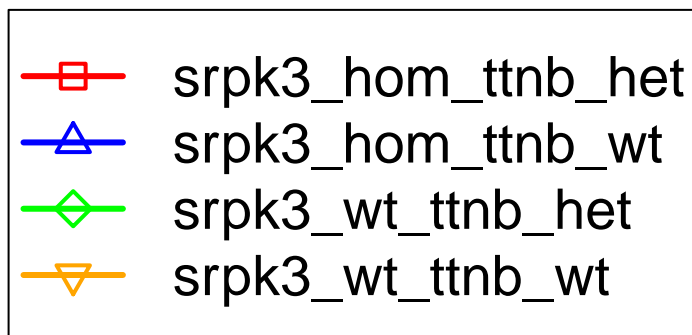
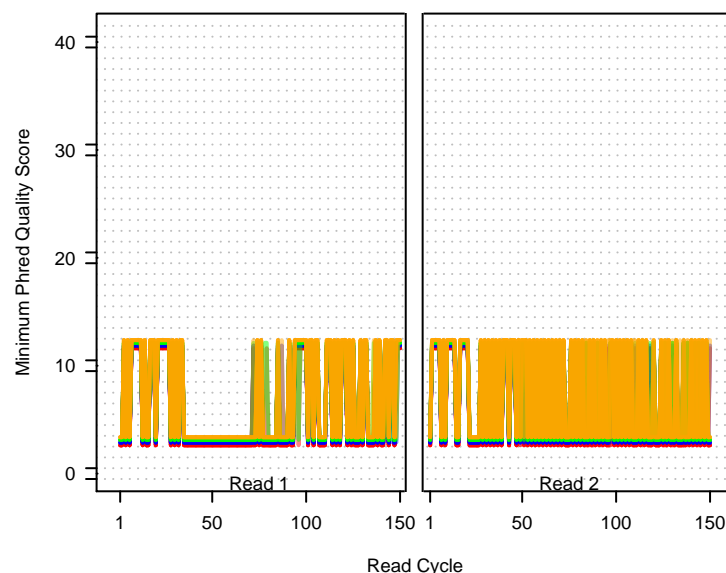


