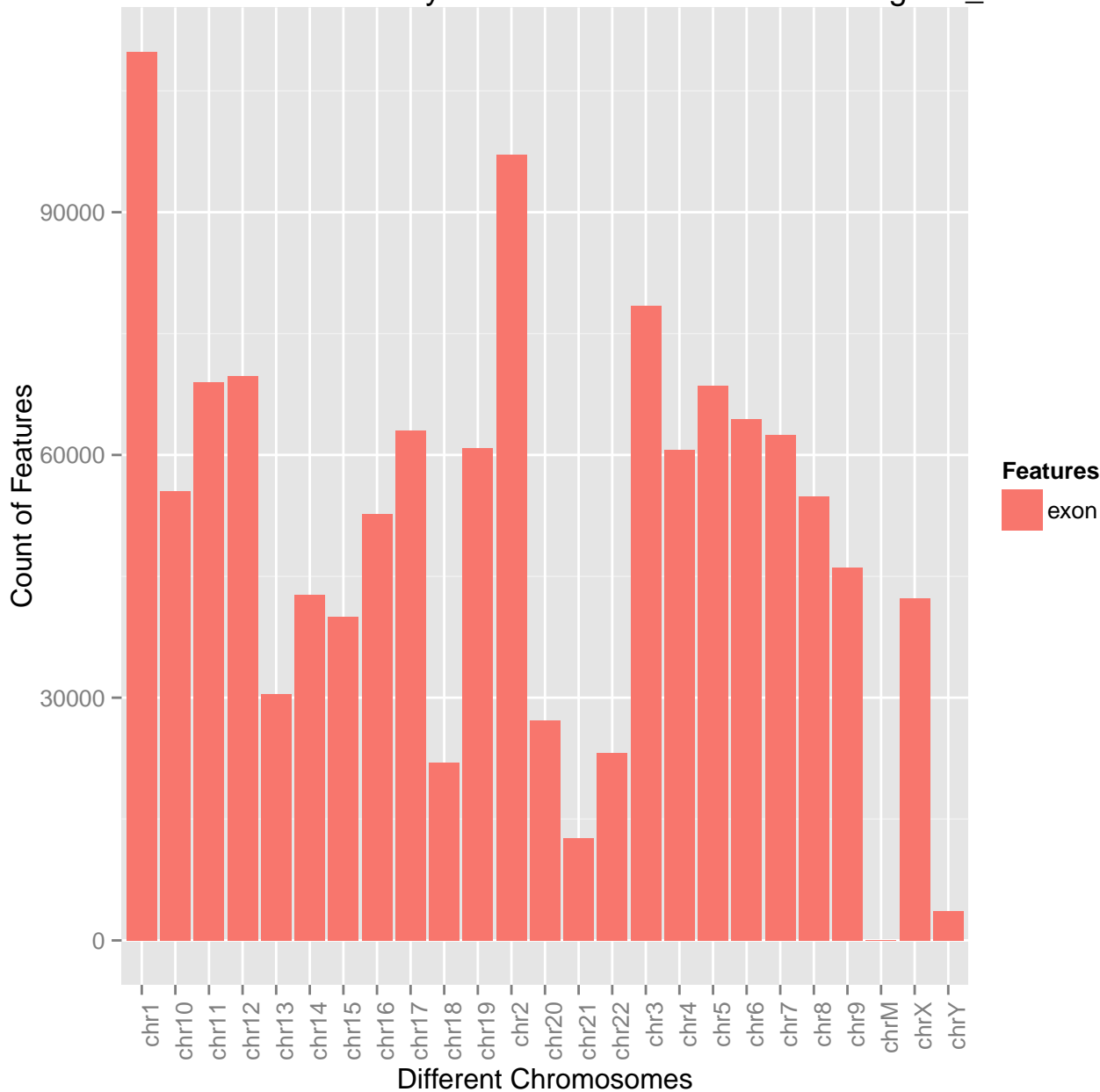
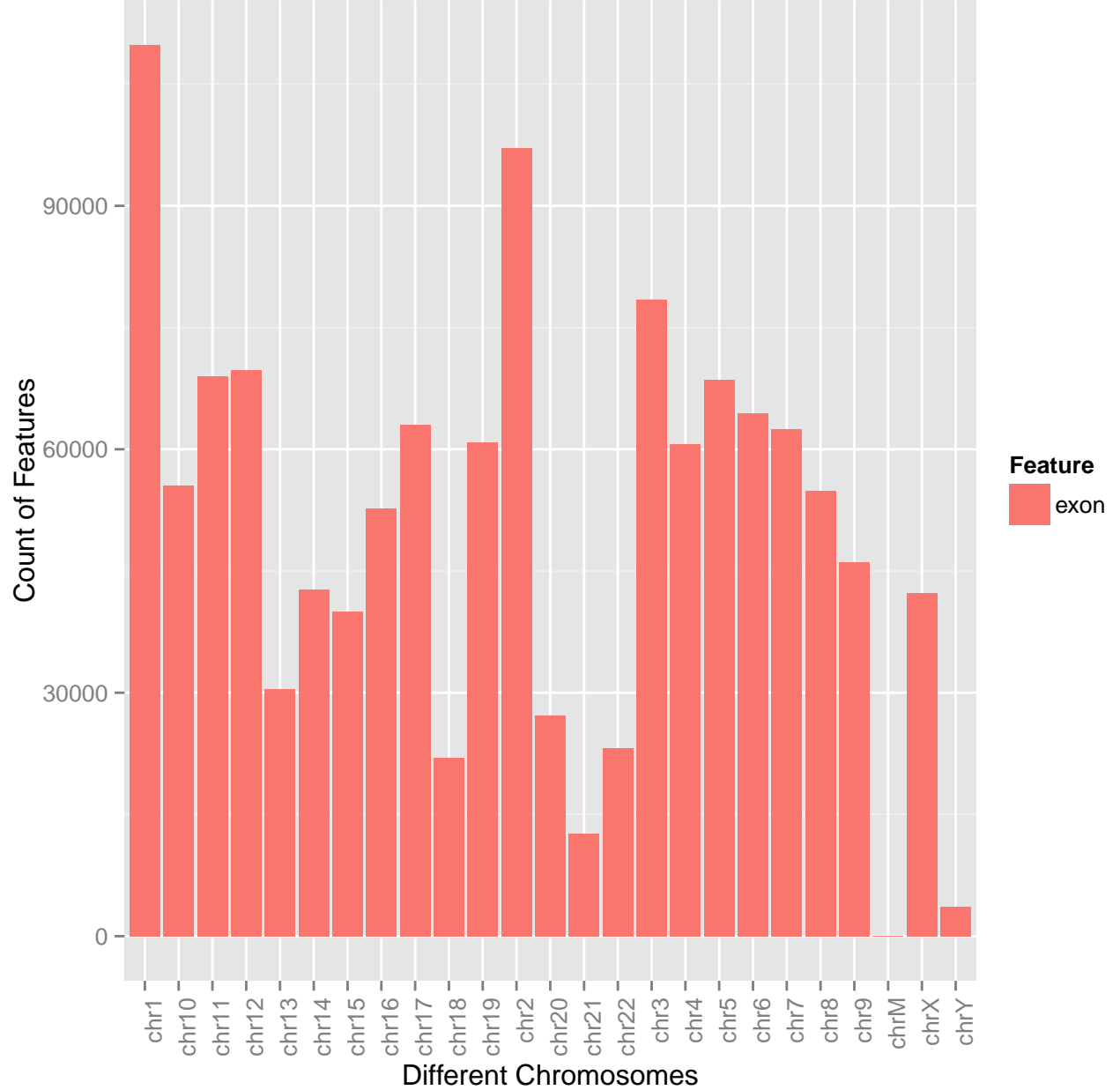


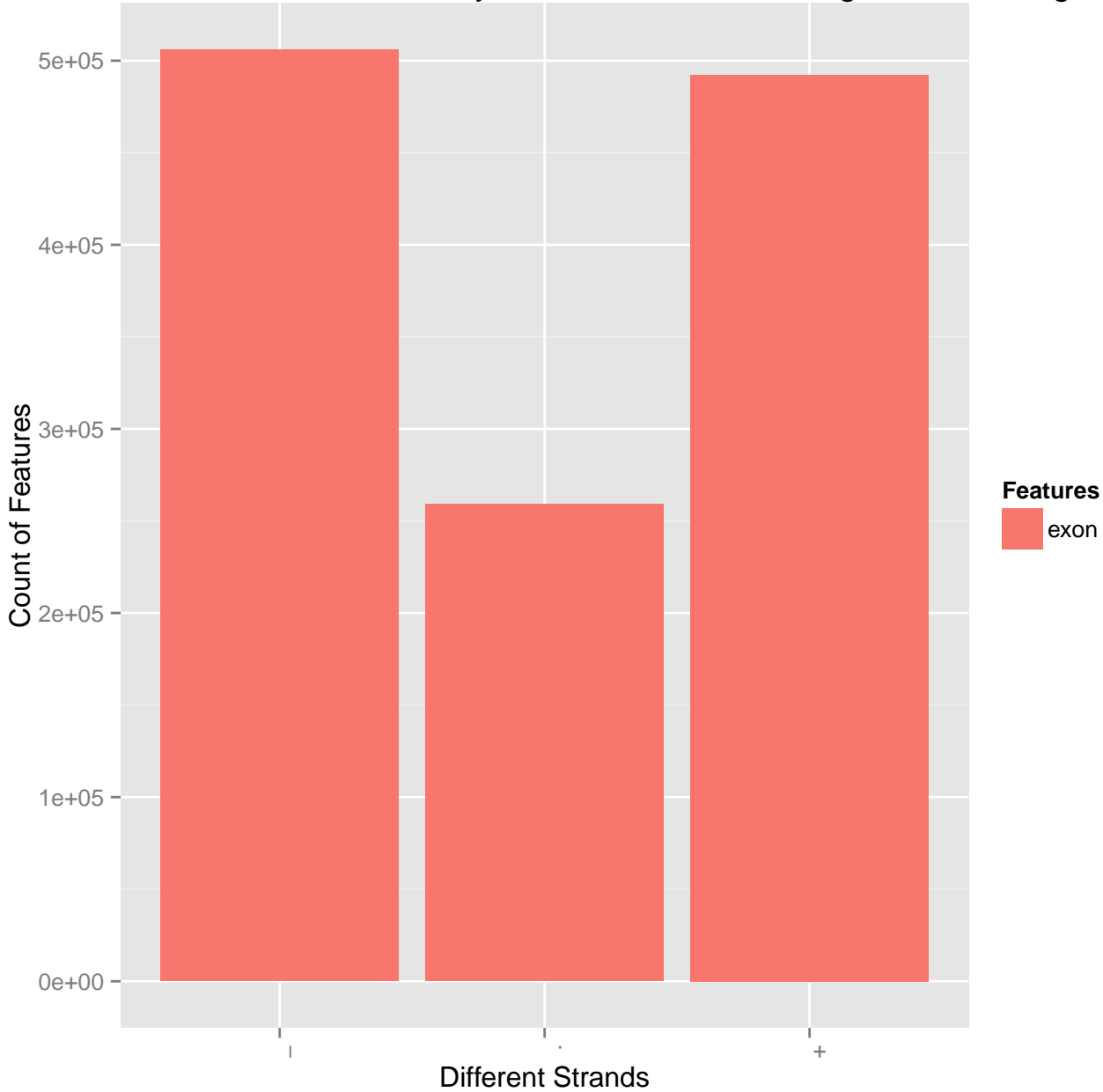
Counts of Features breakdown