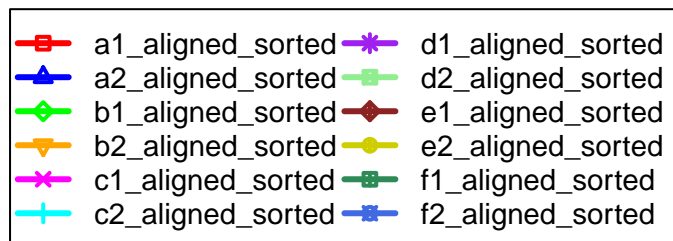


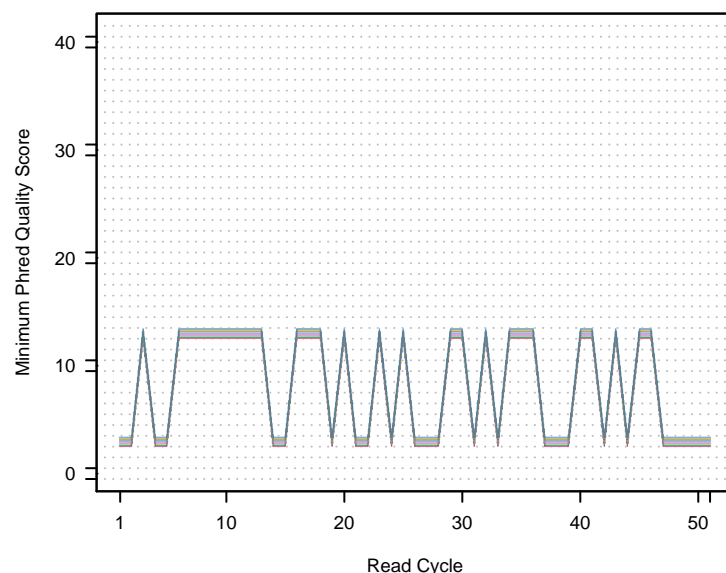
a

Summary Plots, By Sample



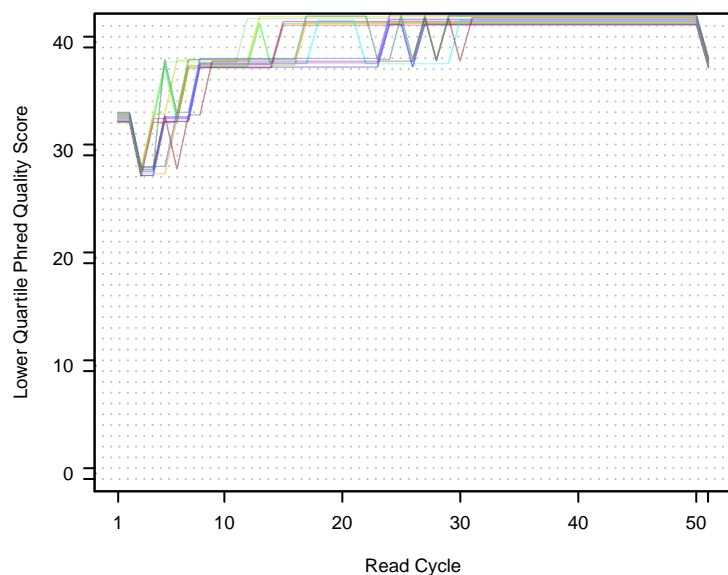
b

**Minimum Phred Quality Score
Colored by Sample**



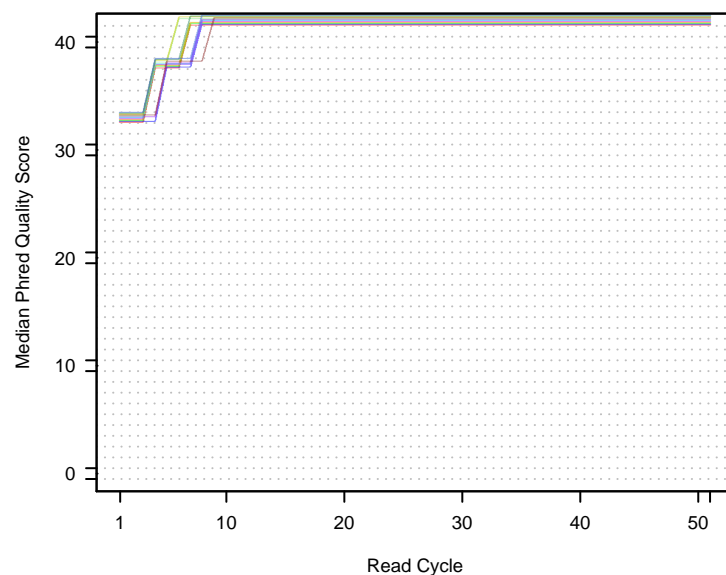
