VIETNAMESE-GERMAN UNIVERSITY COMPUTER SCIENCE AND ENGINEERING



Compulsory Elective Subject: Data Analysis in High Dimensions

Group 4 Project Report

Hierarchical Cluster Analysis: Movie Genres Preferences

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Member list & Workload

Our team has 4 members and we together decide to split the total workload into proper parts equally. Each member is responsible for the work as below:

No.	Full name	ID	Percentage of work
1	Vu Hoang Tuan Anh	18812	100%
2	Tran Kim Hoan	18810	100%
3	Ba Nguyen Quoc Anh	17965	100%
4	Nguyen Hoang Hai Nam	17035	100%

• Vu Hoang Tuan Anh (ID: 18812)

- Collecting and preparing data
- Applying hierarchical cluster model with dummy variables
- Analysising results, plotting dendrogram and histogram, distribution diagram
- Using Gap Statistics method to find the optimal number of clusters
- Writing the report (Section 4. Hierarchical clustering with Dummy variables)

• Tran Kim Hoan (ID: 18810)

- Conducting the survey
- Collecting and preparing data
- Applying hierarchical cluster model with dummy variables
- Analysising results, plotting dendrogram, histogram
- Writing and finalizing the report (Section 3. Data)

• Ba Nguyen Quoc Anh (ID: 17965)

- Collecting data
- Applying hierarchical clustering model with Gower's distance
- Analysising results, plotting dendrogram, histogram
- Writing the report (Section 5. Hierarchical clustering with mixed-type variables + Section 6. Conclusion)

• Nguyen Hoang Hai Nam (ID: 17035)

- Conducting the survey
- Collecting data
- Applying hierarchical cluster model with dummy variables
- Writing the report (Section 1. Project Objective + Section 2. Overview of Hierarchical clustering methodology)

Group 4 Project Report Page 2/37

${\bf Contents}$

1	Pro	ject Objective	4
2	erview of Hierarchical clustering methodology	4	
3	Dat	za	6
	3.1	Attributes Identification and Selection	6
	3.2	Data Collection	6
	3.3	Data Preparation	8
4	Hie	rarchical clustering with Dummy variables	10
	4.1	Problem	10
	4.2	Dummy variables	10
	4.3	Applying the Hierarchical clustering model	11
	4.4	Finding the optimal number of clusters	13
		4.4.1 Gap statistics method	13
	4.5	Result	14
		4.5.1 Dendrogram	14
		4.5.2 Histogram and observing patterns	15
	4.6	Advantages and Disadvantages: Dummy variables	20
		4.6.1 Advantages	20
		4.6.2 Disadvantages	20
5	Hie	rarchical clustering with mixed-type variables	21
	5.1	Problem	21
	5.2	Gower's Distance	21
	5.3	Code Explaination	21
	5.4	Results	24
		5.4.1 Dendrogram	24
		5.4.2 Histograms and observing patterns	25
	5.5	Shortcomings	28
6	Cor	nclusion	29
	6.1	Accomplishment of Project Objectives	29
	6.2	Future Work	29
7	App	pendix	30
	7.1	Project Repository on GitHub	30
	7.2	Survey Questions	30
	7.3	R Code	30

Group 4 Project Report Page 3/37

1 Project Objective

We would like to group together users with similar viewing patterns in order to recommend similar content (genres). By asking different questions about age, genre, hobbies, we can collect data and conclude some insights.

2 Overview of Hierarchical clustering methodology

Hierarchical clustering is an algorithm that groups similar objects into clusters. The clusters are distinct from each other, and the objects within each cluster are broadly similar to each other.

Hierarchical clustering is an unsupervised learning technique. This means that a model does not have to be trained, and there is no need for a "target" variable.

There are two common categories of Hierarchical clustering, agglomerative and divisive.

 Agglomerative clustering starts in it individual clusters, then merges pair of clusters and continuing until all clusters have been merge in one huge clusters.

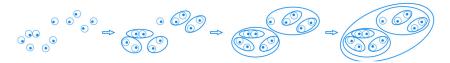


Figure 1: Agglomerative clustering

• Divisive clustering is the inverse approach of agglomerative clustering which starts in one cluster then splits into smaller cluster base on their difference.

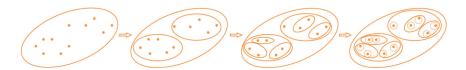


Figure 2: Divisive clustering

In our project, we only focus on agglomerative clustering.

The method of hierarchical clustering starts by treating each data point as a separate cluster and then iteratively combines the closest clusters until a stopping criterion is reached. The clusters are visually represented in a hierarchical tree called a dendrogram.

Group 4 Project Report Page 4/37

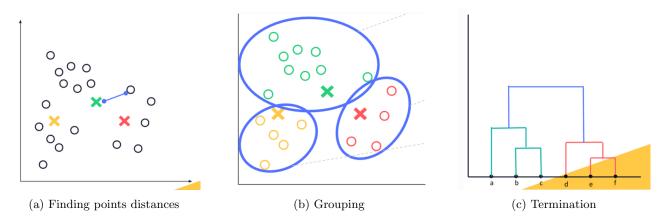


Figure 3: Hierarchical clustering steps

- 1. **Finding points distances:** Calculate the distance between each point of observation. There are various mathematical algorithms to apply: "euclidean", "maximum", "manhattan", "canberra", "binary", "minkowski"
- 2. **Grouping points (clustering)** Calculate the distance between each cluster. There are various mathematical algorithms to apply: "Ward", "single", "complete", "average", "mcquitty", "median", "centroid". After that, choose the optimal number of clusters.
- 3. **Termination** Plotting the final dendrogram and starting the analysis process based on attributes of each cluster.

Group 4 Project Report Page 5/37

3 Data

According to the objective of our project, we require data for our analysis. Our data is derived from our online survey conducted over a week. All collected data is accessible only to authorized research team members.

3.1 Attributes Identification and Selection

Initially, we brainstorm and identify key attributes which will be included in our analysis such as user demographics, user behavior and user preferences. After discussing their relevance and potential impact on the user's movie genre preferences, we eventually determine seven key attributes essential for our analysis: age, gender, working/learning area, preferred film genre, factor influencing genre choice, frequency of movie watching, source of film viewing.

3.2 Data Collection

• Step 1: Create a survey

We create a survey titled 'Movie Genre Preferences' to collect data via Google Forms. In the form, we focus on seven key attributes that we determined before. (as discussed in Section 2.1).

