# Read Maps and Size Distributions



Read counts

Read sizes / Nucleotide coordinates

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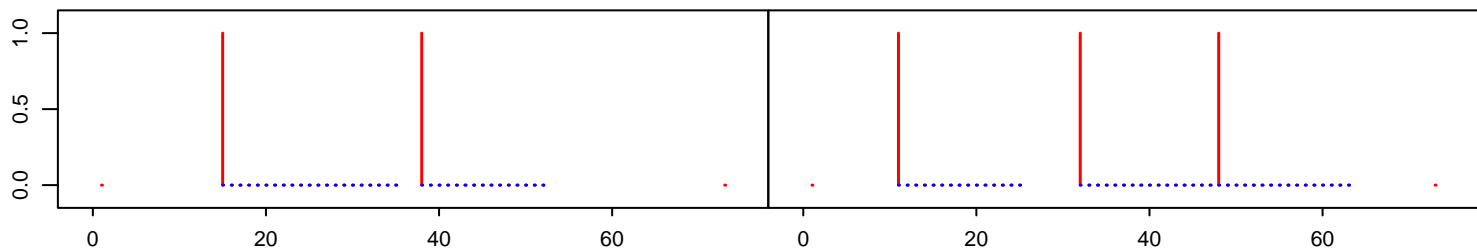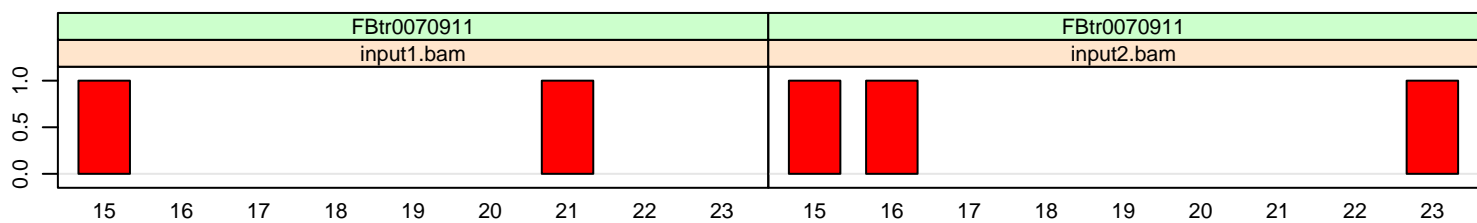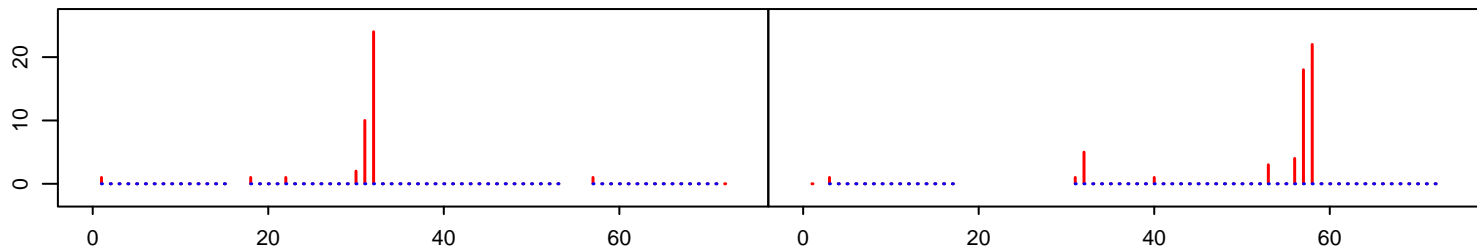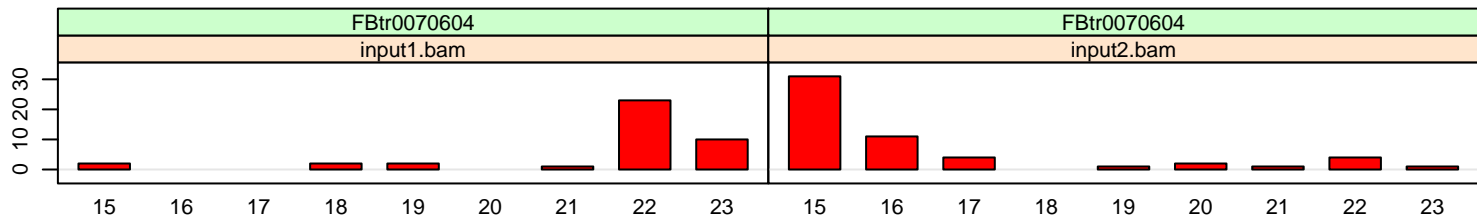