c

**Lower Quartile Phred Quality Score
Colored by Sample**



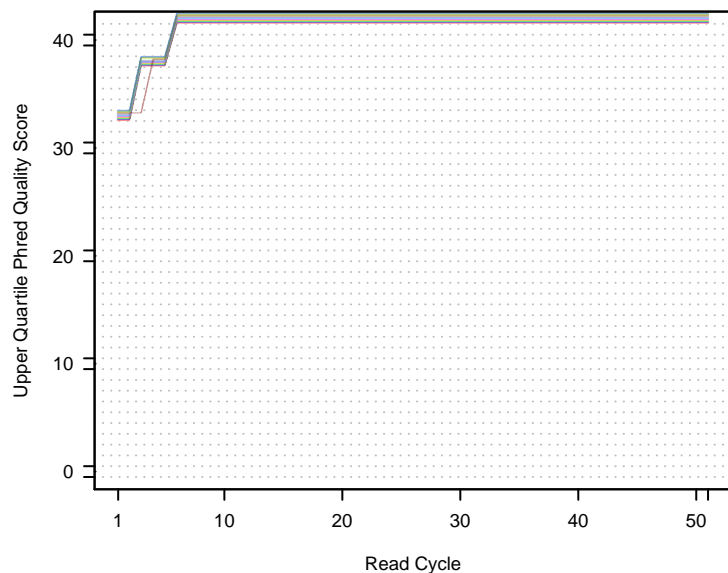
d

**Median Phred Quality Score
Colored by Sample**



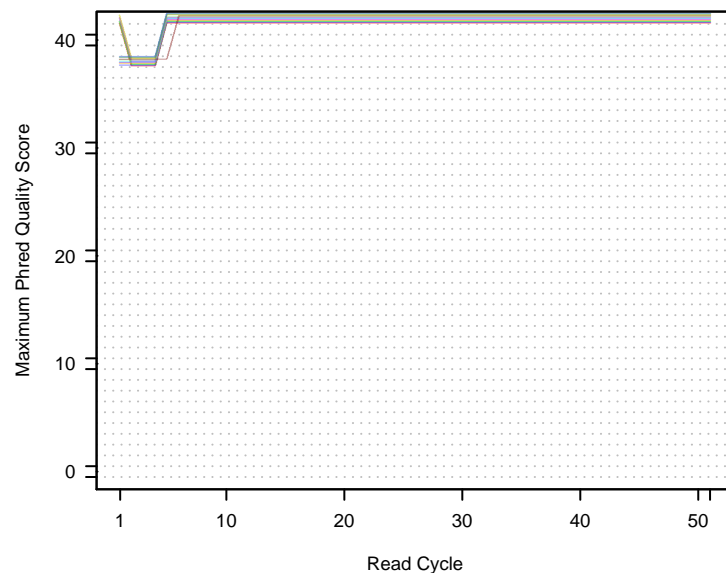
e

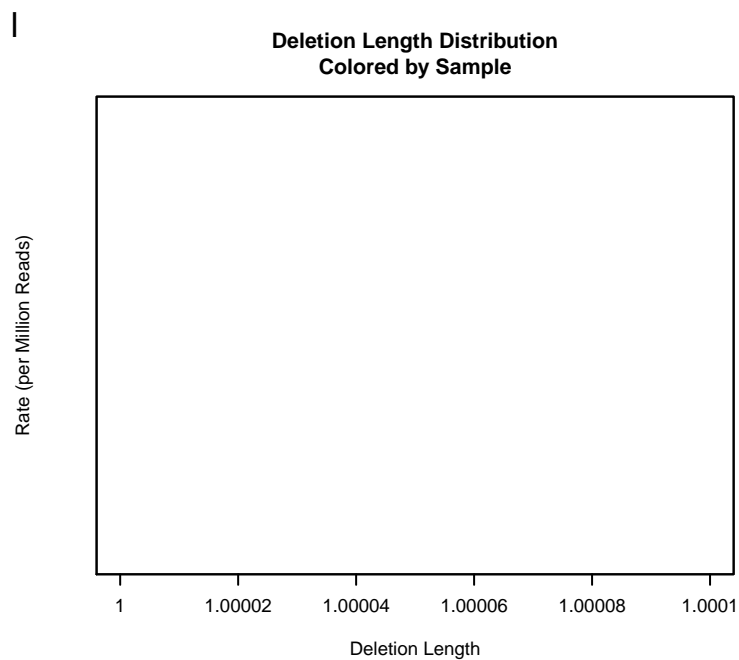
