

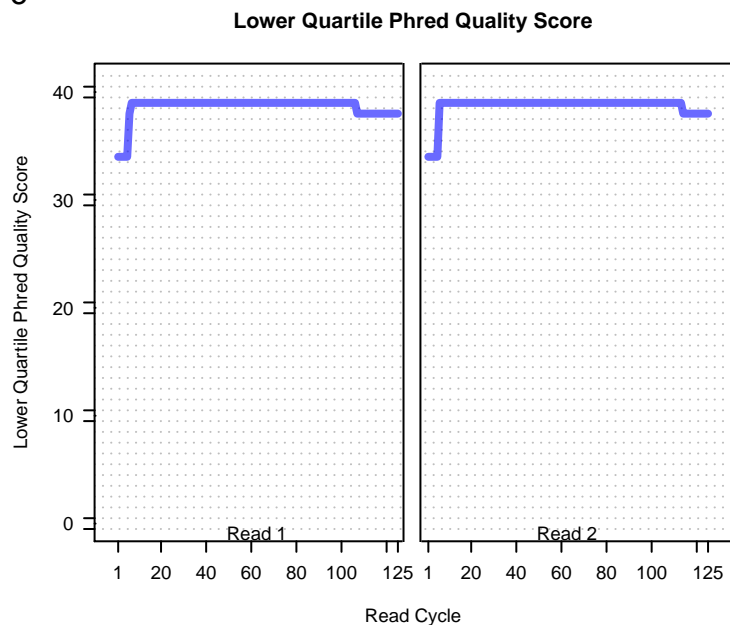
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All Quality Control Summary Plots

