

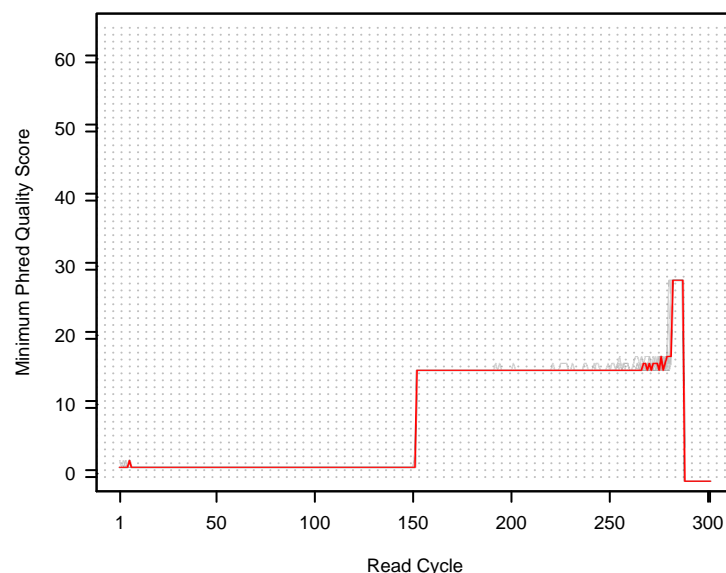
a

Sample Highlight: ps38_S5 Colored by Lane



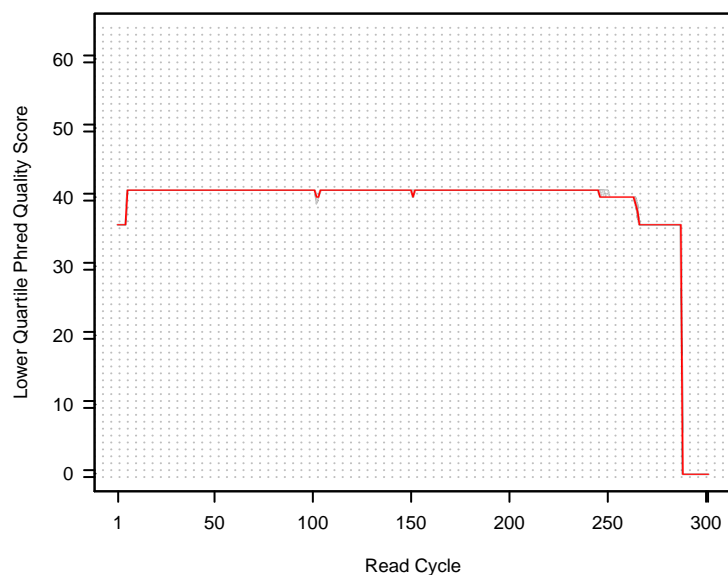
b

Minimum Phred Quality Score
With Sample ps38_S5 Colored by lane



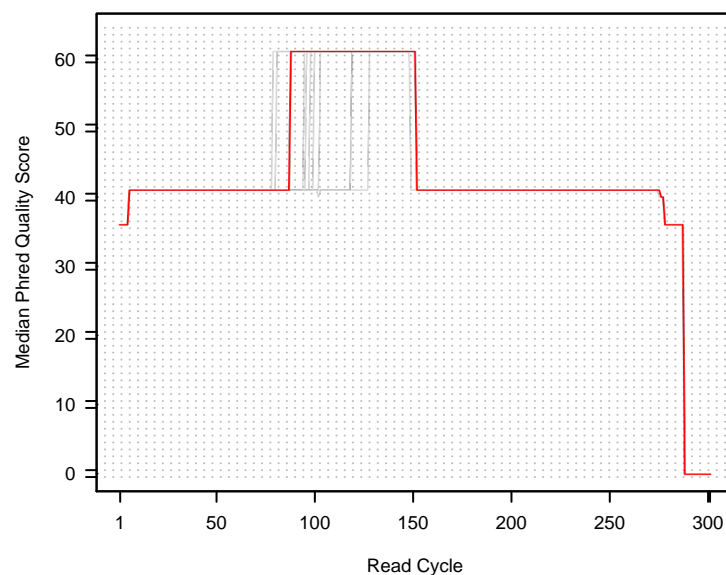
c

Lower Quartile Phred Quality Score
