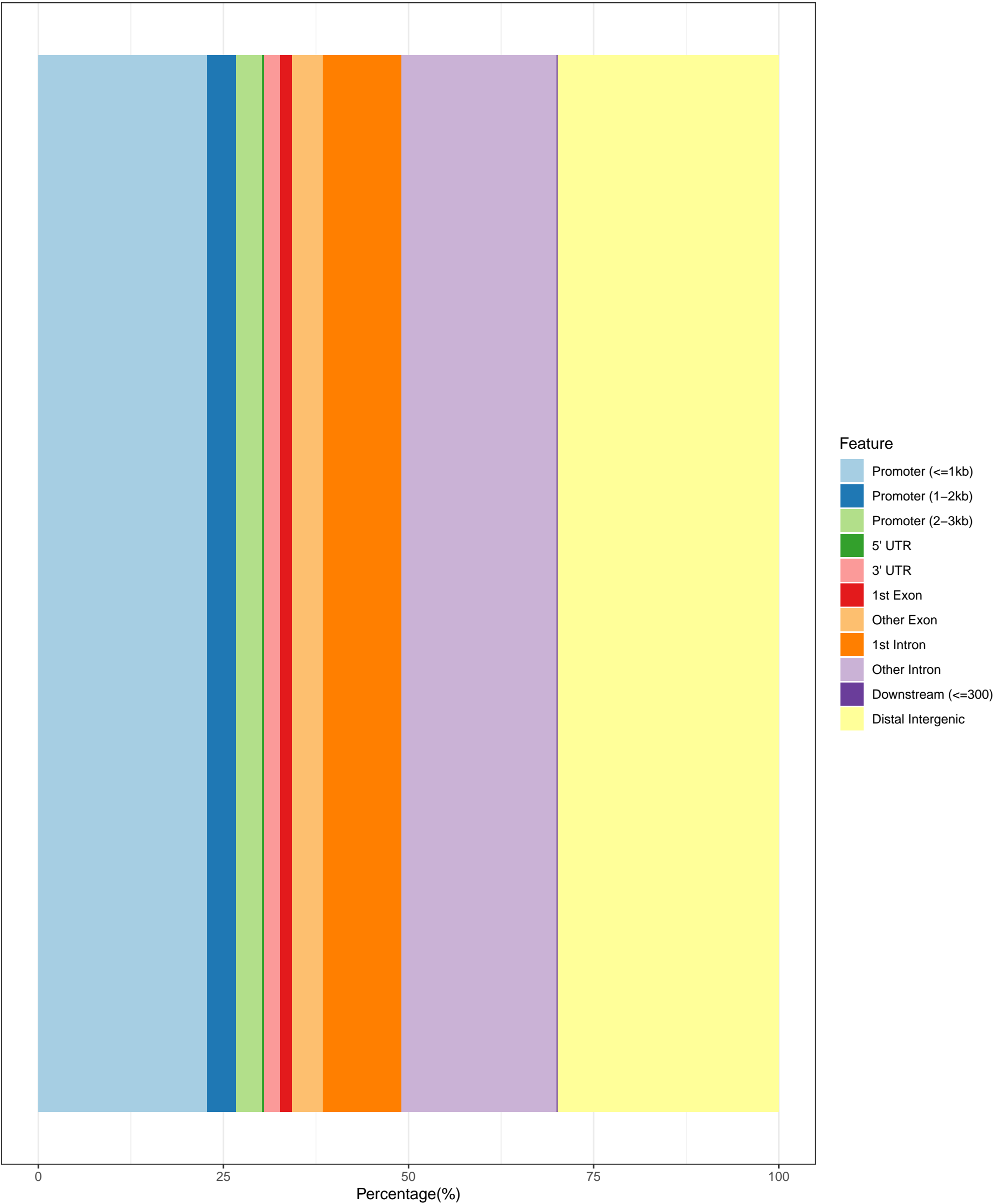
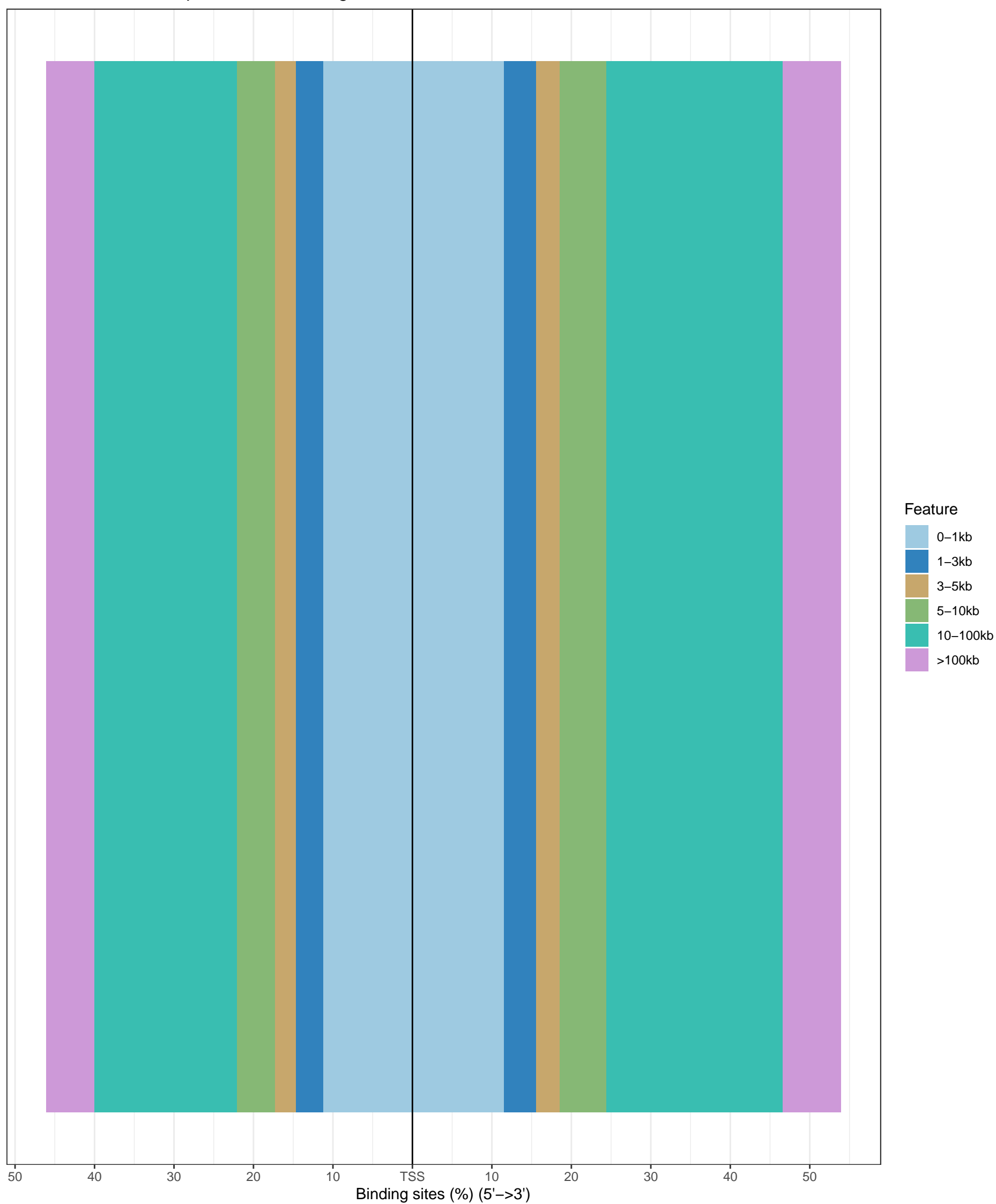
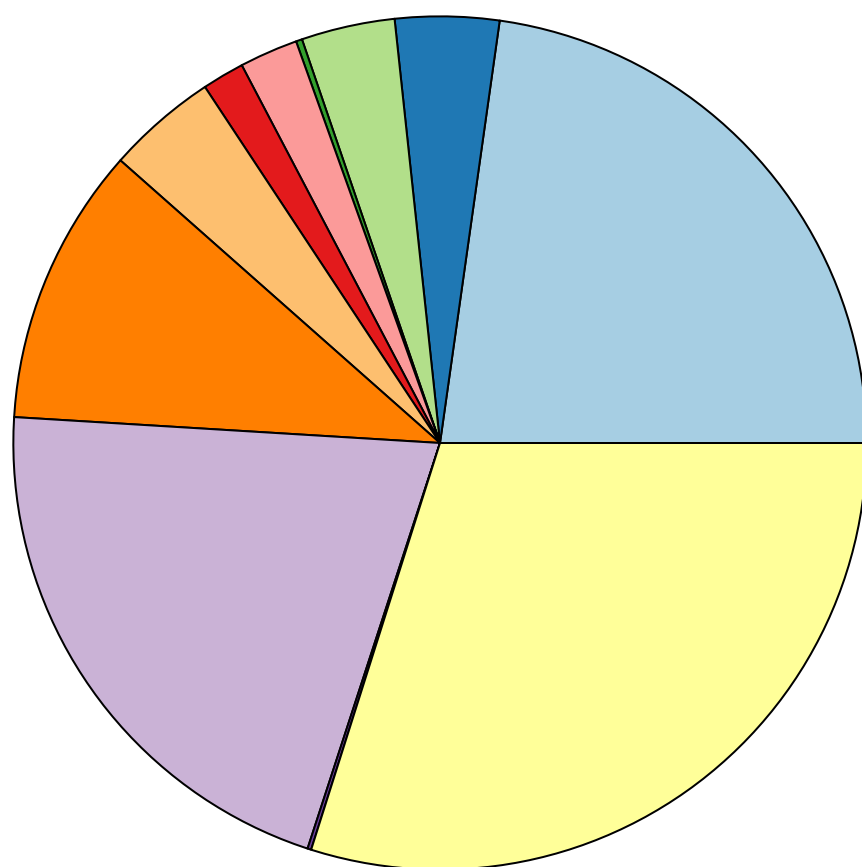


Feature Distribution



Distribution of transcription factor-binding loci relative to TSS





- Promoter (≤ 1 kb) (22.76%)
- Promoter (1–2kb) (3.93%)
- Promoter (2–3kb) (3.55%)
- 5' UTR (0.23%)
- 3' UTR (2.21%)
- 1st Exon (1.6%)
- Other Exon (4.19%)
- 1st Intron (10.56%)
- Other Intron (20.94%)
- Downstream (≤ 300) (0.14%)
- Distal Intergenic (29.88%)