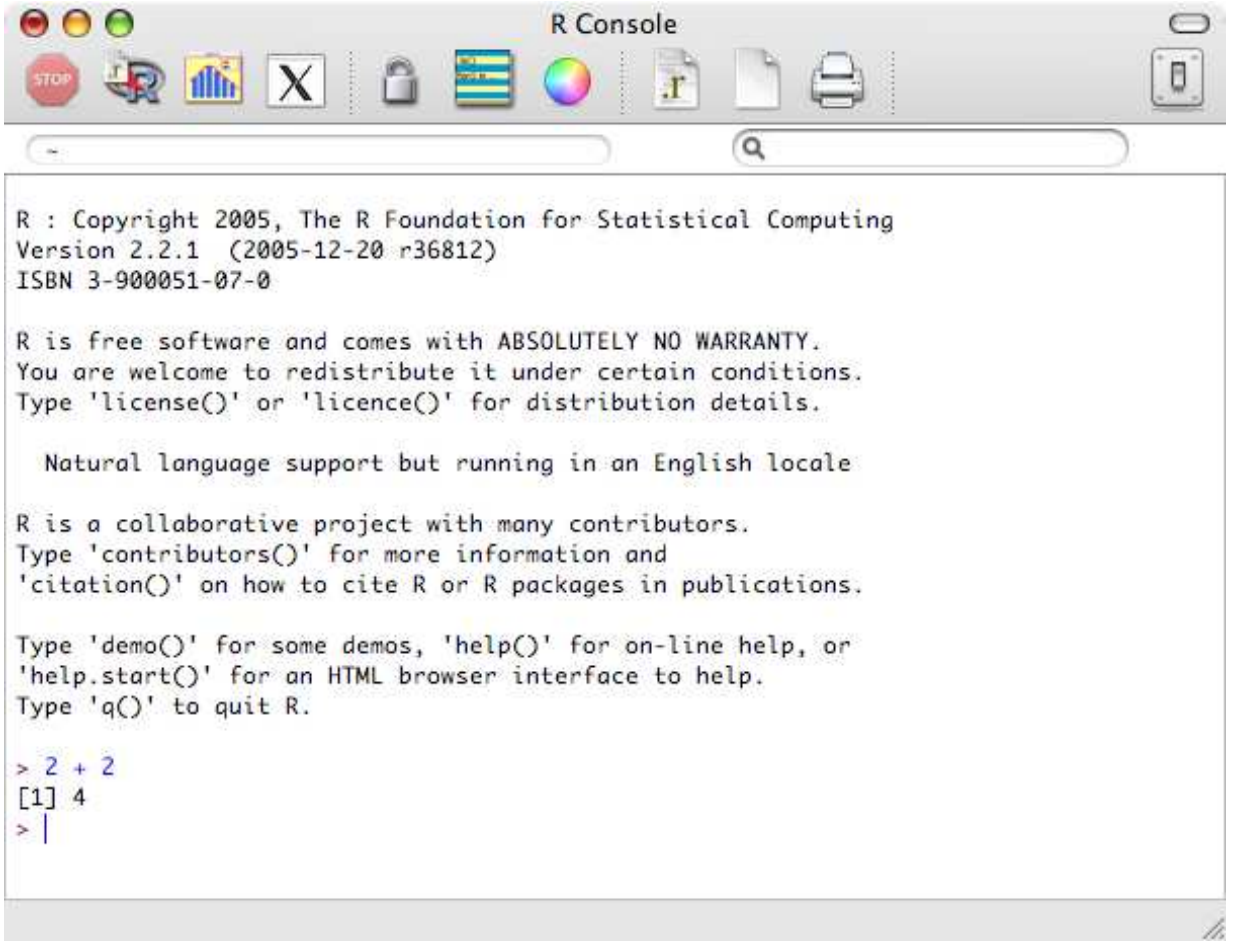
Statistical Computing in R

R is a programming language designed to support data analysis and model building.

- All traditional programming constructs such as expressions, assignments, conditionals, loops, and functions are present.
- A straight-forward object system that supports high-level constructs such as statistical models with all their parameters etc. very nicely.
- Vector arithmetic (very powerful and the preferred way of accomplishing things in R).
- Graphics engine supporting graphical techniques (automatic scatter plots, histograms, etc.)
- Many, many extension modules implementing everything from basic statistics to micro array analysis...and in particular support vector machines.

