a

Sample Highlight:
f1_aligned_sorted
Colored by Lane

—□—	UNKNOWN
⋯○⋯	Other Samples

b

Minimum Phred Quality Score
With Sample f1_aligned_sorted Colored by lane
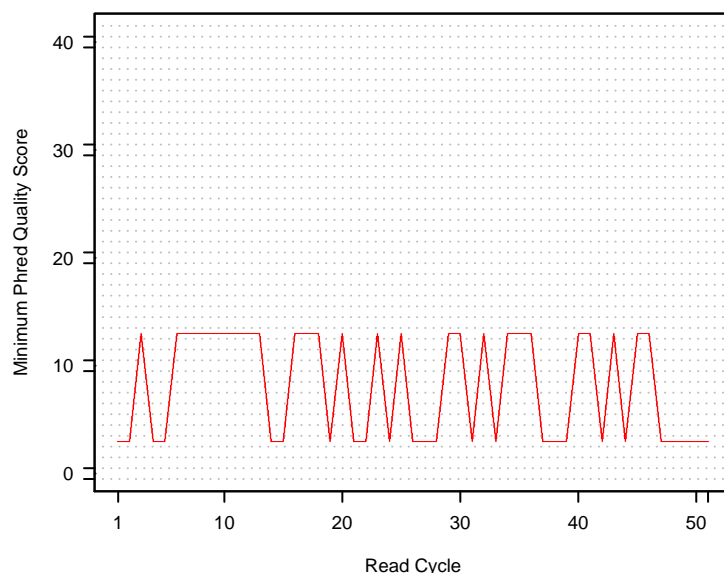
c

Lower Quartile Phred Quality Score
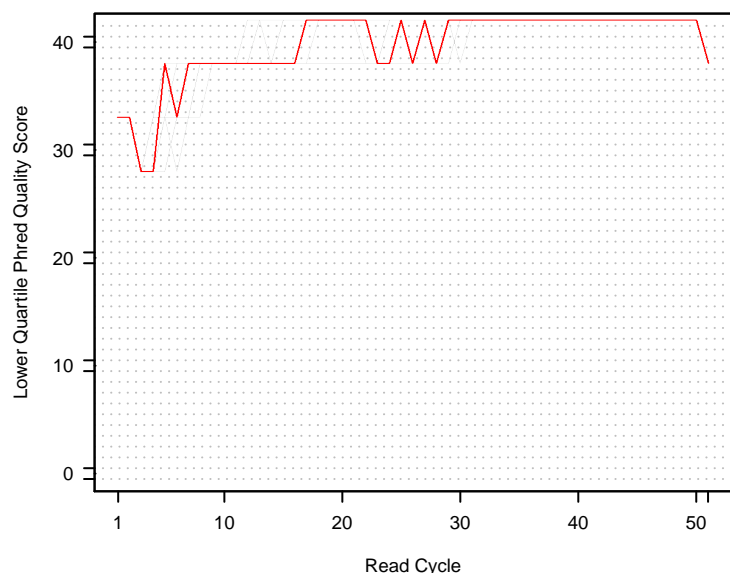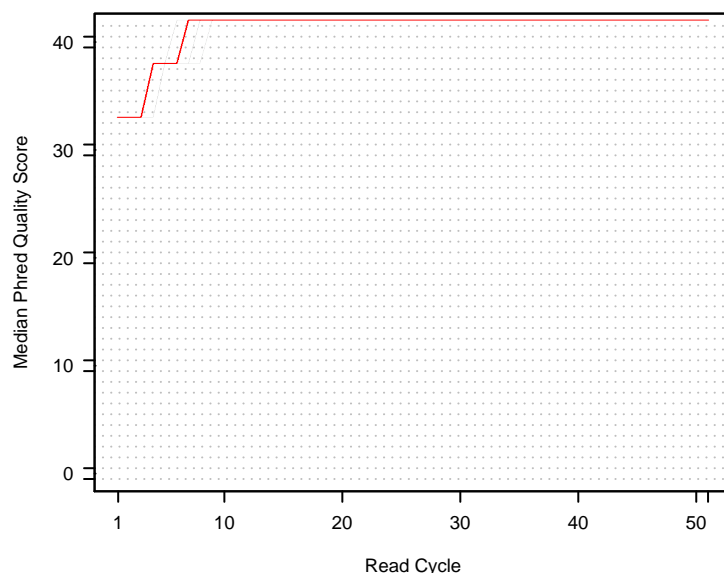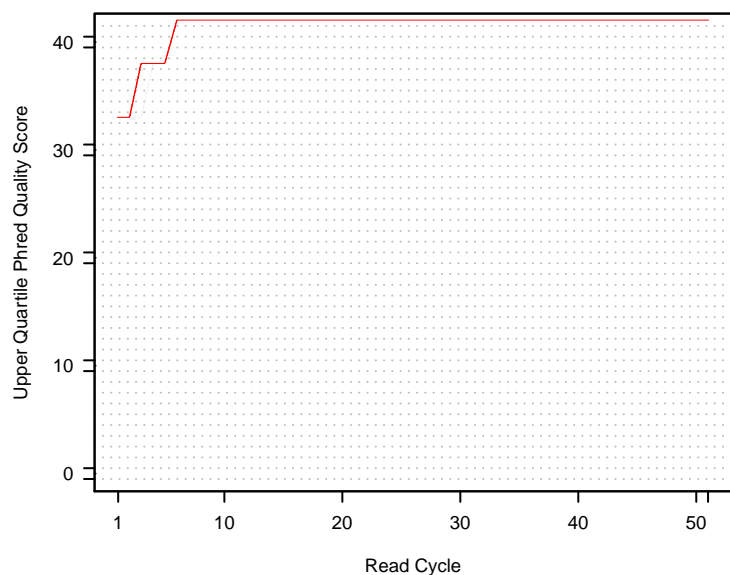**With Sample f1_aligned_sorted Colored by lane**

