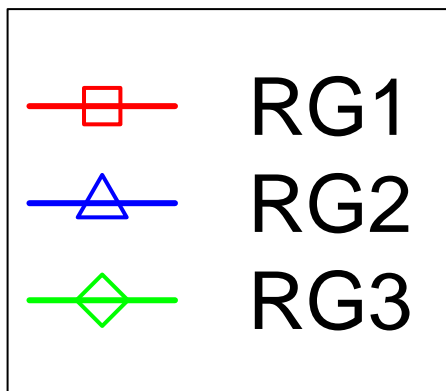
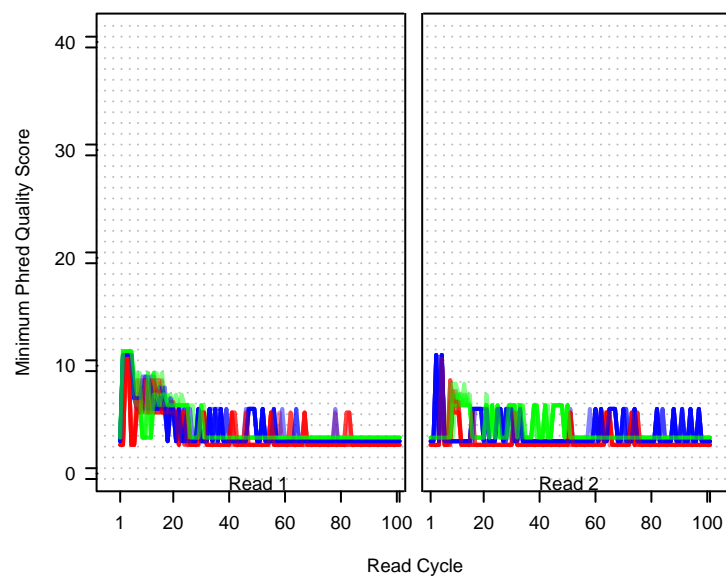


