

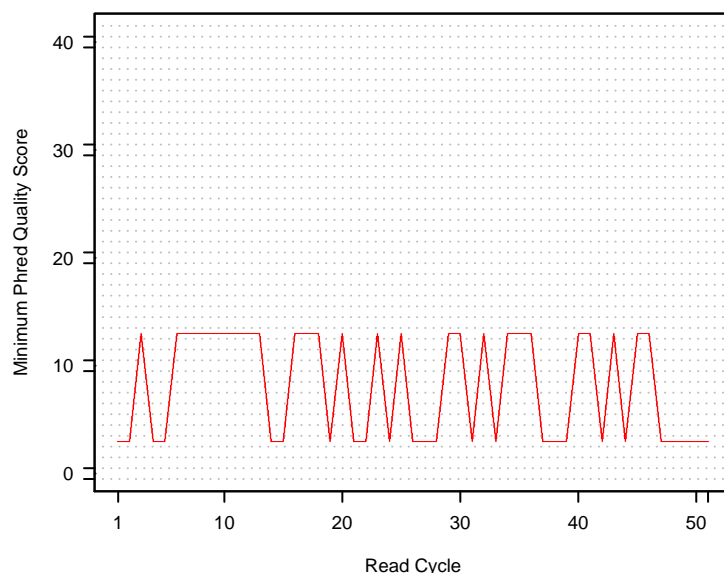
a

Sample Highlight:
a2_aligned_sorted
Colored by Lane



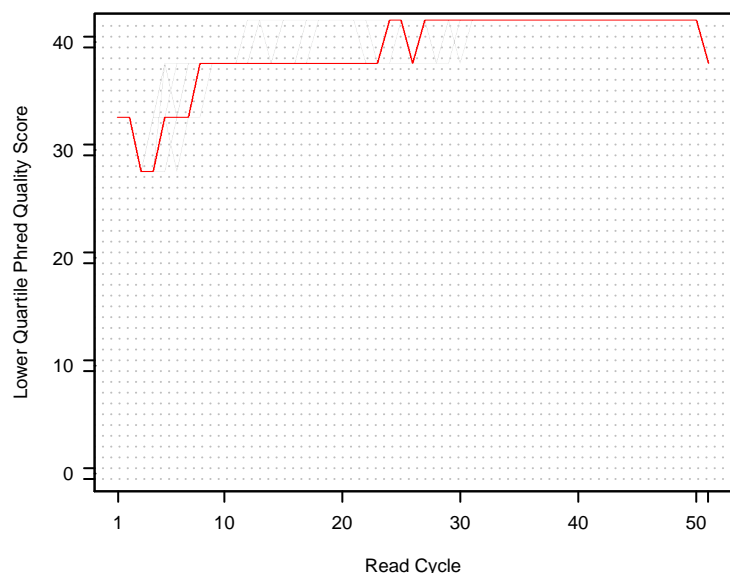
b

Minimum Phred Quality Score
With Sample a2_aligned_sorted Colored by lane



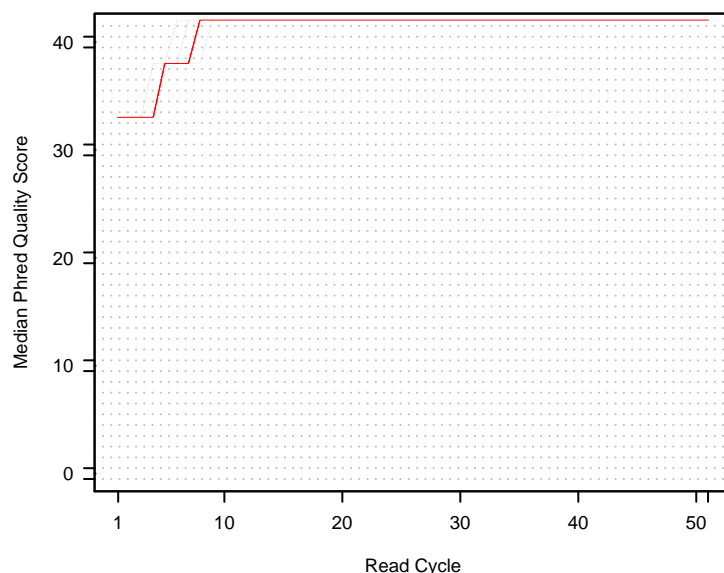
c

Lower Quartile Phred Quality Score
