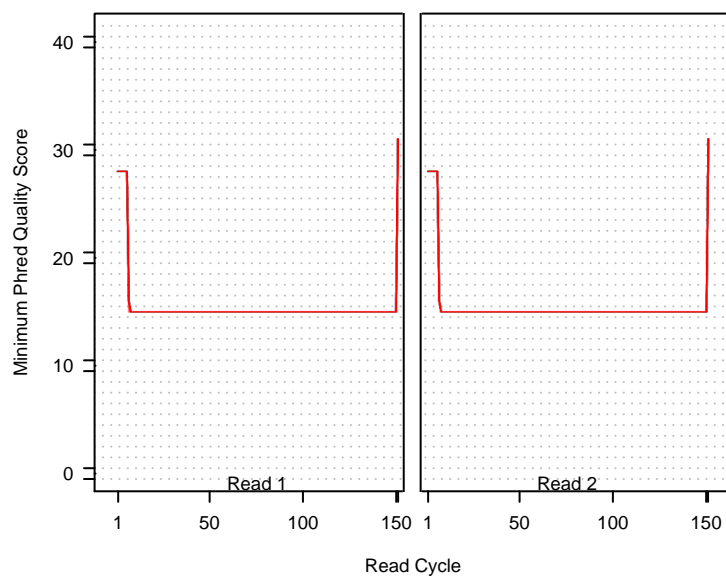


a

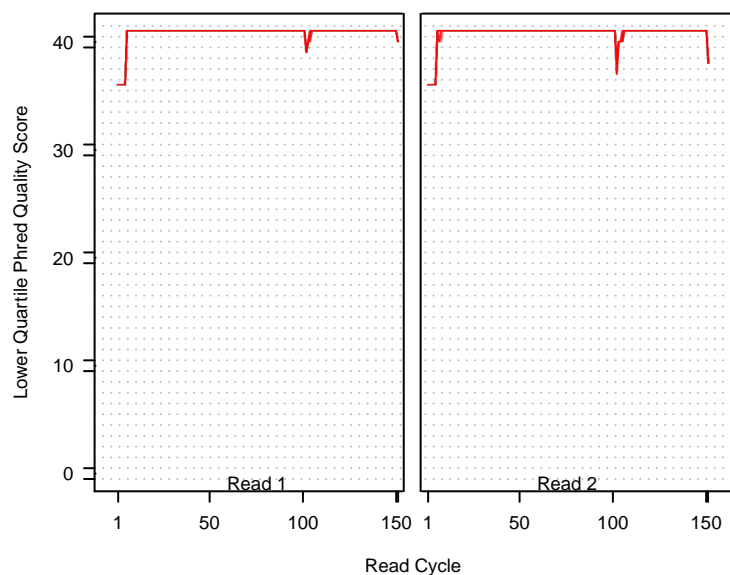
Summary Plots, By Lane



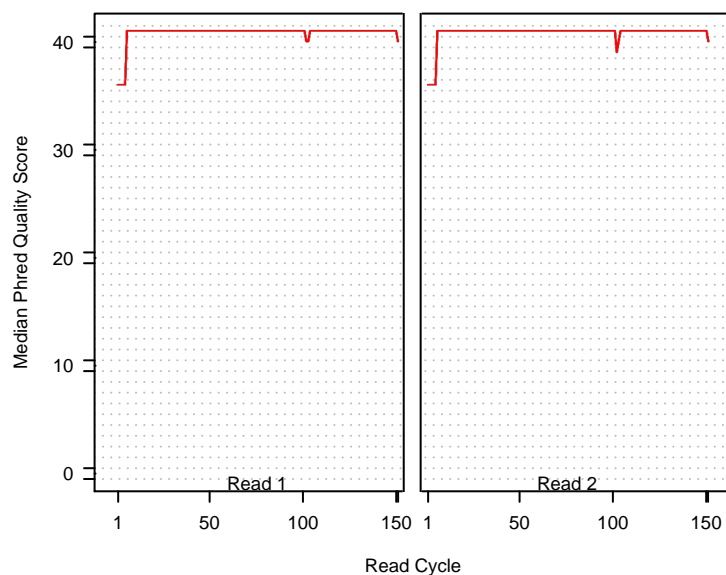
b

Minimum Phred Quality Score
Colored by Lane

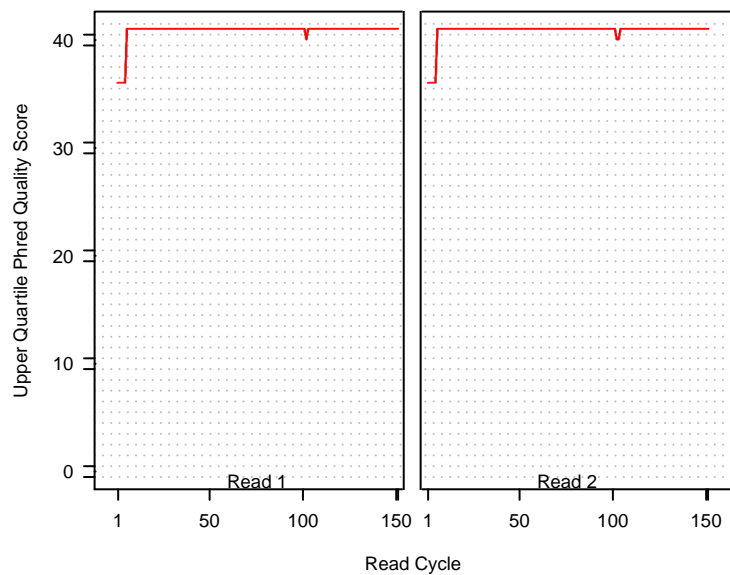
c

Lower Quartile Phred Quality Score
Colored by Lane

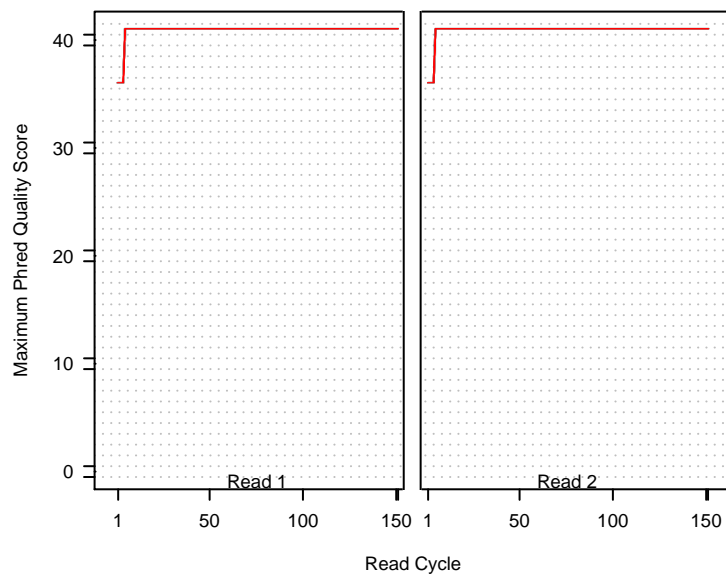
