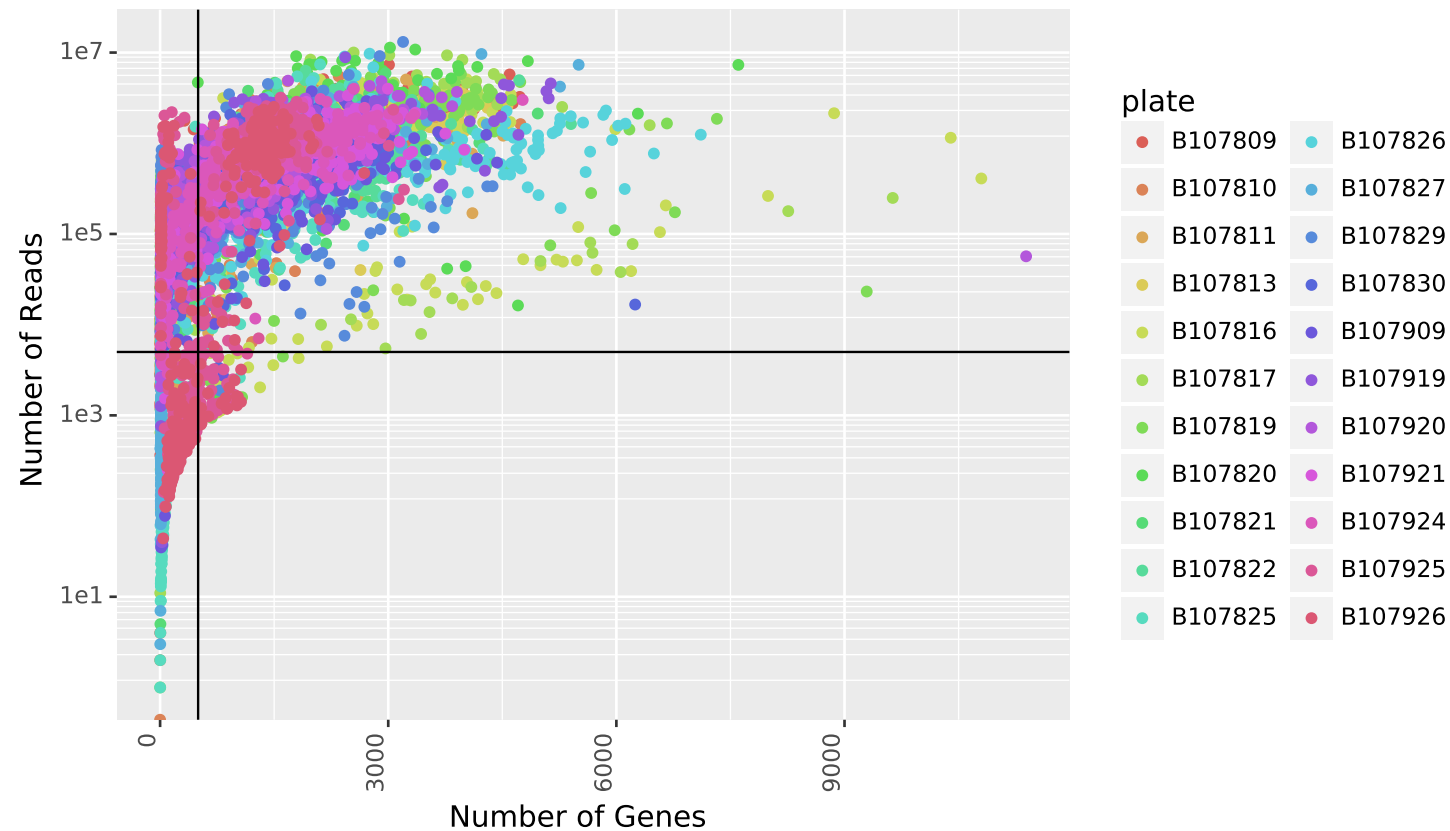
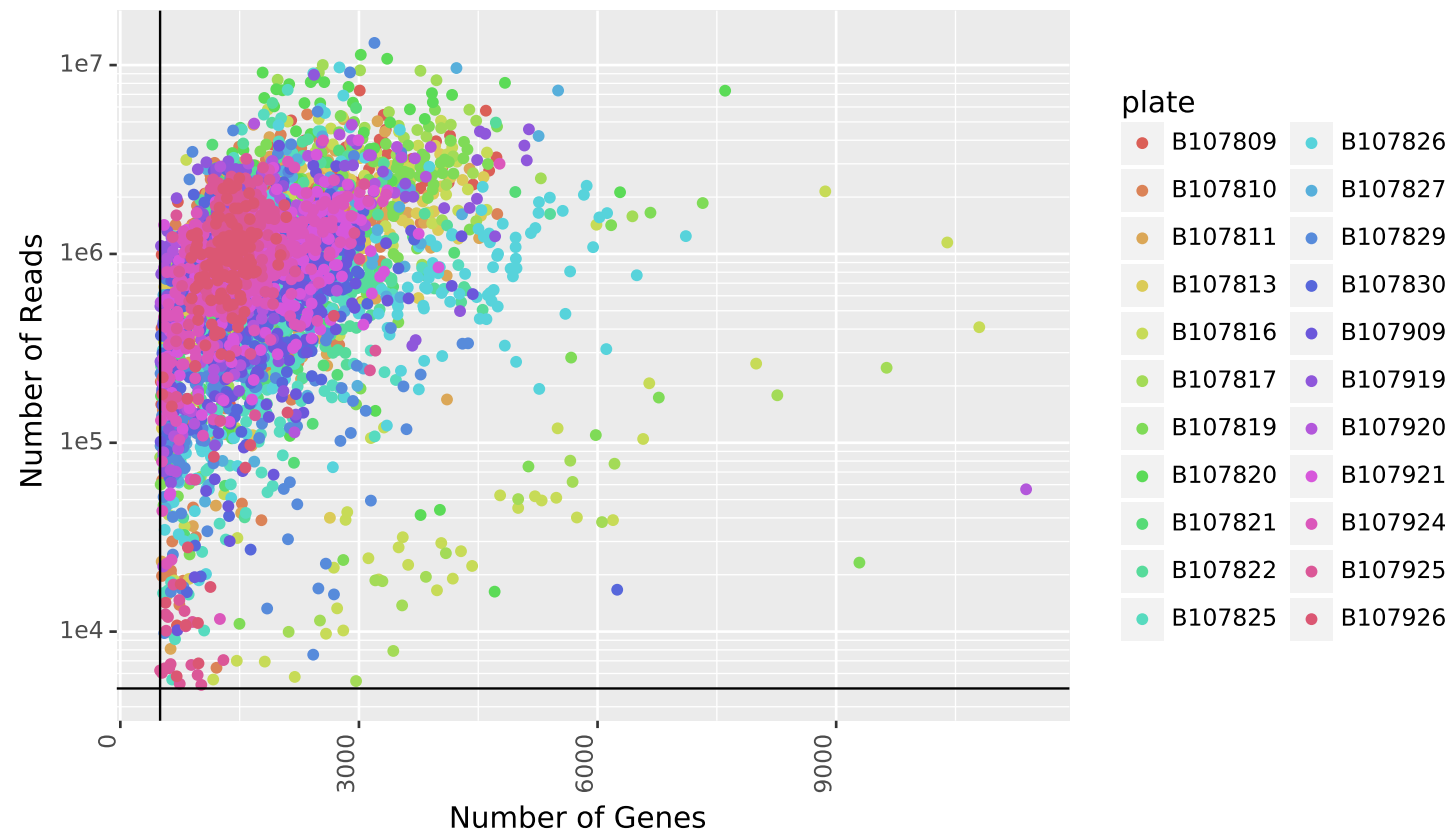


