Lab 02 – Statistical programming

ENVX1002 Handbook

Semester 1, 2025

Q Learning Outcomes

At the end of this practical students should be able to:

- Acknowledge the importance of academic integrity
- Import, subset and inspect data in R
- Calculate simple summary statistics using both R and Excel
- Generate dynamic reports in Quarto using inline R code
- Understand how to debug R code

Before we begin

Please create a new Quarto document in your project folder to practice R code and complete lab exercises. We will go through the process in Exercise 1.

The following packages are required for this practical:

- tidyverse a collection of R packages designed for data science
- readxl a package for reading Excel files
- modeest a package for estimating the mode of a distribution
- lubridate a package for working with dates and times

If you have not already installed these packages, you can do so by adding the following code into a code chunk in your Quarto document:

```
library(tidyverse)
library(readxl)
library(modeest)
library(lubridate)
```

Finally, please download the data file used in this lab here: soil.xlsx.

Academic integrity

This exercise encourages students to discuss academic integrity, and in particular the grey areas often present. Your demonstrator will provide you with a number of scenarios to discuss with each other in smaller groups, and then with the class.

If you are interested in more information on Academic Integrity at the University of Sydney, see the following link: Academic Integrity. Also ensure you have completed the Academic Honesty Education Module (AHEM). This must be complete before your first assessment is due (next week for ENVX1002).

Exercise 1: Setting up (walk-through)

Getting Started - Important!

Before starting each lab session, you will need to create a new Quarto document to work on the exercises. Follow these steps to set up your lab environment:

1. Open RStudio Project

- Any of the following should work:
 - a. In RStudio, click on the project name in the top-right corner and select your project from a recent list. (It might already be open, otherwise it would show "Project: (None)")
 - b. In RStudio, use File > Open Project to navigate to your . Rproj file
 - c. Double-click the .Rproj file directly from your file explorer

Your project name should appear in the top-right corner of RStudio.

2. Create a New Quarto Document

- Several ways to do it, the easiest is by clicking the "New File" button in the toolbar and selecting "Quarto Document..."
- Give your document a meaningful name (e.g., Lab 2 Exercises), leave other options as default and click "Create"

3. Save and Render Your Document

- You must save your document now. This acts as a simple check to see that you are working in an environment that is not restricted.
- Click File > Save or press Ctrl+S (Windows) / Cmd+S (Mac)
- Name it something meaningful like lab02 practice.qmd
- Click the "Render" button in the editor toolbar to generate the HTML output

Note

If you encounter any issues, please ask your demonstrator for help.

Follow our lead

Important

We've created this section to make sure you don't get lost during the lab. Remember to read "A brief R guide for surviving ENVX1002" available in the Tool Kit section on Canvas for a better understanding of how to use R in this course.

Two golden rules

- 1. Text are formatted in Markdown (like this text)
- 2. Code must be written inside code chunks

A code chunk is what makes your document "dynamic" as it can execute code and perform all sorts of tasks.

To create a code chunk:

- Quick way: Use keyboard shortcuts
 - ► Windows/Linux: Ctrl + Alt + I
 - ► Mac: Cmd + Option + I
- Manual way: Type three backticks followed by {r}, then end with three backticks

Here's what a code chunk looks like:

```
```{r}
mean(c(1, 2, 3))
```

Note: the "fence" that we use (the three backticks plus the {r}) will only be visible in Source view. Your demonstrator will show you how to switch between Source and Preview views.

## **Understanding Functions**

This is a function:

```
Input → Process → Output
mean(c(2, 4, 6)) # Takes numbers, calculates average, returns 4
```

```
[1] 4
```

You can recognise it because it has:

- A name (mean)
- Parentheses () to hold input data
- Input data inside the parentheses (e.g., c(2, 4, 6))

A function will almost always return an output, which you can use in your code. In this case, the output is 4. Functions are the reason why we can do so much with R – they are like actions that someobe else has written for us to use so that we can complete our tasks with minimal programming.

Let's look at three common functions for central tendency, but at the same time look at what it means to run functions on data we often call "objects":

1. mean():

```
Create a vector of numbers and store it in an object called "scores"
scores <- c(85, 92, 78, 95, 8)
Calculate the mean of the scores
mean(scores)</pre>
```

```
[1] 71.6
```

2. median():

```
heights <- c(160, 165, 168, 170, 185)
median(heights)
```

```
[1] 168
```

3. mode():