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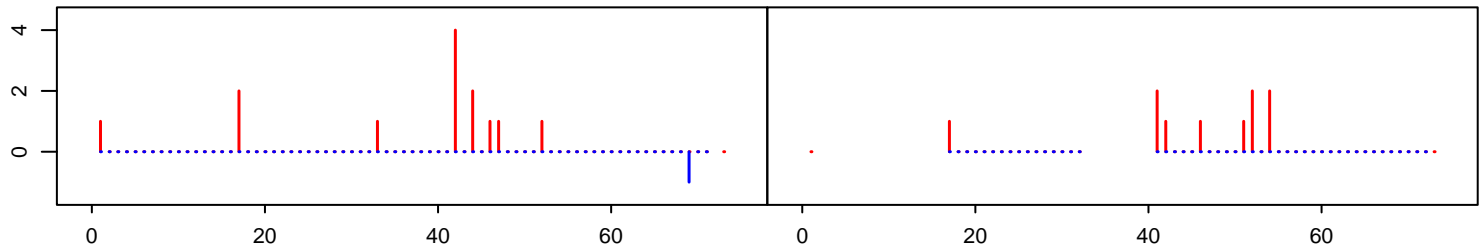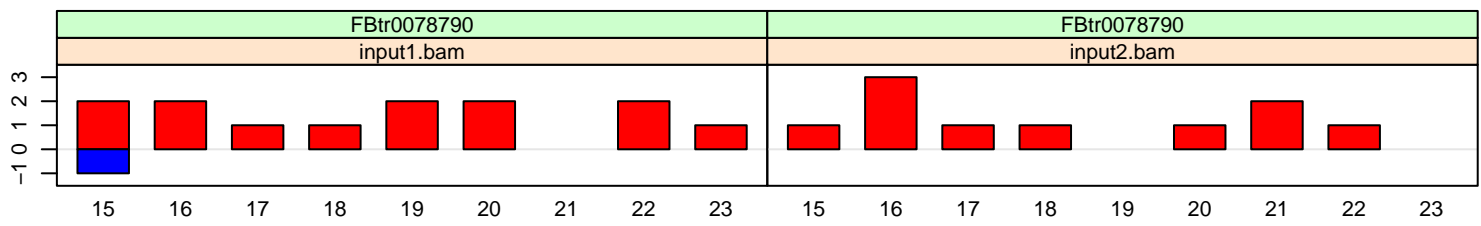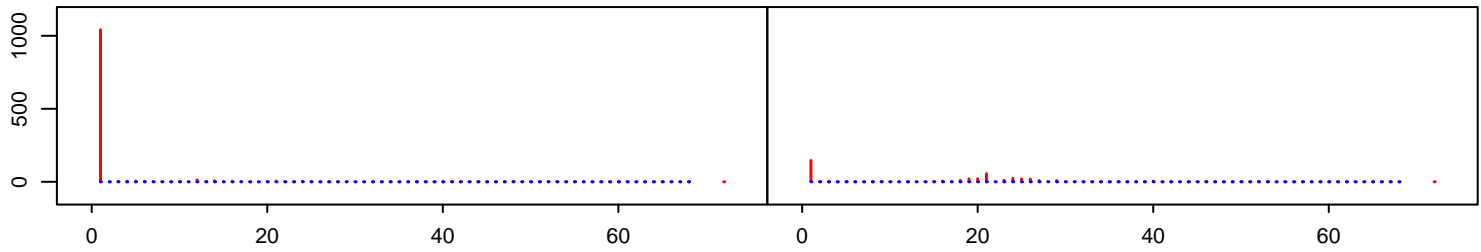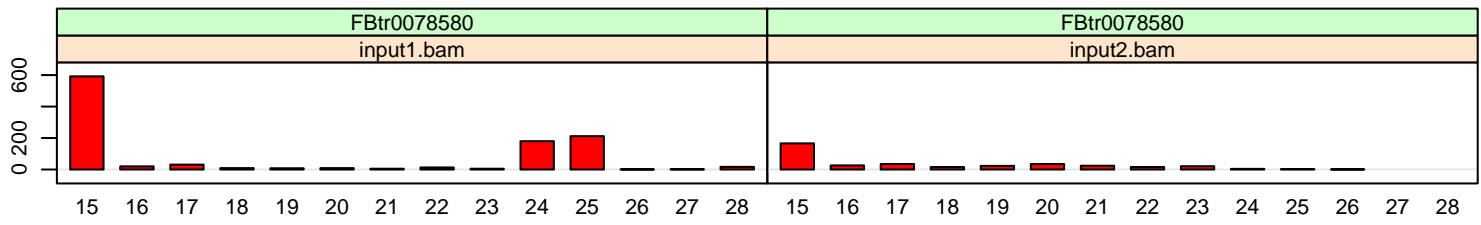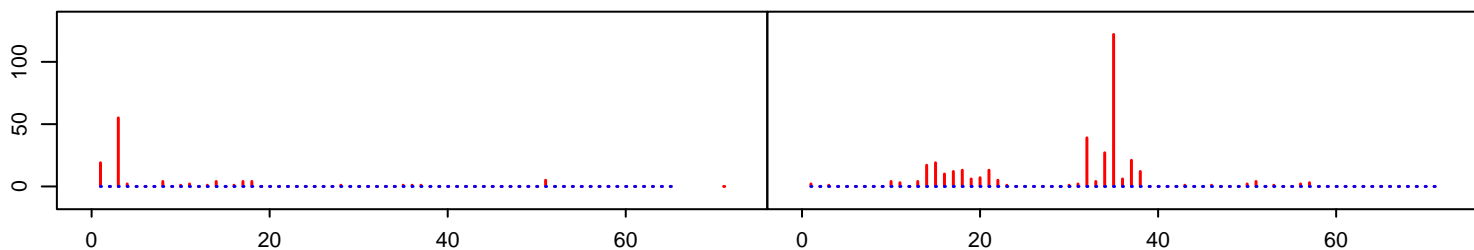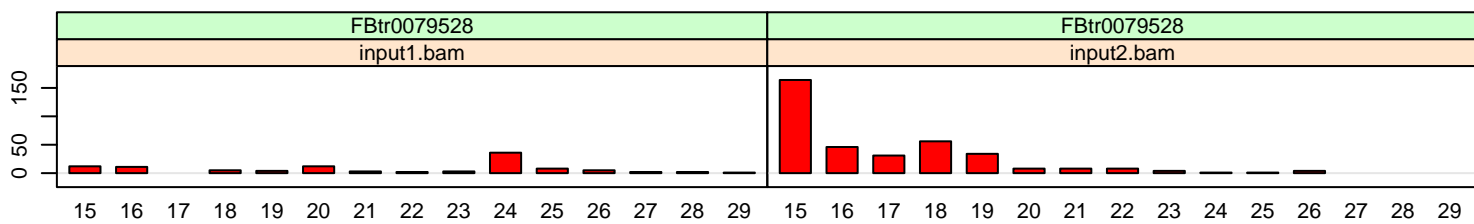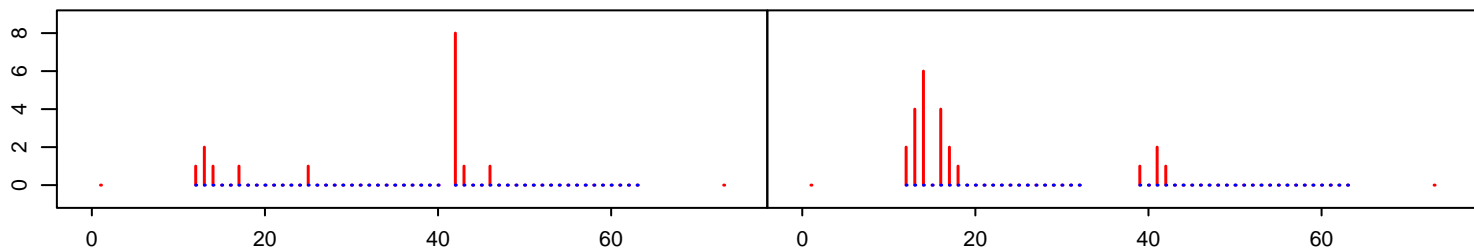