a

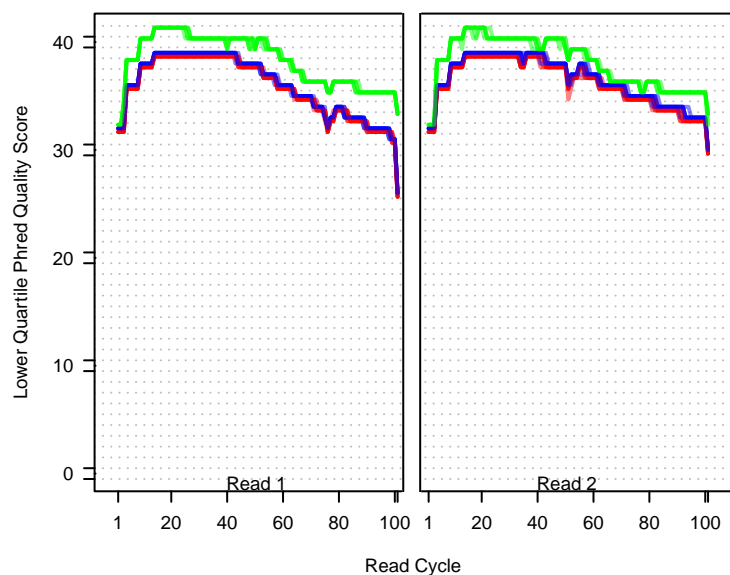
Summary Plots, By Lane



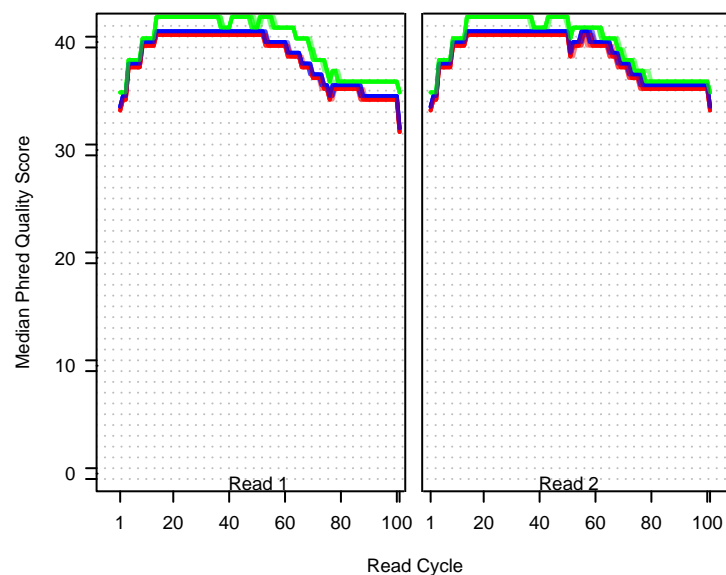
b

Minimum Phred Quality Score
Colored by Lane

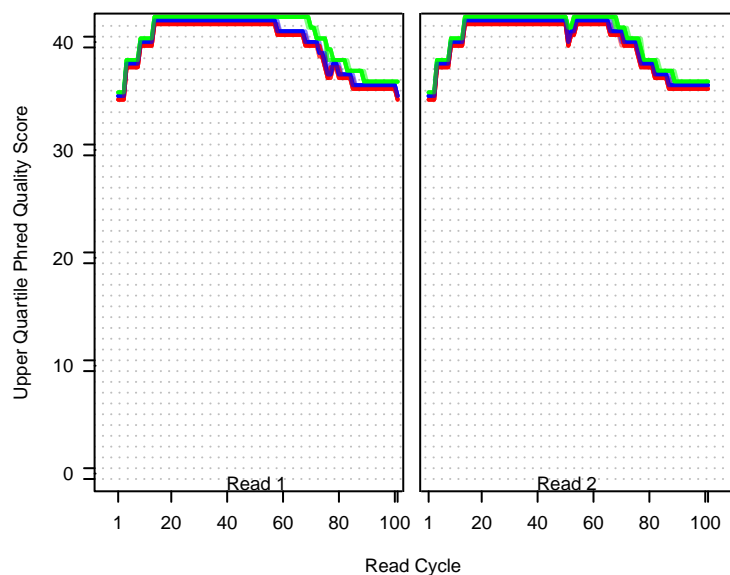
c

Lower Quartile Phred Quality Score
Colored by Lane

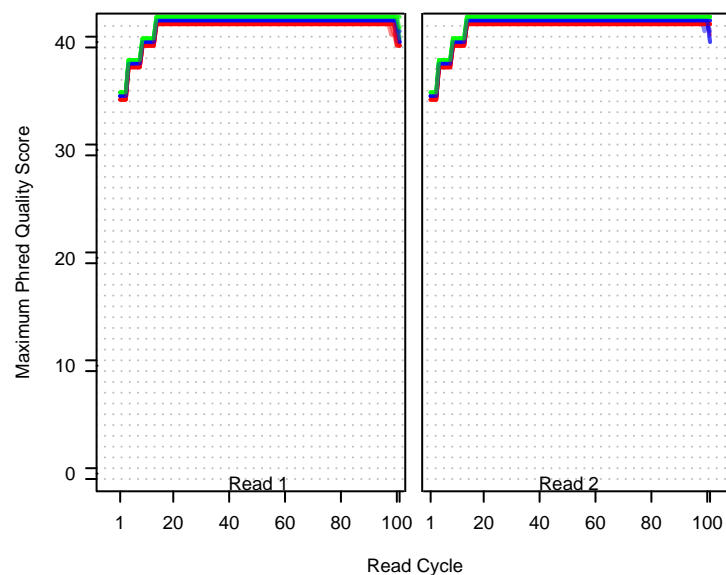