With Sample ps38_S5 Colored by lane



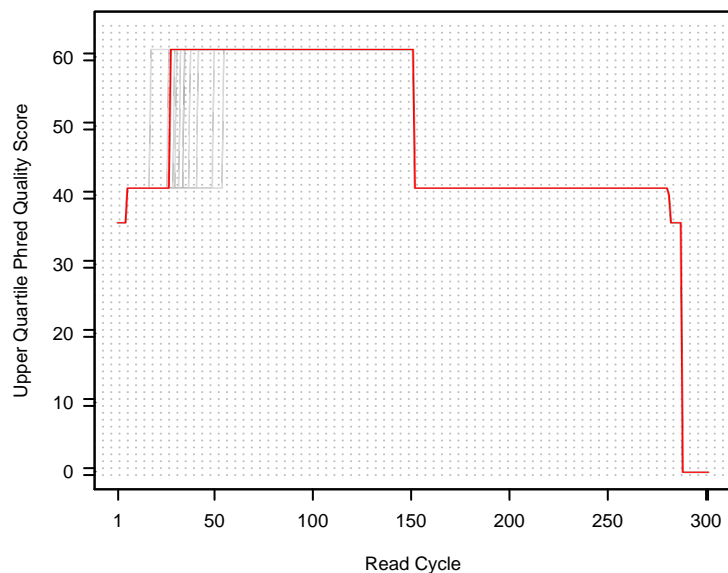
d

Median Phred Quality Score
With Sample ps38_S5 Colored by lane



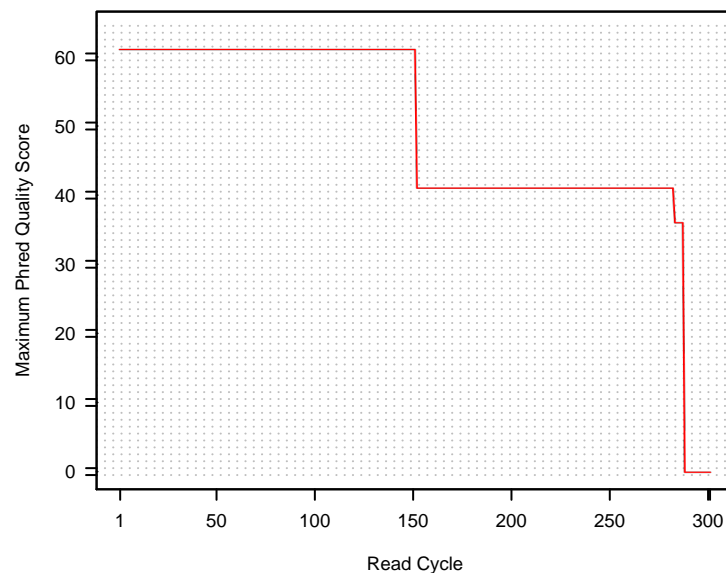
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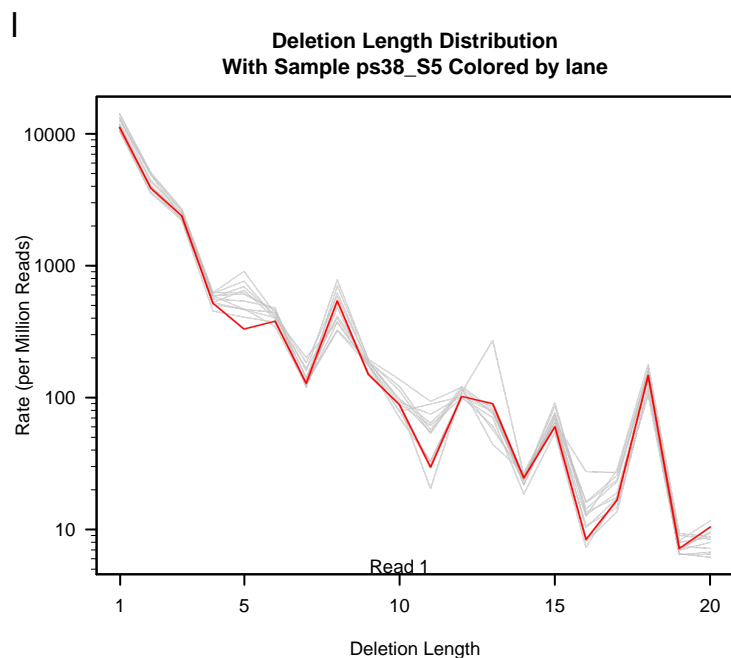
