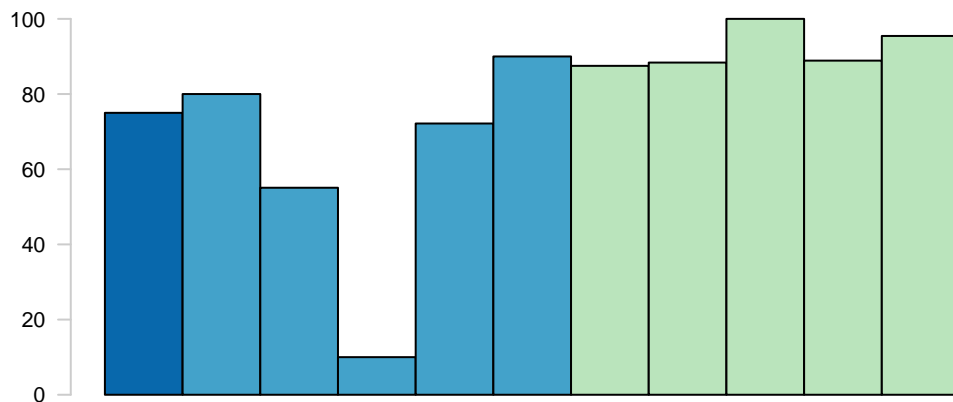
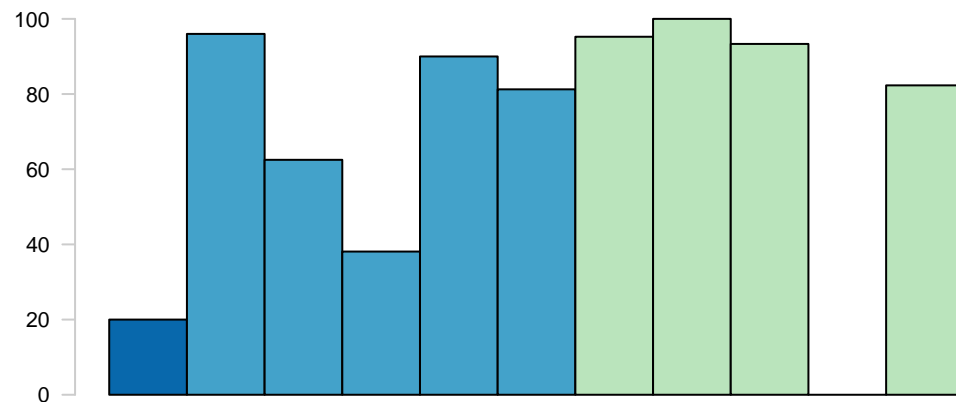


