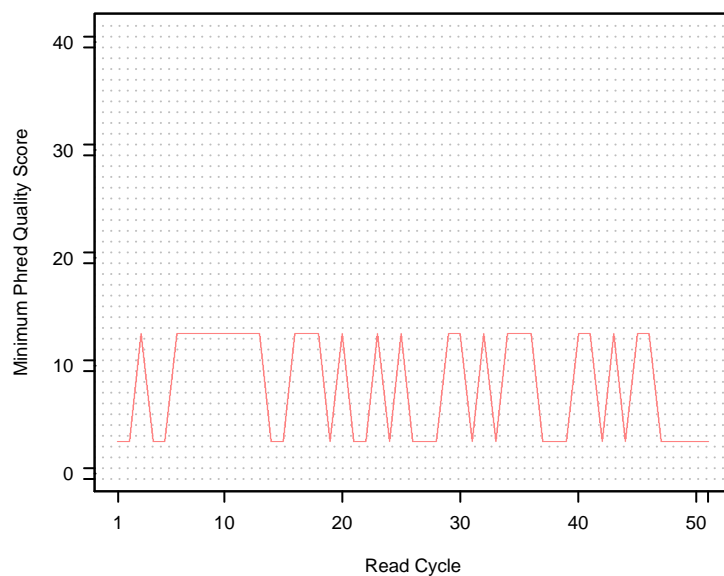


a

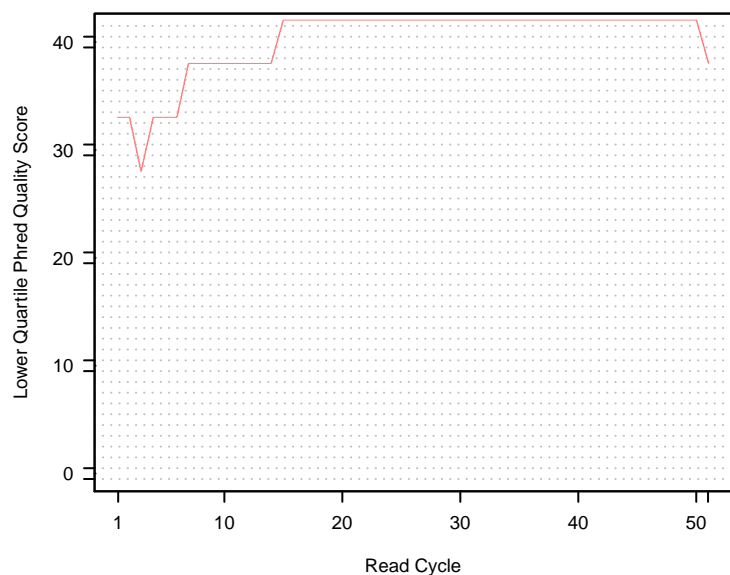
Summary Plots, By Lane



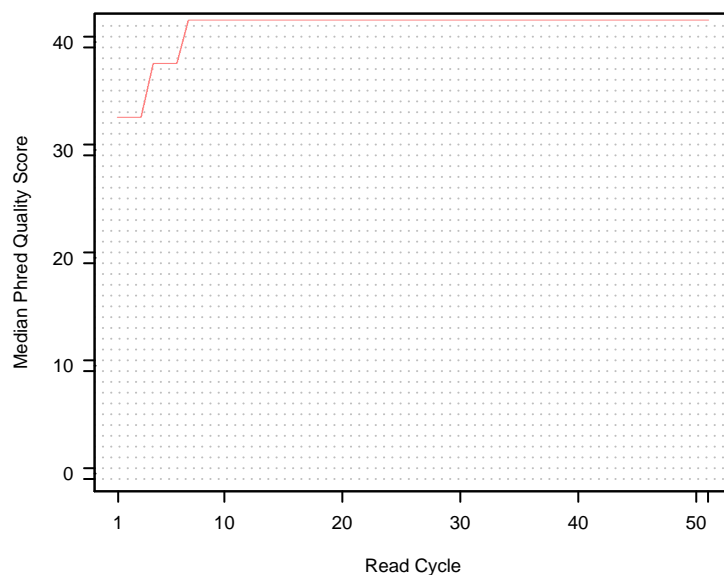
b

Minimum Phred Quality Score
Colored by Lane