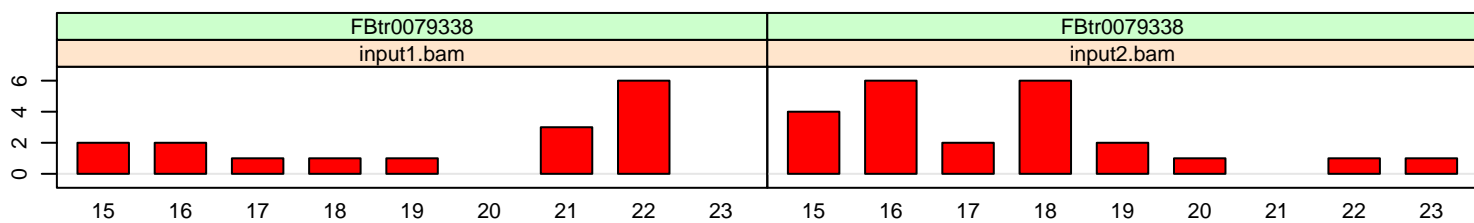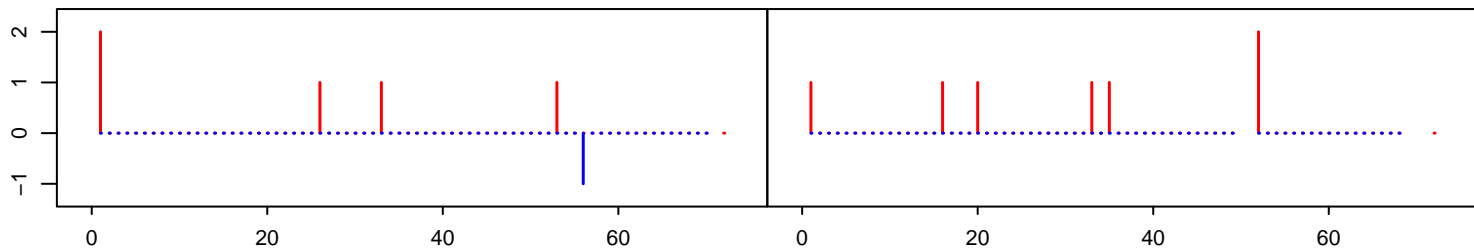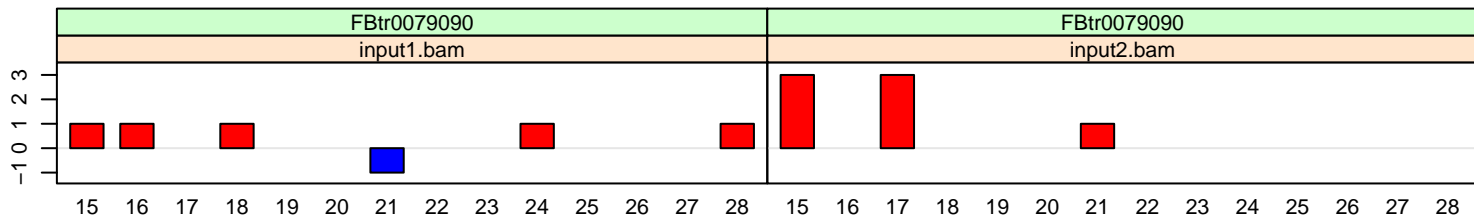


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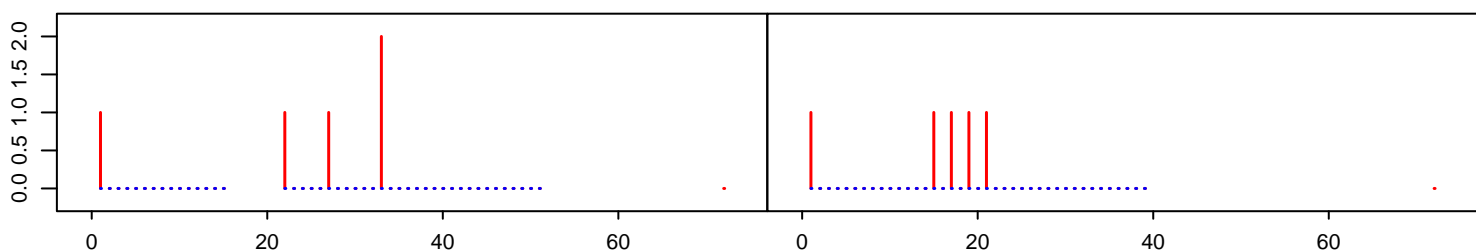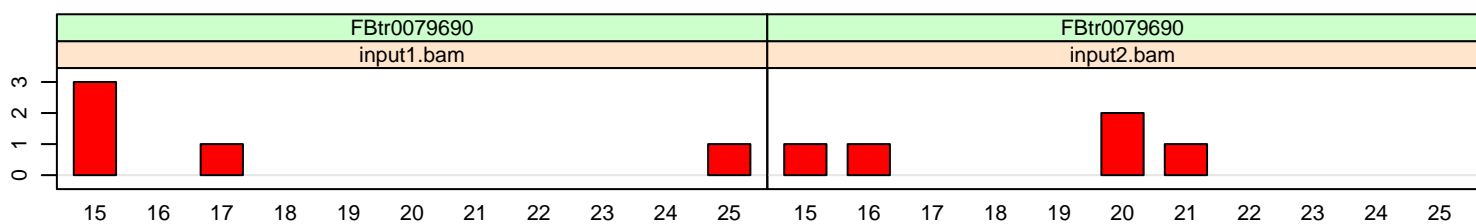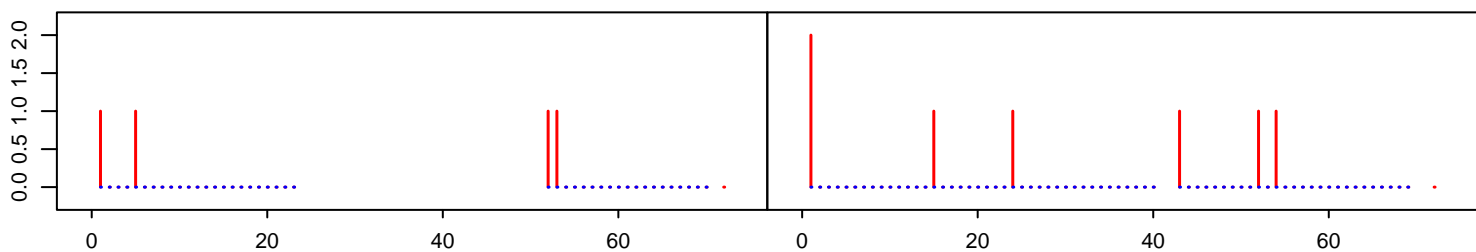