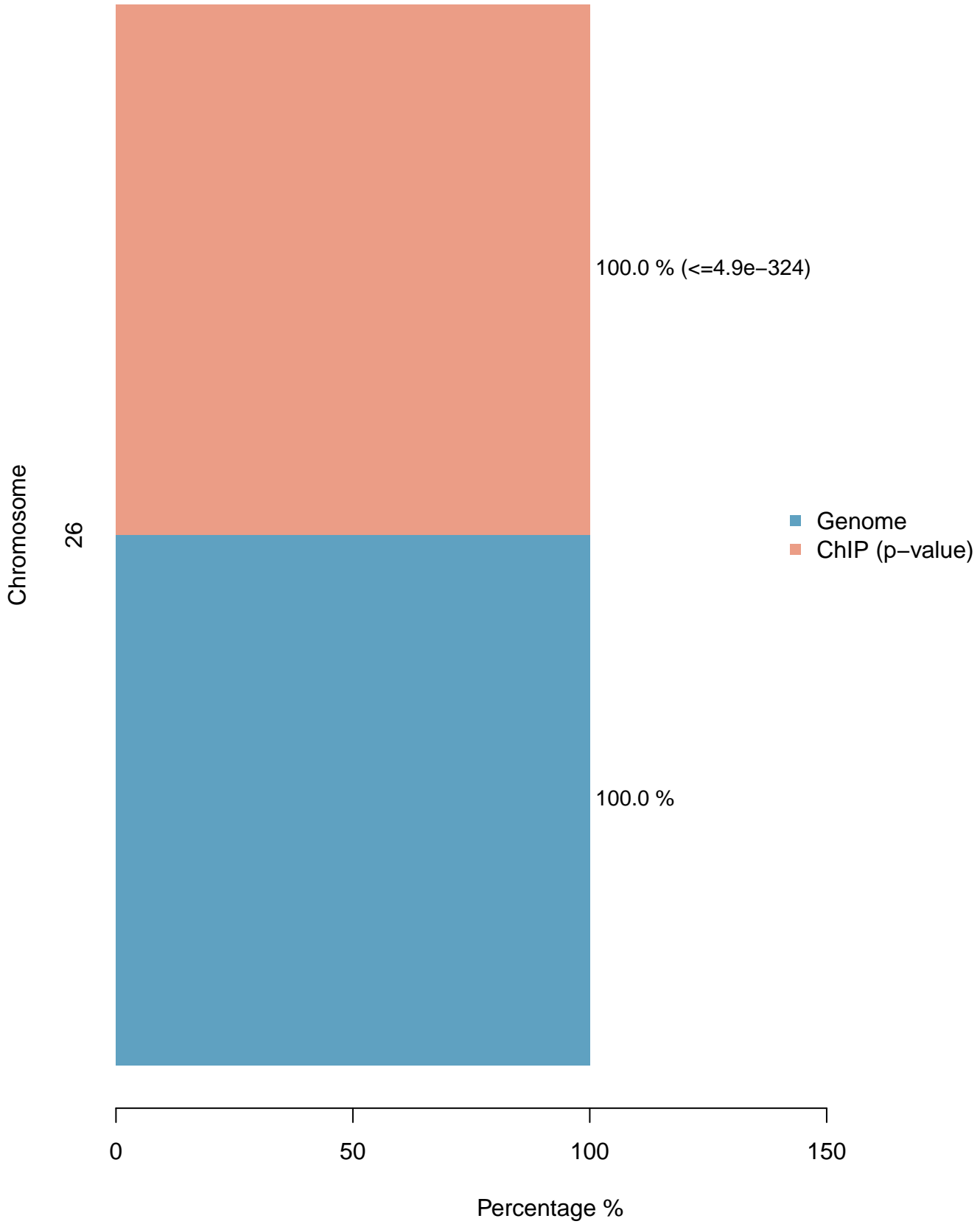
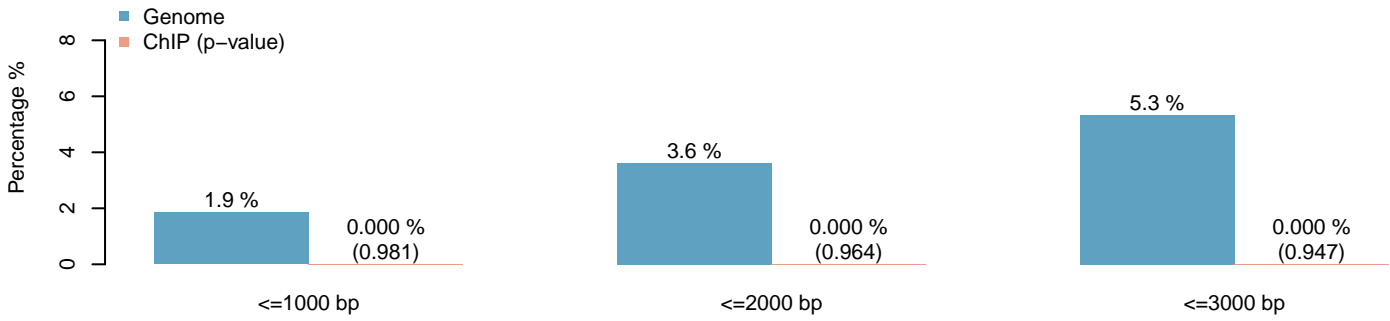


