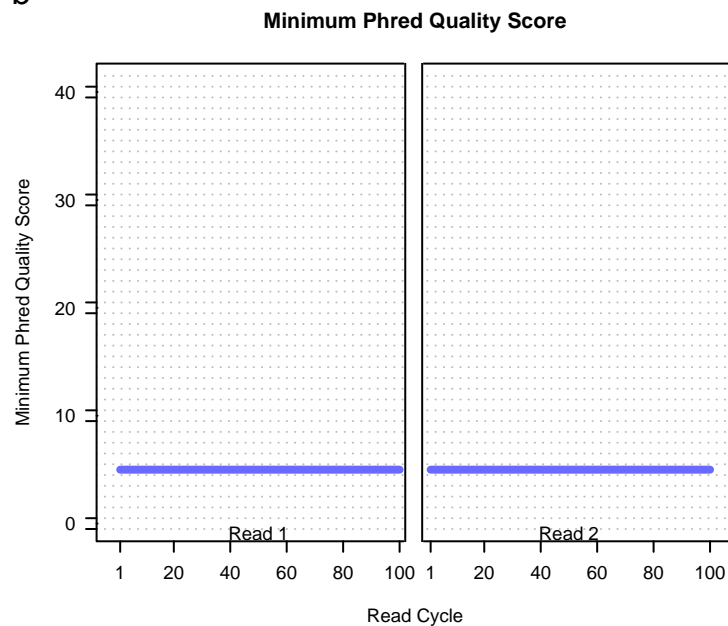


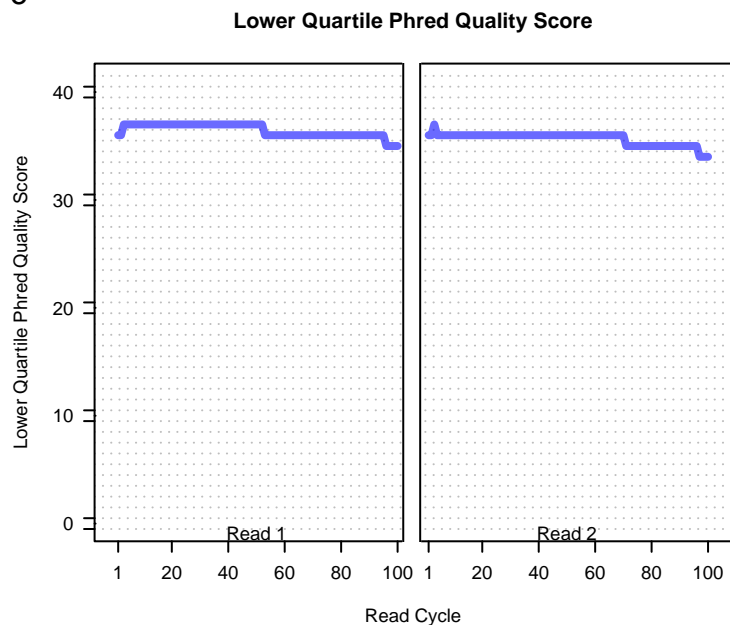
a

# All Quality Control Summary Plots

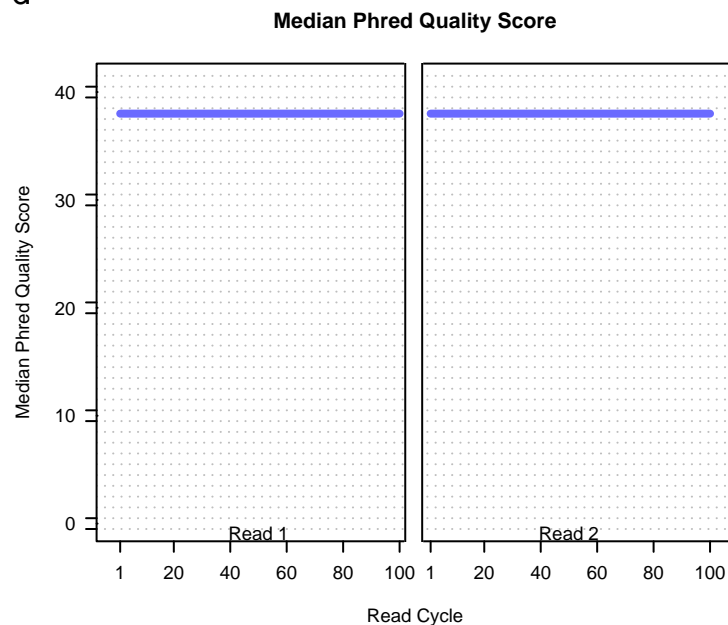
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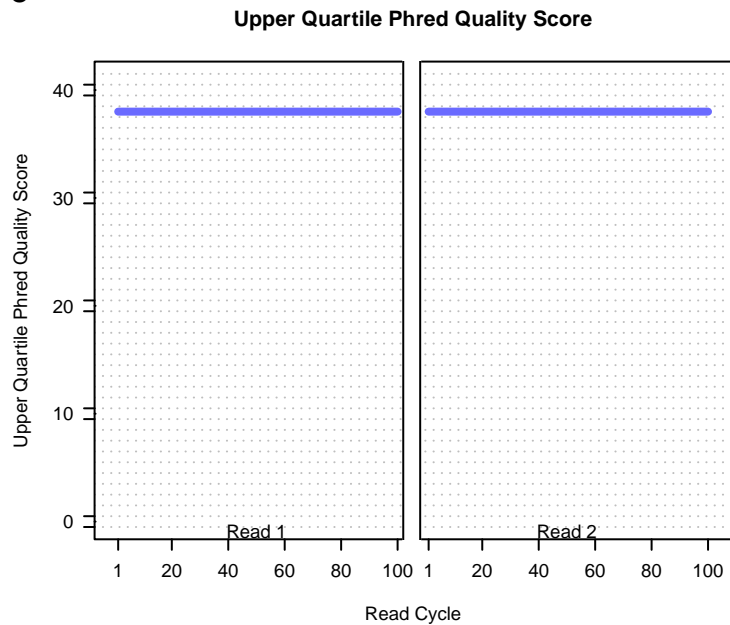
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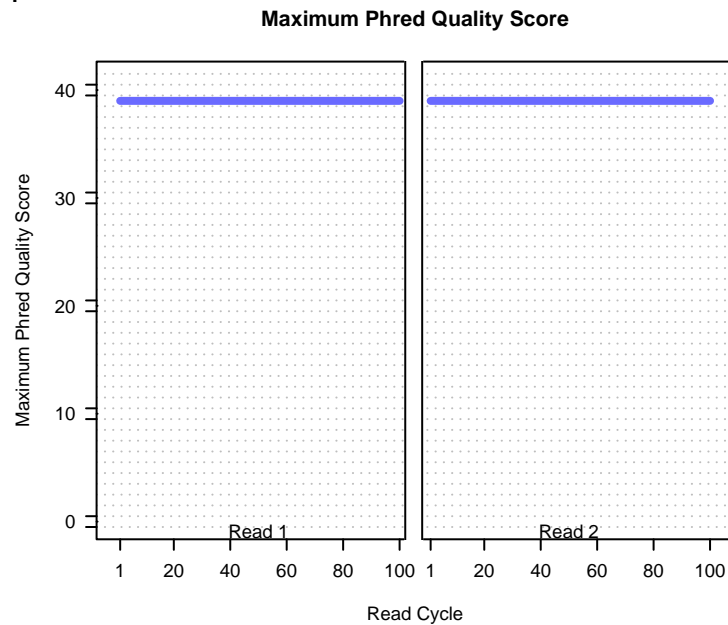
