

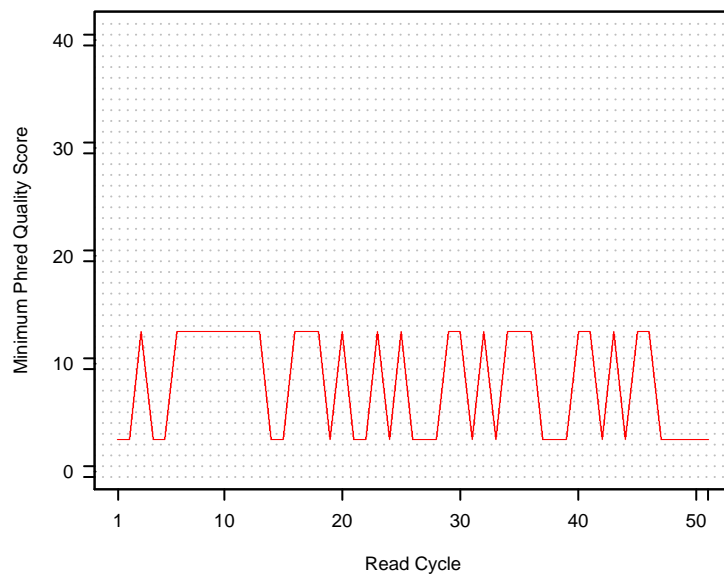
a

Sample Highlight: Untitled Colored by Lane



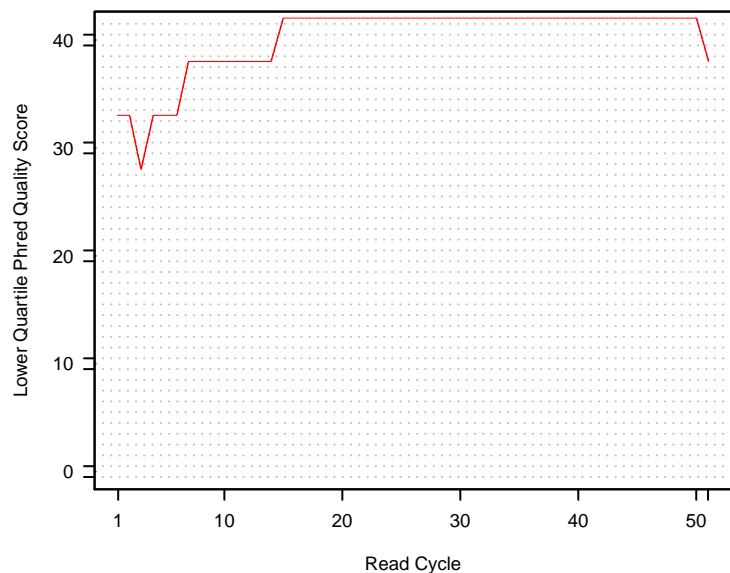
b

Minimum Phred Quality Score
With Sample Untitled Colored by lane



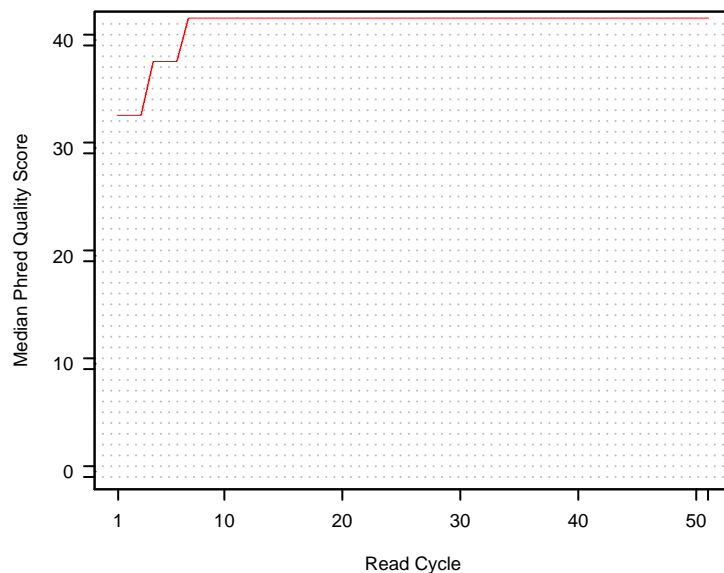
c

Lower Quartile Phred Quality Score
