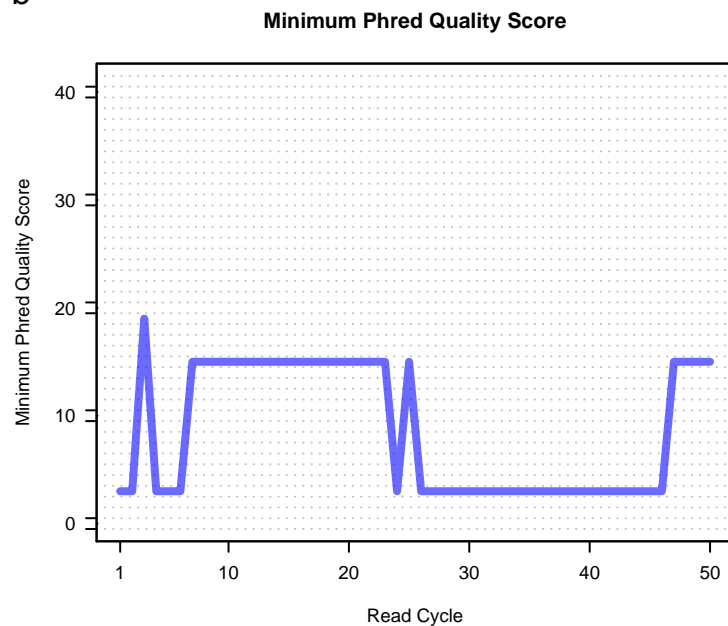


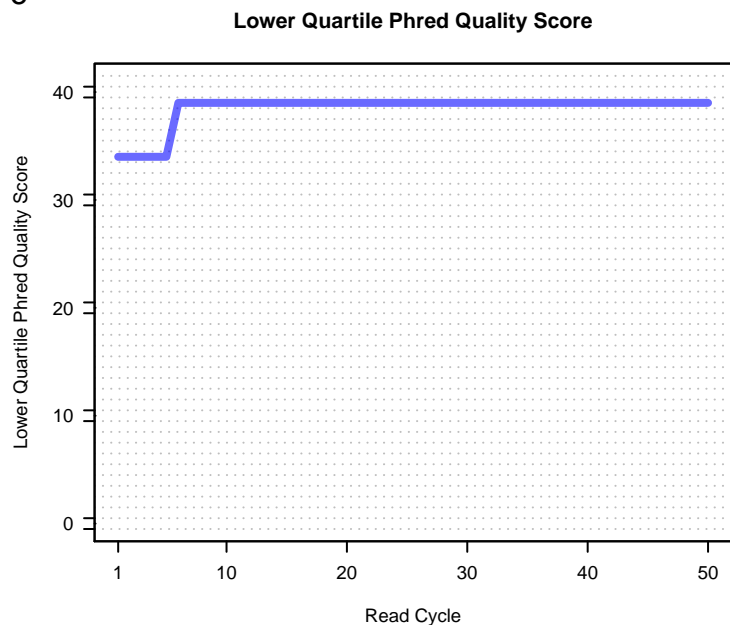
a

# All Quality Control Summary Plots

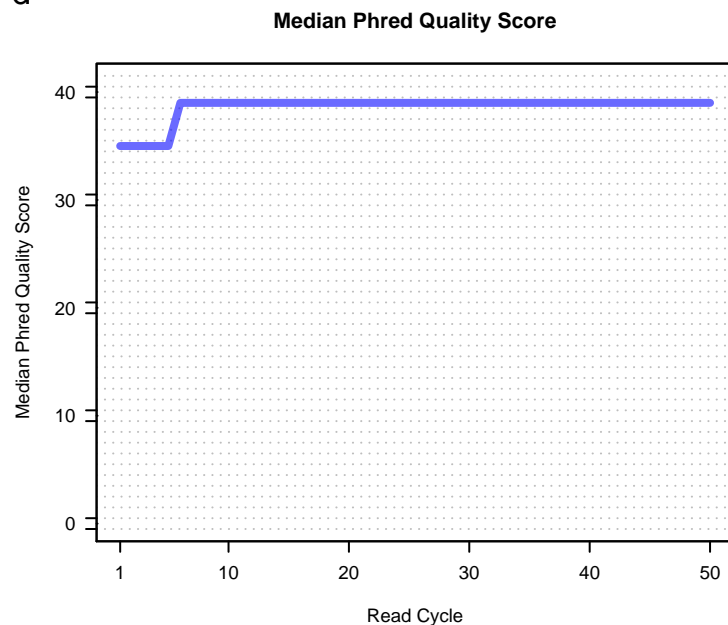
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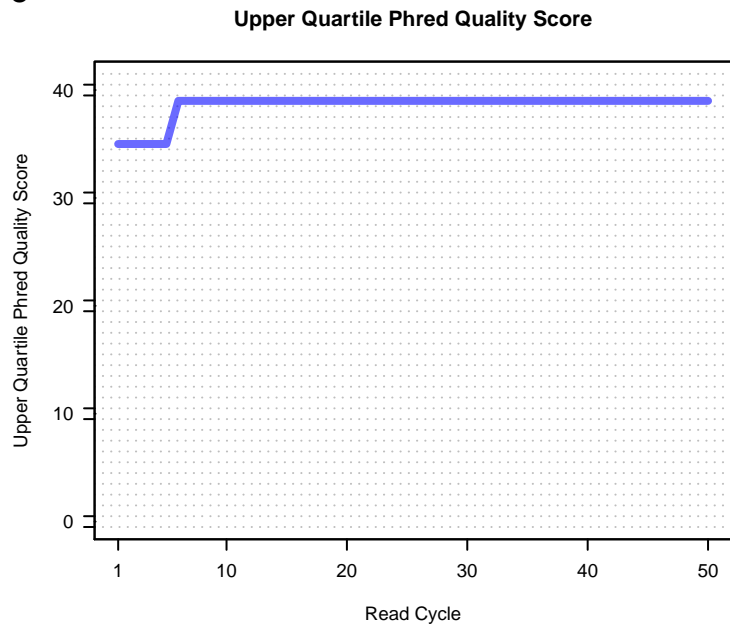
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d

