

# Build a MS Word report

## Step-by-step correction

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This page provides a step-by-step correction for completing Exercise 3. To get the most out of your learning experience, attempt to solve the exercise on your own before looking at this correction. Trial and error is an essential part of the learning process, helping to strengthen your understanding and build confidence. Remember, simply copying and pasting the solution without trying to solve the exercise by yourself first will limit your long-term growth. Take your time, experiment, and learn actively!

### 0.1 Setup basic elements of the Quarto document

All these basic elements are part of the YAML header.

1

---

2

title: My outbreak report

3

author: Helene Langet

4

date: "2023-12-31"

5

①

②

③

```

6 format: docx
7 echo: false
8 warning: false
9 ---

```

- ① Update the title of the Quarto document.
- ② Put your name as author of the Quarto document.
- ③ Add the date 2023-12-31 to the Quarto document.
- ④ Change the output format to generate a MS Word document.
- ⑤ Configure the Quarto document to hide code in the rendered MS Word document.
- ⑥ Configure the Quarto document to hide warnings in the rendered MS Word document.

## 0.2 Improve navigation and readability

```

1 ---
2 title: My outbreak report
3 author: Helene Langet
4 date: "2023-12-31"
5
6 format:
7   docx:
8     toc: true
9     number-sections: true
10    toc-depth: 2
11 echo: false
12 warning: false
13 ---

```

- ⑦ Insert a table of contents to the Quarto document.
- ⑧ Automatically number the different sections of the Quarto document.
- ⑨ Configure the table of contents to only display two levels of section headings.

### ! YAML indentation

YAML is a whitespace-sensitive language where indentation determines the structure; tabs are not recognised for indentation. The recommended practice is therefore to use **two spaces** per indentation level to ensure consistency and avoid errors.

## 0.3 Implement dynamic calculations

### 0.3.1 In the YAML header

```
1 ---
2 title: My outbreak report
3 author: Helene Langet
4 date: last-modified
5 date-format: long
6
7 format:
8   docx:
9     toc: true
10    number-sections: true
11    toc-depth: 2
12 echo: false
13 warning: false
14 ---
```

⑩

⑪

⑩ Change the date 2023-12-31 to the date at which the Quarto document was last modified.

⑪ Format this date to display it with the format `December 31, 2023`.

### 0.3.2 In the Quarto notebook

- ☐ Replace the placeholder text in bold with the automated calculation of the outbreak start and end dates ;

*“The outbreak ran from **date** to **date**”*

- ☐ Replace the placeholder text in bold with the automated calculation of the number of cases, confirmed cases and deaths.

*“Over the studied period, there were **N** cases, including **N** confirmed cases and **N** confirmed deaths.”*

## 0.4 Create and reference publication-ready tables

You can use packages such as `gtsummary`

To install the package, run the following command in your console.

```

1 ```{r}
2 install.packages(gtsummary)
3 ```

```

```

1 ```{r}
2 library(gtsummary)
3 ```

```

```

1 ```{r}
2 #| label: tbl-1 # <14>
3 #| tbl-cap: Population characteristics # <13>
4
5 subdf |> # <12>
6   dplyr::select(age,
7                 sex,
8                 bmi,
9                 confirmed,
10                death) |>
11   gtsummary::tbl_summary()
12 ```

```

- ⑫ Create a table summarising the demographic characteristics and outcome frequency of all cases.
- ⑬ Add a caption to the table.
- ⑭ Assign a label to the table.

Table 1: Population characteristics

Characteristic	N = 65,669
age	50 (35, 65)
sex	
1	33,114 (50%)
2	32,555 (50%)
bmi	29 (21, 38)
confirmed	
0	13,235 (20%)
1	52,434 (80%)
death	
0	64,455 (98%)
1	1,214 (1.8%)

☒ Replace the placeholder text in bold with a cross-reference to the table ;

@tbl-1 provides a summary of the demographic characteristics and the outcome proportion for the overall population.

