

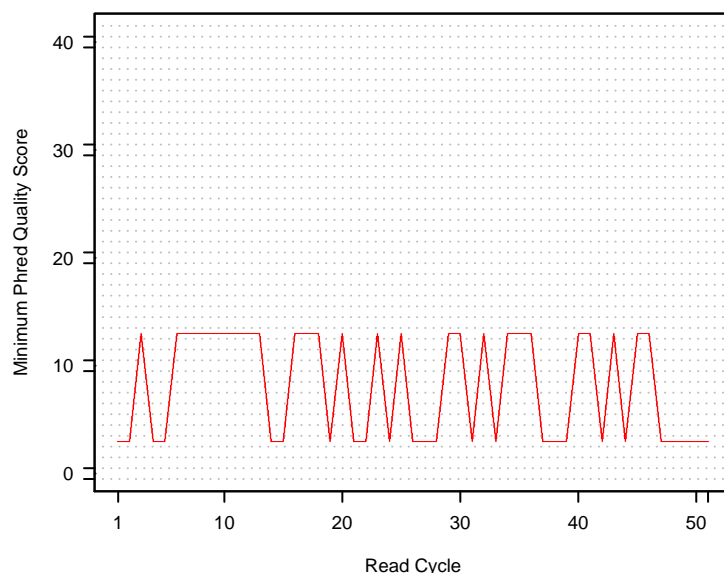
a

Sample Highlight:
d1_aligned_sorted
Colored by Lane



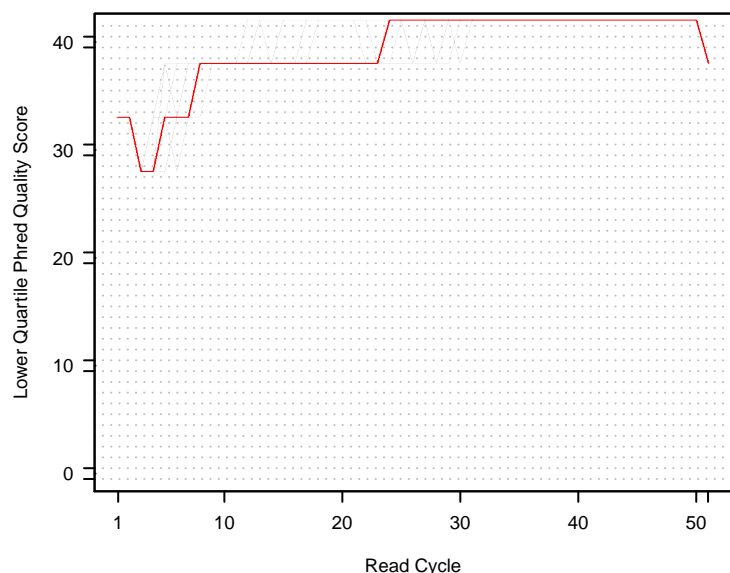
b

Minimum Phred Quality Score
With Sample d1_aligned_sorted Colored by lane



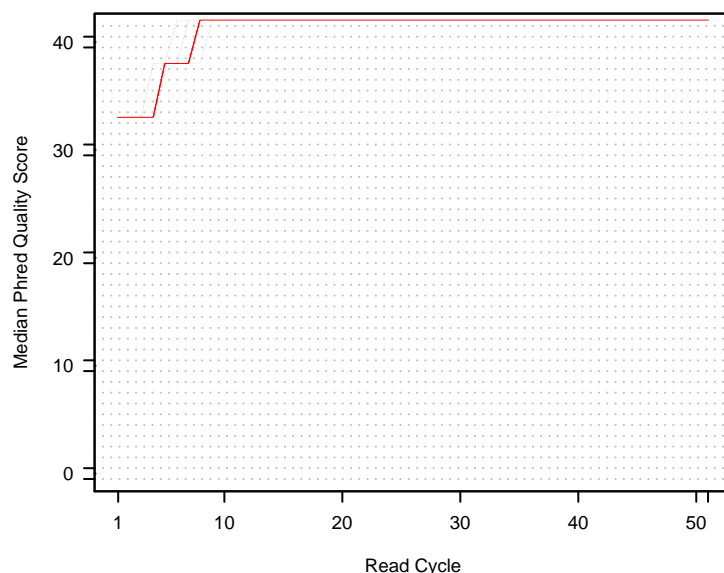
c

Lower Quartile Phred Quality Score