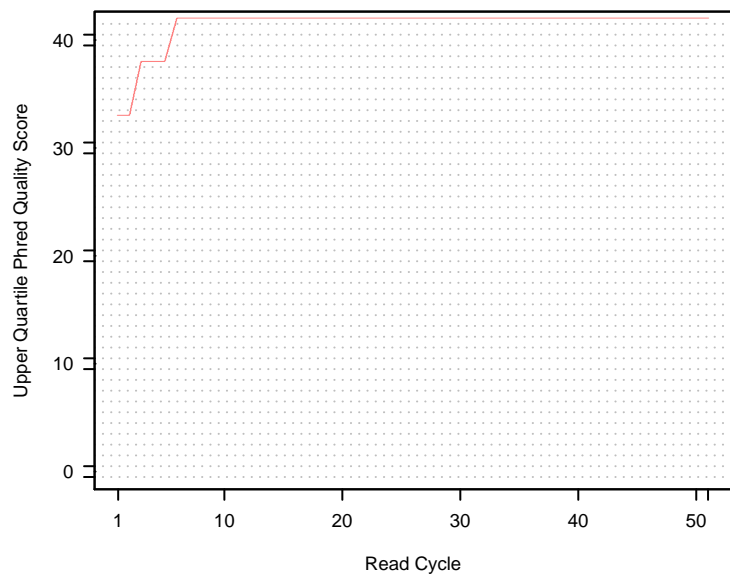
c

Lower Quartile Phred Quality Score
Colored by Lane

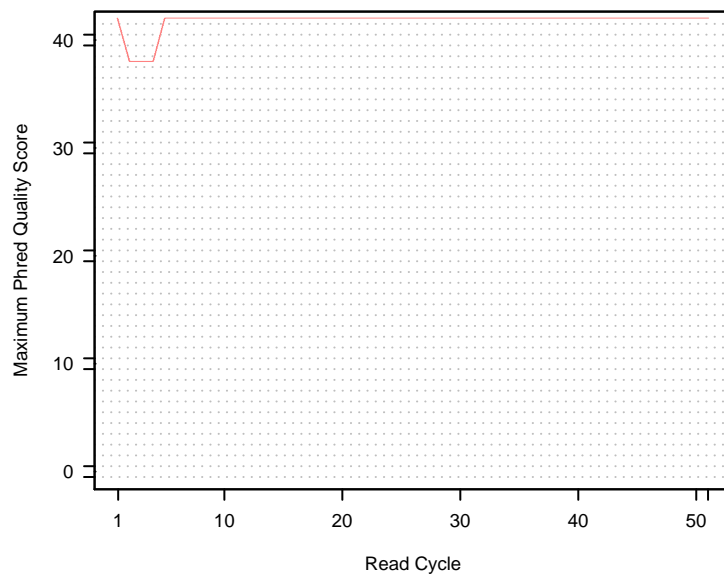
d

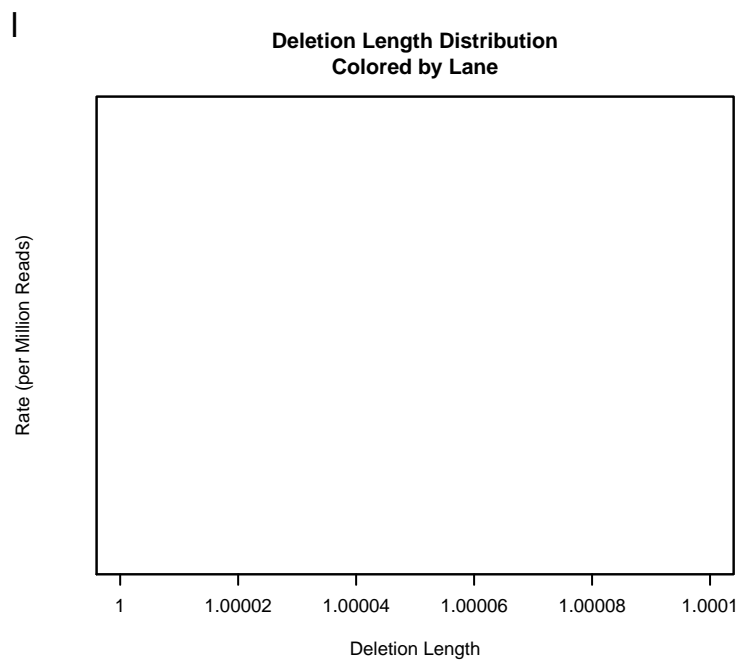
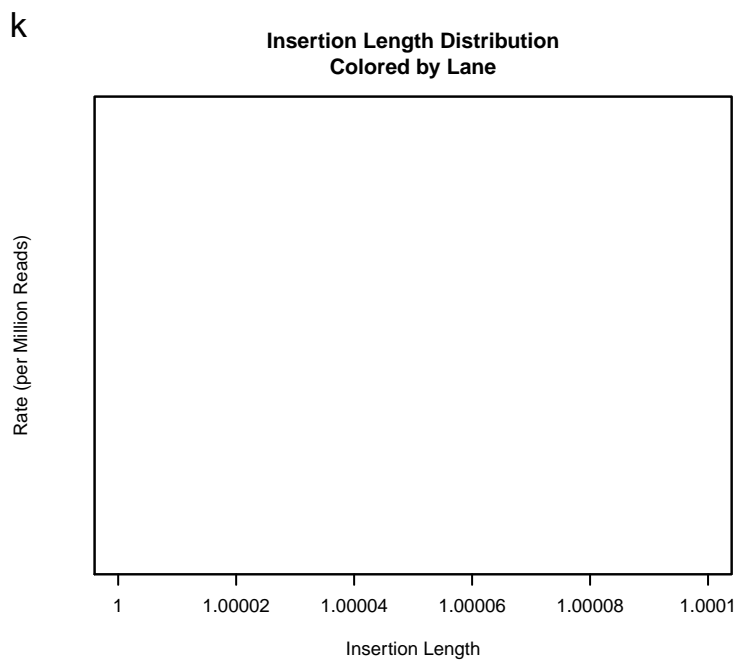
