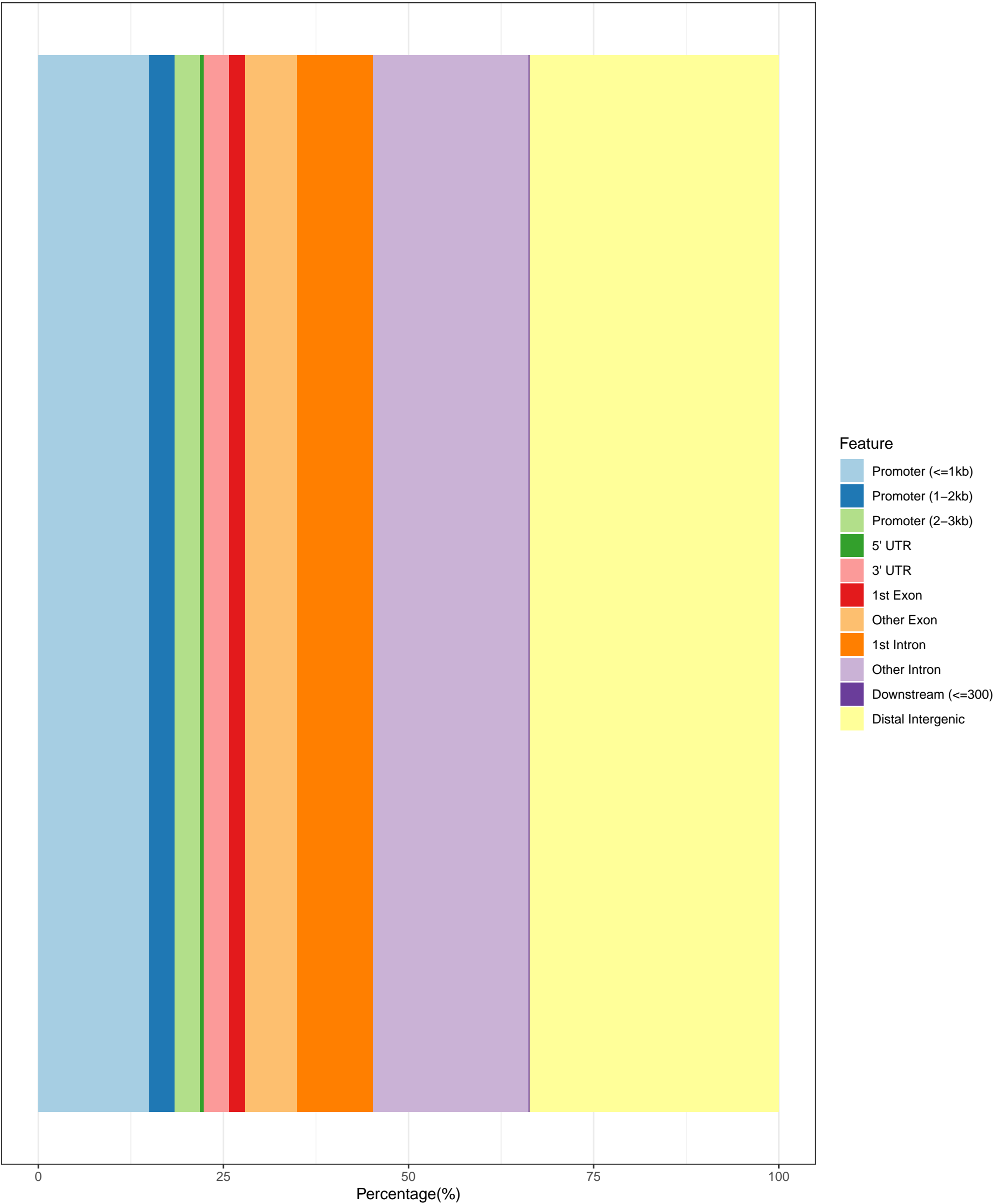
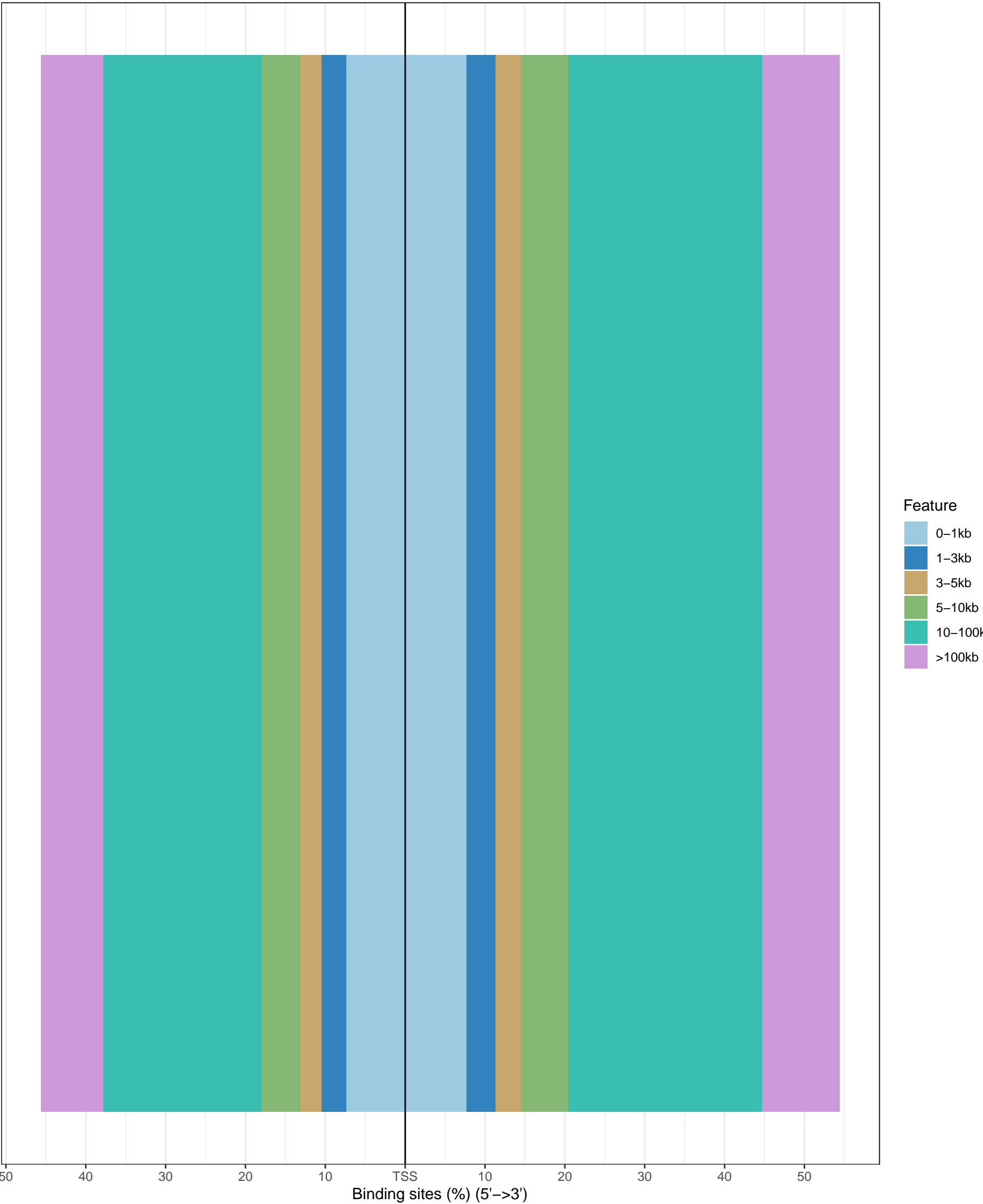
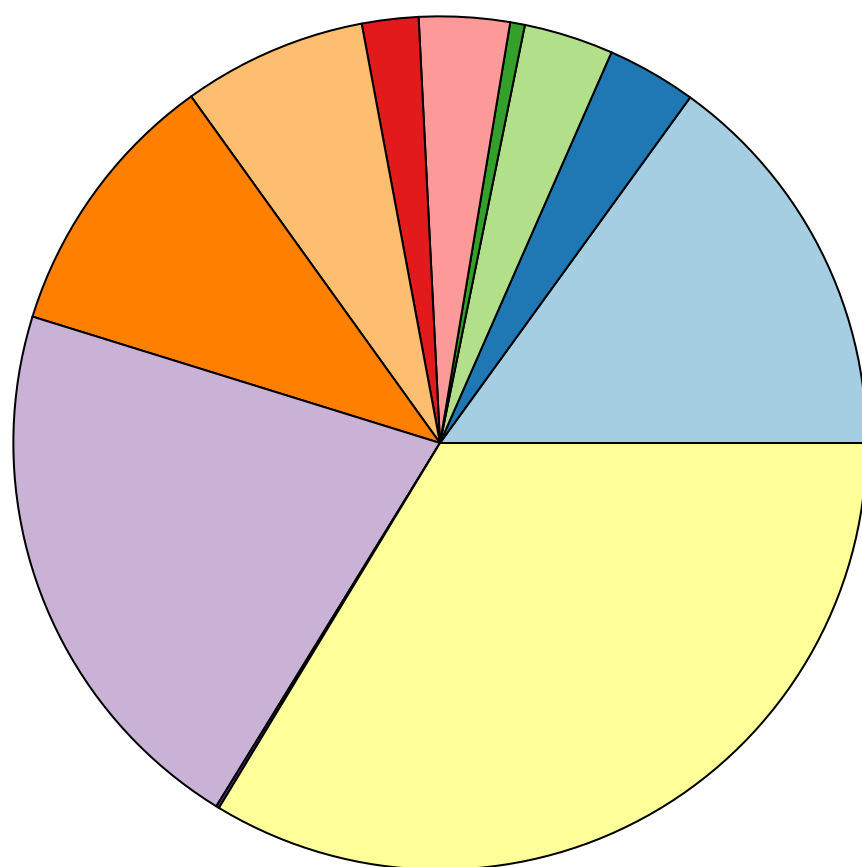


Feature Distribution



Distribution of transcription factor-binding loci relative to TSS





- Promoter (<=1kb) (15.03%)
- Promoter (1-2kb) (3.39%)
- Promoter (2-3kb) (3.39%)
- 5' UTR (0.56%)
- 3' UTR (3.43%)
- 1st Exon (2.14%)
- Other Exon (6.97%)
- 1st Intron (10.3%)
- Other Intron (21%)
- Downstream (<=300) (0.11%)
- Distal Intergenic (33.67%)