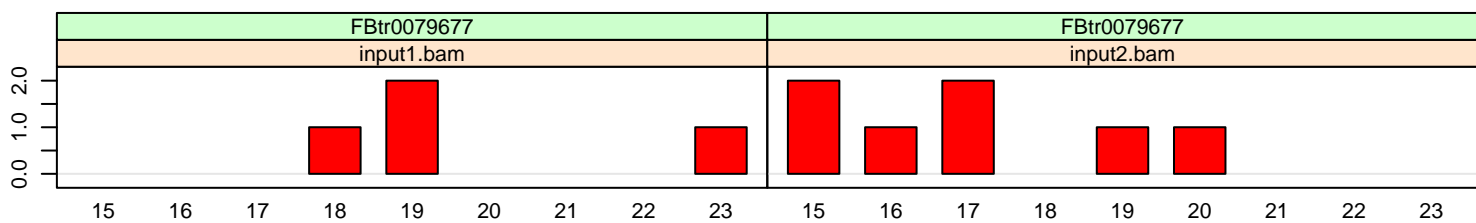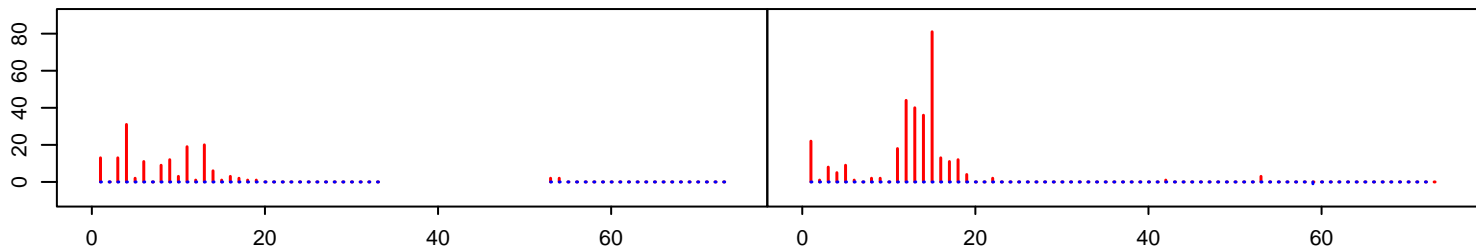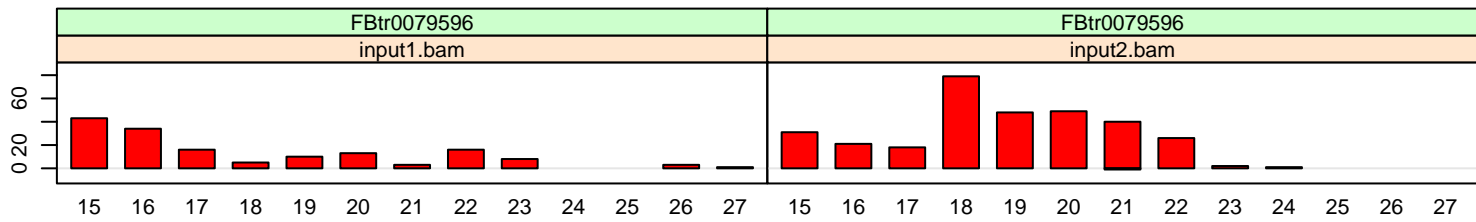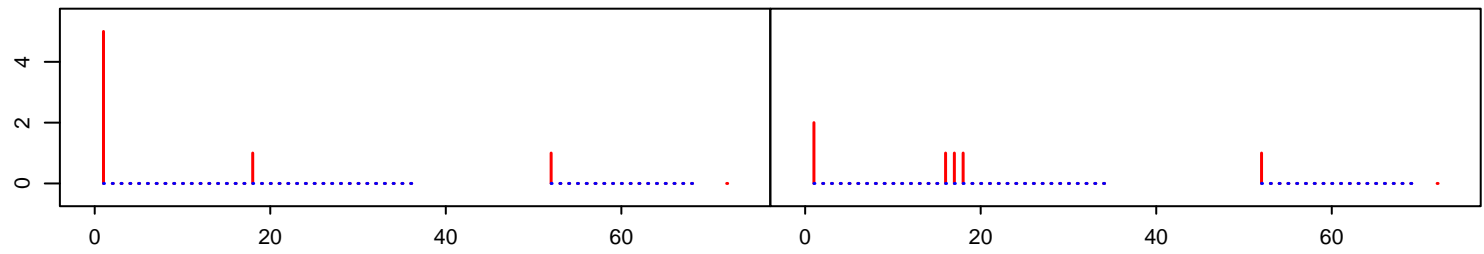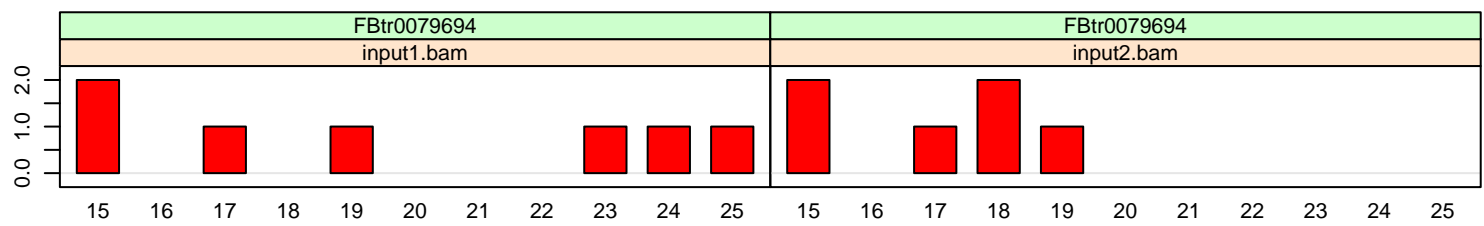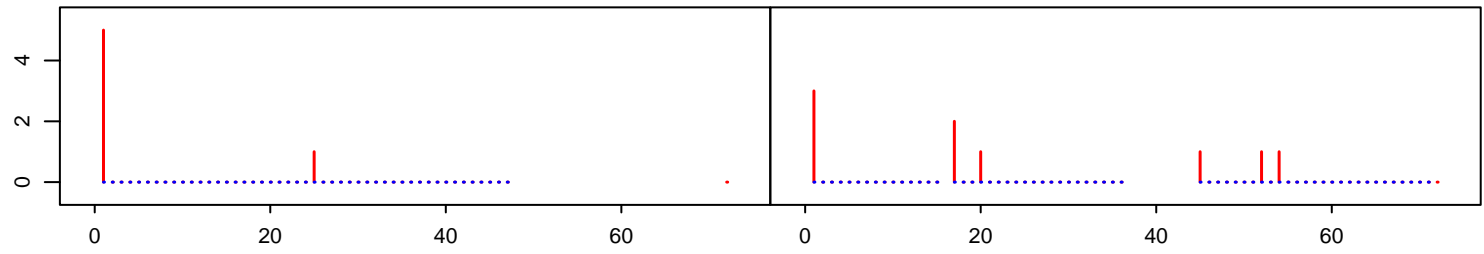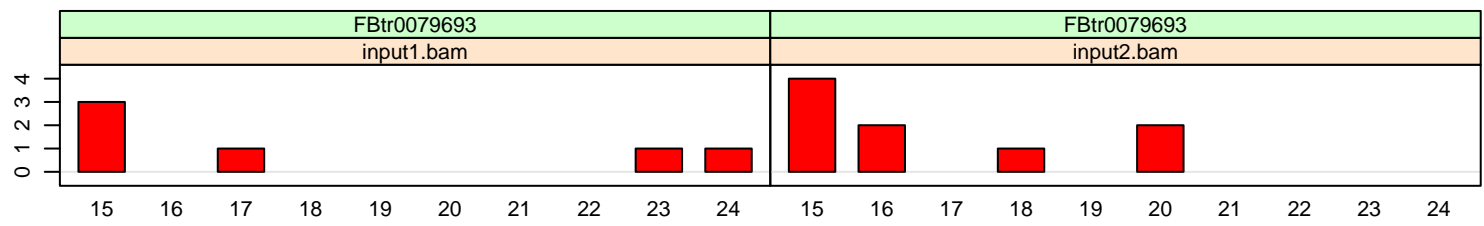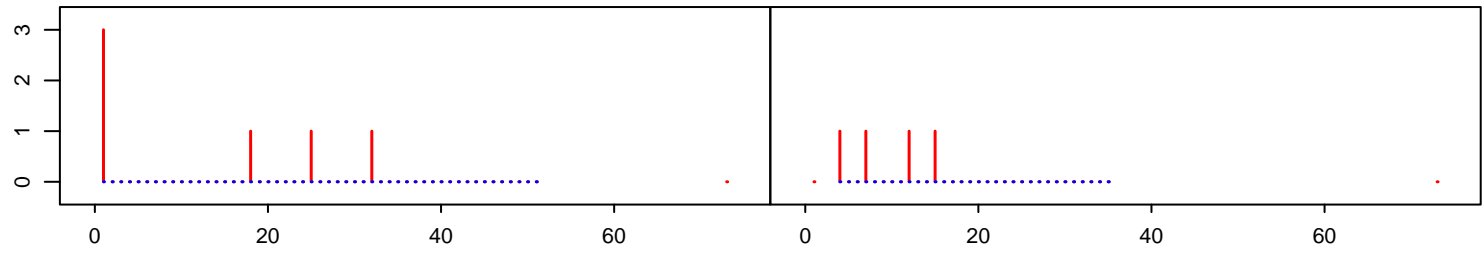Read counts