With Sample Untitled Colored by lane



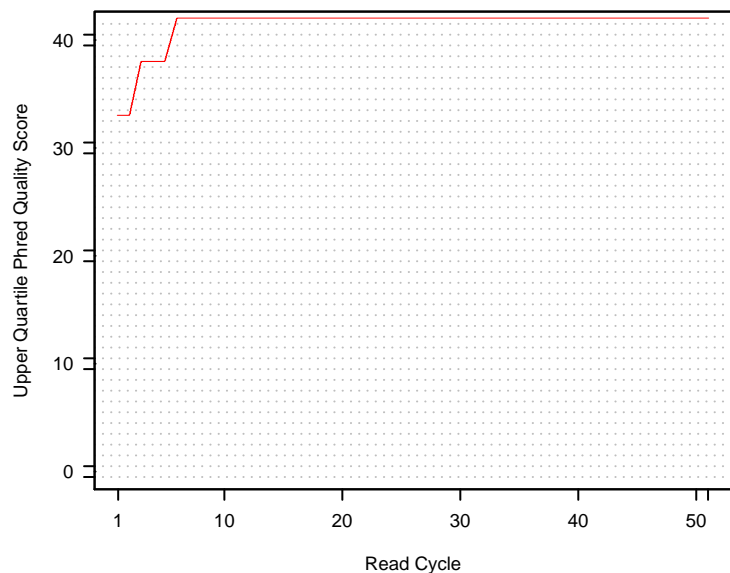
d

Median Phred Quality Score
With Sample Untitled Colored by lane



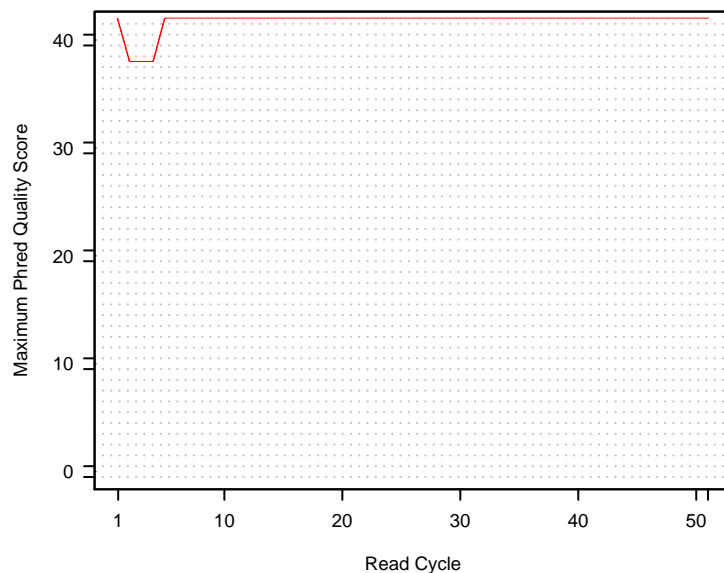
e

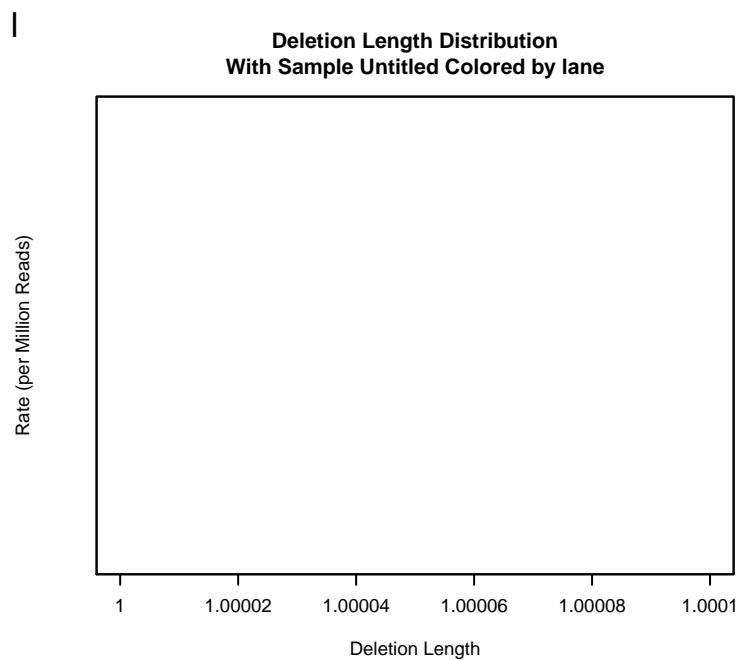
