

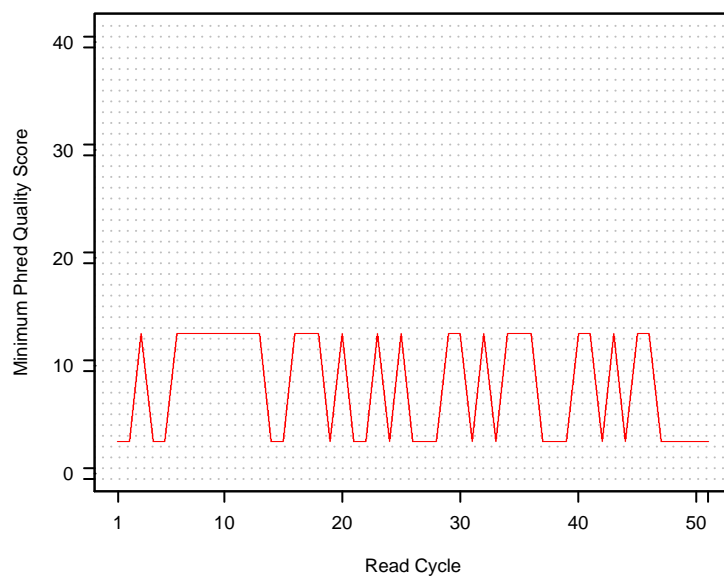
a

Sample Highlight: Untitled



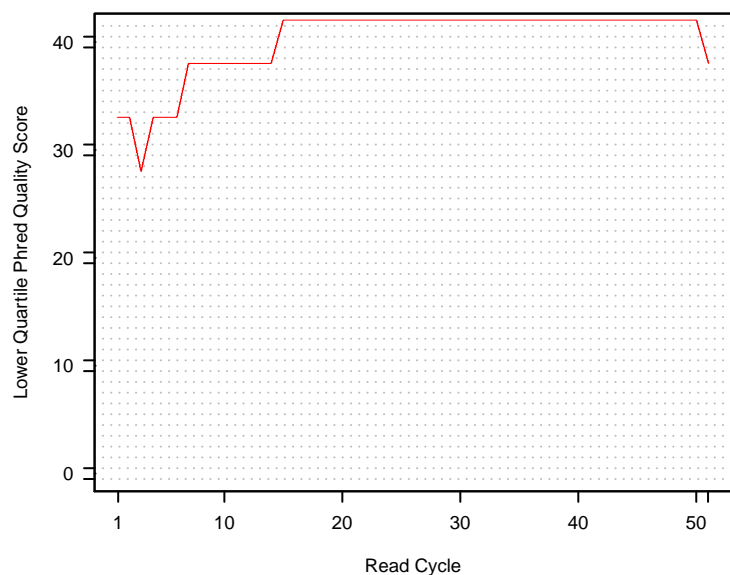
b

Minimum Phred Quality Score
With Sample Untitled in red



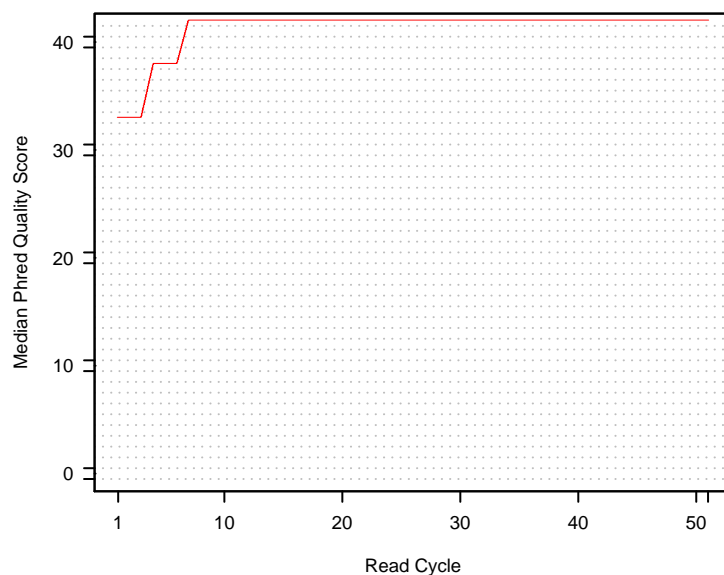
c

Lower Quartile Phred Quality Score