d

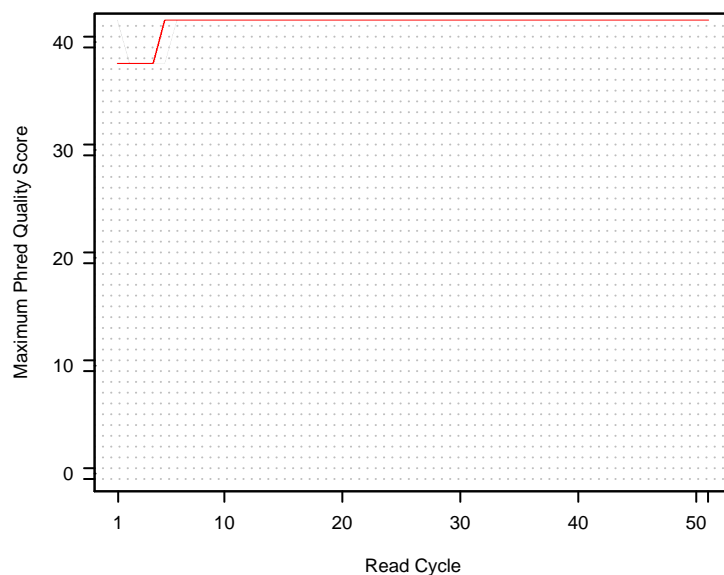**Median Phred Quality Score**
With Sample f1_aligned_sorted Colored by lane