Interactive R Session



R Programming

```
> x <- 2
> 2 * x
[1] 4
```

```
> v <- c(1,2,3)
> v
[1] 1 2 3
```

```
> v + 1
[1] 2 3 4
```

```
> w <- v + 1
> q <- w + v
> q
[1] 3 5 7
```

```
> add1 <- function(x) { x + 1; }
> add1
function(x) { x + 1; }
> add1(1)
[1] 2
```

R Programming

```
> addv1
function(v)
   {
      y <- c()
      for (x in v) {
         x1 <- x + 1
         y <- c(y,x1)
      }
      y
   }
> w
[1] 2 3 4
> addv1(w)
[1] 3 4 5
```

This function performs the same operation as the vector operation `w + 1`. From a performance point of view it is always desirable to use the vector operations, explicit iteration over vector elements is SLOW!

R Data

R has many different ways to represent data:

- vectors
- lists
- arrays/matrices

The most important one (for our purposes) is the *data frame*. A data frame is a two-dimensional data matrix with additional structure.

```
> df <- data.frame(v,w)
> df
  v w
1 1 2
2 2 3
3 3 4
> df$v
[1] 1 2 3
> df$w
[1] 2 3 4
```

Loading Data Frames

We can read comma-separated-value (CSV) files directly into an R data frame.

Here is our mammal training data set represented as a CSV file:

```
Legs, Wings, Fur, Feathers, Mammal
4, no, yes, no, true
2, yes, no, yes, false
4, no, no, no, false
4, yes, yes, no, true
3, no, no, no, false
```

Assume that we saved this into a file called “mammals.csv”, in a directory called “datasets”.

Loading Data Frames

```
> setwd("datasets")
> mammals.df <- read.csv("mammals.csv")
> mammals.df
  Legs Wings  Fur Feathers Mammal
1    4    no  yes       no   true
2    2   yes   no      yes  false
3    4    no   no       no  false
4    4   yes  yes       no   true
5    3    no   no       no  false
> summary(mammals.df)
      Legs      Wings     Fur    Feathers    Mammal
 Min.   :2.0    no :3    no :3    no :4    false:3
 1st Qu.:3.0    yes:2    yes:2    yes:1    true :2
 Median :4.0
 Mean   :3.4
 3rd Qu.:4.0
 Max.   :4.0
```

R Built-in Data Frames

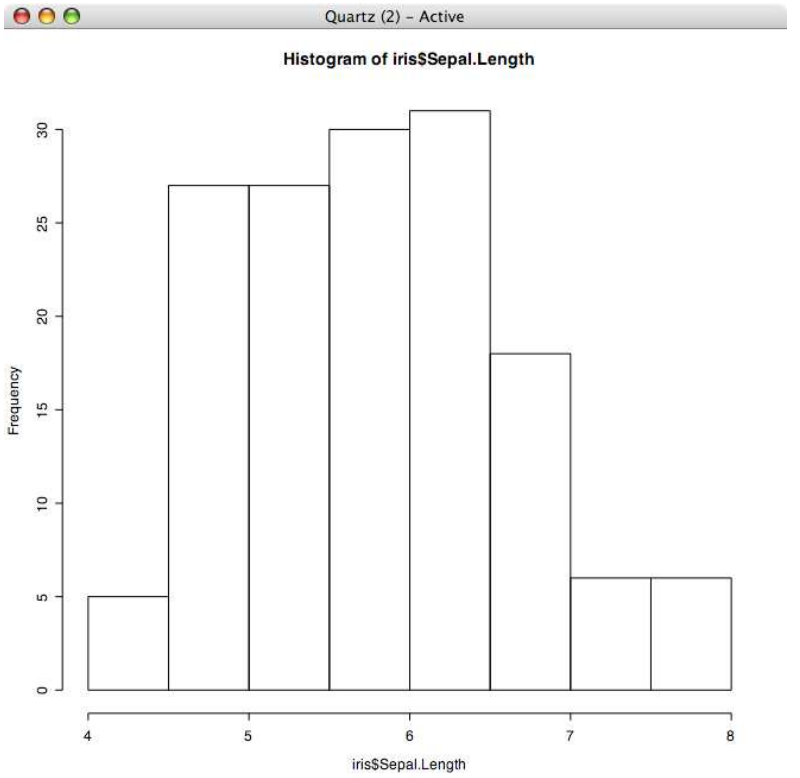
For convenience sake, R comes with a number of predefined data frames. One such predefined data frame is the *iris data set*.

