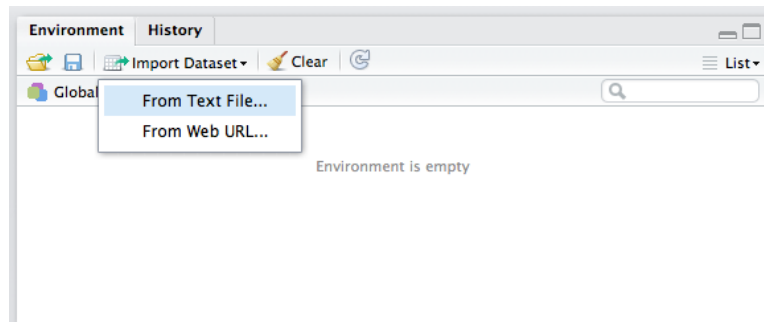


1/15/14 Class Notes

I. Step-by-Step: Using R

- a. Open the pre-downloaded files: File → Open File
 - i. Make sure that the file is an R file (Titanic.R)
 - ii. R files are scripts that may contain directions and commands to be used in the assignment
- b. Install R packages (“effects” & “mosaic”). Packages need to only be installed once. Once installed, they need to be loaded each time you open R
 - i. At the right-hand side of R Studio, look for the tab “Packages”
 - ii. Choose “Install Packages”
 - iii. In “Packages” section, type “mosaic”
 - iv. To reload, type `>library(mosaic)` in the console
 - v. Type the above command into the console OR highlight this command in the script and press CTRL+Enter (for PC) or Command+Enter (for MACs)
- c. To import data to R



- i. If dataset can be downloaded in .csv format:
 1. Environment→Import Dataset→From Text File
- ii. If dataset is accessed through a web link:
 1. Environment→Import Dataset→From Web URL

II. Categorical v. Quantitative Data

a. **Categorical Data**

- i. TitanicSurvival.R: Who survived and who died during the sinking of the Titanic?
 1. Data can be grouped into those who “survived” and “died”

b. **Quantitative/Numerical Data**

- i. Data that can be measured numerically (height, length, etc.)

III. Basic Commands (This example uses the TitanicSurvival.R file)

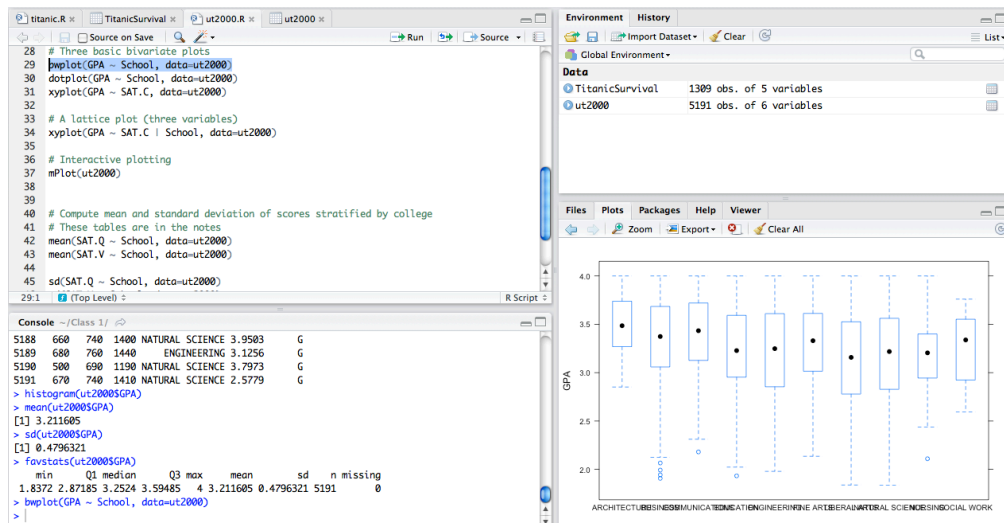
- a. `>names(TitanicSurvival)` returns variable names. This reminds you of the spelling and capitalization of each variable within the dataset
- b. `>head(TitanicSurvival)` returns the first 6 lines of the data set
- c. `>tail(TitanicSurvival)` returns the last 6 lines of the data set
- d. `>xtabs(~survived + sex, data=TitanicSurvival)` gives you a cross tabulation of variables “survived” and “sex” from the data retrieved by TitanicSurvival.csv file
 - i. “~” or the tilde symbol means stratify by or model by

