# Additional Material for Merging and Importing Data

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### Built-in functions for exploring a data frame

We will use built-in dataset iris to explore some of the useful functions in base package of R language. In order to know the dimensions of iris, we use dim function. The output of dim function is a vector, in which the elements represent the number of rows and number of columns, respectively.

```
dim(iris)
## [1] 150 5
```

We can also use nrow and ncol to get the number of rows and number of columns, respectively.

```
nrow(iris)
## [1] 150
ncol(iris)
```

## [1] 5

Thus, iris has 150 rows and 2 columns, which can also be verified by using str function. It also returns many useful pieces of information, including the above information and the types of data for each column.

```
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 0.2 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 ...
```

The first row in the output indicates that this dataset is a data frame with 150 observations of 5 variables. Also, num denotes that the variables Sepal.Length, Sepal.Width, Petal.Length and Petal.Width are numeric. Factor denotes that the variable Species is categorical with 3 levels (setosa, versicolor, virginica).

To know the range of values inside iris, we use summary function. In particular, this function provides a number of useful statistics including range, median and mean (Andrew Shaughnessy 2018).

#### summary(iris)

```
##
     Sepal.Length
                      Sepal.Width
                                       Petal.Length
                                                        Petal.Width
                                              :1.000
##
           :4.300
                             :2.000
                                                       Min.
                                                               :0.100
##
    1st Qu.:5.100
                     1st Qu.:2.800
                                      1st Qu.:1.600
                                                       1st Qu.:0.300
##
    Median :5.800
                     Median :3.000
                                      Median :4.350
                                                       Median :1.300
##
    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
                             :4.400
                                              :6.900
                                                               :2.500
                     Max.
                                      Max.
                                                       Max.
##
          Species
##
   setosa
               :50
    versicolor:50
    virginica:50
```

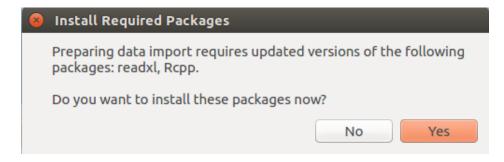


Figure 1: Installing readxl and Rcpp

## ## ##

We use head to obtain the first n observations and tail to obtain the last n observations; by default, n = 6. These are good commands for obtaining an intuitive idea of what the data look like without revealing the entire dataset, which could have millions of rows and thousands of columns (Cai 2013).

```
head(iris, 2)
##
     Sepal.Length Sepal.Width Petal.Length Petal.Width Species
## 1
               5.1
                           3.5
                                                      0.2
## 2
               4.9
                           3.0
                                         1.4
                                                      0.2
                                                           setosa
tail(iris, 2)
##
       Sepal.Length Sepal.Width Petal.Length Petal.Width
## 149
                 6.2
                              3.4
                                           5.4
                                                        2.3 virginica
## 150
                 5.9
                              3.0
                                           5.1
                                                        1.8 virginica
```

## Dependencies for reading datasets in R

In order to read XML files in R, we need to install XML package. However, the Ubuntu package libxml2-dev needs to be installed beforehand (Overflow 2013). On Linux operating system, open the terminal and type the following commands.

```
sudo apt-get update
sudo apt-get install libxml2-dev
```

Similarly, while importing Excel data in R, we need to install readx1 and Rcpp. If these packages are not installed and you try importing Excel data, a pop-up message as shown in Figure 1 will be generated. By clicking Yes to this message, these packages can be installed.

### References

Andrew Shaughnessy, Elizabeth Hasenmueller, Christopher Prener. 2018. "Exploring Data in R." https://cran.r-project.org/web/packages/driftR/vignettes/ExploringData.html.

Cai, Eric. 2013. "Exploratory Data Analysis: Useful R Functions for Exploring a Data Frame." https://chemicalstatistician.wordpress.com/2013/08/19/exploratory-data-analysis-useful-r-functions-for-exploring-a-data-frame/.

Overflow, Stack. 2013. "Unable to install R package in Ubuntu 11.04." https://stackoverflow.com/questions/

7765429/unable-to-install-r-package-in-ubuntu-11-04.