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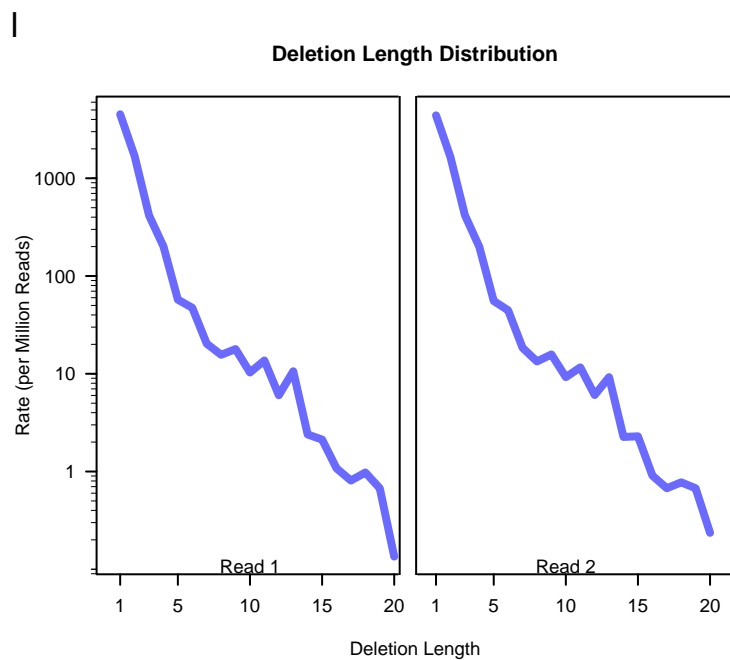
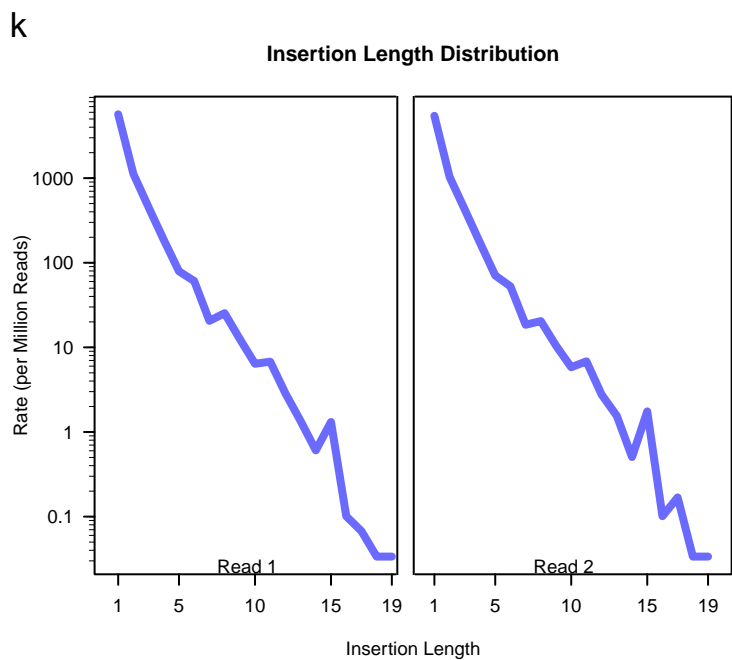
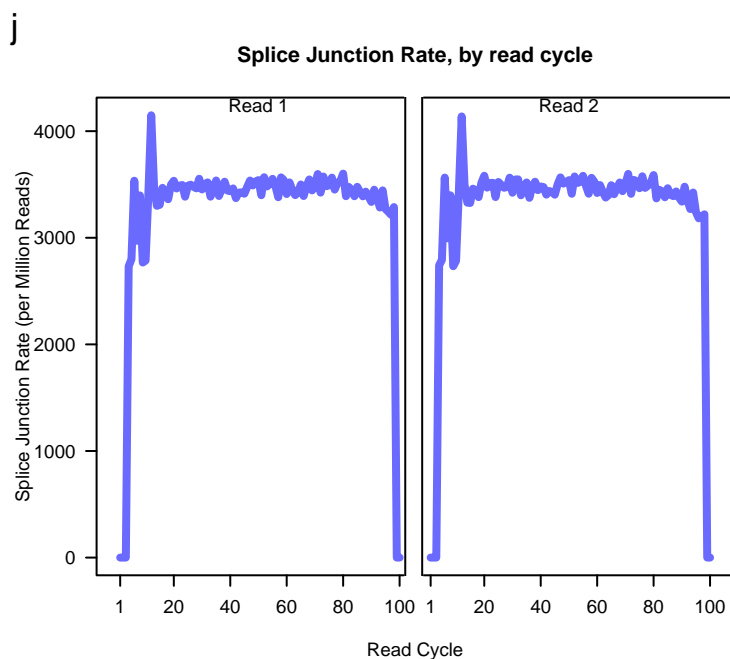
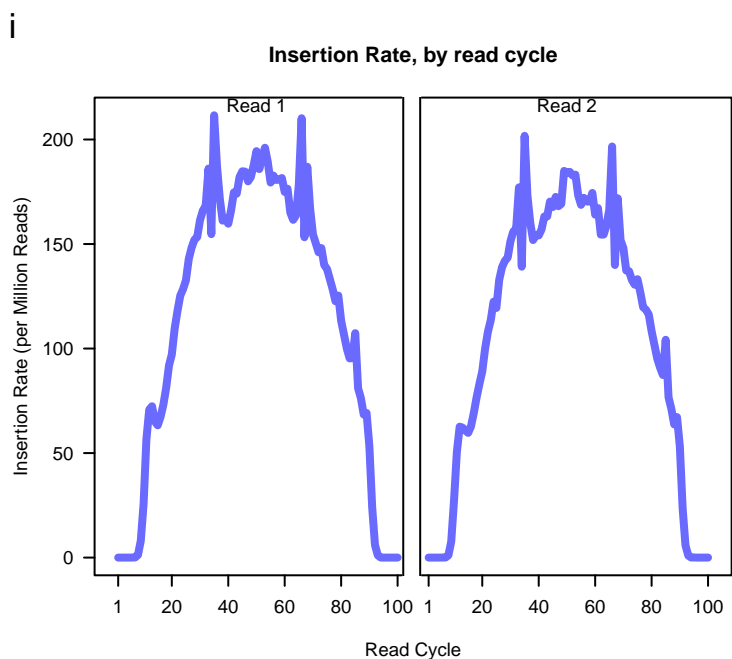
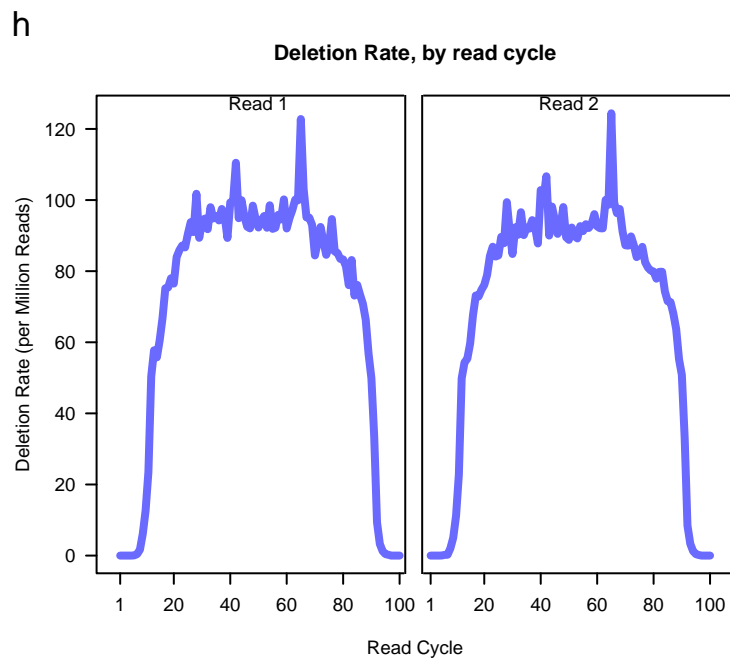
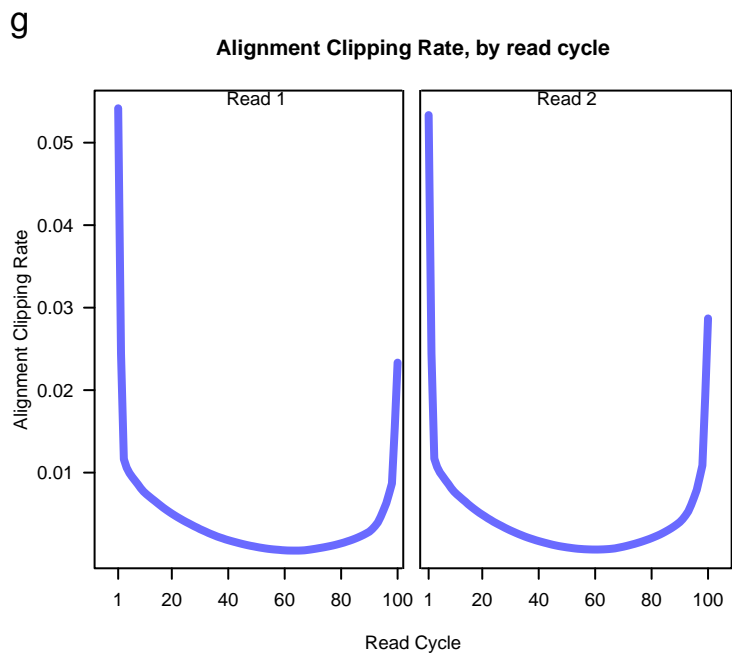