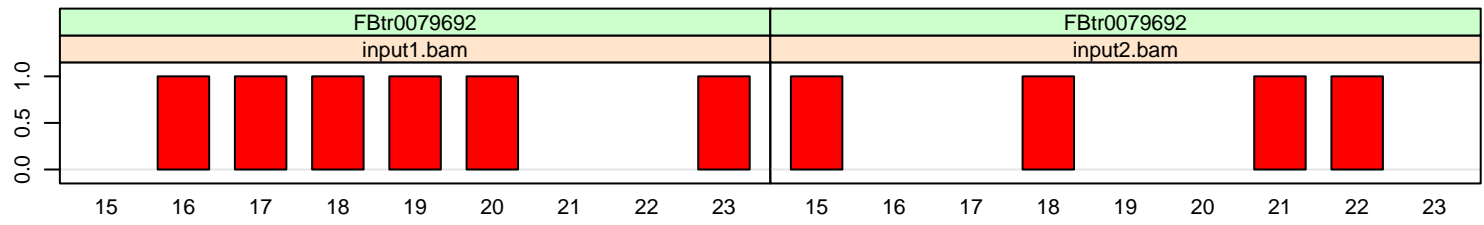Read sizes / Nucleotide coordinates

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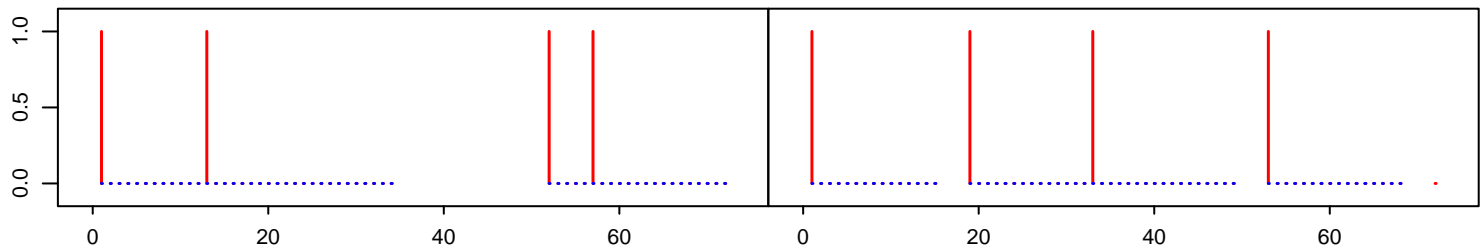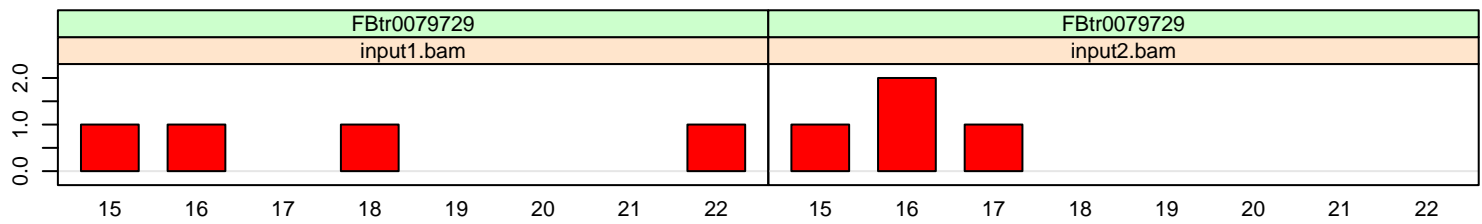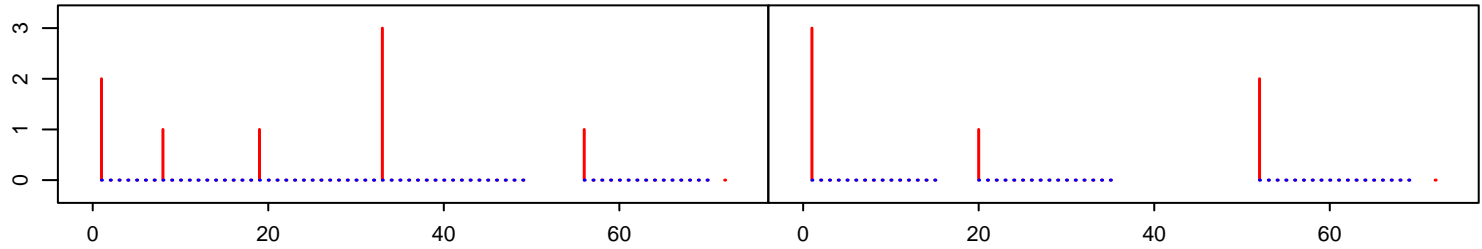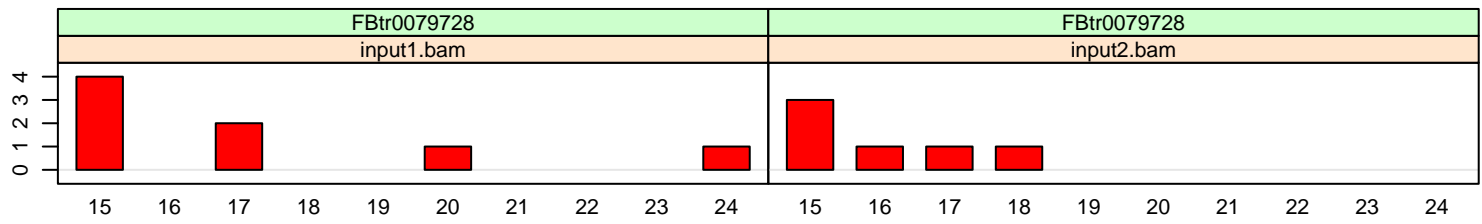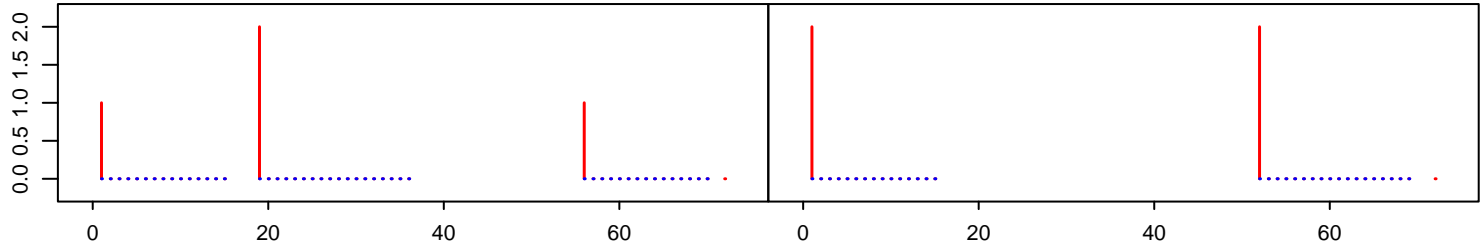