*“Table 1 provides a summary of the demographic characteristics and the outcome proportion for the overall population”*

```

1  ```{r}
2  #| label: tbl-2 # <17>
3  #| tbl-cap: Demographic characteristics of deceased vs. alive # <16>
4
5  subdf |> # <15>
6    dplyr::select(sex,
7                  age,
8                  bmi,
9                  death) |>
10   gtsummary::tbl_summary(by = death) |>
11   gtsummary::add_overall()
12  ```

```

- ⑮ Create a table summarising the demographic characteristics of individuals who died versus those who are still alive.
- ⑯ Add a caption to the table.
- ⑰ Assign a label to the table.

Table 2: Demographic characteristics of deceased vs. alive

Characteristic	Overall, N = 65,669	0, N = 64,455	1, N = 1,214
sex			
1	33,114 (50%)	32,504 (50%)	610 (50%)
2	32,555 (50%)	31,951 (50%)	604 (50%)
age	50 (35, 65)	50 (35, 65)	52 (37, 67)
bmi	29 (21, 38)	29 (21, 38)	34 (28, 41)

☐ Replace the placeholder text in bold with a cross-reference to the table.

@tbl-2 compares the demographic characteristics of individuals who died versus those who are still alive.

*“..., while Table 2 compares the demographic characteristics of individuals who died versus those who are still alive.”*

## 0.5 Customise figures

```
1  ```{r}
2  #| label: fig-1 # <19>
3  #| fig-cap: Weekly count of all cases, confirmed cases and deaths # <18>
4  #| fig-width: 8 # <20>
5  #| fig-height: 4 # <20>
6
7  # Aggregate the data to get the weekly count of all cases, confirmed cases and deaths
8  weekly_data <- subdf |>
9    dplyr::group_by(week) |>
10    dplyr::summarise(count = dplyr::n(),
11                      confirmed_count = sum(confirmed == "1"),
12                      death_count = sum(death == "1"))
13
14  # Plot the weekly cases, confirmed cases and deaths
15  ggplot2::ggplot(weekly_data, ggplot2::aes(x = week)) +
16    ggplot2::geom_line(ggplot2::aes(y = count,
17                                     color = "All cases"),
18                       size = 1) +
19    ggplot2::geom_line(ggplot2::aes(y = confirmed_count,
20                                     color = "Confirmed cases"),
21                       size = 1) +
22    ggplot2::geom_line(ggplot2::aes(y = death_count,
23                                     color = "Confirmed deaths"),
24                       size = 1) +
25    ggplot2::labs(x = "Week",
26                  y = "Count",
27                  color = "Legend") +
28    ggplot2::scale_color_manual(values = c("All cases" = "#440e54",
29                                           "Confirmed cases" = "#f8766d",
30                                           "Confirmed deaths" = "#128984")) +
31    ggplot2::theme_minimal() +
32    ggplot2::theme(panel.grid.major.y = ggplot2::element_line(linewidth = 0.5, linetype = "dashed"),
33                  panel.grid.minor.y = ggplot2::element_line(linewidth = 0.5, linetype = "dashed"),
34                  panel.grid.major.x = ggplot2::element_blank(),
35                  panel.grid.minor.x = ggplot2::element_blank())
36  ```
```

- ⑩ Add a caption to the figure.
- ⑪ Assign a label to the figure.
- ⑫ Adjust the dimensions of the figure until you are happy with it

