

# Project guidelines

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## General notes

- MATH246 students have a separate project. This guidelines only apply to **MATH245** projects.
- **You only have to do one project.** There are three projects to choose from to offer you a range of topics. The three projects are **similar in difficulty**, but they use **different algorithms and methods**. My suggestion is that you choose a project that you find **interesting!** See more details on the project topics below.
- You can do your project in R or **Python**.
- I strongly recommend that you use the **libraries we discussed in the module**. In particular, the plotting libraries (**ggplot2 or plotnine**). Some of the questions are *more difficult* if they are done using **other plotting libraries**.

## Deadline

- All projects are due on **April 23rd 2024, 2pm**.

## Submission

- You **must** submit your projects in **Quarto and PDF format**.
- The PDF output of your project **must not exceed 20 pages** (including the **question statements**).
- **Quarto templates** for R and Python have been **provided**.
- Your Quarto submission **must** produce the submitted PDF, and should run **in a folder exactly as the project folder** in Moodle. In particular, note that **all paths to files and folders** must be relative to the **qmd file location**, and that the data files will be **located exactly as in the Moodle folders**.

## Marking

- A detailed marking scheme will be published before the submission deadline.
- 25 marks will be allocated to coding style (comments, correct use of loops, variable names, functions). Some of this marks will be removed if the Quarto document cannot be compiled.
- Plots must be correctly labeled. Please look at Chapter 5 of the lecture notes for details.

## On the project topics

- The DNA project is about the information content of DNA and a method to visualise it. Although it is “inspired” by biology, it is very much a project about how information is encoded in a string of symbols, and how do we measure information. The data provided is the DNA sequence of one chromosome of a species of yeast (*S. Pombe*)
- The Earthquakes project is about predicting secondary earthquakes. It is a statistical analysis of real earthquake data, and how to use simple features to fit a statistical model.
- The COVID-19 project is about using ordinary differential equations to model infectious diseases. It uses data from the Omicron variant of COVID-19.

## Final notes

- Questions are independent (you might do Question 2 even if you cannot complete Question 1). In some cases, extra data files are provided for that.
- “Roughly”, the questions go from easy to challenging.