DSA2101

Essential Data Analytics Tools: Data Visualization

Yuting Huang

AY23/24 Semester 2

Course Policies

The teaching team

Instructor:

► Dr. Huang Yuting (yhuang@nus.edu.sg)

▶ Office: S16 04-01

▶ Office hour: In-person and by appointment

Teaching assistants (TAs):

- ► Consultation: By appointment via Zoom/MS Teams.
- ► Contacts of the TAs will be announced before Week 3.

Topics covered

Here is a list of topics we plan to cover this semester:

- 1. R programming
- 2. Importing data into R
- 3. Data manipulation with R
- 4. Principles of data visualization
- 5. The grammar of graphics
- 6. Exploring data through visualization

Tentative teaching plan

- ► Weeks 1–2: Basics in R programming
- ► Weeks 3–4: Importing data into R
- ► Week 5: Data manipulation
- ► Week 6: Tidy data
- ▶ Week 7: Relational data
- ▶ Week 8: In-class midterm test
- ▶ Week 9: Principles of data visualization
- ▶ Weeks 10–11: The grammar of graphics
- ▶ Weeks 12-13: Exploring data through visualization

Lectures

- ▶ Mondays and Wednesdays from 8 to 9:30 am at LT32.
- ▶ Although lectures will be recorded and web-cast, we encourage you to attend the live lectures in person.

Tutorials begin in Week 3.

- ► Attendance is compulsory.
- ▶ You will also need to bring your own laptop.
- ► Tutorial worksheets will be released one week in advance. You are expected to work on the worksheet beforehand.

Evaluation components

► Tutorial	5%
▶ DataCamp assignments	10%
► Group project	15%
▶ Midterm test	30%
► Final exam	40%

Tutorial

Tutorial attendance is compulsory - worth 5% of your final grade.

- ▶ Please adhere to the tutorial session you selected.
- ► In case of illness, send your medical certificate (MC) to your TA. You will be excused from attending the tutorial on that day.

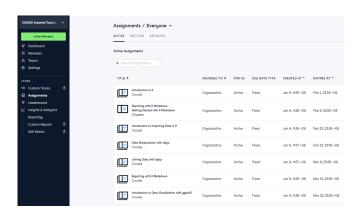
DataCamp assignments

As part of the class, you are required to complete DataCamp assignments (10% of final grade).

- ► The activation link is available on Canvas. Use it to activate your DataCamp Classroom account.
 - ► Sign in with your NUS email (with domain @u.nus.edu).
- ► After activation, you will have access to courses there for 6 months. Do make full use of it!

For each DataCamp assignment, you will receive a binary grade – either full or none.

► They are graded based on completion by the due date, not by XP points.



Group project

Project guideline will be released after the midterm exam.

- ▶ You will work as a team of $3 \sim 5$ members.
- ▶ More details will be announced in due course.

Late submission policy

- ▶ Late submissions will not be graded.
- ► Email submissions will not be graded.
- ▶ Only submissions through Canvas will be graded.

Software

For this class, we will be using the following software.

- 1. R
- 2. RStudio
- 3. Examplify

Make sure that you have a laptop (Windows or Mac), and the latest versions of software above.

Main references

R for Data Science

- ➤ Online textbook: https://r4ds.hadley.nz/preface-2e
- ► Great resource for R using the tidyverse syntax.
- ▶ Lots of examples and exercises.



Due to time constraint, we won't be able to cover all the chapters. If you are serious about the field of data science, we encourage you to read through the book as carefully as you can.

How to reach us?

You can talk to us after lectures and tutorials.

▶ Or email us using your NUS email account.

How to ask a coding question?

- ▶ Copy and paste your code and the entire error message.
- A code that allows us to reproduce the error (i.e., a minimal working example) is ideal.
- ▶ Do not send screenshots.

R-Bloggers: How to ask good questions that prompt useful answers?

BAD ways of asking questions.

```
The stop include ALSI

Like Syr (Sandar (1-4-xiv))

The stop in 100.(1, as. inpital(neal), as. inpital(nea), ...);

Error in 100.(1, as. inpital(local), as. inpital(l
```



- Backtrace:
- dplyr::mutate(starwars_small, height_m = height/100, BMI = mass/(height_m^2))
- dplyr:::mutate.data.frame(starwars_small, height_m = height/100, BMI = mass/(height_m^2))
- dplyr:::mutate_cols(.data, dplyr_quosures(...), by)
- dplyr:::mutate_col(dots[[i]], data, mask, new_columns)
- 6. maskSeval_all_mutate(quo)
- 7. dplyr (local) eval()

Error in mutate(starwars_small, height_m = height/100, BMI = mass/(height_m'2)) : 1 thow Traceback

Caused by error:
| object 'height_m'2' not found

GOOD way of asking questions.

I have an Rscript file (let's call it main.r) which has a reference to another file, using the below code:

```
source("functions.R")
```

But, when I run the RScript file, it complains with the below error:

```
Error in file(filename, "r", encoding = encoding):
cannot open the connection
In addition: Warning message:
In file(filename, "r", encoding = encoding):
cannot open file 'functions.R': No such file or directory
```

I am sure, my main.R file is next to functions.R in the same directory. I can call the functions.R in the Rmd (RMarkdown) file which exist in the same directory

Good ways of asking questions

- 1. Be clear and concise.
 - ▶ What are you trying to achieve? What is the issue you are encountering?
- 2. Provide context, show the code
 - ► Copy and paste the actual line of code that is causing the problem.
- 3. Explain what you have tried.
 - ▶ Mention any attempts to solve the problem.
- 4. Be specific about the error.
 - ▶ Include error message as text rather than screenshots.