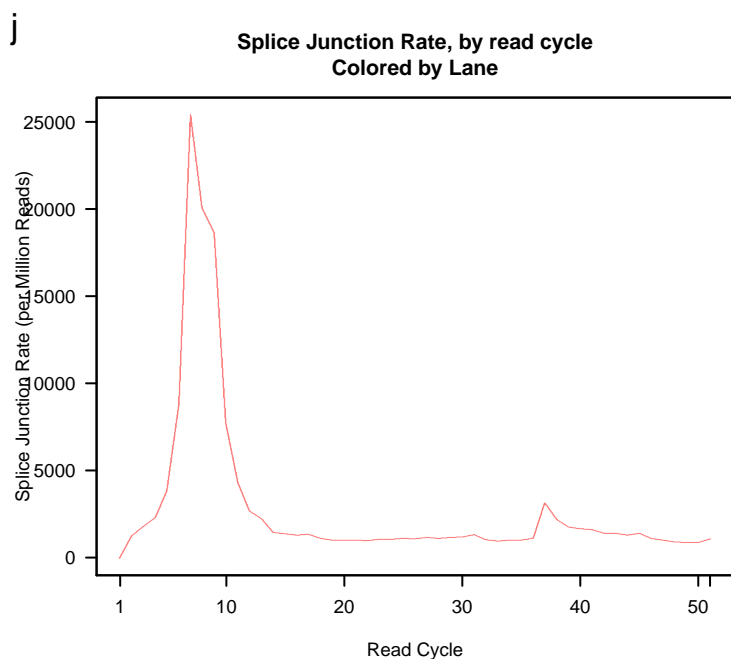
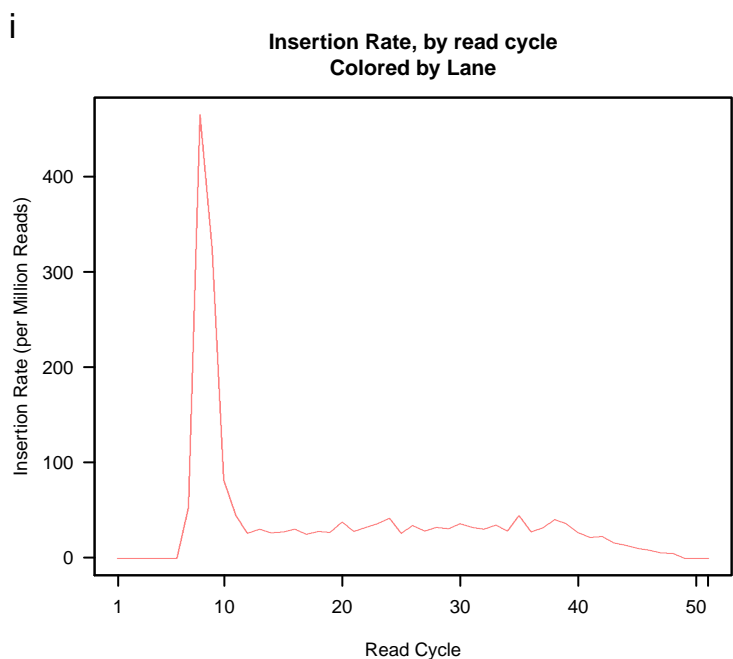
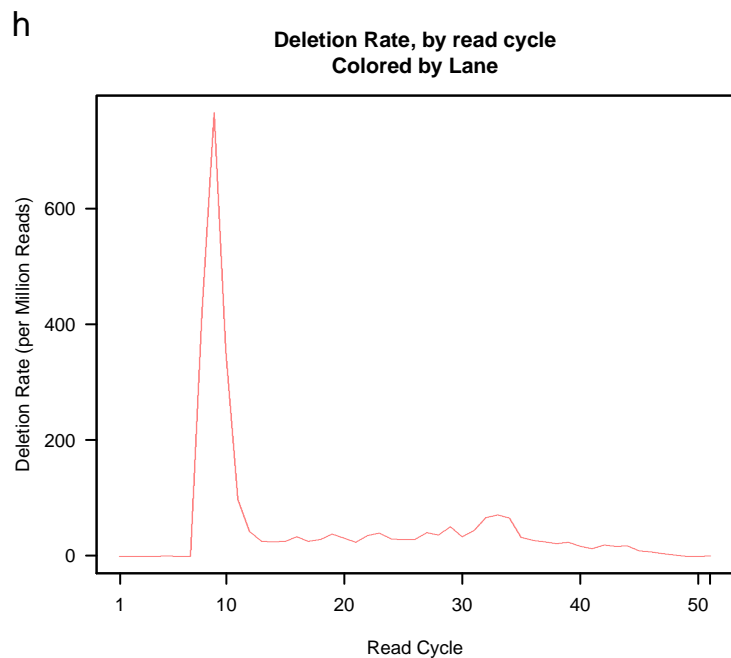
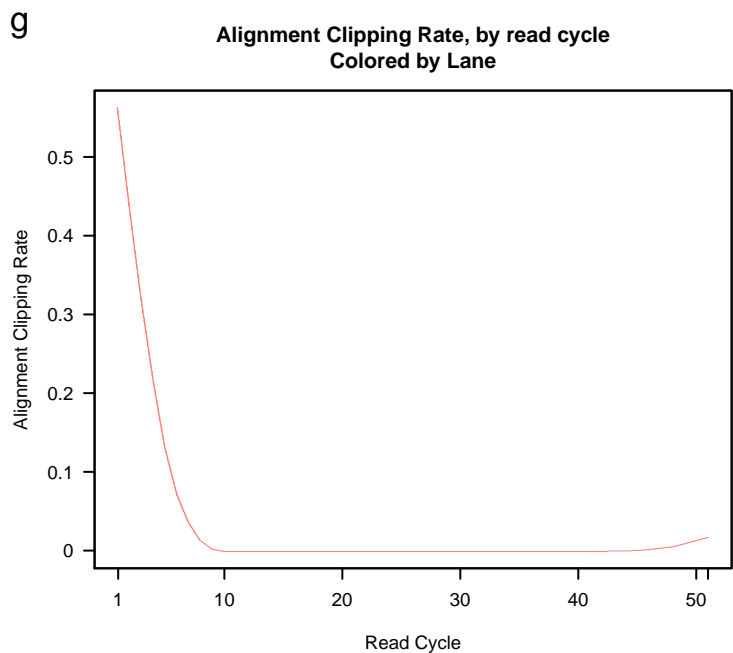
Median Phred Quality Score
Colored by Lane

