

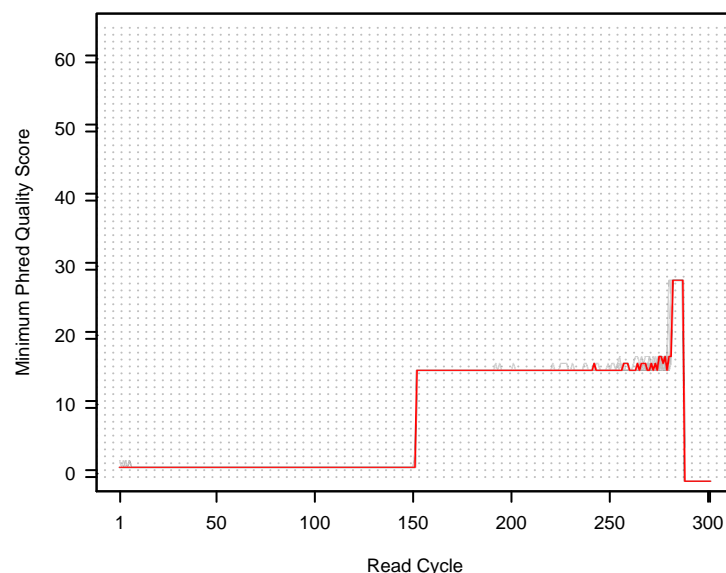
a

Sample Highlight:
ps11_S2
Colored by Lane



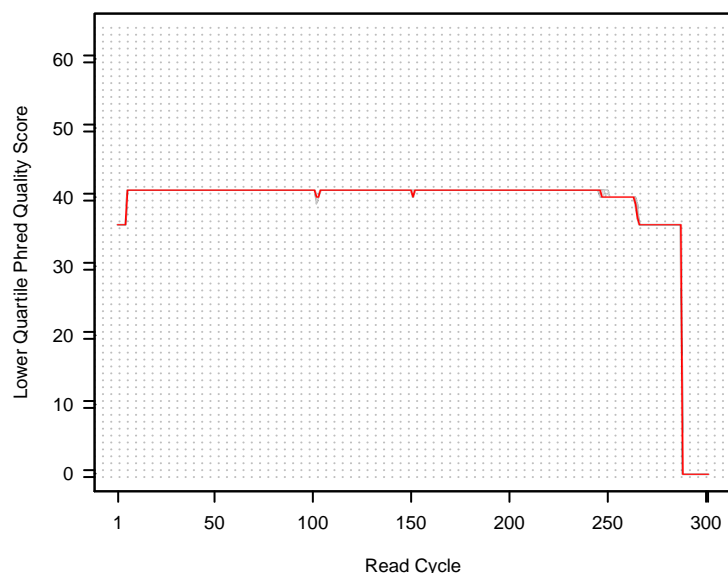
b

Minimum Phred Quality Score
With Sample ps11_S2 Colored by lane



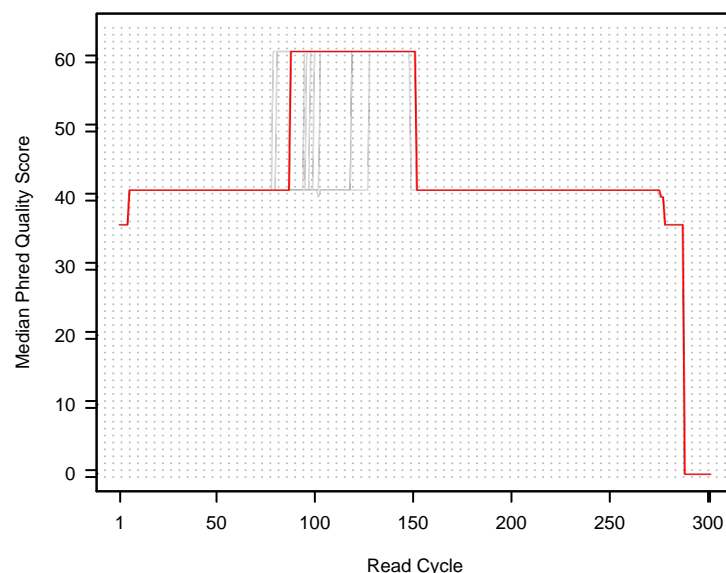
c

Lower Quartile Phred Quality Score
