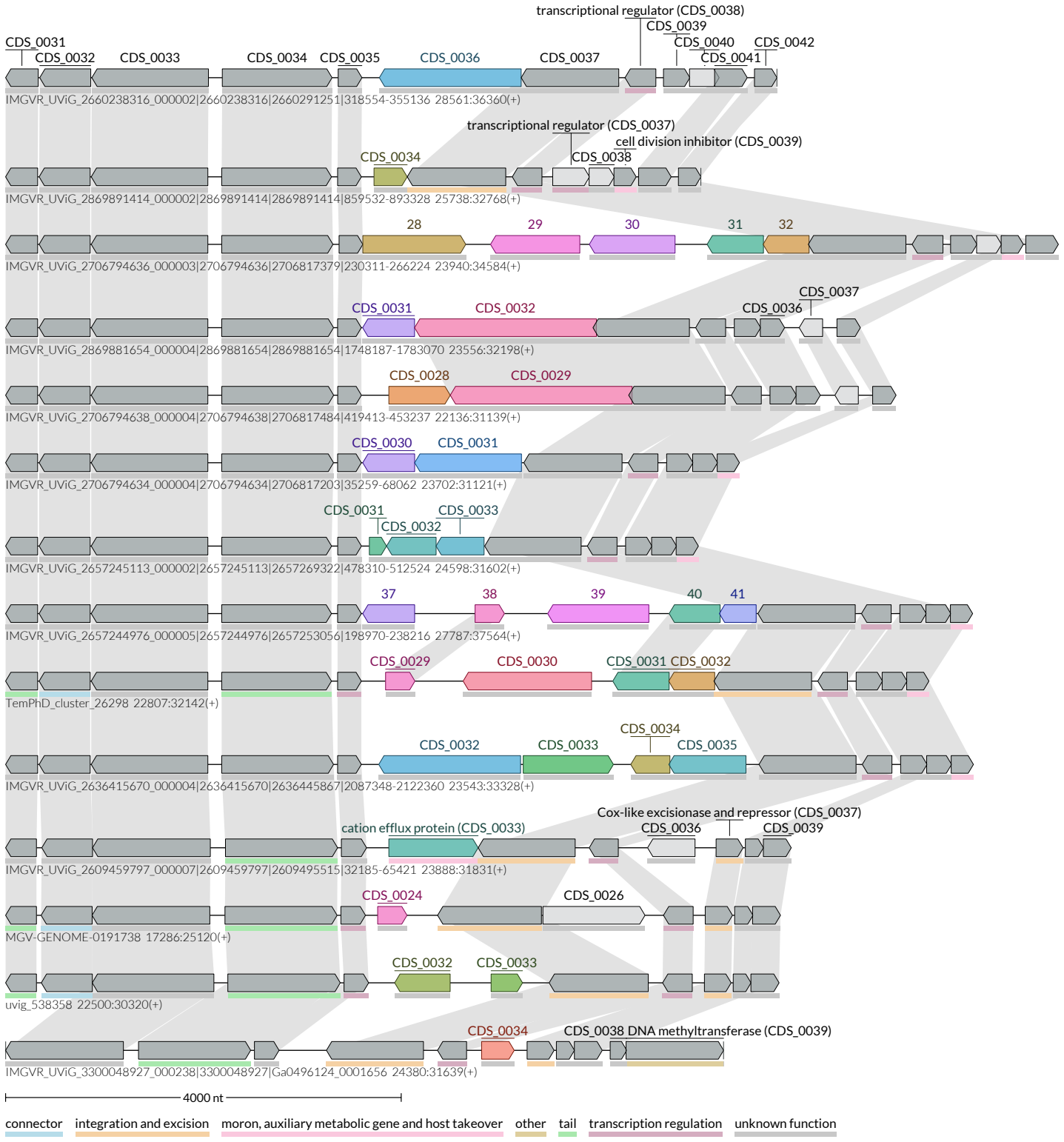
transcriptional regulator (CDS_0038)
CDS_0039
CDS_0031
CDS_0032
CDS_0033
CDS_0034
CDS_0035
CDS_0036
CDS_0037
CDS_0040
CDS_0041
CDS_0042
IMGVR_UViG_2660238316_000002|2660238316|2660291251|318554-355136 28561:36360(+)
transcriptional regulator (CDS_0037)
cell division inhibitor (CDS_0039)
CDS_0034
CDS_0038
IMGVR_UViG_2869891414_000002|2869891414|2869891414|859532-893328 25738:32768(+)
28
29
30
31
32
IMGVR_UViG_2706794636_000003|2706794636|2706817379|230311-266224 23940:34584(+)
CDS_0037
CDS_0031
CDS_0032
CDS_0036
IMGVR_UViG_2869881654_000004|2869881654|2869881654|1748187-1783070 23556:32198(+)
CDS_0028
CDS_0029
IMGVR_UViG_2706794638_000004|2706794638|2706817484|419413-453237 22136:31139(+)
CDS_0030
CDS_0031
IMGVR_UViG_2706794634_000004|2706794634|2706817203|35259-68062 23702:31121(+)
CDS_0031
CDS_0033
CDS_0032
IMGVR_UViG_2657245113_000002|2657245113|2657269322|478310-512524 24598:31602(+)
37
38
39
40
41
IMGVR_UViG_2657244976_000005|2657244976|2657253056|198970-238216 27787:37564(+)
CDS_0029
CDS_0030
CDS_0031
CDS_0032
TemPhD_cluster_26298 22807:32142(+)
CDS_0034
CDS_0032
CDS_0033
CDS_0035
IMGVR_UViG_2636415670_000004|2636415670|2636445867|2087348-2122360 23543:33328(+)
Cox-like excisionase and repressor (CDS_0037)
cation efflux protein (CDS_0033)
CDS_0036
CDS_0039
IMGVR_UViG_2609459797_000007|2609459797|2609495515|32185-65421 23888:31831(+)
CDS_0024
CDS_0026
MGV-GENOME-0191738 17286:25120(+)
CDS_0032
CDS_0033
uvig_538358 22500:30320(+)
CDS_0034
CDS_0038 DNA methyltransferase (CDS_0039)
IMGVR_UViG_3300048927_000238|3300048927|Ga0496124_0001656 24380:31639(+)
4000 nt
connector
integration and excision
moron, auxiliary metabolic gene and host takeover
other
tail
transcription regulation
unknown function