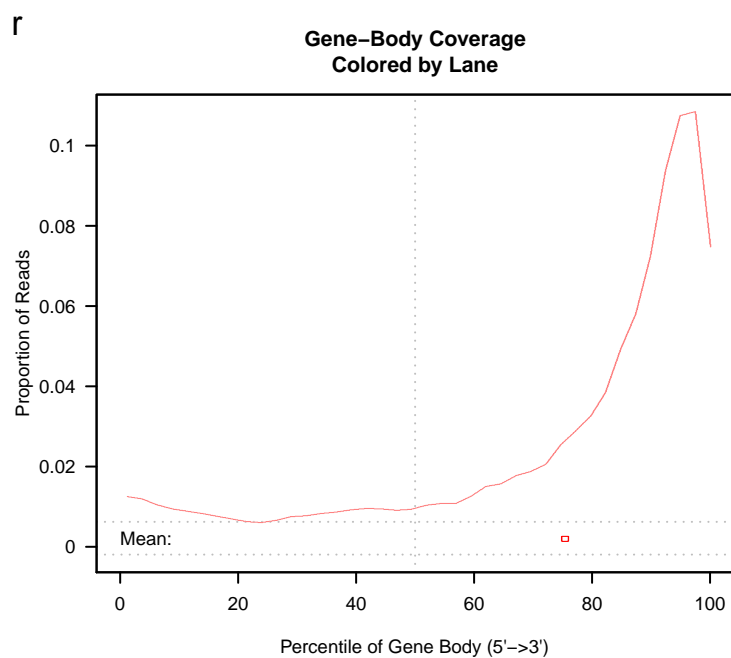
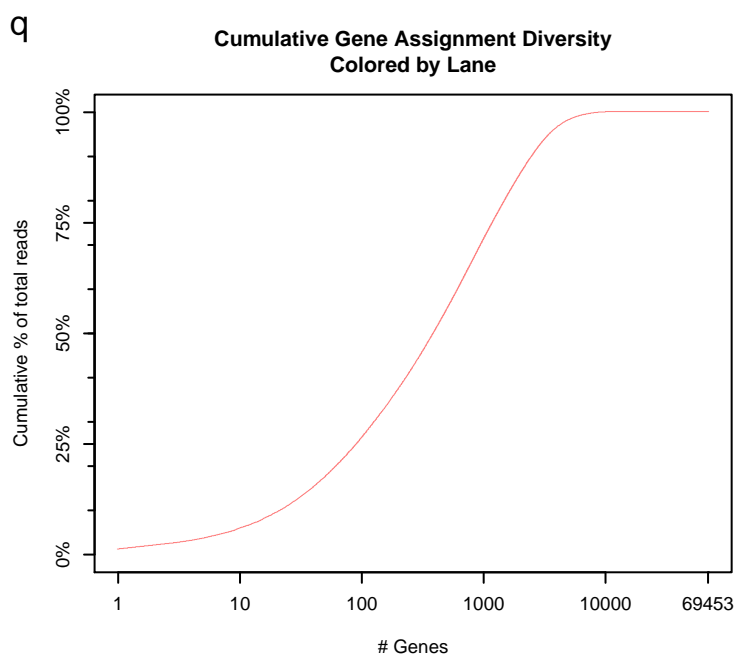
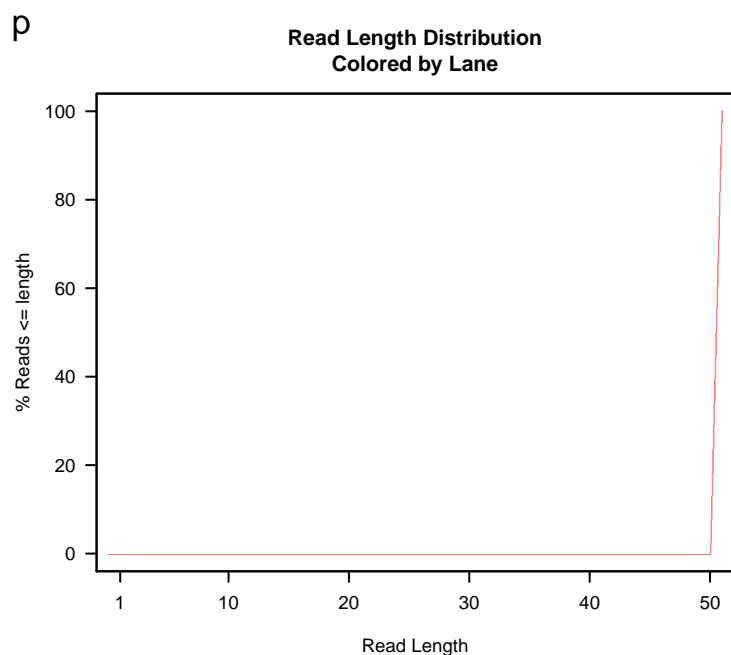