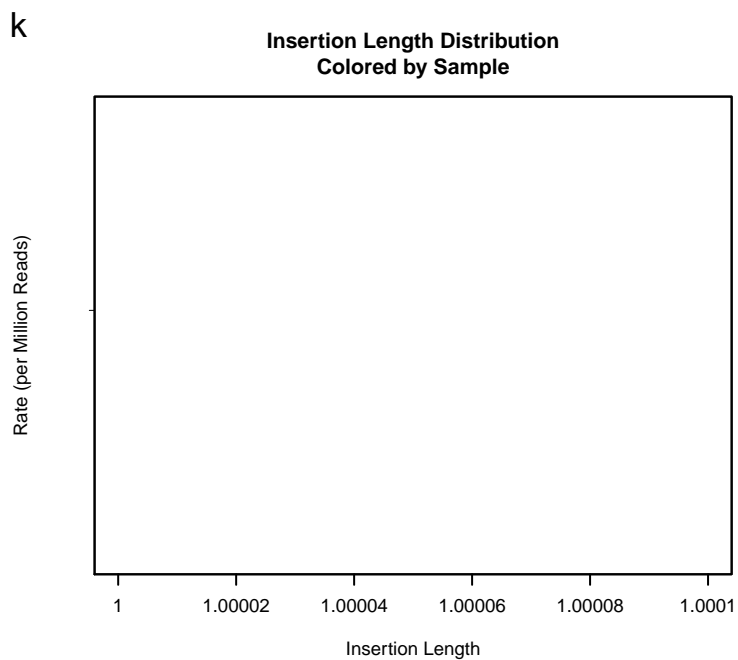
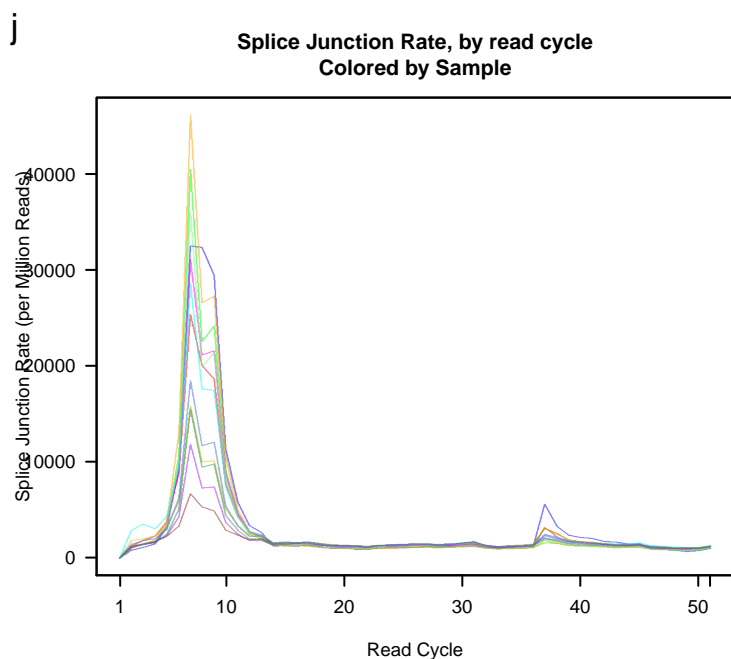
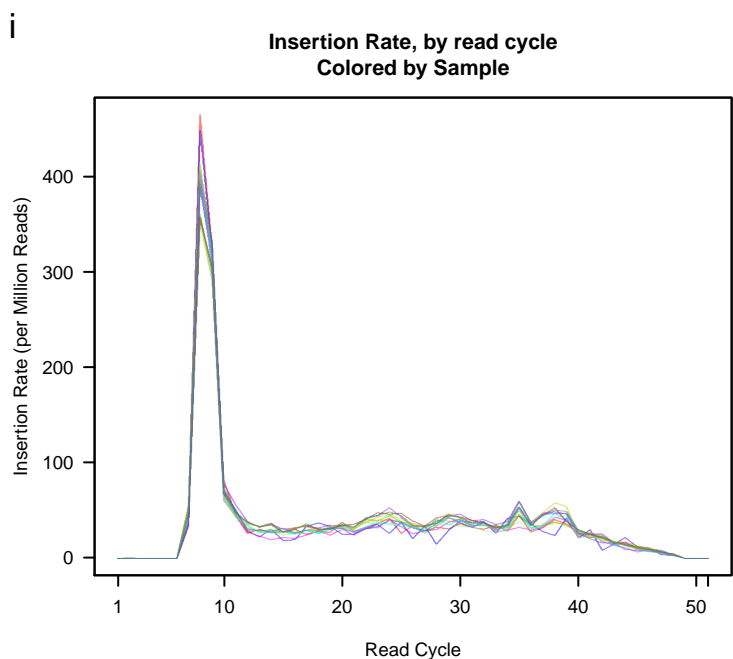
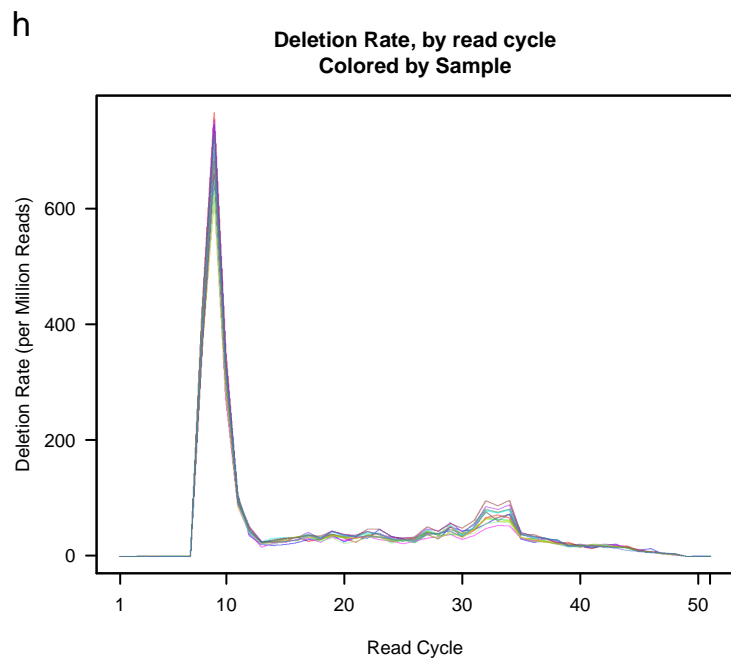
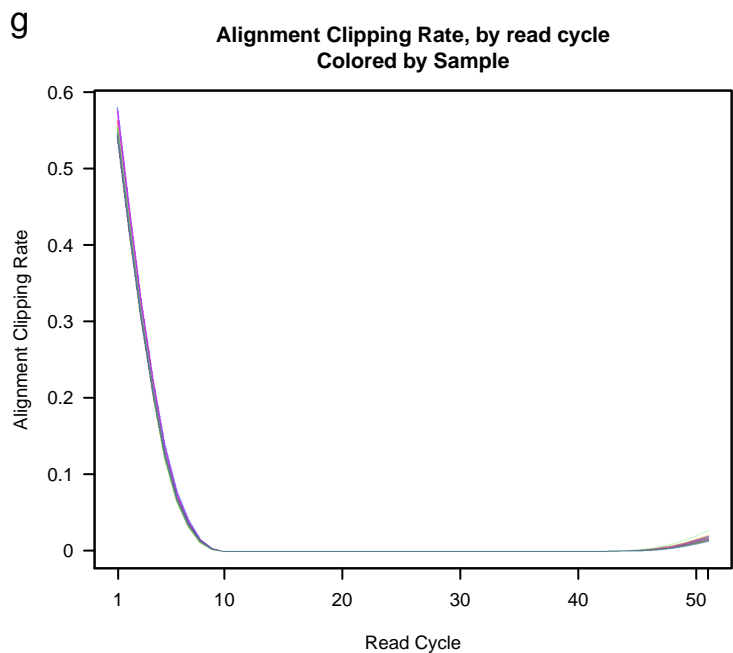
**Upper Quartile Phred Quality Score
Colored by Sample**