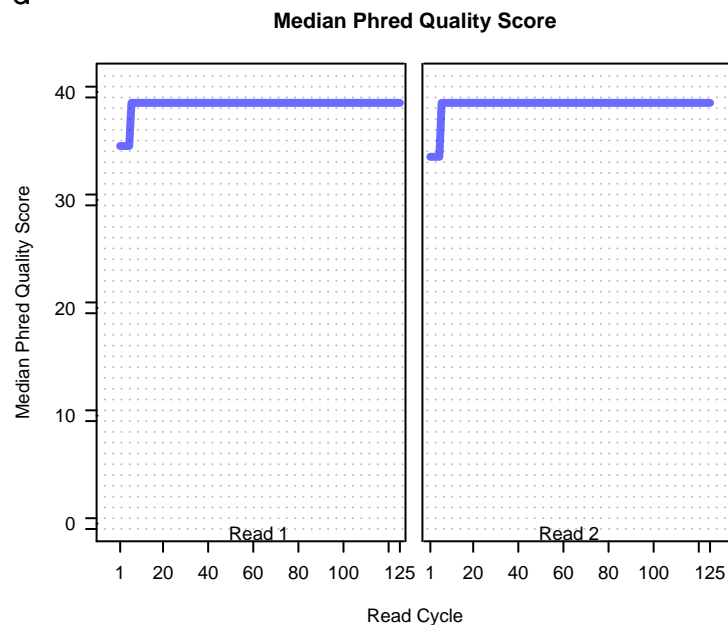
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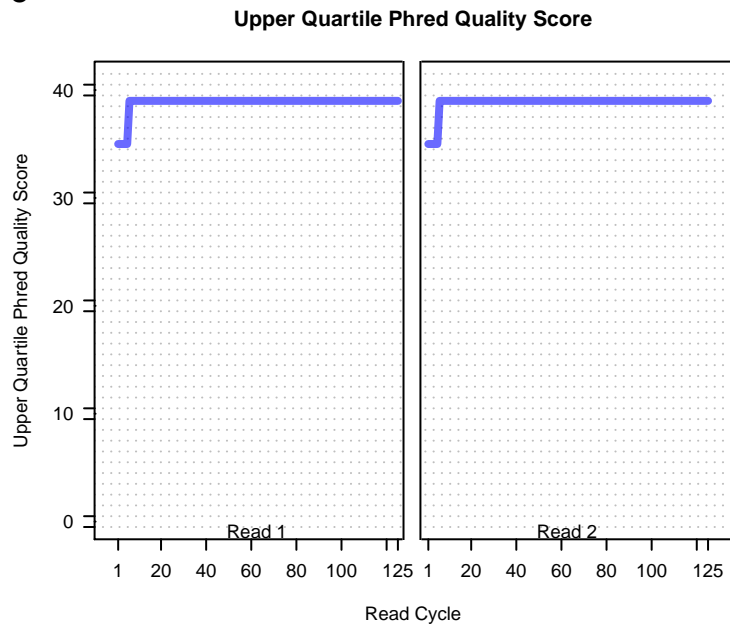
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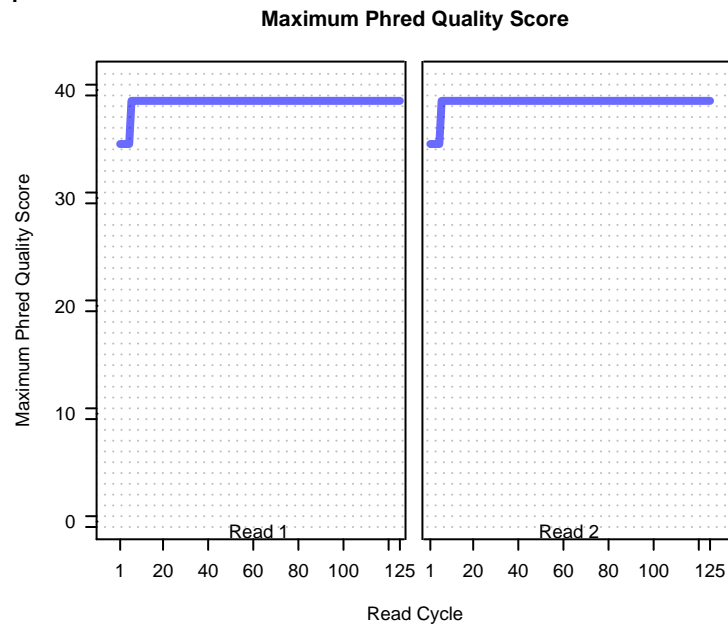
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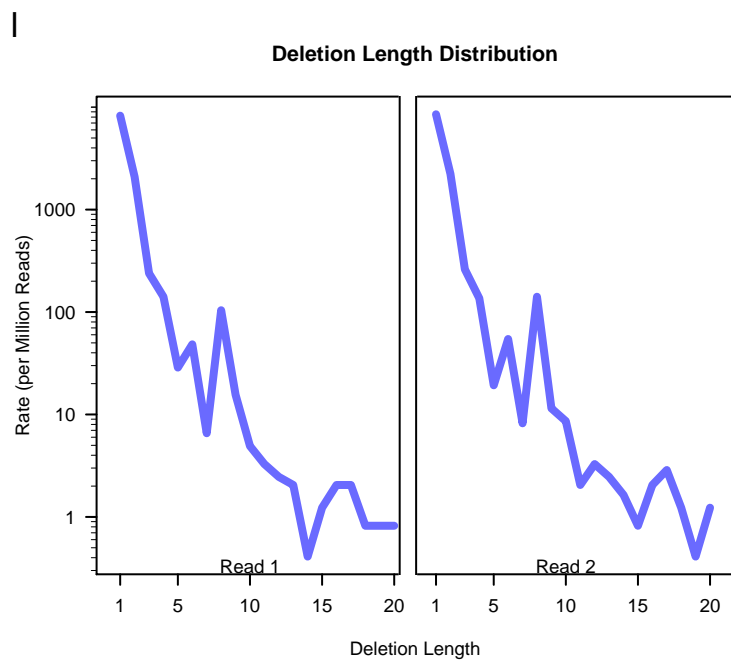
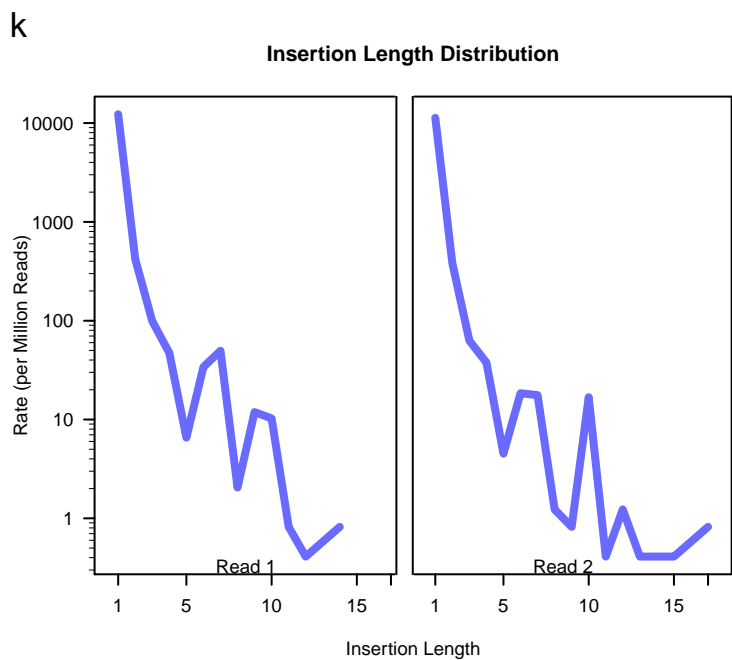
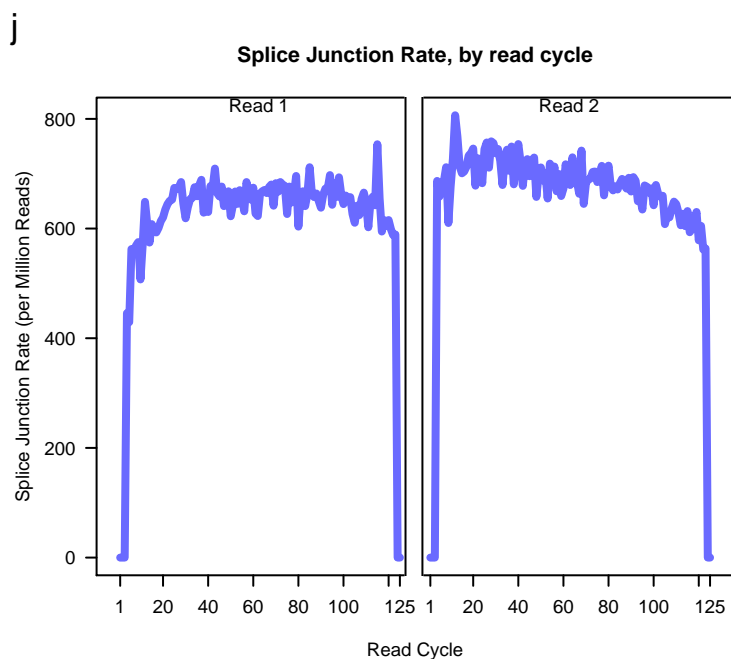
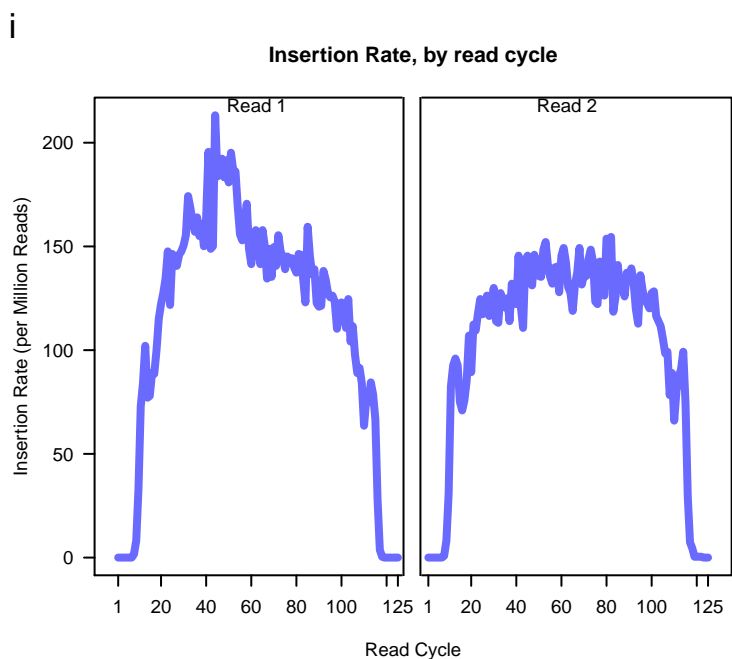
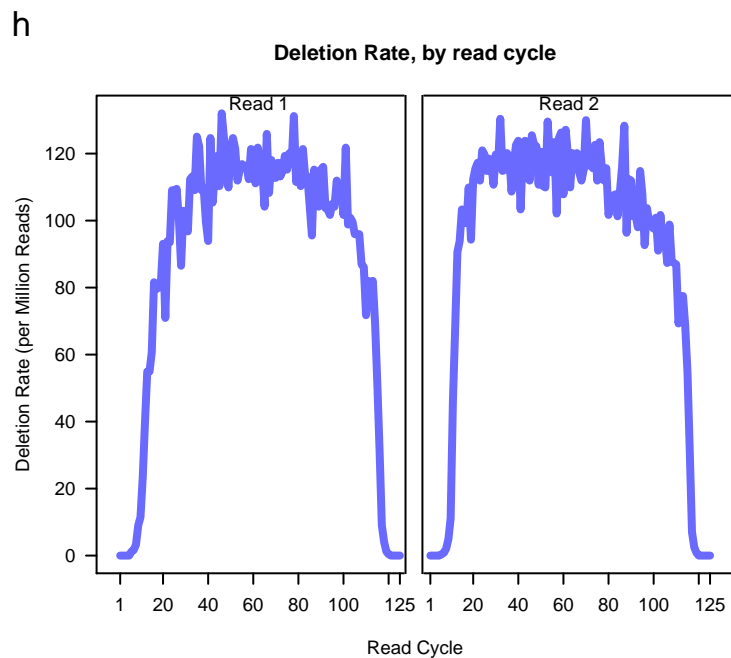
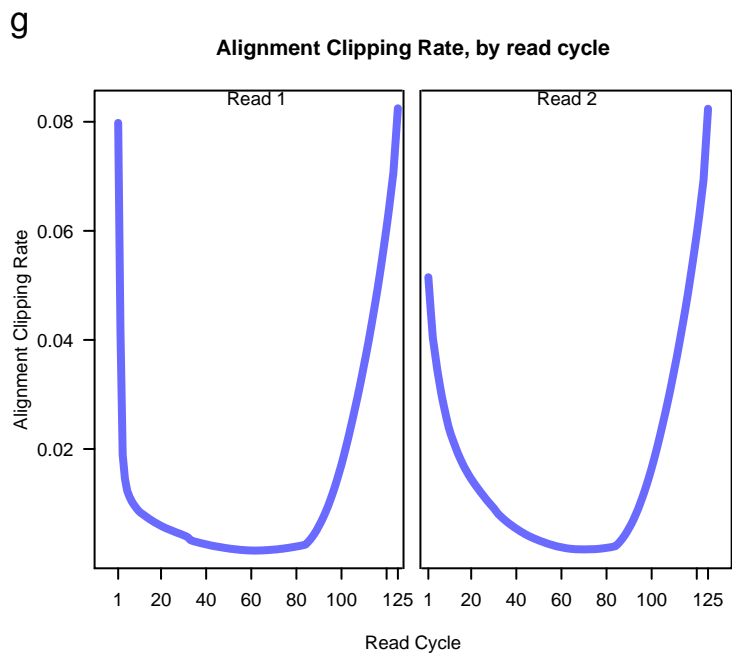