With Sample a2_aligned_sorted Colored by lane



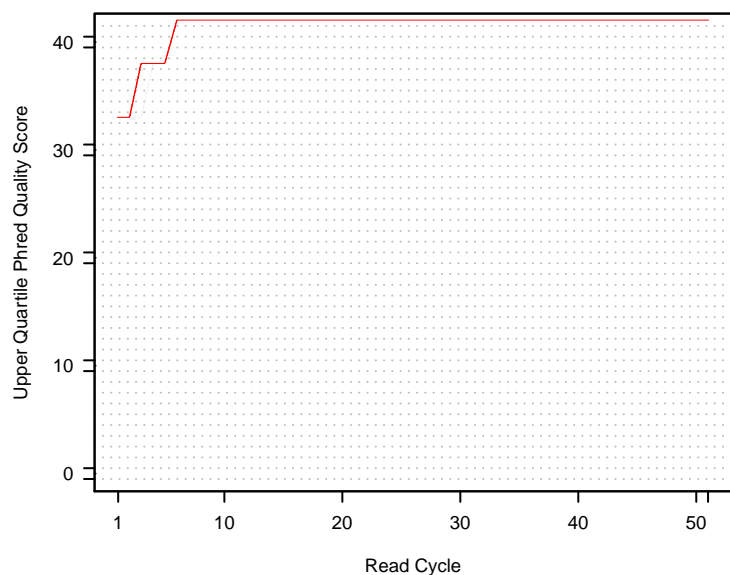
d

Median Phred Quality Score
With Sample a2_aligned_sorted Colored by lane



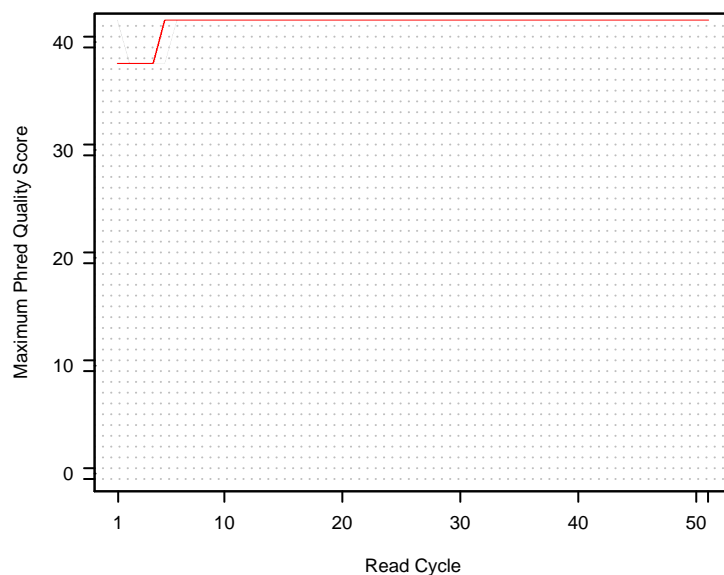
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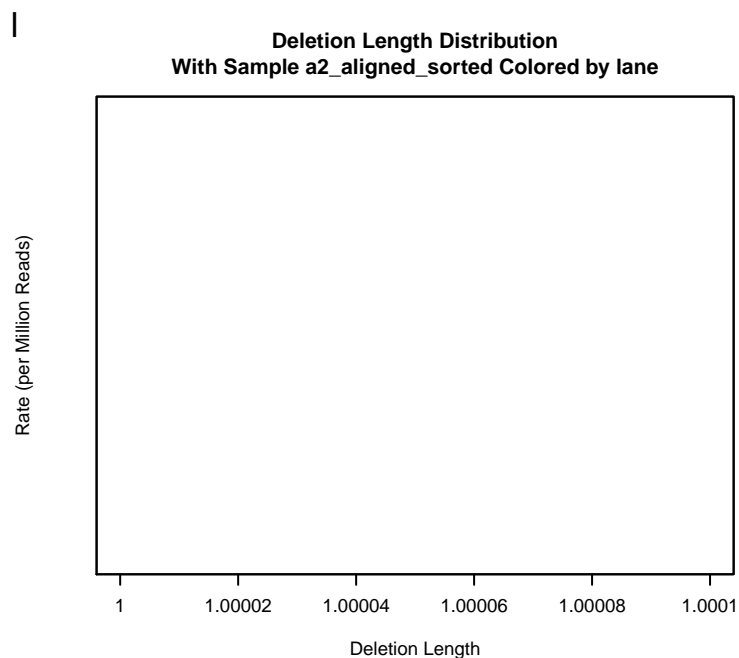