by Chromosomes in the file- mergedX_nmm10.gtf



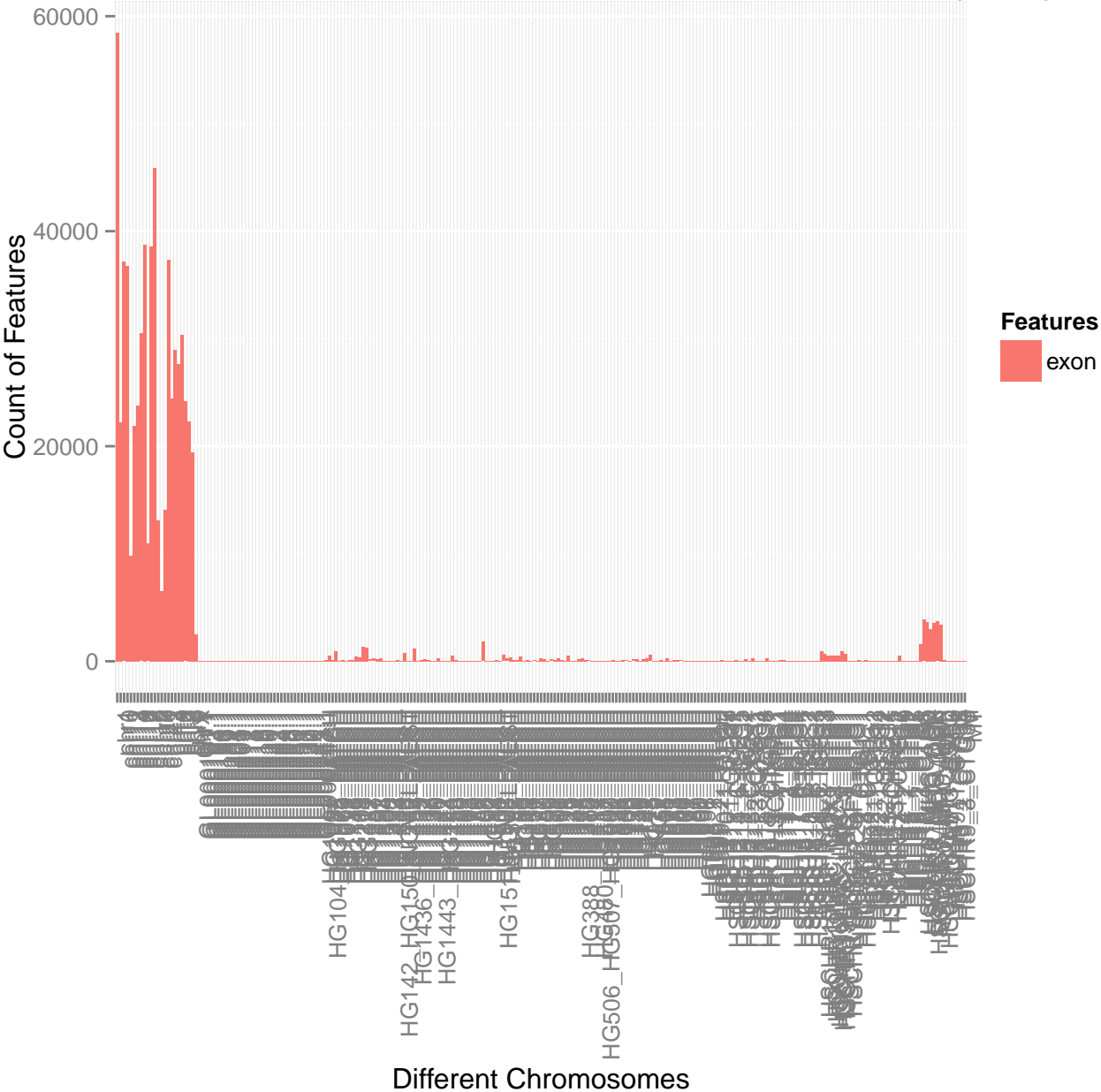
Counts of Features by Chromosomes in the file– mergedX_nmm10.gtf



