**Lab Members: Ignore! This is automated by the boot script!**

Prerequisites:

**If running for the first time run:**

install.packages("data.table")  
install.packages("RColorBrewer")

install.packages("shiny")

install.packages("gplots")

**Change working directory, then run:**

**setwd("R:/Medicine/Hematology-Oncology/Yu\_Lab/Nate/scripts\_and\_tools/YuLab-SU2C-TCGA-HeatmapApp")**

**library(shiny)**

**runApp("YuLab-SU2C-TCGA-HeatmapApp")**