a

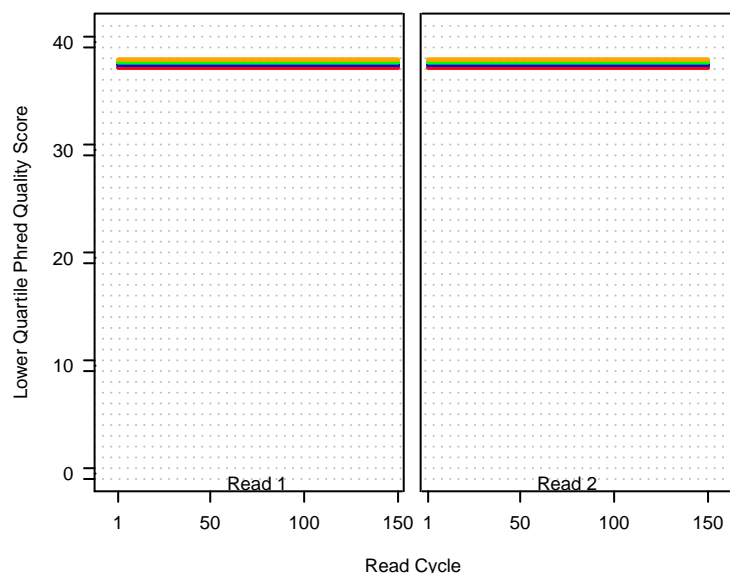
## Summary Plots, By Group



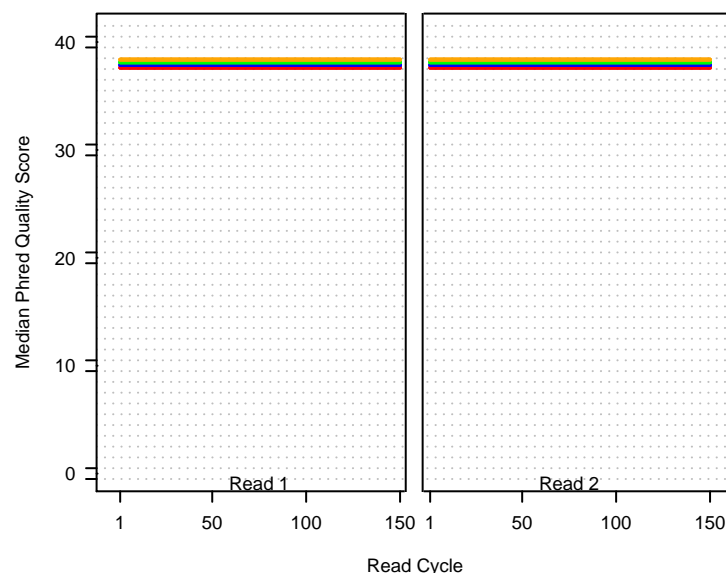
b

Minimum Phred Quality Score  
Colored by Group

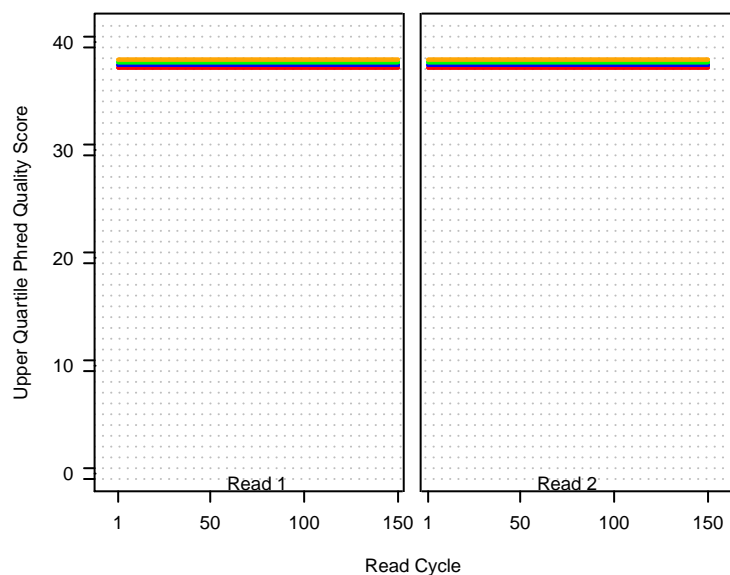
c

Lower Quartile Phred Quality Score  
Colored by Group

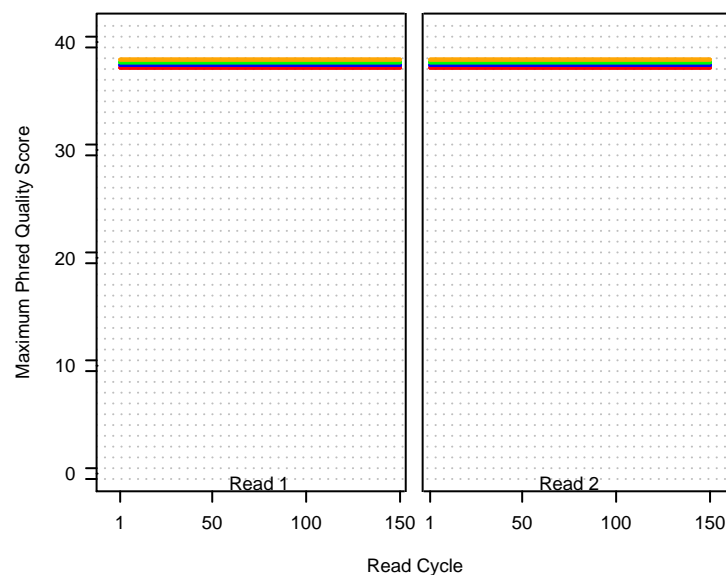