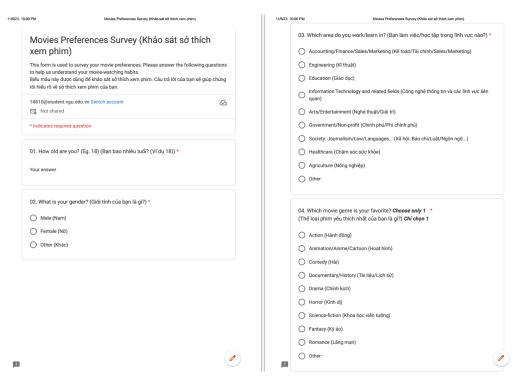


Figure 4: Movie genre preferences survey image (1)

Group 4 Project Report Page 6/37

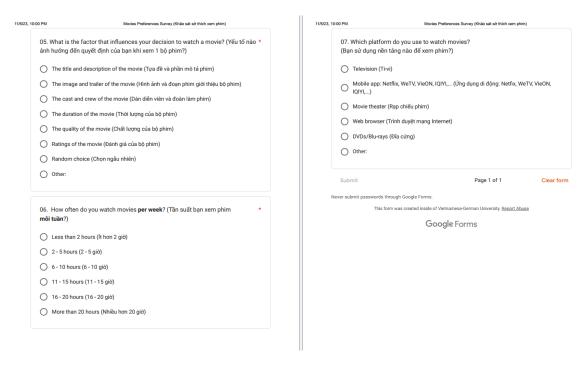


Figure 5: Movie genre preferences survey image (2)

• Step 2: Publish survey and receive responses

After completing the form, we distribute it to individuals, specifically targeting those aged 16 and above. Subsequently, we received a total of 118 responses from the survey. Here is an image of the responses list:

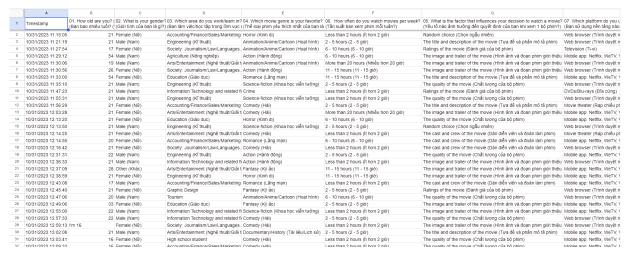


Figure 6: Image of the responses list

Group 4 Project Report Page 7/37

3.3 Data Preparation

This process is crucial in minimizing the potential for errors and inaccuracies that may arise during our data processing stage. By ensuring the data is accurately prepared and processed, we can enhance the reliability and validity of our analysis.

With a dataset of 118 responses, we come to the first crucial step for our data analysis: cleaning data.

• In terms of attributes, we convert them from question format to word/phrase format:

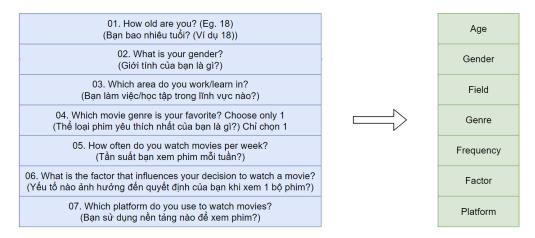


Figure 7: Image of attributes before and after

• We also substitute long words/phrases with shorter ones:



Figure 8: Image of the data before and after substitution

• Regarding working/learning area, we decided to combine "real estate" as a part of "business":

23	Female	Information Technology	23	Female	Information Technology
27	Female	Real estate	27	Female	Business
35	Male	Real Estate	35	Male	Business
20	Female	Arts/Entertainment	20	Female	Arts/Entertainment

Figure 9: Image of the data before and after combination

Group 4 Project Report Page 8/37

• We adjust responses that are not in the correct format into our anticipated format:

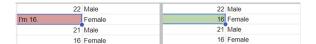


Figure 10: Image of the data before and after

• We remove responses that deviate from our expected range of values:



Figure 11: Image of the data example before and after elimination

• The entry "housewife" does not correspond to a specific work or learning area. Therefore, we exclude it from our dataset:

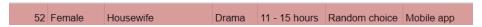


Figure 12: Image of the entry "housewife"

After the data cleaning process, we obtain the final dataset. This dataset has the correct format, concise words/phrases, and values within an appropriate range for all attributes, making it suitable for our analysis.



Figure 13: Image of the final data after data cleaning process

In the next step, we apply the hierarchical clustering method to our final dataset, using dummy variables and mixed-type variables.

Group 4 Project Report Page 9/37

4 Hierarchical clustering with Dummy variables

4.1 Problem

Regarding the objective of the project, users with similar viewing patterns need to be grouped together in a cluster, and in the previous section, we know that the dataset consists of both numerical and categorical data. Hence, it is impossible to calculate the distance between any 2 entries of the data frame by applying mathematical algorithms (Euclidean, Manhattan, Canberra, etc) directly since all variables are not numerical. To handle this situation, an encode is needed to convert variables from nominal to numerical type. (The conversion from ordinal variables to numerical variables is redundant since ordinal variables could be labeled with levels and then be scaled to behavior as numerical variables). That encoding is called **Dummy variables**, and this section will discuss the implementation of dummy variables for hierarchical clustering.

4.2 Dummy variables

- <u>Definition</u>: A Dummy (or Indicator) variable is an artificial variable created to represent a categorical variable with two or more distinct categories or levels.
- <u>Usage:</u> With dummy variables, all nominal variables could be converted to numerical. Hence, it is possible to apply mathematical algorithms to start the analysis process.
- Example: A nominal variable named "Genre" always takes one of 3 values "Action", "Comedy", and "Fantasy". After converting it into the dummy variable:
 - If the value of the variable is "Action", the indicator for "Genre.Action" attribute will be 1, and other attributes are 0's.
 - If the value of the variable is "Comedy", the indicator for "Genre.Comedy" attribute will be 1, and other attributes are 0's.
 - If the value of the variable is "Fantasy", the indicators for all attributes are 0's.

	Genre
ĺ	Action
ĺ	Comedy
ĺ	Fantasy

(a) Original "Genre" variable

	Genre.Action	Genre.Comedy
Action	1	0
Comedy	0	1
Fantasy	0	0

(b) "Genre" Dummy variable

Table 1: Using Dummy variables

 \rightarrow A variable which has a total of n different values will have new n-1 attributes after converting to a dummy variable, and the last value has the indicators for all attributes are 0's.