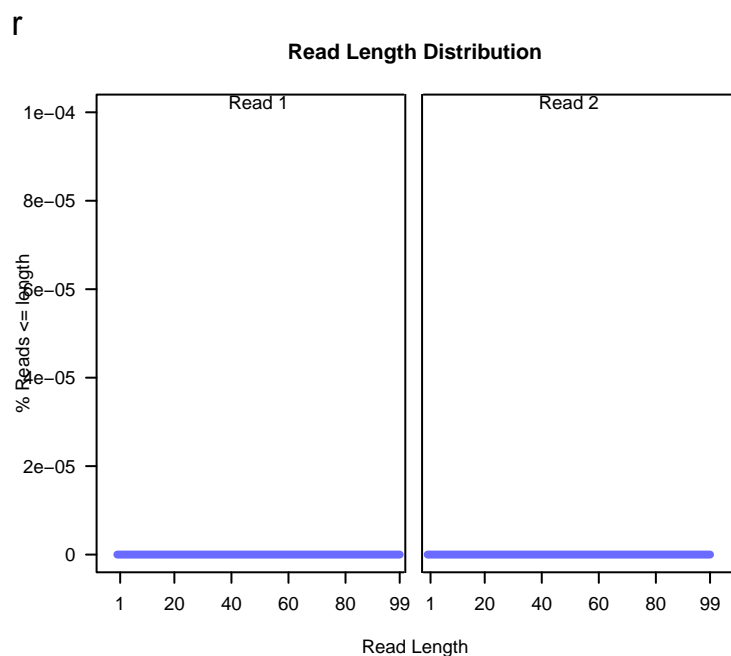
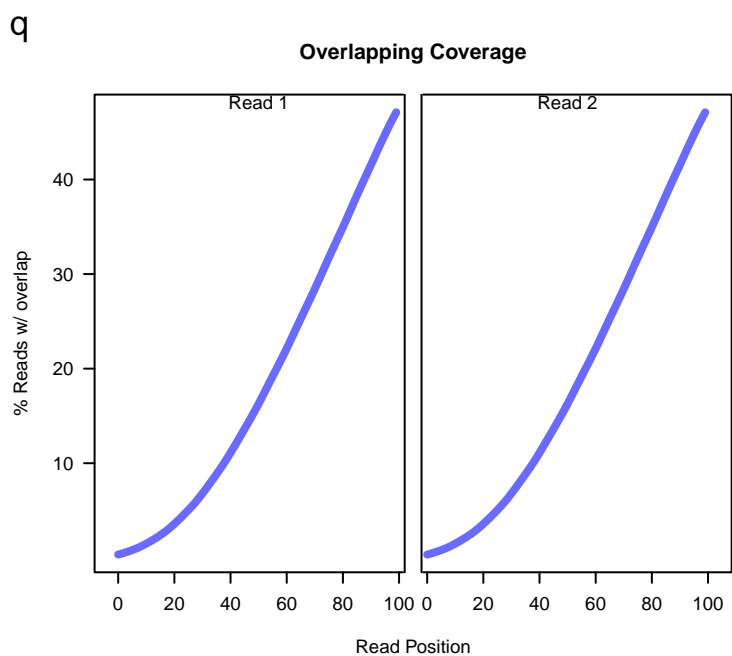
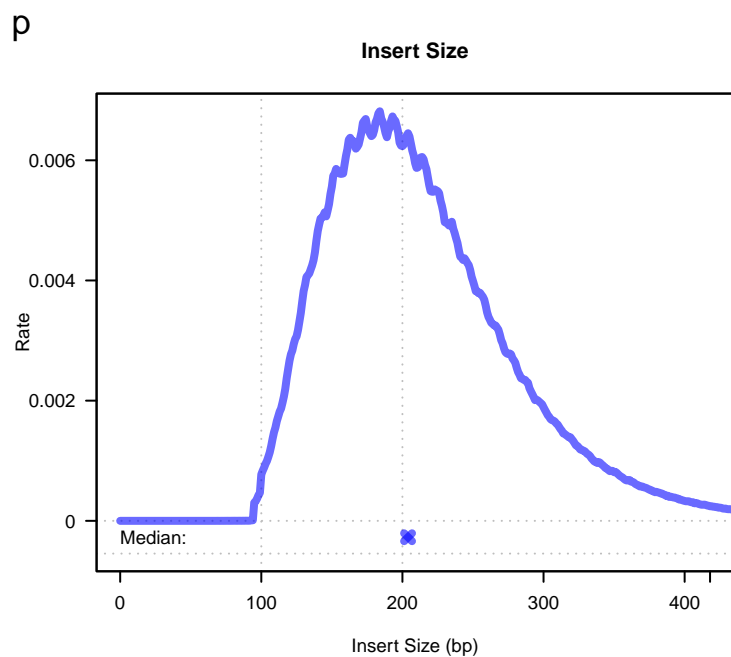
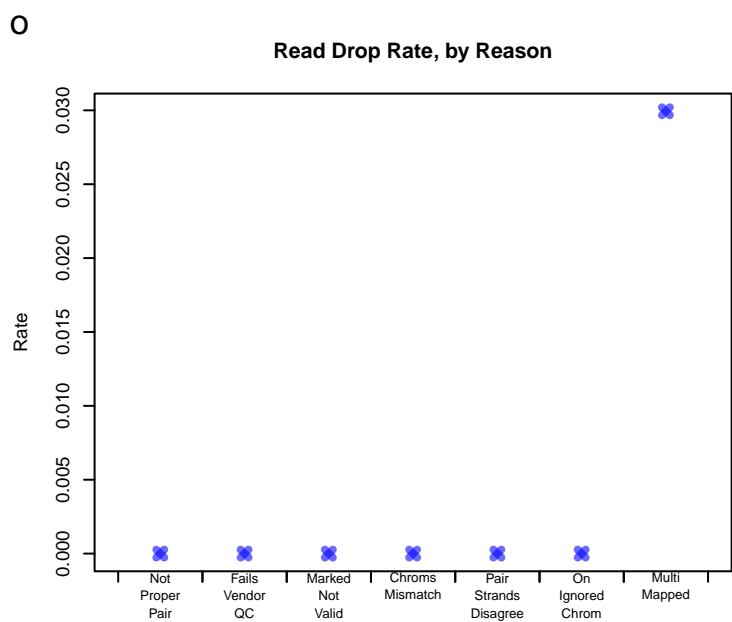
