

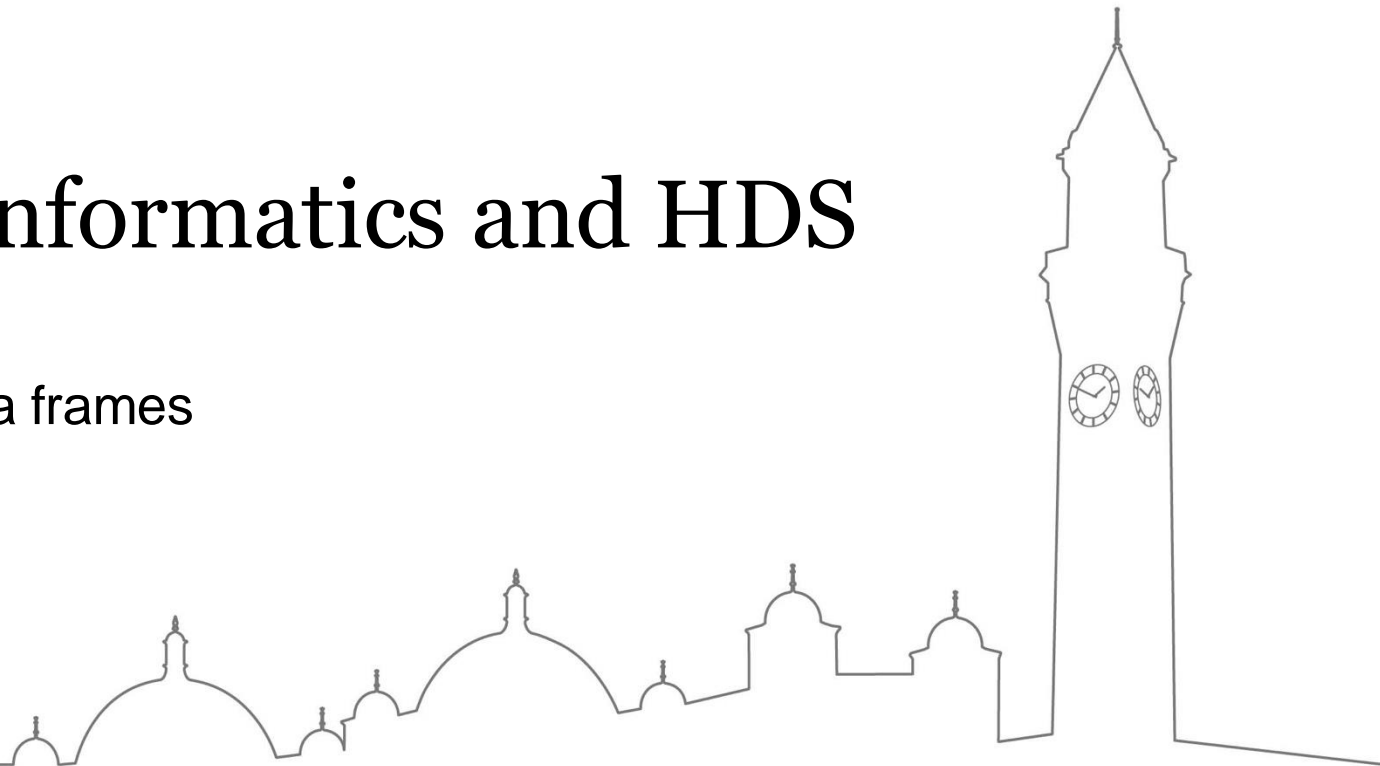


UNIVERSITY OF
BIRMINGHAM

R for Bioinformatics and HDS

R & RStudio:

Working with Data frames



Vasileios Panagiotis Lenis
Laura Bravo Merodio

Course Structure

- Introduction to R & RStudio
- Syntax, Comments, Variables, Data Types and Operators
- Conditions, Loops, Functions and Data Structures
- Working with Data frames



Getting started

- The best way to learn how to work with data frames is to do something useful, so this session is built around a common scientific task: data analysis.
- To do that, we will use the following case scenario, borrowed by the Carpentries:

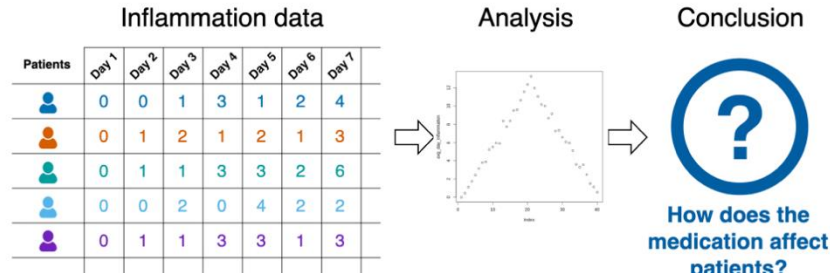


Arthritis Inflammation

- We are studying **inflammation in patients** who have been given a new treatment for arthritis.
- There are 60 patients, who had their inflammation levels recorded for 40 days. We want to analyse these recordings to study the effect of the new arthritis treatment.
- To see how the treatment is affecting the patients in general, we would like to:
 1. Calculate the average inflammation per day across all patients.
 2. Plot the result to discuss and share with colleagues.



UNIVERSITY OF
BIRMINGHAM



Loading the inflammation data

- To begin processing data, we need to load them.
- Loading our inflammation data

```
read.csv(file = "data/inflammation-01.csv", header = FALSE)
```

- **Don't forget!** Store them into a variable to reuse them

```
data <- read.csv(file = "data/inflammation-01.csv", header = FALSE)
```



What is the structure of the "data" object?

- **Data frame:**
- Think of this structure as a spreadsheet in MS Excel that many of us are familiar with.
- Data frames are very useful for storing data and you will use them frequently when programming in R.
- A typical data frame of experimental data contains individual observations in **rows** and variables in **columns**.



Data Inspection

- Let's see the data

```
data
```

- What's the type of our data

```
class(data)
```

- What's the shape? (How many days? How many patients?)

```
dim(data)
```



Data Navigation

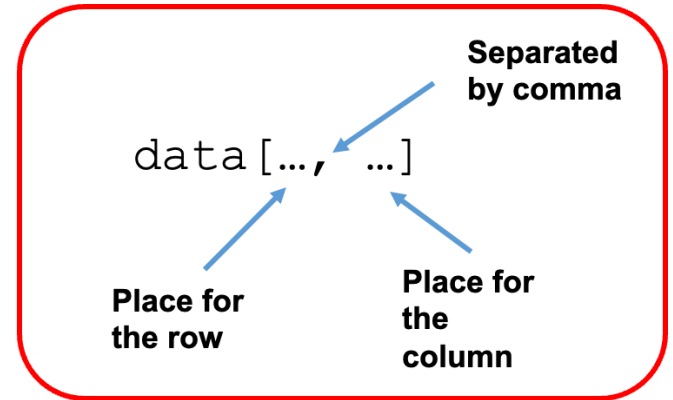
- What is the inflammation value of the first patient in the first day?

Remember! We start counting from “1”!

```
data[1, 1]
```

- What about the middle one?

```
data[30, 20]
```



Slicing data

- We can take more than one values from the dataset

The inflammation scores of the first 4 patients across the first 10 days:

```
data[1:4, 1:10]
```

Or

The inflammation scores of 5 -10 patients across the first 10 days:

```
print(data[5:10, 1:10])
```



Range



UNIVERSITY OF
BIRMINGHAM

Analyzing Data

- We use functions (“canned scripts” that automate something complicated)
- Functions take inputs as arguments and return values as outputs (not in all cases!)
- How can we find the average of all the inflammation scores:

```
mean(data)
```

We use the function “mean” and our data as input



UNIVERSITY OF
BIRMINGHAM

Analyzing data – Descriptive functions

- We can apply many descriptive statistics
maxval, minval, stdval = max(data), min(data), sd(data)
- We can focus on the stats of one patient, or of one day
Like:

```
max(data[1,] #The maximum inflammation of patient 1  
min(data[,1] #The minimum inflammation of day 1  
median(data[,7]) #The median inflammation of day 7
```



Analysing data - summary

- We can have a summary of the descriptive statistics of all our data

```
summary(data[, 1:4]) # Summarize function summary
```

