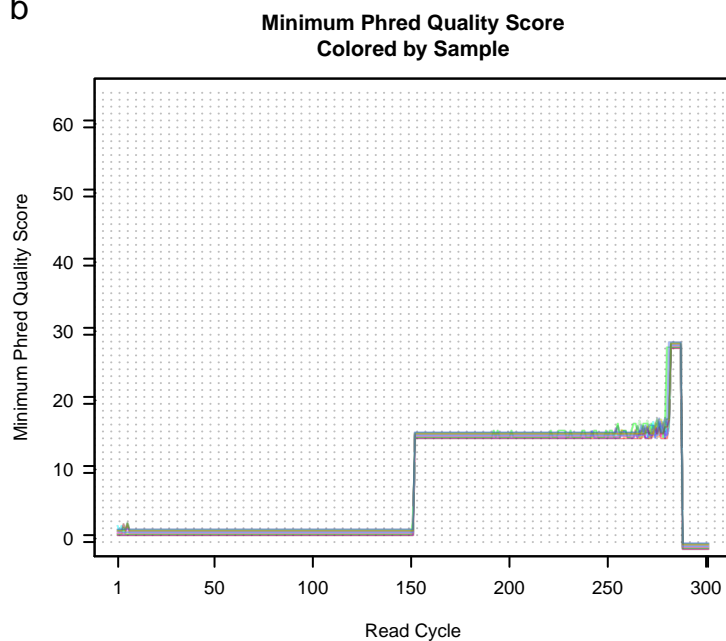


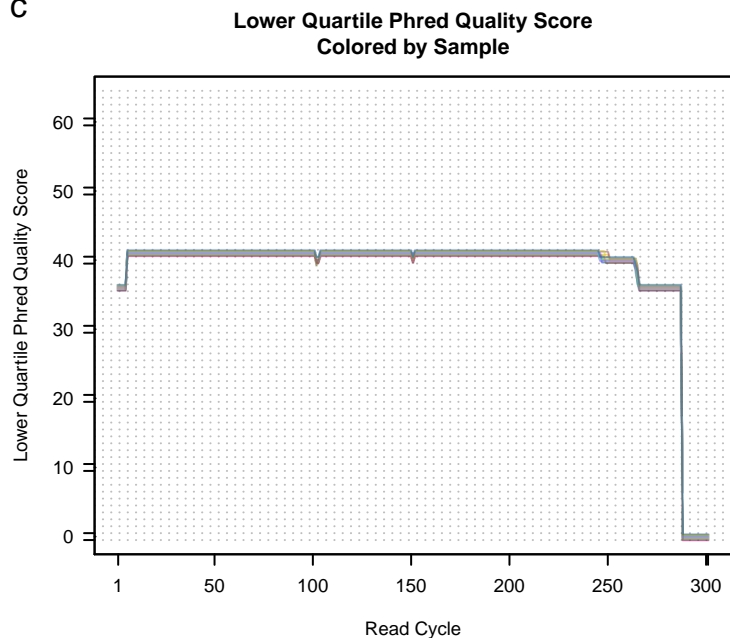
a

All Quality Control Summary Plots

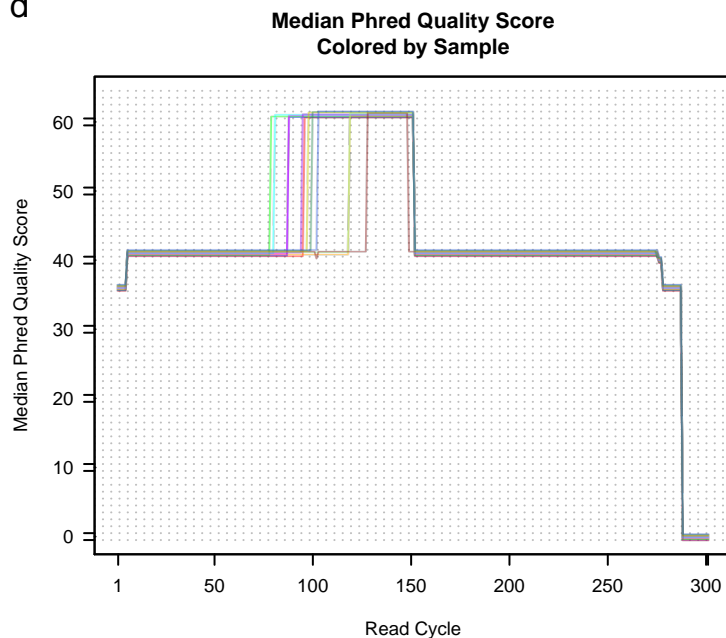
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