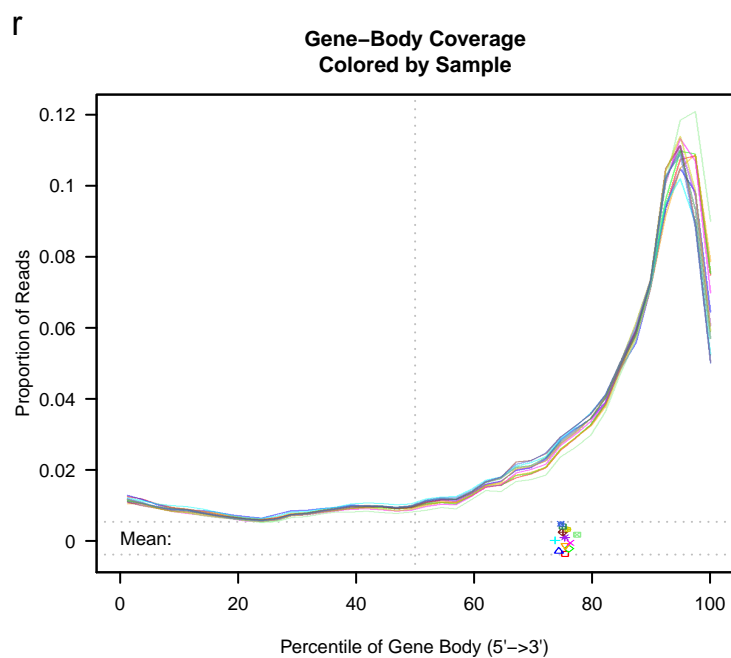
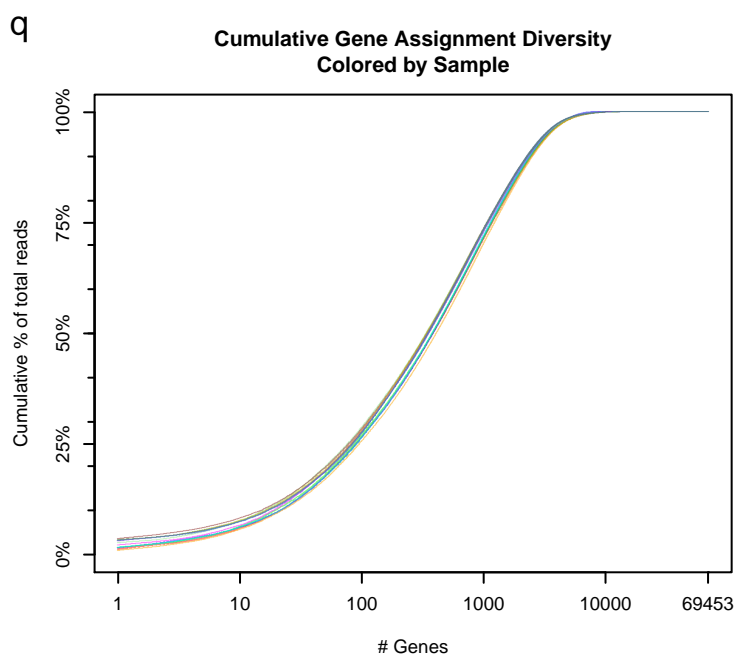
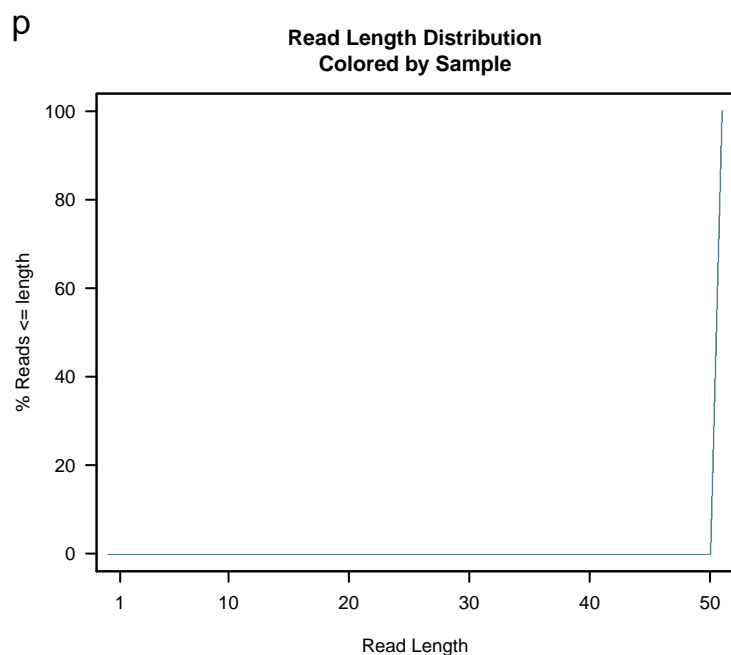
