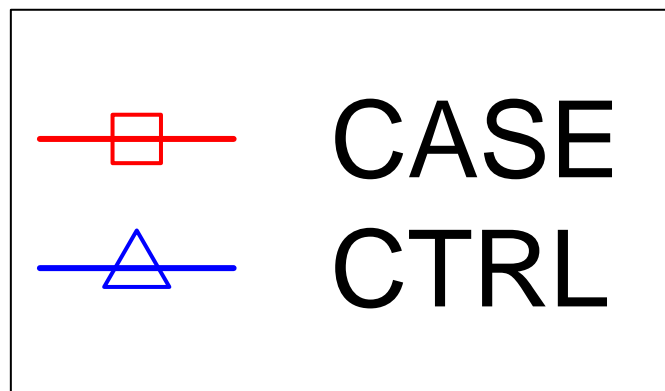
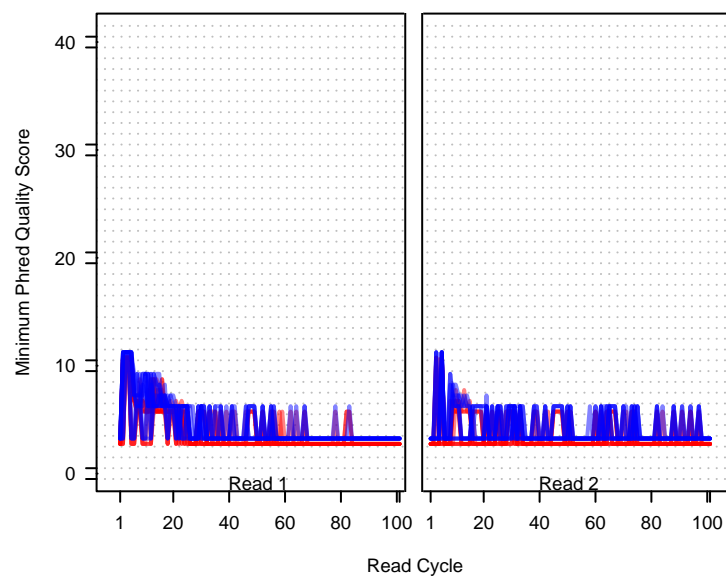