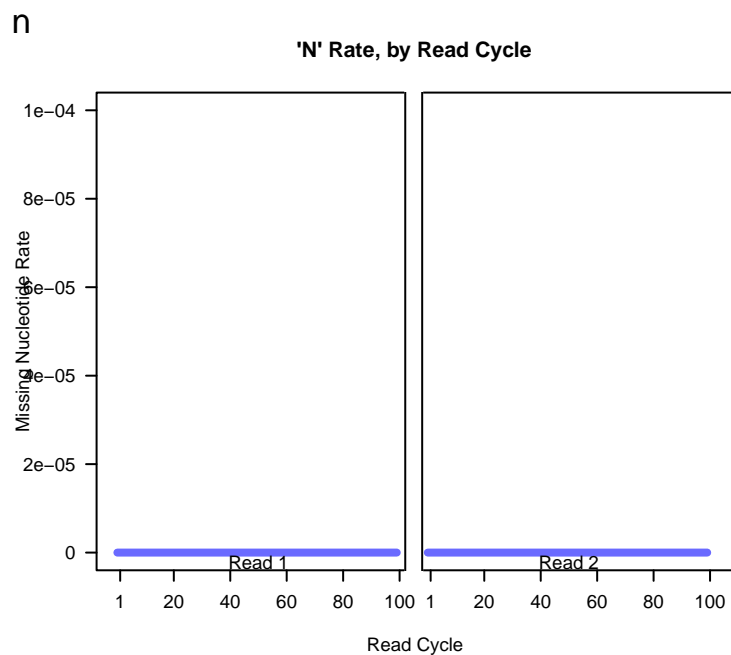
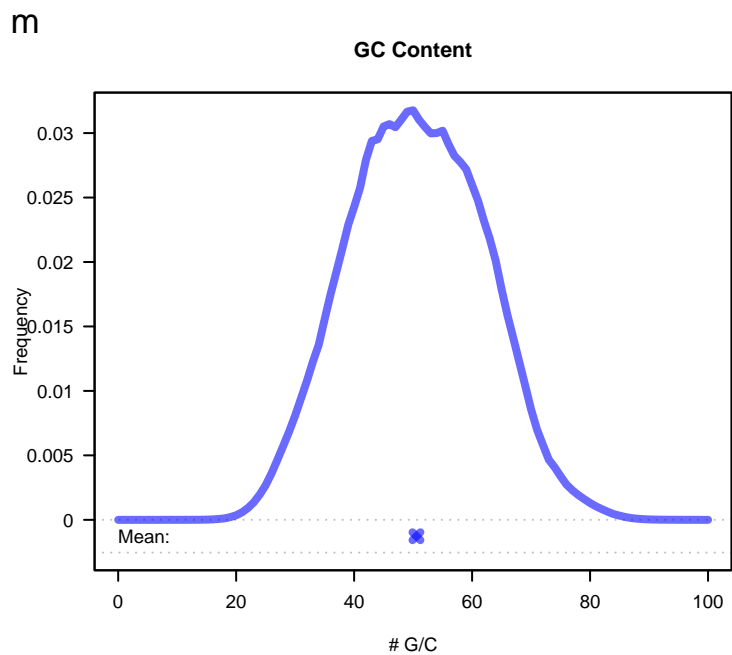
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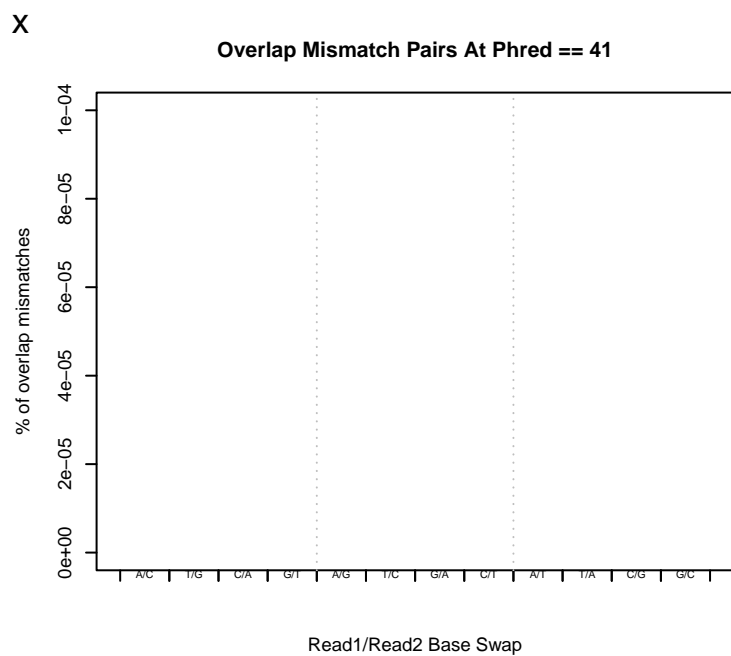
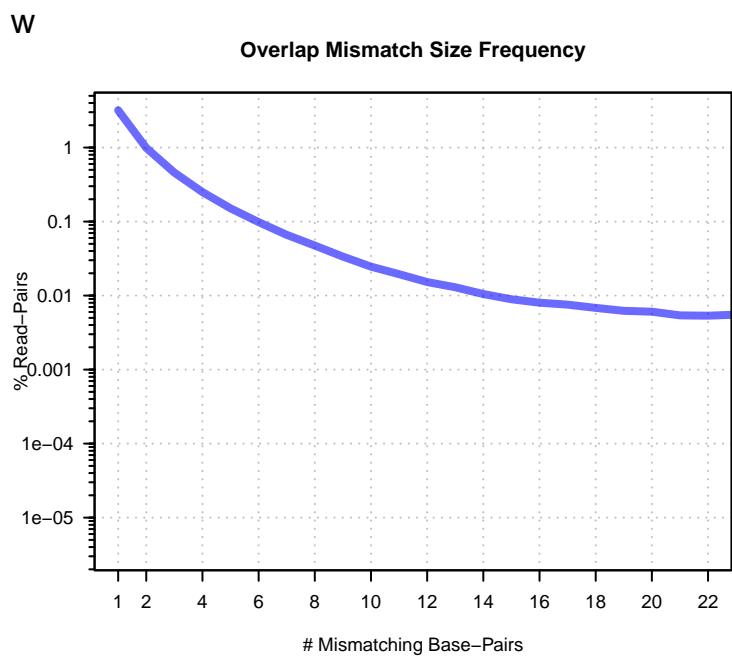
