

Read Counts

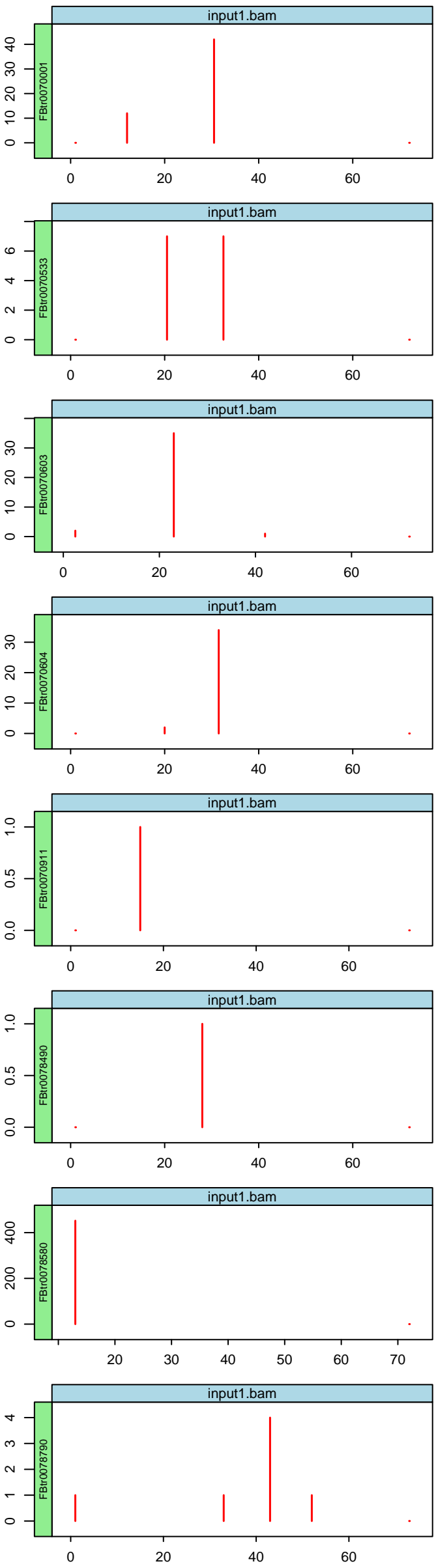
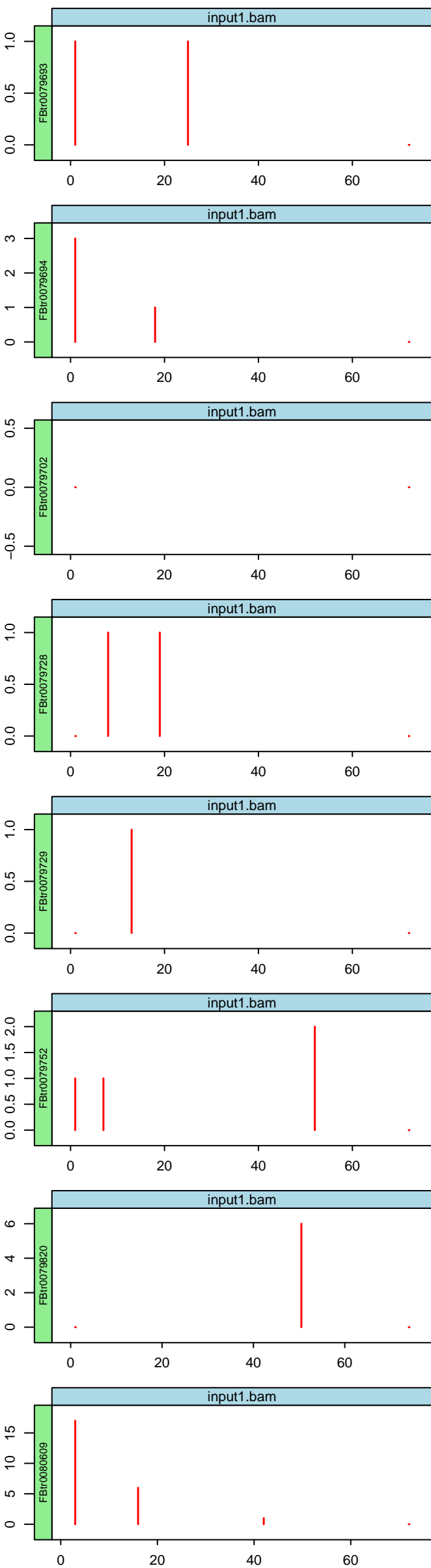


Figure 1 displays eight vertically stacked plots showing the distribution of read lengths for different BAM files. Each plot has a light blue header 'input1.bam' and a green vertical bar on the left with a file name. The y-axis represents frequency, and the x-axis represents read length (0 to 70). Red vertical lines indicate the distribution of read lengths. The files are: FBtr0079064, FBtr0079090, FBtr0079338, FBtr0079528, FBtr0079596, FBtr0079677, FBtr0079690, and FBtr0079692.

