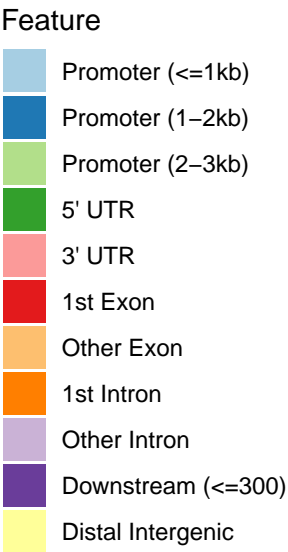
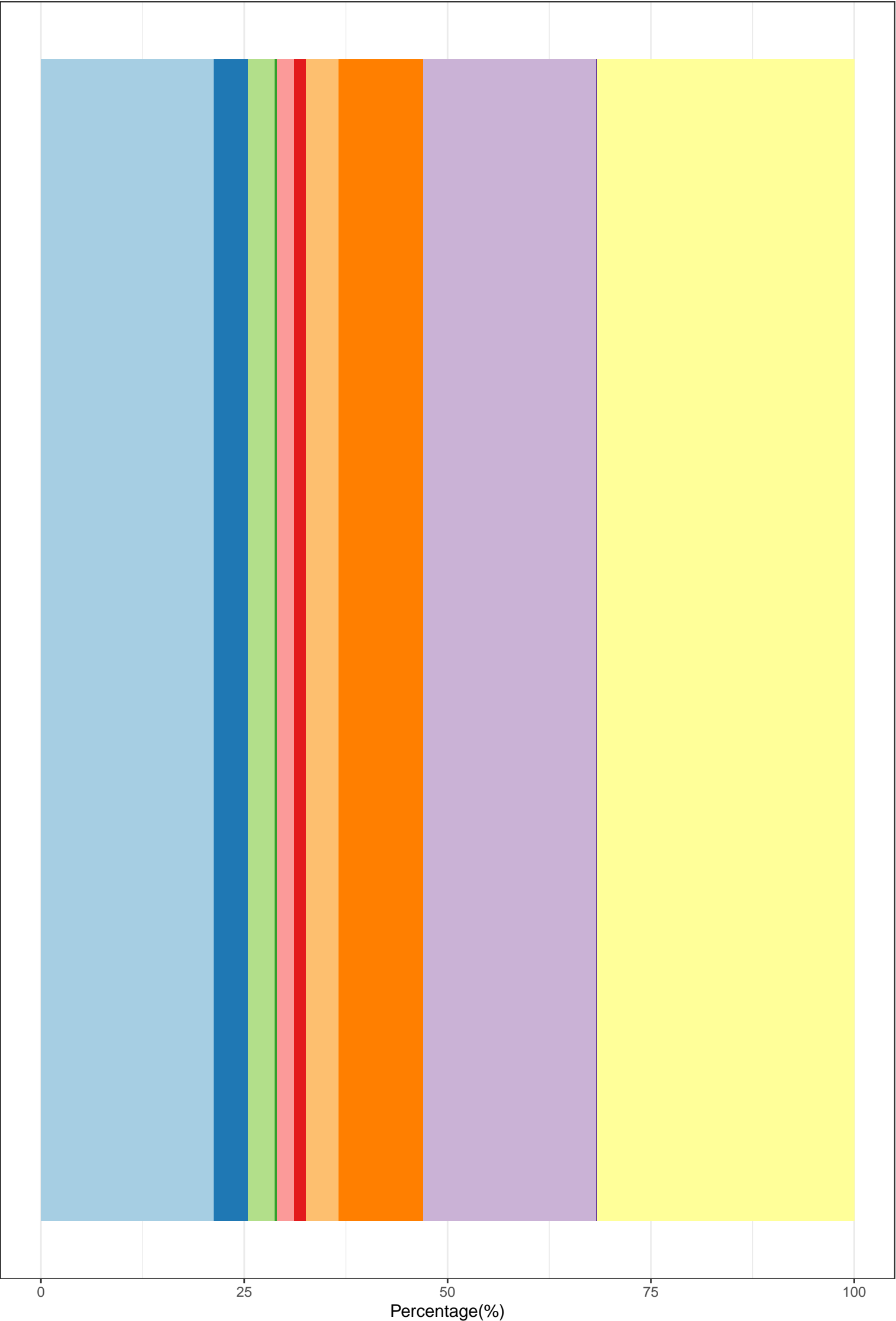
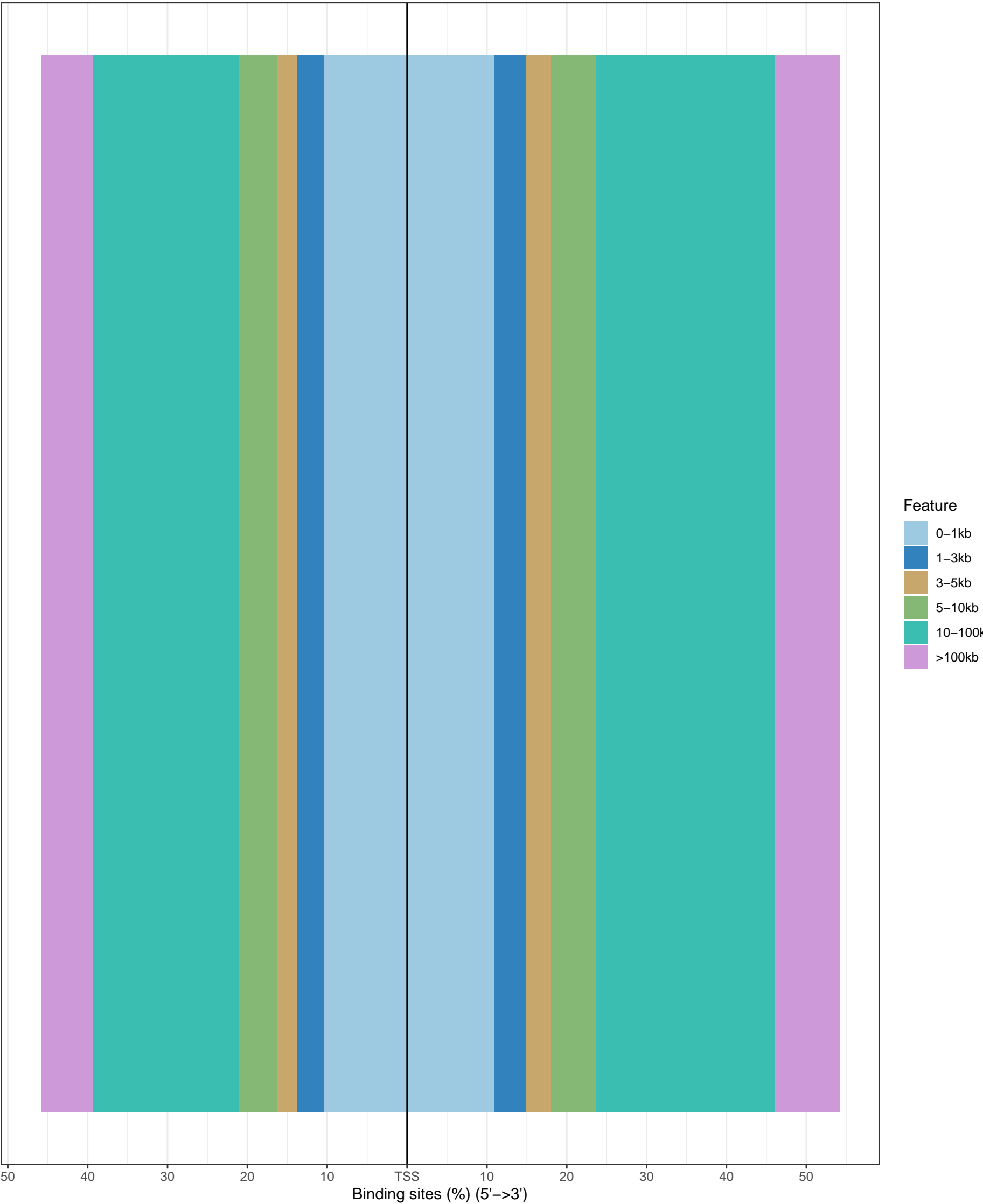
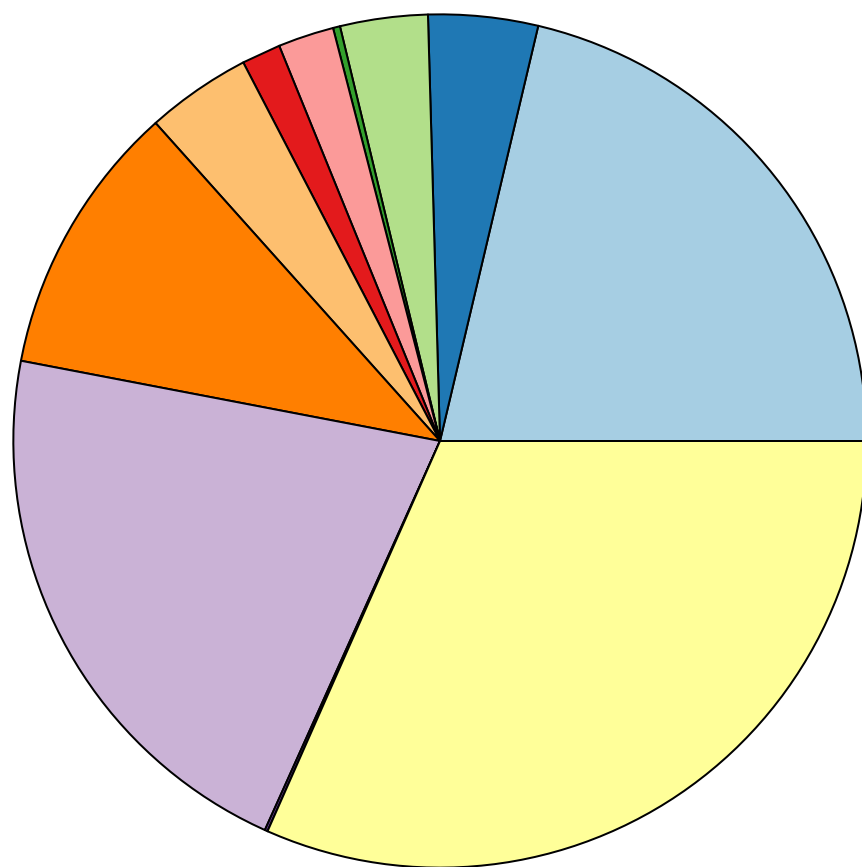


Feature Distribution



Distribution of transcription factor-binding loci relative to TSS





- Promoter (<=1kb) (21.3%)
- Promoter (1-2kb) (4.15%)
- Promoter (2-3kb) (3.32%)
- 5' UTR (0.27%)
- 3' UTR (2.12%)
- 1st Exon (1.48%)
- Other Exon (3.99%)
- 1st Intron (10.37%)
- Other Intron (21.28%)
- Downstream (<=300) (0.11%)
- Distal Intergenic (31.62%)