e

Upper Quartile Phred Quality Score
With Sample f1_aligned_sorted Colored by lane

f

Maximum Phred Quality Score
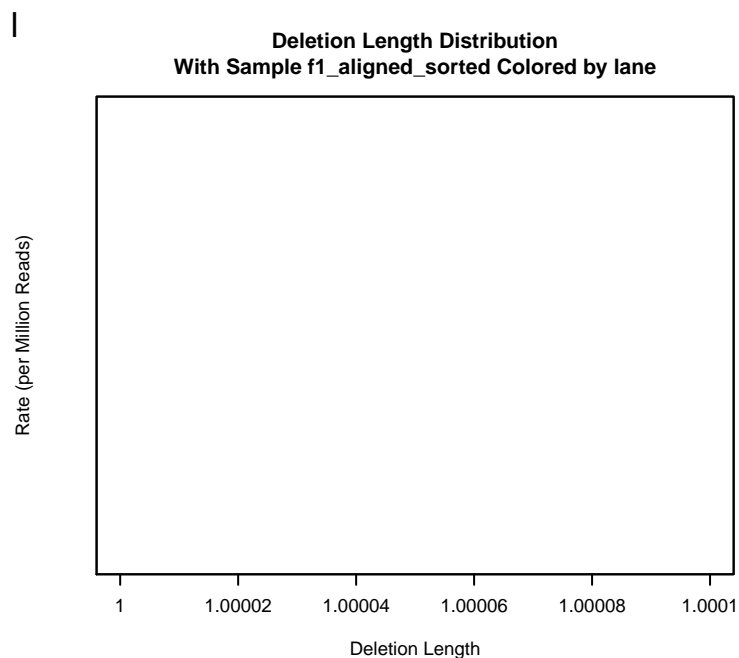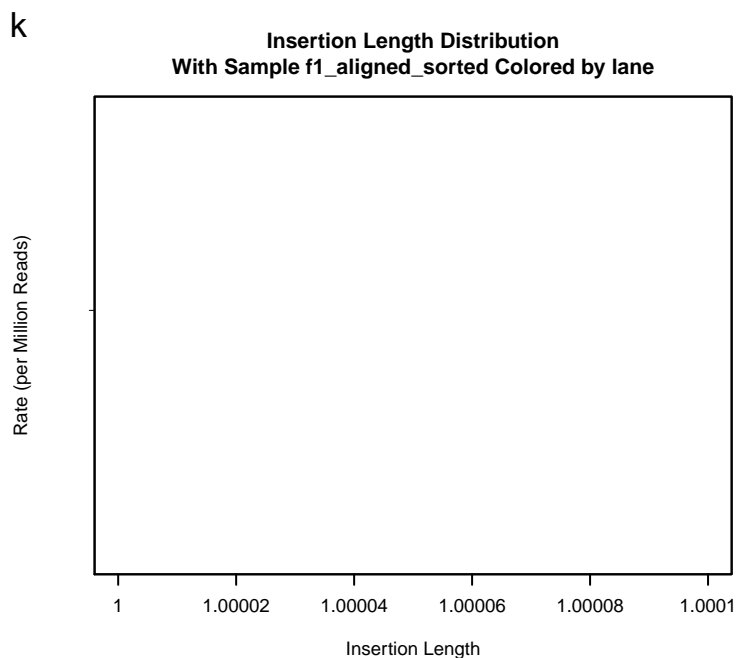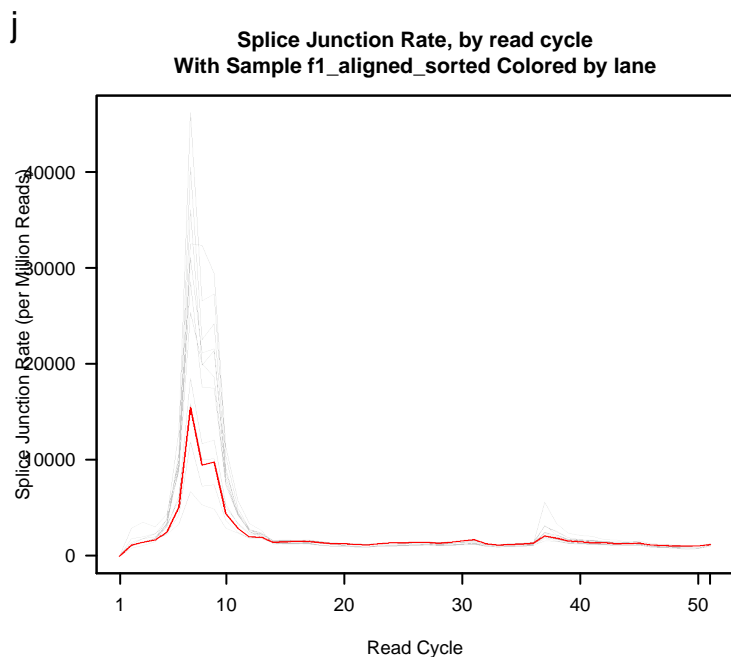