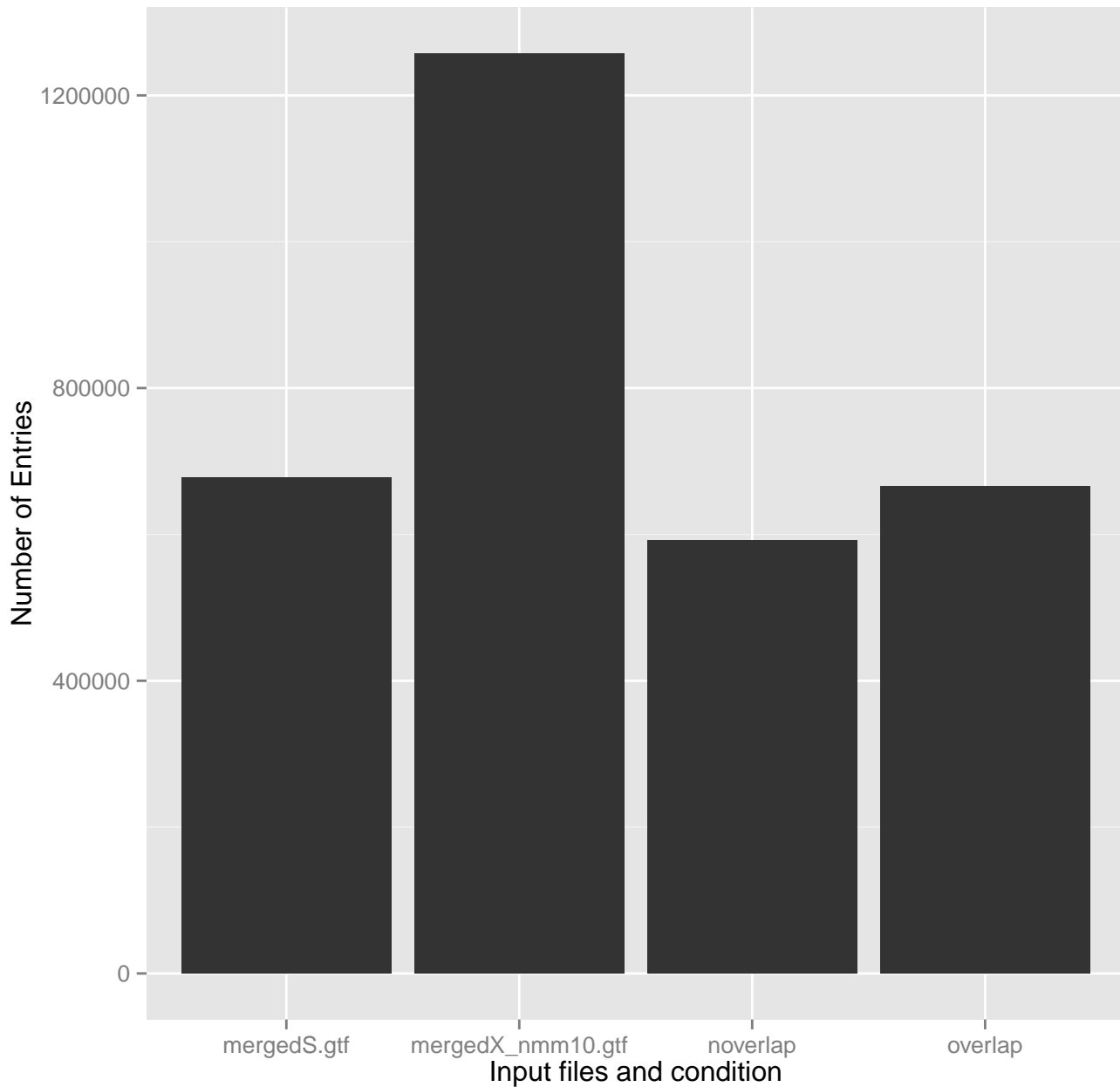
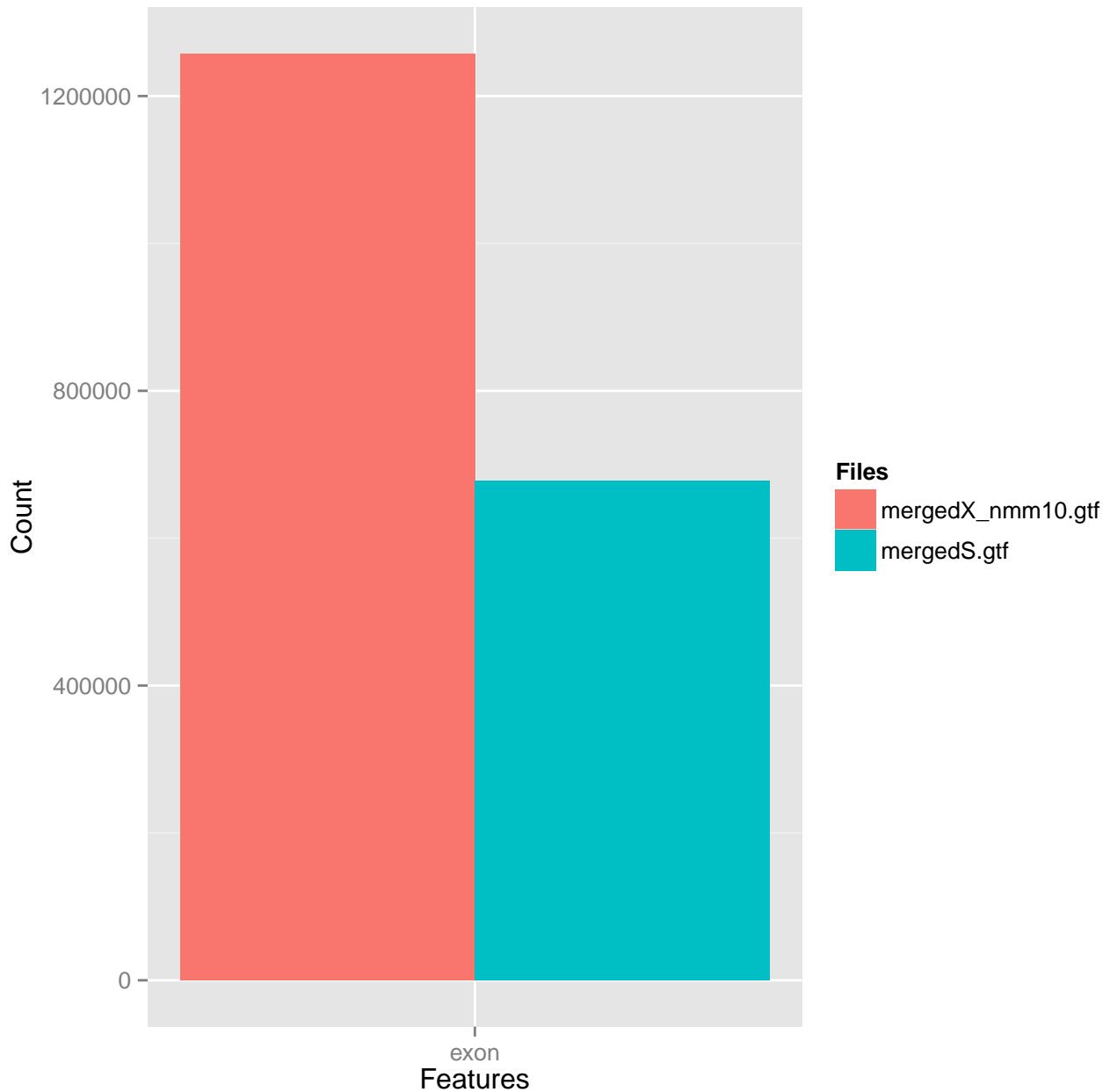


