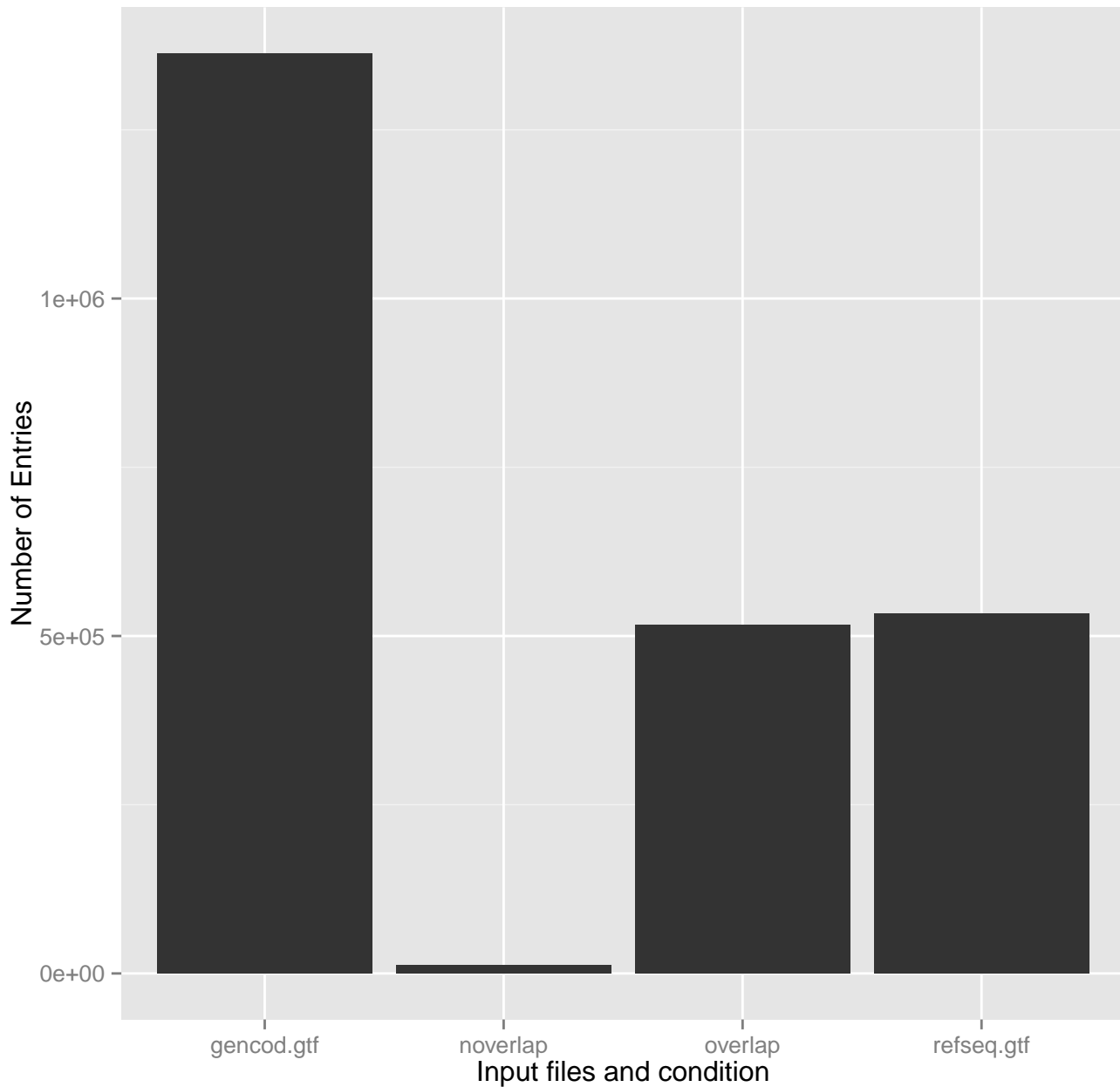
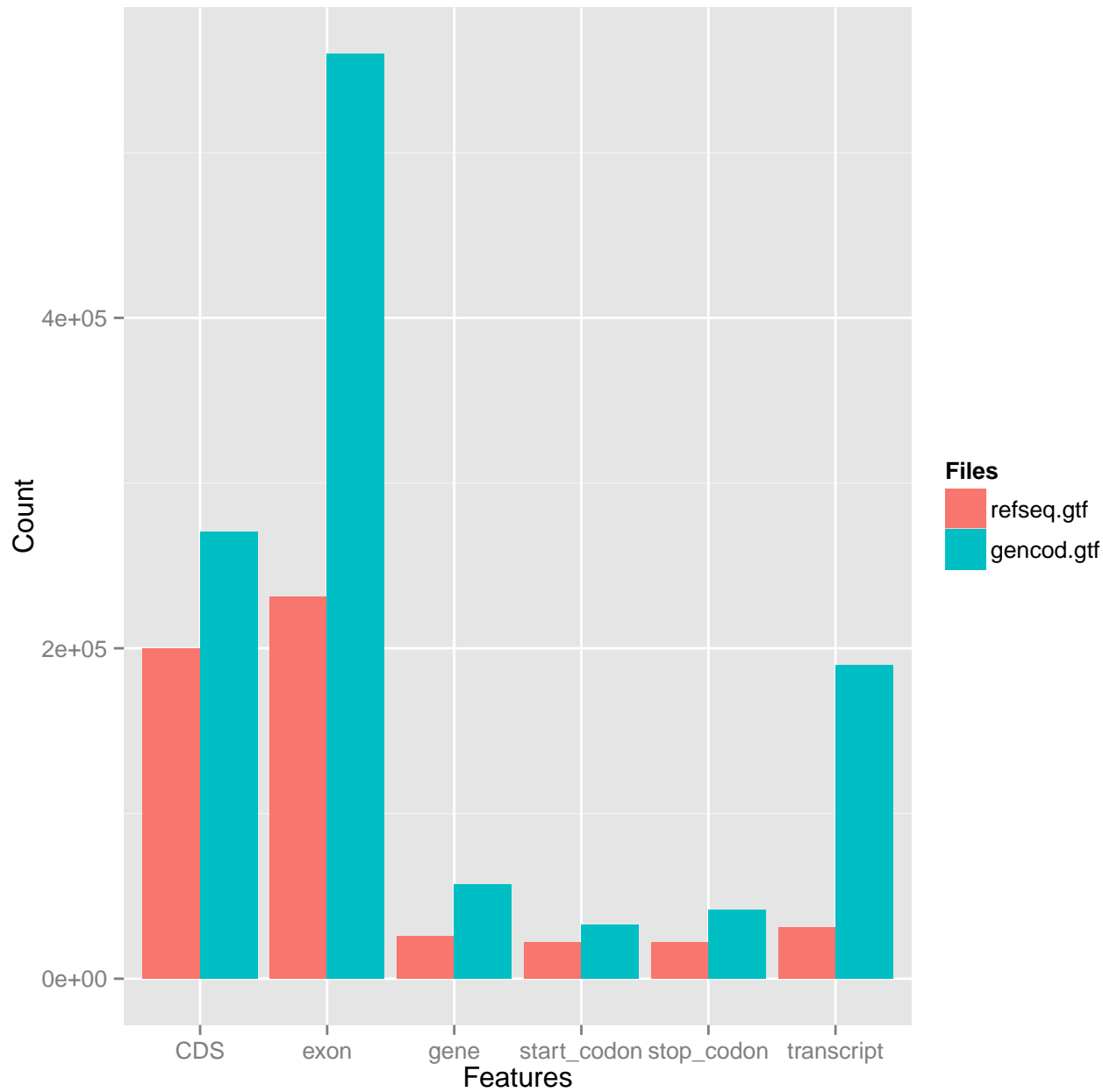