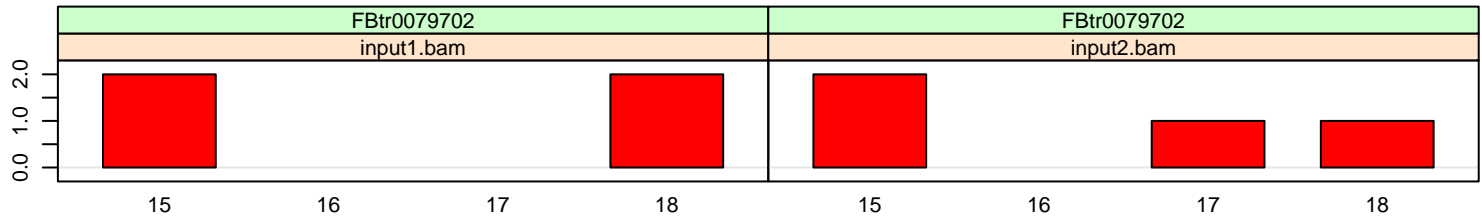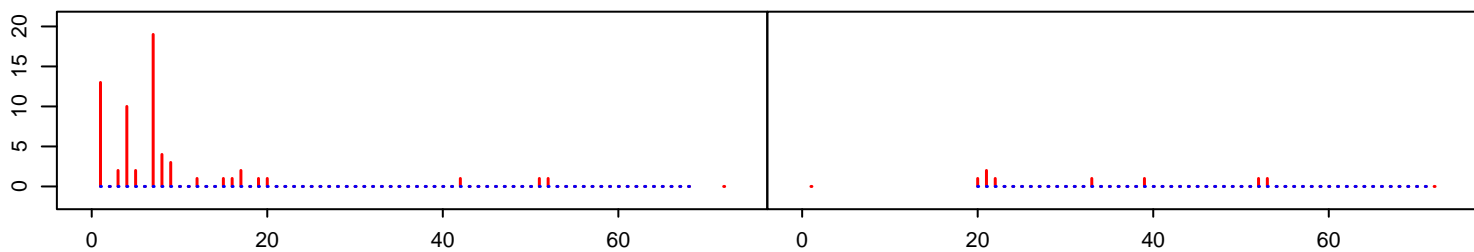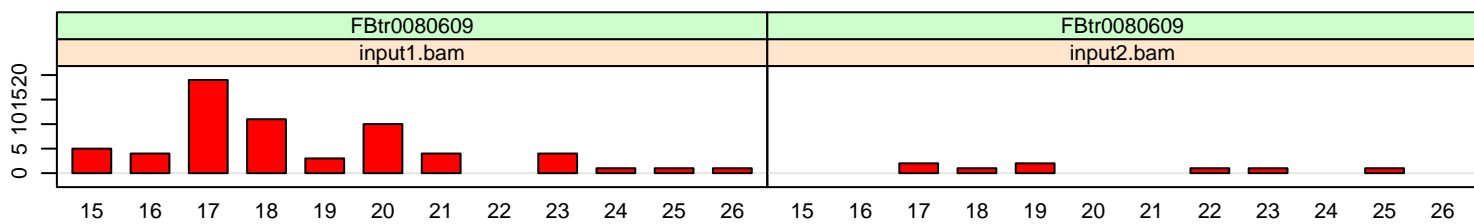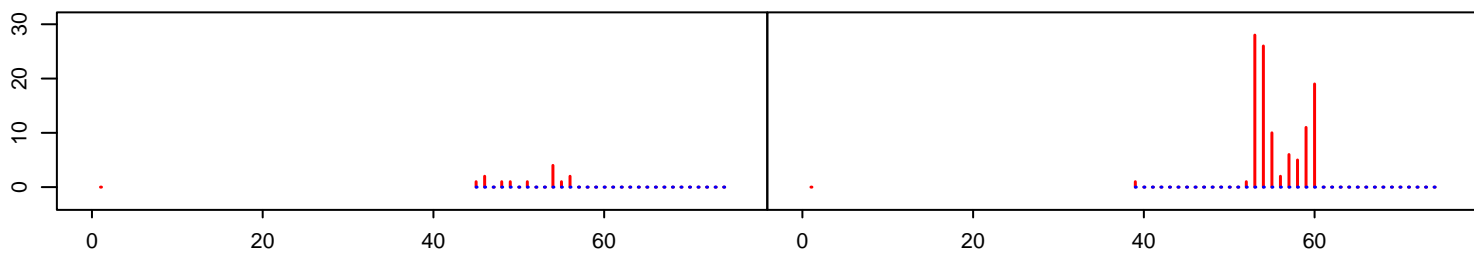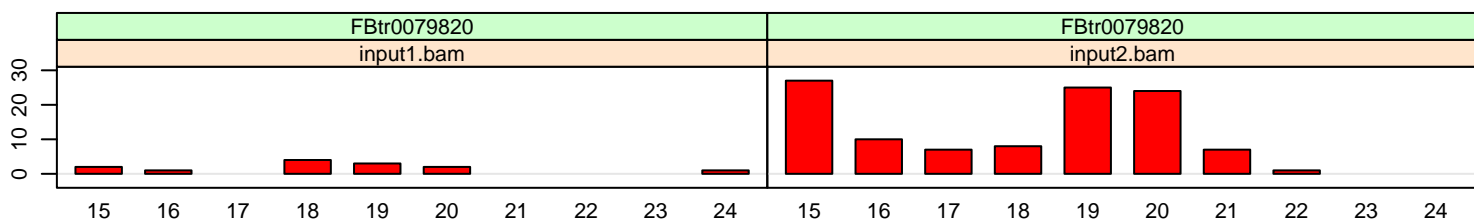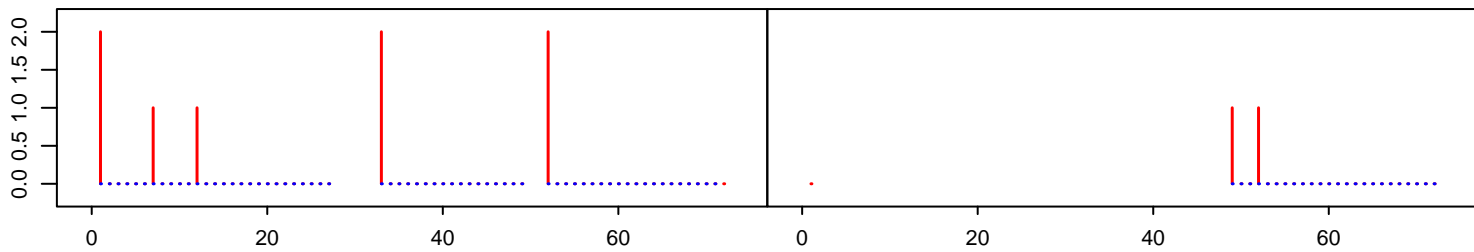Read counts

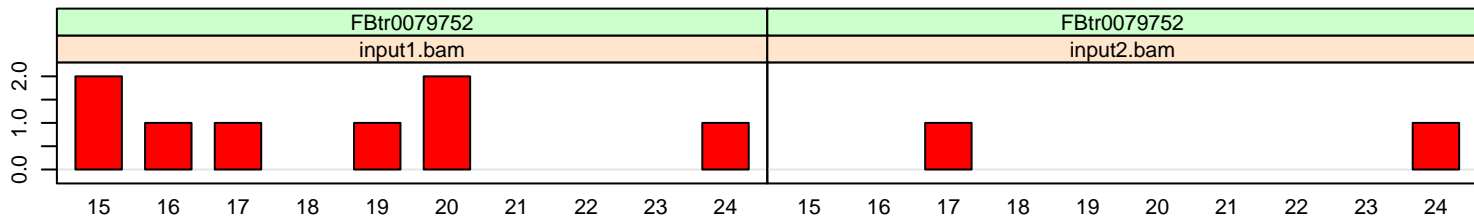