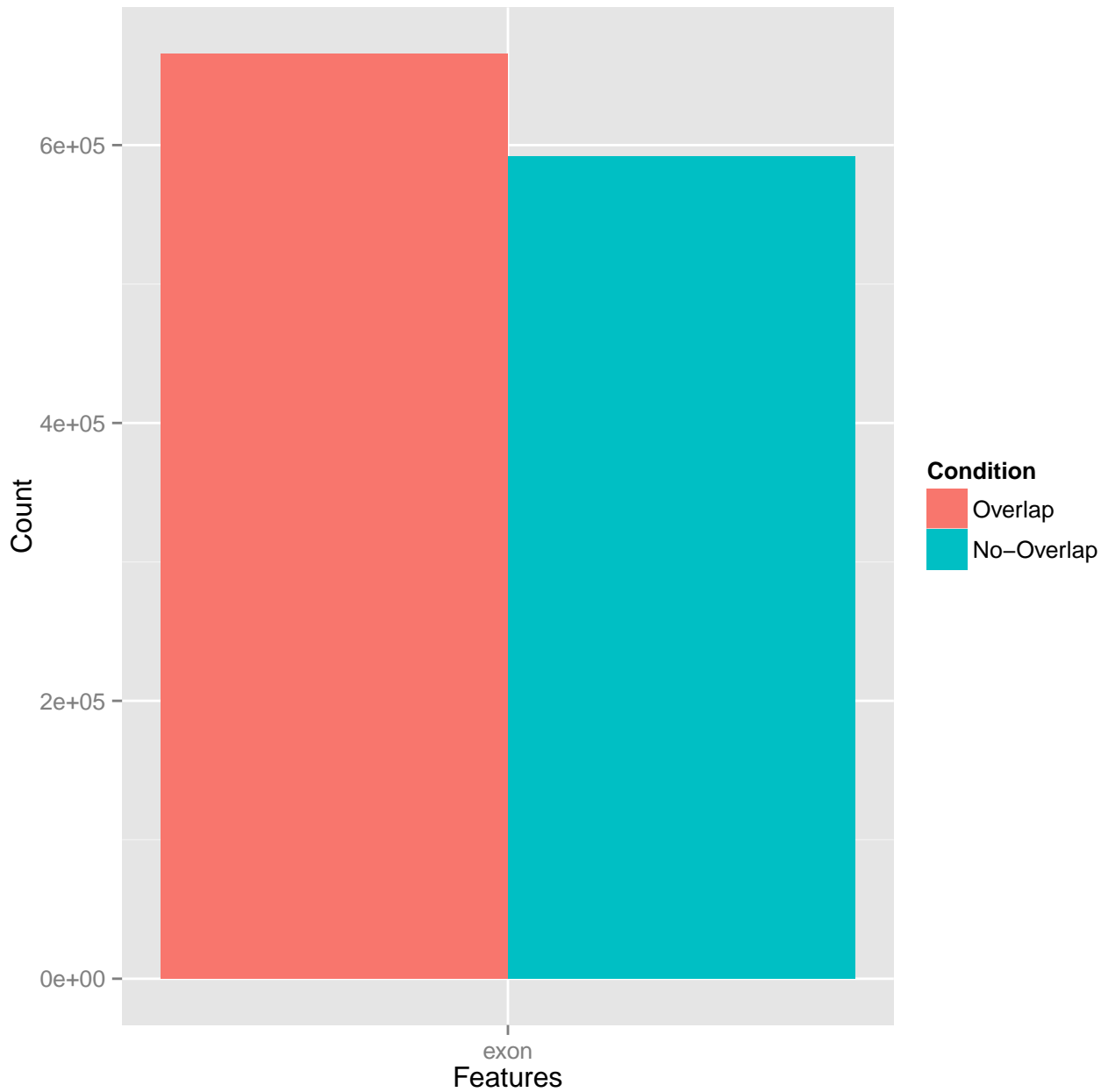
Total Entries



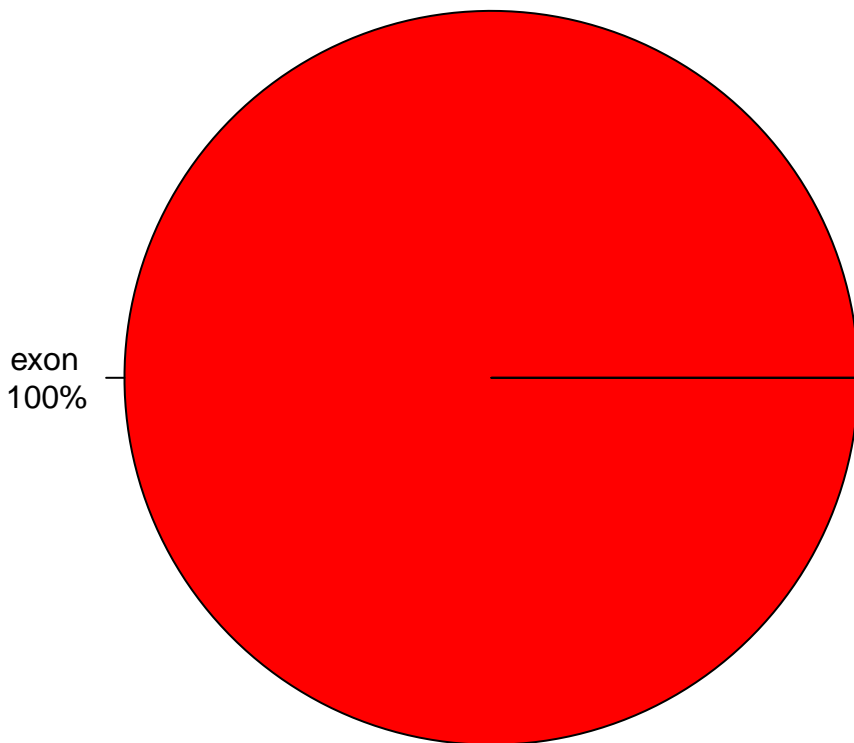
Counts of Different Features



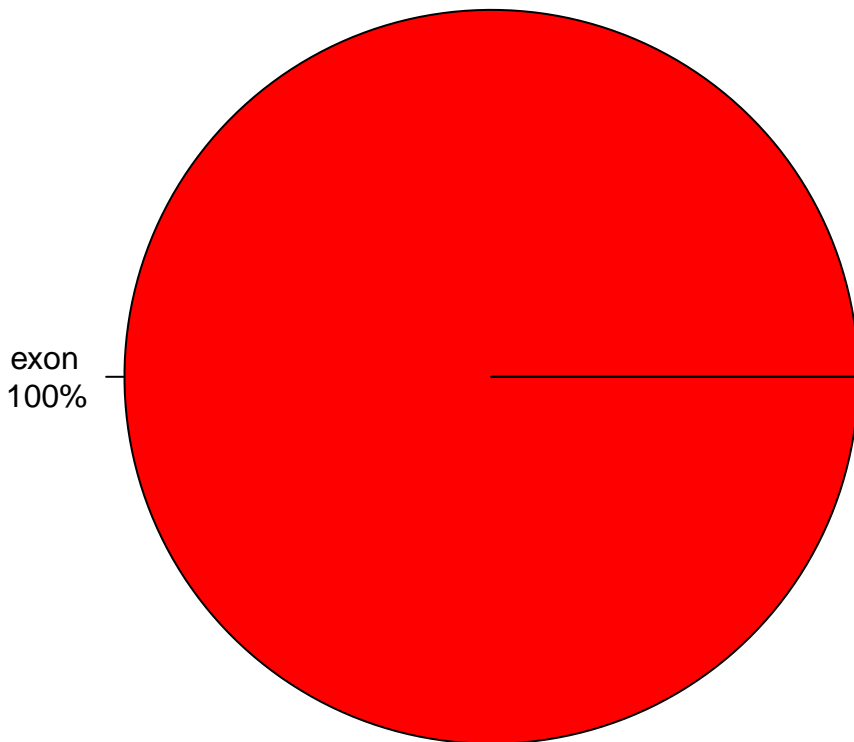
Counts of Different Features



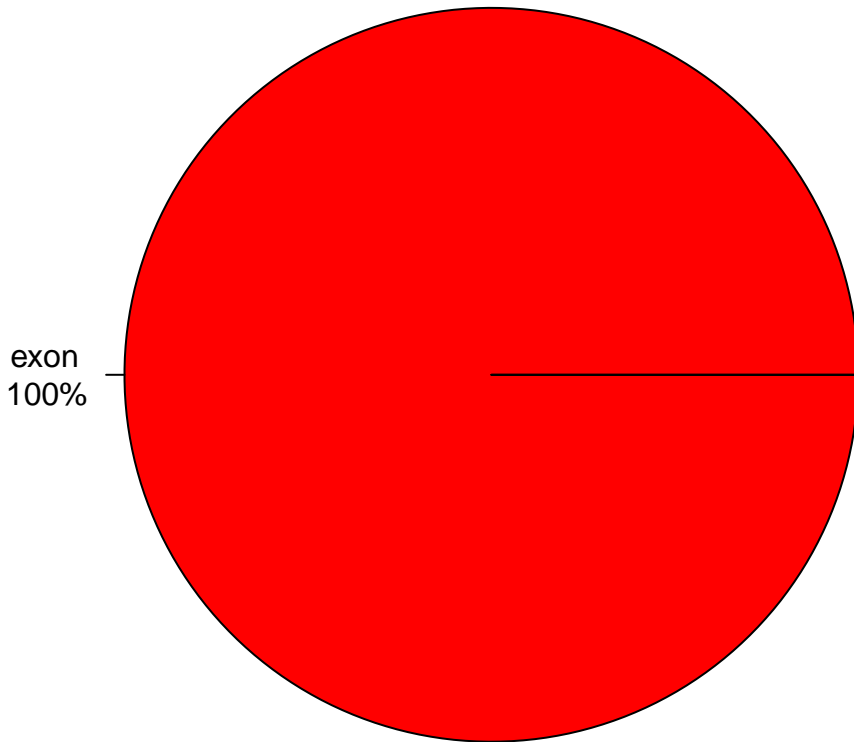
## mergedX\_nmm10.gtf – Feature Distribution