With Sample ps11_S2 Colored by lane



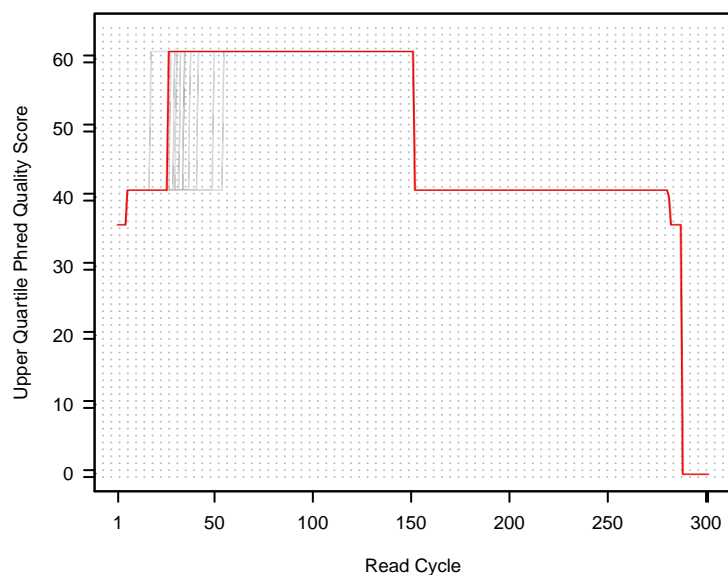
d

Median Phred Quality Score
With Sample ps11_S2 Colored by lane



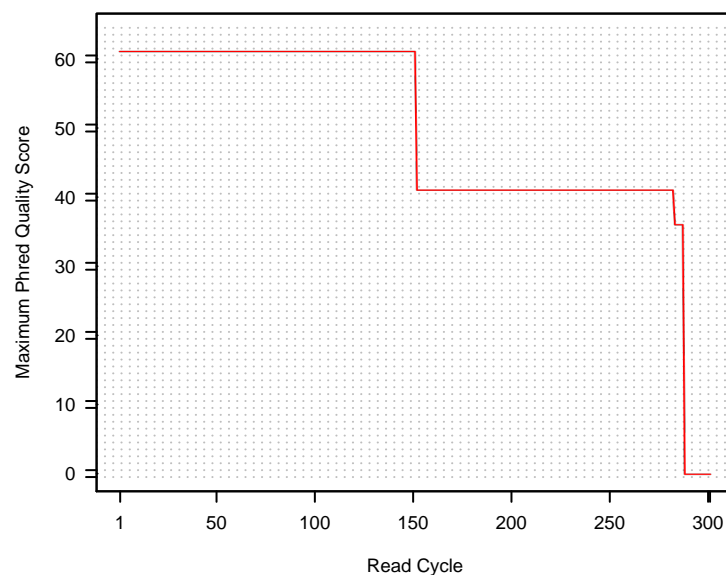
e

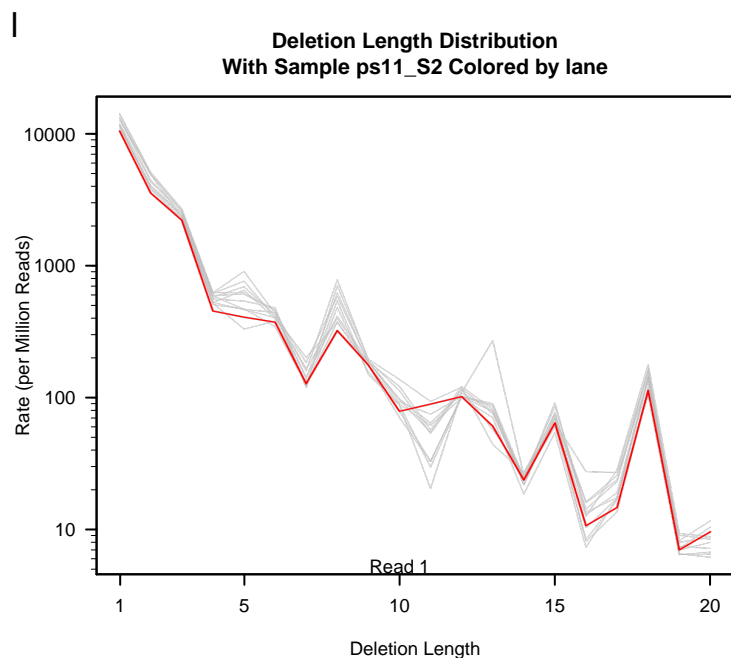