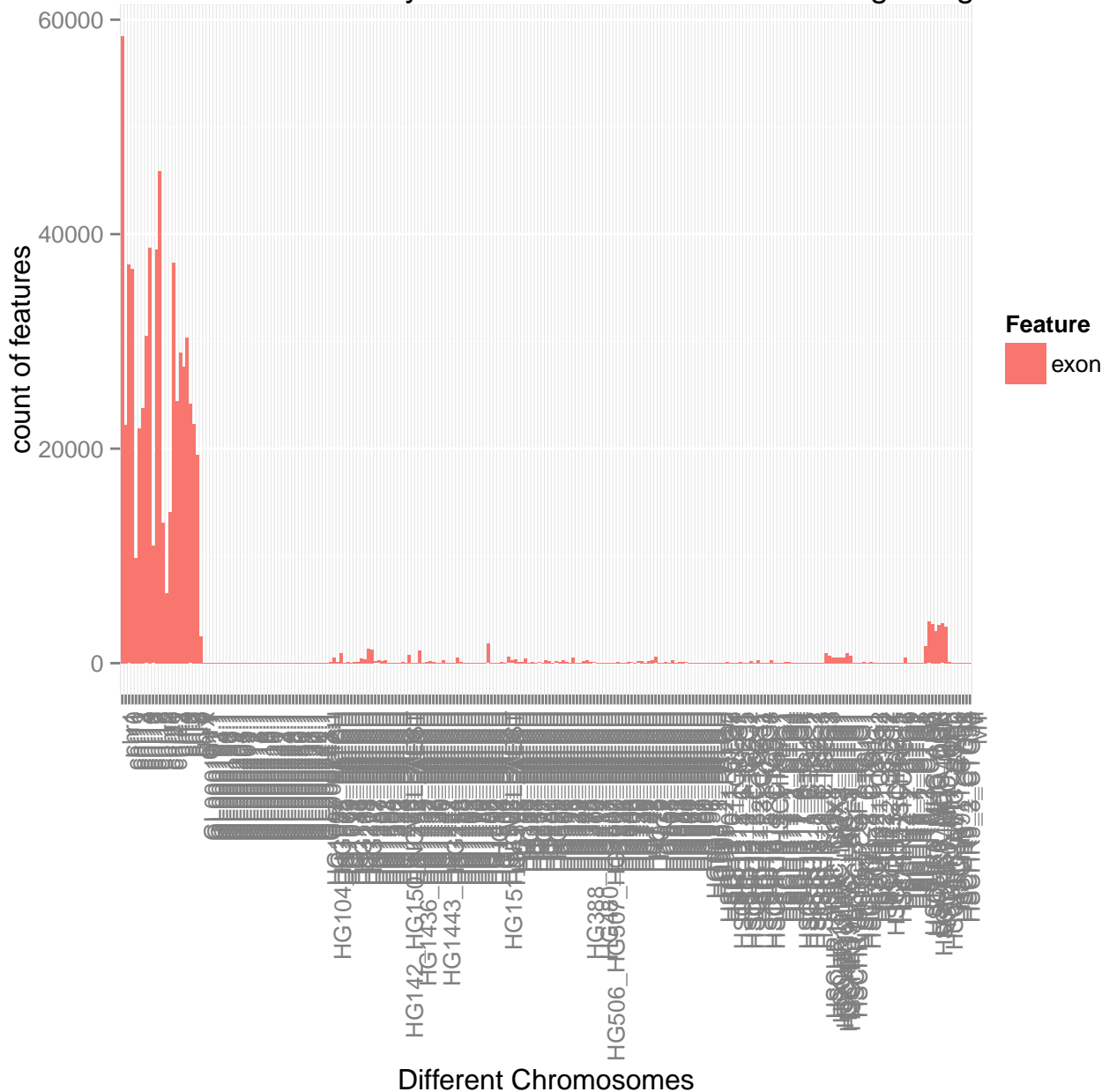
Counts of Features breakdown by Strands in the file– mergedX_nmm10.gtf



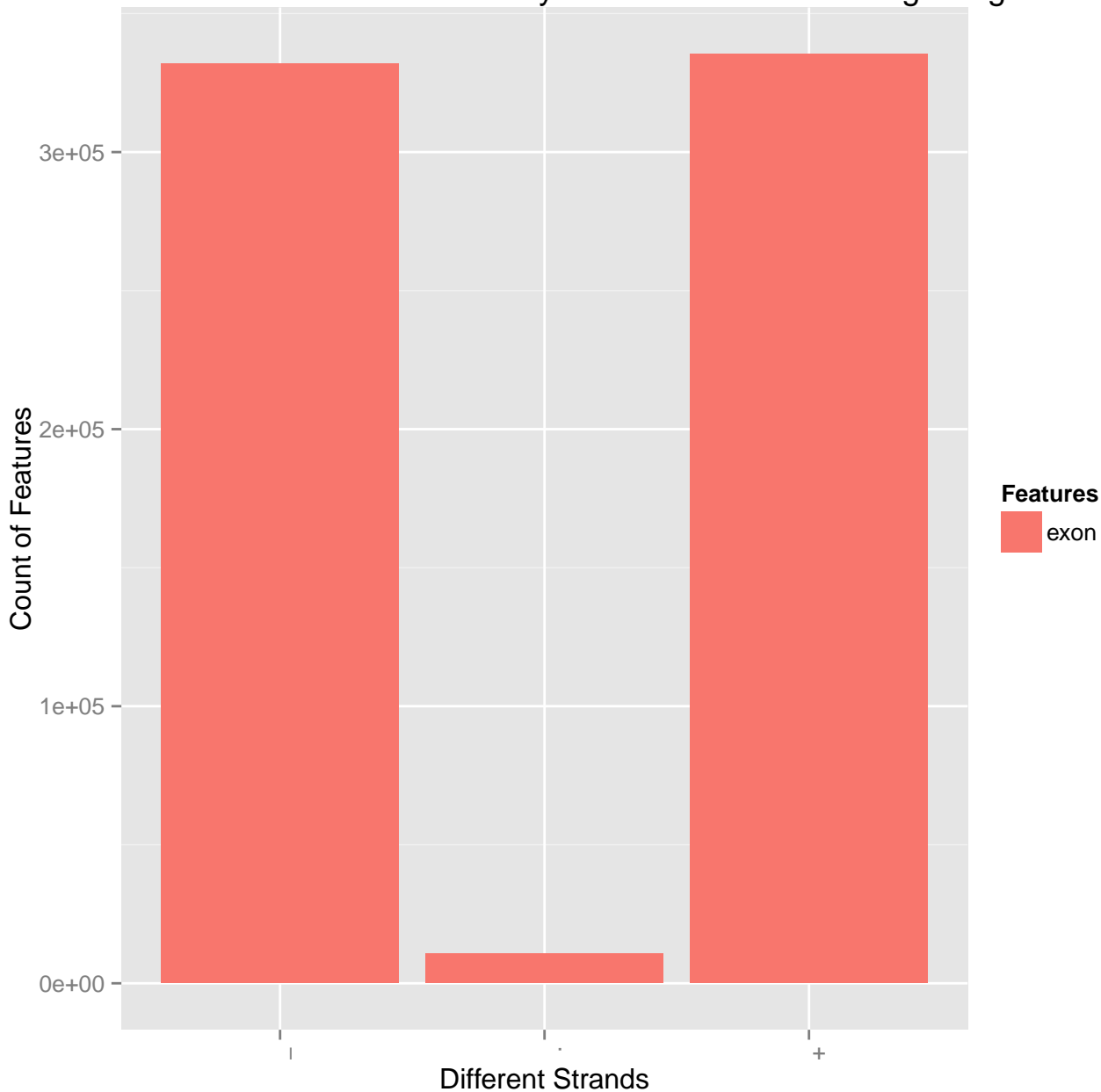