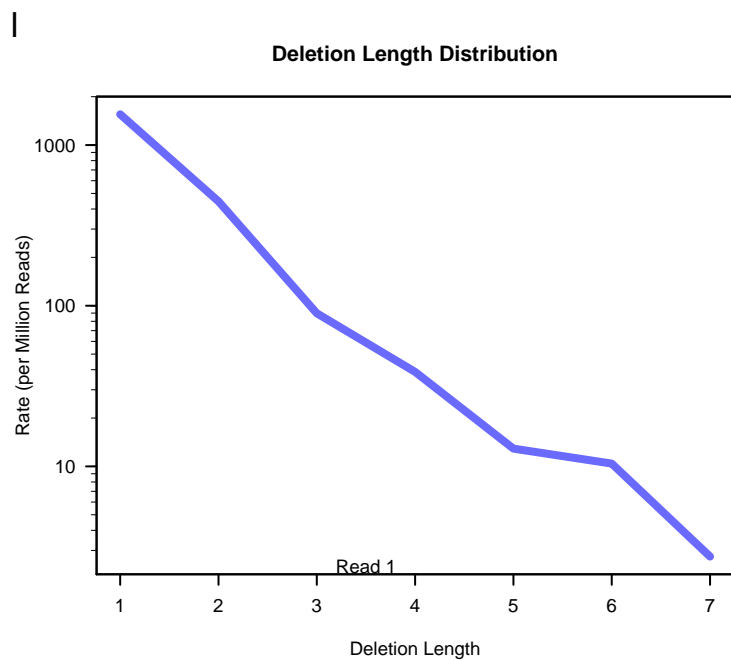
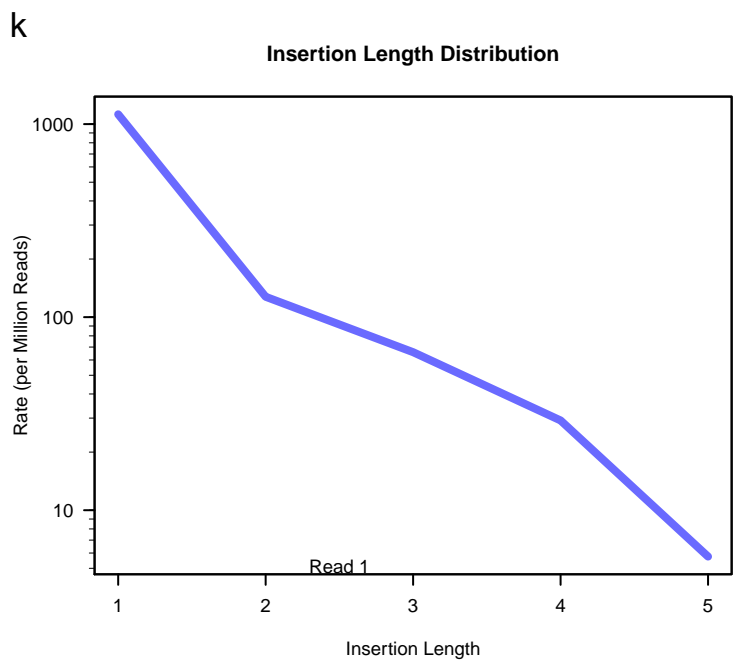
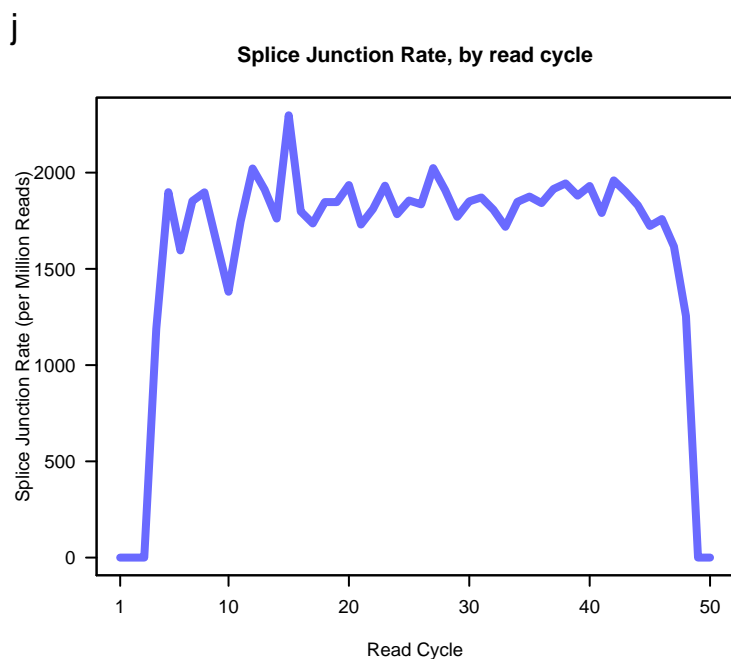