With Sample d1_aligned_sorted Colored by lane



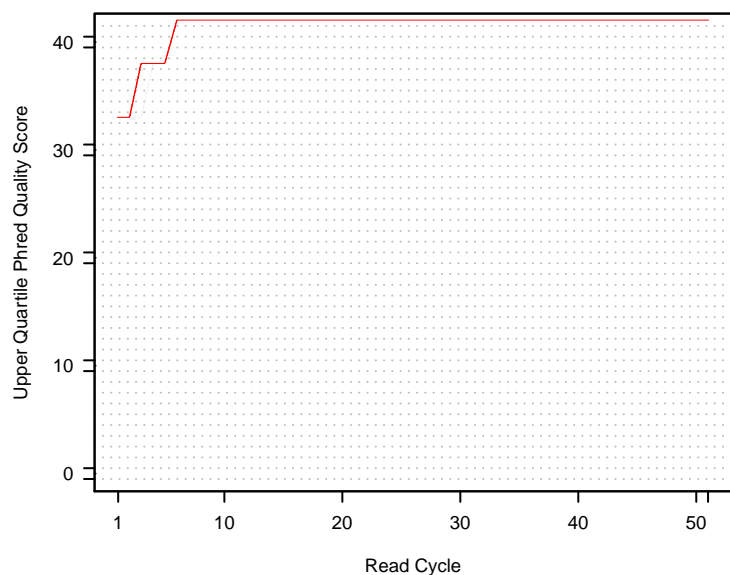
d

Median Phred Quality Score
With Sample d1_aligned_sorted Colored by lane



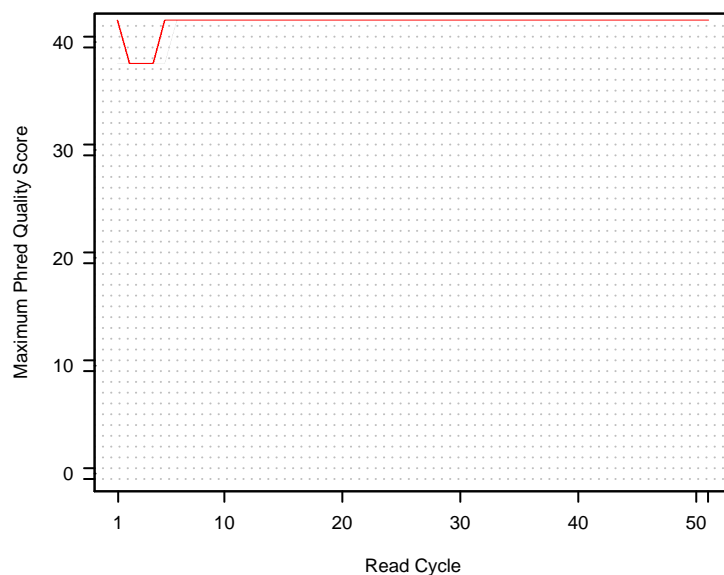
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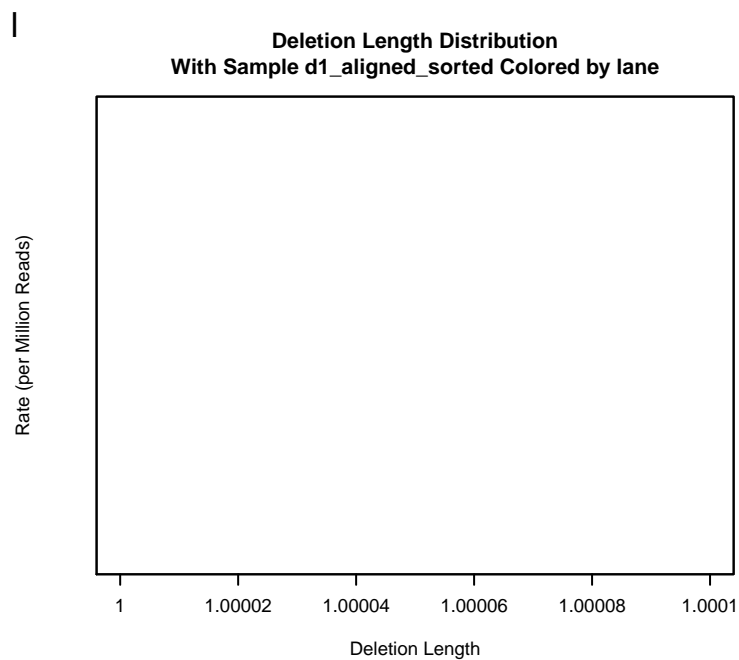