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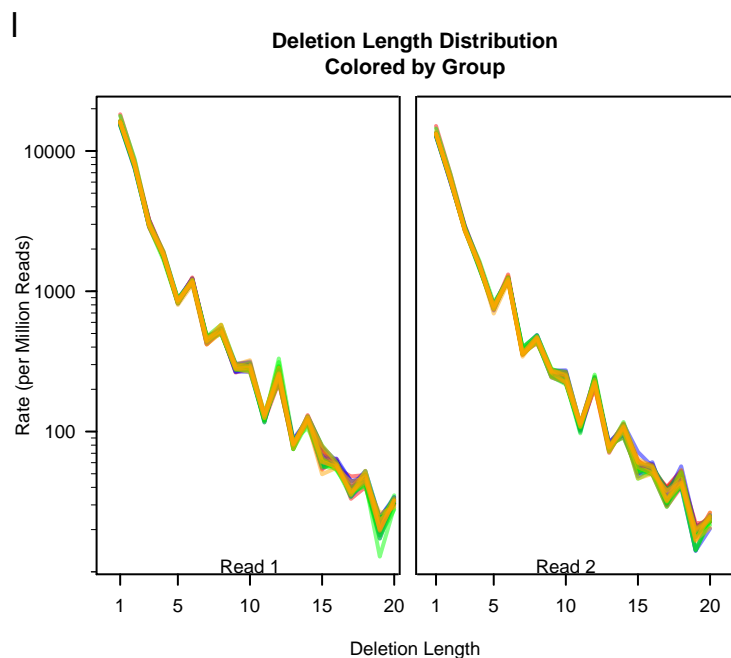
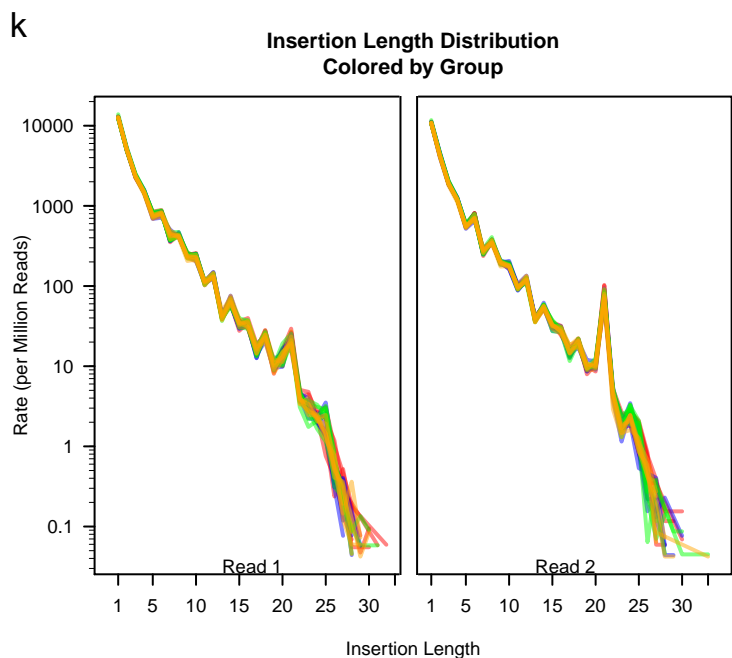
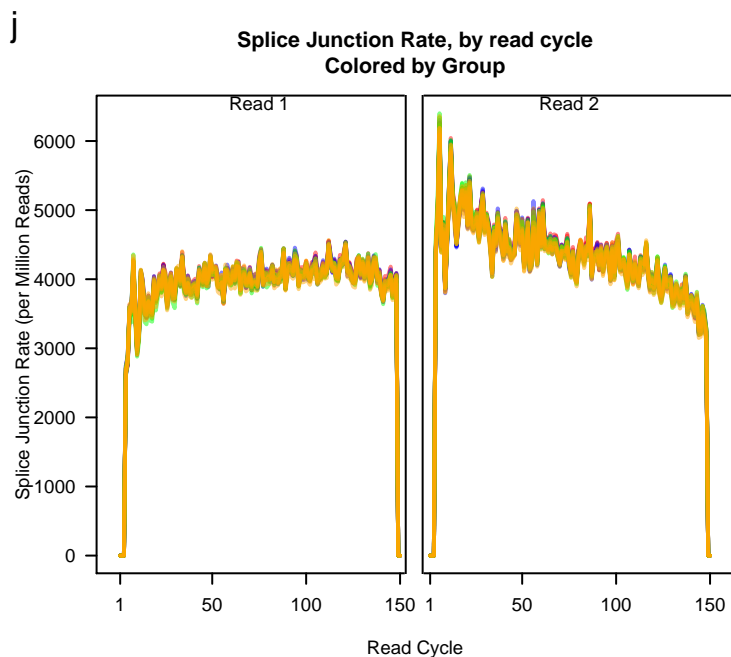
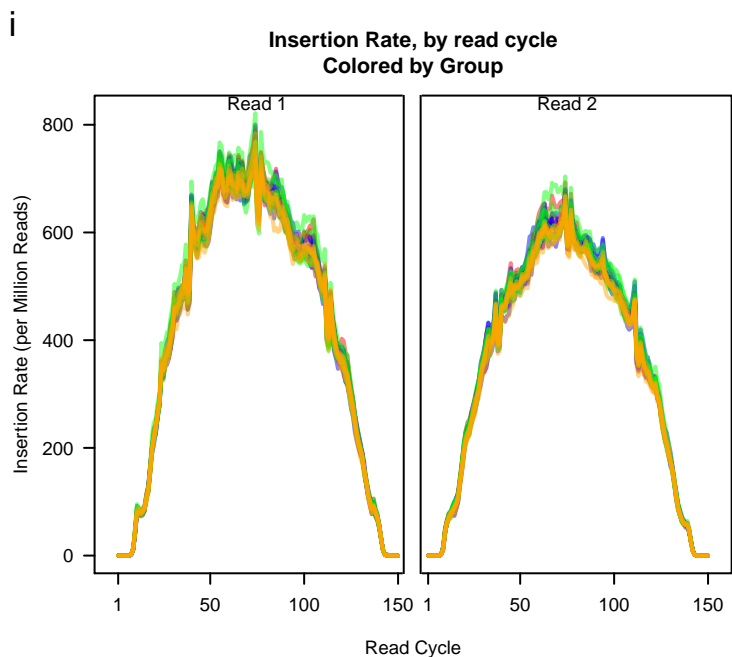
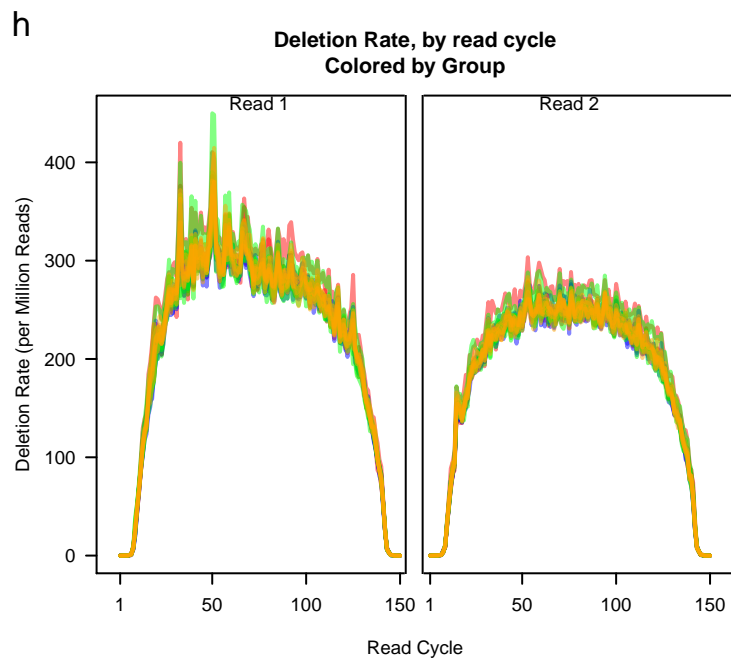
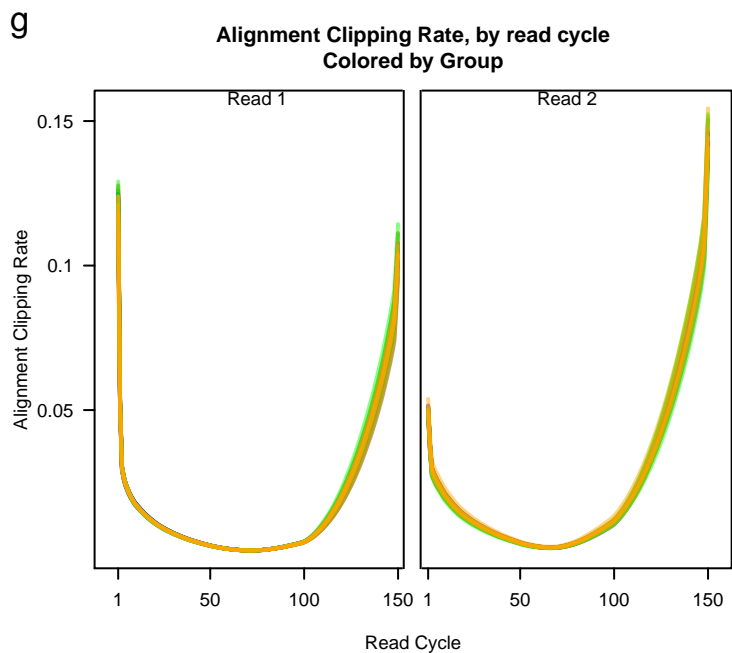
Median Phred Quality Score  
Colored by Group