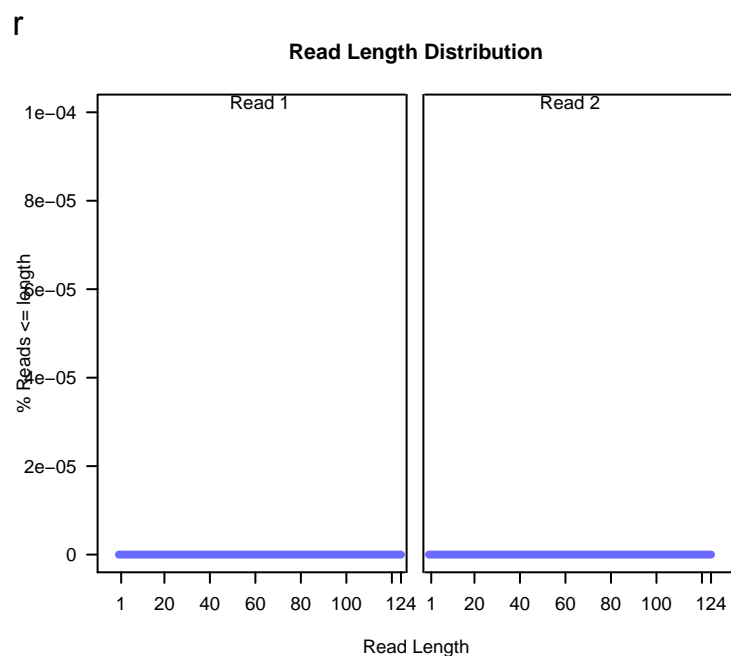
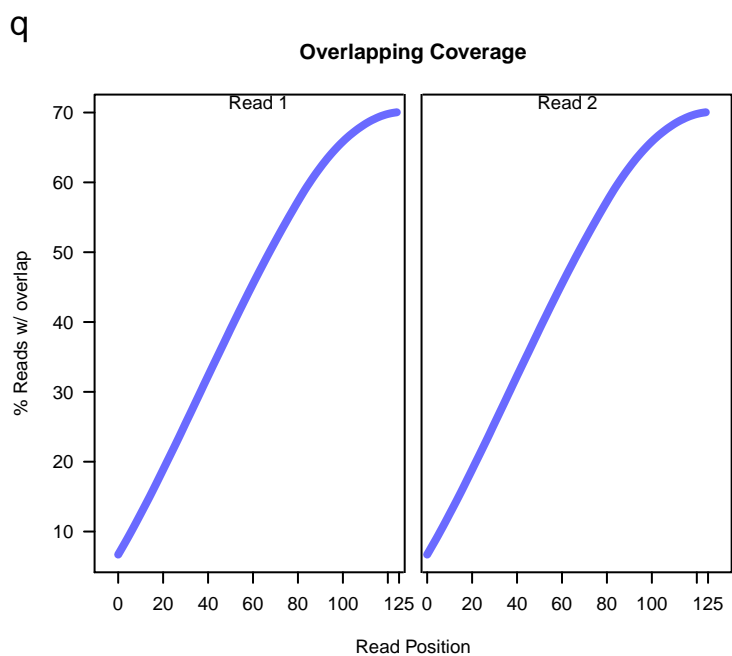
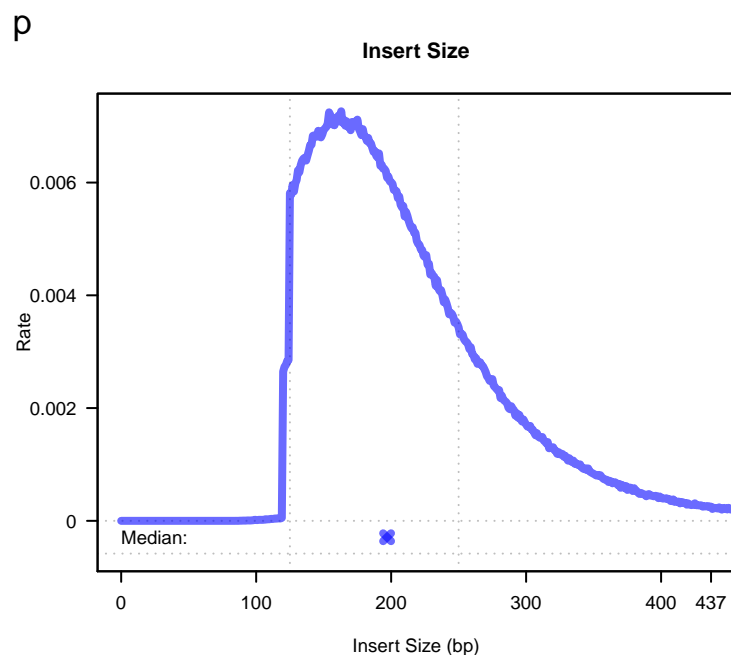
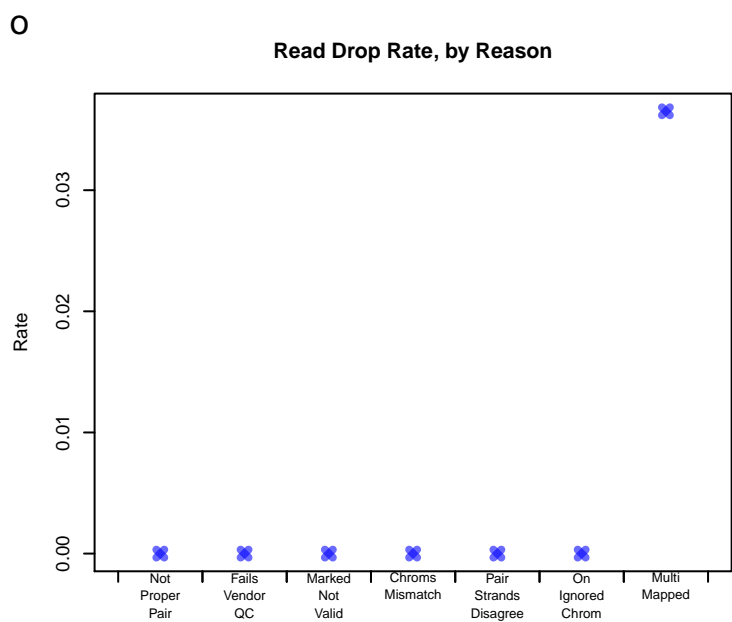
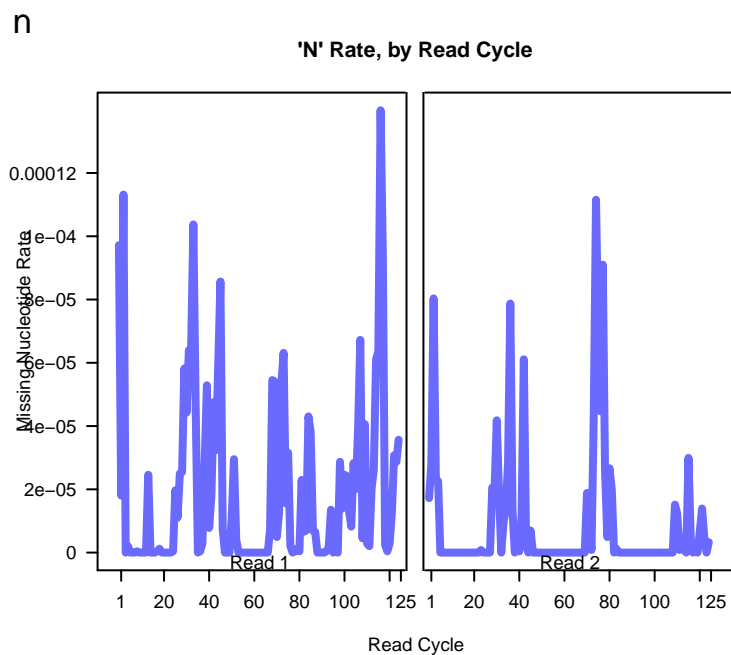
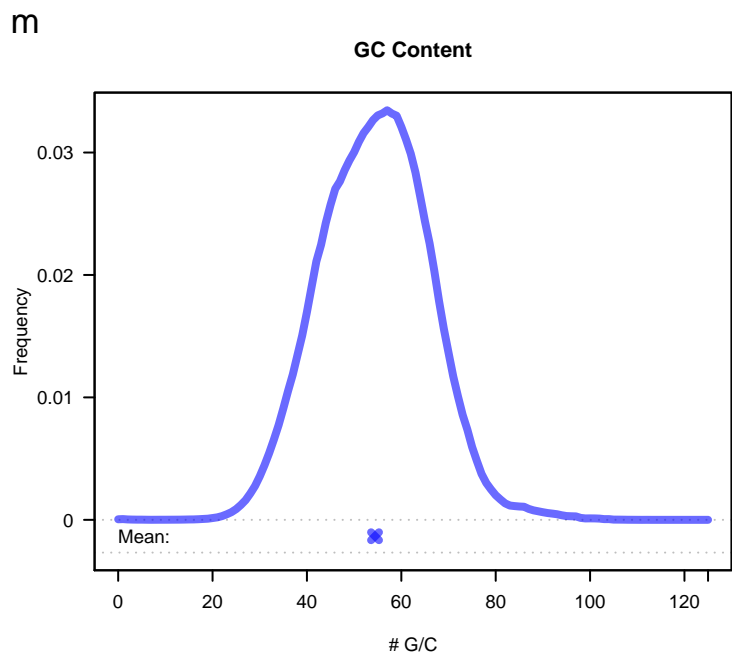
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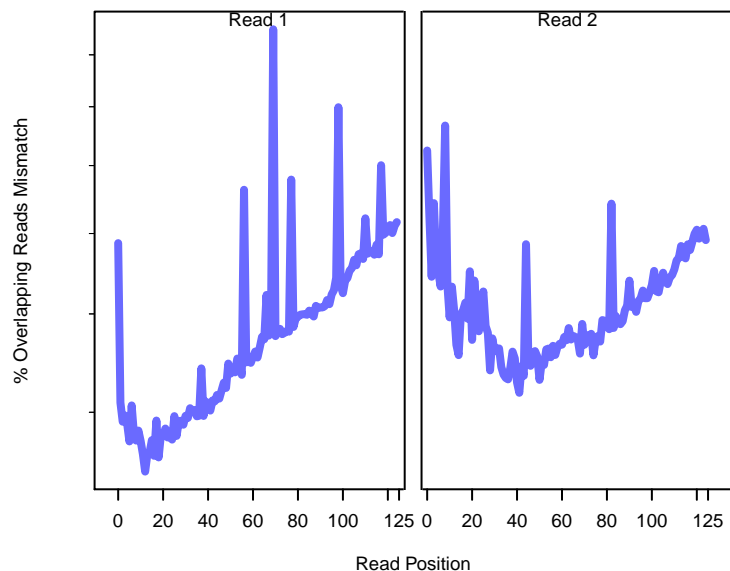