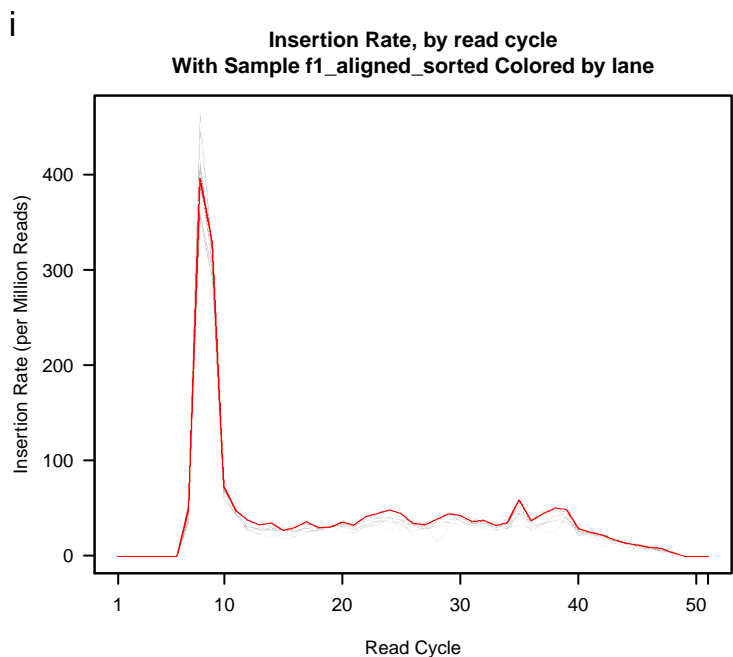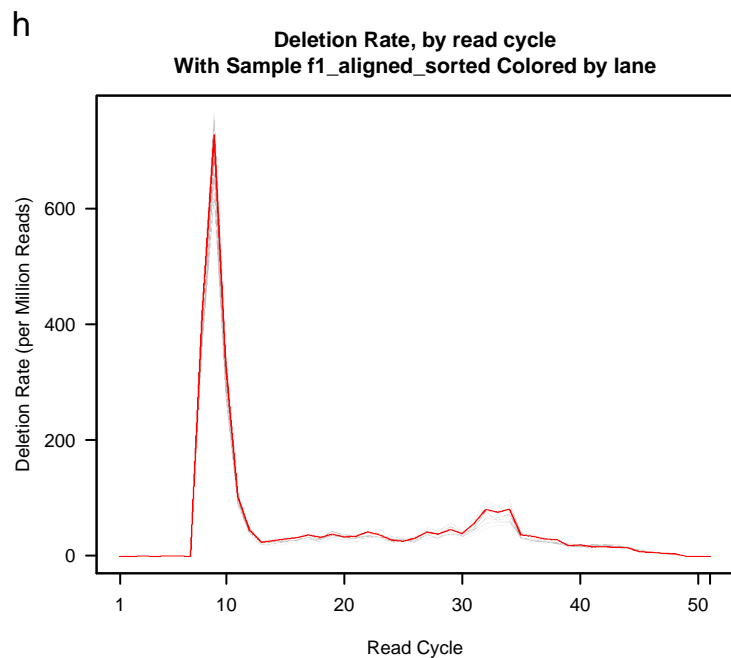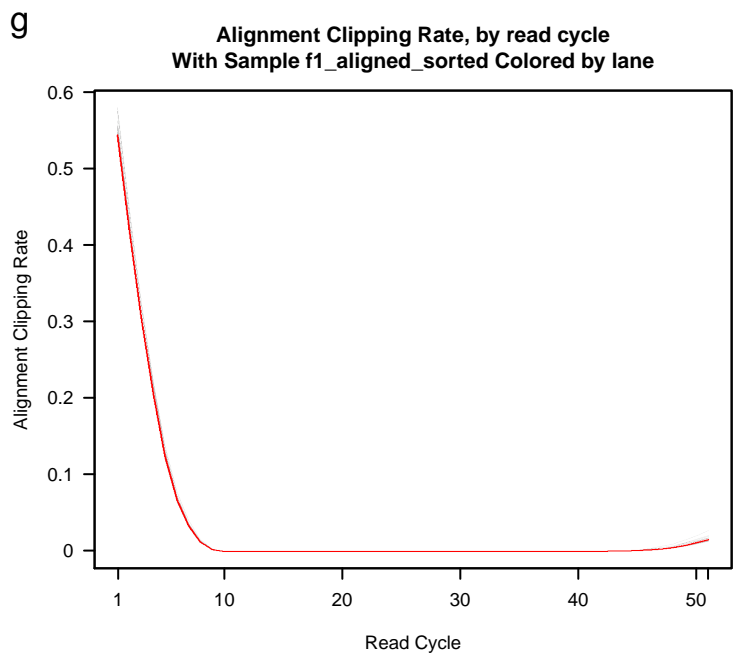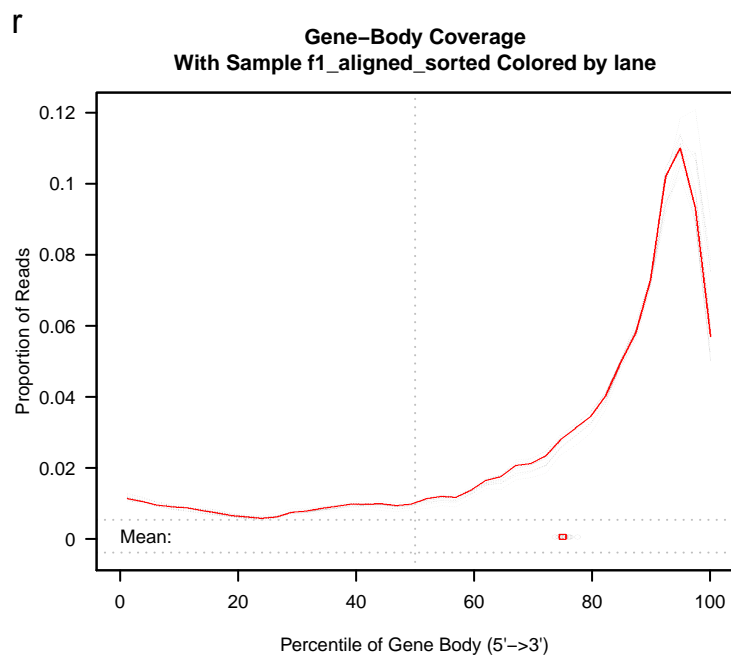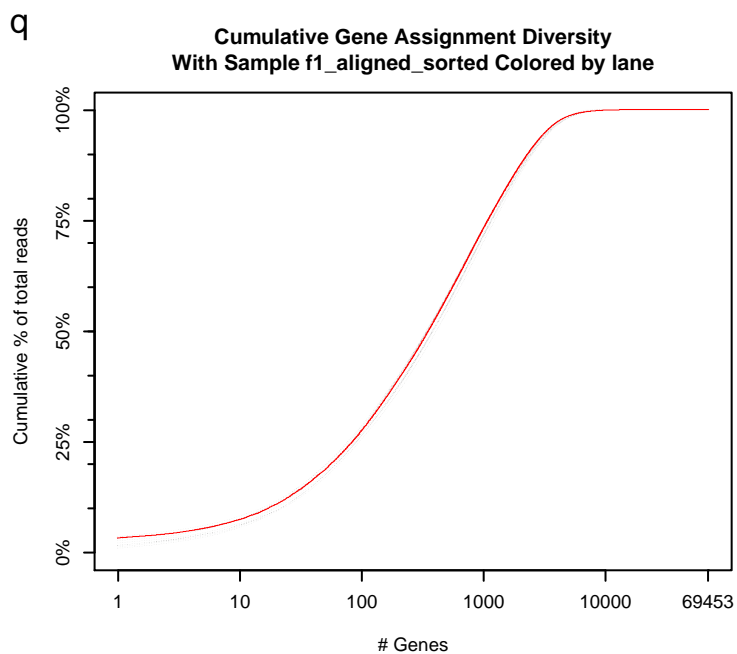**With Sample f1_aligned_sorted Colored by lane**

