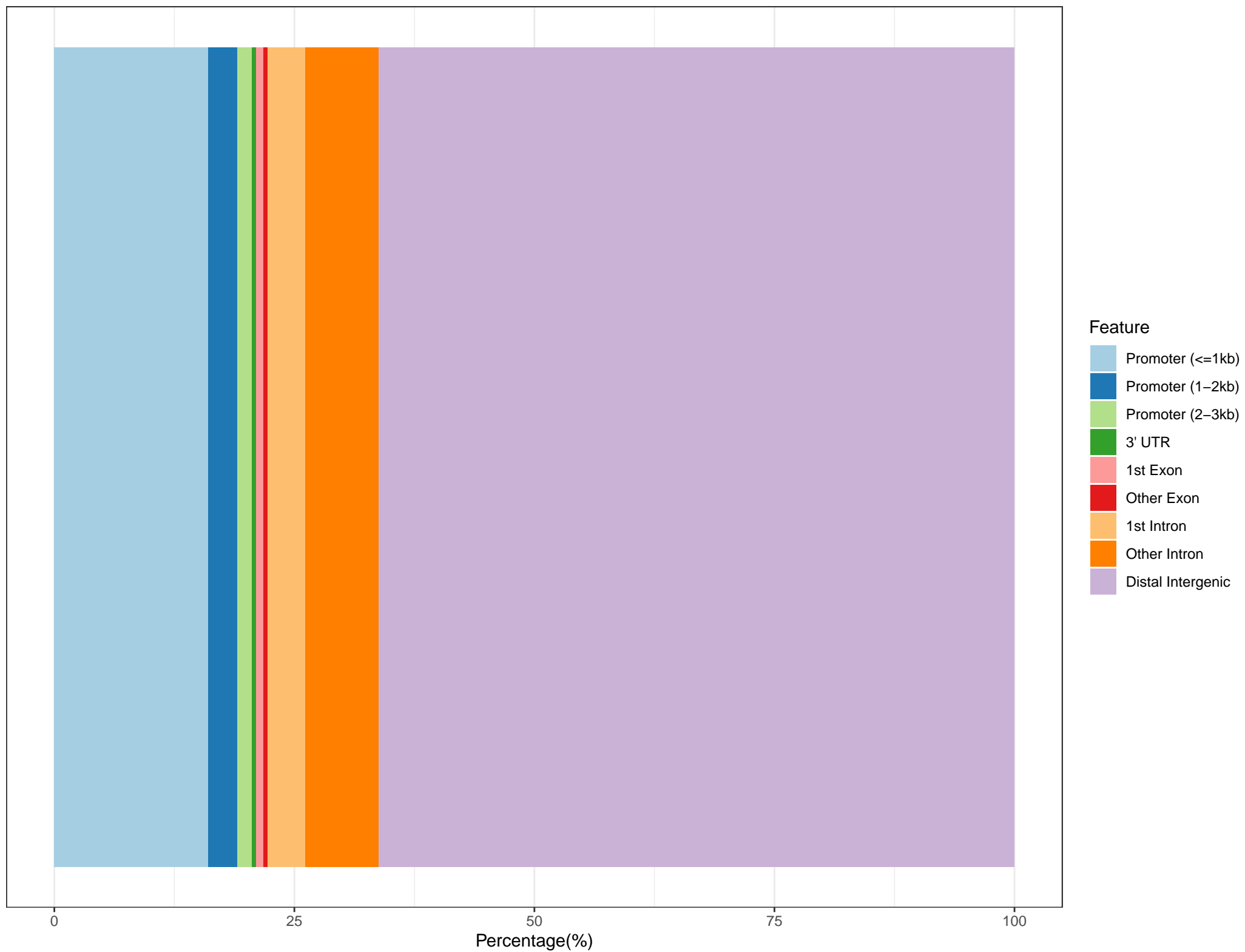
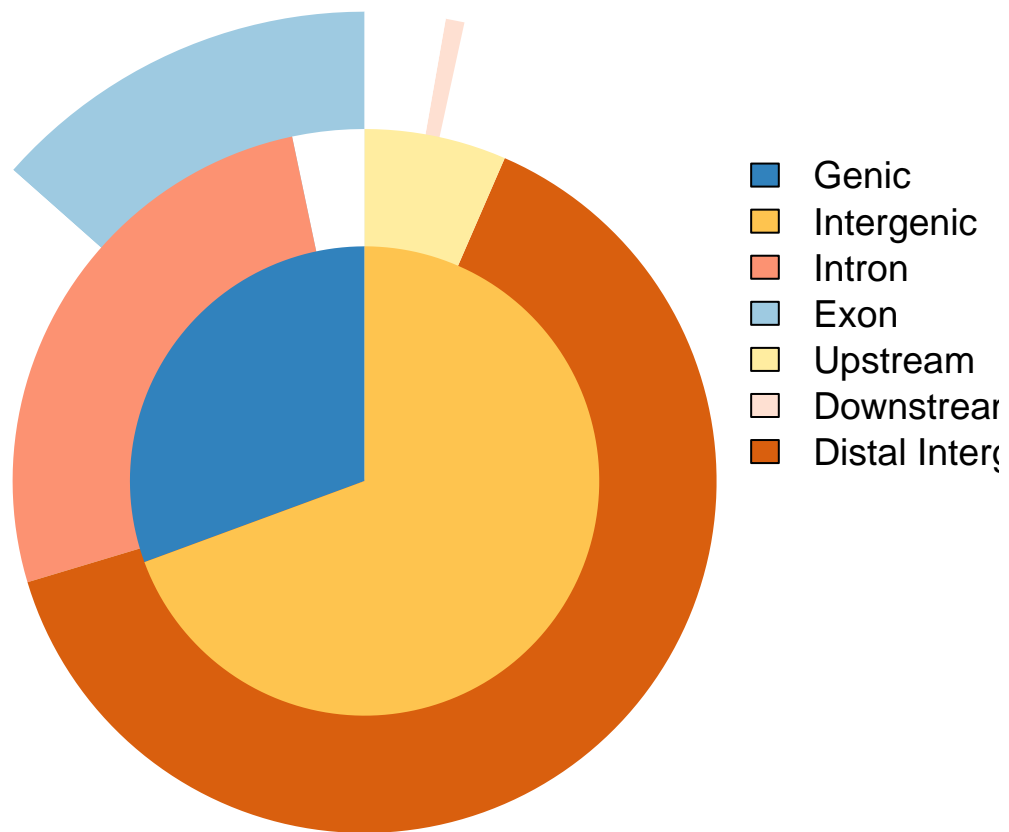


- Promoter (<=1kb) (16.03%)
- Promoter (1-2kb) (3.06%)
- Promoter (2-3kb) (1.48%)
- 3' UTR (0.46%)
- 1st Exon (0.74%)
- Other Exon (0.46%)
- 1st Intron (3.89%)
- Other Intron (7.69%)
- Distal Intergenic (66.17%)

Feature Distribution





Distribution of transcription factor–binding loci
relative to TSS