d

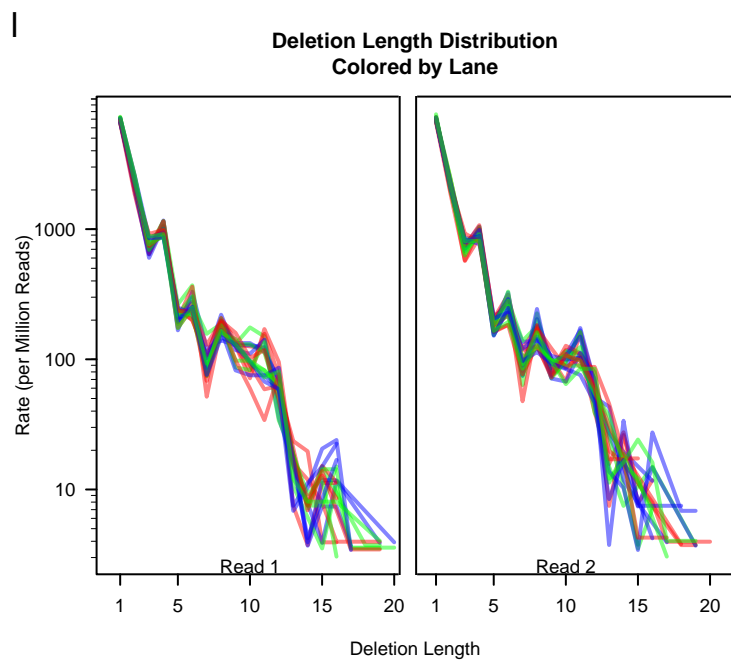
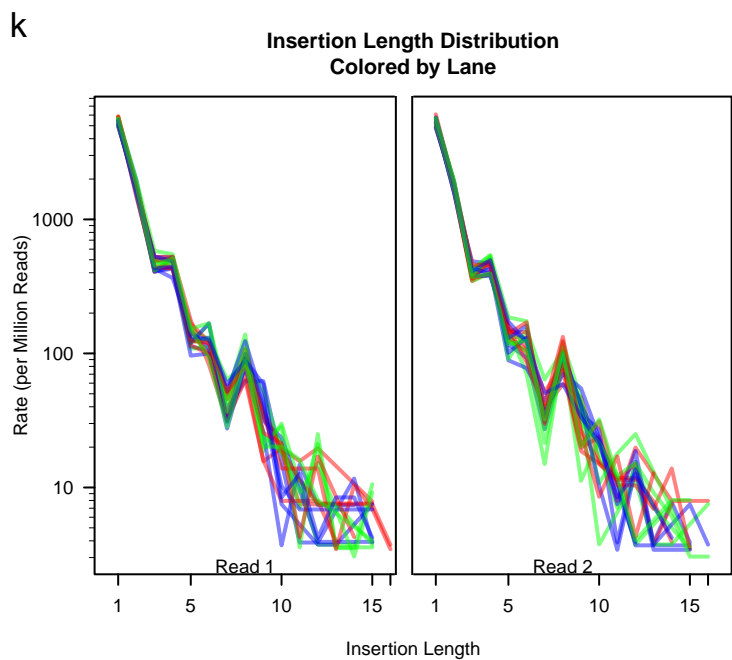
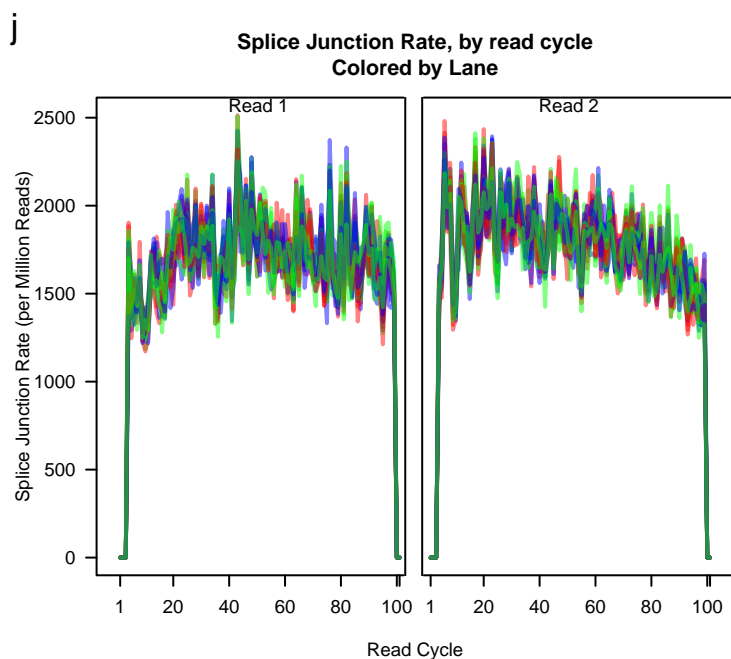
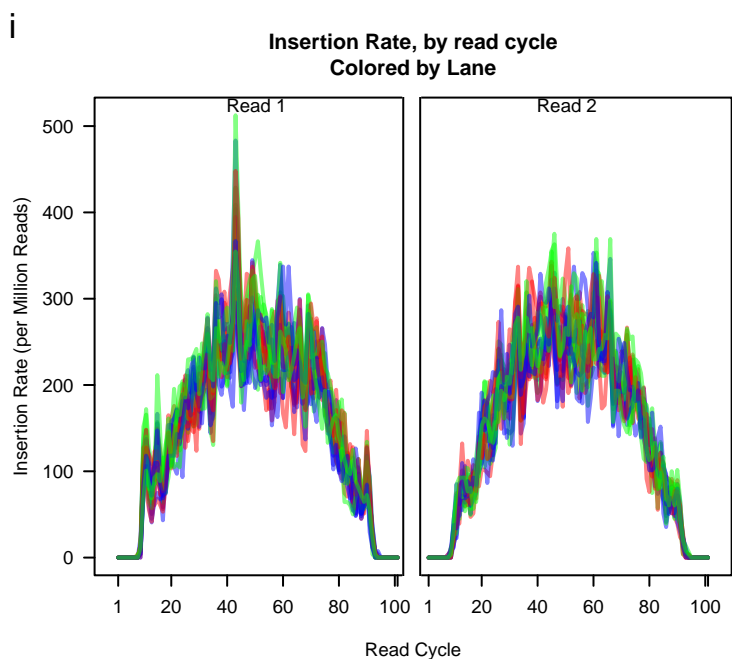
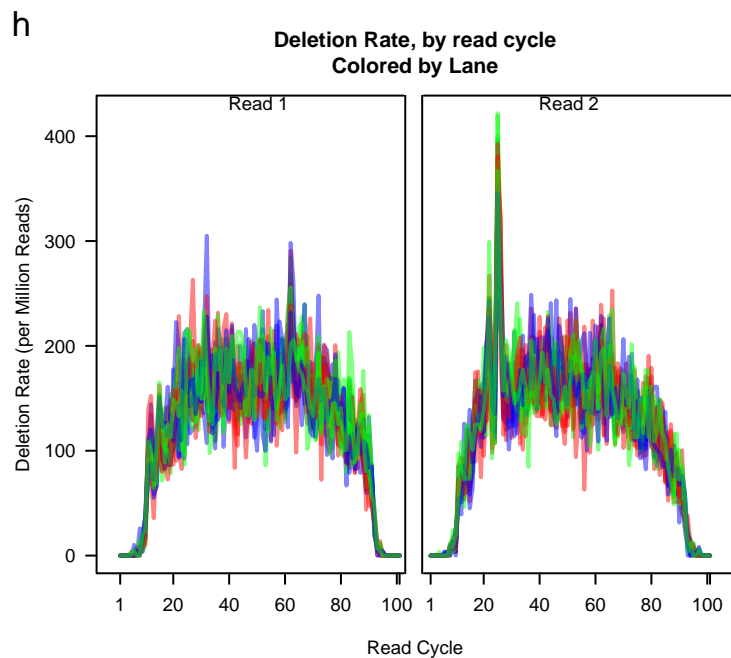
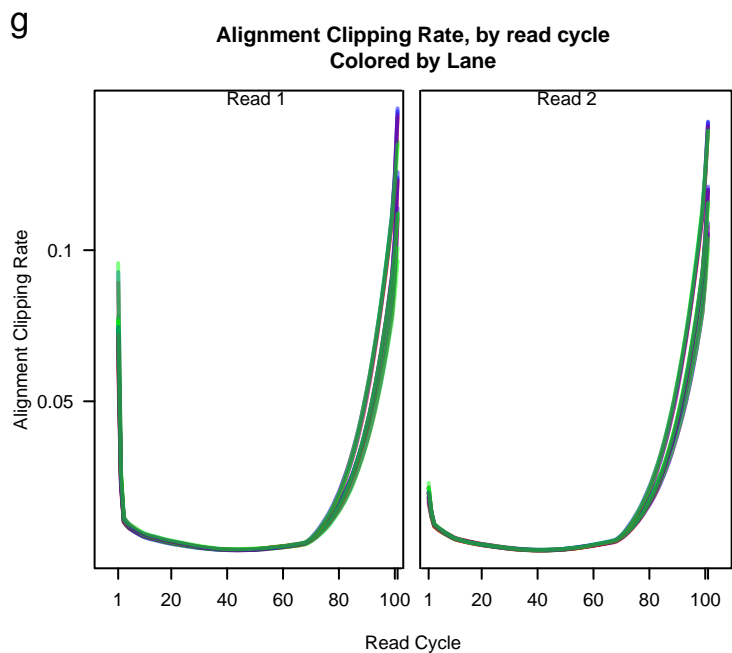
Median Phred Quality Score
Colored by Lane