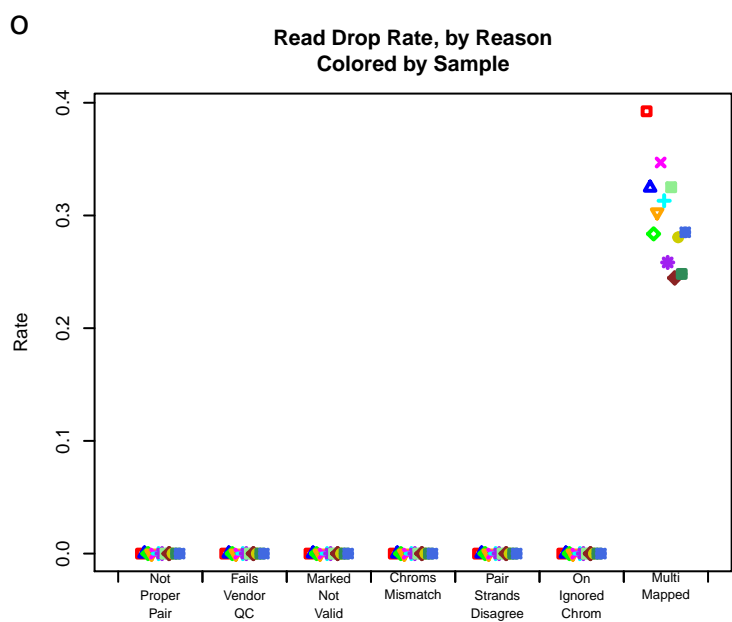
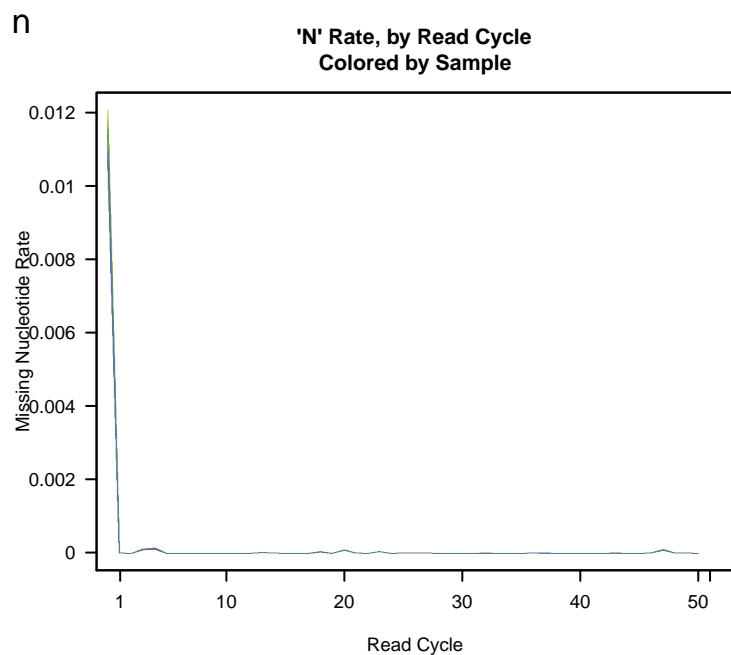
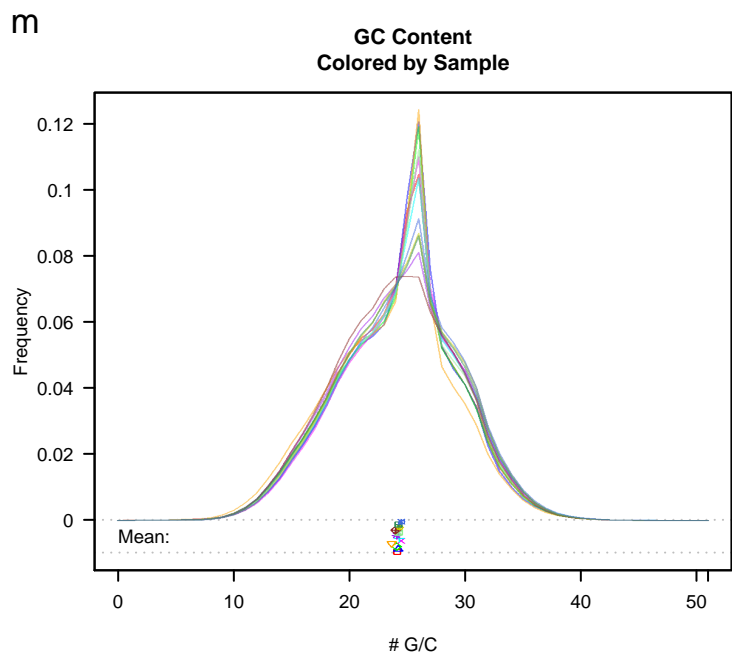