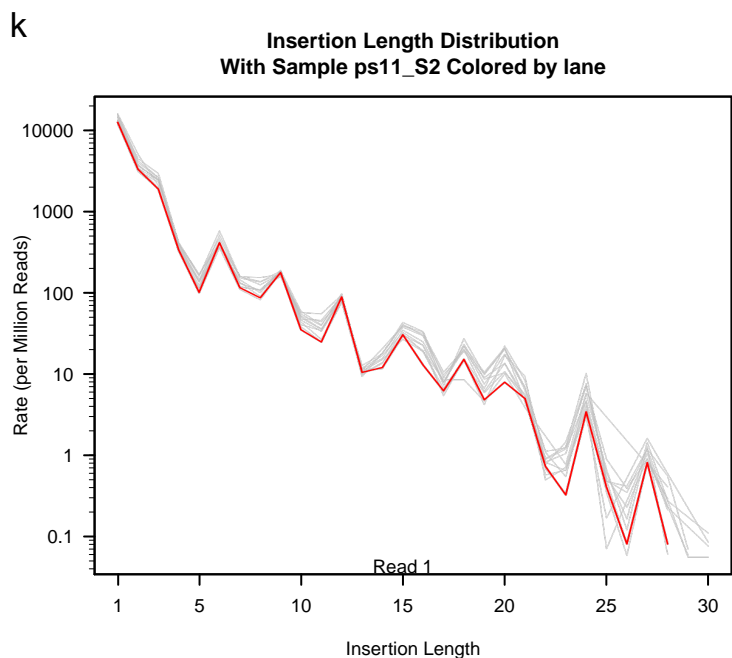
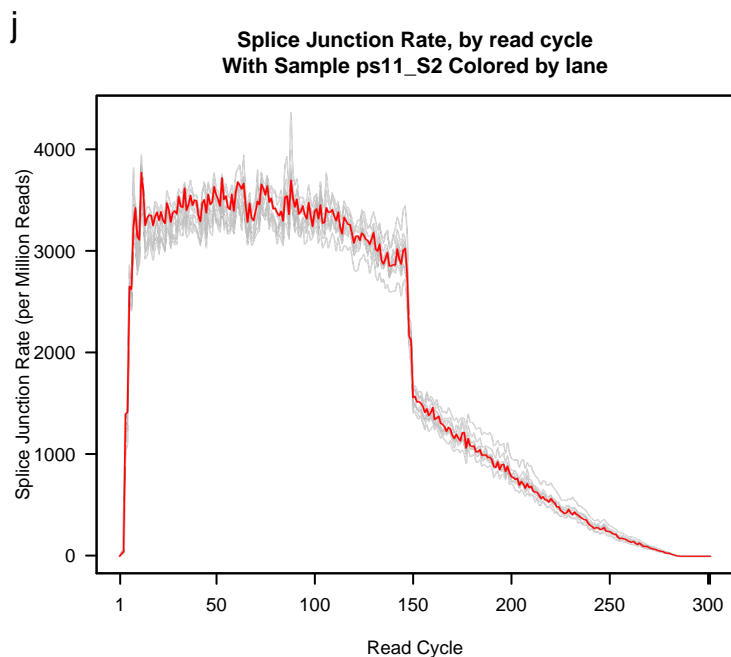
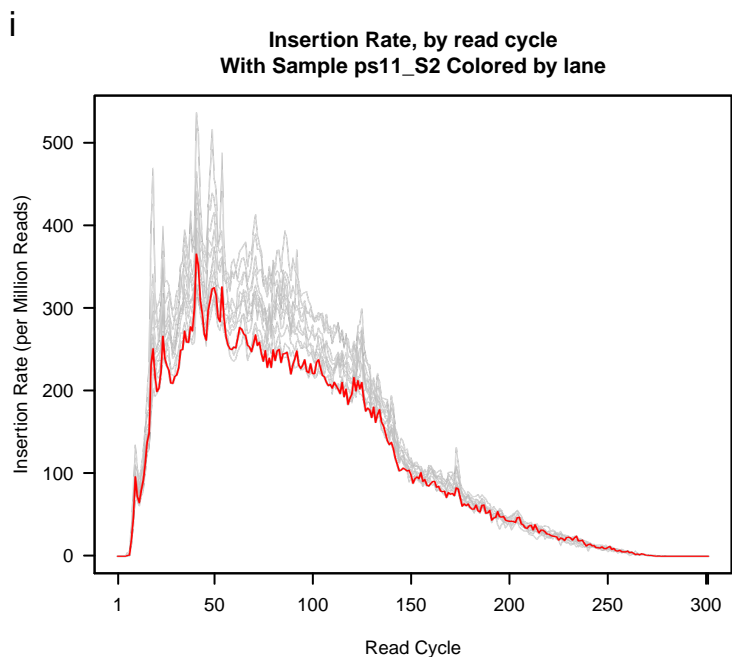
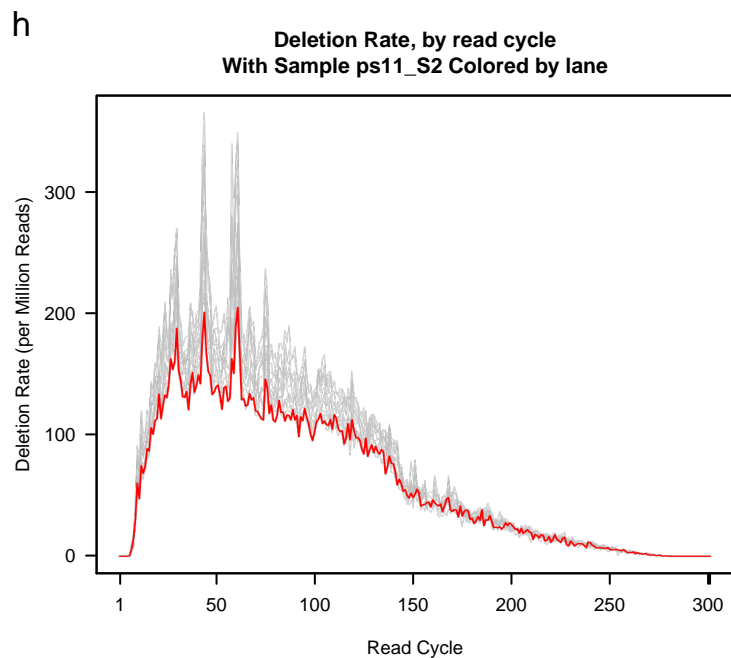
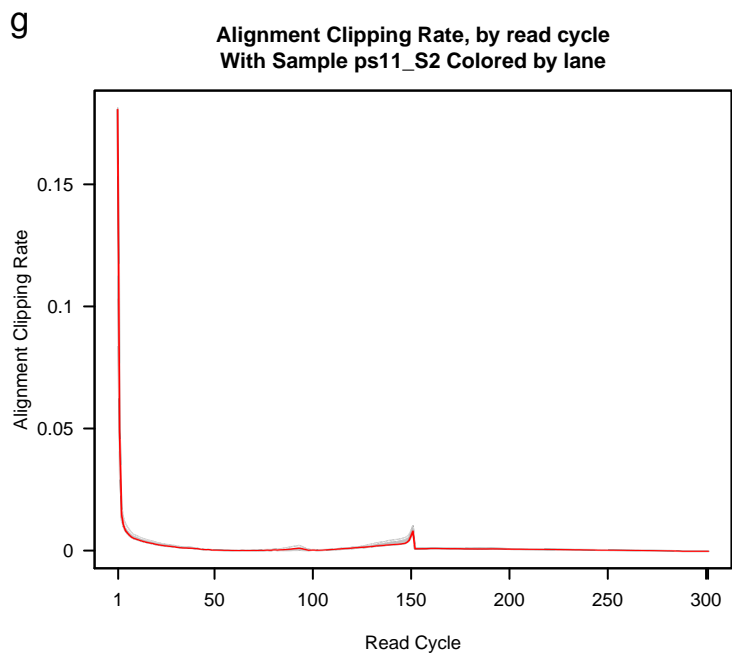
Upper Quartile Phred Quality Score
With Sample ps11_S2 Colored by lane