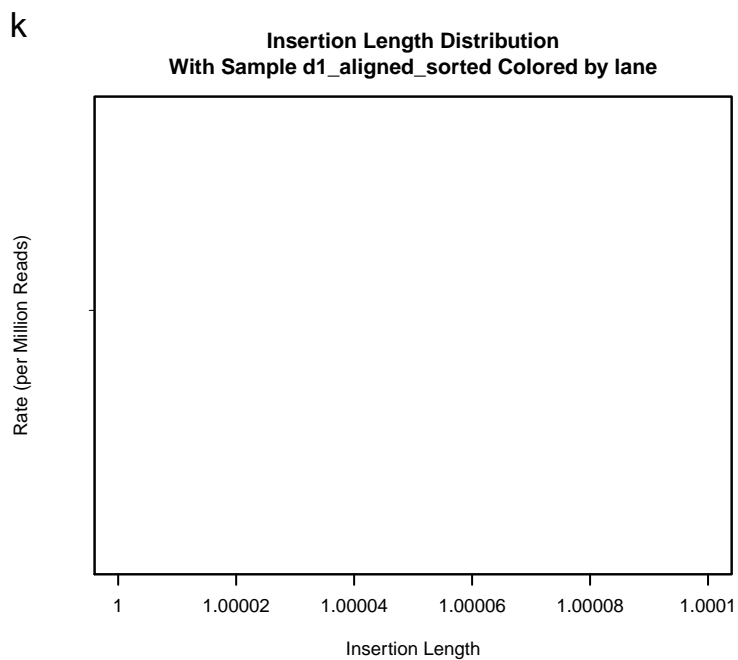
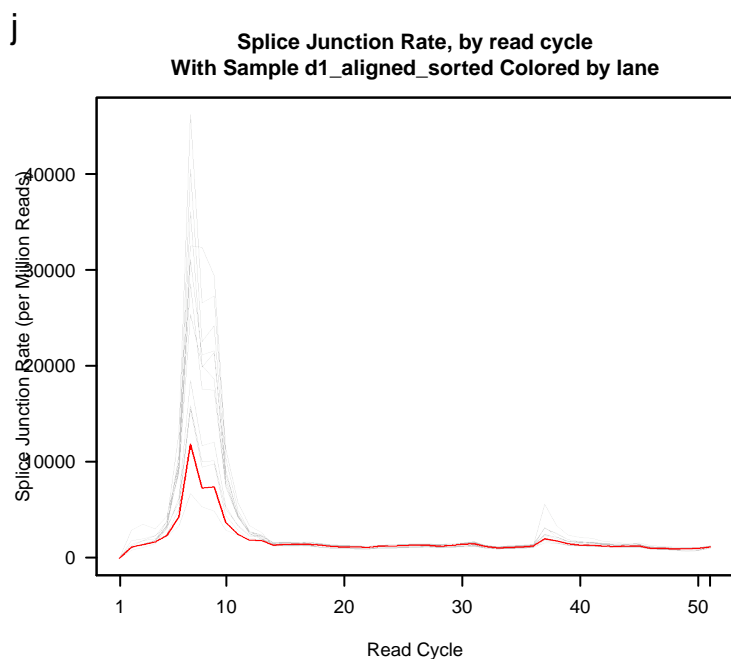
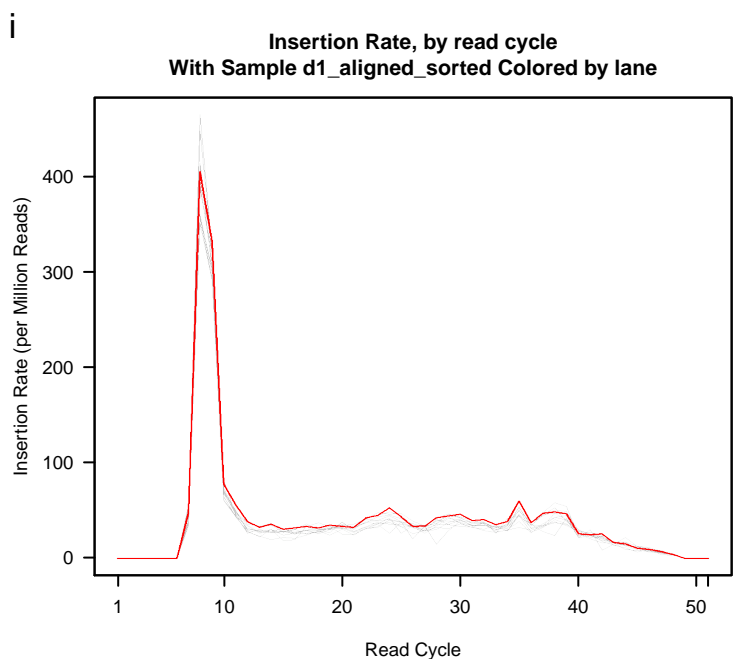
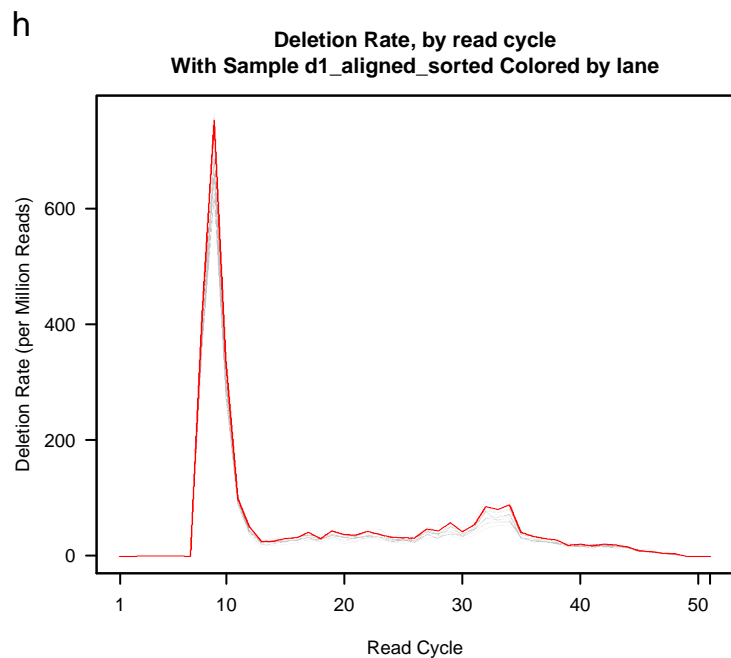
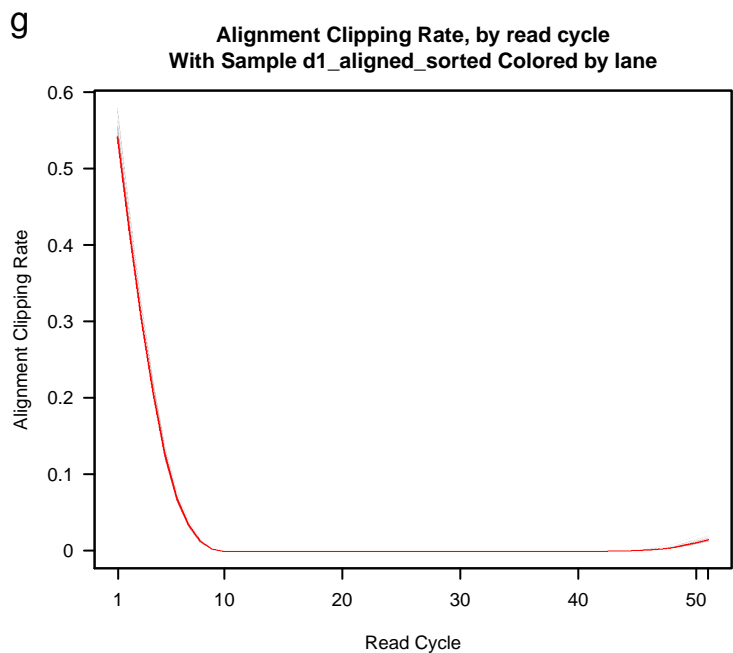
Upper Quartile Phred Quality Score
With Sample d1_aligned_sorted Colored by lane

