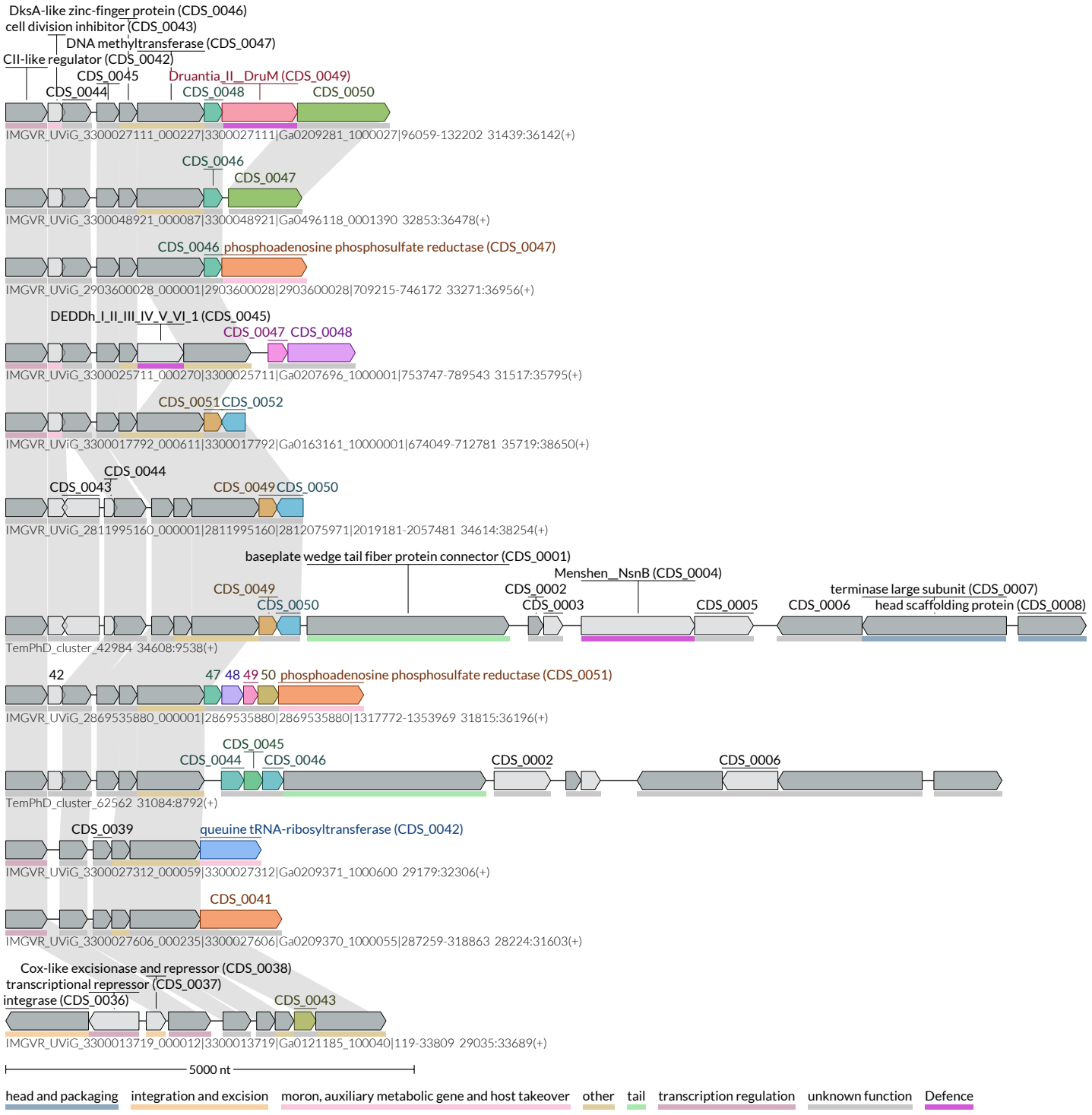
DksA-like zinc-finger protein (CDS_0046)
cell division inhibitor (CDS_0043)
DNA methyltransferase (CDS_0047)
CII-like regulator (CDS_0042)
CDS_0045
Druantia_II__DruM (CDS_0049)
CDS_0044
CDS_0048
CDS_0050
IMGVR_UViG_3300027111_000227|3300027111|Ga0209281_1000027|96059-132202 31439:36142(+)
CDS_0046
CDS_0047
IMGVR_UViG_3300048921_000087|3300048921|Ga0496118_0001390 32853:36478(+)
CDS_0046 phosphoadenosine phosphosulfate reductase (CDS_0047)
IMGVR_UViG_2903600028_000001|2903600028|2903600028|709215-746172 33271:36956(+)
DEDDh_I_II_III_IV_V_VI_1 (CDS_0045)
CDS_0047 CDS_0048
IMGVR_UViG_3300025711_000270|3300025711|Ga0207696_1000001|753747-789543 31517:35795(+)
CDS_0051 CDS_0052
IMGVR_UViG_3300017792_000611|3300017792|Ga0163161_10000001|674049-712781 35719:38650(+)
CDS_0044
CDS_0043
CDS_0049 CDS_0050
IMGVR_UViG_2811995160_000001|2811995160|2812075971|2019181-2057481 34614:38254(+)
baseplate wedge tail fiber protein connector (CDS_0001)
Menshen__NsnB (CDS_0004)
CDS_0049
CDS_0002
terminase large subunit (CDS_0007)
CDS_0050
CDS_0003
CDS_0005
CDS_0006
head scaffolding protein (CDS_0008)
TemPhD_cluster_42984 34608:9538(+)
42
47 48 49 50 phosphoadenosine phosphosulfate reductase (CDS_0051)
IMGVR_UViG_2869535880_000001|2869535880|2869535880|1317772-1353969 31815:36196(+)
CDS_0045
CDS_0044
CDS_0046
CDS_0002
CDS_0006
TemPhD_cluster_62562 31084:8792(+)
CDS_0039
queuine tRNA-ribosyltransferase (CDS_0042)
IMGVR_UViG_3300027312_000059|3300027312|Ga0209371_1000600 29179:32306(+)
CDS_0041
IMGVR_UViG_3300027606_000235|3300027606|Ga0209370_1000055|287259-318863 28224:31603(+)
Cox-like excisionase and repressor (CDS_0038)
transcriptional repressor (CDS_0037)
integrase (CDS_0036)
CDS_0043
IMGVR_UViG_3300013719_000012|3300013719|Ga0121185_100040|119-33809 29035:33689(+)
5000 nt
head and packaging
integration and excision
moron, auxiliary metabolic gene and host takeover
other
tail
transcription regulation
unknown function
Defence