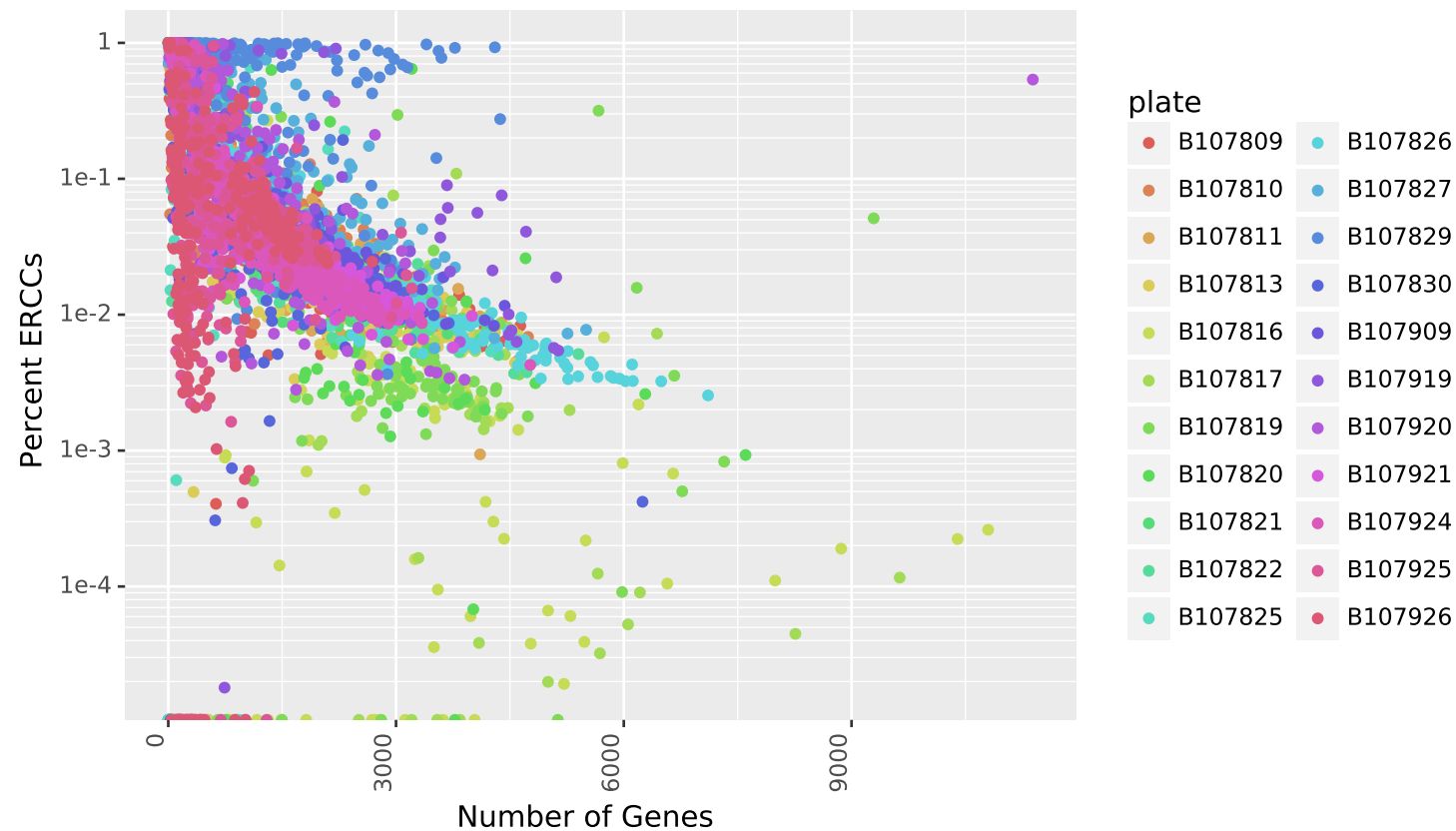
Number of Reads versus Number of Genes (Unfiltered)



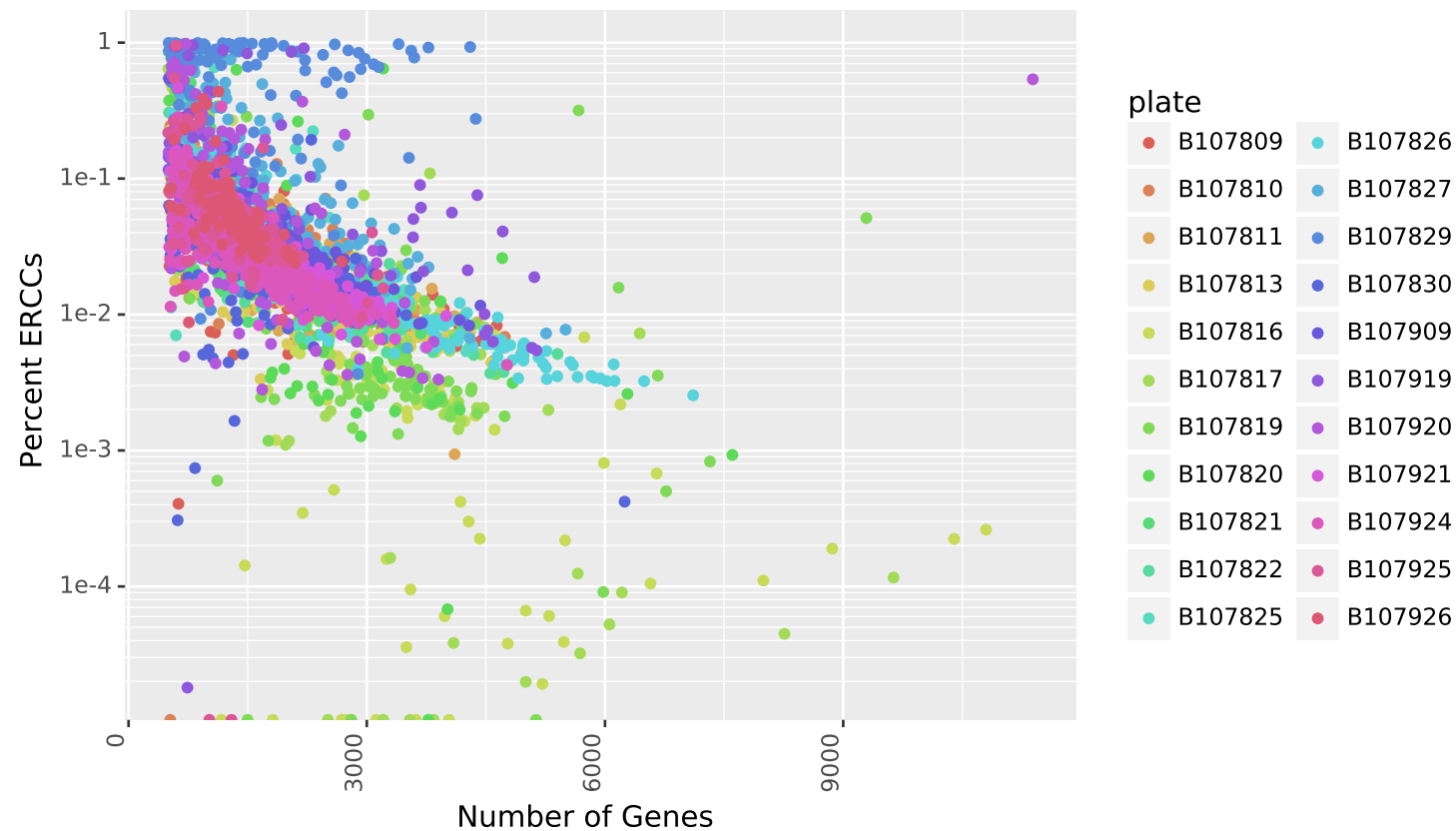
Number of Reads versus Number of Genes (Filtered)