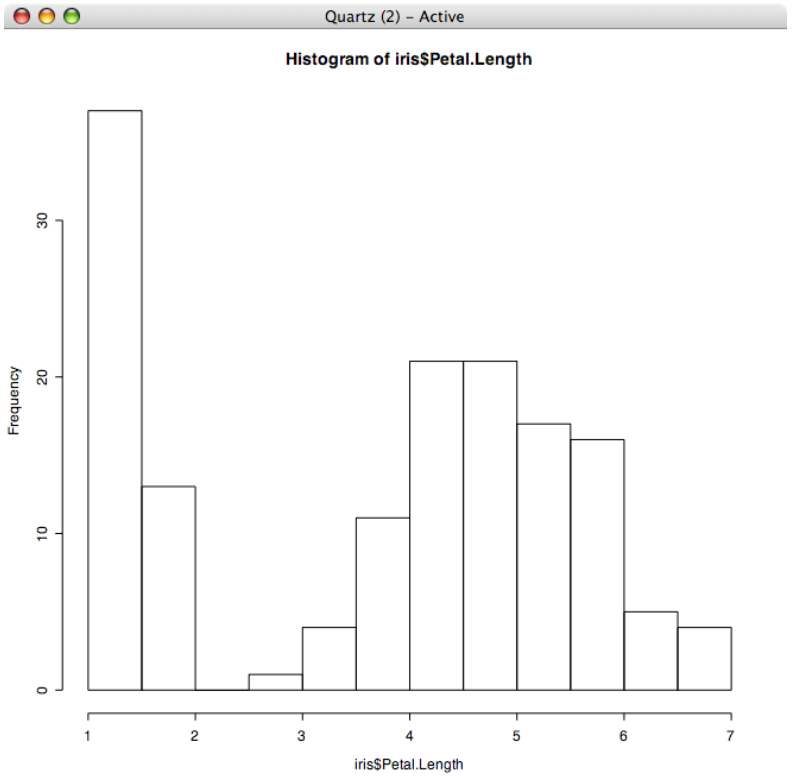
```
> data(iris)
> summary(iris)
  Sepal.Length    Sepal.Width     Petal.Length    Petal.Width          Species
 Min.   :4.300   Min.   :2.000   Min.   :1.000   Min.   :0.100   setosa    :50
 1st Qu.:5.100   1st Qu.:2.800   1st Qu.:1.600   1st Qu.:0.300   versicolor:50
 Median :5.800   Median :3.000   Median :4.350   Median :1.300   virginica :50
 Mean   :5.843   Mean   :3.057   Mean   :3.758   Mean   :1.199
 3rd Qu.:6.400   3rd Qu.:3.300   3rd Qu.:5.100   3rd Qu.:1.800
 Max.   :7.900   Max.   :4.400   Max.   :6.900   Max.   :2.500
```

We might wish to inspect the data distributions visually as well:

```
> hist(iris$Sepal.Length)
> hist(iris$Petal.Length)
```