With Sample Untitled in red



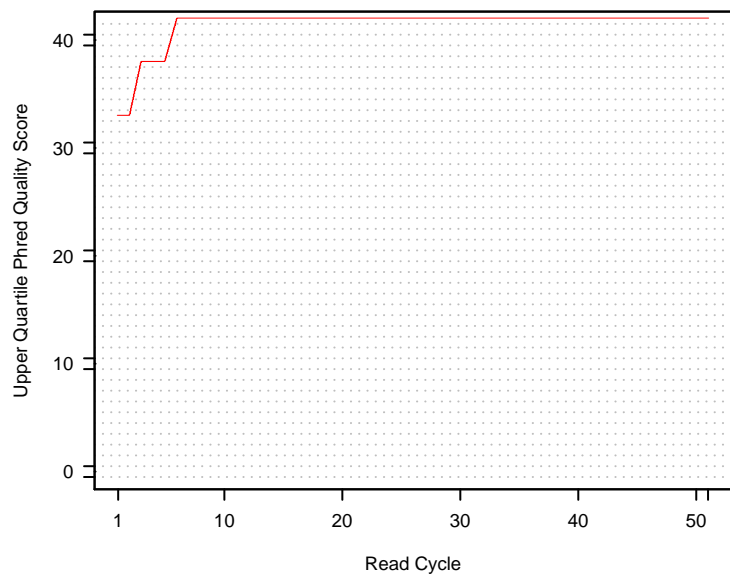
d

Median Phred Quality Score
With Sample Untitled in red



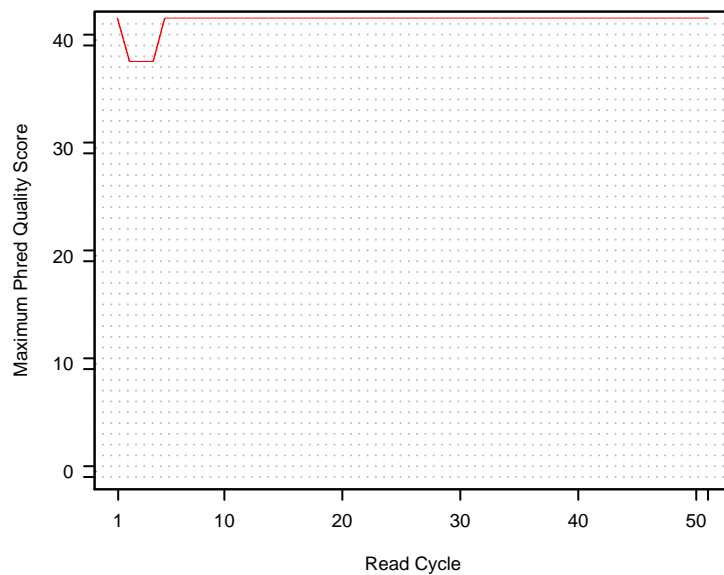
e

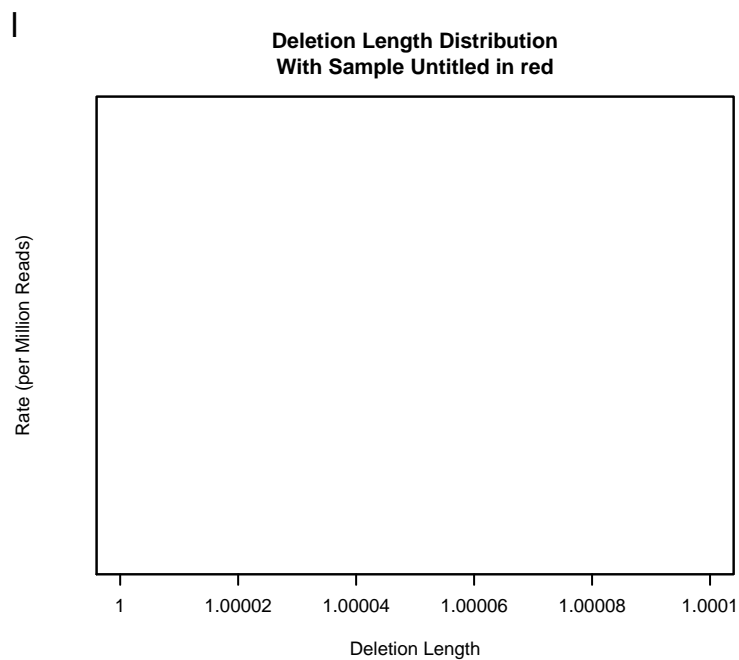
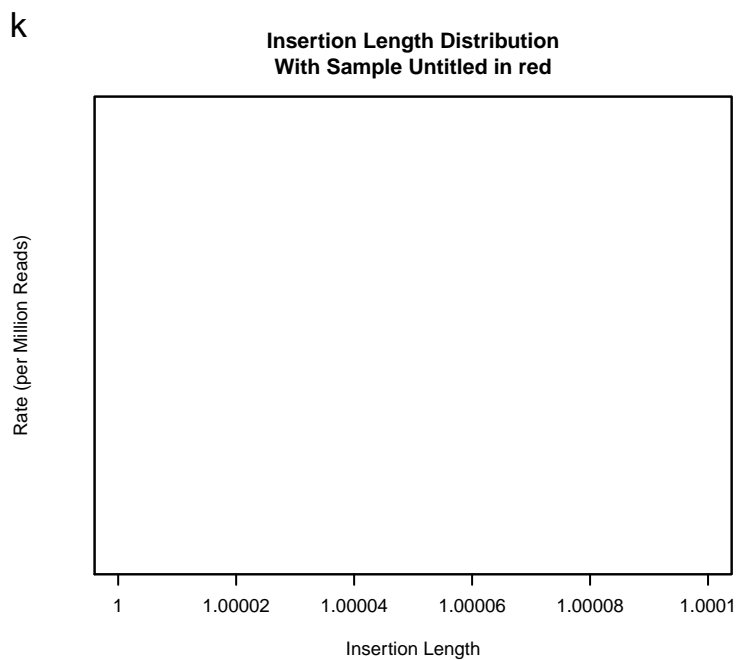