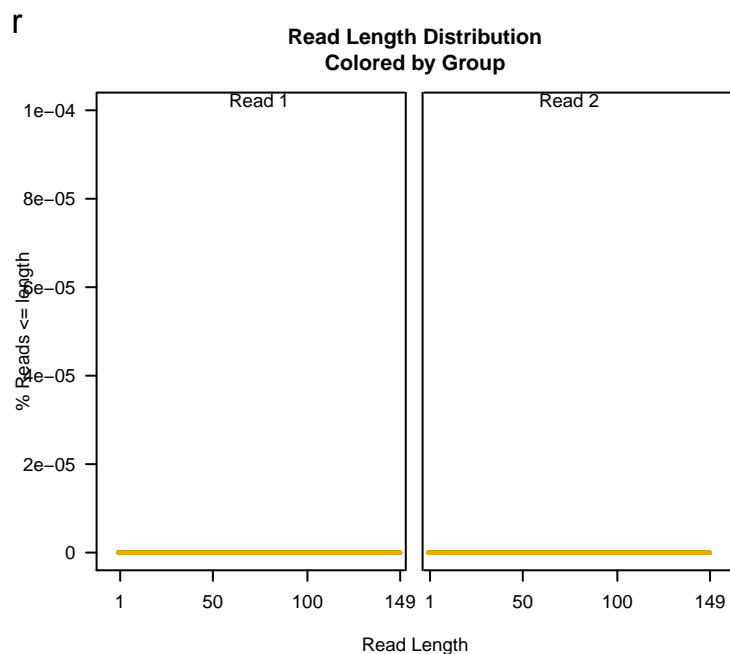
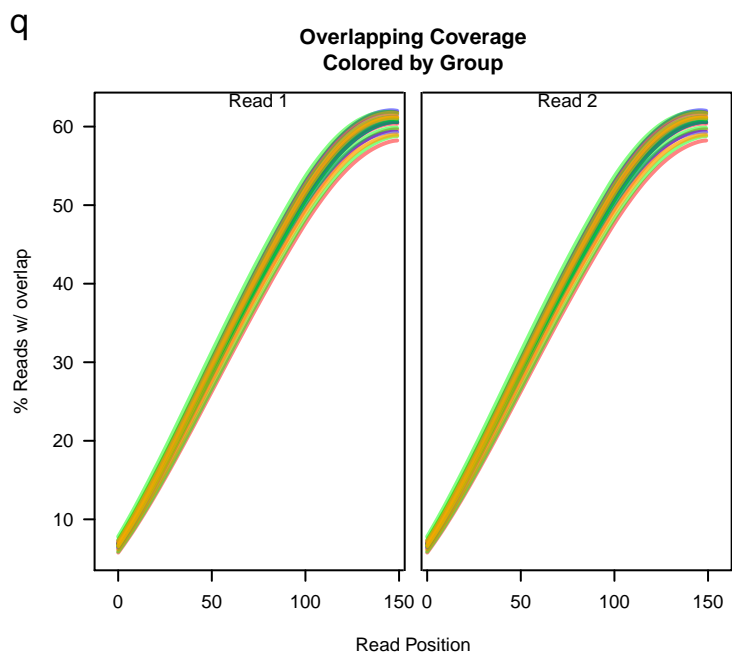
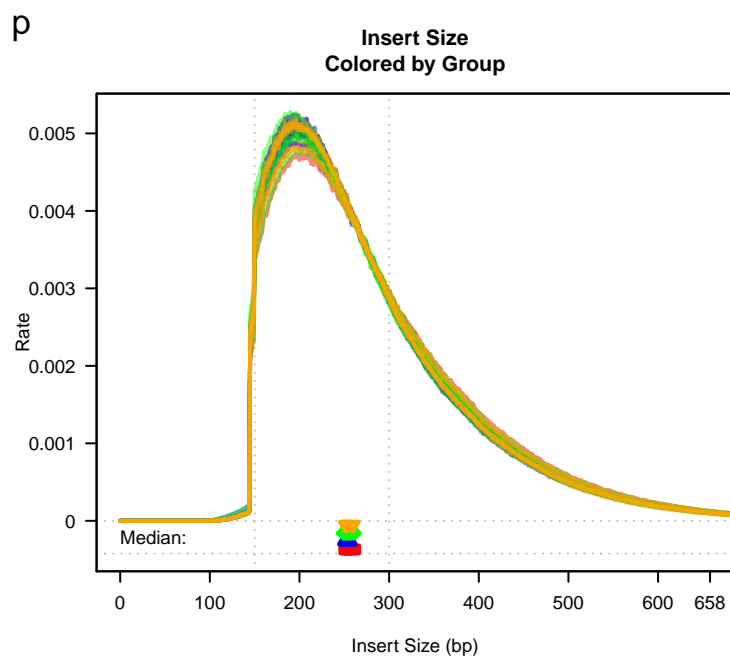
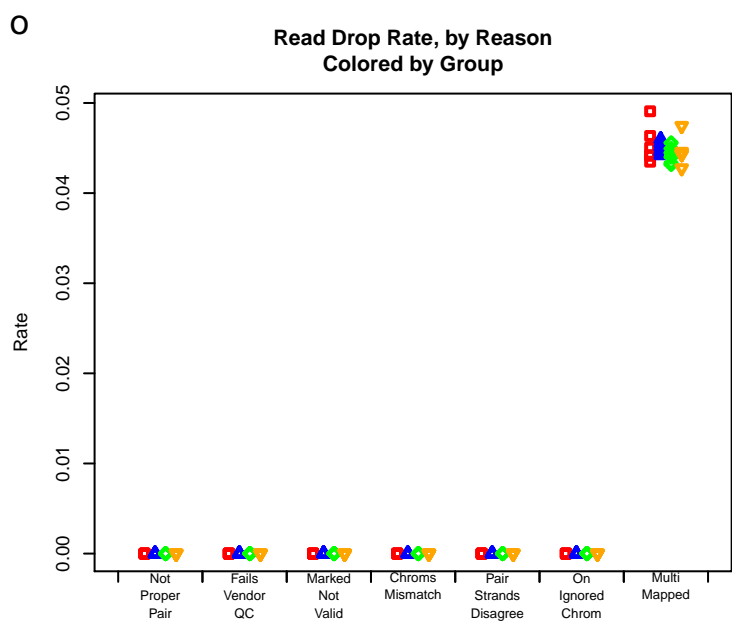
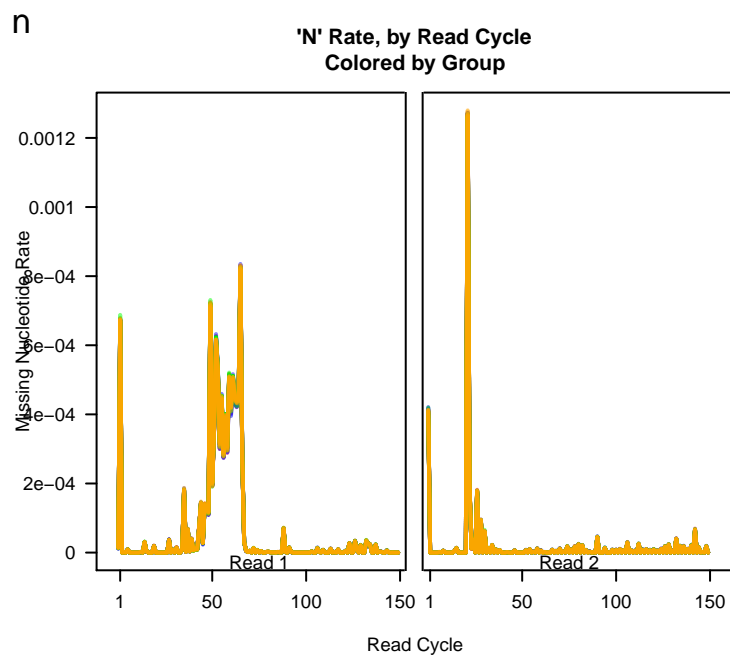
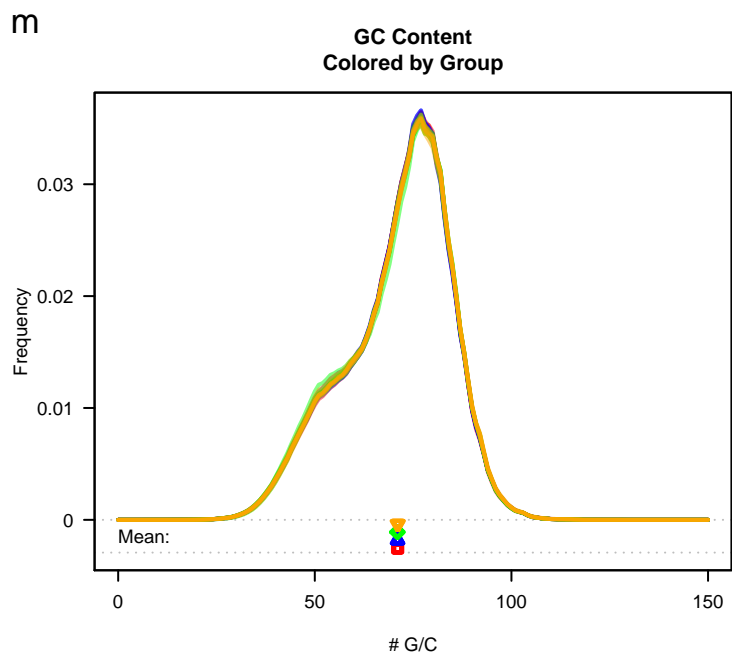
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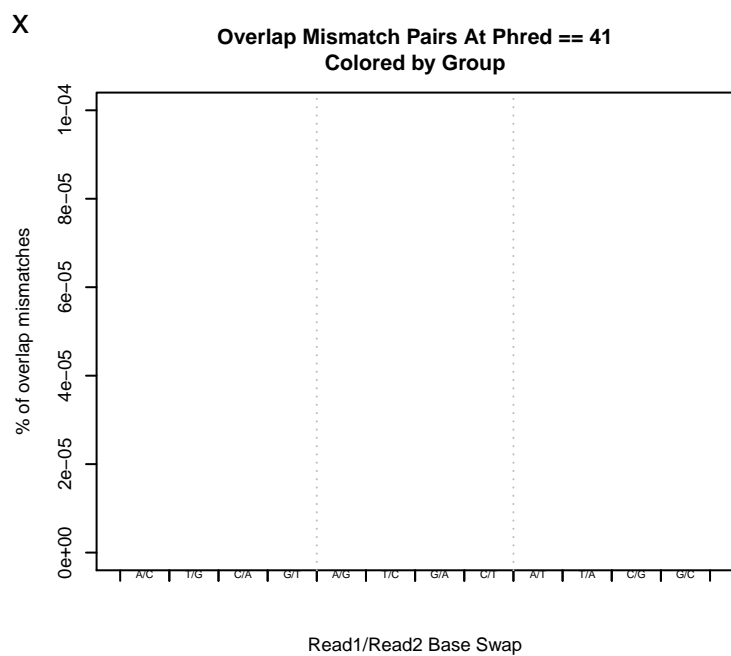