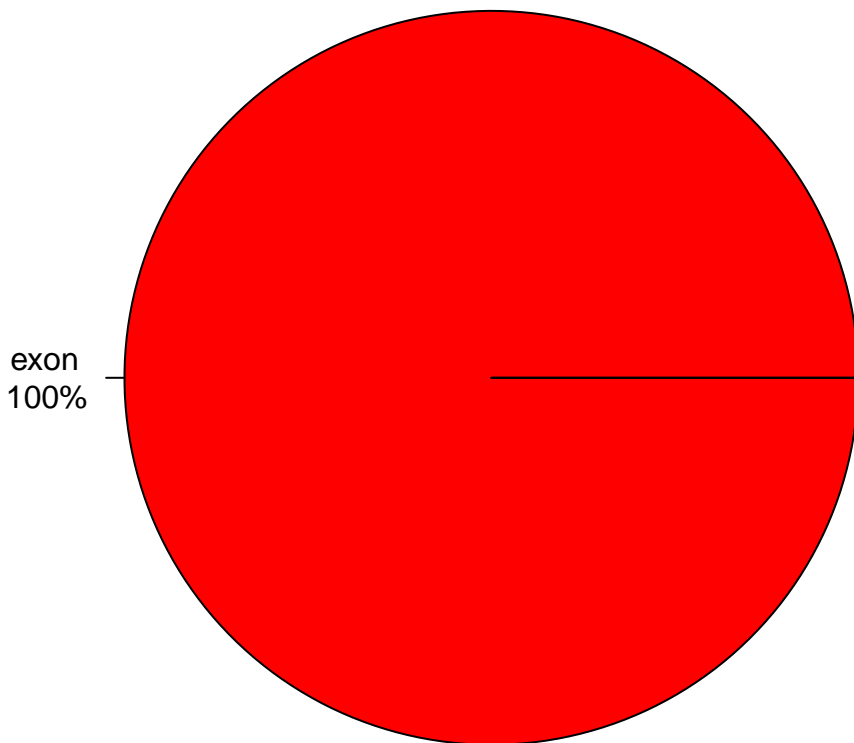
## mergedS.gtf – Feature Distribution

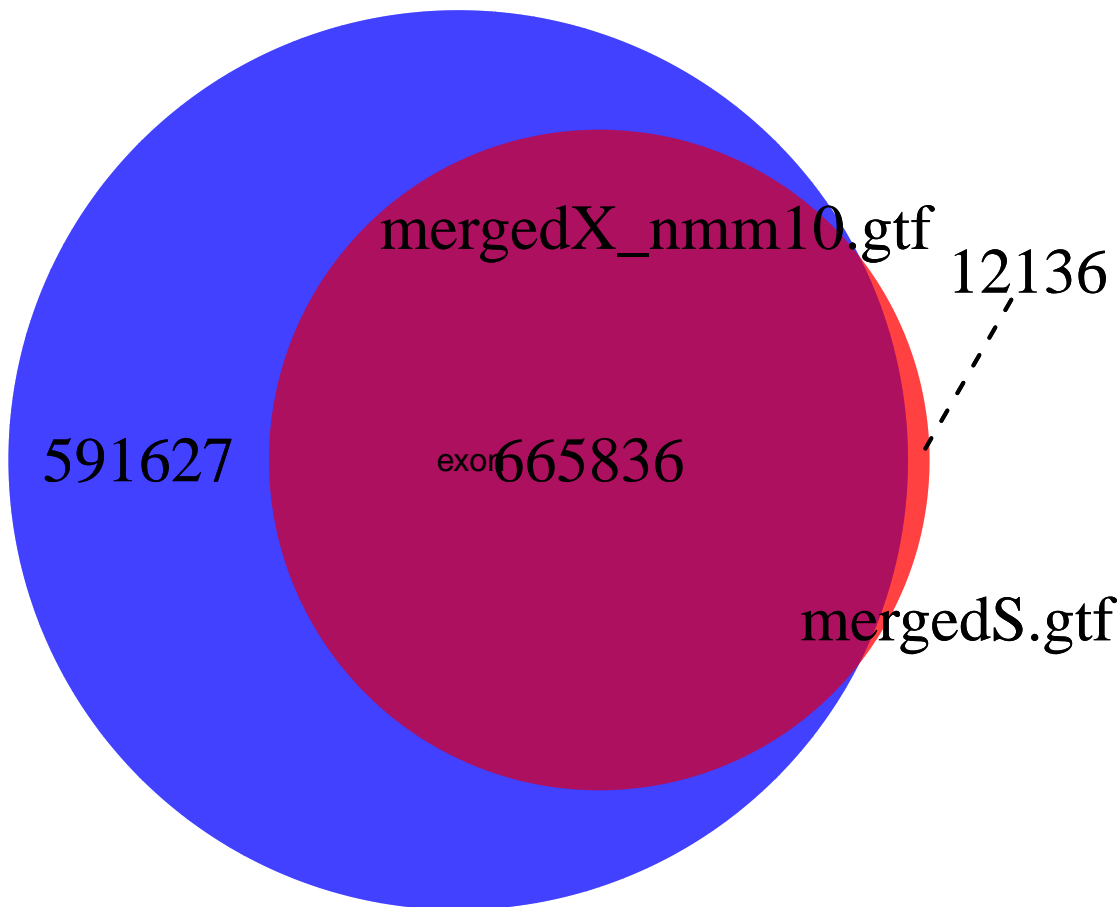


## Feature Distribution of Total Overlaps Found

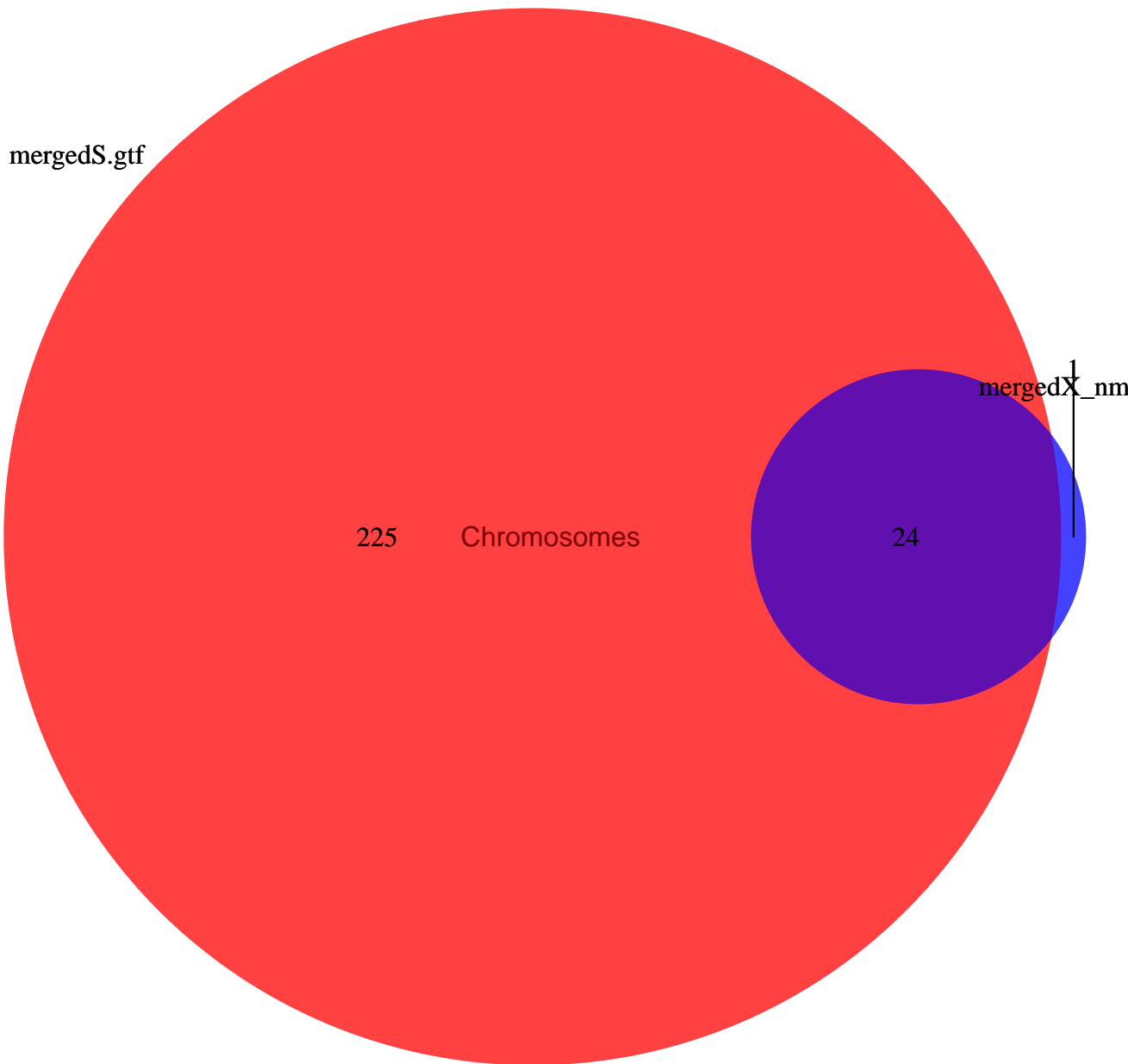