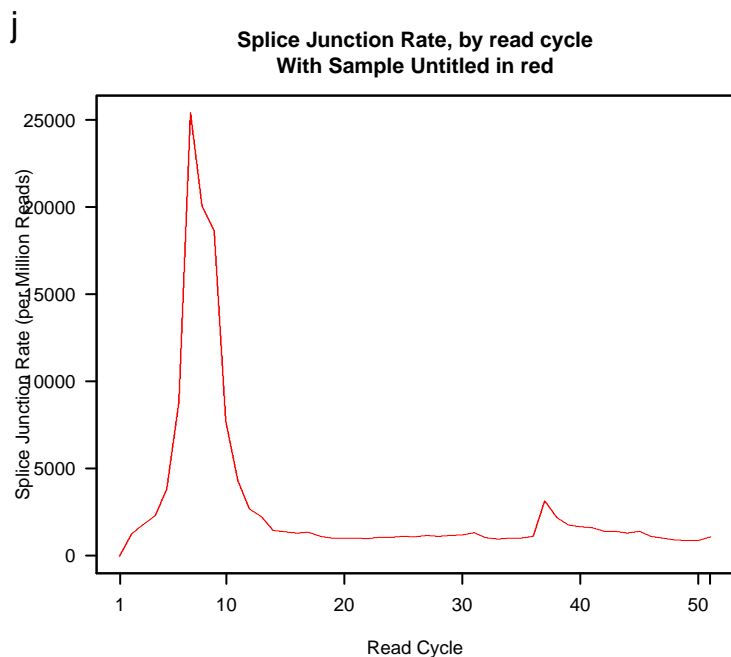
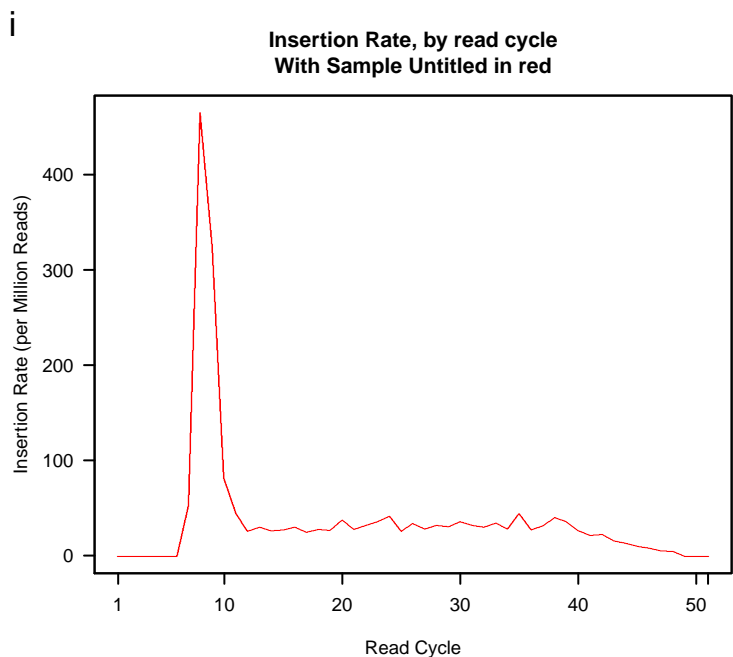
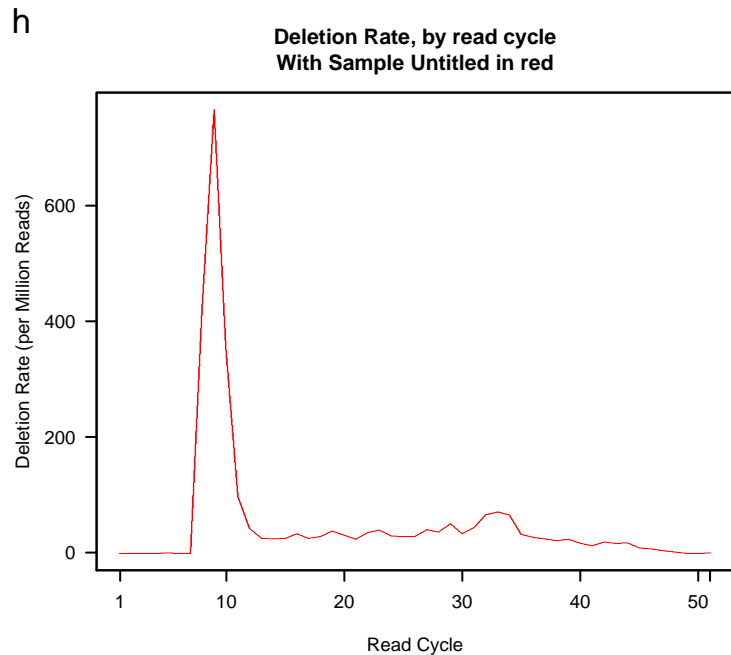
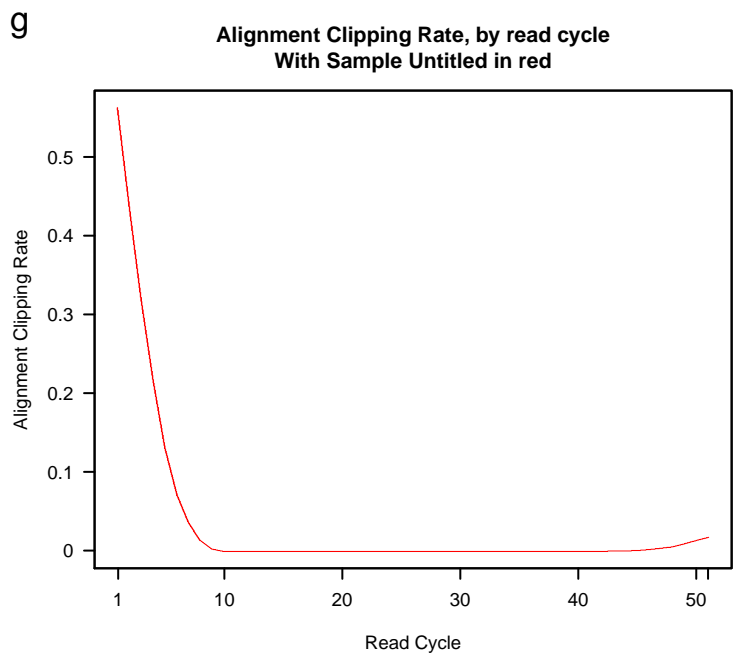
Upper Quartile Phred Quality Score
With Sample Untitled in red