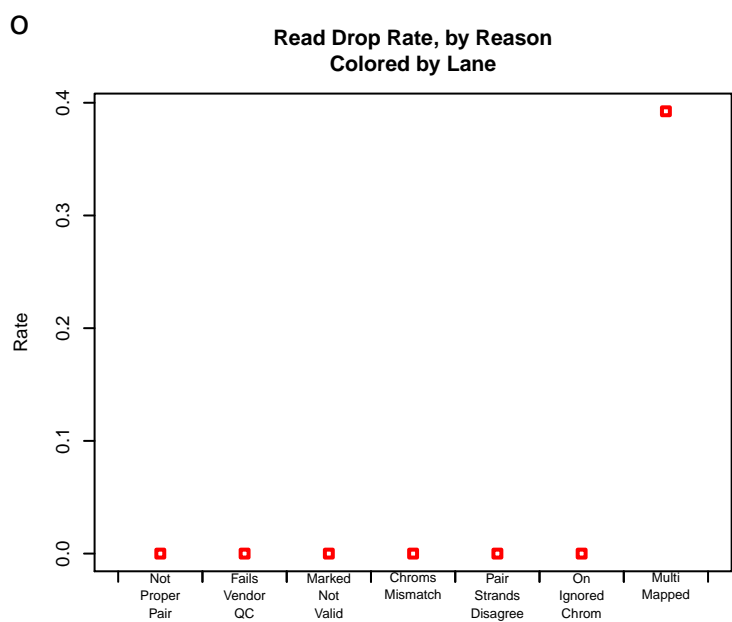
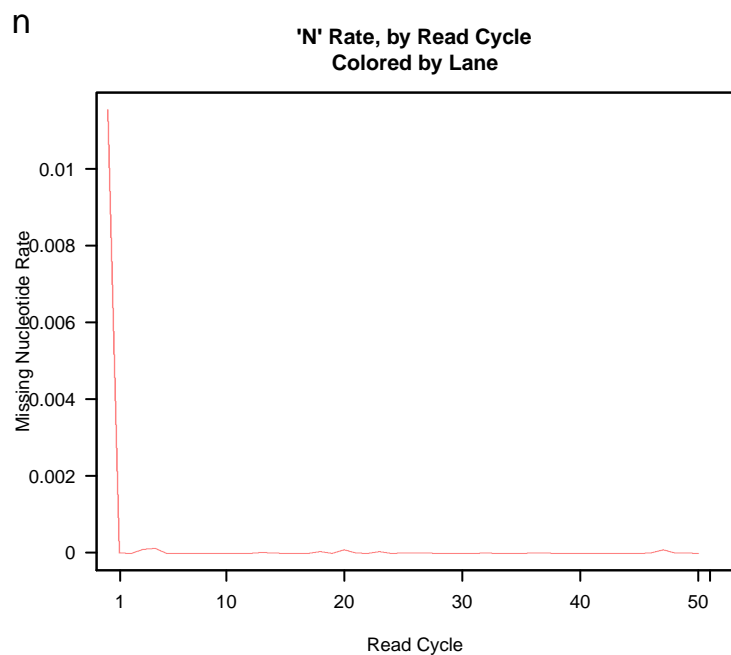
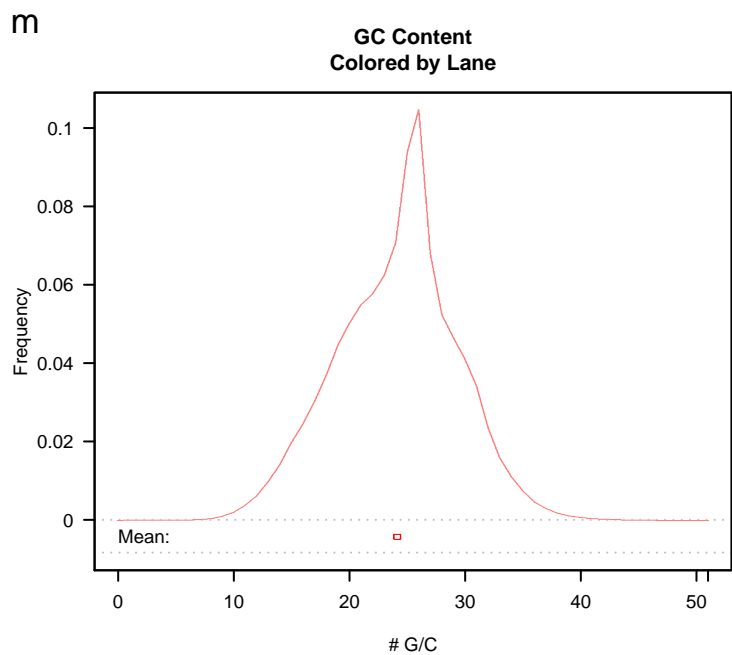
e

