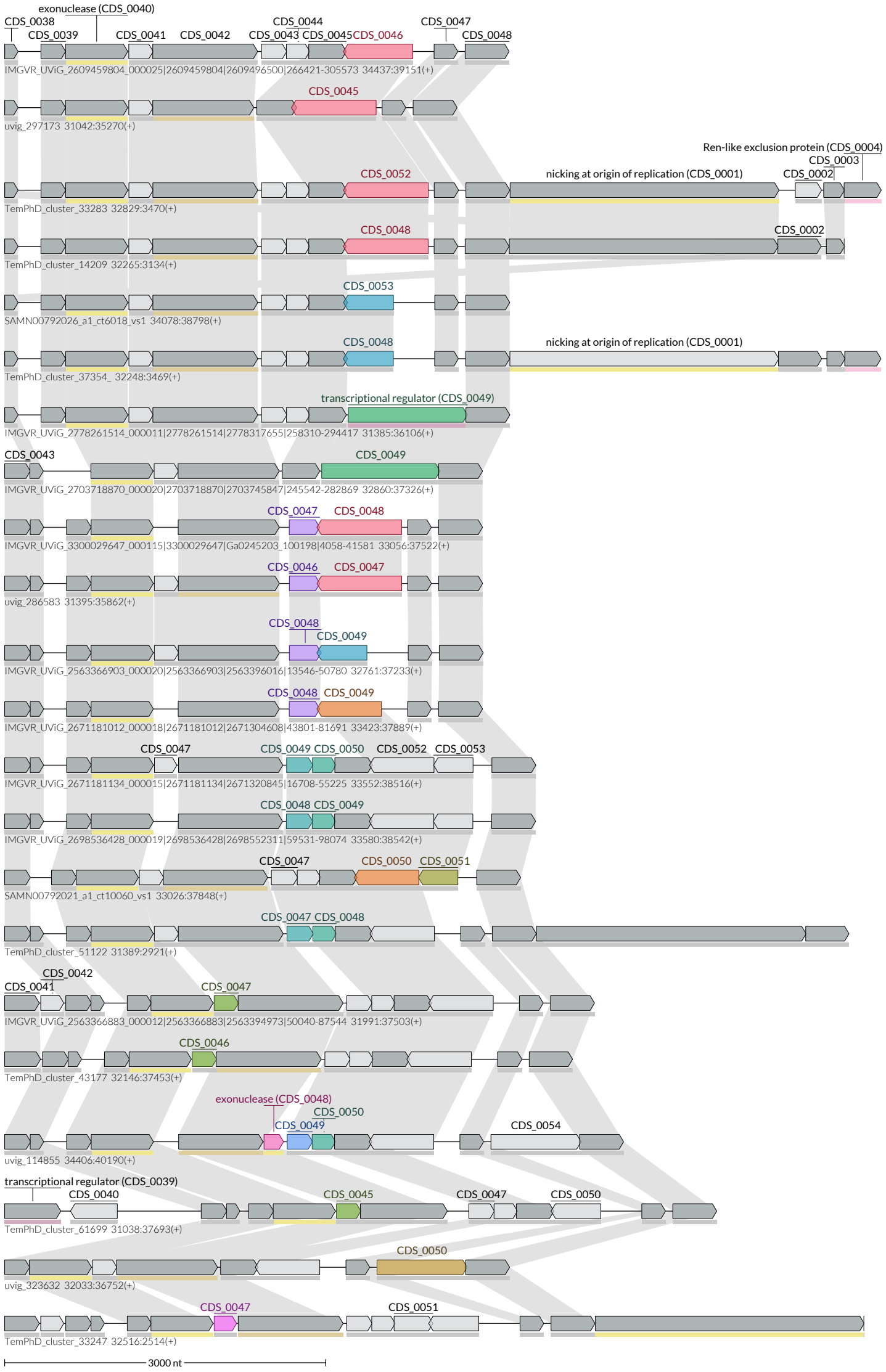
exonuclease (CDS_0040)
CDS_0038
CDS_0039
CDS_0041
CDS_0042
CDS_0044
CDS_0043
CDS_0045
CDS_0046
CDS_0047
CDS_0048
IMGVR_UViG_2609459804_000025|2609459804|2609496500|266421-305573 34437:39151(+)
CDS_0045
uvig_297173 31042:35270(+)
Ren-like exclusion protein (CDS_0004)
CDS_0003
CDS_0052
nicking at origin of replication (CDS_0001)
CDS_0002
TemPhD_cluster_33283 32829:3470(+)
CDS_0048
CDS_0002
TemPhD_cluster_14209 32265:3134(+)
CDS_0053
SAMN00792026_a1_ct6018_vs1 34078:38798(+)
CDS_0048
nicking at origin of replication (CDS_0001)
TemPhD_cluster_37354_ 32248:3469(+)
transcriptional regulator (CDS_0049)
IMGVR_UViG_2778261514_000011|2778261514|2778317655|258310-294417 31385:36106(+)
CDS_0043
CDS_0049
IMGVR_UViG_2703718870_000020|2703718870|2703745847|245542-282869 32860:37326(+)
CDS_0047
CDS_0048
IMGVR_UViG_3300029647_000115|3300029647|Ga0245203_100198|4058-41581 33056:37522(+)
CDS_0046
CDS_0047
uvig_286583 31395:35862(+)
CDS_0048
CDS_0049
IMGVR_UViG_2563366903_000020|2563366903|2563396016|13546-50780 32761:37233(+)
CDS_0048
CDS_0049
IMGVR_UViG_2671181012_000018|2671181012|2671304608|43801-81691 33423:37889(+)
CDS_0047
CDS_0049
CDS_0050
CDS_0052
CDS_0053
IMGVR_UViG_2671181134_000015|2671181134|2671320845|16708-55225 33552:38516(+)
CDS_0048
CDS_0049
IMGVR_UViG_2698536428_000019|2698536428|2698552311|59531-98074 33580:38542(+)
CDS_0047
CDS_0050
CDS_0051
SAMN00792021_a1_ct10060_vs1 33026:37848(+)
CDS_0047
CDS_0048
TemPhD_cluster_51122 31389:2921(+)
CDS_0042
CDS_0041
CDS_0047
IMGVR_UViG_2563366883_000012|2563366883|2563394973|50040-87544 31991:37503(+)
CDS_0046
TemPhD_cluster_43177 32146:37453(+)
exonuclease (CDS_0048)
CDS_0050
CDS_0049
CDS_0054
uvig_114855 34406:40190(+)
transcriptional regulator (CDS_0039)
CDS_0040
CDS_0045
CDS_0047
CDS_0050
TemPhD_cluster_61699 31038:37693(+)
CDS_0050
uvig_323632 32033:36752(+)
CDS_0047
CDS_0051
TemPhD_cluster_33247 32516:2514(+)
3000 nt
DNA, RNA and nucleotide metabolism
moron, auxiliary metabolic gene and host takeover
other
transcription regulation
unknown function