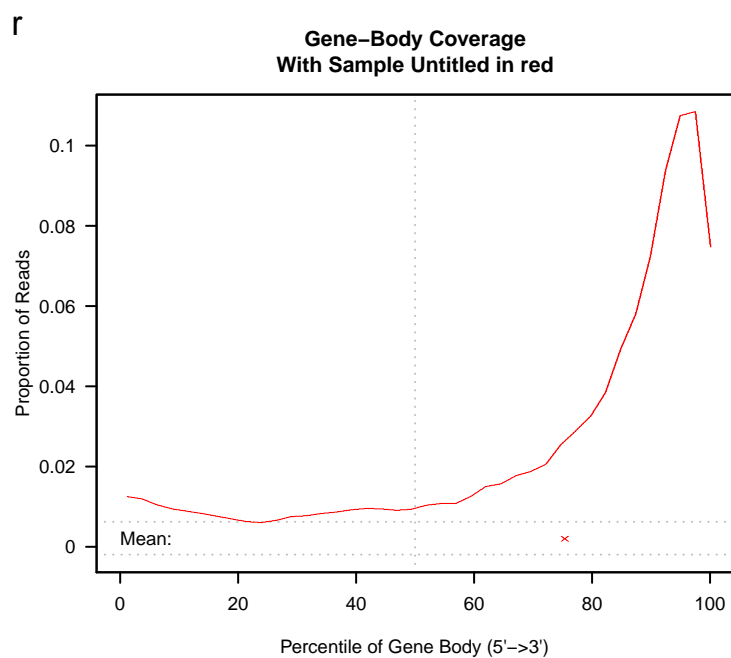
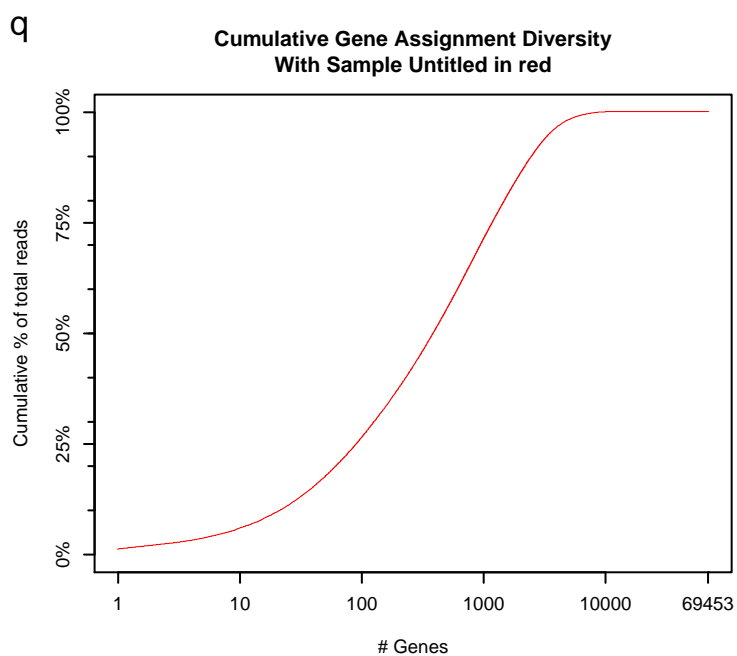
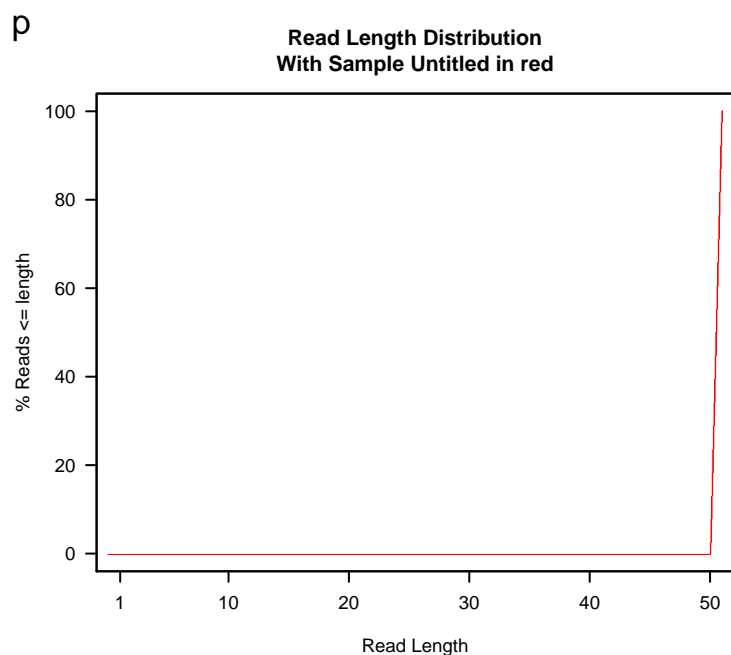
