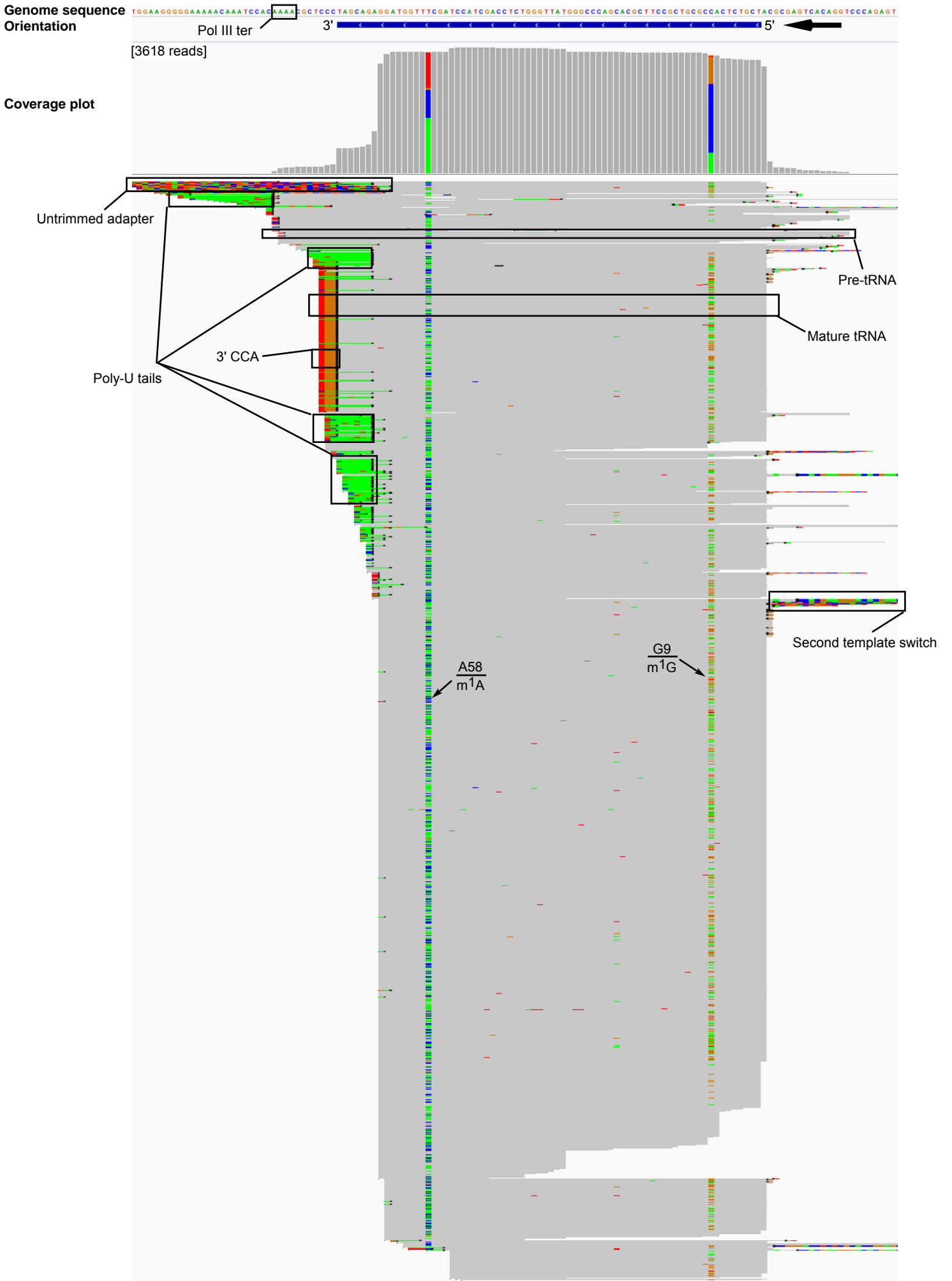
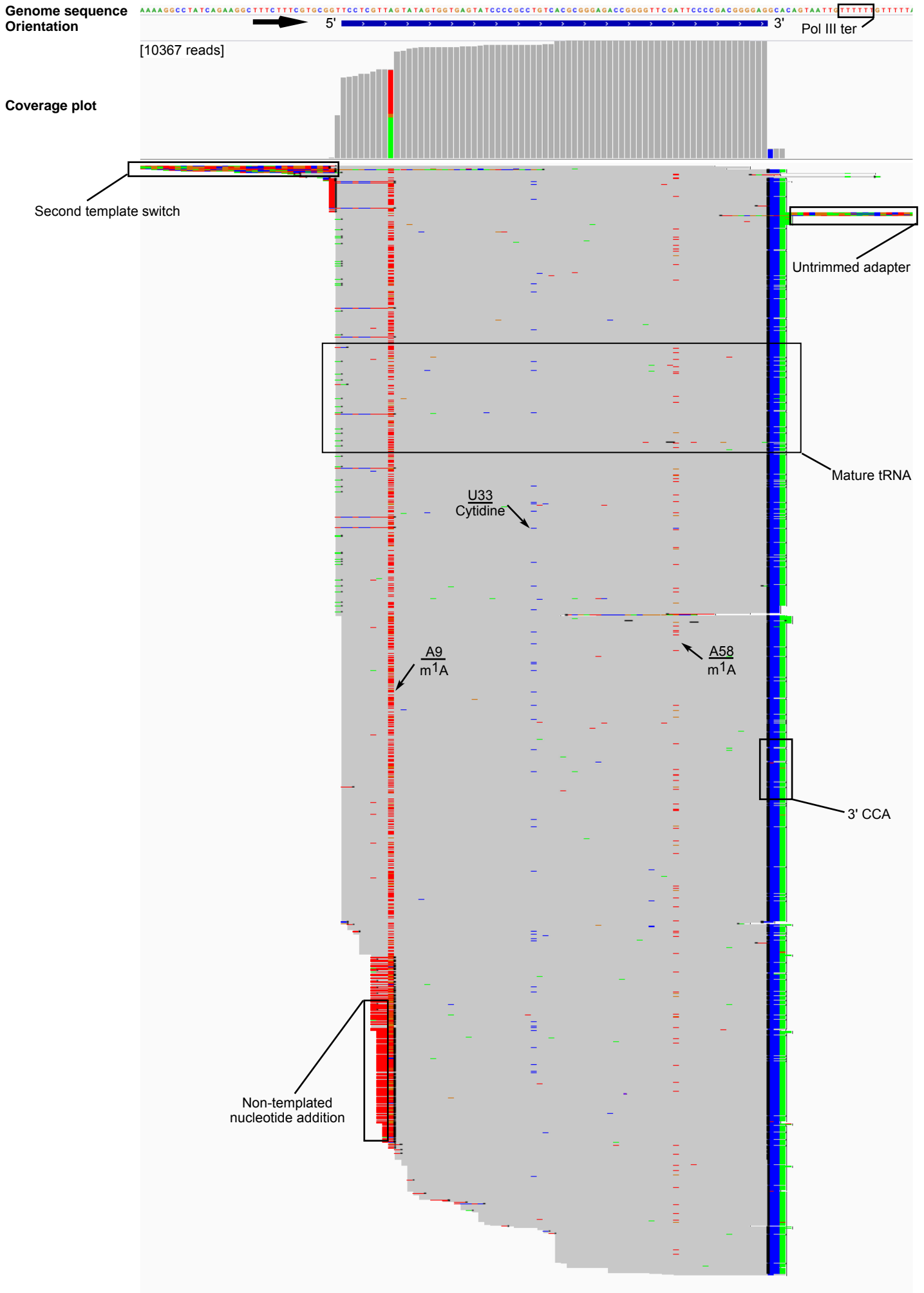