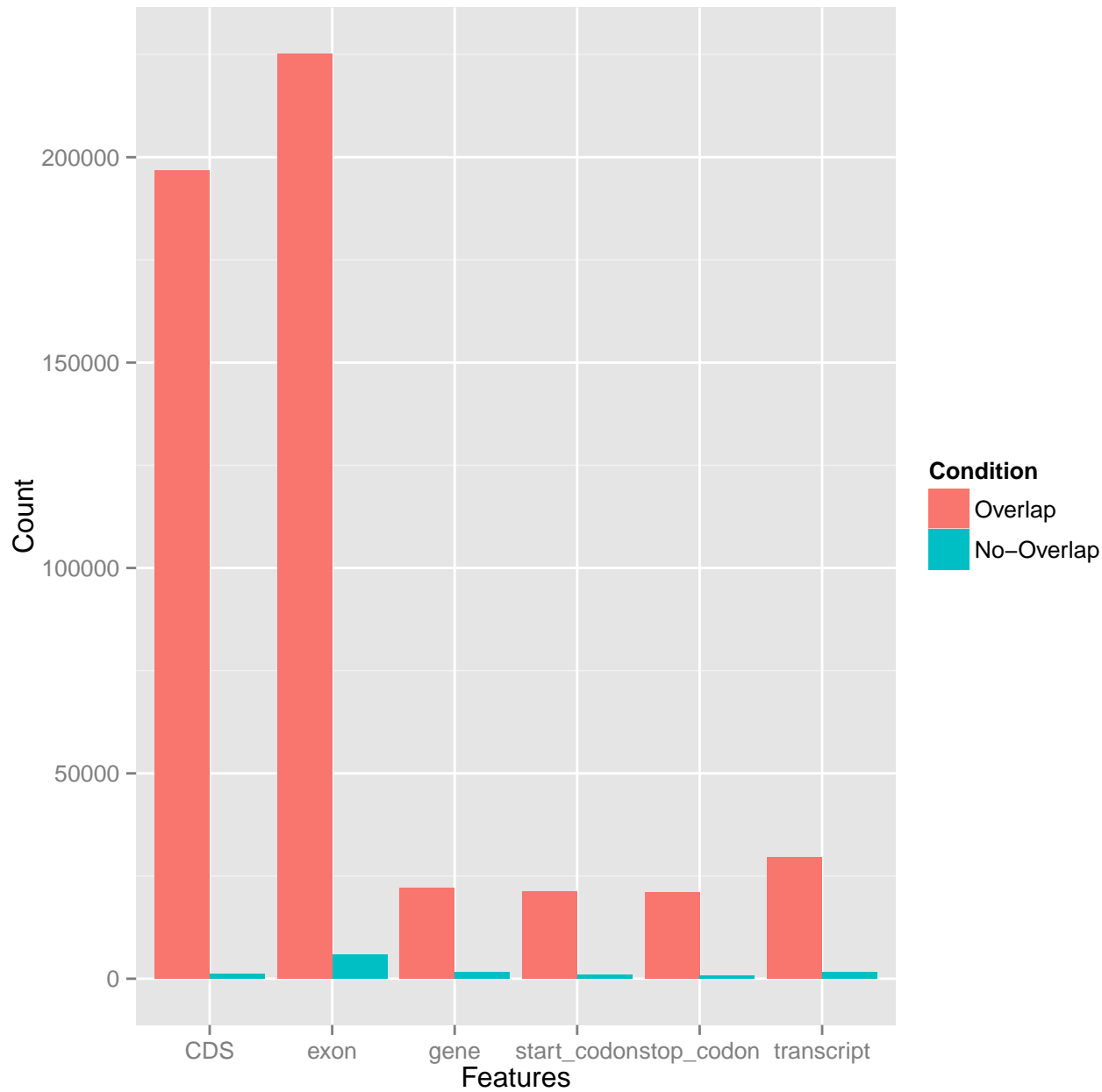
Total Entries



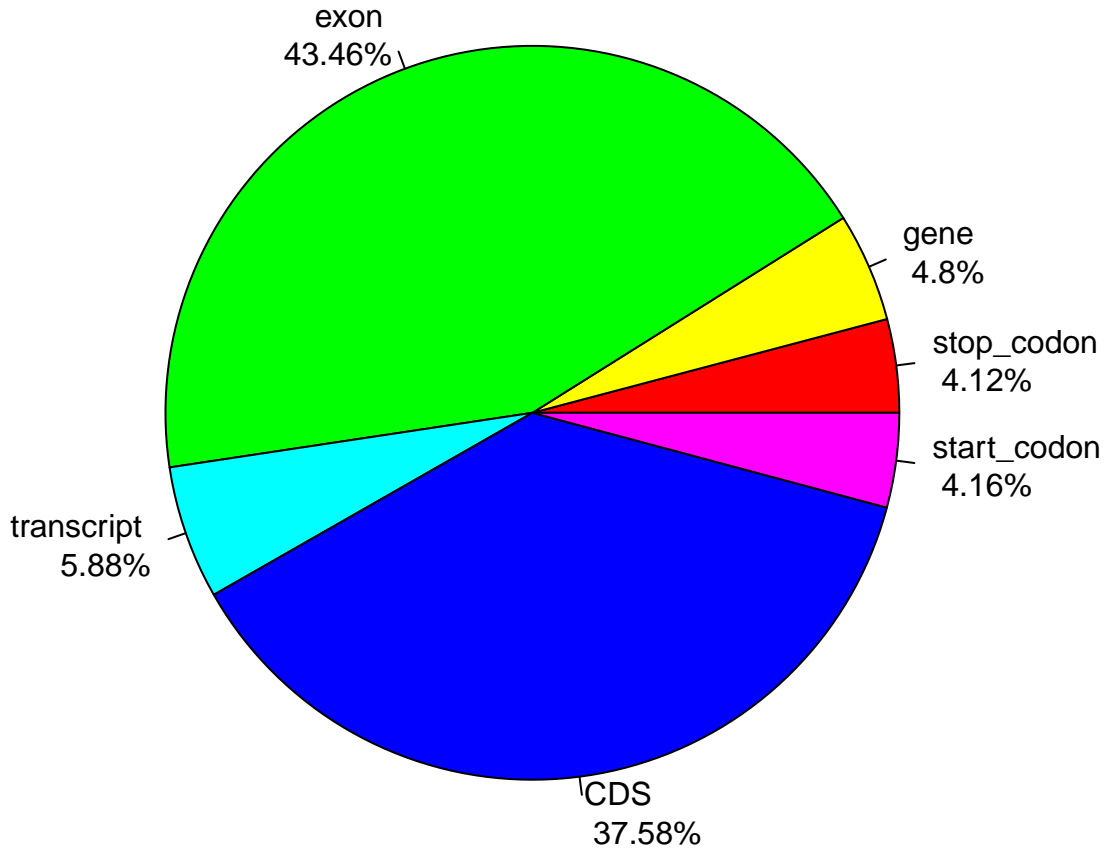
Counts of Different Features



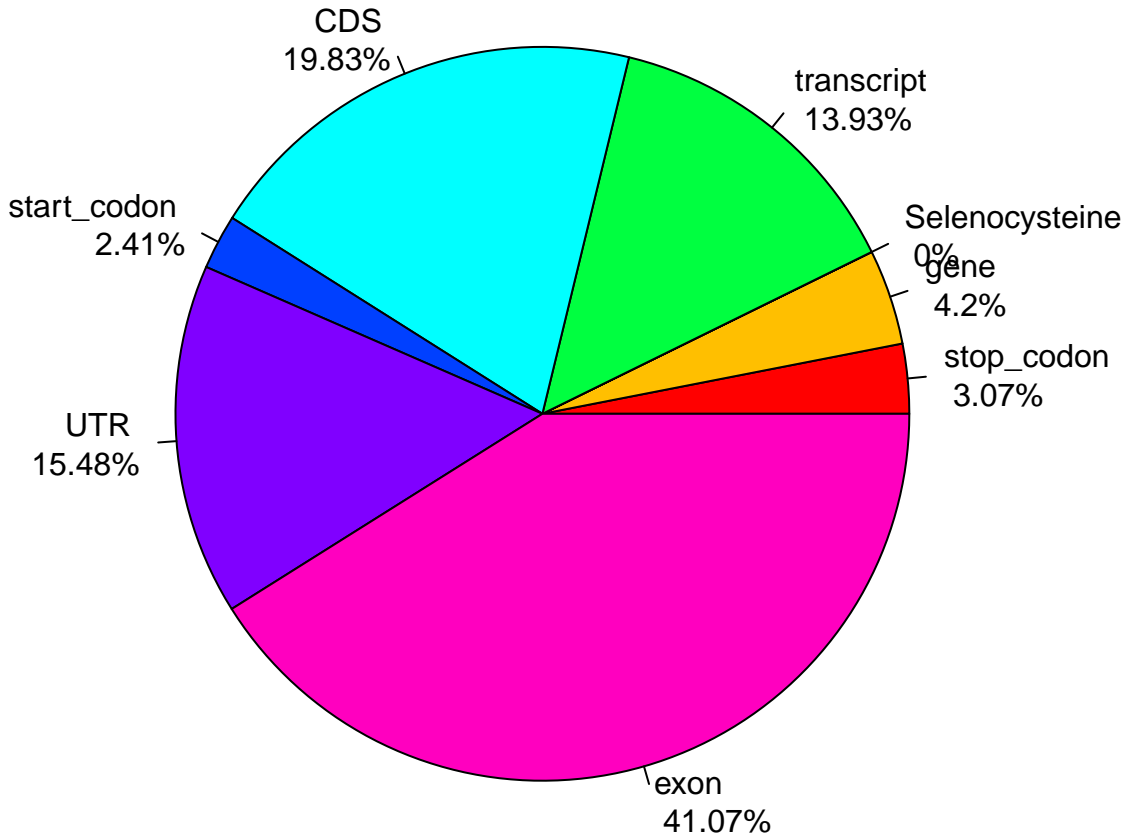
Counts of Different Features



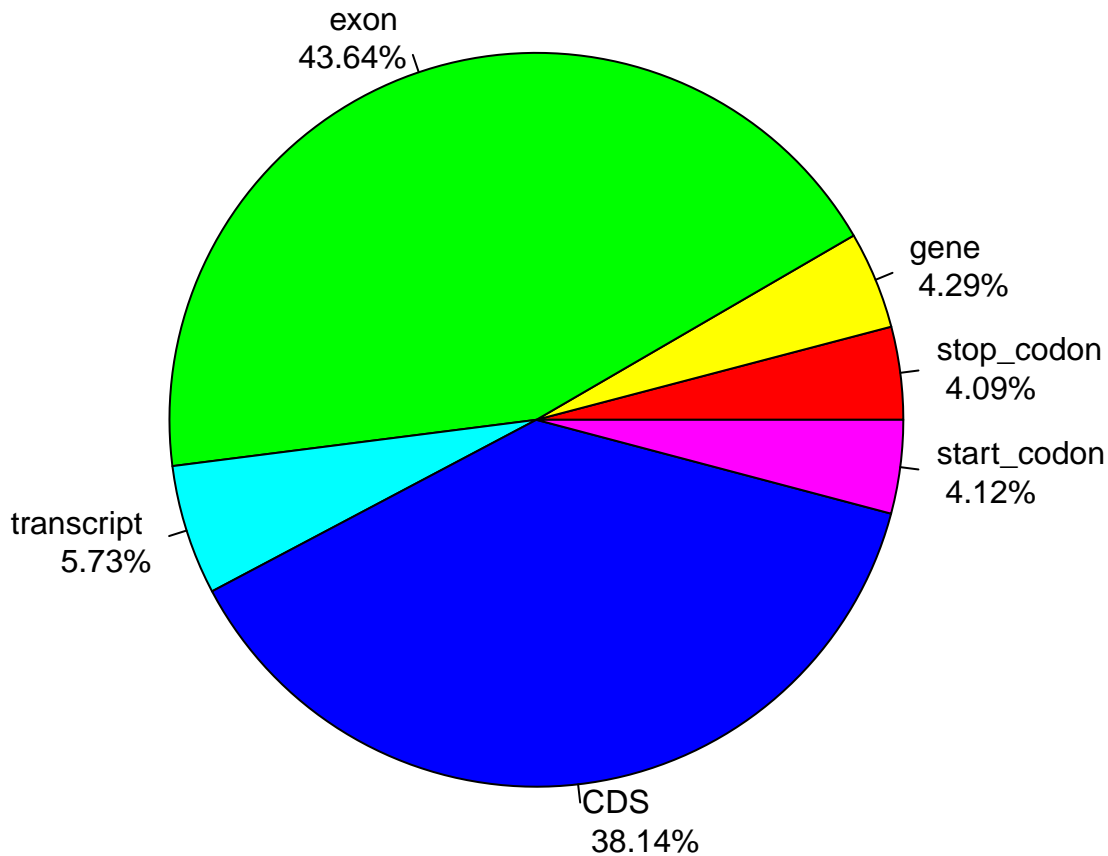
refseq.gtf – Feature Distribution



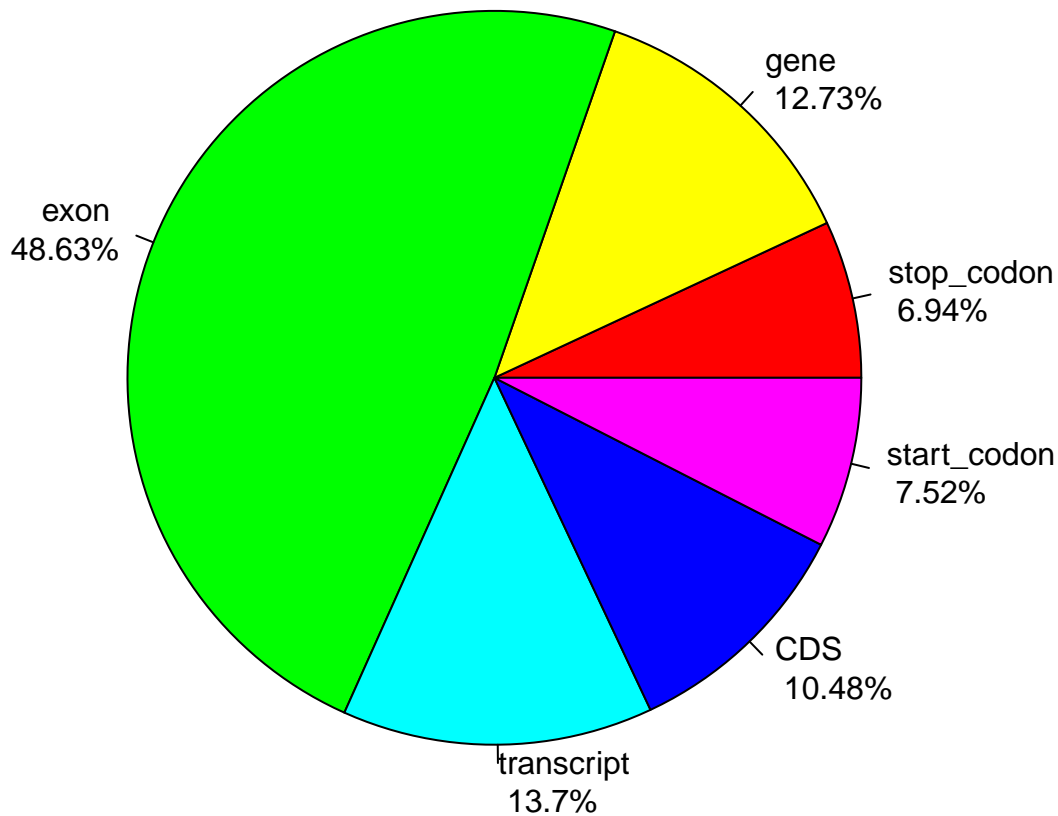
gencode.gtf – Feature Distribution



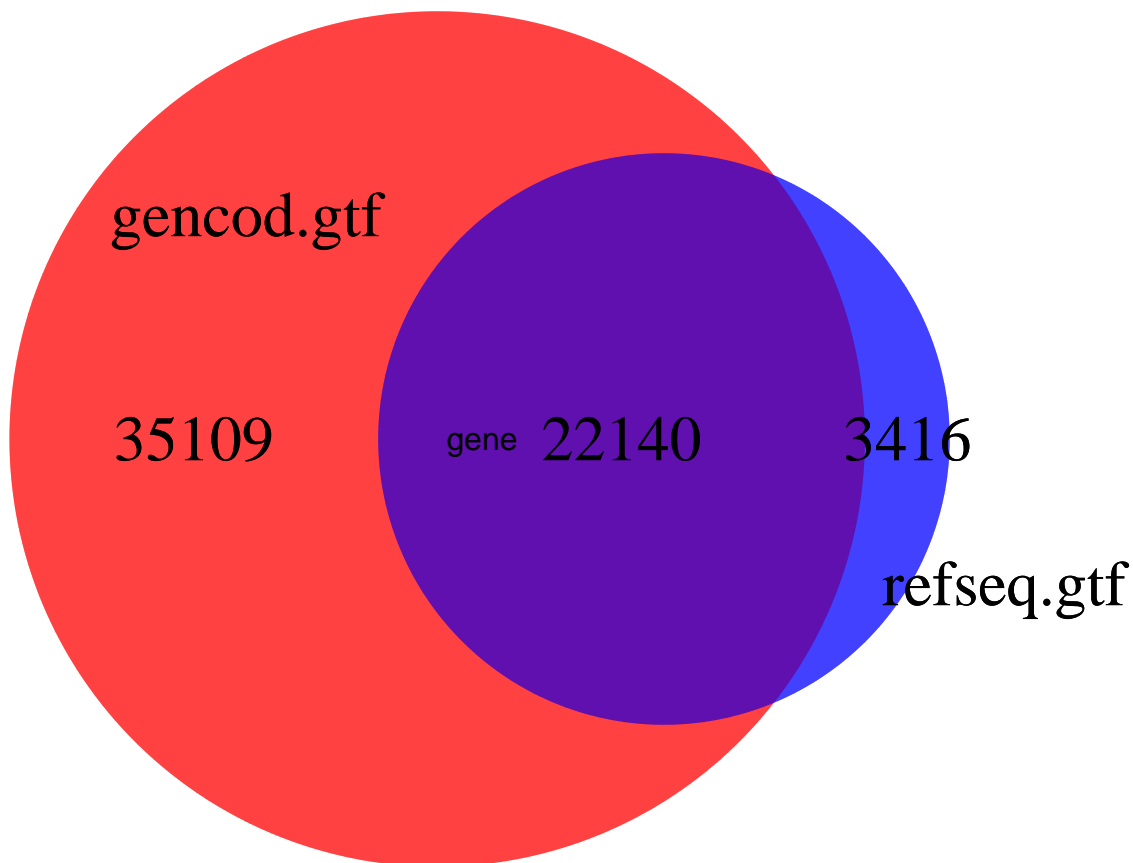
Feature Distribution of Total Overlaps Found