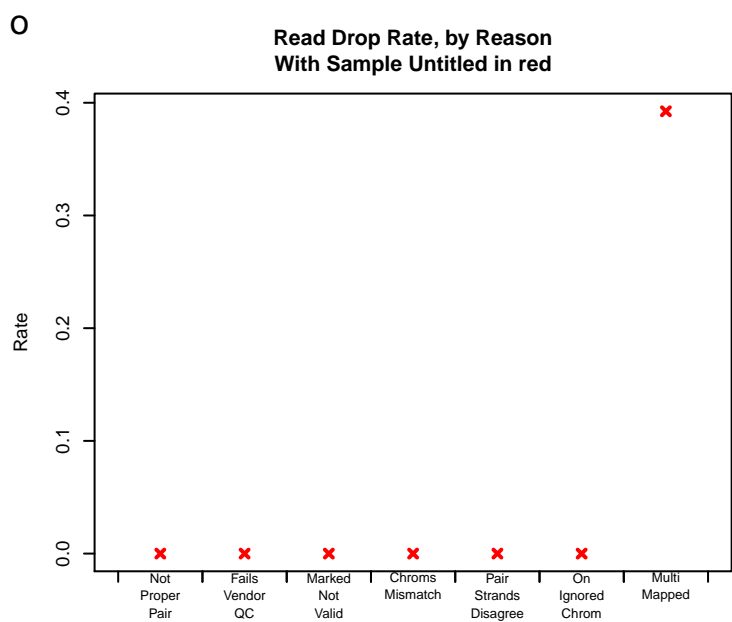
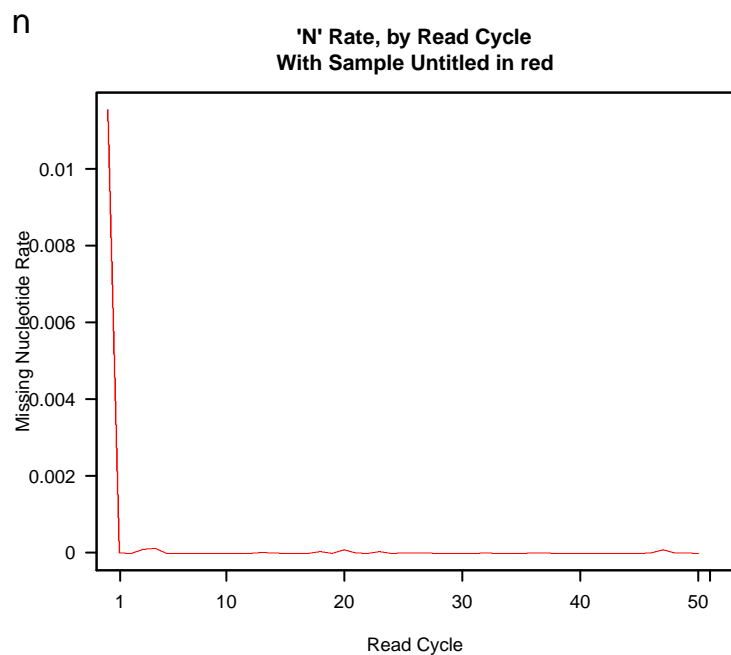
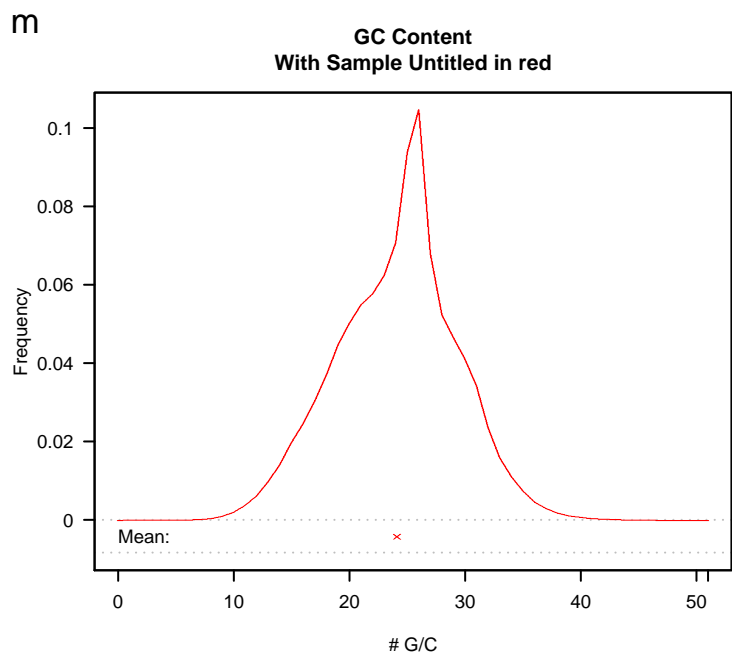


