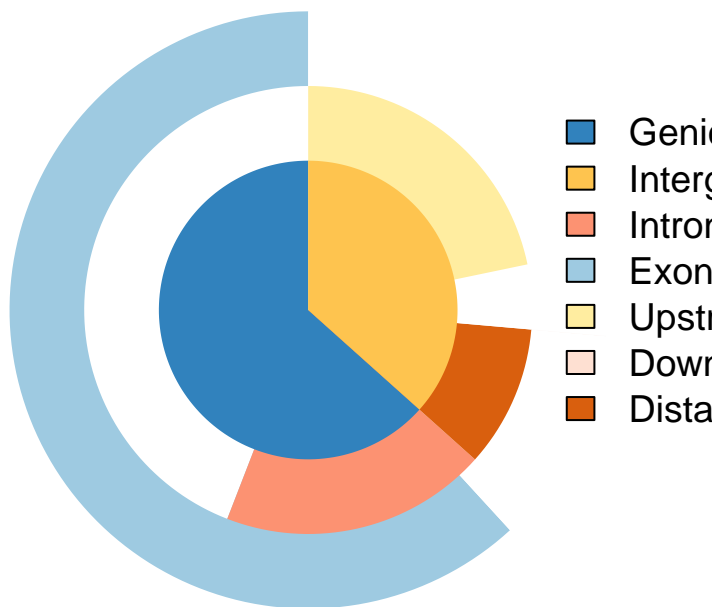
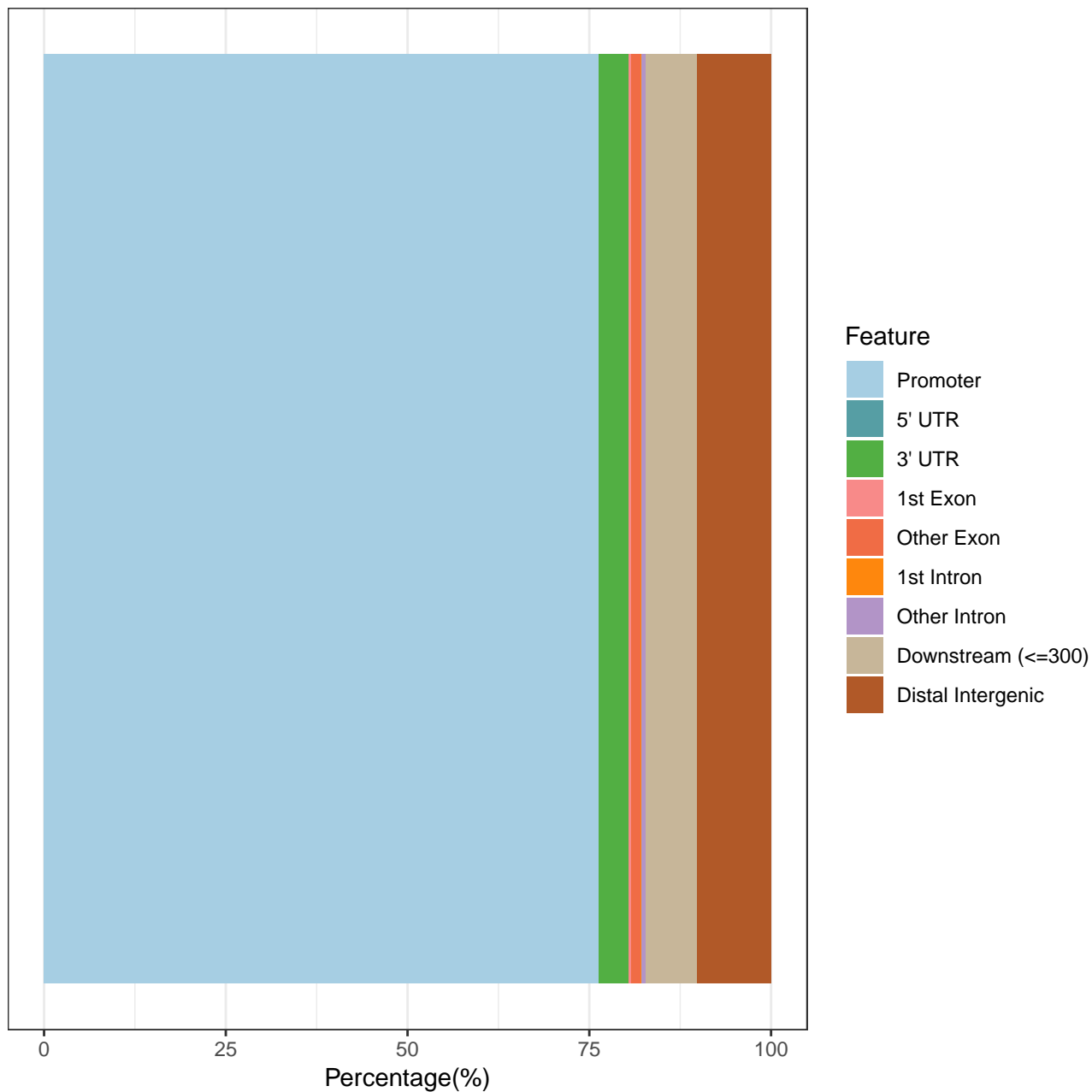


- Promoter (76.28%)
- 5' UTR (0.05%)
- 3' UTR (4.1%)
- 1st Exon (0.3%)
- Other Exon (1.39%)
- 1st Intron (0.12%)
- Other Intron (0.48%)
- Downstream (<=300) (7.05%)
- Distal Intergenic (10.23%)



Feature Distribution



Distribution of genomic loci relative to TSS

