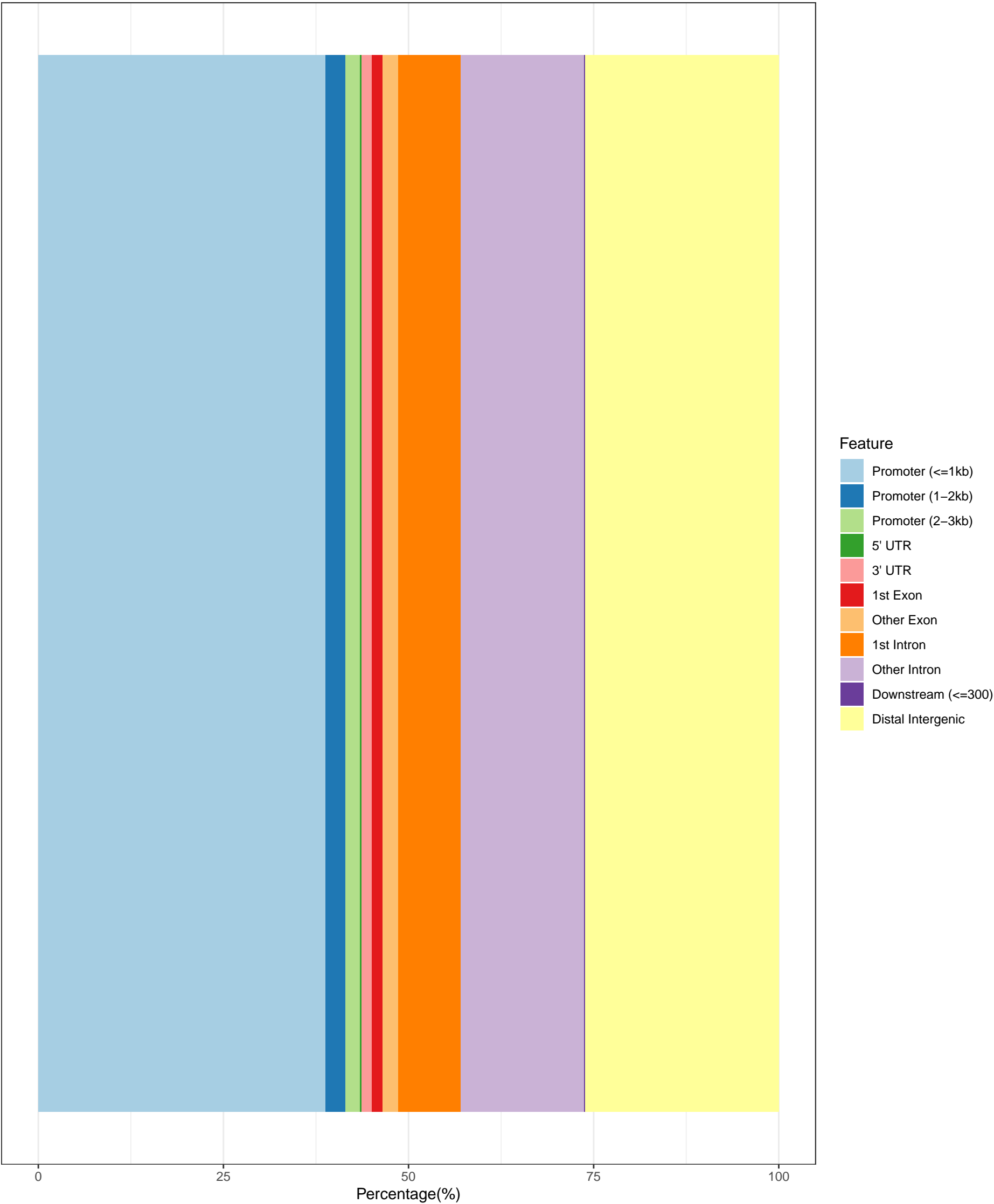
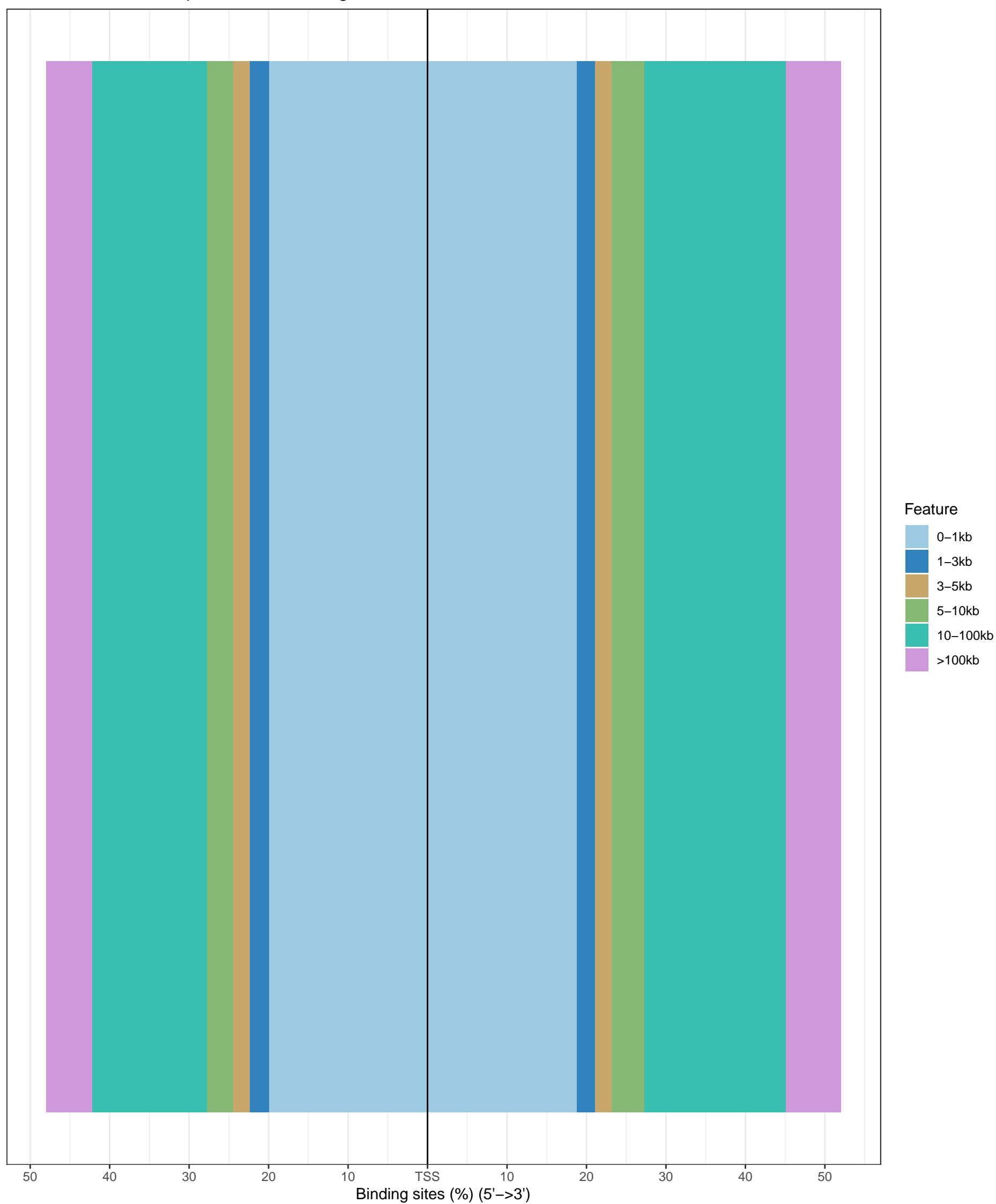
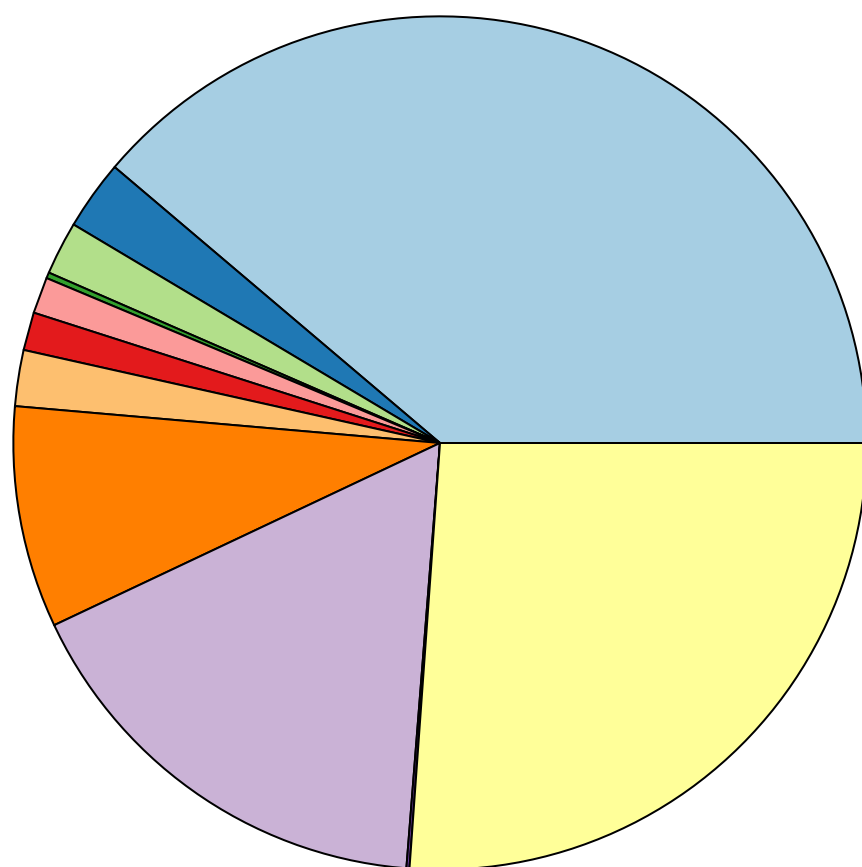


Feature Distribution



Distribution of transcription factor-binding loci relative to TSS





- Promoter (<=1kb) (38.81%)
- Promoter (1-2kb) (2.64%)
- Promoter (2-3kb) (2.02%)
- 5' UTR (0.22%)
- 3' UTR (1.38%)
- 1st Exon (1.44%)
- Other Exon (2.12%)
- 1st Intron (8.4%)
- Other Intron (16.72%)
- Downstream (<=300) (0.12%)
- Distal Intergenic (26.14%)