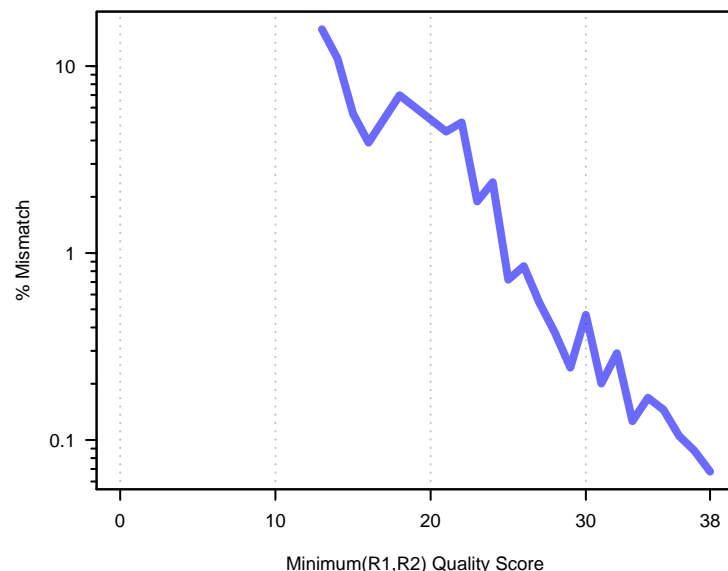
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Overlap Mismatch by Read Cycle



