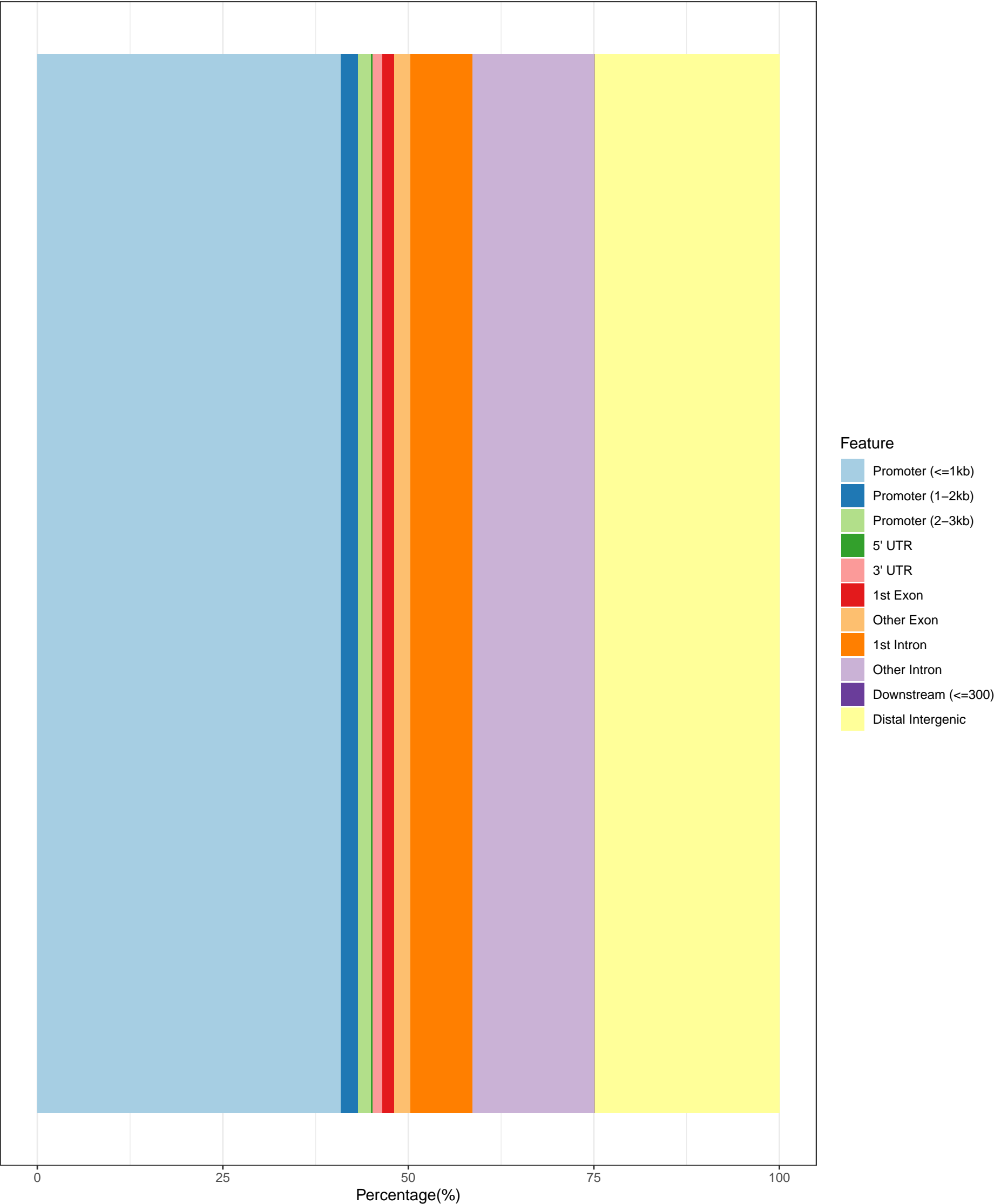
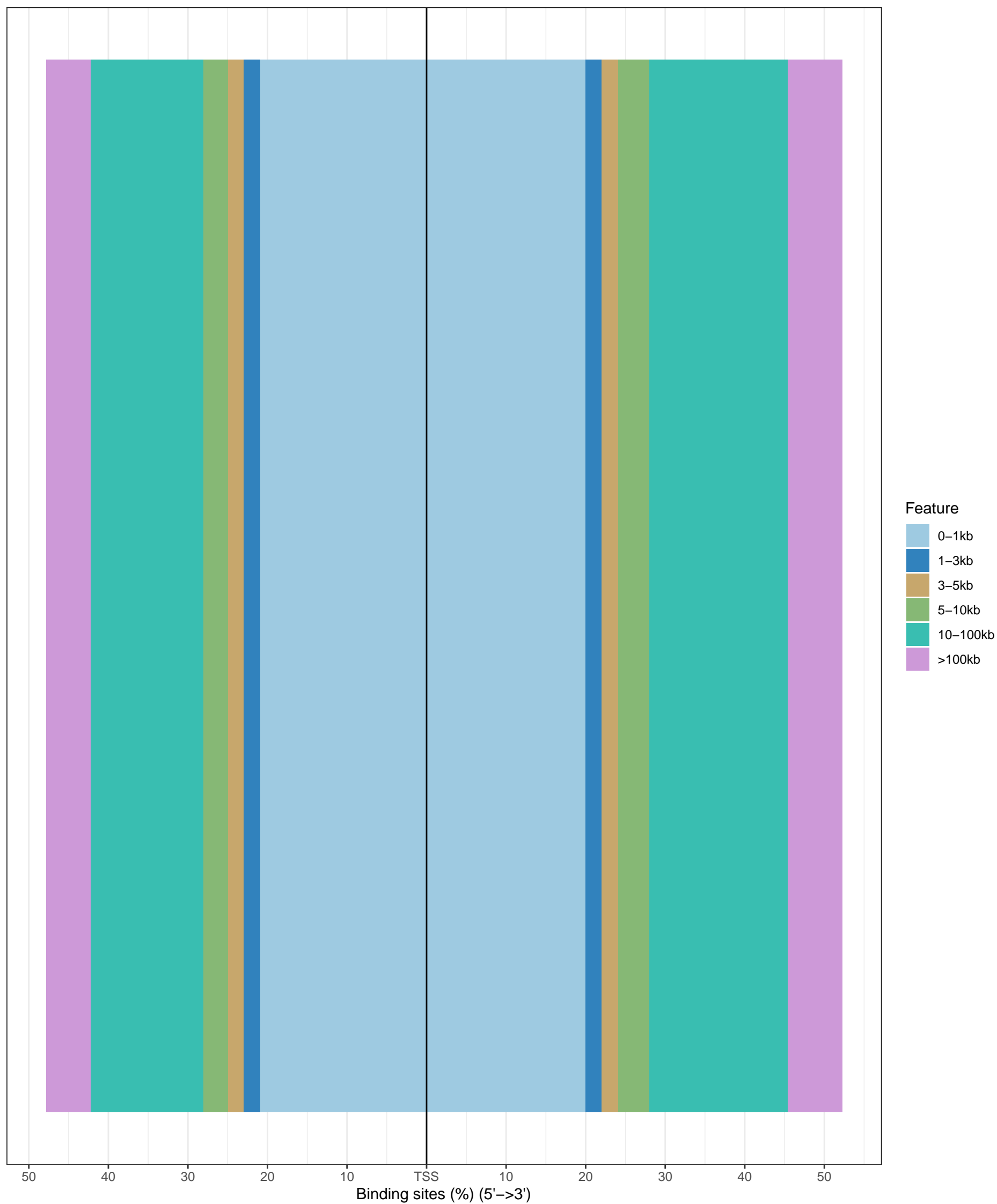
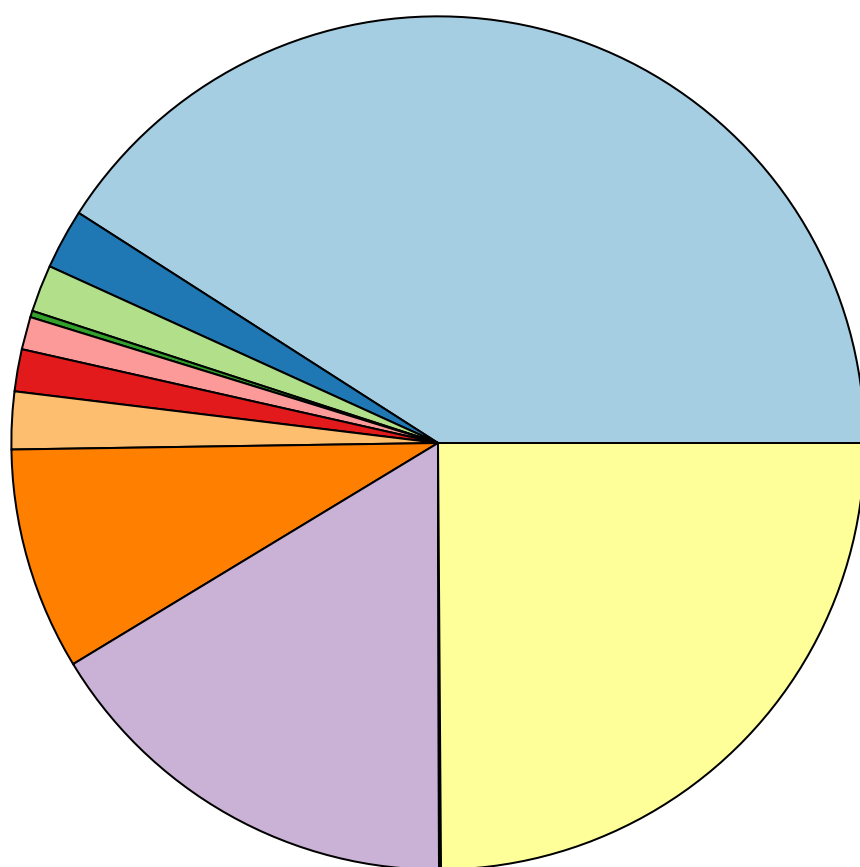


Feature Distribution



Distribution of transcription factor-binding loci relative to TSS





- Promoter (<=1kb) (40.93%)
- Promoter (1-2kb) (2.27%)
- Promoter (2-3kb) (1.78%)
- 5' UTR (0.24%)
- 3' UTR (1.24%)
- 1st Exon (1.6%)
- Other Exon (2.18%)
- 1st Intron (8.43%)
- Other Intron (16.37%)
- Downstream (<=300) (0.08%)
- Distal Intergenic (24.89%)