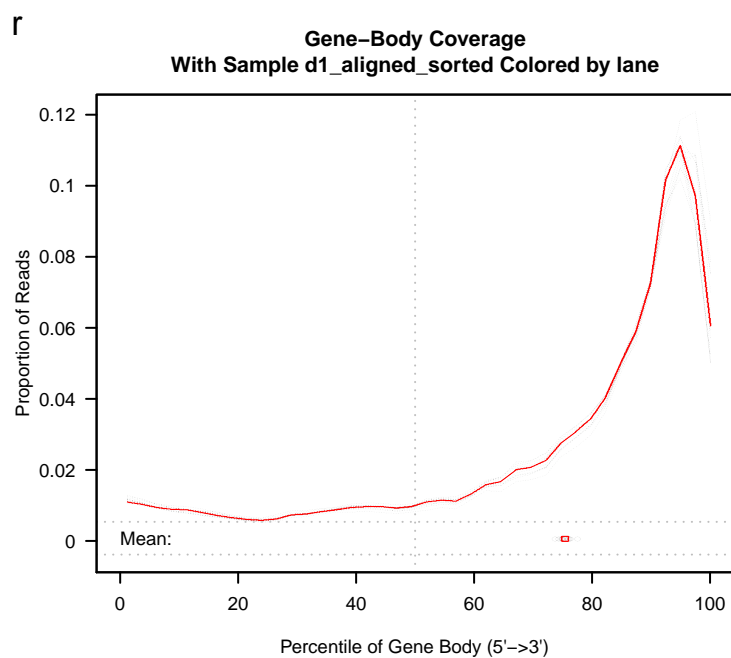
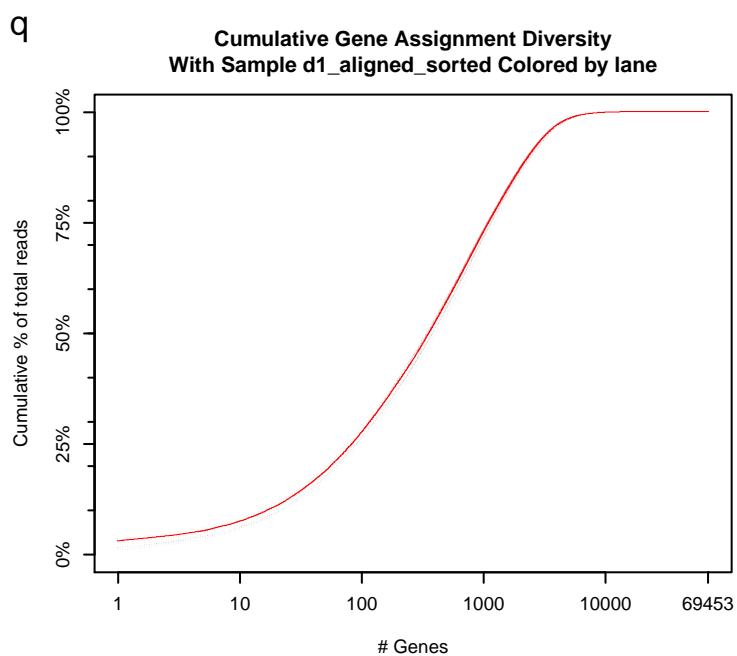
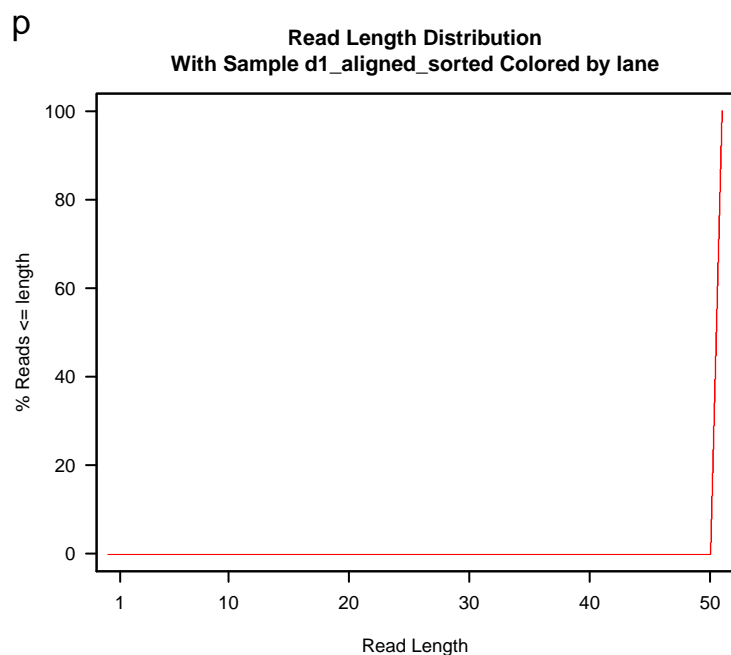
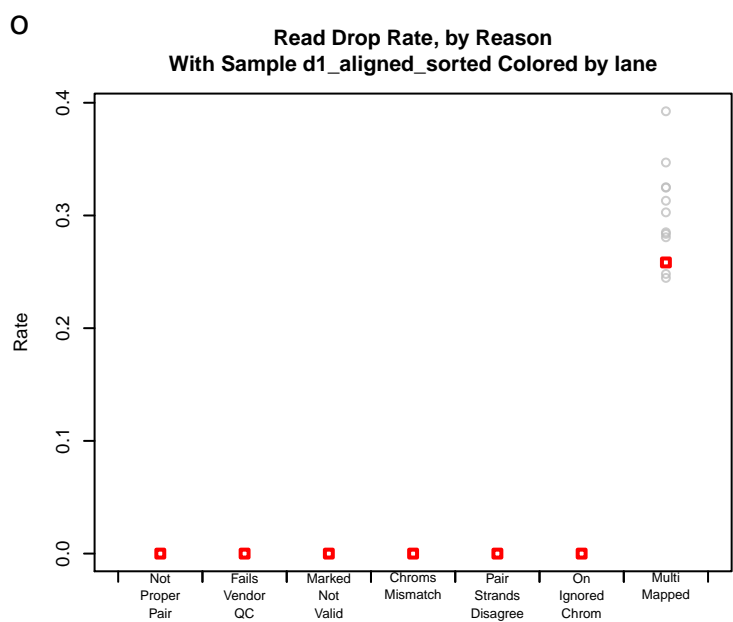
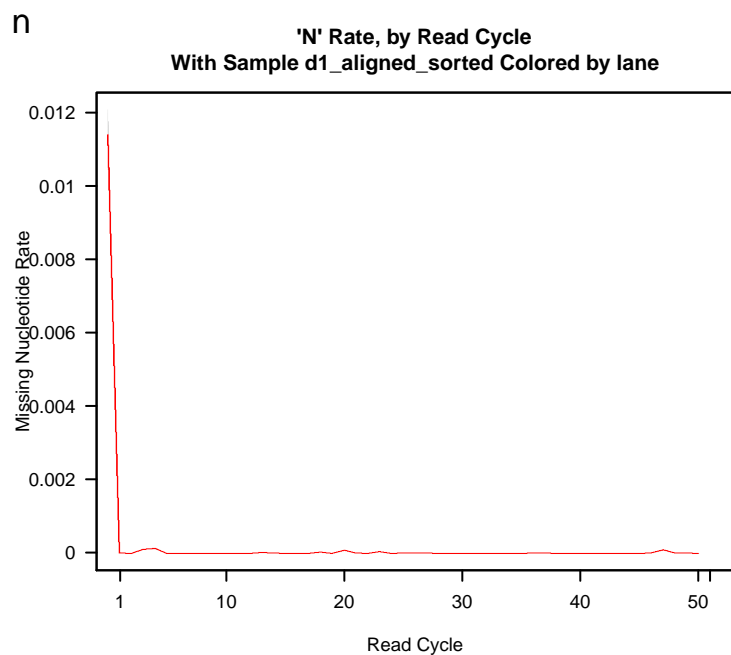
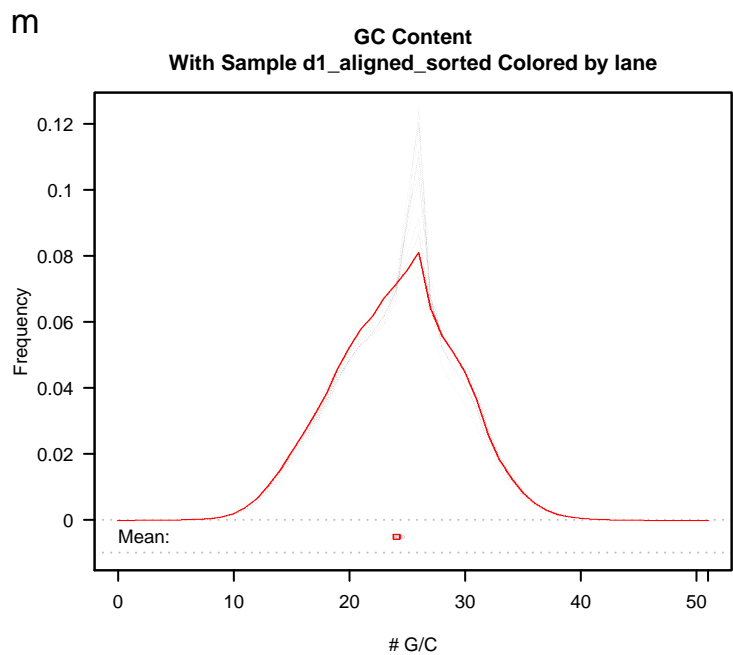