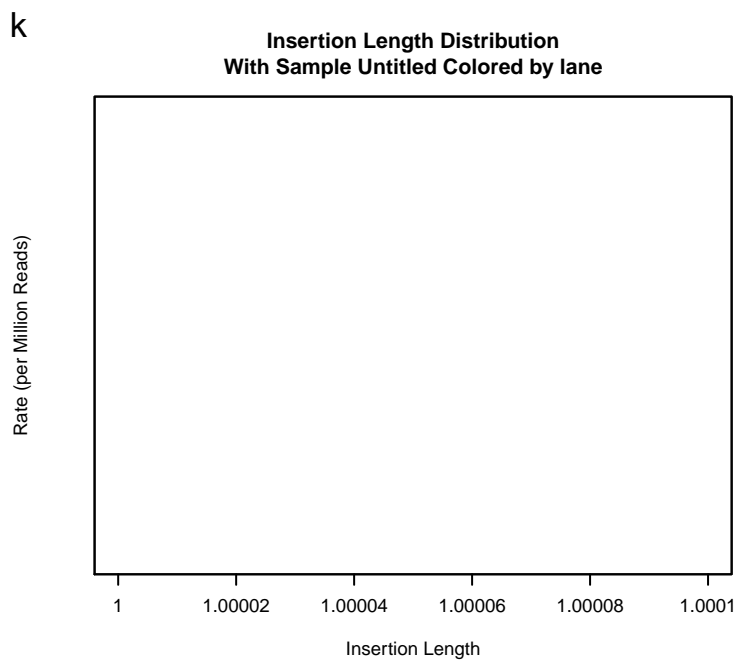
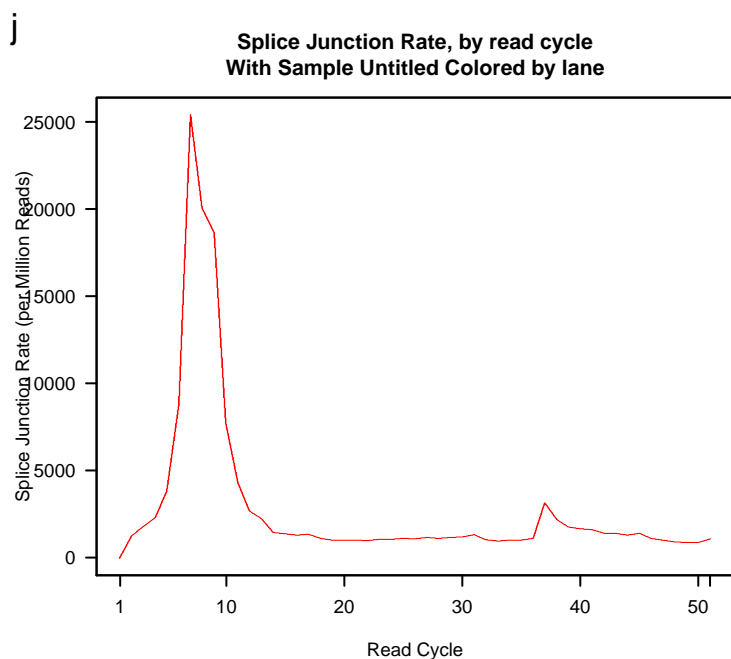
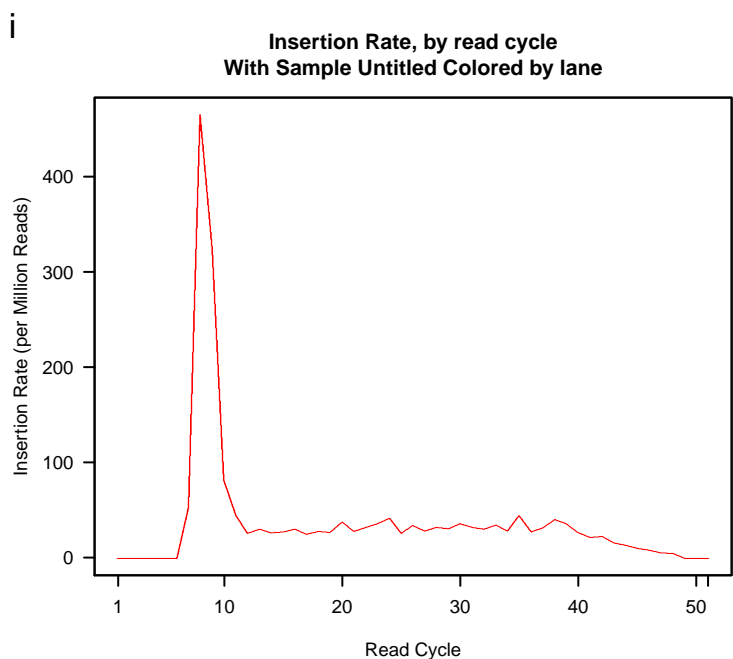
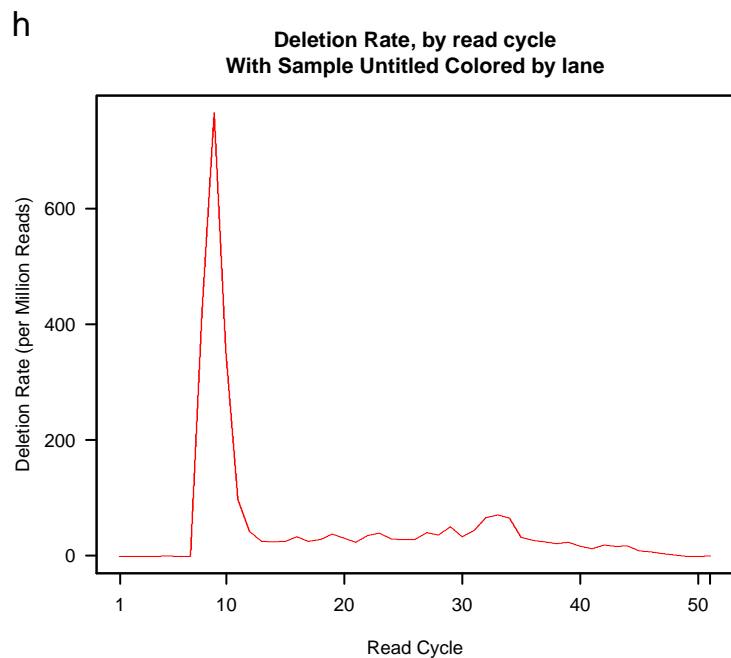
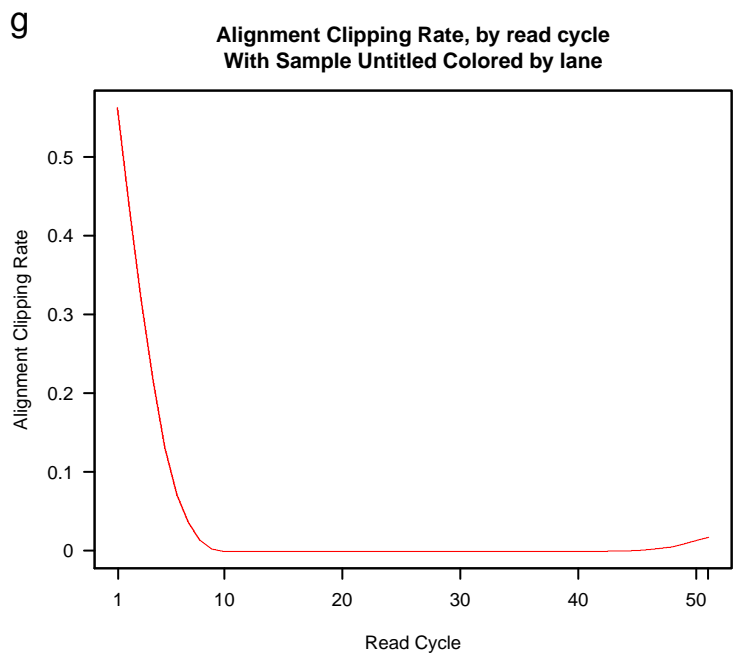
Upper Quartile Phred Quality Score
With Sample Untitled Colored by lane

