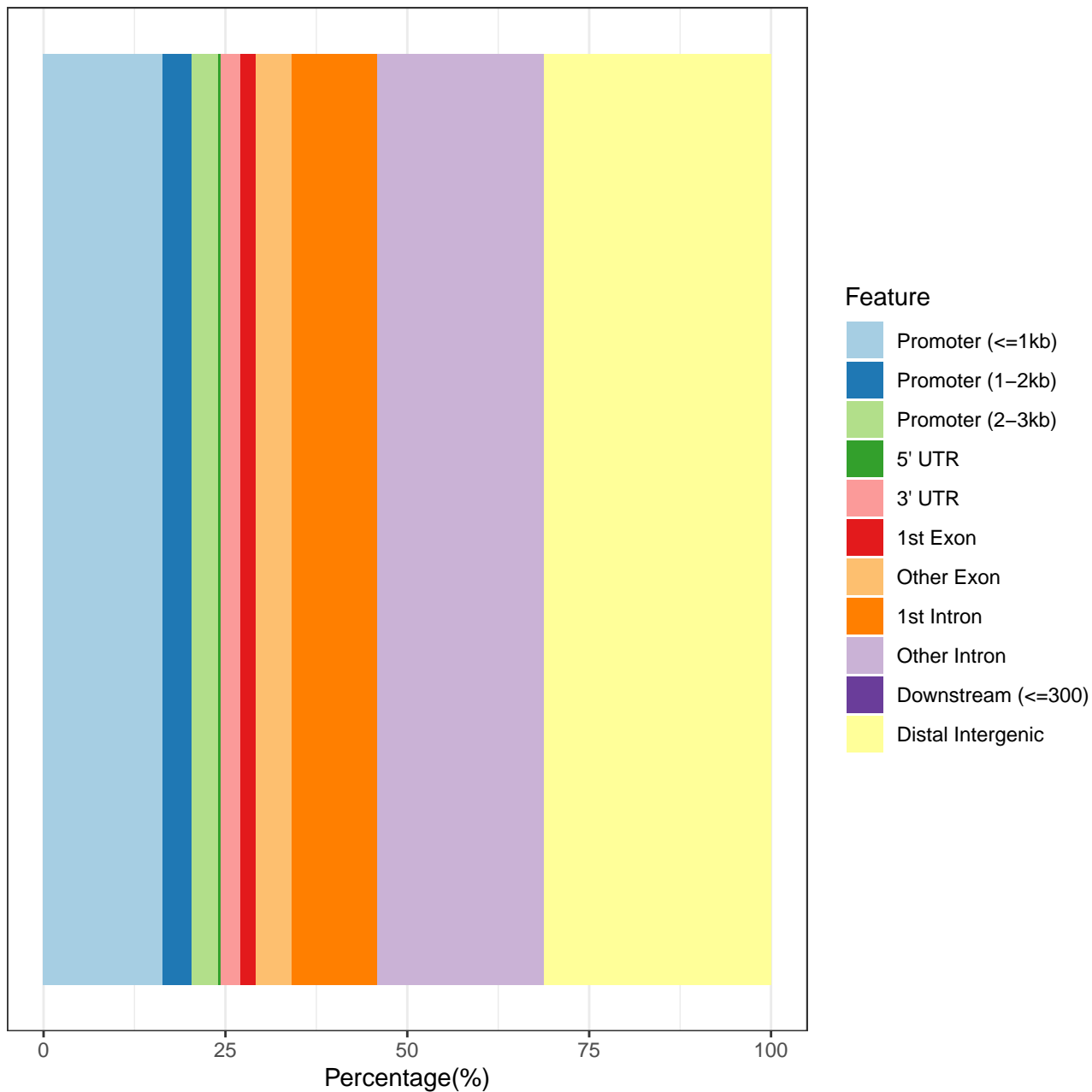
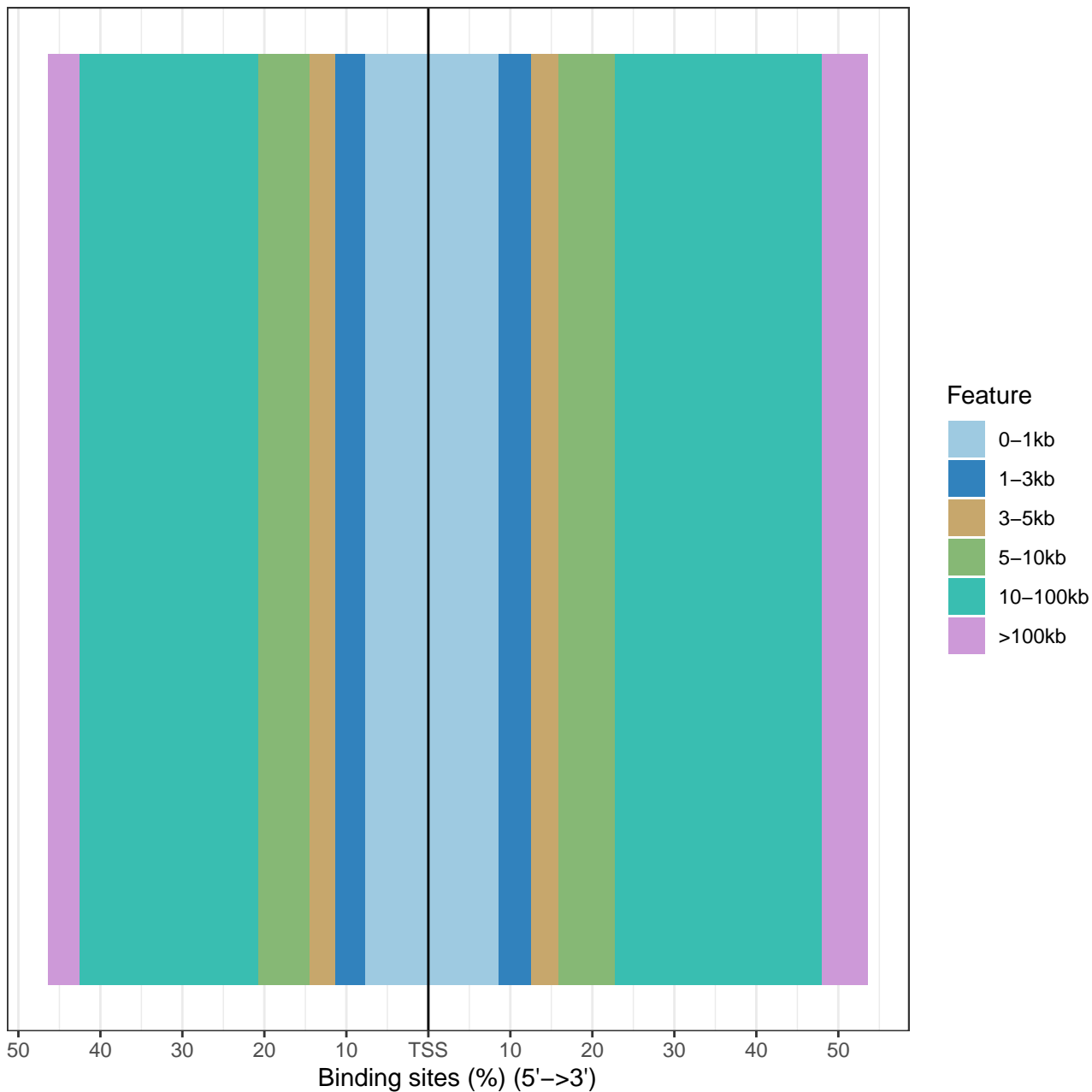
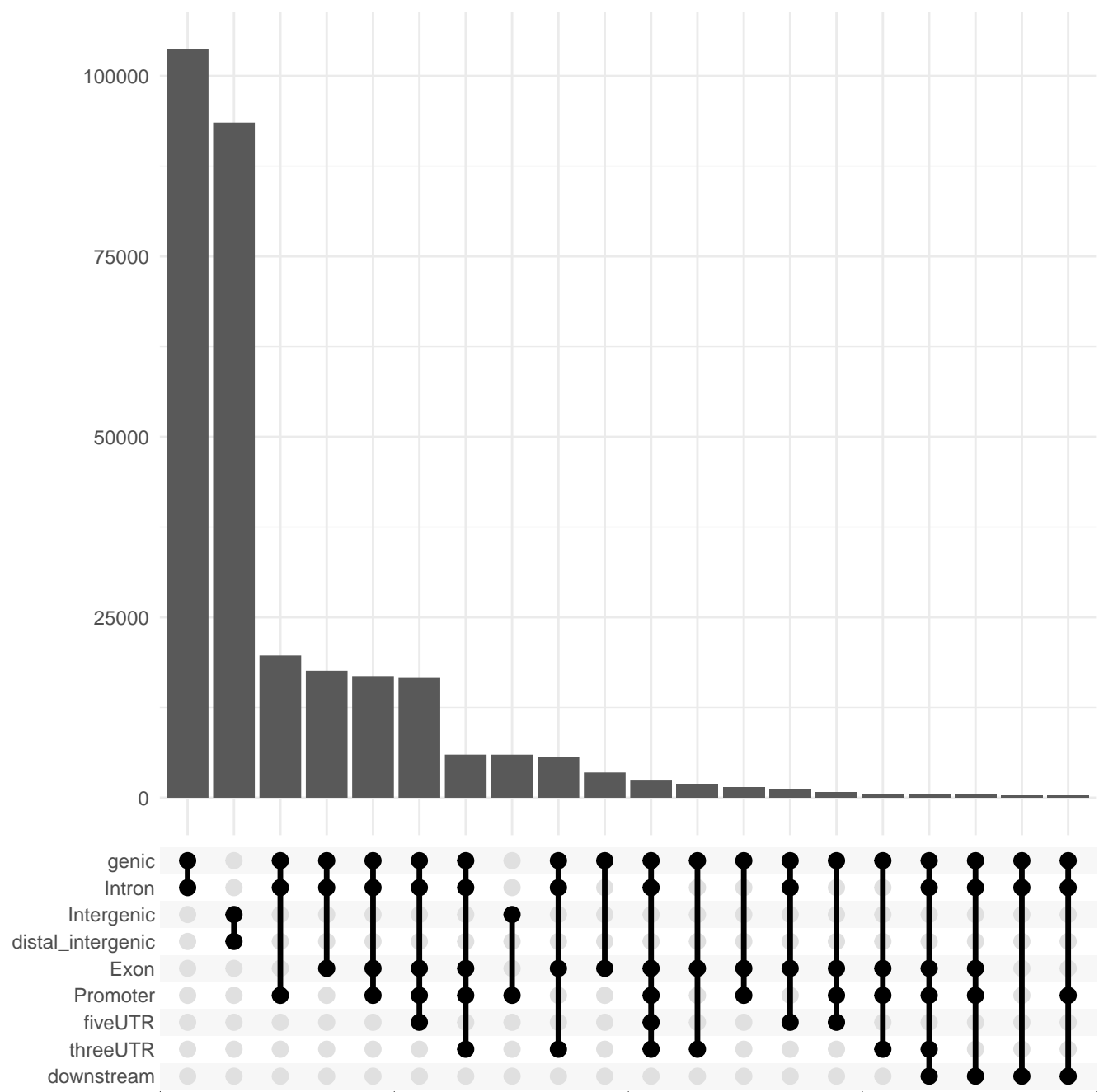


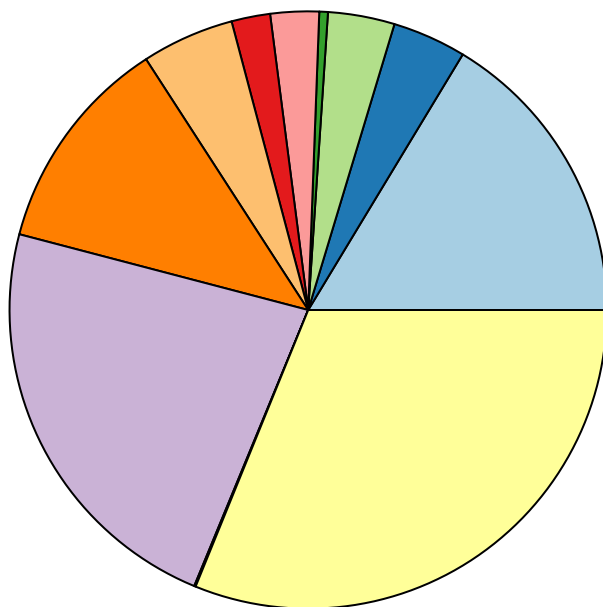
# Feature Distribution



Distribution of transcription factor–binding loci relative to TSS







- Promoter (<=1kb) (16.34%)
- Promoter (1-2kb) (3.99%)
- Promoter (2-3kb) (3.61%)
- 5' UTR (0.46%)
- 3' UTR (2.63%)
- 1st Exon (2.09%)
- Other Exon (5.01%)
- 1st Intron (11.78%)
- Other Intron (22.85%)
- Downstream (<=300) (0.08%)
- Distal Intergenic (31.15%)