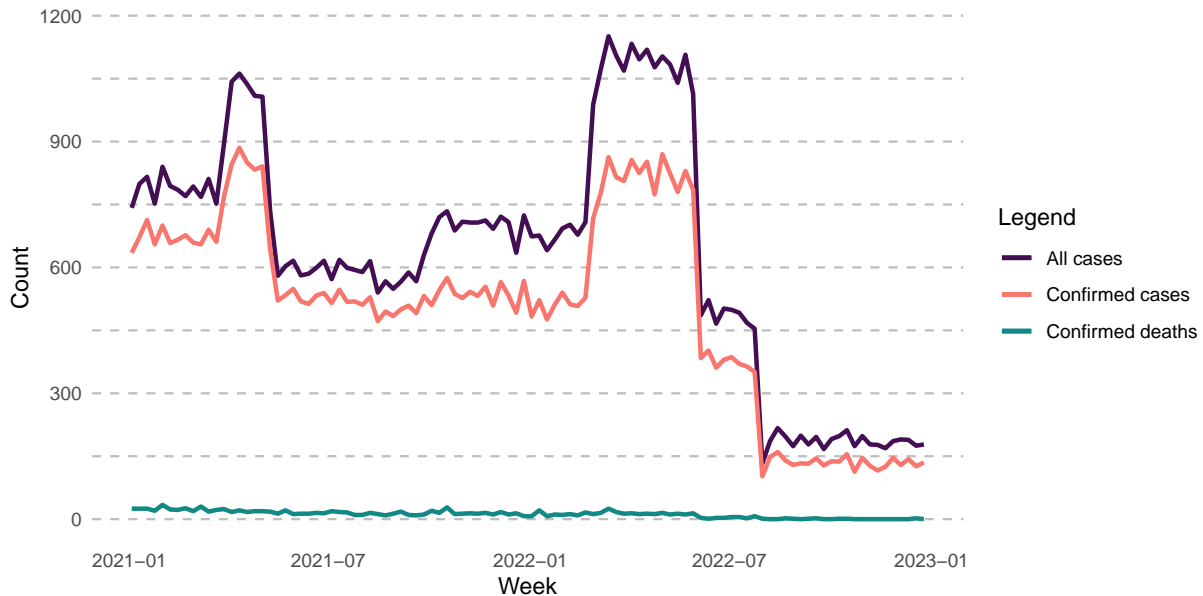


Figure 1: Weekly count of all cases, confirmed cases and deaths

- ☐ Replace the placeholder text in bold with a cross-reference to the figure.

Figure 1 illustrates the outbreak's progression, which can be divided into distinct phases.

*“**cross-reference** illustrates the outbreak's progression, which can be divided into distinct phases.”*

21. Beautify the plot using ggplot options

## 0.6 Code

- ☐ Display the code chunk for your R implementation of the logistic regression (and only this code chunk) in the rendered MS Word document ;
- ☐ Add a caption to the code chunk ;
- ☐ Assign a label to the code chunk ;
- ☐ Replace the placeholder text in bold with a cross-reference to the code chunk.

The logistic regression model uses `death` as the response variable, and `bmi` and `age` as predictor variables. It is implemented in R as shown in the code chunk referenced by Listing 1

22. Implement a logistic regression model based on description in the Quarto document.

The results of the logistic regression model are summarized in the formatted regression table, which is presented in Table 3.

---

**Listing 1** R code

---

```
1 ```{r}
2 #| lst-label: lst-test
3 #| lst-cap: R code
4
5 coeffs <- glm(death ~ bmi + age, # <22>
6               subdf |> dplyr::filter(confirmed == "1"),
7               family = binomial)
8 ```
```

---

```
1 ```{r}
2 #| label: tbl-3 # <25>
3 #| tbl-cap: Formatted regression table # <24>
4
5 gtsummary::tbl_regression(coeffs, exponentiate = TRUE) # <23>
6 ```
```

- ②③ Create a table summarizing the odds ratios from the logistic regression model.
- ②④ Add a caption to the table.
- ②⑤ Assign a label to the table.