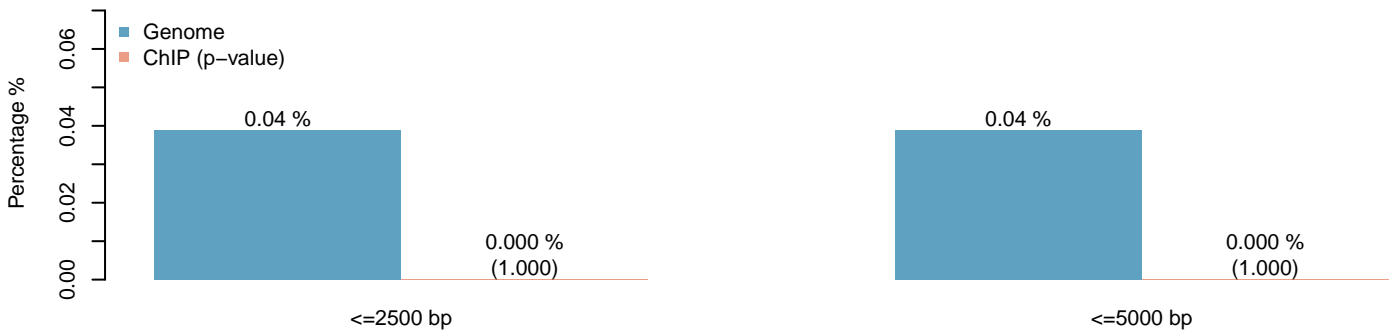
Chromosomal Distribution of ChIP Regions