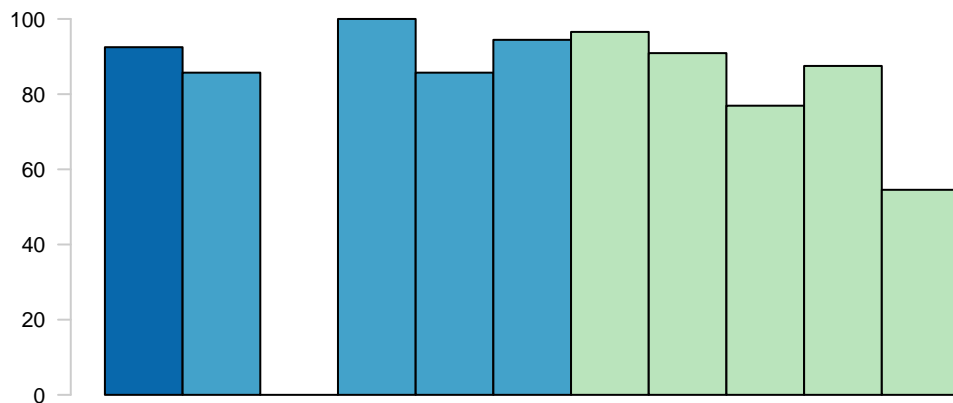
a. Putative promoters



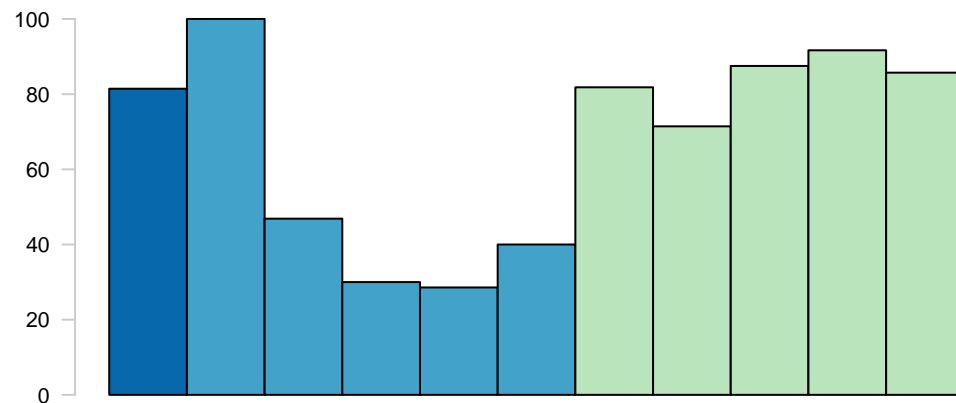
b. UTR



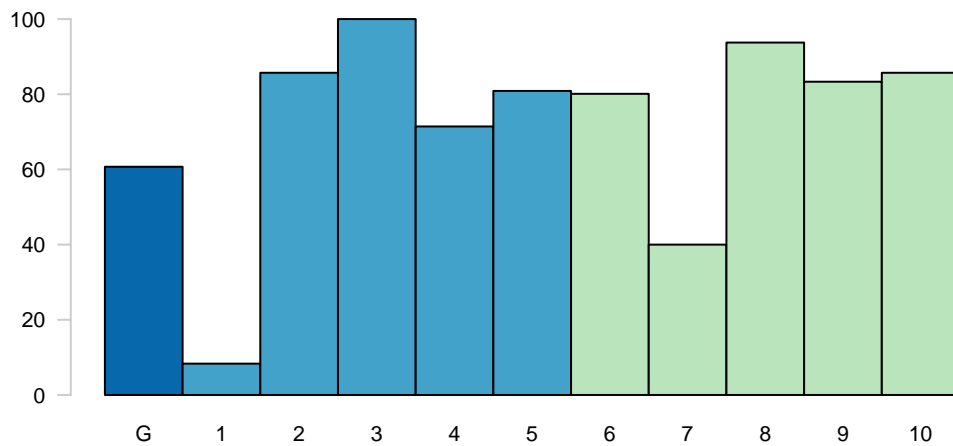
c. Exons



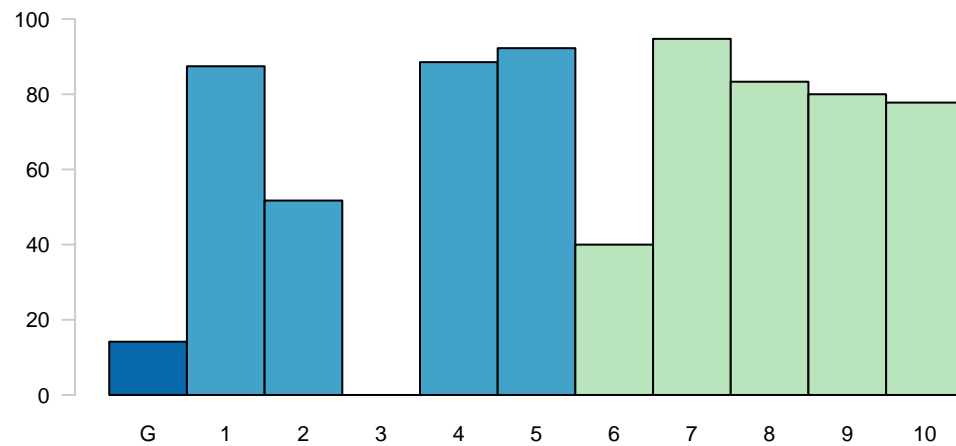
d. Introns



e. Transposable elements



f. Intergenic regions



Methylation (%)

G 1 2 3 4 5 6 7 8 9 10

G 1 2 3 4 5 6 7 8 9 10