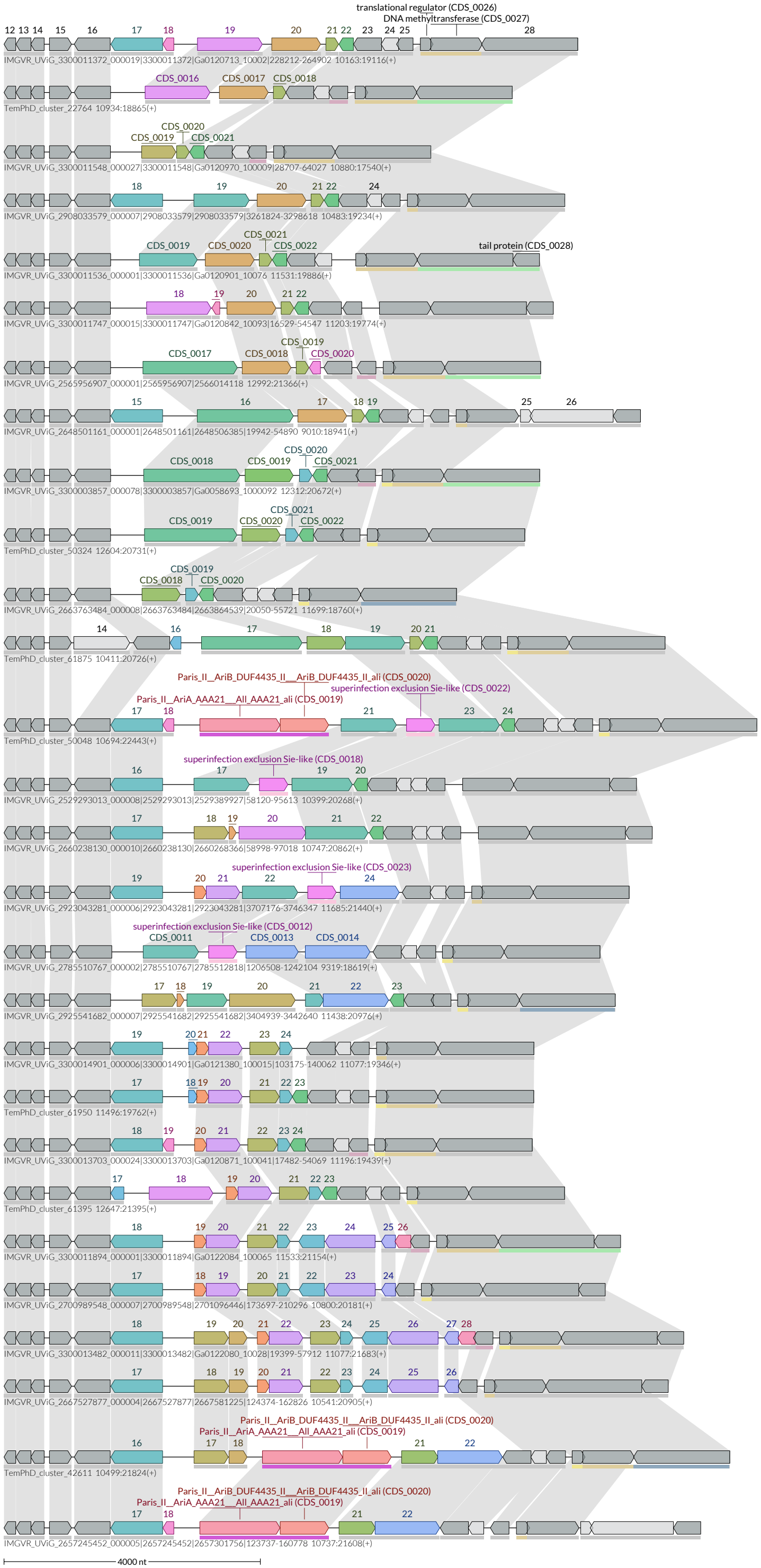
translational regulator (CDS_0026)
DNA methyltransferase (CDS_0027)
IMGVR_UViG_3300011372_000019|3300011372|Ga0120713_10002|228212-264902 10163:19116(+)
TemPhD_cluster_22764 10934:18865(+)
IMGVR_UViG_3300011548_000027|3300011548|Ga0120970_100009|28707-64027 10880:17540(+)
IMGVR_UViG_2908033579_000007|2908033579|2908033579|3261824-3298618 10483:19234(+)
tail protein (CDS_0028)
IMGVR_UViG_3300011536_000001|3300011536|Ga0120901_10076 11531:19886(+)
IMGVR_UViG_3300011747_000015|3300011747|Ga0120842_10093|16529-54547 11203:19774(+)
IMGVR_UViG_2565956907_000001|2565956907|2566014118 12992:21366(+)
IMGVR_UViG_2648501161_000001|2648501161|2648506385|19942-54890 9010:18941(+)
IMGVR_UViG_3300003857_000078|3300003857|Ga0058693_1000092 12312:20672(+)
TemPhD_cluster_50324 12604:20731(+)
IMGVR_UViG_2663763484_000008|2663763484|2663864539|20050-55721 11699:18760(+)
TemPhD_cluster_61875 10411:20726(+)
Paris_II__AriB_DUF4435_II___AriB_DUF4435_II_ali (CDS_0020)
superinfection exclusion Sie-like (CDS_0022)
Paris_II__AriA_AAA21___All_AAA21_ali (CDS_0019)
TemPhD_cluster_50048 10694:22443(+)
superinfection exclusion Sie-like (CDS_0018)
IMGVR_UViG_2529293013_000008|2529293013|2529389927|58120-95613 10399:20268(+)
IMGVR_UViG_2660238130_000010|2660238130|2660268366|58998-97018 10747:20862(+)
superinfection exclusion Sie-like (CDS_0023)
