

# Lab 02 Week 01 Worksheet

## R Functions Glossary

This glossary provides an overview of key R functions used in **Lab 09**, explaining their **purpose** and **general use** in data processing and manipulation.

### Data Import & Management

| Function                                    | Description                        | Example                                  |
|---|------------------------------------|--|
| <code>read_csv()</code>                     | Reads a CSV file into a tibble     | <code>read_csv("data.csv")</code>        |
| <code>setwd()</code> / <code>getwd()</code> | Sets or gets the working directory | <code>setwd("path/to/folder")</code>     |
| <code>write_csv()</code>                    | Writes a data frame to a CSV file  | <code>write_csv(df, "output.csv")</code> |

### Data Wrangling (dplyr)

| Function   | Description                        | Example   |
|--|------------------------------------|---|
| <code>filter()</code>                              | Filters rows based on condition(s) | <code>filter(unit_price &gt; 0)</code>  |
| <code>mutate()</code>                              | Adds or transforms variables       | <code>mutate(total = unit_price * unit_quantity)</code>                             |
| <code>arrange()</code>                             | Sorts rows by variables            | <code>arrange(shopper_id)</code>  |
| <code>drop_na()</code>                             | Removes rows with missing values   | <code>drop_na()</code>  |
| <code>group_by()</code> + <code>summarize()</code> | Groups data and summarizes it      | <code>group_by(shopper_id) %&gt;% summarize(avg_items = mean(unit_quantity))</code> |
| <code>distinct()</code>                            | Selects distinct rows              | <code>distinct(shopper_id, store_id)</code>   |
| <code>select()</code>                              | Selects columns                    | <code>select(total_spent, avg_items)</code>   |

### Joins

| Function                  | Description                               | Example                                       |
|---------------------------|---|---|
| <code>inner_join()</code> | Keeps only matching rows from both tables | <code>inner_join(df1, df2, by = "key")</code> |

| Function                  | Description                         | Example                                       |
|---------------------------|-------------------------------------|---|
| <code>left_join()</code>  | Keeps all rows from the left table  | <code>left_join(df1, df2, by = "key")</code>  |
| <code>right_join()</code> | Keeps all rows from the right table | <code>right_join(df1, df2, by = "key")</code> |
| <code>full_join()</code>  | Combines all rows from both tables  | <code>full_join(df1, df2, by = "key")</code>  |

## Summary Statistics

| Function                        | Description                            | Example  |
|---------------------------------|--|--|
| <code>datasummary_skim()</code> | Summary of numeric or categorical data | <code>datasummary_skim(df, type = "numeric")</code>          |
| <code>datasummary()</code>      | Custom summary table                   | <code>datasummary(var1 + var2 ~ Mean + SD, data = df)</code> |

## Visualizations

| Function                    | Description                           | Example  |
|-----------------------------|---------------------------------------|--|
| <code>ggpairs()</code>      | Pairwise scatter/density plots        | <code>ggpairs(df %&gt;% select(x, y, z))</code>                |
| <code>fviz_nbclust()</code> | Plots to determine number of clusters | <code>fviz_nbclust(scaled_data, kmeans, method = "wss")</code> |

## Clustering

| Function              | Description                 | Example   |
|-----------------------|-----------------------------|---|
| <code>scale()</code>  | Standardizes variables      | <code>scale(df)</code>                              |
| <code>kmeans()</code> | Performs k-means clustering | <code>kmeans(data, centers = 3, nstart = 25)</code> |

## Other Tools & Utilities

| Function                      | Description   | Example                                       |
|-------------------------------|---|---|
| <code>length(unique())</code> | Counts distinct elements  | <code>length(unique(df\$shopper_id))</code>   |
| <code>quantile()</code>       | Returns quantiles   | <code>quantile(df\$total_spent, 0.999)</code> |
| <code>set.seed()</code>       | Use this with <code>kmeans()</code> so you can reproduce your results | <code>set.seed(123)</code>                    |

## Helpful Tips

- Use `left_join()` when you want to *keep all rows* from your base dataset.
- Use `group_by()` with `summarise()` to collapse and summarize grouped data.
- Standardize your variables before clustering using `scale()`.
- Use `fviz_nbclust()` to help determine the best number of clusters.
- Always inspect your summary statistics and visualize your data before running models.

**Remember:** Each step in data cleaning, joining, and clustering depends on your research question. Document your decisions clearly!