

a

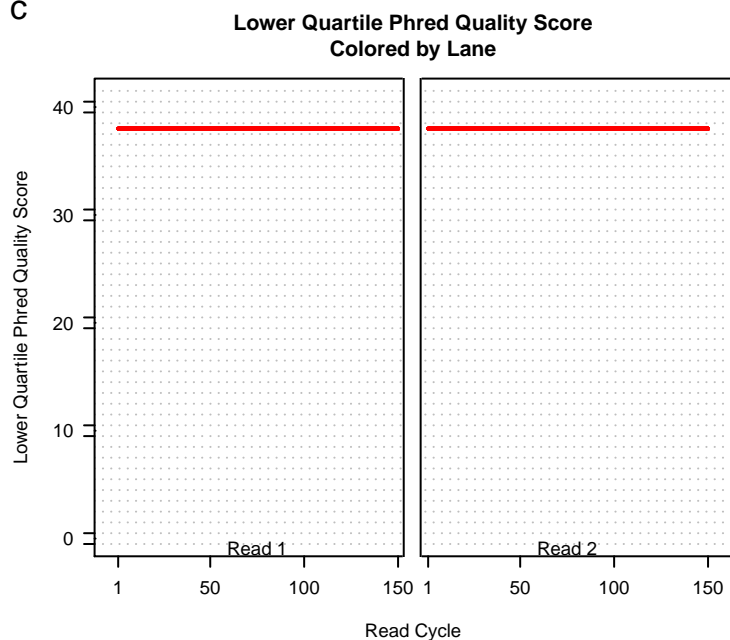
Summary Plots, By Lane



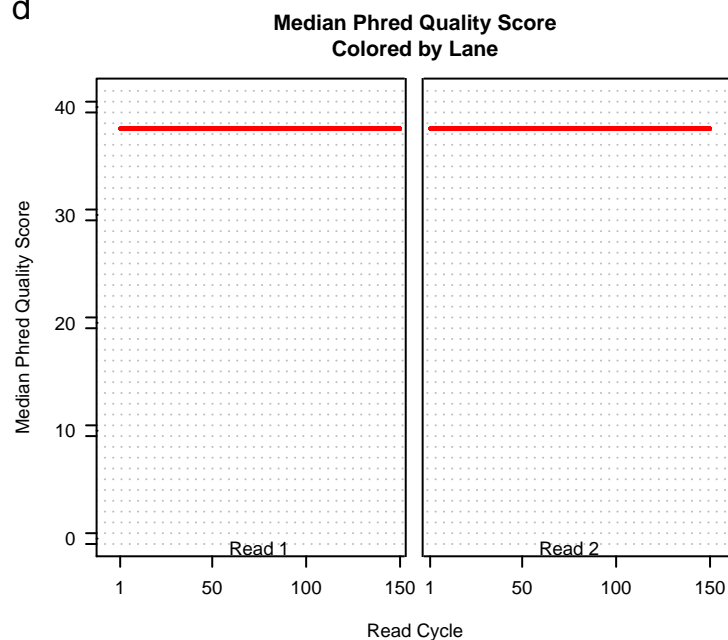
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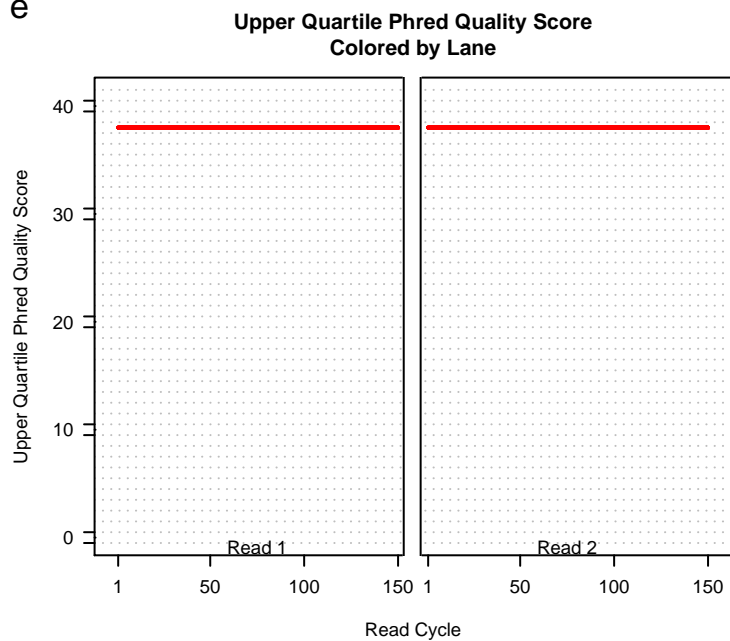
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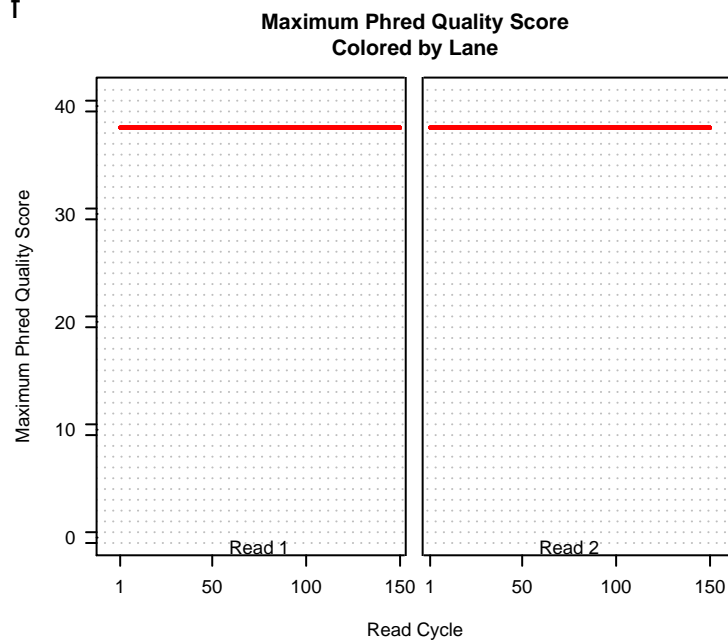
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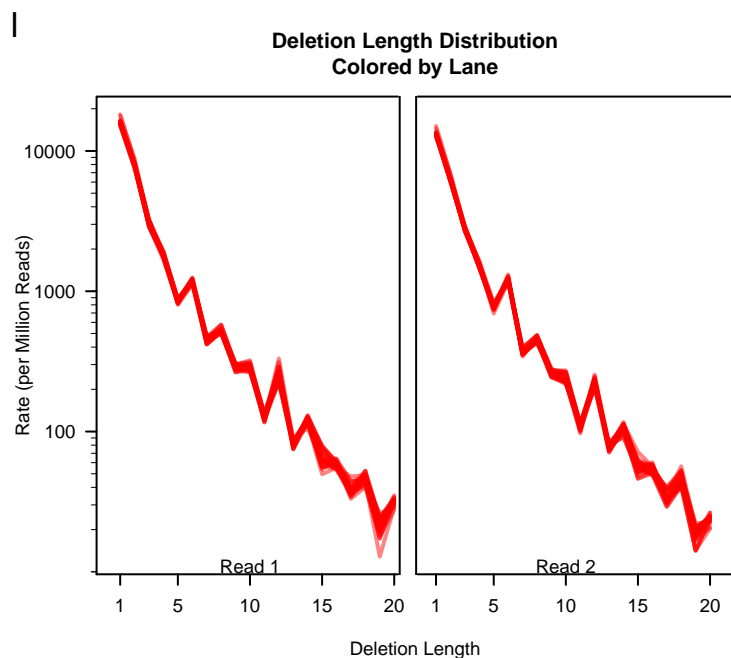
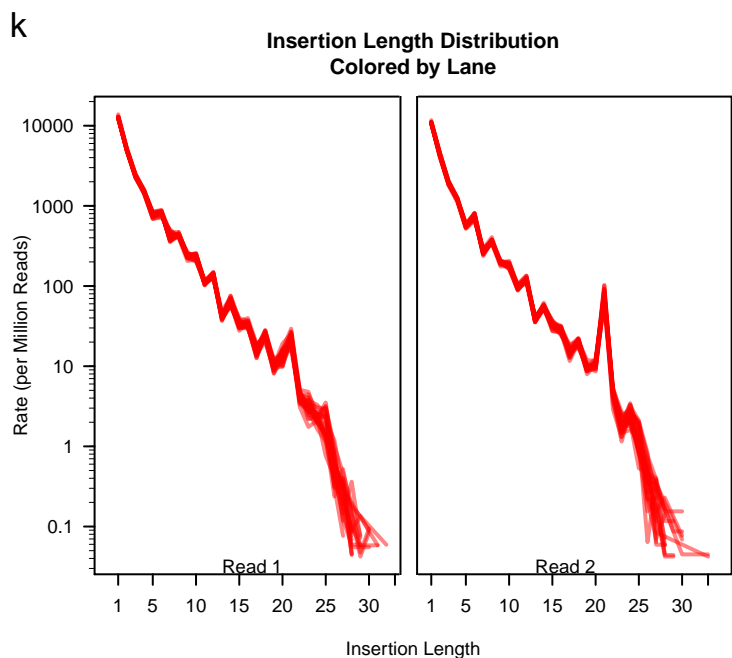
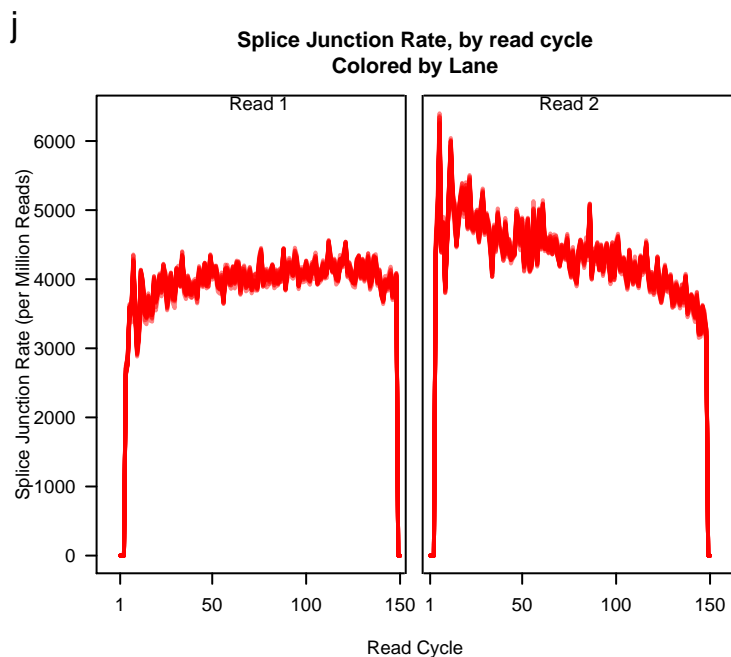
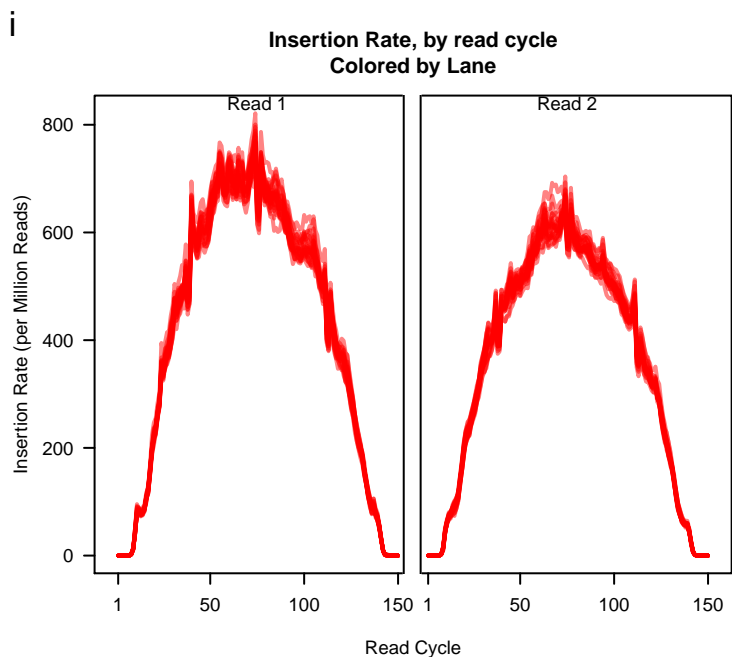
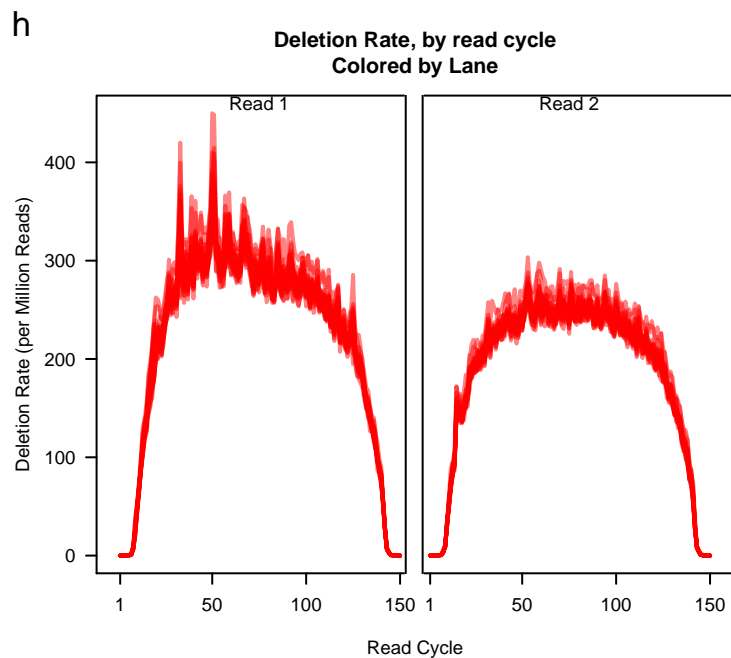
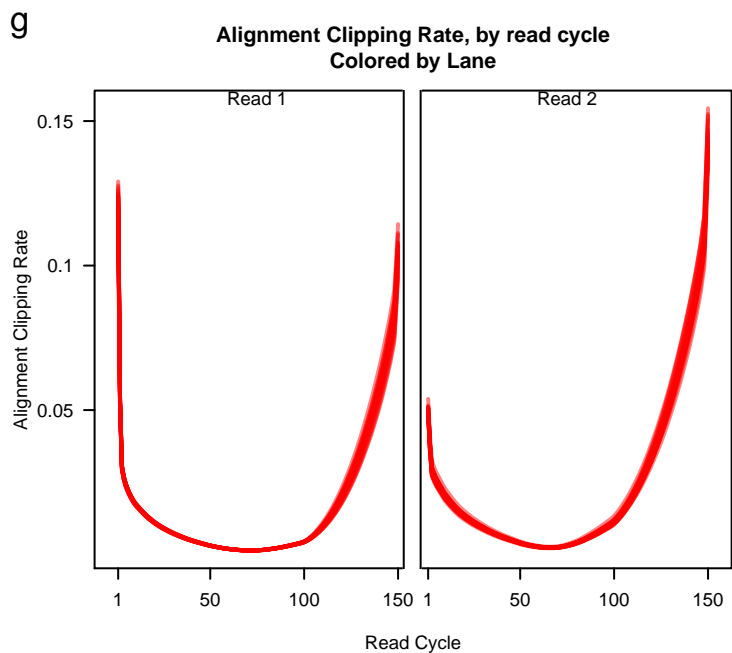