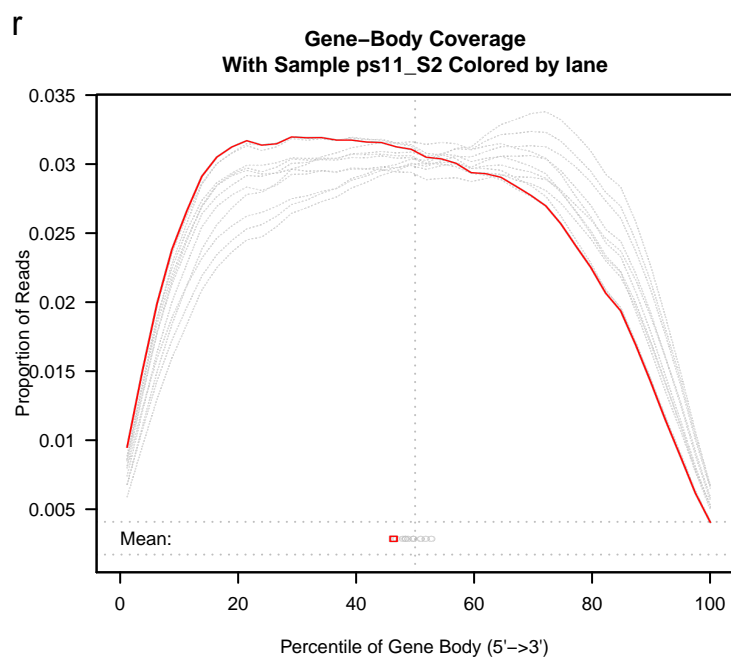
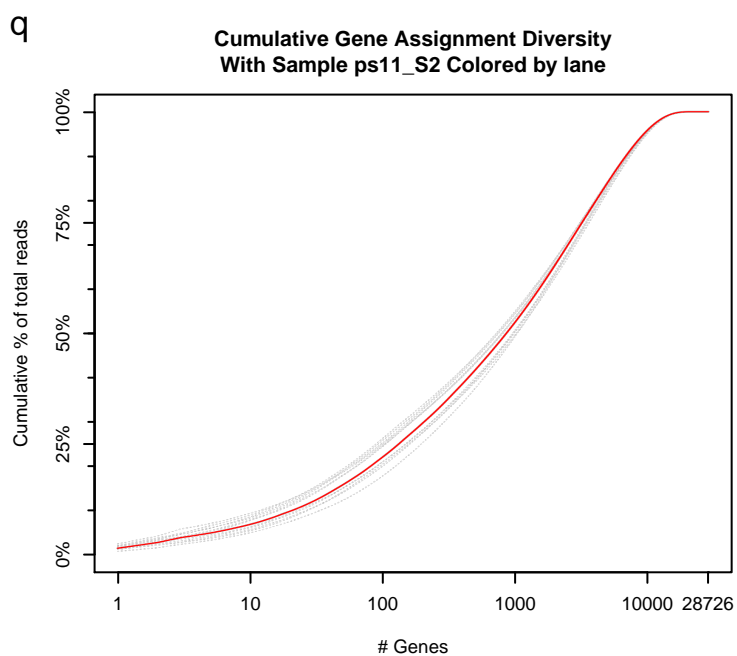
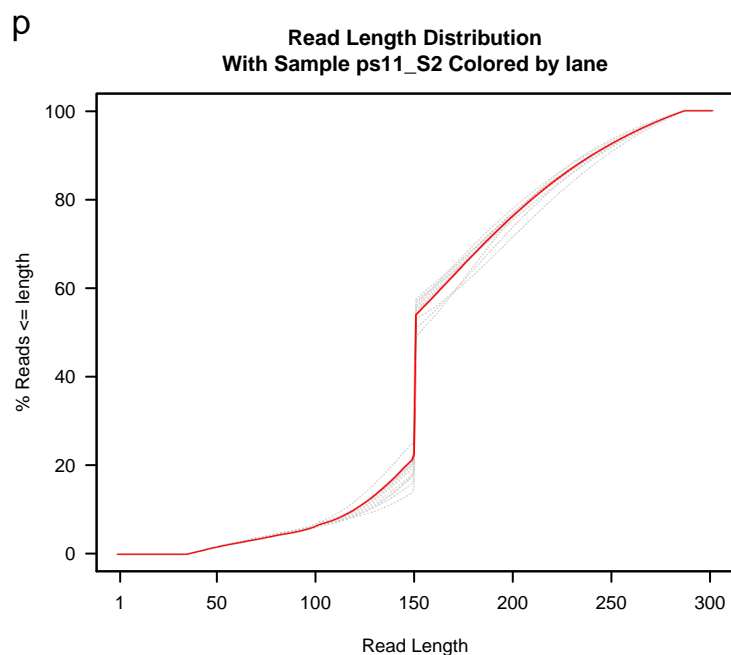