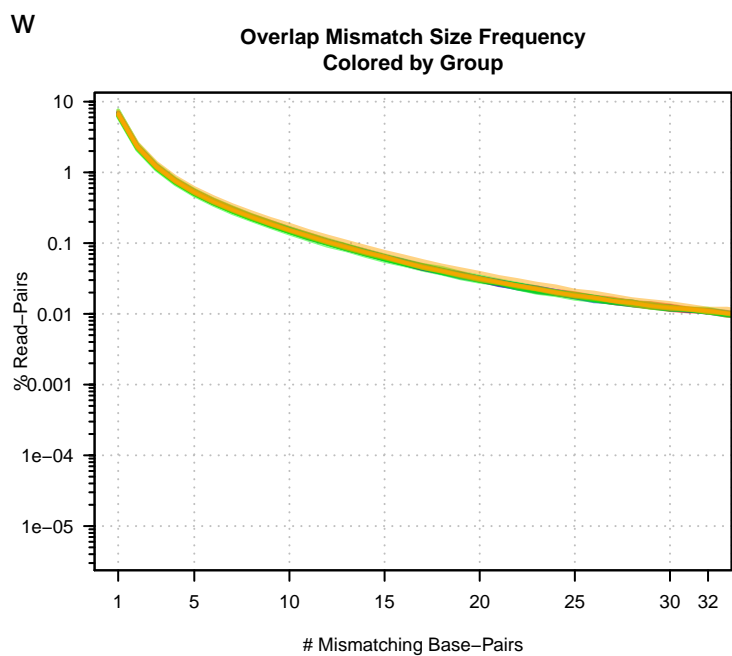
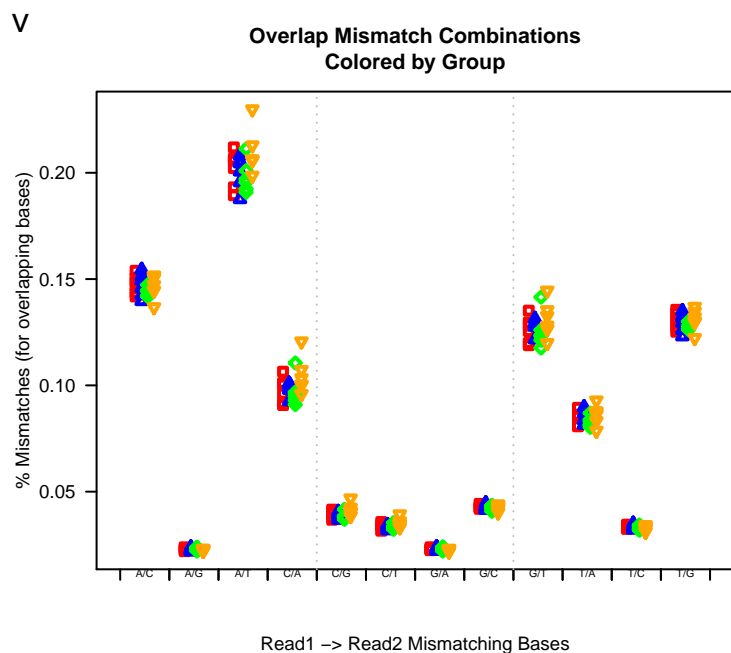
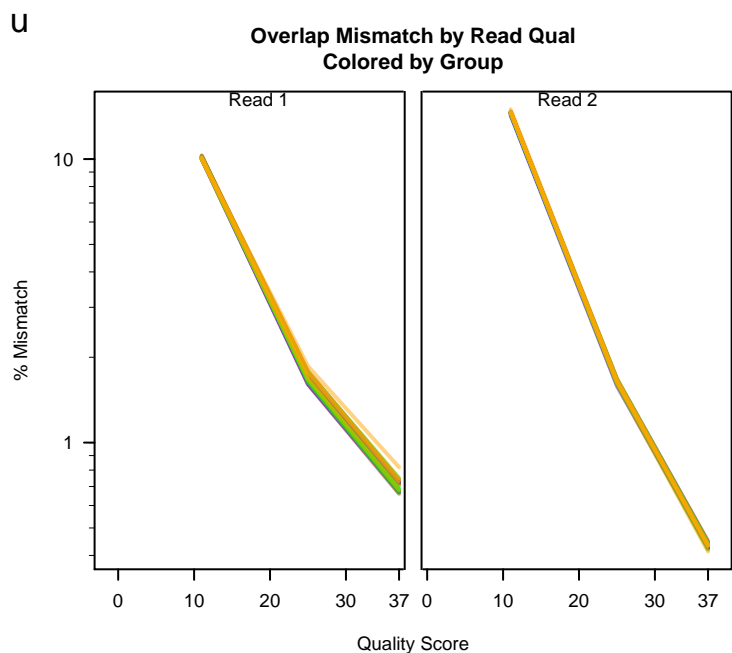
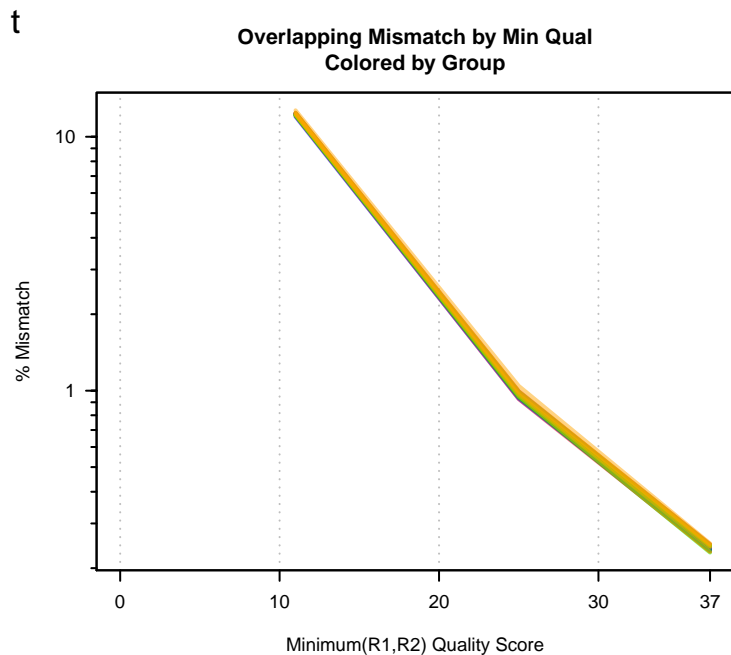
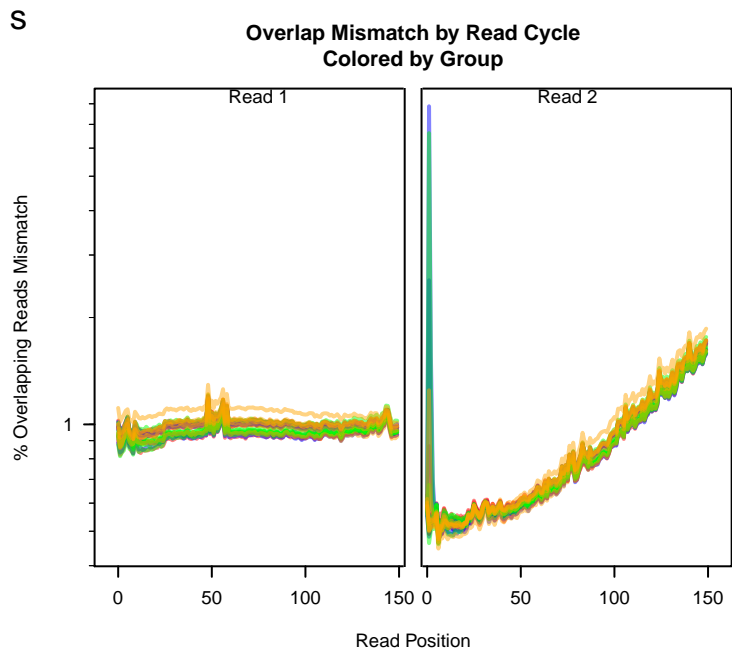
Upper Quartile Phred Quality Score  
Colored by Group