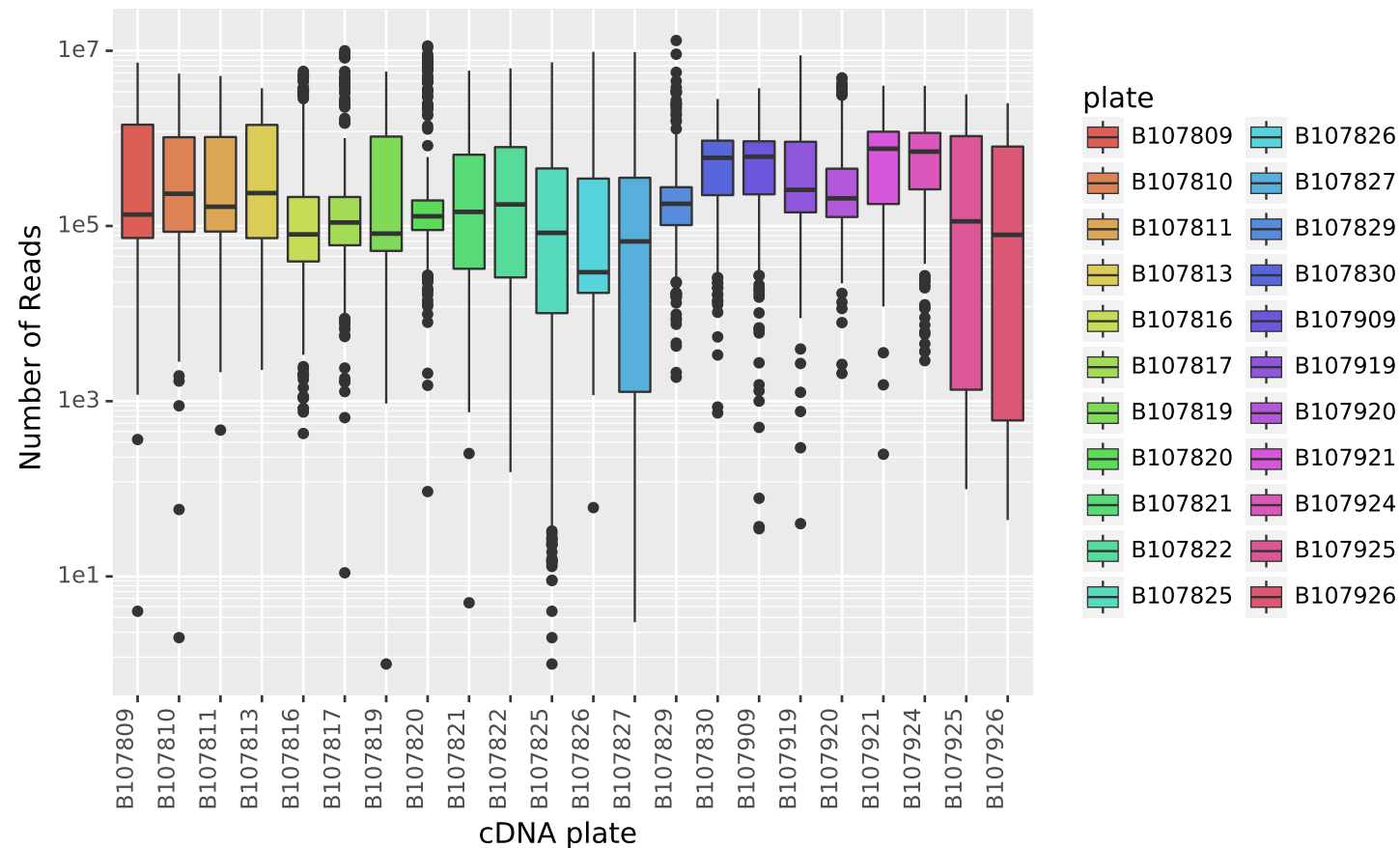
Percent ERCCs versus Number of Genes (Unfiltered)