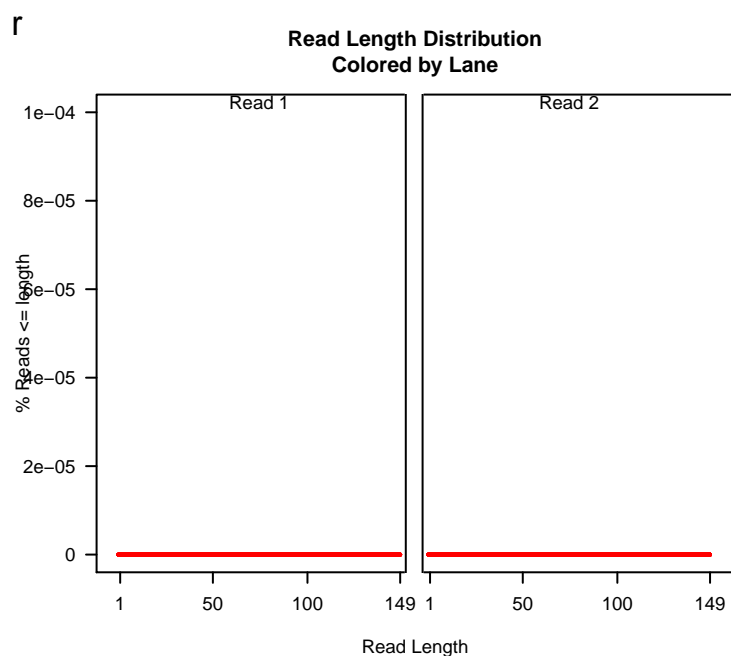
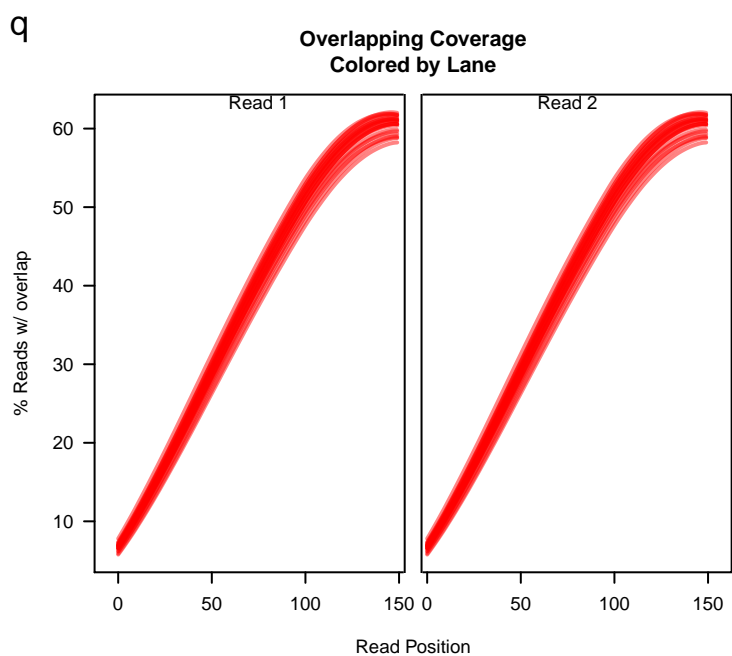
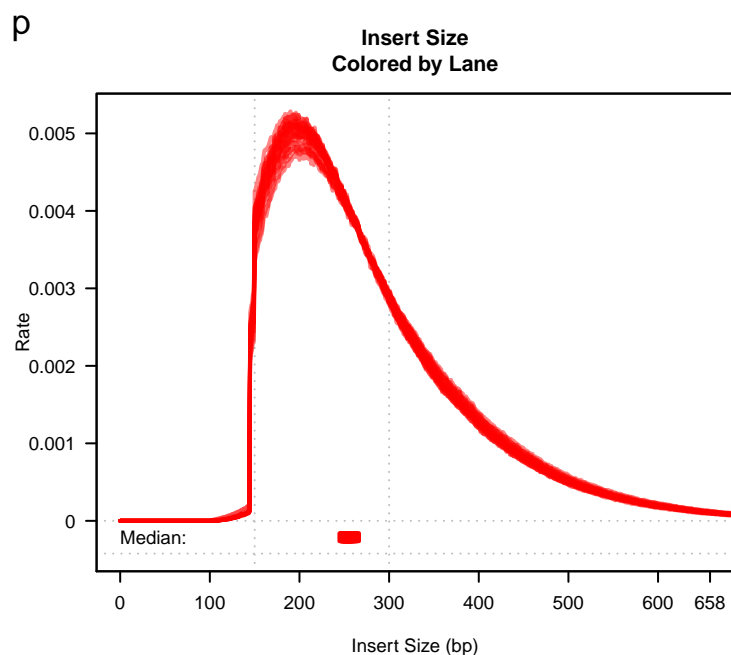
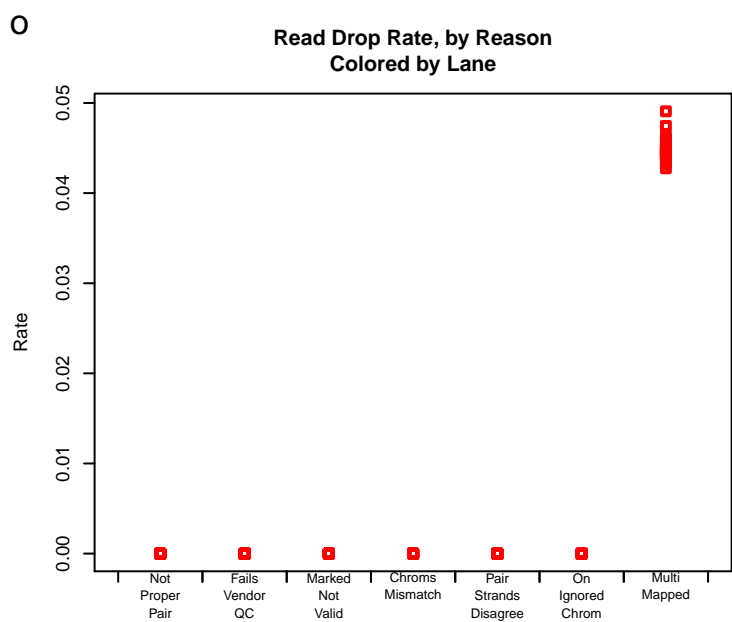
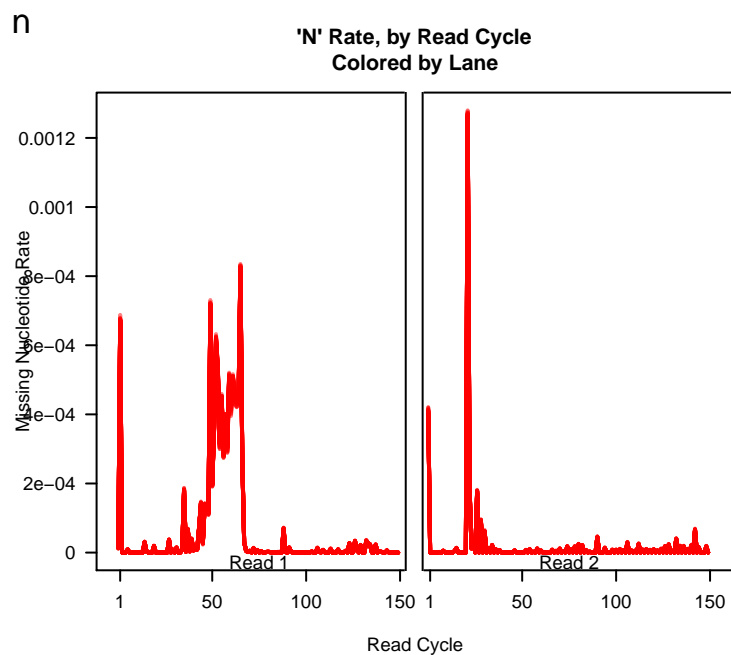
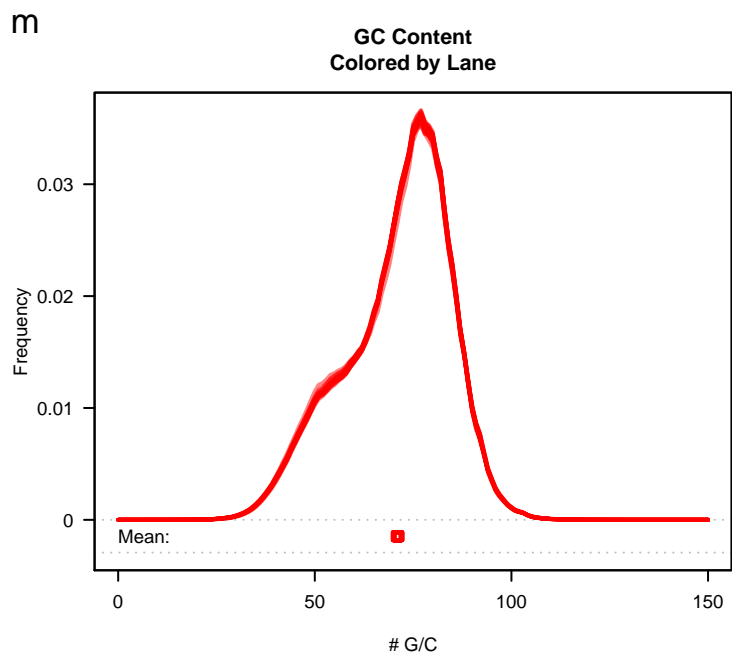
