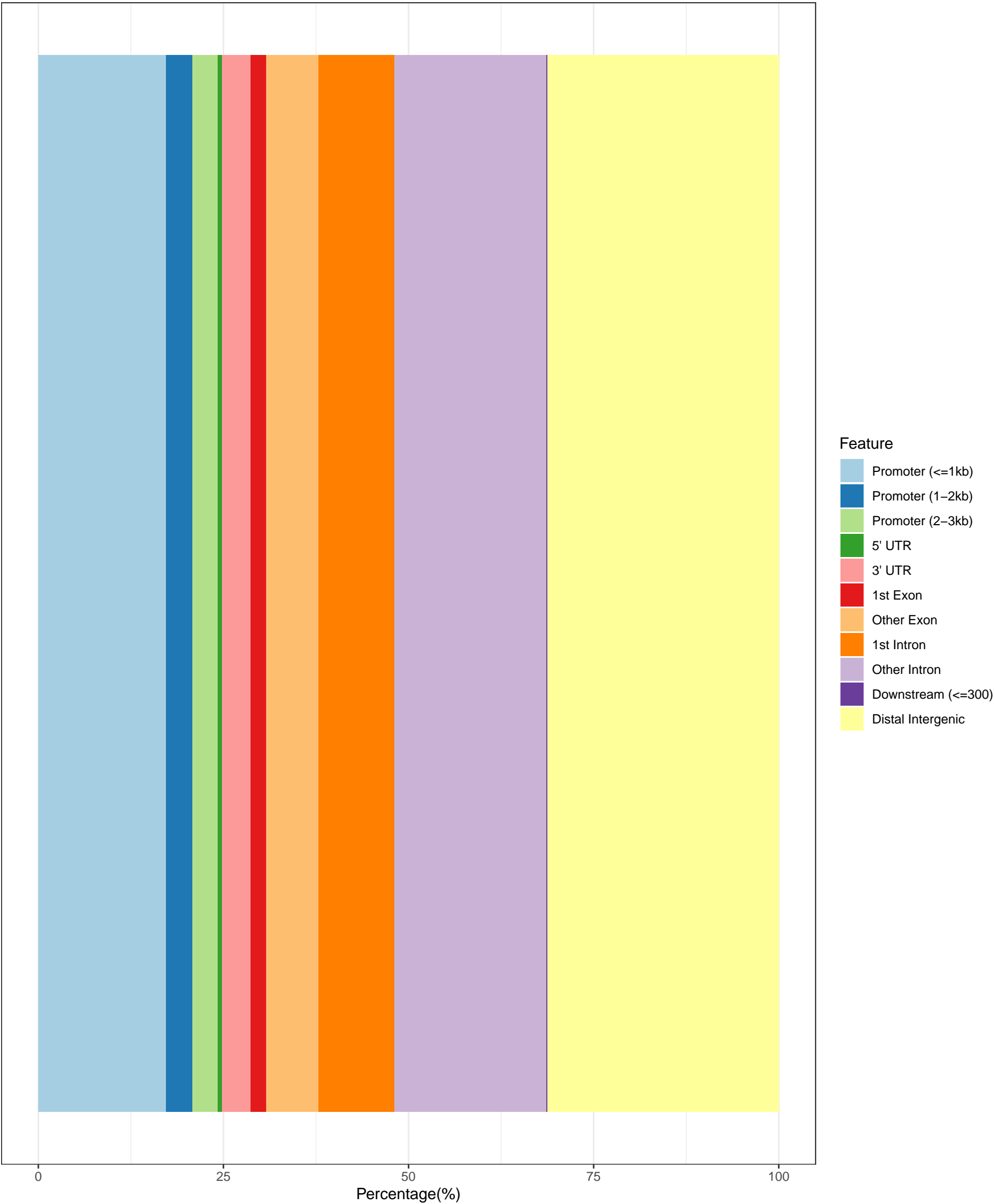
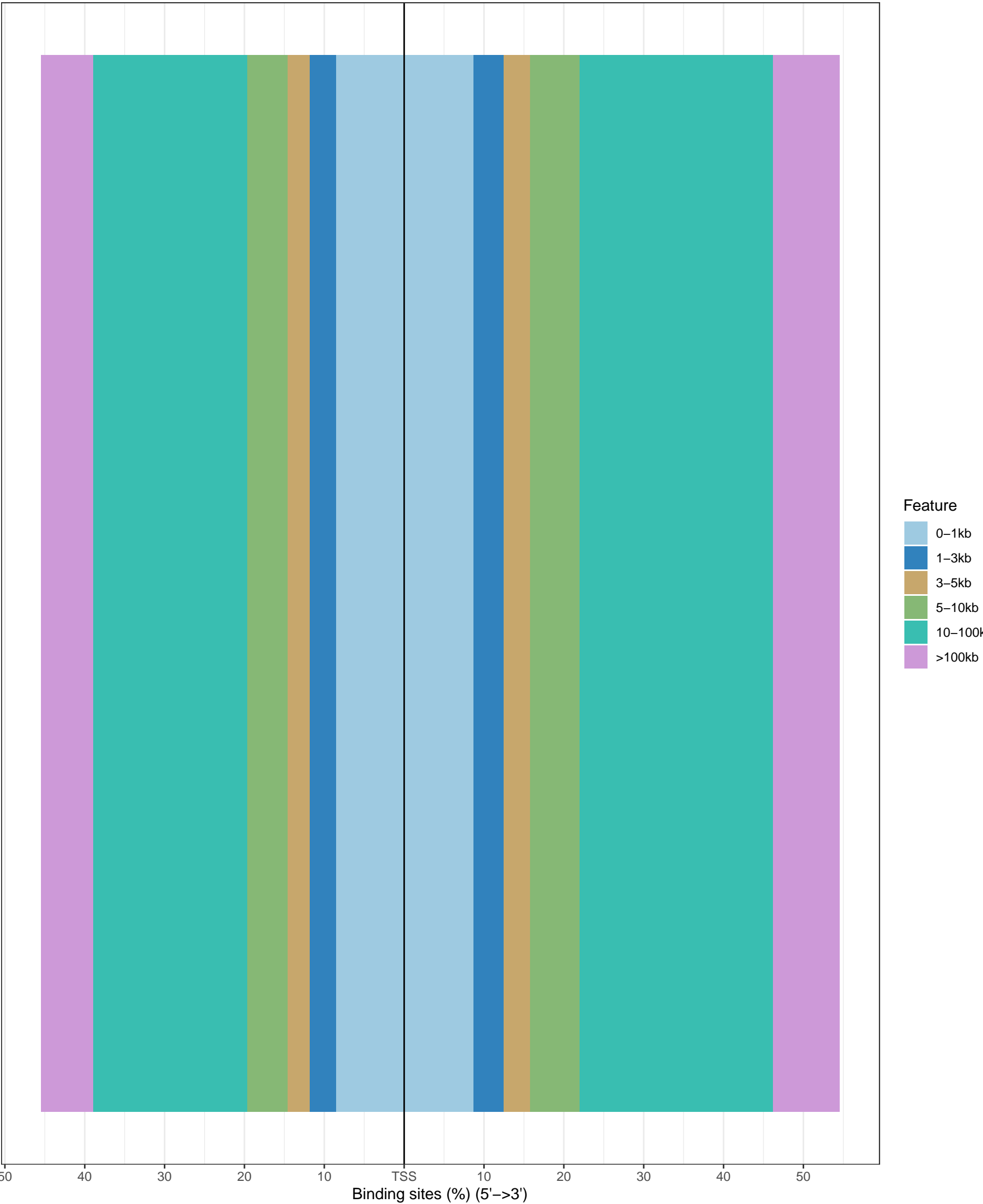
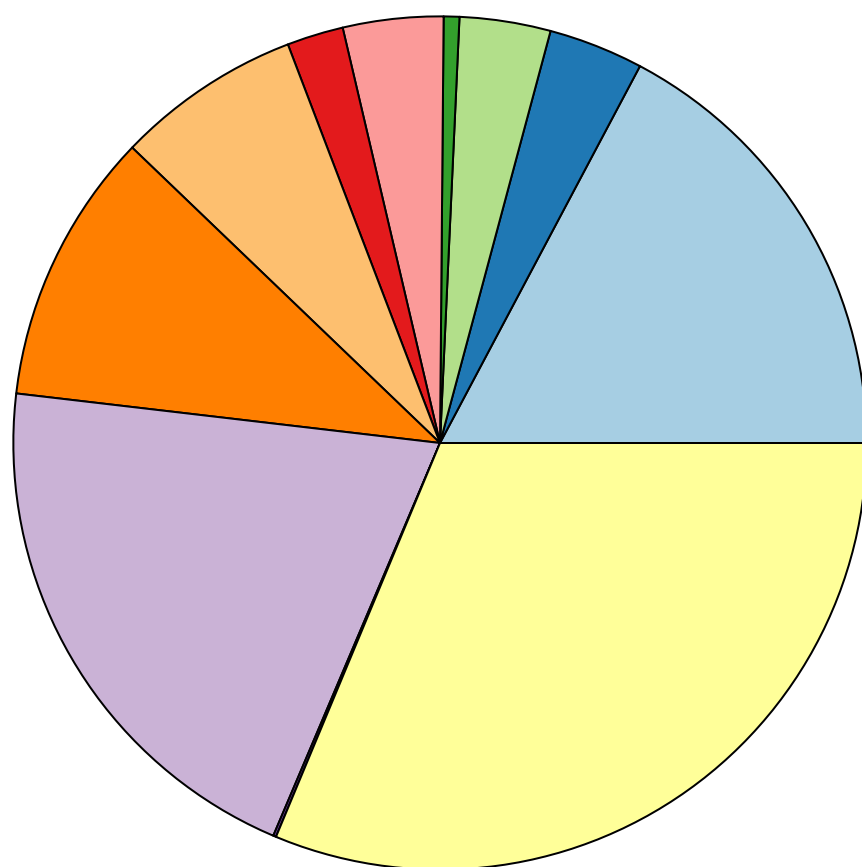


Feature Distribution



Distribution of transcription factor–binding loci relative to TSS





- Promoter (<=1kb) (17.24%)
- Promoter (1-2kb) (3.59%)
- Promoter (2-3kb) (3.44%)
- 5' UTR (0.59%)
- 3' UTR (3.8%)
- 1st Exon (2.14%)
- Other Exon (7.03%)
- 1st Intron (10.32%)
- Other Intron (20.46%)
- Downstream (<=300) (0.11%)
- Distal Intergenic (31.28%)