Group 4 Project Report Page 10/37

4.3 Applying the Hierarchical clustering model

- In this section, R programming language is used as the main language to implement the hierarchical clustering model.
- Dependencies (List of packages used):

Package name	Description	Version
base	base The R Base Package	
cluster	Finding Groups in Data	$\geq 2.1.4$
factoextra	clustering visualization	$\geq 1.0.7$
gplots	plotting data, dendrogram	$\geq 3.1.3$
ggplot2	draw distribution graph	$\geq 3.4.4$

(Source code link: https://github.com/vhtuananh020402/Group4_Data_analysis/blob/main/dummy_canberra_ward.r)

To start applying the hierarchical clustering model, first, the dataset is imported and stored as a data frame in R. The function na.omit() is necessary for omitting the missing value of the imported data frame.

```
# Read the data frame
df <- read.csv("data/clean_data_v2.csv")

# Omit the NA values of the data frame
df <- na.omit(df)</pre>
```

Then, we standardize (scale) the "Age" (numerical) variables. Since the "Frequency" variable is ordinal, it first needs to be ordered and labeled with levels, then converted to a numerical variable using **as.numeric()** function. After that, the "Frequency" (numerical) variable is scaled.

```
# Standardize the Age variable
df$Age_std <- scale(df$Age)

# Standardize the Frequency variable
df$Frequency <- factor(
df$Frequency,
order = TRUE,
levels = c("Less than 2 hours", "2 - 5 hours", "6 - 10 hours", "11 - 15 hours",
"16 - 20 hours", "More than 20 hours"))</pre>
```

Group 4 Project Report Page 11/37

```
df$Frequency_numeric <- as.numeric(factor(df$Frequency))
df$Frequency_std <- scale(df$Frequency_numeric)
```

The function *model.matrix()* is used to create a data frame matrix consisting of all numerical variables (all nominal variables "Genre", "Field", "Factor", "Gender", "Platform" are converted to dummy variables)

```
# Turn the nominal variables into dummy variables
df_dummy <- model.matrix(~ Age_std + Genre + Frequency_std + Field + Factor + Gender +
Platform, data = df)
```

Here we use Canberra algorithms to calculate the points distance and Ward's method to calculate the clusters distance. Then we can visualize the hierarchical clusters by using the function plot() to plot a dendrogram.

```
# Calculate the points distance
point_dist <- dist(df_dummy, method = "canberra")  # Using Canberra distance

# Hierarchical cluster analysis on the data frame
the c- helust(point_dist, method = "ward.D")  # Using Ward's method

# Plot the dendrogram
dend <- as.dendrogram(he)
plot(dend)</pre>
```

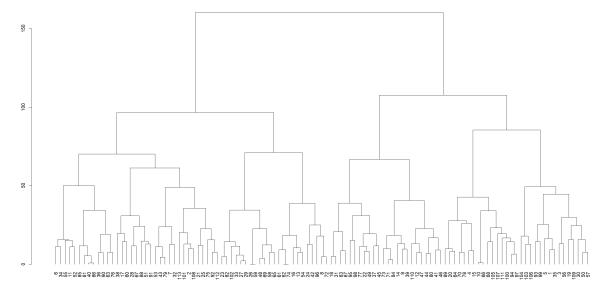


Figure 14: Dendrogram visualization

Group 4 Project Report Page 12/37

4.4 Finding the optimal number of clusters

After dendrogram visualization, as can be clearly seen that there are many possible numbers of clusters to choose from for analysis. Each different number of clusters will bring different outcomes, thus we need an exact optimal number of clusters, and there are some methods that help us to find it, **Gap Statistics** is one of the most well-known and easily understandable methods to find the optimal number of clusters, so this section will focus on it.

4.4.1 Gap statistics method

- This approach can be applied to any clustering method.
- The estimate of the optimal clusters will be the value that maximizes the gap statistic. This means that the clustering structure is far away from the random uniform distribution of points.
- R code implementation:

```
# ====== Gap statistic method to find the optimal number of cluster
   # Calculate Within-Cluster Dispersion (WCD) for the original data
   wss <- sum(hc$height)
   # Generate Random Data for Comparison
   set.seed(123)
                    # Set seed for reproducibility
   B <- 100
                    # Number of random datasets
   random_datasets <- lapply(1:B, function(i) matrix(runif(length(df_dummy)), ncol =</pre>
   ncol(df_dummy)))
   # Cluster the Random Data
   random_hcs <- lapply(random_datasets, function(random_data) {</pre>
11
     dist_matrix <- dist(random_data, method = dist_method[4])</pre>
12
     hclust(dist_matrix, method = hc_method[1])
   })
14
15
   # Calculate Within-Cluster Dispersion for Random Data
   wss_random <- sapply(random_hcs, function(random_hc) sum(random_hc$height))
17
18
   # Calculate Gap Statistic
   gap <- (log(wss_random) - log(wss)) + mean(log(wss_random) - log(wss))</pre>
20
21
   # Determine the Optimal Number of Clusters
   num_clusters <- 1 : 10 # Desired number of cluster domain</pre>
23
```

Group 4 Project Report Page 13/37

Gap Statistic Plot

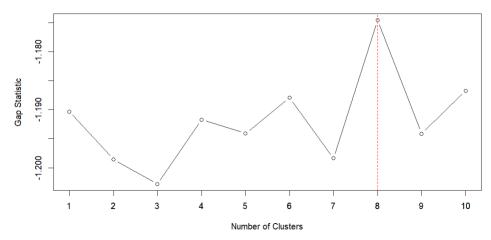


Figure 15: Gap Statistics diagram

→ From the above Gap Statistics diagram (range 1 : 10), the highest point which indicates the optimal number of clusters for the dendrogram is 8. Hence, we chose the number 8 for the number of clusters for the dendrogram.

4.5 Result

4.5.1 Dendrogram

```
# Draw the rectangle around each cluster in k clusters
k <- 8
rect.hclust(hc, k, border = 2:8)
```

Here is the final dendrogram with 8 clusters, each cluster has a rectangle around it. The observations in each cluster have a close relationship, which will be discussed right after this dendrogram.

Group 4 Project Report Page 14/37

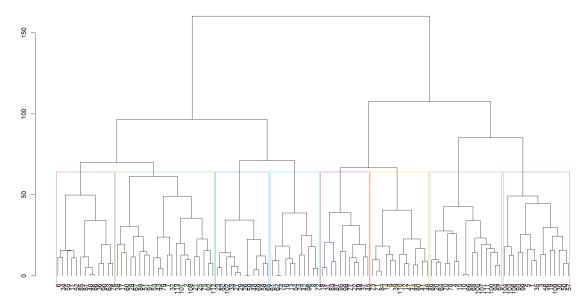


Figure 16: Age Distribution in each cluster

4.5.2 Histogram and observing patterns

"Age", "Genre", "Field", "Frequency", and "Factor" are considered the main attributes of the movie genre recommender system, then we observe the patterns of those attributes by histograms.

