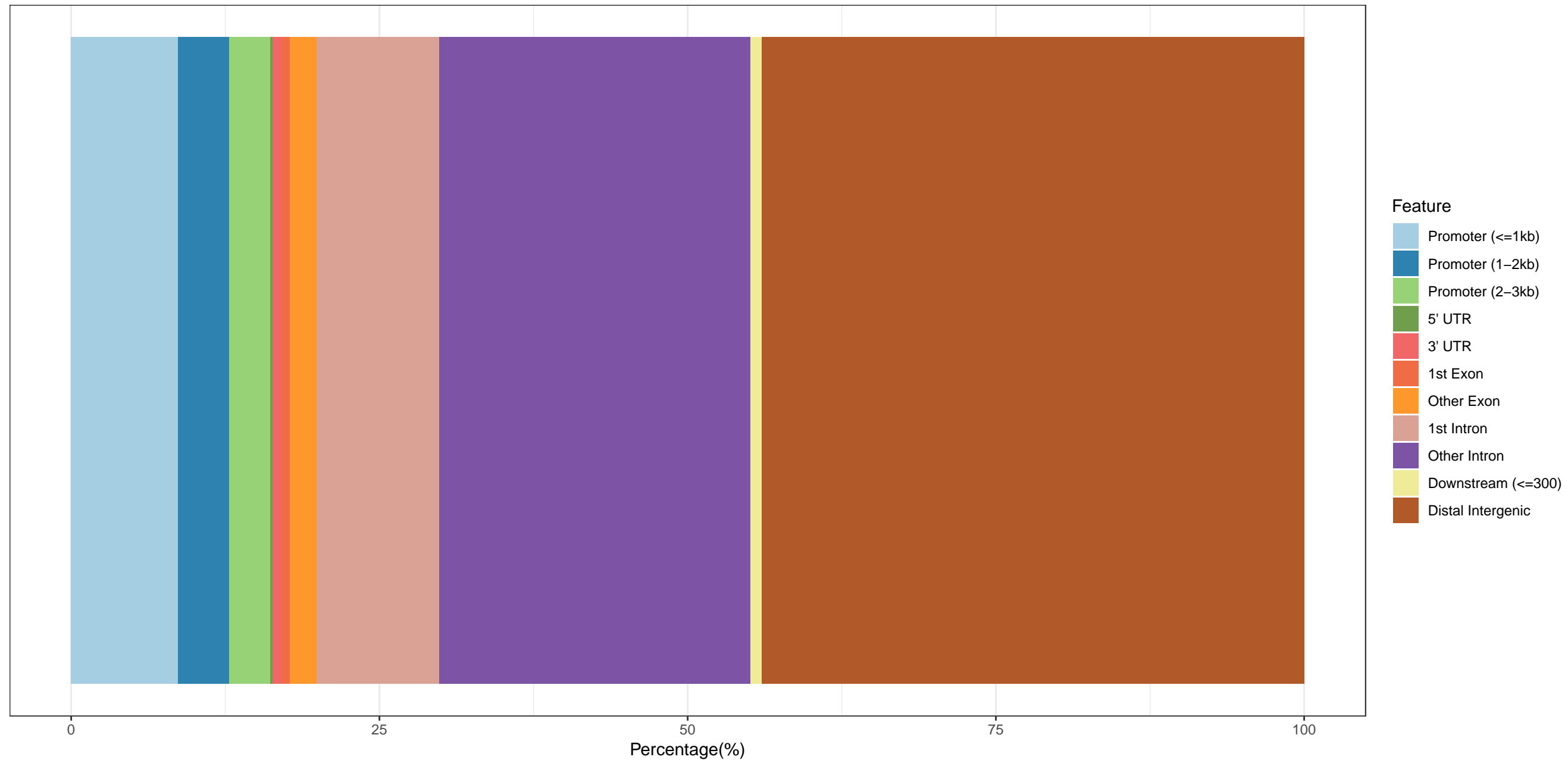
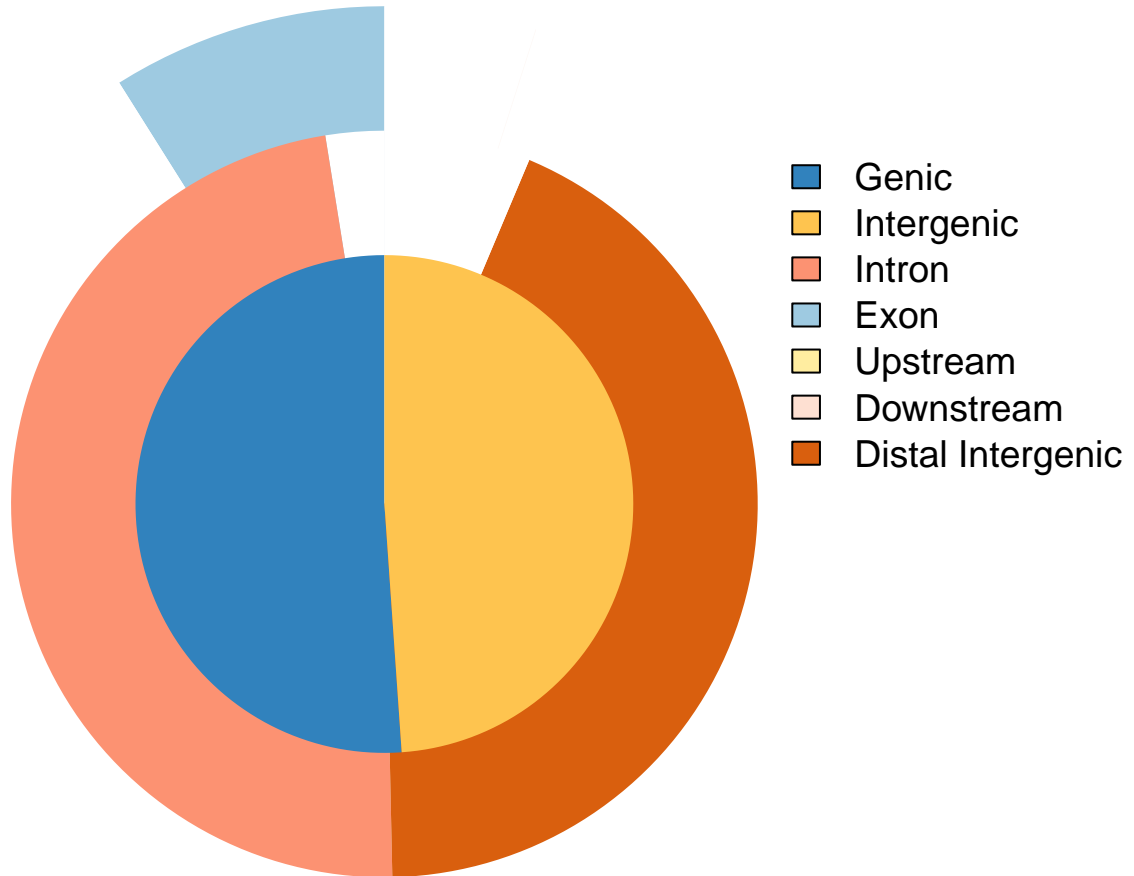
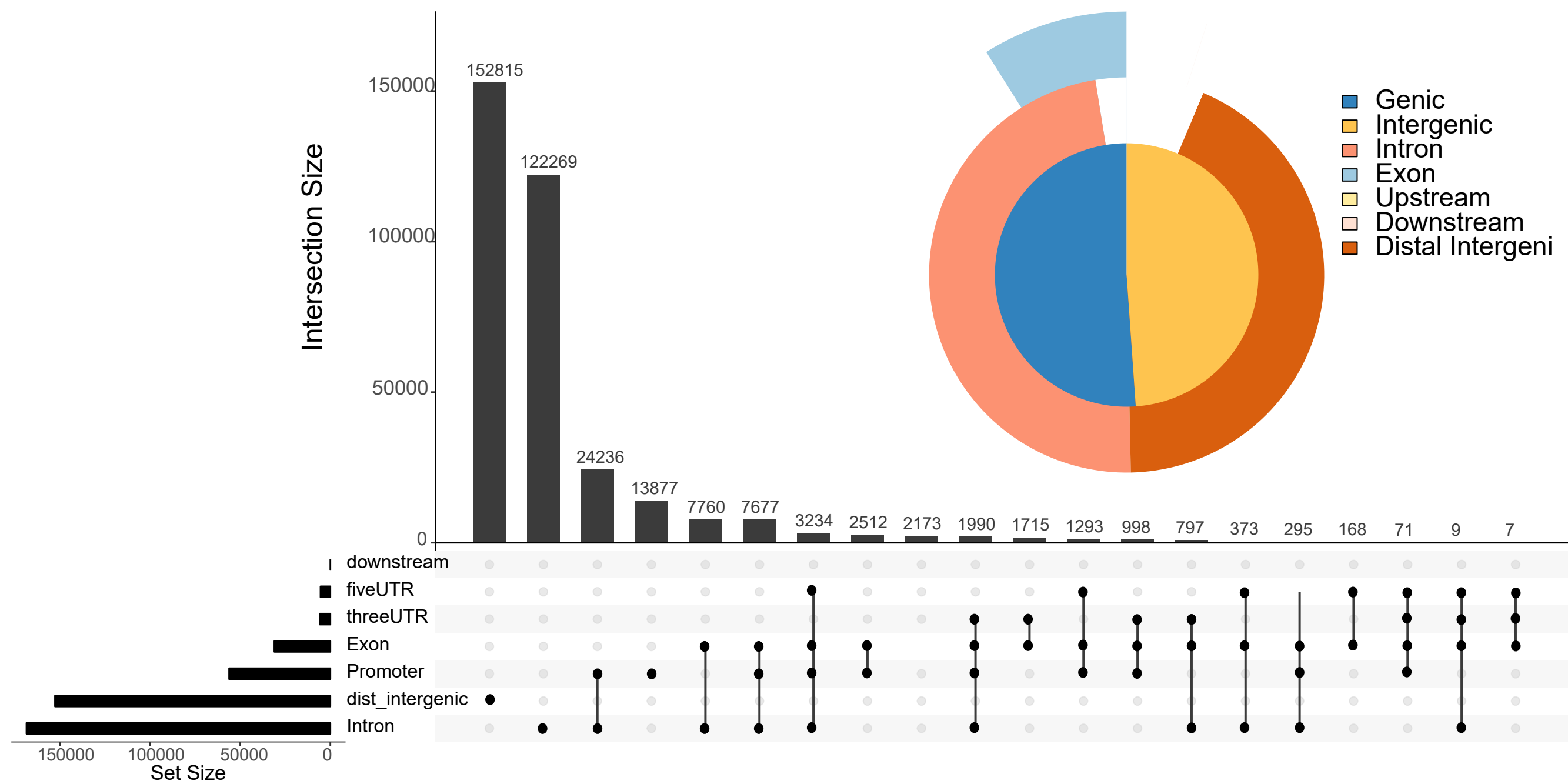


- Promoter (<=1kb) (8.67%)
- Promoter (1-2kb) (4.15%)
- Promoter (2-3kb) (3.35%)
- 5' UTR (0.16%)
- 3' UTR (0.72%)
- 1st Exon (0.72%)
- Other Exon (2.14%)
- 1st Intron (9.93%)
- Other Intron (25.26%)
- Downstream (<=300) (0.92%)
- Distal Intergenic (43.98%)

Feature Distribution







Distribution of transcription factor-binding loci relative to TSS