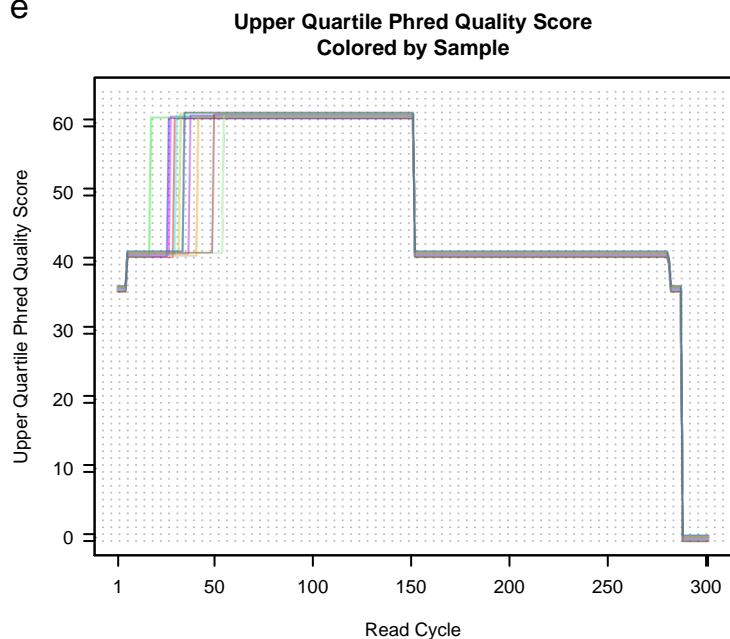
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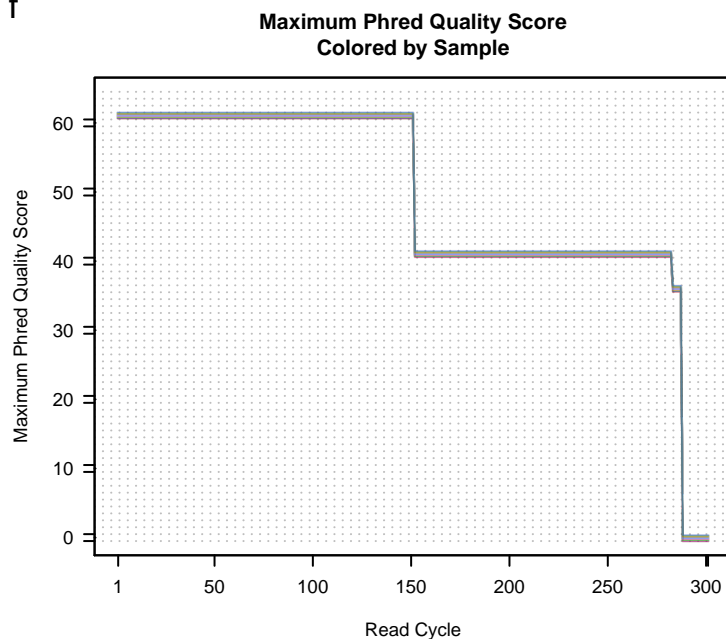
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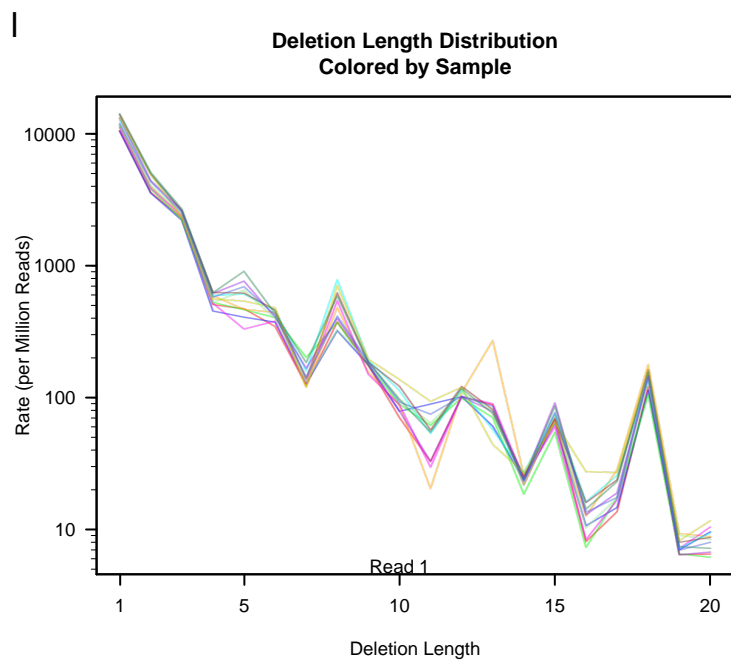