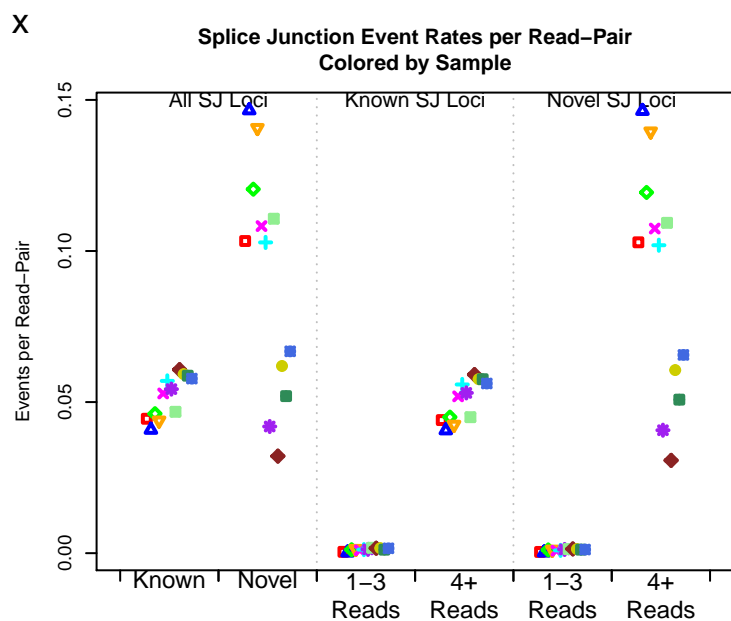
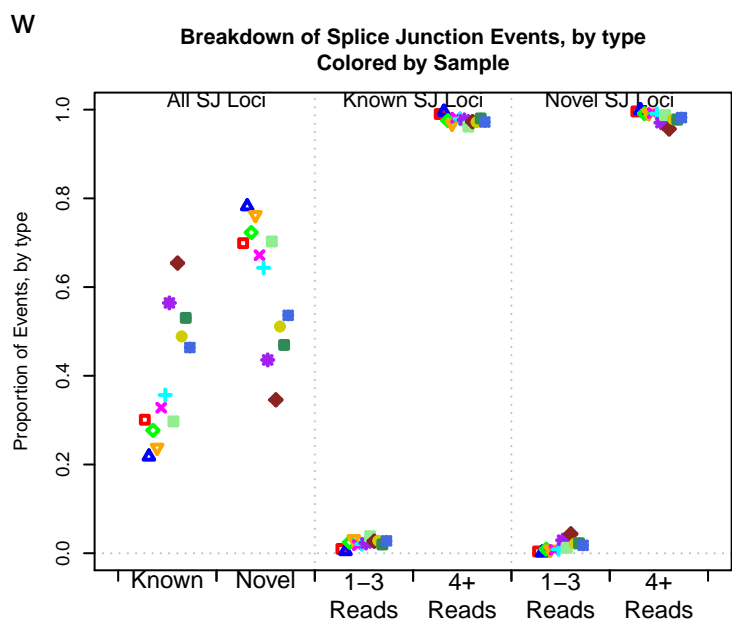
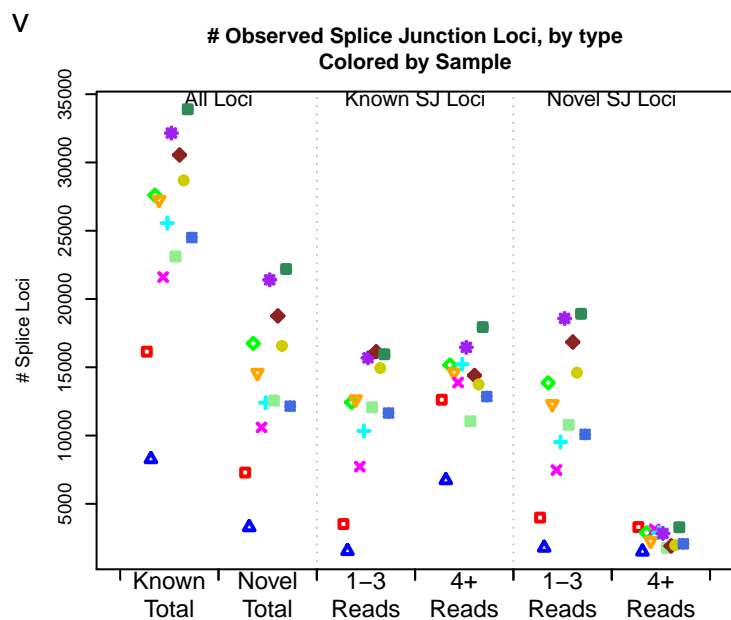
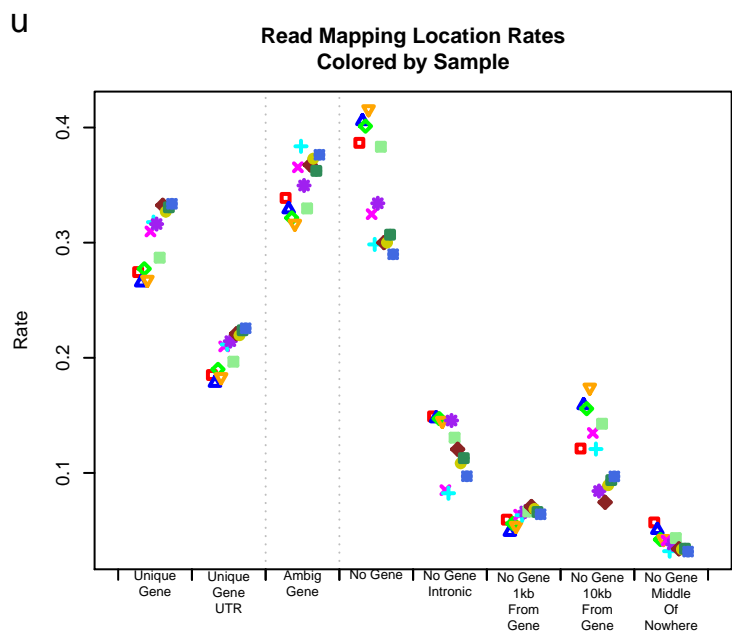
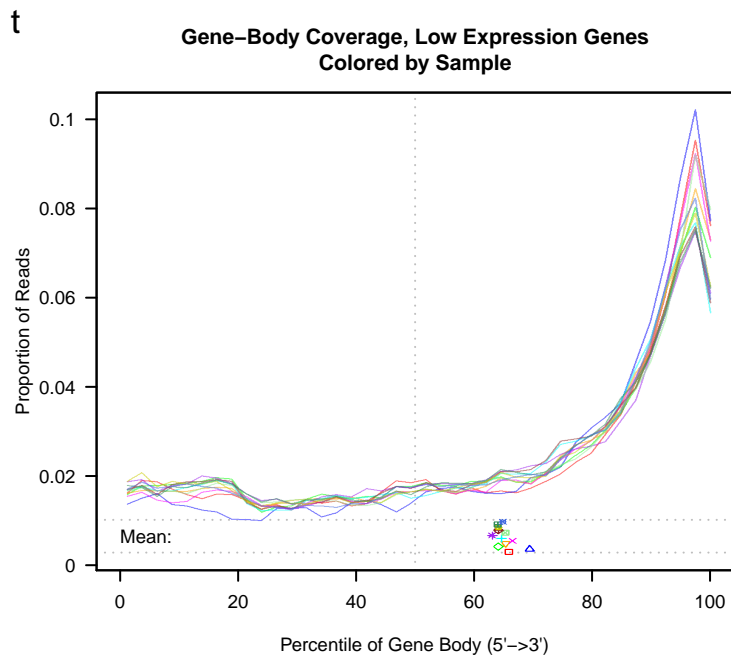
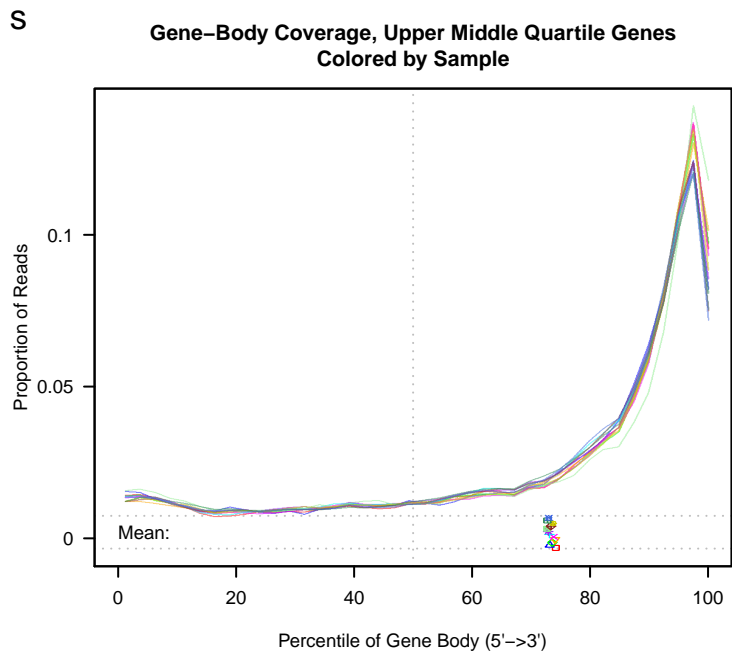
f