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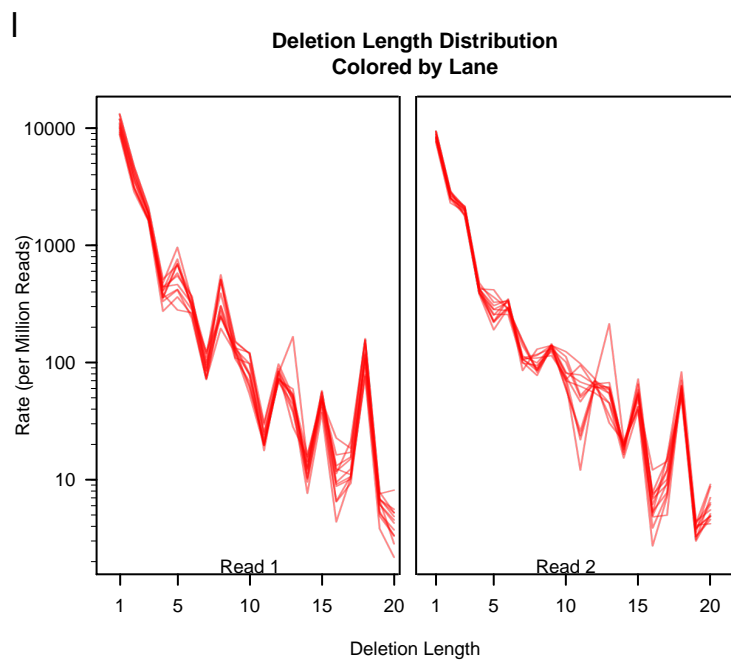
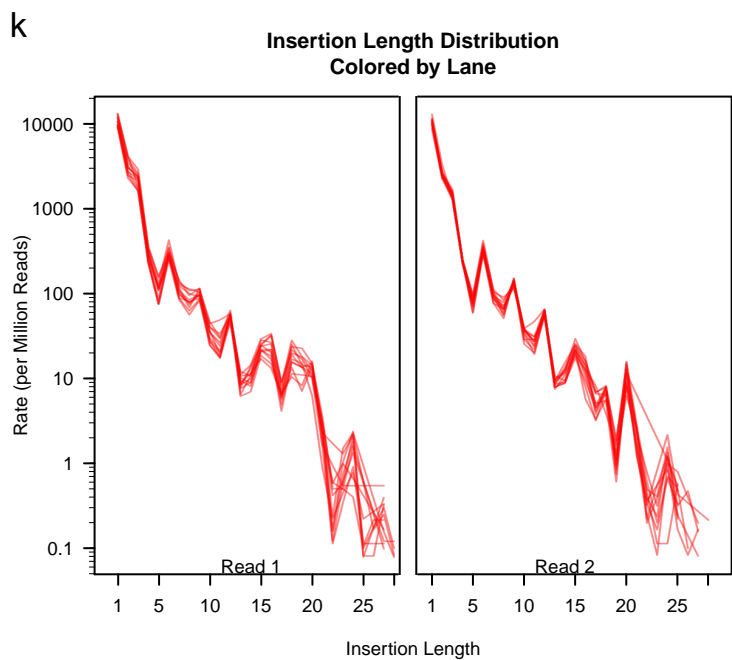
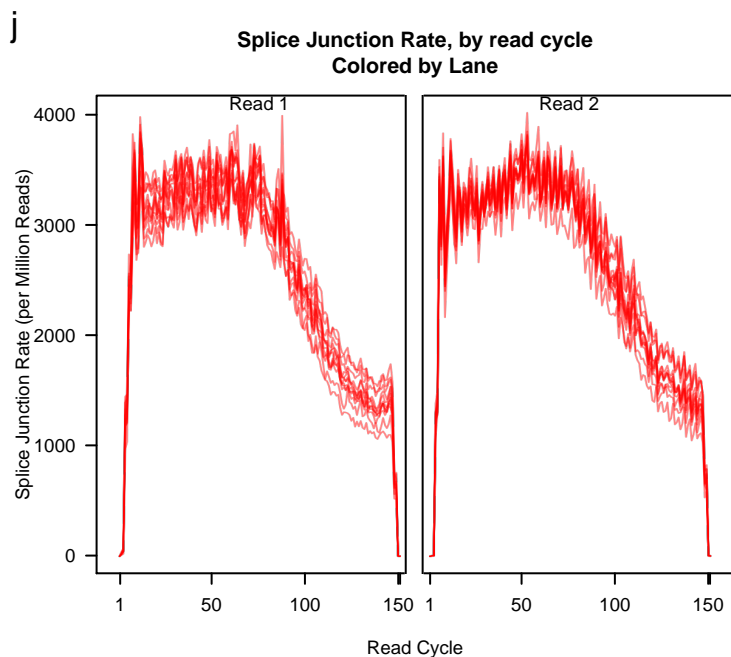
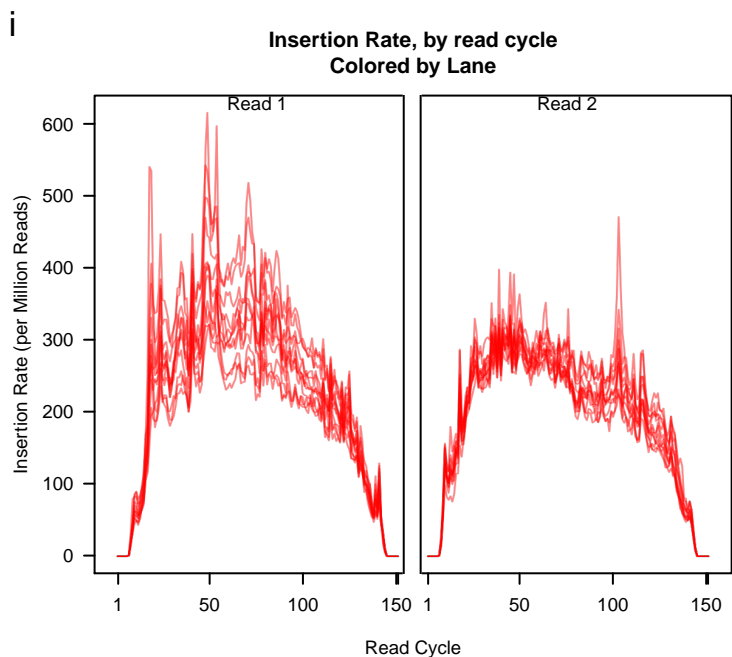
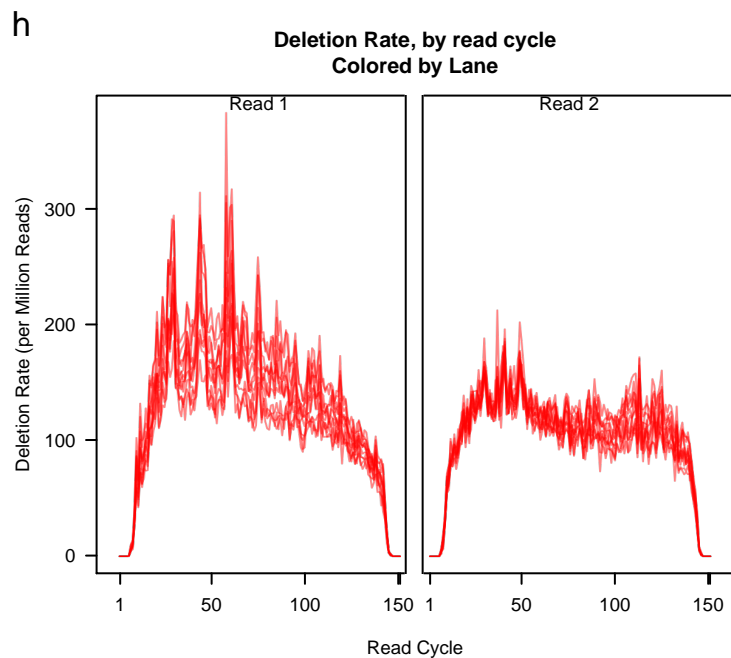
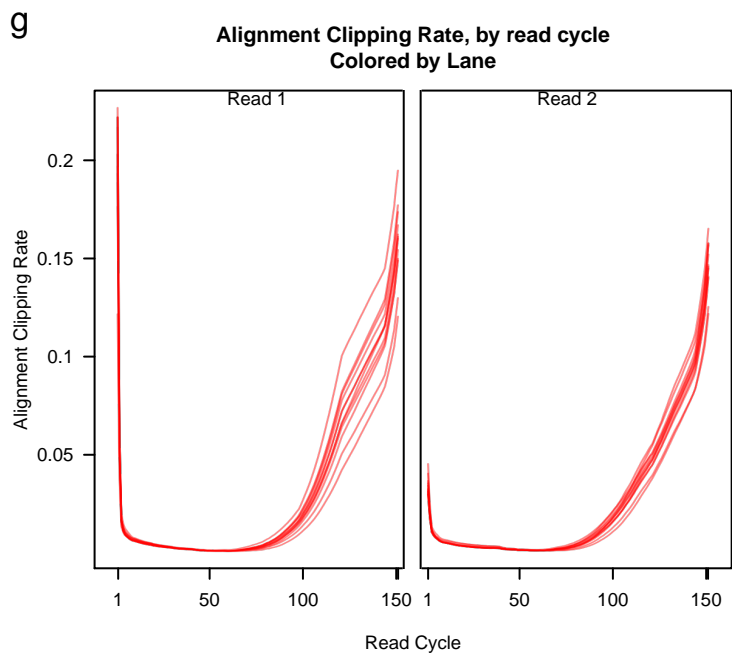
Median Phred Quality Score
Colored by Lane