g

Alignment Clipping Rate, by read cycle
With Sample f1_aligned_sorted Colored by lane

h

Deletion Rate, by read cycle
With Sample f1_aligned_sorted Colored by lane

i

Insertion Rate, by read cycle
With Sample f1_aligned_sorted Colored by lane

j

Splice Junction Rate, by read cycle
With Sample f1_aligned_sorted Colored by lane

k

Insertion Length Distribution
With Sample f1_aligned_sorted Colored by lane

l

Deletion Length Distribution
With Sample f1_aligned_sorted Colored by lane

m GC Content
With Sample f1_aligned_sorted Colored by lane

n 'N' Rate, by Read Cycle
With Sample f1_aligned_sorted Colored by lane

o Read Drop Rate, by Reason
With Sample f1_aligned_sorted Colored by lane

p Read Length Distribution
With Sample f1_aligned_sorted Colored by lane

q Cumulative Gene Assignment Diversity
With Sample f1_aligned_sorted Colored by lane

r Gene−Body Coverage
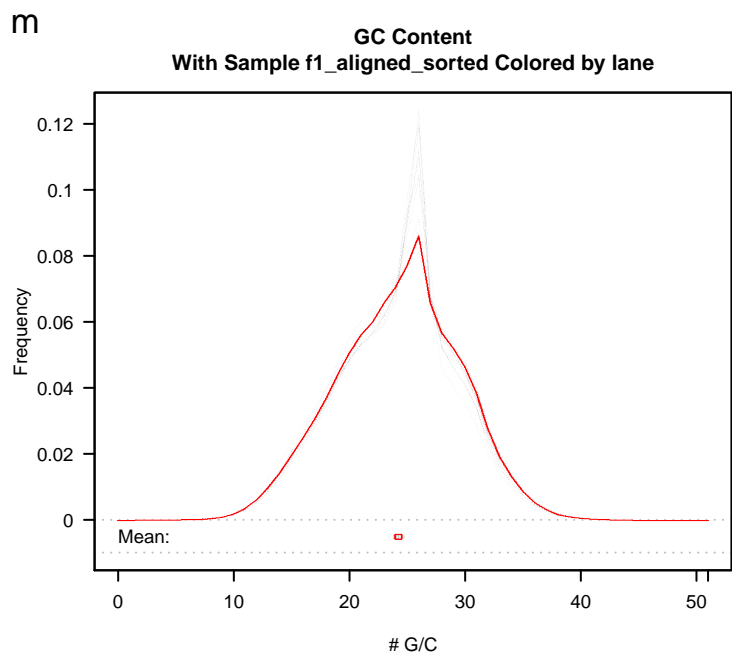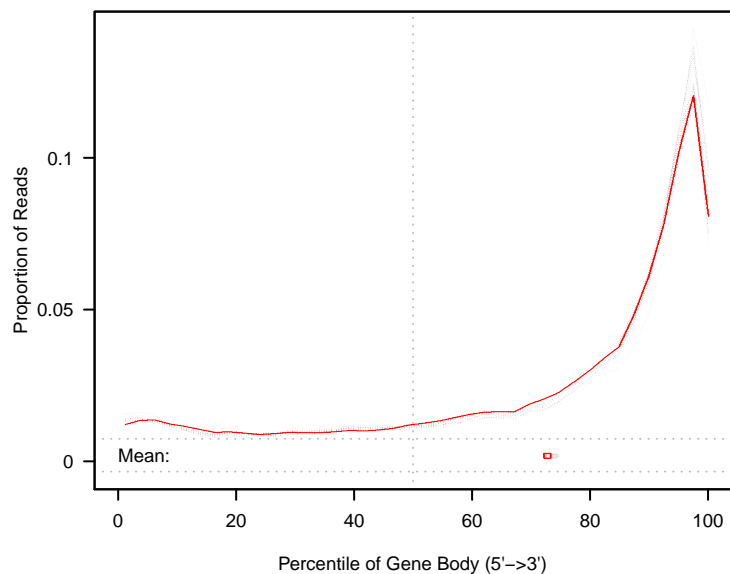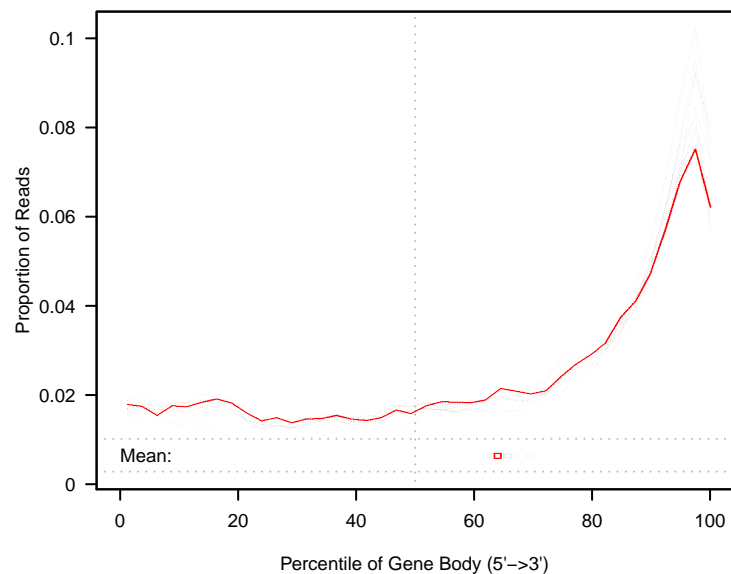With Sample f1_aligned_sorted Colored by lane