R code to plot the histograms for each cluster:

```
# Add the cluster assignments to the data frame
   df$Cluster <- factor(clusters)</pre>
   # Create a histogram of the Genre distribution in each cluster
   ggplot(df, aes(x = Genre)) +
     geom_histogram(stat = "count", fill = "lightblue", color = "black", linewidth = 0.8) +
     facet_wrap(~ Cluster) +
     theme(axis.text.x = element_text(angle = 45, hjust = 1)) +
     labs(title = "Genre Distribution in Each Cluster", x = "Genre", y = "Count")
10
   # Create a histogram of the Age distribution in each cluster
11
   ggplot(df, aes(x = Age)) +
     geom_histogram(stat = "count", fill = "orange", color = "black", linewidth = 0.8) +
13
     facet_wrap(~ Cluster) +
14
     theme(axis.text.x = element_text(angle = 45, hjust = 1)) +
     labs(title = "Age Distribution in Each Cluster", x = "Age", y = "Count")
16
```

Group 4 Project Report Page 15/37

```
17
    # Create a histogram of the Field distribution in each cluster
18
   ggplot(df, aes(x = Field)) +
19
     geom_histogram(stat = "count", fill = "red", color = "black", linewidth = 0.8) +
     facet_wrap(~ Cluster) +
21
     theme(axis.text.x = element_text(angle = 45, hjust = 1)) +
22
     labs(title = "Field Distribution in Each Cluster", x = "Field", y = "Count")
23
24
    # Create a histogram of the Frequency distribution in each cluster
25
   ggplot(df, aes(x = Frequency)) +
26
     geom_histogram(stat = "count", fill = "darkgrey", color = "black", linewidth = 0.8) +
27
     facet_wrap(~ Cluster) +
28
     theme(axis.text.x = element_text(angle = 45, hjust = 1)) +
29
     labs(title = "Frequency Distribution in Each Cluster", x = "Frequency", y = "Count")
30
31
   # Create a histogram of the Factor distribution in each cluster
32
   ggplot(df, aes(x = Factor)) +
33
     geom_histogram(stat = "count", fill = "lightgreen", color = "black", linewidth = 0.8)
34
     facet_wrap(~ Cluster) +
     theme(axis.text.x = element_text(angle = 45, hjust = 1)) +
36
     labs(title = "Factor Distribution in Each Cluster", x = "Factor", y = "Count")
37
```

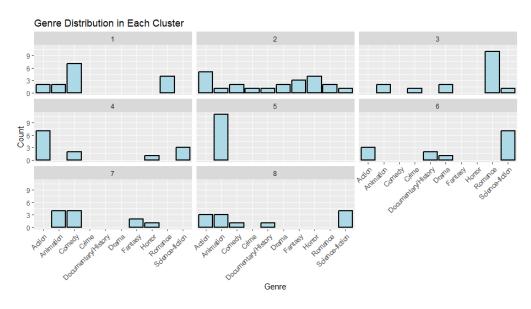


Figure 17: Genre Distribution in each cluster

Group 4 Project Report Page 16/37

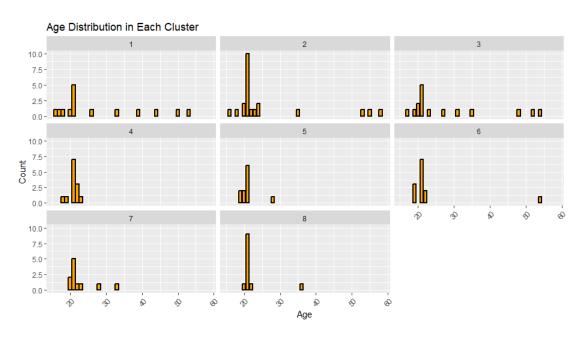


Figure 18: Age Distribution in each cluster

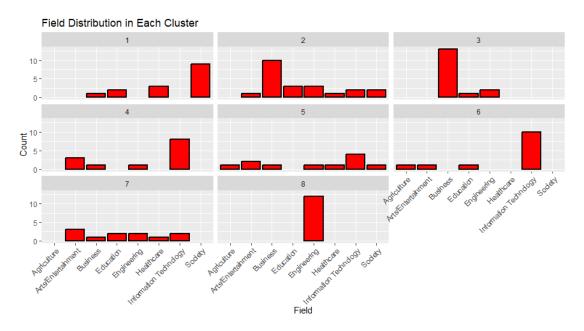


Figure 19: Field Distribution in each cluster

Group 4 Project Report Page 17/37

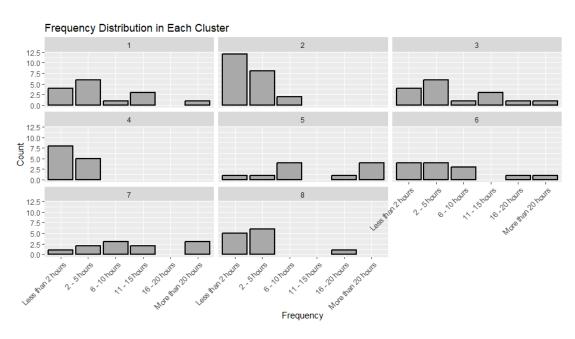


Figure 20: Frequency Distribution in each cluster

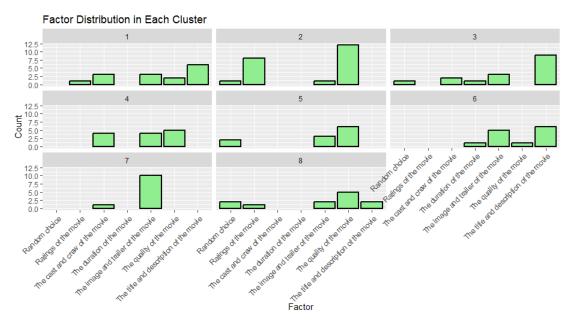


Figure 21: Factor Distribution in each cluster

From the histograms, here are conclusions for each cluster:

• <u>Cluster 1:</u> Comedy is the most favorite genre followed by Romance, with the majority of viewers being 21 years old and a small number ranged from 22 to 55 years old. Watching time is from 2 to 5 hours per

Group 4 Project Report Page 18/37

week. Society field is the highest choice here. "Title/Description" is the most popular reason for choosing and watching.

