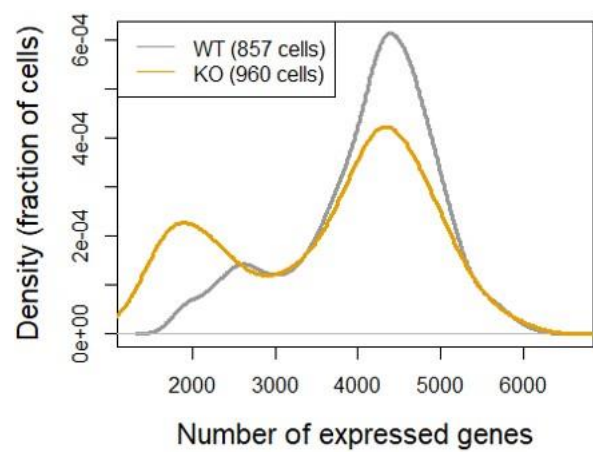


Supplementary Figure 1



Supplementary Figure 1. Characteristics of scRNA-seq with density plot of number of expressed genes per cell. 960 and 857 transcriptomes (filtered and normalized with Seurat) from *Srf* KO and WT iPSCs, respectively, were analyzed.