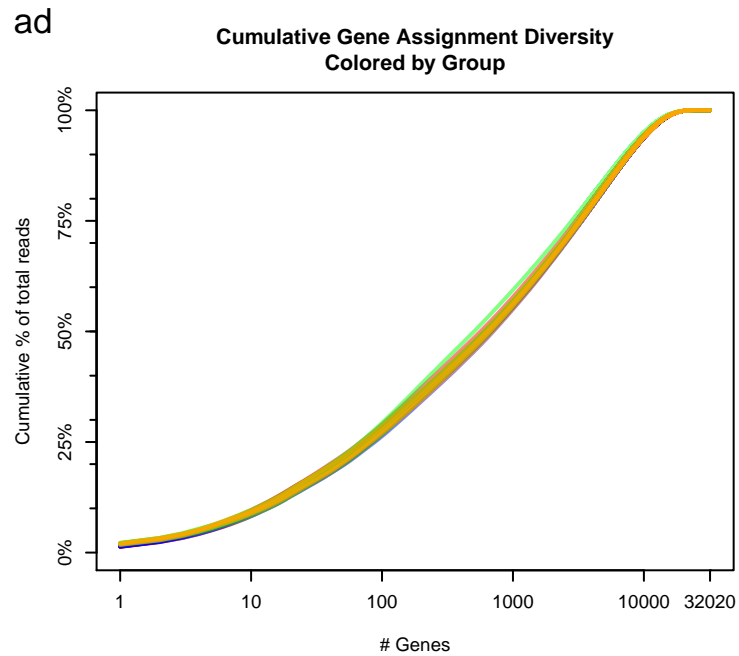
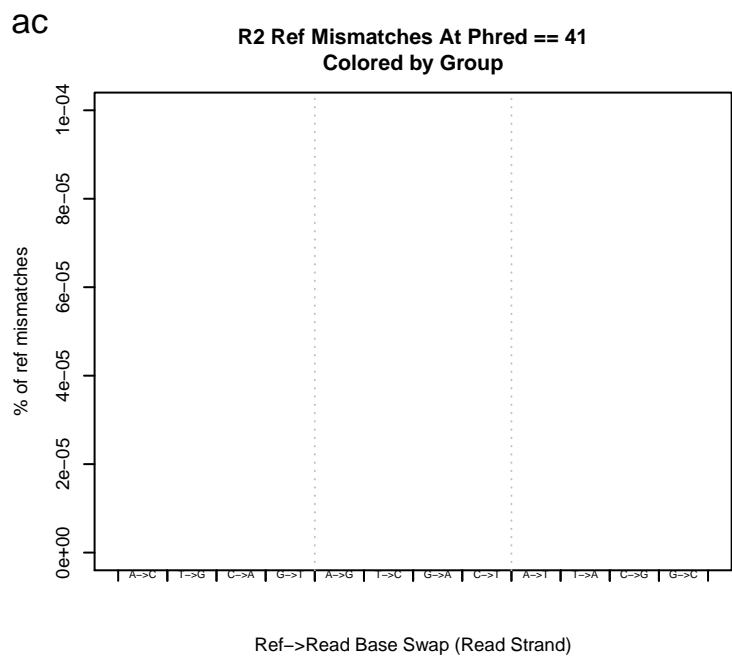
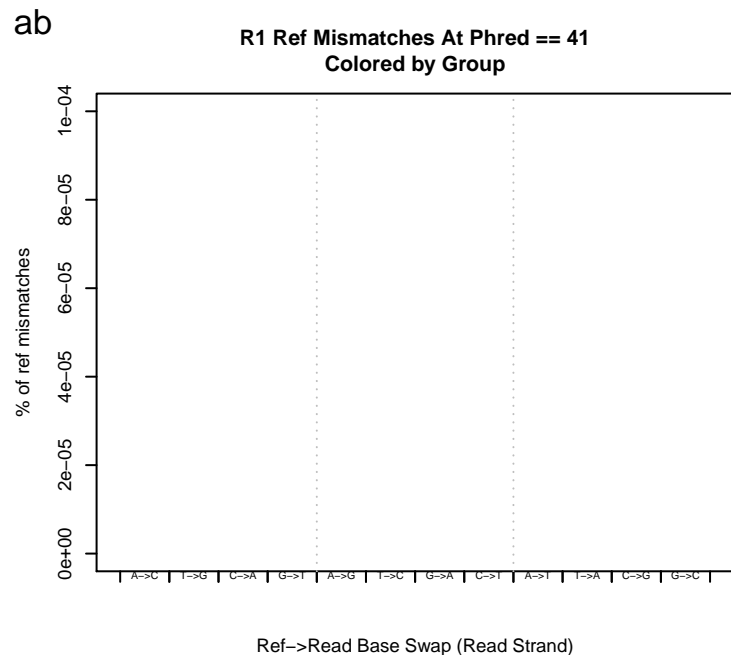
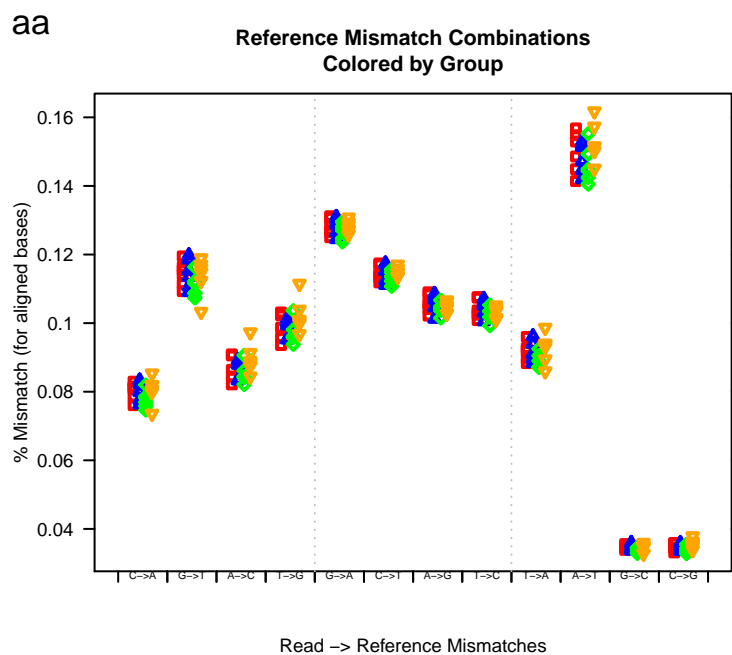
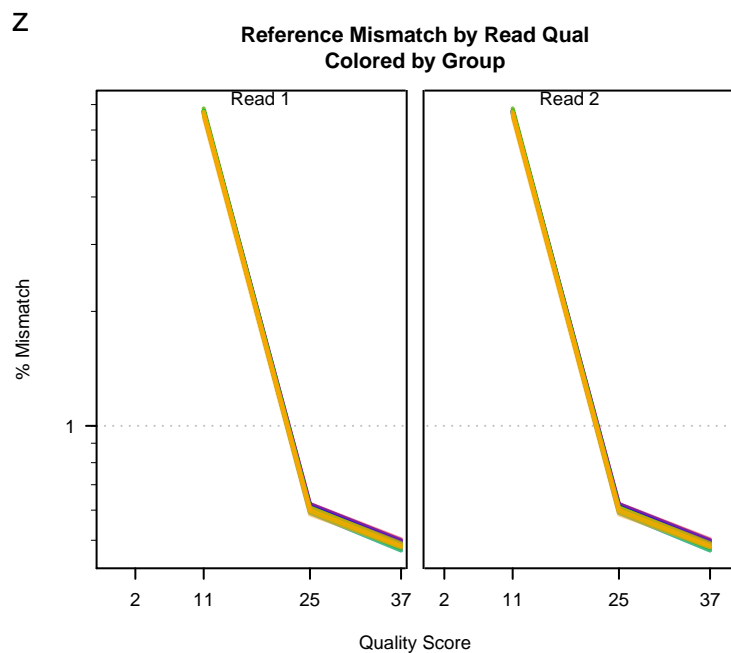
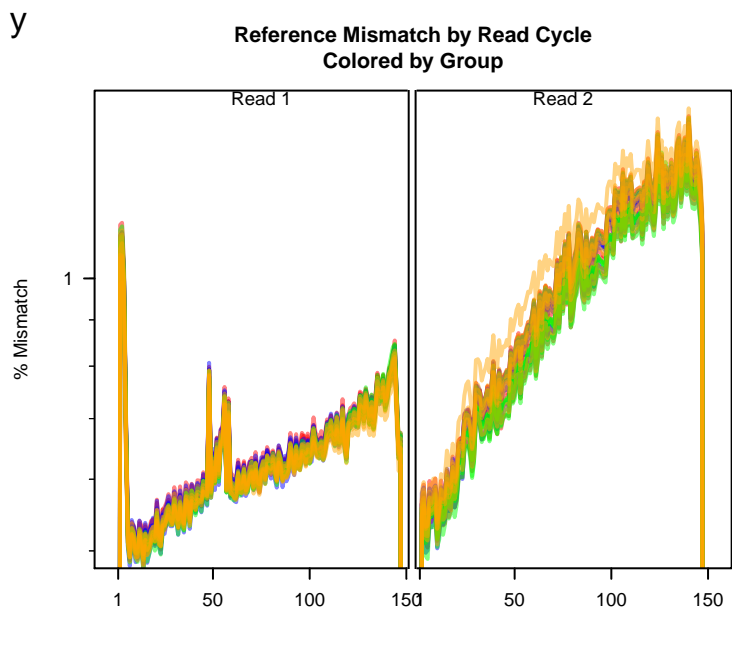
f