- Cluster 2: Action, Horror, and Fantasy are top-viewed genres (Romance and Drama are also considered). The viewers are mostly under 25 years old. The average watching time falls between "Less than 2 hours" and "2 5 hours" per week. The number is largely come from Business and evenly distributed in the Education and Engineering fields. "Ratings" and "Quality" are the two most important factors for picking movies.
- <u>Cluster 3:</u> Romance is the top-most choice of movie genre in this cluster. The viewers are divided into 2 main age groups: around 25 and above 45 years old. Watching times are "2 5 hours" and "11 15 hours". Business is the highest-picked field. "Title/Descriptions" is the most important factor.
- <u>Cluster 4:</u> Action and Science-fiction are the most favorite genre, with the majority of viewers being under 21 years old. Watching time is "Less than 5 hours" per week. IT field is the highest choice here, followed by Arts/Entertainment. "Cast/Crew", "Image/Trailer", and "Quality" are the most popular reasons for choosing and watching.
- <u>Cluster 5:</u> It is clearly seen that **Animation** is the only choice of movie genre in this cluster. The viewers are mostly 19 21 years old, and a small number is from 26 27 years old. The average watching time is pretty high and falls between "6 11 hours" and "More than 20 hours" per week. The number is largely come from IT and evenly distributed in the Agriculture, Arts/Entertainment, Business, Engineering, Healthcare, and Society fields. "Image/Trailer" and "Quality" are the two most important factors for picking movies.
- <u>Cluster 6</u>: Action and Science-fiction are the top-most choices of movie genres. Viewers are mostly around 20 and a small one is above 50 years old. Total watching time is less than 10 hours per week. IT is the highest-picked field. "Image/Trailer" and "Title/Descriptions" are the most important factor.
- Cluster 7: Animation along with Comedy is the most favorite genre followed by Fantasy and Horror, with the majority of viewers being under 35 years old. Watching time varies from less than 15 hours and more than 20 hours per week. The agriculture field is the highest choice here, and there exists evenly distribution of Education, Engineering, and IT fields. "Image/Trailer" is the most popular reason for choosing and watching.
- <u>Cluster 8:</u> Action, Animation, and Science-fiction are top-viewed genres. Viewers are mostly under 25 years old. The average watching time falls between "Less than 2 hours" and "2 5 hours" per week. The number is all come from Engineering fields. "Quality" and "Title/Description" are the two most important factors for picking movies.

Group 4 Project Report Page 19/37

4.6 Advantages and Disadvantages: Dummy variables

4.6.1 Advantages

- Nominal Data Handling: Hierarchical clustering algorithms typically work with numerical distances or similarities between data points. Dummy variables allow us to represent nominal variables numerically, enabling their incorporation into the clustering process.
- Information Preservation: Dummy variables can help preserve information about the nature of the data. By creating separate binary variables for different categories, we maintain the distinctions between nominal variables during clustering.

4.6.2 Disadvantages

- Unequal Distances: Dummy variables assume equal distances between points, which might not reflect the actual dissimilarities between them.
- Increased Noise: If the nominal variable has a large number of attributes with limited observations in each attribute, the dummy variables may introduce noise into the clustering process, leading to less reliable results.

Group 4 Project Report Page 20/37

5 Hierarchical clustering with mixed-type variables

5.1 Problem

One important aspect of processing information is to deal with different types of data, because not every attribute is of the same nature.

Given an example: A survey is conducted about a person's daily life. The considered factors are: Age, Gender, Job, and how Satisfactory they are. A quick observation shows that these values are not the same type. Age is measured in integer, which is a numerical value; Gender can be used as a binary value, with 0 (false) being female, and 1 (true) being male; Job is shown with many options (engineer, businessman, storekeeper, homemaker, ...), therefore it is a nominal value; and a scale of 1 to 10 to rate the person's satisfaction, which is an ordinal value. This is just some common types of data that we usually deal with, so knowing how to process mixed data can benefit us with a more realistic model.

This section will discuss Gower's Distance, which is one of the methods to deal with this problem.

5.2 Gower's Distance

Gower's Distance is a method for computing distances between two data points. The strength of this method comes from the fact that it is usable for other types of data beside numerical, which is more flexible than the common methods (Euclidean distance, Manhattan distance, ...). Another advantage is that Gower's Distance scales the ranges of data into between 0 and 1, with addition of allowing a user-defined weighting scheme. But for this project, an unweighted model is constructed.

The basic calculation of Gower's Distance is as follow. Data will be separated into two types: numerical and non-numerical.

With numerical, we can compute the data by the formular: |Difference| / Range

- Difference = Data[i] Data[i+1];
- Range is the difference between the maximum and the minimum data points.

With non-numerical values, we can compute by compare the data points. If they are identical, the distance will be 0, if not then the distance will be 1.

There are some packages that uses or have the option for Gower's calculation, and in this project "cluster" package is used, which contains function "daisy" that allows Gower as an option.

5.3 Code Explaination

(Source code link: https://github.com/vhtuananh020402/Group4_Data_analysis/blob/main/completed_gower_ward.r)

First, libraries and data frame will be imported. The function na.omit(df) is necessary for omitting missing values in the data frame.

Group 4 Project Report Page 21/37

```
library(factoextra) # clustering visualization
library(ggplot2) # draw distribution graph

# read data frame
df <- read.csv("data/clean_data_v2.csv")
# remove missing values in data frame
df <- na.omit(df)</pre>
```

Then, data will be processed by splitting into 3 types: numerical, nominal, and ordinal. Nominal is defined by using function lapply(df[nom_attr], as.factor) to change value type into factor. Ordinal is defined by using function factor(), which has the option order = TRUE, and level is from lowest to highest. Finally, process_dataset will contain the complete dataset for analysing steps.

```
# --- Data Preparation --- #
    # add into numerical value
   num_attr <- c("Age")</pre>
    # add into nominal value
   nom_attr <- c("Field", "Genre", "Factor")</pre>
   df[cat_attr] <- lapply(df[cat_attr], as.factor)</pre>
    # add into ordinal value
   ord_attr <- c("Frequency")</pre>
10
   df$Frequency <- factor(df$Frequency,</pre>
11
                                   order = TRUE,
                                   level = c("Less than 2 hours",
13
                                                "2 - 5 hours",
14
                                                "6 - 10 hours",
                                                "11 - 15 hours",
16
                                                "16 - 20 hours",
17
                                                "More than 20 hours"))
19
    # put everything into a complete data set
20
   process_dataset <- df %>% select(num_attr, ord_attr, cat_attr)
21
22
   head(process_dataset)
23
```

Function daisy() will calculate the dissimilarity matrix, with process_dataset as input, and gower is used as the calculation metric. After that, the hierarchical clustering model is built using hclust() function. Clustering

Group 4 Project Report Page 22/37

method ward.D is chosen because it brings the best result when plotting dendrogram.

```
# --- Calculation --- #

# calculate Gower's distance

gower_dist <- daisy(process_dataset, metric="gower")

# hierarchical clustering, using ward.D method

gower_hcl <- hclust(gower_dist, method = "ward.D")

# --- DENDROGRAM ---- #

# plot dendrogram

plot(gower_hcl, cex = 0.6)

# draw borders for the individual clusters

rect.hclust(gower_hcl, k = 7, border = 2:7)</pre>
```