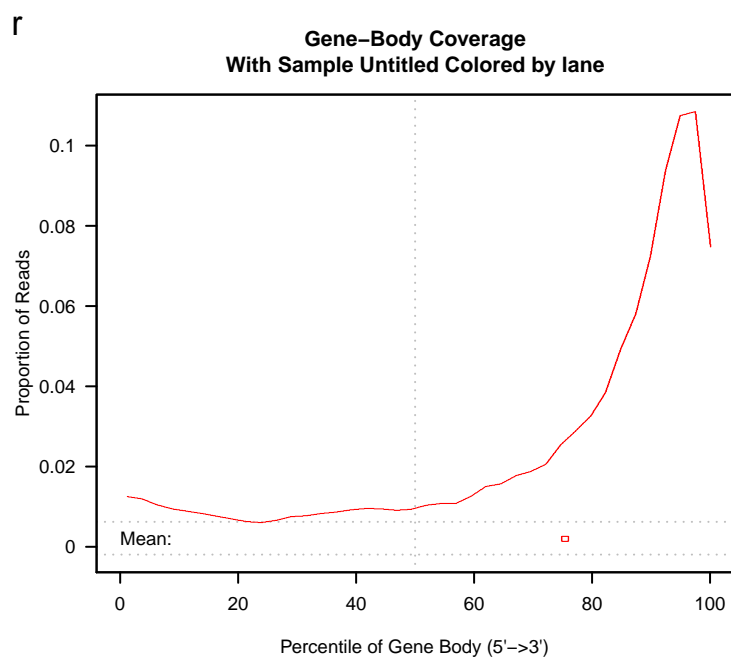
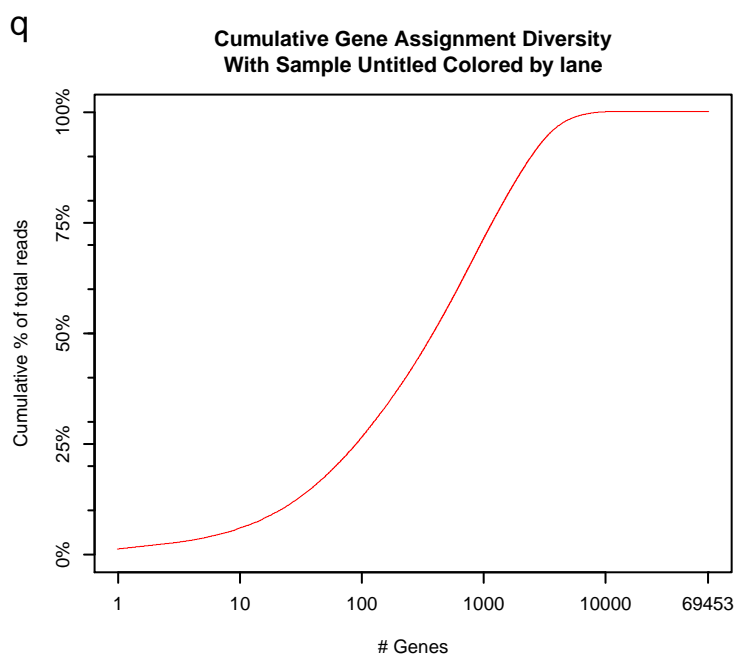
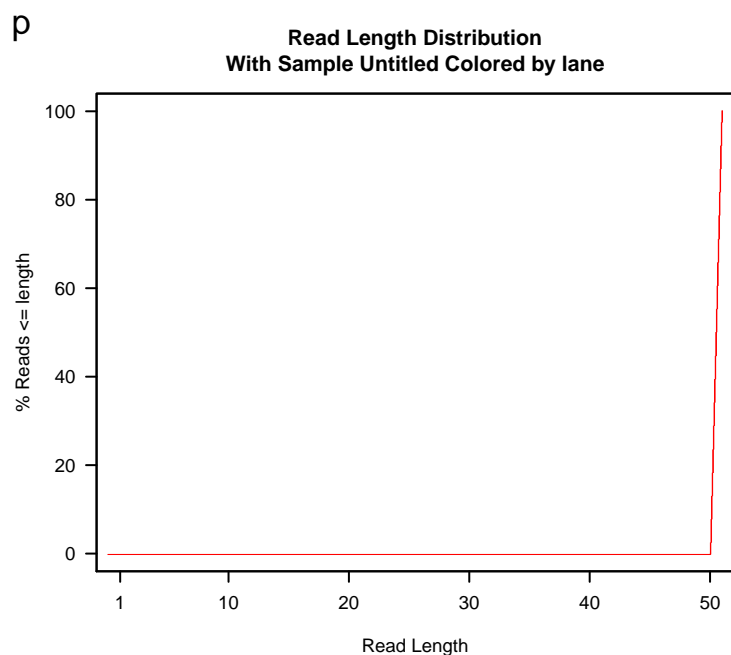