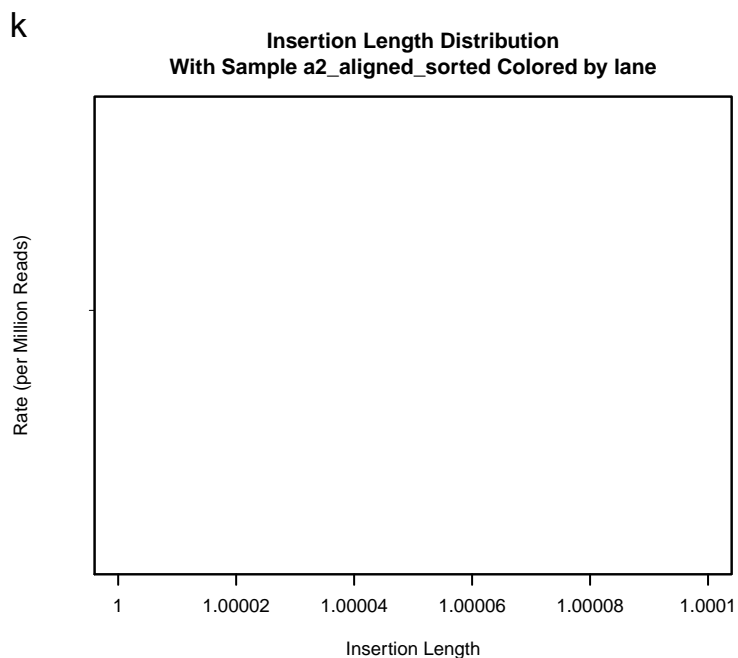
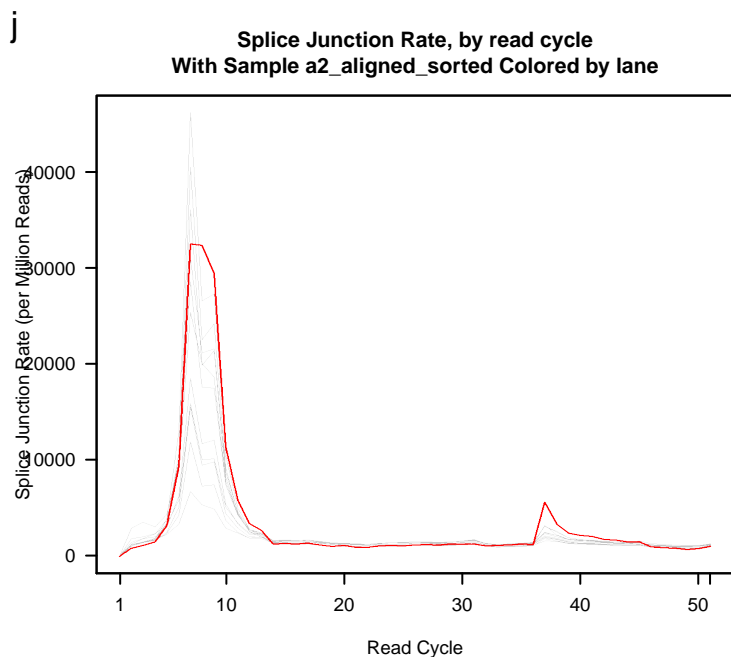
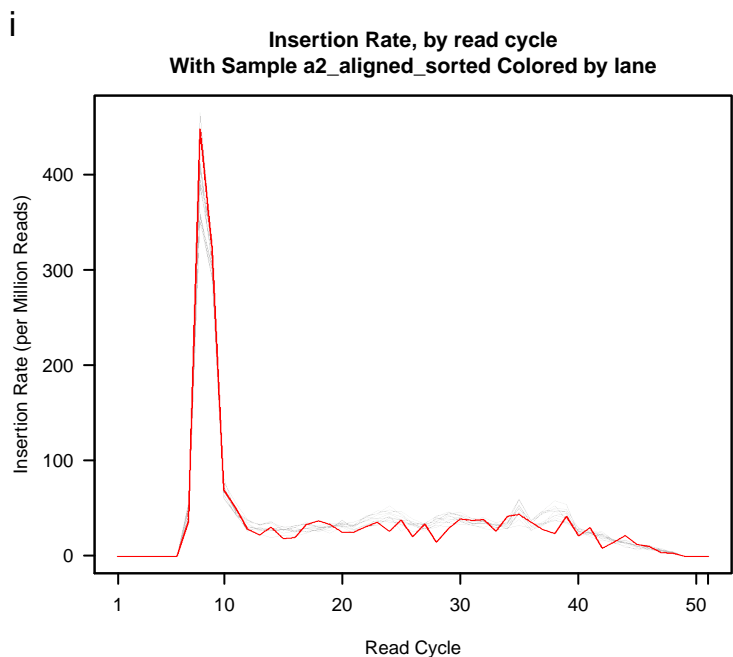
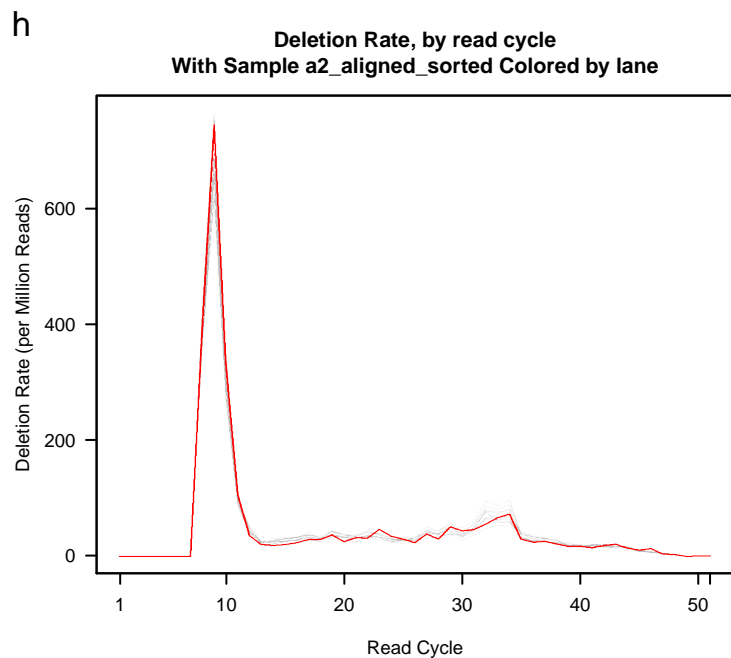
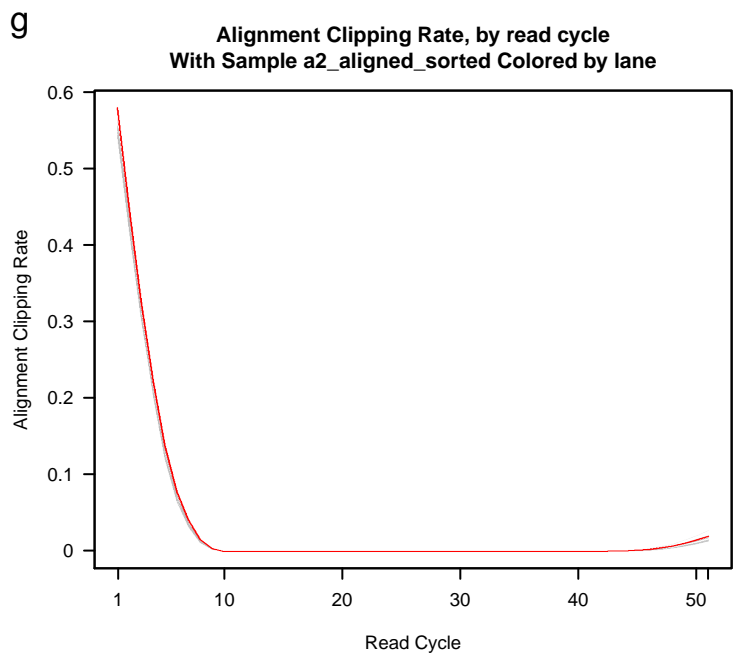
Upper Quartile Phred Quality Score
With Sample a2_aligned_sorted Colored by lane