Histograms are used to present the distribution of each attribute in each cluster. "k" represents the number of clusters that we want. The decision k=7 is made according to the previous method of building hierarchical clustering using Dummy Variable. For some unknown reasons, the functions that find the optimal number of clusters is unusable in this section.

```
# --- HISTOGRAM --- #
   # cut into k clusters
   k <- 7
   clusters <- cutree(gower_hcl, k)</pre>
   # add the cluster assignments to the data frame
   df\$Cluster <- factor(clusters)</pre>
   # histogram of Genre distribution in each cluster
9
   ggplot(df, aes(x = Genre)) +
10
       geom_histogram(stat = "count", fill = "lightblue", color = "black", linewidth = 0.8)
11
       facet_wrap(~ Cluster) +
12
       theme(axis.text.x = element_text(angle = 45, hjust = 1)) +
       labs(title = "Genre Distribution in Each Cluster", x = "Genre", y = "Count")
14
15
   # histogram of Field distribution in each cluster
   ggplot(df, aes(x = Field)) +
```

Group 4 Project Report Page 23/37

```
geom_histogram(stat = "count", fill = "red", color = "black", linewidth = 0.8) +
       facet_wrap(~ Cluster) +
19
       theme(axis.text.x = element_text(angle = 45, hjust = 1)) +
20
       labs(title = "Field Distribution in Each Cluster", x = "Field", y = "Count")
22
   # histogram of Factor distribution in each cluster
23
   ggplot(df, aes(x = Factor)) +
24
       geom_histogram(stat = "count", fill = "lightgreen", color = "black", linewidth =
25
       0.8) +
       facet_wrap(~ Cluster) +
26
       theme(axis.text.x = element_text(angle = 45, hjust = 1)) +
27
       labs(title = "Factor Distribution in Each Cluster", x = "Factor", y = "Count")
28
29
    # histogram of Age distribution in each cluster
30
   ggplot(df, aes(x = Age)) +
31
       geom_histogram(stat = "count", fill = "orange", color = "black", linewidth = 0.8) +
32
       facet_wrap(~ Cluster) +
33
       theme(axis.text.x = element_text(angle = 45, hjust = 1)) +
34
       labs(title = "Age Distribution in Each Cluster", x = "Age", y = "Count")
36
   # histogram of Frequency distribution in each cluster
37
   ggplot(df, aes(x = Frequency)) +
38
       geom_histogram(stat = "count", fill = "darkgrey", color = "black", linewidth = 0.8)
39
       facet_wrap(~ Cluster) +
40
       theme(axis.text.x = element_text(angle = 45, hjust = 1)) +
       labs(title = "Frequency Distribution in Each Cluster", x = "Frequency", y = "Count")
42
```

5.4 Results

5.4.1 Dendrogram

The dendrogram shows the relations between all of the observations, with each cluster is shown by colored lines. With many tries and many different methods, we decided to keep this result because this is more evenly distributed model than other versions.

Group 4 Project Report Page 24/37

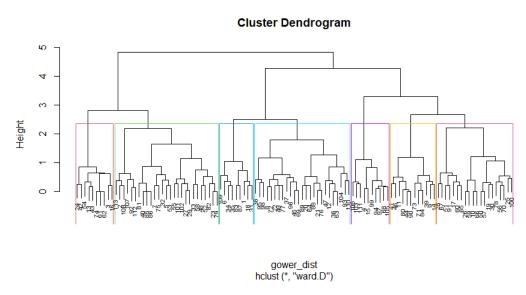


Figure 22: Result visualized by Dendrogram

5.4.2 Histograms and observing patterns

We only obverse the histograms of the following attributes: Age, Genre, Field, Frequency and Factor. Gender and Platform are useful for unifying the dataset, but those are not considered to be the attribute that contributing to the recommended system in real-world practices. Therefore, we omit them when graphing the histograms.

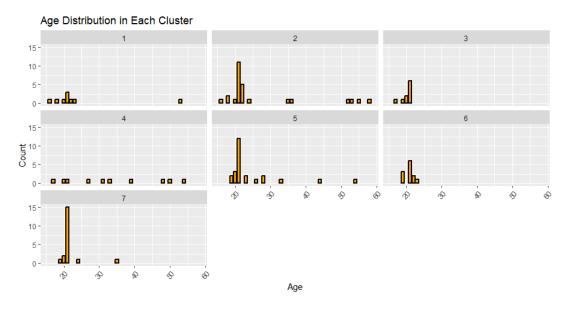


Figure 23: Age distribution in each cluster visualized by Histogram

Group 4 Project Report Page 25/37

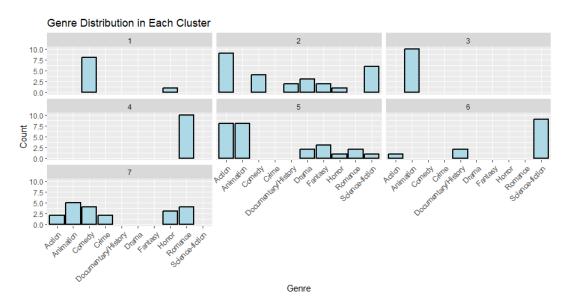


Figure 24: Genre distribution in each cluster visualized by Histogram

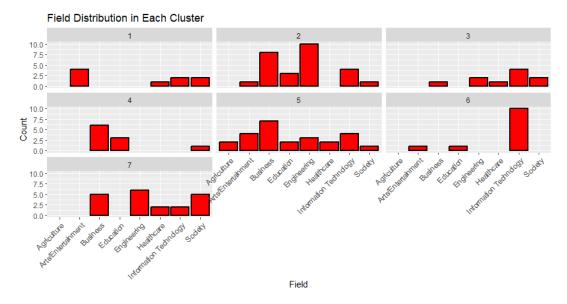


Figure 25: Field distribution in each cluster visualized by Histogram

Group 4 Project Report Page 26/37

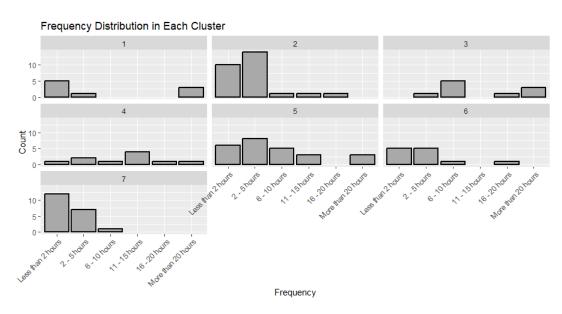


Figure 26: Frequency distribution in each cluster visualized by Histogram

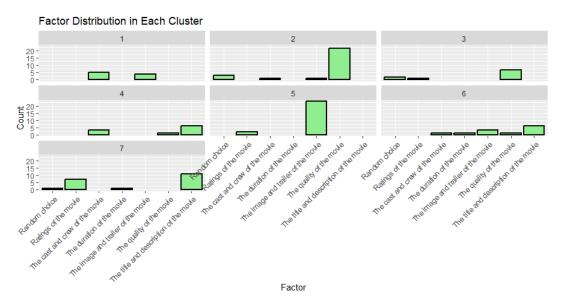


Figure 27: Factor distribution in each cluster visualized by Histogram

The followings are concluded from the histogram for each cluster:

Cluster 1: Mostly from under 25 years old. Comedy is the most watched genre, with watching time falls between "Less than 2 hours" and "Over 20 hours". Art/Entertainment field being the highest choice here, with IT and Society being the lesser. "Cast/Crew" and "Image/Trailer" are the reasons for choosing and watching.