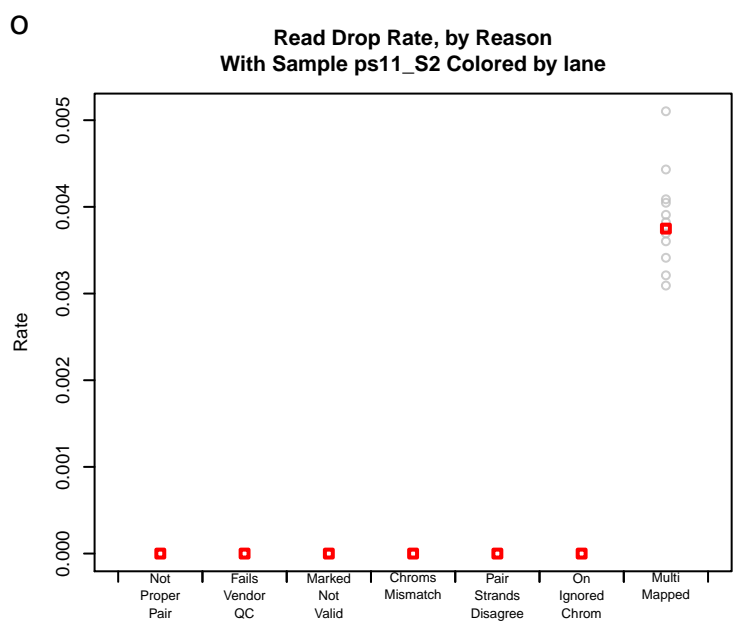
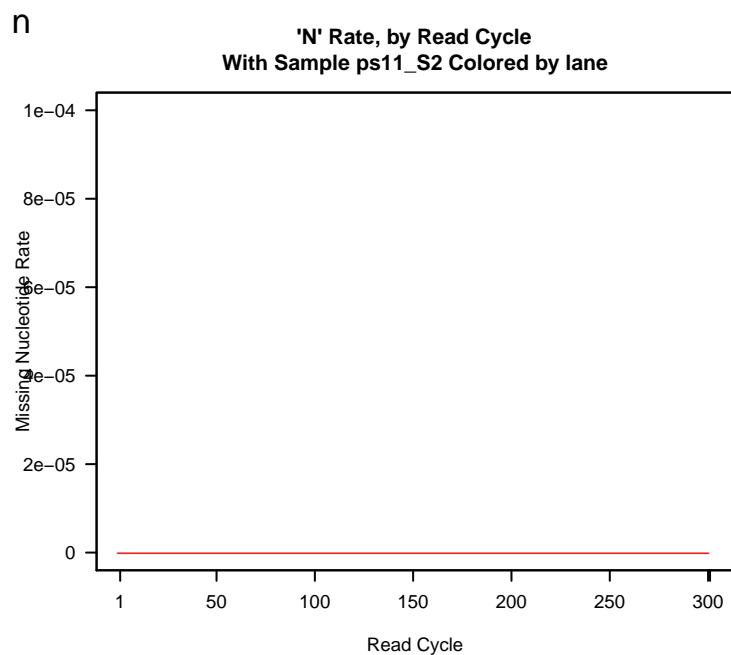
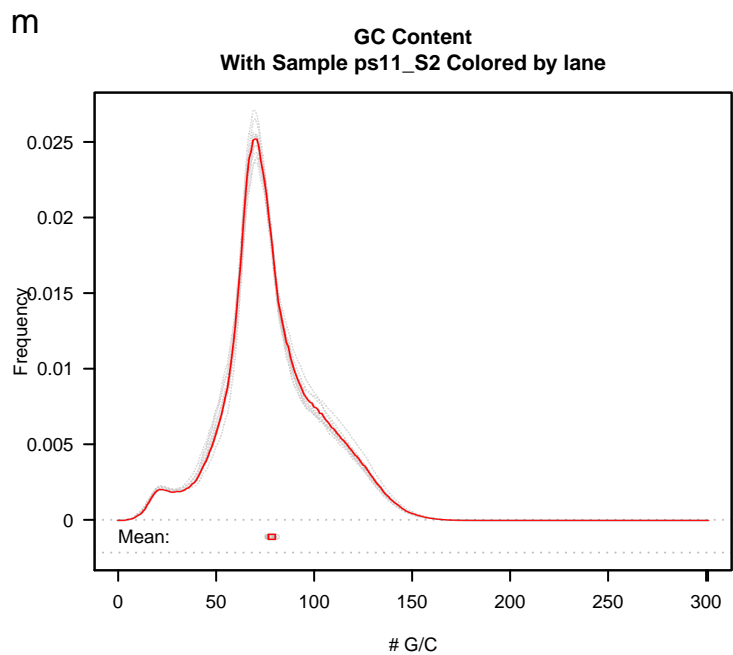