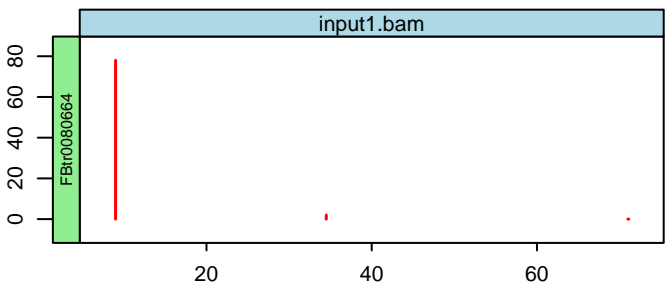
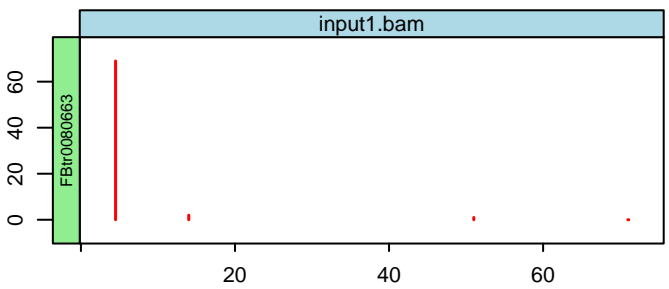
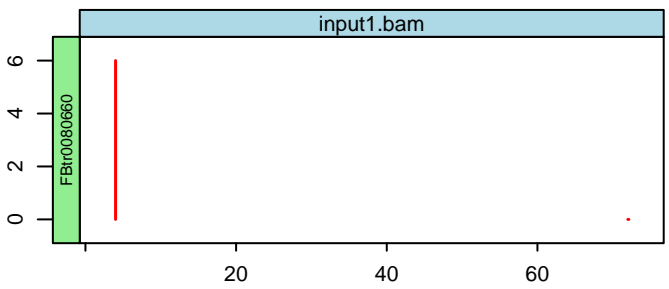
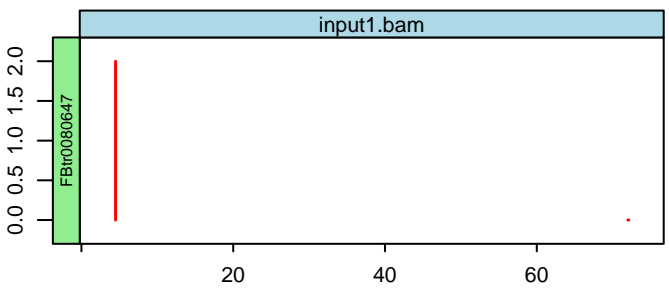
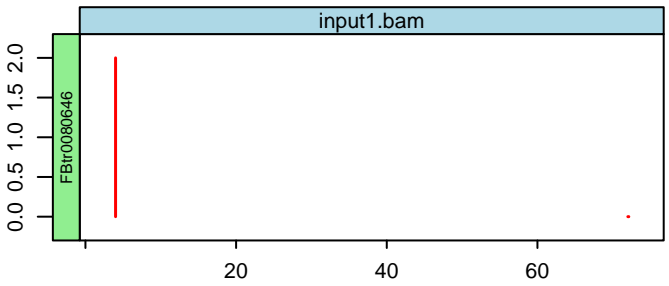
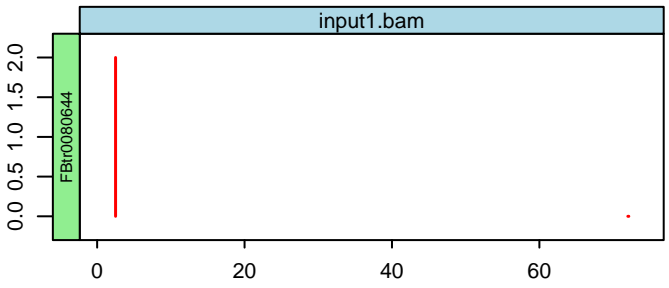
Coordinates (nucleotides)

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Coordinates (nucleotides)