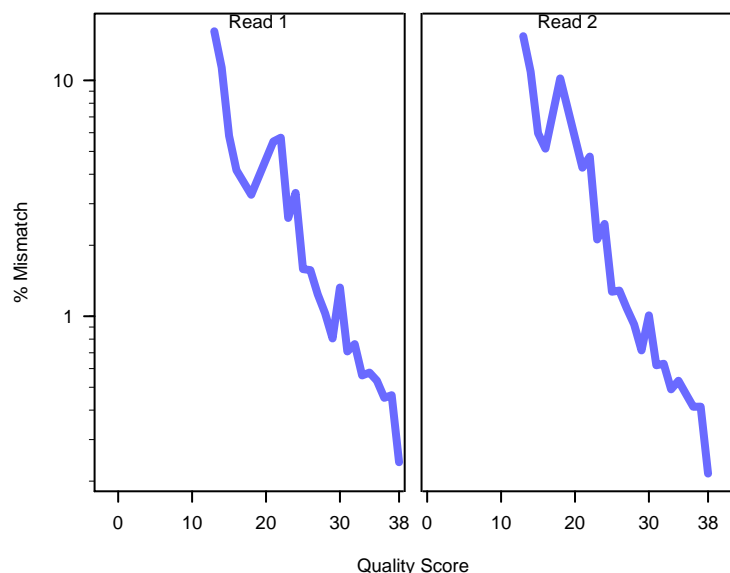
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Overlapping Mismatch by Min Qual



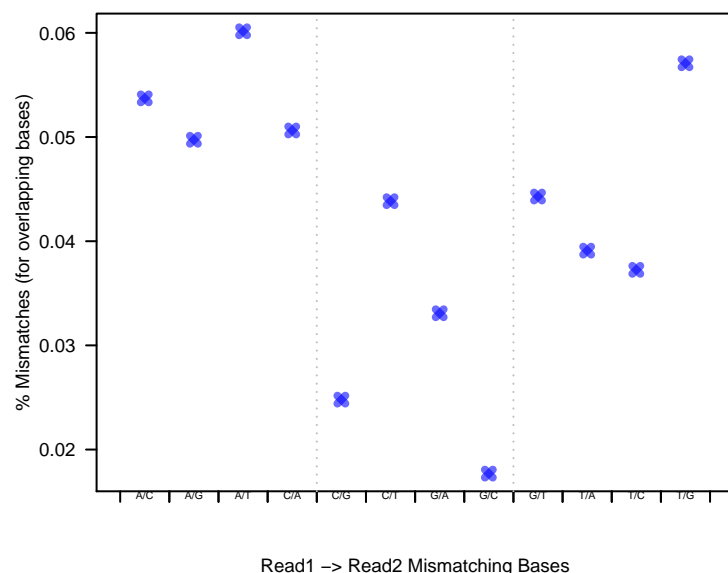
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Overlap Mismatch by Read Qual



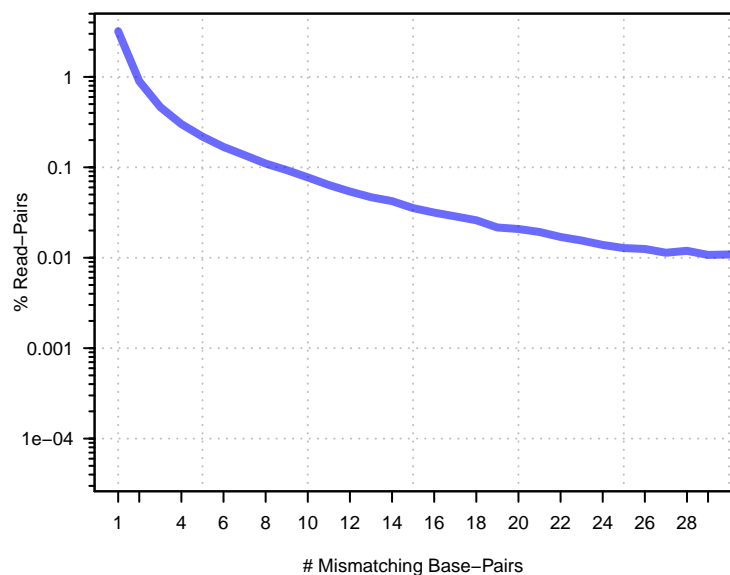
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Overlap Mismatch Combinations