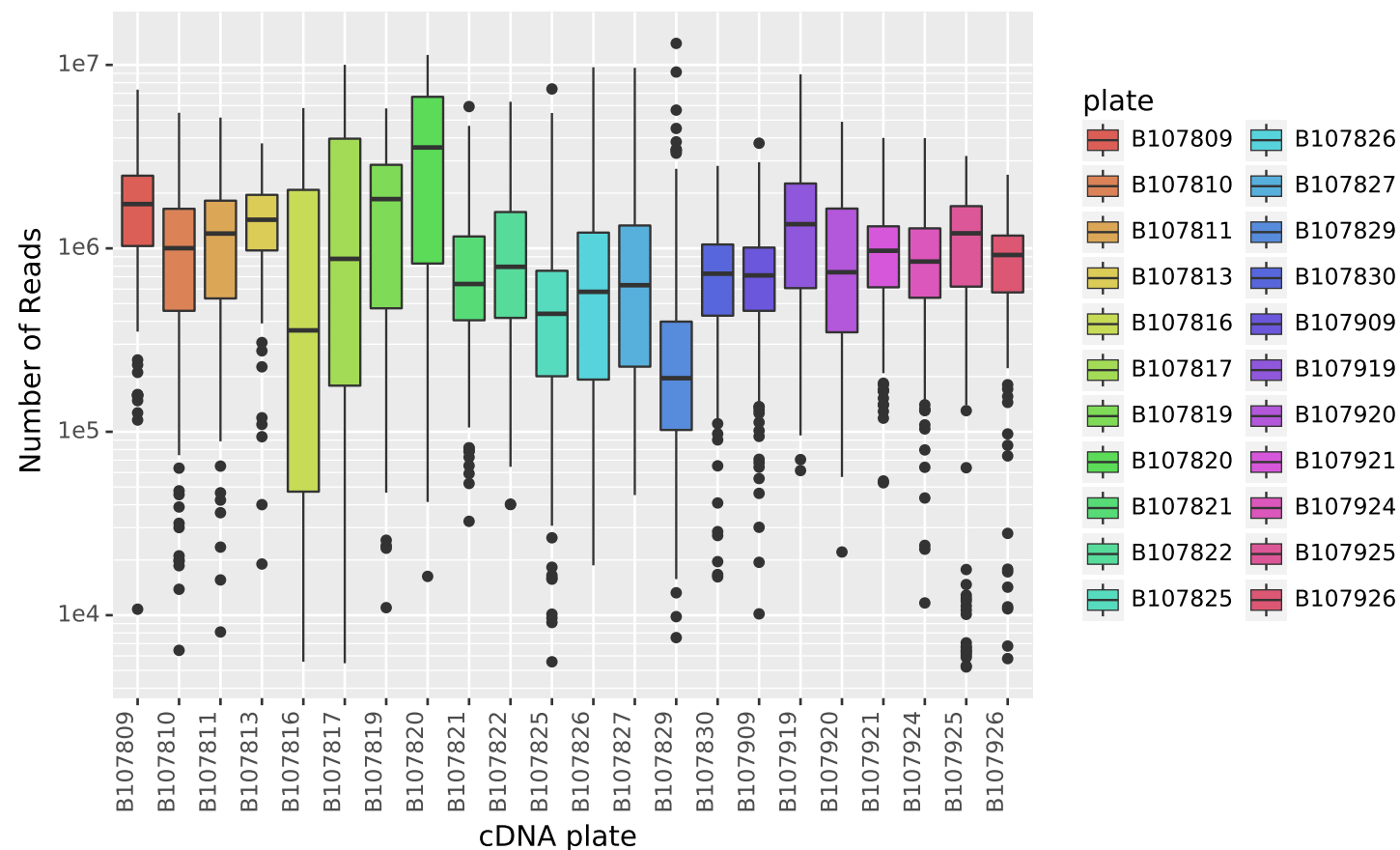
Percent ERCCs versus Number of Genes (Filtered)



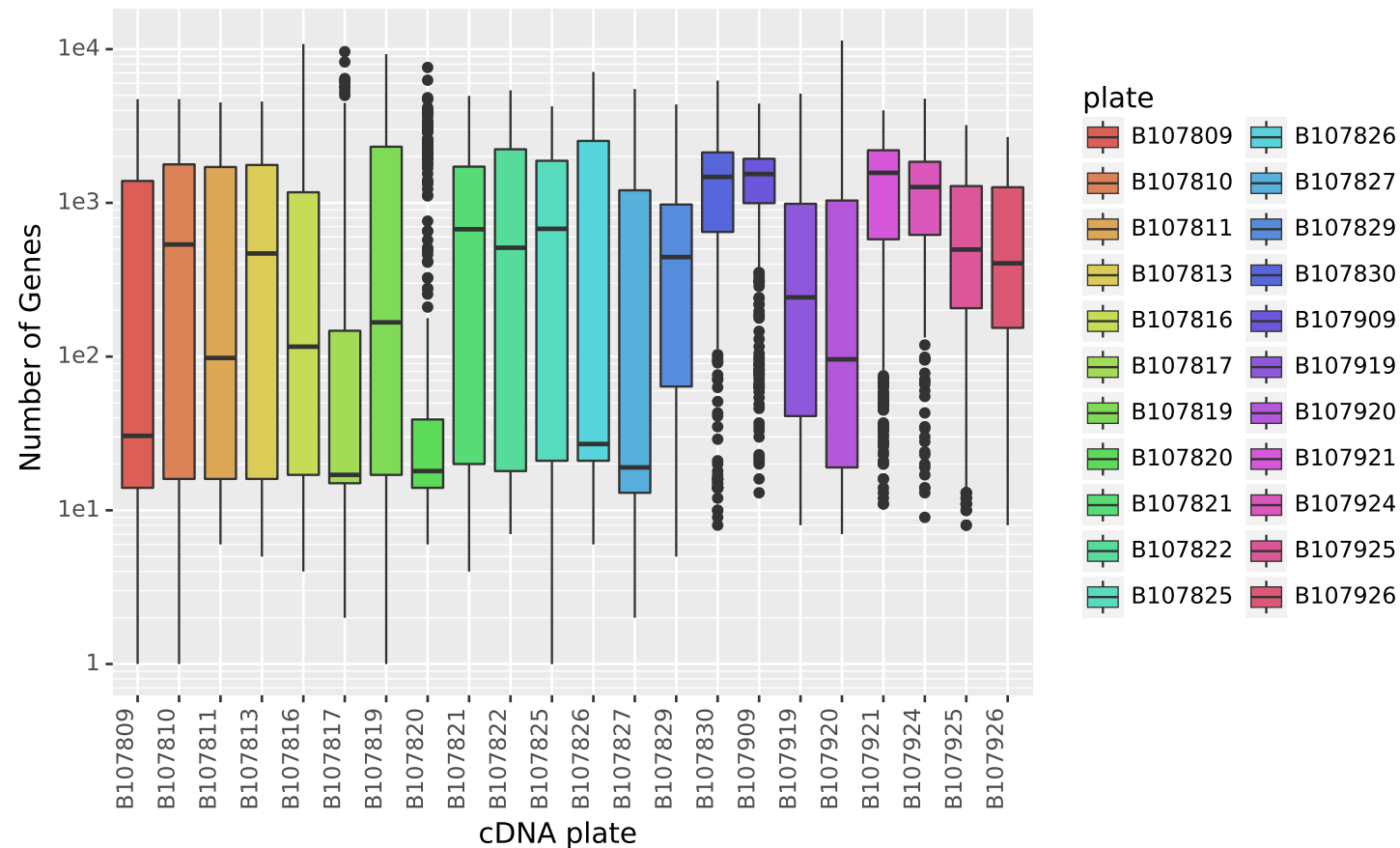
Number of Reads for Each cDNA Plate (Unfiltered)