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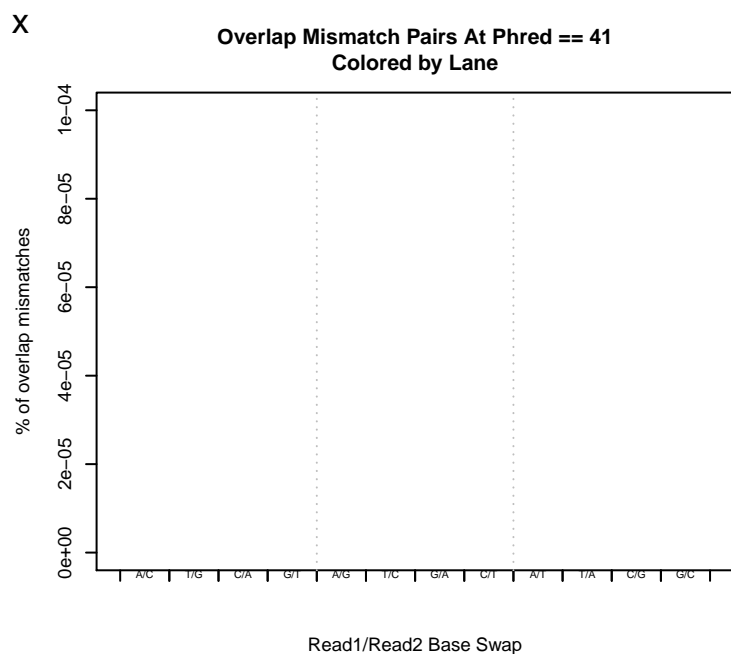
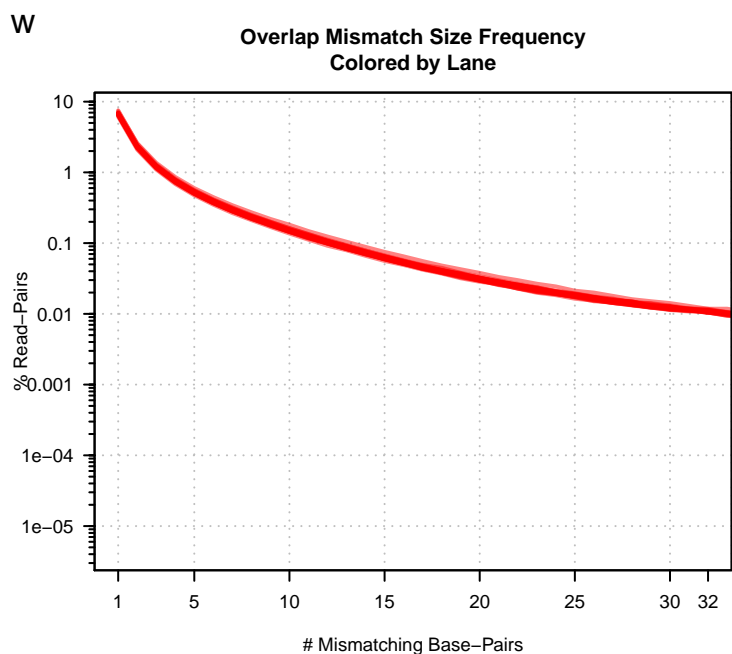
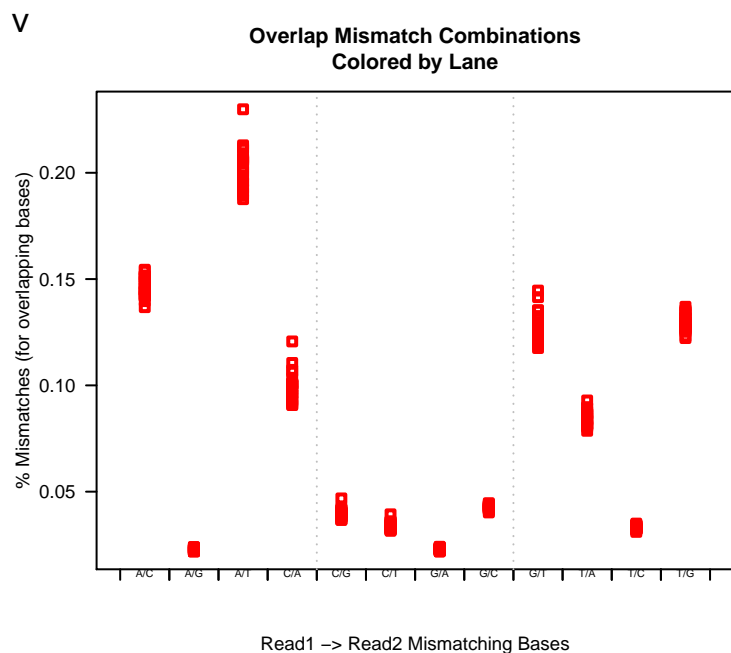
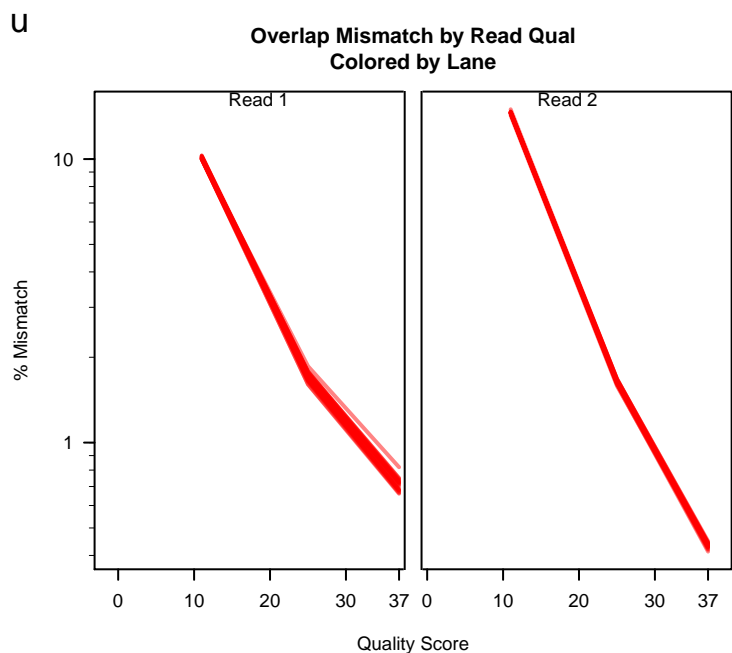
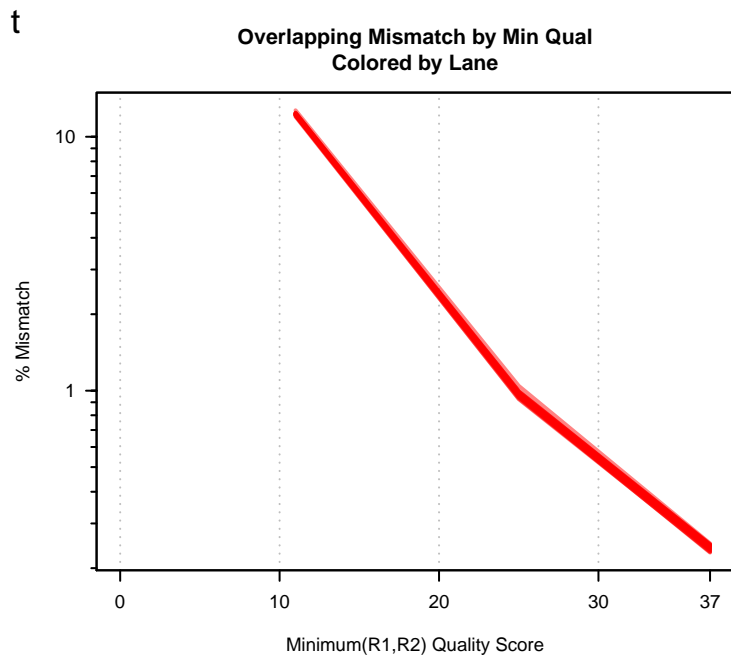
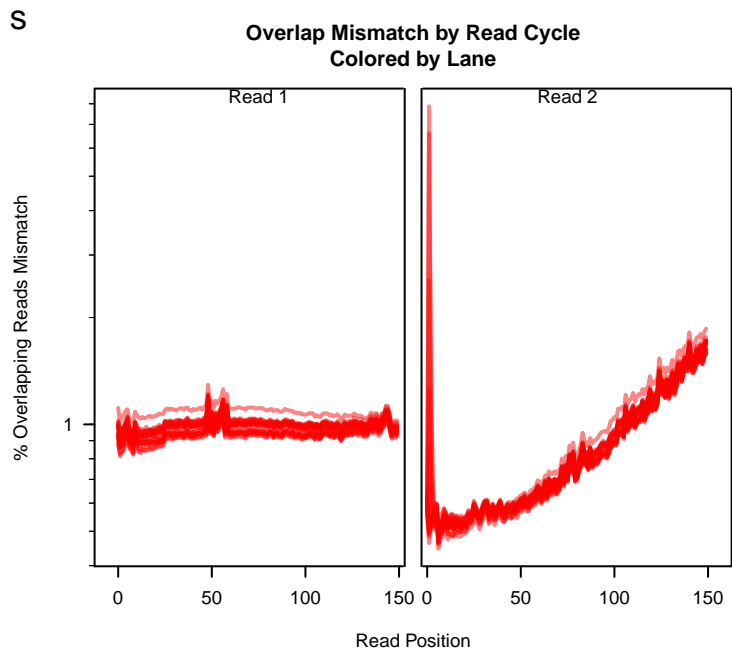


