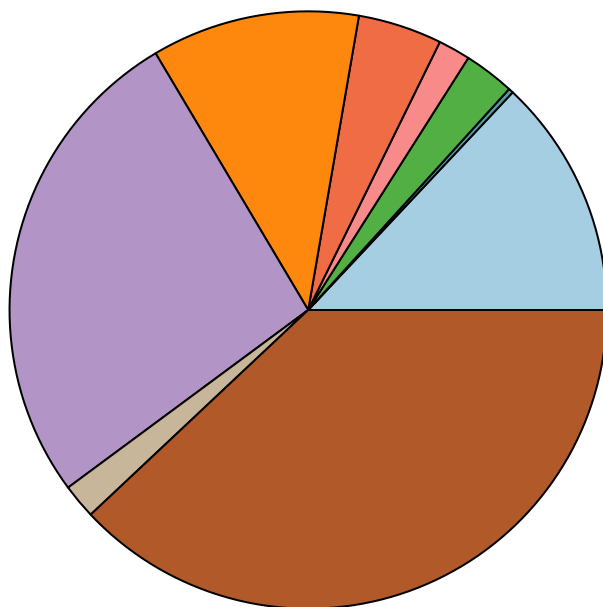
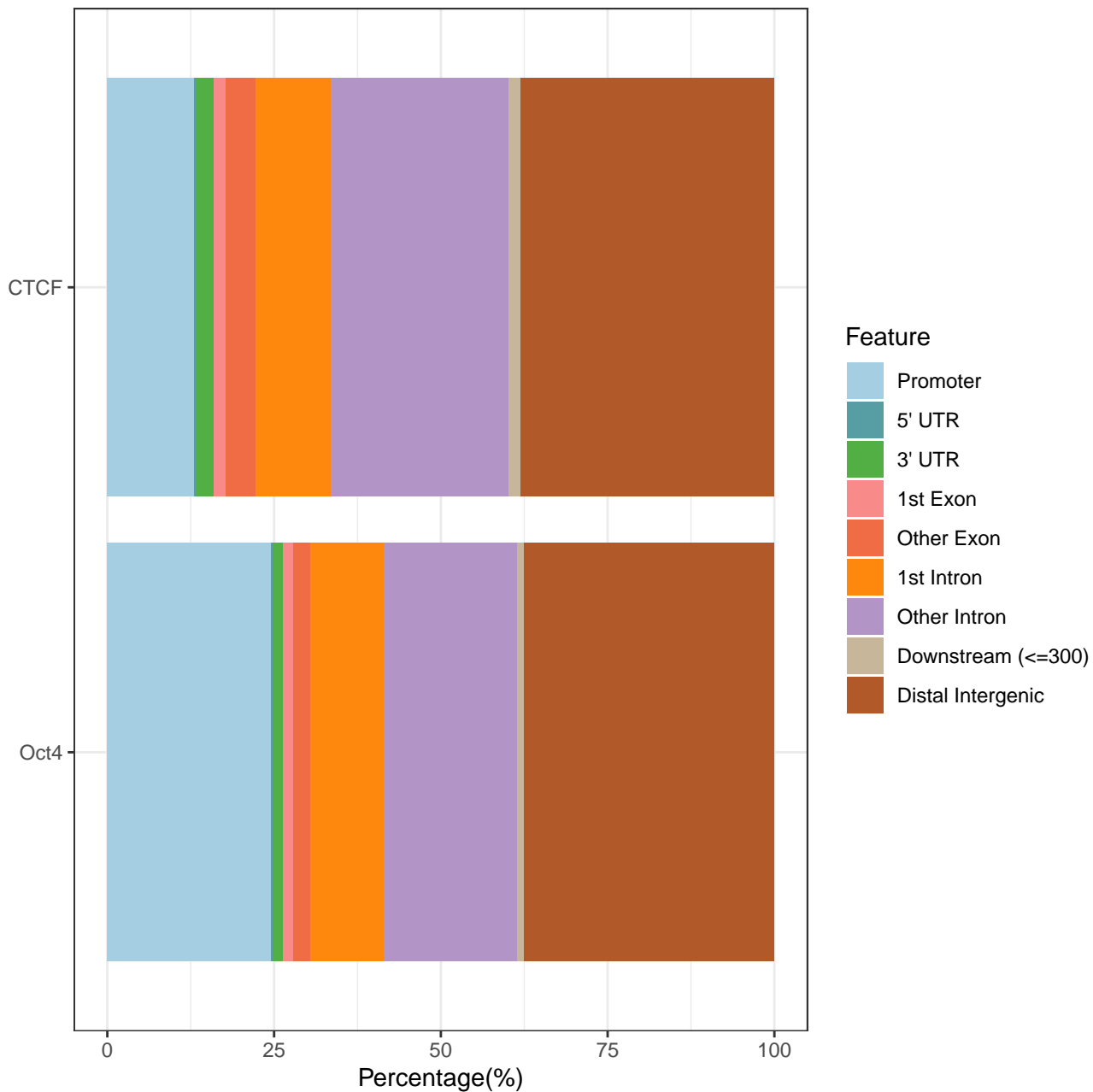


- Promoter (24.56%)
- 5' UTR (0.22%)
- 3' UTR (1.58%)
- 1st Exon (1.48%)
- Other Exon (2.65%)
- 1st Intron (10.99%)
- Other Intron (19.95%)
- Downstream (<=300) (1.08%)
- Distal Intergenic (37.49%)



- Promoter (13%)
- 5' UTR (0.26%)
- 3' UTR (2.74%)
- 1st Exon (1.74%)
- Other Exon (4.55%)
- 1st Intron (11.26%)
- Other Intron (26.6%)
- Downstream (<=300) (1.88%)
- Distal Intergenic (37.99%)

# Feature Distribution



# Distribution of transcription factor-binding loci relative to TSS

