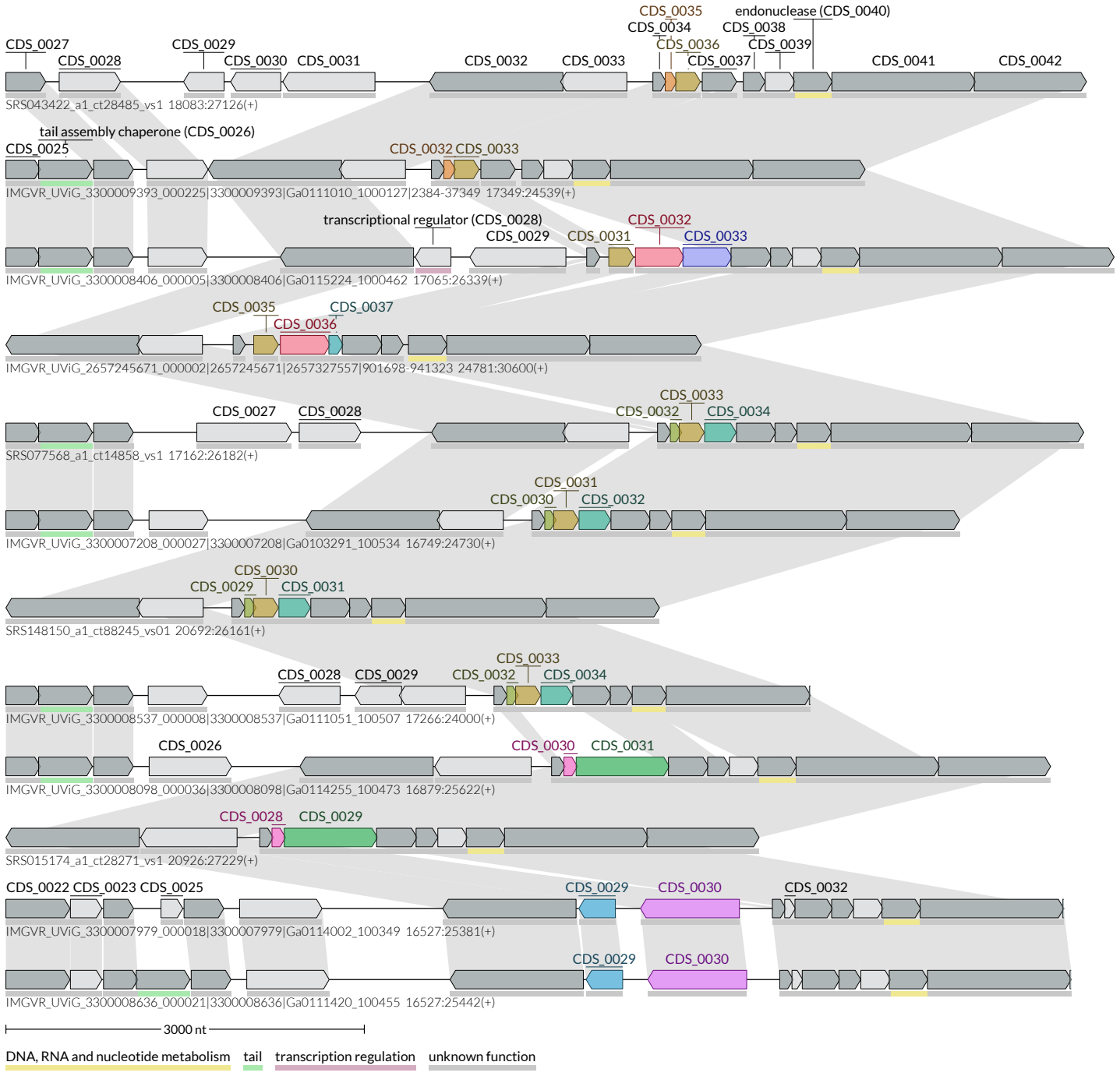
CDS_0035
endonuclease (CDS_0040)
CDS_0034
CDS_0038
CDS_0036
CDS_0039
CDS_0027
CDS_0029
CDS_0028
CDS_0030
CDS_0031
CDS_0032
CDS_0033
CDS_0037
CDS_0041
CDS_0042
SRS043422_a1_ct28485_vs1 18083:27126(+)
tail assembly chaperone (CDS_0026)
CDS_0025
CDS_0032 CDS_0033
IMGVR_UViG_3300009393_000225|3300009393|Ga0111010_1000127|2384-37349 17349:24539(+)
transcriptional regulator (CDS_0028)
CDS_0032
CDS_0029
CDS_0031
CDS_0033
IMGVR_UViG_3300008406_000005|3300008406|Ga0115224_1000462 17065:26339(+)
CDS_0035
CDS_0037
CDS_0036
IMGVR_UViG_2657245671_000002|2657245671|2657327557|901698-941323 24781:30600(+)
CDS_0033
CDS_0027
CDS_0028
CDS_0032
CDS_0034
SRS077568_a1_ct14858_vs1 17162:26182(+)
CDS_0031
CDS_0030
CDS_0032
IMGVR_UViG_3300007208_000027|3300007208|Ga0103291_100534 16749:24730(+)
CDS_0030
CDS_0029
CDS_0031
SRS148150_a1_ct88245_vs01 20692:26161(+)
CDS_0033
CDS_0028
CDS_0029
CDS_0032
CDS_0034
IMGVR_UViG_3300008537_000008|3300008537|Ga0111051_100507 17266:24000(+)
CDS_0026
CDS_0030
CDS_0031
IMGVR_UViG_3300008098_000036|3300008098|Ga0114255_100473 16879:25622(+)
CDS_0028
CDS_0029
SRS015174_a1_ct28271_vs1 20926:27229(+)
CDS_0022 CDS_0023 CDS_0025
CDS_0029
CDS_0030
CDS_0032
IMGVR_UViG_3300007979_000018|3300007979|Ga0114002_100349 16527:25381(+)
CDS_0029
CDS_0030
IMGVR_UViG_3300008636_000021|3300008636|Ga0111420_100455 16527:25442(+)
3000 nt
DNA, RNA and nucleotide metabolism
tail
transcription regulation
unknown function