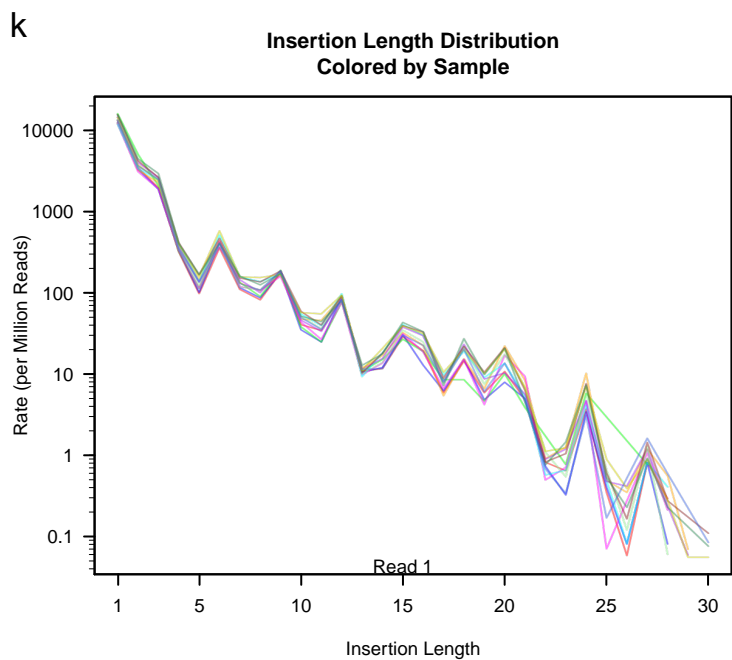
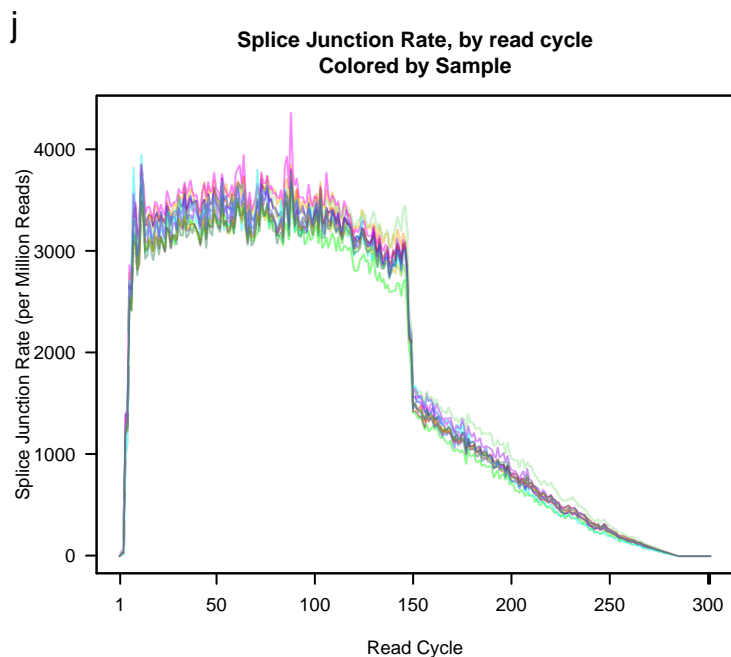
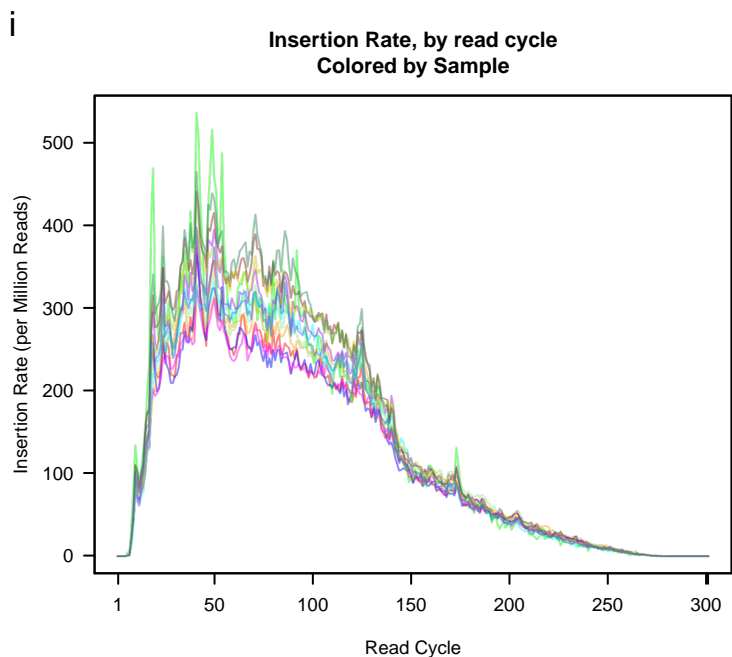
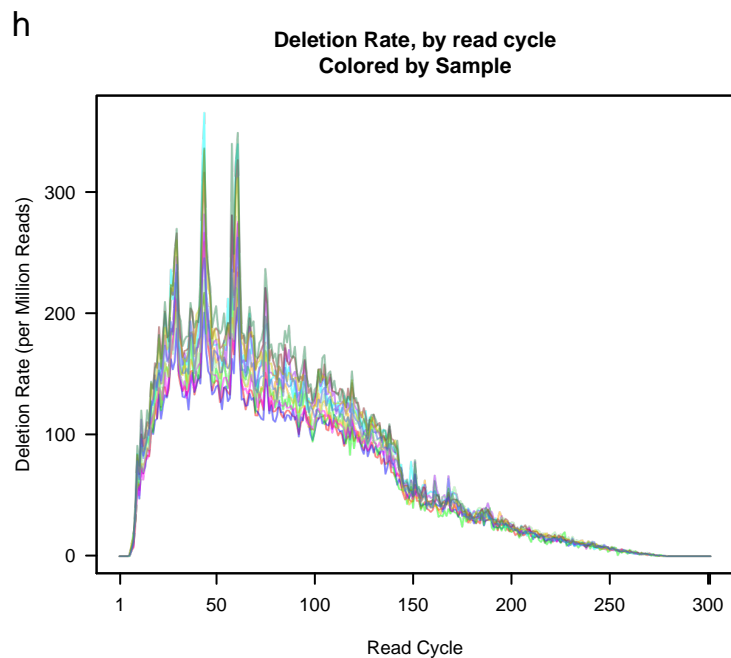
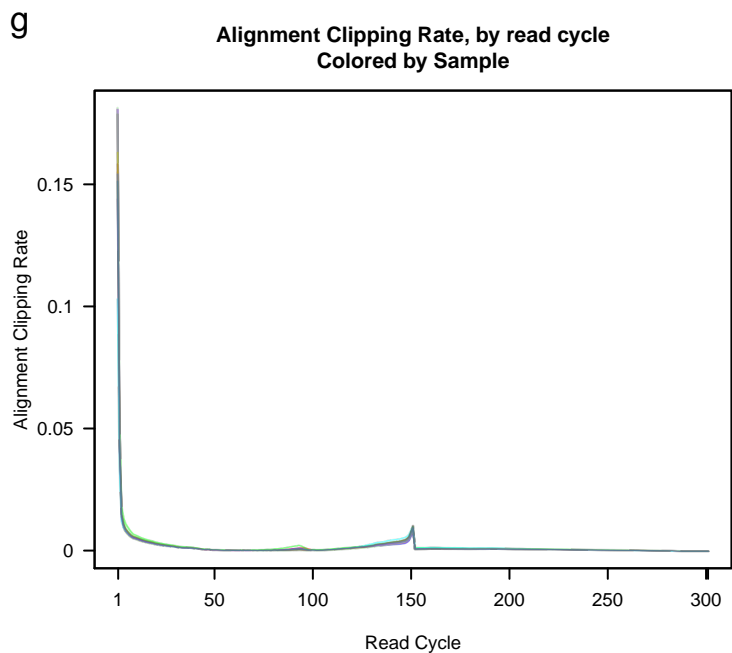