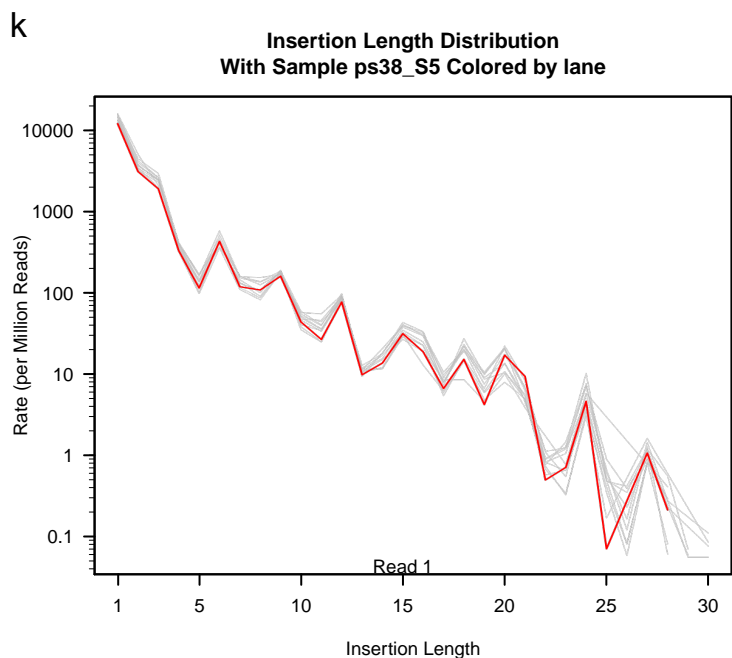
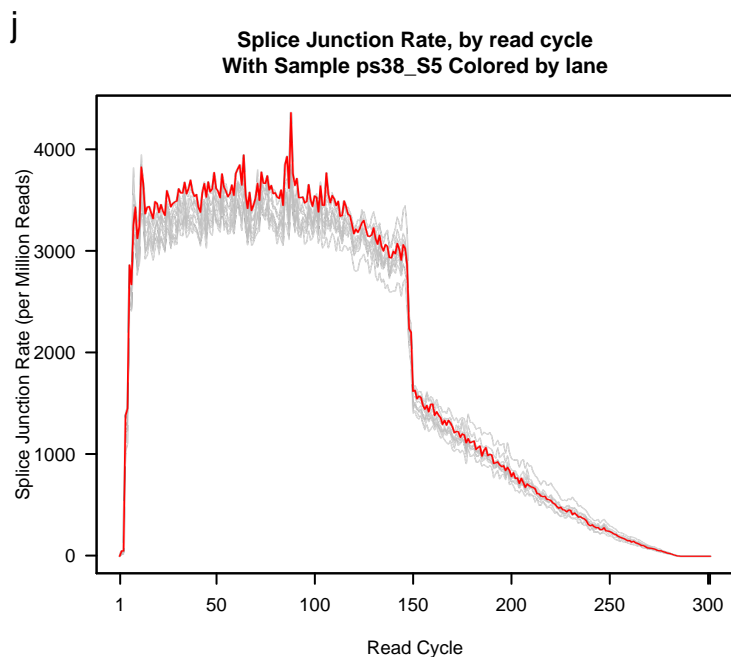
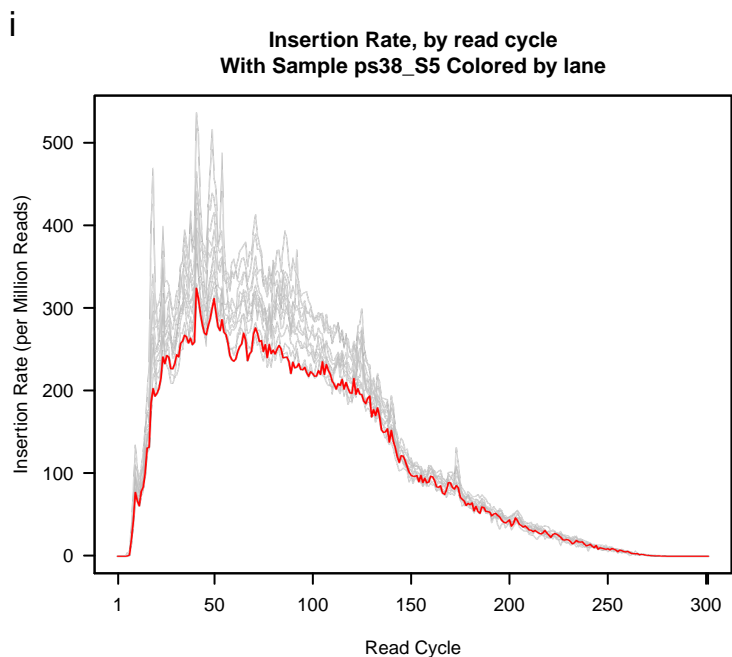
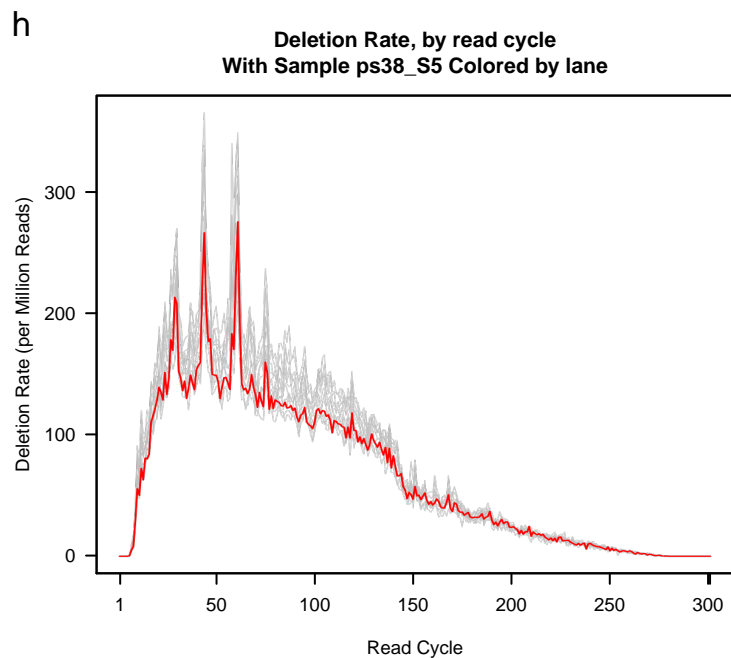
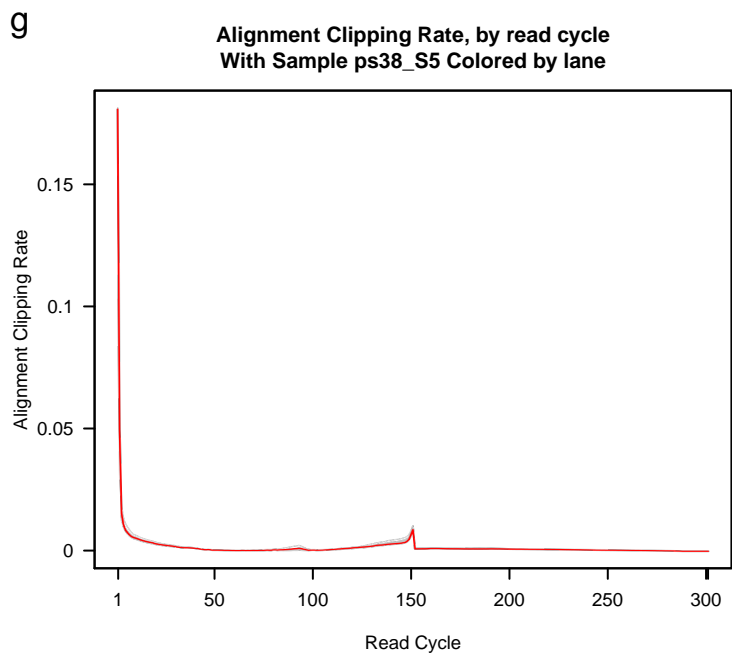
Upper Quartile Phred Quality Score
With Sample ps38_S5 Colored by lane