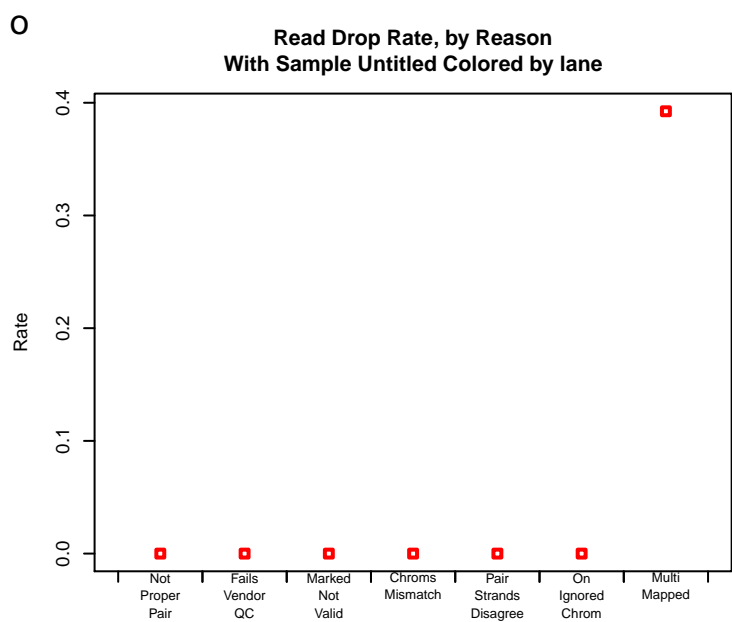
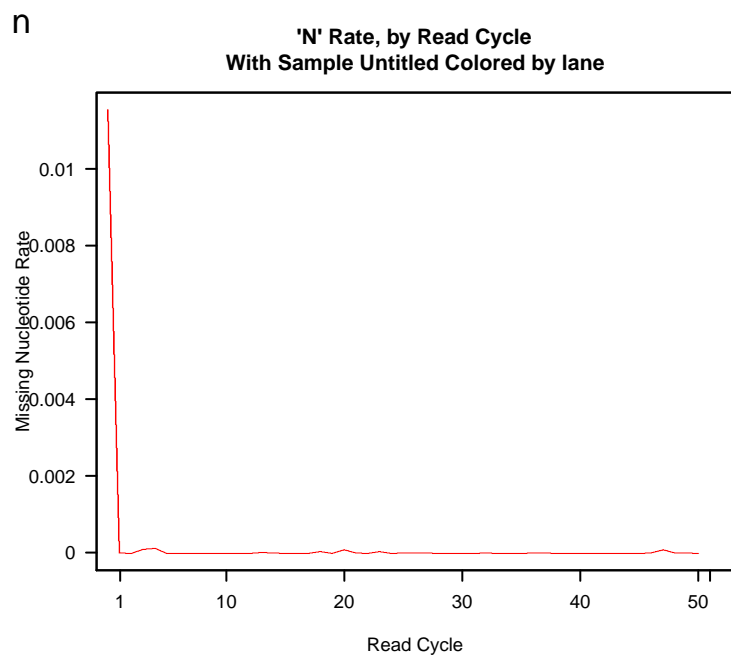
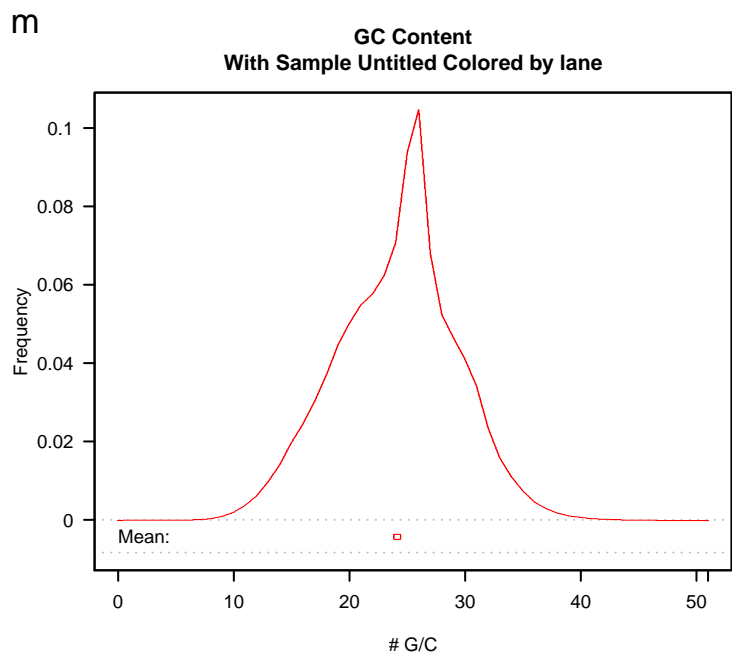