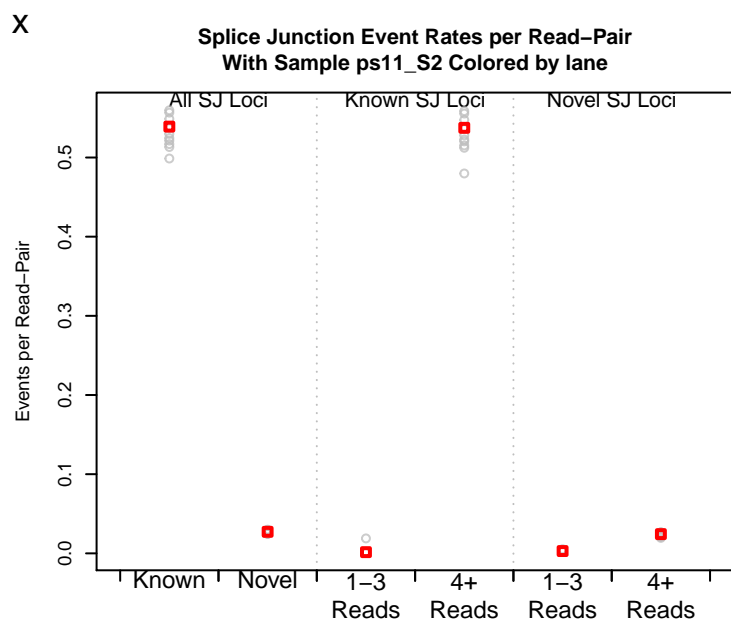
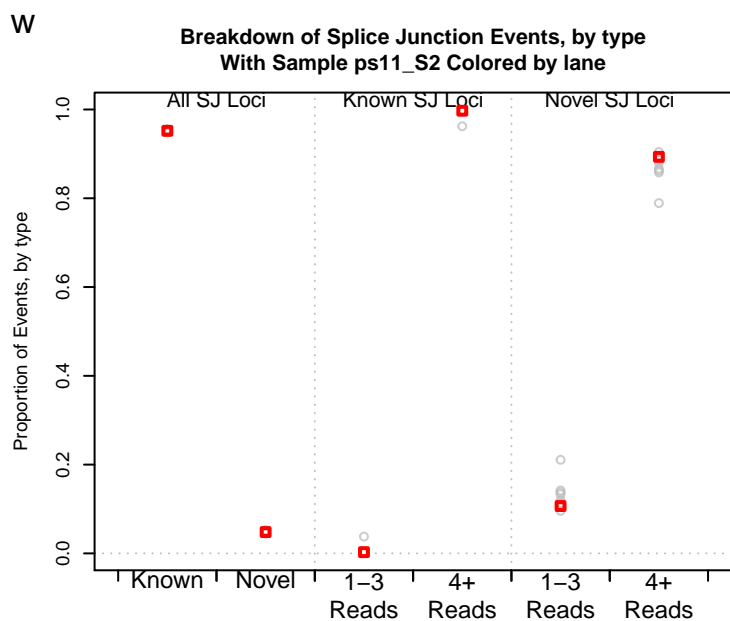
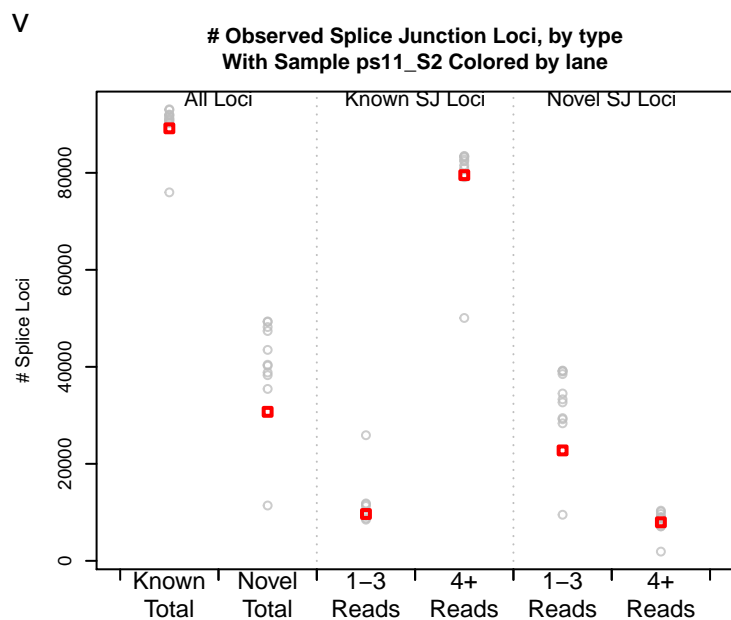
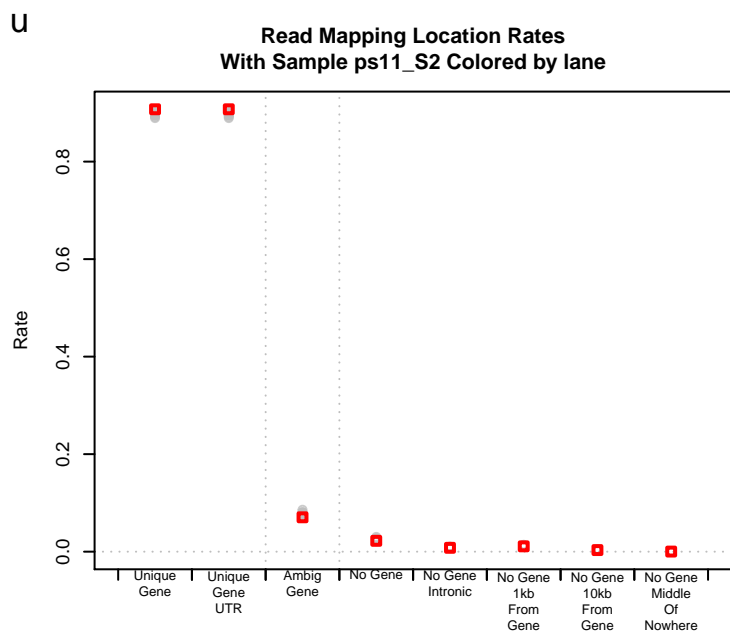
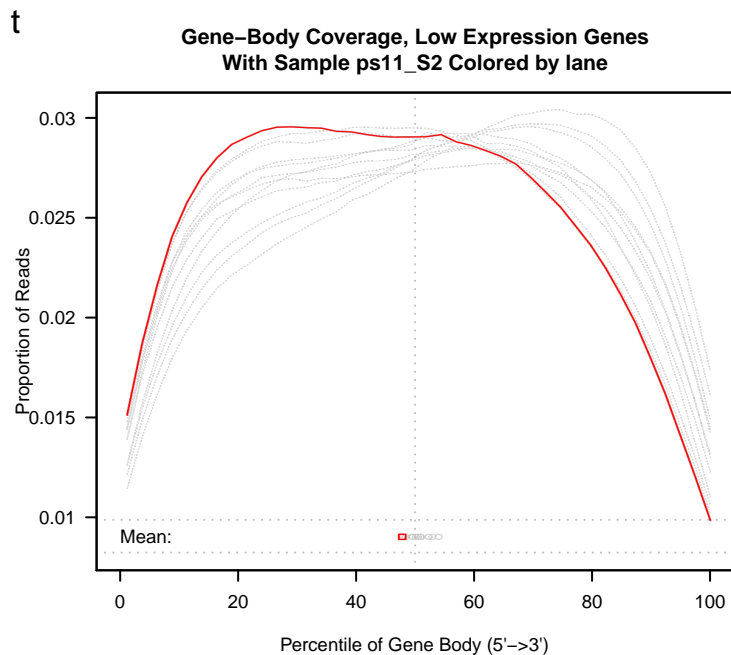
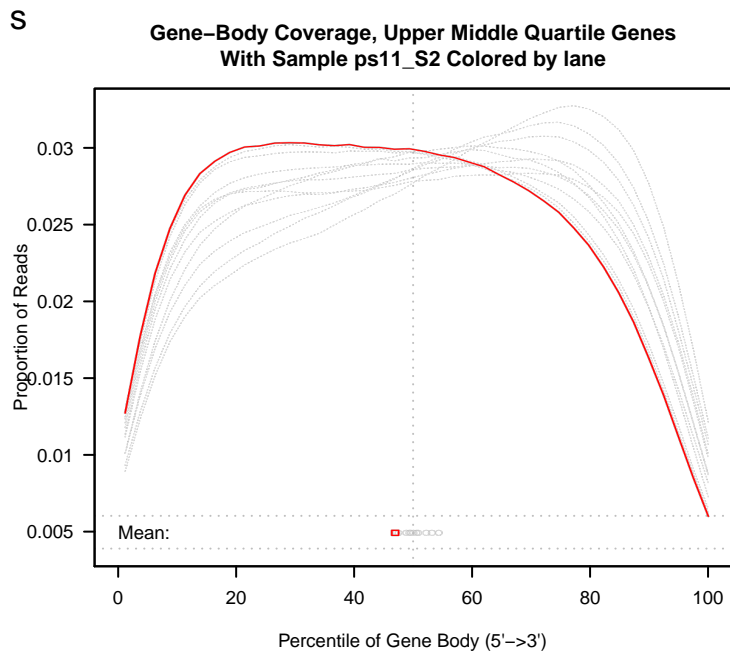
f