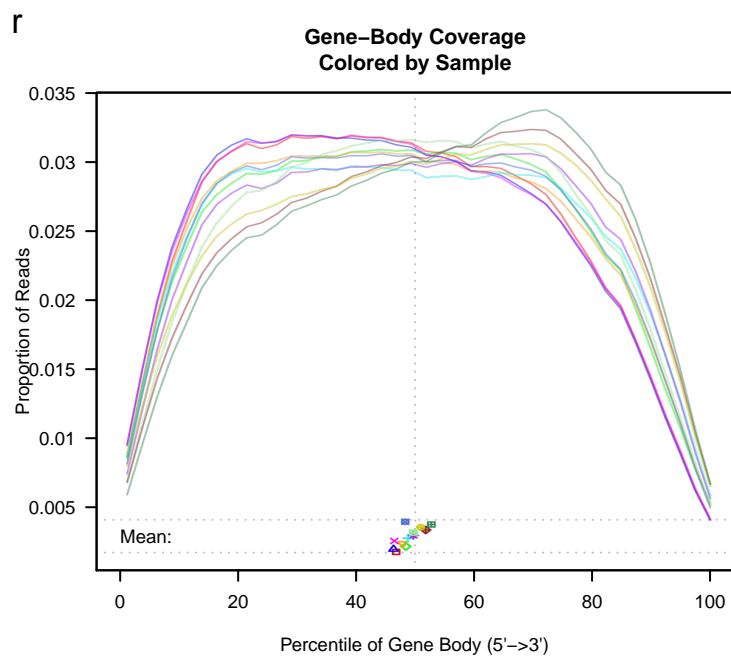
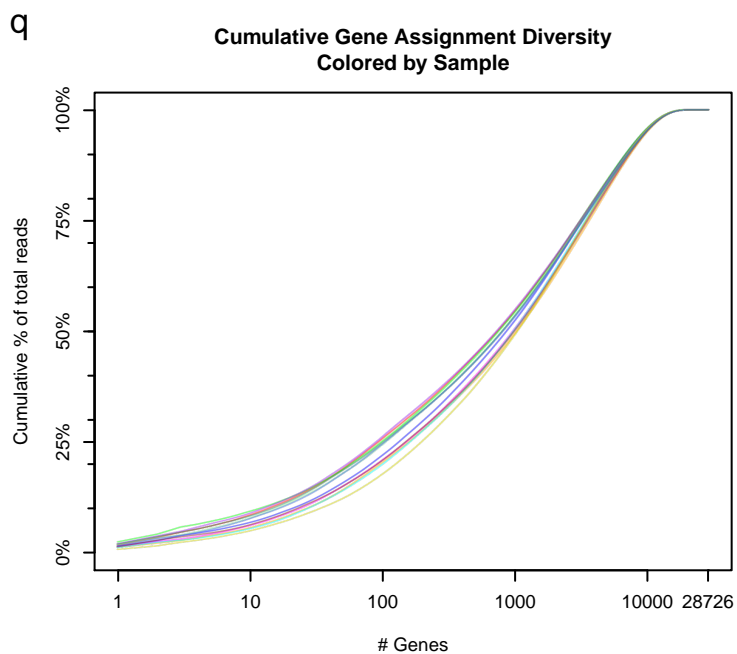
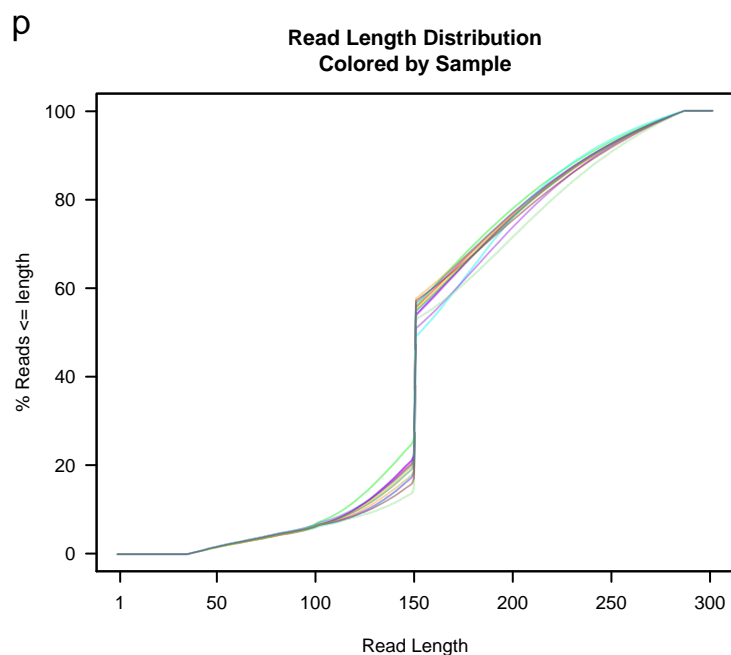
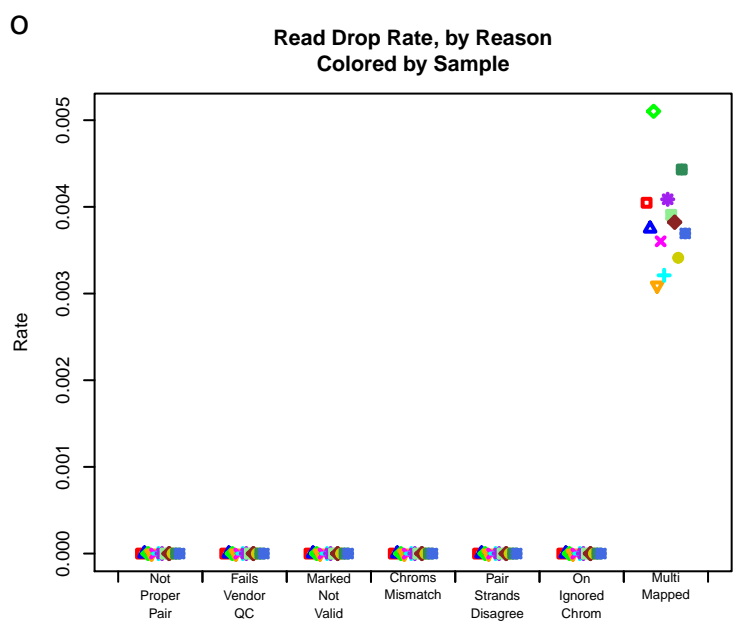