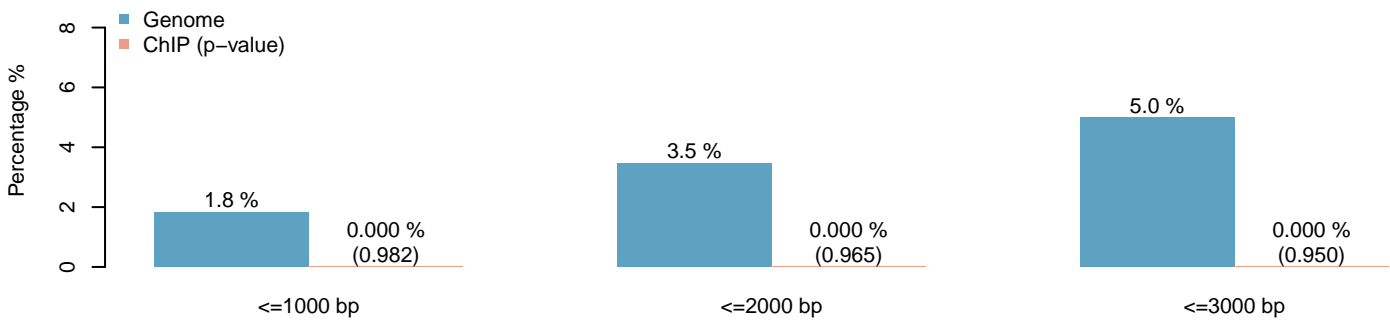
Promoter



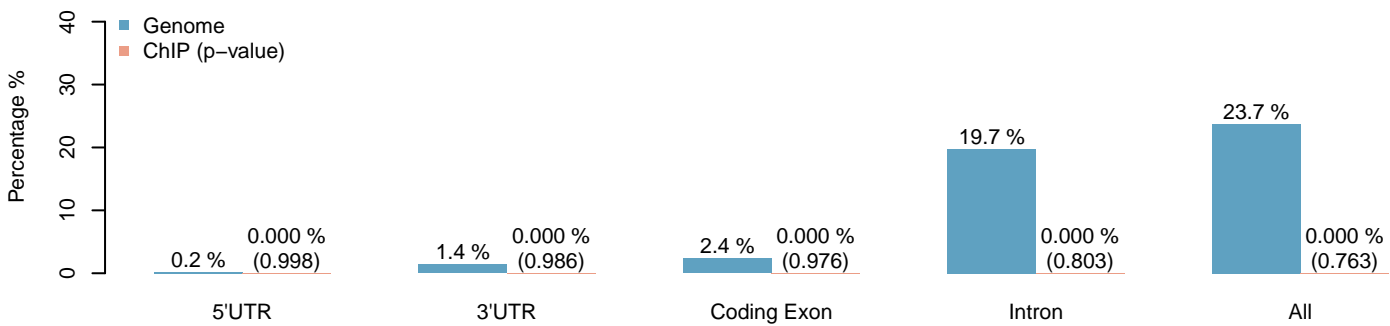
Bidirectional Promoter



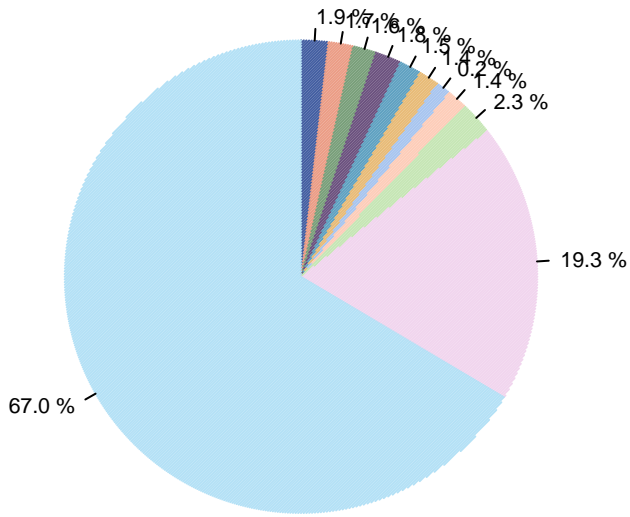
Downstream



Gene

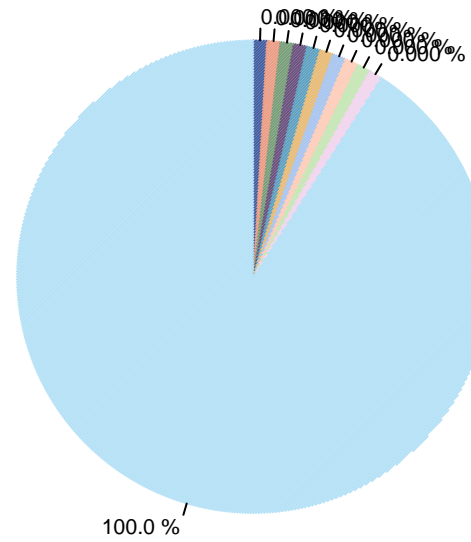


Genome



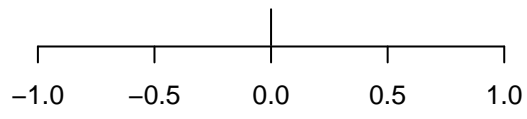
- Promoter (<=1000 bp): 1.9 %
- Promoter (1000–2000 bp): 1.7 %
- Promoter (2000–3000 bp): 1.6 %
- Downstream (<=1000 bp): 1.8 %
- Downstream (1000–2000 bp): 1.5 %
- Downstream (2000–3000 bp): 1.4 %
- 5'UTR: 0.2 %
- 3'UTR: 1.4 %
- Coding exon: 2.3 %
- Intron: 19.3 %
- Distal intergenic: 67.0 %