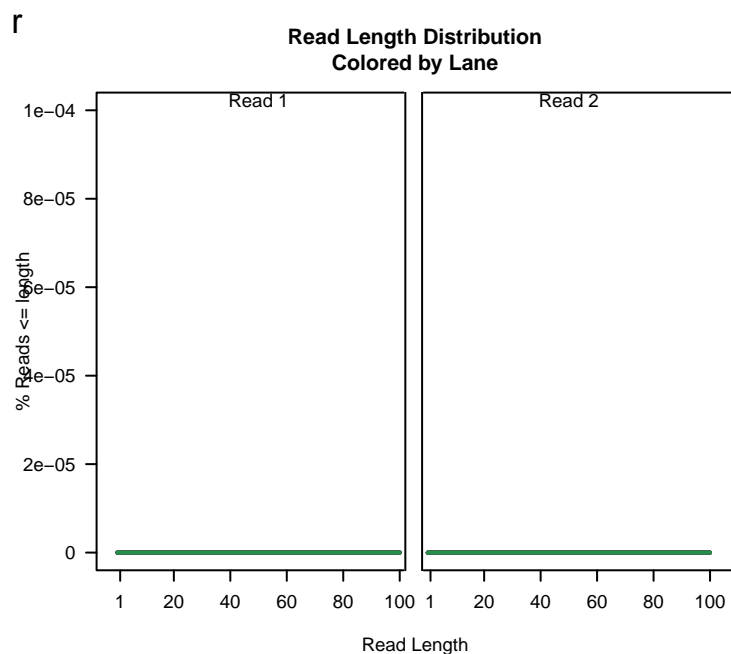
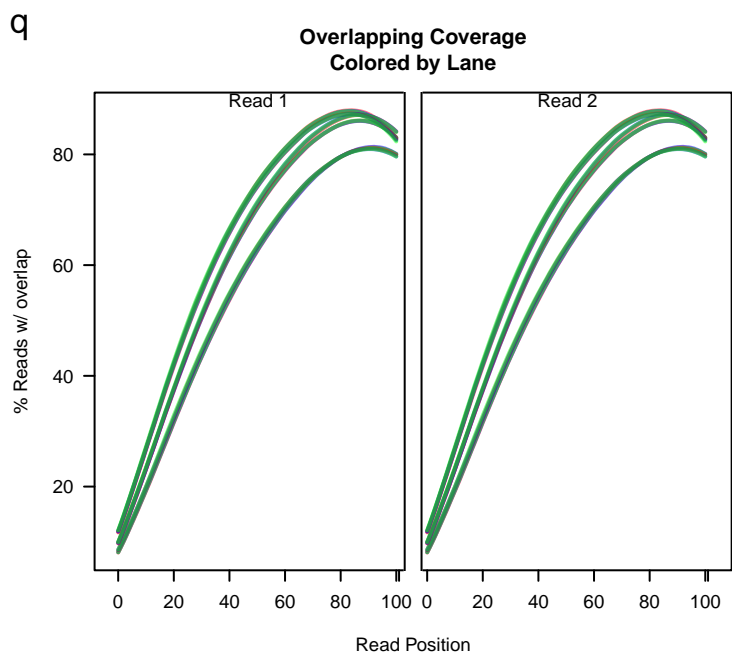
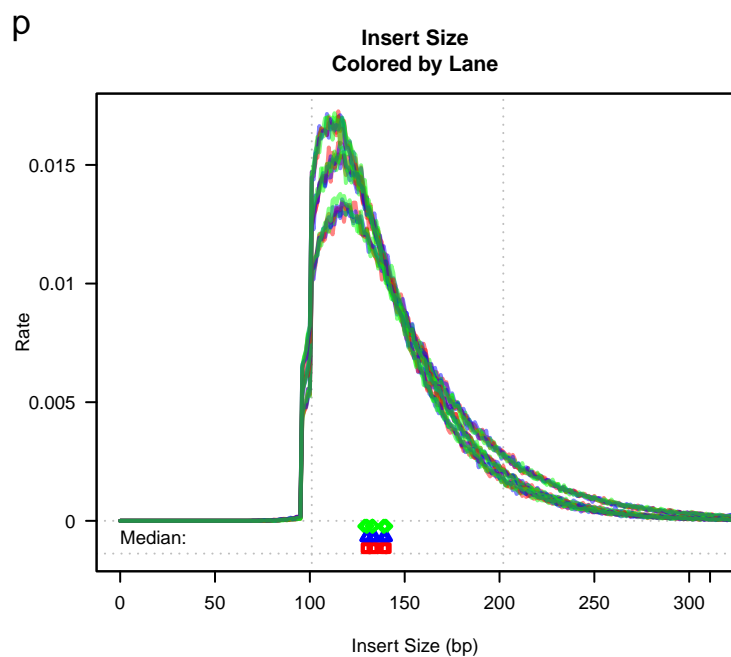
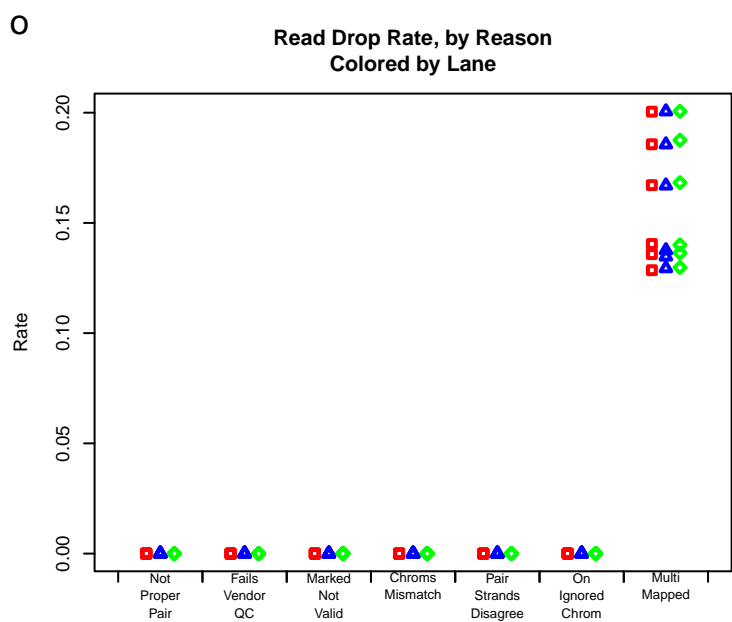
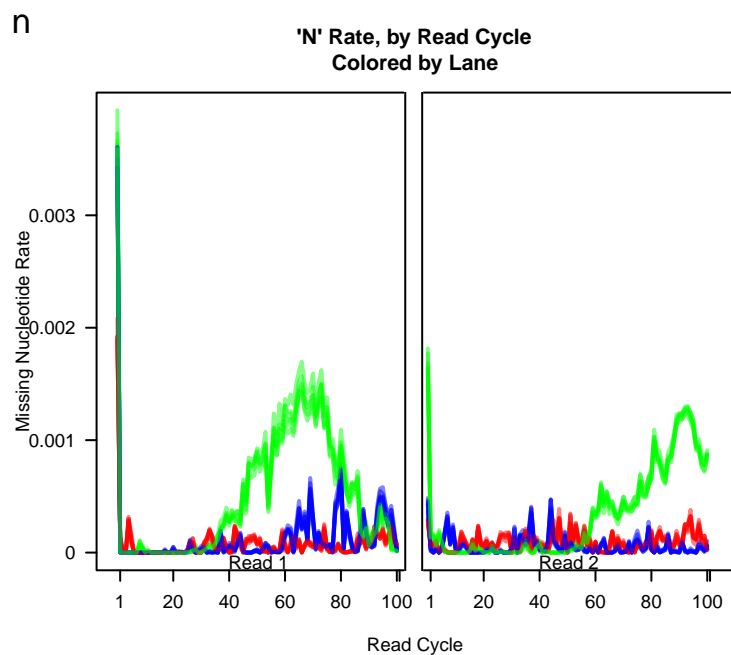
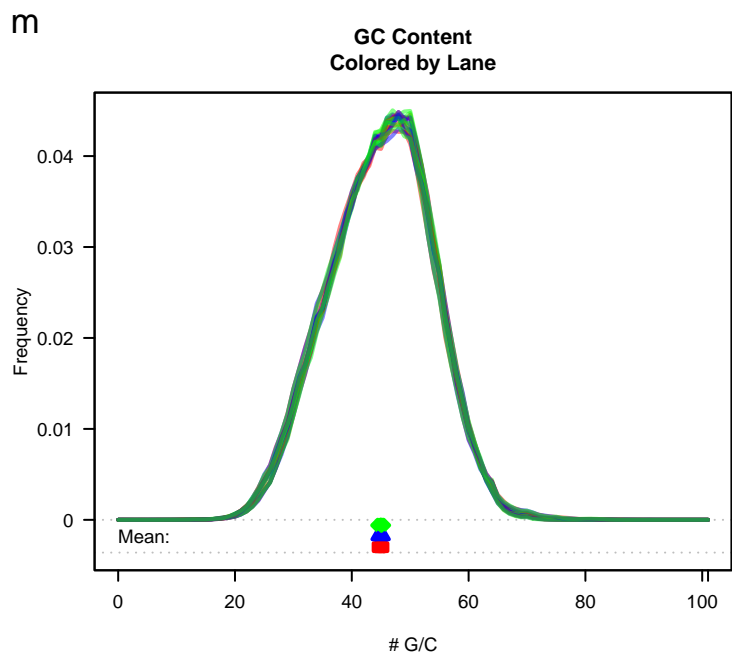
e