Table 3: Formatted regression table

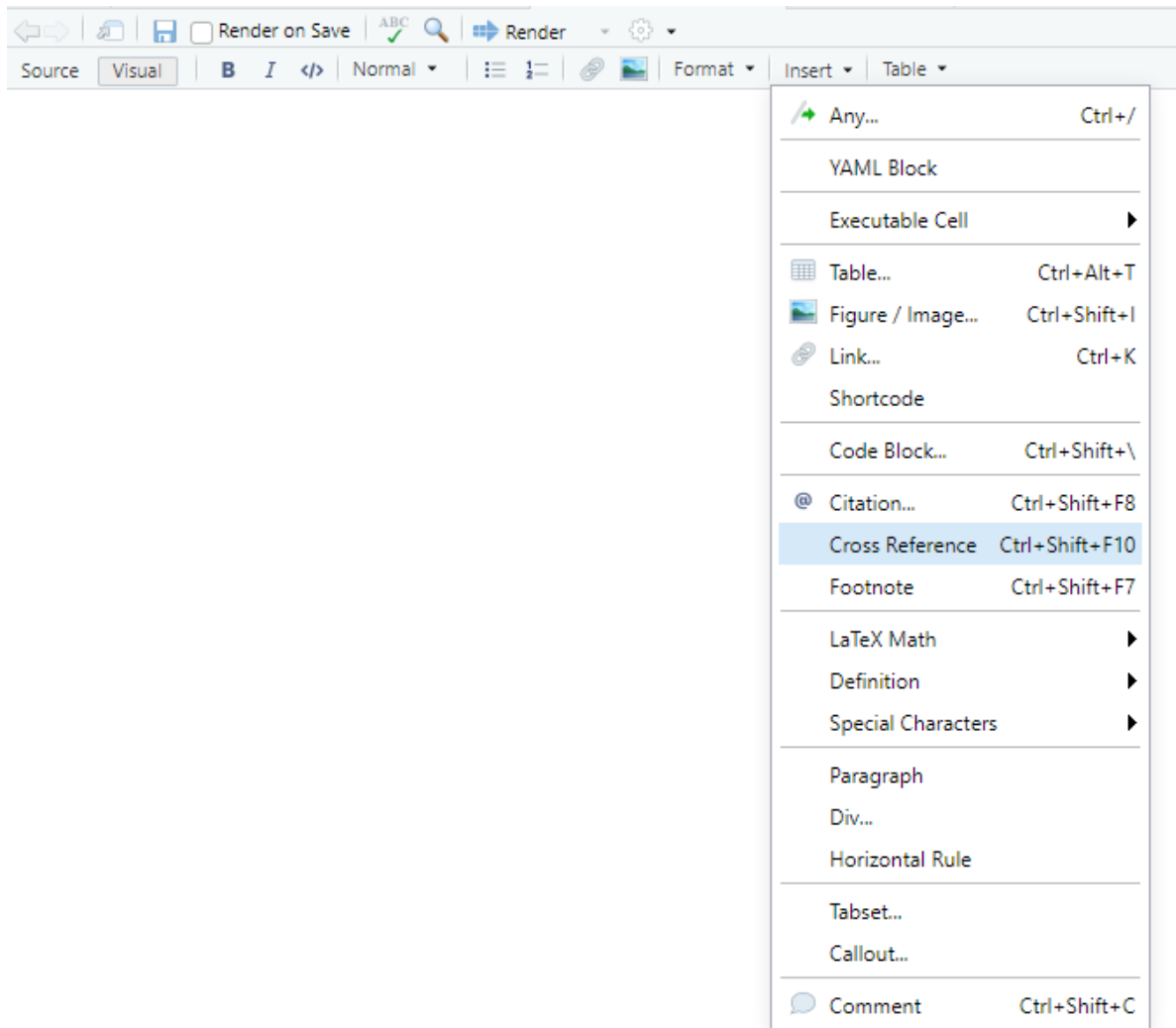
Characteristic	OR	95% CI	p-value
bmi	1.04	1.03, 1.04	<0.001
age	1.00	1.00, 1.01	0.003