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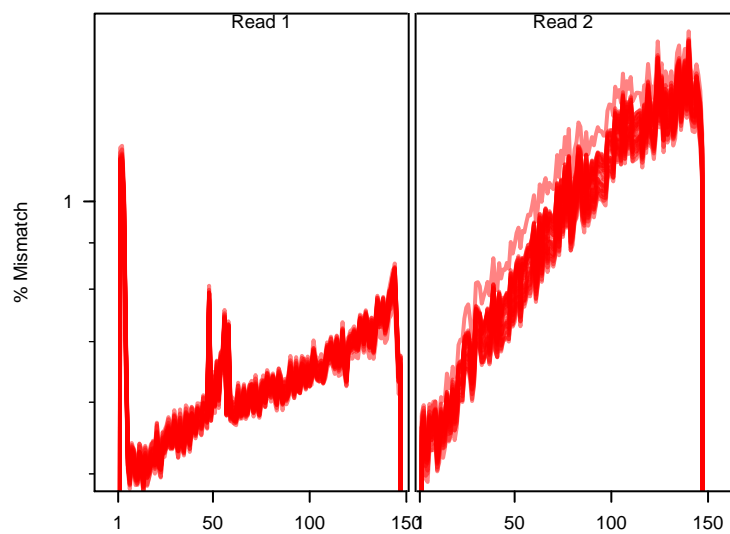