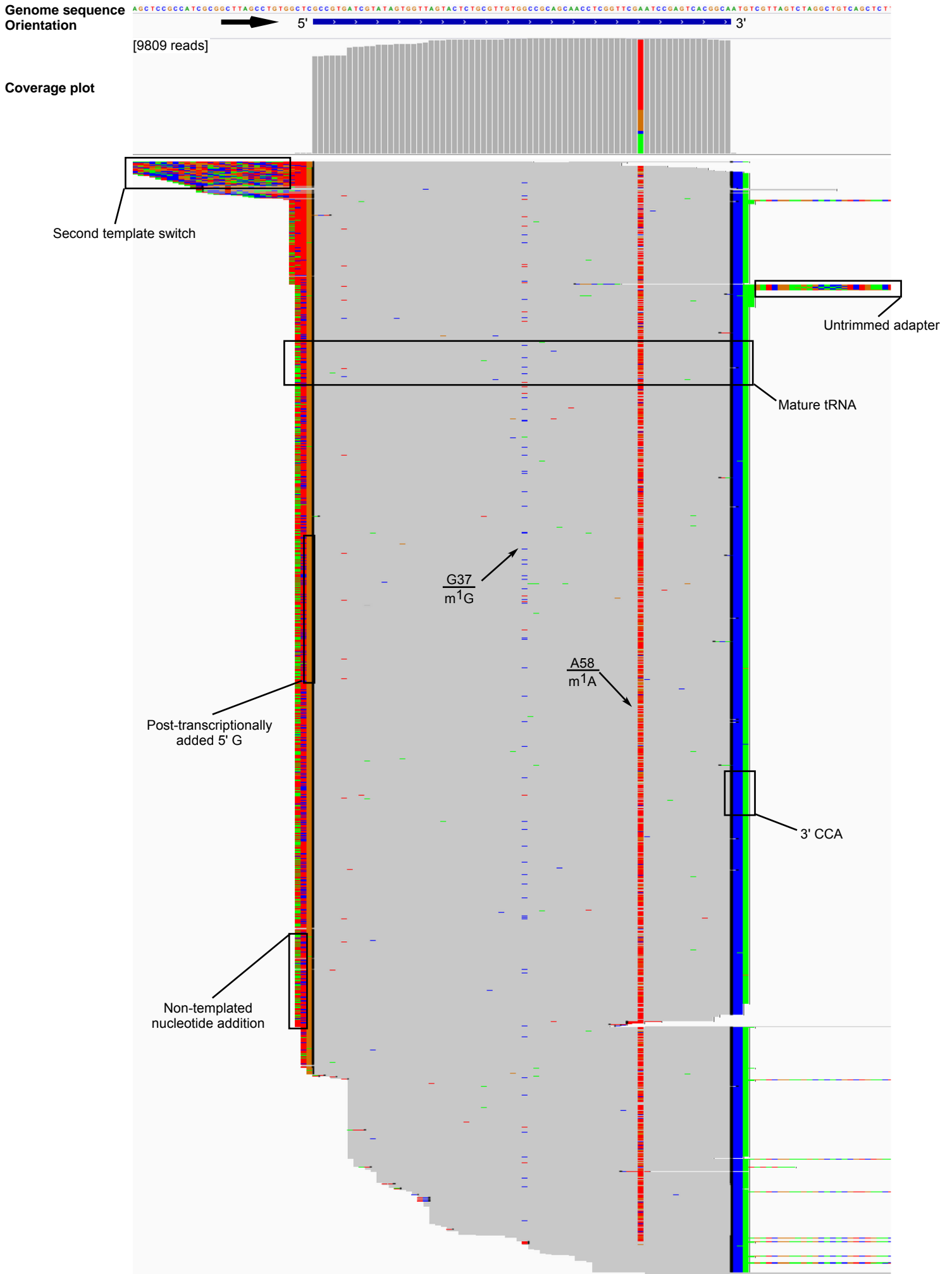
# chr6.tRNA169-iMetCAT



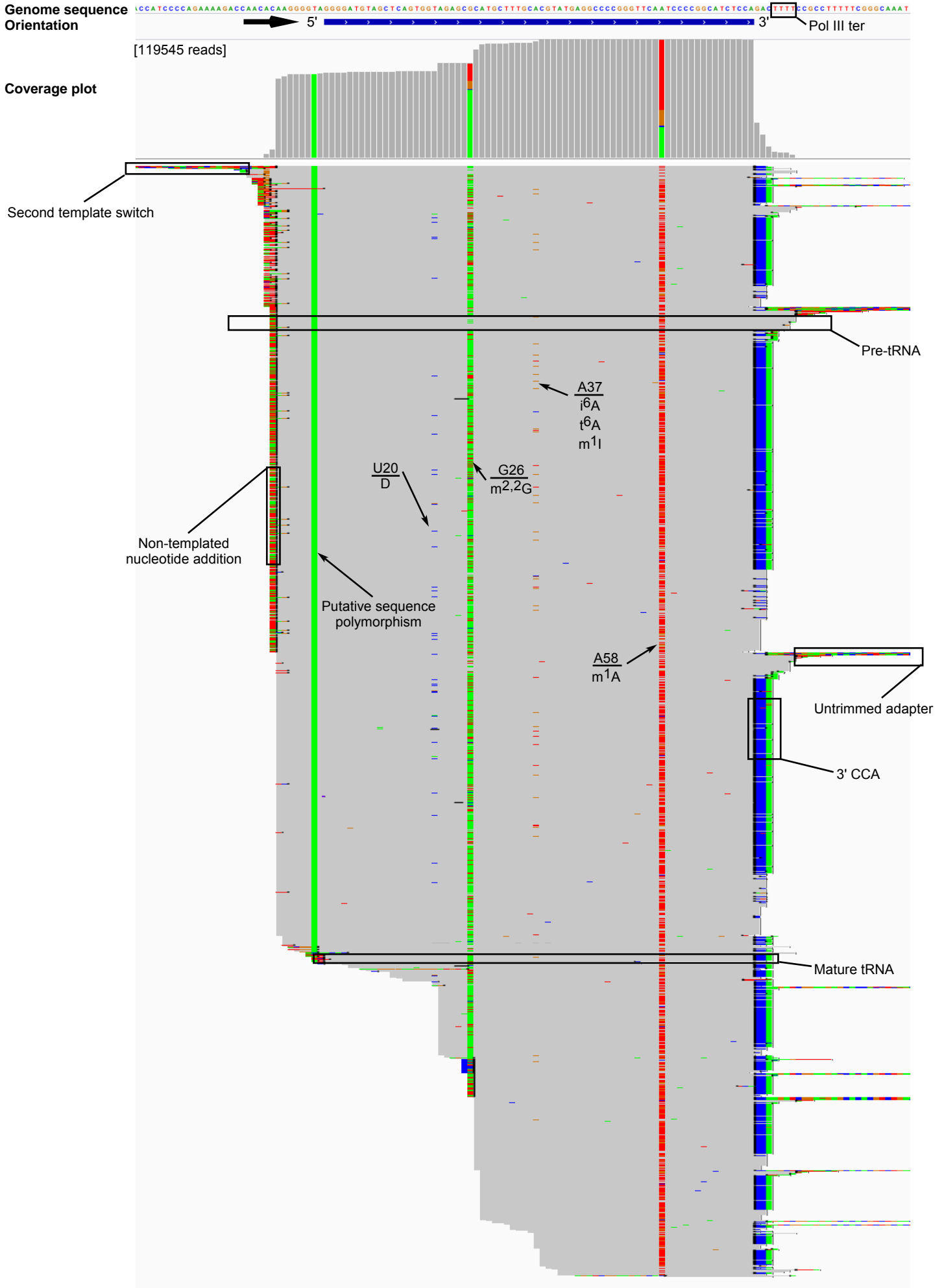