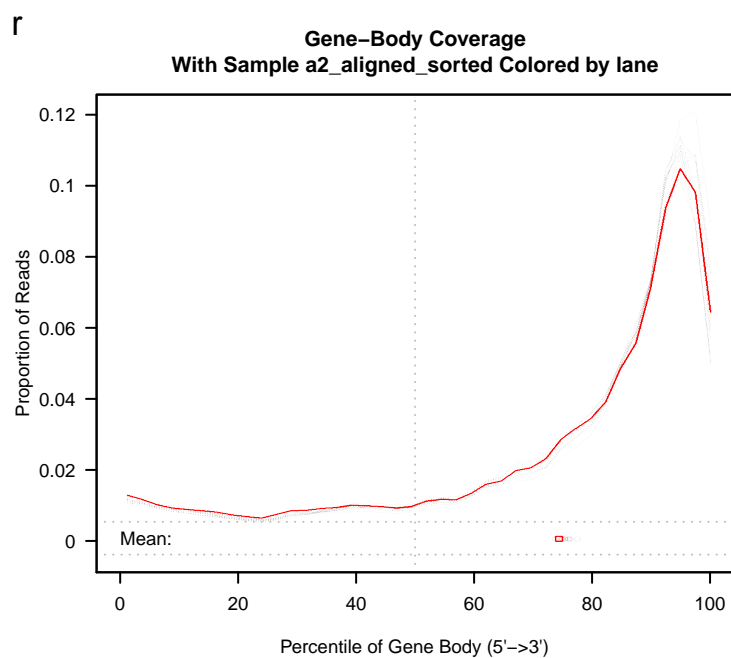
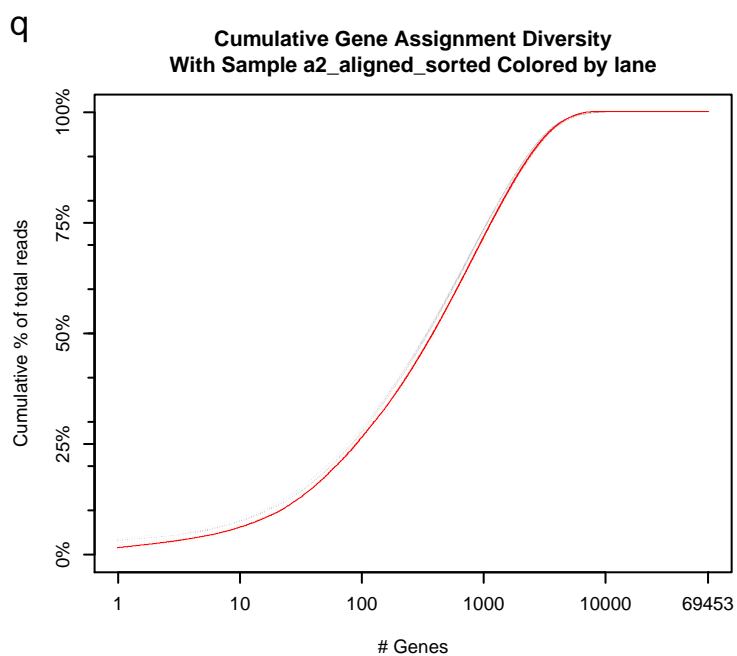
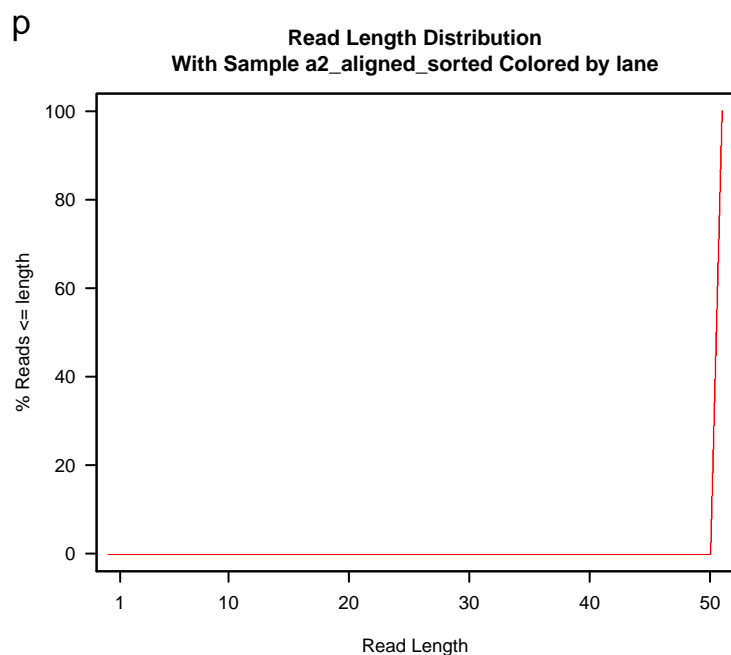