Counts of Features breakdown by Chromosomes in the file– mergedS.gtf



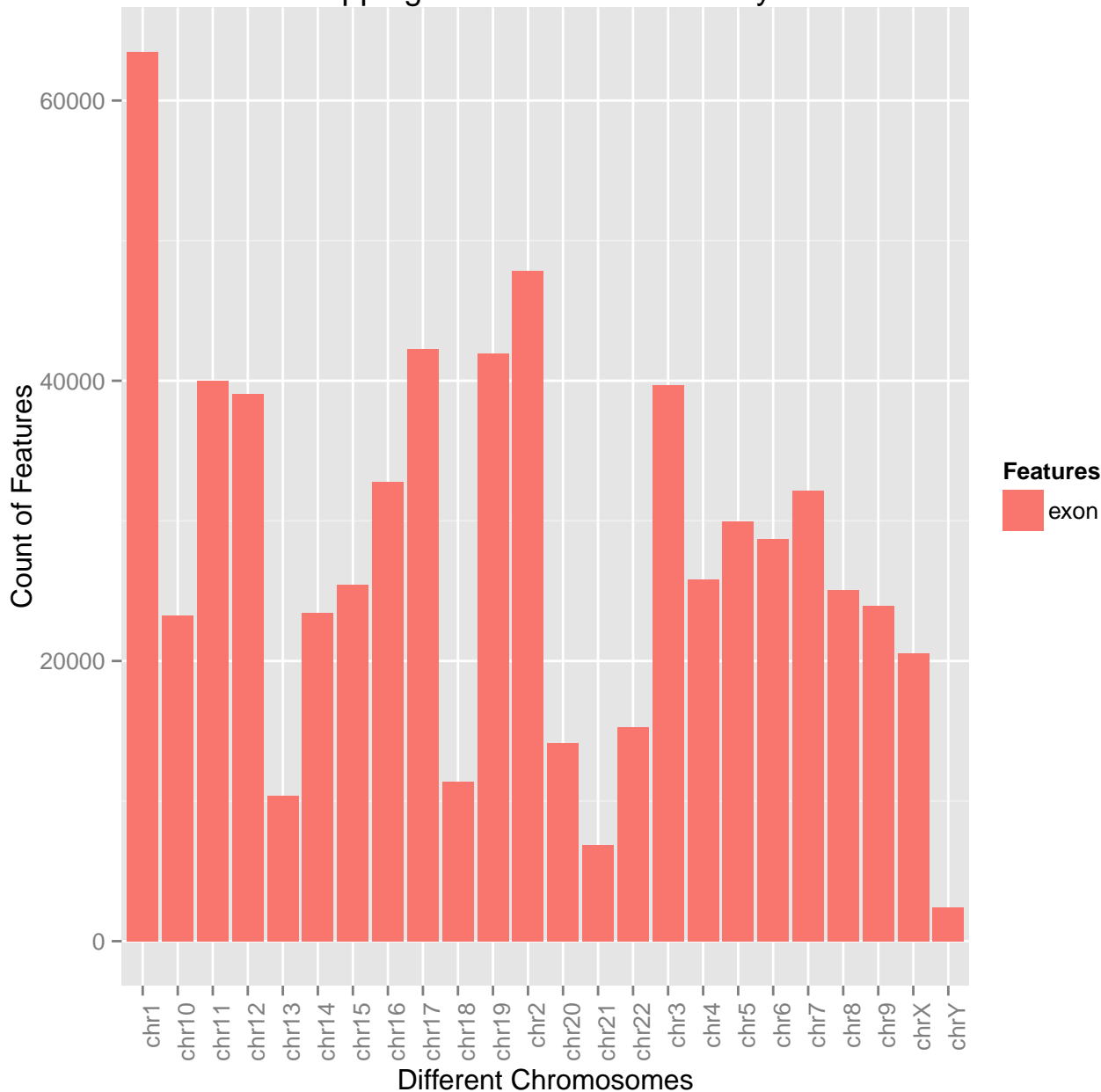
Counts of Features by Chromosomes