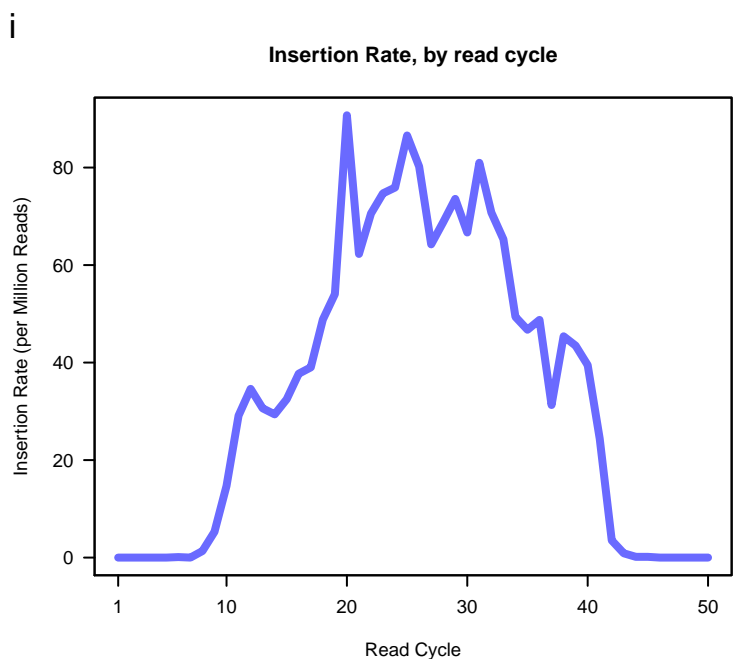
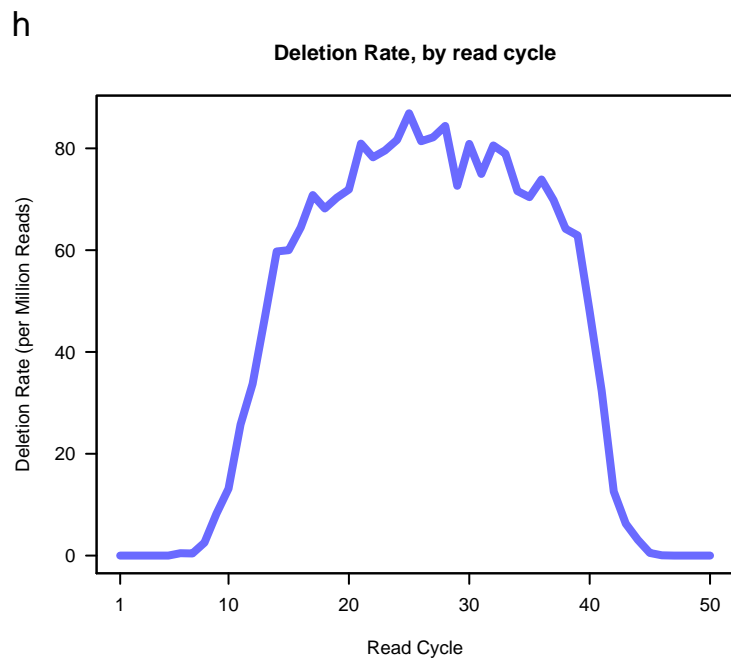
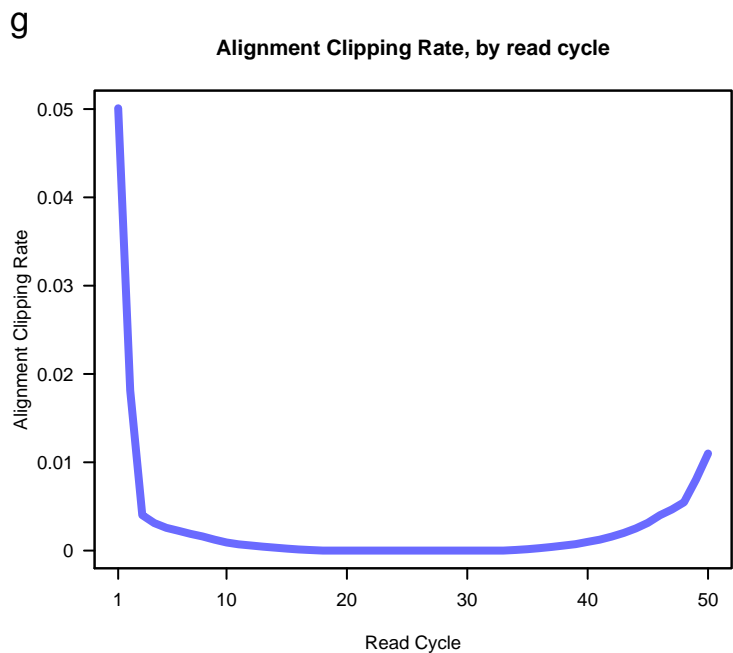