```
votes <- c("yes", "no", "yes", "maybe", "yes")
mfv(votes)</pre>
```

```
[1] "yes"
```

Remember: Every function needs:

- Parentheses () to work
- Input data (inside the parentheses)
- Sometimes extra options (like na.rm = TRUE to handle missing values)

#### **Exercise 2: Water quality**

Sulphate ( $SO_4$ ) is a key indicator in water quality monitoring and can be used to assess environmental impacts from industry and agriculture. In this exercise, we will explore a dataset of  $SO_4$  concentrations (mg/kg) in water samples.

The data is stored in a sheet called "SO4" in the MS Excel file, soil.xlsx.

```
so4 <- read_excel("data/soil.xlsx", sheet = "S04")</pre>
```

Try writing and running the code chunks in your own Quarto document to see their outputs. Results will appear below each chunk. See Lab 01 for more information on inserting and running code chunks in Quarto.

When we load data into R for the first time, it is important to check what it looks like (and whether it loaded correctly). The str() function is a good way to quickly inspect the data:

```
str(so4)
```

```
tibble [39 × 1] (S3: tbl_df/tbl/data.frame)
$ S04: num [1:39] 50.6 55.4 56.5 57.5 58.3 63 66.5 64.5 63.4 58.4 ...
```

# Question 1

What does the output of str() tell us about the data? You may want to look at the documentation ?str or search online for more information, but ask your demonstrator if you're still unsure about why we use this function. Relate this to the data you are working with.

## Answer 1

The str() function provides a concise summary of the data frame structure. It tells us the number of observations (rows) and variables (columns), as well as the data type of each variable. This is useful for understanding the data we are working with. For this particular dataset, we see that it contains 30 observations of one variable, S04, which is a numeric variable inside a tibble object. All this can be inferred from the output of str():

```
tibble [39 × 1] (S3: tbl_df/tbl/data.frame)
$ S04: num [1:39] 50.6 55.4 56.5 57.5 58.3 63 66.5 64.5 63.4 58.4 ...
```

Since the data is a data frame object, we have a good idea of what functions we can use to explore it. Let's examine the first few rows of our data:

```
head(so4)
```

Let's calculate some basic descriptive statistics. Read the code and try to understand every part of it, including the text in #.

```
Calculate mean, median, and mode
mean_so4 <- mean(so4$S04)
median_so4 <- median(so4$S04)
mode_so4 <- mfv(so4$S04)[1] # Most frequent value using modeest package

Calculate measures of spread
range_so4 <- range(so4$S04)
iqr_so4 <- IQR(so4$S04)
var_so4 <- var(so4$S04)
sd_so4 <- sd(so4$S04)</pre>
```

#### Reporting your data

When reporting statistics in a Quarto document, there are two approaches we could take:

## 1. Basic R output:

```
mean_so4

[1] 61.92308

sd_so4

[1] 5.241558

median_so4

[1] 62.1
```

## 2. Inline reporting (recommended):

Consider the following reporting:

The mean  $SO_4$  concentration in our samples is 61.92 mg/kg, with a standard deviation of 5.24 mg/kg. The median value is 62.1 mg/kg.

Using inline R code (approach 2) has several advantages:

- Statistics are seamlessly integrated into your text
- Numbers are automatically updated if your data changes
- Results are presented in a reader-friendly format

To create inline R code, use backticks with r, like this:

```
The mean S0~4~ concentration in our samples is `{r} round(mean_so4, 2)` mg/kg...
```

Try to recreate the report above in your Quarto document, or use other objects like mode\_so4 and var\_so4 in your report.

# **Exercise 3: Using MS Excel**

Why use Excel when you have R? Well, Excel is an ubiquitous tool in many industries and can be a useful complement to R for quick data exploration and analysis. Sometimes it may even be easier to use Excel for simple tasks.

Let's calculate the same statistics using Excel to compare approaches. First:

- 1. Open the soil.xlsx file in Excel
- 2. Navigate to the "SO4\_excel" sheet (we'll use this sheet to avoid modifying the original data)
- 3. Ensure the data is properly displayed in columns

MS Excel remains a popular tool for quick and dirty data analysis and can be a valuable resource in just about any field.

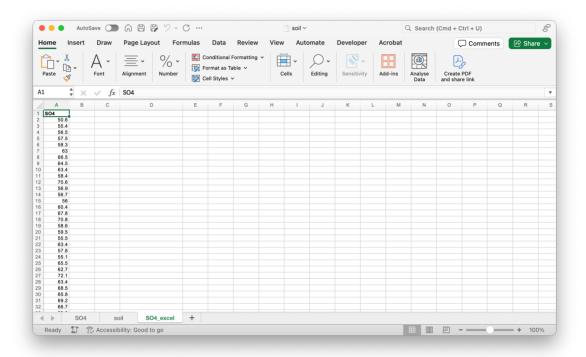


Figure 1: Imported data in Excel. Make sure to use the "SO4\_excel" sheet.

For any calculation in Excel:

- 1. Select an empty cell
- 2. Type "=" followed by the function name
- 3. Select your data range
- 4. Press Enter

For example, to calculate the mean:

- 1. Click an empty cell
- 2. Type =AVERAGE(
- 3. Select all SO4 values
- 4. Type ) and press Enter

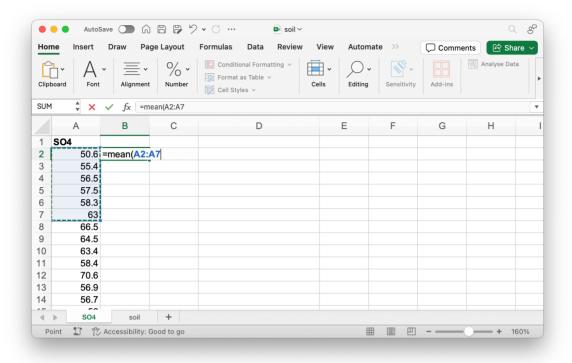
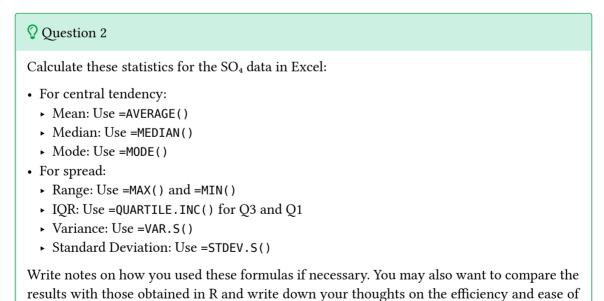


Figure 2: Using the =AVERAGE() formula in Excel. Note that this screenshot demonstrates the selection of a specific range of data.



use for each method.

#### Answer 2

This exercise should be straightforward. Ask your demonstrator if you encounter any issues.

## Exercise 4: Soil data

In agricultural science, soil characteristics are essential for understanding soil health and fertility. The soil sheet in the soil.xlsx file contains data on soil properties at different depths, as well as lithology and land use information.

In this exercise, we'll explore different ways to subset data in R using the soil characteristics data. First, let's load the data:

```
soil <- read_excel("data/soil.xlsx", sheet = "soil")</pre>
```

## Understanding data structure

Let's examine our data structure:

```
str(soil)
```

```
head(soil)
```

```
A tibble: 6 \times 7
 id
 clay0 clay60
 ec0 ec60 lithology
 land use
 <chr> <dbl> <dbl> <dbl> <dbl> <chr>
 <chr>
1 AT1 21.3
 30.4 52.4 32.6 Siliceous Mid
 Grazing
 38.2 34.8 25.1 Siliceous Mid
2 AT2 21.4
 Grazing
3 AT3
 4.52 42.6 52.8 38.5 Siliceous Lower Grazing
4 BM1 19.4 24.6 35.2 26.4 Siliceous Lower Grazing
5 BM2 40.6 75.6 46.4 35.2 Mafic
 Grazing
6 BM3 42.7
 75.1 50.0 59.7 Mafic
 Grazing
```

# Question 3

What do you notice about the data structure that is different from the SO<sub>4</sub> data? How does this affect the way we subset the data?

## Answer 3

From str() we can observe that the soil data frame contains multiple columns, each representing a different soil property. This means we can subset the data based on different variables, such as clay content, electrical conductivity, lithology, and land use. The head() function shows the first few rows of the data, which can help us understand the data structure and content.

The soil data frame contains the following columns:

- clay0: Clay content at 0 cm depth
- clay60: Clay content at 60 cm depth
- ec0: Electrical conductivity at 0 cm depth
- ec60: Electrical conductivity at 60 cm depth
- lithology: Type of soil composition
- land\_use: Land usage type

#### Basic subsetting in R

There are two main ways to subset data in R:

- 1. Using the \$ operator to select columns
- 2. Using square brackets [] to select rows and columns

#### Using the \$ operator

The \$ operator allows us to select a specific column from our data frame. For example:

```
Get the land use column
soil$land_use
```

```
[1] "Grazing"
 "Grazing"
 "Grazing"
 "Grazing"
 "Grazing"
 "Grazing"
 "Grazing"
 "Grazing"
 "Grazing"
 "Grazing"
 "Grazing"
 [7] "Grazing"
 "Grazing"
 "Grazing"
[13] "Grazing"
 "Grazing"
 "Grazing"
 "Grazing"
[19] "Grazing" "Grazing"
 "Grazing"
 "Cropping" "Cropping"
 "Grazing"
[25] "Cropping" "Cropping"
 "Grazing"
 "Grazing"
 "Grazing"
 "Grazing"
[31] "Grazing"
 "Grazing"
 "Cropping" "Cropping"
 "Cropping" "Grazing"
[37] "Cropping" "Grazing"
 "Grazing"
 "Grazing"
 "Cropping" "Grazing"
 "Cropping" "Grazing"
[43] "Grazing"
 "Grazing"
 "Grazing"
 "Grazing"
 "Natural" "Natural"
[49] "Grazing"
 "Grazing"
 "Grazing"
 "Cropping"
[55] "Cropping"
```

Combined with functions, the \$ operator can be used to calculate statistics on specific columns.

```
Calculate the mean clay content at 0 cm depth
mean_clay0 <- mean(soil$clay0)
mean_clay0</pre>
```

```
[1] 23.198
```

Notice how we saved the result of the mean() function in a new object mean\_clay0. This is useful for storing results and using them later in your code (e.g. for inline reporting). However, it means that the result will not be displayed in the console unless you explicitly print it by typing mean\_clay0 again in a new line.

#### i Note

The \$ operator is particularly useful when you want to:

- · Access a single column quickly
- Use column values in calculations
- Create plots with specific variables

## Using square brackets []

Square brackets allow more flexible subsetting using the format: dataframe[rows, columns]. It also works with vectors, lists, and matrices. Try the following examples:

```
soil[1:5,] # First 5 rows, all columns
```

```
A tibble: 5 × 7
id clay0 clay60 ec0 ec60 lithology land_use
<chr> <dbl> <dbl> <dbl> <dbl> <chr> < 30.4 52.4 32.6 Siliceous Mid Grazing
2 AT2 21.4 38.2 34.8 25.1 Siliceous Mid Grazing
3 AT3 4.52 42.6 52.8 38.5 Siliceous Lower Grazing
4 BM1 19.4 24.6 35.2 26.4 Siliceous Lower Grazing
5 BM2 40.6 75.6 46.4 35.2 Mafic Grazing
```

```
soil[, c("clay0", "clay60")] # Clay columns
```

```
A tibble: 55 × 2
clay0 clay60
<dbl> <dbl>
1 21.3 30.4
```

```
2 21.4
 38.2
3 4.52 42.6
4 19.4
 24.6
5 40.6
 75.6
6 42.7
 75.1
7 28.1
 50.5
8 41.3
 72.6
9 48.7
 78.7
10 26.3
 50.5
i 45 more rows
```

```
soil[1:3, c("ec0", "ec60")] # First 3 rows, electrical conductivity columns
```

```
A tibble: 3 × 2
 ec0 ec60
 <dbl> <dbl>
1 52.4 32.6
2 34.8 25.1
3 52.8 38.5
```

When using [], leaving the row or column section empty (with just a comma) means "select all"

**Basic subsetting**: Use square brackets to extract soil samples at 0cm depth that have clay content greater than 40%.

```
soil[soil$clay0 > 40,]
```

**Multiple conditions**: Find samples where clay content at 60cm is greater than 30% AND electrical conductivity at 0cm is less than 0.5.

