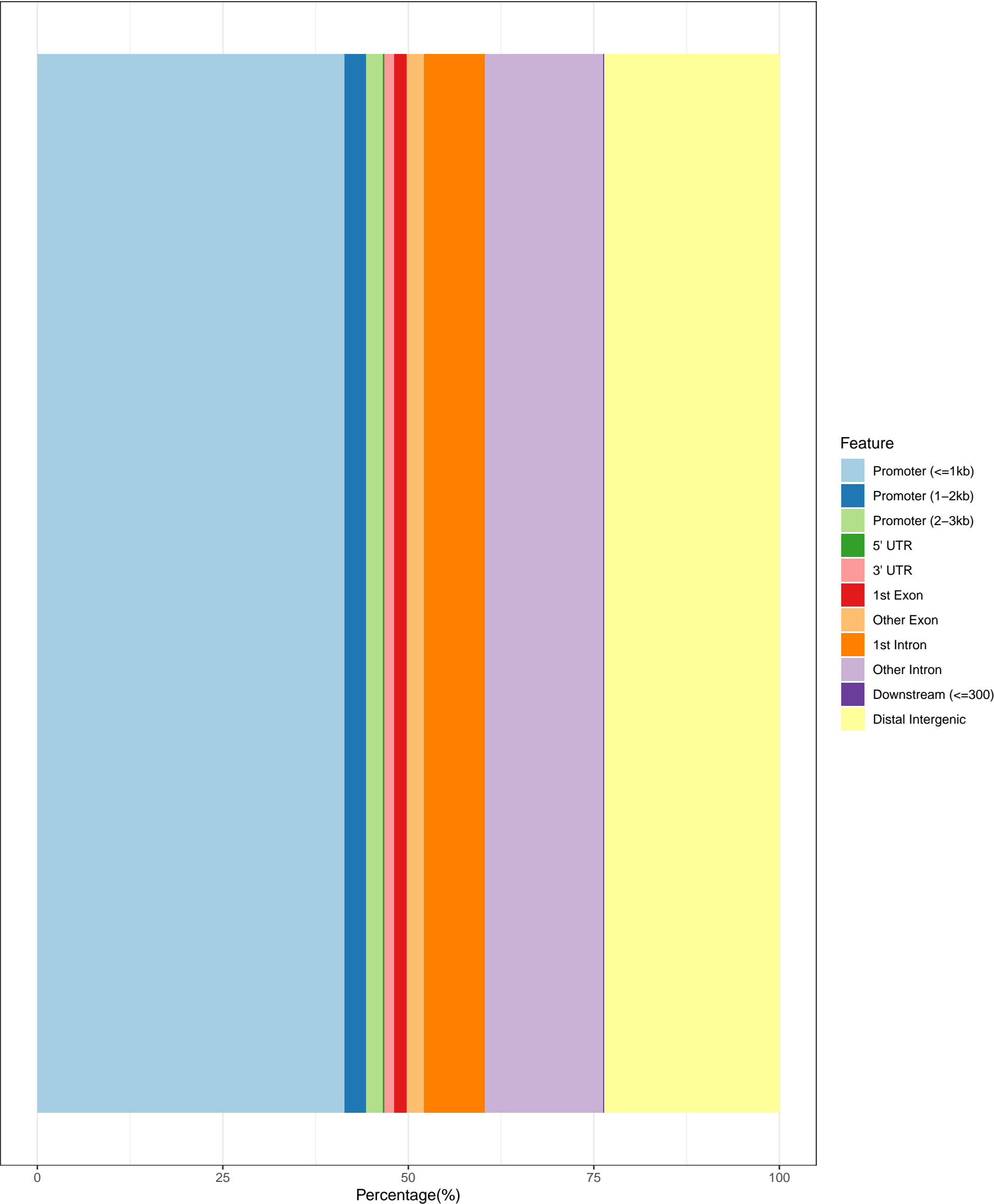
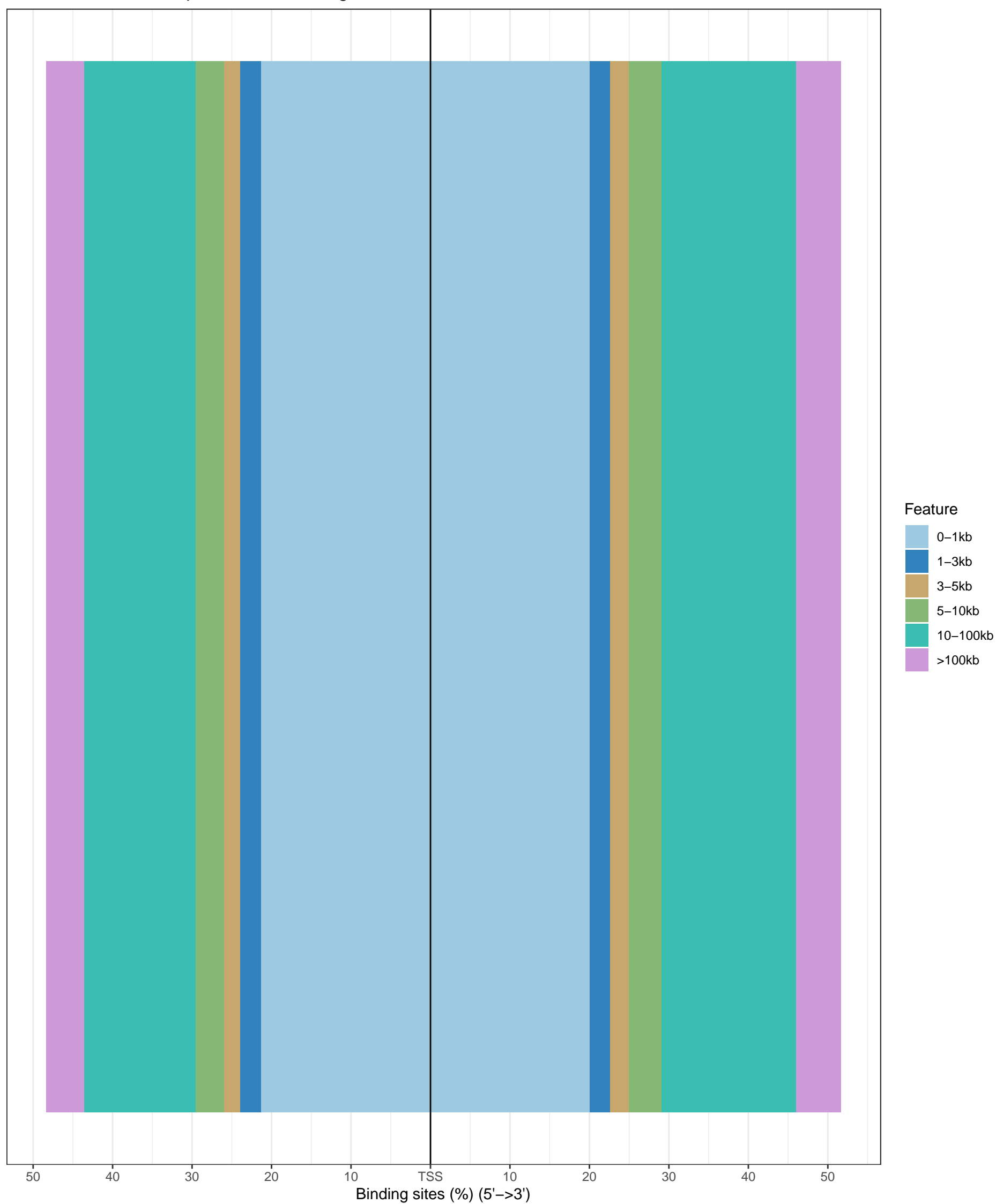
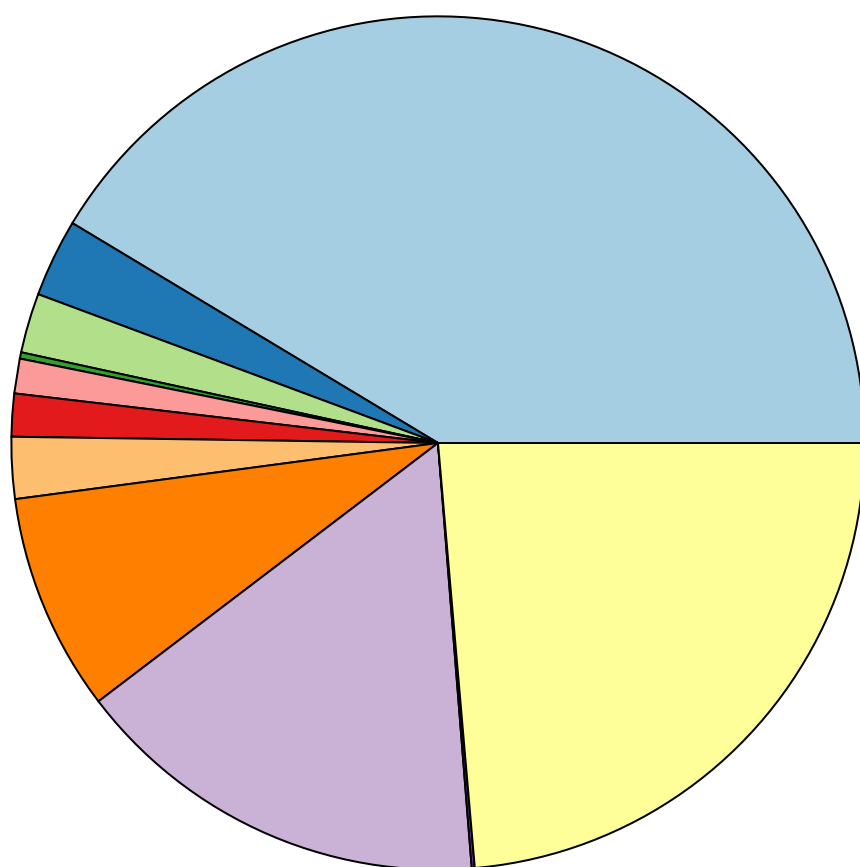


Feature Distribution



Distribution of transcription factor-binding loci relative to TSS





- Promoter (<=1kb) (41.38%)
- Promoter (1-2kb) (2.95%)
- Promoter (2-3kb) (2.26%)
- 5' UTR (0.24%)
- 3' UTR (1.32%)
- 1st Exon (1.62%)
- Other Exon (2.34%)
- 1st Intron (8.25%)
- Other Intron (15.91%)
- Downstream (<=300) (0.11%)
- Distal Intergenic (23.62%)