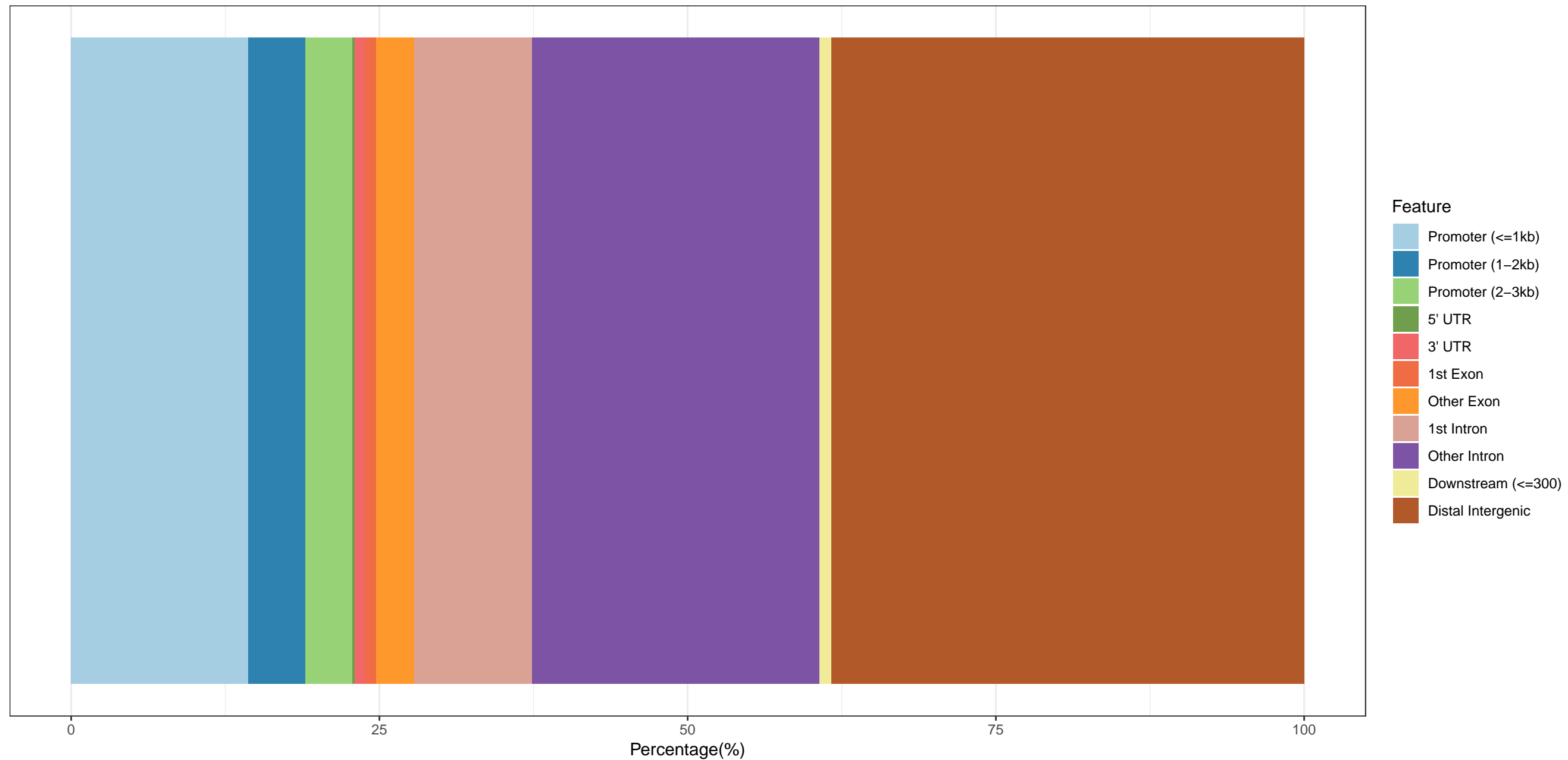
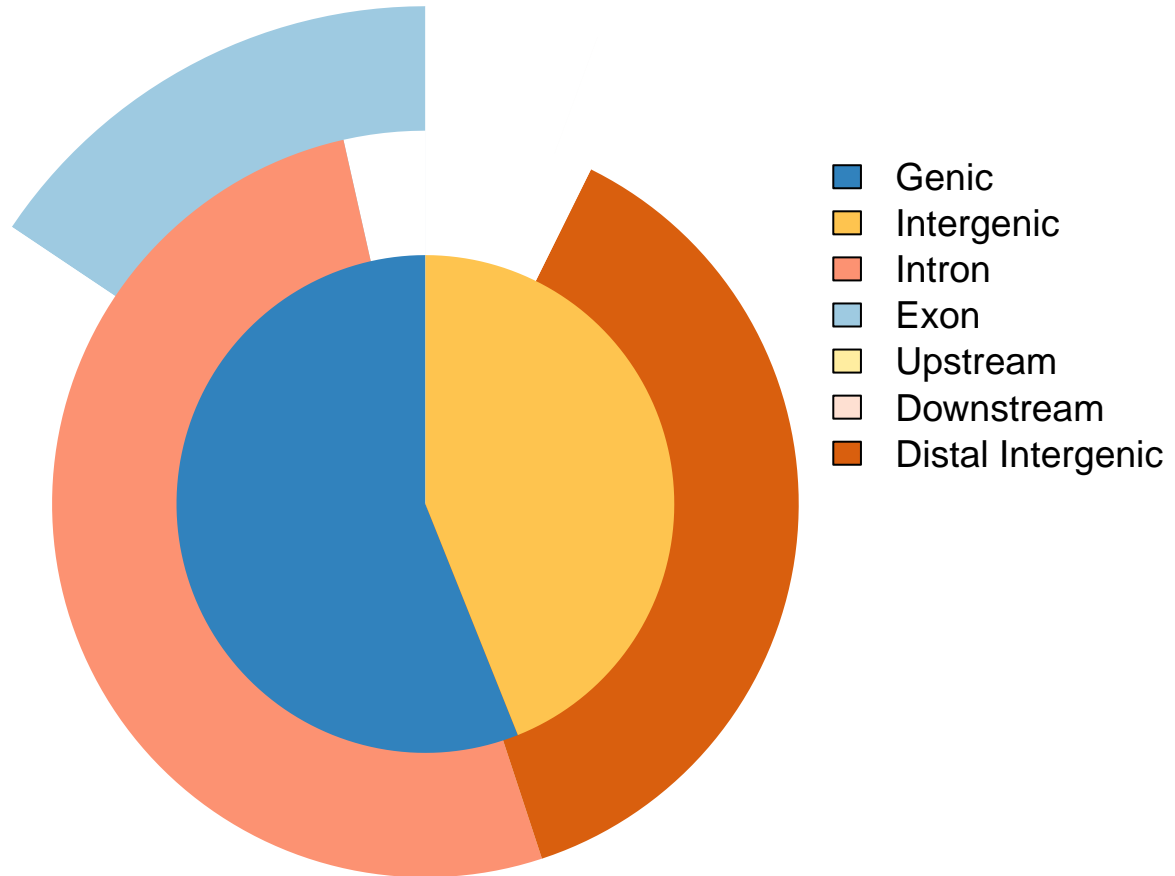
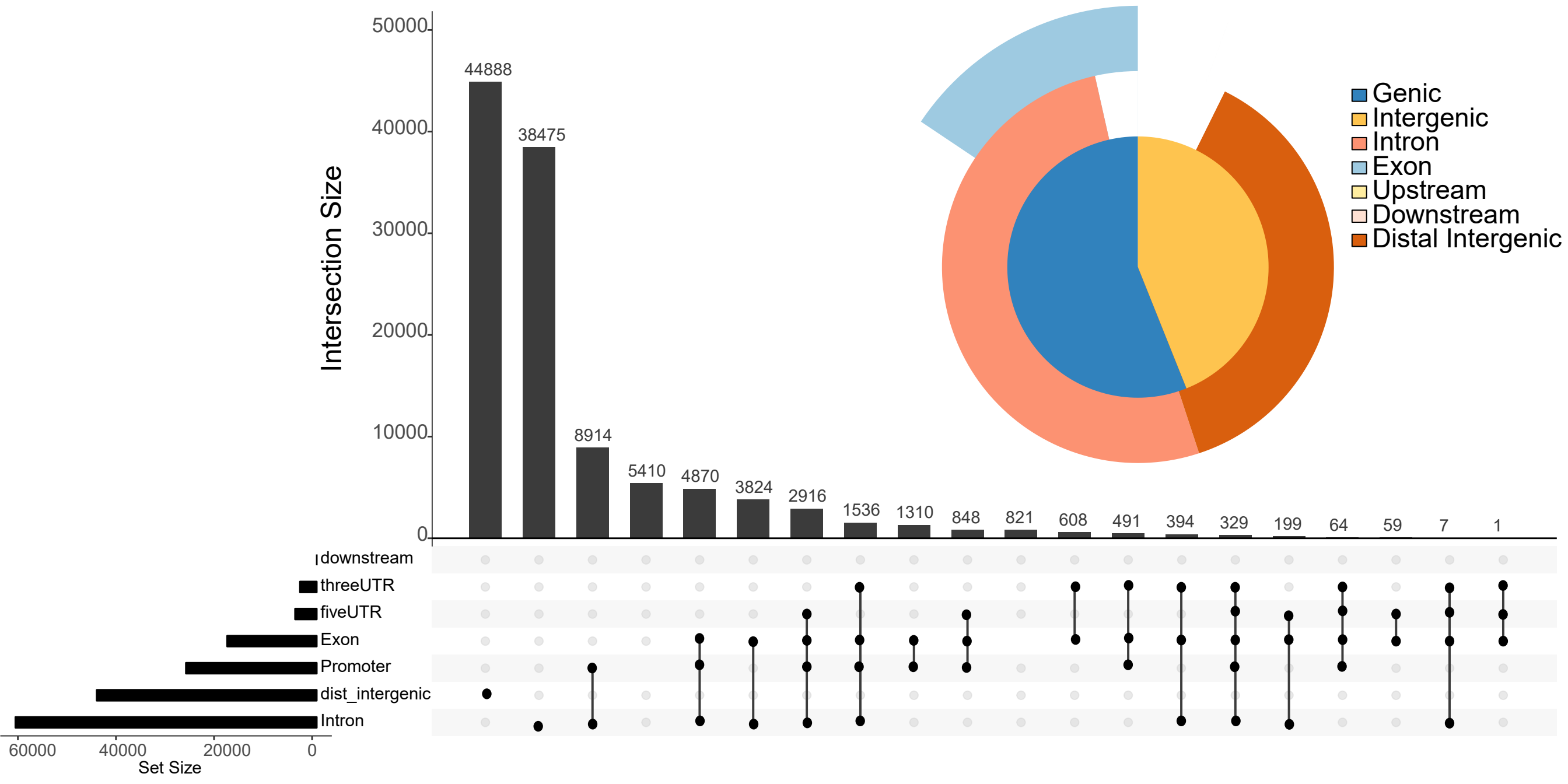


- Promoter (<=1kb) (14.35%)
- Promoter (1-2kb) (4.64%)
- Promoter (2-3kb) (3.79%)
- 5' UTR (0.23%)
- 3' UTR (0.86%)
- 1st Exon (0.86%)
- Other Exon (3.11%)
- 1st Intron (9.57%)
- Other Intron (23.28%)
- Downstream (<=300) (1%)
- Distal Intergenic (38.32%)

Feature Distribution







Distribution of transcription factor-binding loci
relative to TSS