s — Gene−Body Coverage, Upper Middle Quartile Genes
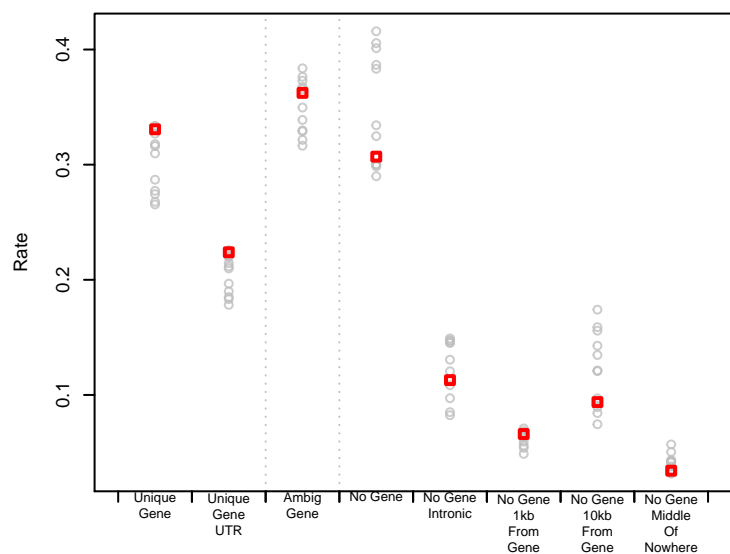With Sample f1_aligned_sorted Colored by lane

t — Gene−Body Coverage, Low Expression Genes
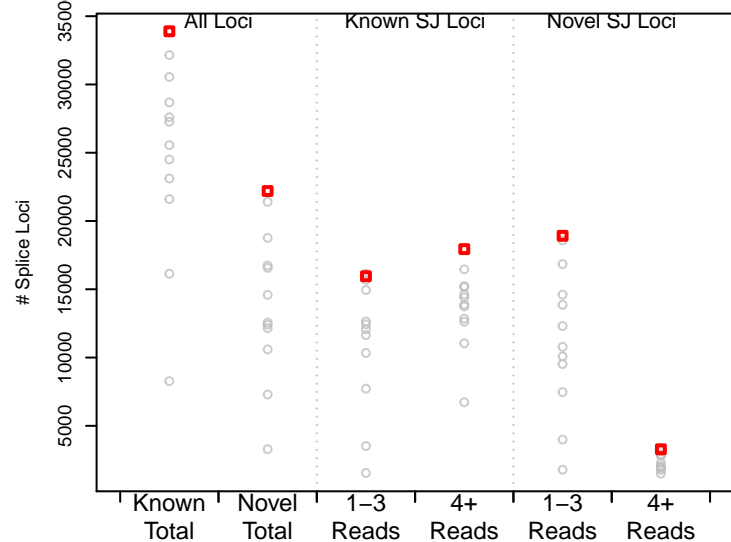With Sample f1_aligned_sorted Colored by lane

u — Read Mapping Location Rates
With Sample f1_aligned_sorted Colored by lane
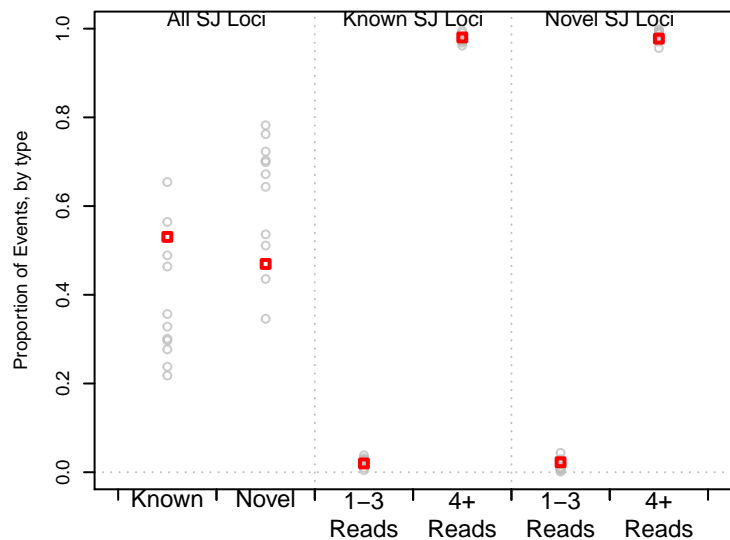
v — # Observed Splice Junction Loci, by type
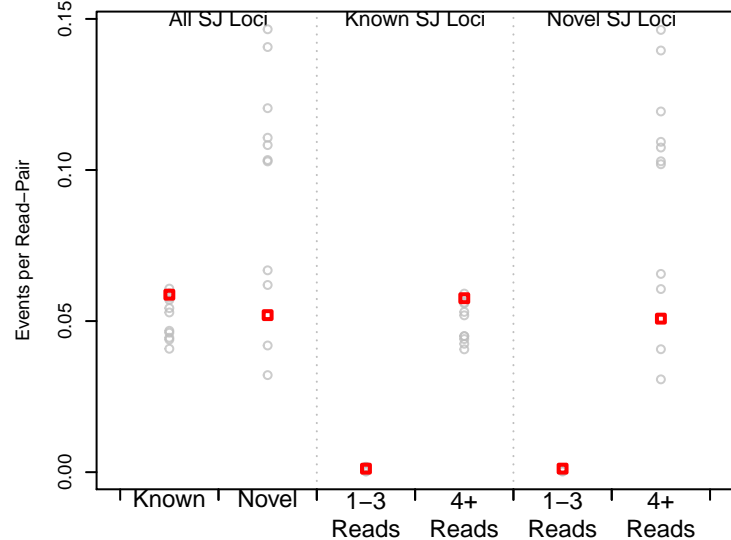With Sample f1_aligned_sorted Colored by lane