```
soil[soil$clay60 > 30 & soil$ec0 < 0.5,]
```

# Question 4

**Practice**: Subset the data to find samples where:

- clay content at 0cm depth is less than 50%
- electrical conductivity at 60cm depth is between 25 and 30, inclusive
- clay content at 0cm is less than 10% OR electrical conductivity at 60cm is greater than 25

#### Answer 4

Clay content at 0cm depth is less than 50%

```
soil[soil$clay0 < 50,]
```

```
A tibble: 54 \times 7
 id
 clay0 clay60
 ec0 ec60 lithology
 land use
 <chr> <dbl> <dbl> <dbl> <dbl> <chr>
 <chr>
 21.3
1 AT1
 30.4 52.4 32.6 Siliceous Mid
 Grazing
2 AT2
 21.4
 38.2 34.8 25.1 Siliceous Mid
 Grazing
3 AT3
 4.52
 42.6 52.8 38.5 Siliceous Lower
 Grazing
 4 BM1
 19.4
 24.6 35.2 26.4 Siliceous Lower
 Grazing
5 BM2
 40.6
 75.6 46.4 35.2 Mafic
 Grazing
6 BM3
 42.7
 75.1 50.0 59.7 Mafic
 Grazing
 7 BM4
 28.1
 50.5 66.9 47.6 Mafic
 Grazing
8 BS1
 41.3
 72.6 100. 396. Intermediate Lower Grazing
9 BS2
 48.7
 78.7 83.3 447. Intermediate Lower Grazing
10 C1
 26.3
 50.5 80.2 39.8 Mafic
 Grazing
i 44 more rows
```

Electrical conductivity at 60cm depth is between 25 and 30, inclusive

```
soil[soil\$ec60 >= 25 \& soil\$ec60 <= 30,]
```

```
A tibble: 7 \times 7
 id
 clay0 clay60
 ec0 ec60 lithology
 land_use
 <chr>
 <chr> <dbl> <dbl> <dbl> <dbl> <chr>
 38.2 34.8 25.1 Siliceous Mid
1 AT2
 21.4
 Grazing
2 BM1
 19.4
 24.6 35.2 26.4 Siliceous Lower Grazing
3 HEN1 11.0
 53.1 38.6 28.8 Siliceous Lower Grazing
 56.2 29.5 28.6 Siliceous Lower Grazing
4 HEN4
 9.65
5 JB3
 30.8
 48.2 57.2 27.2 Mafic
 Cropping
6 <NA>
 NA
 NA
 NA
 NA
 <NA>
 <NA>
7 R0M7
 17.5
 36.1 42.6 25.2 Siliceous Lower Grazing
```

Clay content at 0cm is less than 10% OR electrical conductivity at 60cm is greater than 25

```
soil[soil$clay0 < 10 | soil$ec60 > 25,]
```

```
A tibble: 46 \times 7
 clay0 clay60
 ec0 ec60 lithology
 land_use
 id
 <chr> <dbl> <dbl> <dbl> <dbl> <chr>
 <chr>
1 AT1
 21.3
 30.4 52.4 32.6 Siliceous Mid
 Grazing
 38.2 34.8 25.1 Siliceous Mid
2 AT2
 21.4
 Grazing
 38.5 Siliceous Lower
 3 AT3
 4.52
 42.6 52.8
 Grazing
 4 BM1
 19.4
 24.6 35.2 26.4 Siliceous Lower
 Grazing
 5 BM2
 40.6
 75.6 46.4
 35.2 Mafic
 Grazing
 6 BM3
 42.7
 75.1 50.0 59.7 Mafic
 Grazing
 7 BM4
 28.1
 50.5 66.9 47.6 Mafic
 Grazing
8 BS1
 41.3
 72.6 100. 396. Intermediate Lower Grazing
 78.7 83.3 447. Intermediate Lower Grazing
 9 BS2
 48.7
```

## **Exercise 5: Soil statistics**

We will continue to work on the soil dataset and practice calculating some basic summary statistics.

# **Question** 5

- 1. Calculate the mean, median, and mode for clay content at 0cm depth
- 2. Calculate the range, IQR, variance, and standard deviation for electrical conductivity at  $60\mathrm{cm}$  depth
- 3. Report these statistics in your Quarto document using inline R code

#### Answer 5

Notice how in all the calculations, we used the na.rm = TRUE argument to handle missing values. This is important when calculating statistics to avoid errors.

1. Calculate the mean, median, and mode for clay content at 60cm depth

```
mean_clay0 <- mean(soil$clay60, na.rm = TRUE)
mean_clay0</pre>
```

[1] 46.63815

```
median_clay0 <- median(soil$clay60, na.rm = TRUE)
median_clay0</pre>
```

[1] 46.91

