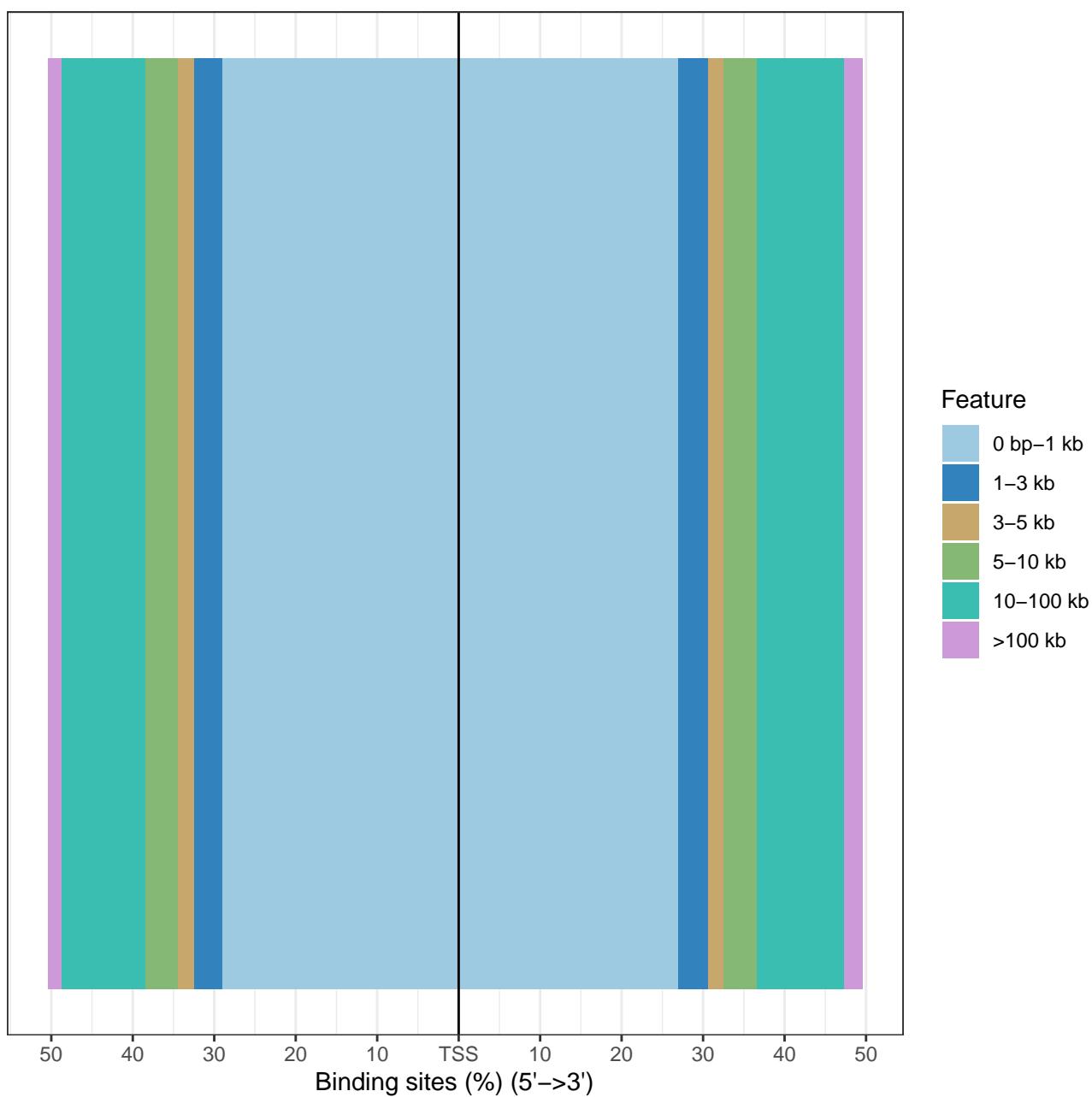


- Promoter (<=1kb) (56.01%)
- Promoter (1–2kb) (4.3%)
- Promoter (2–3kb) (2.78%)
- 5' UTR (0.04%)
- 3' UTR (0.62%)
- 1st Exon (0.76%)
- Other Exon (1.33%)
- 1st Intron (7.21%)
- Other Intron (10.83%)
- Downstream (<=300) (0.1%)
- Distal Intergenic (16.03%)

Distribution of transcription factor–binding loci relative to TSS



Feature Distribution