# chr12.tRNA4-AspGTC



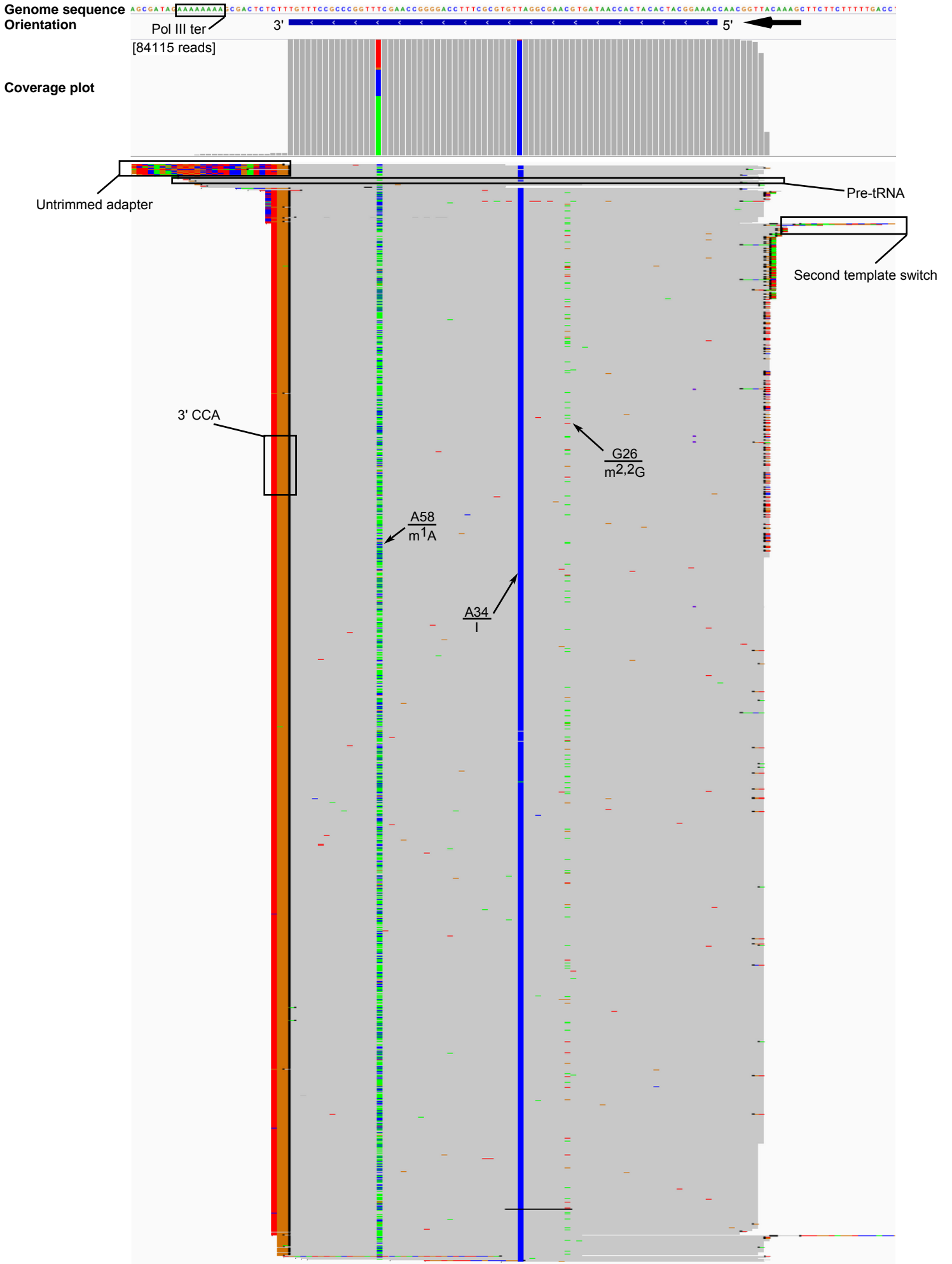