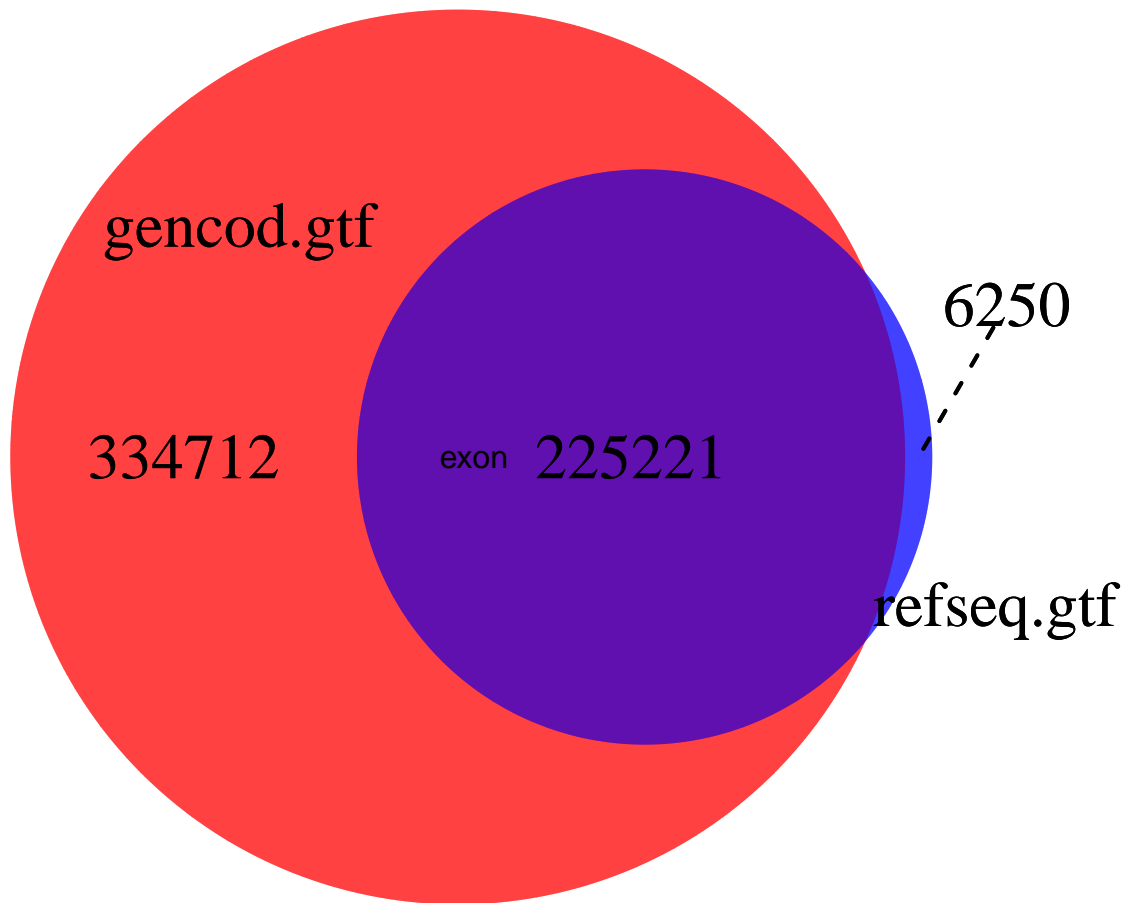


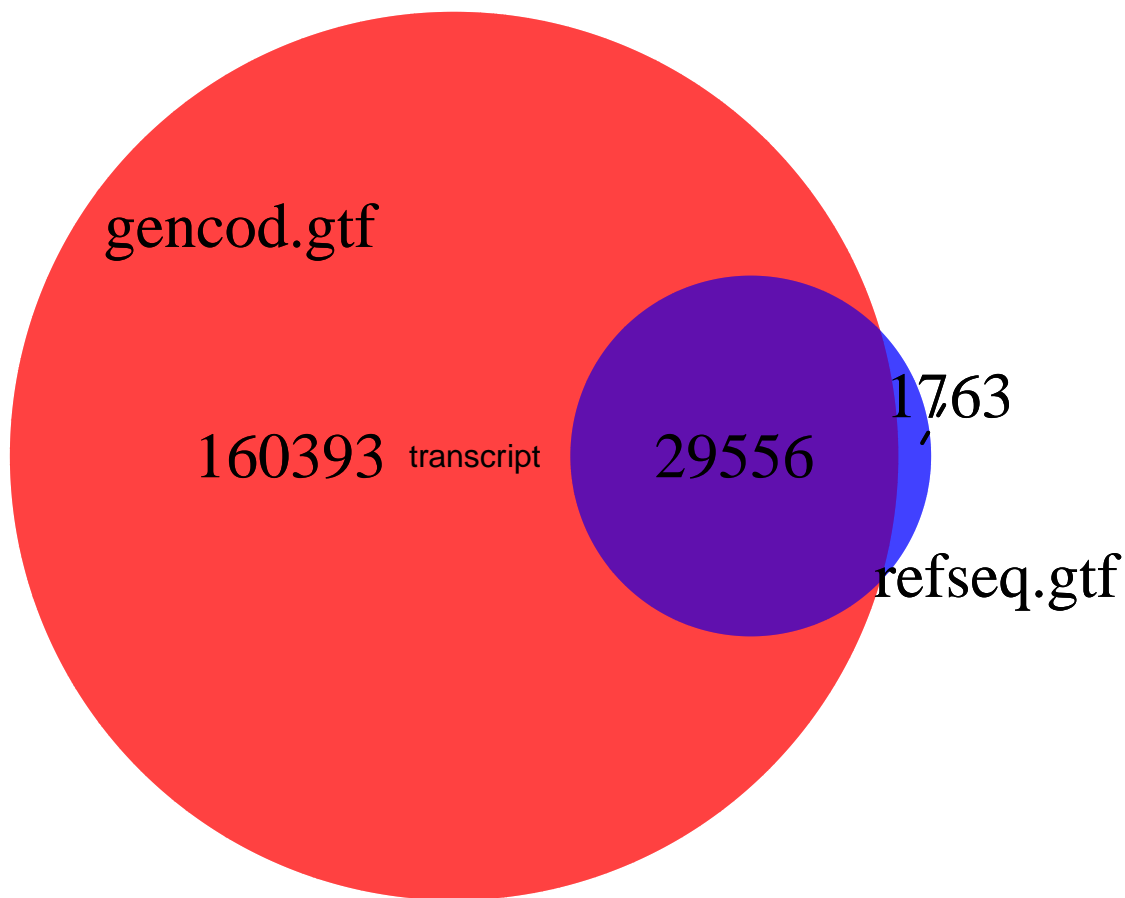
Feature Distribution of Total Non-Overlaps Found