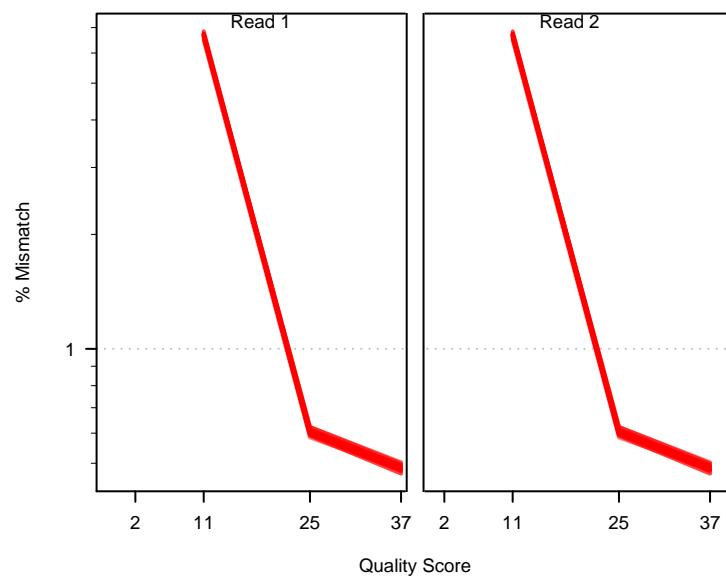
y

**Reference Mismatch by Read Cycle
Colored by Lane**



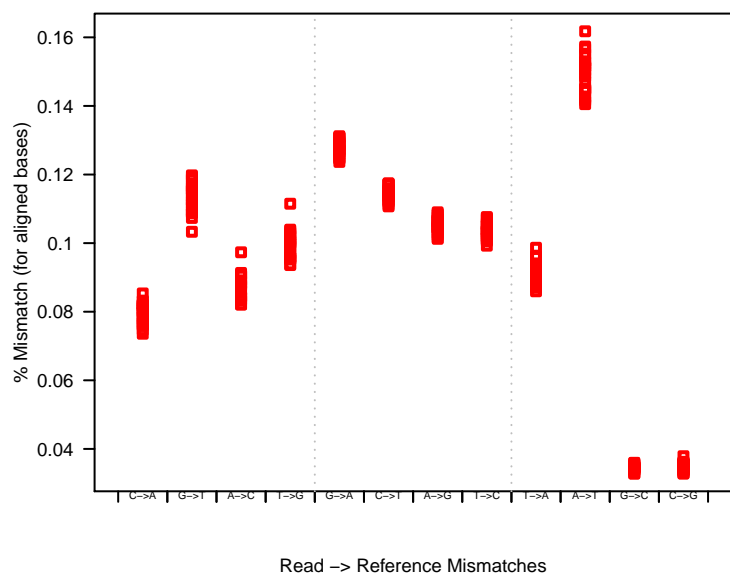
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**Reference Mismatch by Read Qual
Colored by Lane**



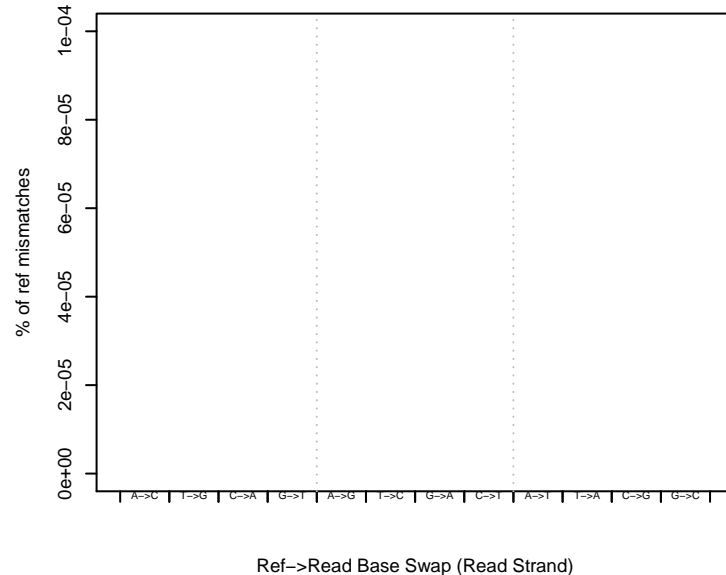
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**Reference Mismatch Combinations
Colored by Lane**