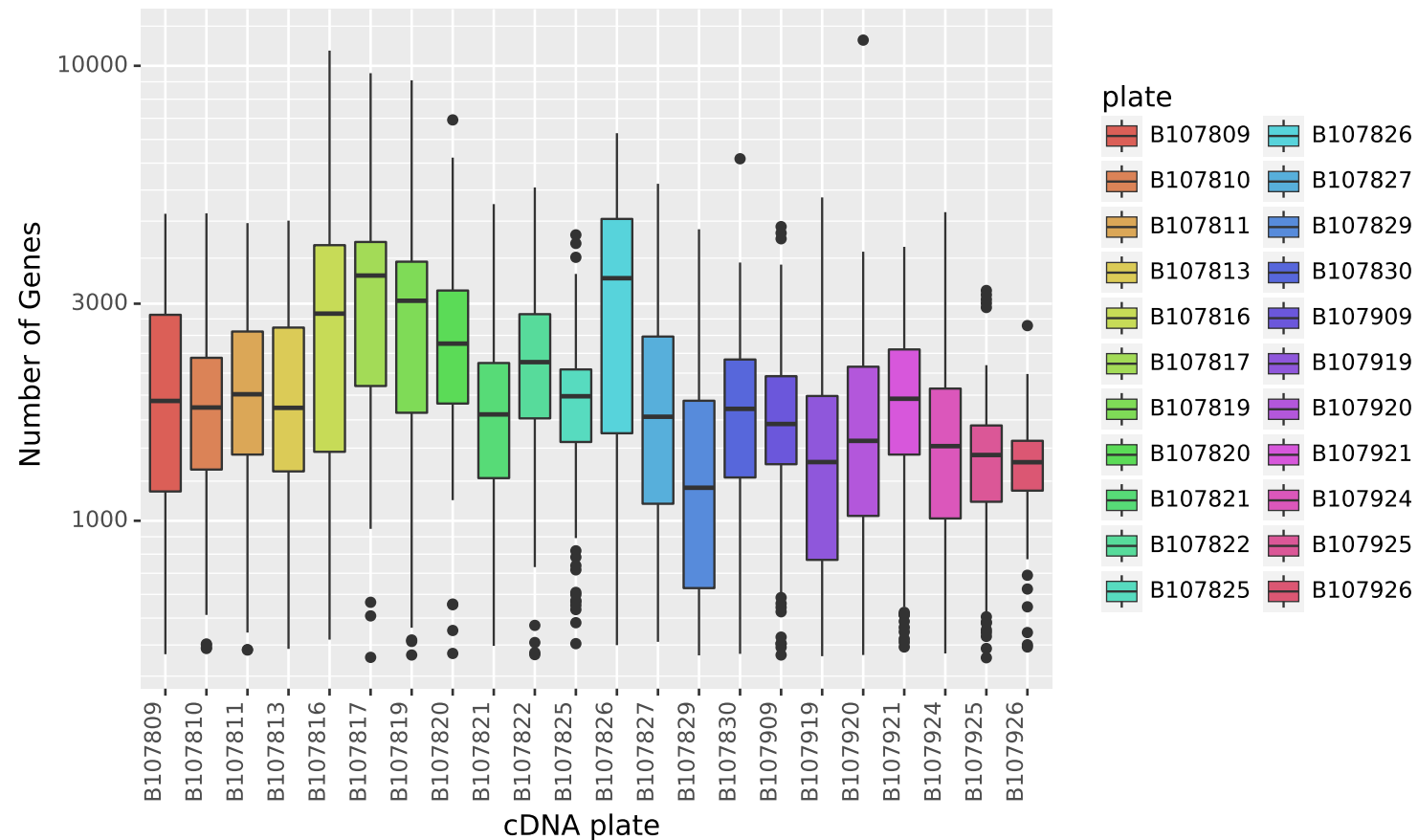
Number of Reads for Each cDNA Plate (Filtered)



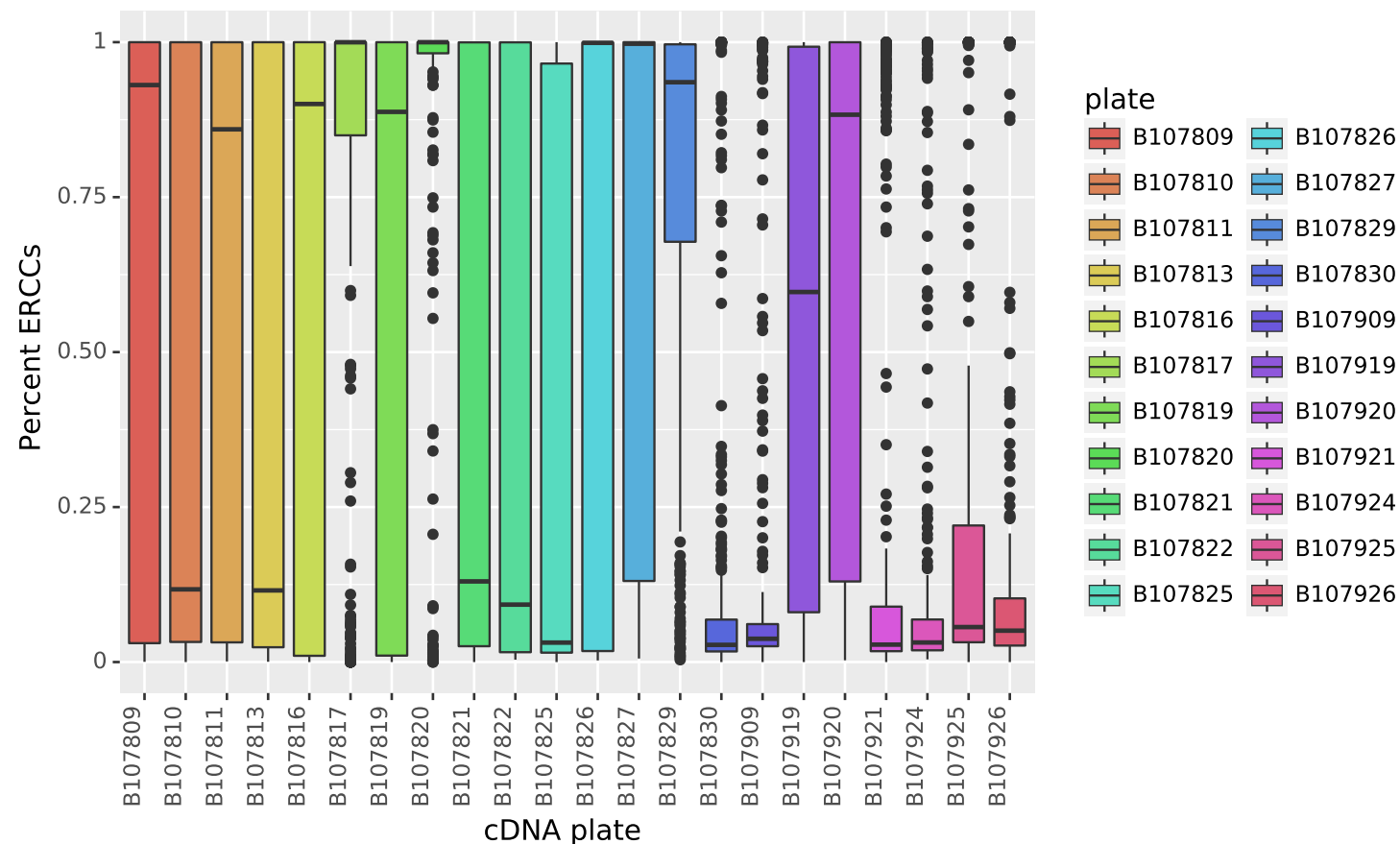
Number of Genes for Each cDNA Plate (Unfiltered)