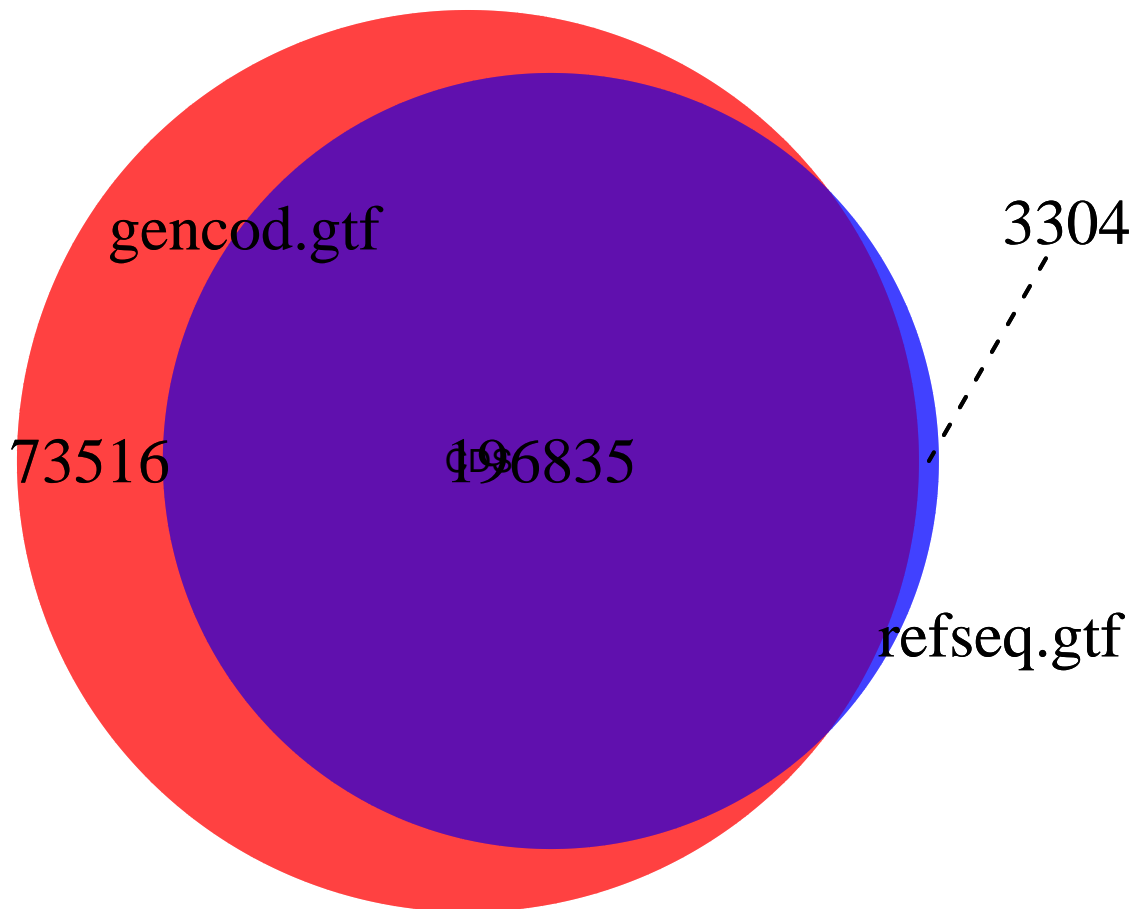


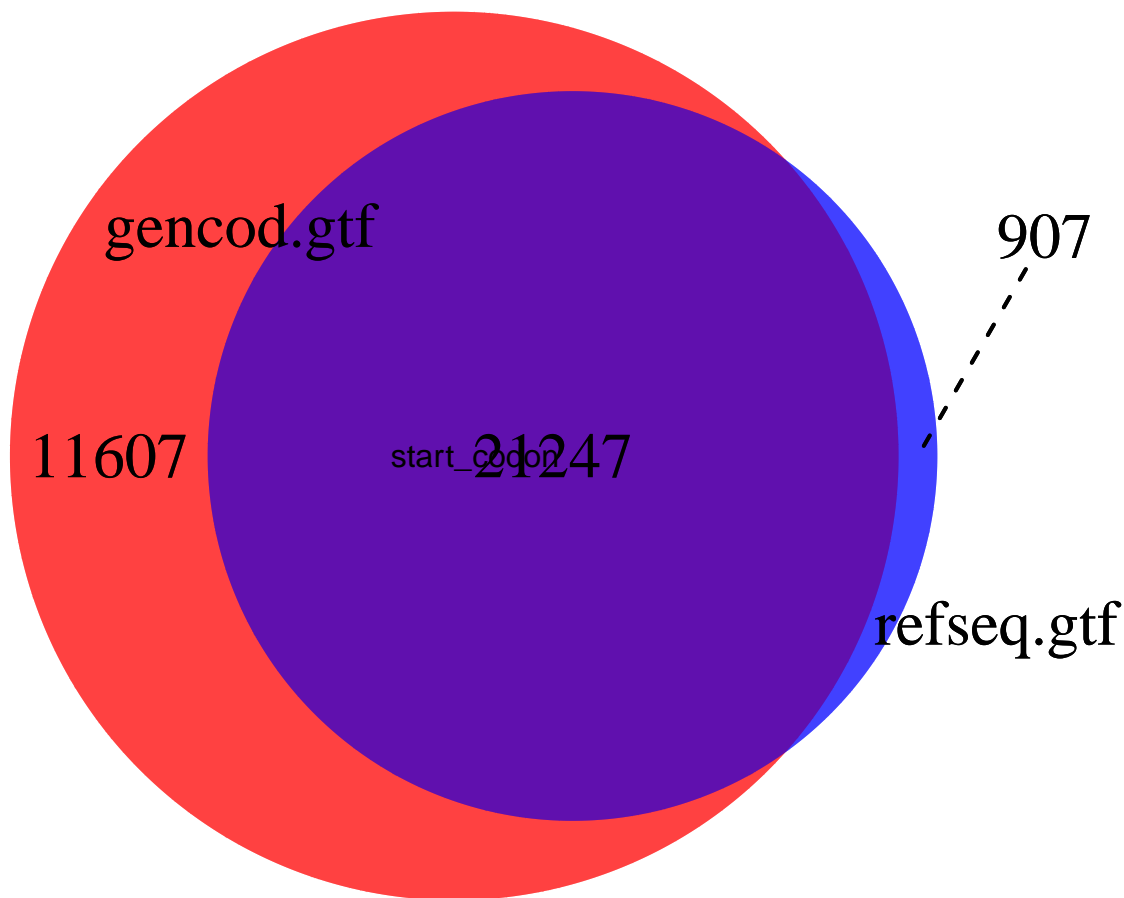


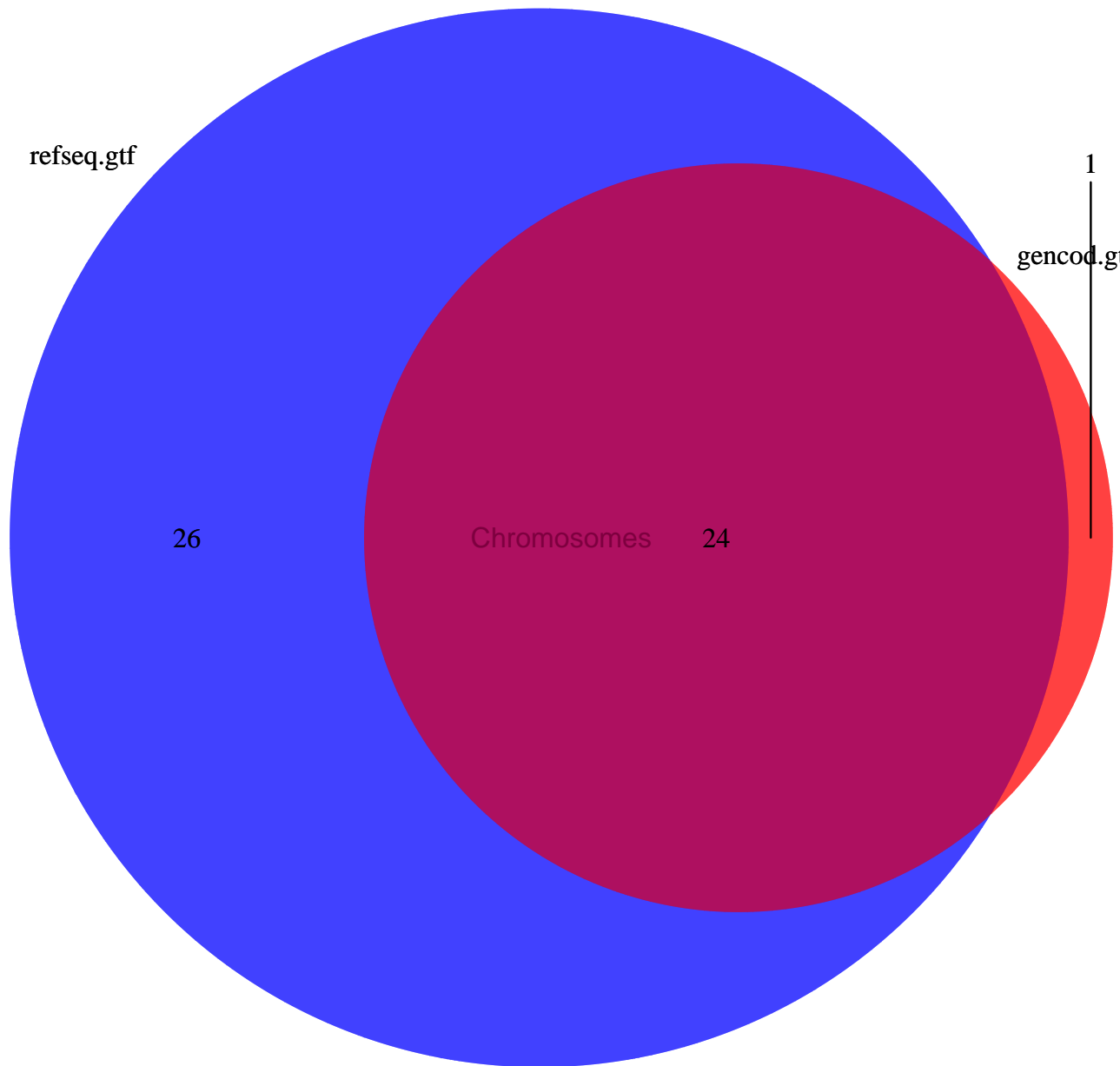




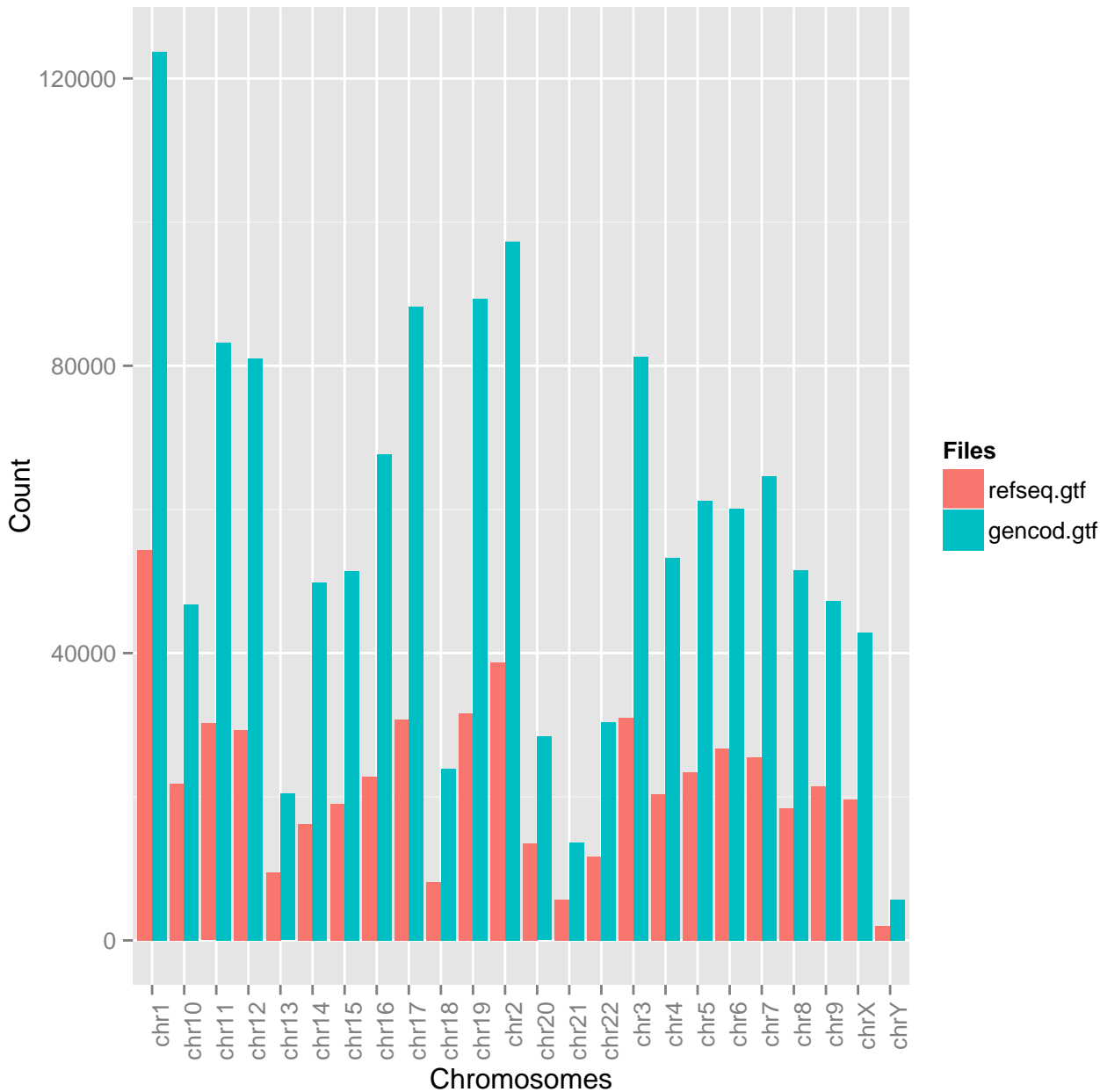




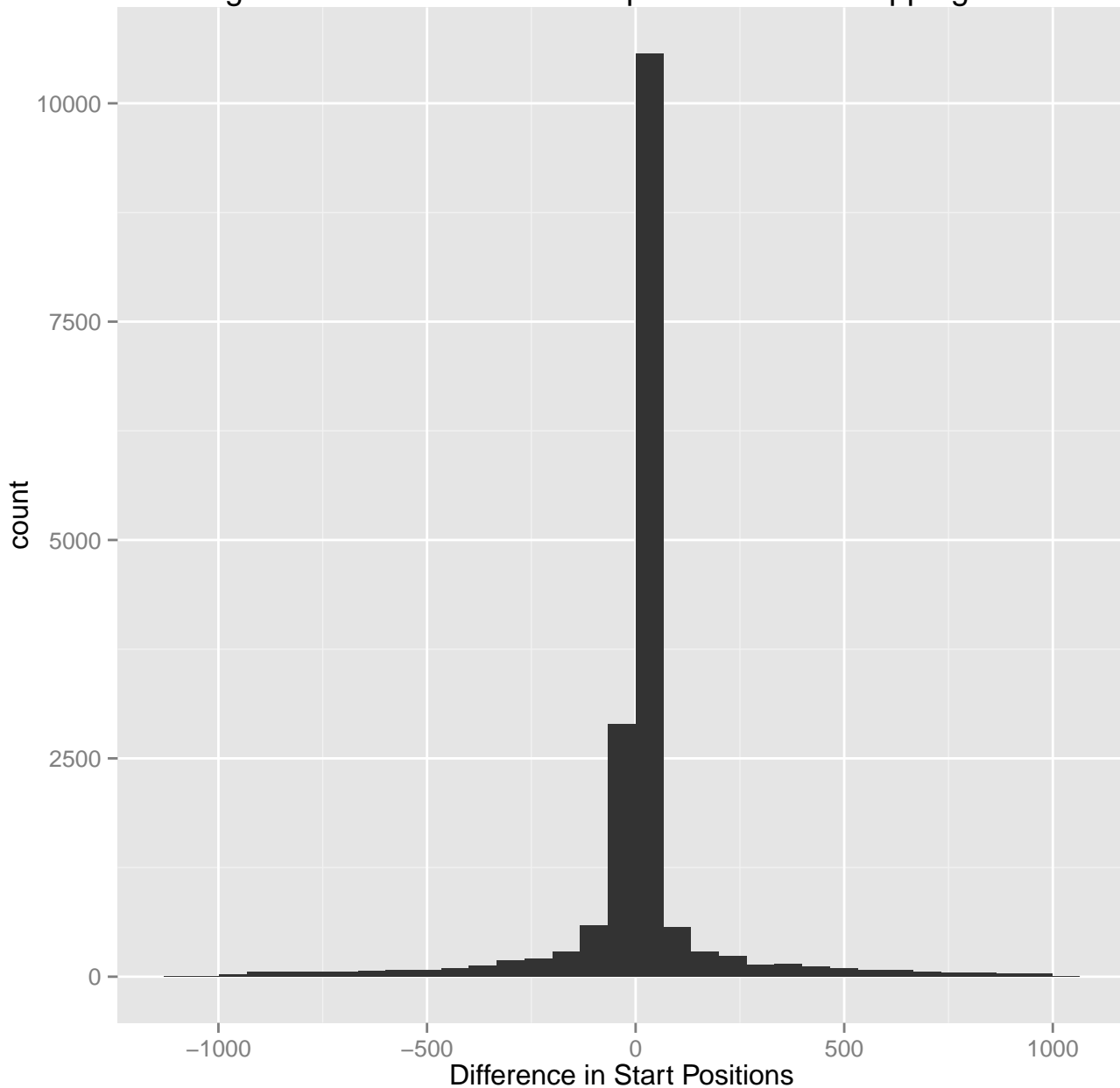




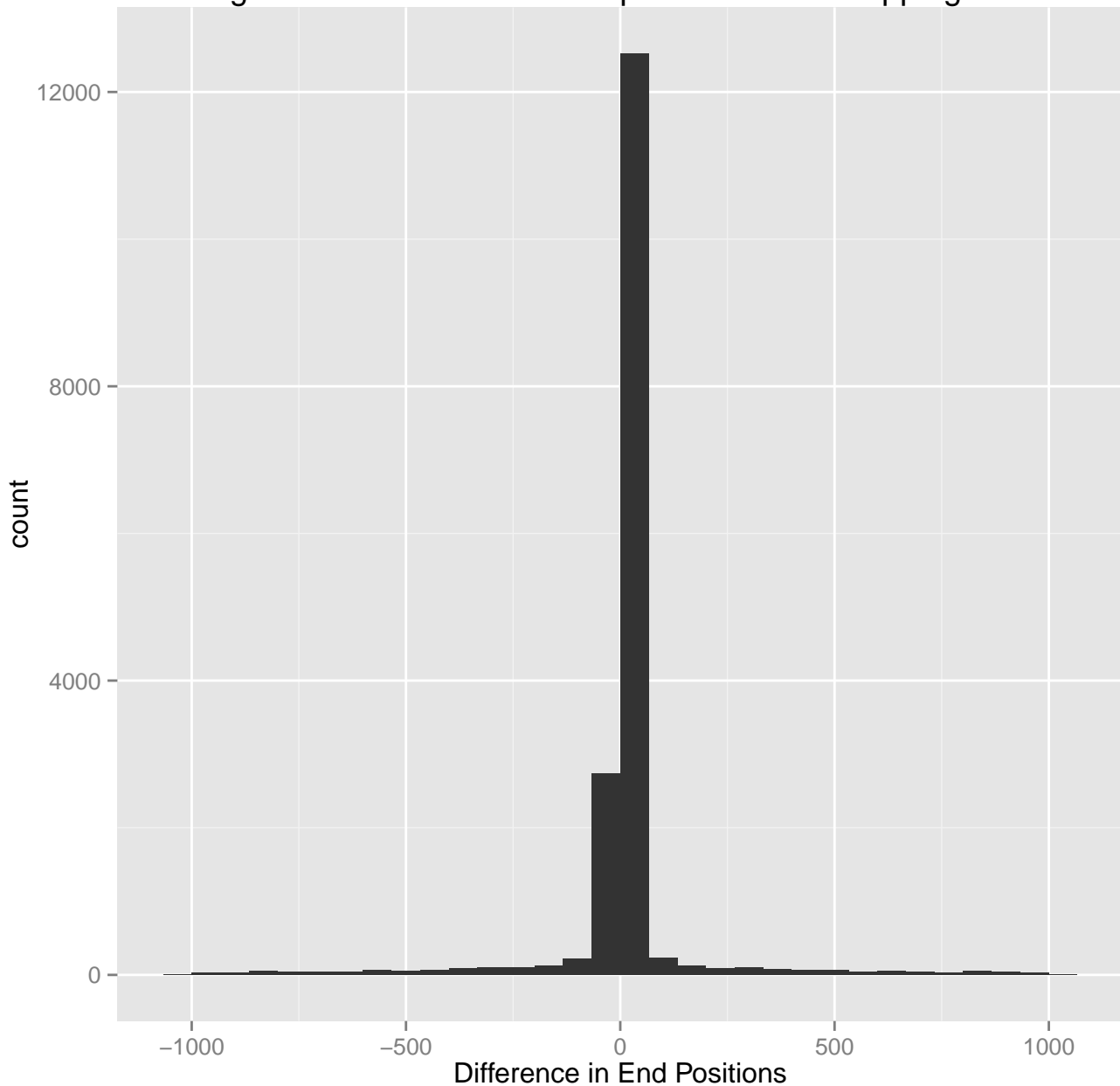
Counts of Common Chromosomes



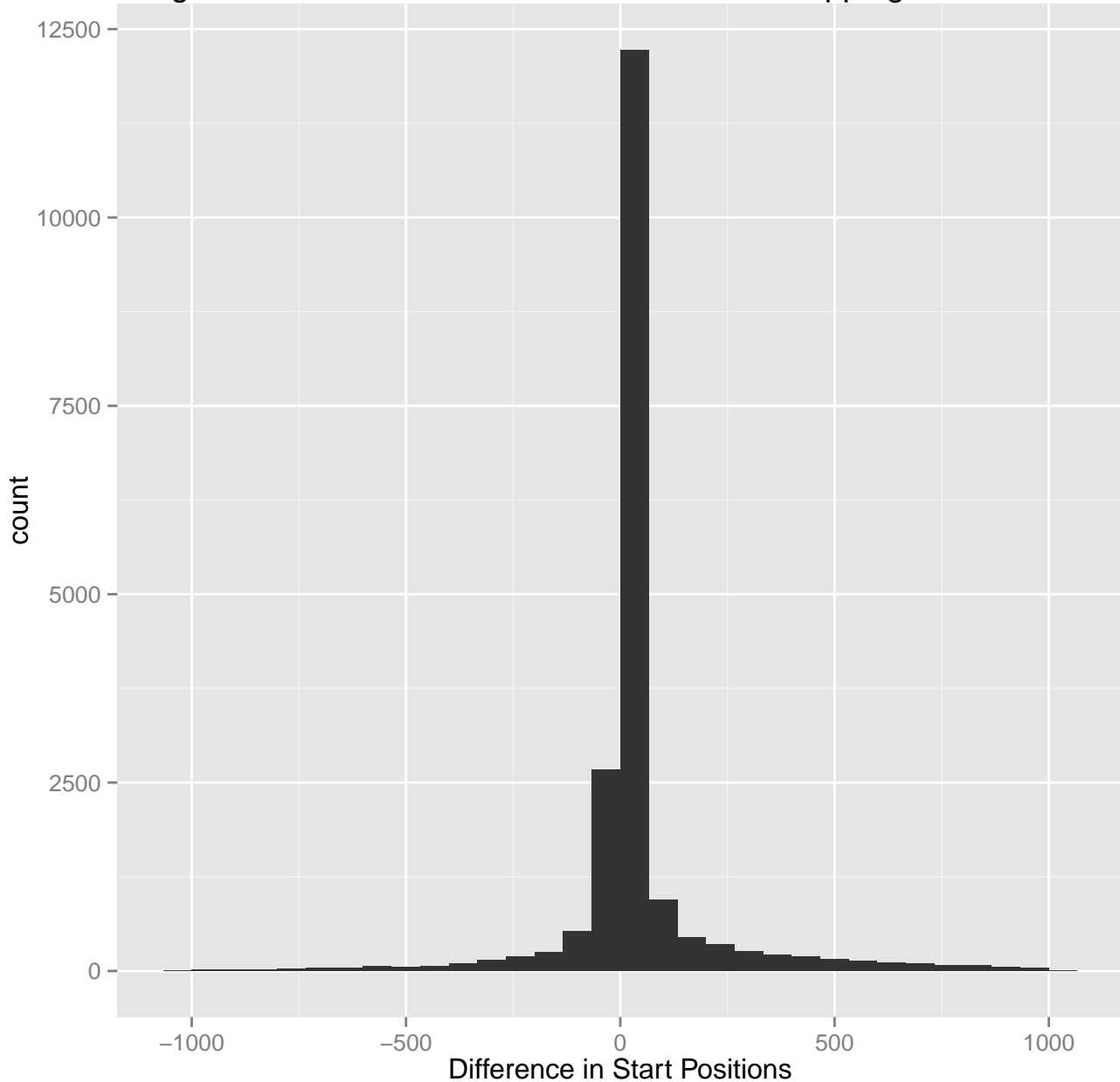
Histogram of Differences in start positions of overlapping GENES