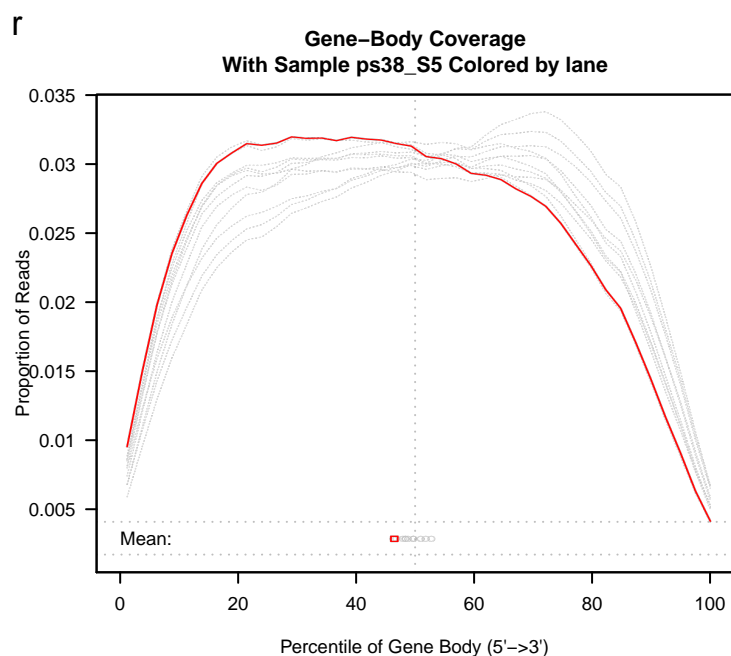
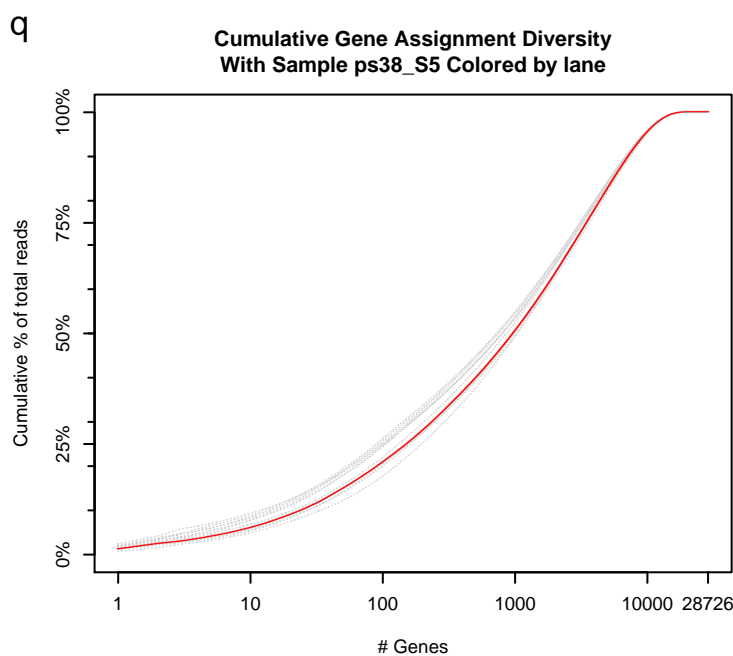
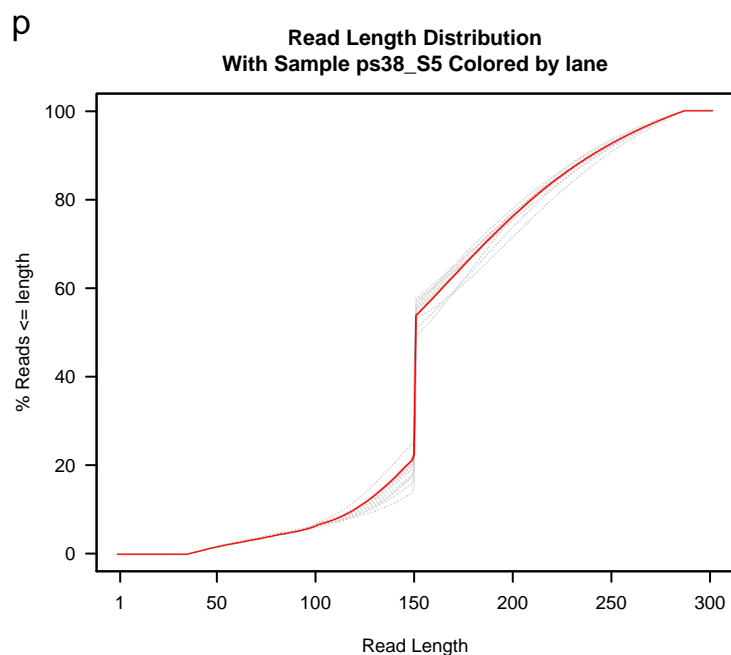
