Task Title:

Clustering and Dimensionality Reduction in Machine Learning

Objective:

To explore and implement unsupervised learning techniques including clustering and dimensionality reduction, using a real-world dataset.

Dataset Used:

I used the **Wholesale Customers Dataset** for this project. This dataset contains information on the annual spending habits of wholesale clients across different product categories, including:

- > Fresh
- > Milk
- Grocery
- > Frozen
- > Detergents Paper
- > Delicassen

Additionally, the dataset includes two categorical columns: **Channel** (1 = Horeca, 2 = Retail) and **Region** (1 = Lisbon, 2 = Oporto, 3 = Other)

Steps Performed:

1-Importing Libararies

I began the task by importing essential Python libraries required for data handling, preprocessing, visualization, clustering, and evaluation:

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns

from sklearn.preprocessing import StandardScaler
from sklearn.decomposition import PCA from
sklearn.manifold import TSNE
from sklearn.cluster import KMeans, DBSCAN, AgglomerativeClustering
from sklearn.metrics import silhouette_score, davies_bouldin_score
```

2-Loading Dataset

The dataset was uploaded to **Google Colab** using the file upload feature. After uploading:

I explored the first few rows using df.head() to understand its structure.

```
df = pd.read csv('/content/Wholesale customers data.csv')
df.head()
{"summary":"{\n \"name\": \"df\",\n \"rows\": 440,\n \"fields\": [\
n {\n \"column\": \"Channel\",\n \"properties\": {\n \"dtype\": \"number\",\n \"std\": 0,\n \"min\": 1,\n \"max\": 2,\n \"num_unique_values\": 2,\n \"samples\": [\n 1,\n 2\n ],\n \"semantic_type\": \"\",\n \"description\": \"\"\n }\n {\n \"}
\"column\": \"Region\", \n \"properties\": {\n \"dtype\":
\"number\",\n \"std\": 12647,\n \"min\": 3,\n \"max\": 112151,\n \"num_unique_values\": 433,\n \"samples\": [\n 21117,\n 20398\n ],\n \"semantic_type\": \"\",\n \"description\": \"\"\n }\
n },\n {\n \"column\": \"Milk\",\n \"properties\": {\n \"dtype\": \"number\",\n \"std\": 7380,\n \"min\": 55,\n
\"max\": 73498,\n \"num_unique_values\": 421,\n \"samples\": [\n 8384,\n 7184\n ],\n \"semantic_type\": \"\",\n \"description\": \"\"\n }\
                                                                                                                                                                                        ],\n
n },\n {\n \"column\": \"Grocery\",\n \"properties\":