a

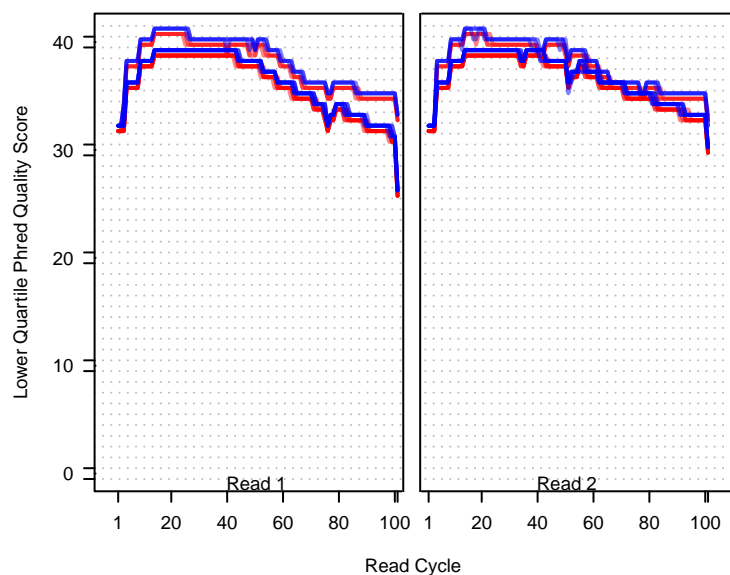
Summary Plots, By Group



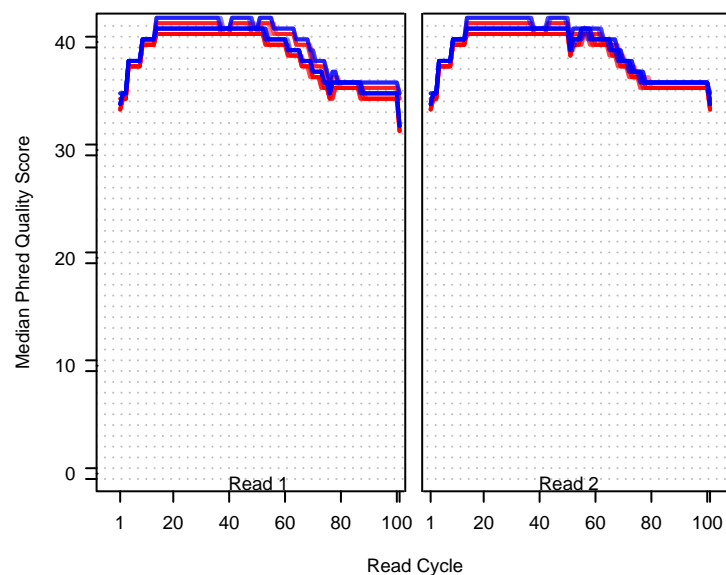
b

Minimum Phred Quality Score
Colored by Group

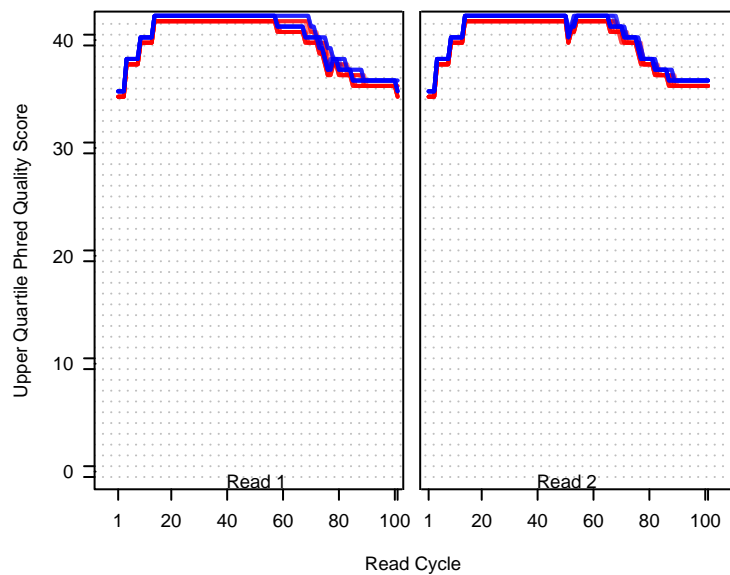
c

Lower Quartile Phred Quality Score
Colored by Group

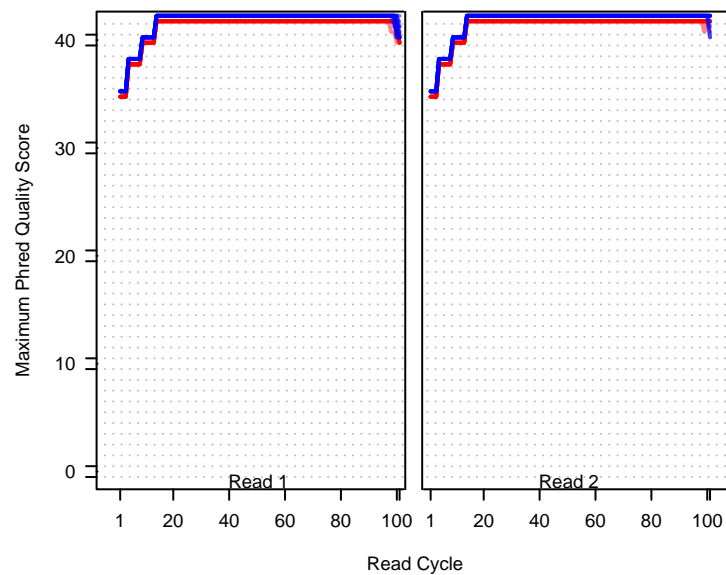