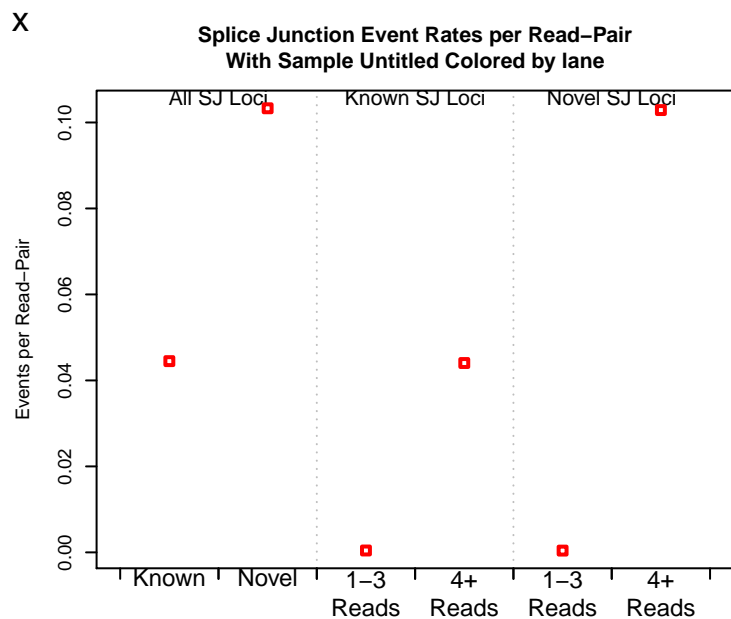
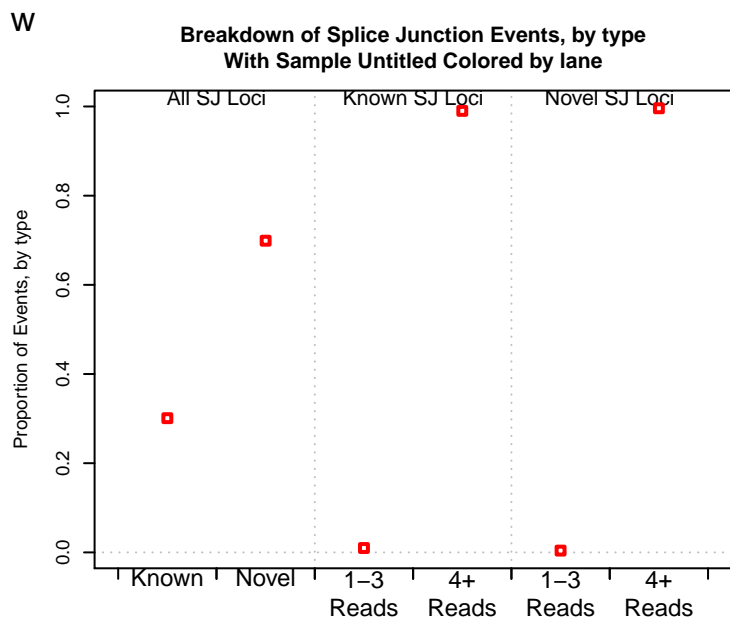
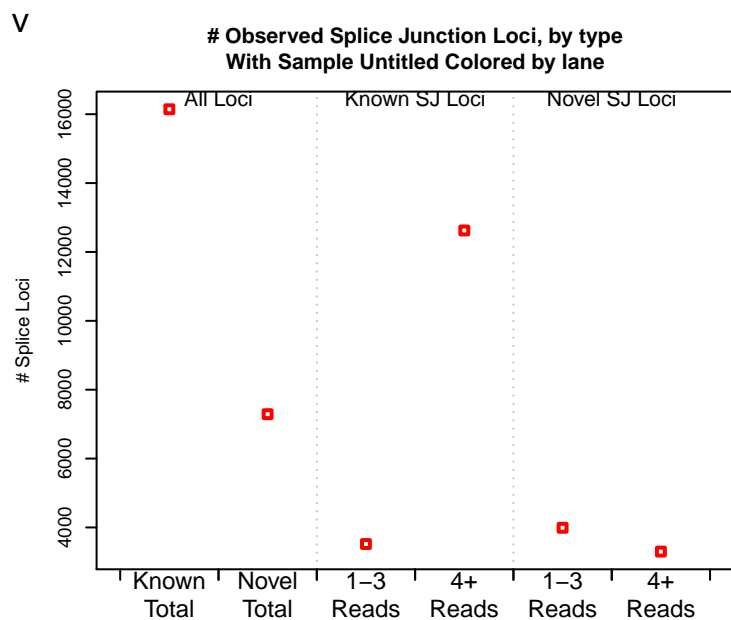