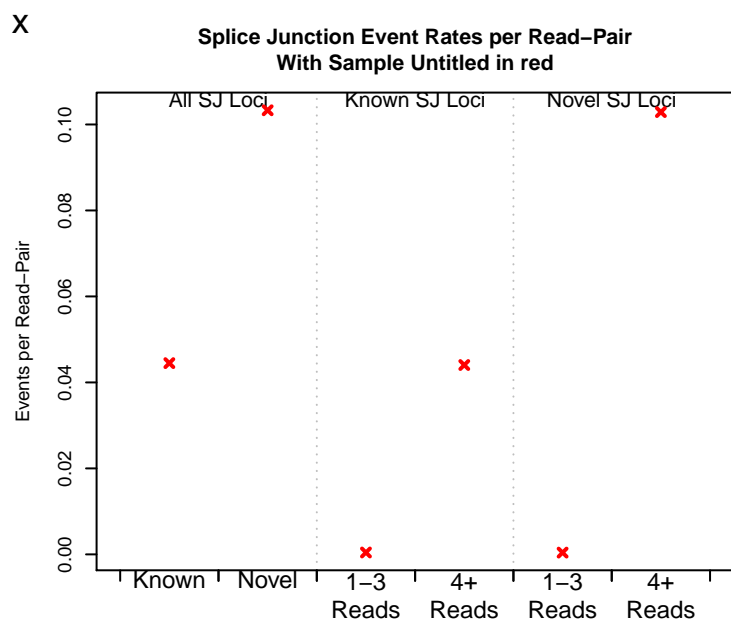
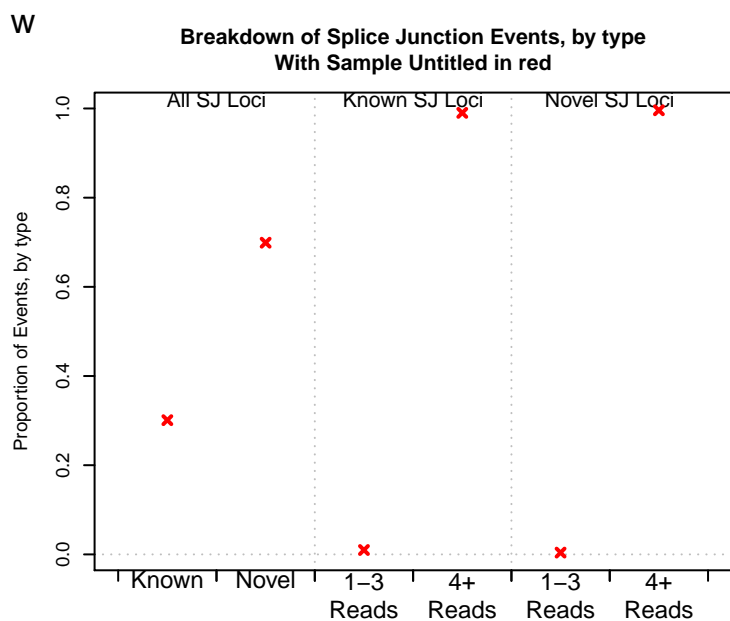
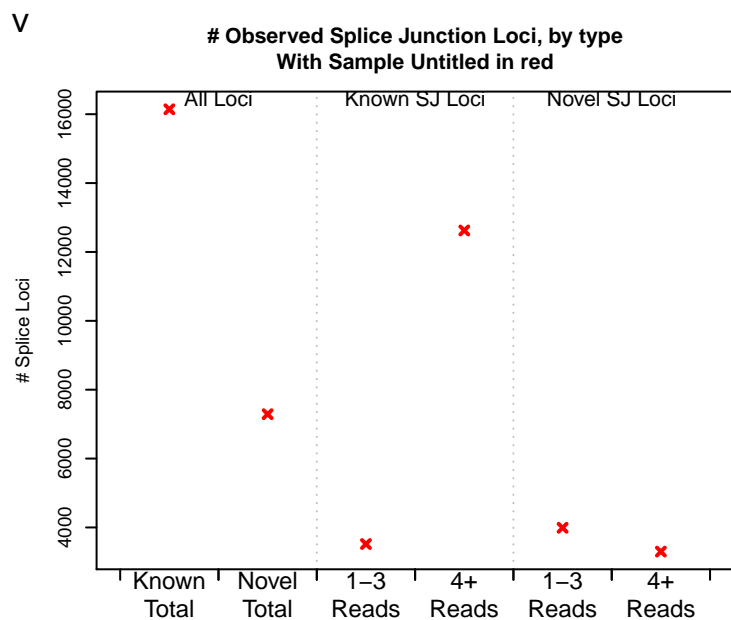