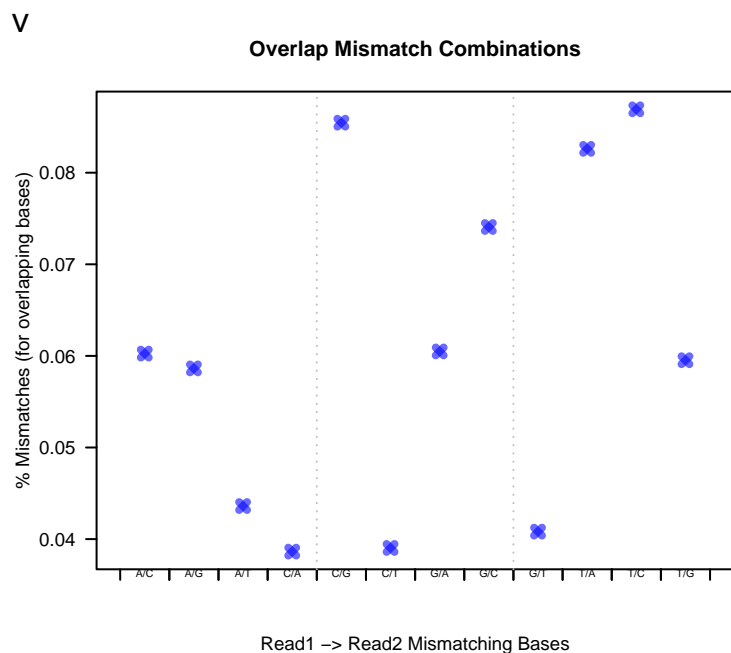
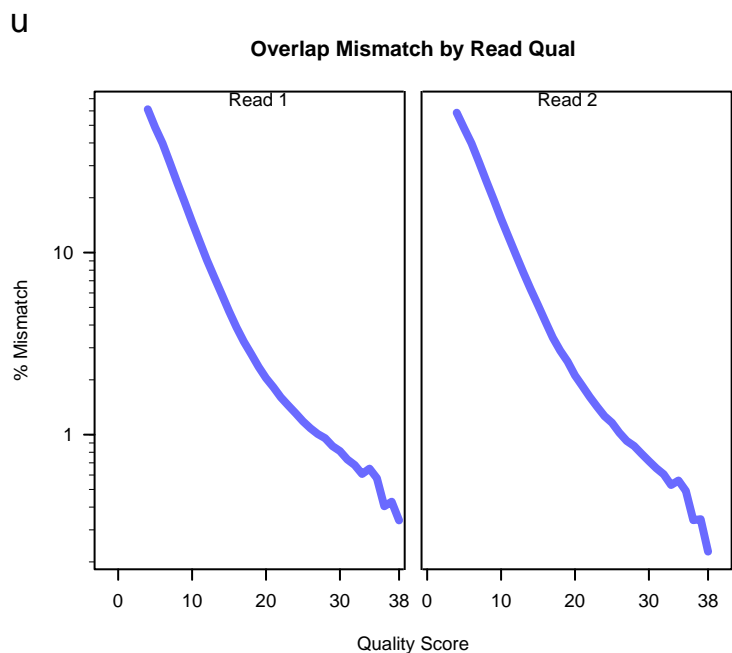
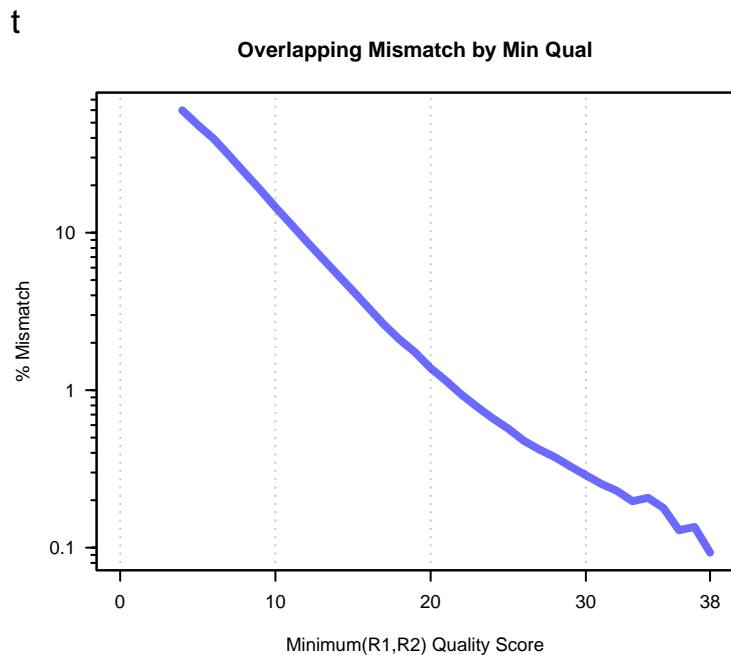
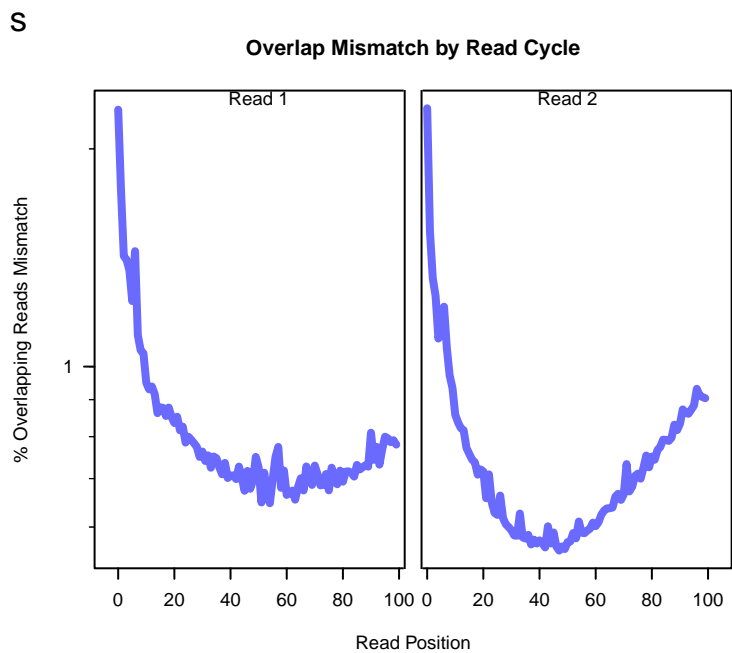