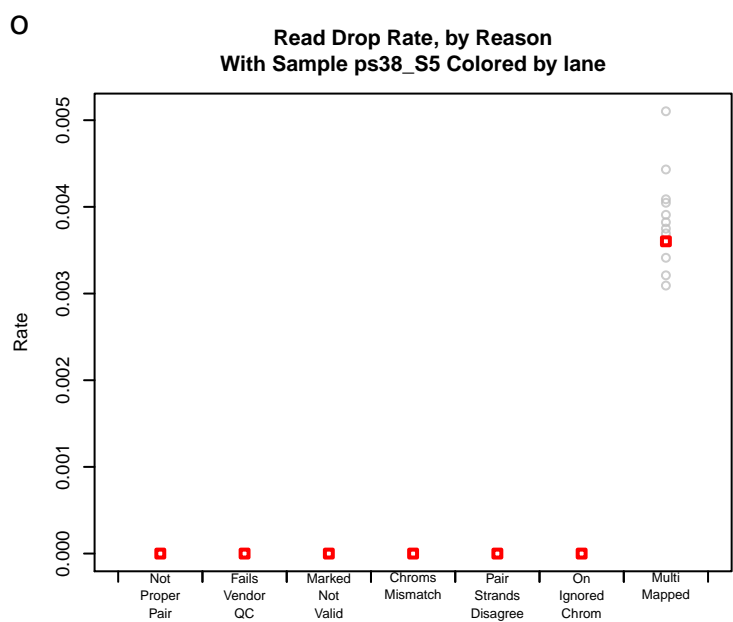
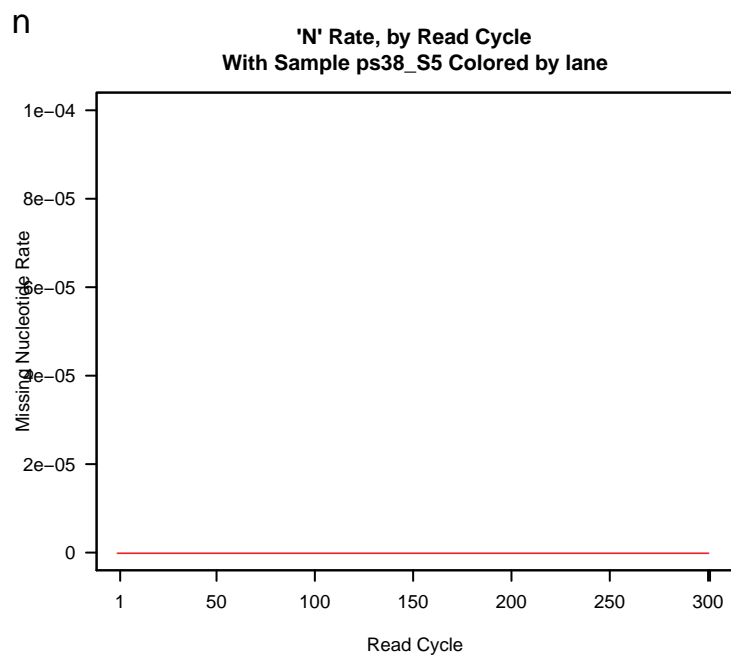
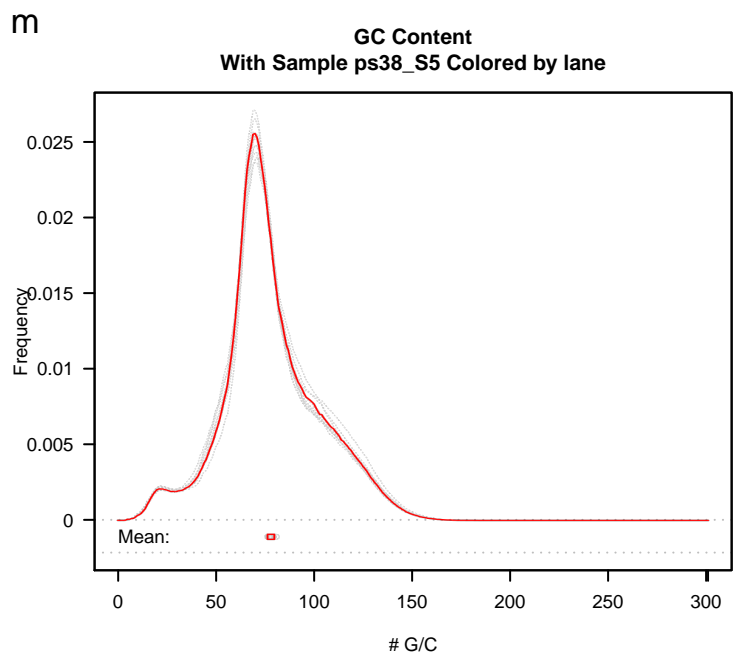


