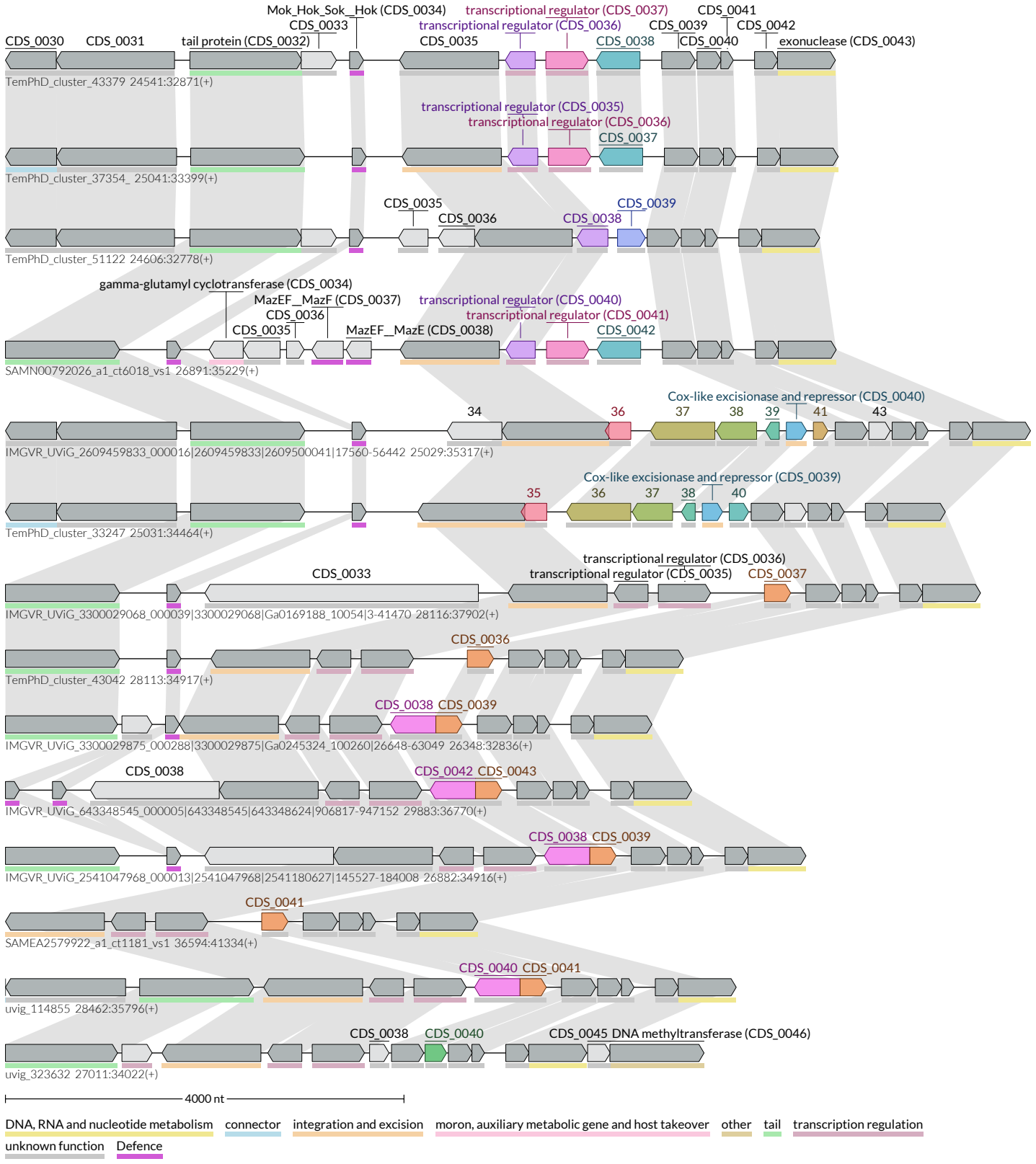
Mok_Hok_Sok__Hok (CDS_0034)
transcriptional regulator (CDS_0037)
CDS_0041
CDS_0033
transcriptional regulator (CDS_0036)
CDS_0039
CDS_0042
CDS_0030
CDS_0031
tail protein (CDS_0032)
CDS_0035
CDS_0038
CDS_0040
exonuclease (CDS_0043)
TemPhD_cluster_43379 24541:32871(+)
transcriptional regulator (CDS_0035)
transcriptional regulator (CDS_0036)
CDS_0037
TemPhD_cluster_37354_ 25041:33399(+)
CDS_0035
CDS_0039
CDS_0036
CDS_0038
TemPhD_cluster_51122 24606:32778(+)
gamma-glutamyl cyclotransferase (CDS_0034)
MazEF__MazF (CDS_0037)
transcriptional regulator (CDS_0040)
CDS_0036
transcriptional regulator (CDS_0041)
CDS_0035
MazEF__MazE (CDS_0038)
CDS_0042
SAMN00792026_a1_ct6018_vs1 26891:35229(+)
Cox-like excisionase and repressor (CDS_0040)
34
36
37
38
39
41
43
IMGVR_UViG_2609459833_000016|2609459833|2609500041|17560-56442 25029:35317(+)
Cox-like excisionase and repressor (CDS_0039)
35
36
37
38
40
TemPhD_cluster_33247 25031:34464(+)
transcriptional regulator (CDS_0036)
CDS_0033
transcriptional regulator (CDS_0035)
CDS_0037
IMGVR_UViG_3300029068_000039|3300029068|Ga0169188_10054|3-41470 28116:37902(+)
CDS_0036
TemPhD_cluster_43042 28113:34917(+)
CDS_0038 CDS_0039
IMGVR_UViG_3300029875_000288|3300029875|Ga0245324_100260|26648-63049 26348:32836(+)
CDS_0038
CDS_0042 CDS_0043
IMGVR_UViG_643348545_000005|643348545|643348624|906817-947152 29883:36770(+)
CDS_0038 CDS_0039
IMGVR_UViG_2541047968_000013|2541047968|2541180627|145527-184008 26882:34916(+)
CDS_0041
SAMEA2579922_a1_ct1181_vs1 36594:41334(+)
CDS_0040 CDS_0041
uvig_114855 28462:35796(+)
CDS_0038
CDS_0040
CDS_0045 DNA methyltransferase (CDS_0046)
uvig_323632 27011:34022(+)
4000 nt
DNA, RNA and nucleotide metabolism
connector
integration and excision
moron, auxiliary metabolic gene and host takeover
other
tail
transcription regulation
unknown function
Defence