f

Maximum Phred Quality Score
With Sample d1_aligned_sorted Colored by lane

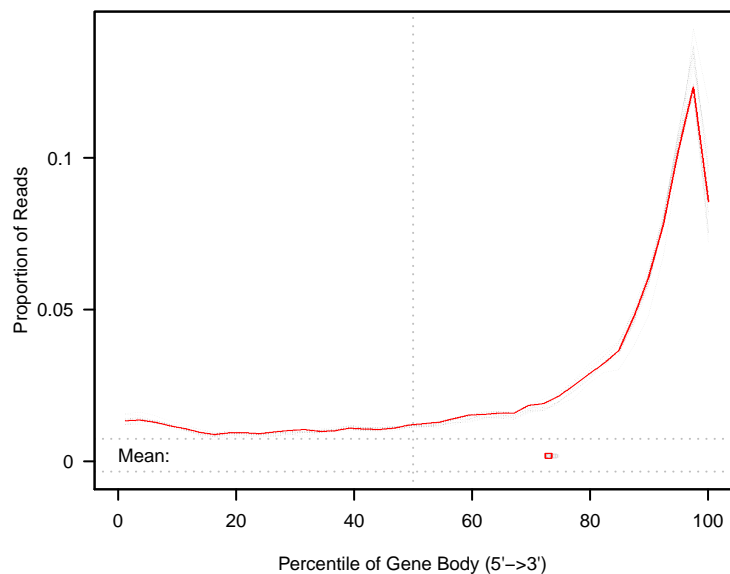






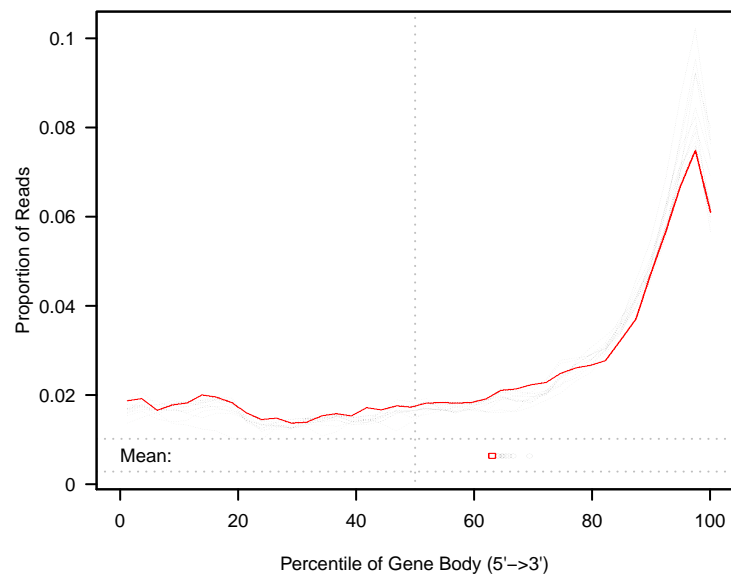
S

Gene–Body Coverage, Upper Middle Quartile Genes
With Sample d1_aligned_sorted Colored by lane



