2571149897|162416-214009 10755:31047(+)
FS_HsdR_like__HdrR (CDS_0031)
TemPhD_cluster_57653 10755:32095(+)
DNA invertase (CDS_0026)
29_transposase (CDS_0030)
uvig_298831 10321:33403(+)
CDS_0028
IMGVR_UViG_2571042761_000001|2571042761|2571145467|165817-222927 10319:31987(+)
IMGVR_UViG_2877560816_000004|2877560816|2877560816|4035951-4086752 10765:30516(+)
CDS_0022
CDS_0029
TemPhD_cluster_46718 10319:31289(+)
exonuclease VII (CDS_0034)
TemPhD_cluster_57730 10319:36136(+)
transposase (CDS_0022)
DNA invertase (CDS_0021)
Norman_141_NODE_2_length_52817_cov_2466.372655 10319:33034(+)
tail fiber protein (CDS_0024)
tail protein (CDS_0026)
TemPhD_cluster_57466 10747:31358(+)
tail fiber protein (CDS_0021)
IMGVR_UViG_2576861100_000003|2576861100|2576869354|786985-841222 10319:30493(+)
Reverse transcriptase (CDS_0028)
Retron__RT_Tot_1 (CDS_0031)
IMGVR_UViG_2687453327_000003|2687453327|2687453860|3050939-3106870 10765:34640(+)
IMGVR_UViG_2690315738_000002|2690315738|2690319877|4012146-4069177 10765:37690(+)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0031)
RecX-like filament modulator (CDS_0025)
DNA binding protein (CDS_0028)
TemPhD_cluster_25294 10320:32089(+)
SAMN00792015_a1_ct6670_vs1 10320:29146(+)
tail fiber protein (CDS_0007)
TemPhD_cluster_57545 1:19092(-), 53110:55570(-)
uvig_233967 10759:31521(+)
CDS_0033
CDS_0034
CDS_0014
CDS_0015
CDS_0013
CDS_0022
DNA invertase (CDS_0023)
TemPhD_cluster_39497_ 9416:26591(+)
CDS_0027
TemPhD_cluster_58300 15600:26801(+)
10000 nt
DNA, RNA and nucleotide metabolism
head and packaging
integration and excision
moron, auxiliary metabolic gene and host takeover
tail
unknown function
Defence
Virulence