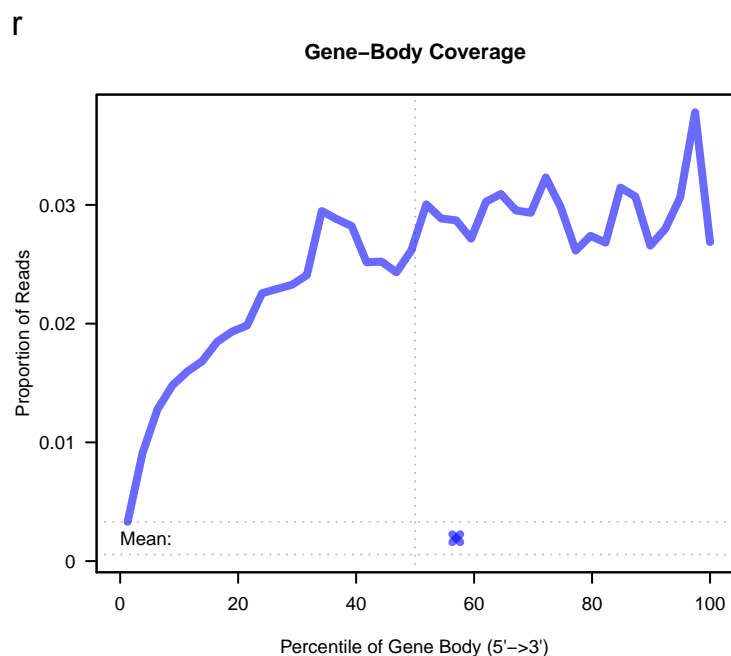
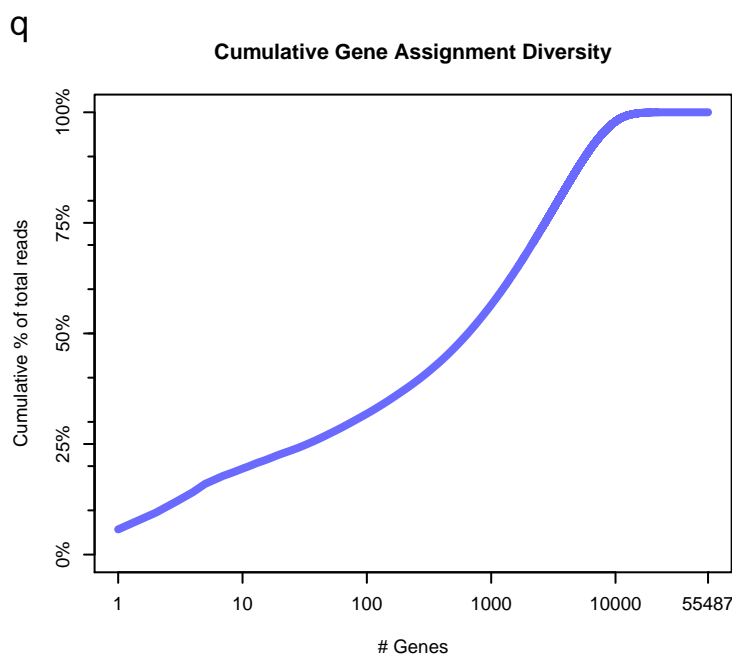
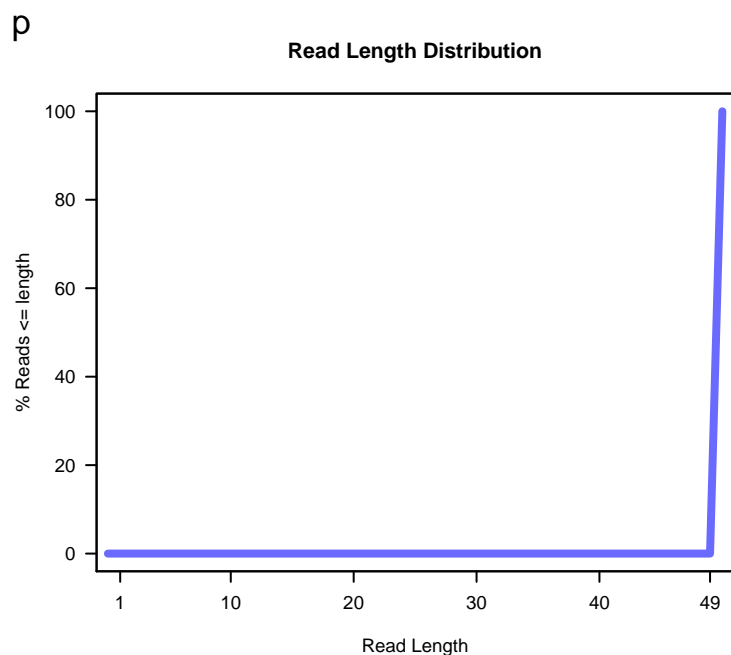
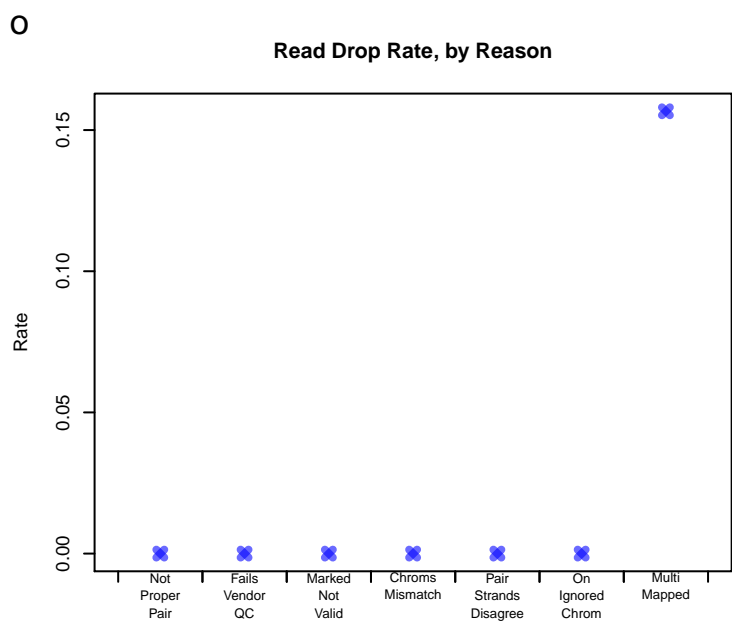
