RSG III: Data Visualization

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Review of Data structures

I. Vectors

- Homogeneous

```
• 1-dimensional
EX
#a vector of numbers
vec_1<-c(1, 19, 34, 76)
vec_1
## [1] 1 19 34 76
#a vector of characters
vec_2<-c('cat', 'dog', 'rat', 'pet')</pre>
vec_2
## [1] "cat" "dog" "rat" "pet"
 II. Lists
  • Heterogeneous
  • 1 - n-dimensional
EX
#a list o numbers
l_1<-list(1, 19, 34, 76)
1_1
## [[1]]
## [1] 1
##
## [[2]]
## [1] 19
##
## [[3]]
## [1] 34
## [[4]]
## [1] 76
#a list of characters
1_2<-list('cat', 'dog', 'rat', 'pet')</pre>
1_2
## [[1]]
## [1] "cat"
##
## [[2]]
## [1] "dog"
```

```
##
## [[3]]
## [1] "rat"
##
## [[4]]
## [1] "pet"
#a list of lists
l_3<-list(l_1, l_2)
1_3
## [[1]]
## [[1]][[1]]
## [1] 1
##
## [[1]][[2]]
## [1] 19
## [[1]][[3]]
## [1] 34
##
## [[1]][[4]]
## [1] 76
##
##
## [[2]]
## [[2]][[1]]
## [1] "cat"
##
## [[2]][[2]]
## [1] "dog"
##
## [[2]][[3]]
## [1] "rat"
## [[2]][[4]]
## [1] "pet"
#lsit of vectors
 III. Dataframes
```

- List of vectors
- 2-dimensional (matrix-like)
- Homogeneous or heterogeneous?

EX

```
#The iris dataframe
head(iris)
```

```
##
     Sepal.Length Sepal.Width Petal.Length Petal.Width Species
## 1
              5.1
                          3.5
                                        1.4
                                                    0.2 setosa
                                                    0.2 setosa
## 2
              4.9
                          3.0
                                        1.4
## 3
              4.7
                          3.2
                                        1.3
                                                    0.2 setosa
## 4
              4.6
                          3.1
                                        1.5
                                                    0.2 setosa
## 5
              5.0
                          3.6
                                        1.4
                                                    0.2 setosa
```

```
## 6
             5.4
                         3.9
                                      1.7
                                                  0.4 setosa
str(iris)
## 'data.frame':
                   150 obs. of 5 variables:
## $ Sepal.Length: num 5.1 4.9 4.7 4.6 5 5.4 4.6 5 4.4 4.9 ...
## $ Sepal.Width : num 3.5 3 3.2 3.1 3.6 3.9 3.4 3.4 2.9 3.1 ...
## $ Petal.Length: num 1.4 1.4 1.3 1.5 1.4 1.7 1.4 1.5 1.4 1.5 ...
## $ Petal.Width : num 0.2 0.2 0.2 0.2 0.4 0.3 0.2 0.2 0.1 ...
                  : Factor w/ 3 levels "setosa", "versicolor", ...: 1 1 1 1 1 1 1 1 1 1 1 ...
## $ Species
head(iris[,3])
## [1] 1.4 1.4 1.3 1.5 1.4 1.7
head(iris$Petal.Length)
## [1] 1.4 1.4 1.3 1.5 1.4 1.7
#The cars dataframe
head(cars)
     speed dist
##
## 1
        4
## 2
         4
            10
## 3
        7
             4
        7
## 4
            22
## 5
        8
            16
        9
## 6
            10
str(cars)
## 'data.frame':
                   50 obs. of 2 variables:
## $ speed: num 4 4 7 7 8 9 10 10 10 11 ...
## $ dist : num 2 10 4 22 16 10 18 26 34 17 ...
head(cars[,2])
## [1] 2 10 4 22 16 10
head(cars$dist)
## [1] 2 10 4 22 16 10
```

Dataframes make plotting easy!

Data Visualization

R has several basic functions for plotting data:

- hist()
- plot()
- boxplot()

These functions are built in – they come with R when you download it.

Various other plotting packages exist.