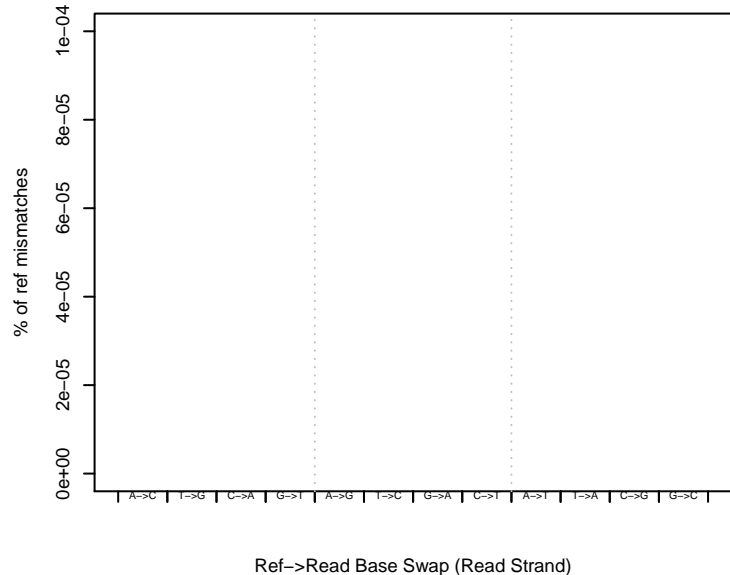
ab

**R1 Ref Mismatches At Phred == 41
Colored by Lane**



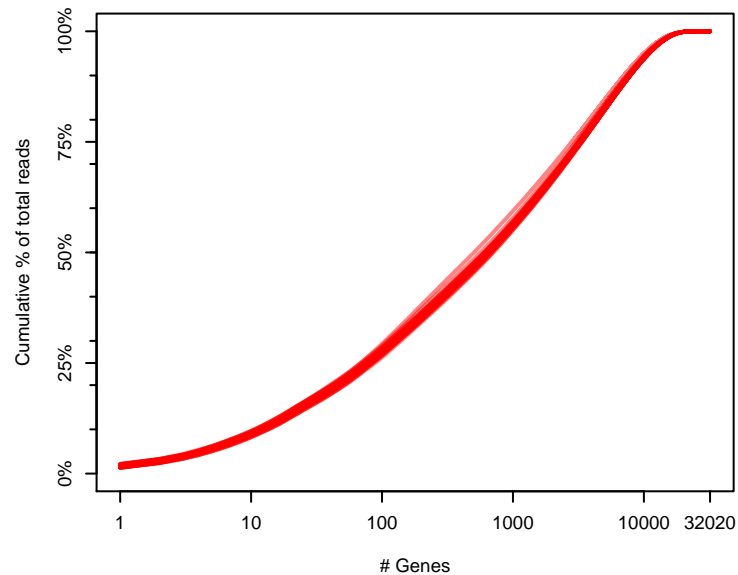
ac

**R2 Ref Mismatches At Phred == 41
Colored by Lane**

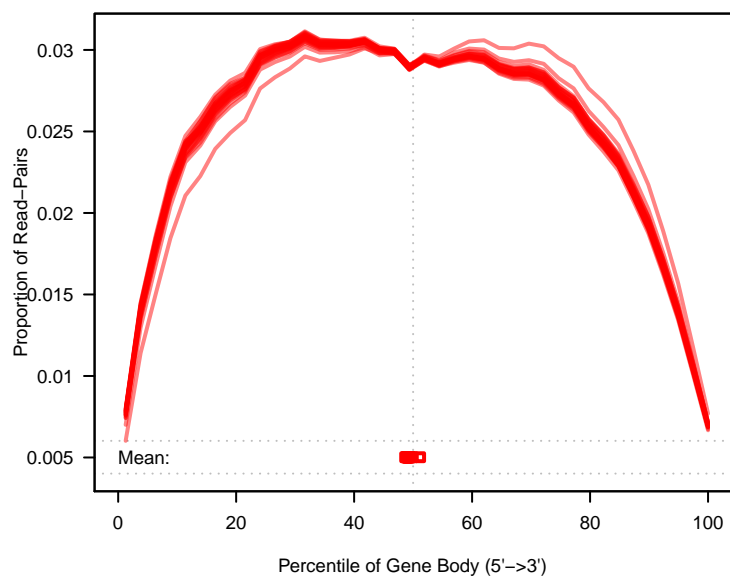


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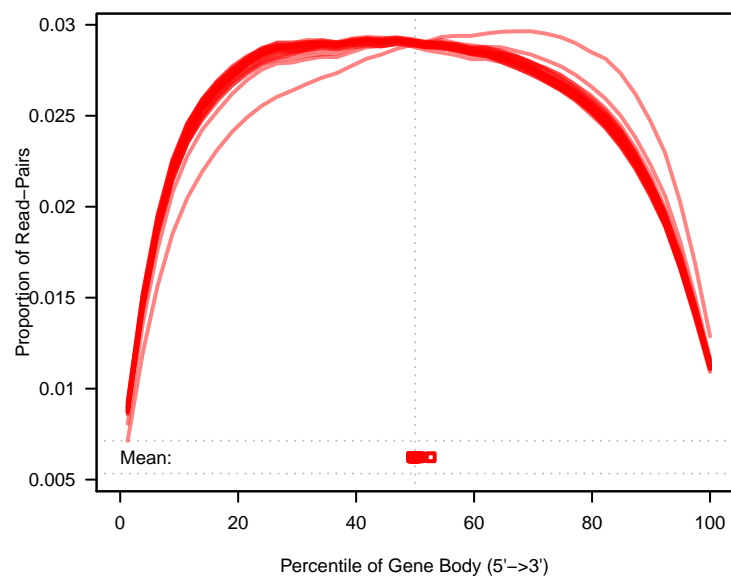
**Cumulative Gene Assignment Diversity
Colored by Lane**