ChIP



- Promoter (<=1000 bp): 0.000 %
- Promoter (1000–2000 bp): 0.000 %
- Promoter (2000–3000 bp): 0.000 %
- Downstream (<=1000 bp): 0.000 %
- Downstream (1000–2000 bp): 0.000 %
- Downstream (2000–3000 bp): 0.000 %
- 5'UTR: 0.000 %
- 3'UTR: 0.000 %
- Coding exon: 0.000 %
- Intron: 0.000 %
- Distal intergenic: 100.0 %

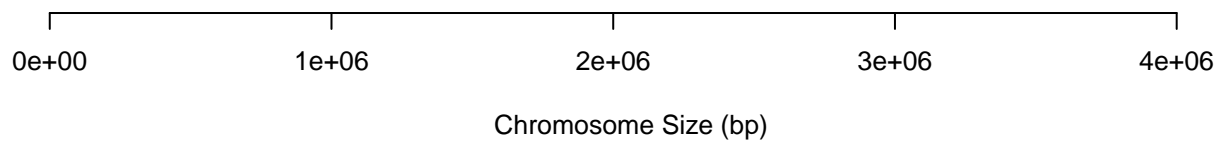
Distribution of Peak Heights



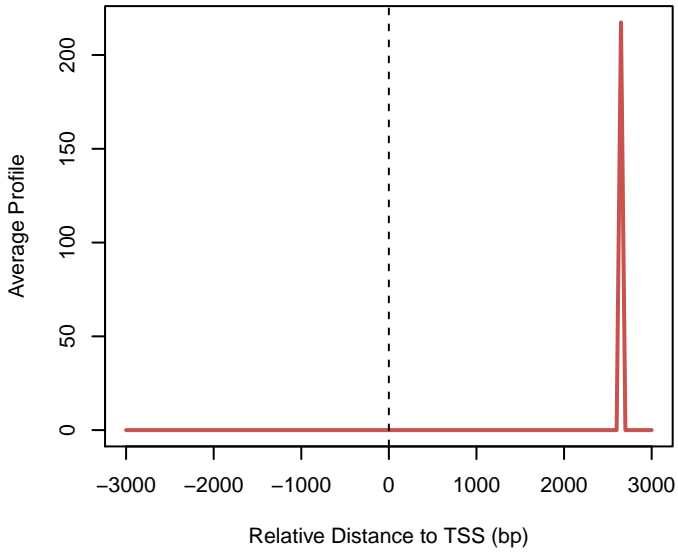
ChIP Regions (Peaks) over Chromosomes

Chromosome

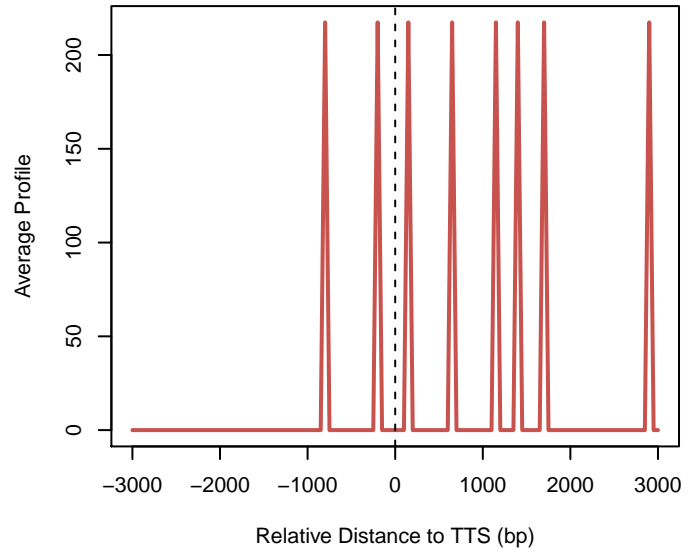
26



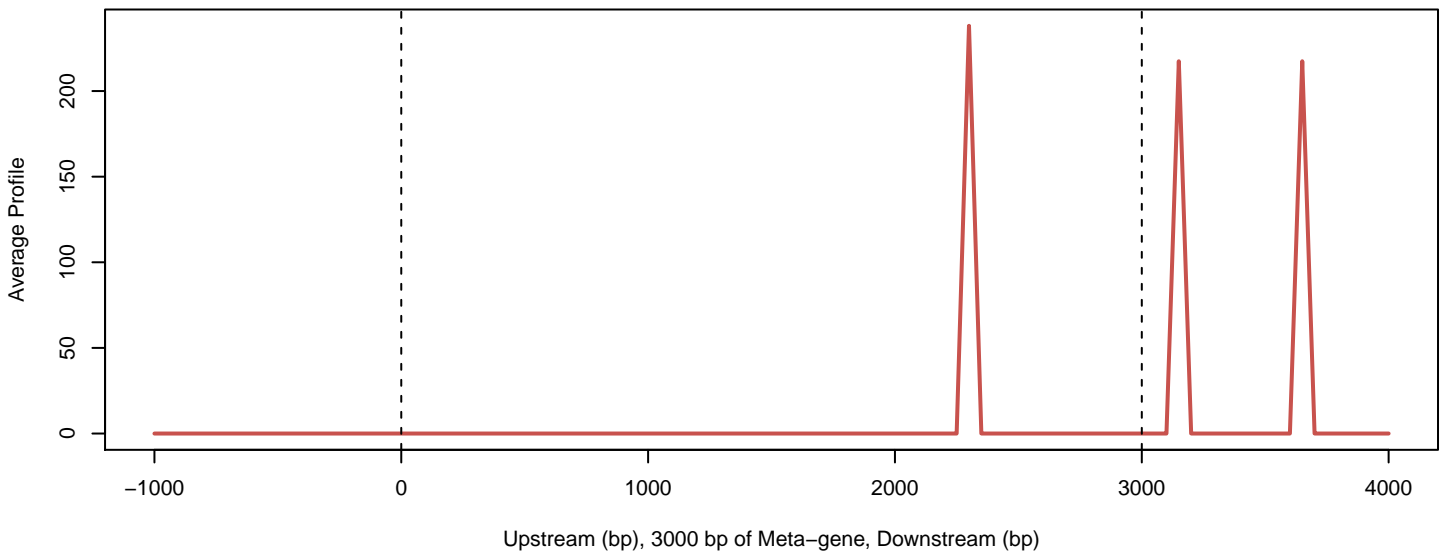
Average Profile near TSS



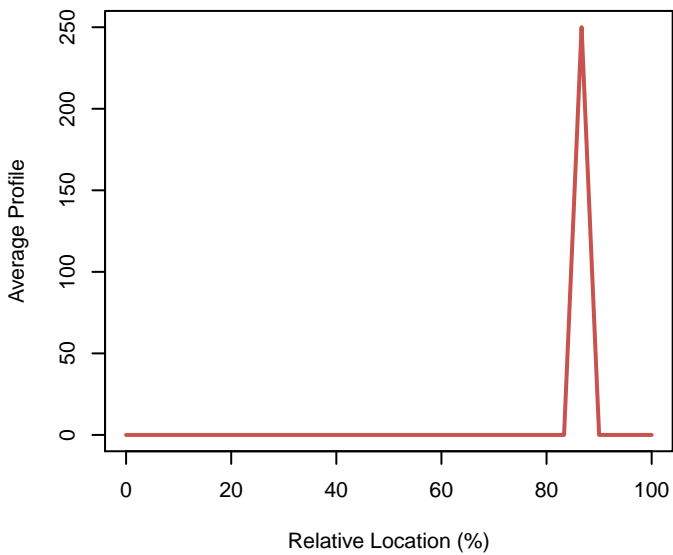
