Box Plots and Scatter Plots

1 Load the Hot Dog Data

Fill in the blanks below to read the file hotdog.csv into R and reorder the levels of the Day factor. You should have this file from the previous workshop. Make sure the file is in your working directory.

Check that the file was read in correctly and that the levels of the Day factor are reordered. The structure of your data should look like this:

```
str(hd)
## 'data.frame': 54 obs. of 5 variables:
## $ Day : Factor w/ 5 levels "Mon","Tue","Wed",..: 1 1 1 1 1 1 1 1 1 1 1 1 1 ...
## $ Brand : Factor w/ 2 levels "Nathans","OscarMayer": 1 1 1 1 1 2 2 2 2 2 2 ...
## $ Type : Factor w/ 3 levels "Beef","Meat",..: 2 3 3 3 3 1 1 1 2 3 ...
## $ Calories: int 175 129 102 135 142 190 153 132 173 152 ...
## $ Sodium : int 507 430 542 426 513 587 401 253 458 588 ...
```

If you can't get this structure, ask me for help.

1.1 Questions

- 1. What do the arguments 'skip=', 'nrows=' and 'header=' do in the read.csv() function?
- 2. Why do we reorder the levels of the Day column?
- 3. **True or False**: We reorder the levels of the Day column by overwriting the original column?

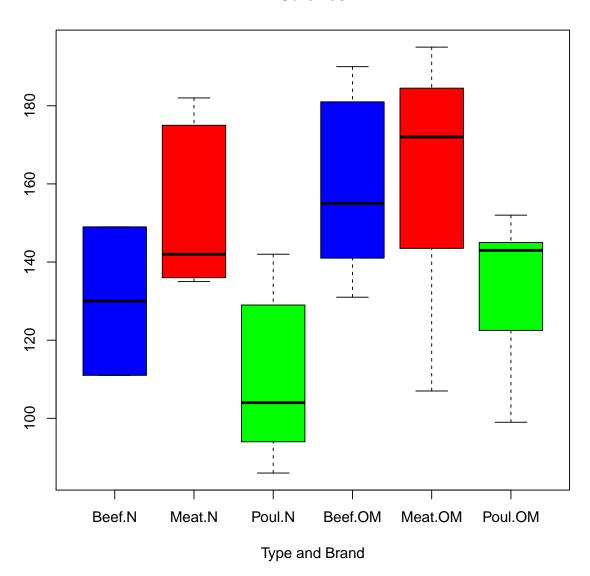
2 Box Plots

Fill in the blank arguments to get the box plot on the next page. Run your code to see if you get the same plot.

2.1 Questions and Challenges

- 1. What does the argument 'axt = "n"' do in the boxplot() function?
- 2. What do the arguments 'side = 1' and 'at = 1:__' do in the axis() function?
- 3. Create a box plot of Sodium by Type. Include the following:
 - A main title
 - Custom x and y axes titles
 - Make the box plots horizontal
 - Different colors for each hot dog type

Calories



3 Scatter Plots

Fill in the blank arguments to get the scatter plot on the next page. Run your code to see if you get the same plot.

```
plot(______,
    data = _____,
    pch = c(15, 16)[_____],
    col = c(___, ___, ___)[hd$Type],
    main = _____)

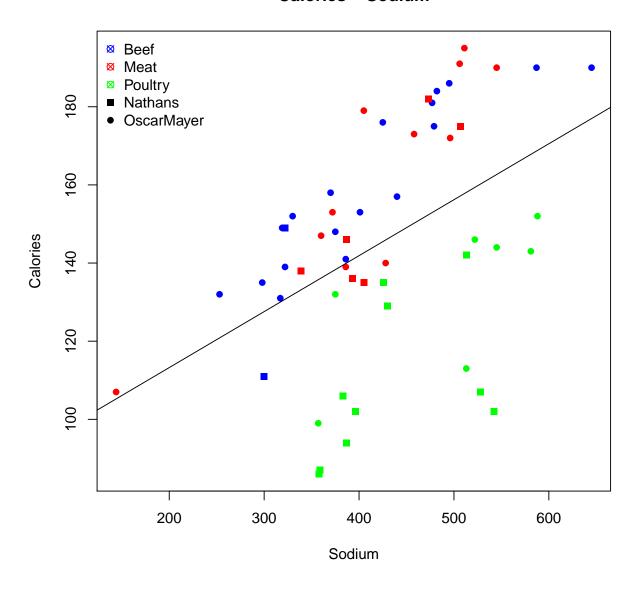
abline(lm(Calories ~ Sodium, data = hd))

legend(x = "top____",
    legend = c(levels(hd$____), levels(hd$____)),
    pch = c(rep(13, 3), 15, 16),
    col = c(____, ___, ____),
    bty = _____)
```

3.1 Questions and Challenges

- 1. What does the argument 'pch=' do in the plot() function?
- 2. What does the command 'c(15, 16)[____]' do?
- 3. Instead of using 'x = "top____" in the legend() function what can you use?
- 4. What does the argument 'bty=' do in the legend() function?
- 5. Create a scatter plot of Sodium and Type. Include the following:
 - Sodium on the y-axis, Calories on the x-axis
 - A main title
 - Different colors for each brand
 - Different shapes for each day (hint: pch)
 - Bonus: Add a line of best fit and a legend

Calories ~ Sodium



R Session Information

```
R version 3.0.2 (2013-09-25)
Platform: x86_64-apple-darwin10.8.0 (64-bit)

locale:
[1] en_US.UTF-8/en_US.UTF-8/en_US.UTF-8/C/en_US.UTF-8/en_US.UTF-8

attached base packages:
[1] stats graphics grDevices utils datasets methods base

other attached packages:
[1] knitr_1.5

loaded via a namespace (and not attached):
[1] digest_0.6.4 evaluate_0.5.1 formatR_0.10 highr_0.3
[5] stringr_0.6.2 tools_3.0.2
```