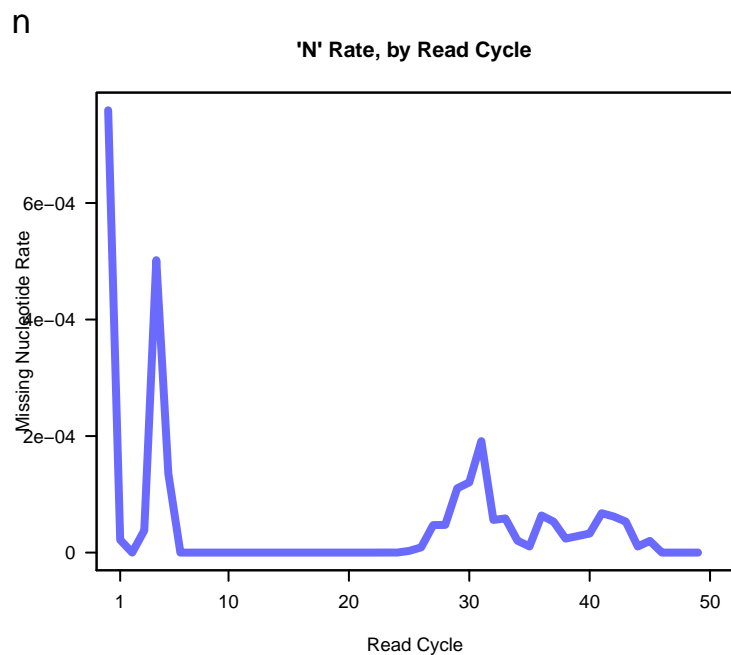
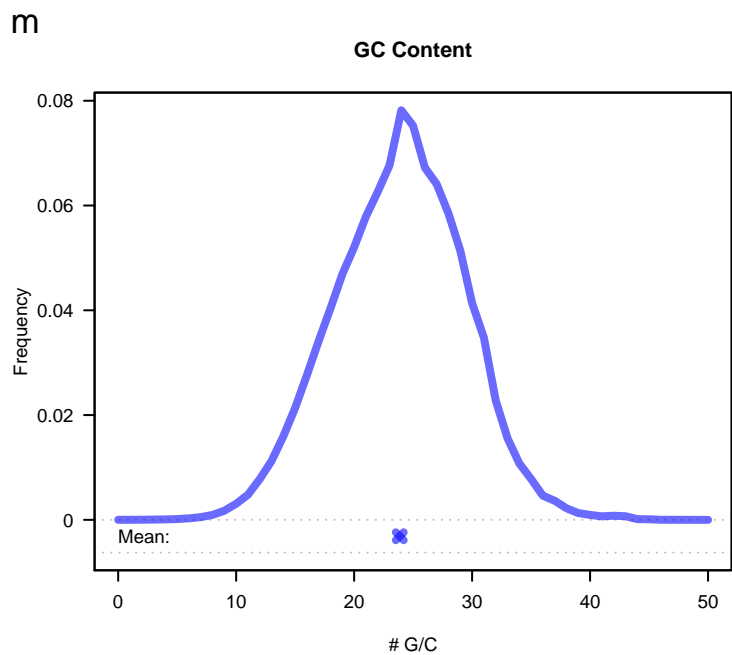
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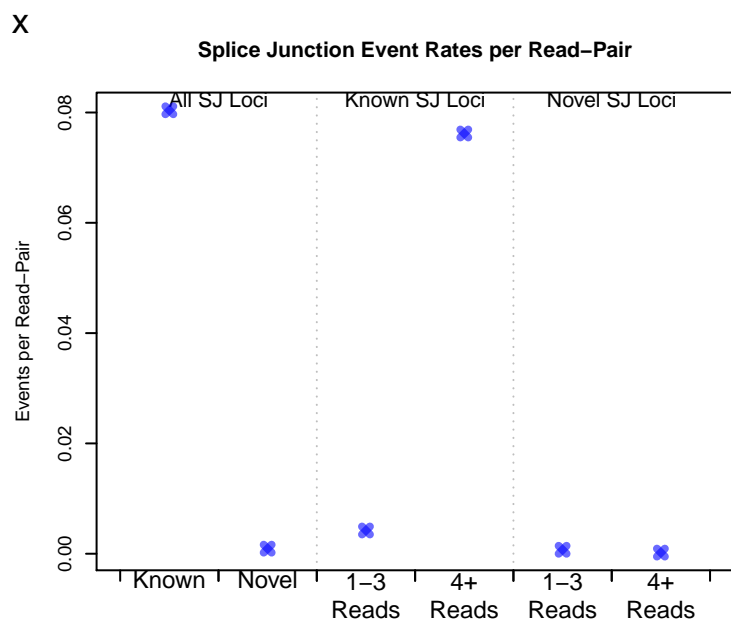
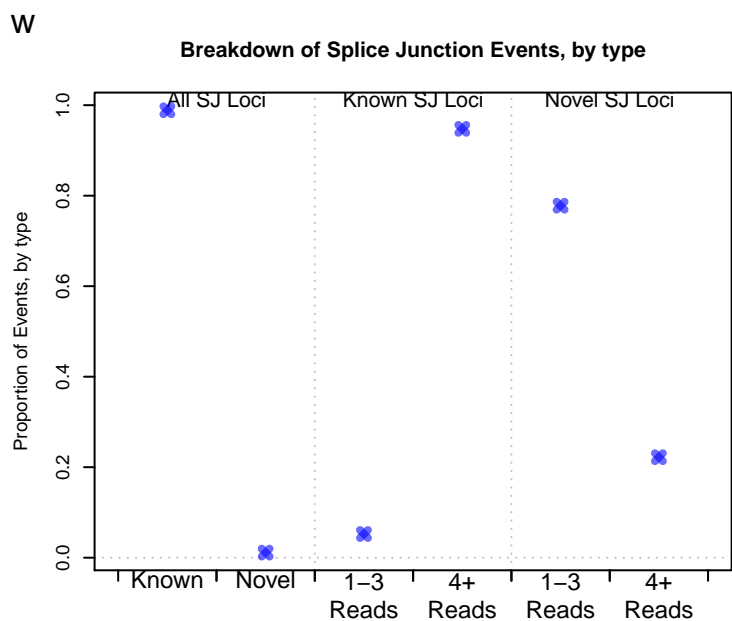