Group 4 Project Report Page 27/37

Cluster 2: The majority are 21 years old, with Action and Sci-fi are the most viewed. Comedy, Documentary/History, Drama, Fantasy and Horror are also considered. Watching time is from 2 to 5 hours. They are largely come from Engineering and Business. "Quality" is the most important factor for picking movies.

Cluster 3: The majority is 21 years old, with Animation being the only choice for genre. "6 - 10 hours" and "more than 20 hours" are the standout choices. Engineering, IT and Society are evenly distributed fields, with IT being the highest choice. "Quality" is the most important factor.

Cluster 4: Ages are evenly distributed across the range. Romance is the only viewed genre here. Watching time is varied, with "11 – 15 hours" being the highest choice. Business is the most picked field here, and they consider "Cast/Crew" and "Title/Description" when choosing movies.

Cluster 5: Mostly 21 years old, with Action and Animation are the highest viewed genre here, also with somewhat evenly distributed between Drama, Fantasy, Romance. Watching time is largely from "Less than 2 hours" to "6 – 10 hours" range. Business is the most choice for field, but the occupation is distributed across the range. "Image/Trailer" is the most important factor.

Cluster 6: Around 20 years of age. Science-fiction is the most viewed genre, with a small percentage coming from Action and Documentary/History. Mostly from "Less than 2 hours" to "2 - 5 hours" range of watching time. The IT field is the highest choice, with "Image/Trailer" and "Title/Description" are the most important factor.

Cluster 7: The majority are 21 to 22 years old. Action, Animation, Comedy, Crime, Horror and Romance shared the genre distribution without much differences. Watching periods mostly come from "Less than 2 hours" to "2 – 5 hours" range. Business, Engineering and Society are the most choices for field of occupation, and "ratings" as well as "title/description" are the most important factors.

5.5 Shortcomings

With the lack of documentation, using libraries that implement Gower's Distance is complicated and troublesome, as it leads to unexpected errors. One example is that functions to find the optimal number of clusters cannot be applied, whether it is Elbow, Silhouette or Gap Statistic method. Because of the limited time frame of this project, the errors cannot be solved yet. Therefore, it is crucial to research thoroughly beforehand if using the existing libraries for this method.

Group 4 Project Report Page 28/37

6 Conclusion

6.1 Accomplishment of Project Objectives

From conducting the survey, preparing data, analyzing input, to visualizing attributes by histograms, our team can identify the relation between viewing patterns (age, film genre, working field, frequency, factor) and we can recommend content based on that insight. For that reason, our team has achieved our project goals and objectives.

6.2 Future Work

Refining clustering algorithms, adding more features, or expanding the recommendation system's capabilities are admirable improvements for this project. Other potential areas can be building the movie recommendation web app, testing our recommendation, accuracy; as well as implementing other methods to analyze data such as non-hierarchical clustering, and comparing the result between two methods.

Group 4 Project Report Page 29/37

7 Appendix

7.1 Project Repository on GitHub

All source codes, data files, presentations, and survey questions of this project can be found in the project repository on GitHub. Here is the link to this project repository on GitHub: https://github.com/vhtuananh020402/Group4_Data_analysis

7.2 Survey Questions

- 1. How old are you?
- 2. What is your gender?
- 3. Which area do you work/learn in?
- 4. Which movie genre is your favorite? Choose only one.
- 5. How often do you watch movies per week?
- 6. What is the factor that influences your decision to watch a movie?
- 7. Which platform do you use to watch movies?

7.3 R Code

• dummy_canberra_ward.r

Group 4 Project Report Page 30/37

```
levels = c("Less than 2 hours", "2 - 5 hours", "6 - 10 hours", "11 - 15 hours", "16 -
      20 hours", "More than 20 hours"))
   df$Frequency_numeric <- as.numeric(factor(df$Frequency))</pre>
   df$Frequency_std <- scale(df$Frequency_numeric)</pre>
19
    # Turn the nominal variables into dummy variables
20
   df_dummy <- model.matrix(~ Age_std + Genre + Frequency_std + Field + Factor + Gender +
21
   Platform, data = df)
22
   # Calculate the points distance
23
   point_dist <- dist(df_dummy, method = "canberra")</pre>
                                                              # Using Canberra distance
24
25
    # Hierarchical cluster analysis on the data frame
26
   hc <- hclust(point_dist, method = "ward.D")</pre>
                                                              # Using Ward's method
27
28
   # Plot the dendrogram
29
   dend <- as.dendrogram(hc)</pre>
30
   plot(dend)
31
32
   \# Draw the rectangle around each cluster in k clusters
34
   rect.hclust(hc, k, border = 2:8)
35
36
37
38
    # ========== Gap statistic method to find the optimal number of cluster
   # Calculate Within-Cluster Dispersion (WCD) for the original data
40
   wss <- sum(hc$height)
42
   # Generate Random Data for Comparison
43
   set.seed(123) # Set seed for reproducibility
   B <- 100 # Number of random datasets
45
   random_datasets <- lapply(1:B, function(i) matrix(runif(length(df_dummy)), ncol =</pre>
   ncol(df_dummy)))
   # Cluster the Random Data
48
   random_hcs <- lapply(random_datasets, function(random_data) {</pre>
```

Group 4 Project Report Page 31/37

```
dist_matrix <- dist(random_data, method = dist_method[4])</pre>
     hclust(dist_matrix, method = hc_method[1])
   })
52
54
   # Calculate Within-Cluster Dispersion for Random Data
55
   wss_random <- sapply(random_hcs, function(random_hc) sum(random_hc$height))</pre>
57
58
   # Calculate Gap Statistic
59
   gap <- (log(wss_random) - log(wss)) + mean(log(wss_random) - log(wss))</pre>
60
61
   # Determine the Optimal Number of Clusters
62
   num_clusters <- 1 : 10 # Change this range according to the problem domain
63
64
   gap <- gap[1:length(num_clusters)]</pre>
65
66
   x_labels <- seq(1, length(num_clusters))</pre>
67
   plot(num_clusters, gap, xlab = "Number of Clusters", ylab = "Gap Statistic",
69
        main = "Gap Statistic Plot", type = "b")
70
71
   # Customize x-axis labels
   axis(1, at = x_labels, labels = num_clusters)
73
   abline(v = 8, col = "red", lty = 2)
74
76
   77
   # Add the cluster assignments to the data frame
   df$Cluster <- factor(clusters)</pre>
   print(clusters)
80
   # Create a histogram of the Genre distribution in each cluster
82
   ggplot(df, aes(x = Genre)) +
83
     geom_histogram(stat = "count", fill = "lightblue", color = "black", linewidth = 0.8) +
     facet_wrap(~ Cluster) +
85
     theme(axis.text.x = element_text(angle = 45, hjust = 1)) +
86
     labs(title = "Genre Distribution in Each Cluster", x = "Genre", y = "Count")
```