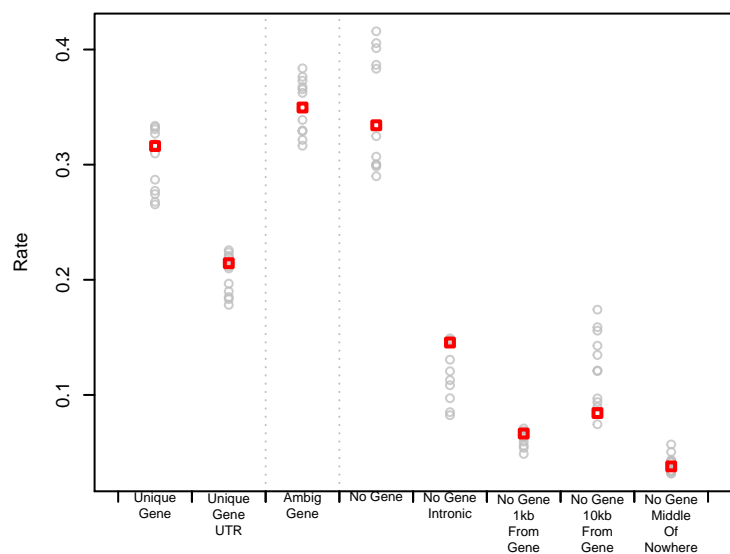
t

Gene–Body Coverage, Low Expression Genes
With Sample d1_aligned_sorted Colored by lane



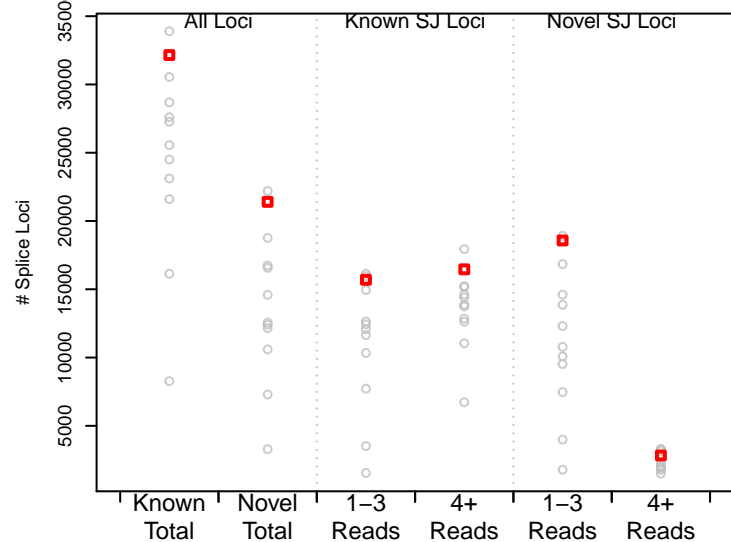
U

Read Mapping Location Rates
With Sample d1_aligned_sorted Colored by lane



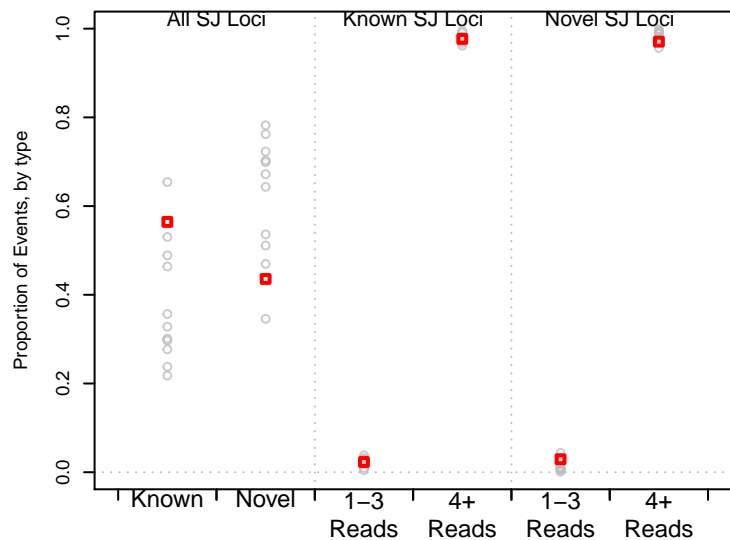
V

Observed Splice Junction Loci, by type
With Sample d1_aligned_sorted Colored by lane



