

# RMarkdown Template File

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April 06, 2025

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## Introduction

This document outlines the methodology for data cleaning, exploration, and visualization. It is structured to ensure transparency and reproducibility of all analyses.

**Methodology** Briefly describe the methods used in the project, including data sources, cleaning steps, and techniques applied to handle missing or inconsistent data.

**Math Equations with LaTeX** Here is an inline equation:  $E = mc^2$ .

And here is a displayed equation:

$$f(x) = \int_{-\infty}^{\infty} e^{-x^2} dx \quad (1)$$

Both inline and block math can be rendered seamlessly with LaTeX.

You can also create multiline equations with alignment:

$$\begin{aligned} a &= b + c \\ d &= e + f \end{aligned} \quad (2)$$

**Setup Packages** The first step in any data analysis is to load the data, options, and packages. This section installs and loads the necessary packages, sets up the output paths, and defines chunk options for the RMarkdown document.

**Loading Data** The following code loads the data from various sources, including CSV files, Excel files, and shapefiles. The `here` package is used to create file paths that are relative to the project directory, ensuring that the code is portable and reproducible.

```
# Example of loading data from a CSV file
# raw_data <- read_csv(here("Data", "Raw", "data_file.csv"))
# raw_data <- read_excel(here("Data", "Raw", "data_file.xlsx"))
# raw_data <- read_sf(here("Data", "Raw", "data_file.shp"))
```

**Cleaning Data** The data cleaning process involves several steps to ensure the data is in a suitable format for analysis. This includes handling missing values, correcting data types, and removing duplicates.

```
# cleaning the data using dplyr and tidyr
```

```
# cleaned_data <- raw_data %>%
# clean_names() %>%
# mutate(column_name = as_factor(column_name)) %>%
# mutate(date_column = as.Date(date_column, format = "%Y-%m-%d")) %>%
# mutate(numeric_column = as.numeric(numeric_column)) %>%
# mutate(accross(everything(), ~str_squish(.))) %>% =
# drop_na()
```

**Data Exploration** Data exploration is a crucial step in understanding the dataset and identifying patterns or anomalies. This section includes summary statistics, visualizations, and any other relevant analyses to gain insights into the data.

```
# Explore the cleaned data using basic summaries:
# glimpse(cleaned_data)
# summary(cleaned_data)
# str(cleaned_data)
```

**Data Visualization** Data visualization is an essential part of data analysis, allowing for the communication of findings in a clear and effective manner. This section includes various plots and charts to illustrate key insights from the data.

```
# Example of creating a summary table
# summary_table <- cleaned_data %>%
# group_by(group_var) %>%
# summarise(mean_value = mean(value_var, na.rm = TRUE)) %>%
# ungroup() %>%
# kable() %>%
# kable_styling(full_width = F, position = "left")

# save_kable(summary_table, file = here(output_path_tables, "summary_table.html"),
#   ),
# bootstrap_options = c("striped", "hover", "condensed"),
# full_width = F, position = "left")

# Example of visualization plot
data("diamonds")

# First Plot (p1) - Scatter plot with smoothing line
p1 <- ggplot(
  subset(diamonds, carat >= 2.2),
  aes(x = table, y = price, colour = cut)
) +
  geom_point(alpha = 0.7) +
```

```
geom_smooth(method = "loess", alpha = 0.05,
linewidth = 1, span = 1, formula = y ~ x) +
theme_bw() + # White background theme
theme(
  plot.title = element_text(hjust = 0.5), # Center the title
  legend.position = "top", # Position the legend at the top
  text = element_text(size = 10), # Set text size
  legend.key.size = unit(0.5, "cm") # Legend key size
)

# Second Plot (p2) - Histogram of depth for specific carat ranges
p2 <- ggplot(
  subset(diamonds, carat > 2.2 & depth > 55 & depth < 70),
  aes(x = depth, fill = cut)
) +
  geom_histogram(colour = "black",
  binwidth = 1, position = "dodge") + # Dodged histogram
  theme_bw() + # White background theme
  theme(
    plot.title = element_text(hjust = 0.5), # Center the title
    legend.position = "top", # Position the legend at the top
    text = element_text(size = 10), # Set text size
    legend.key.size = unit(0.5, "cm") # Legend key size
  )

p1_npg <- p1 + scale_color_npg()
p2_npg <- p2 + scale_fill_npg()
grid.arrange(p1_npg, p2_npg, ncol = 2)

# Example of saving a plot
# ggsave(filename = here(output_path_images, "plot_name.png"),
# plot = last_plot(), width = 6, height = 4)
```

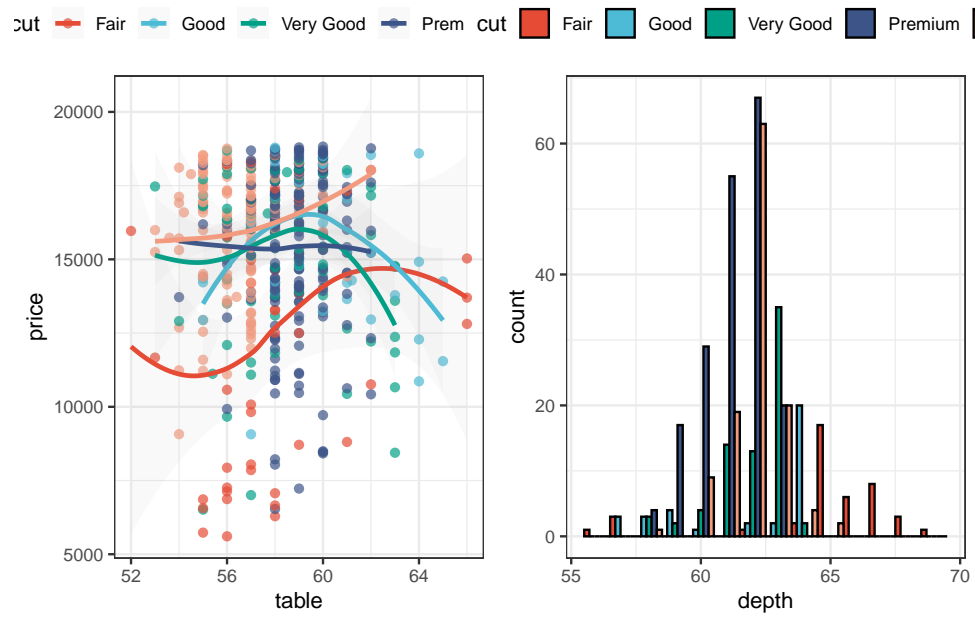


Figure 1: Example of a scatter plot

## Literature Cited

Citing the packages and data used in the analysis is important for reproducibility and transparency. The following code generates a bibliography of all loaded packages. Items can be cited directly within the documentation using the syntax `@key` where key is the citation key in the first line of the entry, e.g., R Core Team (2024), Wickham et al. (2024), Wickham (2023), Müller (2020). To put citations in parentheses, use `[@key]` instead.

Müller, K. (2020). *Here: A simpler way to find your files*. <https://here.r-lib.org/>

R Core Team. (2024). *R: A language and environment for statistical computing*. R Foundation for Statistical Computing. <https://www.R-project.org/>

Wickham, H. (2023). *Tidyverse: Easily install and load the tidyverse*. <https://tidyverse.tidyverse.org>

Wickham, H., Chang, W., Henry, L., Pedersen, T. L., Takahashi, K., Wilke, C., Woo, K., Yutani, H., Dunnington, D., & van den Brand, T. (2024). *ggplot2: Create elegant data visualisations using the grammar of graphics*. <https://ggplot2.tidyverse.org>