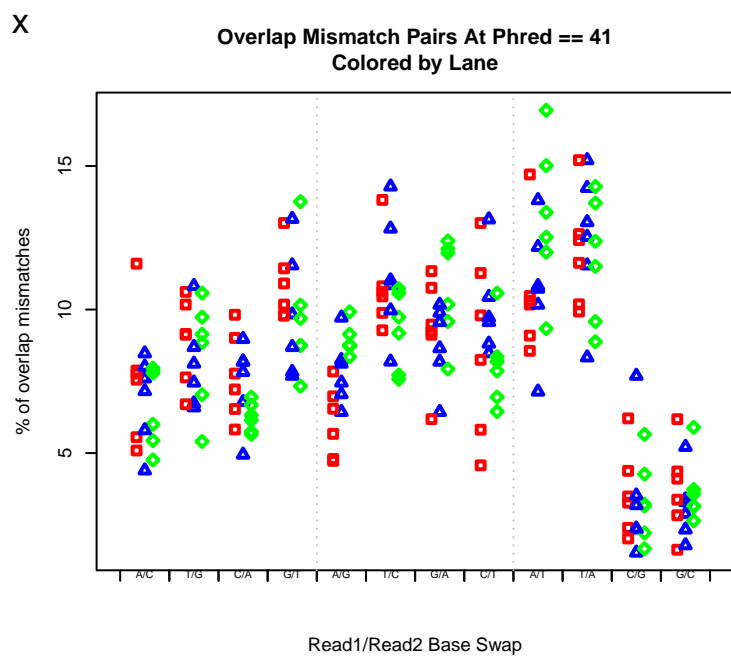
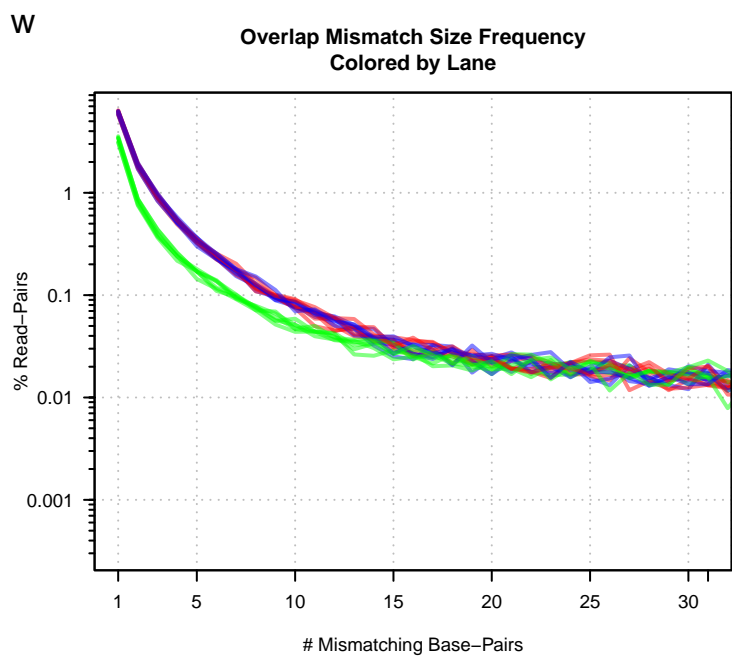
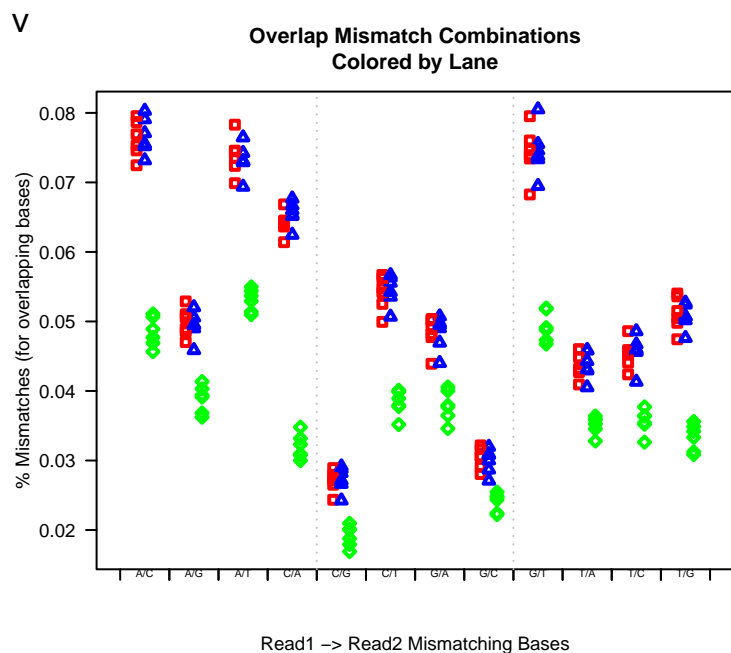
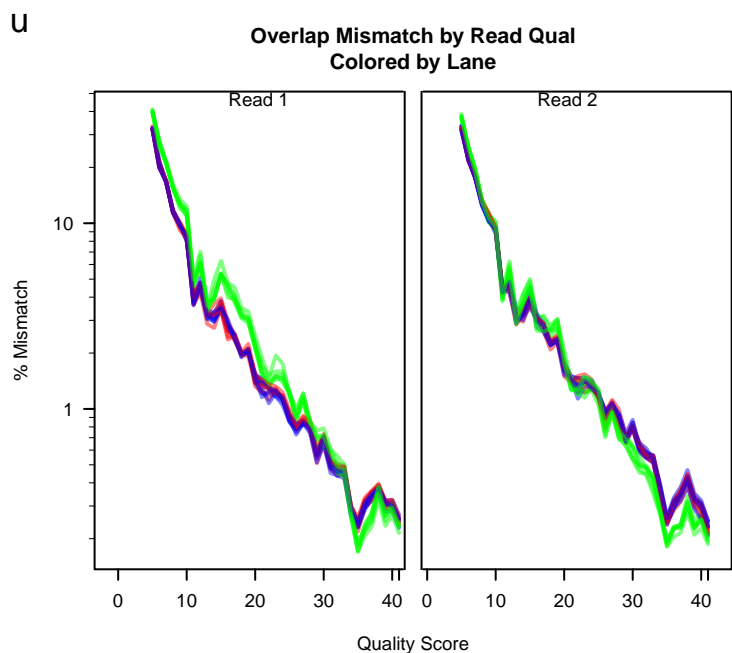
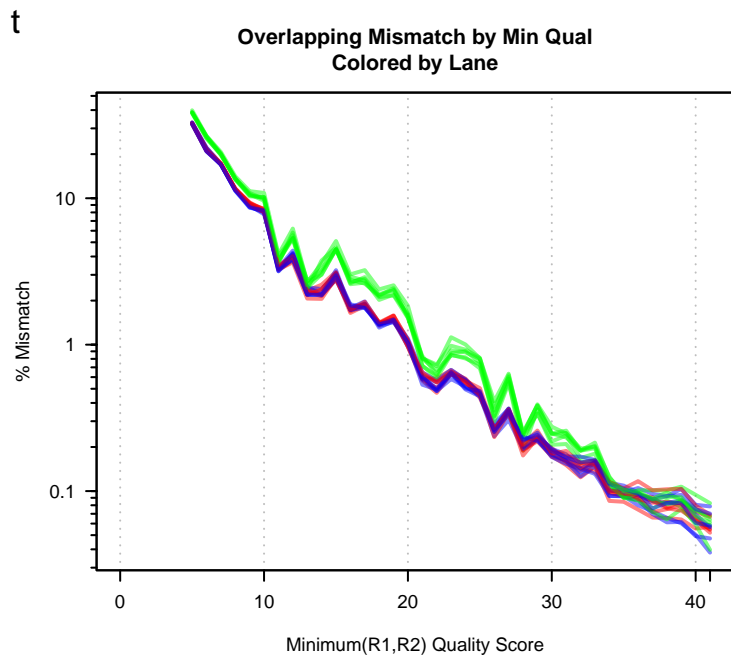
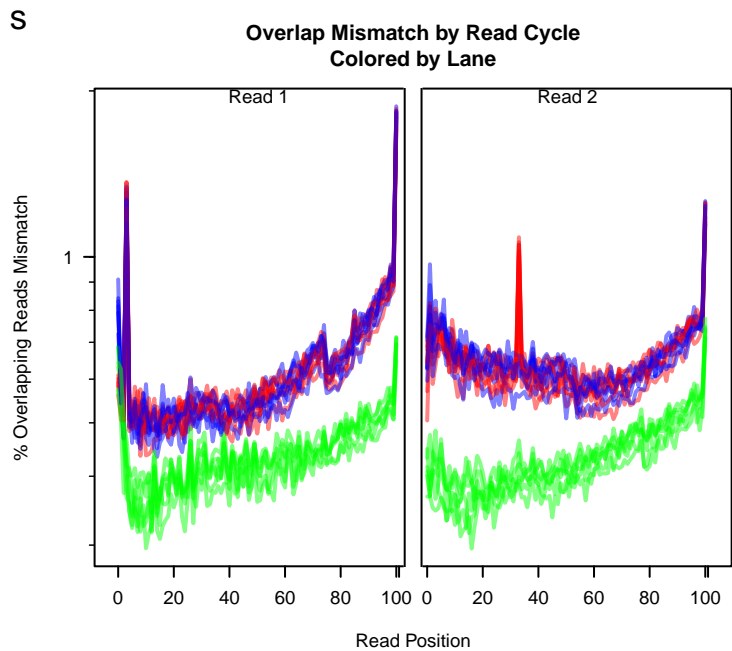
Upper Quartile Phred Quality Score
Colored by Lane