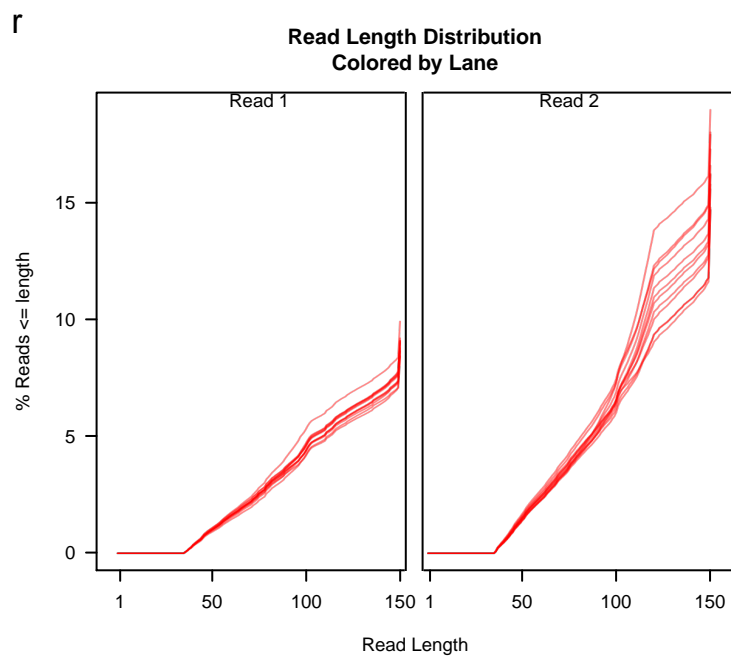
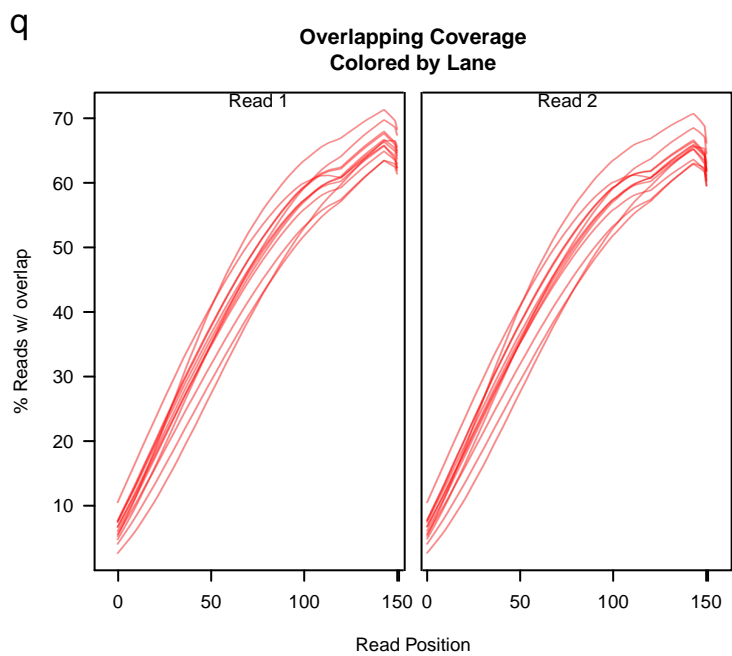
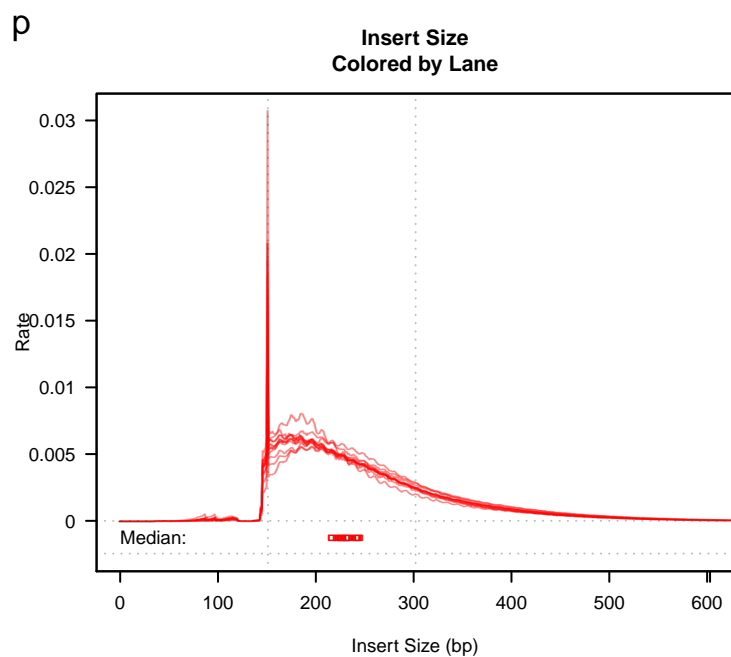
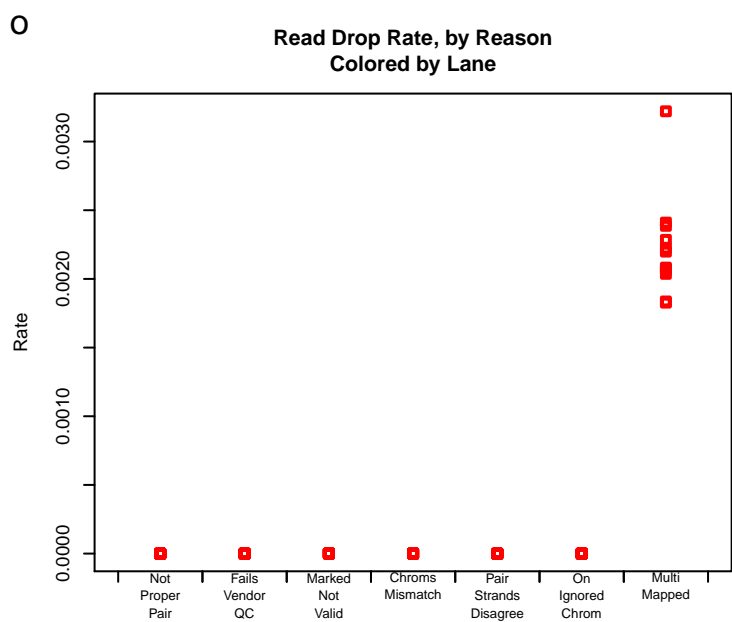
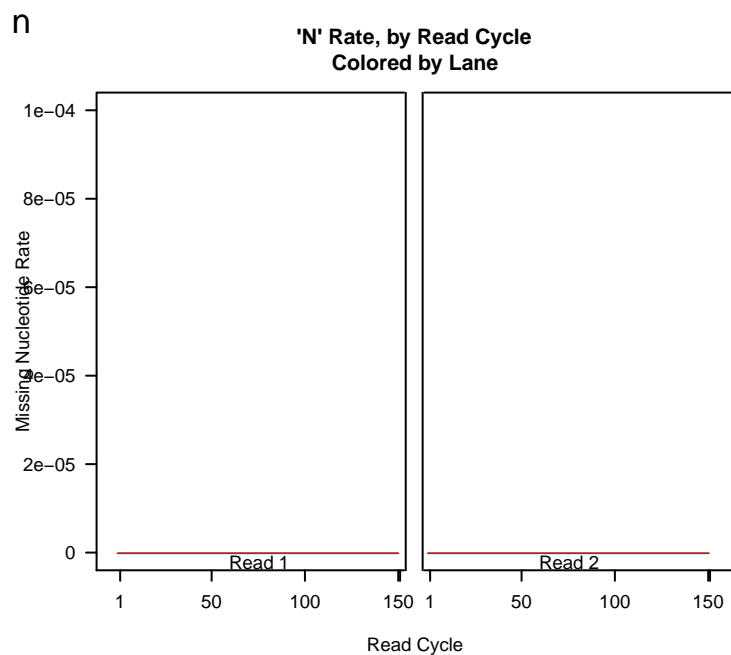