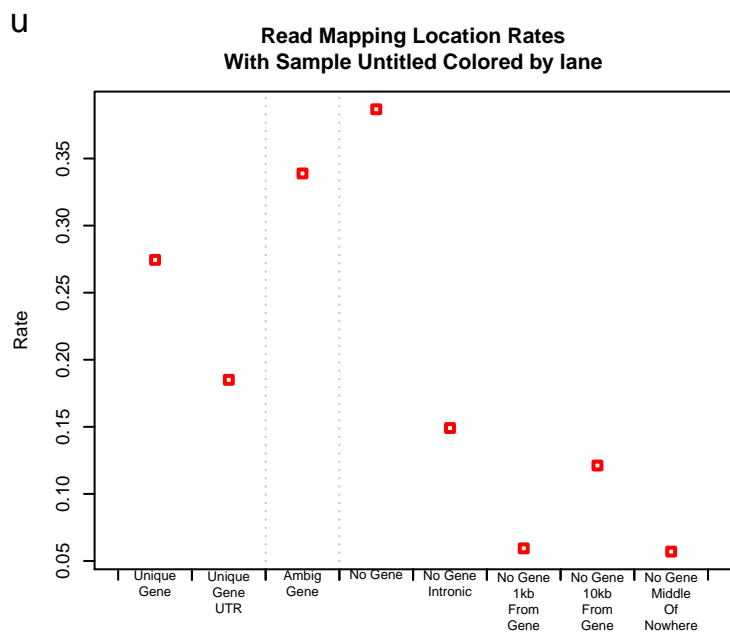
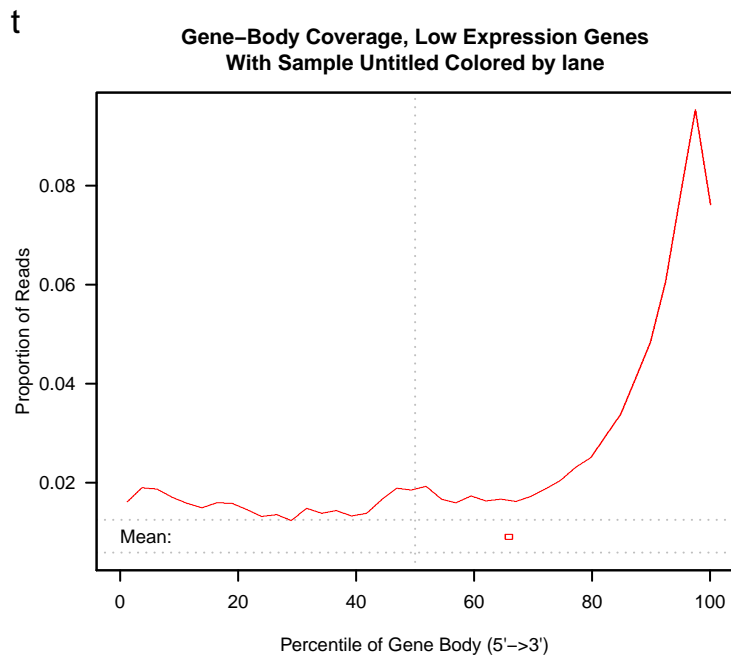
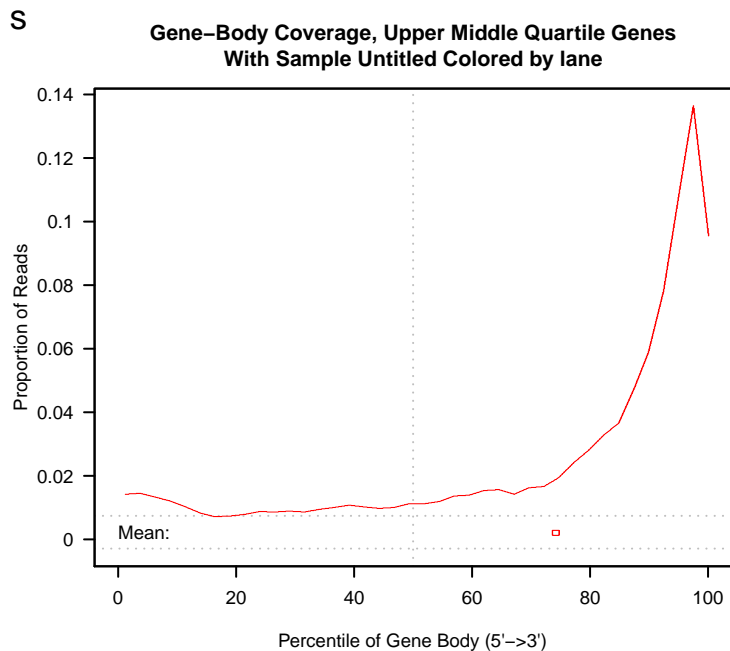
f