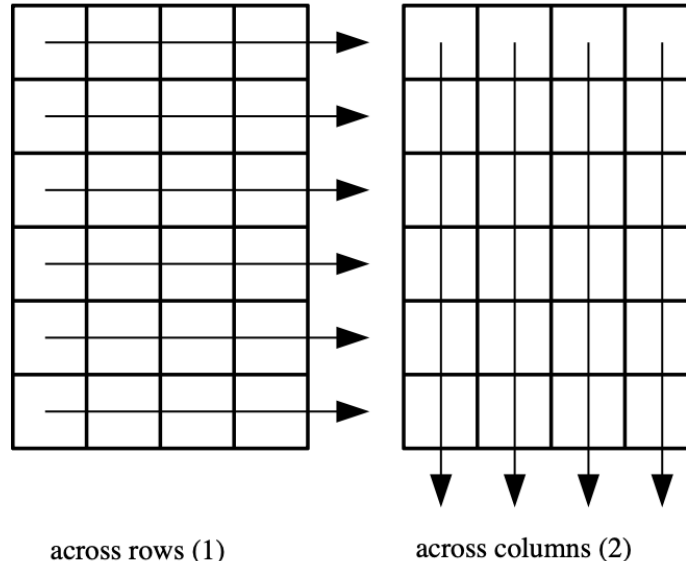
OUTPUT<>

V1	V2	V3	V4
Min. :0	Min. :0.00	Min. :0.000	Min. :0.00
1st Qu.:0	1st Qu.:0.00	1st Qu.:1.000	1st Qu.:1.00
Median :0	Median :0.00	Median :1.000	Median :2.00
Mean :0	Mean :0.45	Mean :1.117	Mean :1.75
3rd Qu.:0	3rd Qu.:1.00	3rd Qu.:2.000	3rd Qu.:3.00
Max. :0	Max. :1.00	Max. :2.000	Max. :3.00



A little bit more advanced calculations....

- What if we need the maximum inflammation for each patient over all days or the average inflammation score for each day?



The *apply()* function

- If we want to calculate the average across the rows

```
apply(data, 1, mean)
```

- If we want to calculate the average across the columns

```
apply(data, 2, mean)
```

- And we can store them to a variable to reuse them...

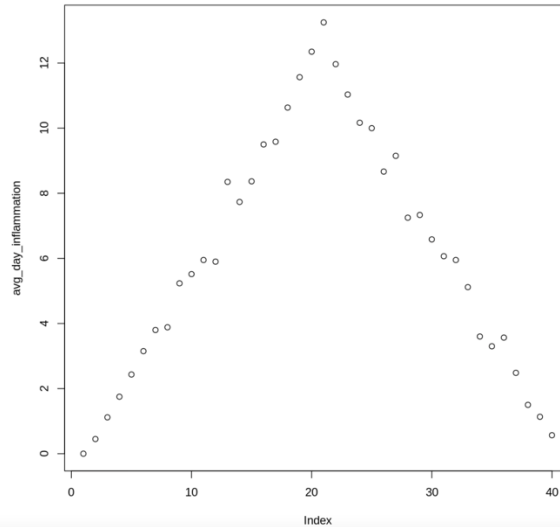
```
avg_patient_inflammation <- apply(data, 1, mean)  
avg_day_inflammation <- apply(data, 2, mean)
```



Let's plot!

- Let's have a look at the average inflammation over time:

```
avg_day_inflammation <- apply(data, 2, mean)  
plot(avg_day_inflammation)
```



10:1 (Top Level) R Script

```
1 #Loading the data
2 data <- read.csv(file = "data/inflammation-01.csv", header = FALSE)
3
4 #Calculating the average inflammation per day (across the columns)
5 avg_day_inflammation <- apply(data, 2, mean)
6
7 #Plotting the results
8 plot(avg_day_inflammation)
9
10
11
```

1
2
3

Save your plot

Files Plots Packages Help Viewer Presentation

Zoom Export Save as Image... Save as PDF... Copy to Clipboard...

