

Excercise3

April 21, 2025

```
[1]: library(tidyverse)
      library(cluster)
      library(gridExtra)
```

Attaching core tidyverse packages

```
tidyverse 2.0.0
  dplyr      1.1.4    readr      2.1.5
  forcats    1.0.0    stringr   1.5.1
  ggplot2    3.5.2    tibble    3.2.1
  lubridate  1.9.4    tidyr     1.3.1
  purrr      1.0.4
```

Conflicts

```
tidyverse_conflicts()
  dplyr::filter() masks stats::filter()
  dplyr::lag()    masks stats::lag()
Use the conflicted package
(<http://conflicted.r-lib.org/>) to force all conflicts to
become errors
```

Attaching package: 'gridExtra'

The following object is masked from 'package:dplyr':

combine

```
[2]: set.seed(1)
```

```
[3]: happiness <- read_csv("/home/asus/content/Notes/Semester 4/FDN Lab/Experiments/
  ↪Experiment 6/archive(11)/2019.csv")
```

Rows: 156 Columns: 9

Column specification

Delimiter: ","

`chr` (1): Country

`dbl` (8): Overall rank, Score, GDP per capita, Social support, Healthy life e...

Use ``spec()`` to retrieve the full column specification for this data.

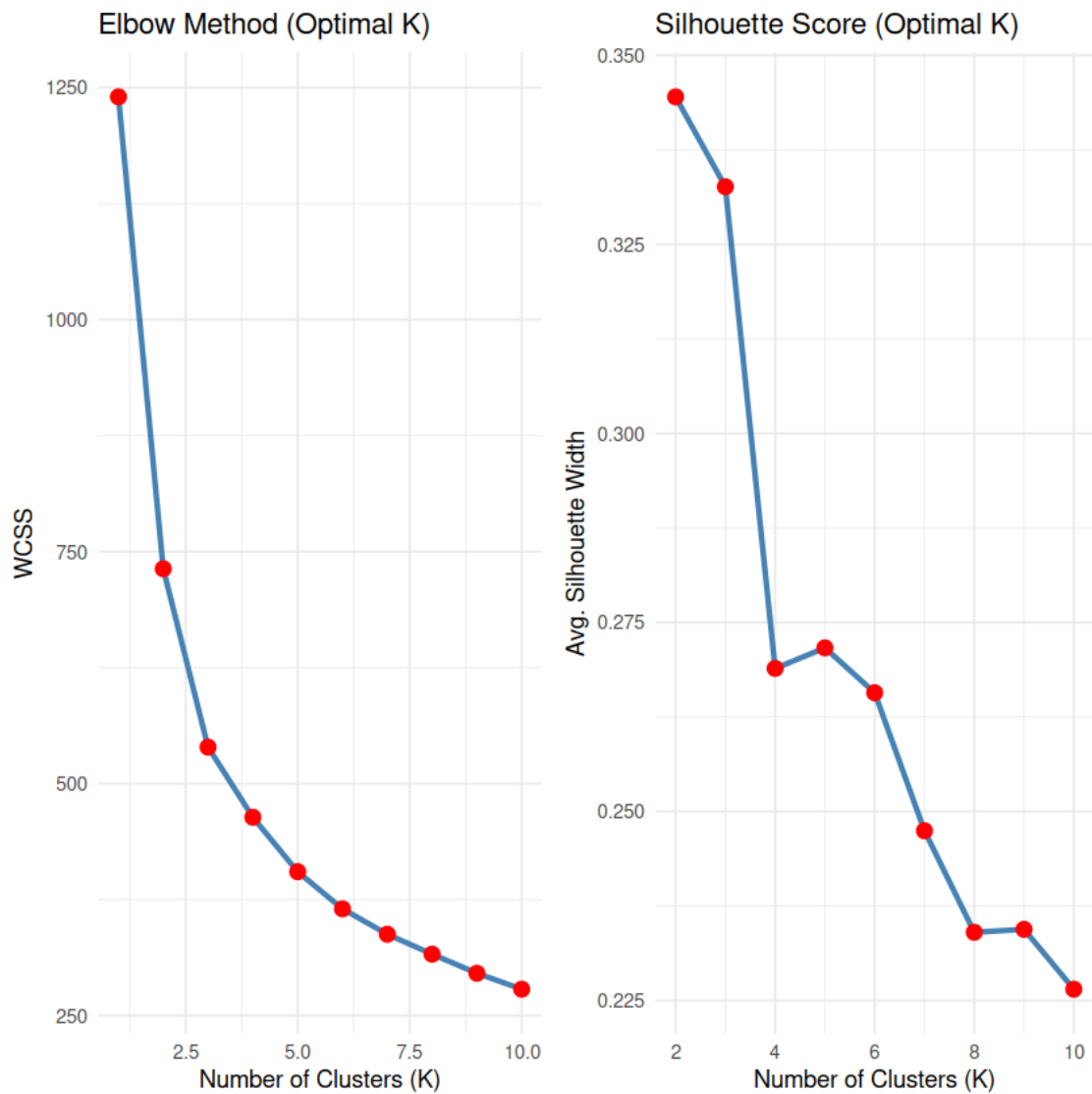
Specify the column types or set ``show_col_types = FALSE`` to quiet this message.

```
[4]: numeric_data <- happiness %>%  
  select(  
    "Overall rank", "Score", "GDP per capita", "Social support", "Healthy life_␣  
    ↪expectancy", "Freedom to make life choices", "Generosity", "Perceptions of_␣  
    ↪corruption"  
  ) %>%  
  scale()  
  
[5]: rownames(numeric_data) <- happiness$`Country`  
  
[6]: wcss <- map_dbl(1:10, ~ kmeans(numeric_data, ., nstart = 25)$tot.withinss)  
  
[7]: avg_sil <- map_dbl(2:10, ~ {  
  km <- kmeans(numeric_data, ., nstart = 25)  
  silhouette_score <- silhouette(km$cluster, dist(numeric_data))  
  mean(silhouette_score[, 3])  
})  
  
[8]: elbow_plot <- ggplot(data.frame(K = 1:10, WCSS = wcss), aes(K, WCSS)) +  
  geom_line(color = "steelblue", size = 1.2) +  
  geom_point(color = "red", size = 3) +  
  labs(title = "Elbow Method (Optimal K)", x = "Number of Clusters (K)", y =_␣  
  ↪"WCSS") +  
  theme_minimal()  
  
silhouette_plot <- ggplot(data.frame(K = 2:10, Silhouette = avg_sil), aes(K,_␣  
  ↪Silhouette)) +  
  geom_line(color = "steelblue", size = 1.2) +  
  geom_point(color = "red", size = 3) +  
  labs(title = "Silhouette Score (Optimal K)", x = "Number of Clusters (K)", y_␣  
  ↪= "Avg. Silhouette Width") +  
  theme_minimal()
```

```
grid.arrange(elbow_plot, silhouette_plot, ncol = 2)
```