Maximum Phred Quality Score
With Sample ps11_S2 Colored by lane



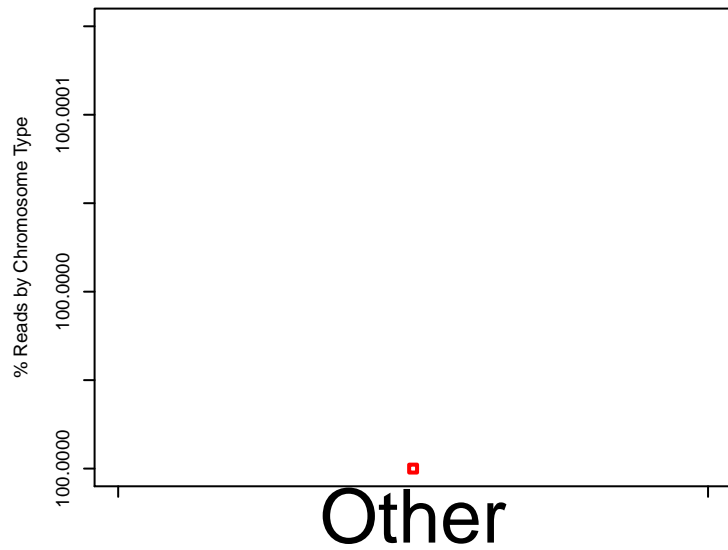






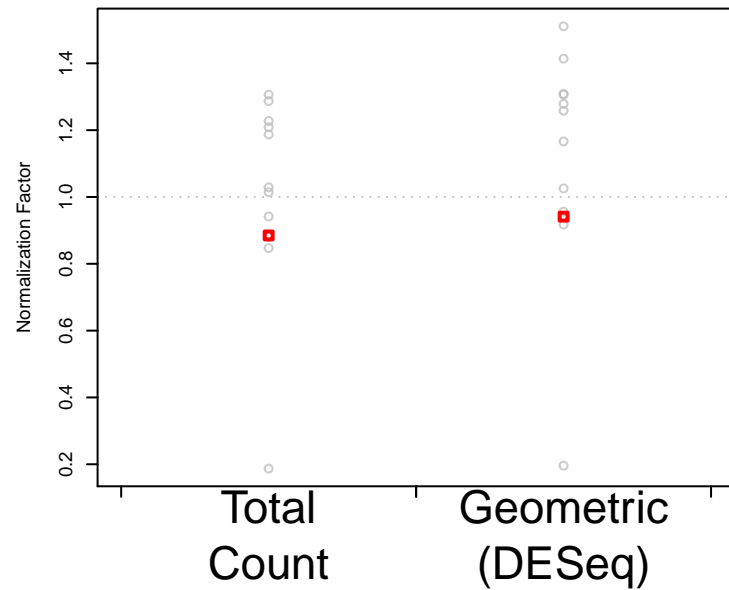
y

Chromosome Distribution (Excluding Autosomes)
With Sample ps11_S2 Colored by lane



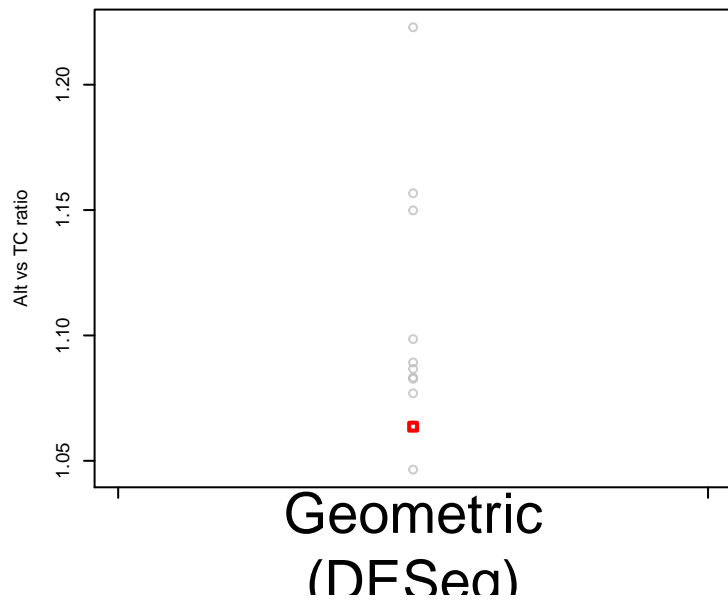
Z

Normalization Factors
With Sample ps11_S2 Colored by lane



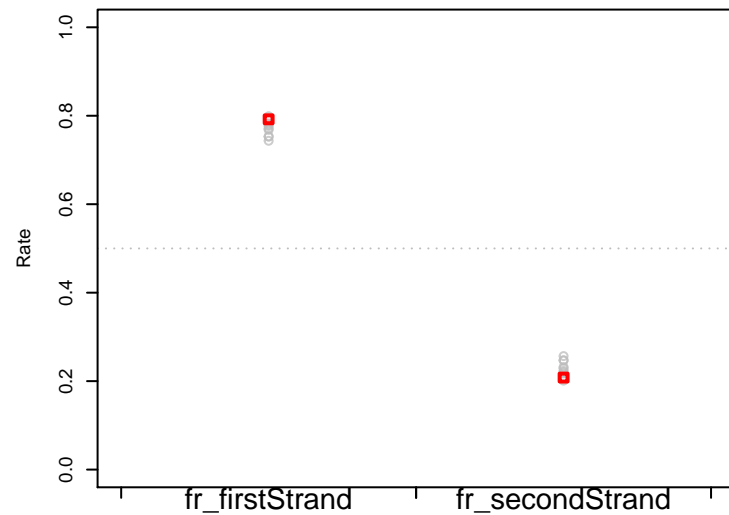
aa

Normalization Factors vs Total-Count Normalization
With Sample ps11_S2 Colored by lane



