# 1 Reading in a CSV file

To read in a csv file, it is convenient to save it to your *working directory*, the default file directory that R uses. Once it is in the working directory, you can use the read.csv() command to load it into the R environment.

```
getwd()
#[1] "C:/Users/Computer5/Documents"

HWO <- read.csv("hw0_data.csv",header = TRUE)</pre>
```

You can change the working directory using the setwd() to the one you require.

```
getwd()
#[1] "C:/Users/Computer5/Documents"

setwd("C:/Users/Computer5/Documents/R")

# > getwd()
# [1] "C:/Users/Computer5/Documents/R"
```

## 2 Inspecting the data set

- 2a) Dimensions
- 2b) Column names (i.e. variables names)
- 2c) structure of data frame

Lets compute the dimensions of the data frame *iris*, and also the length of the *Nile* data set.

```
dim(iris)
nrow(iris)
ncol(iris)
length(Nile)
```

Data frames often have specifically named rows and columns. Lets find out the names of the variables (columns) and cases (rows) for the data sets *iris* and *mtcars*.

```
names(iris)
colnames(iris)
rownames(iris) # simply the case numbers.

names(mtcars)
rownames(mtcars)
colnames(mtcars)
```

### 2.1 The summary() command

The summary() command can be used to extract a short statistical summary (if applicable) from each column of the data frame. If there are missing values, the frequency of missing values will also be listed for each column.

```
> summary(iris)
  Sepal.Length
                                                     Petal.Width
                   Sepal.Width
                                    Petal.Length
                                                                           Species
        :4.300
 Min.
                  Min.
                         :2.000
                                          :1.000
                                                           :0.100
                                   Min.
                                                    Min.
                                                                     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
        :7.900
 Max.
                  Max.
                         :4.400
                                   Max.
                                          :6.900
                                                    Max.
                                                           :2.500
```

## 2.2 Types of Data in Dataframes

What is the data type of each column in the iris data set? To find out, we use the str() command.

```
str(iris)
```

The output of this command is given below. There are four numeric variables, and one factor (i.e. categorical) variable.

```
'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 0.2 0.4 0.3 0.2 0.2 0.1 ...
$ Species : Factor w/ 3 levels "setosa", "versicolor", ..: 1 1 1 1 1 1 1 1 1 1 ...
```

## 3 Reduction and Subsetting

- 3a) The head() and tail() function
- 3b) Accessing a particular row or set of rows
- 3c) subsetting with a relational condition.

#### 3.1 The head() and tail() function

The head and tail functions can be used to access the first six and last six rows of a data frame. If a different number of rows is required, all you have to do is specify that number as an additional argument.

```
head(iris) #First 6 rows
head(iris,2) #First 2 rows

tail(iris) #Last 6 rows
tail(iris,4) #Last 4 rows
```

#### 3.2 Accessing a particular row or set of rows

- Each value in a dataframe can be accessed directly by specifying the row and column i.e. df [r,c].
- To access a particular row, simply specify the row number, while leaving the column number blank i.e. df[r,]
- To access a particular column, simple specify the column number, while leaving the row number blank i.e. df[,c]

```
iris[10,2]
iris[10,]
Formaldehyde[,2]
```

```
> iris[10,2]
[1] 3.1
> iris[10,]
    Sepal.Length Sepal.Width Petal.Length Petal.Width Species
10      4.9      3.1      1.5      0.1 setosa
```

```
> Formaldehyde[,2]
[1] 0.086 0.269 0.446 0.538 0.626 0.782
```

## 4 Missing data

- (4a) Determining the number of missing data items
- (4b) Performing statistical operations removing missing data

As stated previously, the summary() command can be used to determine the number of missing data items in a data frame. The additional argument na.rm=TRUE can also be used with certain functions (see the help file)

```
> X <- c(4,6,3,12,NA,8)
> mean(X)
[1] NA
> summary(X)
   Min. 1st Qu.
                  Median
                             Mean 3rd Qu.
                                                       NA's
                                              Max.
    3.0
             4.0
                     6.0
                              6.6
                                       8.0
                                              12.0
                                                          1
> mean(X ,na.rm = TRUE)
[1] 6.6
```

## 5 Subsetting Data

#### Logical and Relational Operator

- AND The logical operator is &
- OR The logical operator is ||

### Selection using the subset() Function

The subset() function is the easiest way to select variables and observation.

#### Example 1

In the following example we will use the *iris* data set, we select all rows that have a value of sepal length of 6 or more, and determine how many observations there are, and then compute the mean of the petal lengths. (The answer is 5.263, from the summary output).

```
#call the subset iris.2
iris.2 = subset(iris,iris$Sepal.Length >= 6)
dim(iris.2)
summary(iris.2)
```

#### Example 2

There are three types of iris - setosa, versicoloir and virginica. Suppose we wish to compute the median of petal widths for the setosa irises only.

The equality operator is ==.

- > iris.setosa =subset(iris,iris\$Species=="setosa")
- > summary(iris.setosa)

```
Sepal.Length
                                  Petal.Length
                  Sepal.Width
                                                   Petal.Width
       :4.300
Min.
                Min.
                        :2.300
                                 Min.
                                         :1.000
                                                  Min.
                                                          :0.100
1st Qu.:4.800
                 1st Qu.:3.200
                                 1st Qu.:1.400
                                                  1st Qu.:0.200
Median :5.000
                Median :3.400
                                 Median :1.500
                                                  Median :0.200
Mean
       :5.006
                Mean
                        :3.428
                                 Mean
                                         :1.462
                                                  Mean
                                                          :0.246
                 3rd Qu.:3.675
3rd Qu.:5.200
                                 3rd Qu.:1.575
                                                  3rd Qu.:0.300
                        :4.400
                                         :1.900
Max.
       :5.800
                Max.
                                 Max.
                                                  Max.
                                                          :0.600
      Species
```

setosa :50 versicolor: 0 virginica : 0

## Example 3

In the following example we will use the *iris* data set, we select all rows that have a value of sepal length of 6 or more, but have sepal width is at least 2.5.

```
> iris.3 = subset(iris,(iris$Sepal.Length >= 6)&(iris$Sepal.Width >=2.5))
> summary(iris.3)
```

Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
Min. :6.000	Min. :2.500	Min. :4.000	Min. :1.200
1st Qu.:6.300	1st Qu.:2.800	1st Qu.:4.750	1st Qu.:1.500
Median :6.500	Median :3.000	Median :5.300	Median :1.800
Mean :6.641	Mean :3.013	Mean :5.313	Mean :1.848
3rd Qu.:6.900	3rd Qu.:3.200	3rd Qu.:5.750	3rd Qu.:2.150
Max. :7.900	Max. :3.800	Max. :6.900	Max. :2.500

Species

setosa : 0
versicolor:21
virginica :42