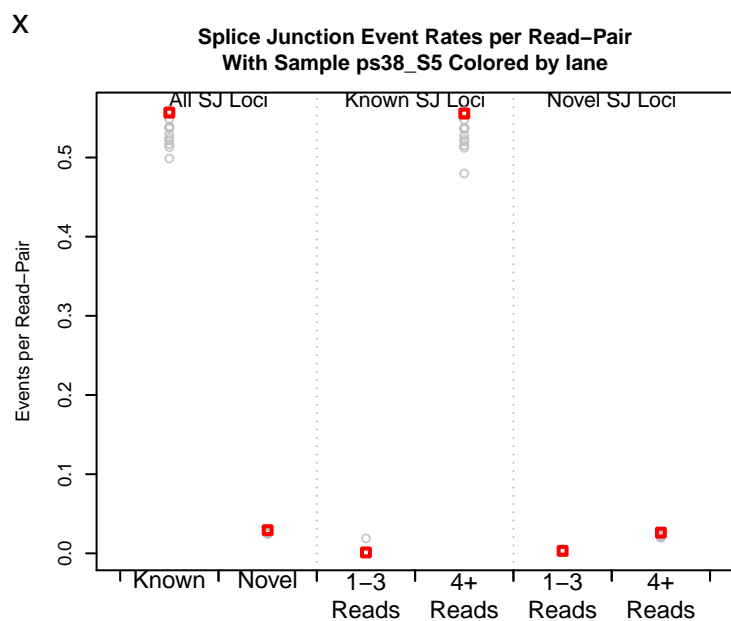
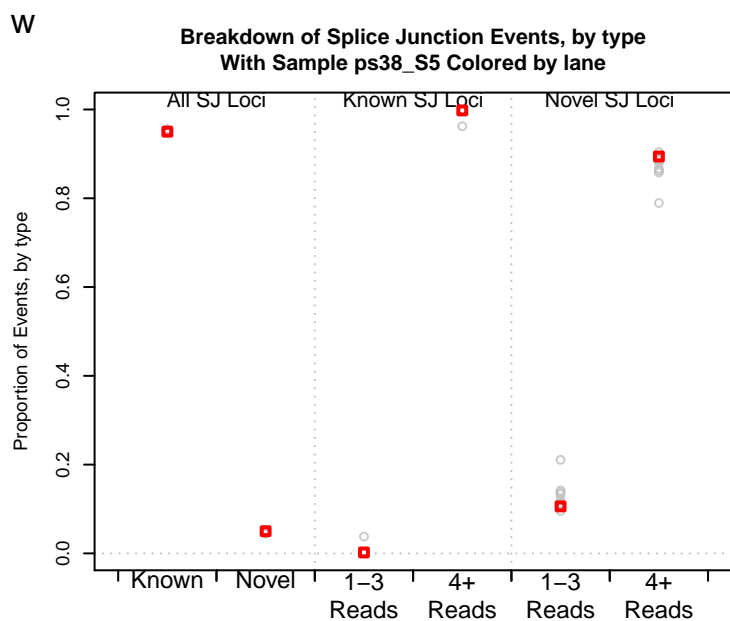
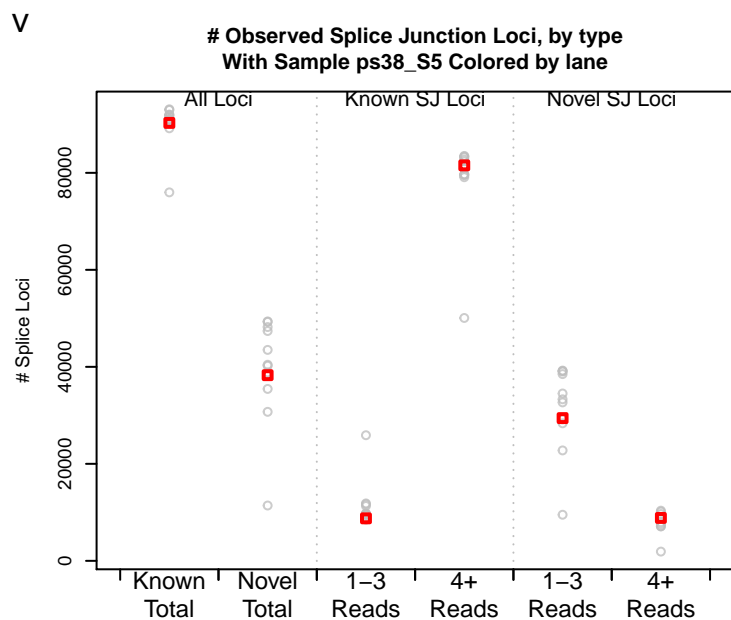
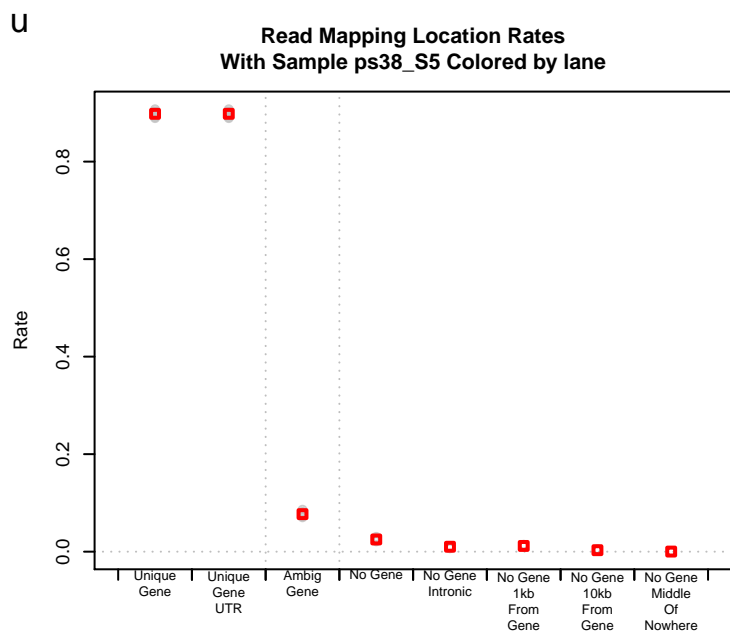
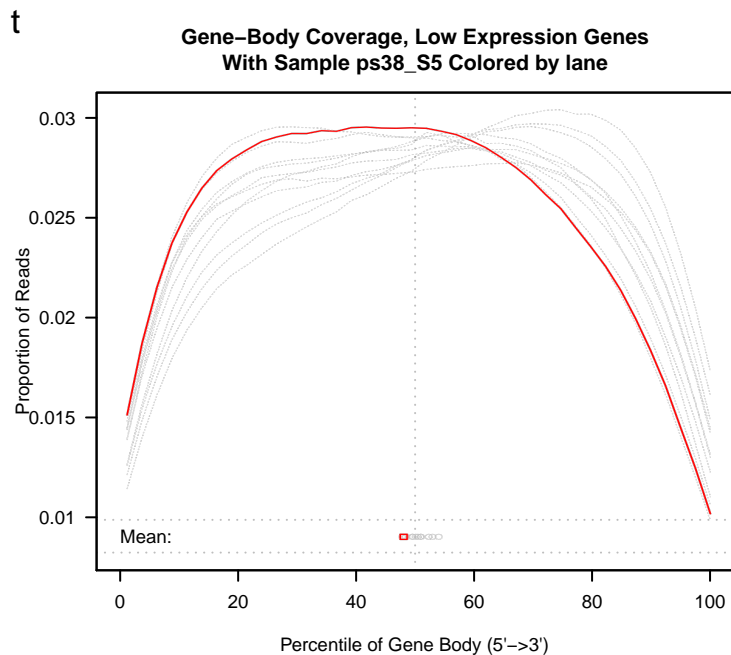
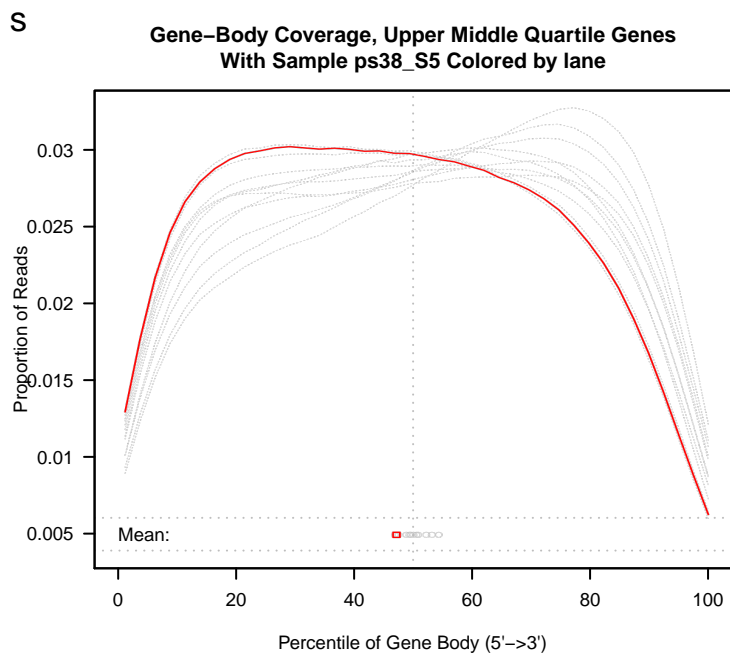
f