**Maximum Phred Quality Score
Colored by Sample**



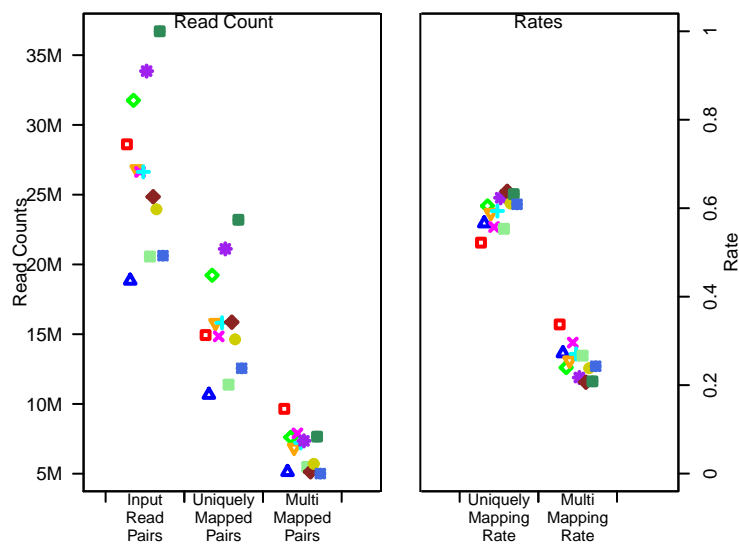






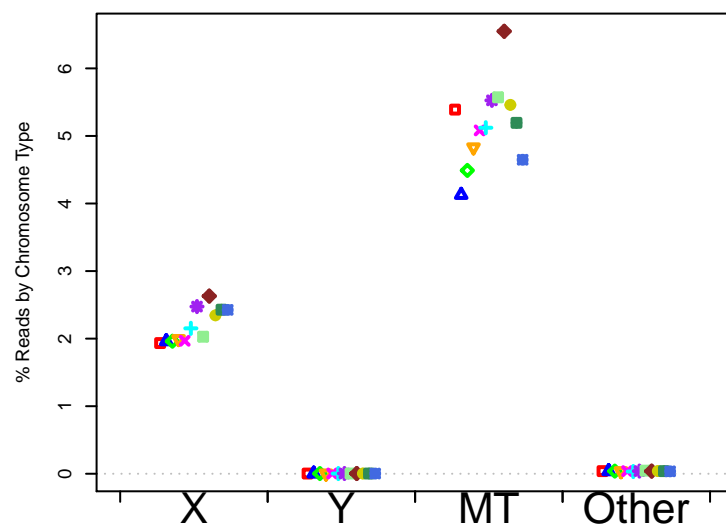
y

Mapping Stats
Colored by Sample



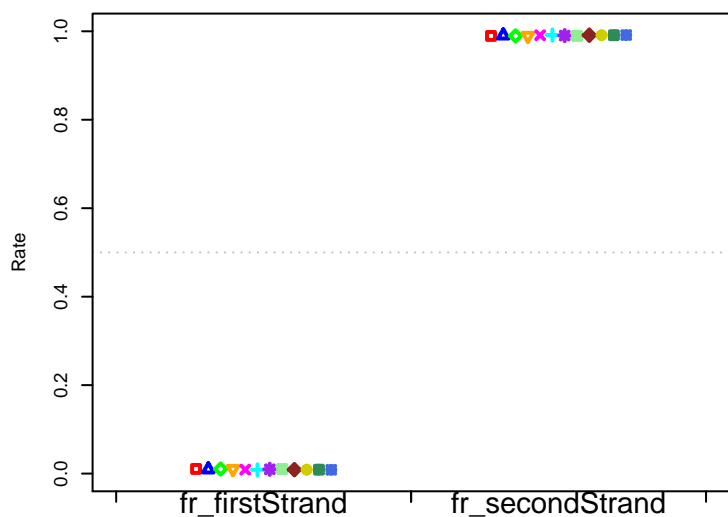
Z

Chromosome Distribution (Excluding Autosomes)
Colored by Sample