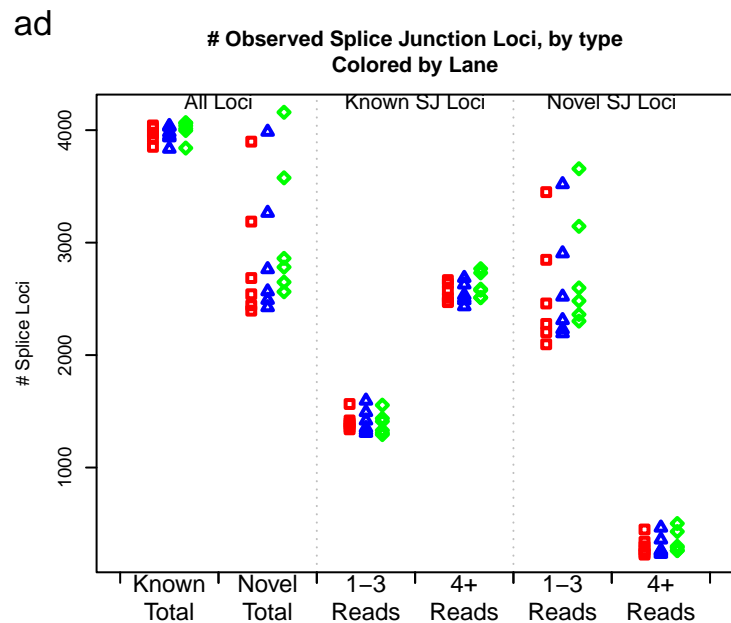
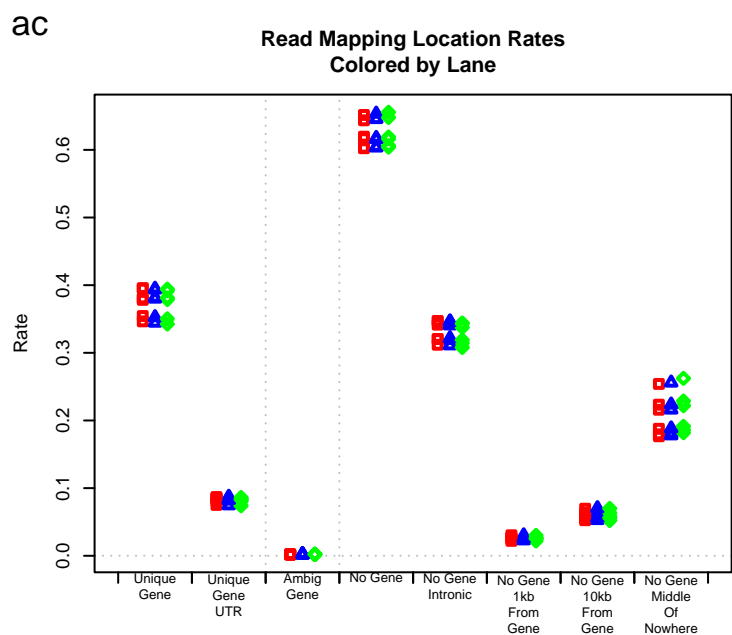
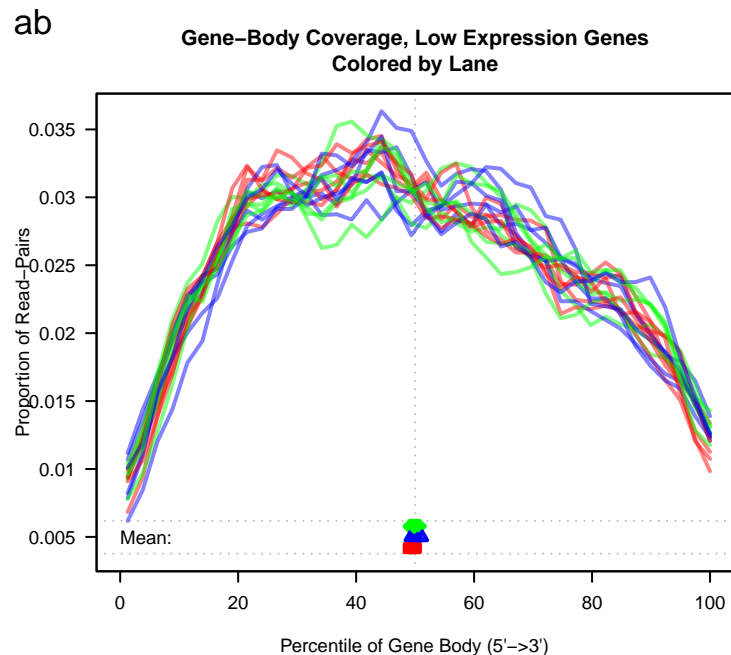
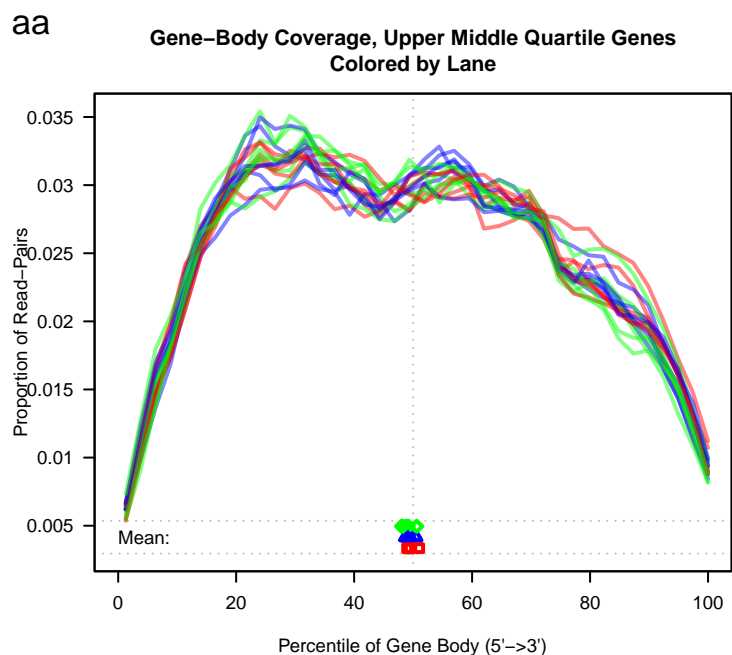
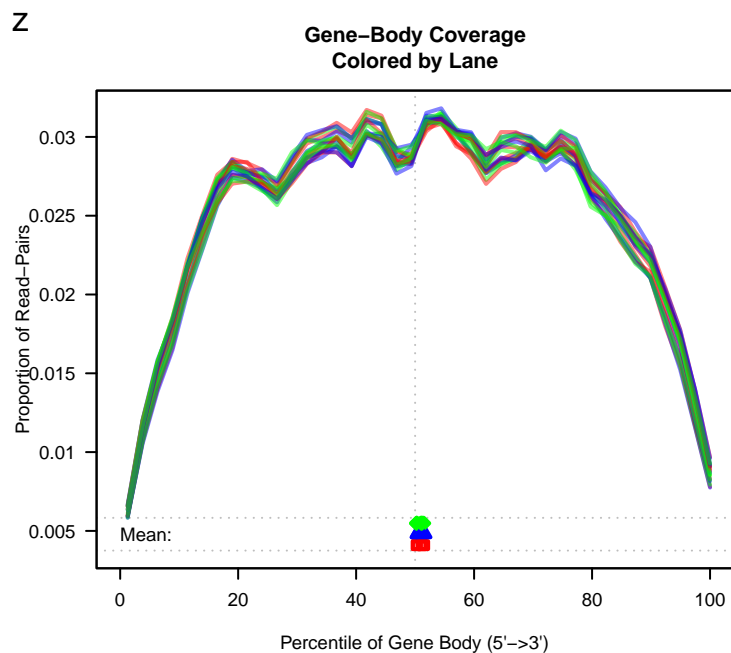
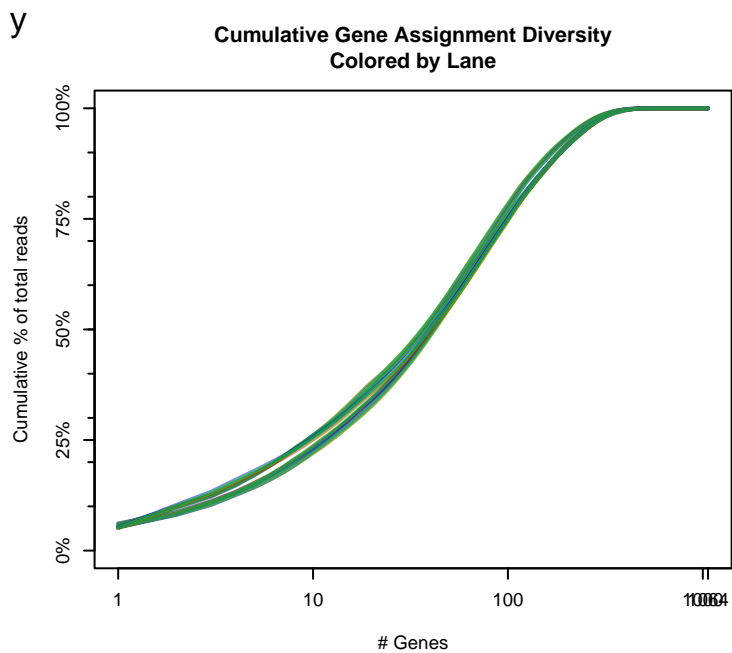
f