d

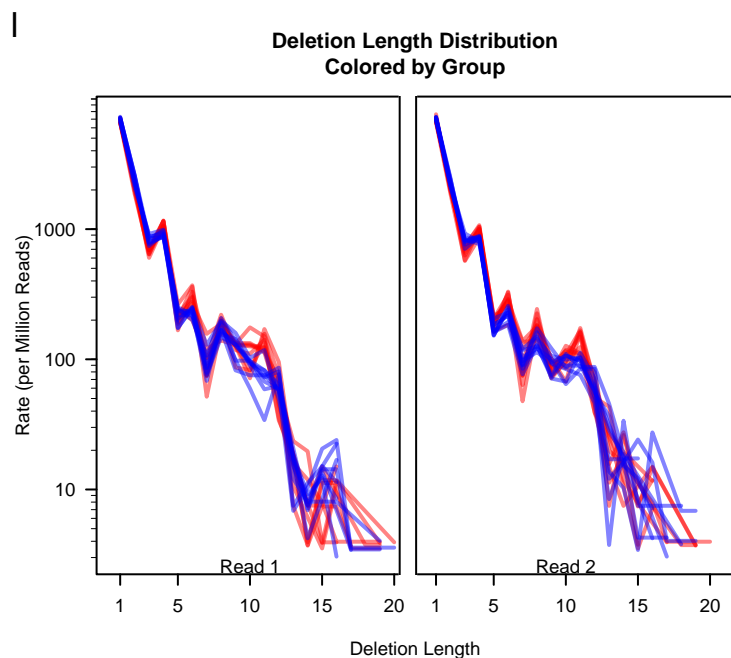
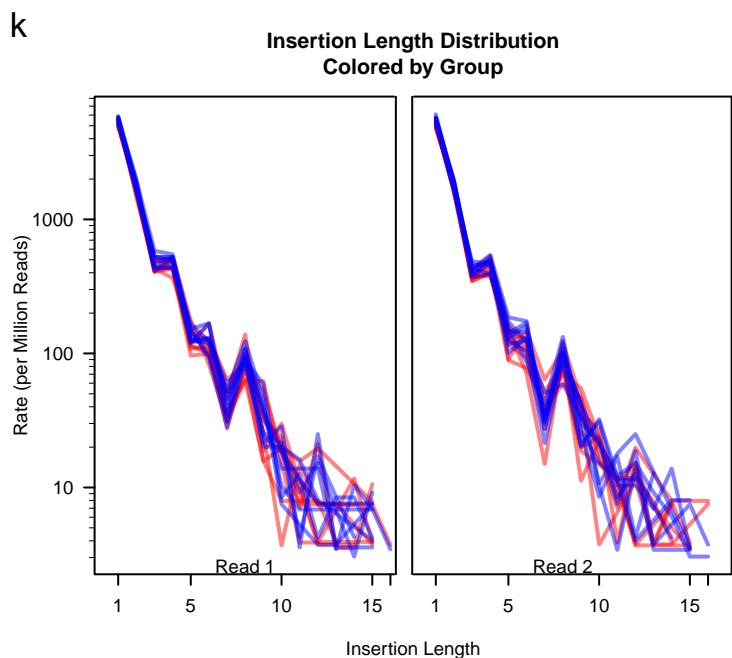
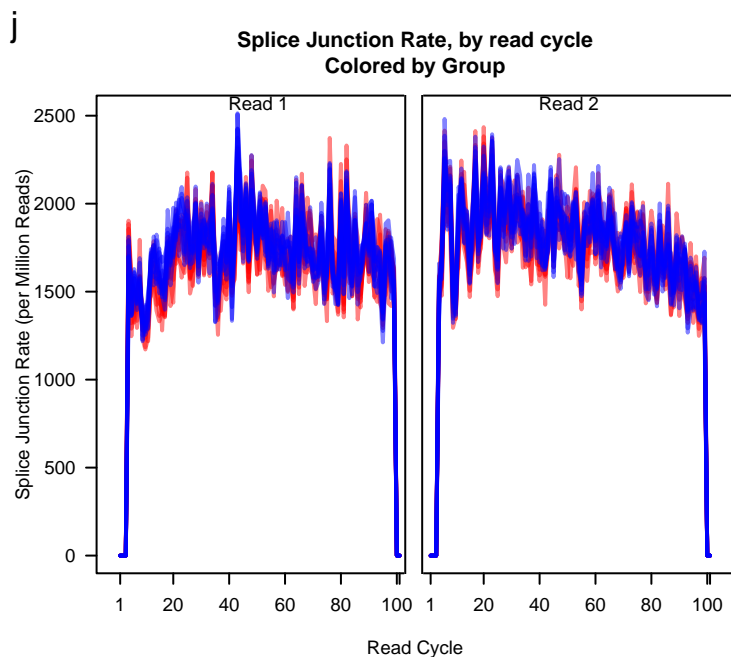
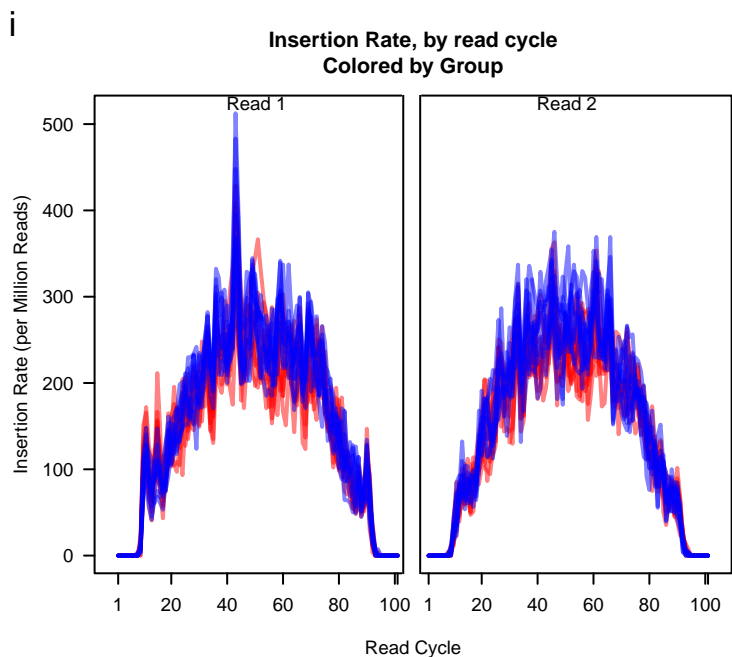
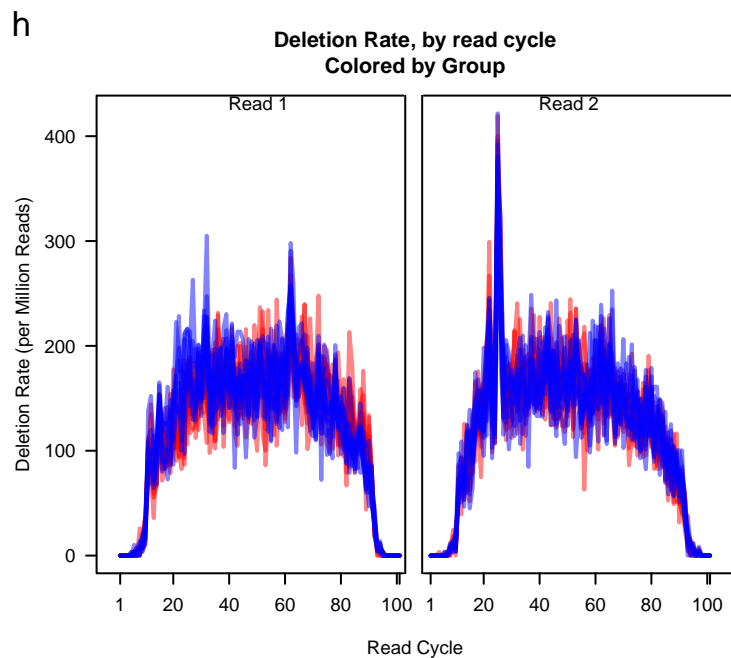
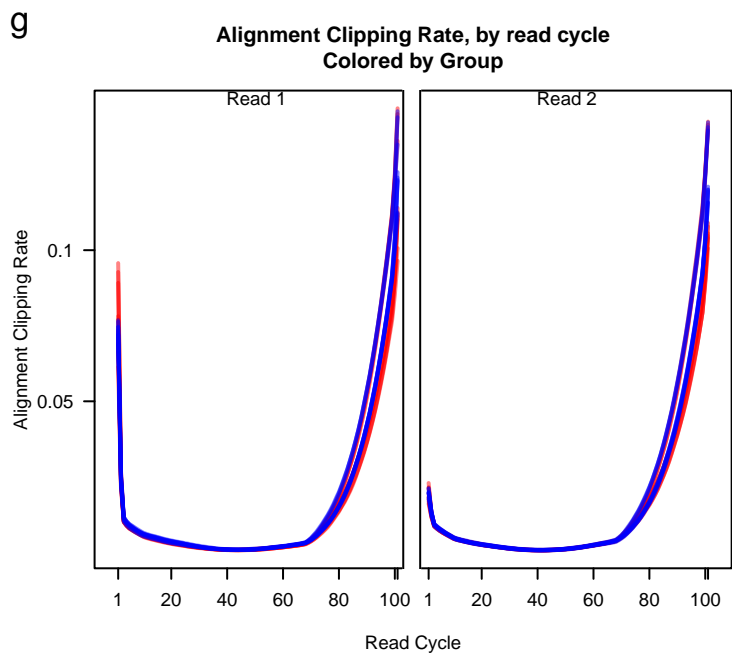
Median Phred Quality Score
Colored by Group