# chr1.tRNA21-HisGTG



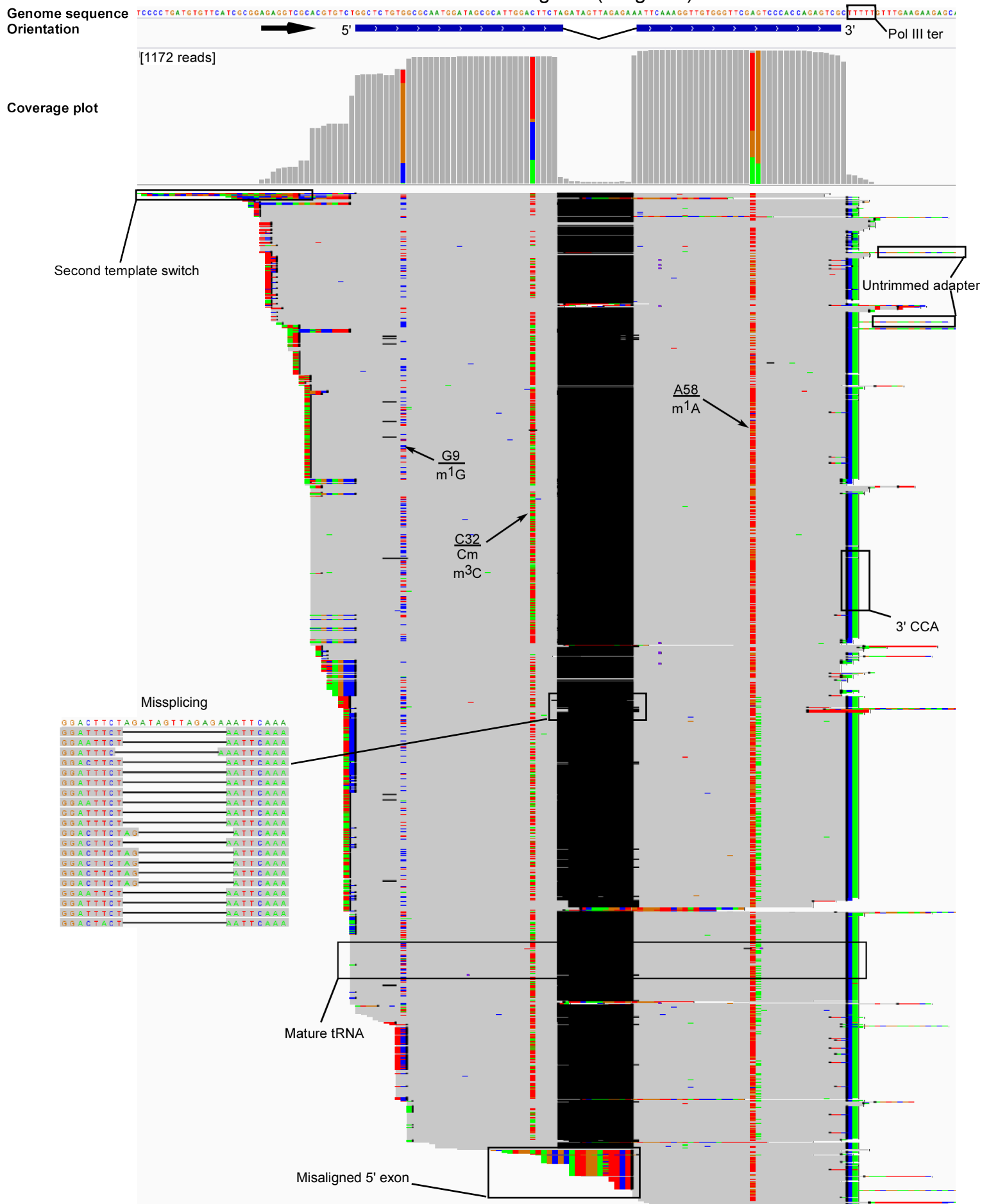
# chr12.tRNA8-AlaTGC