Maximum Phred Quality Score
Colored by Lane

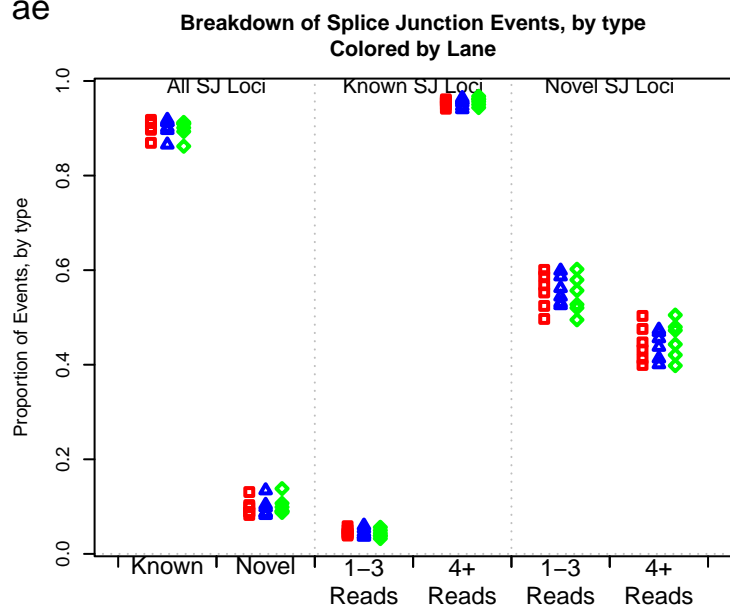




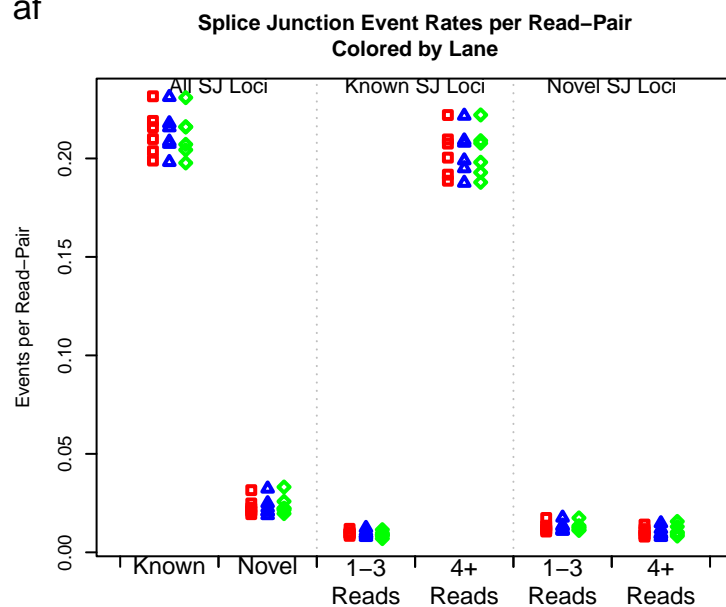




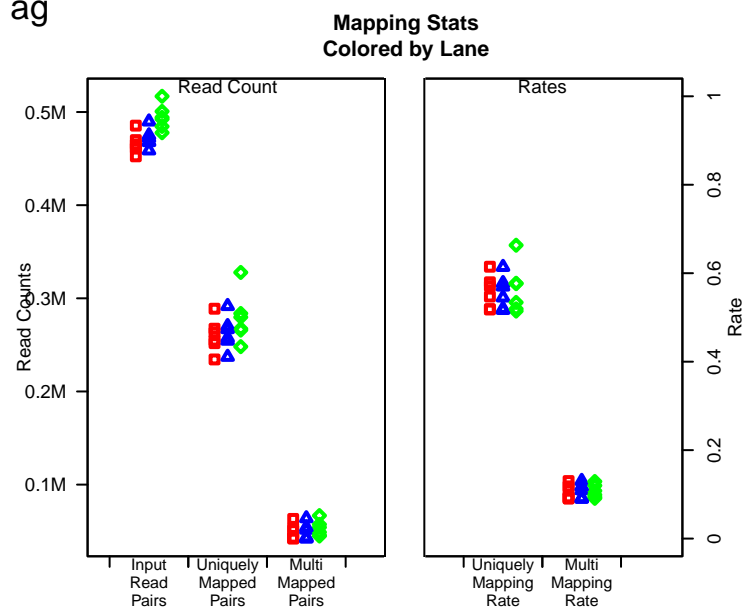
ae



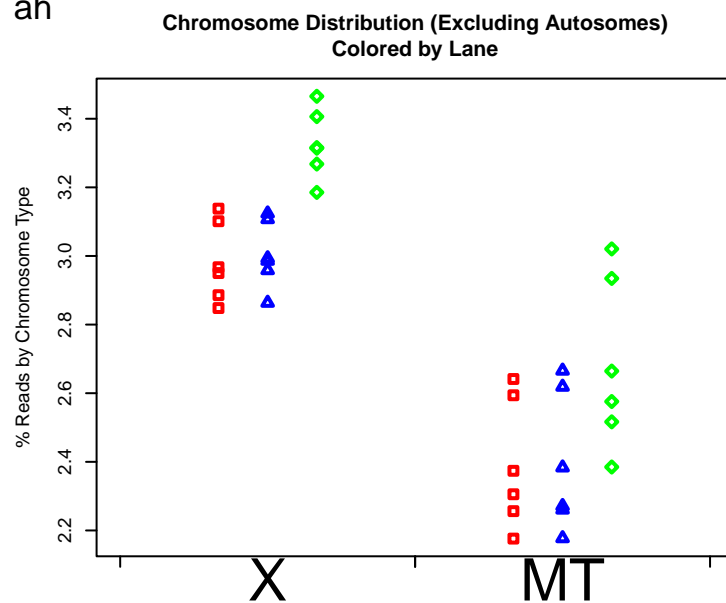
af



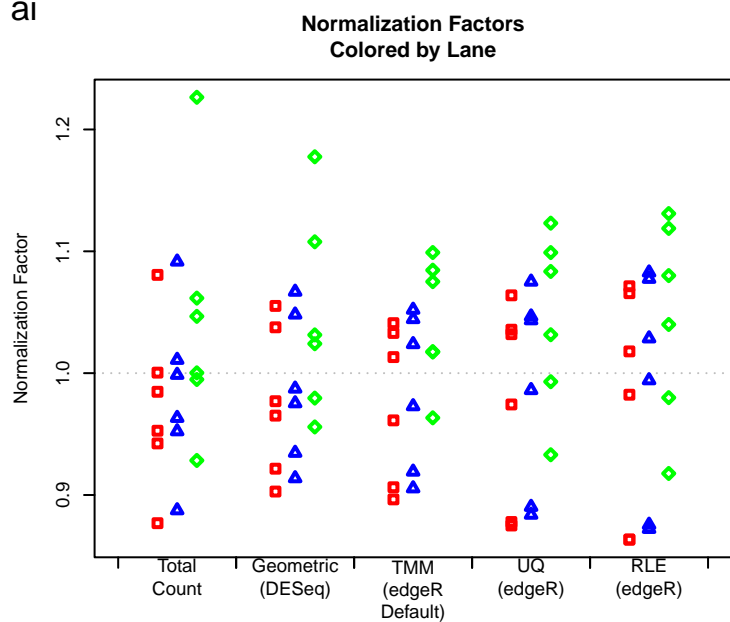
ag



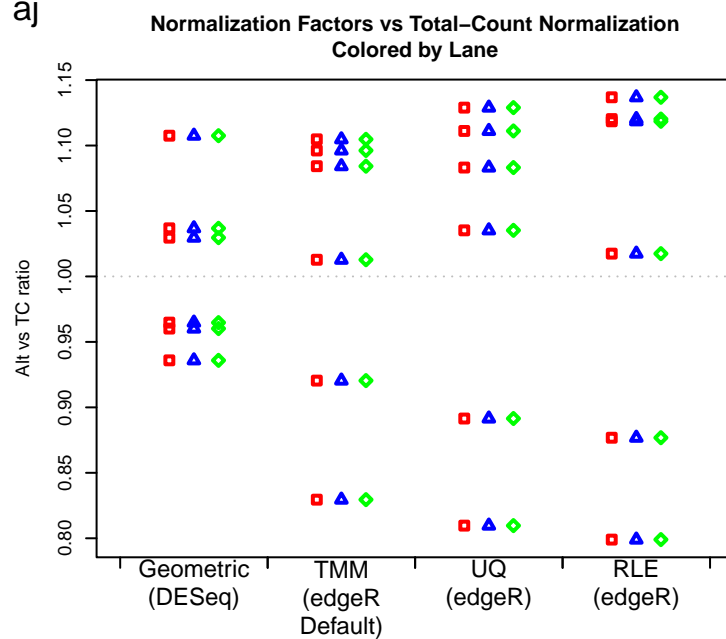
ah



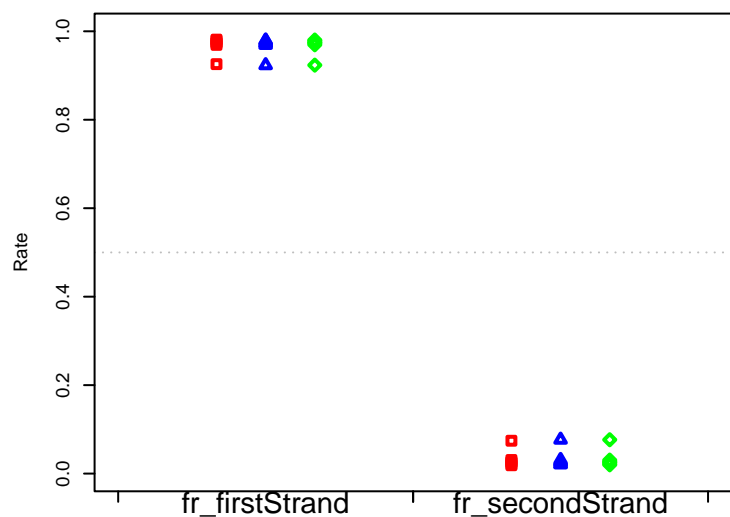
ai



