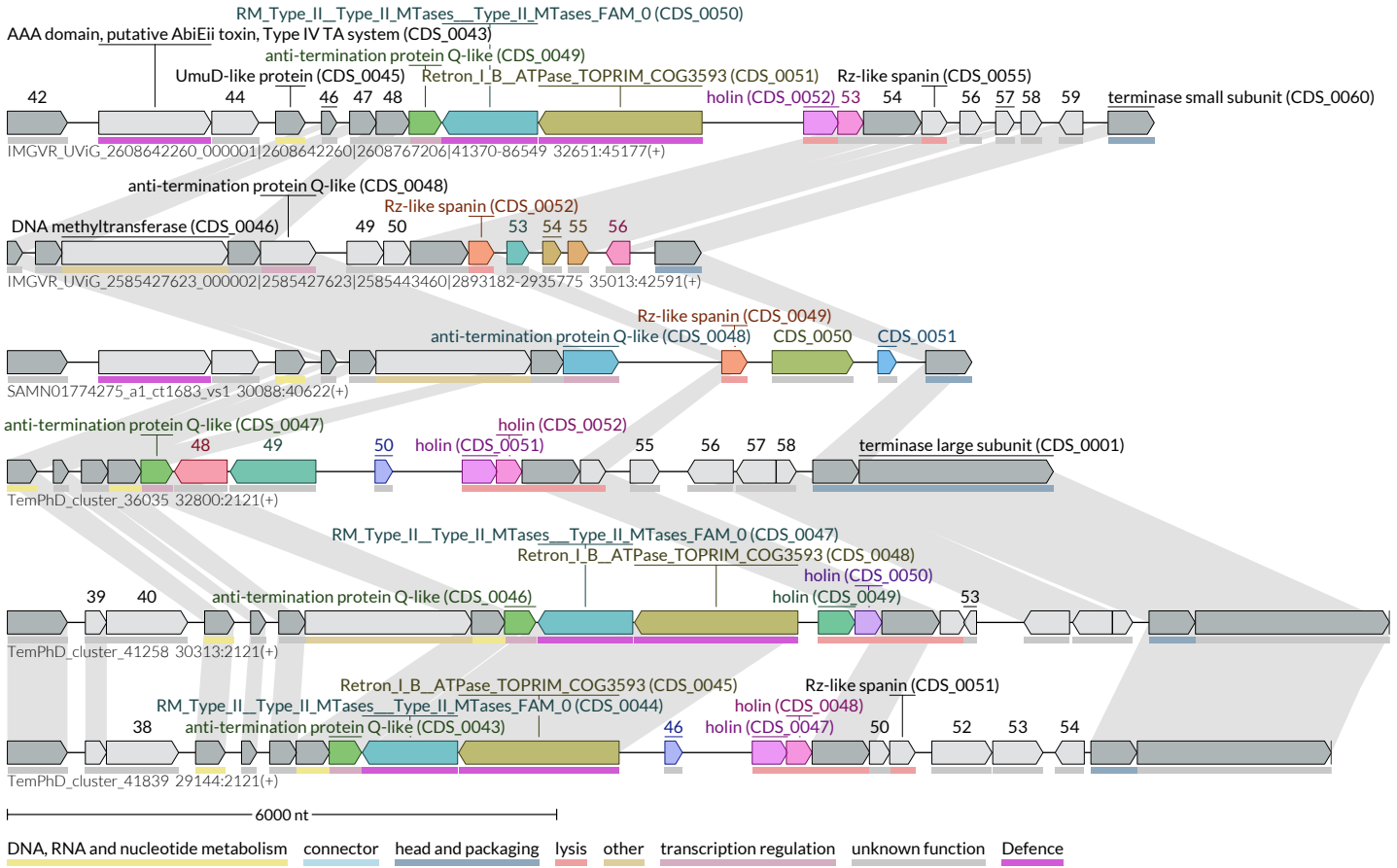
RM_Type_II__Type_II_MTases___Type_II_MTases_FAM_0 (CDS_0050)
AAA domain, putative AbiEii toxin, Type IV TA system (CDS_0043)
anti-termination protein Q-like (CDS_0049)
UmuD-like protein (CDS_0045)
Retron_I_B__ATPase_TOPRIM_COG3593 (CDS_0051)
Rz-like spanin (CDS_0055)
holin (CDS_0052)
terminase small subunit (CDS_0060)
42 44 46 47 48 53 54 56 57 58 59
IMGVR_UViG_2608642260_000001|2608642260|2608767206|41370-86549 32651:45177(+)
anti-termination protein Q-like (CDS_0048)
Rz-like spanin (CDS_0052)
DNA methyltransferase (CDS_0046)
49 50 53 54 55 56
IMGVR_UViG_2585427623_000002|2585427623|2585443460|2893182-2935775 35013:42591(+)
Rz-like spanin (CDS_0049)
anti-termination protein Q-like (CDS_0048)
CDS_0050
CDS_0051
SAMN01774275_a1_ct1683_vs1 30088:40622(+)
anti-termination protein Q-like (CDS_0047)
holin (CDS_0052)
48 49 50
holin (CDS_0051)
55 56 57 58
terminase large subunit (CDS_0001)
TemPhD_cluster_36035 32800:2121(+)
RM_Type_II__Type_II_MTases___Type_II_MTases_FAM_0 (CDS_0047)
Retron_I_B__ATPase_TOPRIM_COG3593 (CDS_0048)
holin (CDS_0050)
39 40
anti-termination protein Q-like (CDS_0046)
holin (CDS_0049)
53
TemPhD_cluster_41258 30313:2121(+)
Retron_I_B__ATPase_TOPRIM_COG3593 (CDS_0045)
Rz-like spanin (CDS_0051)
RM_Type_II__Type_II_MTases___Type_II_MTases_FAM_0 (CDS_0044)
holin (CDS_0048)
38
anti-termination protein Q-like (CDS_0043)
46
holin (CDS_0047)
50 52 53 54
TemPhD_cluster_41839 29144:2121(+)
6000 nt
DNA, RNA and nucleotide metabolism
connector
head and packaging
lysis
other
transcription regulation
unknown function
Defence