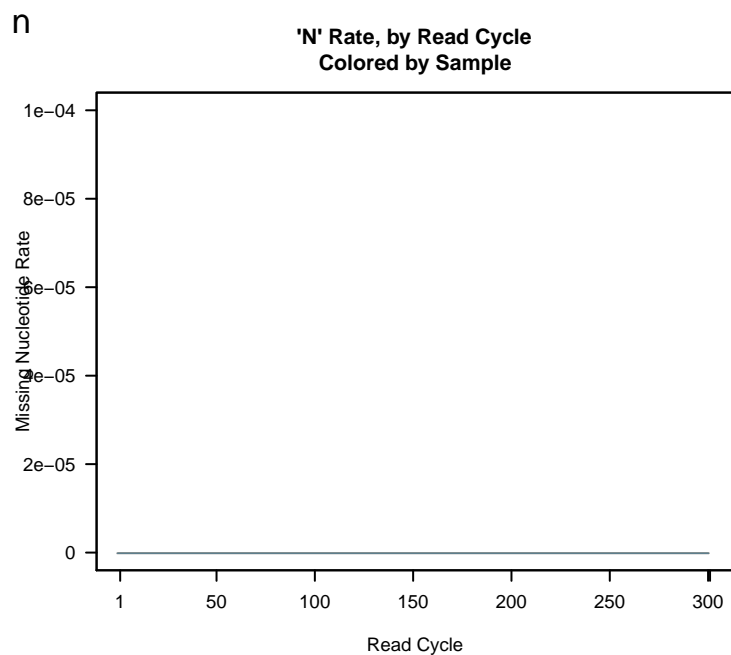
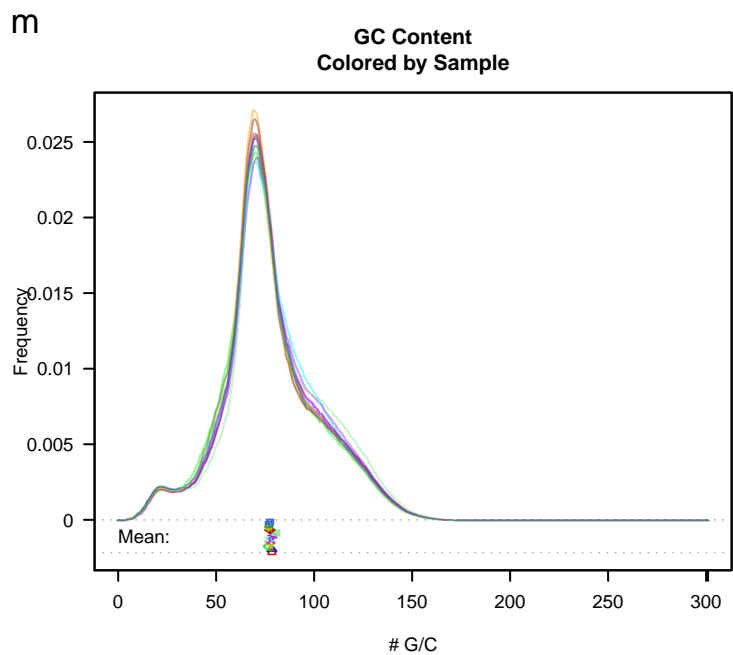
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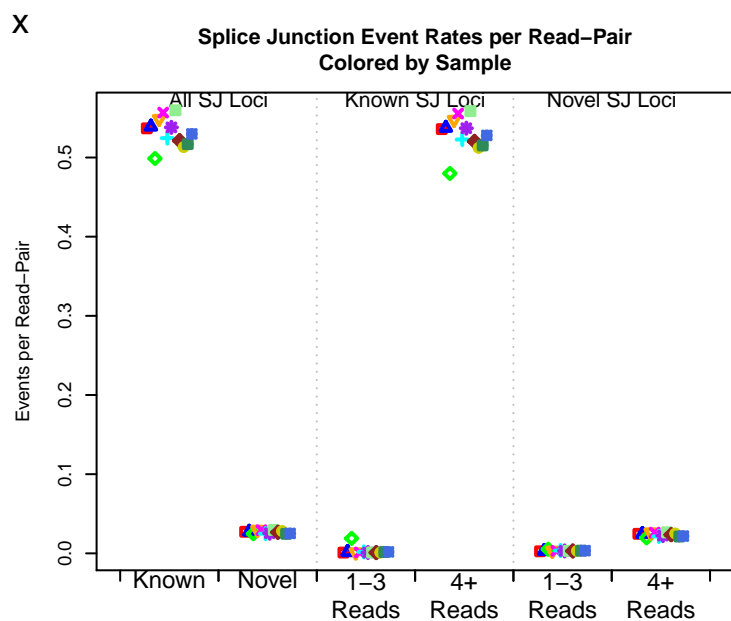
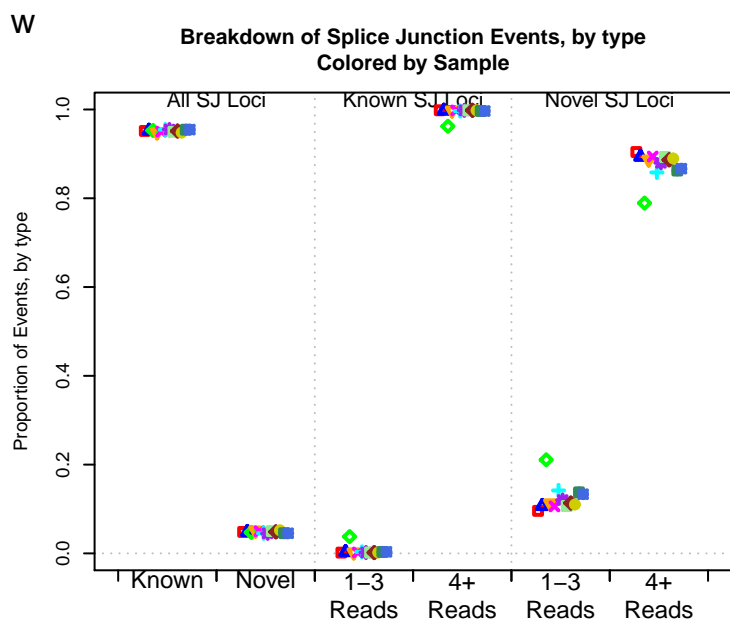
