########################

Rscript to generate SVs barplot

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**df <- read.csv("file\_name.csv", header=T)**

**df**

**df2 <- transform(df, genes = reorder(genes, -Freq))**

**require(ggplot2**)

############### Customized barplot ###############

**ggplot(df2, aes(x=genes, y=Freq)) +geom\_bar(stat="identity", fill="steelblue") + theme(axis.text.x = element\_text(angle=90, vjust=1, hjust=1, size=28, face = "bold"))+ theme(title = element\_text(face = "bold", size = 28))+ xlab("")+ ylab("Frequency")**