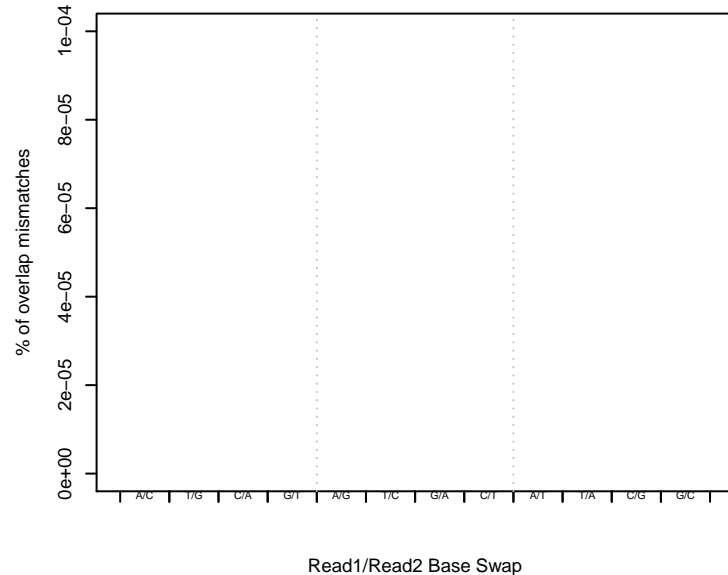
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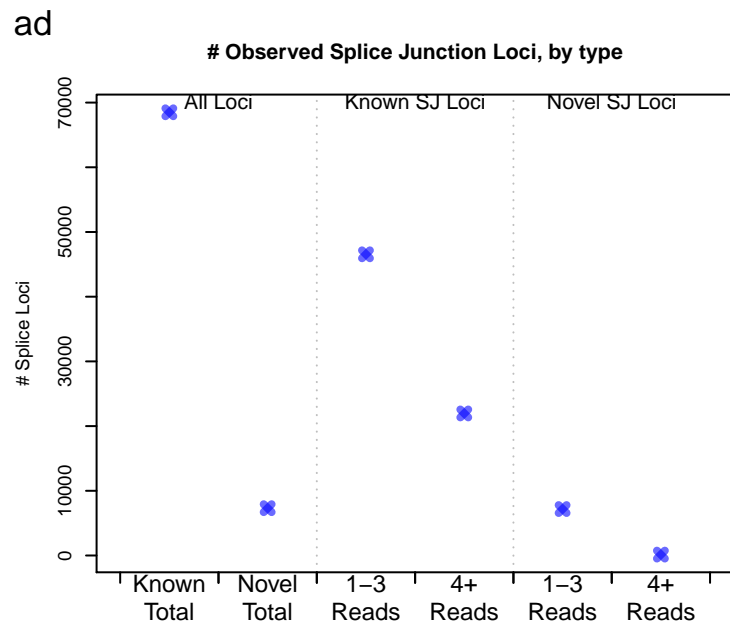
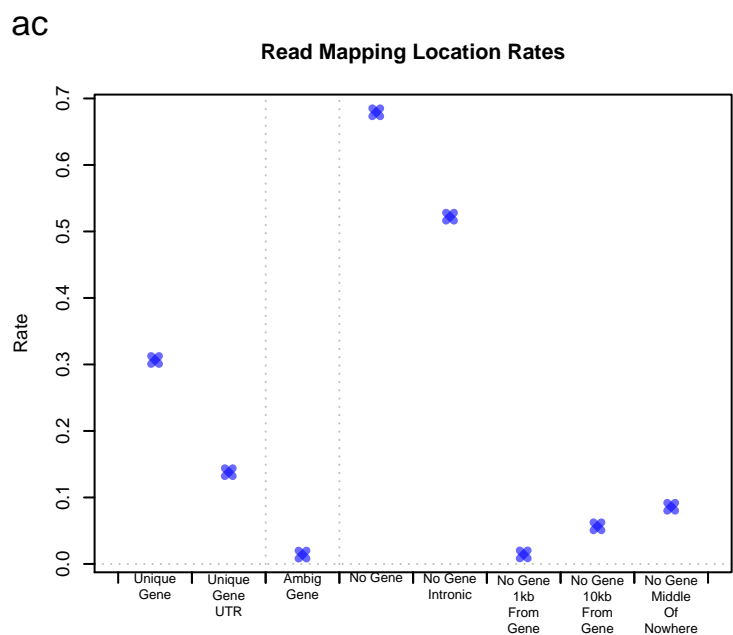
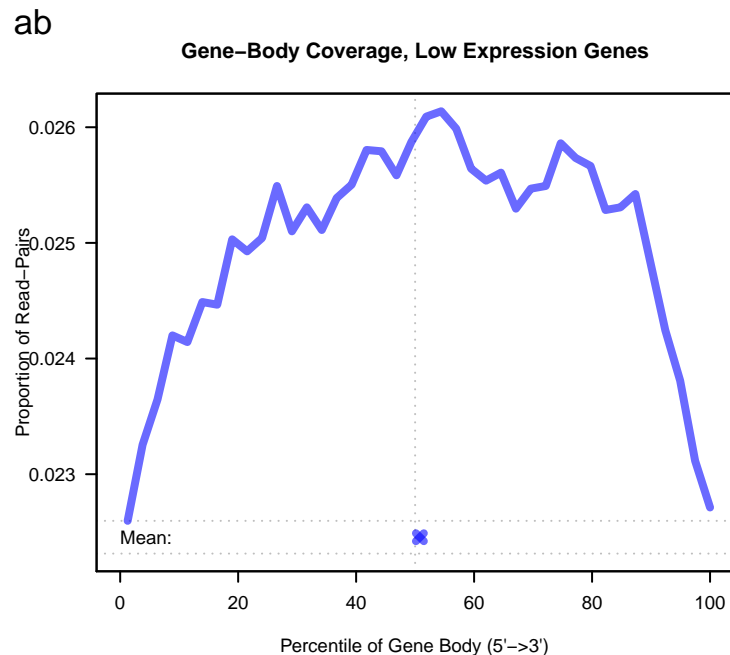
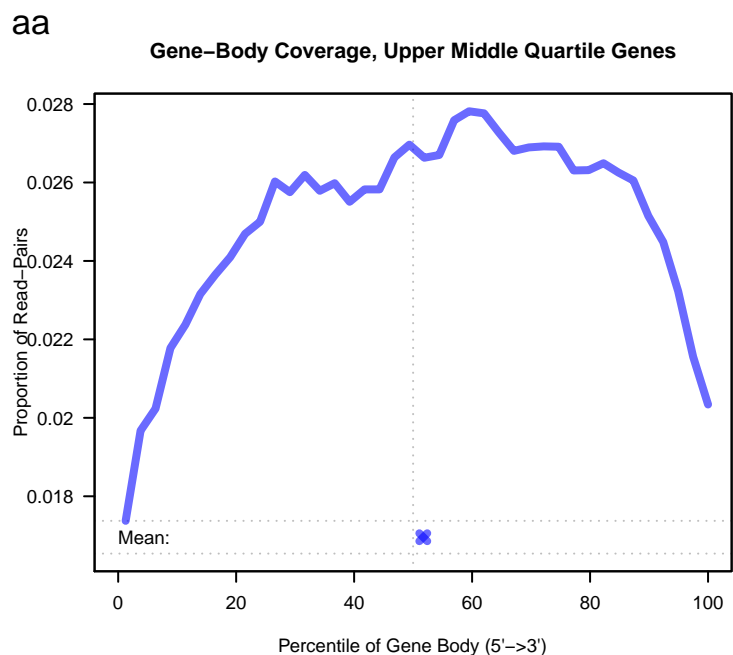
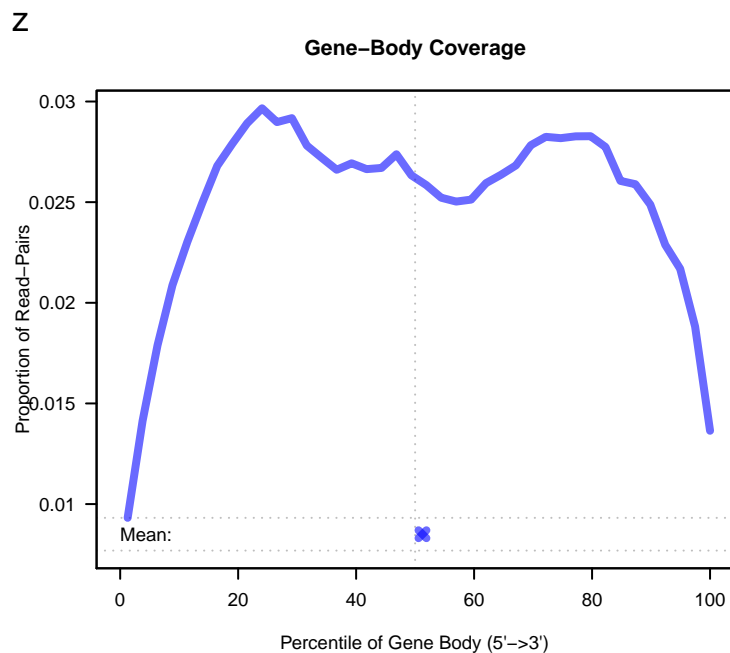
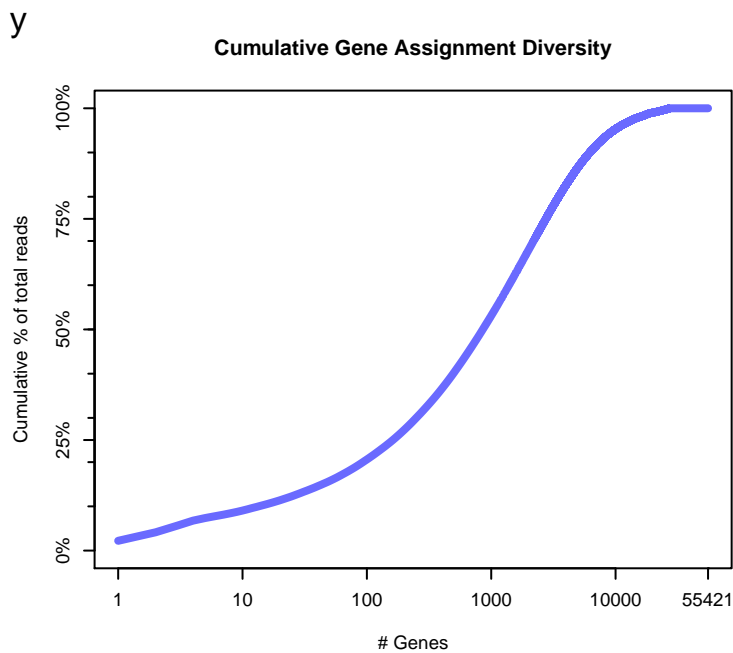
Overlap Mismatch Size Frequency