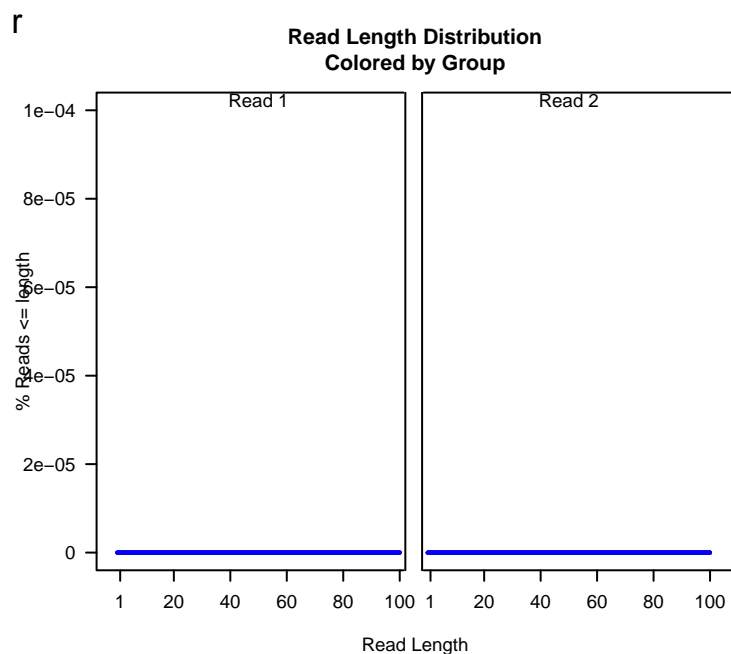
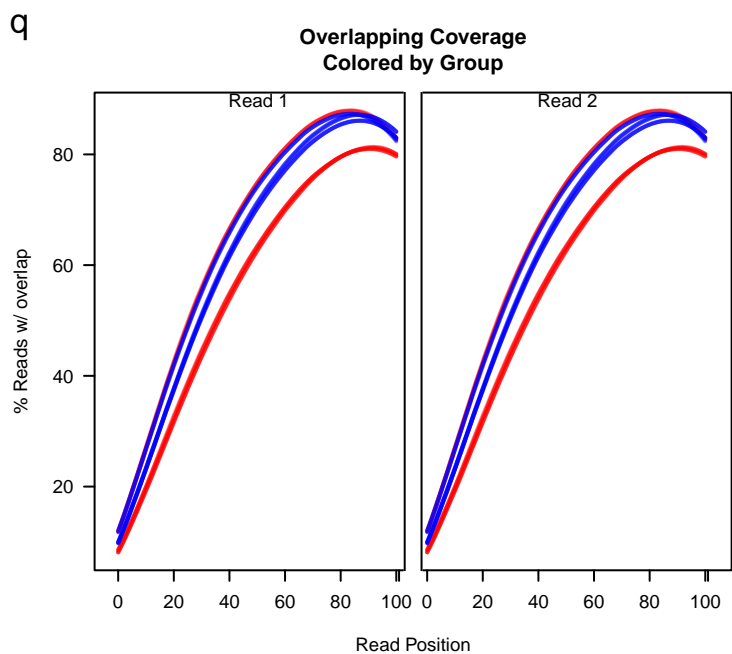
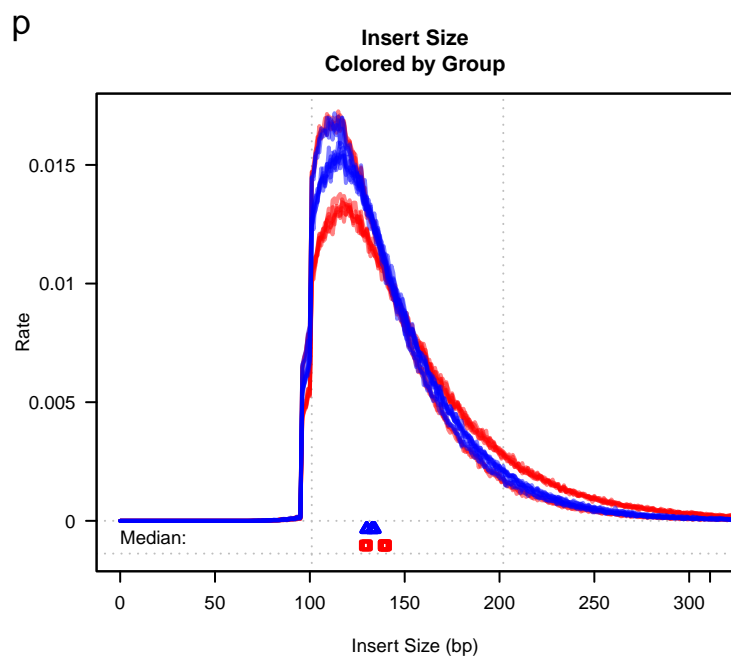
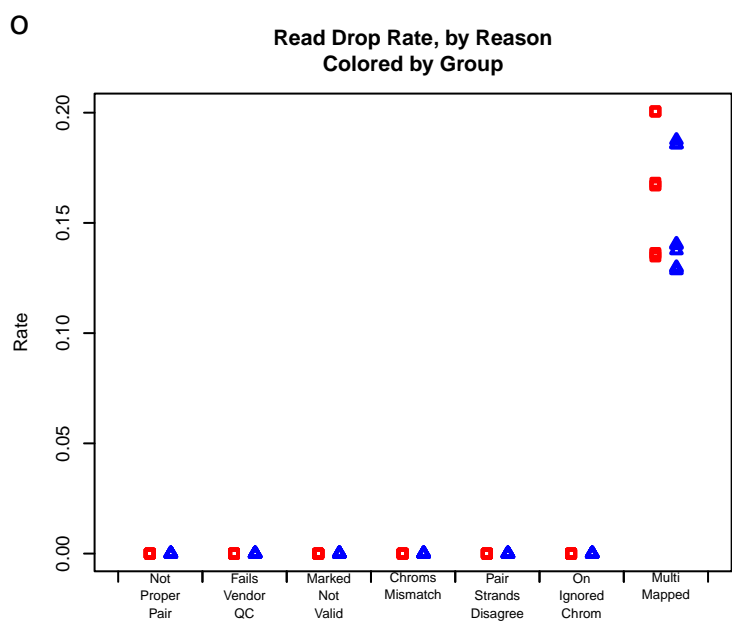
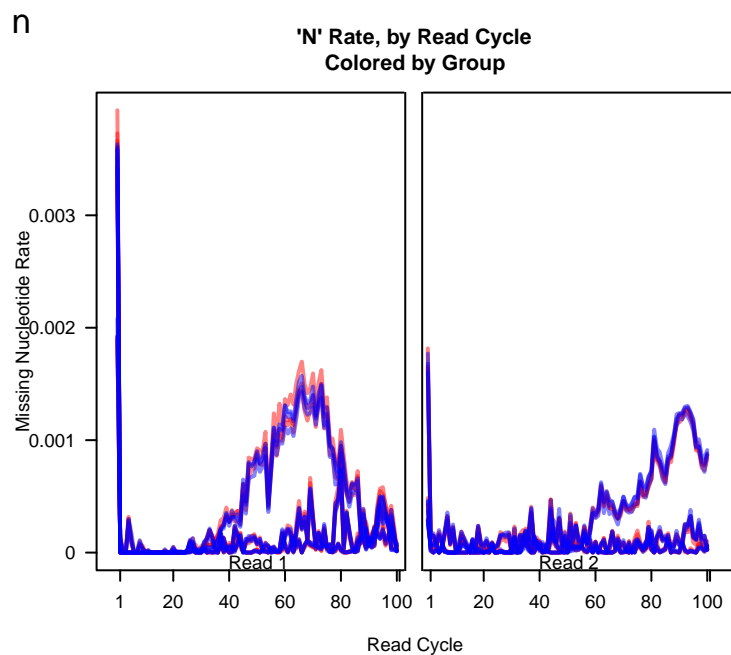
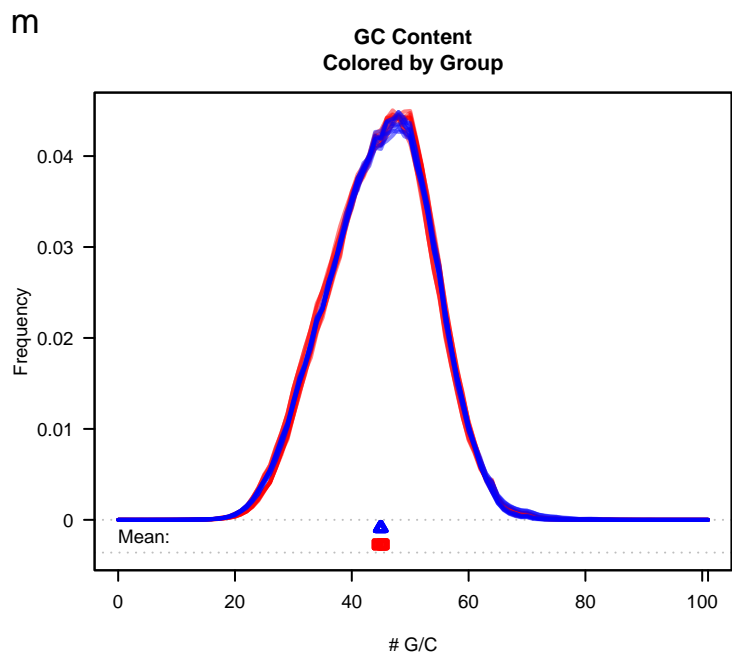
e