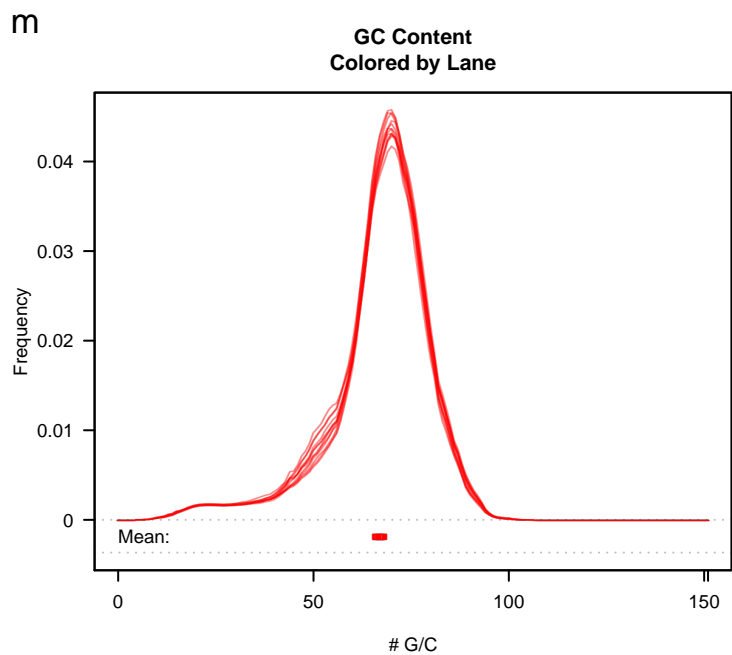
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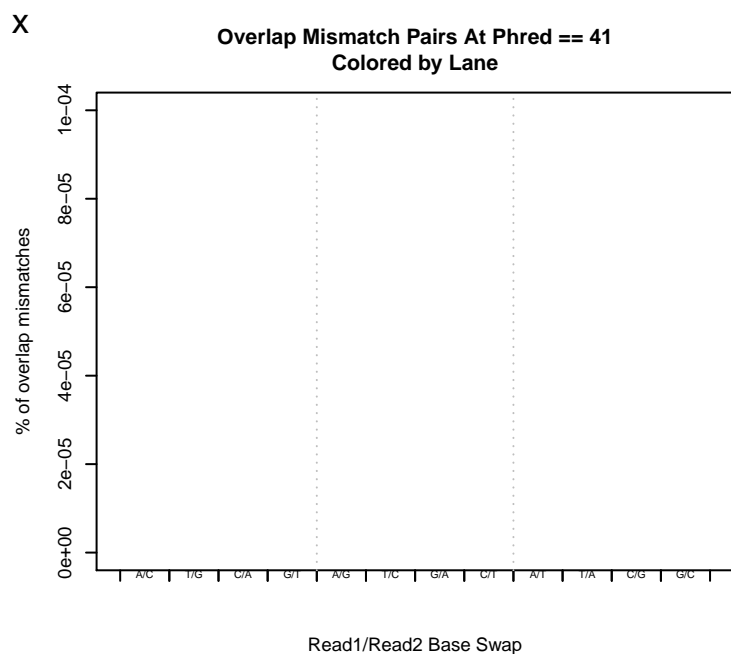
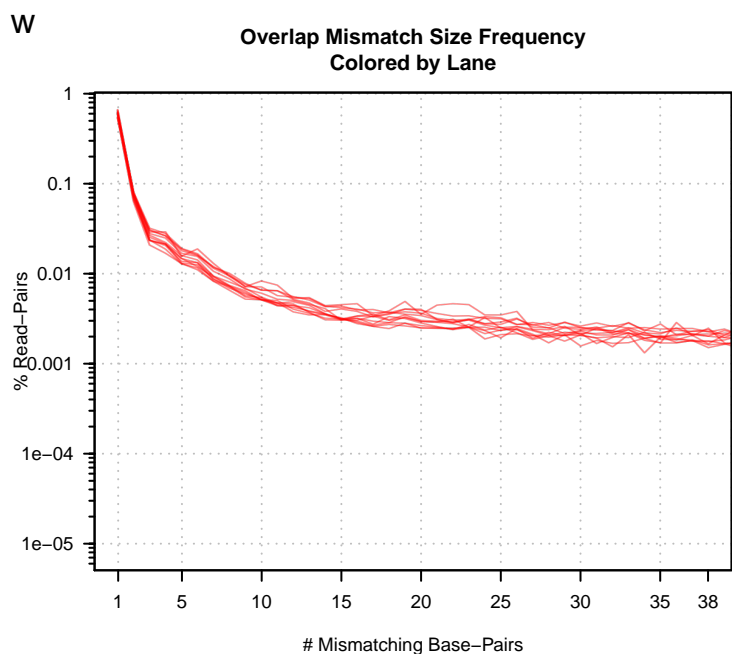