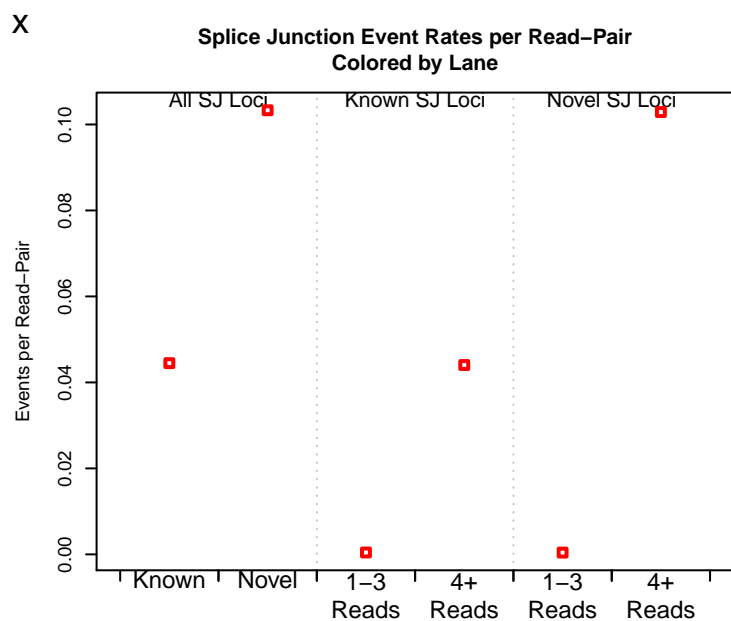
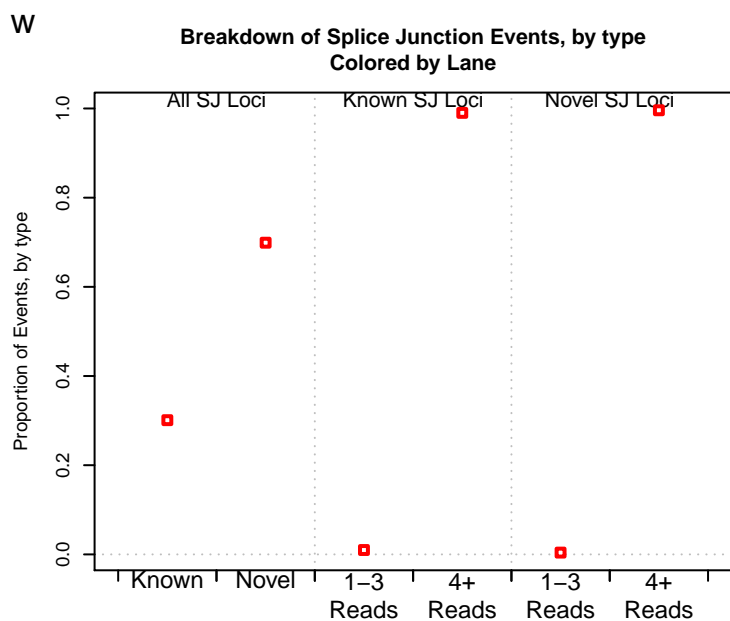
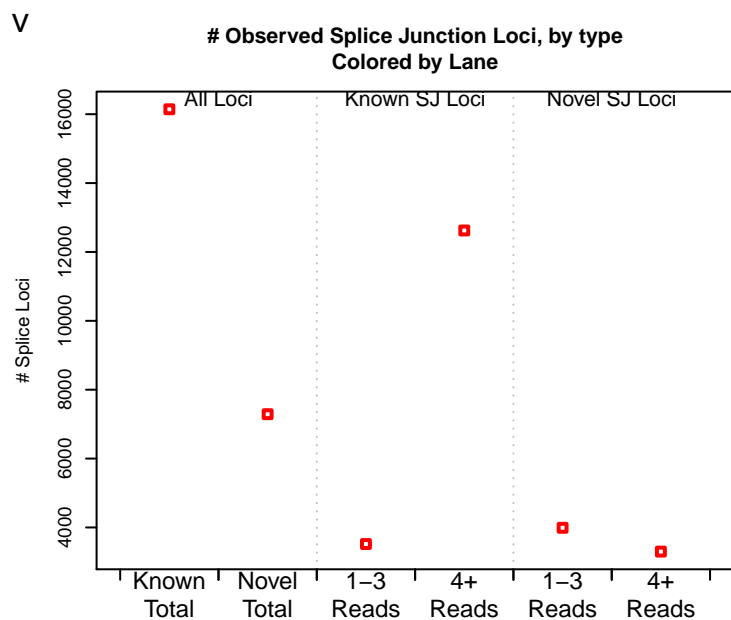
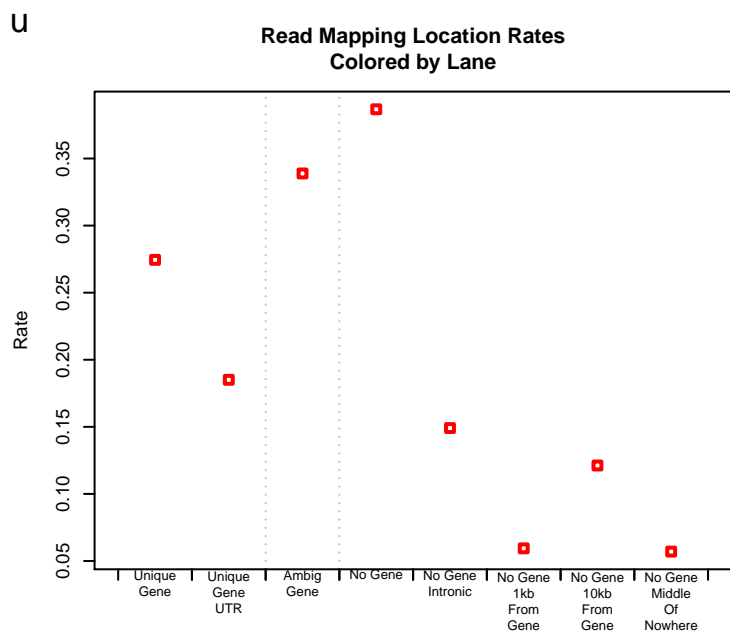
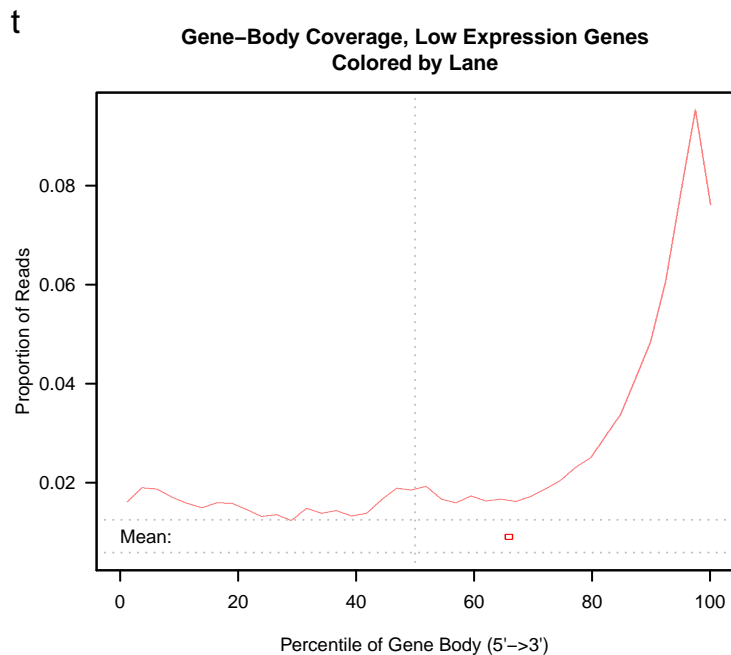
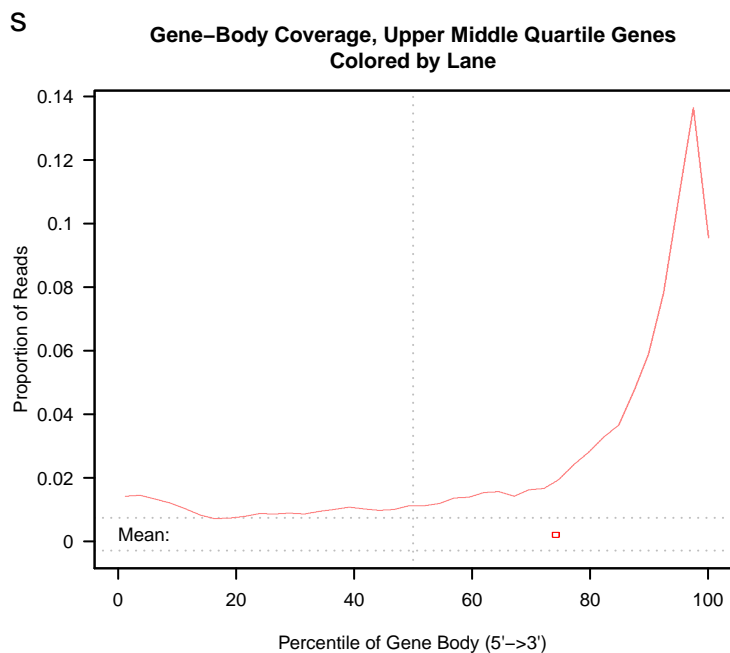
Upper Quartile Phred Quality Score
Colored by Lane

