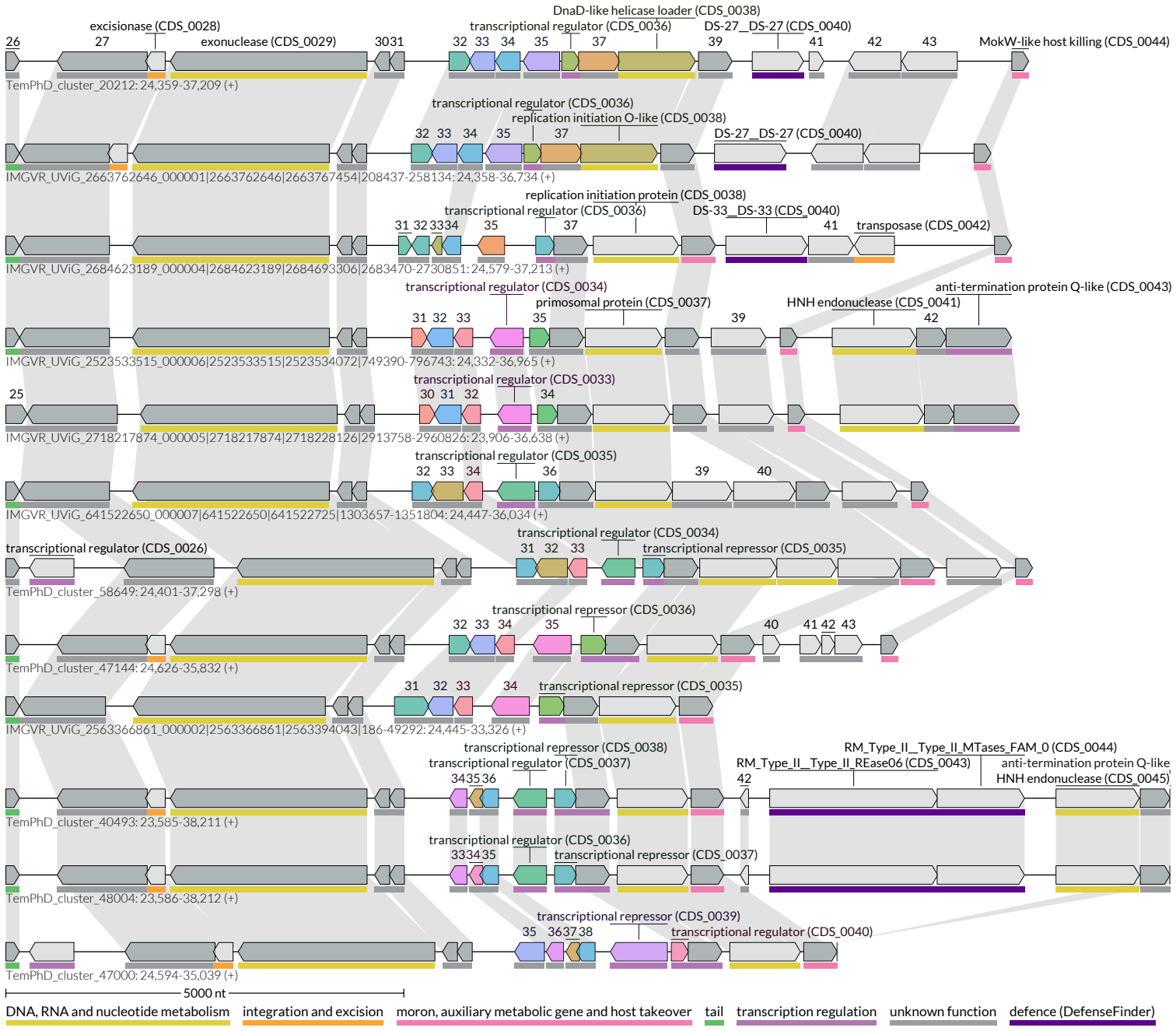
DnaD-like helicase loader (CDS_0038)
excisionase (CDS_0028)
transcriptional regulator (CDS_0036)
DS-27__DS-27 (CDS_0040)
exonuclease (CDS_0029)
MokW-like host killing (CDS_0044)
TemPhD_cluster_20212: 24,359-37,209 (+)
transcriptional regulator (CDS_0036)
replication initiation O-like (CDS_0038)
DS-27__DS-27 (CDS_0040)
IMGVR_UViG_2663762646_000001|2663762646|2663767454|208437-258134: 24,358-36,734 (+)
replication initiation protein (CDS_0038)
transcriptional regulator (CDS_0036)
DS-33__DS-33 (CDS_0040)
transposase (CDS_0042)
IMGVR_UViG_2684623189_000004|2684623189|2684693306|2683470-2730851: 24,579-37,213 (+)
transcriptional regulator (CDS_0034)
anti-termination protein Q-like (CDS_0043)
primosomal protein (CDS_0037)
HNH endonuclease (CDS_0041)
IMGVR_UViG_2523533515_000006|2523533515|2523534072|749390-796743: 24,332-36,965 (+)
transcriptional regulator (CDS_0033)
IMGVR_UViG_2718217874_000005|2718217874|2718228126|2913758-2960826: 23,906-36,638 (+)
transcriptional regulator (CDS_0035)
IMGVR_UViG_641522650_000007|641522650|641522725|1303657-1351804: 24,447-36,034 (+)
transcriptional regulator (CDS_0026)
transcriptional regulator (CDS_0034)
transcriptional repressor (CDS_0035)
TemPhD_cluster_58649: 24,401-37,298 (+)
transcriptional repressor (CDS_0036)
TemPhD_cluster_47144: 24,626-35,832 (+)
transcriptional repressor (CDS_0035)
IMGVR_UViG_2563366861_000002|2563366861|2563394043|186-49292: 24,445-33,326 (+)
transcriptional repressor (CDS_0038)
transcriptional regulator (CDS_0037)
RM_Type_II__Type_II_MTases_FAM_0 (CDS_0044)
RM_Type_II__Type_II_REase06 (CDS_0043)
anti-termination protein Q-like
HNH endonuclease (CDS_0045)
TemPhD_cluster_40493: 23,585-38,211 (+)
transcriptional regulator (CDS_0036)
transcriptional repressor (CDS_0037)
TemPhD_cluster_48004: 23,586-38,212 (+)
transcriptional repressor (CDS_0039)
transcriptional regulator (CDS_0040)
TemPhD_cluster_47000: 24,594-35,039 (+)
5000 nt
DNA, RNA and nucleotide metabolism
integration and excision
moron, auxiliary metabolic gene and host takeover
tail
transcription regulation
unknown function
defence (DefenseFinder)