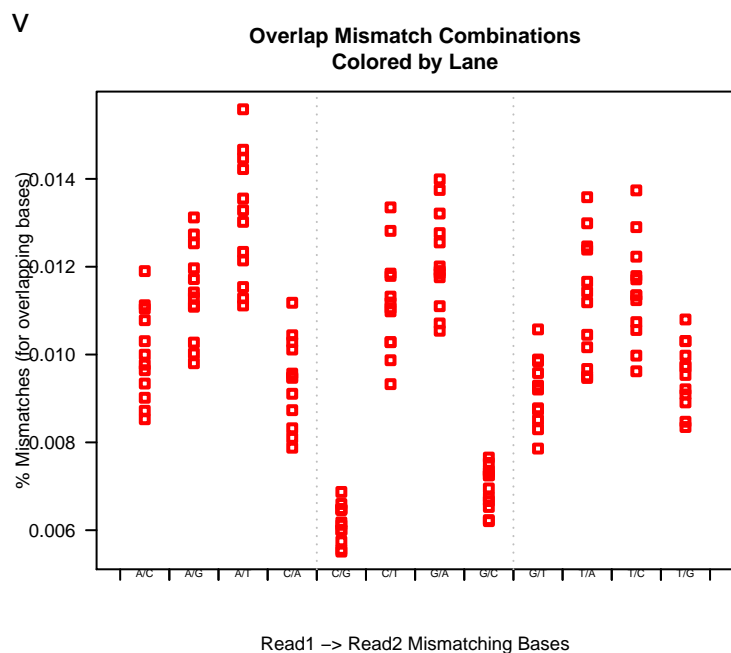
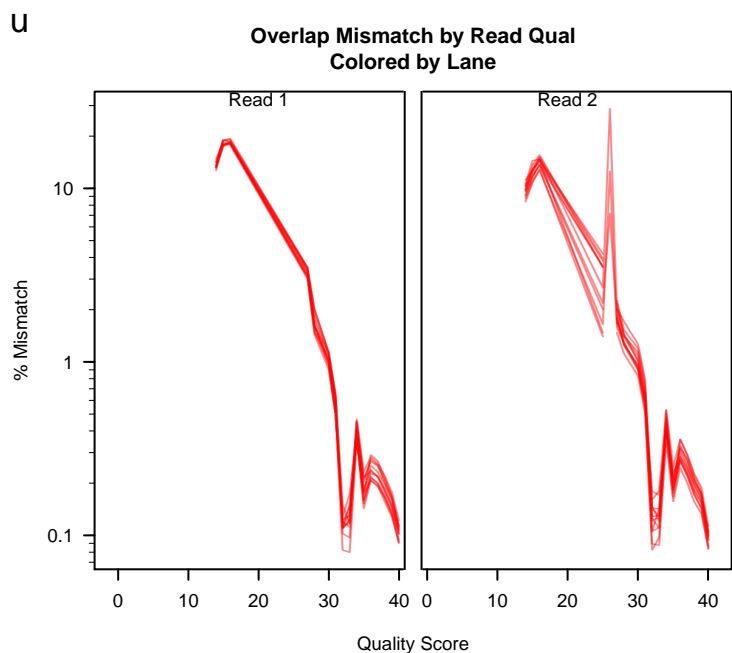
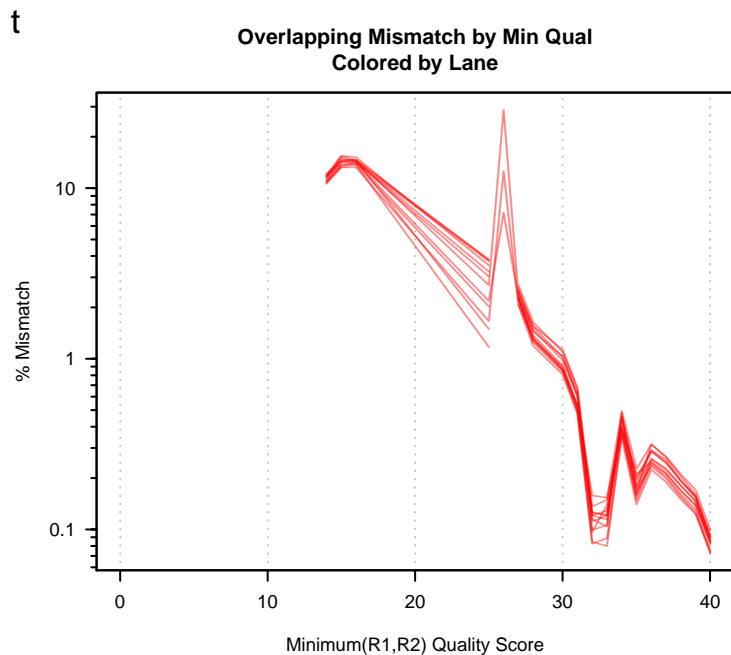
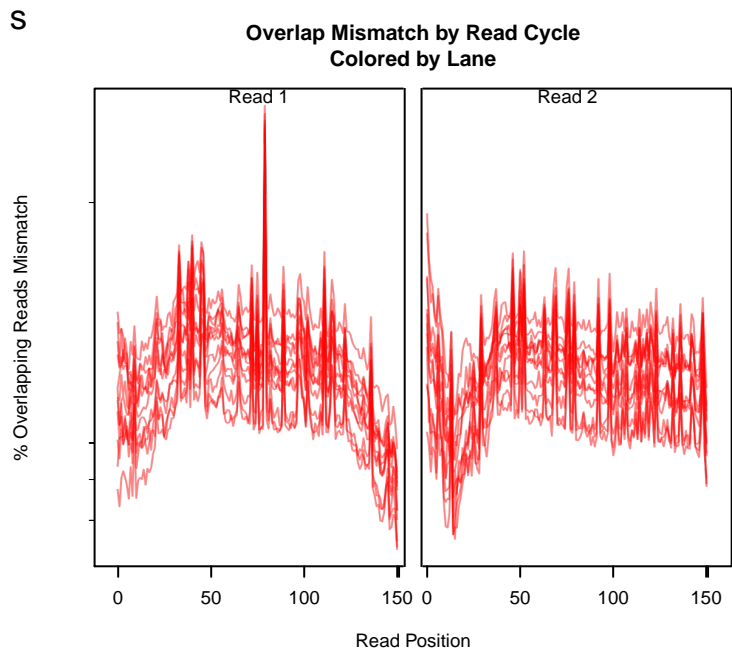
Upper Quartile Phred Quality Score
Colored by Lane