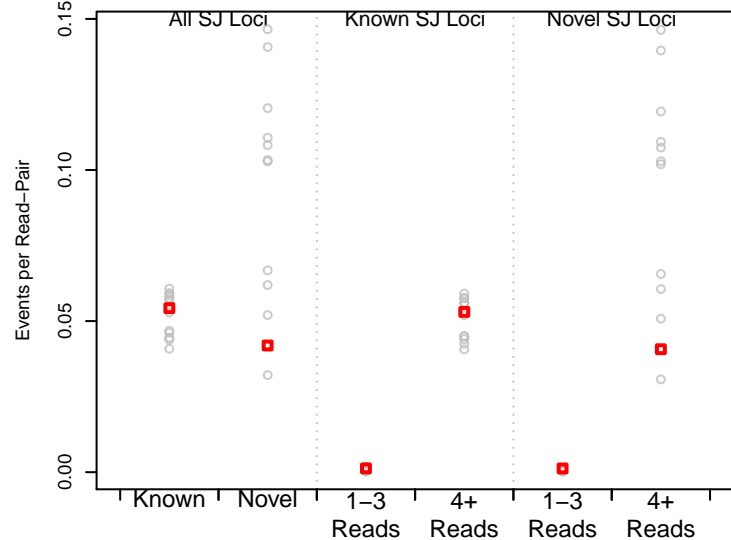
W

Breakdown of Splice Junction Events, by type
With Sample d1_aligned_sorted Colored by lane

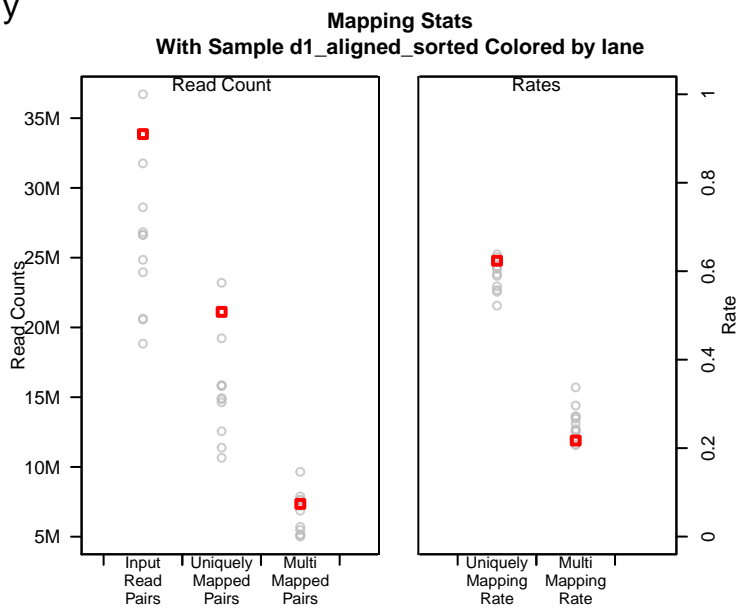


X

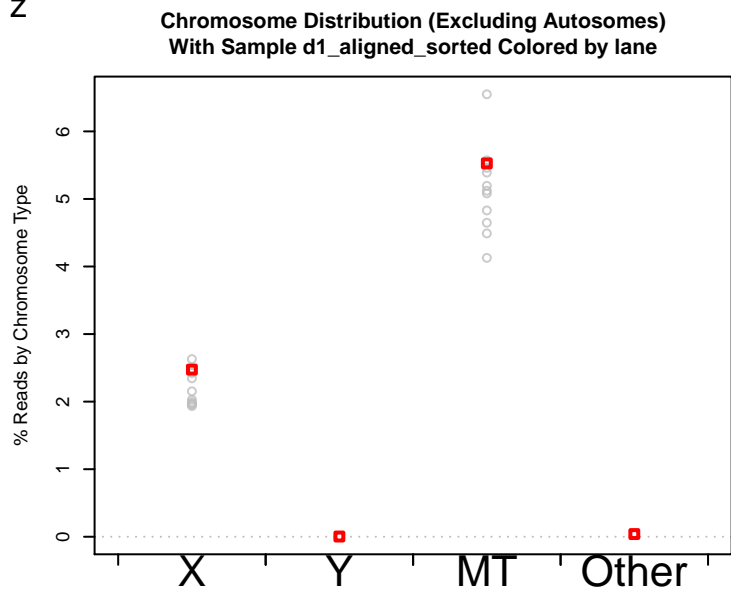
Splice Junction Event Rates per Read–Pair
With Sample d1_aligned_sorted Colored by lane