# chr6.tRNA132-ValAAC



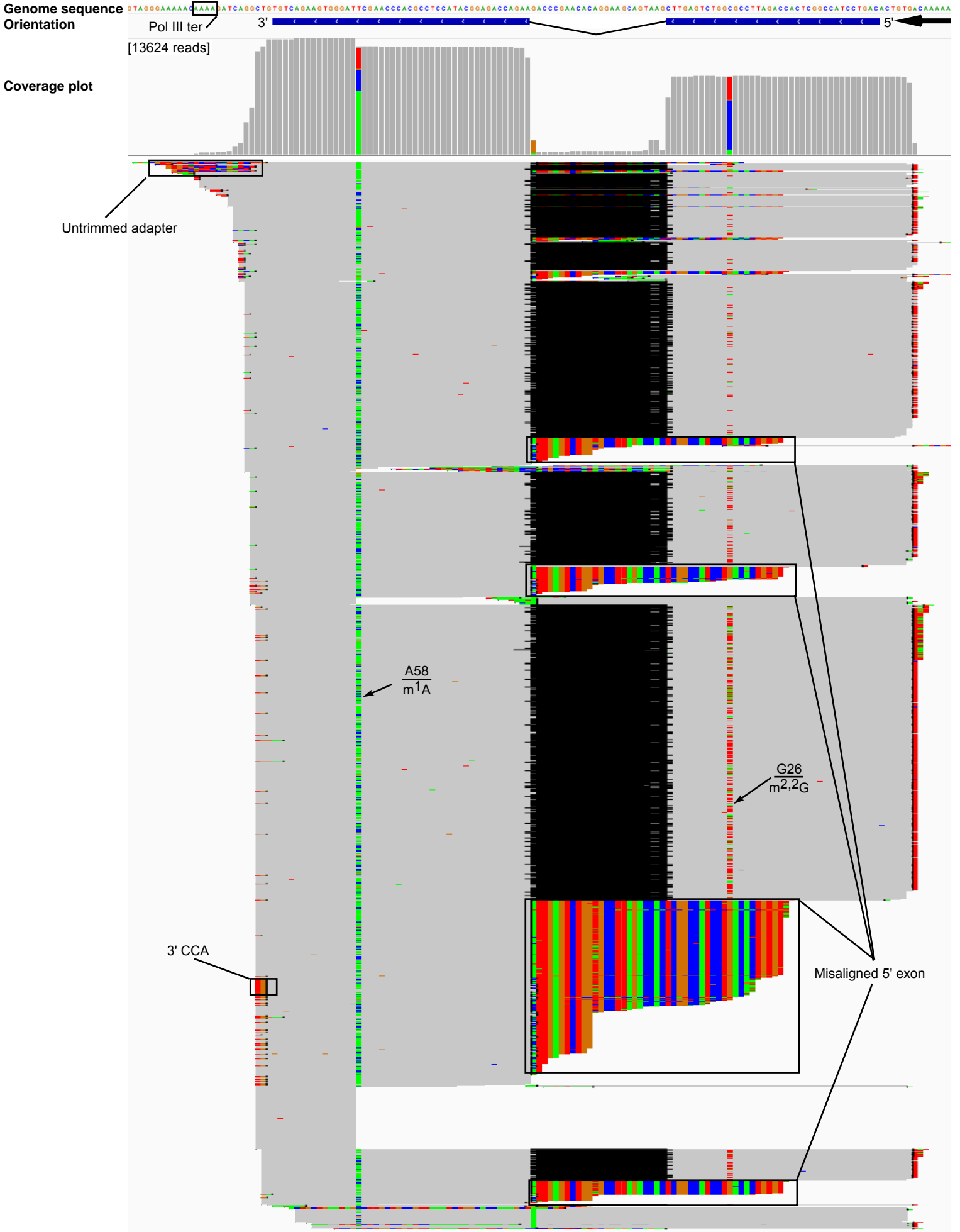
# chr11.tRNA3-ArgTCT ( vs gene )