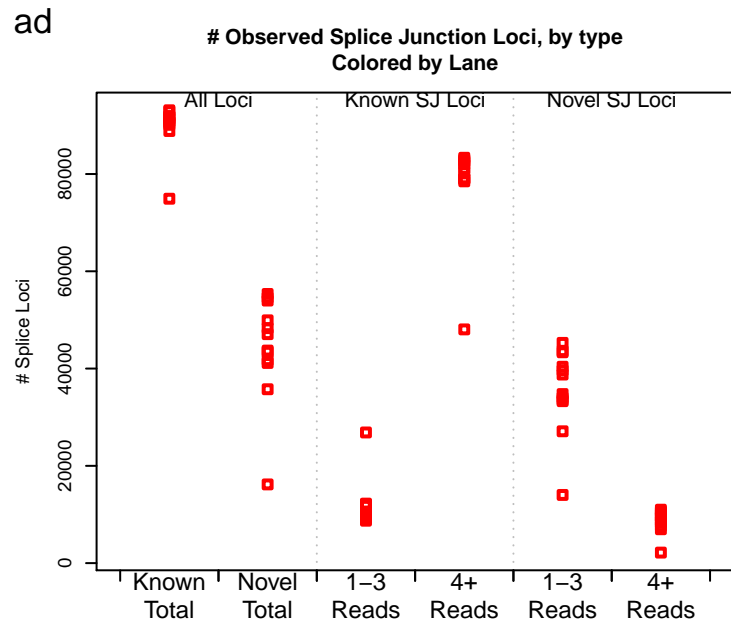
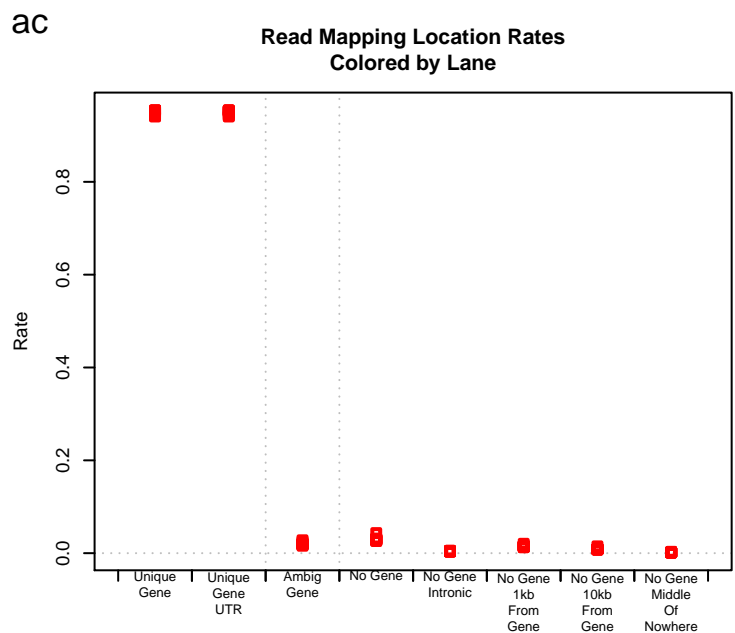
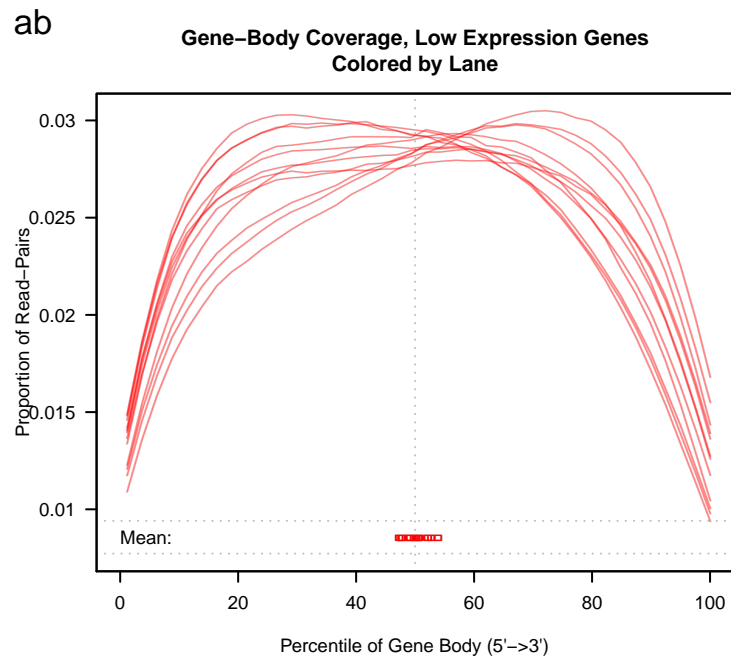
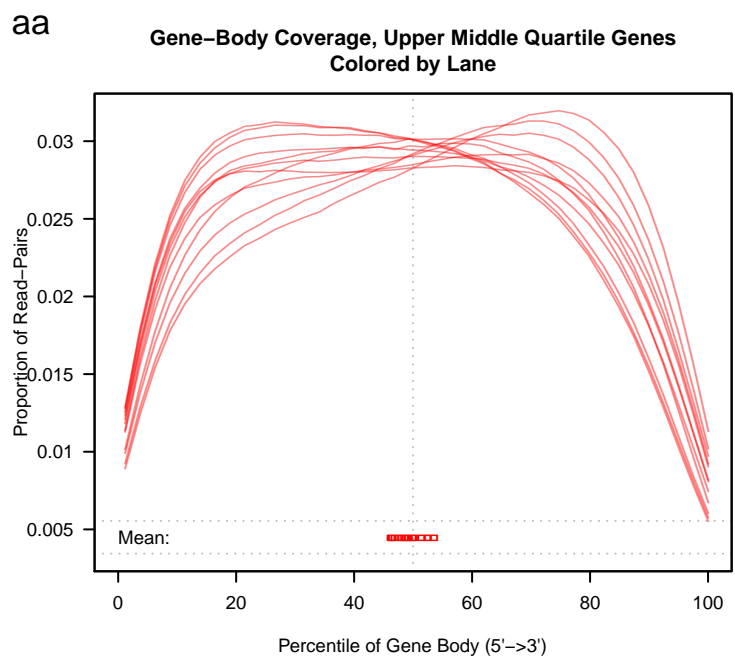
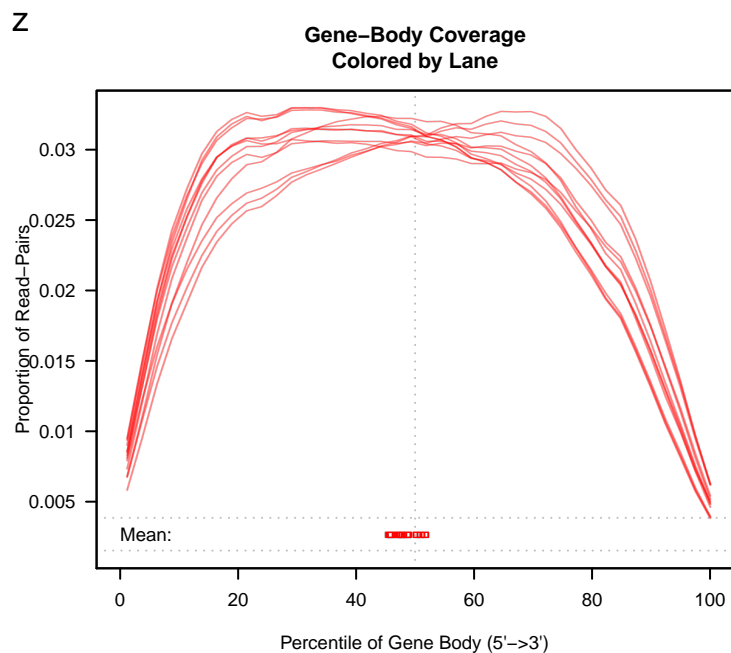
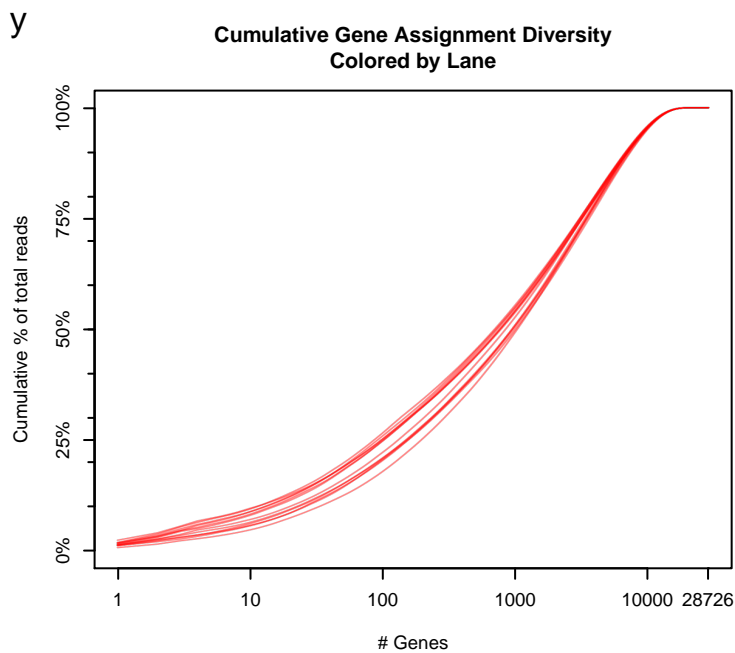
f

