Introduction to R programming for data science – day 6

Dr. Francesca Finotello Medical University of Innsbruck, Austria

Read and write text files

Reading tables with *read.table*

Plain text files in table format can be loaded into R as data.frames using the function read.table

Once you load a table into R, you can check how it looks like using head and tail, and check its dimensions with dim

3/11

Reading tables with *read.delim* and *read.csv*

Depending on the column separator in the input file, read.delim and read.csv can be also used to import files in table format

```
read.delim(file,
    header = TRUE,
    sep = "\t", ...)

read.csv(file,
    header = TRUE,
    sep = ",", ...)
```

The are both based on the read.table function, but they use different paremeter settings (e.g. "sep")

Save data in plain text files

Data.frames and matrices can be saved into text files

```
( DF <- data.frame(name=c("Mary", "John", "Lisa"),
    age=c(19, 30, 20),
    city=c("New York", "Seattle", "New York")) )

## name age    city
## 1 Mary 19 New York
## 2 John 30 Seattle
## 3 Lisa 20 New York

write.table(DF,
    quote = FALSE,
    sep = "\t",
    row.names = FALSE,
    col.names = TRUE,
    file = "../Data/Day3_Friends_table.txt")</pre>
```

5/11

Read and write Excel files

The xlsx package

The xlsx package provides R functions to handle Excel files (97/2000/XP/2003/2007 formats)

It is available on CRAN and can be installed from the "Packages" window or by executing

```
install.packages("xlsx")
```

Once installed, it can be loaded with library, which is the function to load (installed) R packages into R

```
library("xlsx"")
```

7/11

Read Excel files

The read.xlsx function from the xlsx package can be used to read Excel files

```
read.xlsx(file, # File to be read
sheetIndex, # Number of the sheet to be read
sheetName = NULL, # Character indicating the sheet name
startRow = NULL, # First row to be read
endRow = NULL, # Last row to be read
header = TRUE, # Does the first row contain column names?
keepFormulas = FALSE, # Display formulae (instead of results)?
...)
```

Write Excel files

The write.xlsx function from the xlsx package can be used to write Excel files

```
write.xlsx(x,  # Table to be written
file,  # Path to the output file
sheetName = "Sheet1", # Character indicating the sheet name
col.names = TRUE, # Write column names?
row.names = TRUE, # Write row names?
...)
```

9/11

Excel and gene names

Gene name errors are widespread in the scientific literature

```
Mark Ziemann, Yotam Eren and Assam El-Osta 

Genome Biology 2016 17:177

<a href="https://doi.org/10.1186/s13059-016-1044-7">https://doi.org/10.1186/s13059-016-1044-7</a> © The Author(s). 2016

Published: 23 August 2016
```

Abstract

The spreadsheet software Microsoft Excel, when used with default settings, is known to convert gene names to dates and floating-point numbers. A programmatic scan of leading genomics journals reveals that approximately one-fifth of papers with supplementary Excel gene lists contain erroneous gene name conversions.

The *readr* package

The <u>readr</u> package from <u>tidyverse</u> provides functions to read files with different formats:

- read_csv: comma separated (CSV) files
- read_tsv: tab separated files
- · read_delim: general delimited files
- read_fwf: fixed width files
- read_table: tabular files where columns are separated by white-space
- read_log: web log files



11/11