Maximum Phred Quality Score
With Sample Untitled Colored by lane

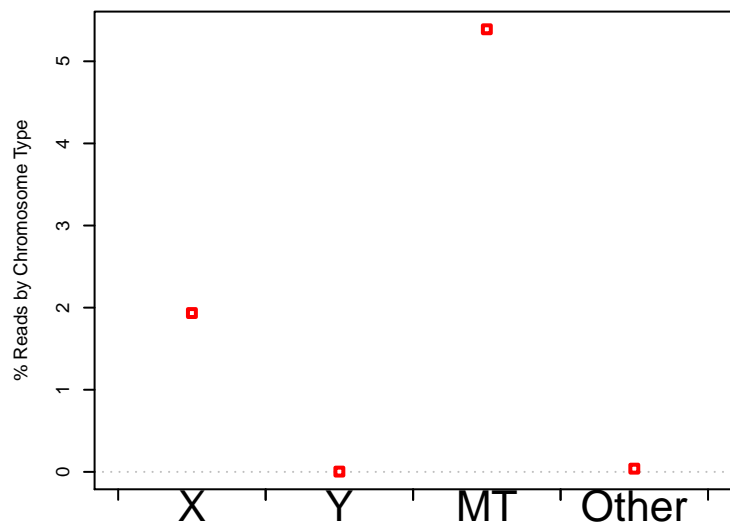




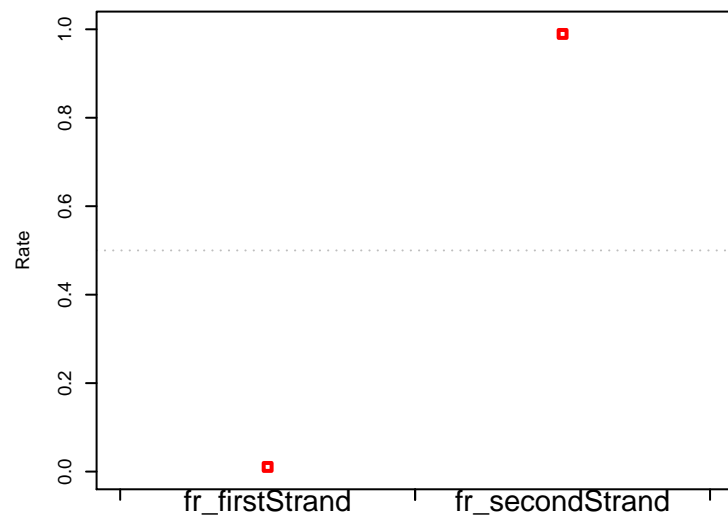




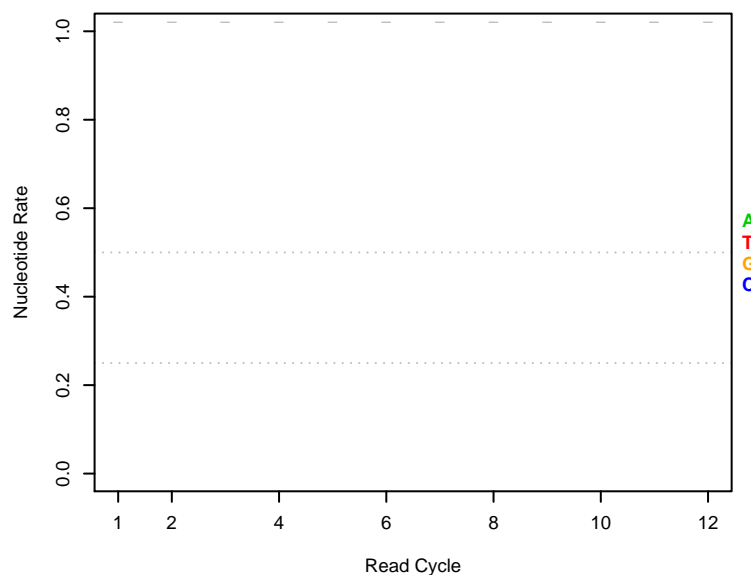
y **Chromosome Distribution (Excluding Autosomes)**
With Sample Untitled Colored by lane



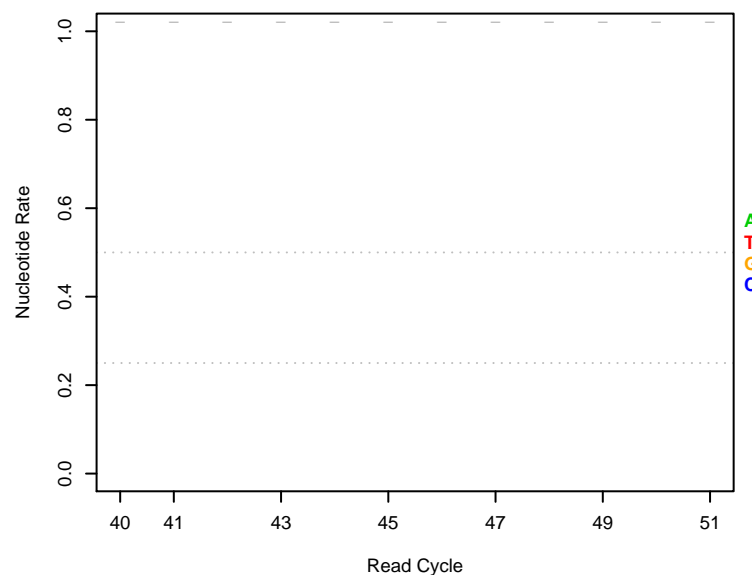
Z **Strandedness Test**
With Sample Untitled Colored by lane



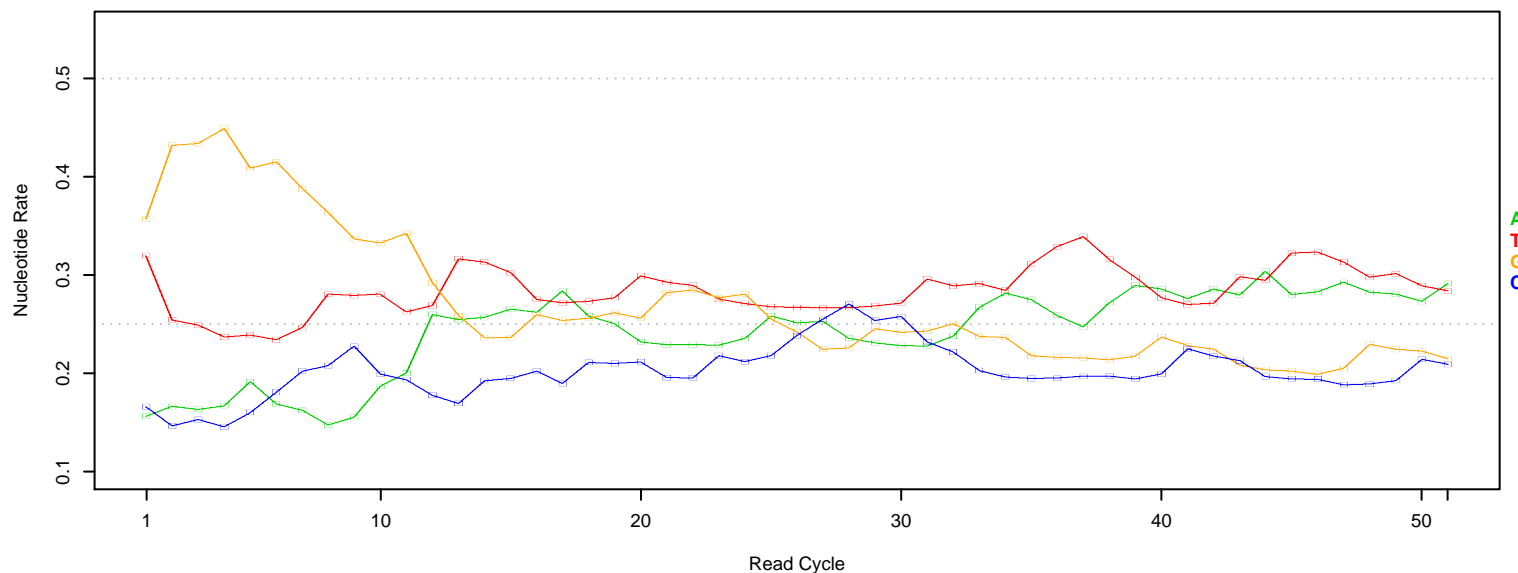
aa **Nucleotide Rate by Cycle, Leading Clipped bases (12)**
With Sample Untitled in Bold



ab **Nucleotide Rate by Cycle, Trailing Clipped bases (12)**
With Sample Untitled in Bold



ac **Raw Nucleotide Rate by Cycle**
With Sample Untitled in Bold



ad

Nucleotide Rate by Cycle, Aligned bases only
With Sample Untitled in Bold