Maximum Phred Quality Score
Colored by Lane



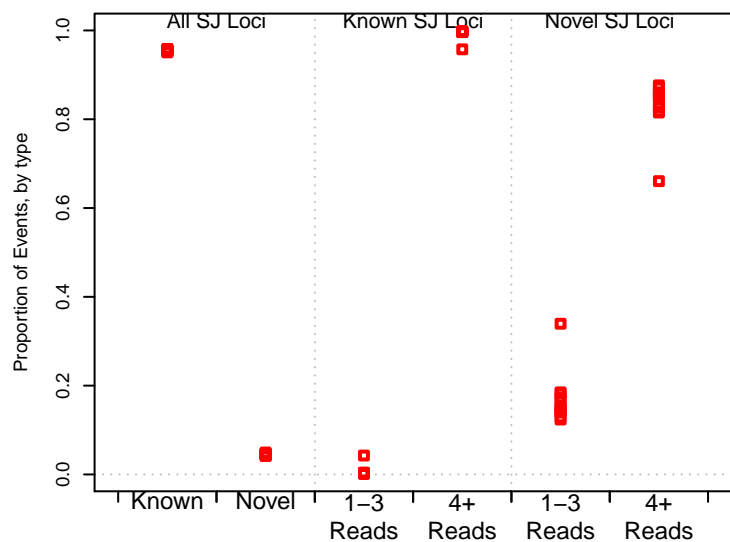






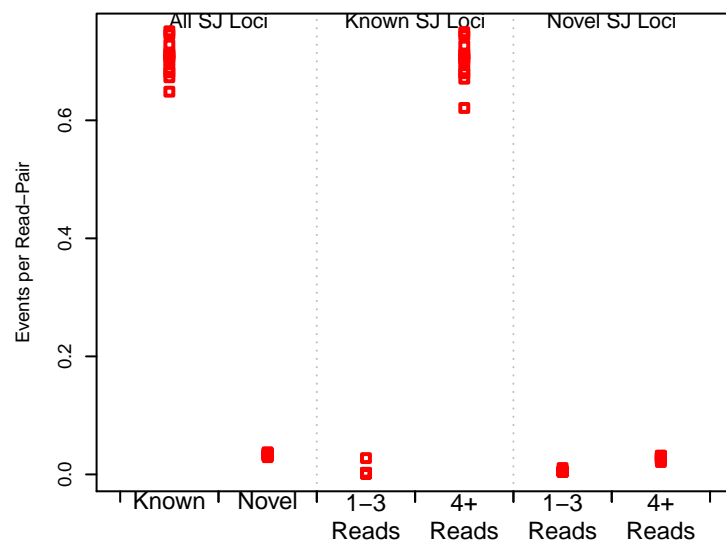
ae

Breakdown of Splice Junction Events, by type
Colored by Lane



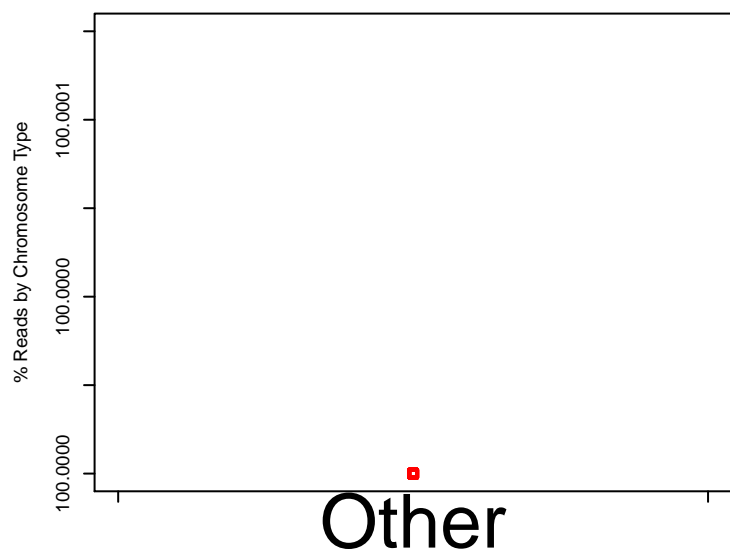
af

Splice Junction Event Rates per Read-Pair
Colored by Lane



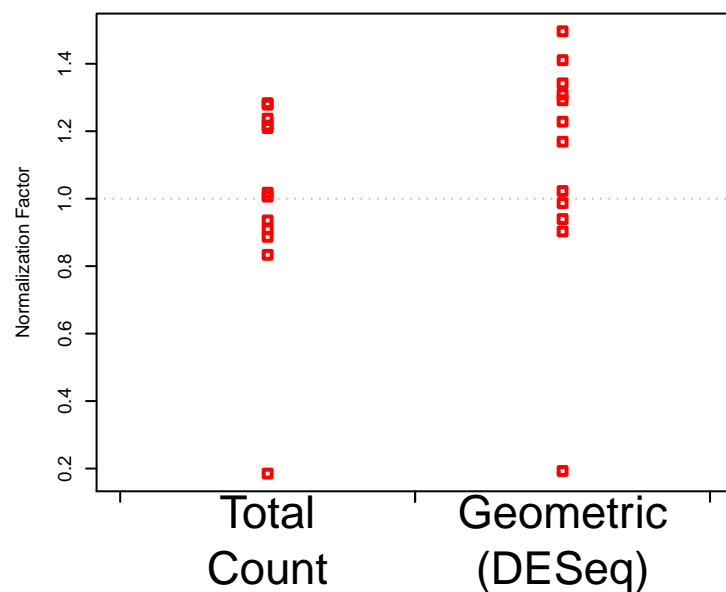
ag

Chromosome Distribution (Excluding Autosomes)
Colored by Lane



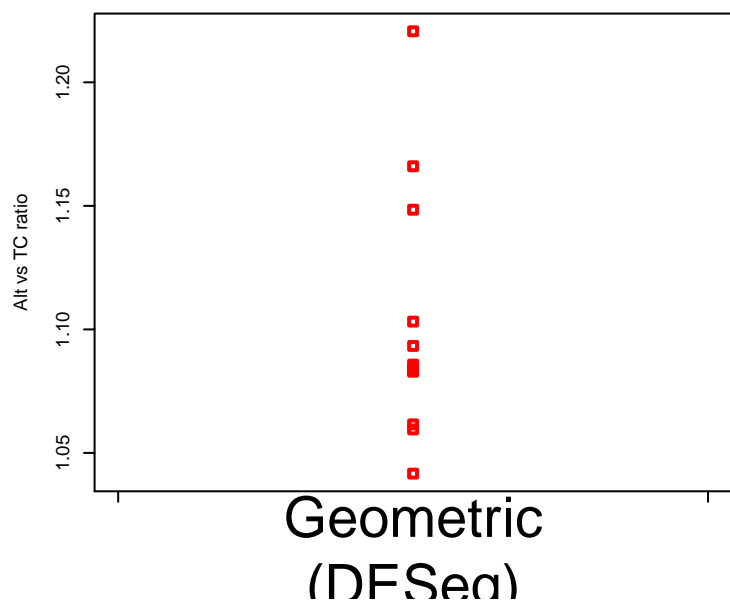
ah

Normalization Factors
Colored by Lane