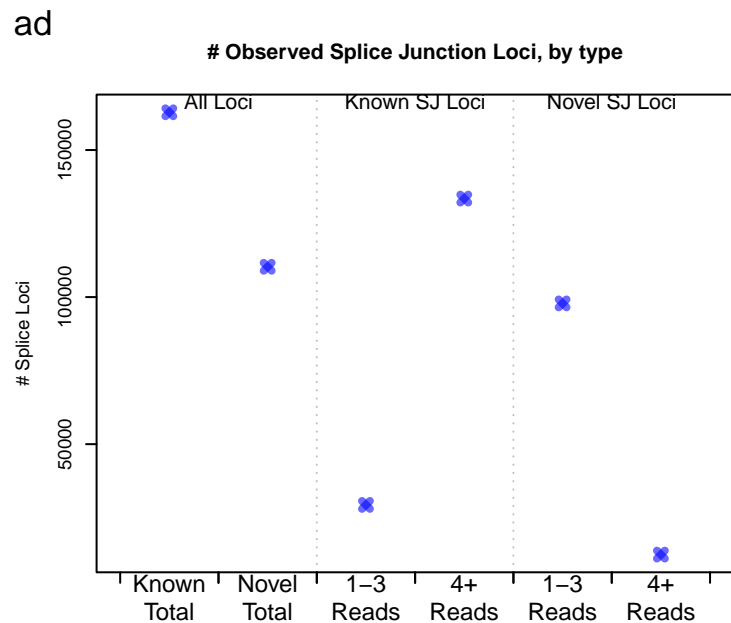
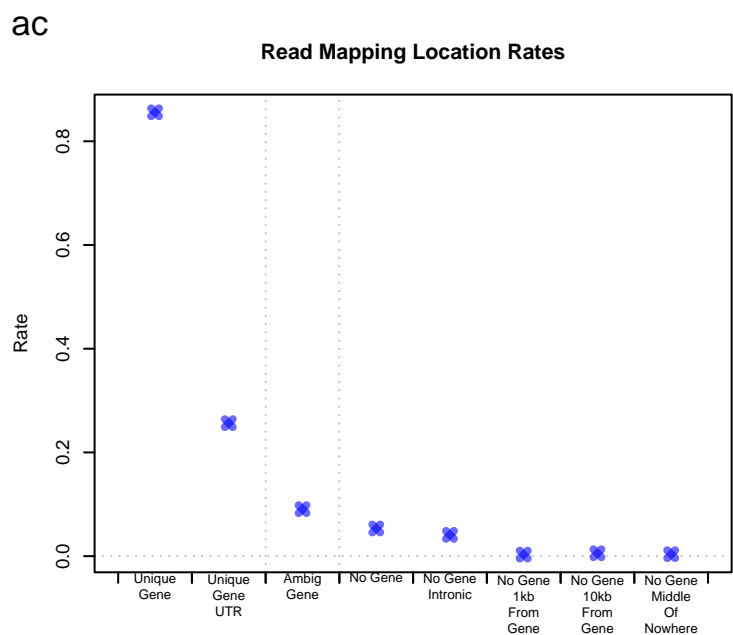
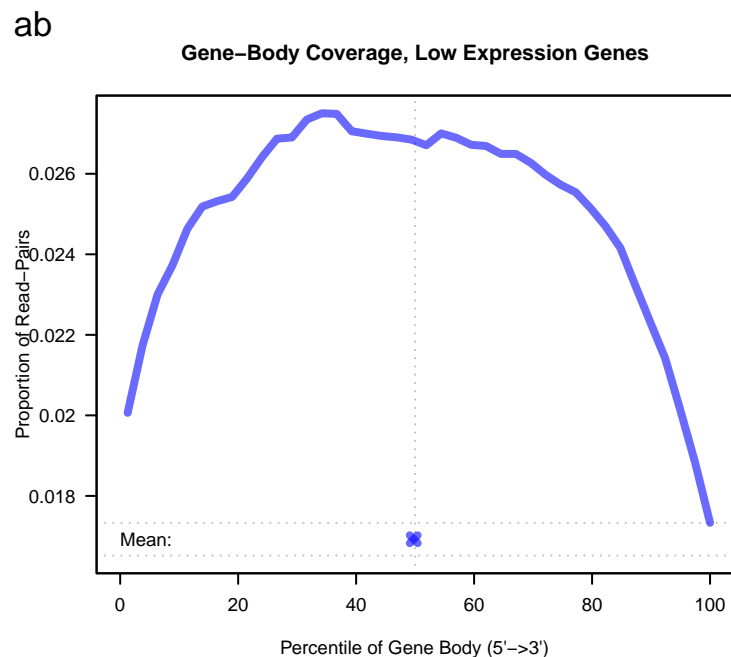
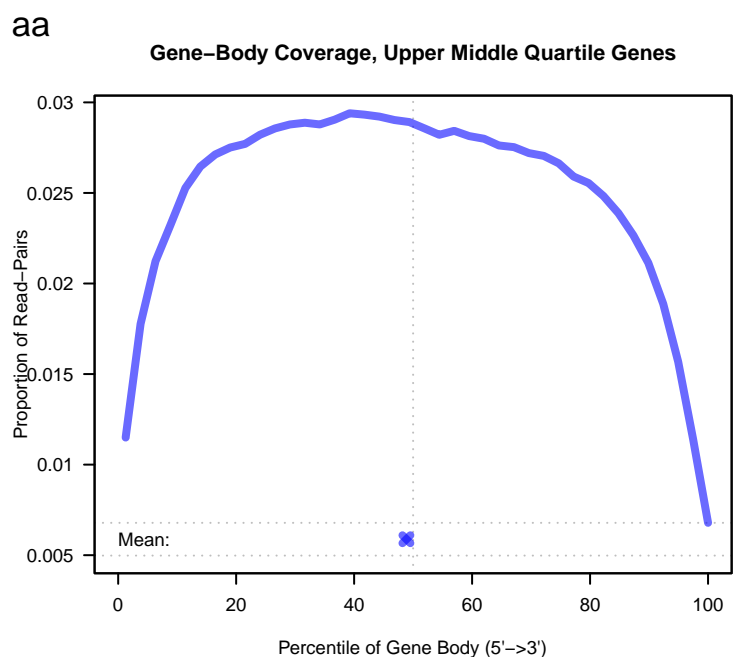
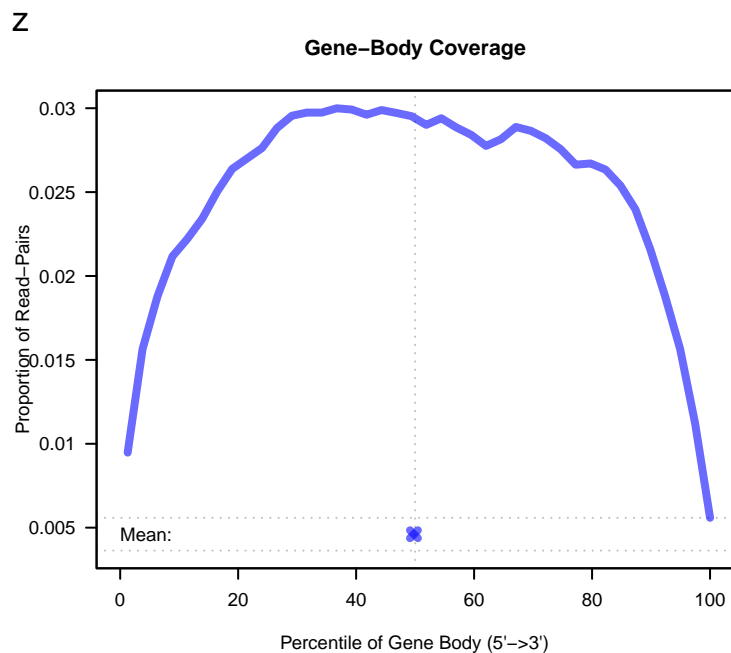
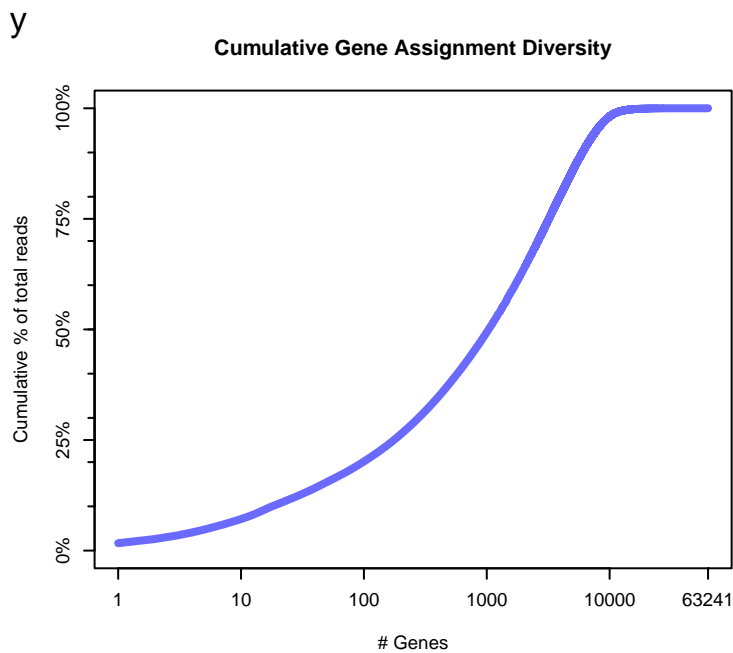
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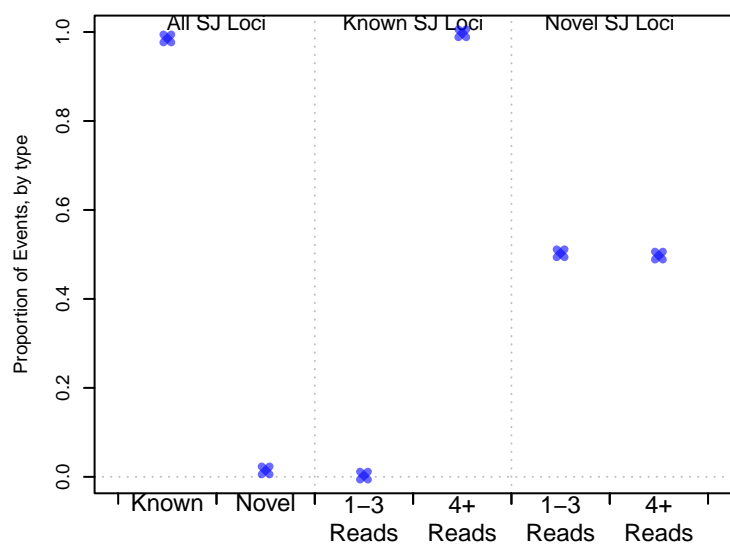