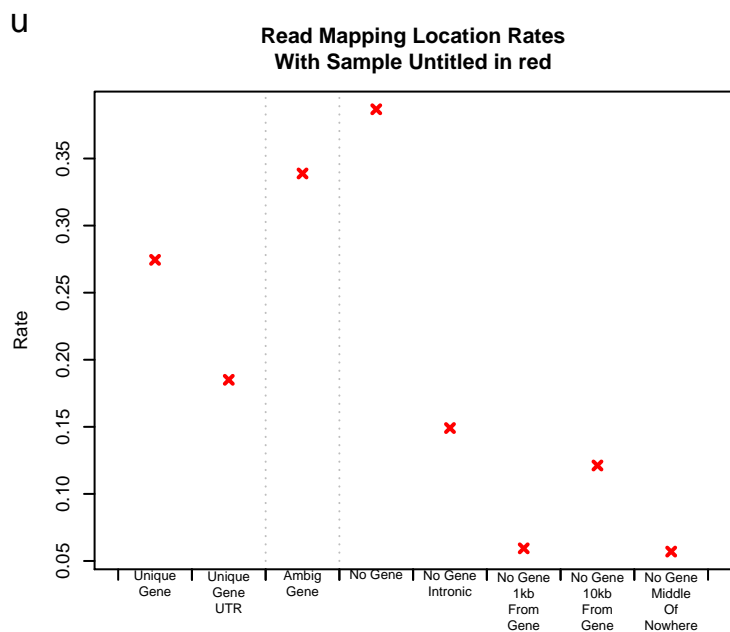
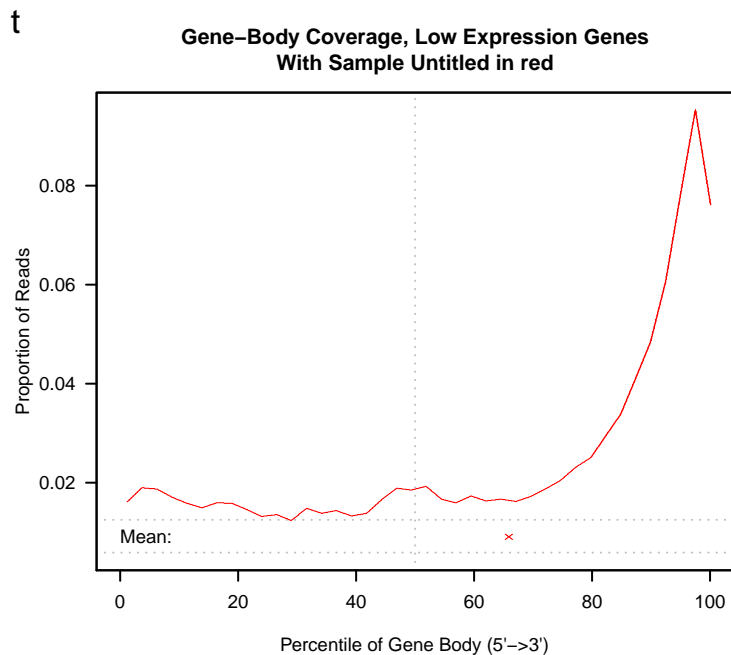
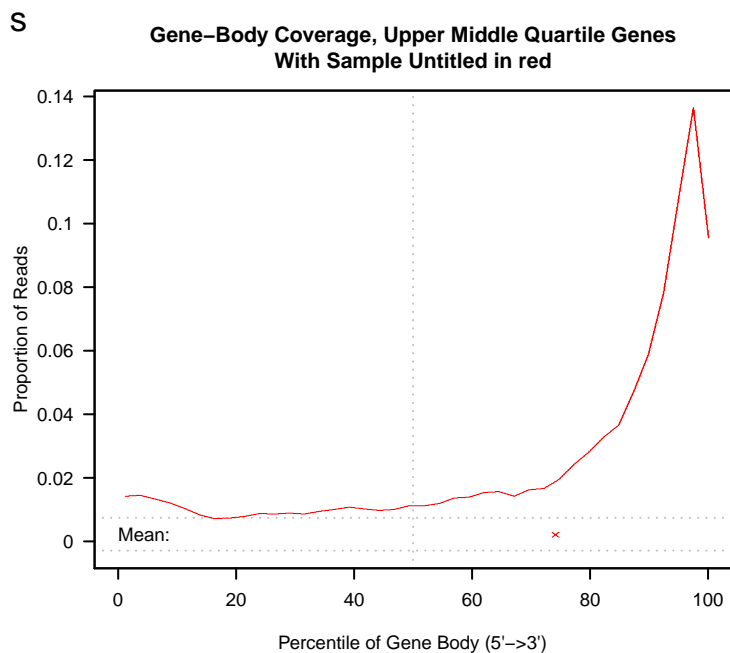
f