# chr11.tRNA3-ArgTCT ( vs spliced tRNA)



chr6.tRNA140-LeuCAA ( vs gene )



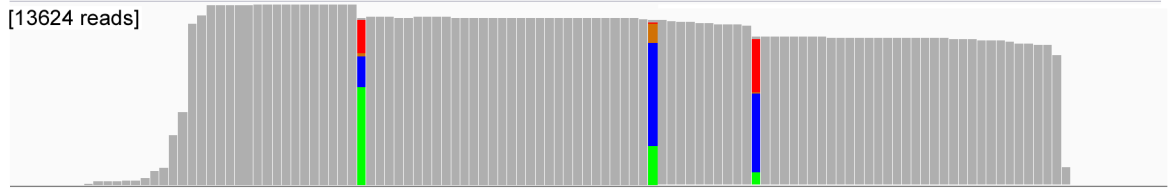


# chr6.tRNA140-LeuCAA ( vs spliced tRNA )

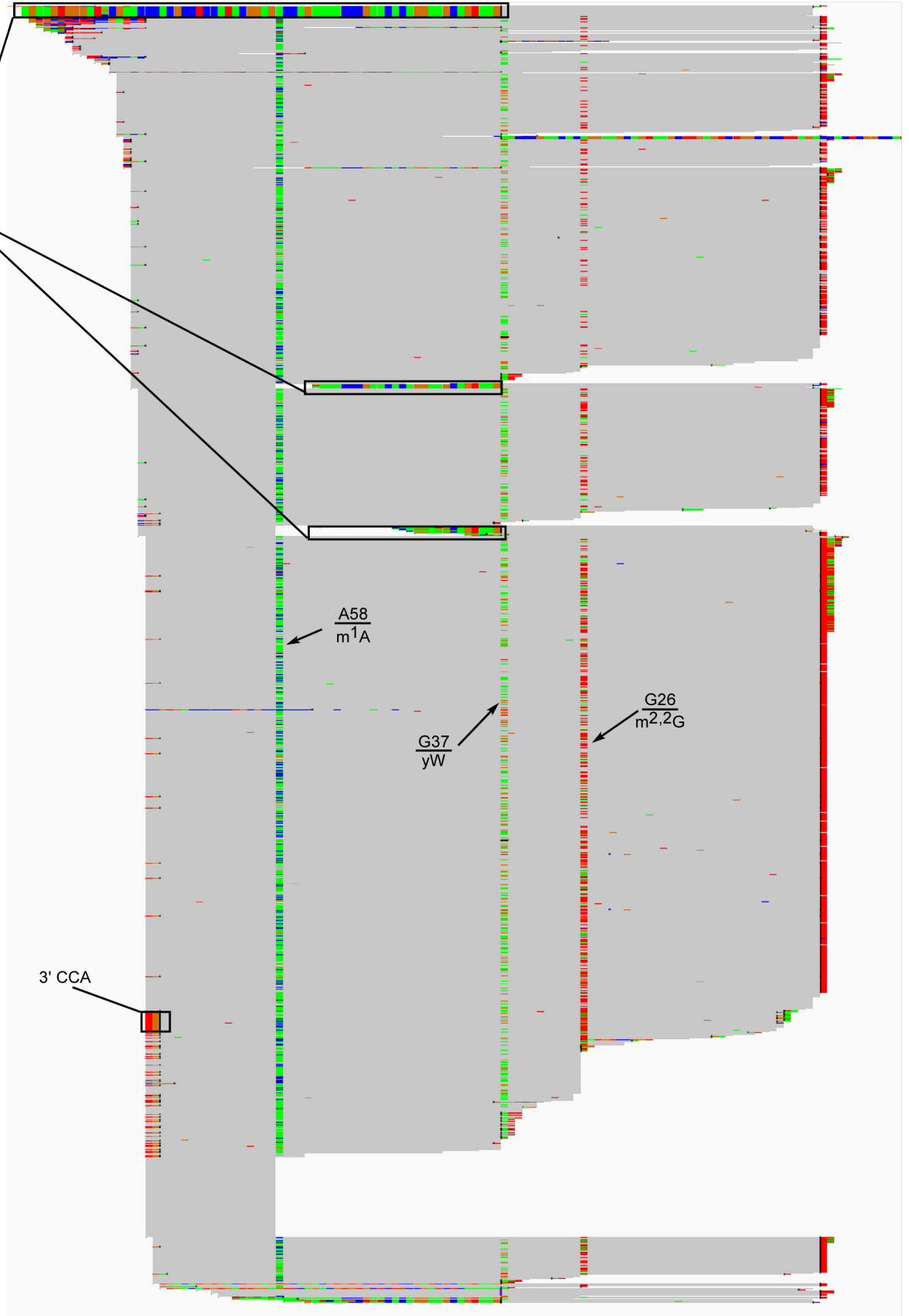
Genome sequence  
Orientation



Coverage plot



Unspliced pre-tRNA  
fragment



A58  
m<sup>1</sup>A

G37  
yW

G26  
m<sup>2,2</sup>G

3' CCA

# chr1.tRNA91-PseudoCCC

Genome sequence