X

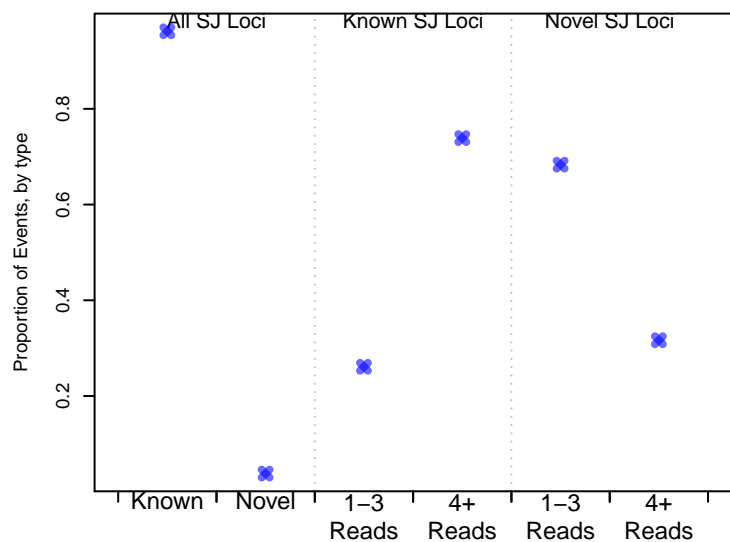
Overlap Mismatch Pairs At Phred == 41





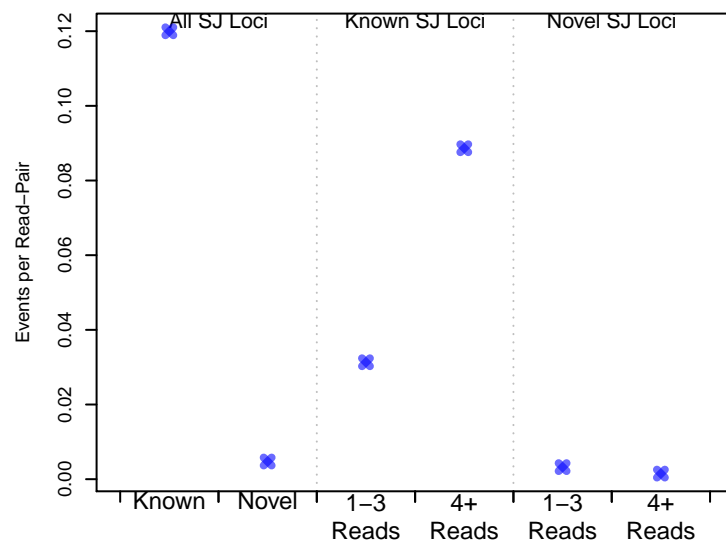
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Breakdown of Splice Junction Events, by type



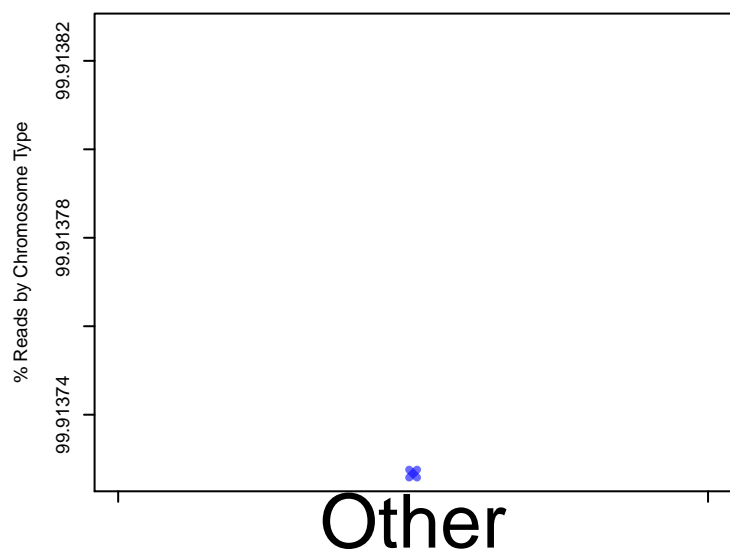
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Splice Junction Event Rates per Read-Pair