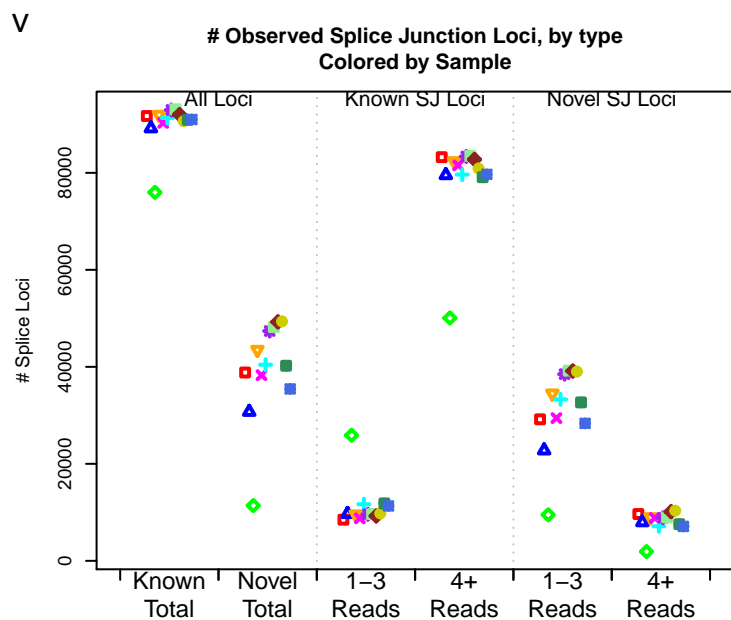
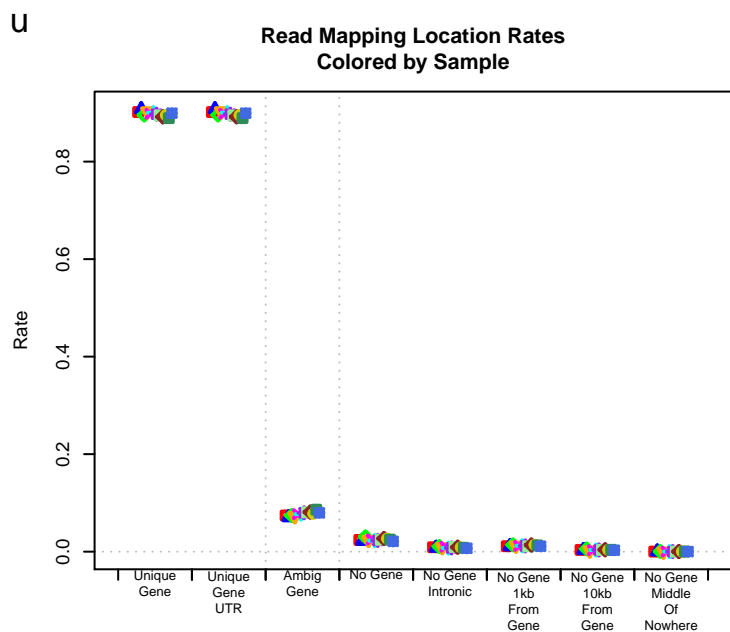
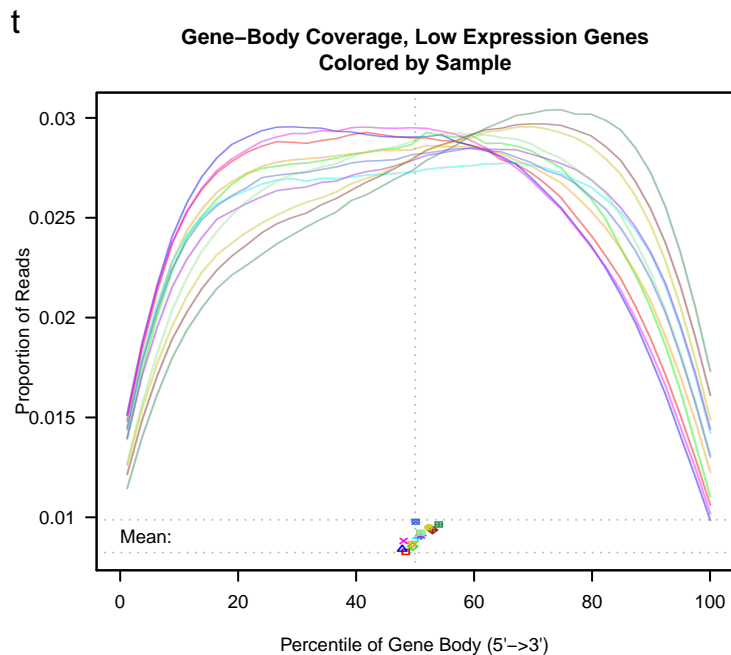
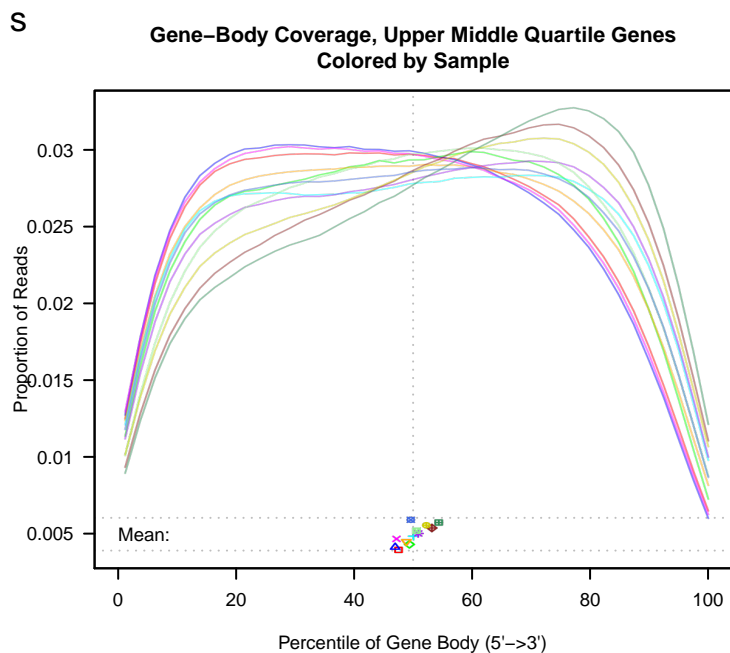