Maximum Phred Quality Score
With Sample ps38_S5 Colored by lane



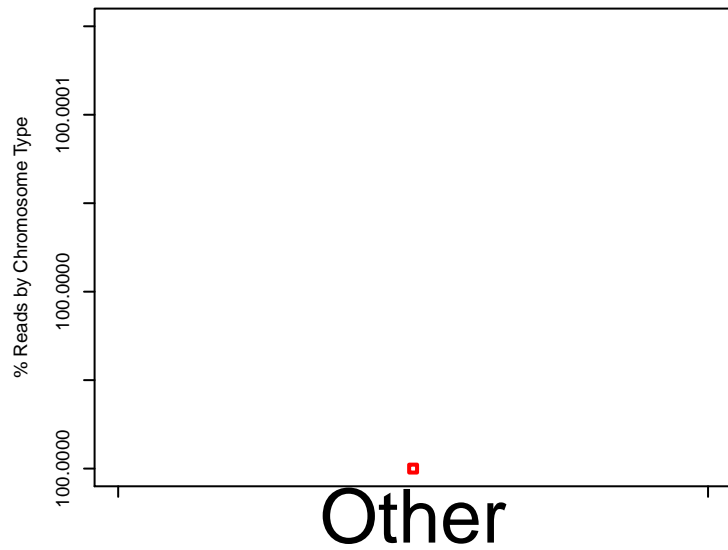






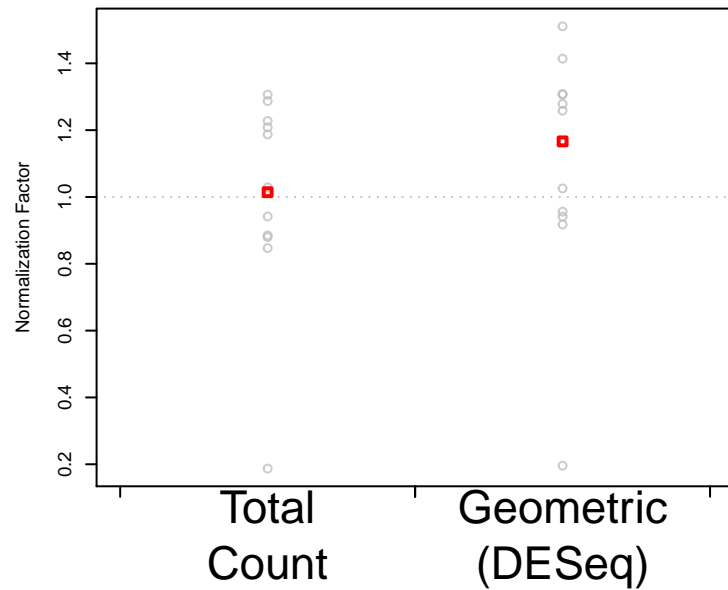
y

Chromosome Distribution (Excluding Autosomes)
With Sample ps38_S5 Colored by lane



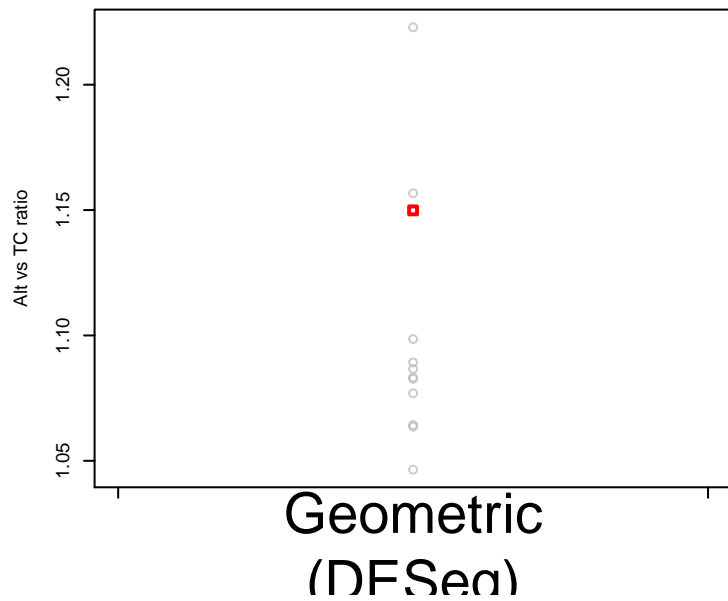
Z

Normalization Factors
With Sample ps38_S5 Colored by lane



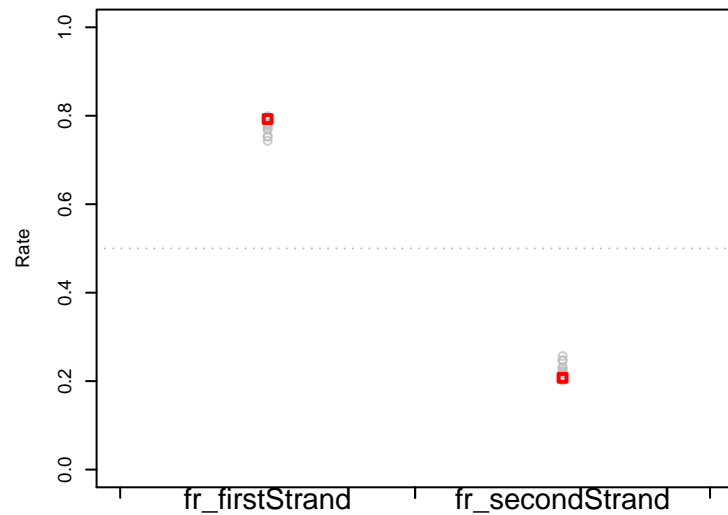
aa

Normalization Factors vs Total-Count Normalization
With Sample ps38_S5 Colored by lane