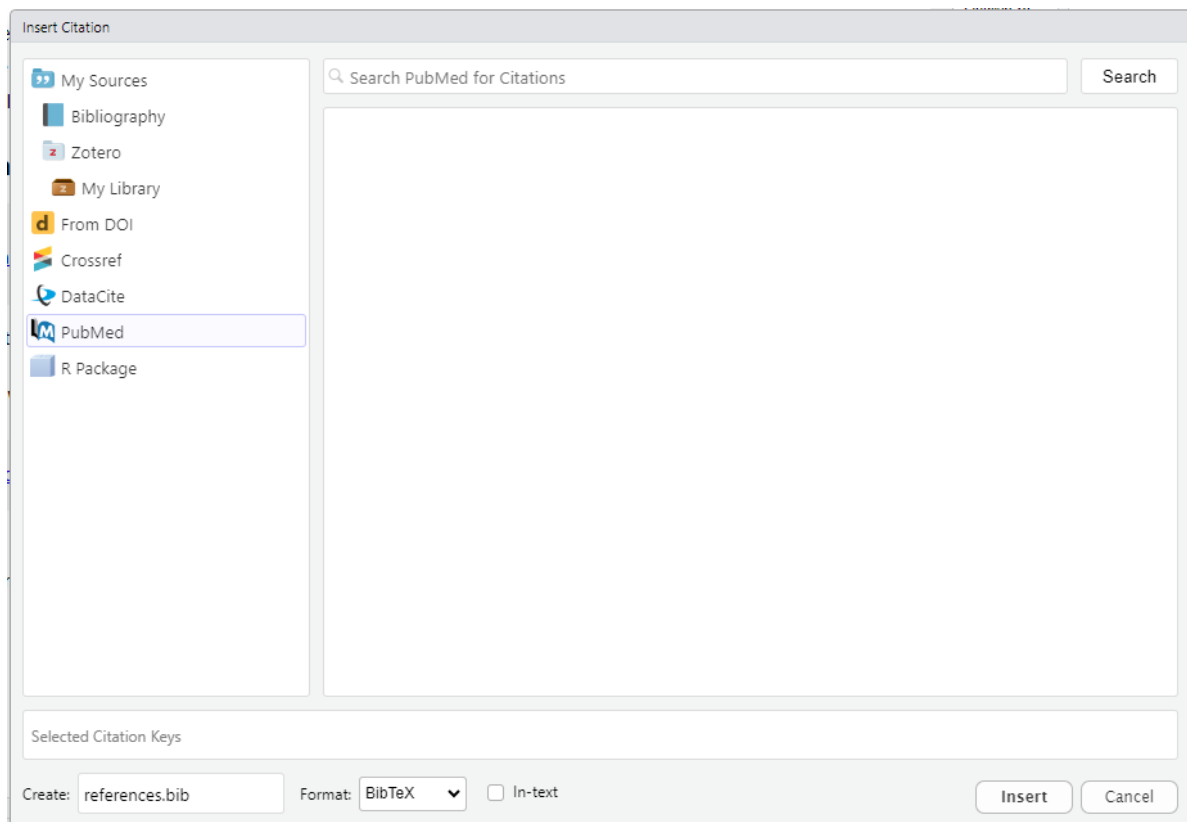
- Replace the placeholder text in bold with a cross-reference to the table ;

“The results of the logistic regression model are summarized in the formatted regression table, which is presented in **cross-reference**.”



## 0.7 Add references





## 0.8 Finalise your MS Word report

```

1  ---
2  title: My outbreak report
3  author: Helene Langet
4  date: last-modified
5  date-format: long
6
7  format:
8    docx:
9      reference-doc: swisstph_template.docx
10     toc: true
11     number-sections: true
12     toc-depth: 2
13  echo: false
14  warning: false
15  ---

```

23

- ☐ Create your own template and apply it to your MS Word rendered report.