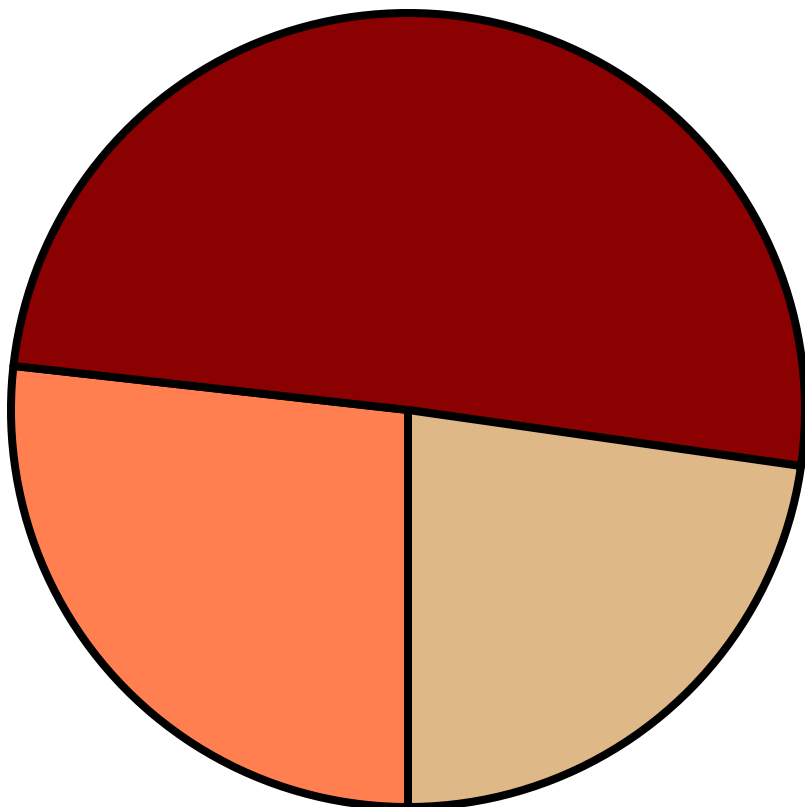# test_1 (56438 peaks)

REGIONS

- PROMOTER (26.8%)
- INTRAGENIC (50.5%)
- INTERGENIC (22.7%)

Figure legend. Distribution of ChIPseq reads across different regions of the genome.

This plot is generated by counting the number of peaks fitting on each class of region.

PROMOTER region is the region within 2.5 Kbp upstream of the TSS and the TSS.

INTRAGENIC regions is the region within the transcripts. INTERGENIC is the rest

of the genome. TSS is the Transcription Start Site.

ChIPseq experiment:

test_1

Number of peaks:

43063

RefSeq transcripts:

36859

Window factor:

10