## Feature Distribution of Total Non-Overlaps Found

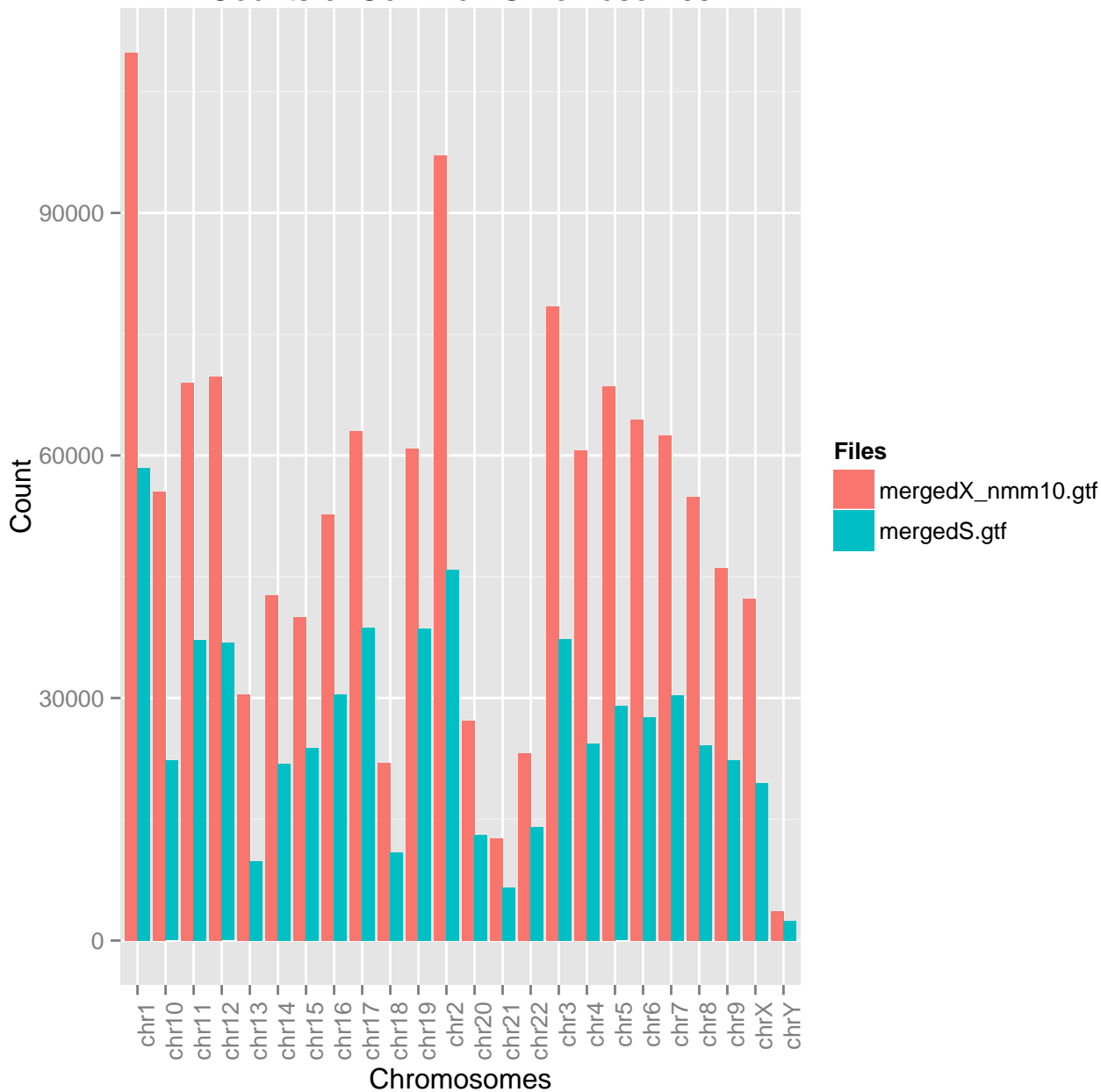




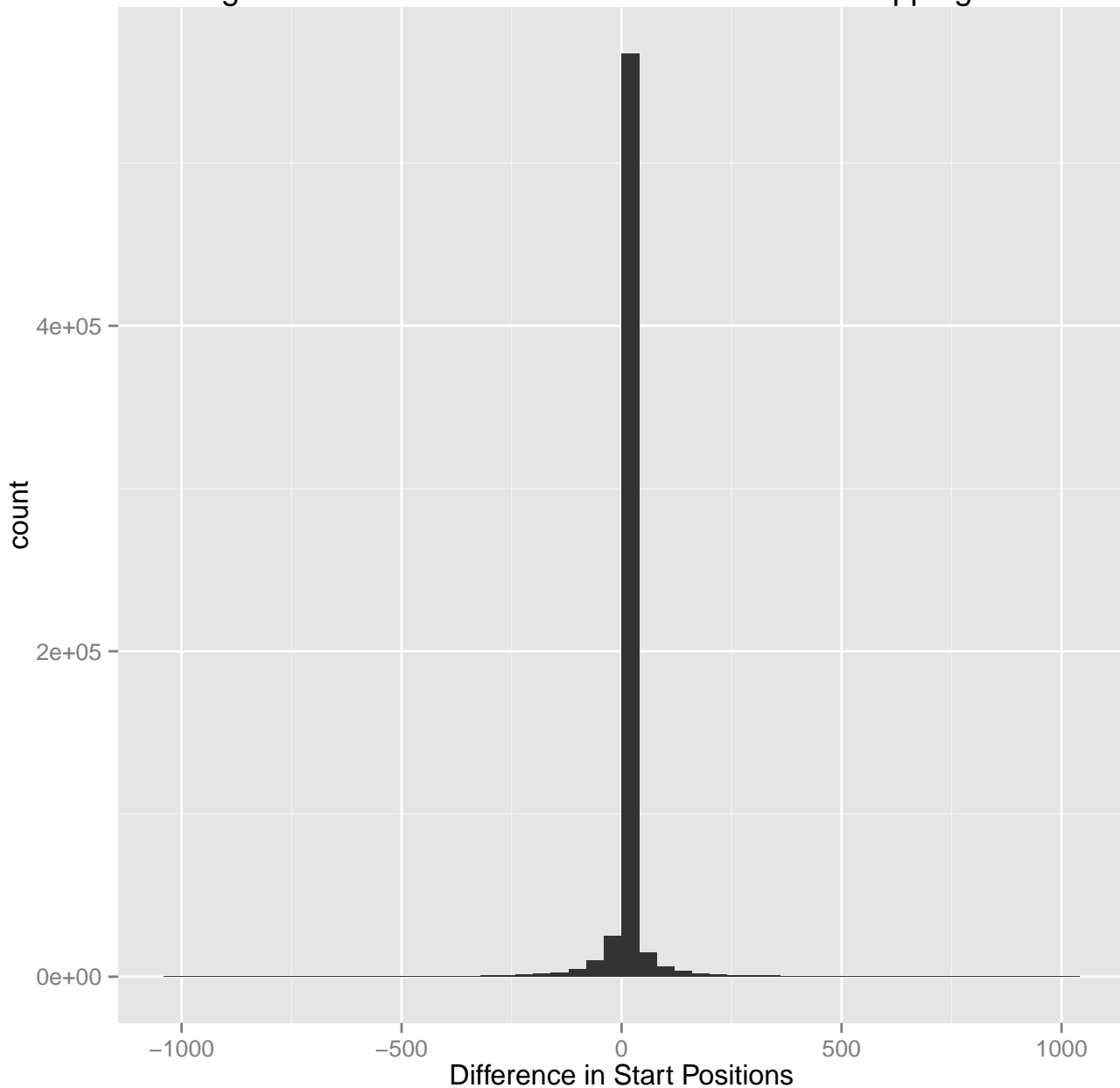




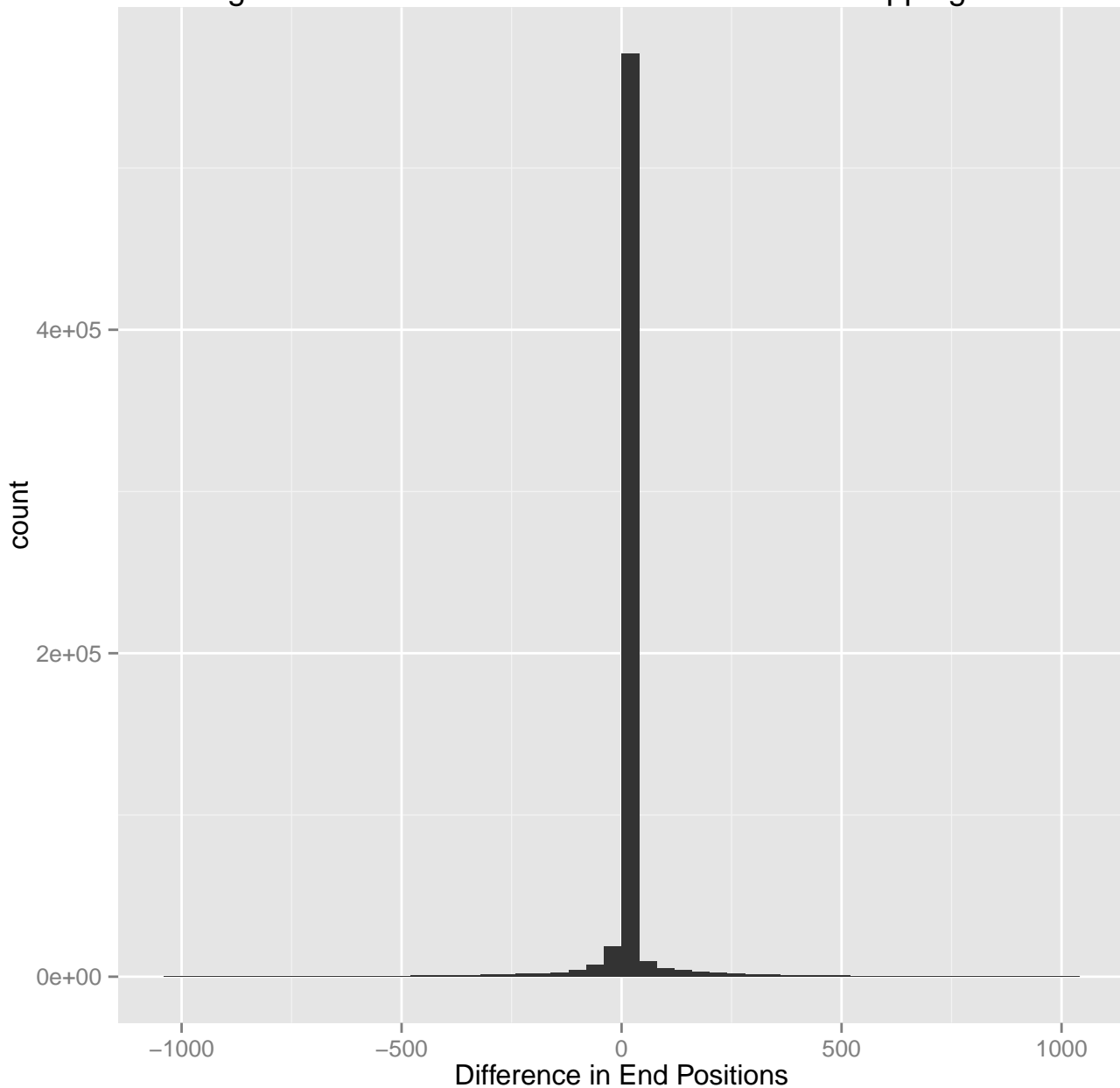
Counts of Common Chromosomes