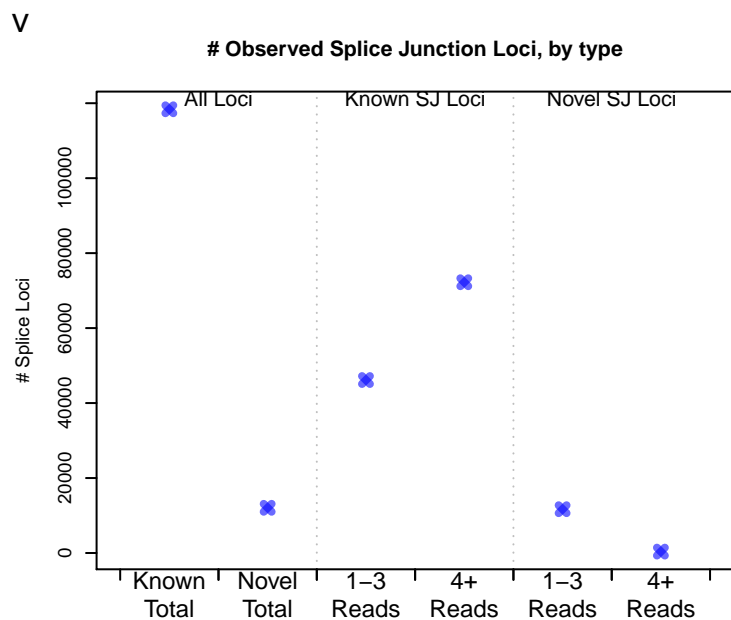
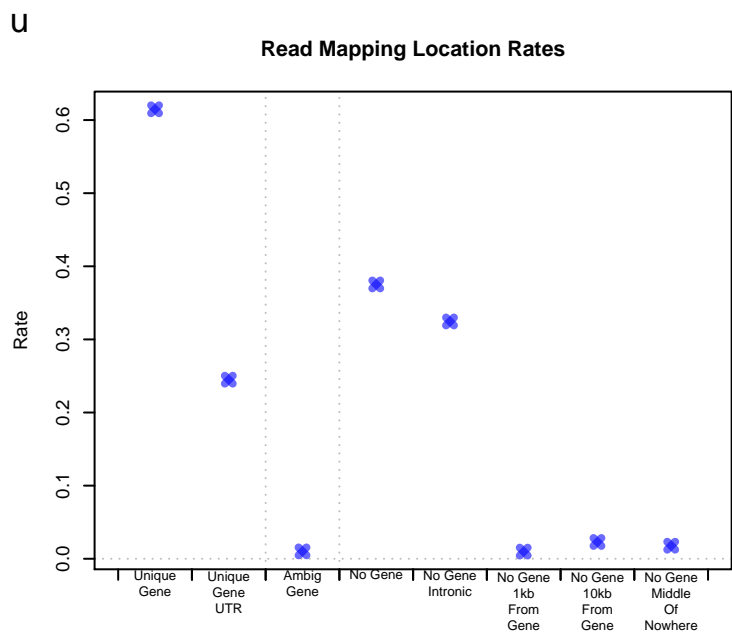
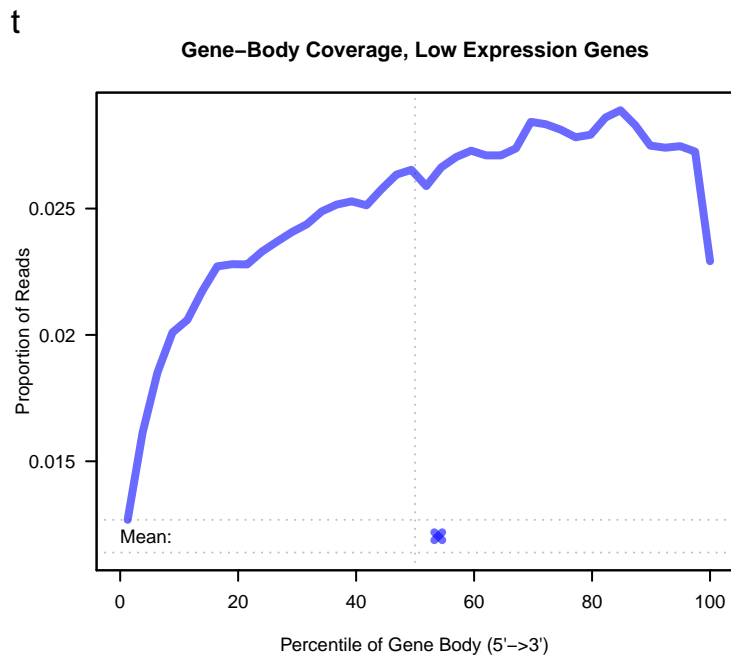
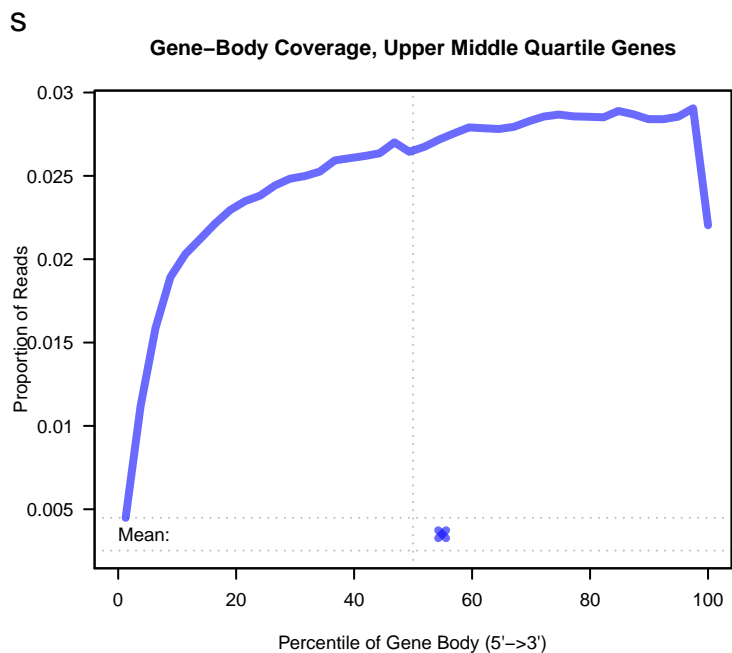