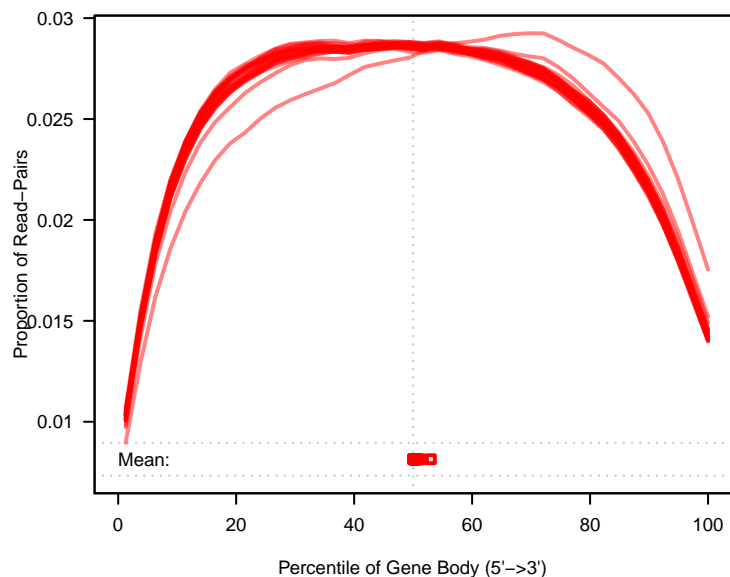
ae

Gene-Body Coverage
Colored by Lane

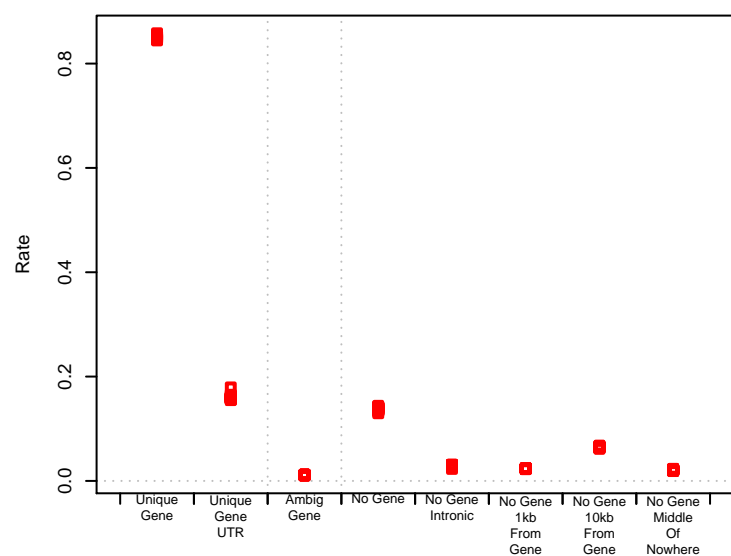
af

Gene-Body Coverage, Upper Middle Quartile Genes
Colored by Lane

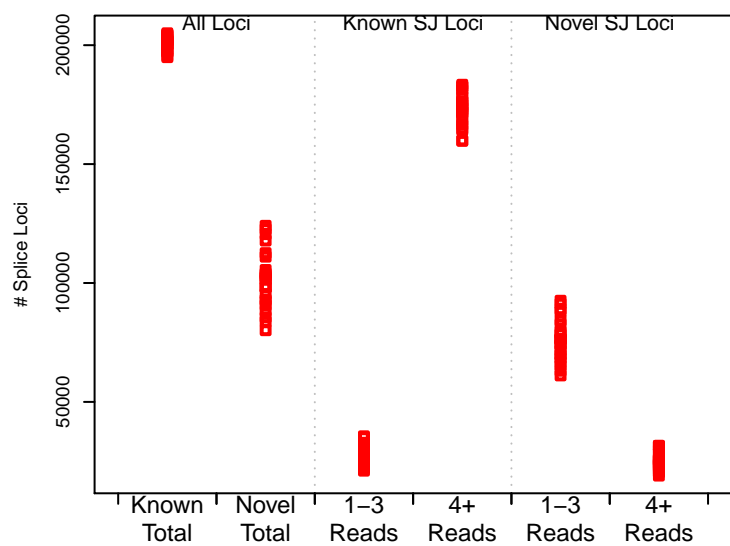
ag

Gene-Body Coverage, Low Expression Genes
Colored by Lane

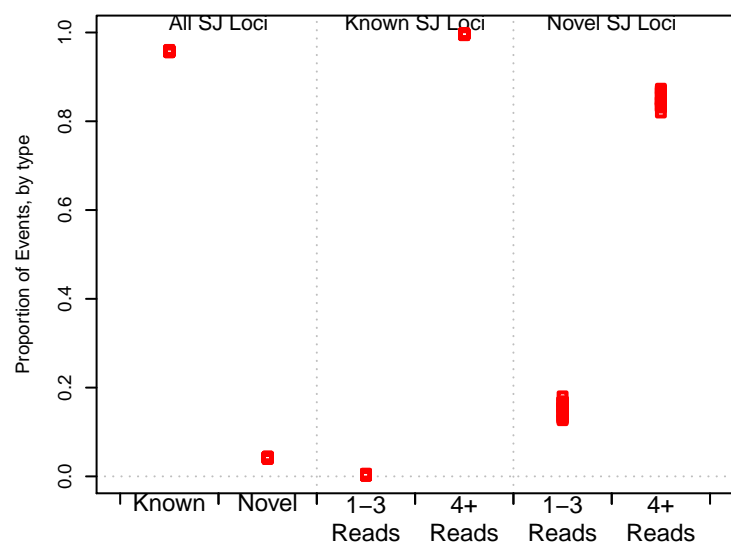
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Read Mapping Location Rates
Colored by Lane

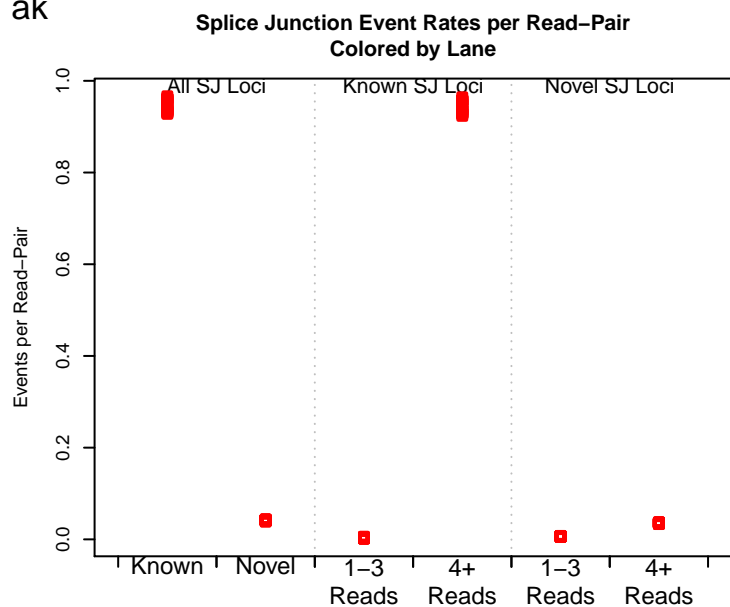
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Observed Splice Junction Loci, by type
Colored by Lane