GGGTAAAGAGCCATAGTGAAGAGCCAAAATACCTCTTATCTTTGACCGGGAATTAAGAACCCGGGCTCCTCCGCGTGGGAGCGGAGAAATTCACCACTGAACCCACCAAGTGCCTCTCTAGCAACCCCTGGGAGA

Orientation

Pol III ter

3'

5'

[2539 reads]

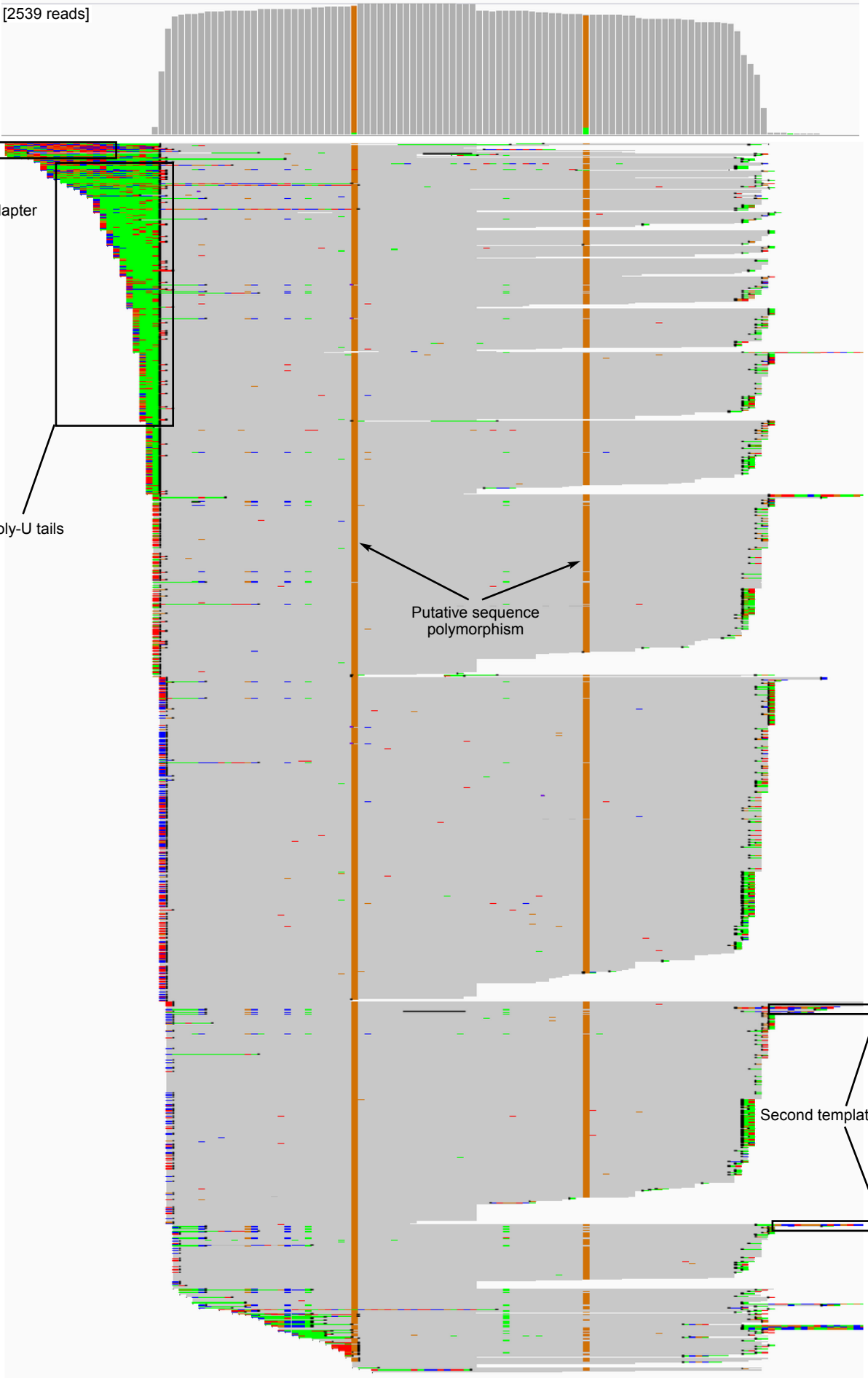
Coverage plot

Untrimmed adapter

Poly-U tails

Putative sequence polymorphism

Second template switch



# SNAR.A-10

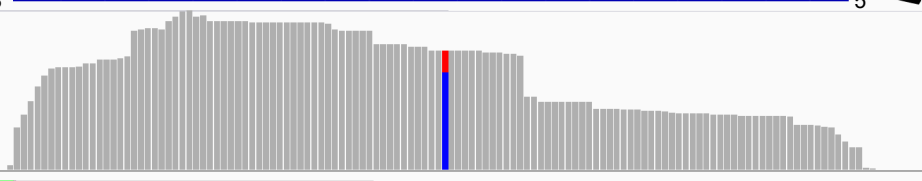
Pol III ter

Genome sequence  
Orientation



[494 reads]