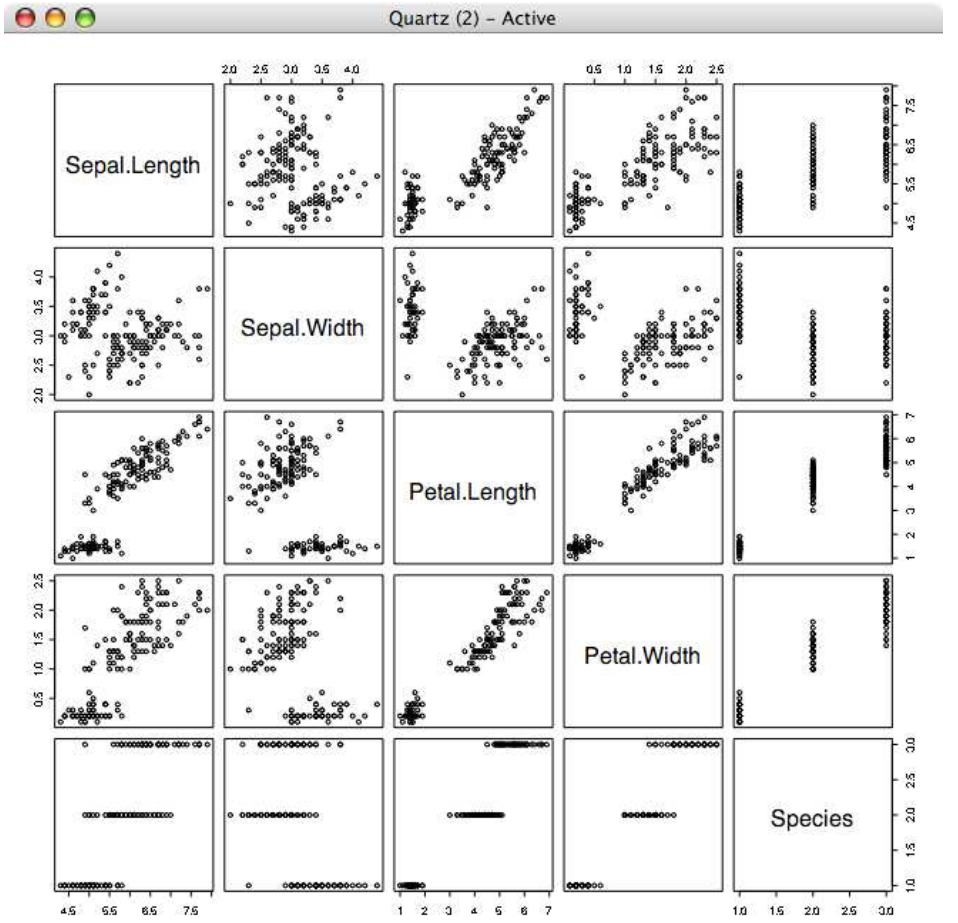
R Built-in Data Frames





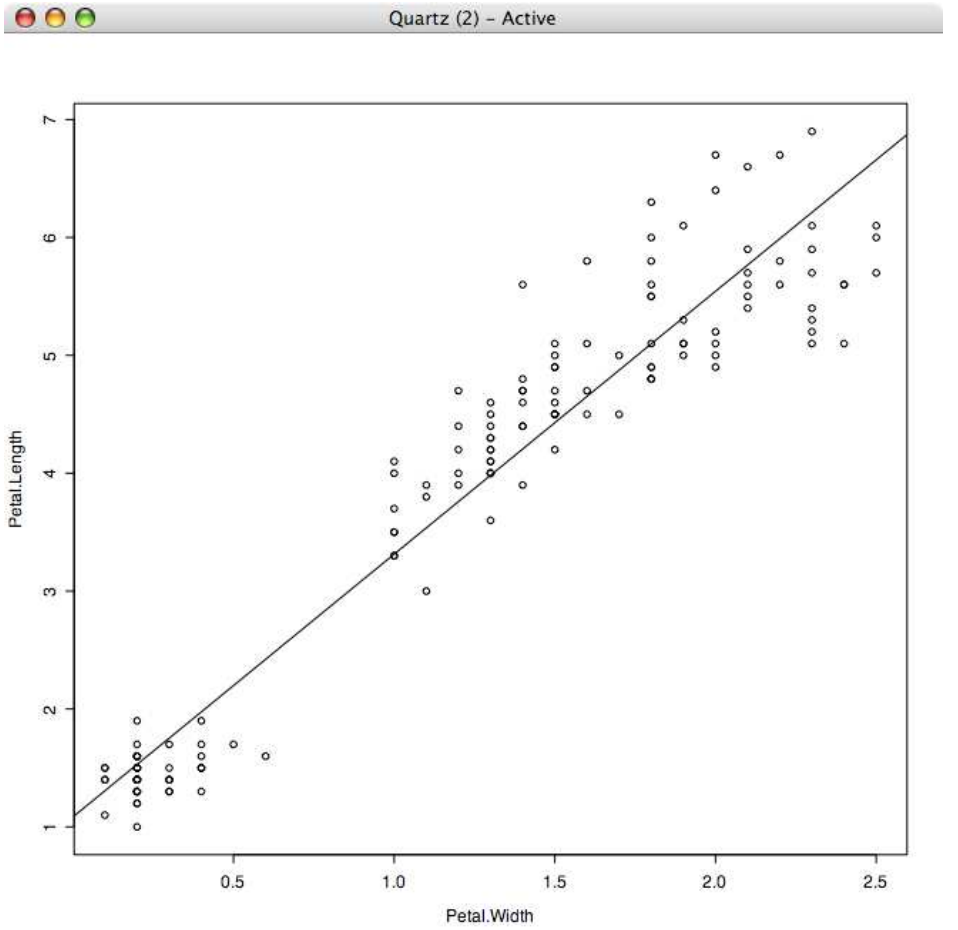
R Built-in Data Frames

```
> plot(iris)
```



Simple Model Building

```
> attach(iris)
> model <- lm(Petal.Length~Petal.Width)
> plot(Petal.Width,Petal.Length)
> abline(model)
```



Homework

Read Chapter 1 and Appendix B

Do Assignment 1, see website.