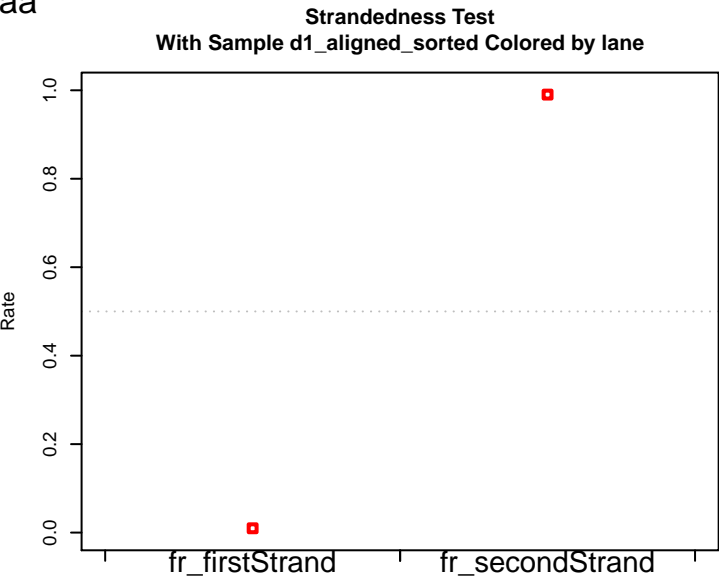
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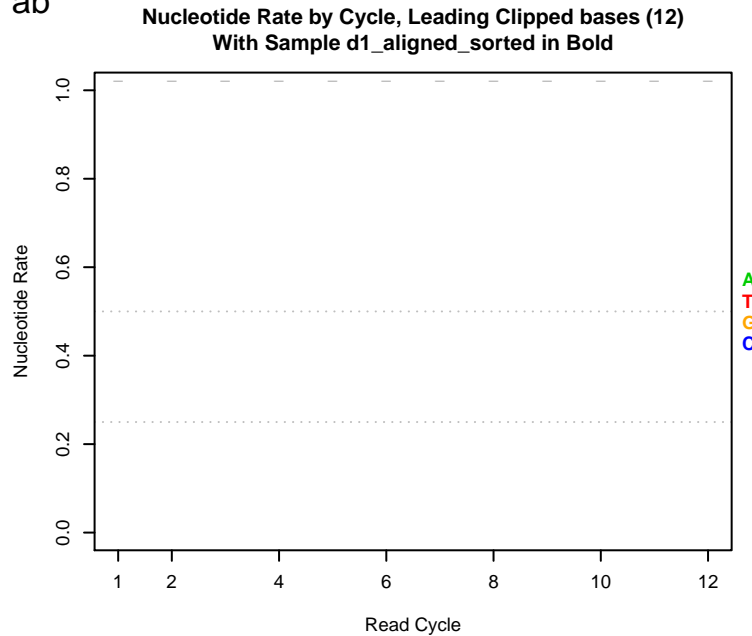
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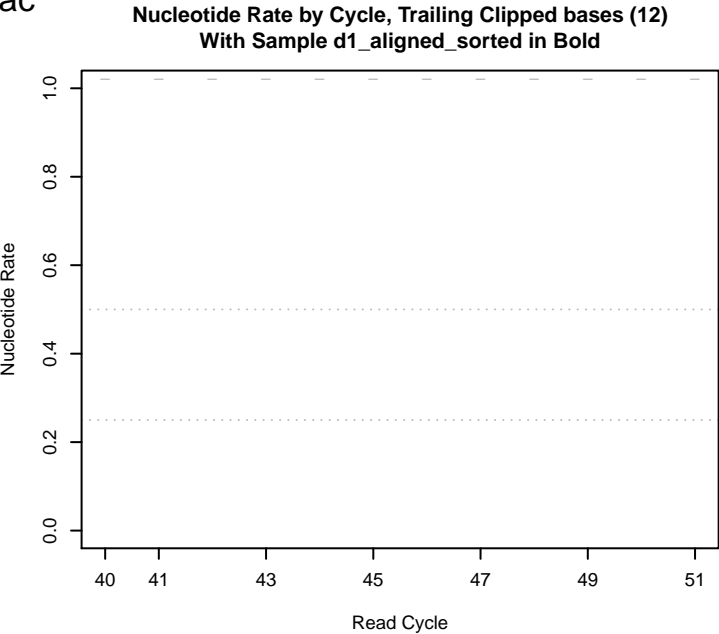
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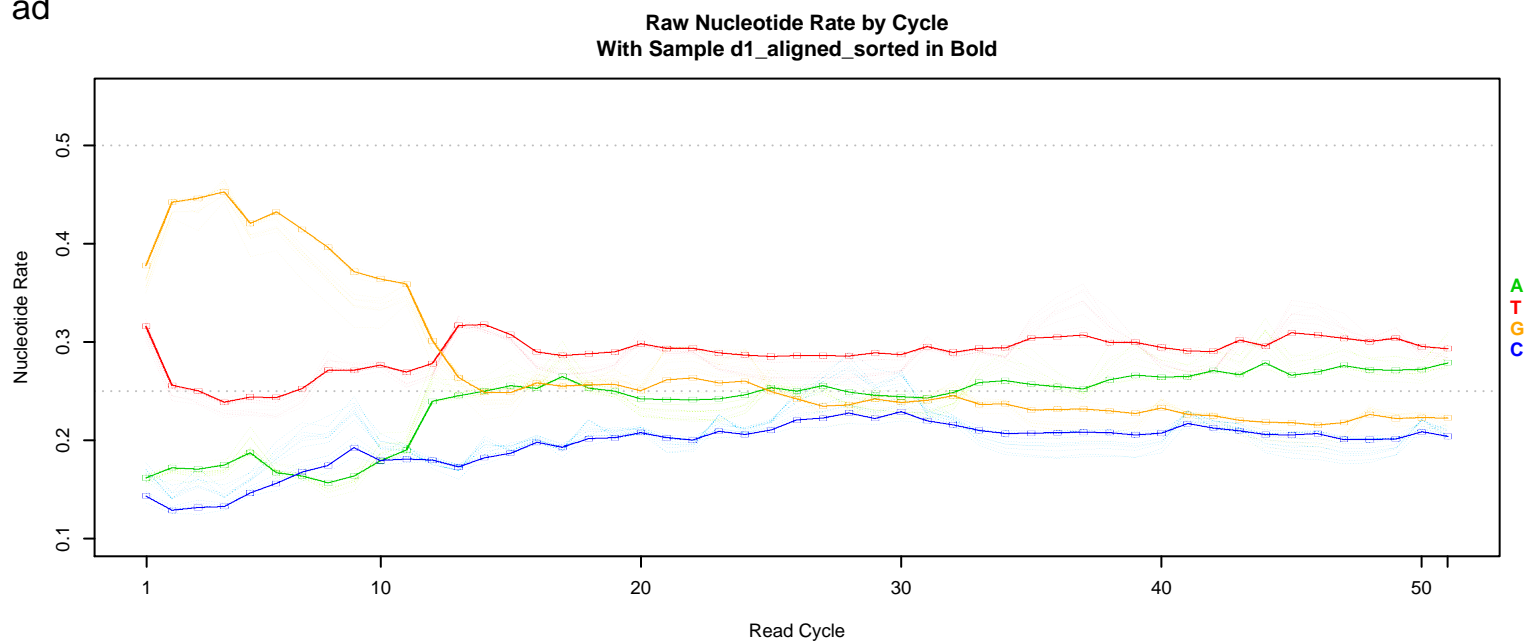
ab



ac



ad



ae

