Additional Material for Merging and Importing Data

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
##
    Min.
            :4.300
                     Min.
                             :2.000
                                      Min.
                                              :1.000
                                                        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
##
            :7.900
                             :4.400
                                              :6.900
                                                                :2.500
    Max.
                     Max.
                                      Max.
                                                        Max.
##
          Species
##
               :50
    setosa
##
    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)

## ## ##	S 1 2	epal.Length 5.1 4.9	Sepal.Width F 3.5 3.0	etal.Length 1.4 1.4	Petal.Width S 0.2 0.2	pecies setosa setosa
tail(iris, 2)						
## ## ##	149 150	Sepal.Lengt 6. 5.	th Sepal.Width 2 3.4 .9 3.0	Petal.Lengt 5. 5.	th Petal.Width 4 2.3 1 1.8	Species virginica 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.

In case you are using Windows OS, you don't need to install these packages.

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.