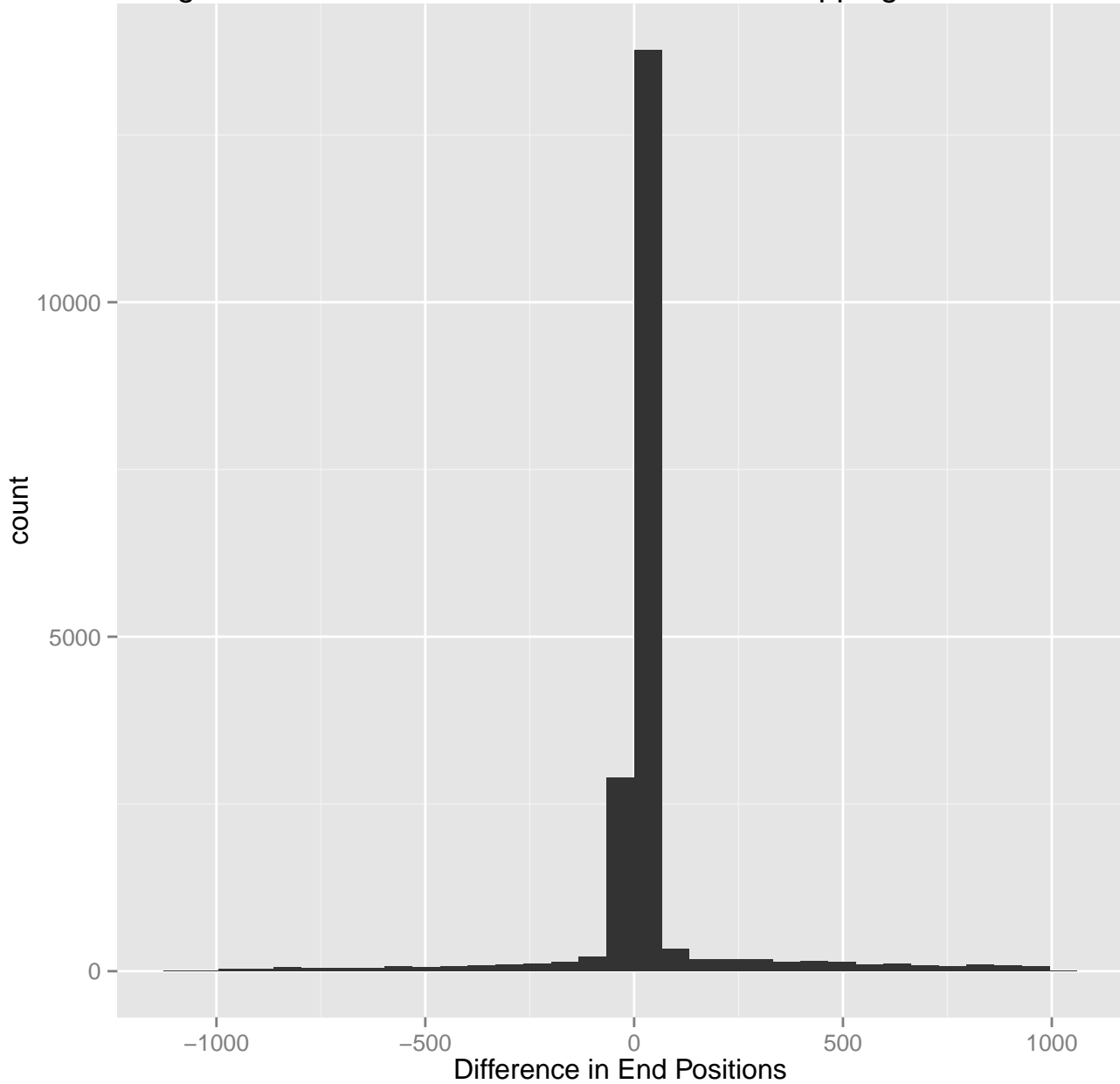
Histogram of Differences in end positions of overlapping GENES



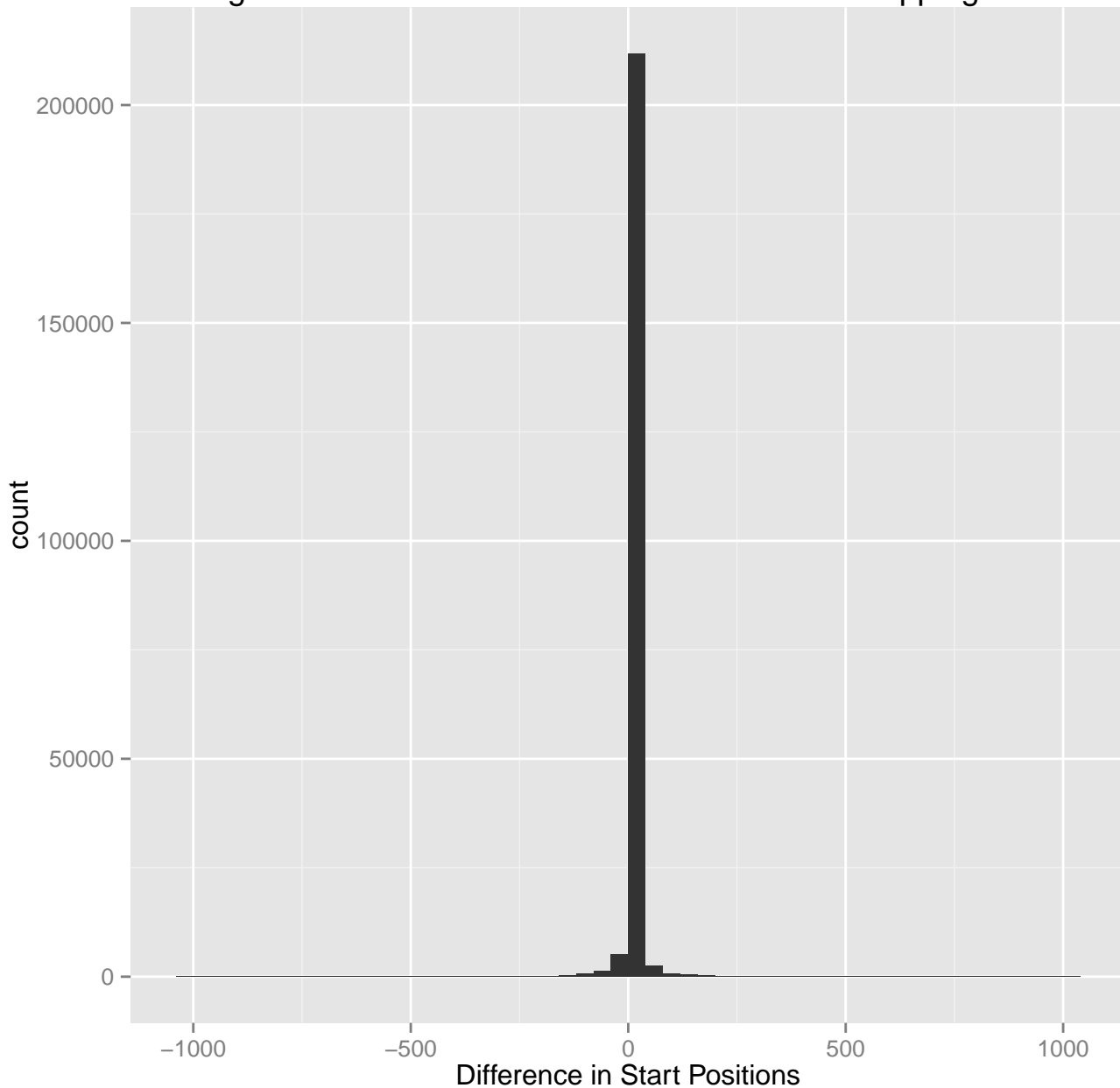
Histogram of Differences in Start Position of overlapping TRANSCRIPTS



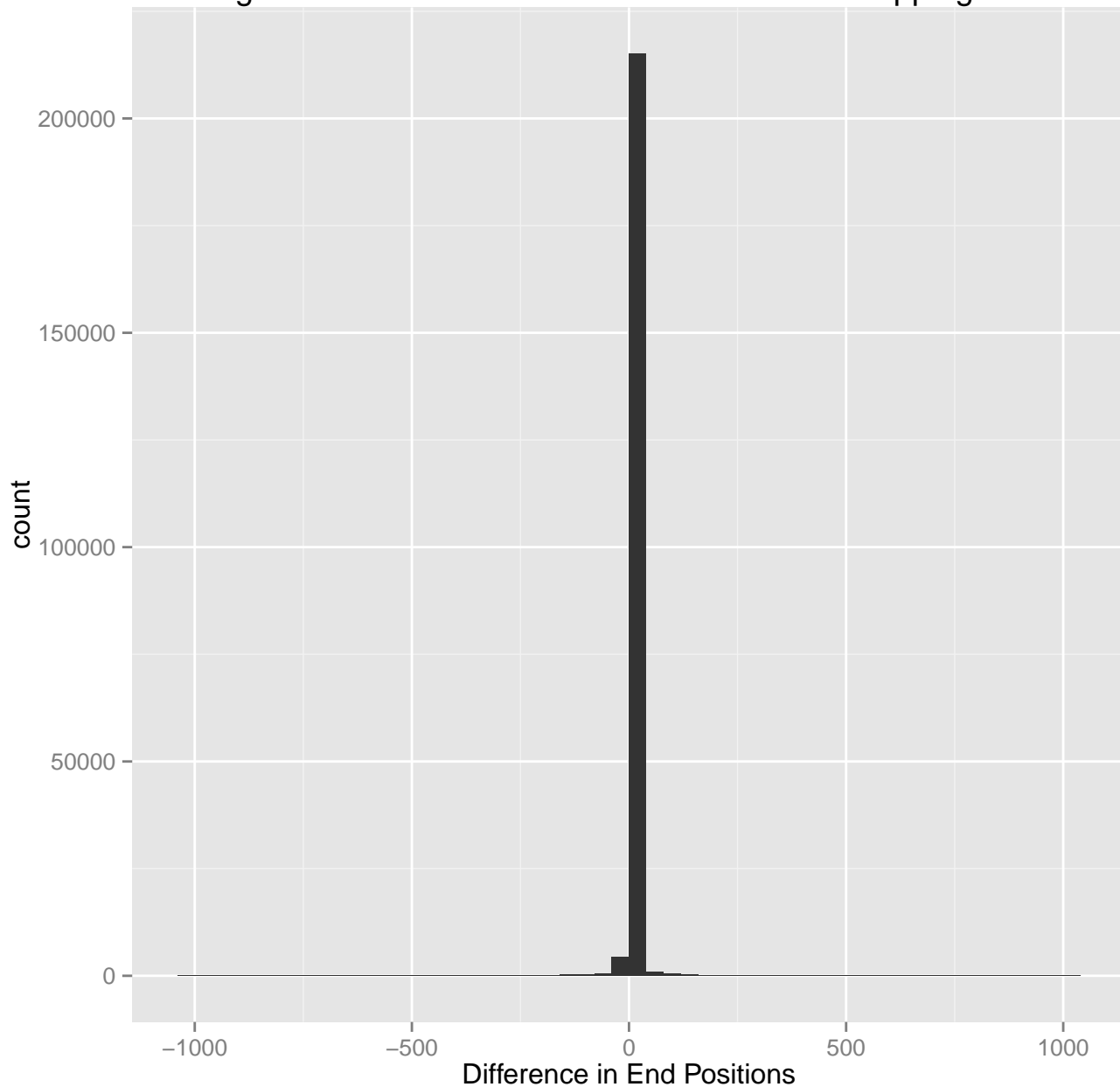
Histogram of Differences in End Position of overlapping TRANSCRIPTS



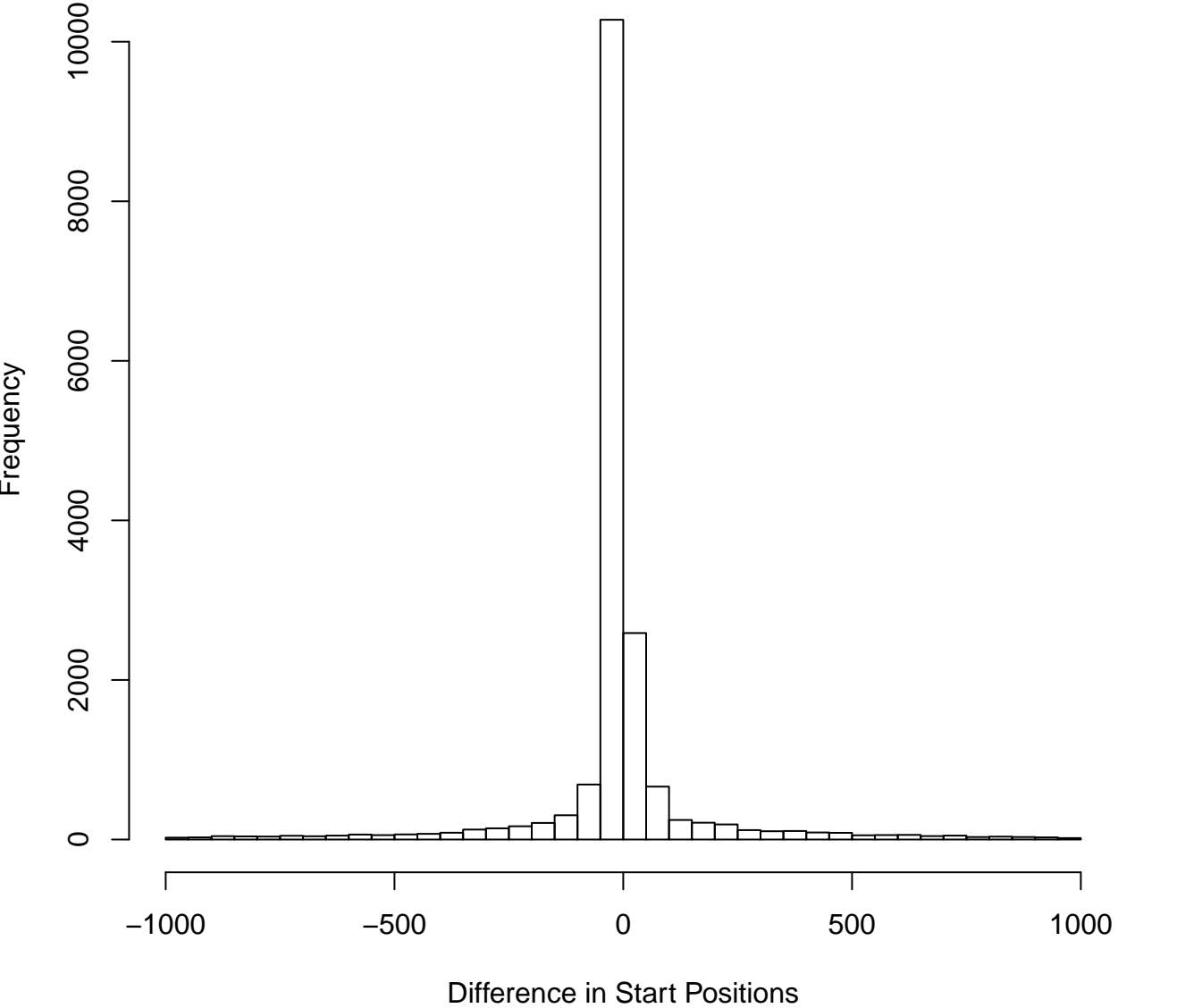
Histogram of Differences in Start Positions of overlapping EXONS