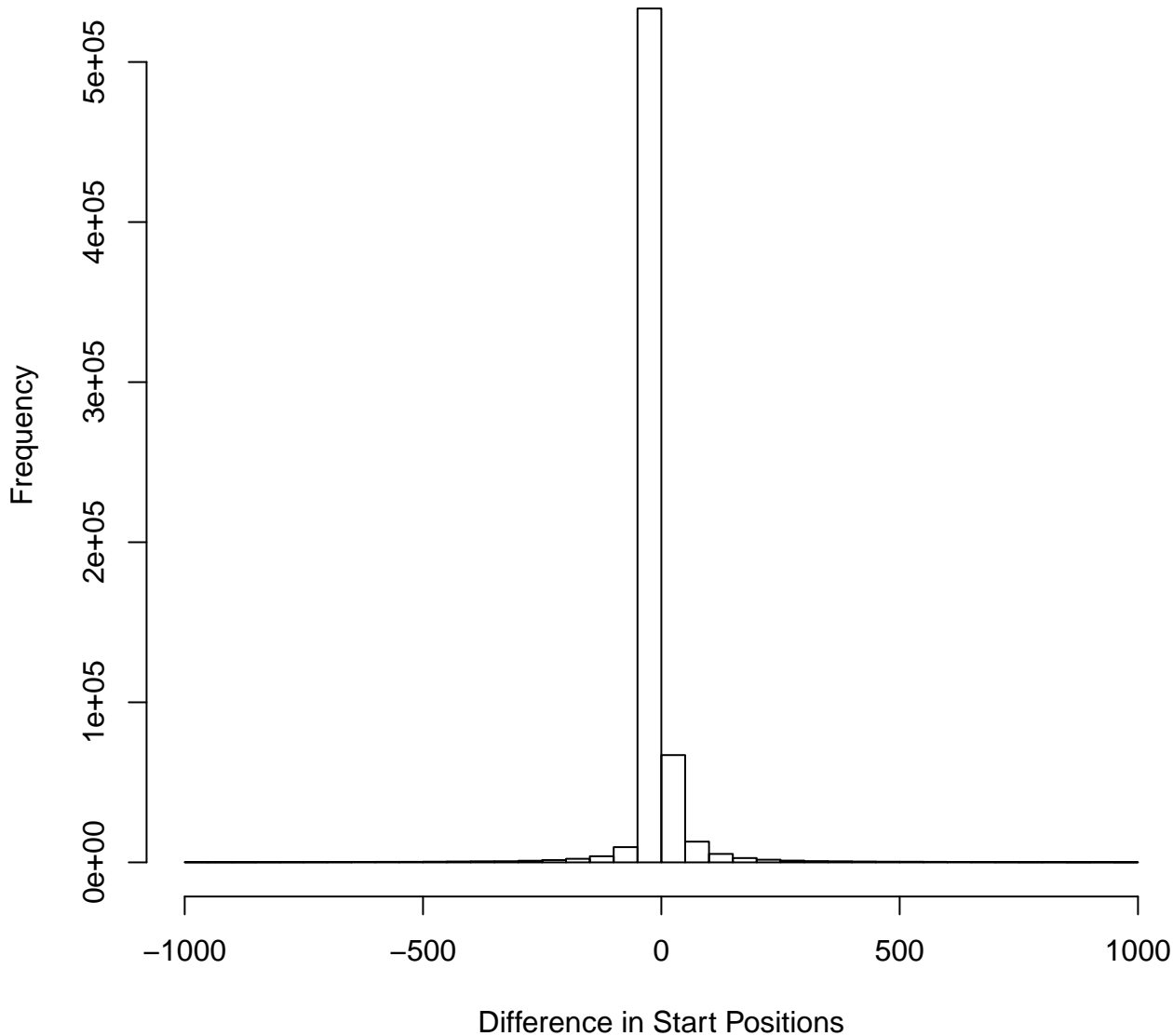
Histogram of Differences in Start Positions of overlapping EXONS



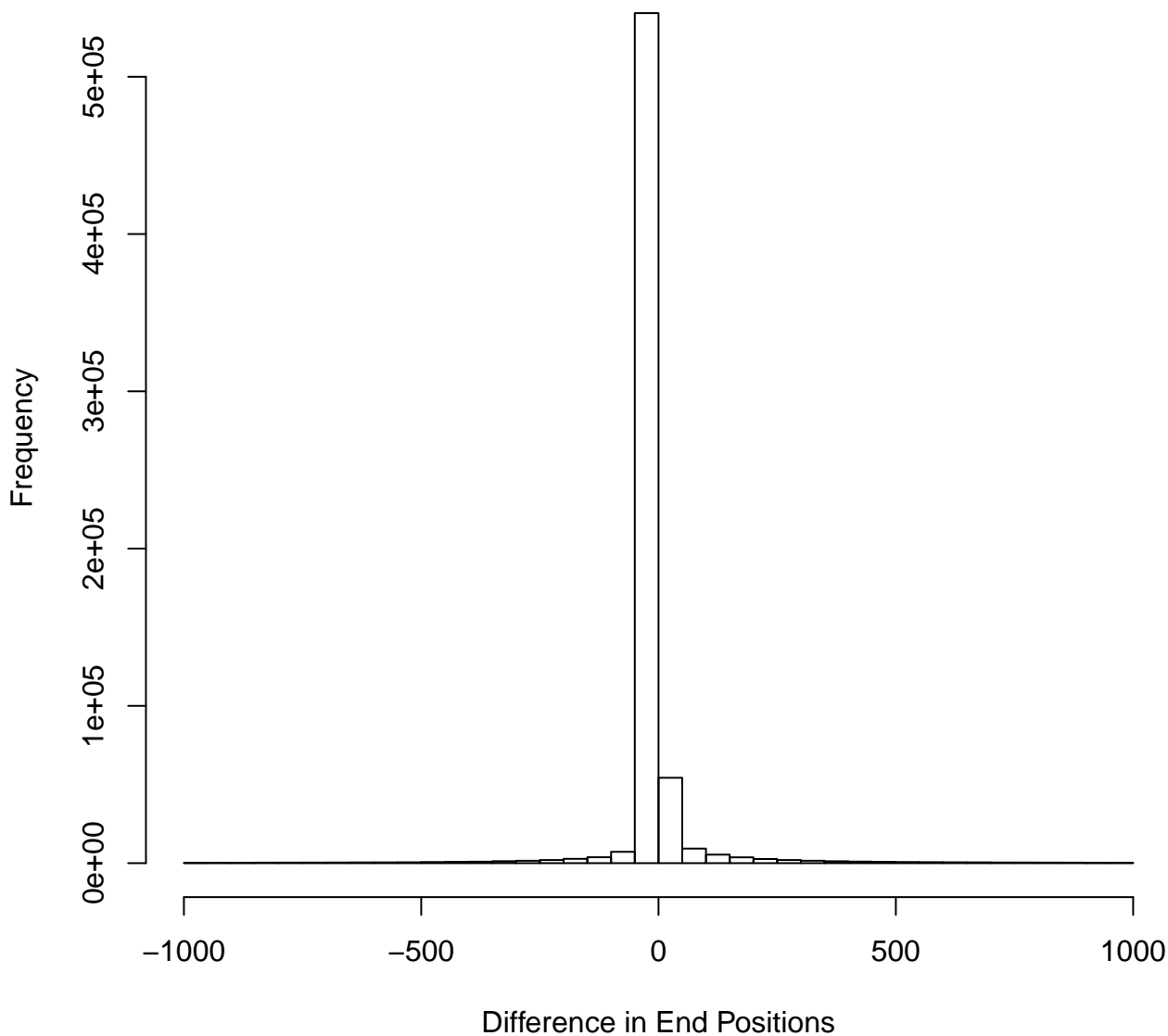
Histogram of Differences in End Positions of overlapping EXONS