Maximum Phred Quality Score  
Colored by Group

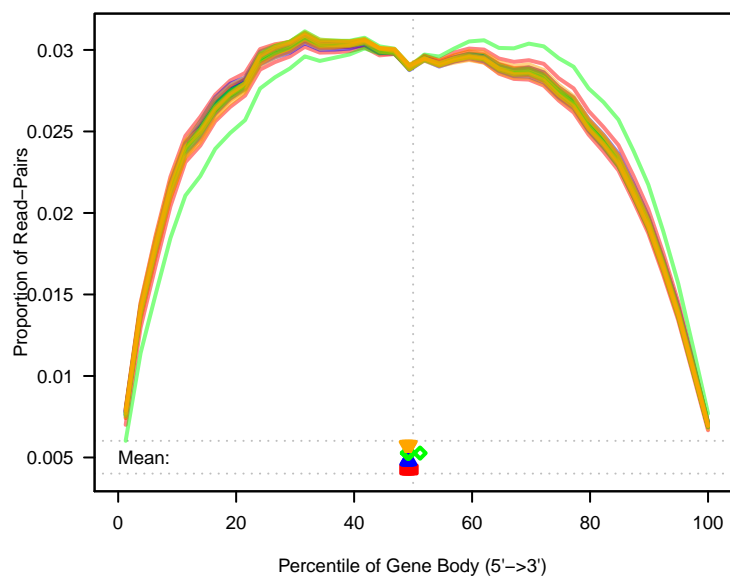




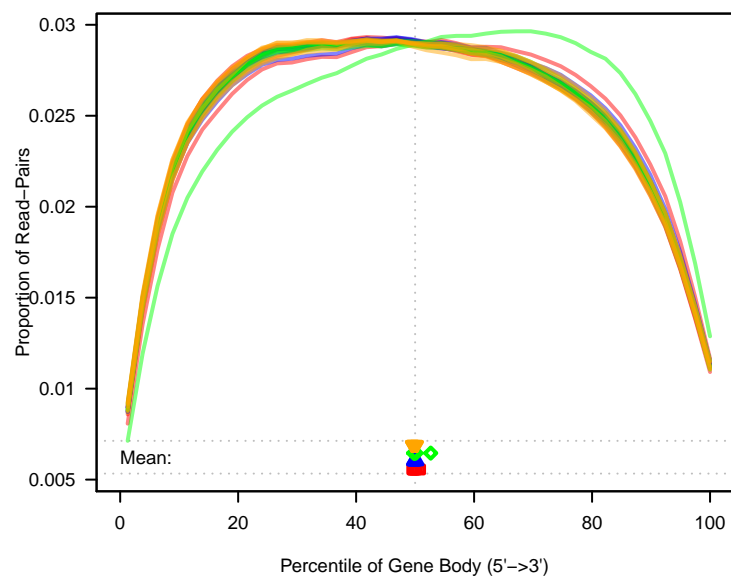




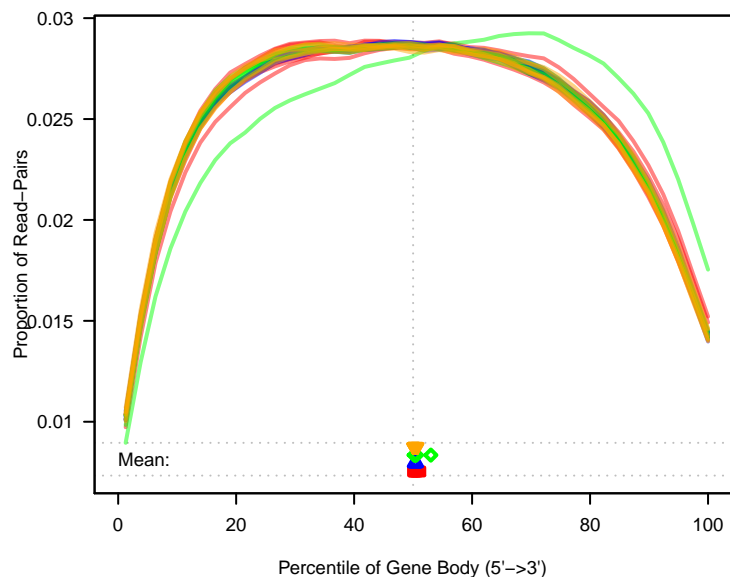
ae

Gene-Body Coverage  
Colored by Group

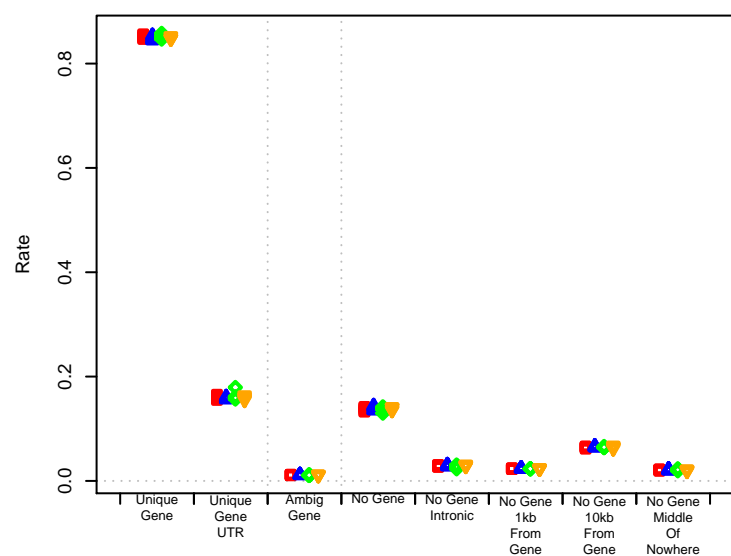
af

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

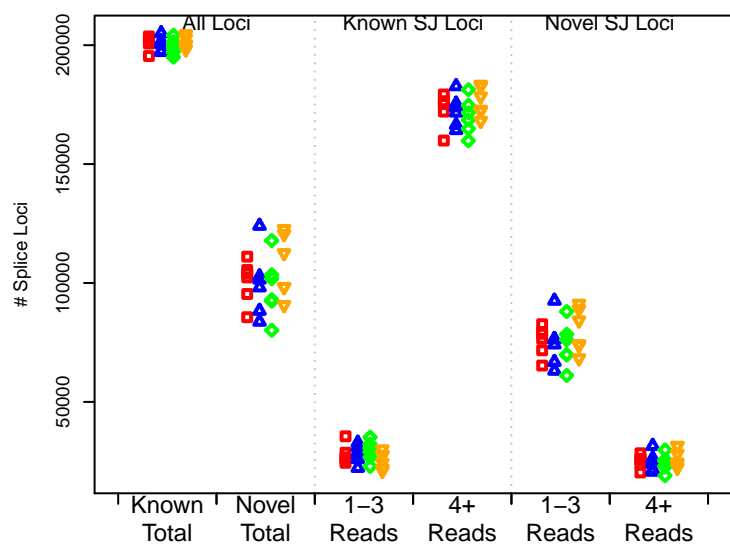
ag

Gene-Body Coverage, Low Expression Genes  
Colored by Group

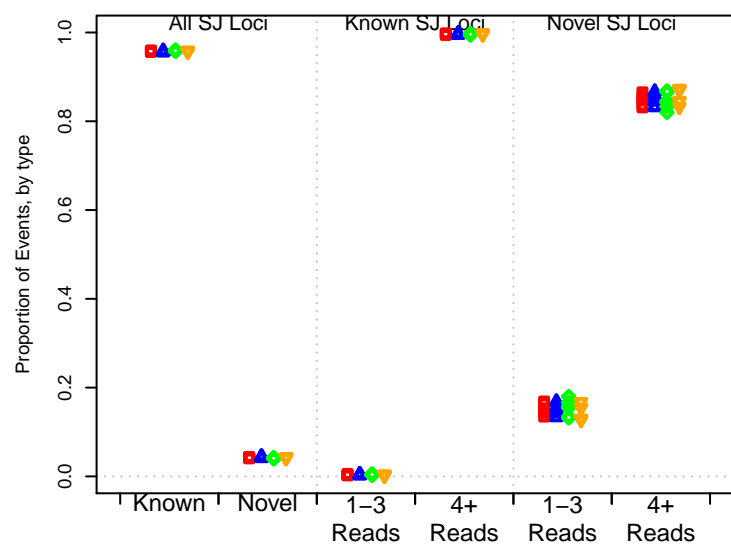
ah

Read Mapping Location Rates  
Colored by Group

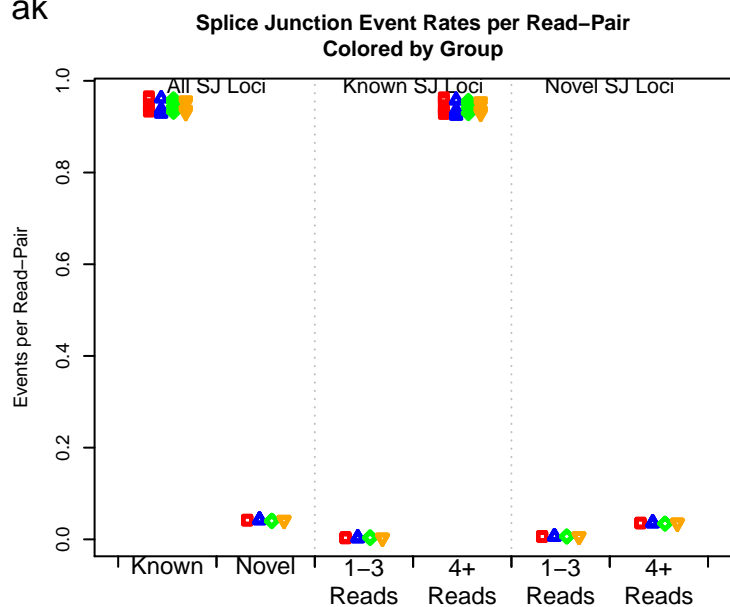
ai

# Observed Splice Junction Loci, by type  
Colored by Group

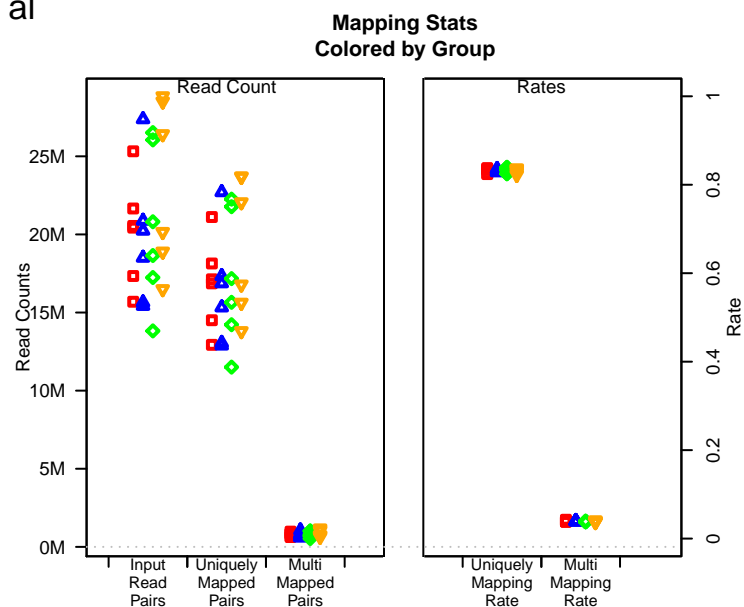
aj

Breakdown of Splice Junction Events, by type  