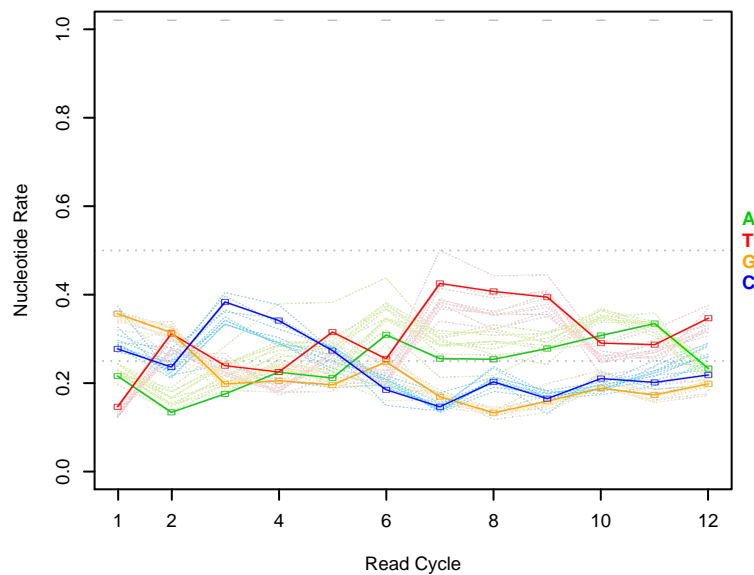
ab

Strandedness Test
With Sample ps11_S2 Colored by lane



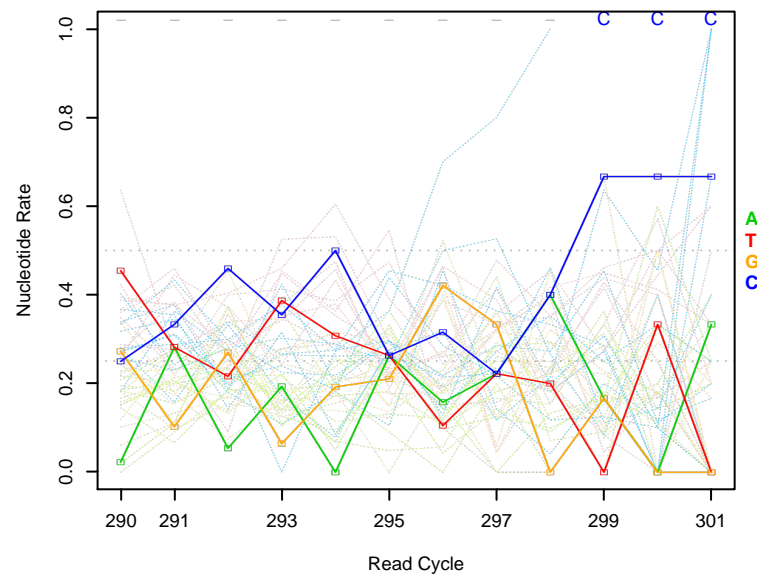
ac

Nucleotide Rate by Cycle, Leading Clipped bases (12)
With Sample ps11_S2 in Bold

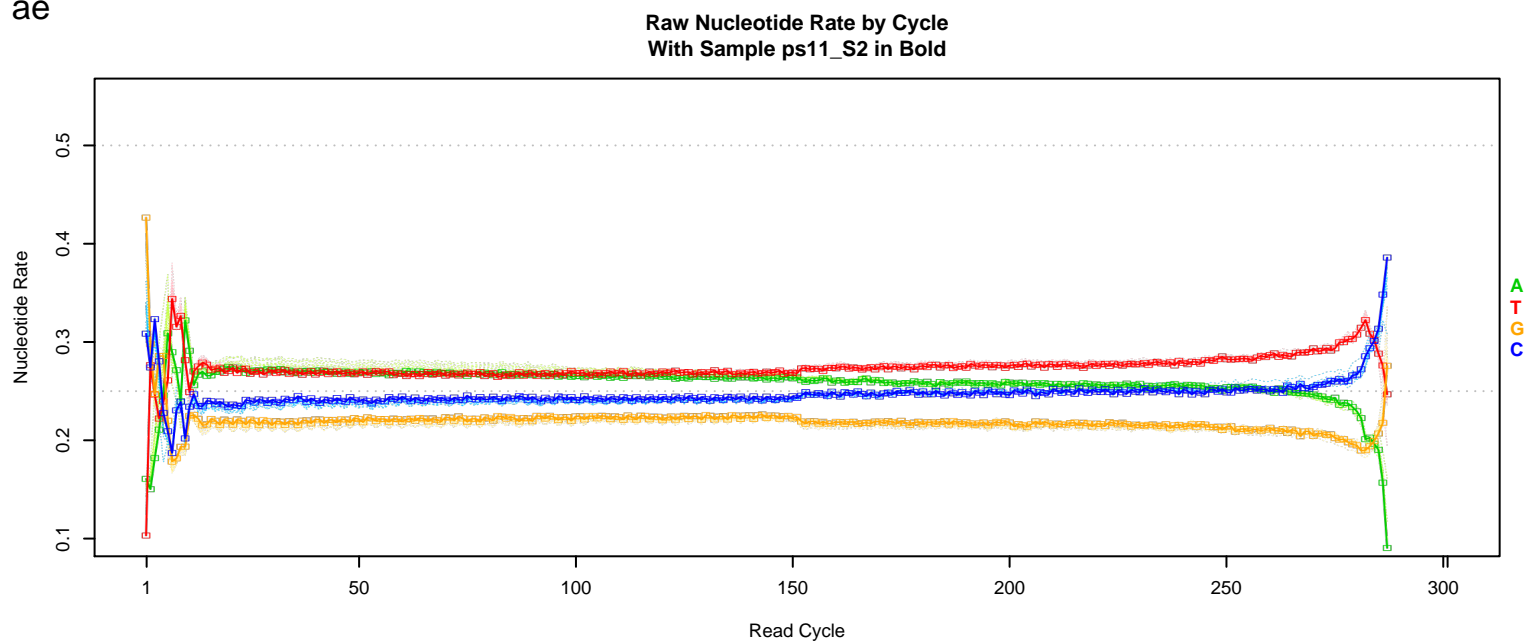


ad

Nucleotide Rate by Cycle, Trailing Clipped bases (12)
With Sample ps11_S2 in Bold



ae



af