- e. `>tally(~survived + sex, data(TitanicSurvival))` gives you the row and column totals for each variable
- f. `>tally(~survived + sex:passengerClass, data=TitanicSurvival)` is used to stratify by two categories
 - i. “:” is used for interaction or cross between factors

IV. Simple Summaries and Graphics (This example uses the ut2000.R file)

- a. Good graphs
 - i. Facilitate comparison
 - ii. Are multivariate
 - iii. Are truthful about magnitude
 - iv. Usually not for small data
- b. `>summary(ut2000)` returns summary statistics. Includes minimum, 1st quartile, median, 3rd quartile, maximum, and mean for quantitative data. For categorical data, this command will return counts for each variable.
- c. `>histogram(ut2000$GPA)` returns density histogram with x axis bins using the GPA variable and y axis being the density
 - i. Histograms help you understand the variability within a data set
- d. `>mean(ut2000$GPA)` returns the mean GPA from the ut2000 data set
- e. `>sd(ut2000$GPA)` returns the standard deviation for GPA from the ut2000 data set
- f. `>favstats(ut2000$GPA)` returns the minimum, quartile 1, quartile 3, maximum, standard deviation, and n missing for the GPA variable from the ut2000 data set

V. Graphs



- a. **Box plots:** bivariate graph that can compare within-group variability and between-group variability
 - i. `>bwplot(GPA ~ school, data=ut2000)` returns a plot for each school with the GPA on the Y axis
- b. **Dot plots:** bivariate graph with each data point representing a dot. Dot plots are better for smaller data sets so you can distinguish one dot from another
 - i. `>dotplot(GPA ~ school, data=ut2000)` returns a dot plot for each school with GPA on the Y axis

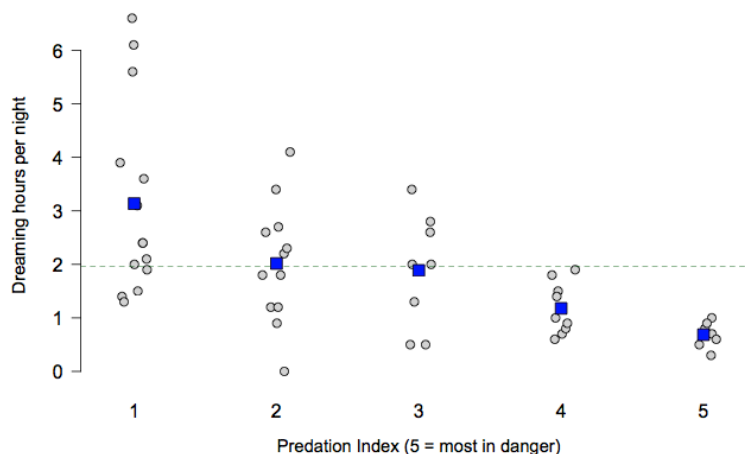
- c. **Scatter plot:** bivariate graph that can accommodate large sets of data
 - i. `>xyplot(GPA ~ SAT.C, data=ut2000)` returns a scatter plot with SAT.C on the x axis and GPA on the Y axis
- d. **Lattice plot:** used for 3 pieces of information
 - i. `>xyplot(GPA ~ SAT.C | school, data=ut2000)` returns scatter plots stratified by college
- e. **Interactive plotting:**
 - i. `mPlot(ut2000)`
 - ii. Another package needed

VI. Means & Standard Deviations

- a. `>mean(SAT.Q ~ school, data=ut2000)` computes the mean SAT.Q stratified by school
- b. `>sd(SAT.Q ~ school, data=ut2000)` computes the standard deviation of SAT.Q stratified by school
- c. `Satqmeans=mean(SAT.Q ~ school, data=ut2000)` stores the results of computations in other variables.
 - i. In this case, “satqmeans” will return the computation of mean SAT.Q stratified by school. When you enter this command, nothing new will appear in the console box. However, if you type `>satqmeans` the computation will appear
 - ii. Useful in intermediate computation to be used in subsequent computation

VII. Fitted Values and Residuals

- a. Actual value = fitted value (predicted by the model) + deviation of that case (from the prediction)
- b. Observed value = model value + residual
- c. $Y_i = Y_i(\text{hat}) + e_i$
 - i. “Hat” is this symbol: $\hat{\cdot}$. It is the generic notation for “predicted”



- d. In the graph above, the blue dot is the group mean (fitted value) and the deviation is the distance between the gray data point and the blue dot