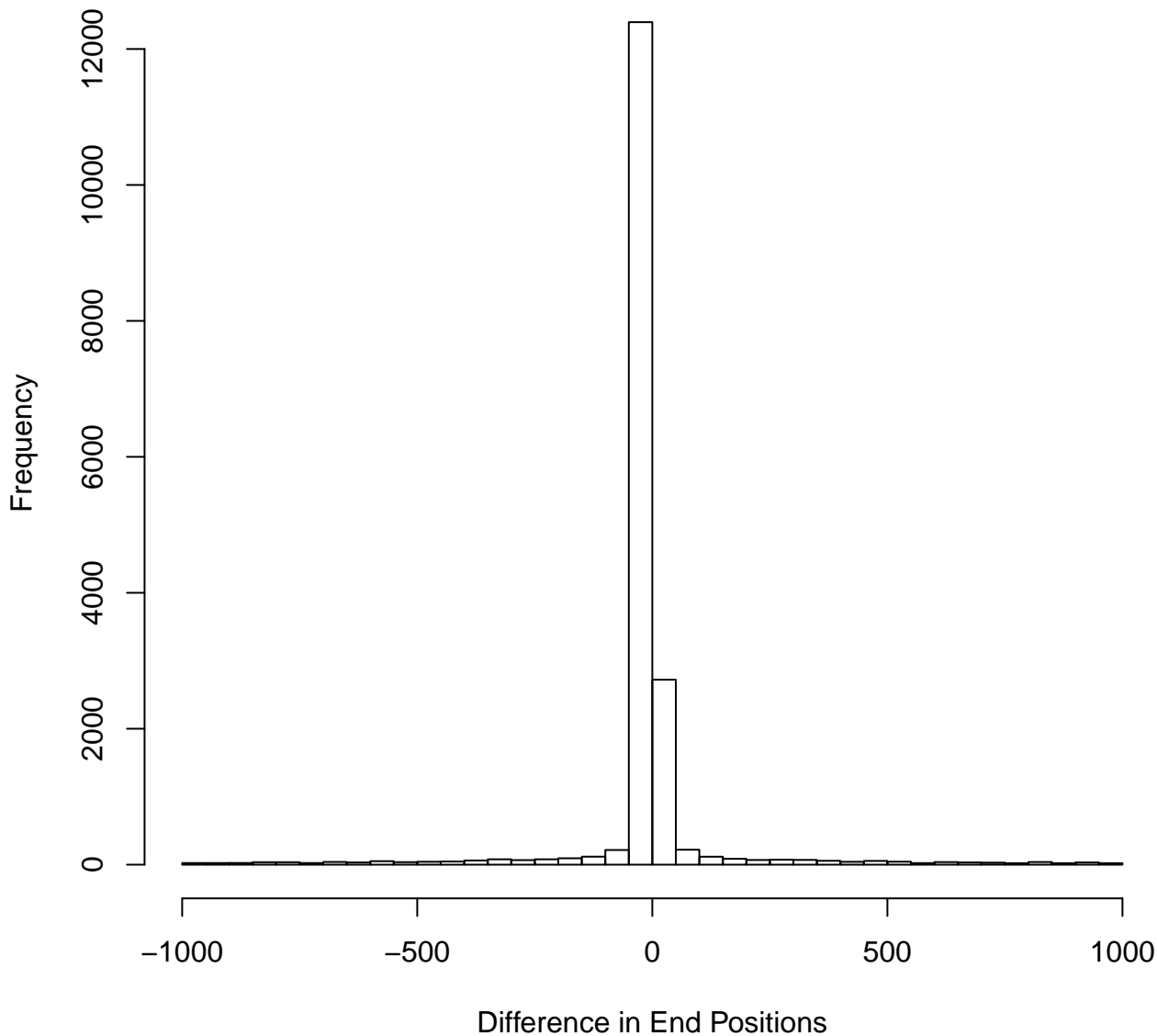
Histogram of Differences in End Positions of overlapping EXONS



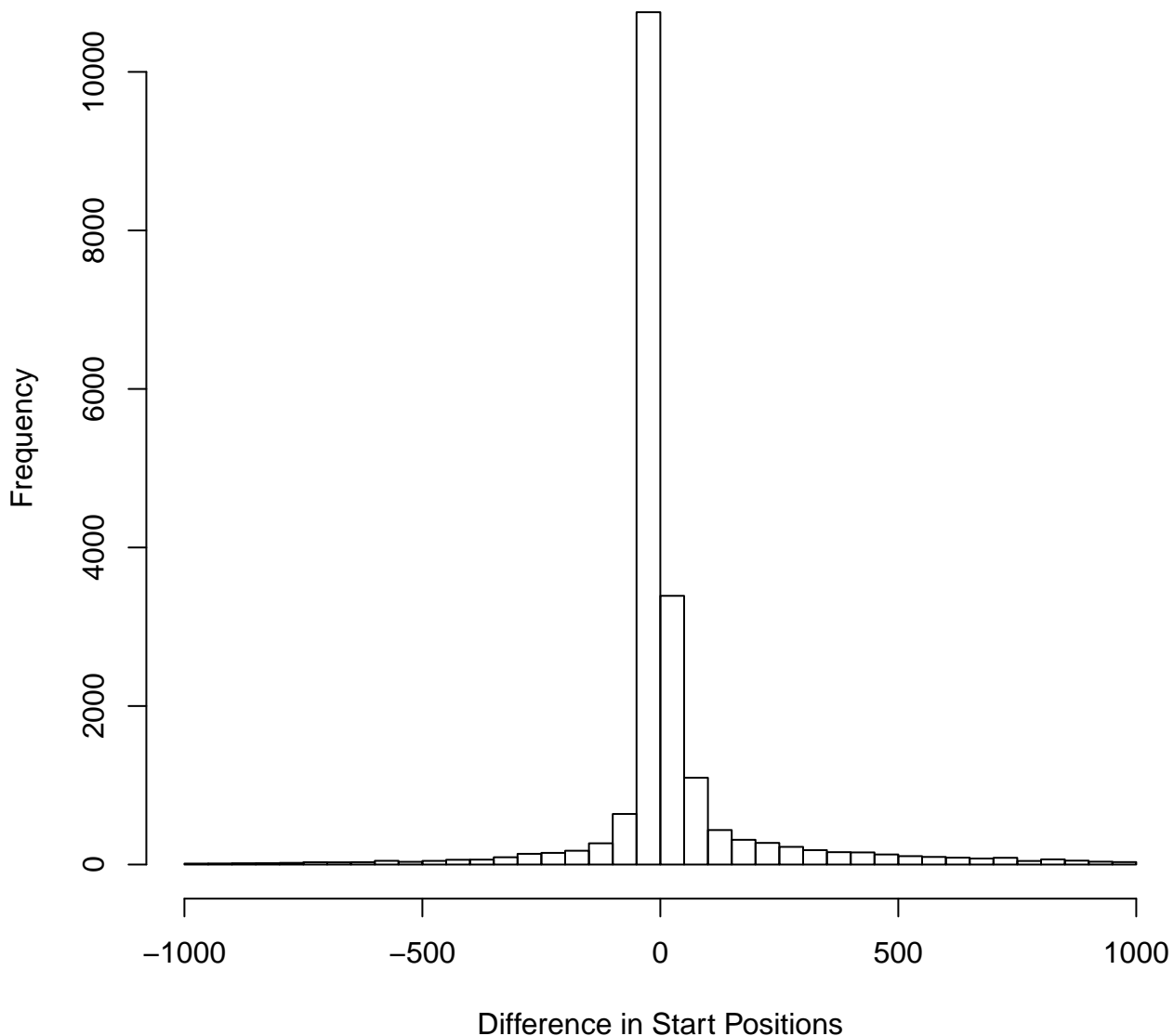
Histogram of Differences in Start Positions of overlapping GENES



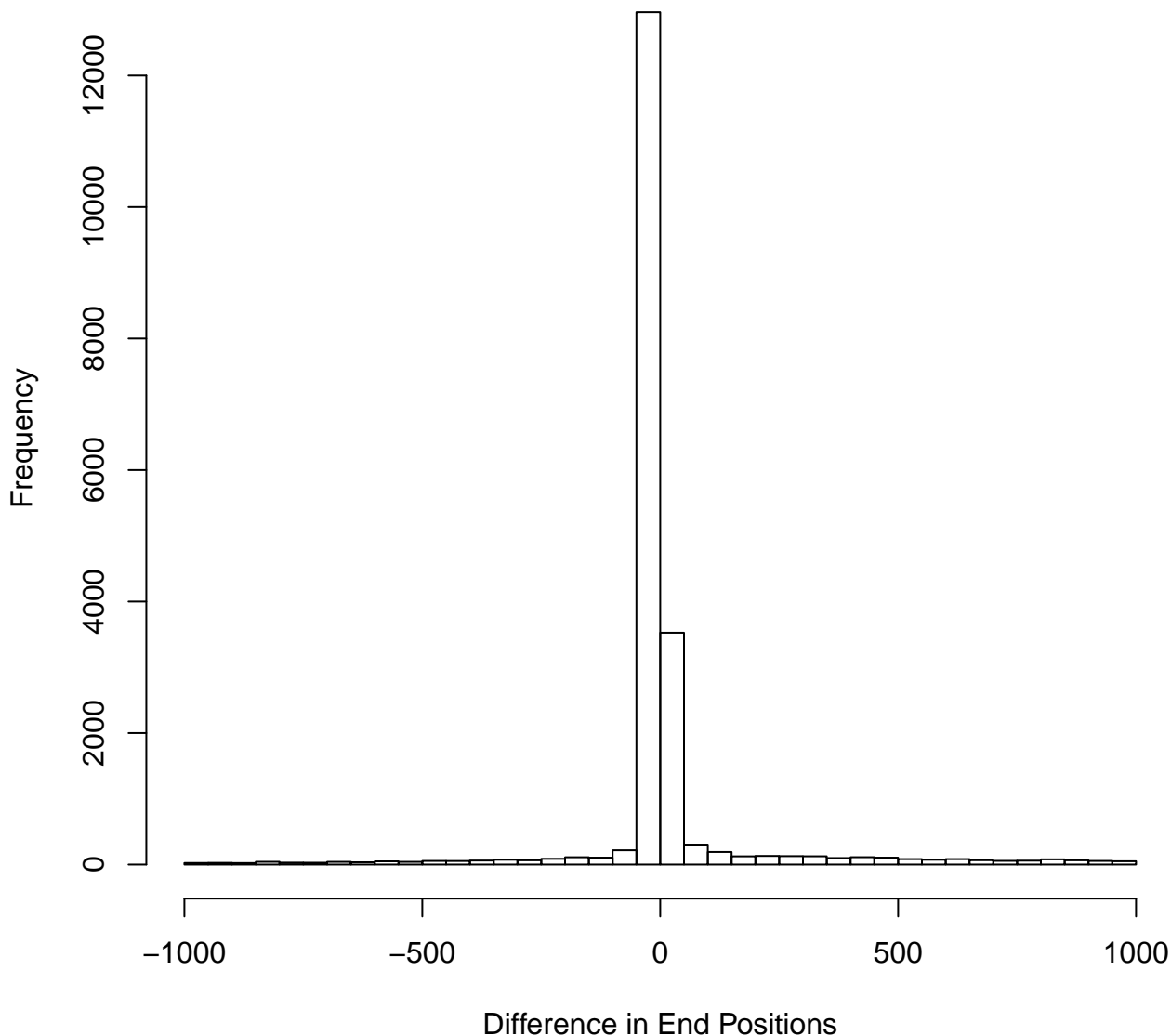
Histogram of Differences in End Positions of overlapping GENES



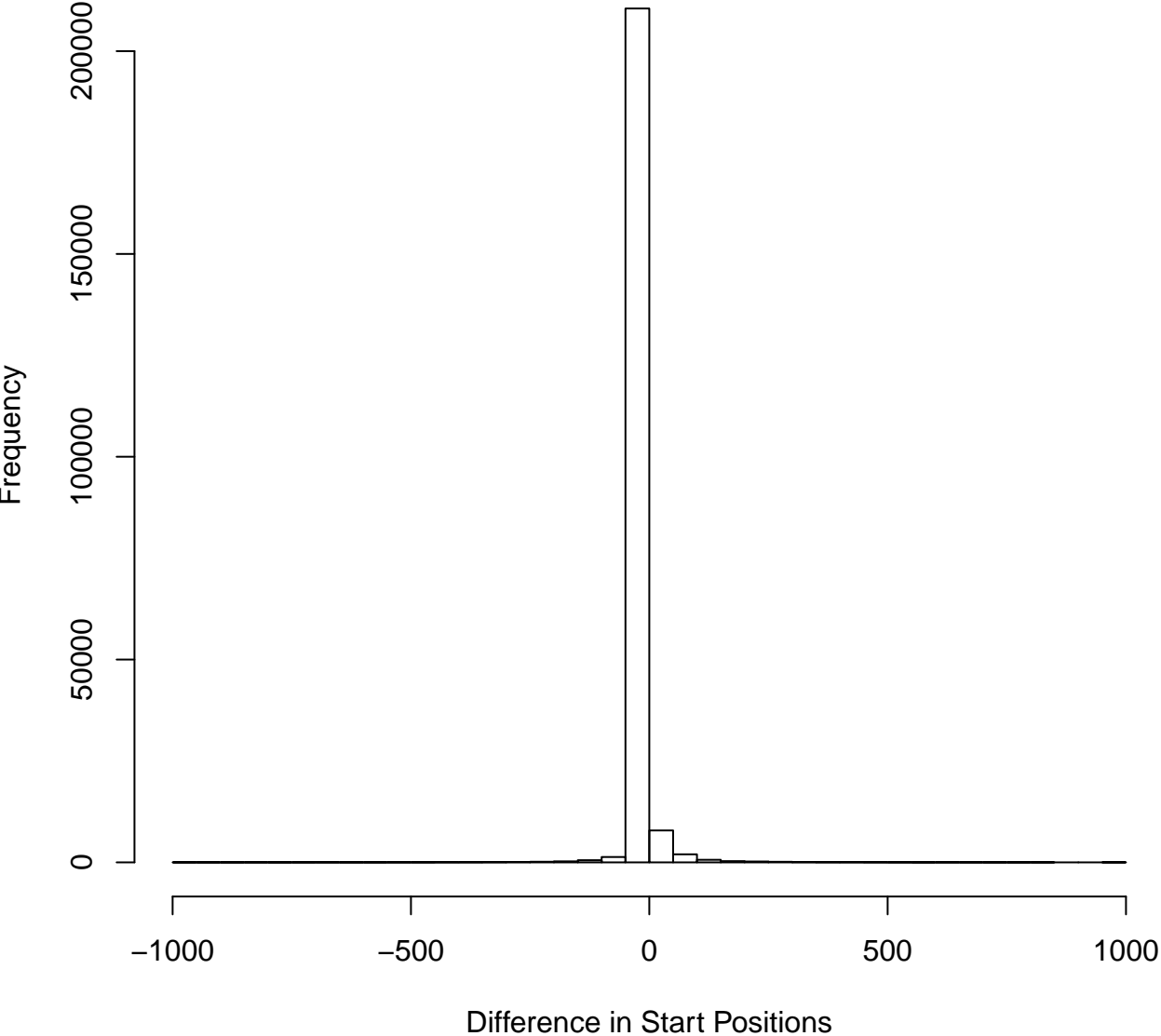
Histogram of Differences in Start Positions of overlapping TRANSCRIPTS