Save the current plot as a PDF file

avg_day_inflammation

Index

Console Terminal Background Jobs

R 4.4.0 · ~/DocumentsLocal/RSessions/

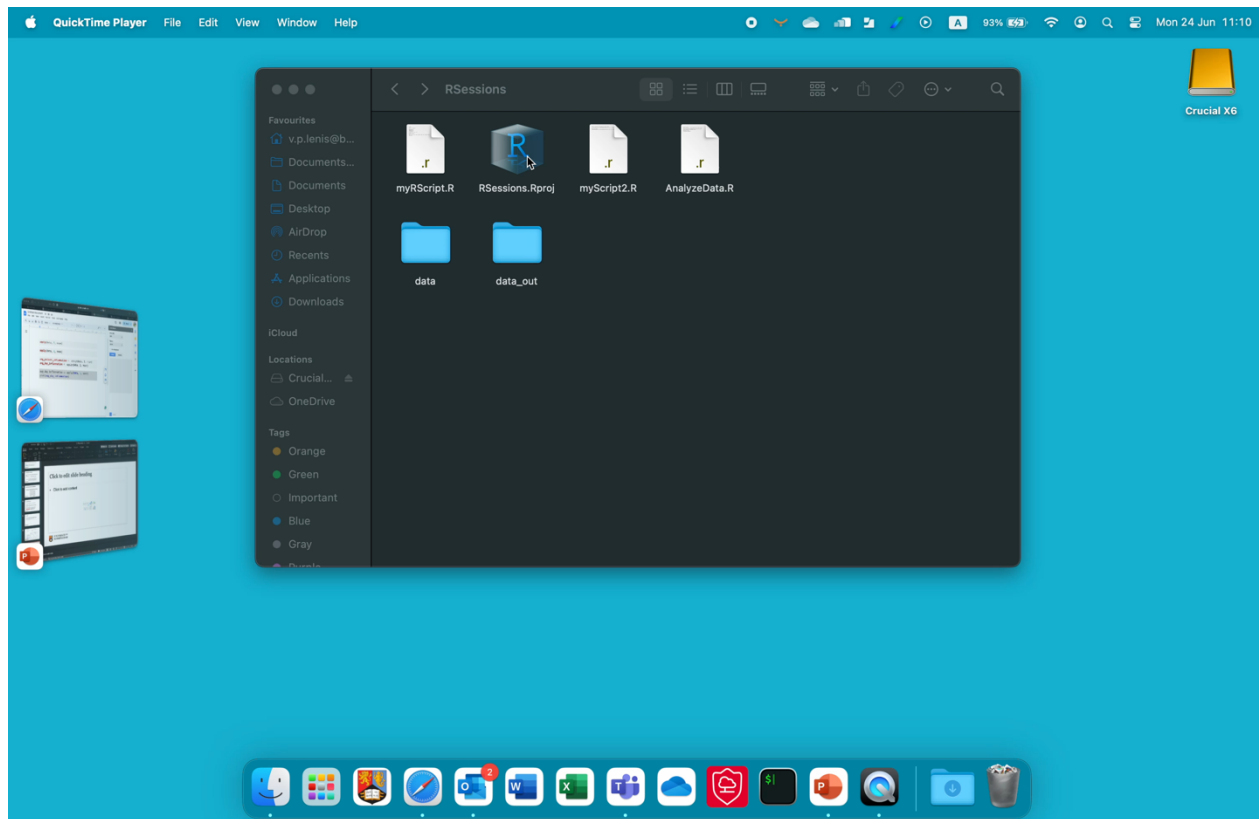
'citation()' on how to cite R or R packages in publications.

Type 'demo()' for some demos, 'help()' for on-line help, or
'help.start()' for an HTML browser interface to help.
Type 'q()' to quit R.

[Workspace loaded from ~/DocumentsLocal/RSessions/.RData]

```
> rm(list = ls())
> #Loading the data
> data <- read.csv(file = "data/inflammation-01.csv", header = FALSE)
> #Calculating the average inflammation per day (across the columns)
> avg_day_inflammation <- apply(data, 2, mean)
> #Plotting the results
> plot(avg_day_inflammation)
>
```





UNIVERSITY OF
BIRMINGHAM



v.p.lenis@bham.ac.uk (Vasilis)
l.bravo@bham.ac.uk (Laura)