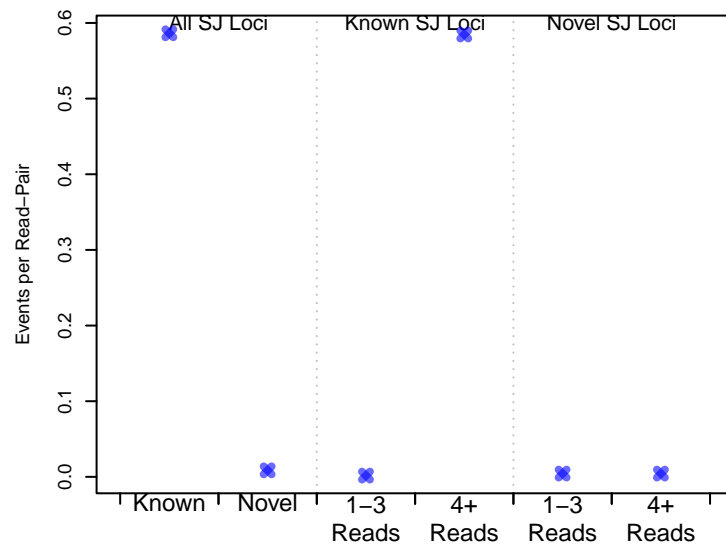
ae

Breakdown of Splice Junction Events, by type



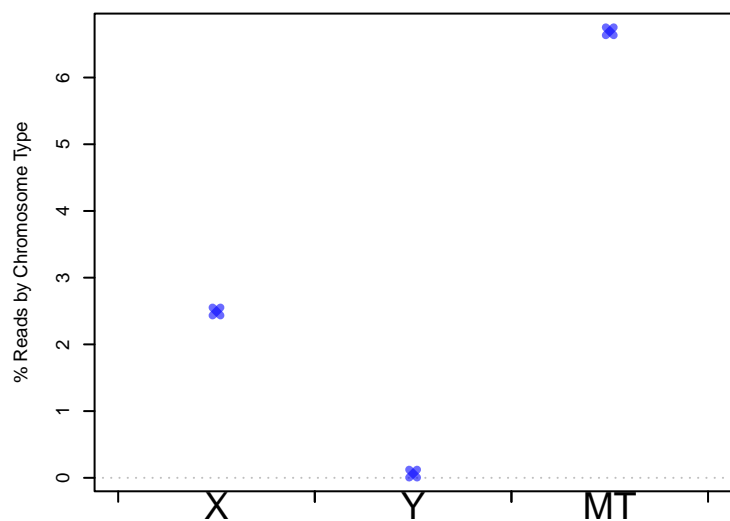
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Splice Junction Event Rates per Read-Pair