f

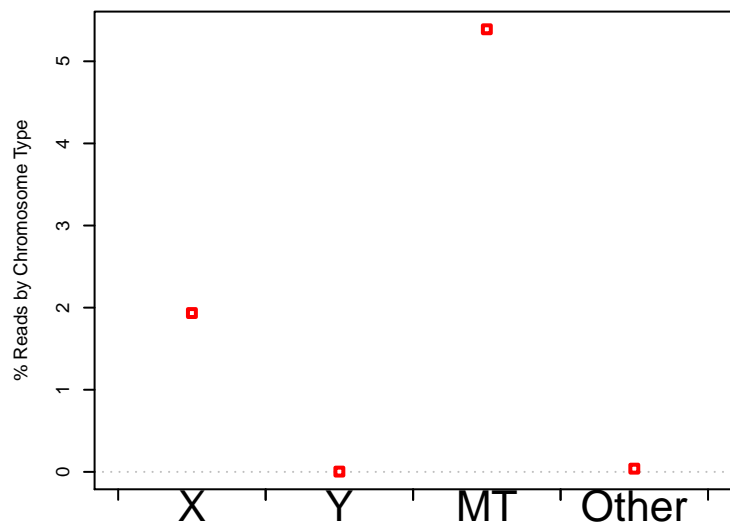
Maximum Phred Quality Score
Colored by Lane