Warning message:

"Using `size` aesthetic for lines was deprecated in ggplot2 3.4.0.
Please use `linewidth` instead."



```
[9]: k2 <- kmeans(numeric_data, centers = 2, nstart = 25)
k3 <- kmeans(numeric_data, centers = 3, nstart = 25)
k4 <- kmeans(numeric_data, centers = 4, nstart = 25)
k5 <- kmeans(numeric_data, centers = 5, nstart = 25)
```

```
[10]: happiness$Cluster_K2 <- as.factor(k2$cluster)
happiness$Cluster_K3 <- as.factor(k3$cluster)
```

```
happiness$Cluster_K4 <- as.factor(k4$cluster)
happiness$Cluster_K5 <- as.factor(k5$cluster)
```

```
[11]: plot_cluster_means <- function(km_result, title) {
  centers <- as.data.frame(km_result$centers)
  centers$Cluster <- factor(rownames(centers))

  centers_long <- centers %>%
    pivot_longer(cols = -Cluster, names_to = "Feature", values_to = "Mean_Value")

  ggplot(centers_long, aes(x = Feature, y = Mean_Value, fill = Cluster)) +
    geom_bar(stat = "identity", position = "dodge") +
    labs(title = title, y = "Standardized Mean Value", x = "") +
    theme_minimal() +
    theme(axis.text.x = element_text(angle = 45, hjust = 1)) +
    scale_fill_brewer(palette = "Set1")
}
```

```
[12]: p2 <- plot_cluster_means(k2, "Cluster Means (k=2)")
p3 <- plot_cluster_means(k3, "Cluster Means (k=3)")
p4 <- plot_cluster_means(k4, "Cluster Means (k=4)")
p5 <- plot_cluster_means(k5, "Cluster Means (k=5)")

grid.arrange(p2, p3, p4, p5, ncol = 2)
```