in the file– mergedS.gtf



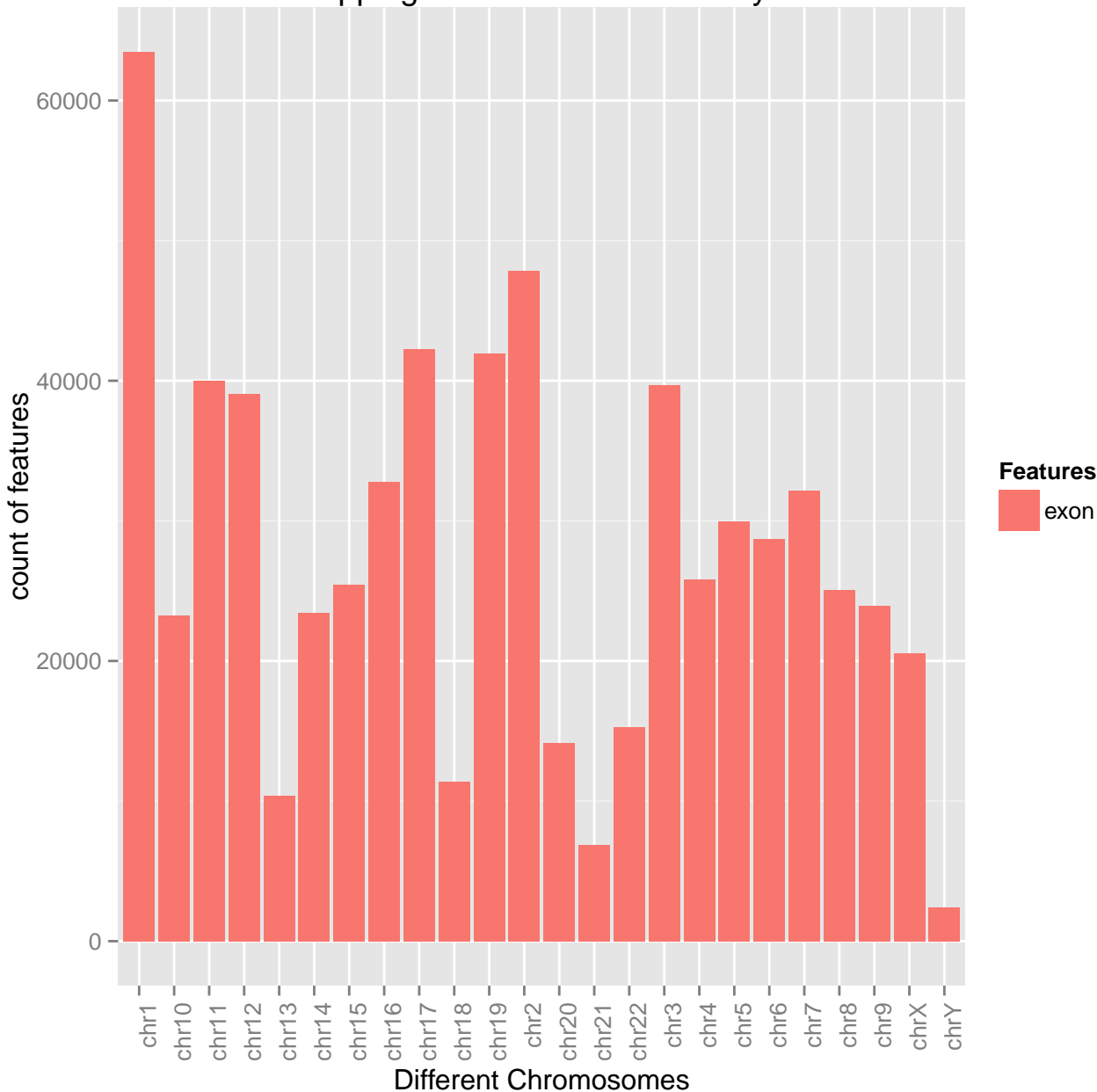
Counts of Features breakdown by Strands in the file– mergedS.gtf



