



SNS COLLEGE OF TECHNOLOGY

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UNIT II

Importing and Exporting Data in R

First, make sure that you are working in the right directory- the directory that you want to be. To see the current working directory, type `getwd()` into the console

```
getwd()
```

```
## [1] "/Users/Ercan/Desktop/R Books/RInto_Ercan"
```

You can change the working directory by typing `setwd()`:

```
setwd("/Users/Ercan/Desktop/Beginning R")
```

```
getwd()
```

Alternatively, just select `Session > Change Working Directory` in Rstudio and navigate to the directory you want to make your new working directory.

Finally, type in `dir()` to see all the documents in the current directory.

The definitive guide for importing data in R is the R Data Import/Export manual available at [R Data Import/Export manual](https://cran.r-project.org/doc/manuals/R-data.pdf) (<https://cran.r-project.org/doc/manuals/R-data.pdf>)

Importing .csv, .txt, .delim files

Read .csv format (comma separated values)

To read data from a CSV file use either `read.table` or `read.csv` functions, the latter being a wrapper around the former. Suppose that you have a data set `health.csv` in *your working directory* and you want to introduce this to R:

```
# use read.table
```

```
health <- read.table(file = "health.csv", header = TRUE, sep = ",")
```

```
# or use read.csv
```

```
health <- read.csv("health.csv")
```

```
# the result is a data.frame
```

```
class(health)
```

```
## [1] "data.frame"  
roster <- read.csv("roster.csv")
```

To learn about the content of a data frame:

```
str(roster)  
## 'data.frame': 13 obs. of 6 variables:  
## $ Jersey : int 0 1 3 5 10 12 15 20 21 33 ...  
## $ Name : Factor w/ 13 levels "Ajukwa, Austin",...: 11 1 8 2 3 6 5 12 7 13 ...  
## $ Position: Factor w/ 3 levels "C","F","G": 3 3 3 2 3 3 2 3 3 2 ...  
## $ Inches : int 74 78 74 79 75 73 80 72 76 80 ...  
## $ Pounds : int 190 205 205 215 200 205 205 165 205 245 ...  
## $ Class : Factor w/ 4 levels "freshman","junior",...: 1 4 2 4 1 3 1 2 3 2 ...
```

To view your data without editing them, you can use the View command:

```
View(roster)
```

Read .txt format

Suppose that you have a data set `health.txt` in *your working directory* and you want read this into R:

```
read.table("health.txt", sep = " ")
```

Importing .xls, .xlsx, .sav, .sas Files

- Use the package `readxl` to read in `.xls` and `.xlsx` files.

```
install.packages("readxl")  
library(readxl)  
read_excel("health.xlsx")
```

- Use the package `haven` to read in **SPSS**, **Stata** and **SAS** files

```
install.packages("haven")  
library(haven)  
read_spss("Dataset.sav")  
read_dta("Dataset.dta")  
read_sas("Dataset.sas7bdat")
```

Importing Data from the Web

Suppose there is some data at `'http://www.something.com/data.csv'` We can import this as

```
getLink <- "http://www.something.com/data.csv"

myData <- read.table(file = getLink, header = TRUE, sep = ",")
```

Unlike `read.table`, `read_excel` cannot read data directly from the Internet, and thus the files must be downloaded first. We could do this by visiting a browser or we can stay within R and use `download.file`.

```
# download the data.
# Note that this will download to the current wd, but you can change it by specifying a
path to "destfile"
download.file(url='http://www.something.com/data.xlsx',
             destfile='excelData.xlsx', method='curl')
# read data
excelData <- read_excel("excelData.xlsx")
```

R Binary Files

When working with other R programmers, a good way to pass around data-or any R objects such as variables and functions—is to use RData files. These are binary files that represent R objects of any kind. They can store a single object or multiple objects and can be passed among Windows, Mac and Linux without a problem.

First, let's create an RData file, remove the object that created it and then read it back into R:

```
health <- read.table(file = "health.csv", header = TRUE, sep = ",")
# save the health.frame to disk
save(health, file="health.rdata")
# remove health from memory
rm(health)
# read it from the rdata file
load("health.rdata")
# check if it exists now
head(health)
## id gender state age health1 health2 health3 health4 health5 health6
## 1 1 M 1 51 1 4 2 1 4 5
```

##	2	2	F	3	35	2	3	3	2	3	4
##	3	3	F	1	29	5	2	4	2	1	3
##	4	4	M	1	21	5	1	5	4	2	1
##	5	5	M	2	56	2	4	2	4	3	3
##	6	6	M	3	72	1	5	4	2	4	5