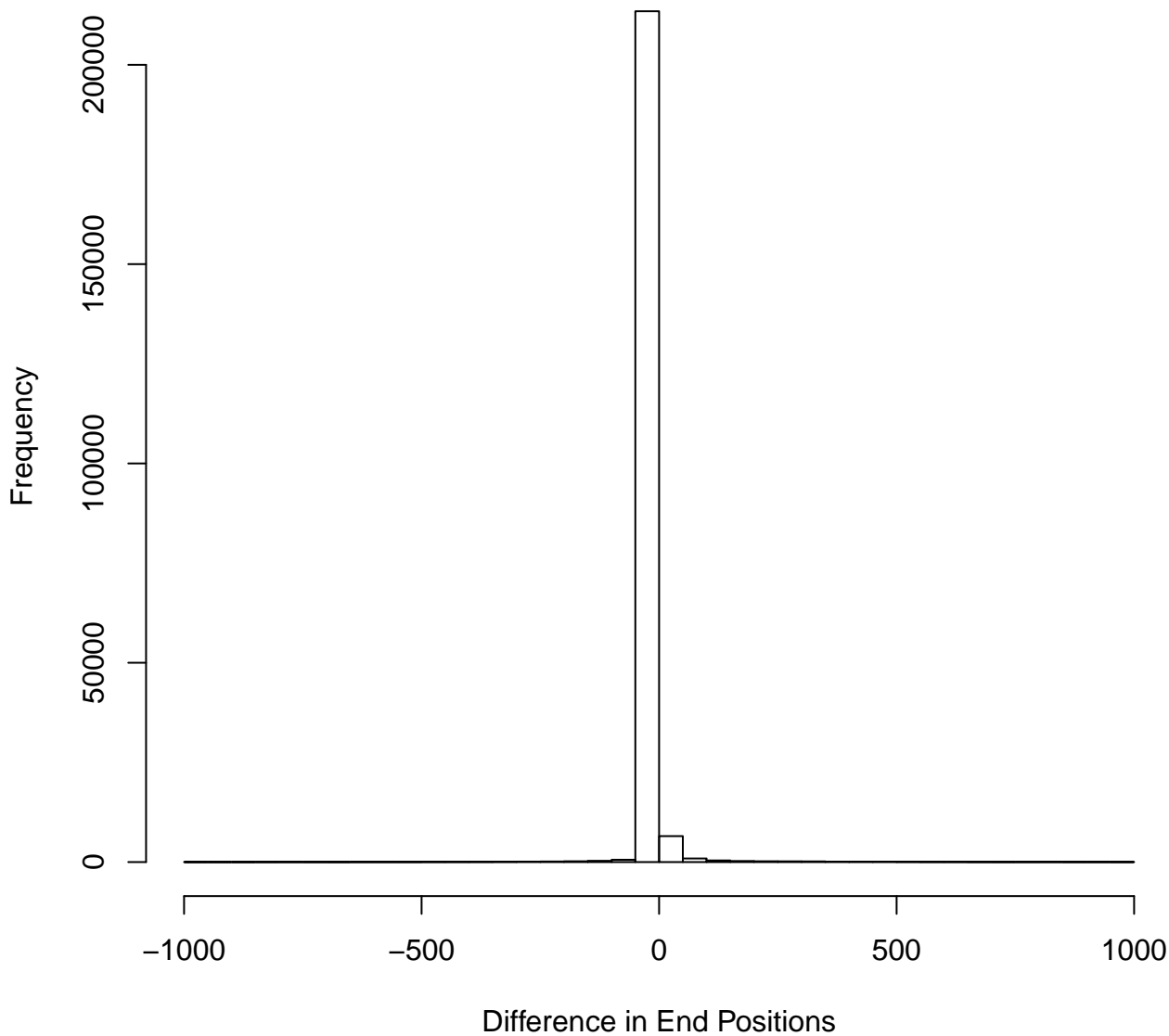
Histogram of Differences in End Positions of overlapping TRANSCRIPTS



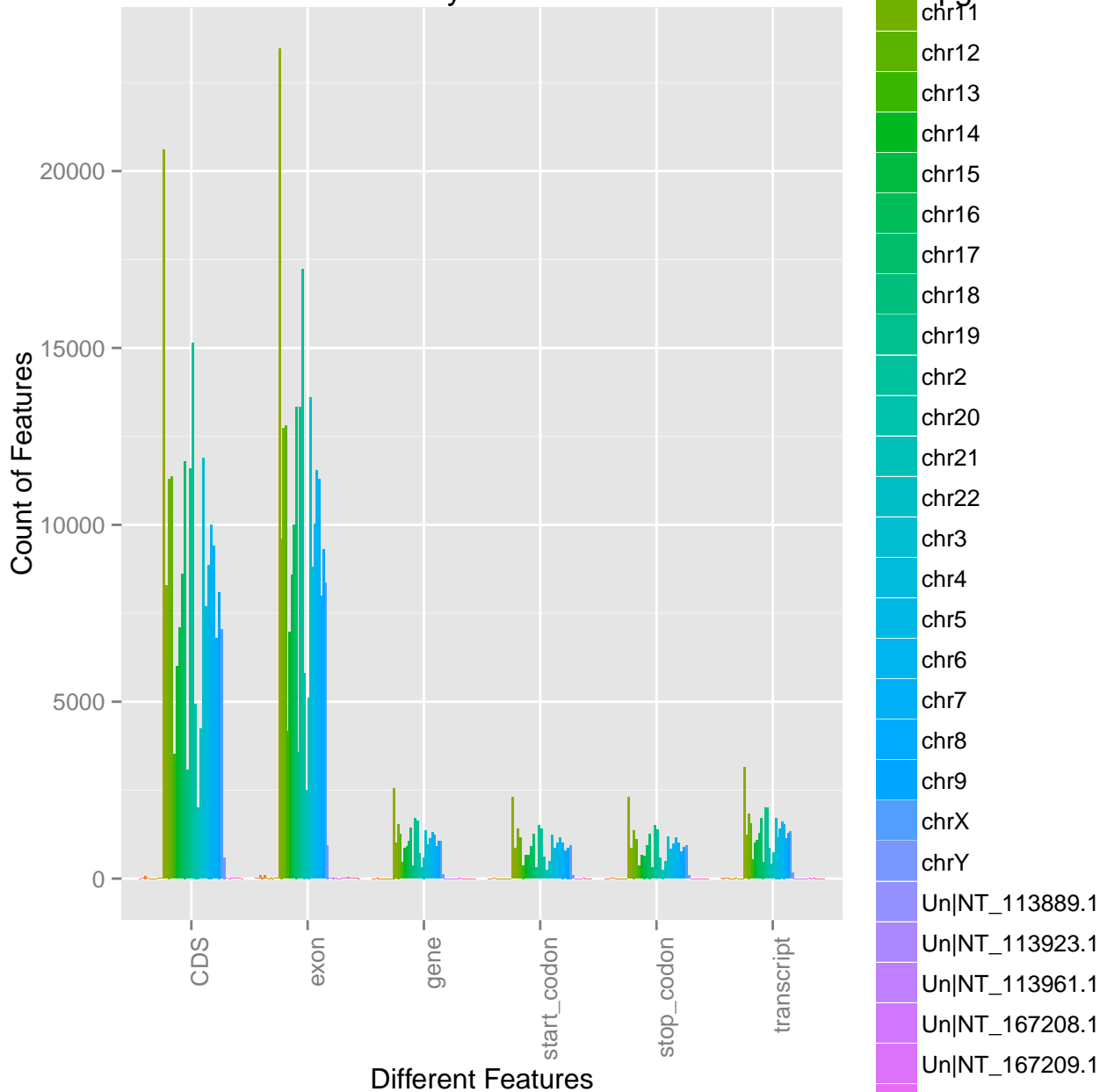
Histogram of Differences in Start Positions of overlapping EXONS



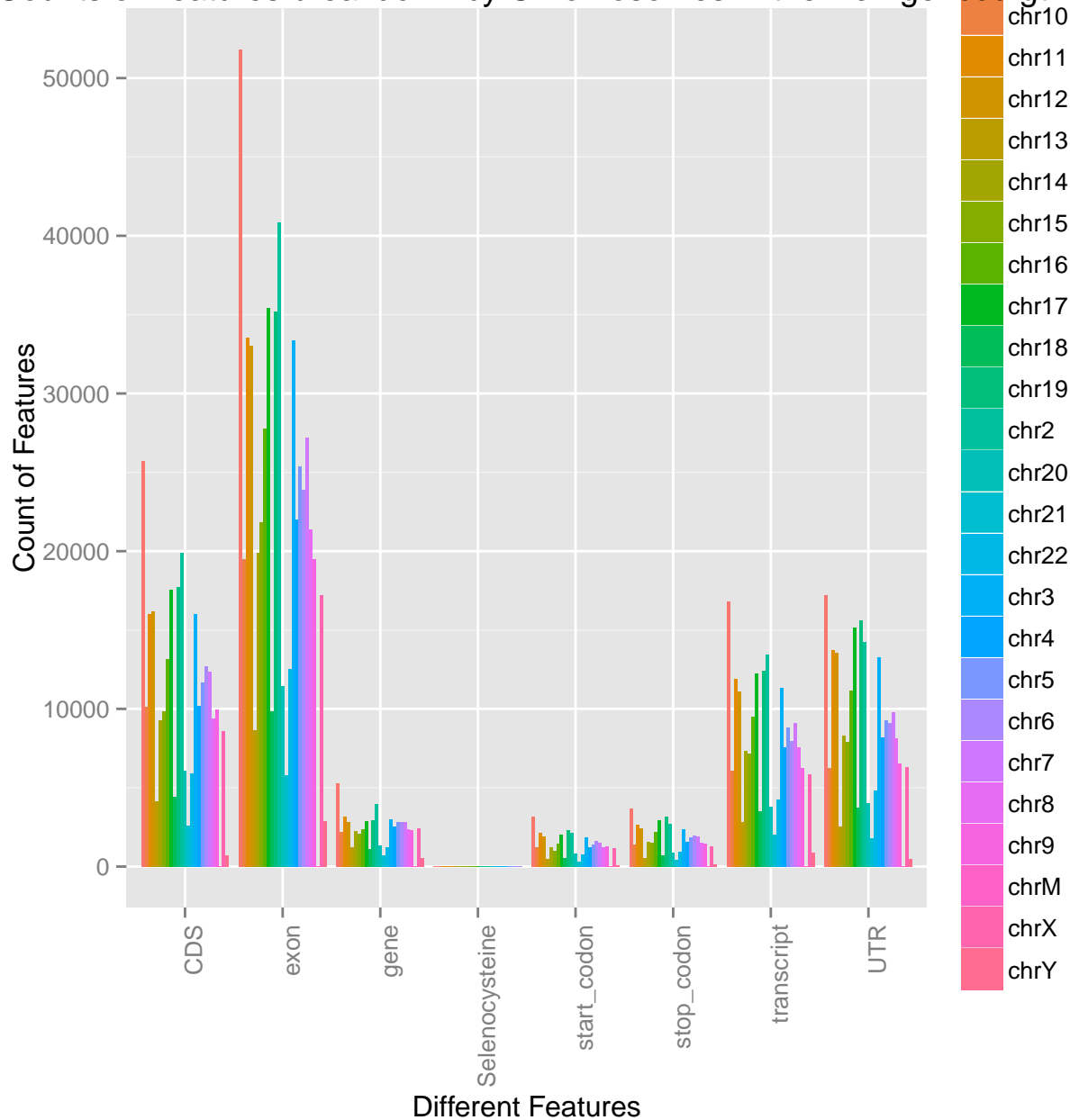
Histogram of Differences in End Positions of overlapping EXONS



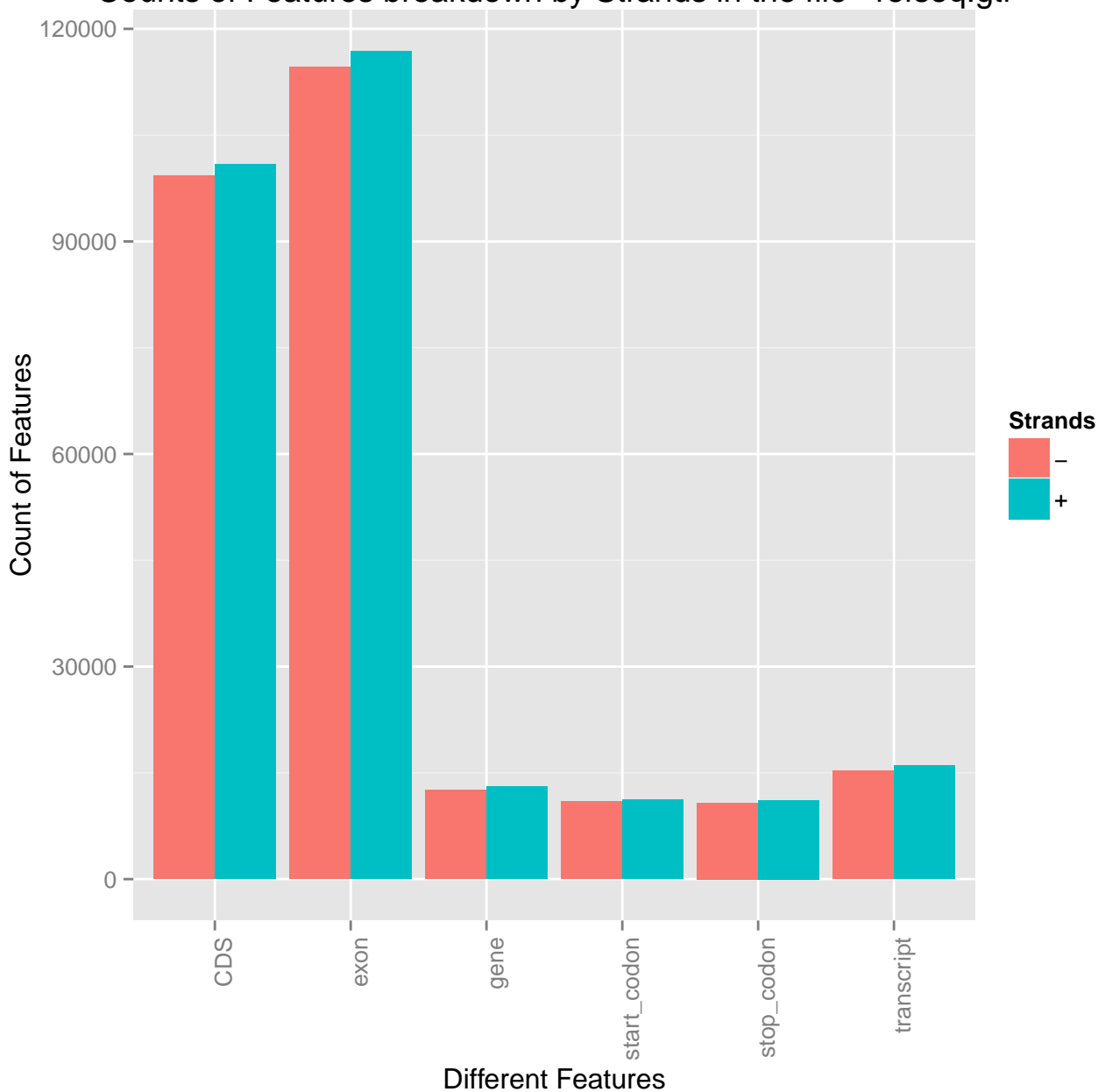
Counts of Features breakdown by Chromosomes in the file– refseq.gtf