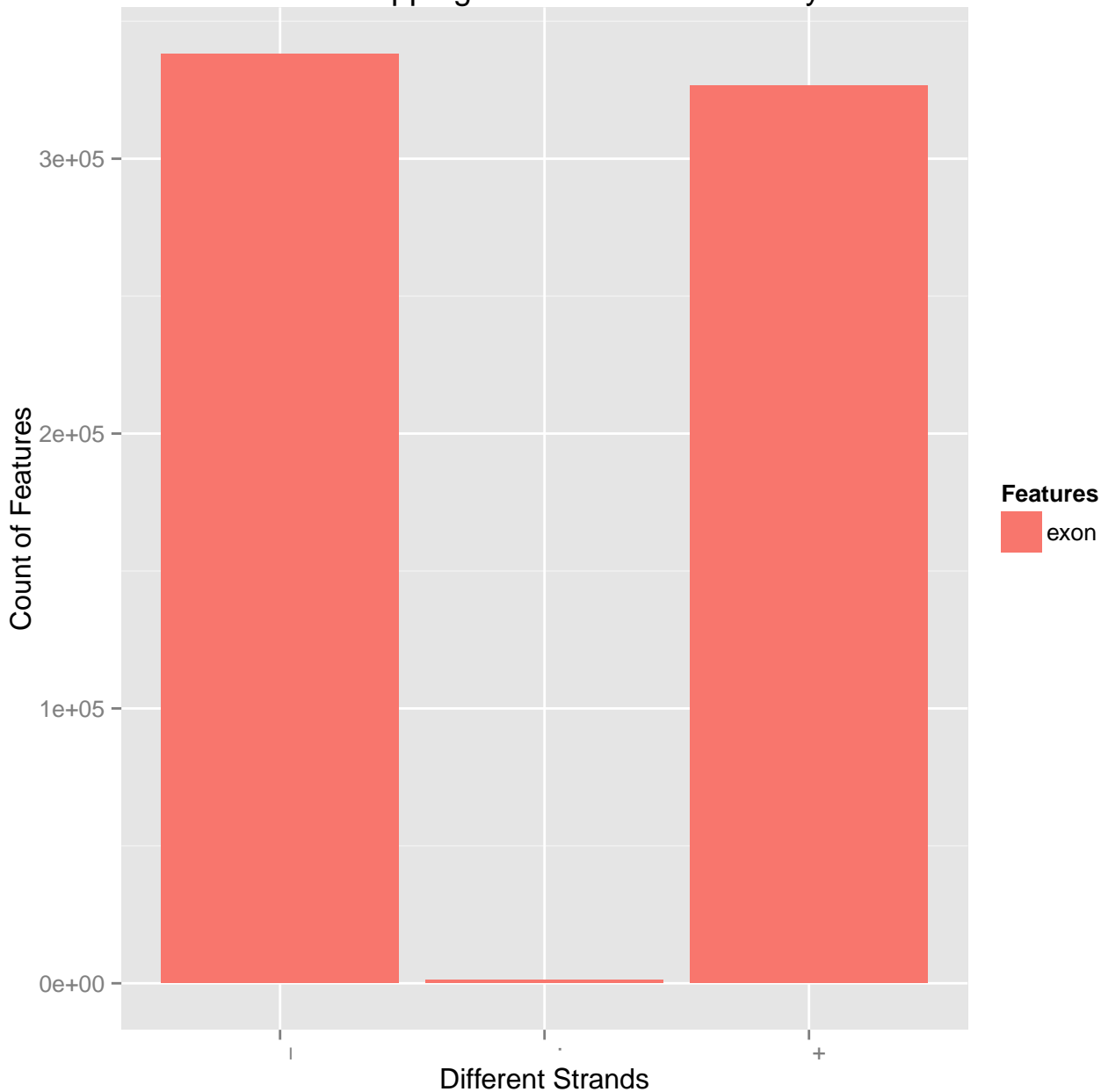
Counts of Overlapping Features breakdown by Chromosomes



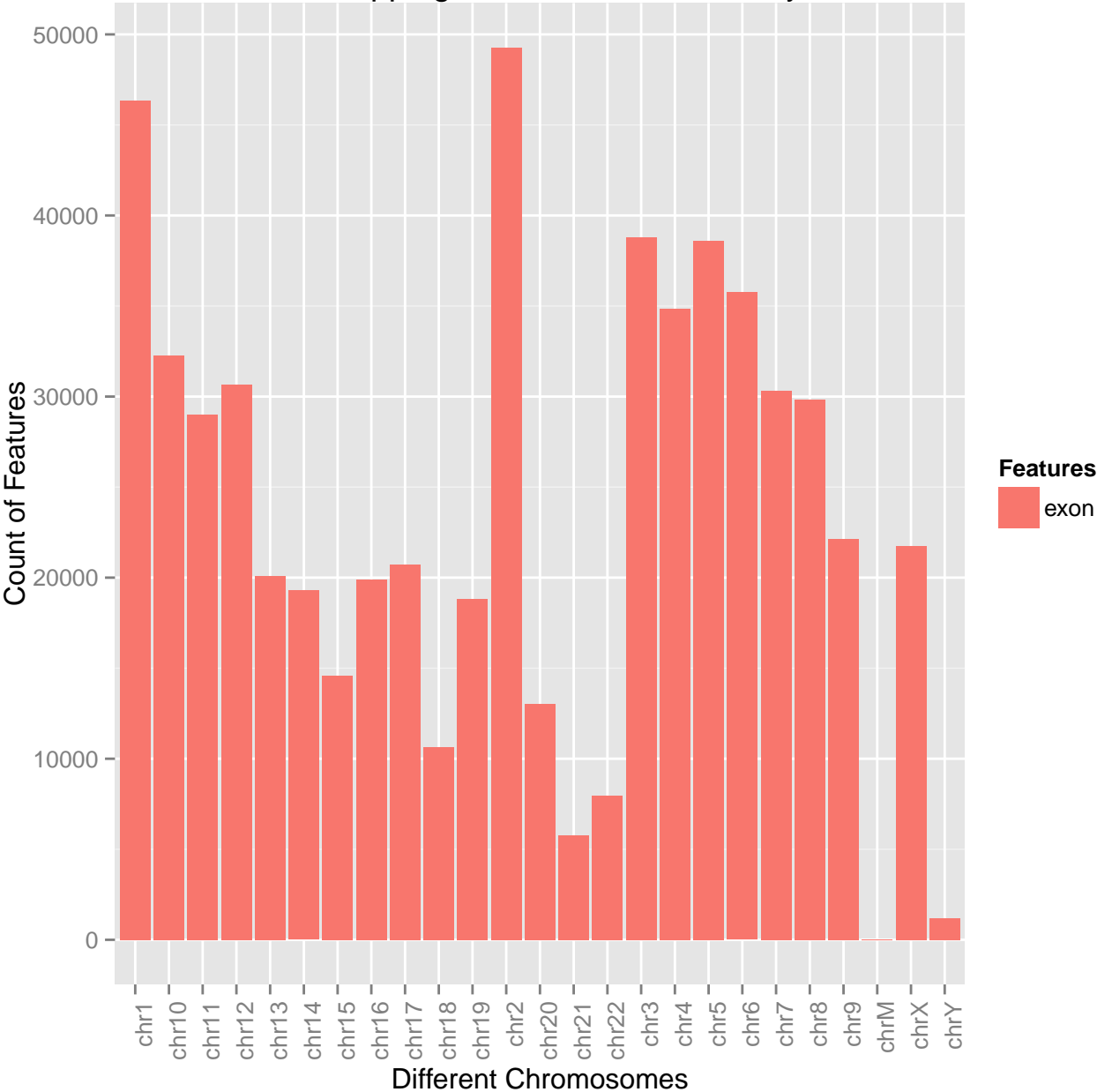
Counts of Overlapping Features