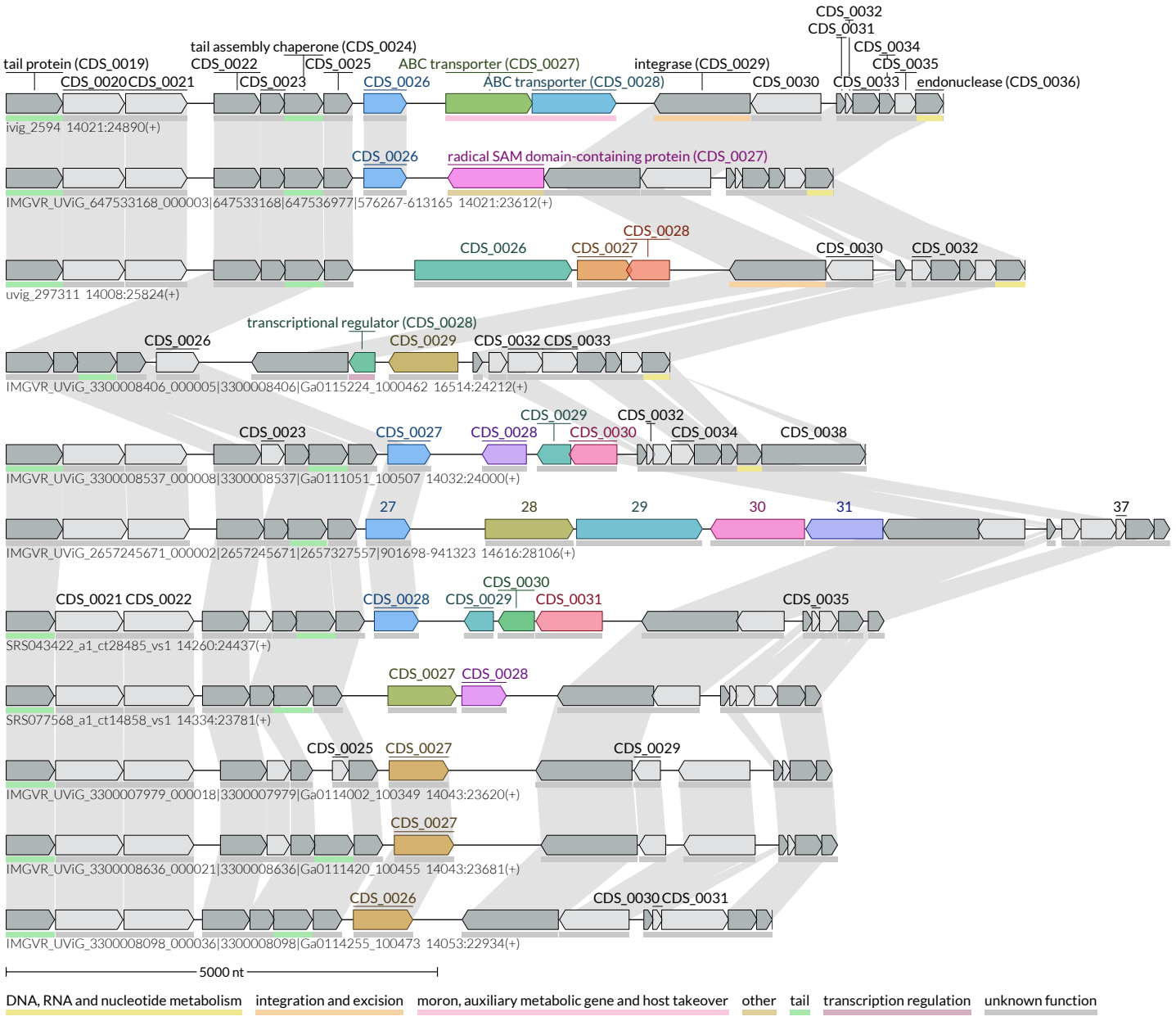
CDS_0032
CDS_0031
CDS_0034
CDS_0035
tail assembly chaperone (CDS_0024)
tail protein (CDS_0019)
CDS_0022
CDS_0025
ABC transporter (CDS_0027)
integrase (CDS_0029)
CDS_0020
CDS_0021
CDS_0023
CDS_0026
ABC transporter (CDS_0028)
CDS_0030
CDS_0033
endonuclease (CDS_0036)
ivig_2594 14021:24890(+)
CDS_0026
radical SAM domain-containing protein (CDS_0027)
IMGVR_UViG_647533168_000003|647533168|647536977|576267-613165 14021:23612(+)
CDS_0028
CDS_0026
CDS_0027
CDS_0030
CDS_0032
uvig_297311 14008:25824(+)
transcriptional regulator (CDS_0028)
CDS_0026
CDS_0029
CDS_0032
CDS_0033
IMGVR_UViG_3300008406_000005|3300008406|Ga0115224_1000462 16514:24212(+)
CDS_0029
CDS_0032
CDS_0023
CDS_0027
CDS_0028
CDS_0030
CDS_0034
CDS_0038
IMGVR_UViG_3300008537_000008|3300008537|Ga0111051_100507 14032:24000(+)
27
28
29
30
31
37
IMGVR_UViG_2657245671_000002|2657245671|2657327557|901698-941323 14616:28106(+)
CDS_0030
CDS_0021
CDS_0022
CDS_0028
CDS_0029
CDS_0031
CDS_0035
SRS043422_a1_ct28485_vs1 14260:24437(+)
CDS_0027
CDS_0028
SRS077568_a1_ct14858_vs1 14334:23781(+)
CDS_0025
CDS_0027
CDS_0029
IMGVR_UViG_3300007979_000018|3300007979|Ga0114002_100349 14043:23620(+)
CDS_0027
IMGVR_UViG_3300008636_000021|3300008636|Ga0111420_100455 14043:23681(+)
CDS_0026
CDS_0030
CDS_0031
IMGVR_UViG_3300008098_000036|3300008098|Ga0114255_100473 14053:22934(+)
5000 nt
DNA, RNA and nucleotide metabolism
integration and excision
moron, auxiliary metabolic gene and host takeover
other
tail
transcription regulation
unknown function