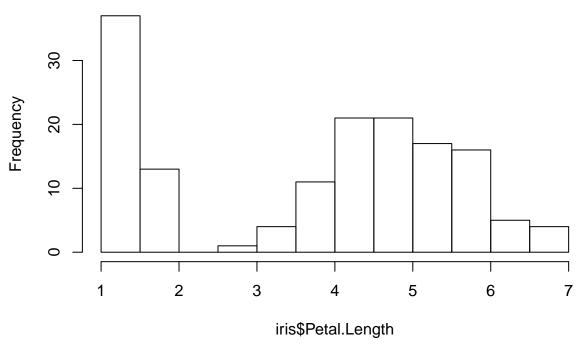
By far, the most popular is ggplot2. We'll get to this today if there's time.

Histograms

- 1) Look up how to use the hist() function using ?hist()
- 2) Use hist() to examine the frequency distribution of petal lengths in thie iris dataset.

hist(iris\$Petal.Length)

Histogram of iris\$Petal.Length



- 3) Subset the iris dataset to plot each species separately
- 4) Partition the plot window to view each of these plots

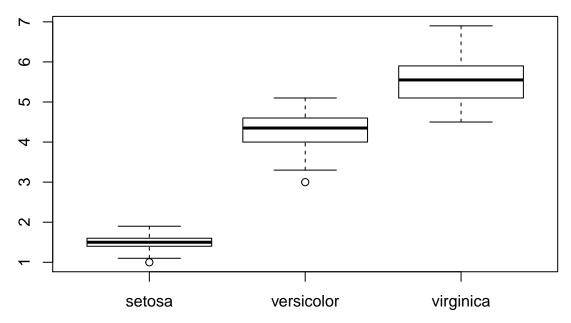
It looks like different species have different distributions of petal length... How to best visualize this?

Boxplots

Boxplots allow for comparison of data across different levels of a factor. Look at the help file for boxplots.

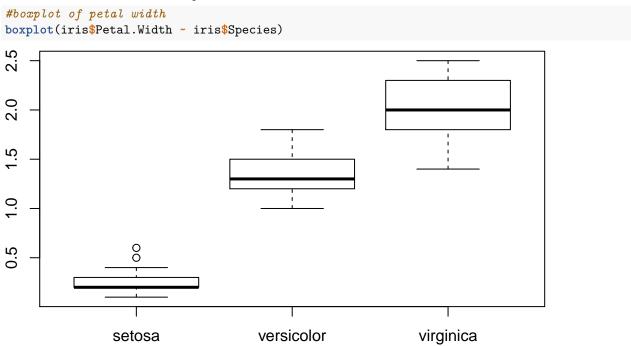
Using this info, construct a boxplot showing petal length vs. species

```
#boxplot of petal length vs. species
boxplot(iris$Petal.Length ~ iris$Species)
```



1st, 2nd and 3rd quartiles (25, 50, 75 quantiles)

Let's look at another variable – petal width



The pattern across species seems similar, are these variables correlated in some way?

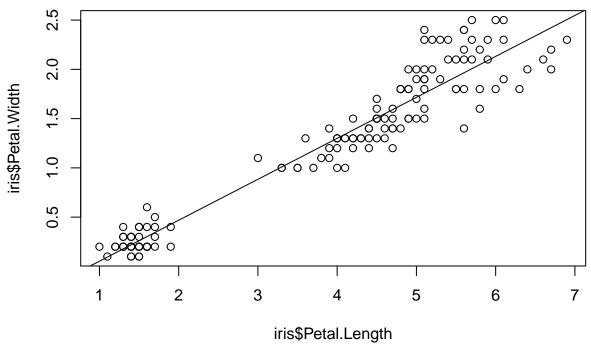
Scatterplots

The simplest scatterplots can be constructed using ${\tt plot}(\tt)$.

Examine the plot() help file to see how it's used. What do you notice about the way arguments can be

supplied?

```
#scatter plot of petal width vs petal length
plot(iris$Petal.Length, iris$Petal.Width)
#add a regression line for fun
abline(lm(iris$Petal.Width ~ iris$Petal.Length))
```



Brief stats foray.

install ggplot2.