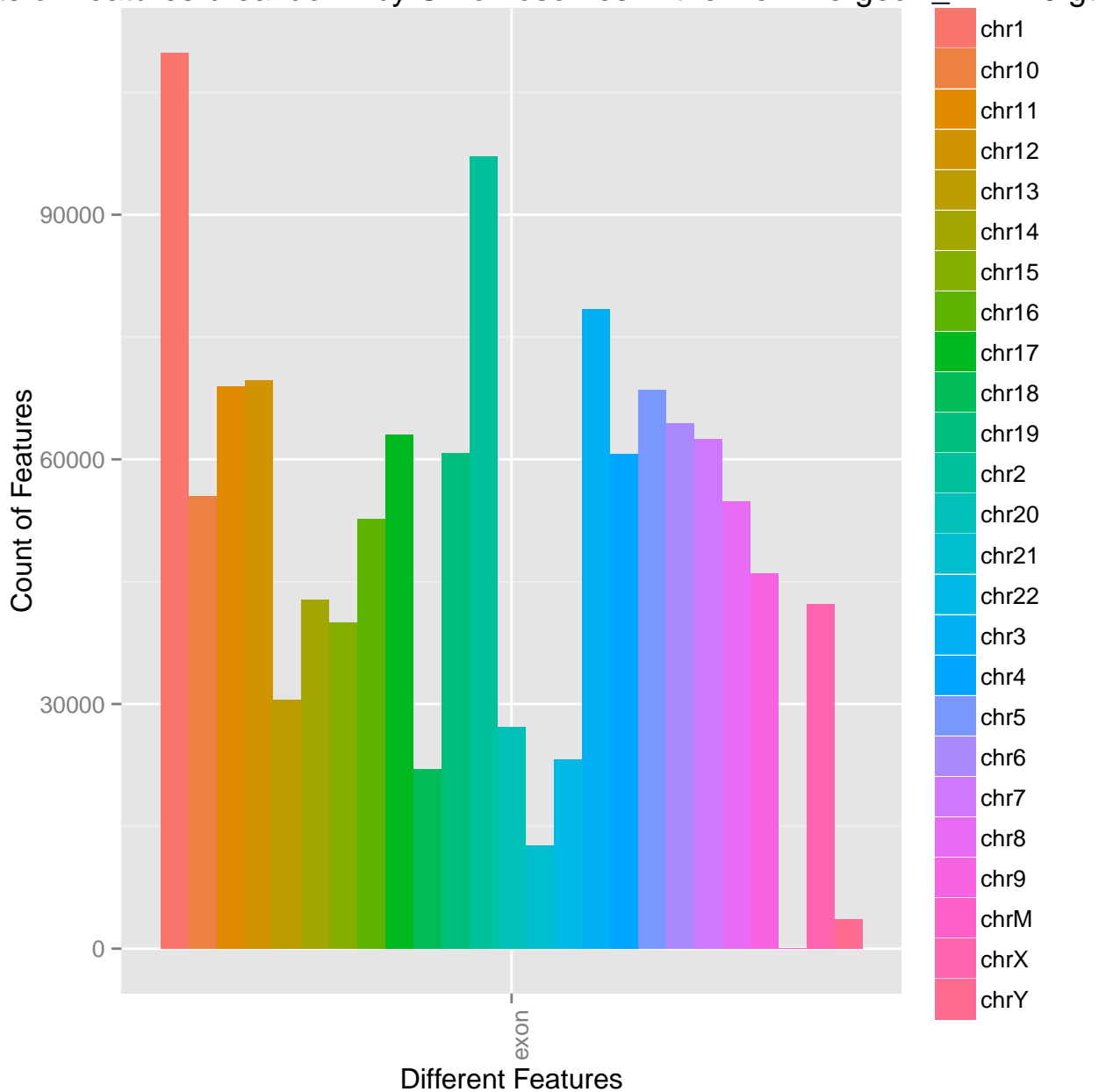
# Histogram of Differences in Start Positions of overlapping EXONS



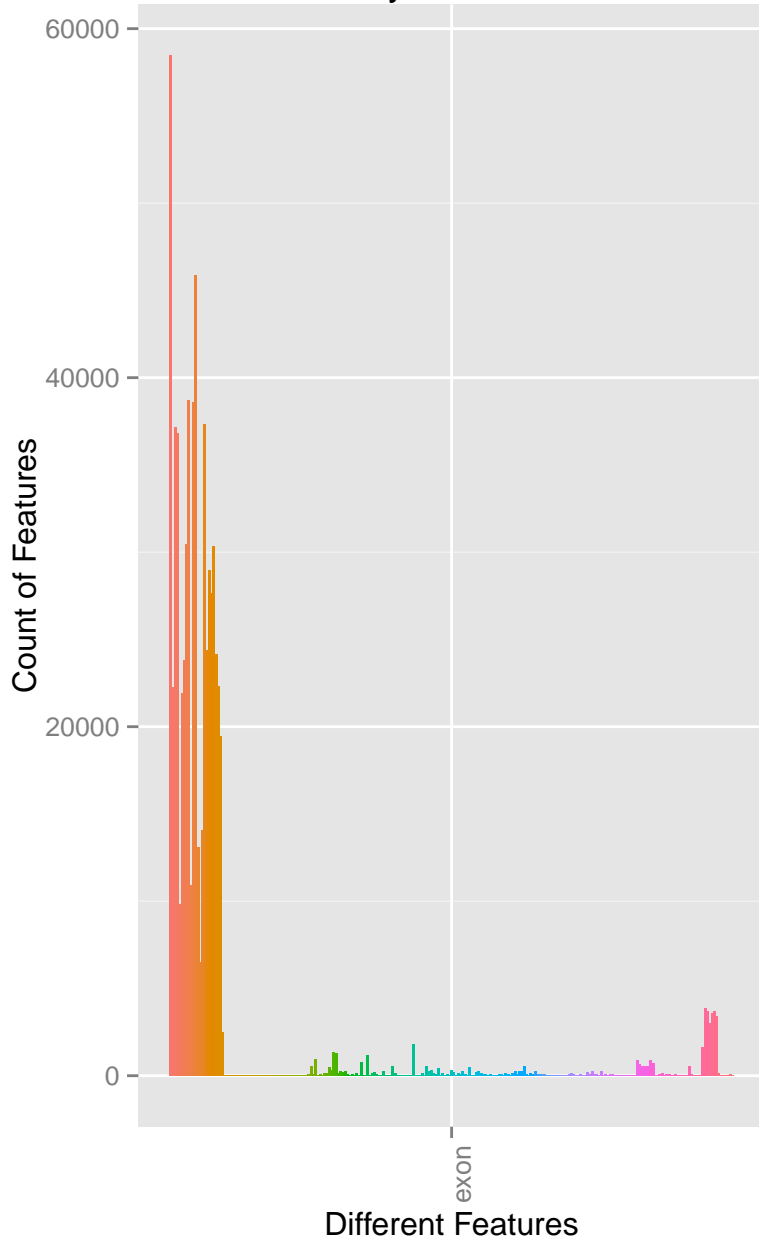
# Histogram of Differences in End Positions of overlapping EXONS