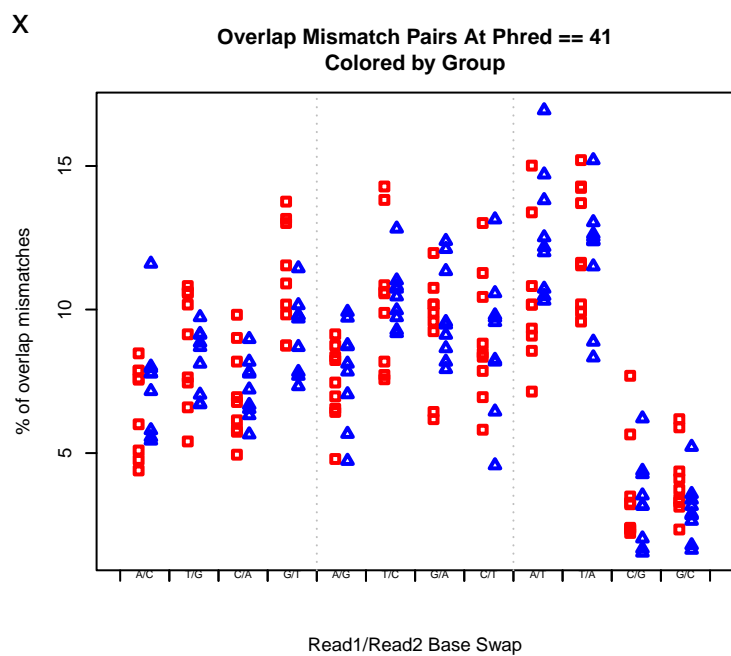
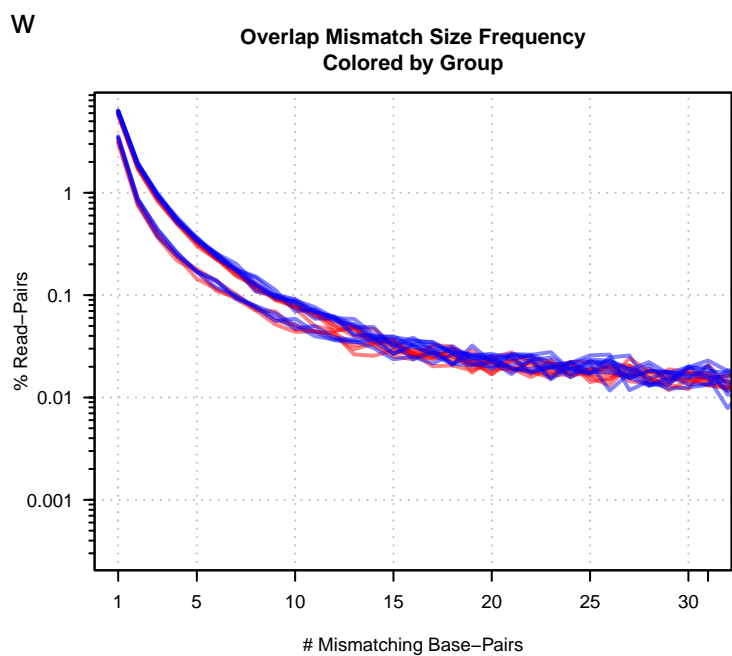
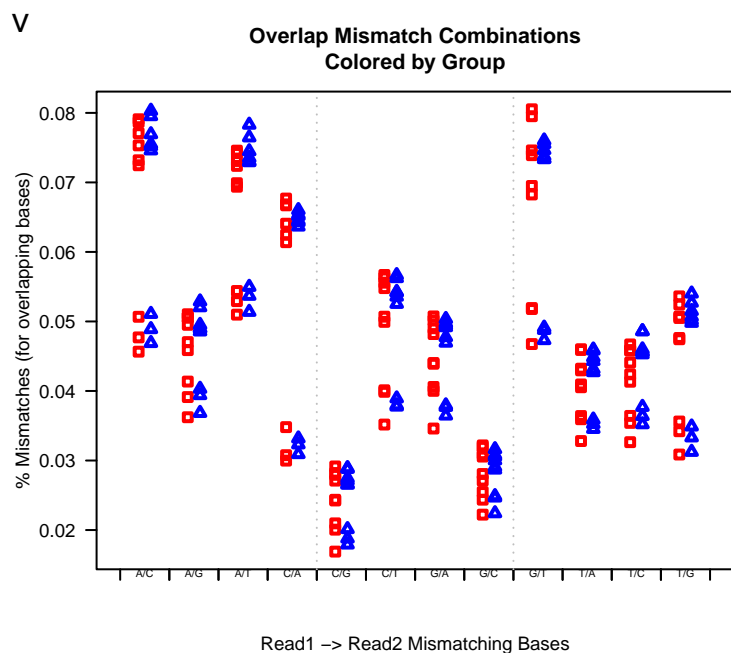
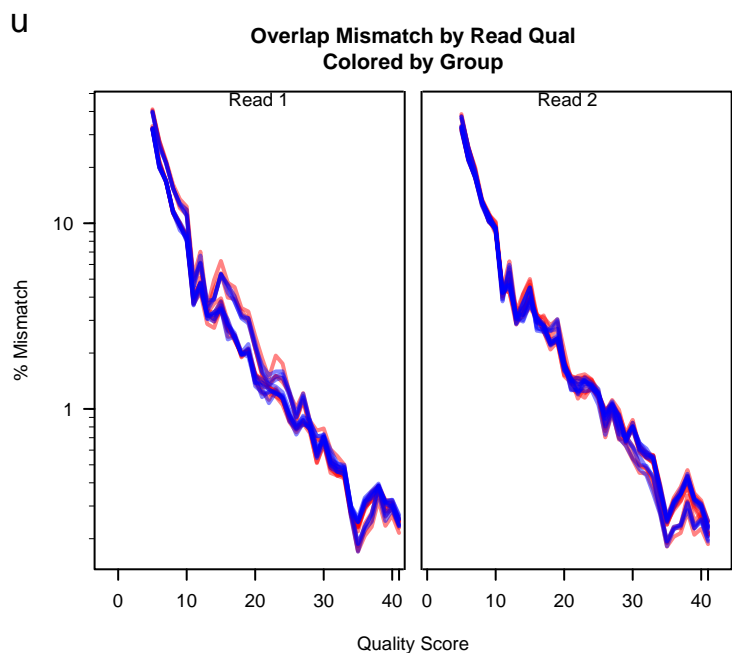
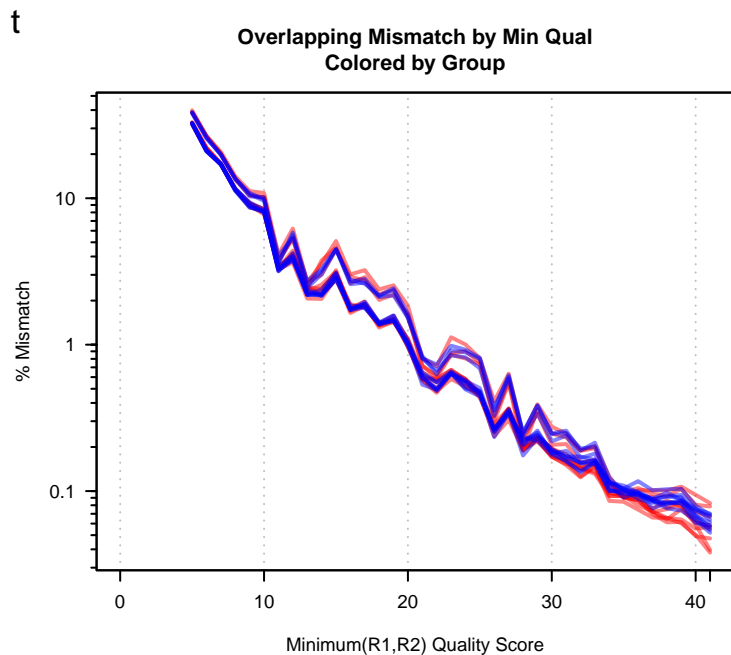
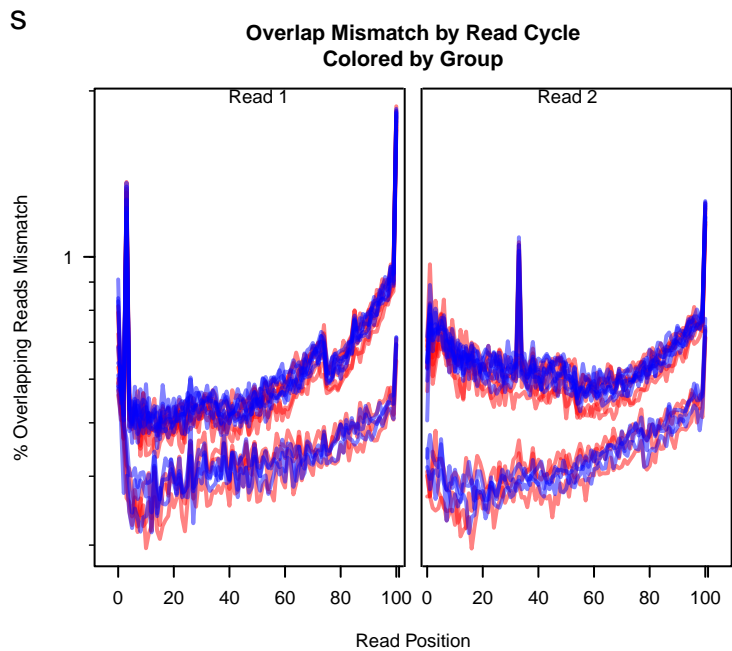
Upper Quartile Phred Quality Score
Colored by Group