Maximum Phred Quality Score
With Sample Untitled in red

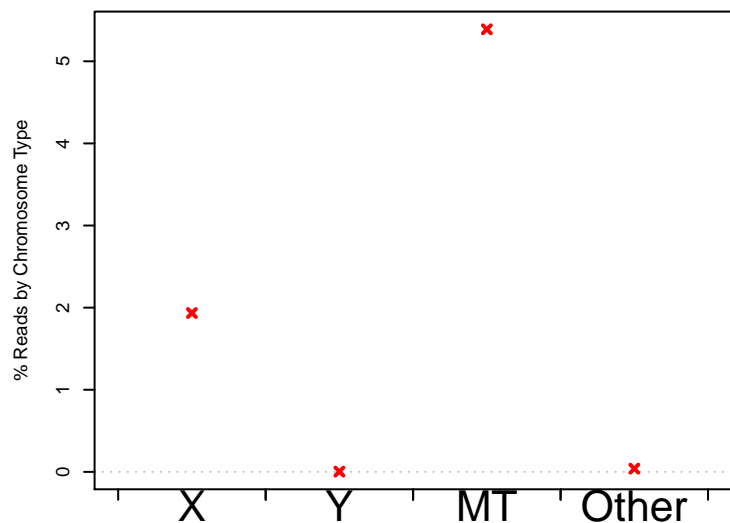




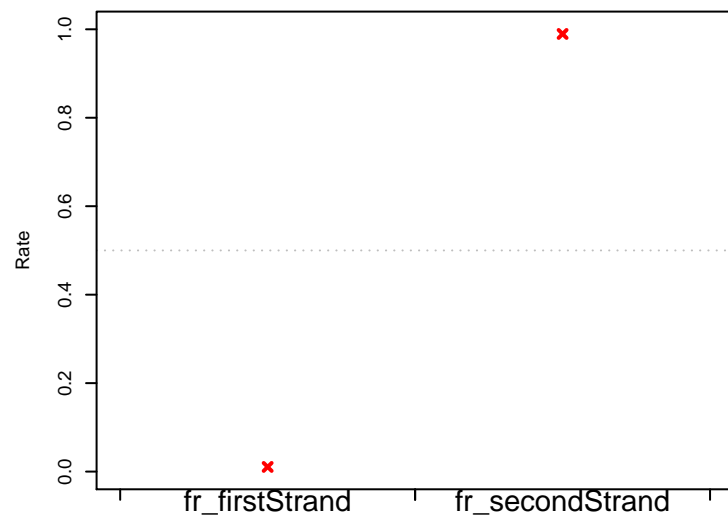




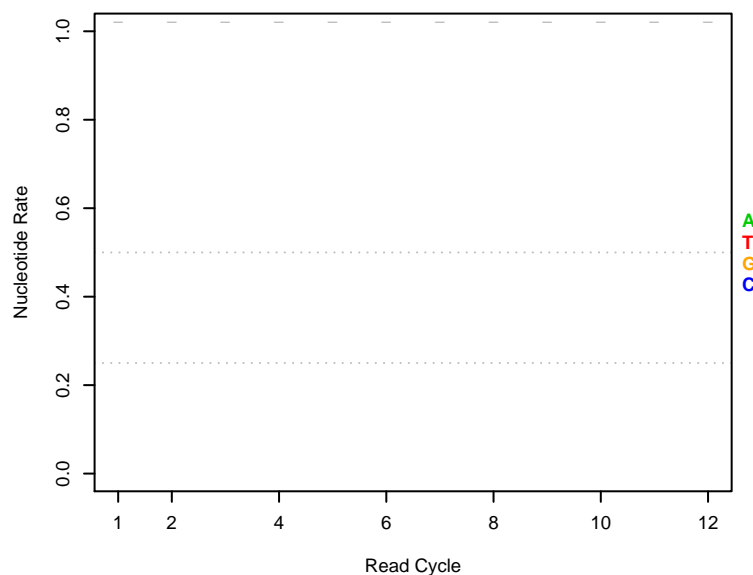
y **Chromosome Distribution (Excluding Autosomes)**
With Sample Untitled in red



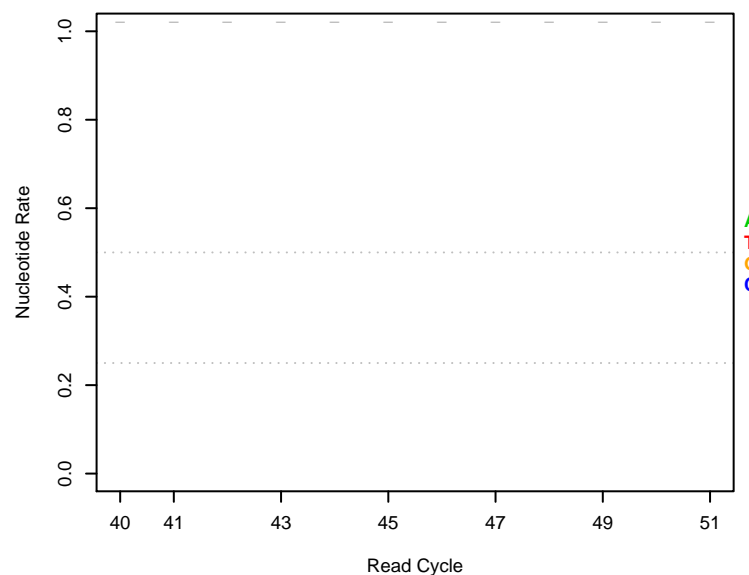
Z **Strandedness Test**
With Sample Untitled in red



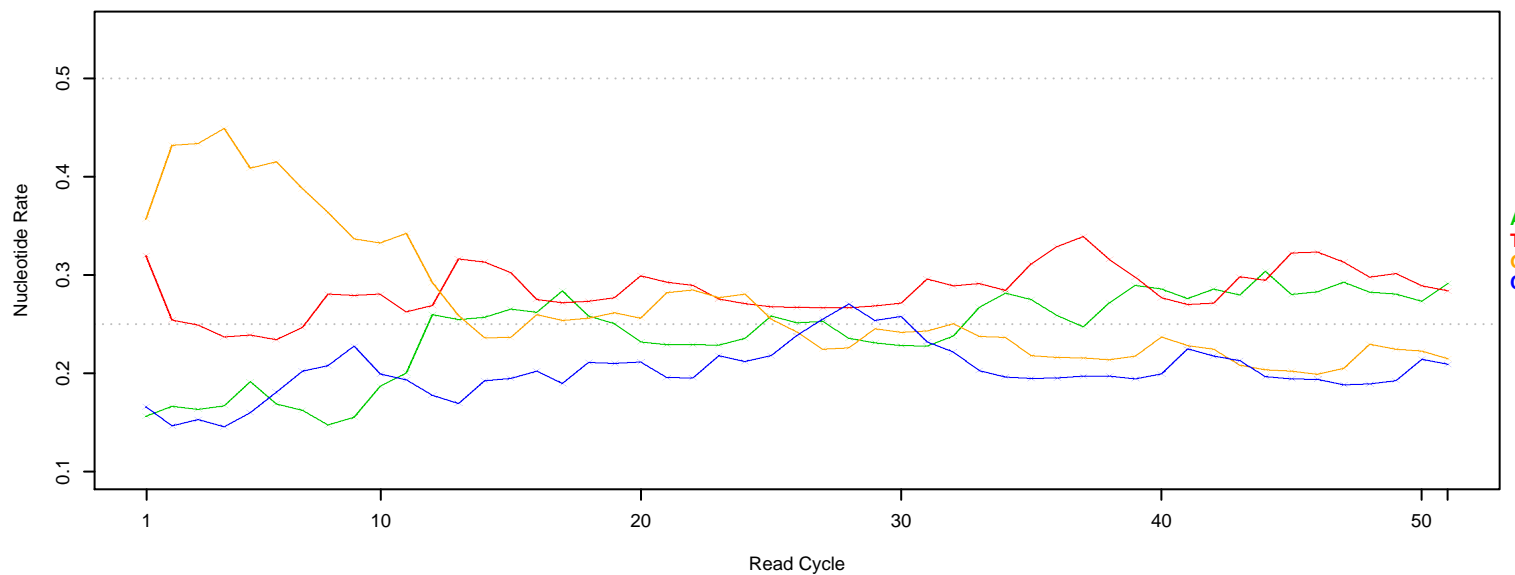
aa **Nucleotide Rate by Cycle, Leading Clipped bases (12)**
With Sample Untitled in red



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



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



ad

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