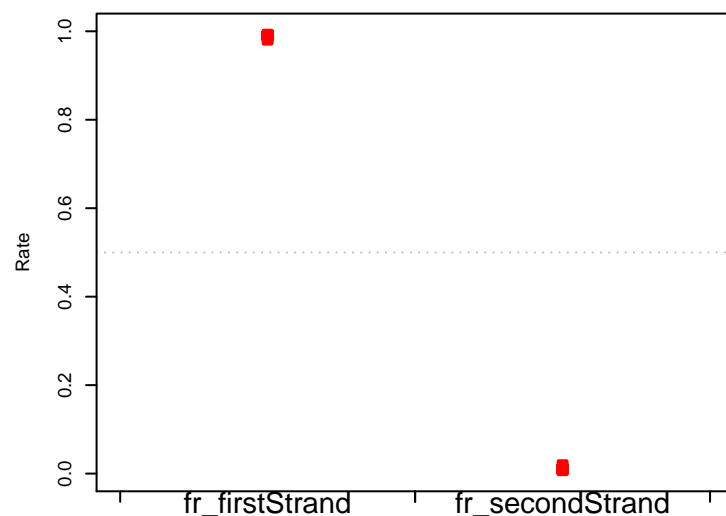
ai

Normalization Factors vs Total-Count Normalization
Colored by Lane



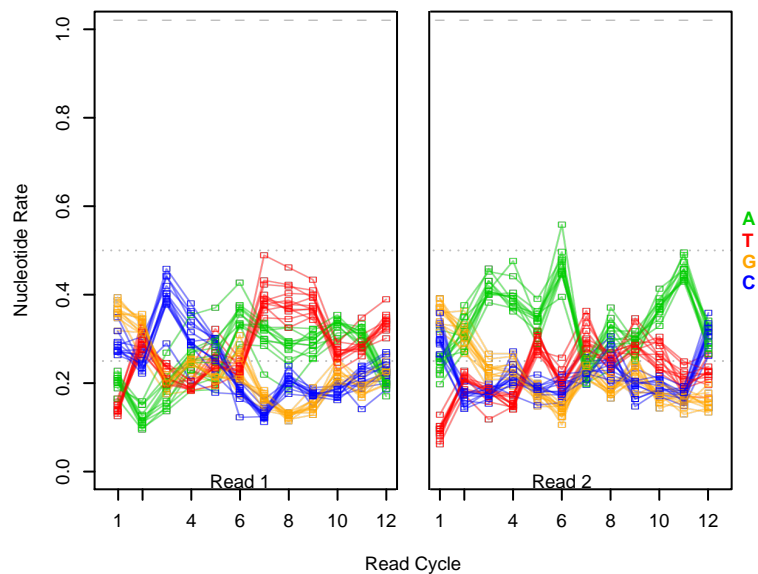
aj

Strandedness Test
Colored by Lane



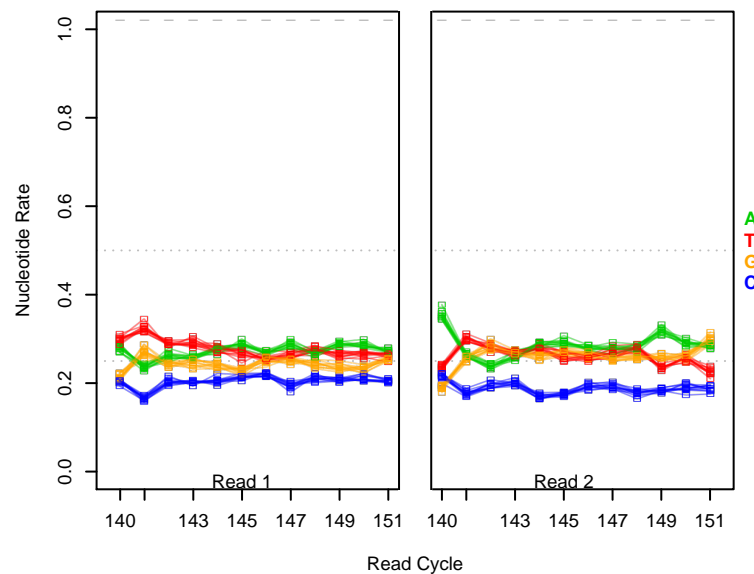
ak

Nucleotide Rate by Cycle, Leading Clipped bases (12)
Marked by Lane



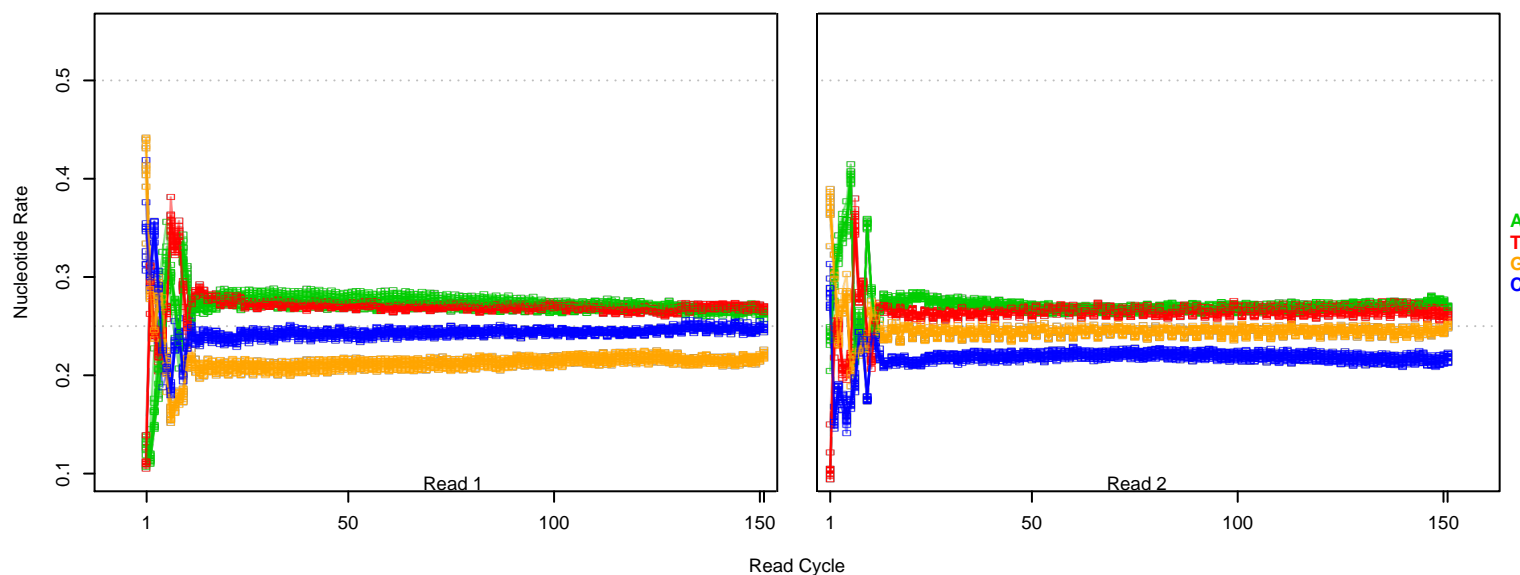
al

Nucleotide Rate by Cycle, Trailing Clipped bases (12)
Marked by Lane



am

Raw Nucleotide Rate by Cycle
Marked by Lane



an

Nucleotide Rate by Cycle, Aligned bases only
Marked by Lane