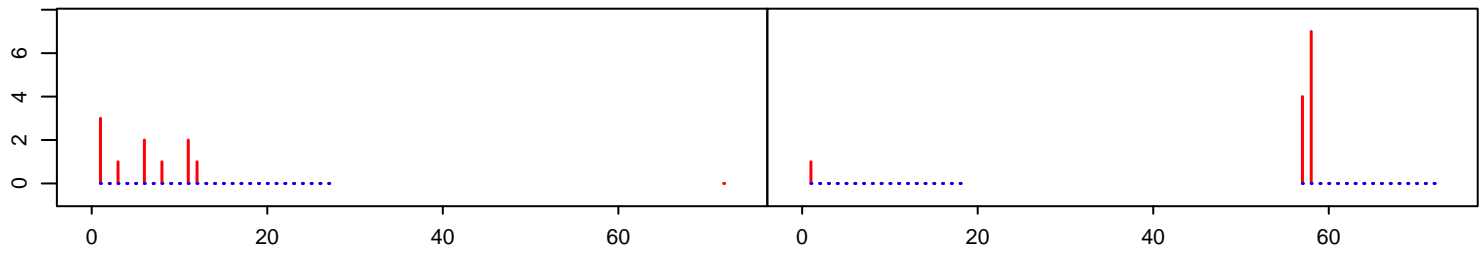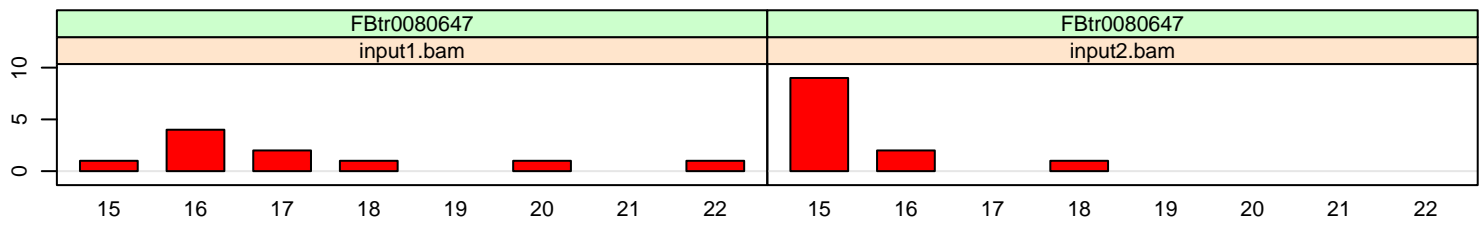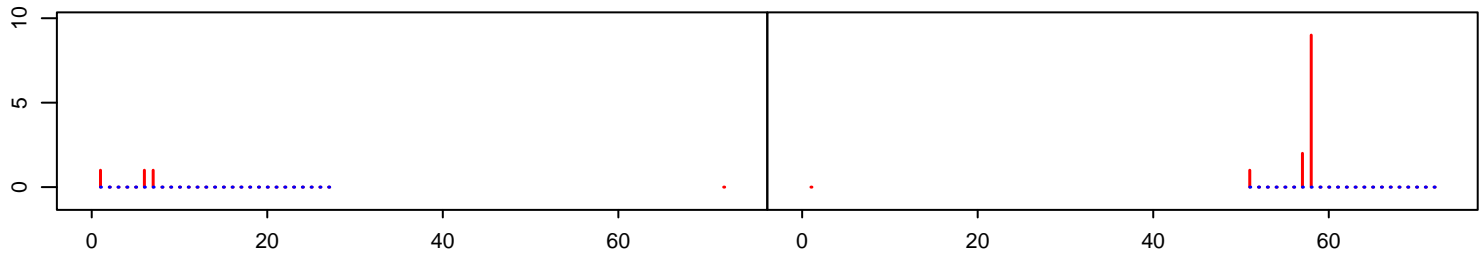Read sizes / Nucleotide coordinates

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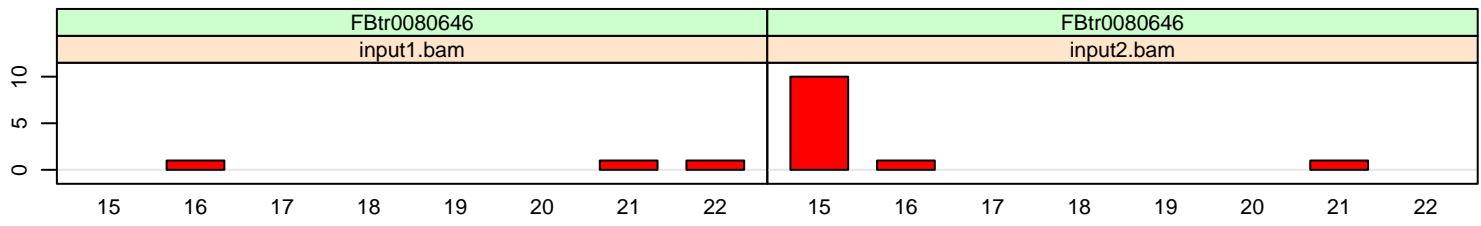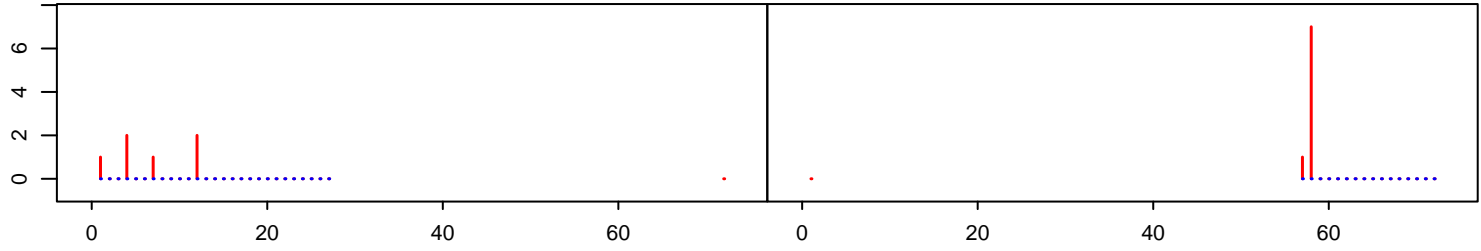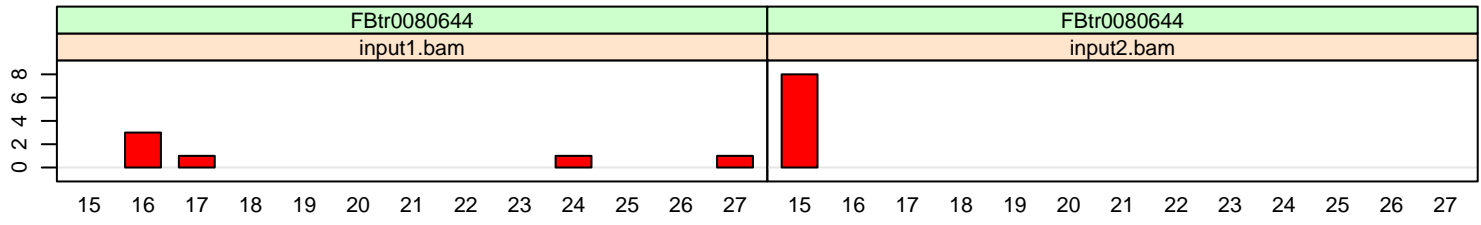