breakdown by Chromosomes



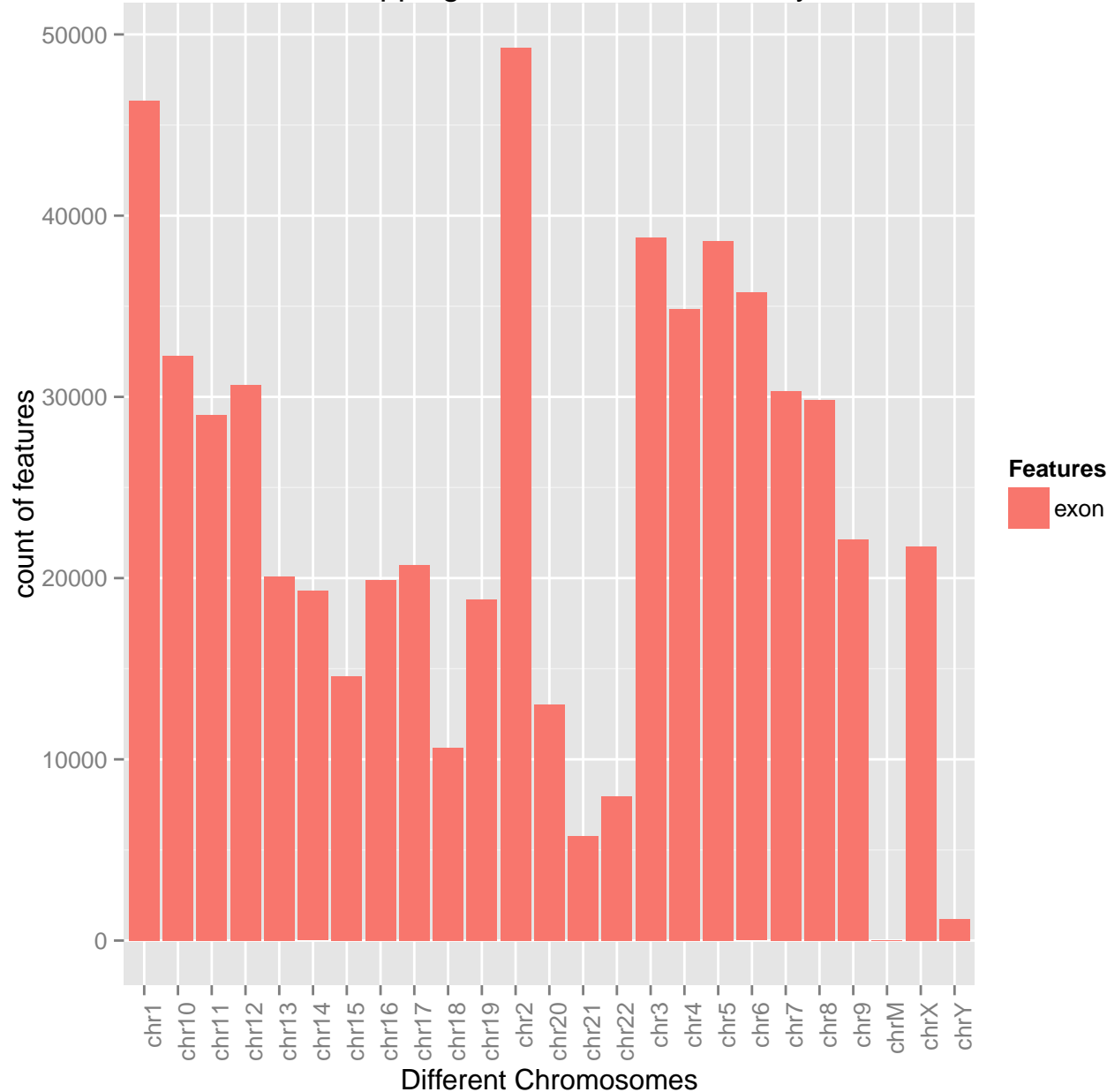
Counts of Overlapping Features breakdown by Strands



Counts of Non Overlapping Features breakdown by Chromosomes



Counts of Non Overlapping Features breakdown by Chromosomes



Counts of Non Overlapping Features breakdown by Strands