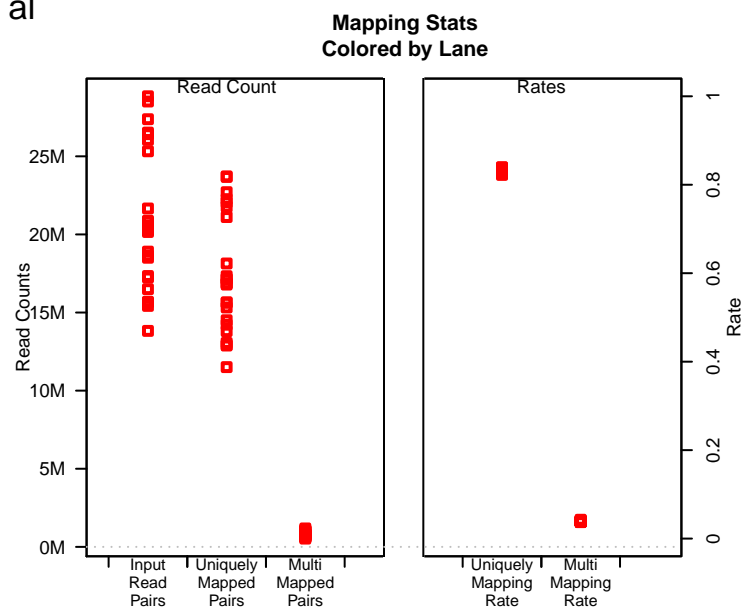
aj

Breakdown of Splice Junction Events, by type
Colored by Lane

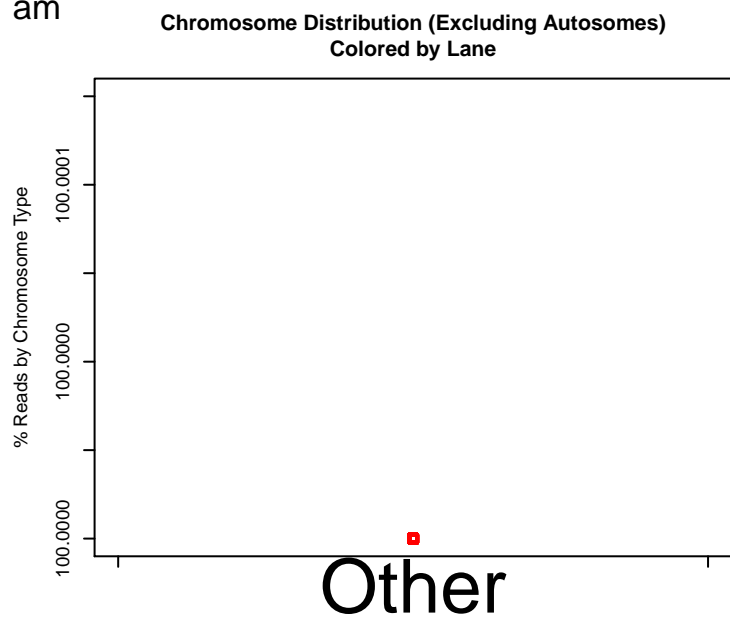
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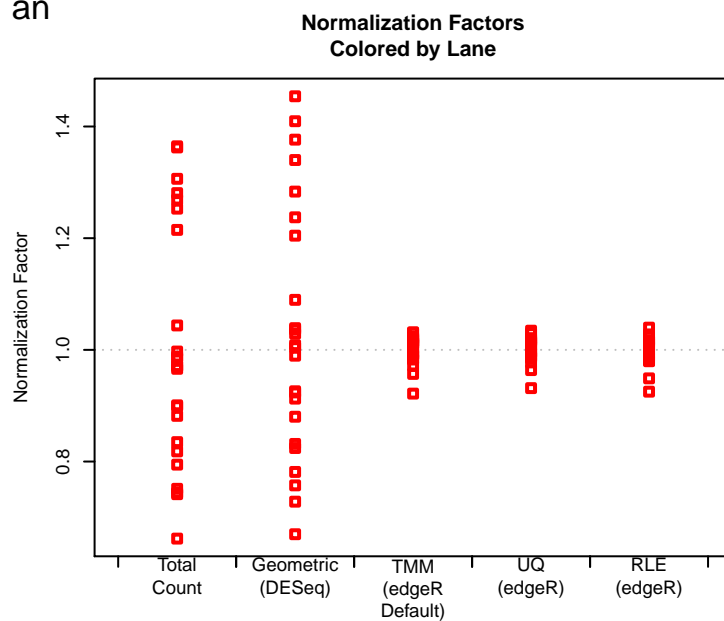
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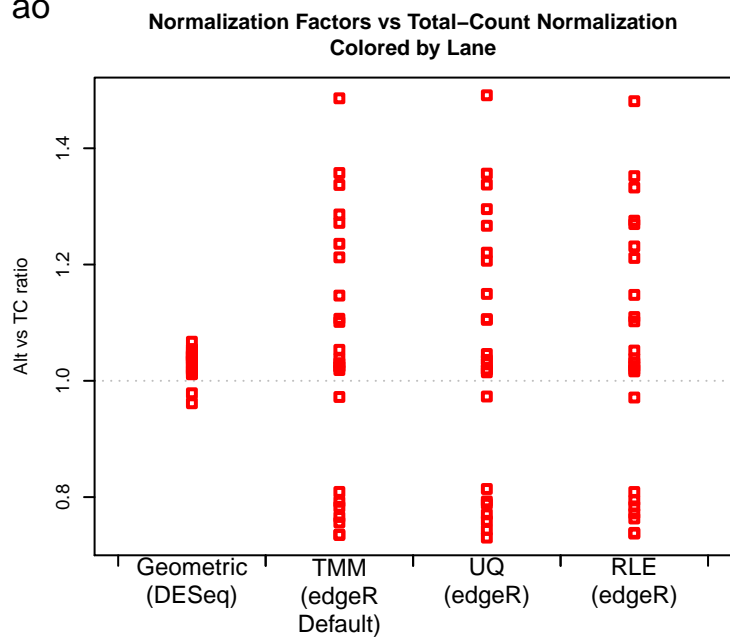
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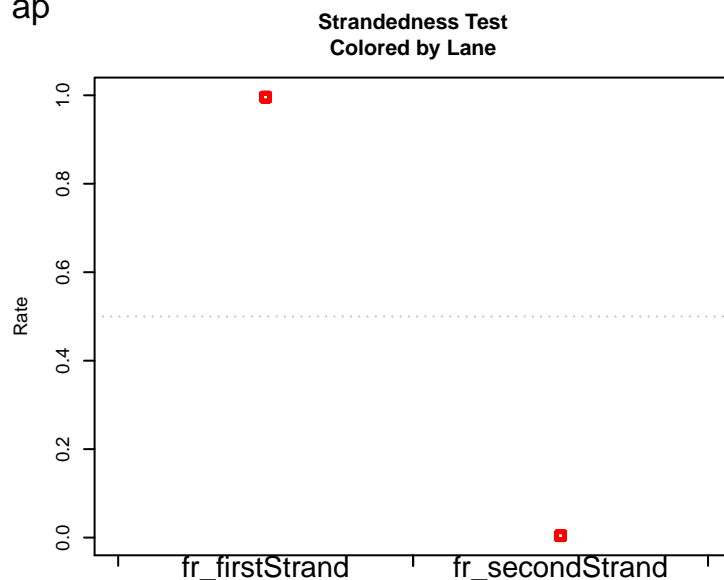
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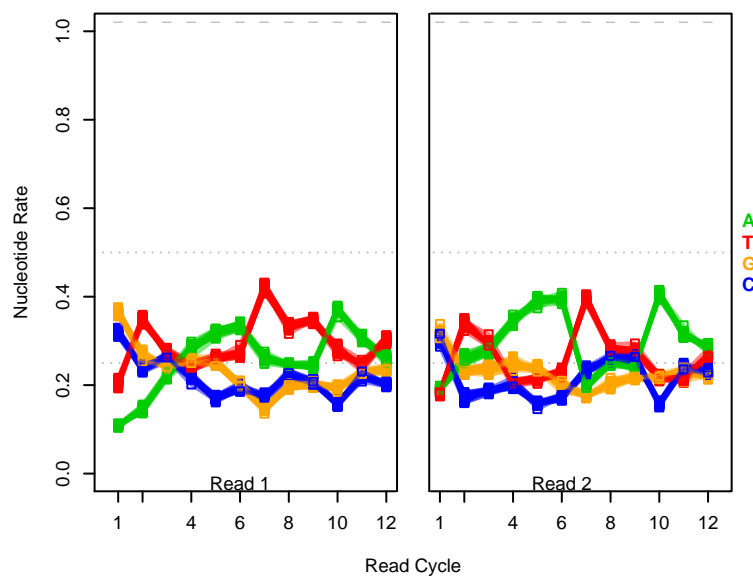


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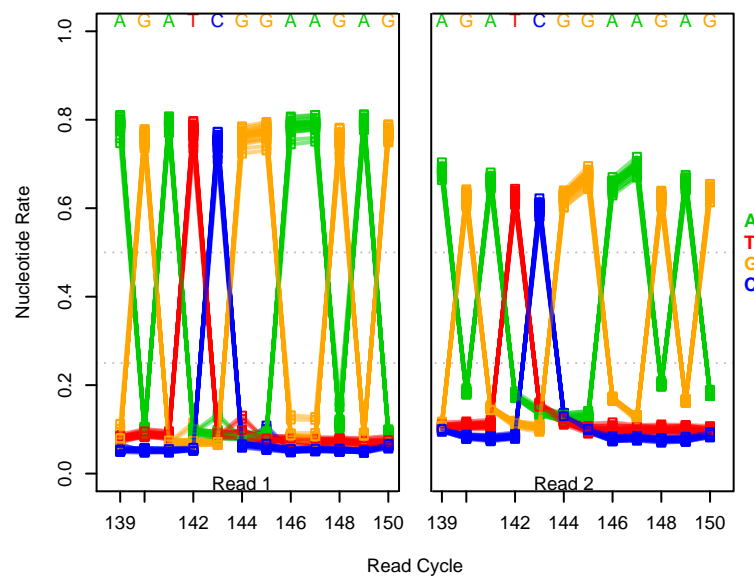
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Nucleotide Rate by Cycle, Leading Clipped bases (12)
Marked by Lane



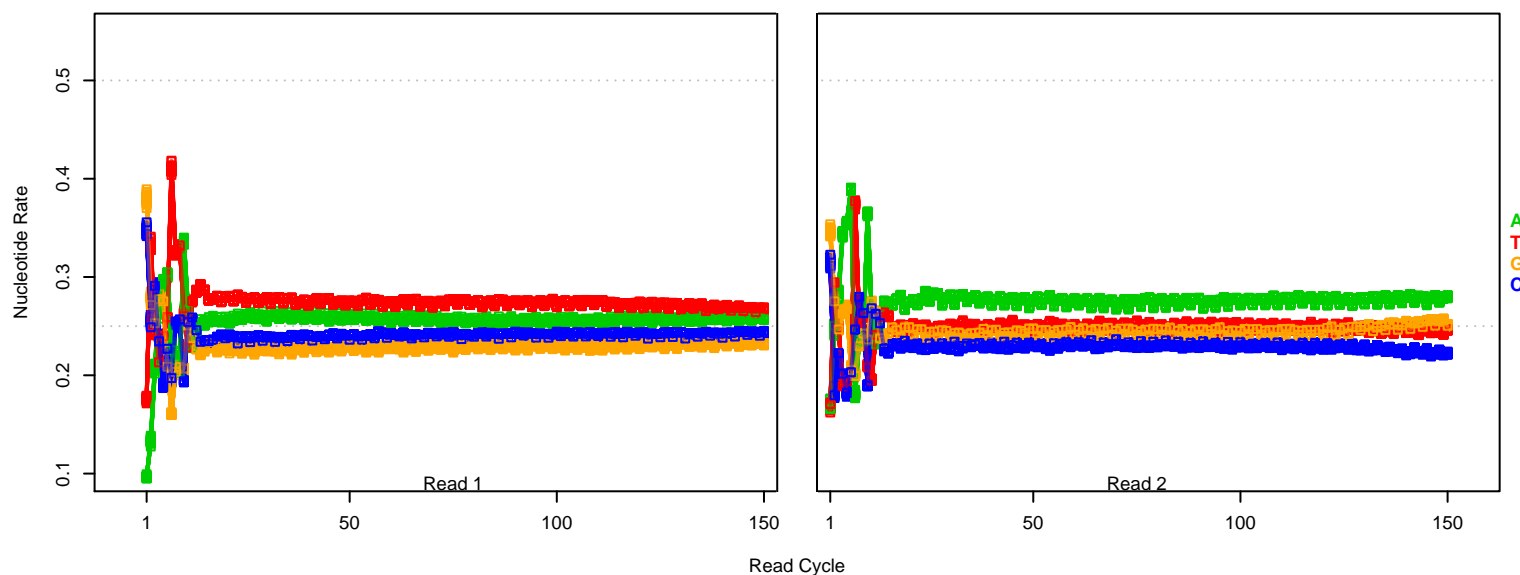
ar

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



as

Raw Nucleotide Rate by Cycle
Marked by Lane



at

Nucleotide Rate by Cycle, Aligned bases only
Marked by Lane