Colored by Group

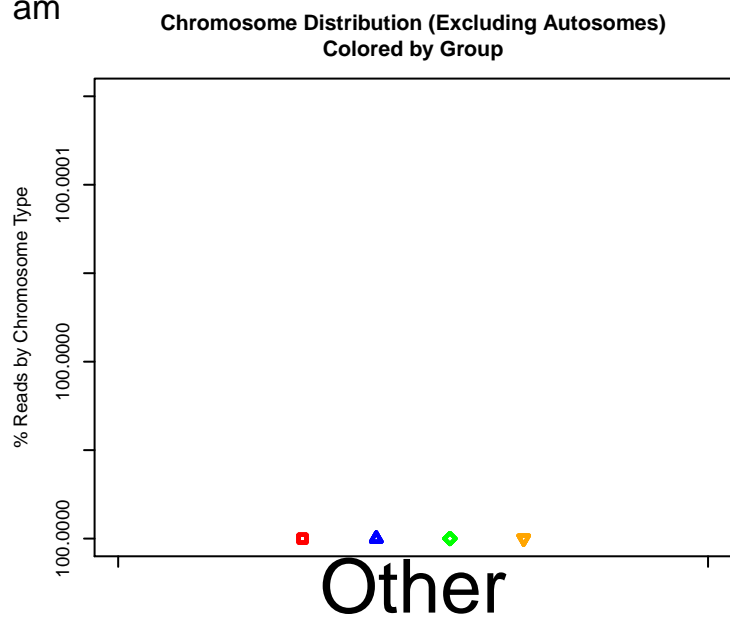
ak



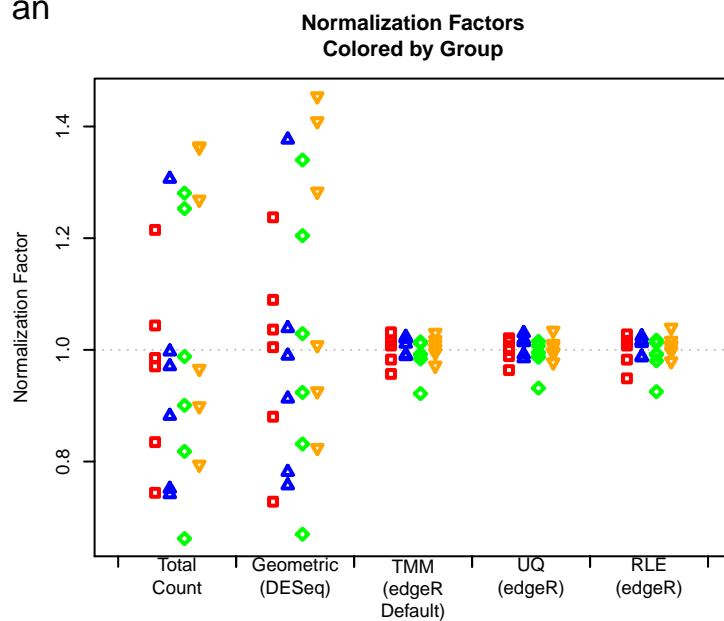
al



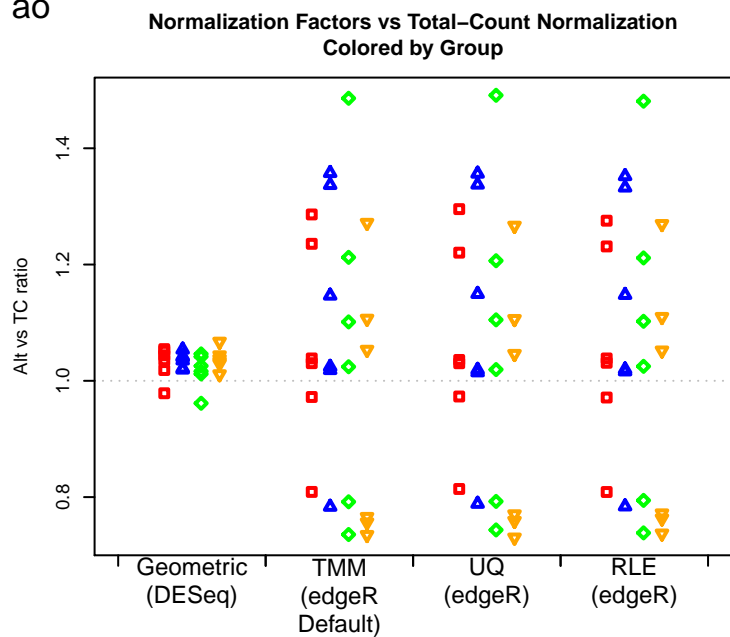
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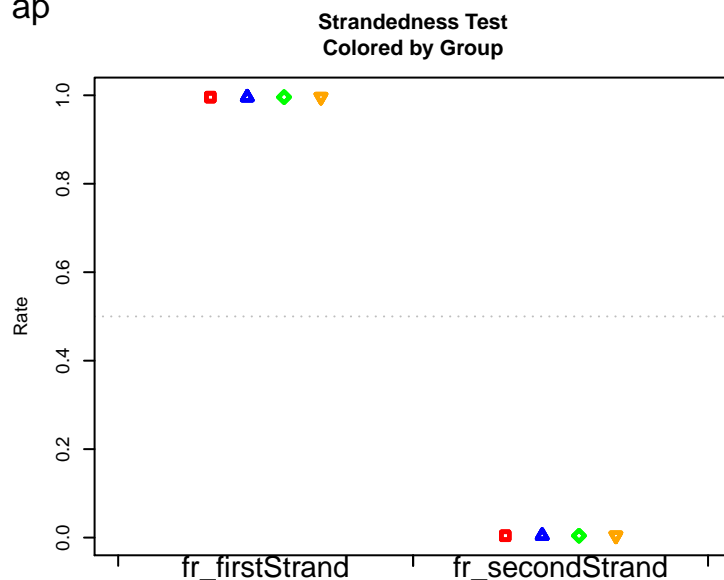
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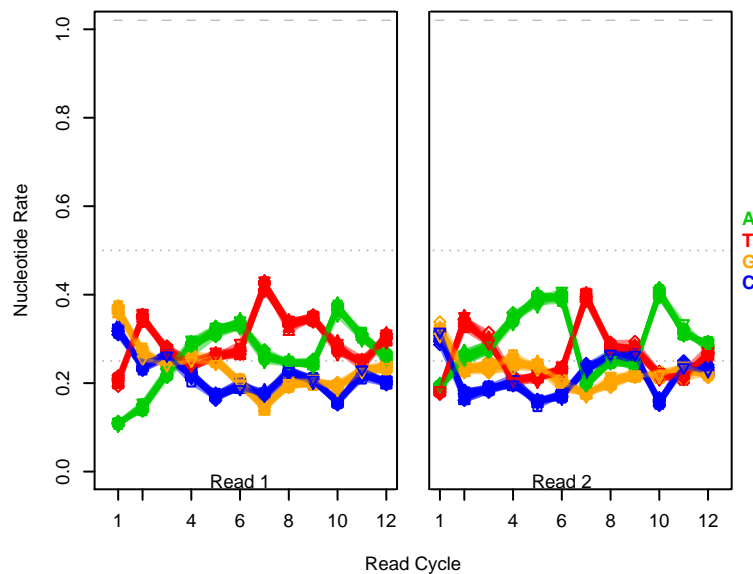


ap



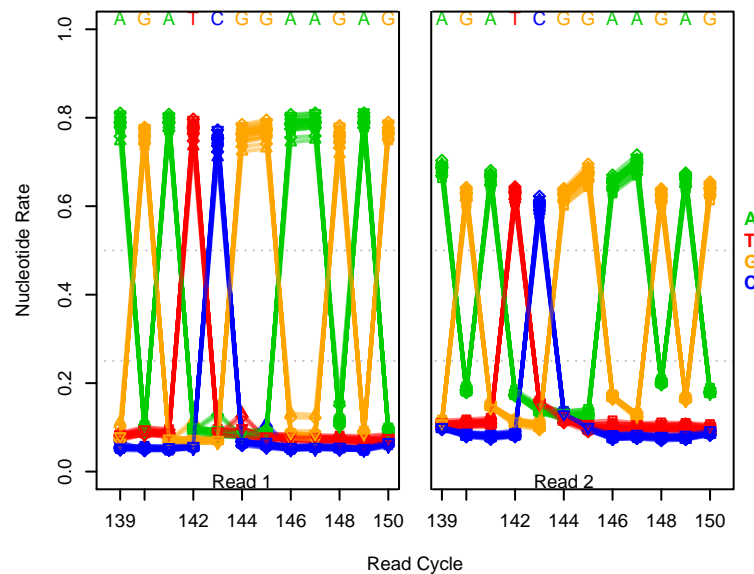
aq

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



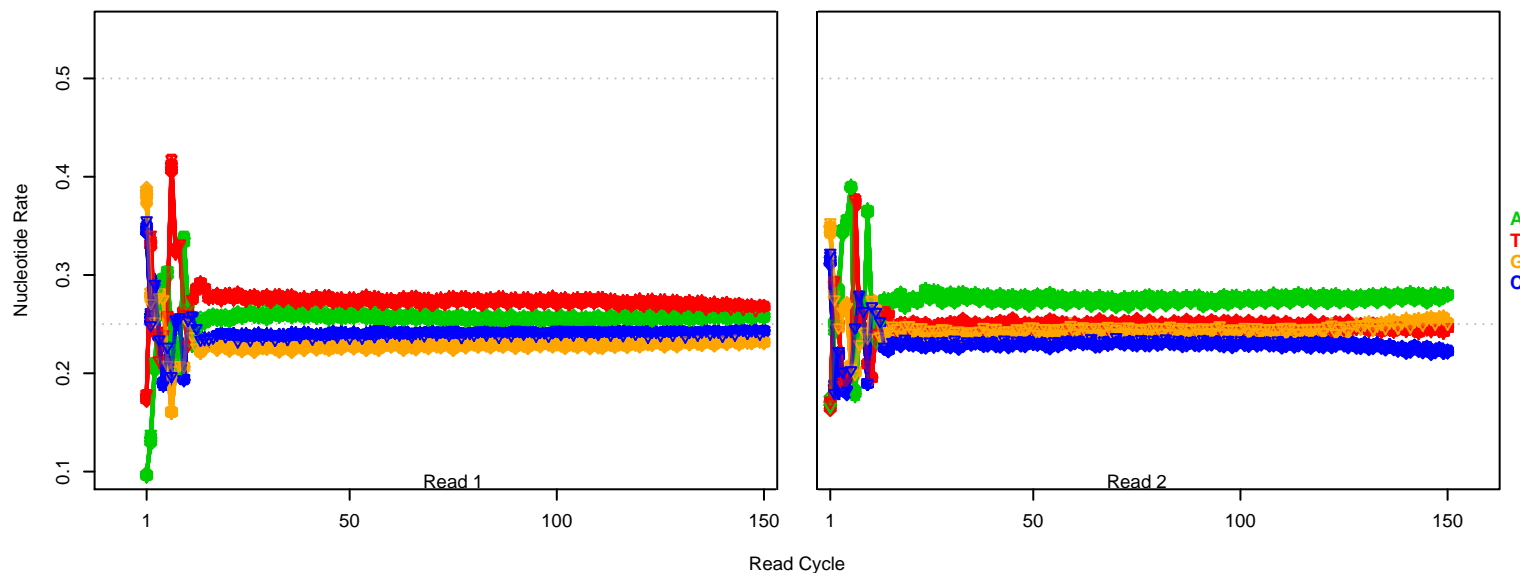
ar

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



as

Raw Nucleotide Rate by Cycle  
Marked by Group



at

Nucleotide Rate by Cycle, Aligned bases only  
Marked by Group

