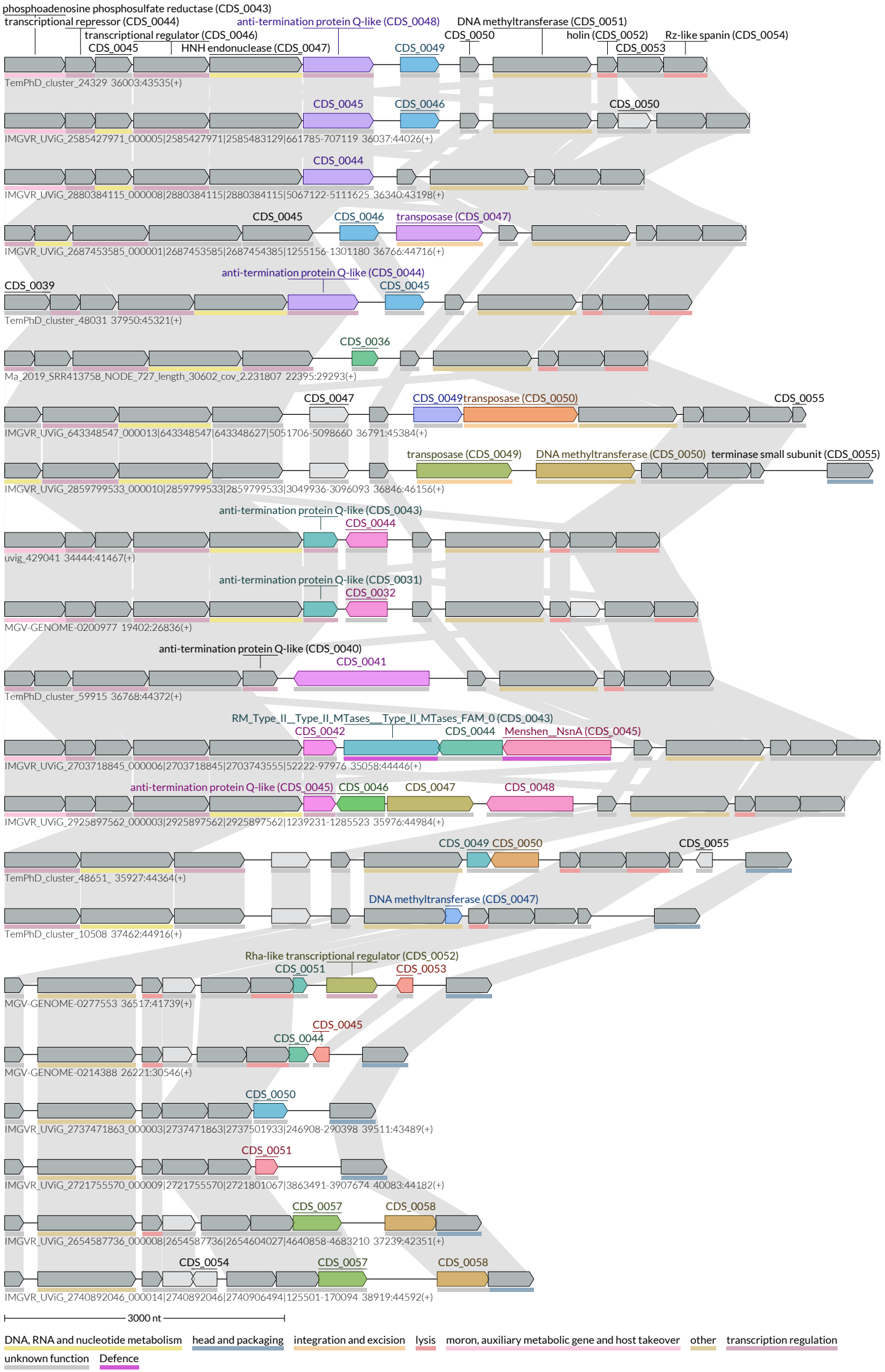
phosphoadenosine phosphosulfate reductase (CDS_0043)
transcriptional repressor (CDS_0044)
anti-termination protein Q-like (CDS_0048)
DNA methyltransferase (CDS_0051)
transcriptional regulator (CDS_0046)
CDS_0050
holin (CDS_0052)
Rz-like spanin (CDS_0054)
CDS_0045
HNH endonuclease (CDS_0047)
CDS_0049
CDS_0053
TemPhD_cluster_24329 36003:43535(+)
CDS_0045
CDS_0046
CDS_0050
IMGVR_UViG_2585427971_000005|2585427971|2585483129|661785-707119 36037:44026(+)
CDS_0044
IMGVR_UViG_2880384115_000008|2880384115|2880384115|5067122-5111625 36340:43198(+)
CDS_0045
CDS_0046
transposase (CDS_0047)
IMGVR_UViG_2687453585_000001|2687453585|2687454385|1255156-1301180 36766:44716(+)
anti-termination protein Q-like (CDS_0044)
CDS_0039
CDS_0045
TemPhD_cluster_48031 37950:45321(+)
CDS_0036
Ma_2019_SRR413758_NODE_727_length_30602_cov_2.231807 22395:29293(+)
CDS_0047
CDS_0049
transposase (CDS_0050)
CDS_0055
IMGVR_UViG_643348547_000013|643348547|643348627|5051706-5098660 36791:45384(+)
transposase (CDS_0049)
DNA methyltransferase (CDS_0050)
terminase small subunit (CDS_0055)
IMGVR_UViG_2859799533_000010|2859799533|2859799533|3049936-3096093 36846:46156(+)
anti-termination protein Q-like (CDS_0043)
CDS_0044
uvig_429041 34444:41467(+)
anti-termination protein Q-like (CDS_0031)
CDS_0032
MGV-GENOME-0200977 19402:26836(+)
anti-termination protein Q-like (CDS_0040)
CDS_0041
TemPhD_cluster_59915 36768:44372(+)
RM_Type_II__Type_II_MTases__Type_II_MTases_FAM_0 (CDS_0043)
CDS_0042
CDS_0044
Menshen__NsnA (CDS_0045)
IMGVR_UViG_2703718845_000006|2703718845|2703743555|52222-97976 35058:44446(+)
anti-termination protein Q-like (CDS_0045)
CDS_0046
CDS_0047
CDS_0048
IMGVR_UViG_2925897562_000003|2925897562|2925897562|1239231-1285523 35976:44984(+)
CDS_0049
CDS_0050
CDS_0055
TemPhD_cluster_48651_ 35927:44364(+)
DNA methyltransferase (CDS_0047)
TemPhD_cluster_10508 37462:44916(+)
Rha-like transcriptional regulator (CDS_0052)
CDS_0051
CDS_0053
MGV-GENOME-0277553 36517:41739(+)
CDS_0045
CDS_0044
MGV-GENOME-0214388 26221:30546(+)
CDS_0050
IMGVR_UViG_2737471863_000003|2737471863|2737501933|246908-290398 39511:43489(+)
CDS_0051
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CDS_0057
CDS_0058
IMGVR_UViG_2654587736_000008|2654587736|2654604027|4640858-4683210 37239:42351(+)
CDS_0054
CDS_0057
CDS_0058
IMGVR_UViG_2740892046_000014|2740892046|2740906494|125501-170094 38919:44592(+)
3000 nt
DNA, RNA and nucleotide metabolism
head and packaging
integration and excision
lysis
moron, auxiliary metabolic gene and host takeover
other
transcription regulation
unknown function
Defence