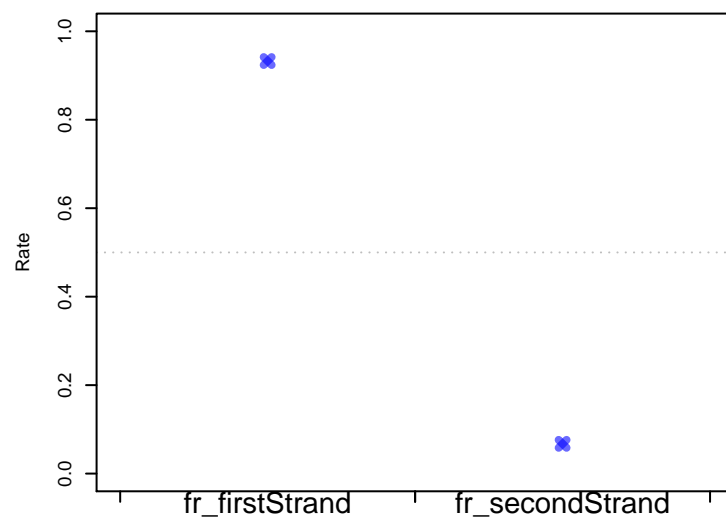
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Chromosome Distribution (Excluding Autosomes)



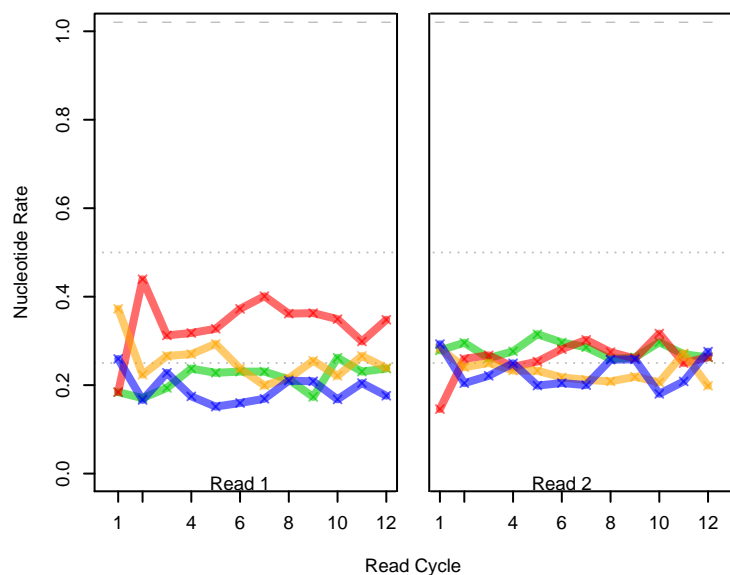
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Strandedness Test



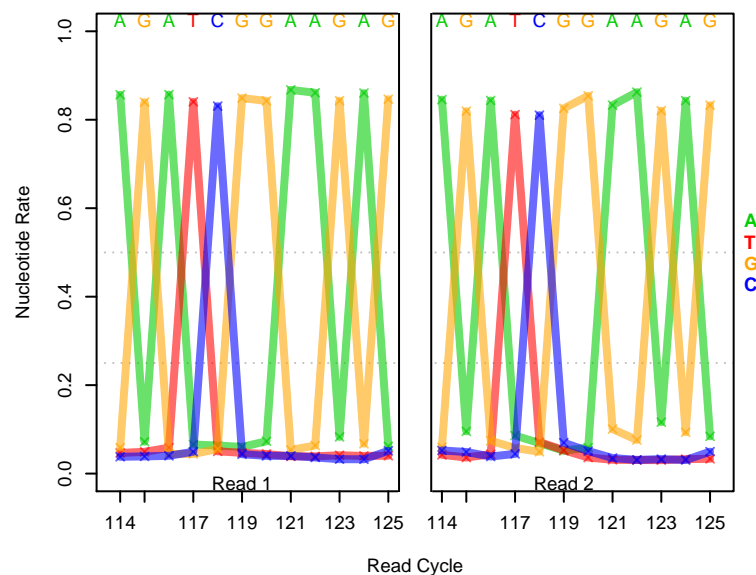
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Nucleotide Rate by Cycle, Leading Clipped bases (12)

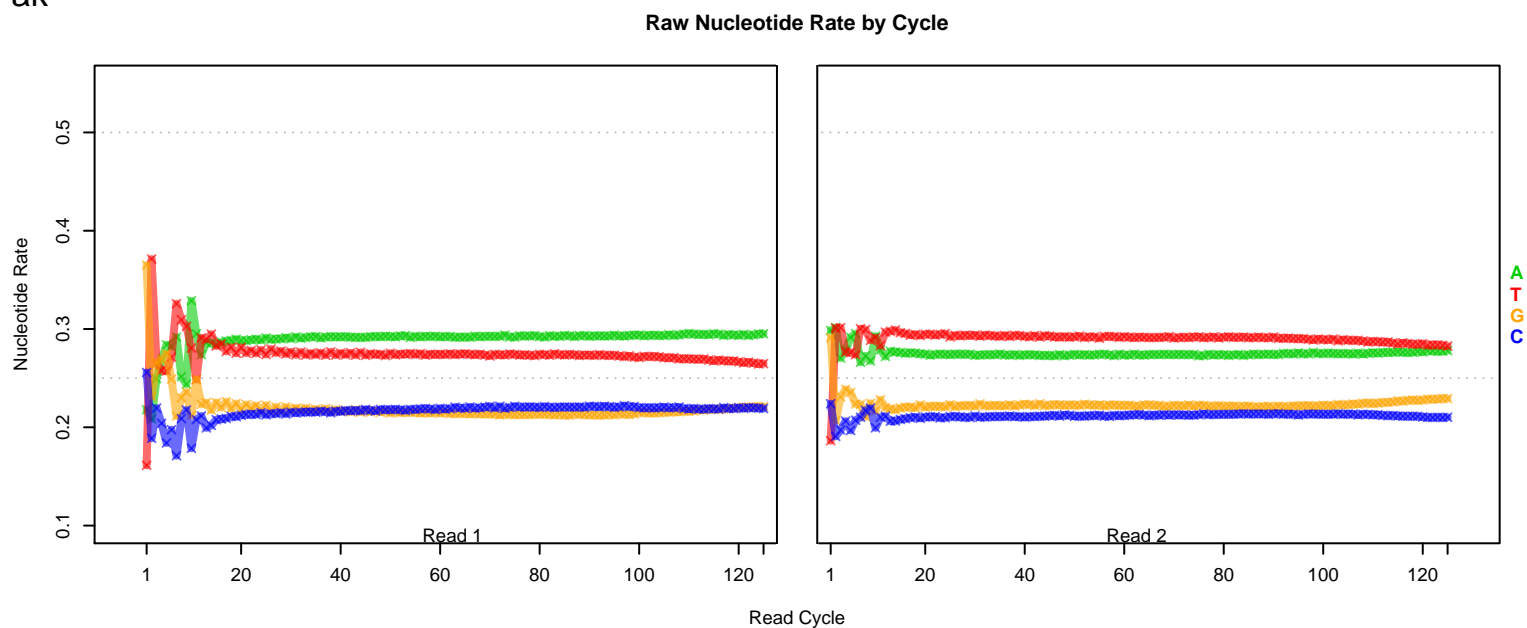


aj

Nucleotide Rate by Cycle, Trailing Clipped bases (12)



ak



al