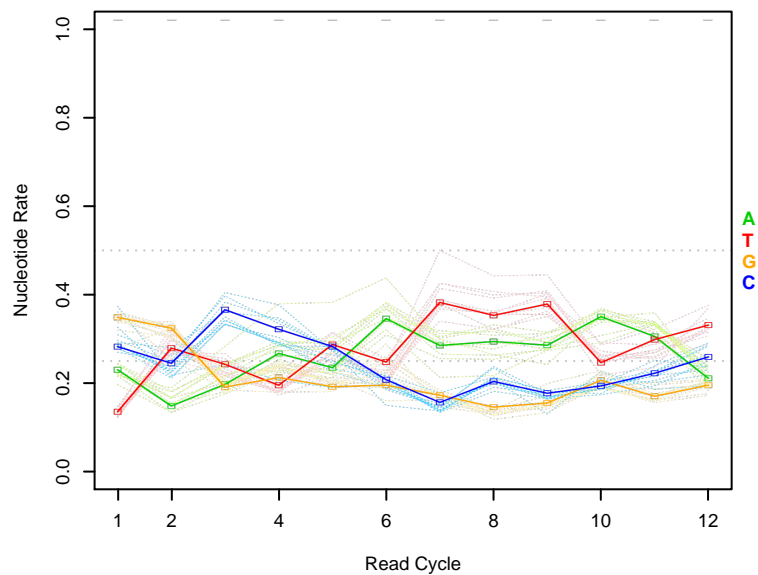
ab

Strandedness Test
With Sample ps38_S5 Colored by lane



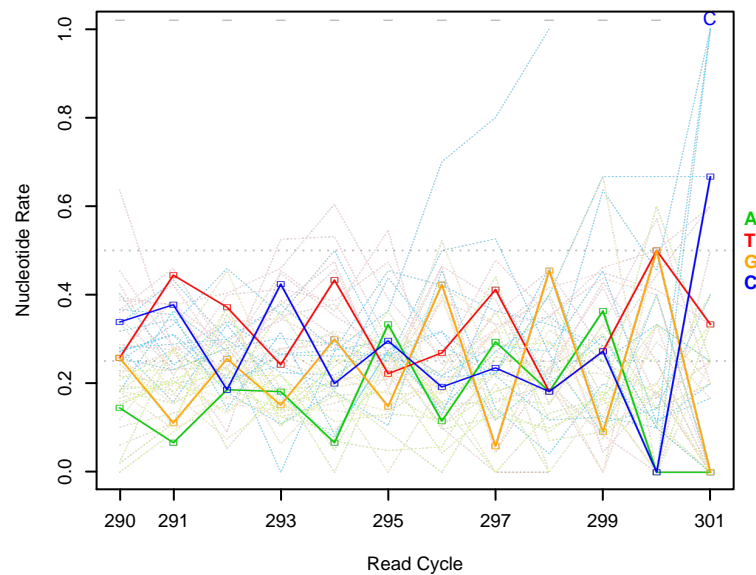
ac

Nucleotide Rate by Cycle, Leading Clipped bases (12)
With Sample ps38_S5 in Bold

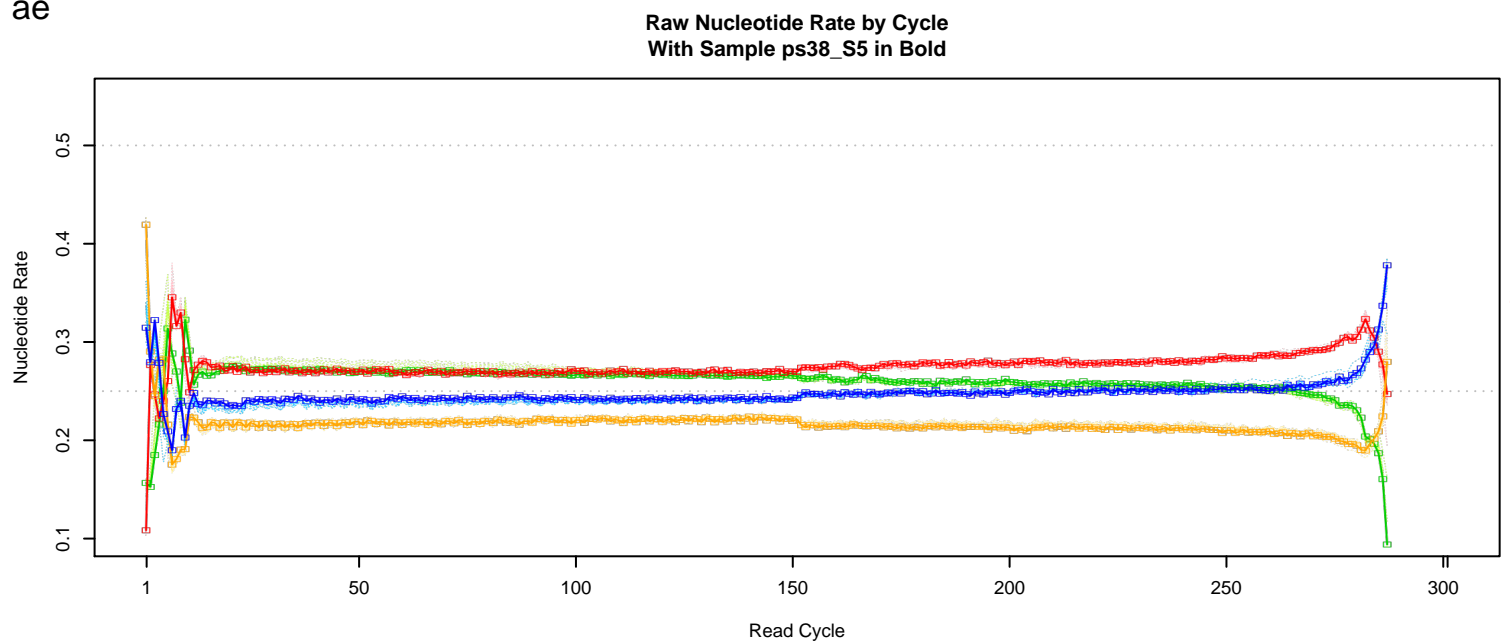


ad

Nucleotide Rate by Cycle, Trailing Clipped bases (12)
With Sample ps38_S5 in Bold



ae



af