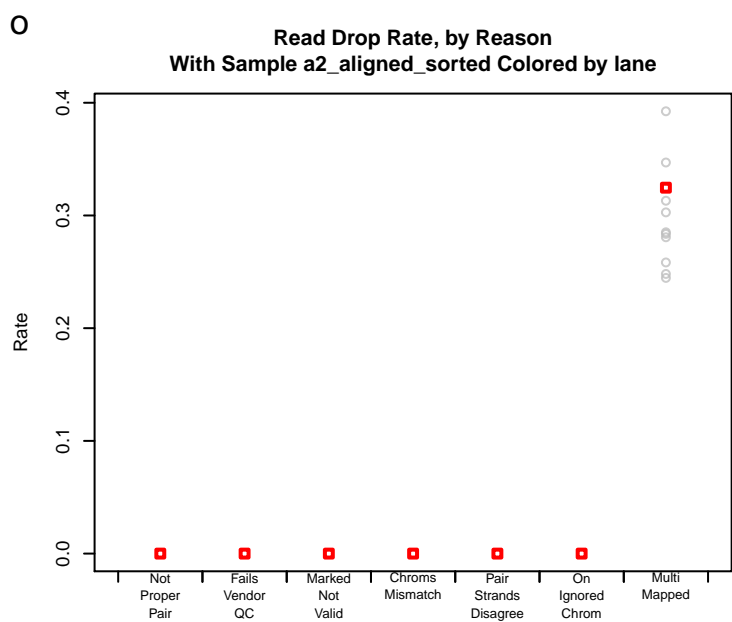
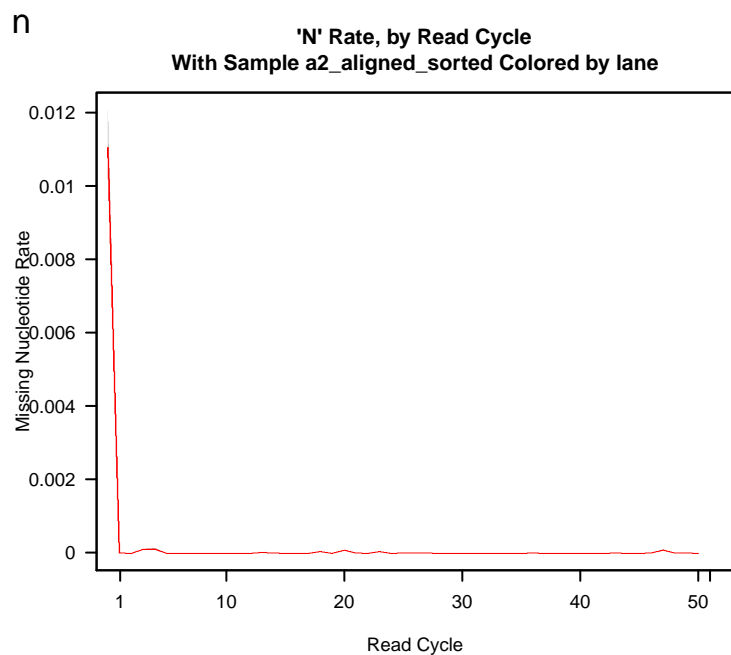
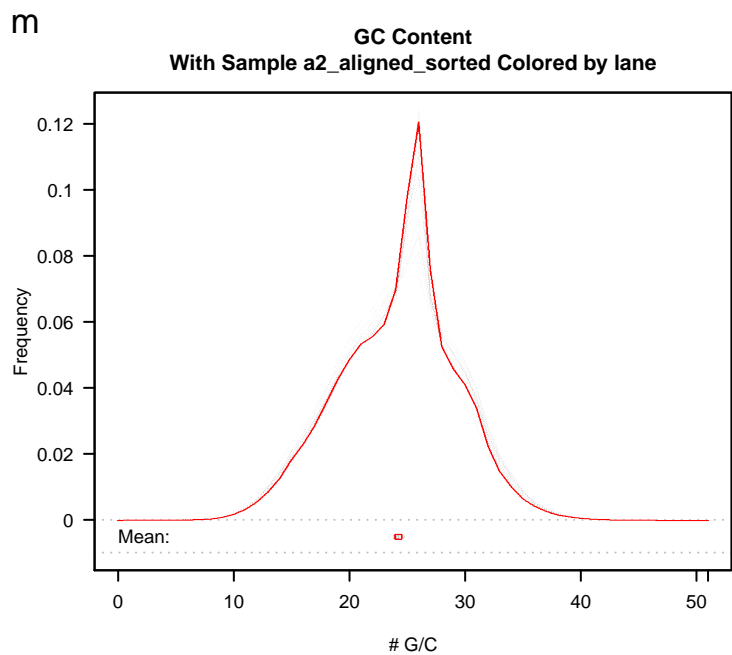