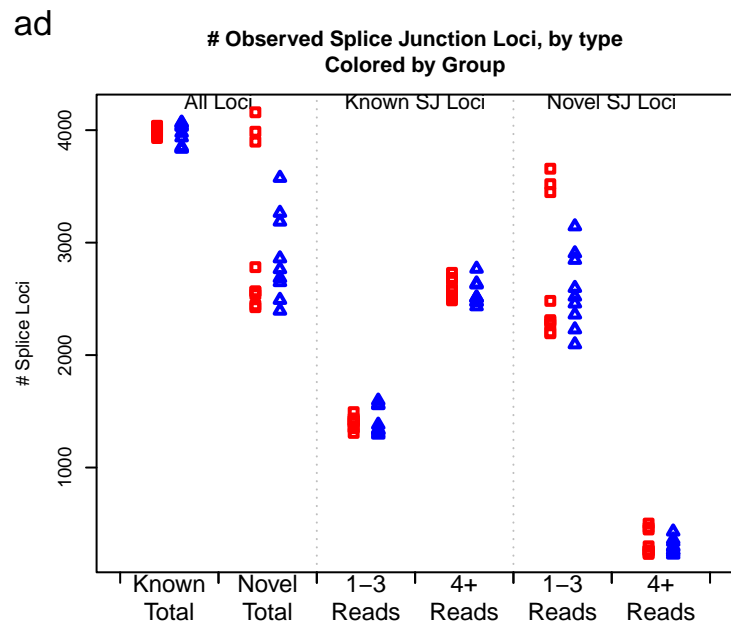
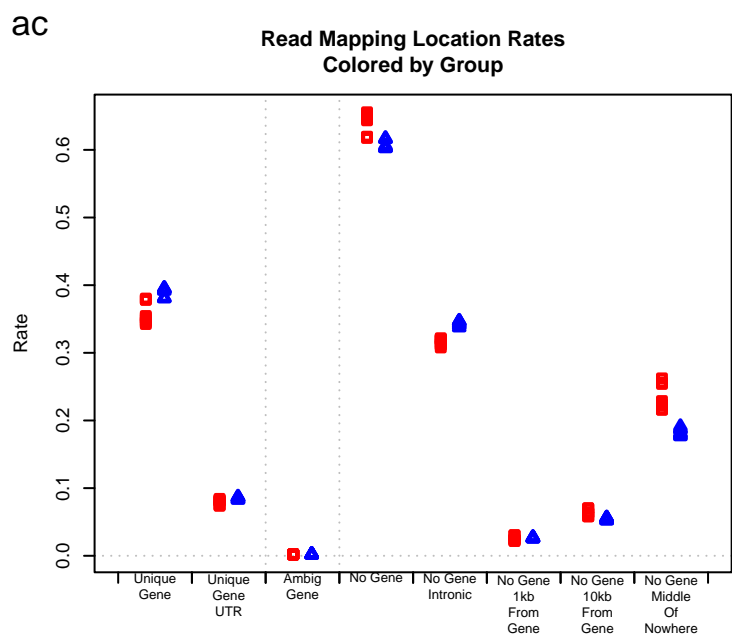
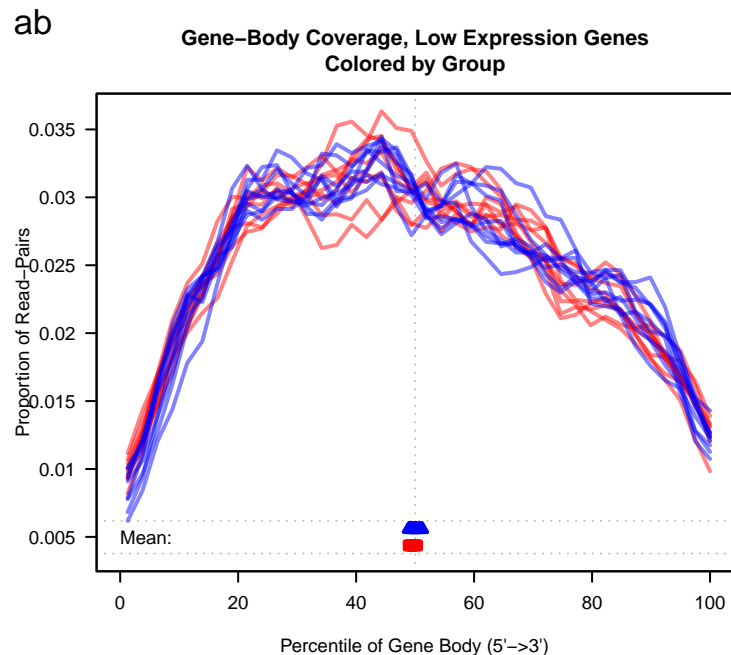
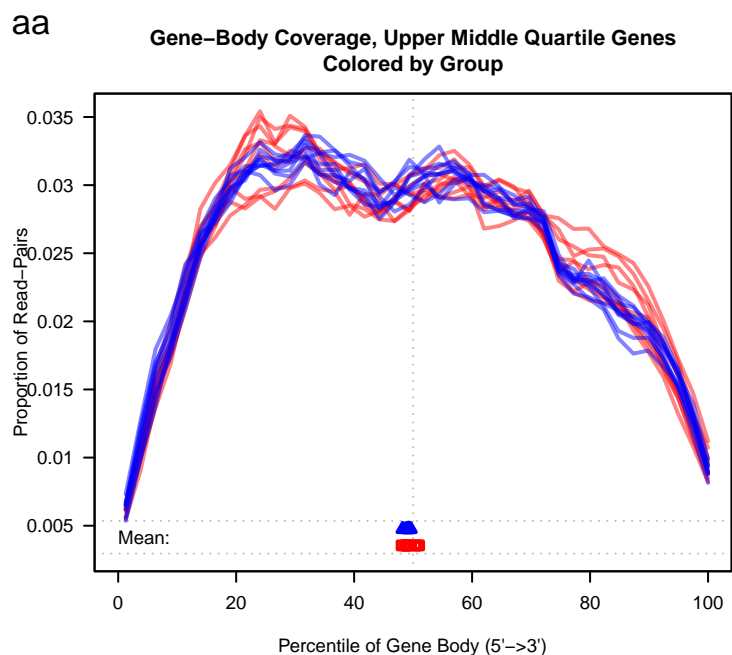
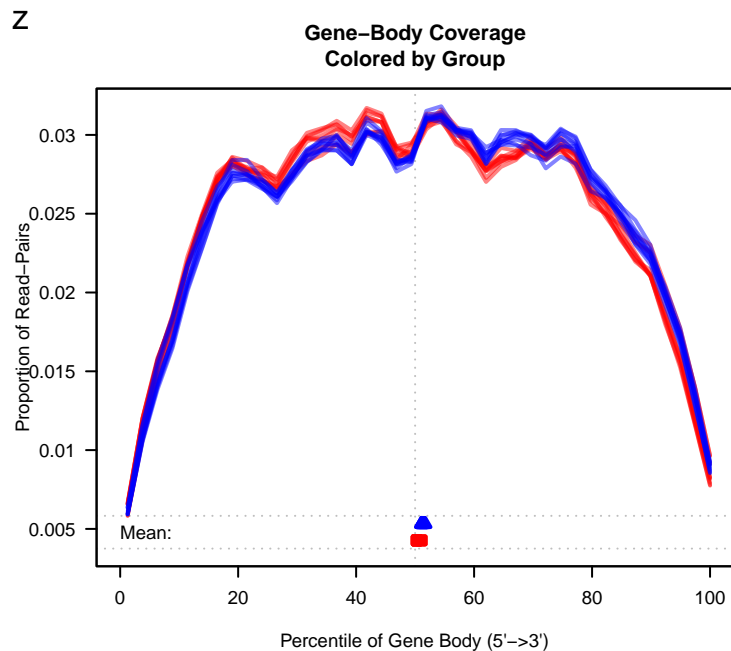
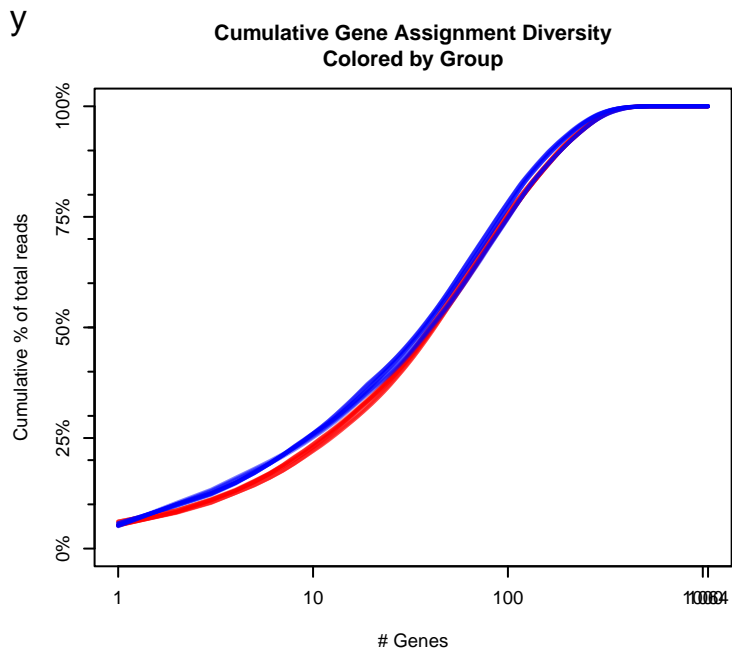
f