Average Profile near TTS



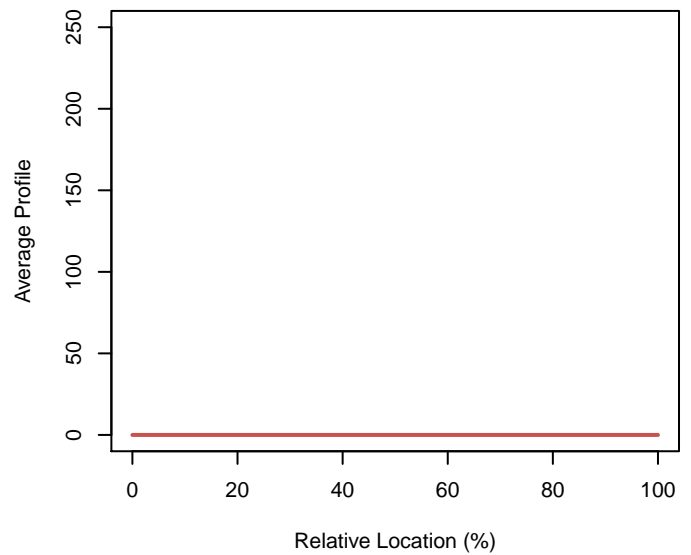
Average Gene Profile



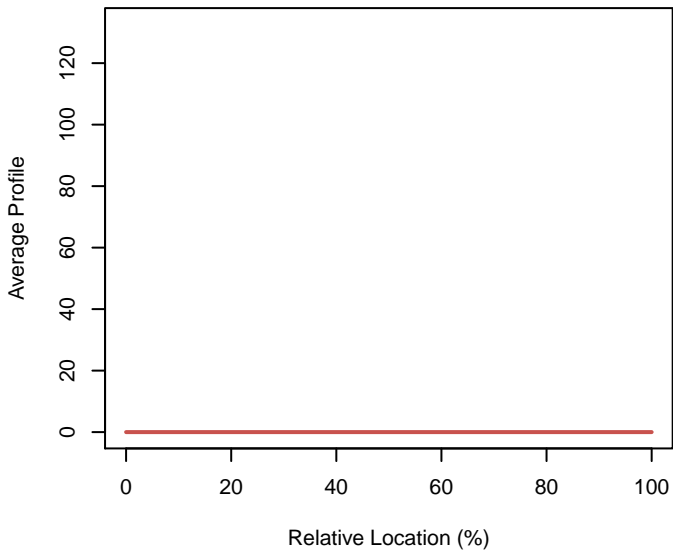
Average Concatenated Exon Profile



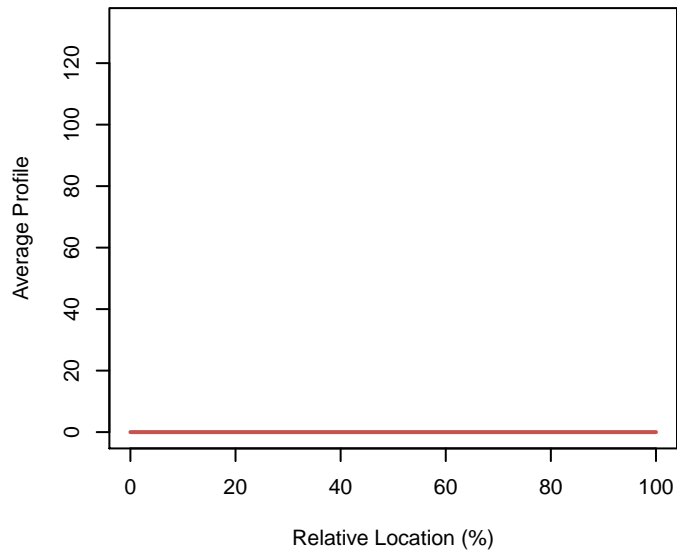
Average Concatenated Intron Profile



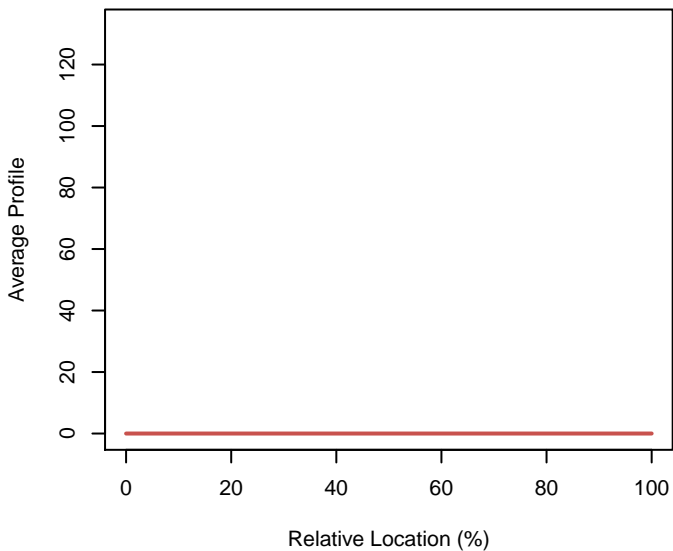
Average Exon Profile
(56 <= length < 109 bp)