IMGVR_UViG_2923043281_000006|2923043281|2923043281|3707176-3746347 11685:21440(+)
superinfection exclusion Sie-like (CDS_0012)
IMGVR_UViG_2785510767_000002|2785510767|2785512818|1206508-1242104 9319:18619(+)
IMGVR_UViG_2925541682_000007|2925541682|2925541682|3404939-3442640 11438:20976(+)
IMGVR_UViG_3300014901_000006|3300014901|Ga0121380_100015|103175-140062 11077:19346(+)
TemPhD_cluster_61950 11496:19762(+)
IMGVR_UViG_3300013703_000024|3300013703|Ga0120871_100041|17482-54069 11196:19439(+)
TemPhD_cluster_61395 12647:21395(+)
IMGVR_UViG_3300011894_000001|3300011894|Ga0122084_100065 11533:21154(+)
IMGVR_UViG_2700989548_000007|2700989548|2701096446|173697-210296 10800:20181(+)
IMGVR_UViG_3300013482_000011|3300013482|Ga0122080_10028|19399-57912 11077:21683(+)
IMGVR_UViG_2667527877_000004|2667527877|2667581225|124374-162826 10541:20905(+)
Paris_II__AriB_DUF4435_II___AriB_DUF4435_II_ali (CDS_0020)
Paris_II__AriA_AAA21___All_AAA21_ali (CDS_0019)
TemPhD_cluster_42611 10499:21824(+)
Paris_II__AriB_DUF4435_II___AriB_DUF4435_II_ali (CDS_0020)
Paris_II__AriA_AAA21___All_AAA21_ali (CDS_0019)
IMGVR_UViG_2657245452_000005|2657245452|2657301756|123737-160778 10737:21608(+)
4000 nt
DNA, RNA and nucleotide metabolism
head and packaging
moron, auxiliary metabolic gene and host takeover
other
tail
transcription regulation
unknown function
Defence