Maximum Phred Quality Score
Colored by Group

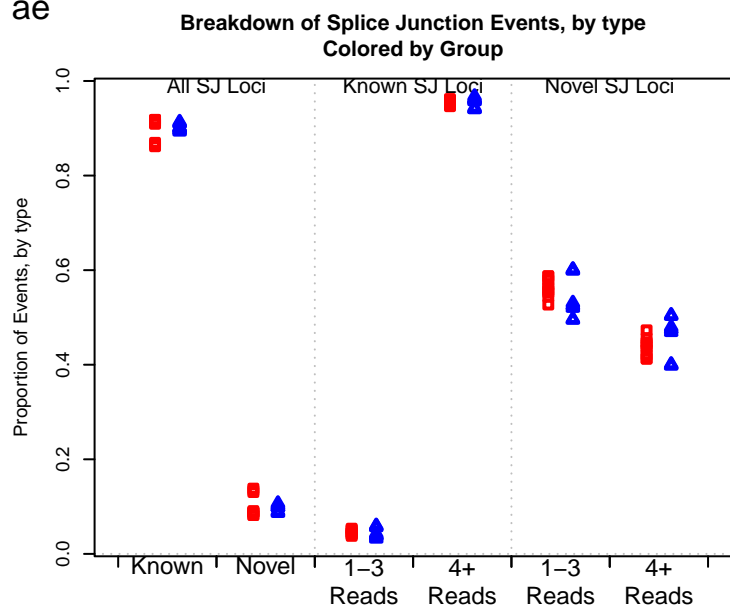




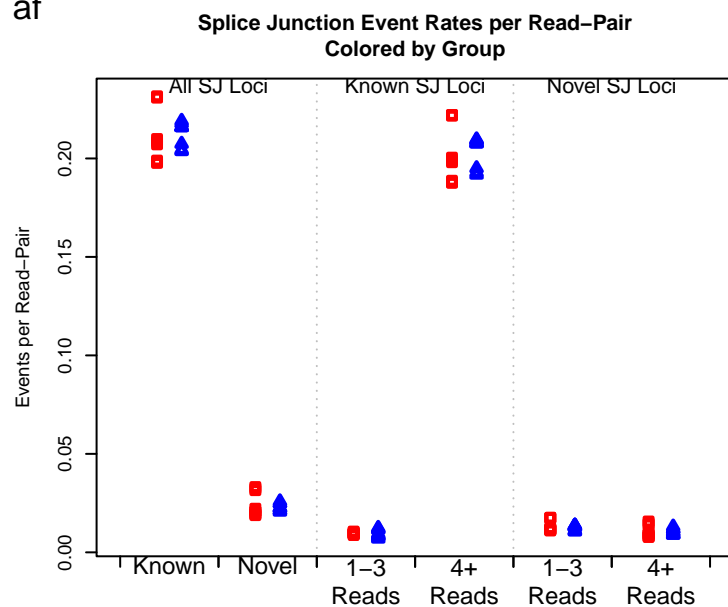




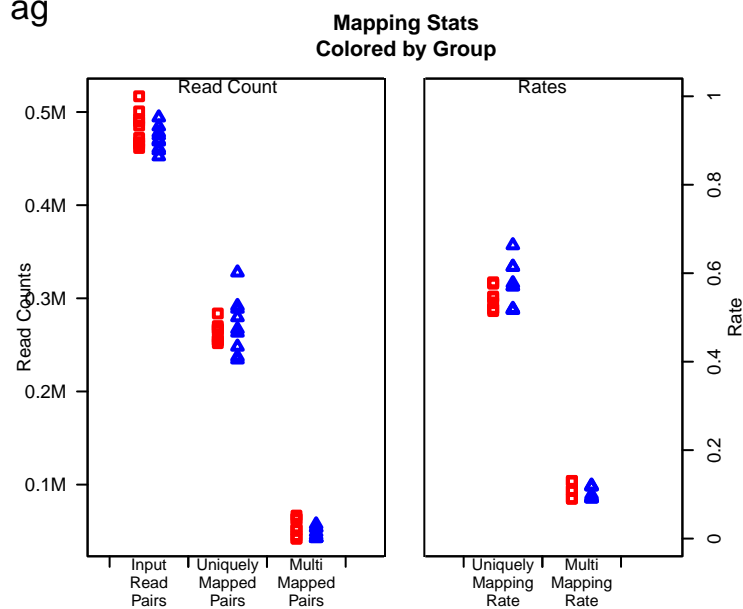
ae



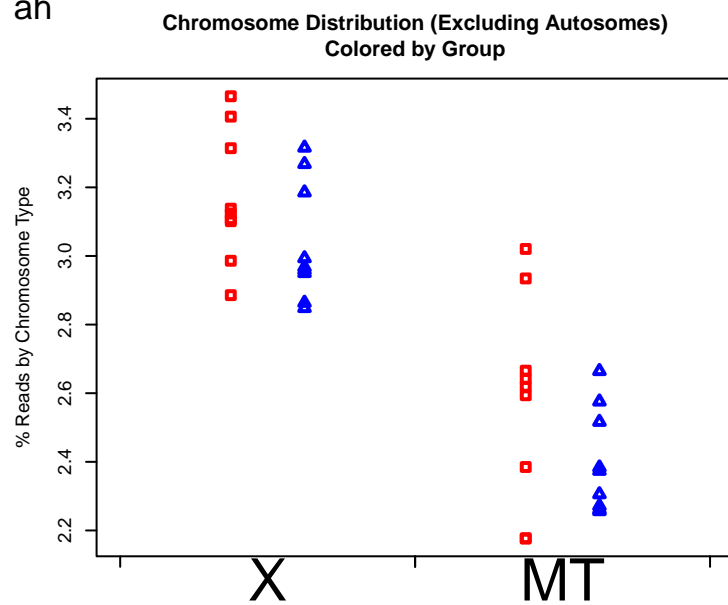
af



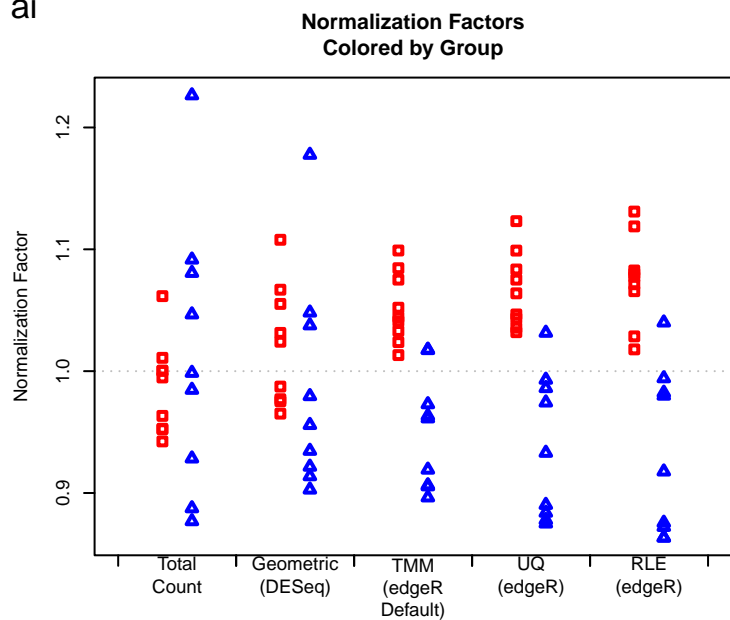
ag



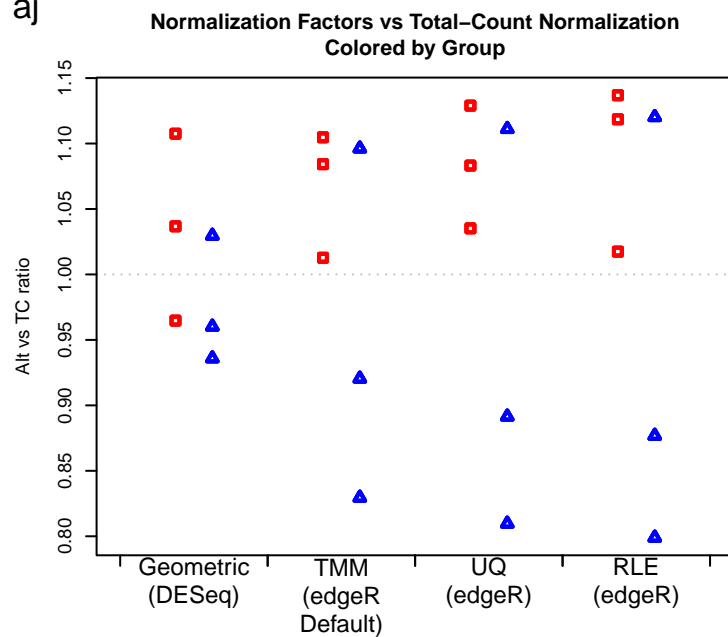
ah



ai

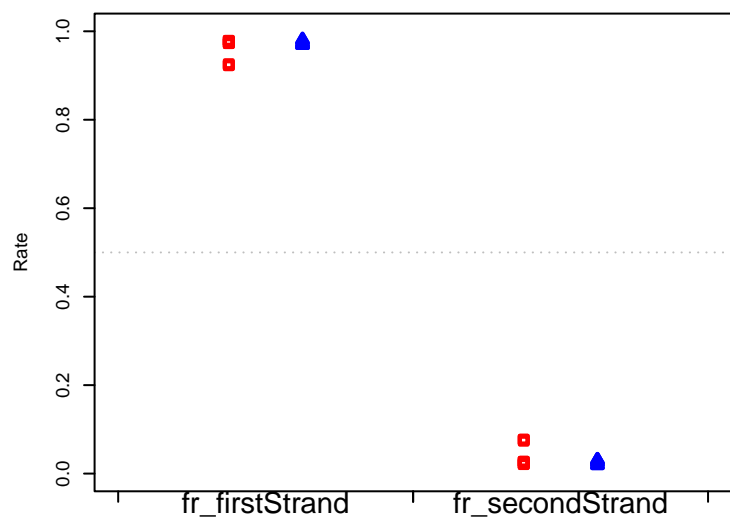


aj