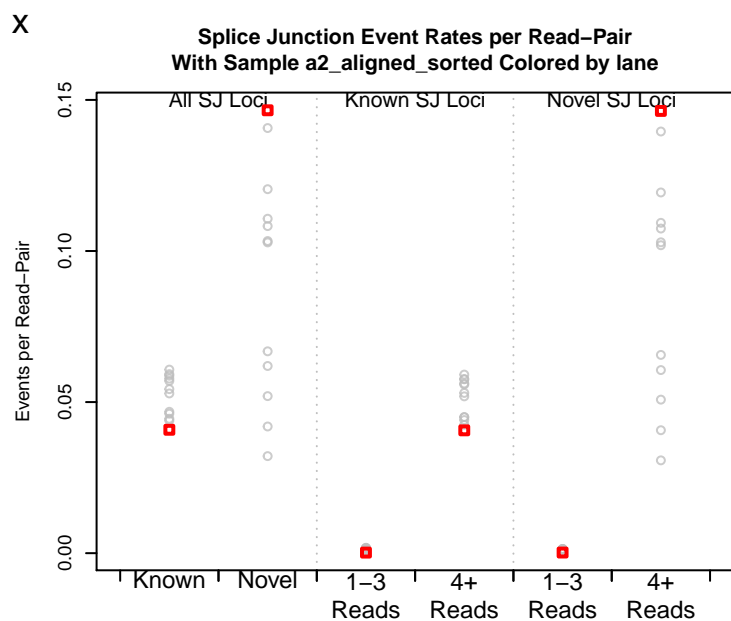
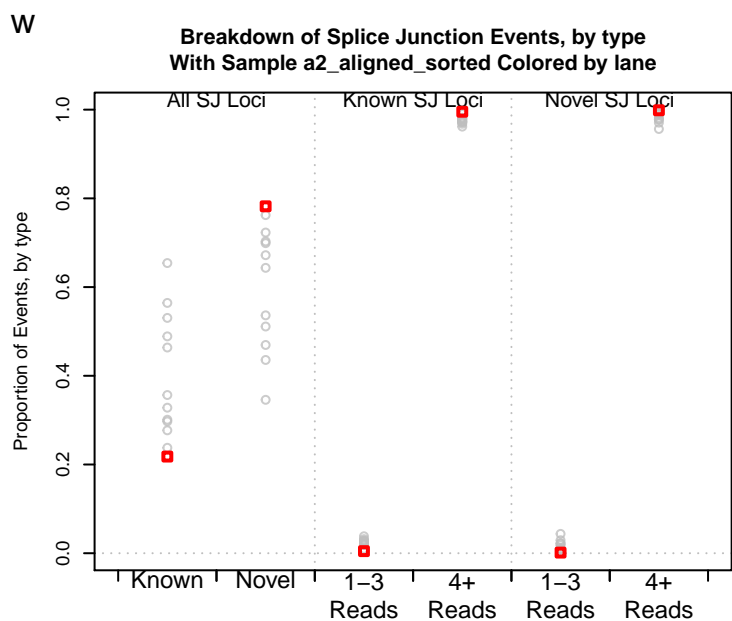
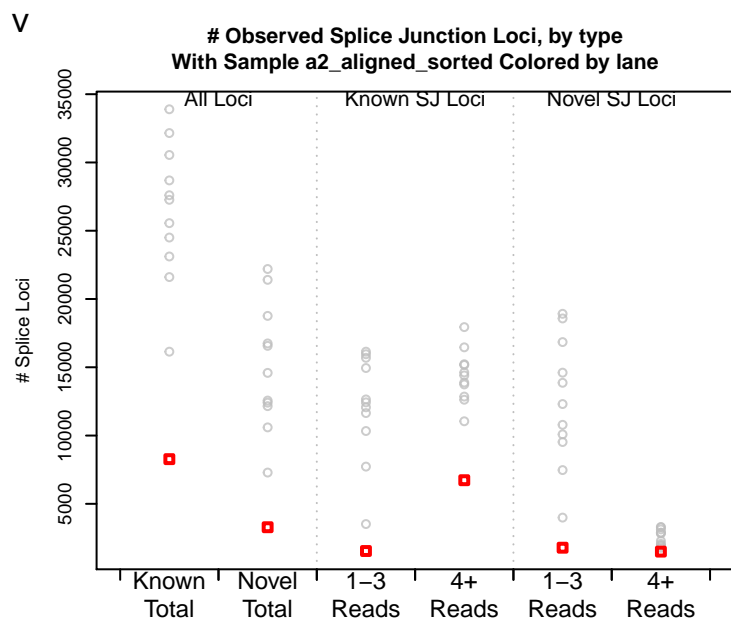
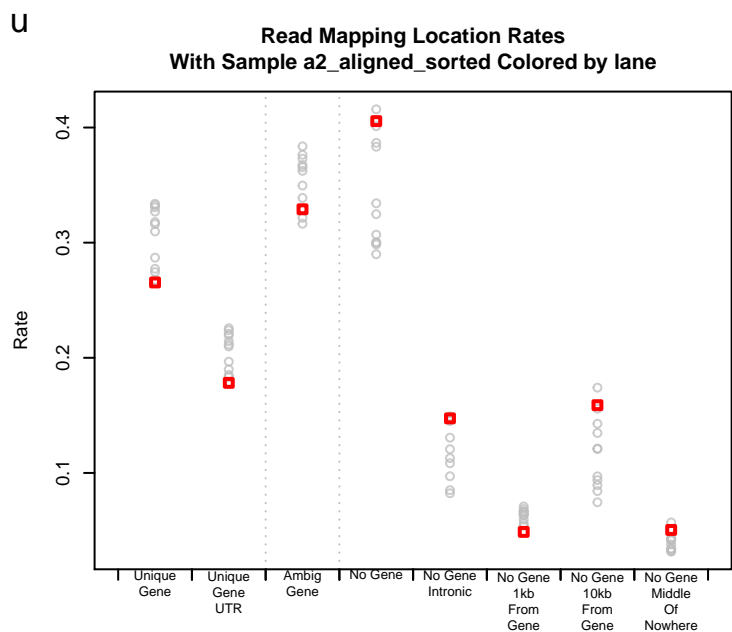
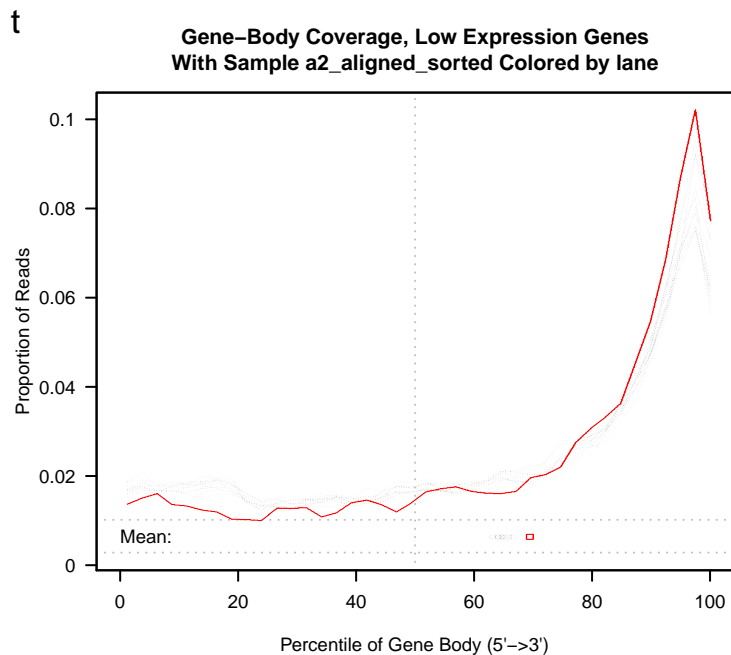
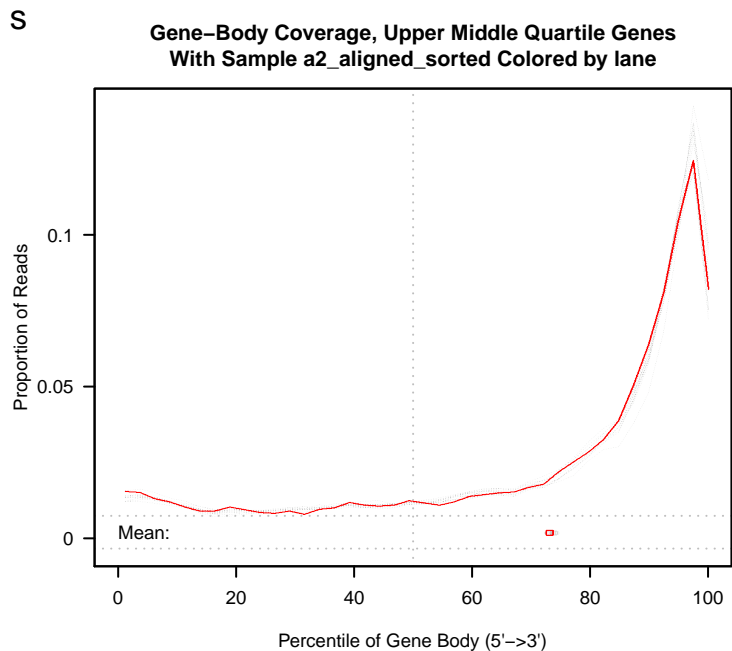
f

