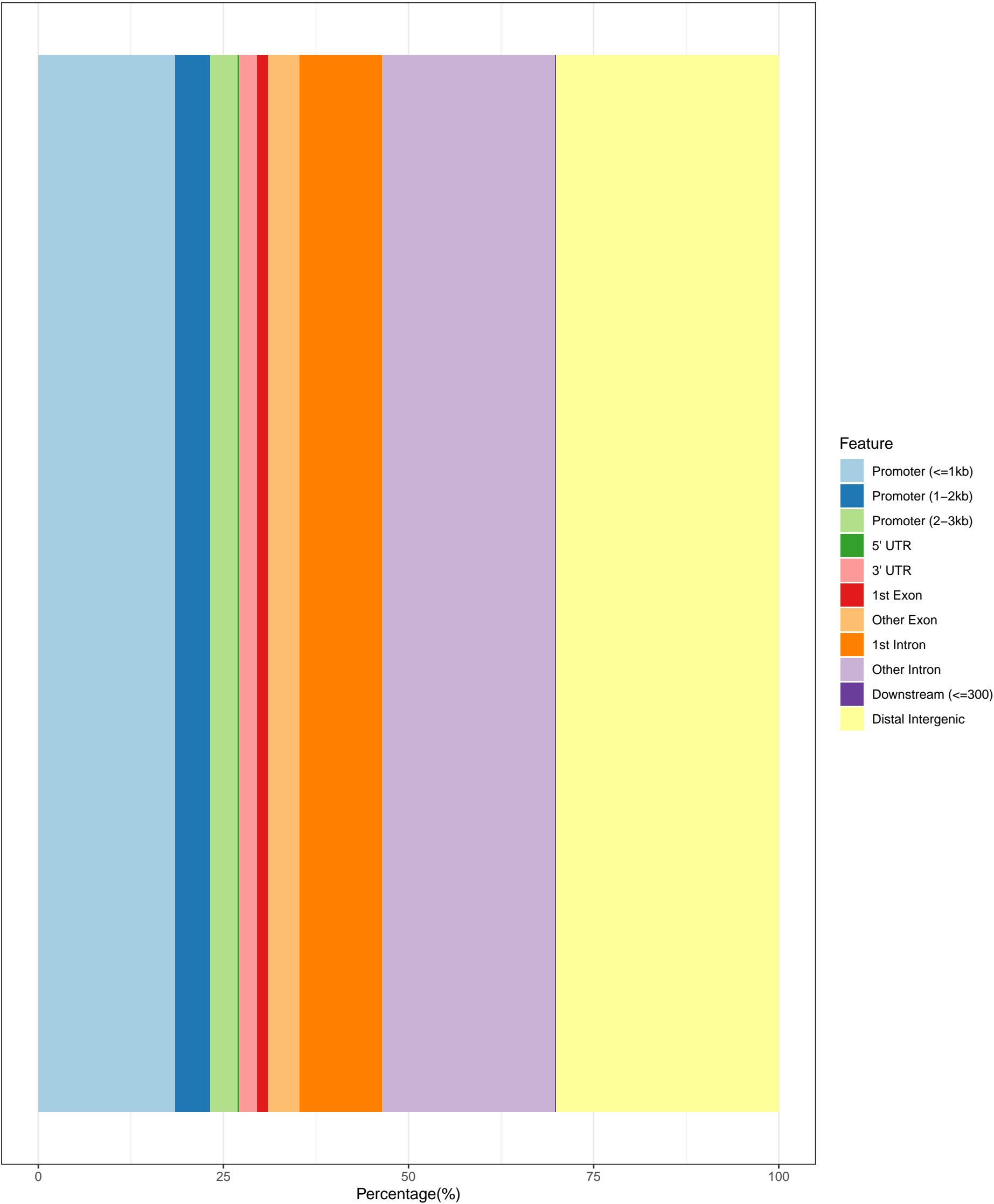
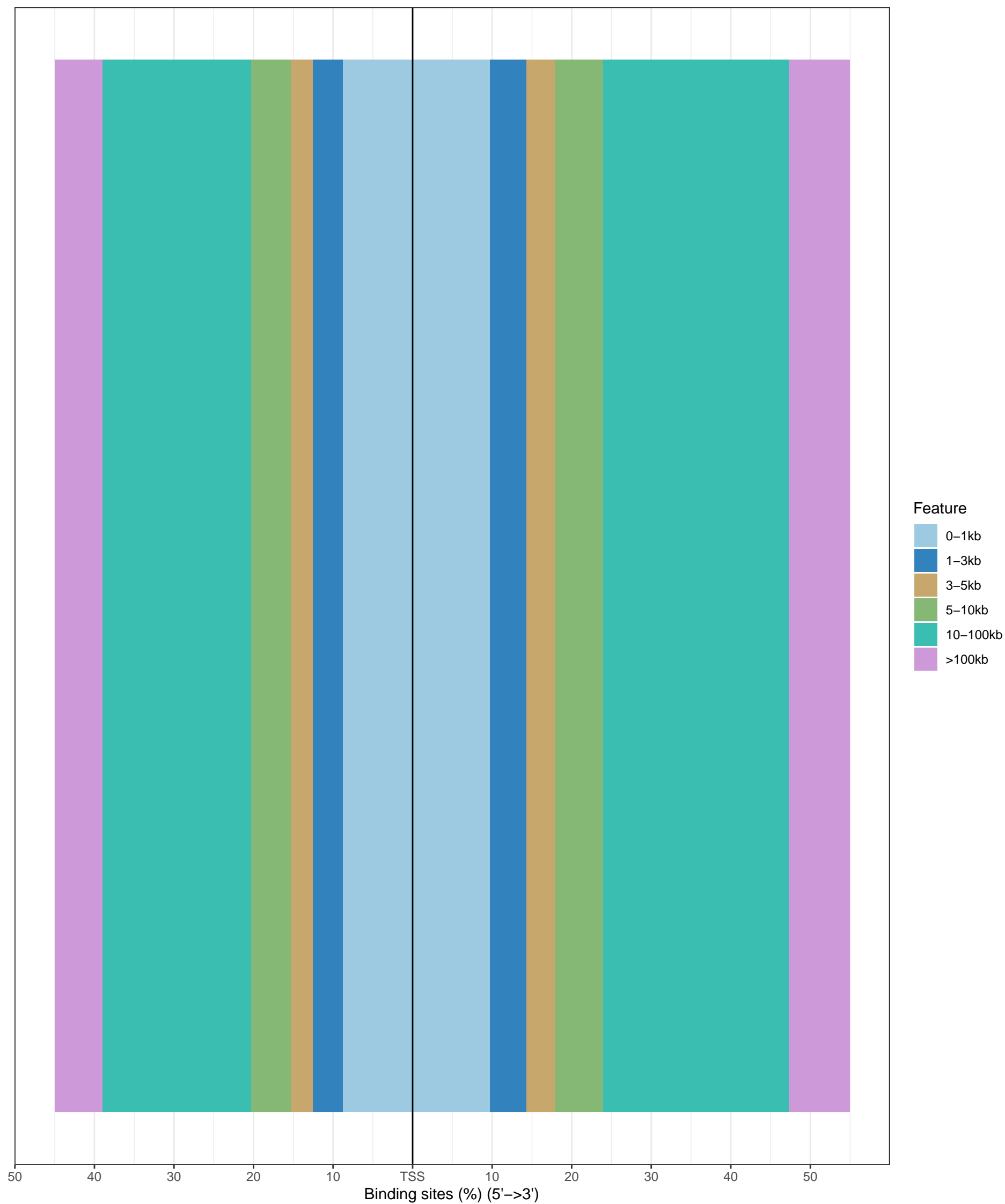
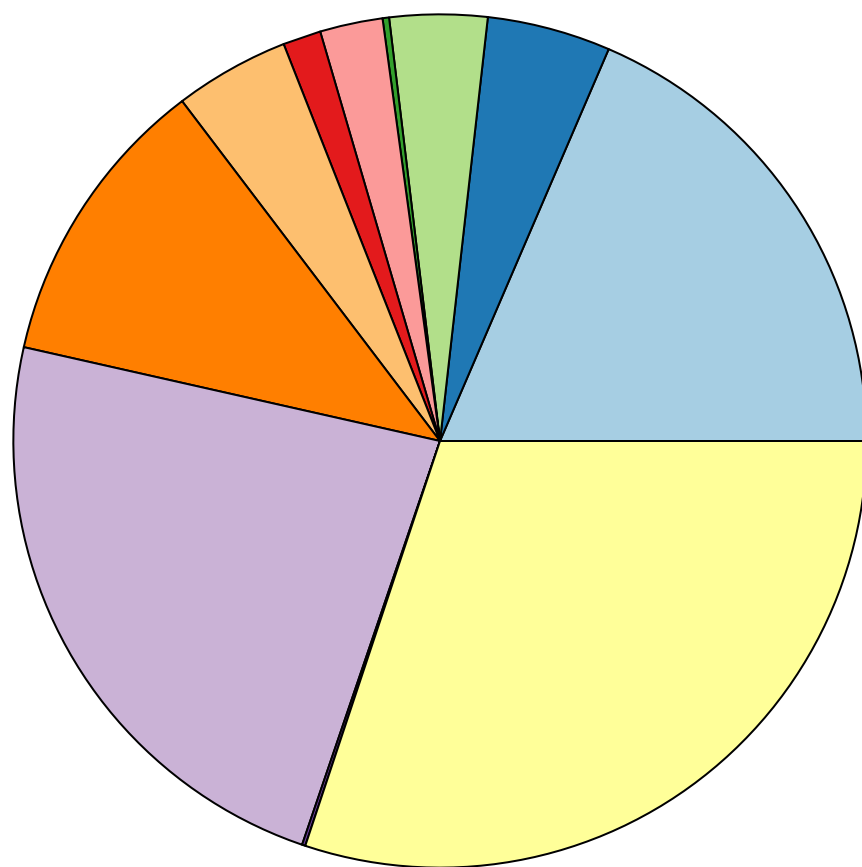


Feature Distribution



Distribution of transcription factor-binding loci relative to TSS





- Promoter (<=1kb) (18.53%)
- Promoter (1-2kb) (4.68%)
- Promoter (2-3kb) (3.7%)
- 5' UTR (0.25%)
- 3' UTR (2.38%)
- 1st Exon (1.44%)
- Other Exon (4.36%)
- 1st Intron (11.14%)
- Other Intron (23.29%)
- Downstream (<=300) (0.12%)
- Distal Intergenic (30.11%)