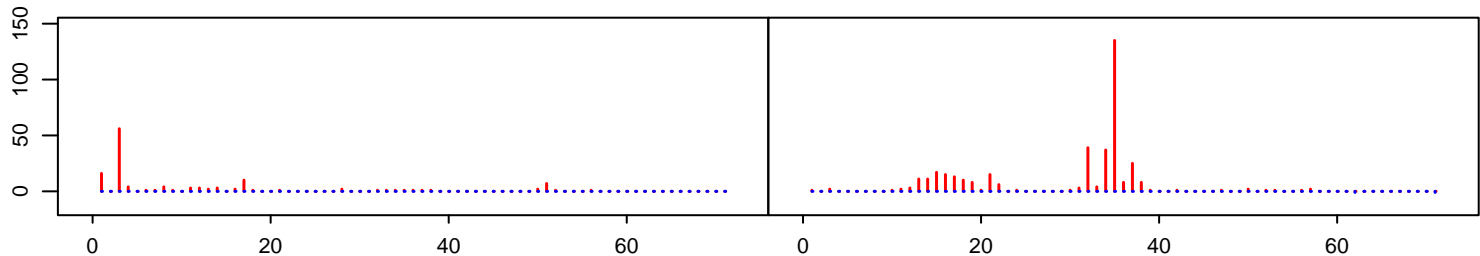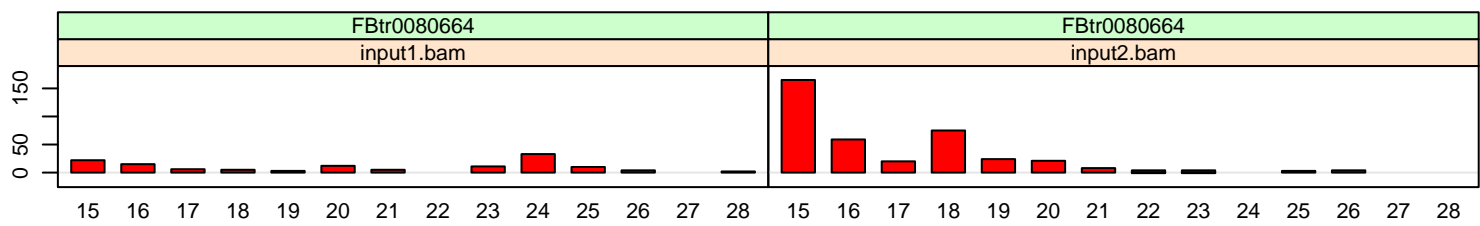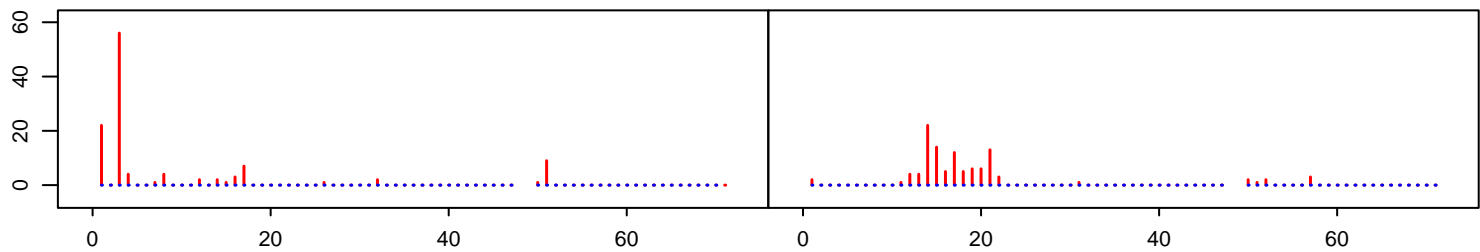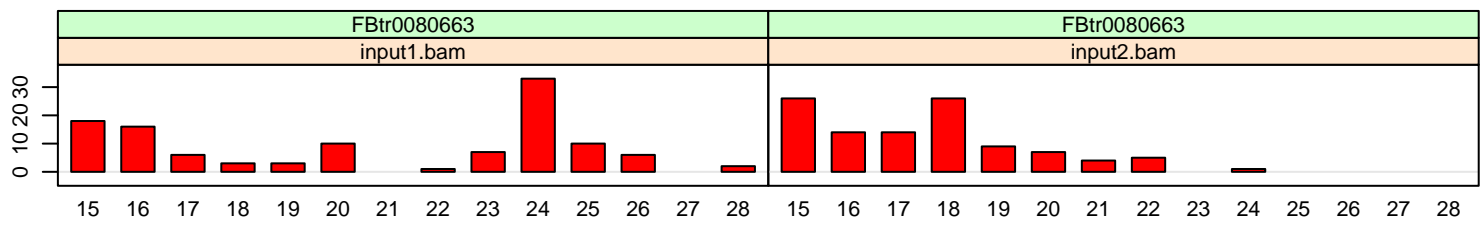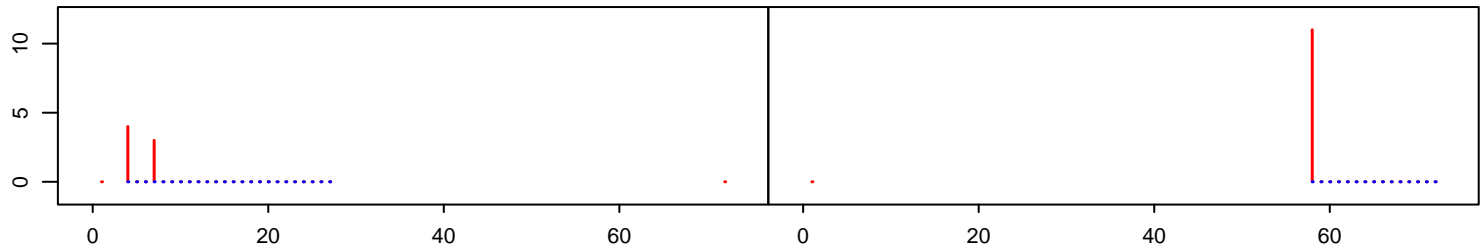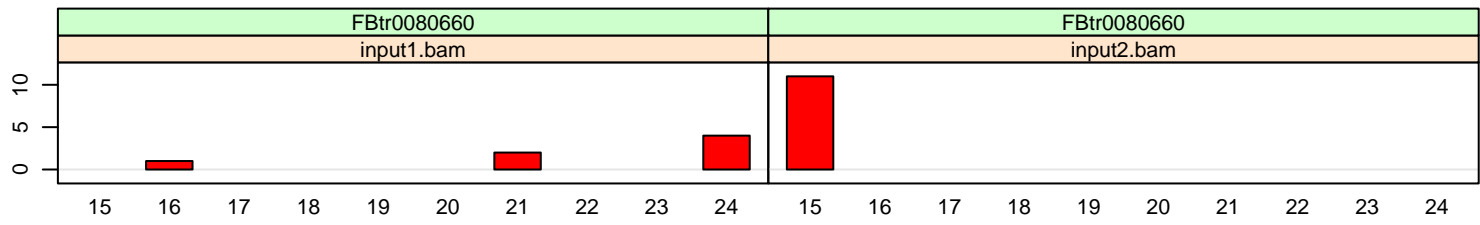


**Read counts** (y-axis)

**Read sizes / Nucleotide coordinates** (x-axis)