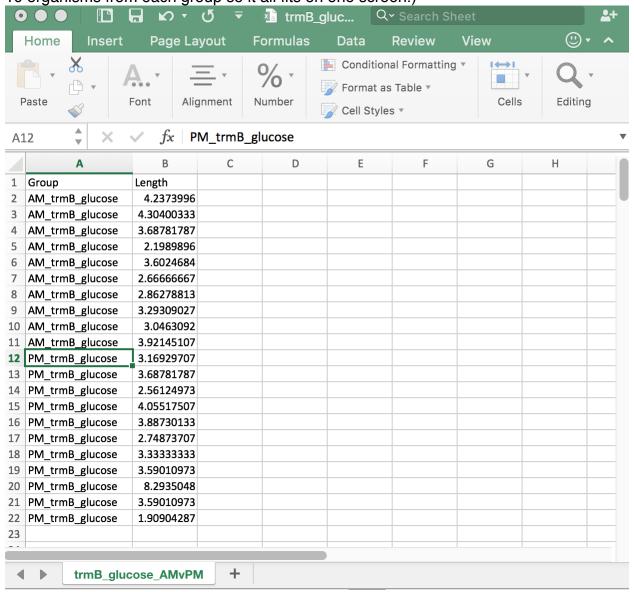
How to Make a Boxplot in R

We can make boxplots to compare two (or more) experimental conditions. First, prepare your data in Excel. You want each organism to have its own row in the sheet and each experimental condition to have a distinct name. In the example below, I am comparing trmB_glucose from the AM group (AM_trmB_glucose) with trmB_glucose from the PM group (PM_trmB_glucose). (I've shrunk the data set in the picture below to only include 10 organisms from each group so it all fits on one screen.)



Save as a .csv file.

Download and install R (https://cloud.r-project.org) and RStudio Desktop (https://www.rstudio.com/products/rstudio/download/)

Open RStudio

```
To install ggplot2:
> install.packages("ggplot2")
To load ggplot2:
> library(ggplot2)
To import your data:
> data2 <- read.csv(file.choose(), header=TRUE)</pre>
To see what the data looks like to R:
> View(data2)
To make a simple box plot:
> ggplot (data=data2, aes(data2$Group, data2$Length)) +
geom_boxplot()
To add labels:
> ggplot (data=data2, aes(data2$Group, data2$Length)) +
geom_boxplot()
     + labs(x = "Group", y = "Length (um)")
To format the plot theme:
panel.grid.minor = element_blank(), panel.background =
element_blank(), axis.line = element_line(color = "black"))
```

Export your plot after doing any other customization you choose.