```
mode_clay0 <- mfv(soil$clay60, na.rm = TRUE)[1]</pre>
```

argument 'na.rm' is soft-deprecated, please start using 'na rm' instead

mode\_clay0

[1] 10.38

2. Calculate the range, IQR, variance, and standard deviation for electrical conductivity at 60cm depth

```
range_ec60 <- range(soil$ec60, na.rm = TRUE)
range_ec60</pre>
```

[1] 15.86 446.72

```
iqr_ec60 <- IQR(soil$ec60, na.rm = TRUE)
iqr_ec60</pre>
```

[1] 31.2675

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#### Done!

This is the end of Lab 02. Remember to save your Quarto document and submit it to Canvas when you're done. If you have any questions, feel free to ask your demonstrator for help.

#### **Bonus: Take-home exercises**

Take-home exercises are optional but recommended for further practice. You can complete these exercises in your own time.

#### **Exercise 5: Mario Kart statistics**

We're not going to go too much about the game – but here is a dataset of character attributes from the popular Mario Kart.

Download it here: mario kart.csv

The data is stored in a CSV file, which MS Excel can open directly. To read it into R, use the read csv() function from the readr package.

The dataset contains the following variables:

- character: Character name
- weight: Character's weight class (1-5 scale)
- speed: Maximum speed rating (1-5 scale)
- acceleration: How quickly they reach top speed (1-5 scale)
- handling: Steering control rating (1-5 scale)
- traction: Grip on different surfaces (1-5 scale)
- drift\_rating: Skill at power-sliding around corners (1-5 scale)

Complete the following exercises using both R and Excel:

#### 1. Data Import and Inspection

- Load the data into R using read\_csv()
- Examine the structure using str()
- View the first few rows using head()
- What do you notice about the data?

#### 2. **Central Tendency** Calculate for the speed attribute:

- Mean
- Median
- Mode

Which measure best represents the "typical" speed rating? Why?

- 3. **Spread Analysis** For the weight attribute, calculate:
  - Range
  - IQR
  - Variance
  - Standard deviation

What do these tell you about the variation in character weights?

## 4. Character Comparison

- Which characters have the highest and lowest acceleration?
- Find all characters with above-average handling
- Identify characters with both speed and weight above 4.0

# 5. Advanced Challenge

- Calculate the mean of all attributes for each character
- Who is the most "well-rounded" character (closest to average in all stats)?
- Create a report comparing Mario and Luigi's stats using inline R code

#### 1. Data Import and Inspection

```
Load the data
mario_kart <- read_csv("data/mario_kart.csv")</pre>
Rows: 12 Columns: 7
 Column
 specification
Delimiter: ","
chr (1): character
dbl (6): weight, speed, acceleration, handling, traction, drift rating
i Use `spec()` to retrieve the full column specification for this data.
i Specify the column types or set `show_col_types = FALSE` to quiet this
message.
Examine structure
str(mario kart)
spc_tbl_ [12 x 7] (S3: spec_tbl_df/tbl_df/tbl/data.frame)
 $ character : chr [1:12] "Mario" "Luigi" "Peach" "Bowser" ...
 $ weight : num [1:12] 3.5 3.4 2.8 4.5 3 2.5 4.3 4.2 2.7 2.8 ...
$ speed : num [1:12] 4.2 4.1 3.8 4.5 4 3.7 4.4 4.3 3.6 3.8 ...
 $ acceleration: num [1:12] 3.8 3.9 4.2 2.8 4.1 4.4 2.9 3 4.3 4.2 ...
 $ handling : num [1:12] 3.5 3.6 4 2.5 3.8 4.2 2.7 2.8 4.1 4 ...
 $ traction : num [1:12] 3.7 3.8 3.5 3.2 3.9 3.6 3.3 3.4 3.7 3.6 ...
 $ drift_rating: num [1:12] 4 3.9 3.7 3.4 4.1 3.8 3.5 3.6 3.9 3.8 ...
 - attr(*, "spec")=
 .. cols(
 character = col_character(),
 .. weight = col double(),
 .. speed = col double(),
 acceleration = col_double(),
 handling = col_double(),
 traction = col double(),
 drift_rating = col_double()
 - attr(*, "problems")=<externalptr>
View first few rows
head(mario_kart)
```

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Mario and Luigi have similar stats, but with some notable differences:

Ona the prince Maritan Ness 3 3 4 2 Pan Lu Inean 4 Cab. Usa 3 4 2 La. to La t

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