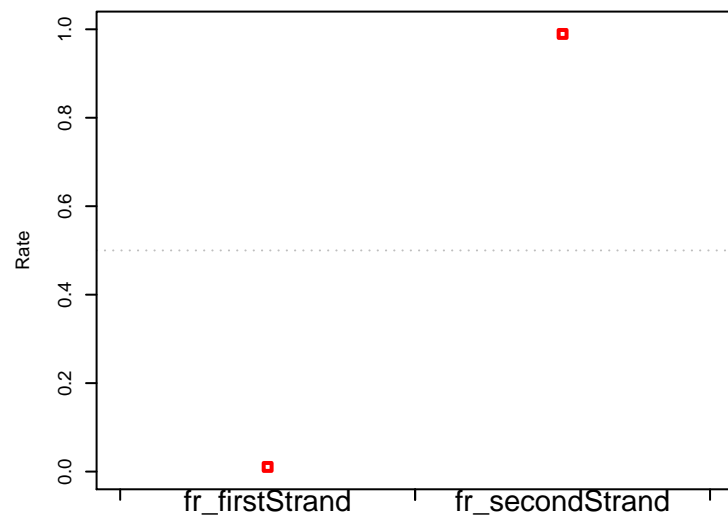




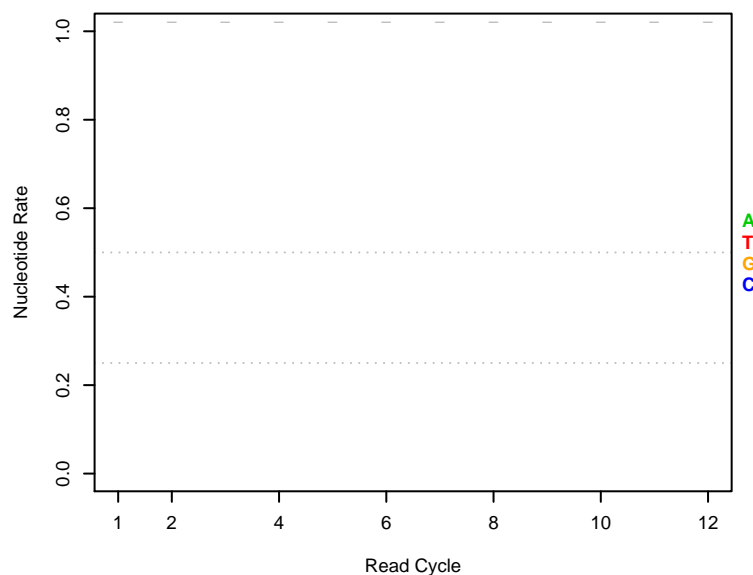
y **Chromosome Distribution (Excluding Autosomes)**
Colored by Lane



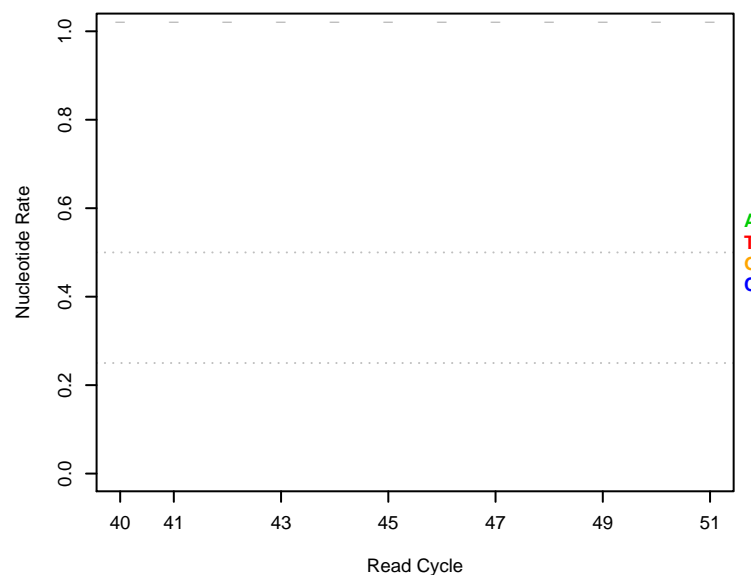
Z **Strandedness Test**
Colored by Lane



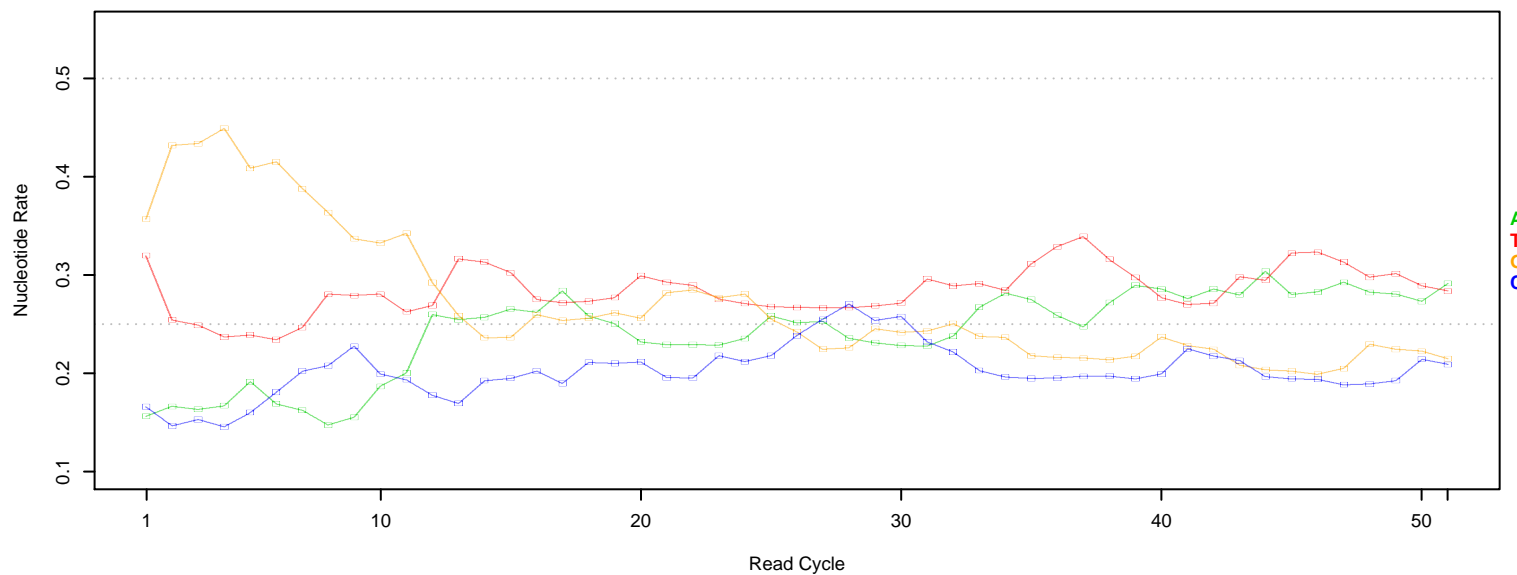
aa **Nucleotide Rate by Cycle, Leading Clipped bases (12)**
Marked by Lane



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



ac **Raw Nucleotide Rate by Cycle**
Marked by Lane



ad

Nucleotide Rate by Cycle, Aligned bases only
Marked by Lane