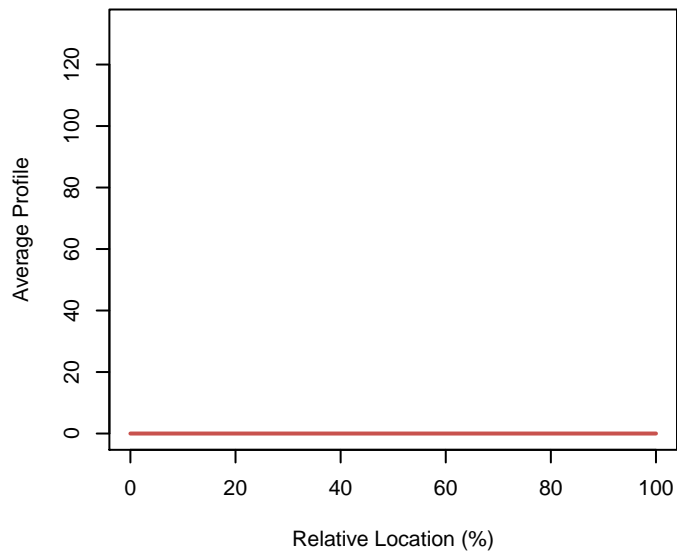
Average Intron Profile
(110 <= length < 345 bp)



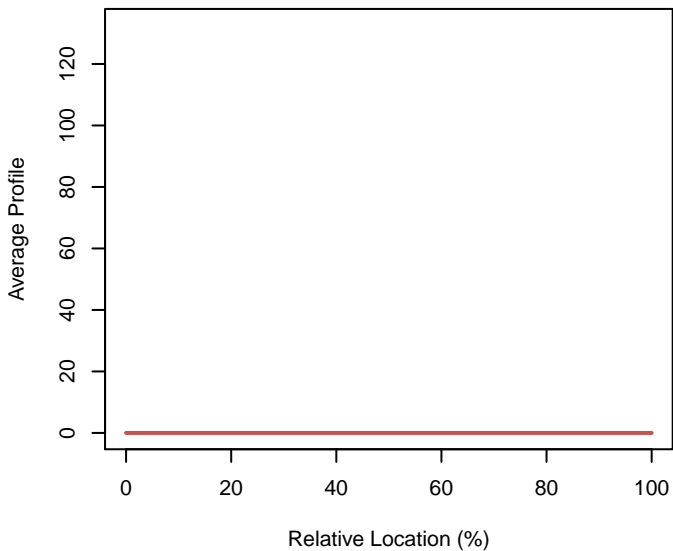
Average Exon Profile
(109 <= length < 160 bp)



Average Intron Profile
(344 <= length < 686 bp)



Average Exon Profile
(160 <= length < 375 bp)



Average Intron Profile
(685 <= length < 2653 bp)