Maximum Phred Quality Score
With Sample a2_aligned_sorted Colored by lane

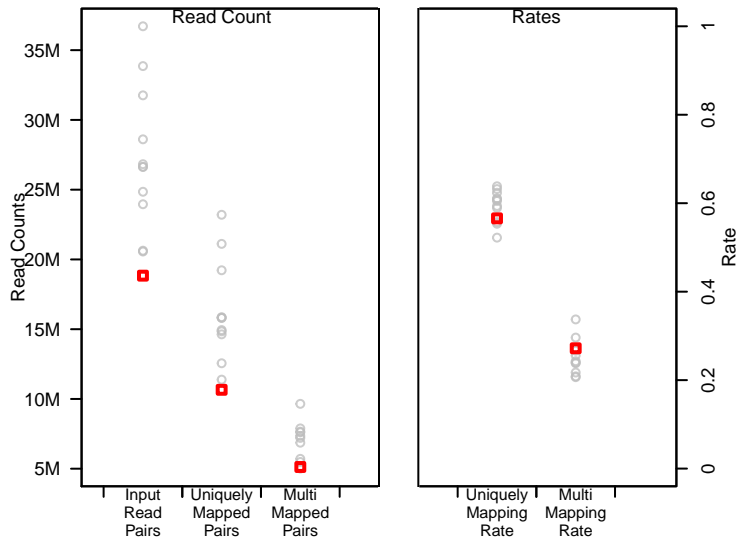




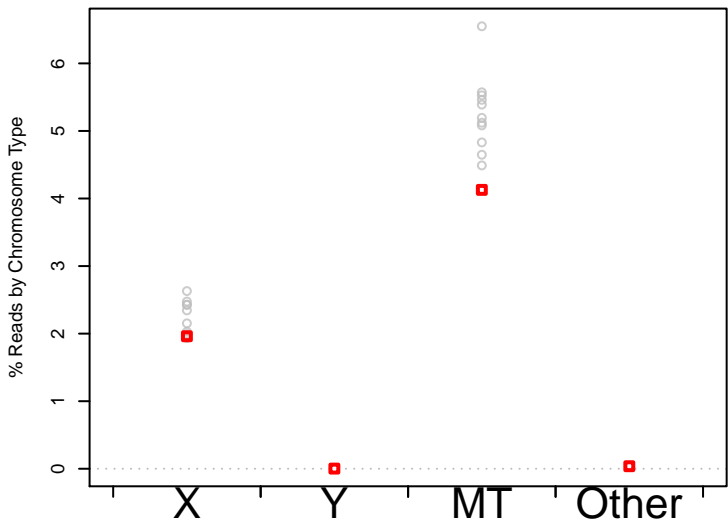




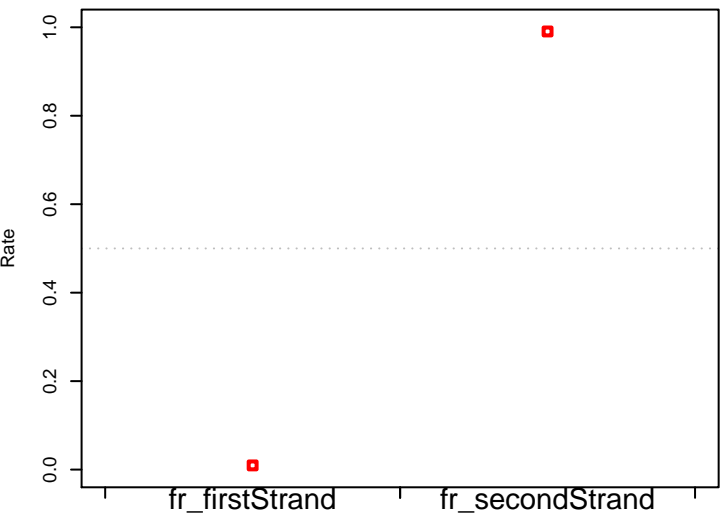
y Mapping Stats
With Sample a2_aligned_sorted Colored by lane