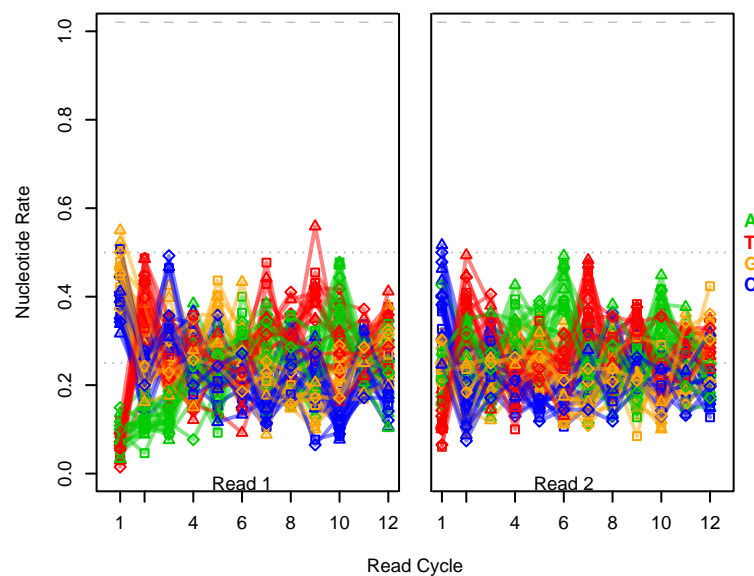
aj



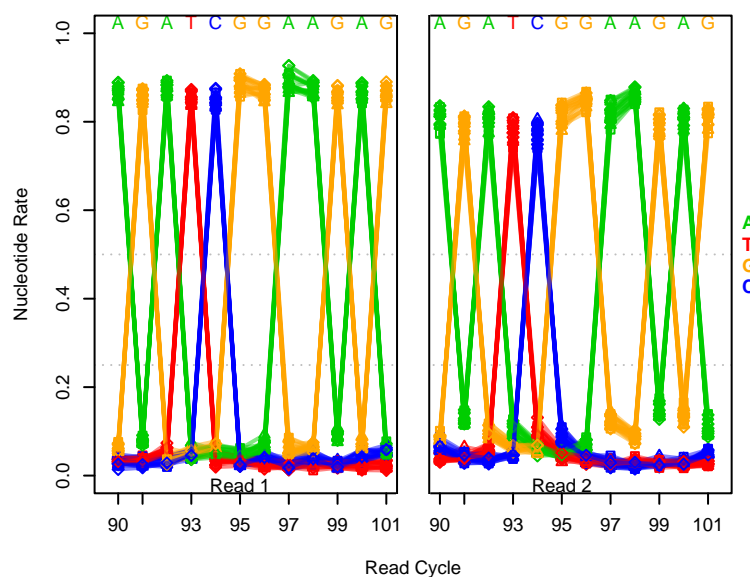
ak

Strandedness Test
Colored by Lane

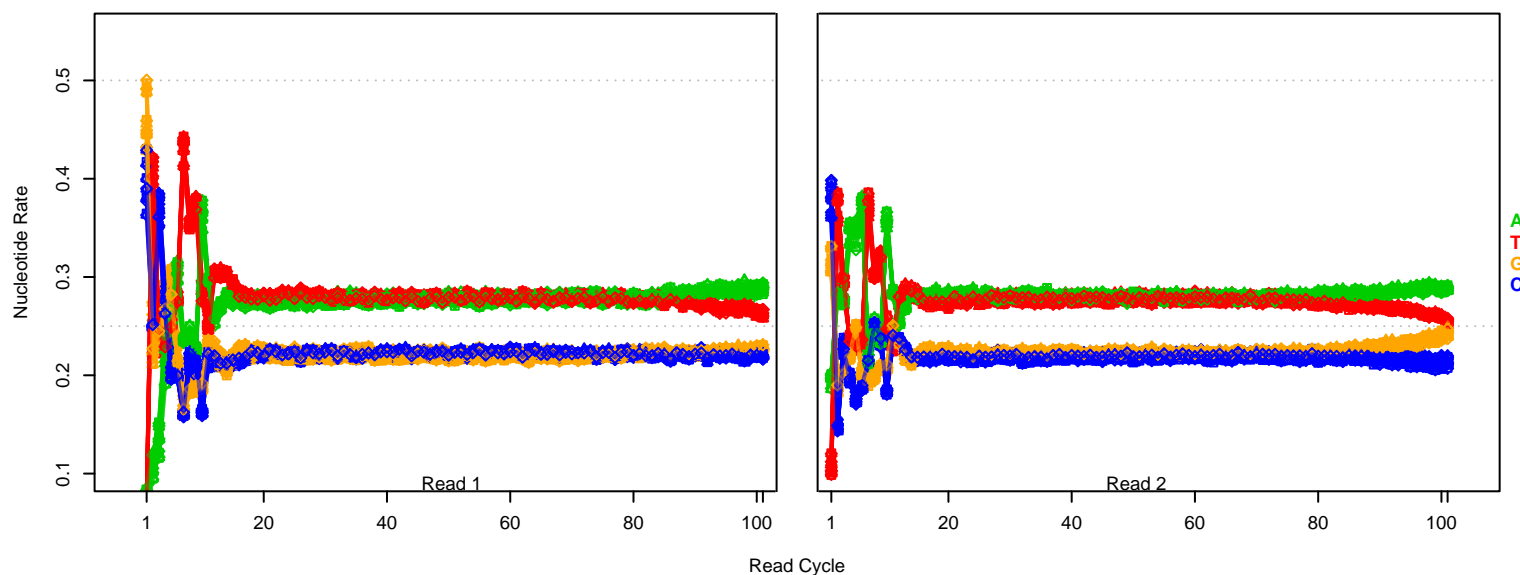
al

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

am

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

an

Raw Nucleotide Rate by Cycle
Marked by Lane

ao

Nucleotide Rate by Cycle, Aligned bases only
Marked by Lane