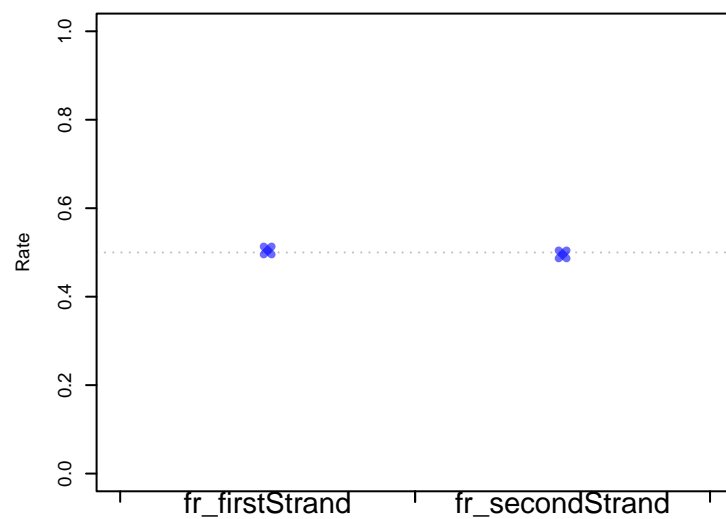
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Chromosome Distribution (Excluding Autosomes)



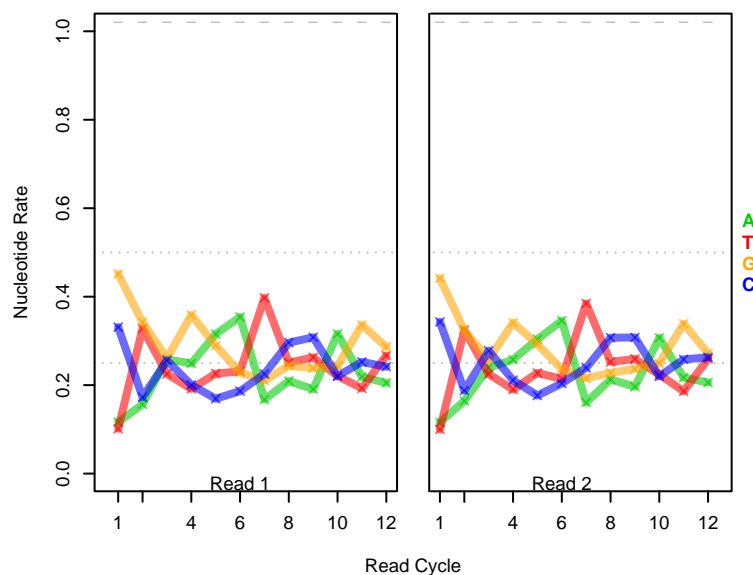
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Strandedness Test



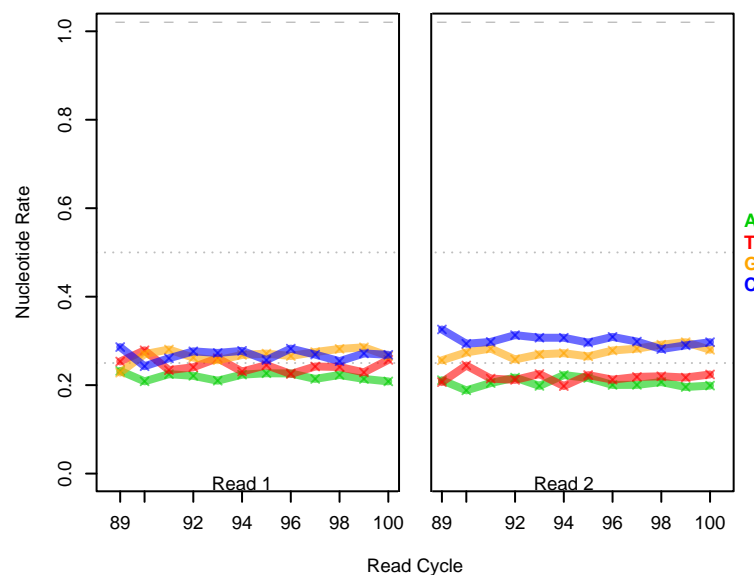
ai

Nucleotide Rate by Cycle, Leading Clipped bases (12)



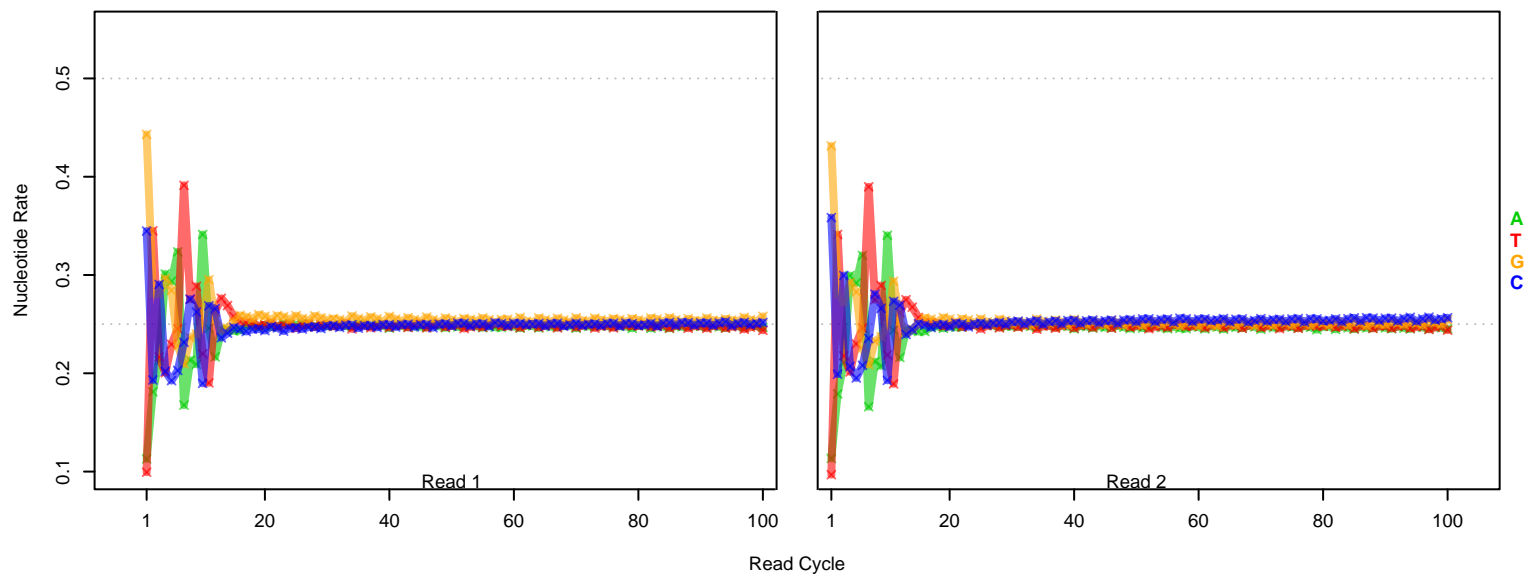
aj

Nucleotide Rate by Cycle, Trailing Clipped bases (12)



ak

Raw Nucleotide Rate by Cycle



al

Nucleotide Rate by Cycle, Aligned bases only