Group 4 Project Report Page 32/37

```
# Create a histogram of the Age distribution in each cluster
89
    ggplot(df, aes(x = Age)) +
90
      geom_histogram(stat = "count", fill = "orange", color = "black", linewidth = 0.8) +
      facet_wrap(~ Cluster) +
92
      theme(axis.text.x = element_text(angle = 45, hjust = 1)) +
93
      labs(title = "Age Distribution in Each Cluster", x = "Age", y = "Count")
94
95
    # Create a histogram of the Field distribution in each cluster
96
    ggplot(df, aes(x = Field)) +
97
      geom_histogram(stat = "count", fill = "red", color = "black", linewidth = 0.8) +
98
      facet_wrap(~ Cluster) +
99
      theme(axis.text.x = element_text(angle = 45, hjust = 1)) +
100
      labs(title = "Field Distribution in Each Cluster", x = "Field", y = "Count")
101
102
    # Create a histogram of the Frequency distribution in each cluster
103
    ggplot(df, aes(x = Frequency)) +
104
      geom_histogram(stat = "count", fill = "darkgrey", color = "black", linewidth = 0.8) +
105
      facet_wrap(~ Cluster) +
106
      theme(axis.text.x = element_text(angle = 45, hjust = 1)) +
      labs(title = "Frequency Distribution in Each Cluster", x = "Frequency", y = "Count")
108
109
    # Create a histogram of the Factor distribution in each cluster
    ggplot(df, aes(x = Factor)) +
111
      geom_histogram(stat = "count", fill = "lightgreen", color = "black", linewidth = 0.8)
112
      facet_wrap(~ Cluster) +
113
      theme(axis.text.x = element_text(angle = 45, hjust = 1)) +
114
      labs(title = "Factor Distribution in Each Cluster", x = "Factor", y = "Count")
```

• completed gower ward.r

```
library(cluster)  # contain function 'daisy'
library(factoextra)  # clustering visualization
library(ggplot2)  # draw distribution graph

# --- Data Preparation --- #

df <- read.csv("data/clean_data_v2.csv")

df <- na.omit(df)  # remove missing values in dataset</pre>
```

Group 4 Project Report Page 33/37

```
# standardize age, then add into numerical value
   df$Age_std <- scale(df$Age)</pre>
   num_attr <- c("Age")</pre>
    # add into categorical value
13
   cat_attr <- c("Field", "Genre", "Factor")</pre>
   df[cat_attr] <- lapply(df[cat_attr], as.factor)</pre>
15
16
    # add into ordinal value
17
   ord_attr <- c("Frequency")</pre>
18
   df$Frequency <- factor(df$Frequency,</pre>
19
                             order = TRUE,
20
                             level = c("Less than 2 hours",
21
                                        "2 - 5 hours",
22
                                        "6 - 10 hours",
23
                                        "11 - 15 hours",
24
                                        "16 - 20 hours",
25
                                        "More than 20 hours"))
26
    # put everything into a complete dataset
28
   process_dataset <- df %>% select(num_attr, ord_attr, cat_attr)
29
   head(process_dataset)
31
32
               # --- Calculation --- #
    # calculate Gower's distance
34
   gower_dist <- daisy(process_dataset, metric="gower")</pre>
35
    # hierarchical clustering, using ward.D method
37
   gower_hcl <- hclust(gower_dist, method = "ward.D")</pre>
38
               # --- DENDROGRAM ---- #
40
    # plot dendrogram
41
   plot(gower_hcl, cex = 0.6)
43
    # draw borders for the individual clusters
44
   rect.hclust(gower_hcl, k = 7, border = 2:7)
```

Group 4 Project Report Page 34/37

```
46
              # --- HISTOGRAM --- #
47
    # cut into k clusters
48
   k <- 7
    clusters <- cutree(gower_hcl, k)</pre>
50
51
    # add the cluster assignments to the data frame
52
   df$Cluster <- factor(clusters)</pre>
53
54
    # histogram of Genre distribution in each cluster
55
    ggplot(df, aes(x = Genre)) +
56
      geom_histogram(stat = "count", fill = "lightblue", color = "black", linewidth = 0.8) +
57
     facet_wrap(~ Cluster) +
58
      theme(axis.text.x = element_text(angle = 45, hjust = 1)) +
59
      labs(title = "Genre Distribution in Each Cluster", x = "Genre", y = "Count")
60
61
    # histogram of Field distribution in each cluster
62
   ggplot(df, aes(x = Field)) +
63
      geom_histogram(stat = "count", fill = "red", color = "black", linewidth = 0.8) +
64
     facet_wrap(~ Cluster) +
      theme(axis.text.x = element_text(angle = 45, hjust = 1)) +
66
      labs(title = "Field Distribution in Each Cluster", x = "Field", y = "Count")
67
    # histogram of Factor distribution in each cluster
69
   ggplot(df, aes(x = Factor)) +
70
      geom_histogram(stat = "count", fill = "lightgreen", color = "black", linewidth = 0.8)
     facet_wrap(~ Cluster) +
72
      theme(axis.text.x = element_text(angle = 45, hjust = 1)) +
      labs(title = "Factor Distribution in Each Cluster", x = "Factor", y = "Count")
74
75
    # create a histogram of Age distribution in each cluster
   ggplot(df, aes(x = Age)) +
77
      geom_histogram(stat = "count", fill = "orange", color = "black", linewidth = 0.8) +
78
     facet_wrap(~ Cluster) +
79
      theme(axis.text.x = element_text(angle = 45, hjust = 1)) +
80
      labs(title = "Age Distribution in Each Cluster", x = "Age", y = "Count")
81
```

Group 4 Project Report Page 35/37

```
# create a histogram of the Genre distribution in each cluster
ggplot(df, aes(x = Frequency)) +
geom_histogram(stat = "count", fill = "darkgrey", color = "black", linewidth = 0.8) +
facet_wrap(~ Cluster) +
theme(axis.text.x = element_text(angle = 45, hjust = 1)) +
labs(title = "Frequency Distribution in Each Cluster", x = "Frequency", y = "Count")
```

Group 4 Project Report Page 36/37

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Group 4 Project Report Page 37/37