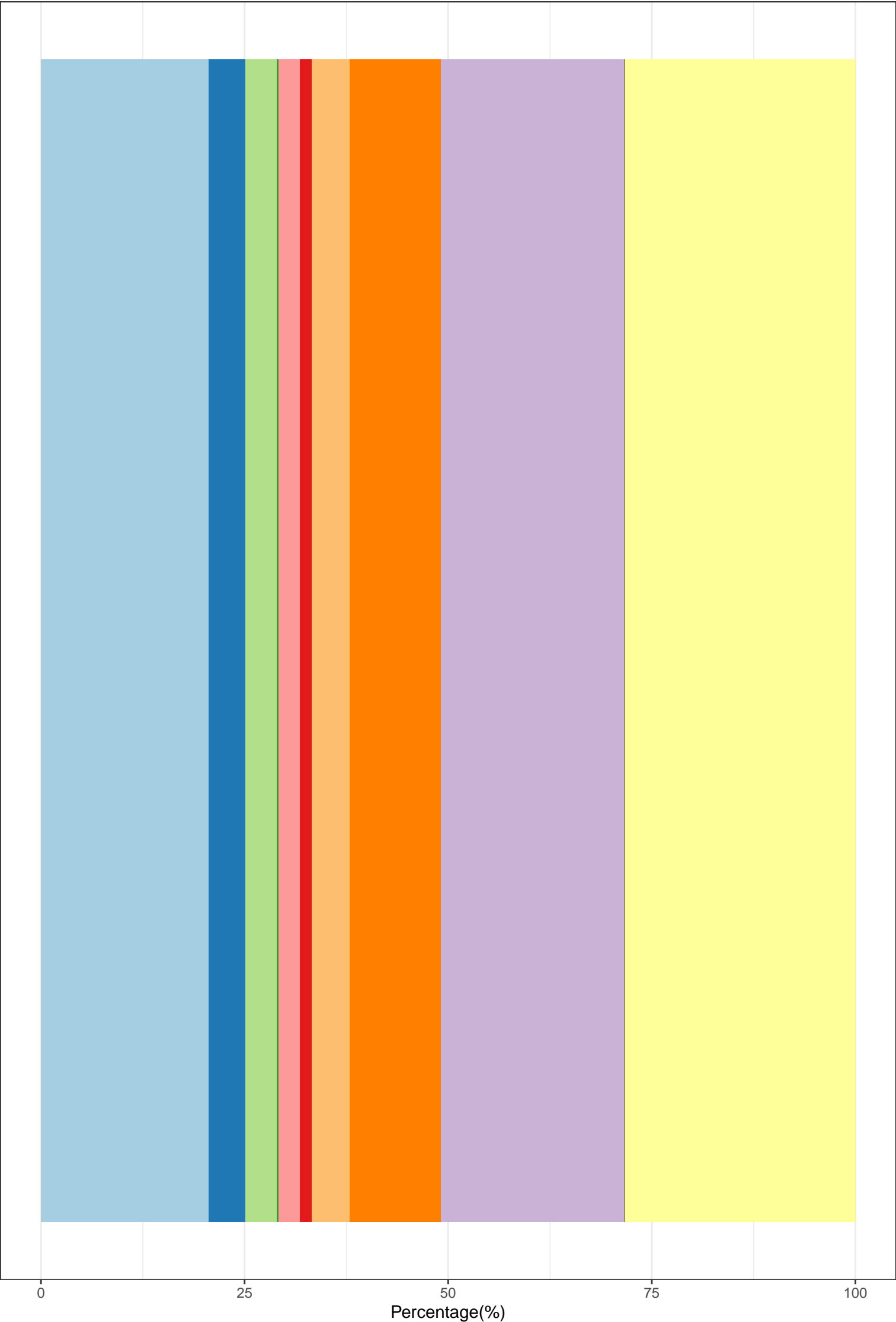
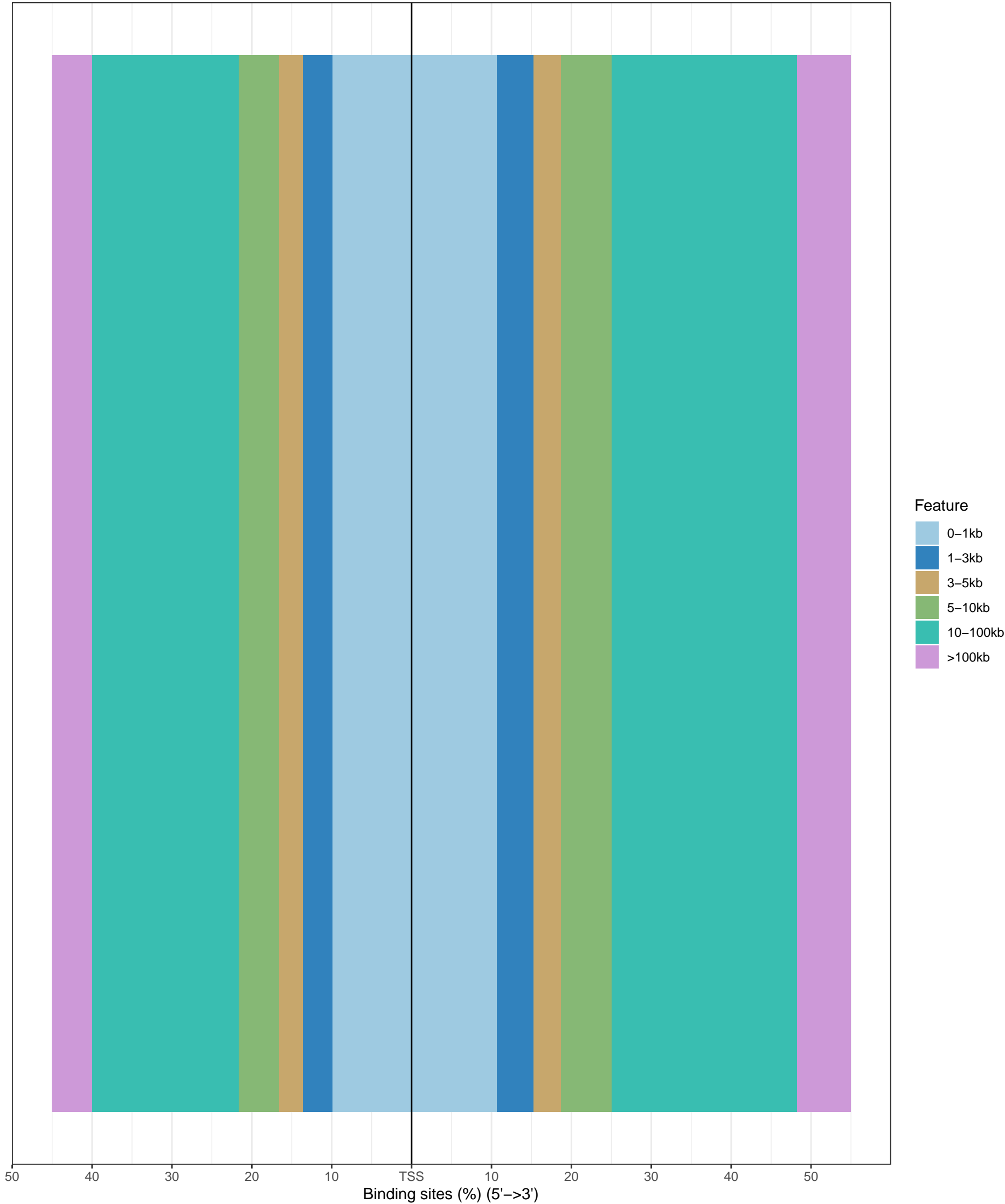
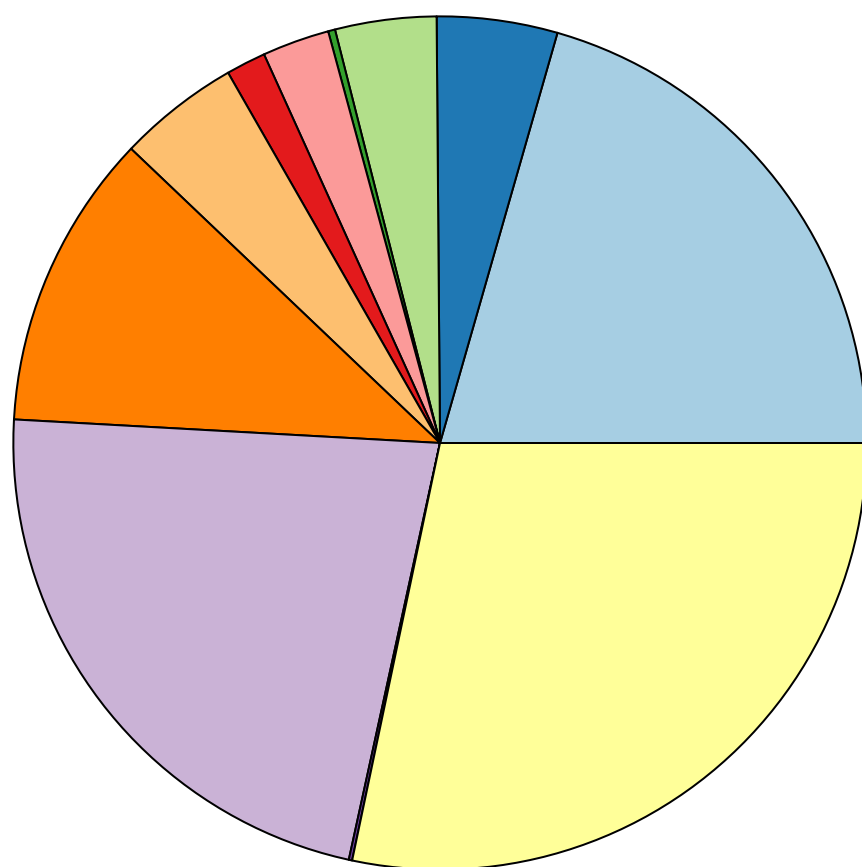


Feature Distribution



Distribution of transcription factor-binding loci relative to TSS





- Promoter (<=1kb) (20.56%)
- Promoter (1-2kb) (4.57%)
- Promoter (2-3kb) (3.83%)
- 5' UTR (0.26%)
- 3' UTR (2.55%)
- 1st Exon (1.51%)
- Other Exon (4.61%)
- 1st Intron (11.23%)
- Other Intron (22.45%)
- Downstream (<=300) (0.13%)
- Distal Intergenic (28.3%)