w — Breakdown of Splice Junction Events, by type
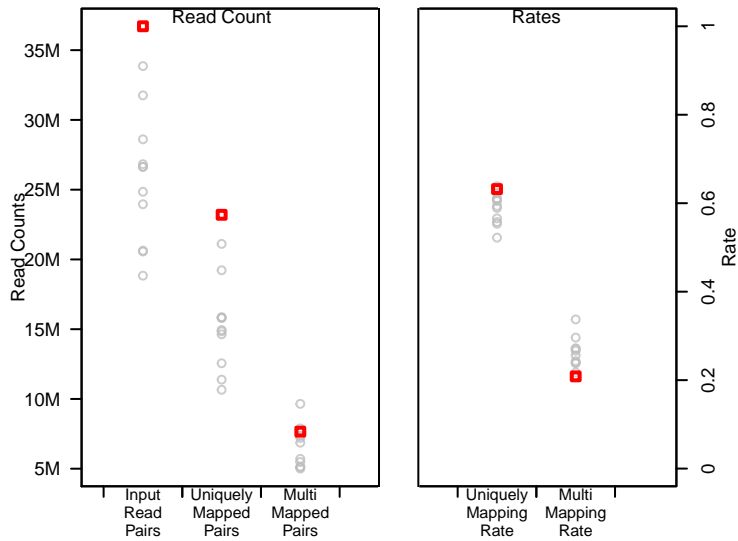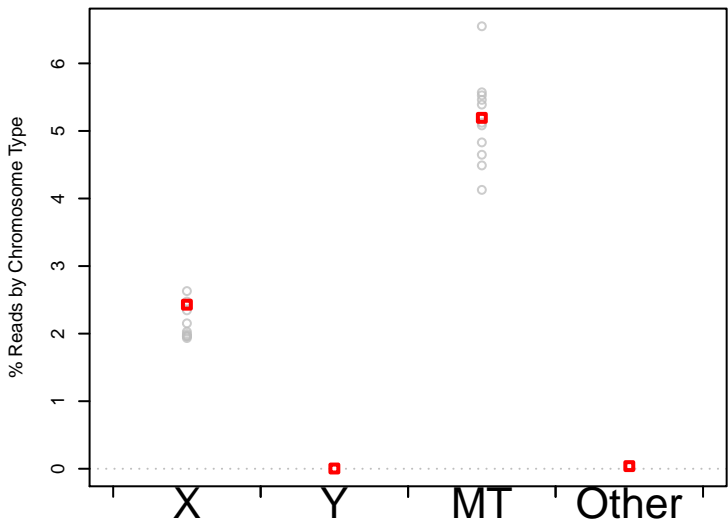With Sample f1_aligned_sorted Colored by lane

x — Splice Junction Event Rates per Read−Pair
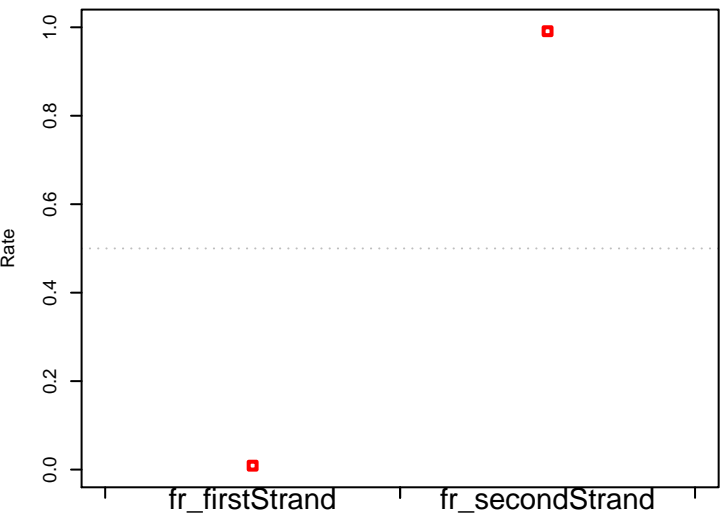With Sample f1_aligned_sorted Colored by lane

y

Mapping Stats
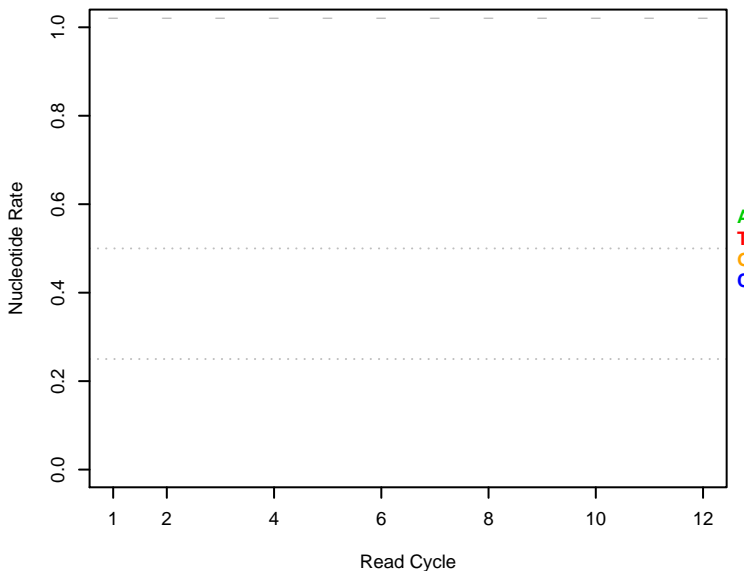**With Sample f1_aligned_sorted Colored by lane**