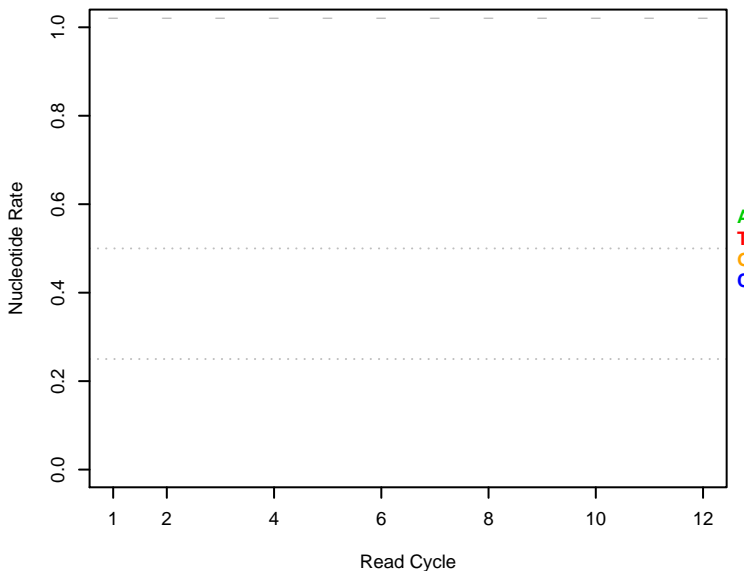
Z Chromosome Distribution (Excluding Autosomes)
With Sample a2_aligned_sorted Colored by lane



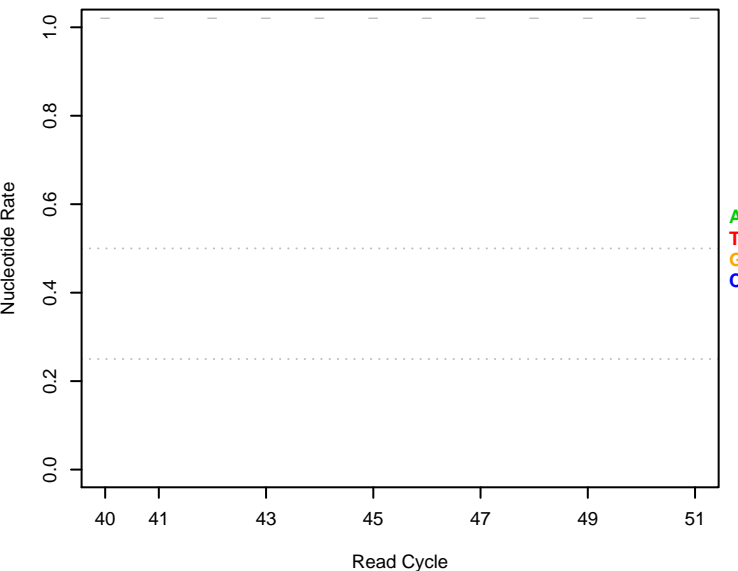
aa Strandedness Test
With Sample a2_aligned_sorted Colored by lane



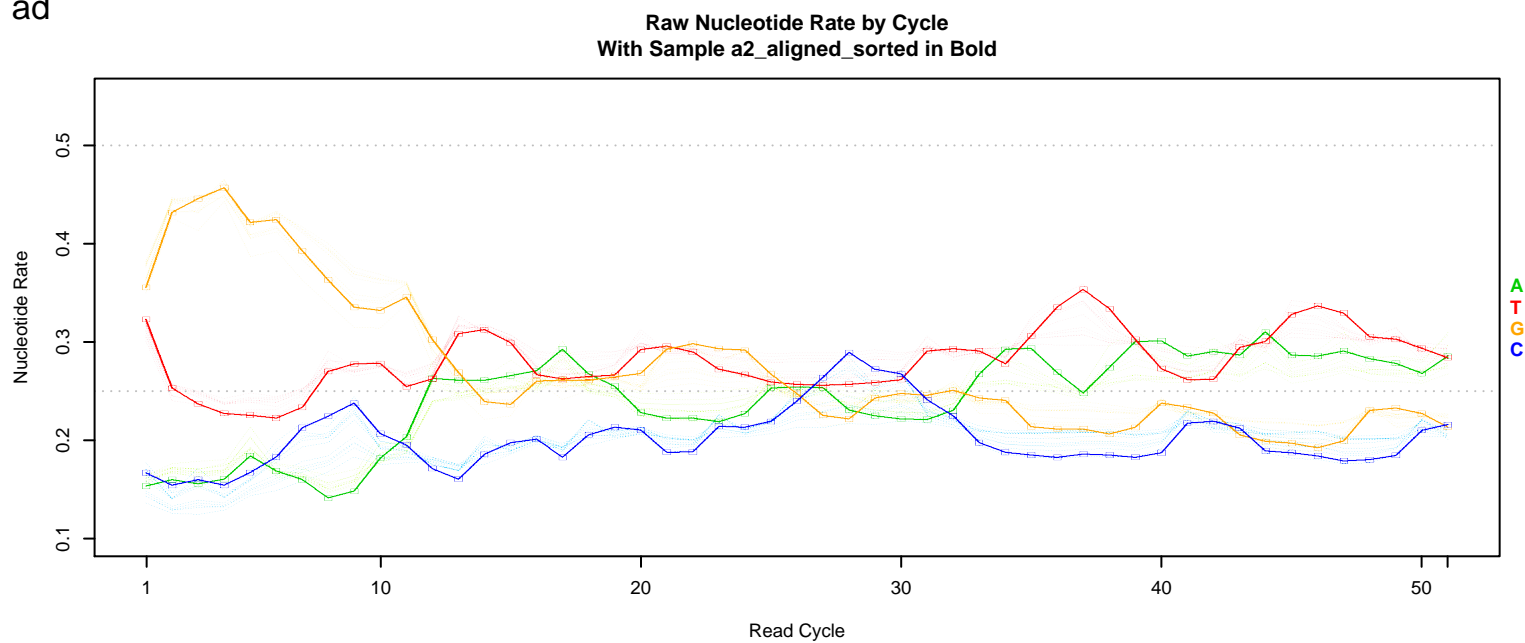
ab Nucleotide Rate by Cycle, Leading Clipped bases (12)
With Sample a2_aligned_sorted in Bold



ac Nucleotide Rate by Cycle, Trailing Clipped bases (12)
With Sample a2_aligned_sorted in Bold



ad



ae