Counts of Features breakdown by Chromosomes in the file- mergedX chromosomes



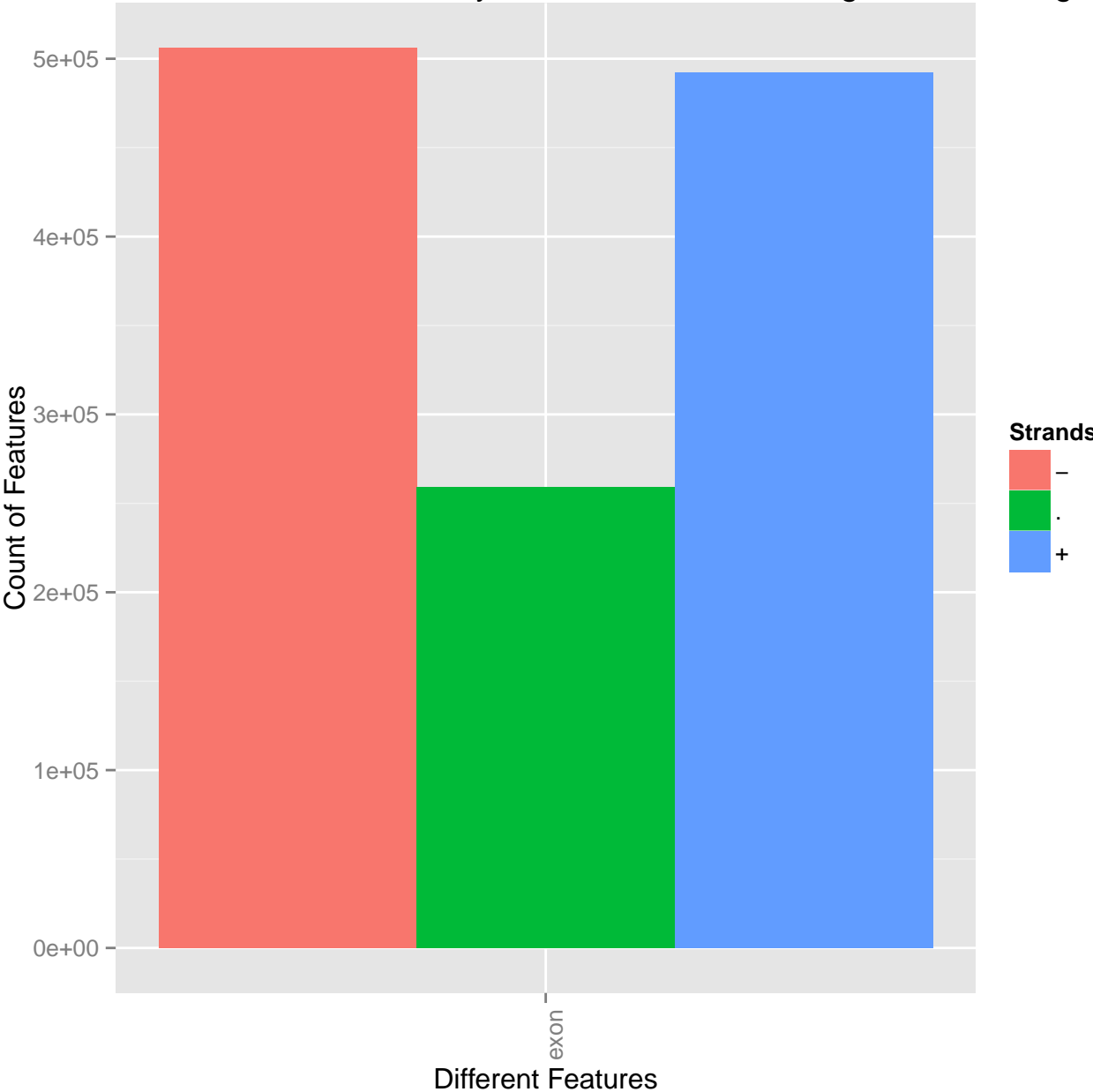
of Features breakdown by Chromosomes in the file – merged Stiff



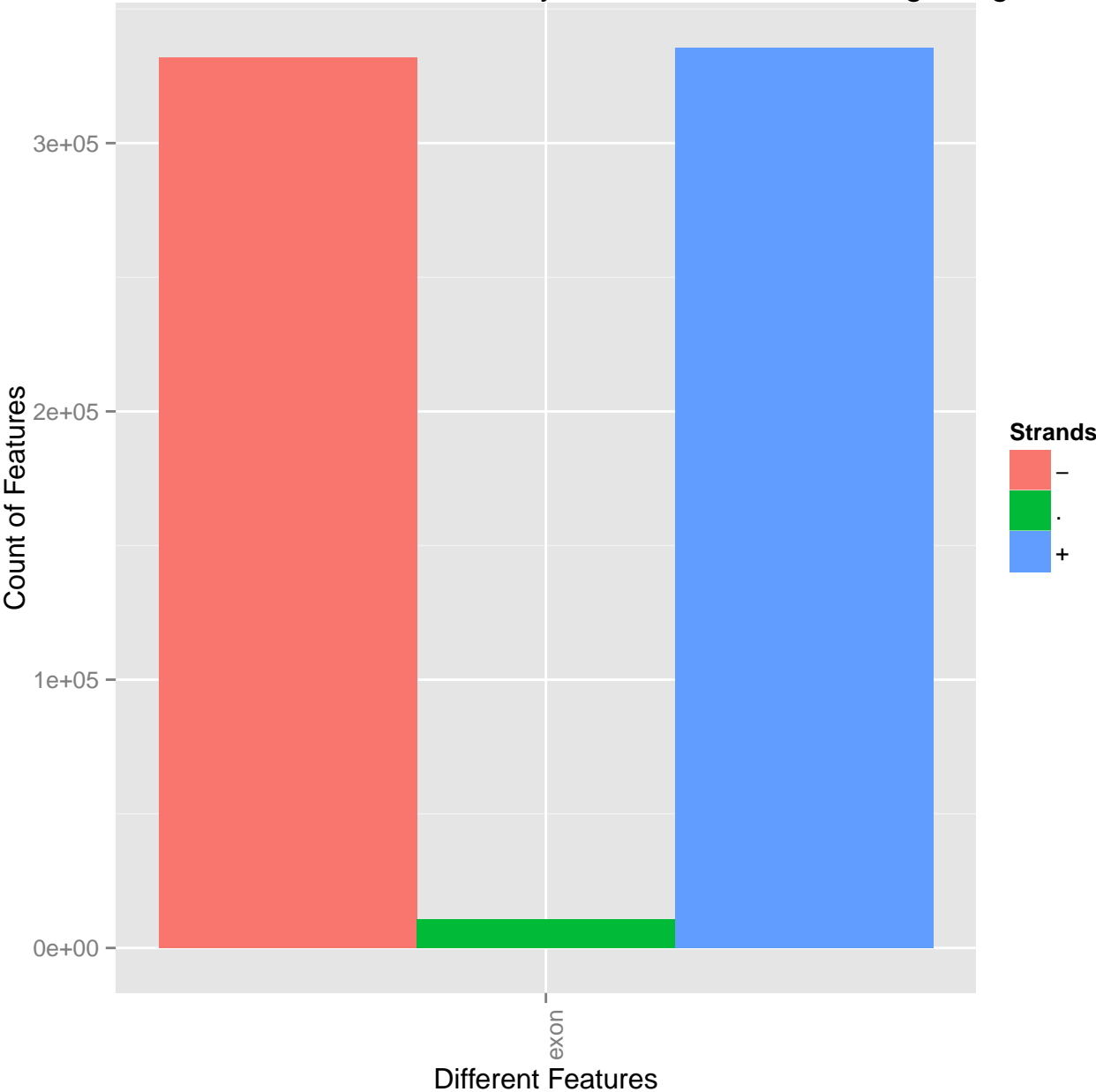
HG1501_PATCH
HG151_NOVEL_TEST
HG1592_PATCH
HG183_PATCH
HG185_PATCH
HG186_PATCH
HG19_PATCH
HG243_PATCH
HG256_PATCH
HG271_PATCH
HG27_PATCH
HG280_PATCH
HG281_PATCH
HG299_PATCH
HG29_PATCH
HG305_PATCH
HG306_PATCH
HG311_PATCH
HG325_PATCH
HG329_PATCH
HG339_PATCH
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HG375_PATCH
HG385_PATCH
HG388_HG400_PATCH
HG414_PATCH
HG417_PATCH
HG419_PATCH



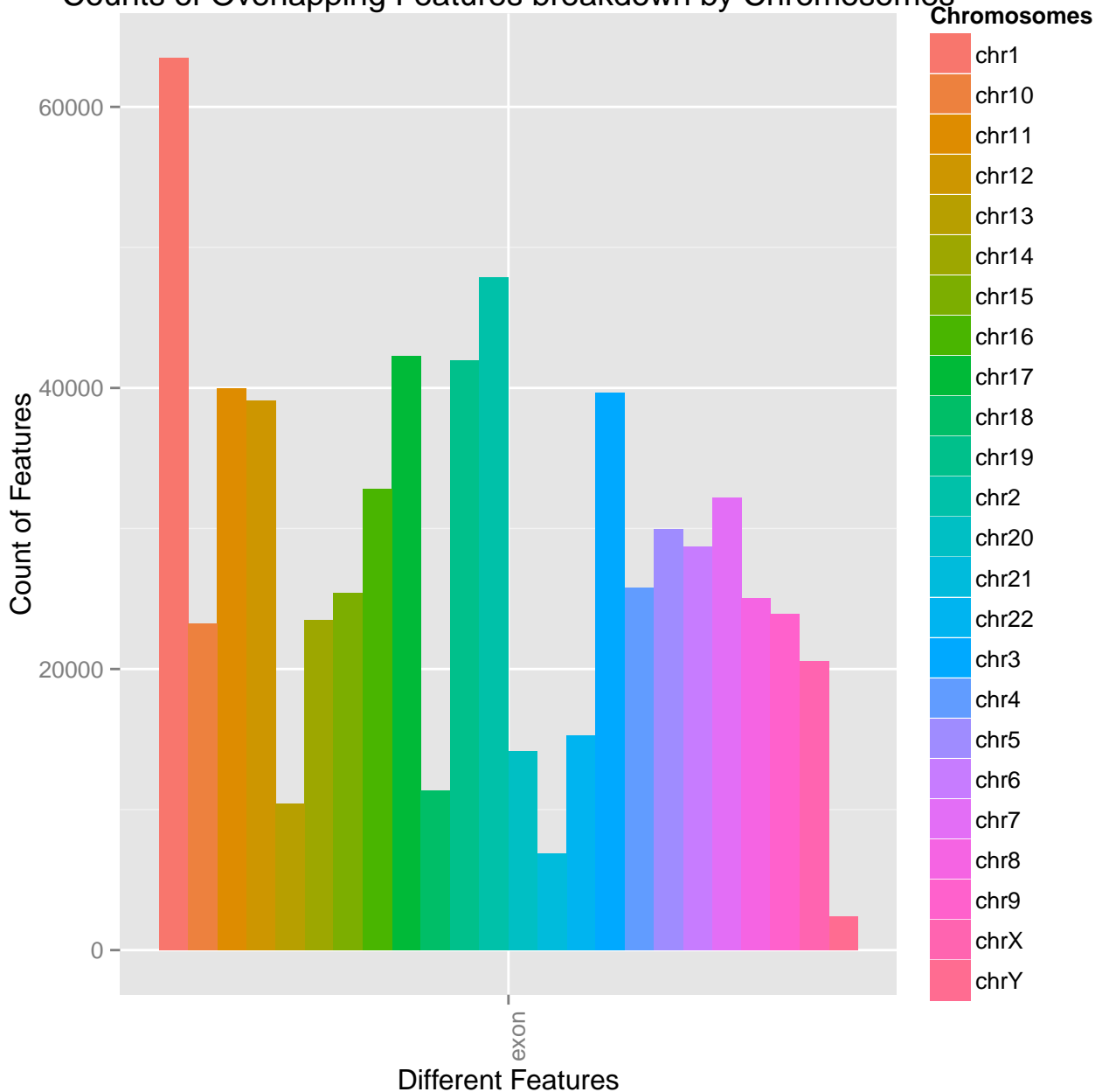
Counts of Features breakdown by Strands in the file– mergedX\_nmm10.gtf