z

Chromosome Distribution (Excluding Autosomes)
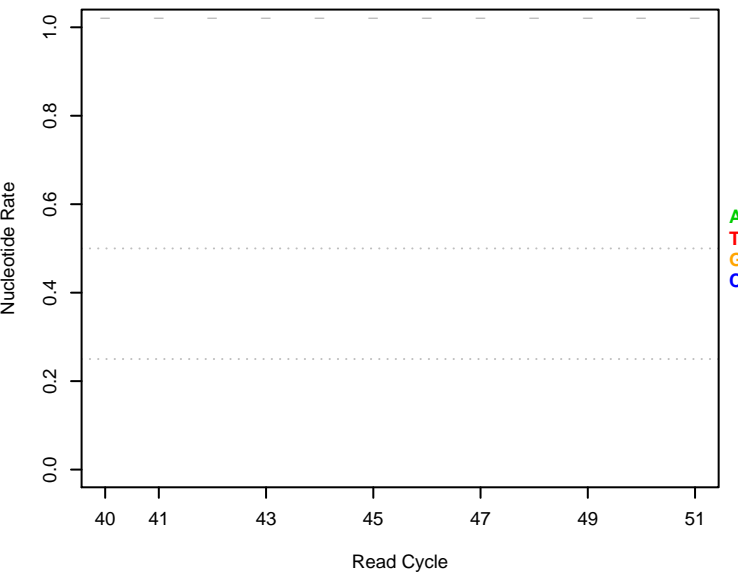**With Sample f1_aligned_sorted Colored by lane**

aa

Strandedness Test
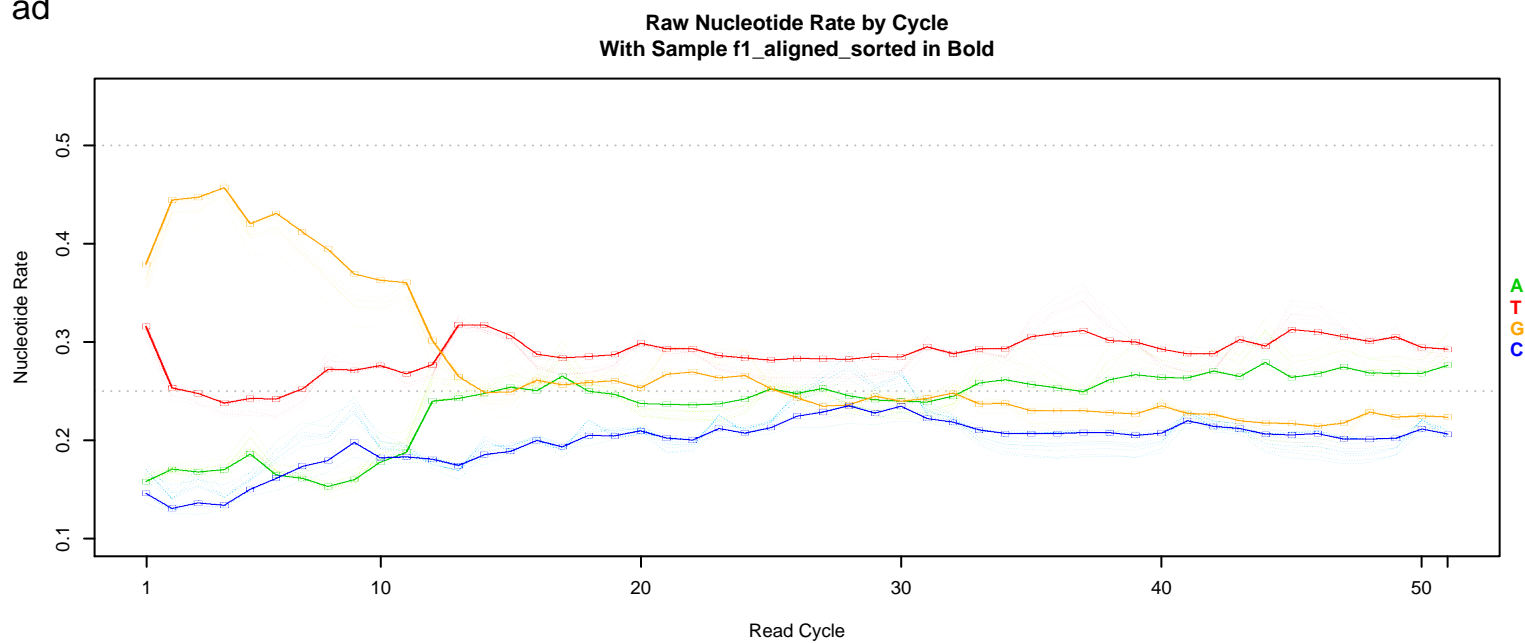**With Sample f1_aligned_sorted Colored by lane**

ab

Nucleotide Rate by Cycle, Leading Clipped bases (12)
With Sample f1_aligned_sorted in Bold

ac

Nucleotide Rate by Cycle, Trailing Clipped bases (12)
With Sample f1_aligned_sorted in Bold

ad

Raw Nucleotide Rate by Cycle
With Sample f1_aligned_sorted in Bold

ae

Nucleotide Rate by Cycle, Aligned bases only
With Sample f1_aligned_sorted in Bold