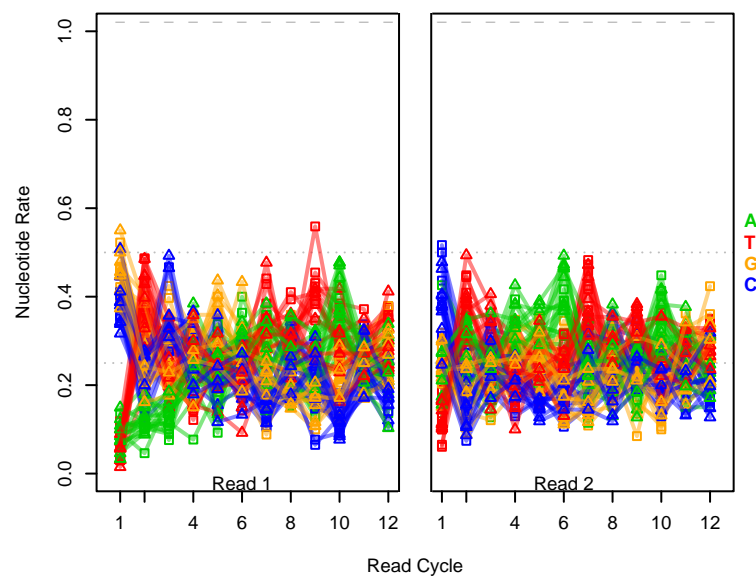
ak

Strandedness Test
Colored by Group



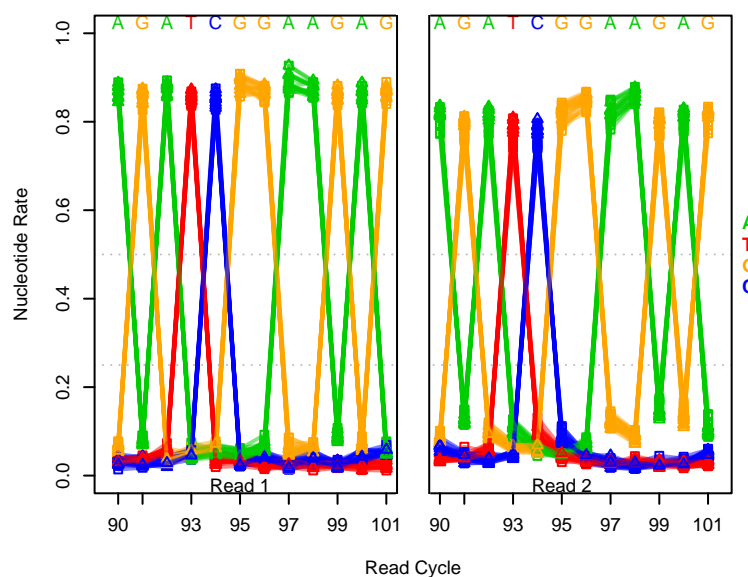
al

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



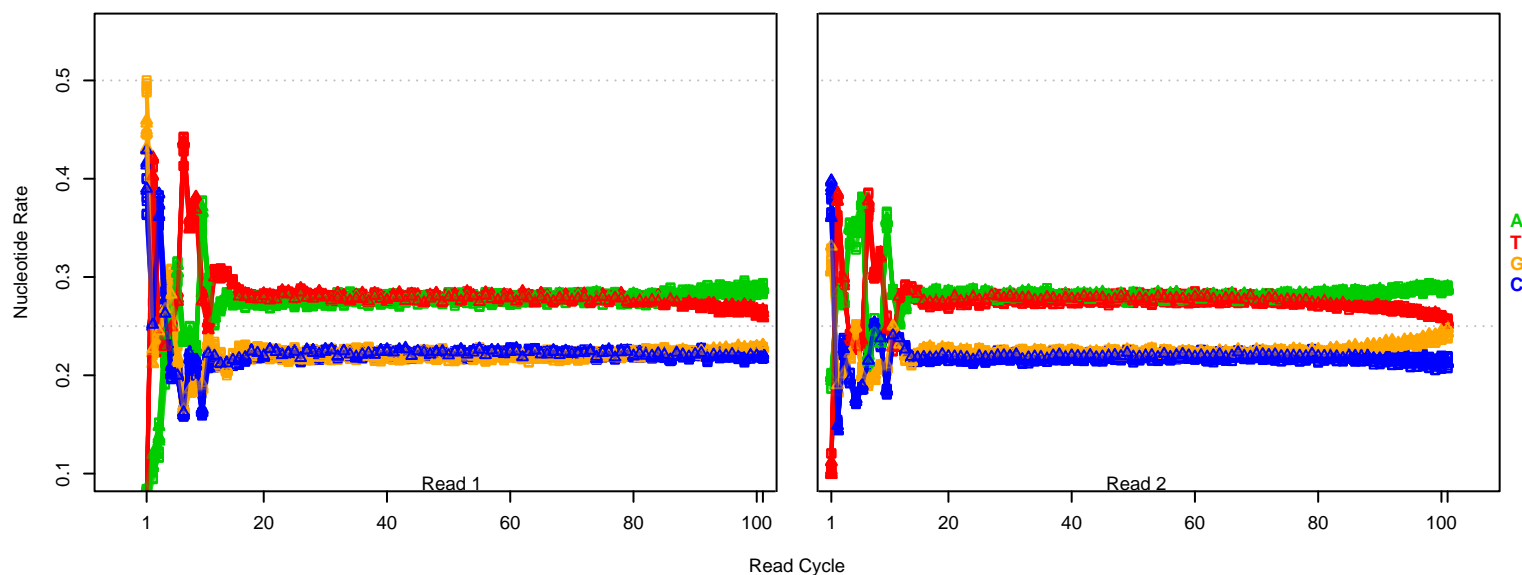
am

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



an

Raw Nucleotide Rate by Cycle
Marked by Group



ao

Nucleotide Rate by Cycle, Aligned bases only
Marked by Group