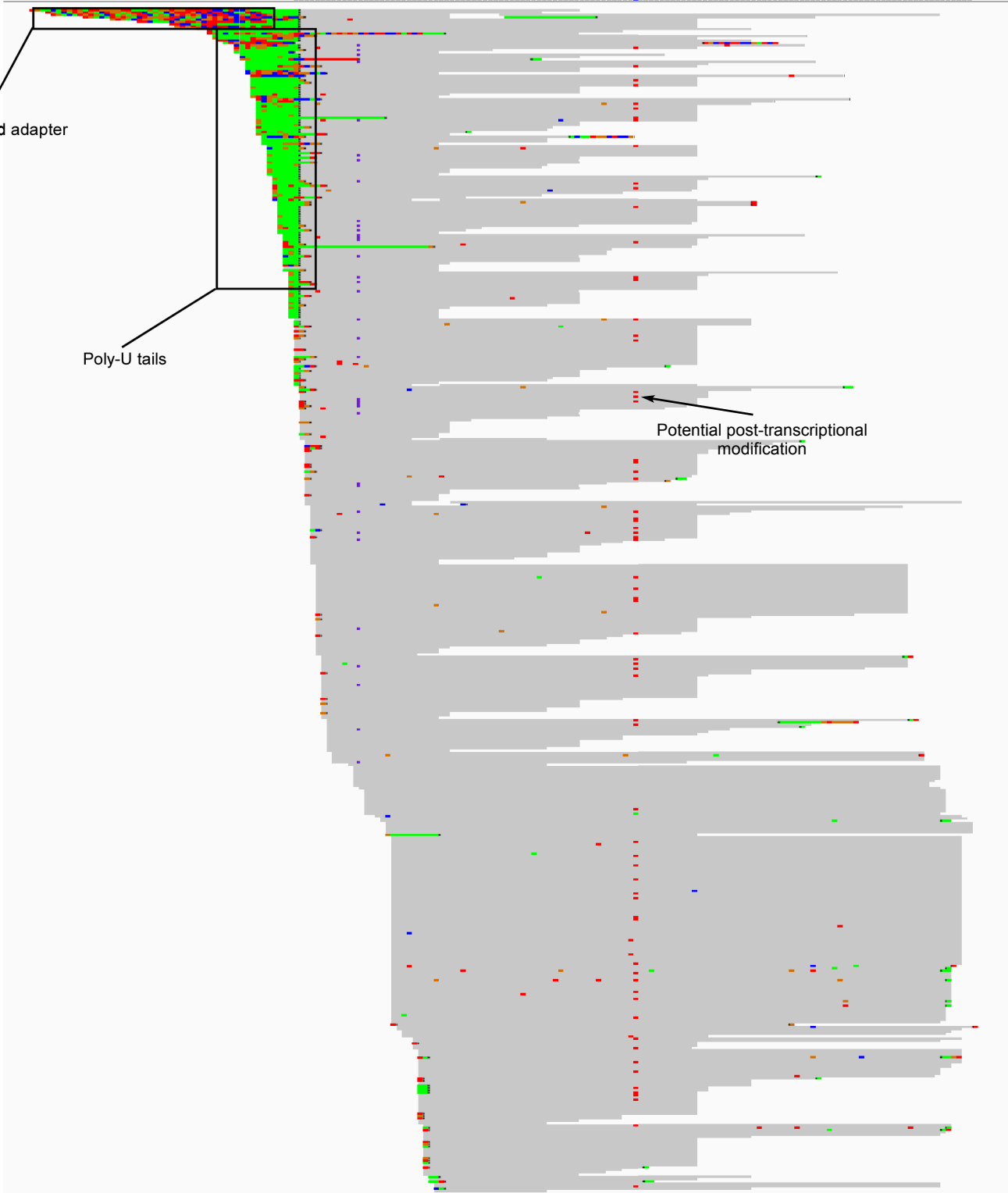
Coverage plot



Untrimmed adapter

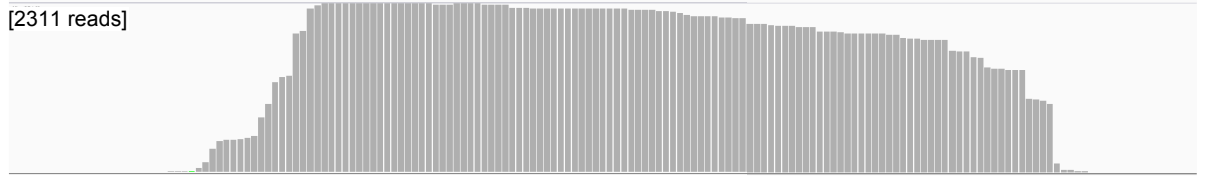
Poly-U tails

Potential post-transcriptional modification



# 5S1.rRNA

Genome sequence  
Orientation



Coverage plot