f



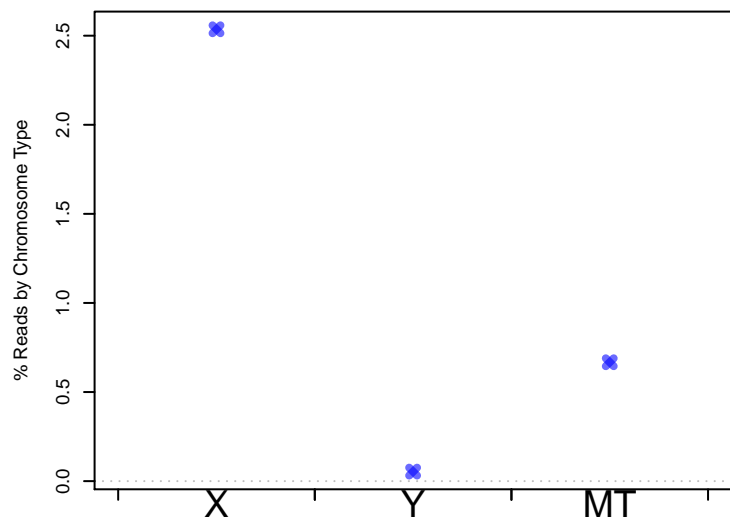






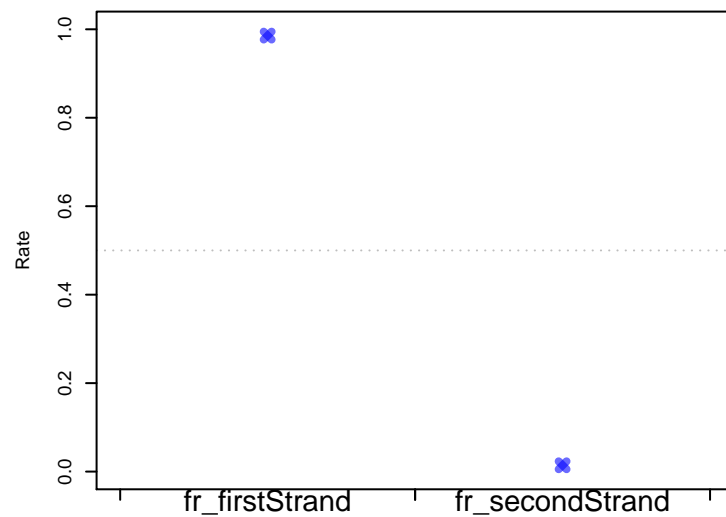
y

Chromosome Distribution (Excluding Autosomes)