{\n \"dtype\": \"number\", \n \"std\": 9503, \n
\"min\": 3, \n \"max\": 92780, \n \"num_unique_values\":
430, \n \"samples\": [\n \ 5160, \n \ 3\\
n \ ], \n \"semantic_type\": \"\", \n
\"description\": \"\"\n \ \\n \\"column\":
\"Frozen\", \n \"properties\": \\n \"dtype\": \"number\", \n
\"std\": 4854, \n \"min\": 25, \n \"max\": 60869, \n
\"num_unique_values\": 426 \n
\"samples\": [\n \"column\": 269 \n
\"num_unique_values\": 426 \n
\"samples\": [\n \"column\": 269 \n
\"samples\": [\n \"column\": 269 \n
\"samples\": [\n \"column\": 269 \n
\"num_unique_values\": 426 \n
\"samples\": [\n \"column\": [\n \n \n]
\"num_unique_values\": 426 \n
\"samples\": [\n \n \n]
\"samples\": [\n \n]
\"samples\": [\n \n]
\"num_unique_values\": 426 \n
\"samples\": [\n \n]
\"num_unique_values\": 426 \n
\"num_unique_values\": [\n \n]
\"num_unique_values\": 426 \n
\"num_unique_values\": [\n \n]
\"num_unique_values\": 426 \n
\"num_unique_values\": [\n \n]
\"num_unique_values\": [\n]
\"num_unique_values\": [\n \n]
\"num_unique
\"num_unique_values\": 426,\n \"samples\": [\n 269,\n
\"number\",\n \"std\": 4767,\n \"min\": 3,\n \"max\": 40827,\n \"num_unique_values\": 417,\n \"samples\": [\n 302,\n 6740\n ],\n
2820,\n \"min\": 3,\n \"max\": 47943,\n \"num_unique_values\": 403,\n \"samples\": [\n 14472,\n 172\n ],\n \"semantic_type\": \"\",\n
\"description\": \"\"\n }\n }\n ]\
n}","type":"dataframe","variable name":"df"}
```

3-Initial Data Exploration

To understand the dataset better, I explored the following:

- Data types and non-null entries with df.info()
- Statistical summary with df.describe()
- Checked for missing values and duplicate rows:

To prepare the dataset for clustering:

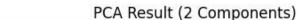
- $\bullet \quad I \ \textbf{removed duplicate entries using } \ \texttt{df.drop_duplicates()}$
- Dropped categorical columns Channel and Region as clustering was applied on numerical data
- Standardized the data using StandardScaler() to bring all features to a similar scale, which is essential before applying PCA and clustering algorithms.

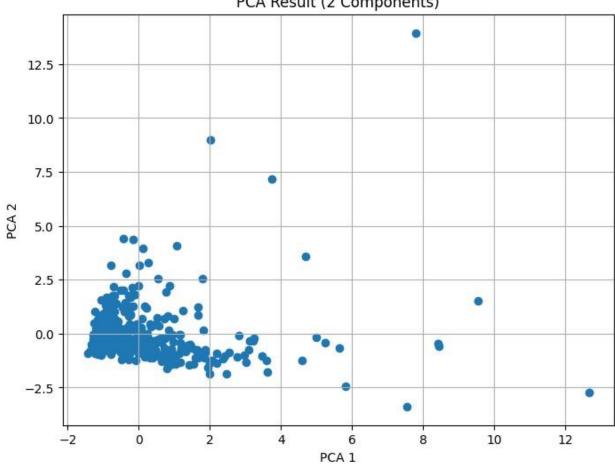
1	0	3	7057	0010	`	0.5.00	1 7	.		2202		
1	2	3	7057	9810	J	9568	17	02		3293		
2	2	3	6353	8808	3	7684	240)5		3516		
3	1	3	13265	1196	5	4221	640	04		507		
4	2			3	2261	.5	541	LO		7198		3915
1777						•						
435	1	3	29703	12051	L î	16027	1313	35		182		
436	1	3	39228	1431	L	764	451	10		93		
427	2	3	1 4 5 2 1	1 - 400		20042	4.1) T		1 4 0 4 1		
437	2	3	14531	15488	5 .	30243	4、	37		14841		
438	1	3			981	22		1038			168	439
1	3	2787	1698	3	2510	(65		47	7		
0	Delica 1338	ssen										
1	1776											
2	7844											
3	1788											
4	5185											
435		220										
436 437		2346 1867										
437		2125										
439		52										
[440 rows x 8 columns]>												

5-PCA for Dimensionality Reduction

I applied **Principal Component Analysis (PCA)** to reduce the high-dimensional dataset into 2 dimensions for easier visualization and analysis:

```
pca = PCA(n_components=2)
pca_result = pca.fit_transform(scaled_data)
print("Explained variance ratio:", pca.explained_variance_ratio_)
plt.figure(figsize=(8, 6))
plt.scatter(pca_result[:, 0], pca_result[:, 1])
plt.xlabel('PCA 1') plt.ylabel('PCA 2')
plt.title('PCA Result (2 Components)')
plt.grid(True) plt.show()
Explained variance ratio: [0.44082893 0.283764]
```

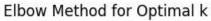


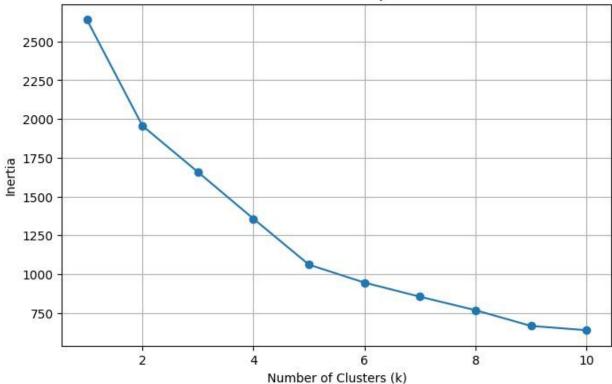


- The first two principal components explained **72.46%** of the variance.
- · The reduced dataset was visualized using a **2D scatter plot**.

6-K-Means Clustering + Elbow Method

I used the **Elbow Method** to determine the optimal number of clusters for K-Means:



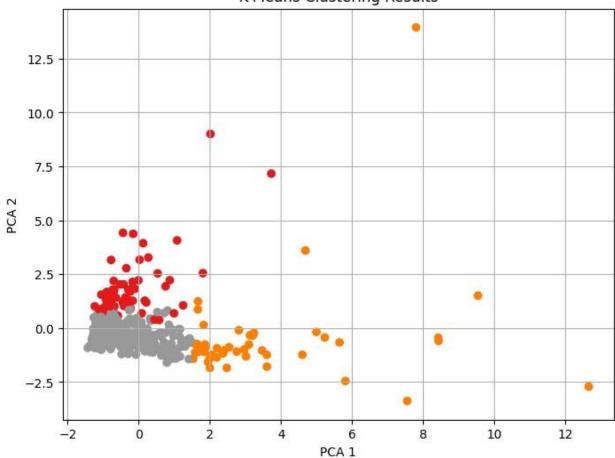


- Based on the elbow point, I chose k = 3 clusters.
- Applied **KMeans** and visualized the clusters on the PCA-reduced data.

7-Apply Clustering Algorithms

```
kmeans = KMeans(n_clusters=3, random_state=0)
kmeans_labels = kmeans.fit_predict(scaled_data)
plt.figure(figsize=(8,6))
plt.scatter(pca_result[:, 0], pca_result[:, 1], c=kmeans_labels,
cmap='Set1')
plt.title('K-Means Clustering Results')
plt.xlabel('PCA 1') plt.ylabel('PCA 2')
plt.grid(True) plt.show()
```

K-Means Clustering Results



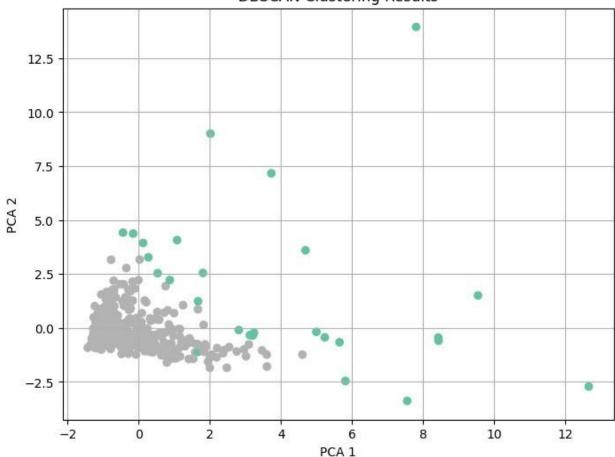
7-DBSCAN

I applied **DBSCAN** (**Density-Based Spatial Clustering**) as a second clustering technique:

```
dbscan = DBSCAN(eps=1.5, min_samples=5)
db_labels = dbscan.fit_predict(scaled_data)

plt.figure(figsize=(8,6))
plt.scatter(pca_result[:, 0], pca_result[:, 1], c=db_labels,
cmap='Set2')
plt.title('DBSCAN Clustering Results')
plt.xlabel('PCA 1') plt.ylabel('PCA
2') plt.grid(True) plt.show()
```

DBSCAN Clustering Results



- DBSCAN was able to identify cluster structure based on density and noise.
- The clusters were visualized using the same 2D PCA-reduced plot.

8-Evaluation

To evaluate the quality of the clusters formed by both KMeans and DBSCAN, I calculated the **Silhouette Score** and **Davies-Bouldin Index**:

K-Means Results:

• Silhouette Score: 0.3916

• Davies-Bouldin Index: 1.2494

DBSCAN Results:

• Silhouette Score: 0.6602

• Davies-Bouldin Index: 1.4808

Insight: DBSCAN produced a higher silhouette score indicating better-defined clusters. However, it had a slightly higher Davies-Bouldin score. Overall, DBSCAN performed better in terms of clustering tightness and separation.