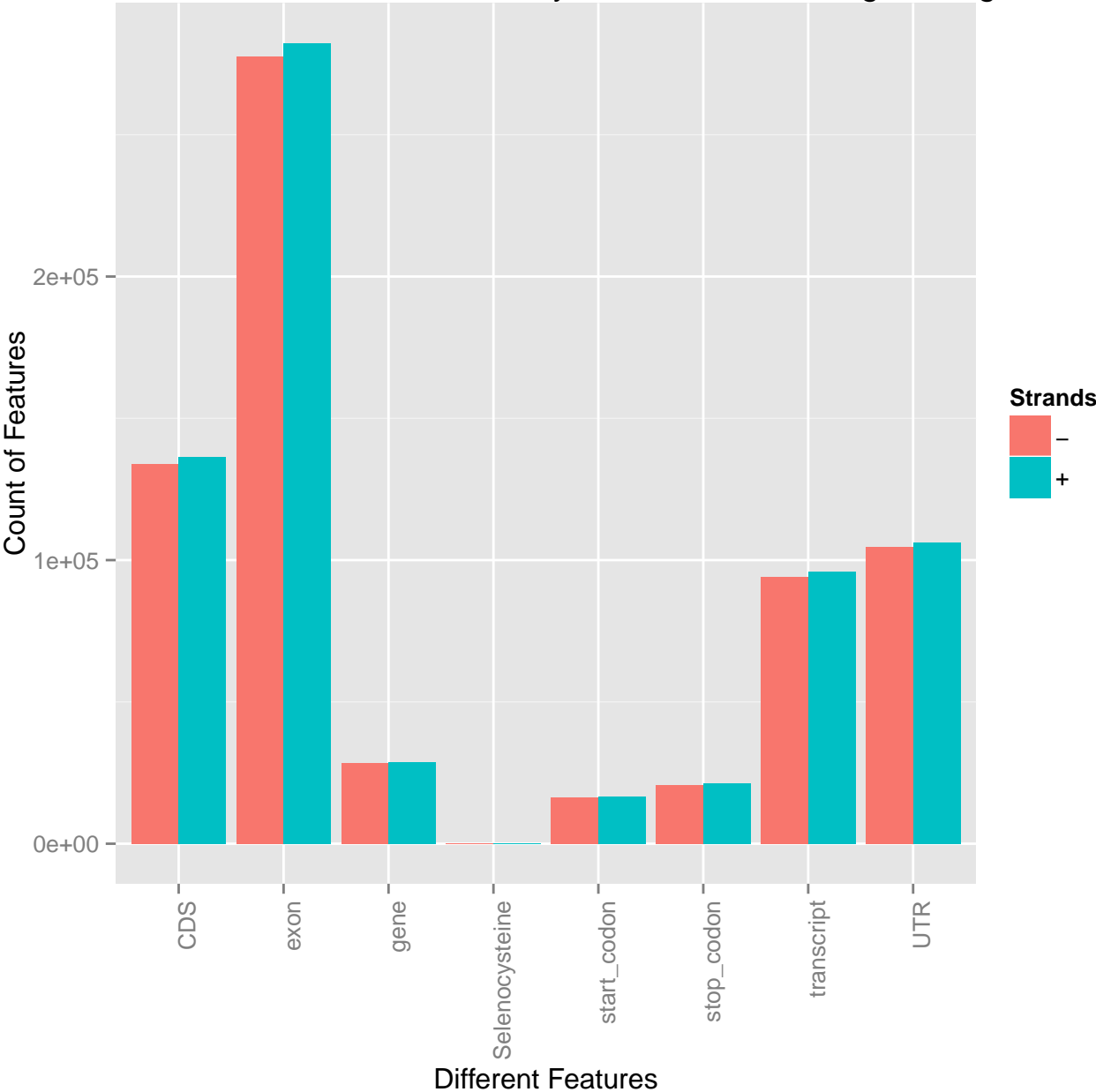
Chromosomes



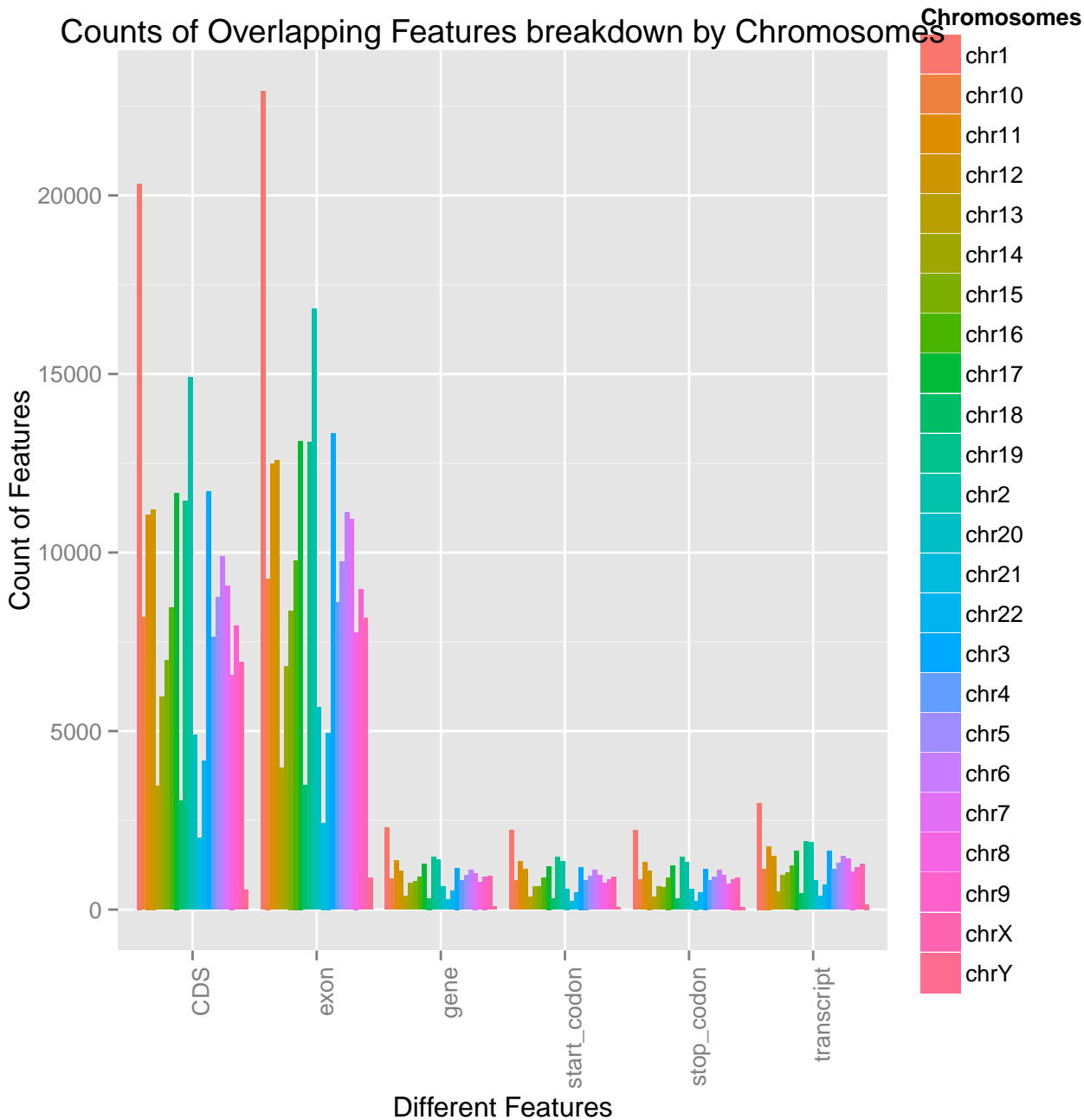
Counts of Features breakdown by Strands in the file– refseq.gtf



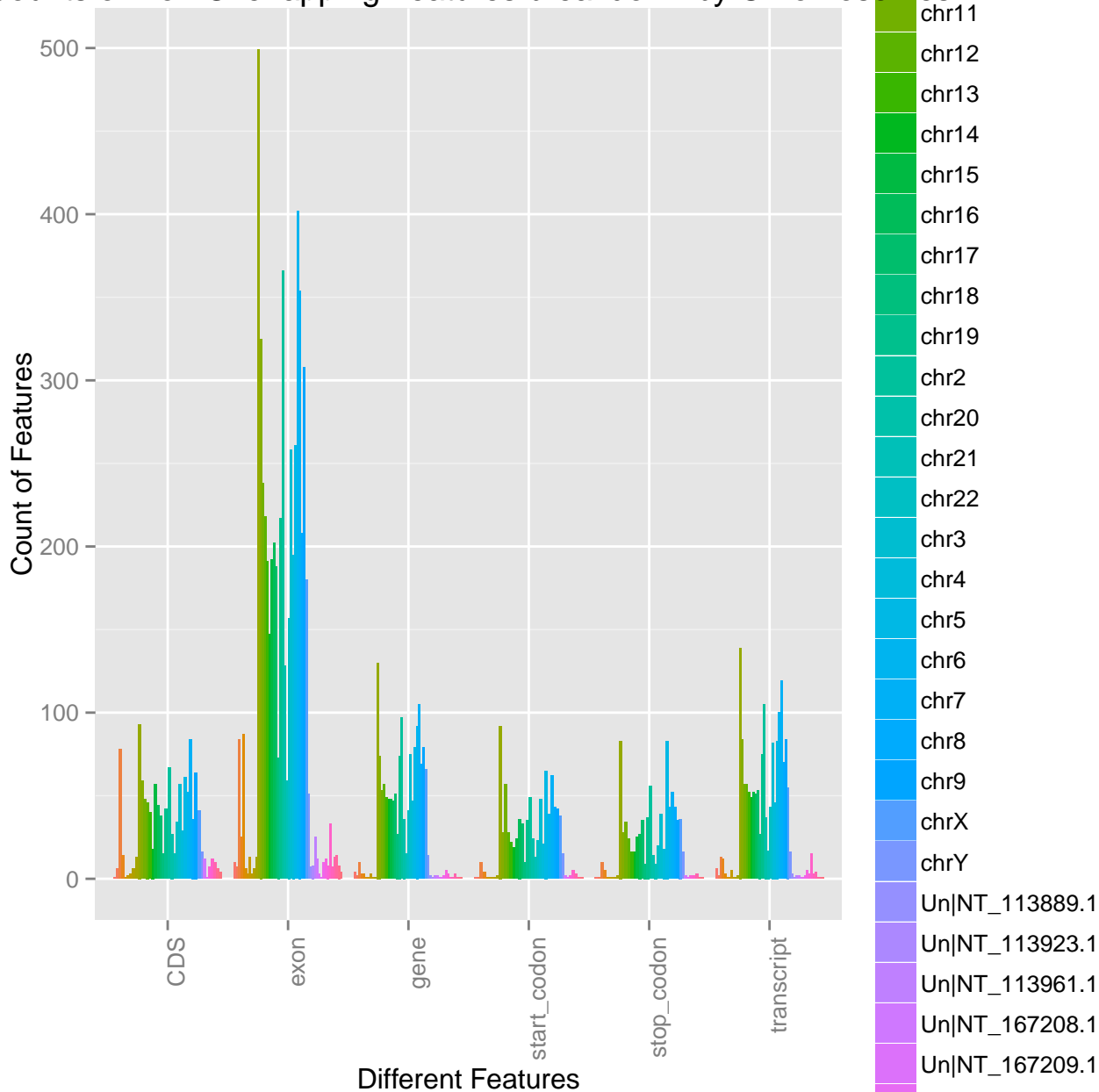
Counts of Features breakdown by Strands in the file– gencode.gtf



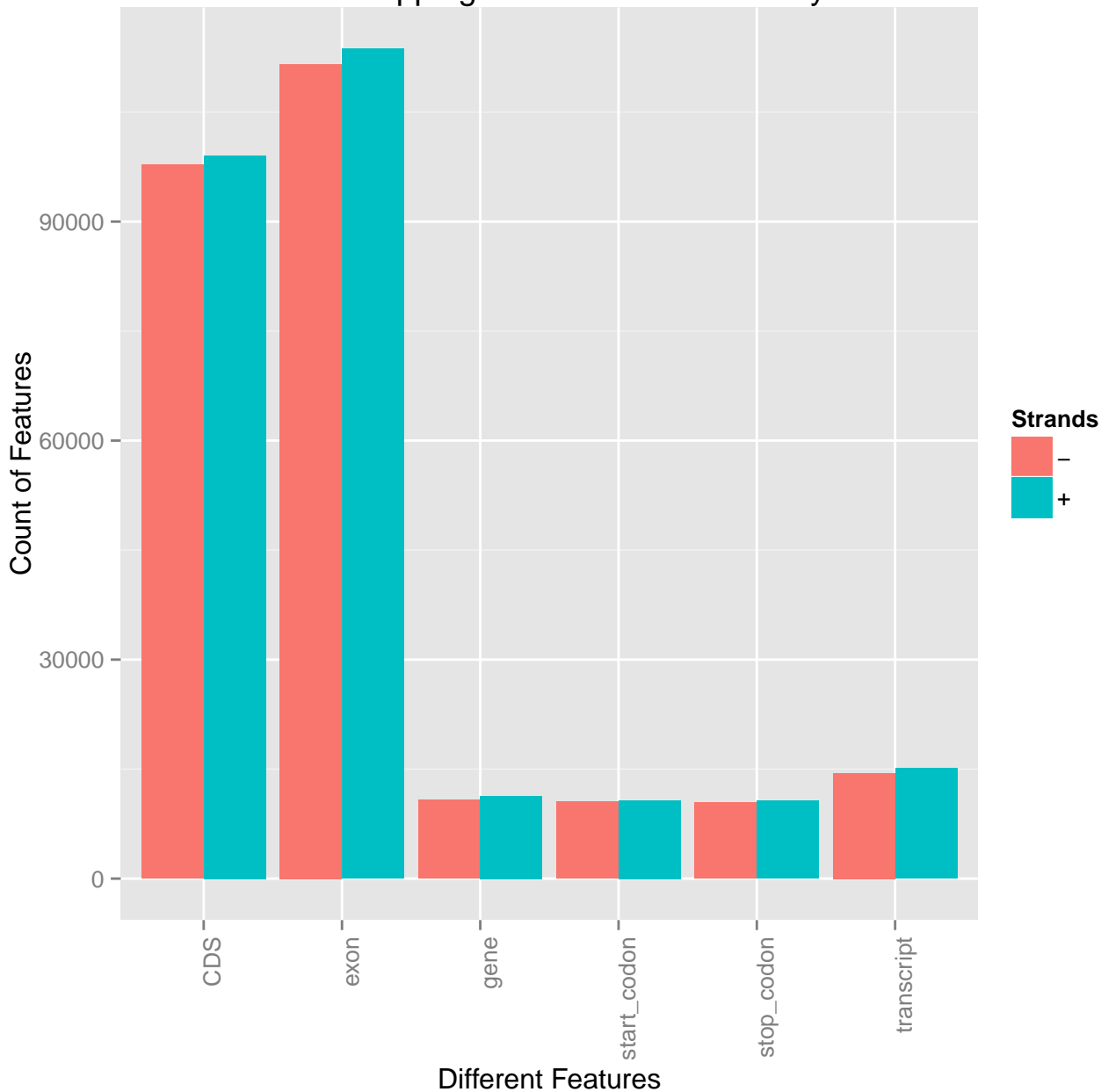
Counts of Overlapping Features breakdown by Chromosomes



Counts of Non Overlapping Features breakdown by Chromosomes



Counts of Overlapping Features breakdown by Strands



Counts of Non Overlapping Features breakdown by Strands