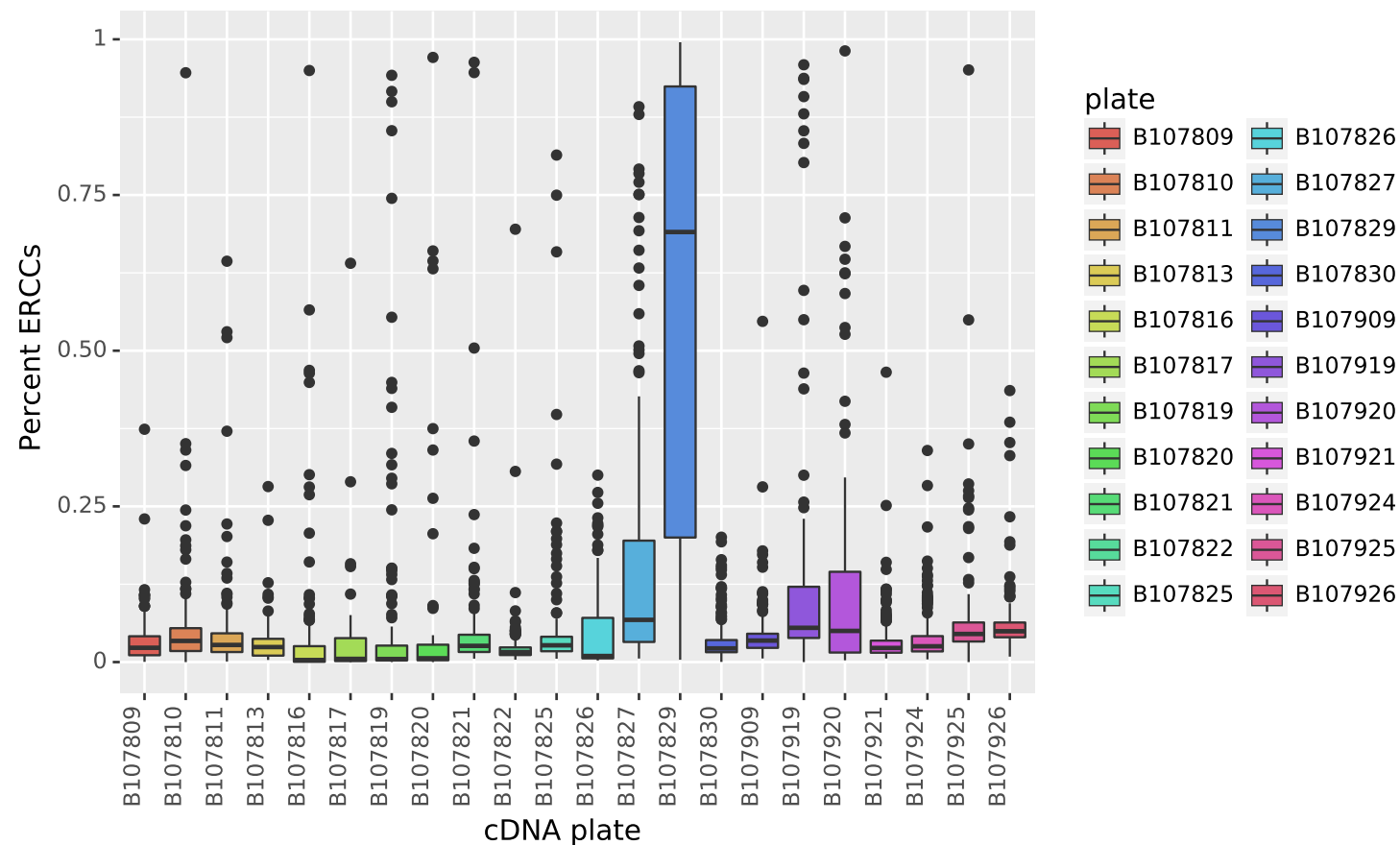
Number of Genes for Each cDNA Plate (Filtered)



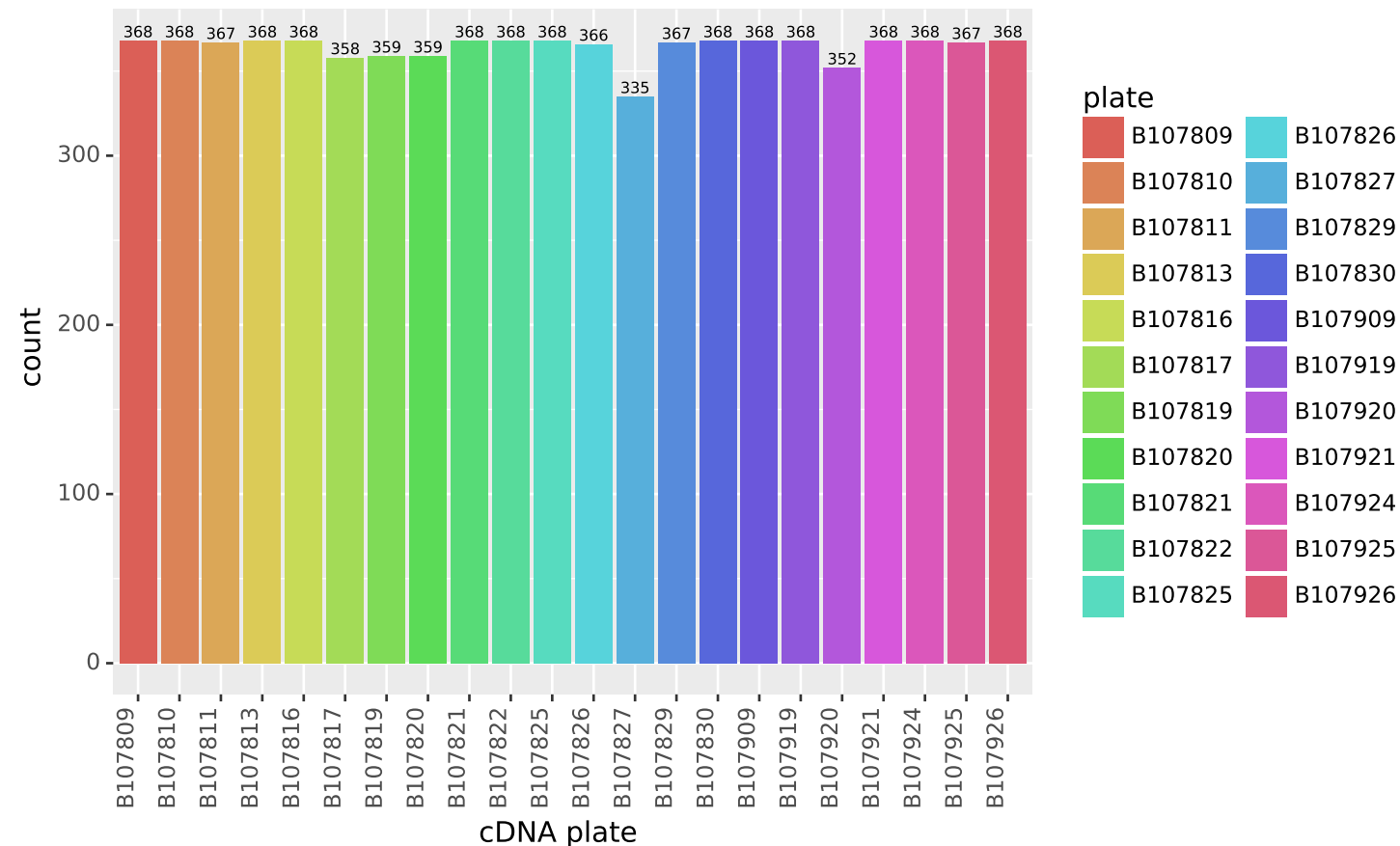
Percent ERCCs for Each cDNA Plate (Unfiltered)