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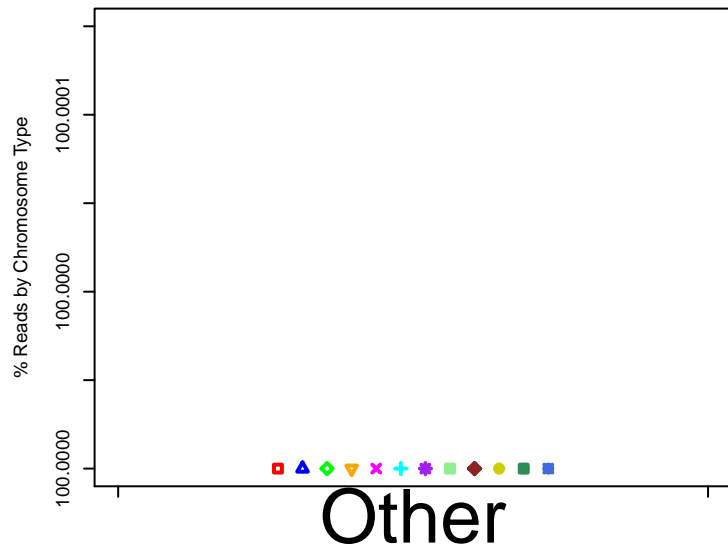