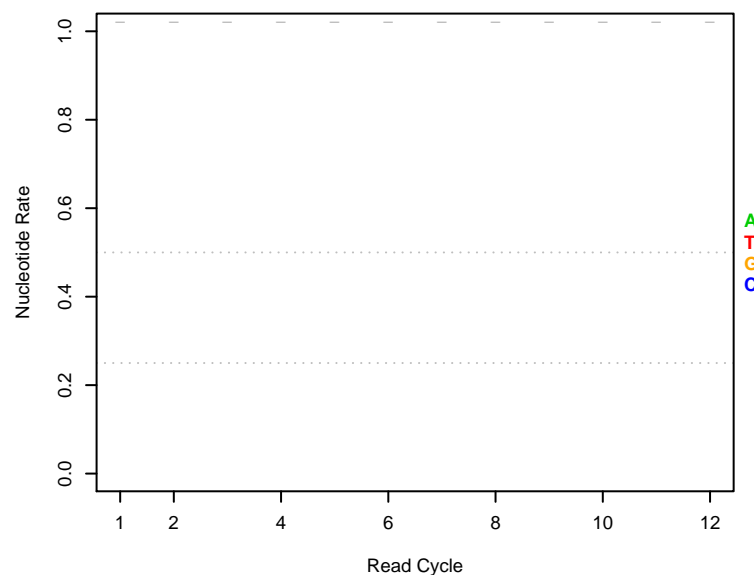
aa

Strandedness Test
Colored by Sample



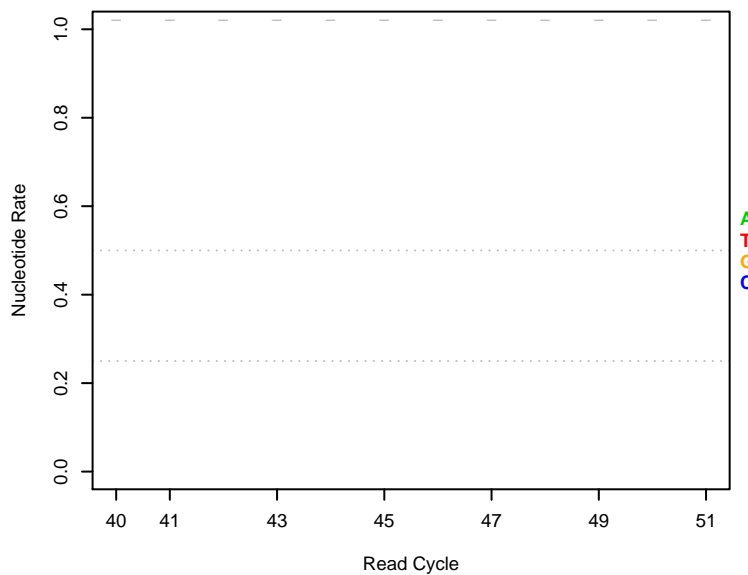
ab

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

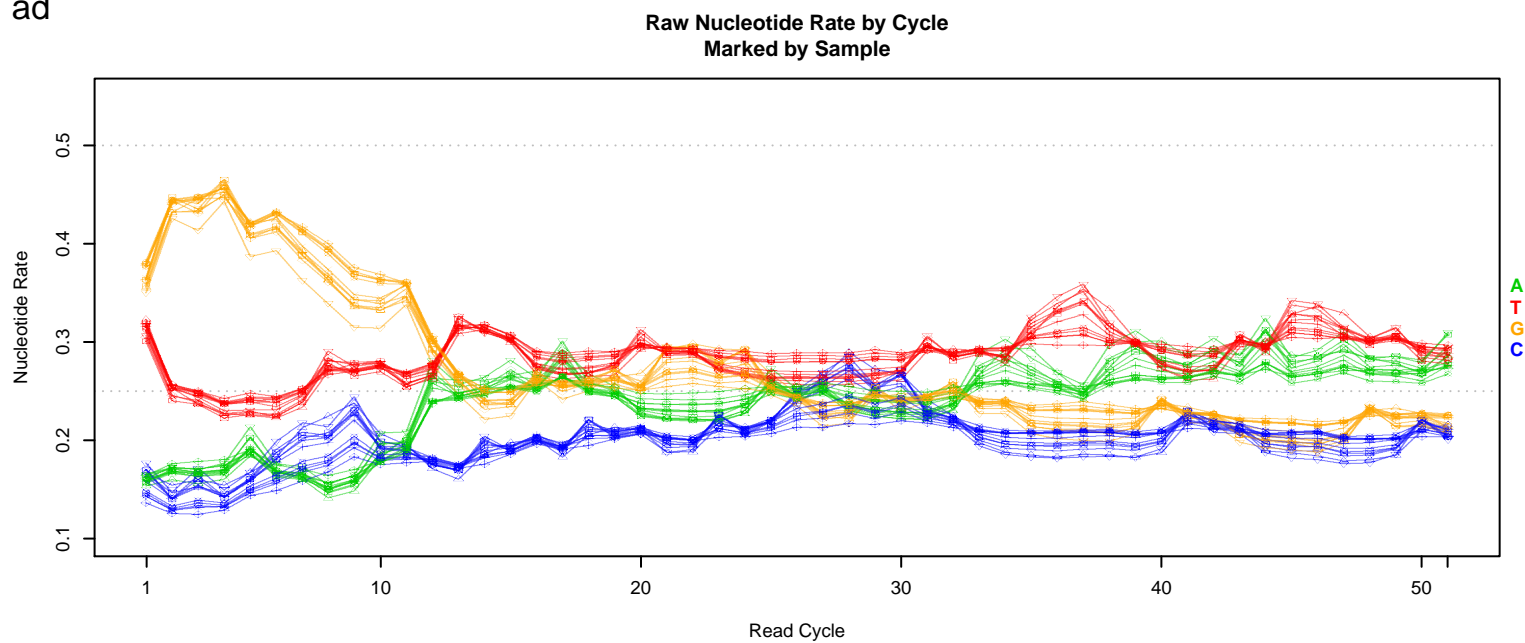


ac

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



ad



ae