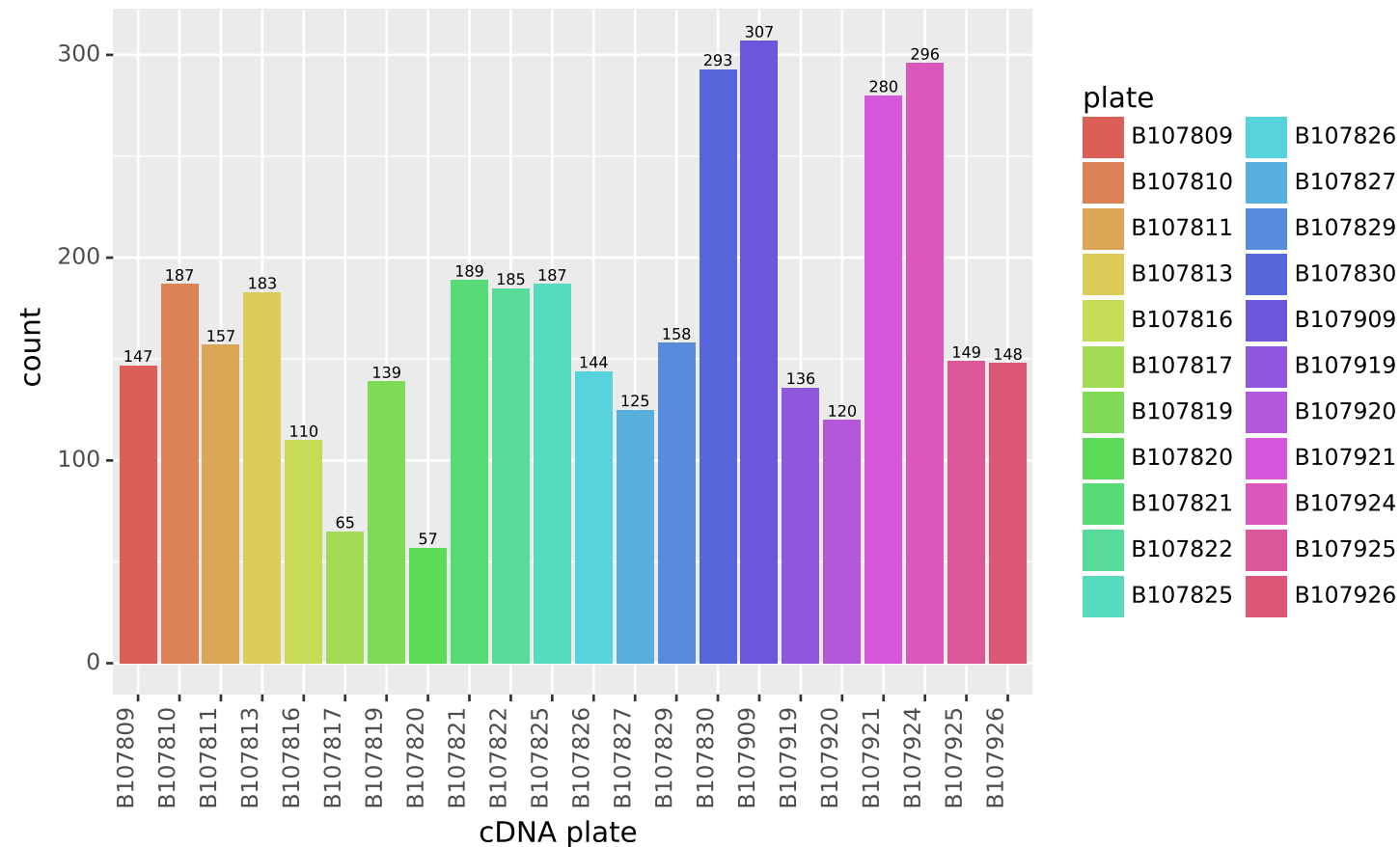
Percent ERCCs for Each cDNA Plate (Filtered)