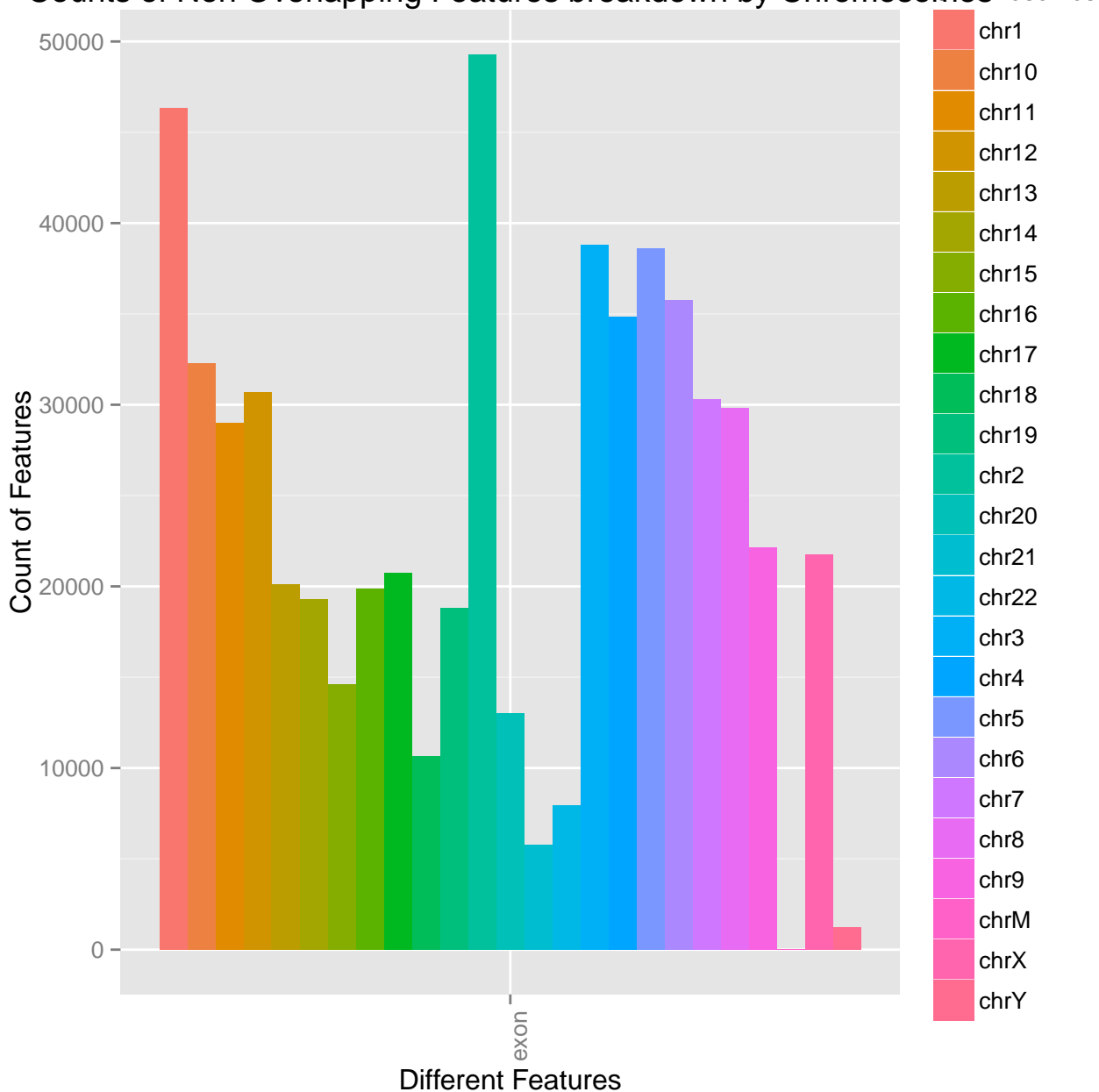
Counts of Features breakdown by Strands in the file– mergedS.gtf



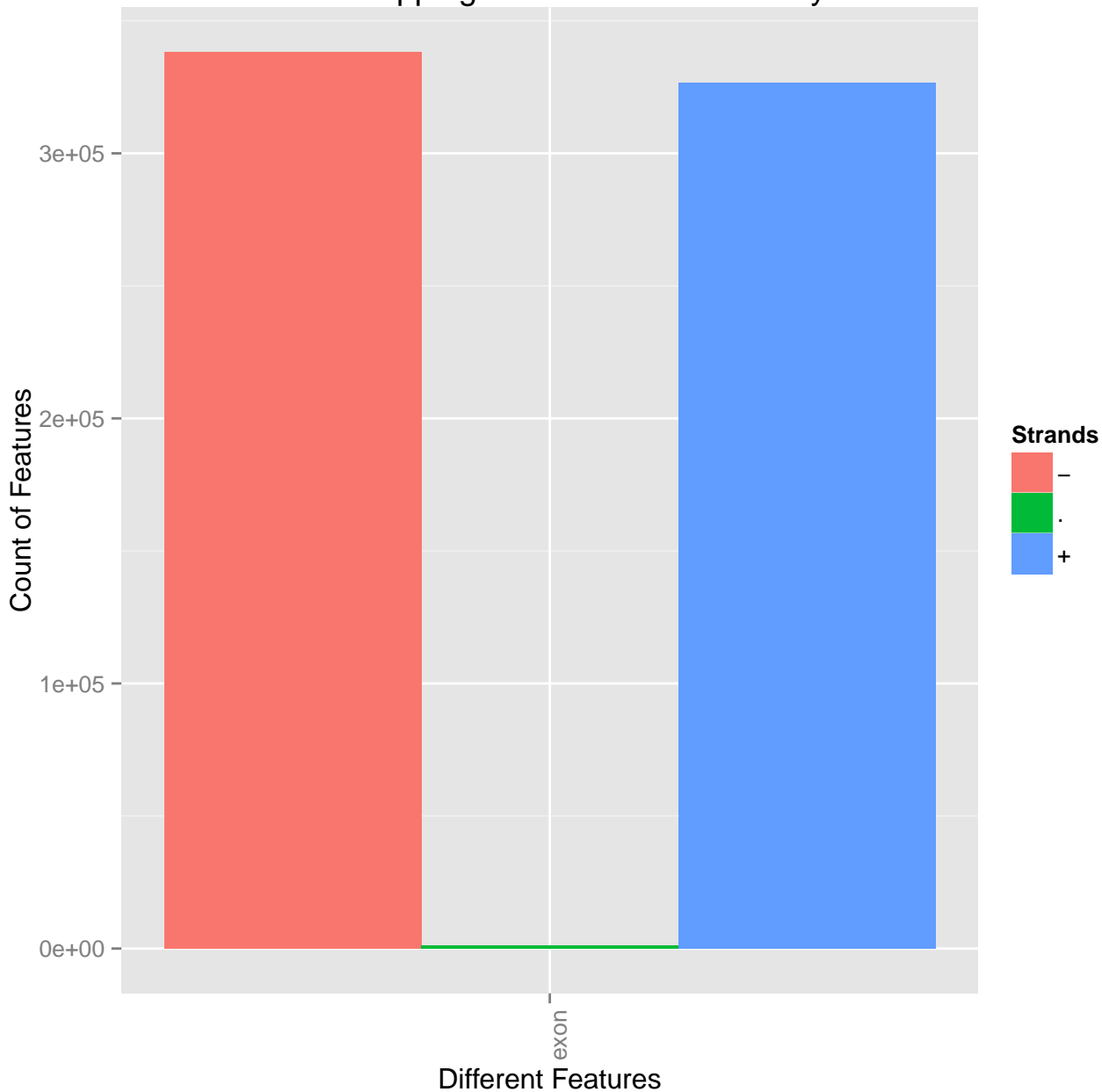
# Counts of Overlapping Features breakdown by Chromosomes



# Counts of Non Overlapping Features breakdown by Chromosomes



Counts of Overlapping Features breakdown by Strands



Counts of Non Overlapping Features breakdown by Strands