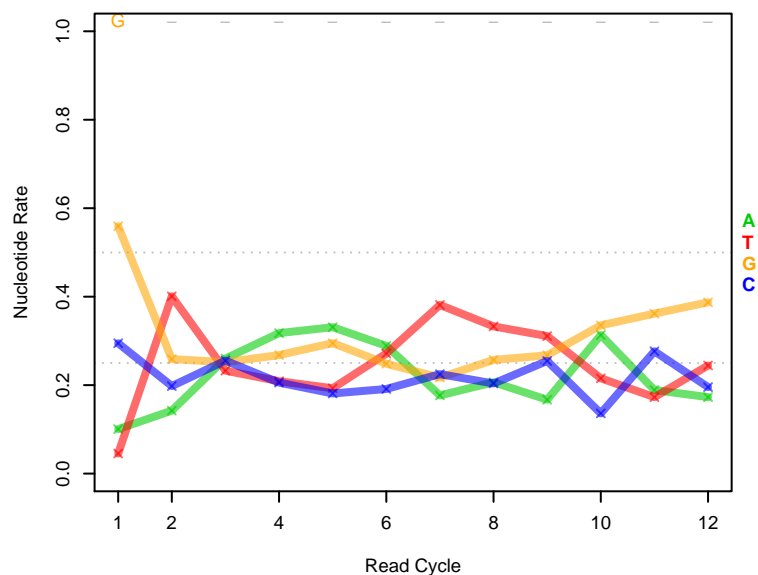
Z

Strandedness Test



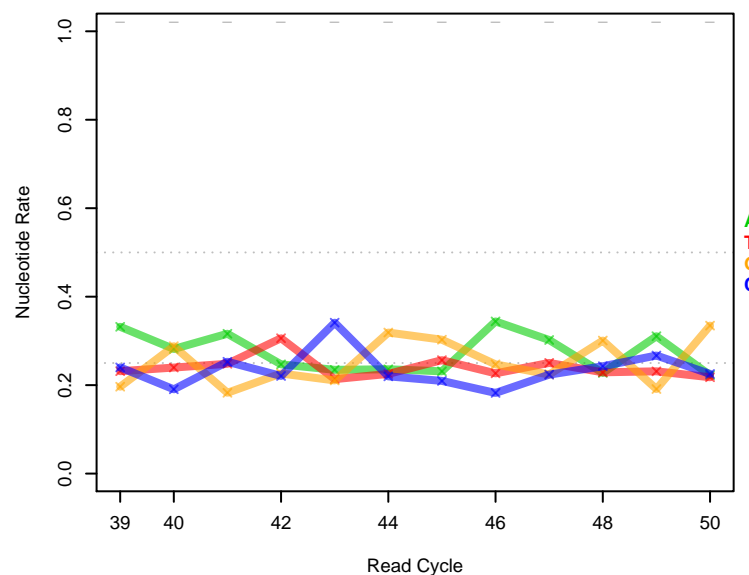
aa

Nucleotide Rate by Cycle, Leading Clipped bases (12)



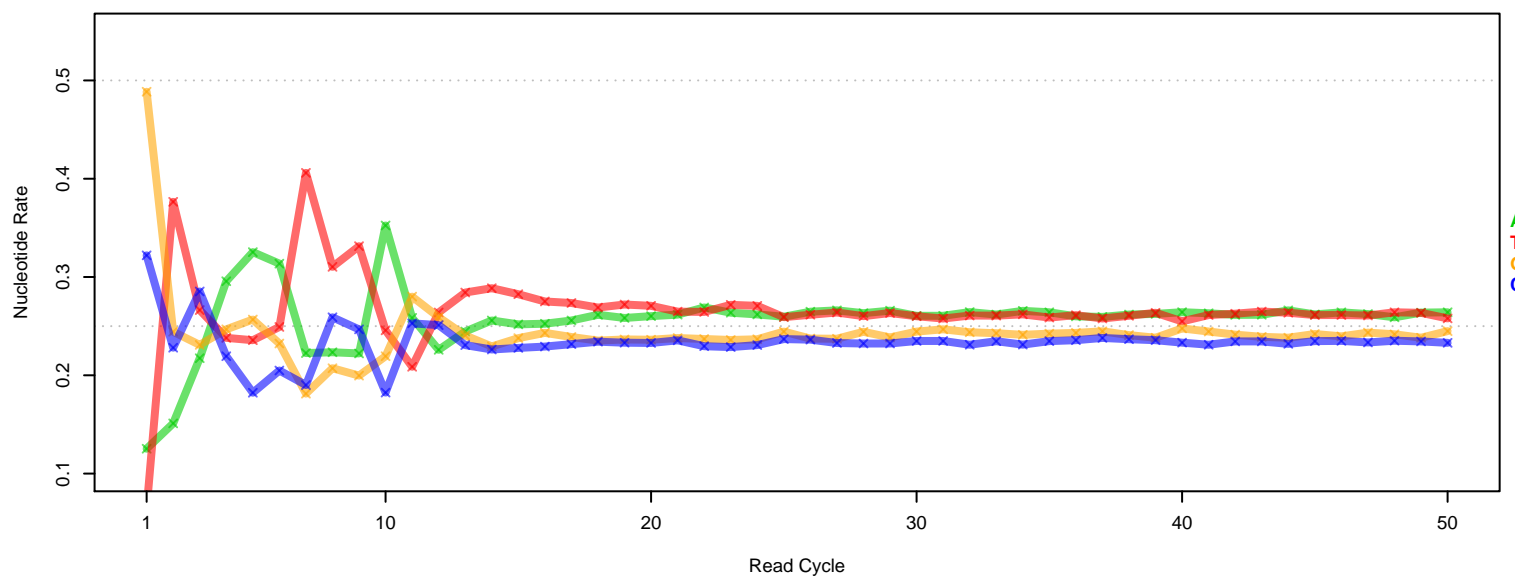
ab

Nucleotide Rate by Cycle, Trailing Clipped bases (12)



ac

Raw Nucleotide Rate by Cycle



Nucleotide Rate by Cycle, Aligned bases only