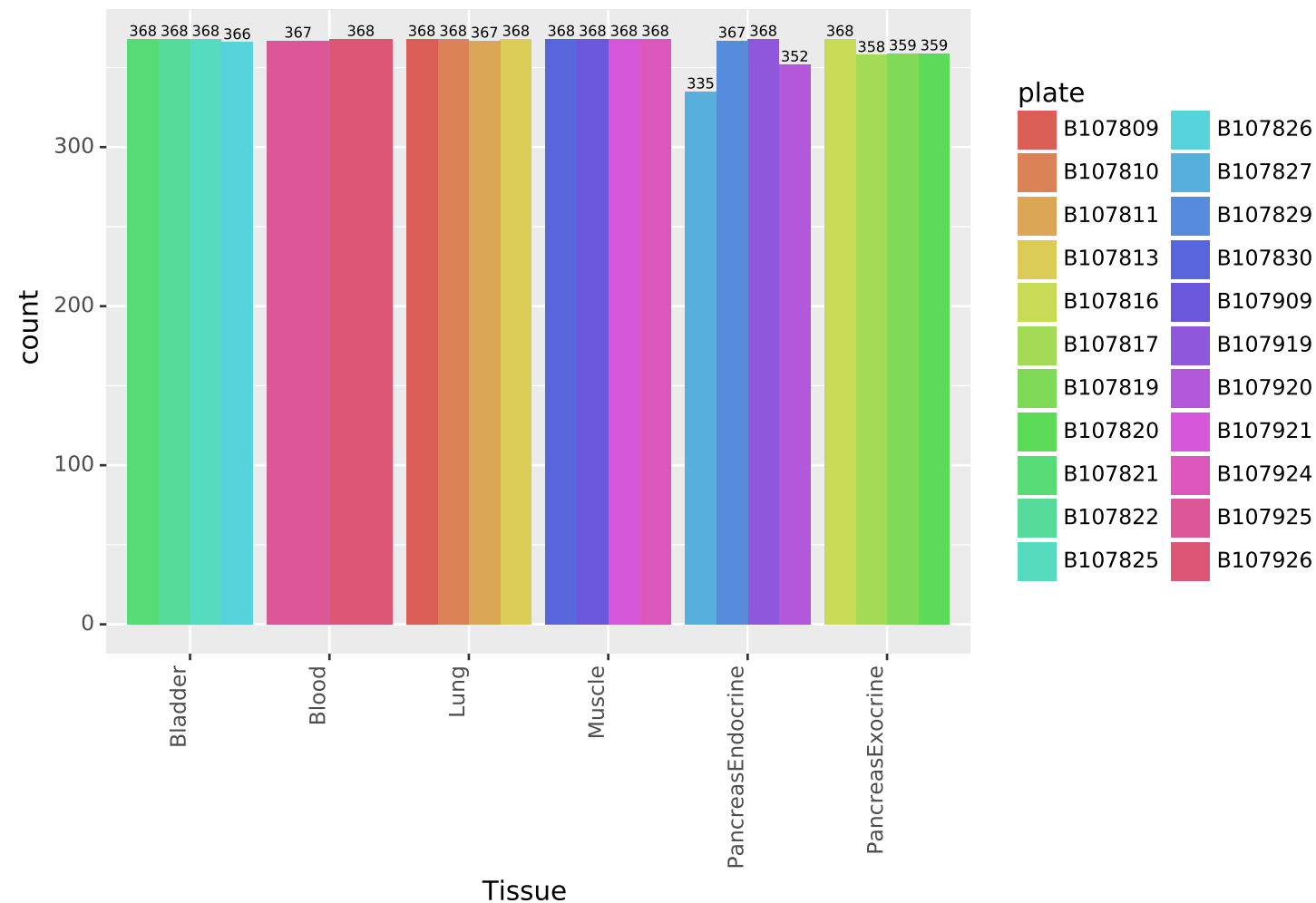
Number of Cells for Each cDNA Plate (Unfiltered)



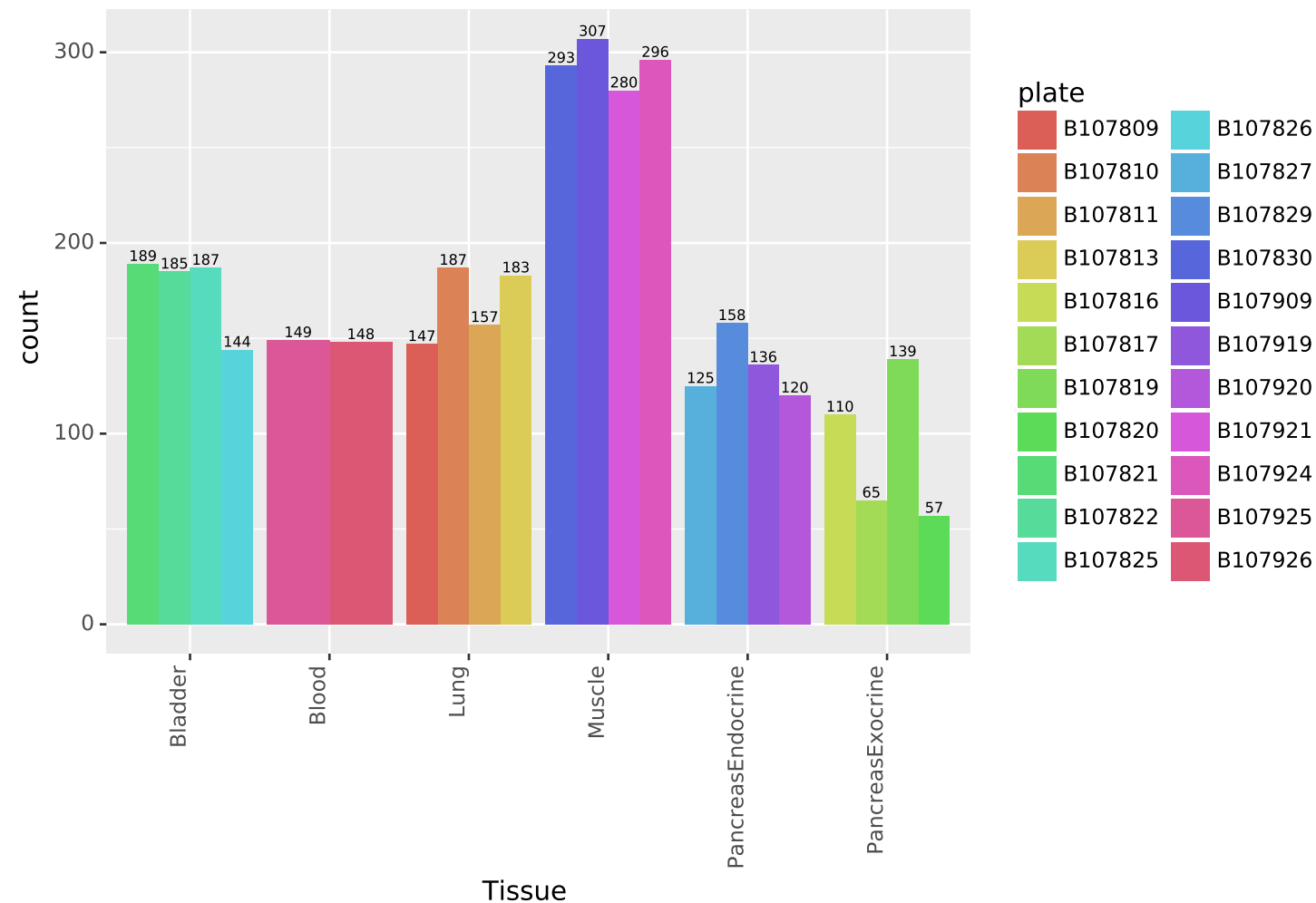
Number of Cells for Each cDNA Plate (Filtered)



Number of Cells for Each Tissue (Unfiltered)



Number of Cells for Each Tissue (Filtered)



Fraction of Cells Passed QC in Each Tissue for Each Plate