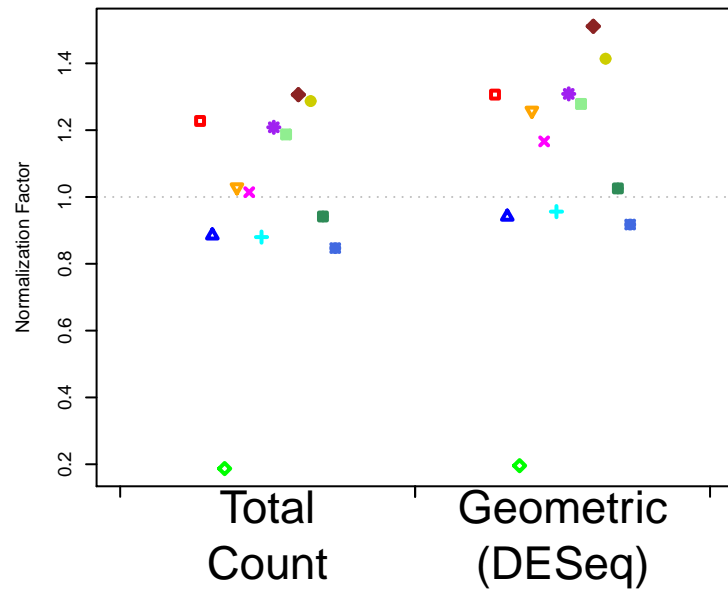
y

Chromosome Distribution (Excluding Autosomes)
Colored by Sample



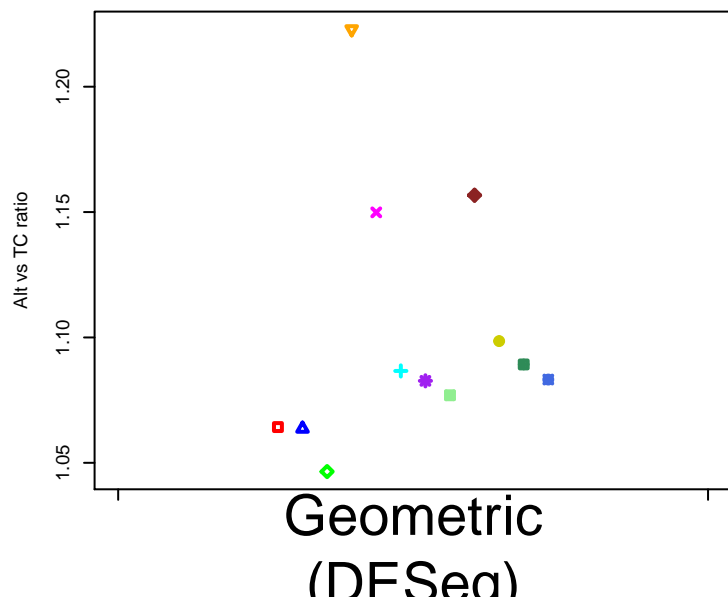
Z

Normalization Factors
Colored by Sample



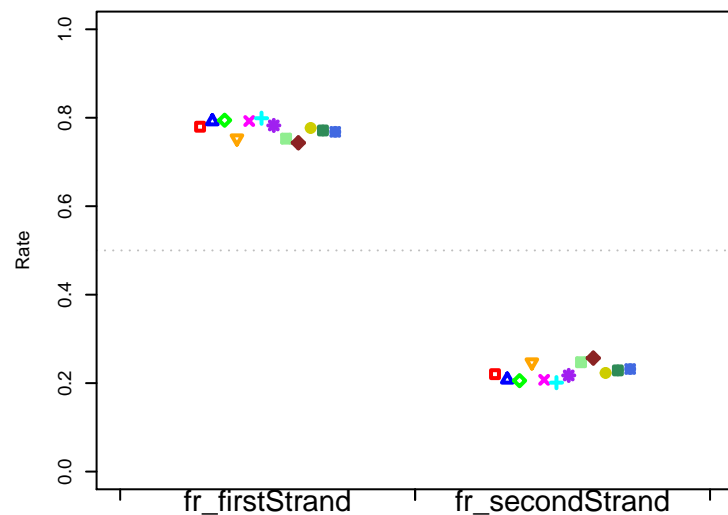
aa

Normalization Factors vs Total-Count Normalization
Colored by Sample



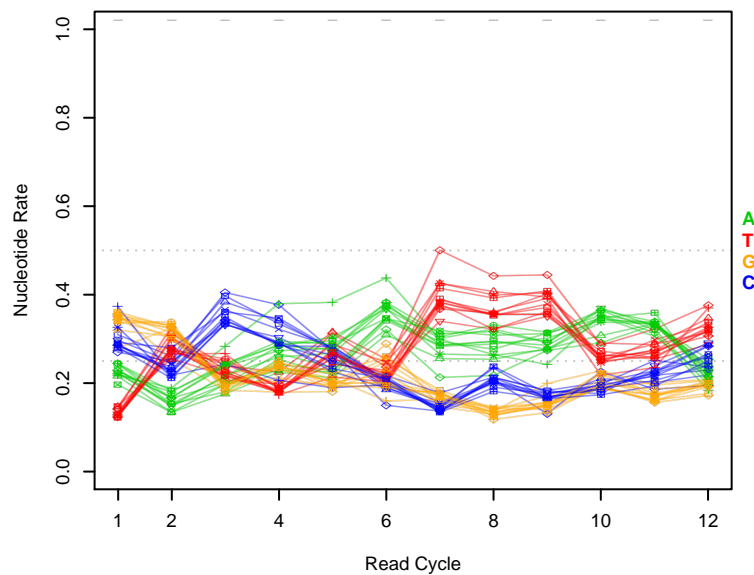
ab

Strandedness Test
Colored by Sample



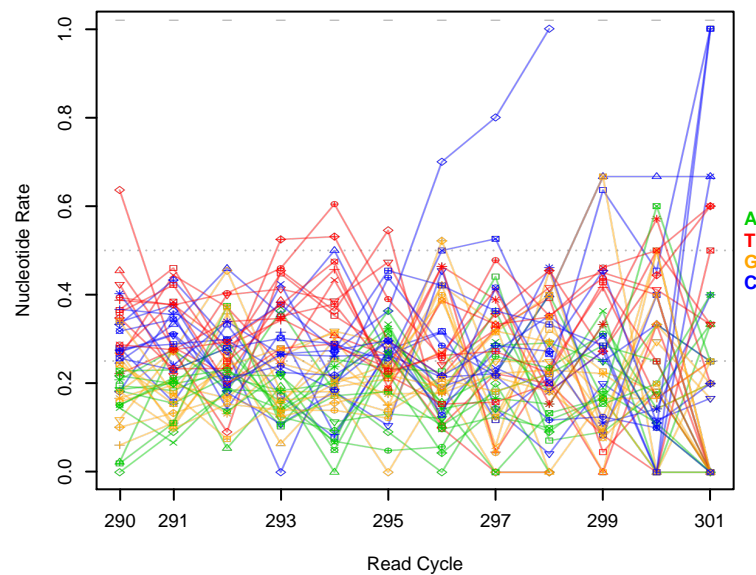
ac

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

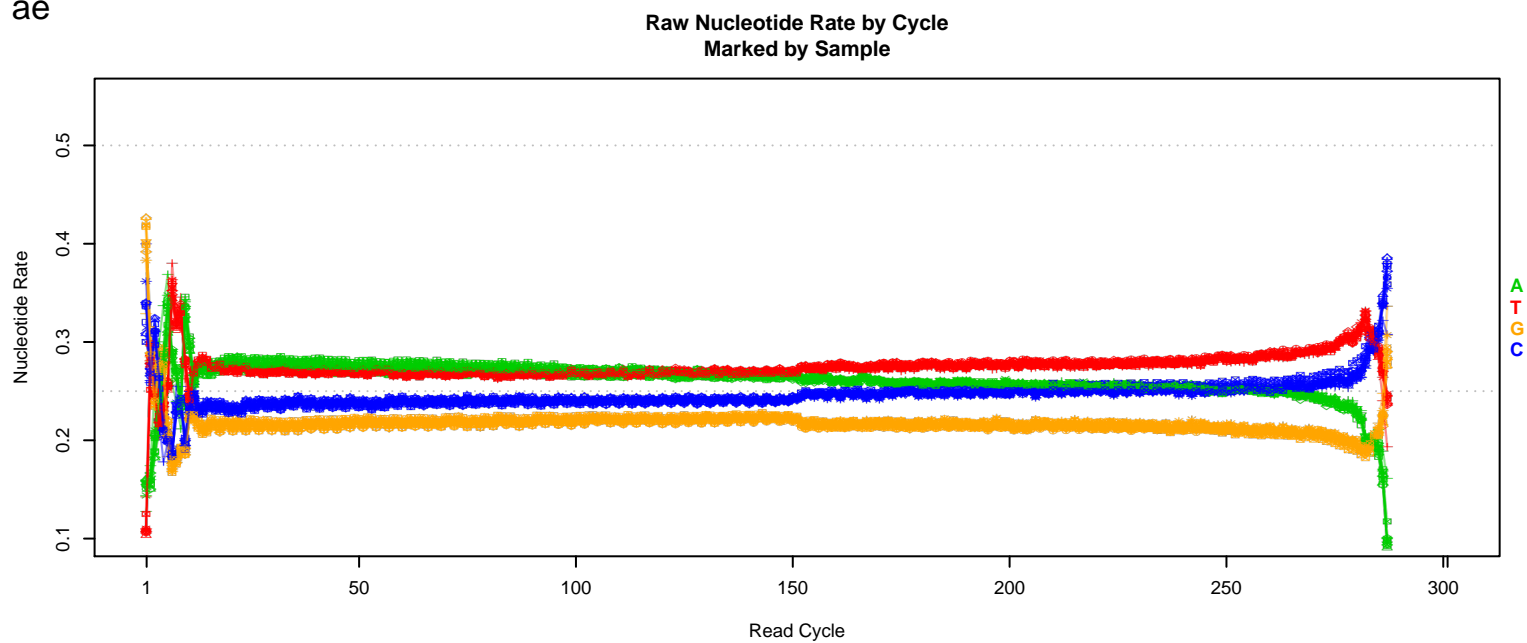


ad

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



ae



af

