Code Implementation & Analysis

Customer Segmentation using Clustering

In today's competitive market, businesses collect large amounts of customer data, but raw data alone does not provide direct value. To make data-driven decisions, it is essential to identify patterns and group customers with similar behaviors, preferences, or demographics.

This project focuses on applying unsupervised machine learning techniques — specifically dustering algorithms — to segment customers into meaningful groups. By understanding these customer segments, businesses can:

- · Personalize marketing strategies
- · Improve customer experience
- · Identify potential high-value customers
- Reduce churn by targeting the right audience with the right message

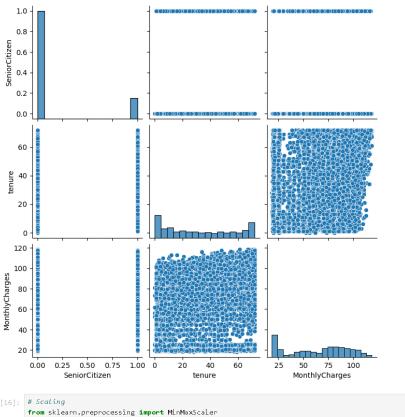
We will use Python along with libraries such as Pandas, NumPy, Matplotlib, and Scikit-learn to:

- · Clean and preprocess the dataset
- Perform exploratory data analysis (EDA) to understand key features
- Apply dustering algorithms (e.g., K-Means)
- · Visualize dusters for better interpretability

Data cleaned. Shape after cleaning: (7043, 21)

```
[6]: # Create config.yaml with absolute path
      import yaml
          "data_source": r"C:\Users\ASUS\OneDrive\Desktop\STELLARA DATASET\WA_Fn-UseC_-Telco-Customer-Churn.xlsx"
      with open("config.yaml", "w") as file:
         yaml.dump(config_data, file)
      print("config.yaml created successfully.")
      config.yaml created successfully.
[8]: # Load configuration & dataset
      import pandas as pd
      with open("config.yaml", "r") as file:
         config = yaml.safe_load(file)
      data_path = config["data_source"]
      df = pd.read_excel(data_path)
      print(" Dataset shape before cleaning:", df.shape)
       Dataset shape before cleaning: (7043, 21)
[10]: # Data Cleaning
      df = df[df.isnull().mean(axis=1) < 0.7] # Remove >70% missing
      df.fillna(df.mean(numeric_only=True), inplace=True) # Fitt with mean
      df.drop_duplicates(inplace=True) # Remove duplicates
      print("Data cleaned. Shape after cleaning:", df.shape)
```

```
[12]: # EDA (Univariate, Bivariate, Multivariate)
        import seaborn as sns
       import matplotlib.pyplot as plt
       df.hist(bins=25, figsize=(15, 10))
       plt.tight_layout()
plt.show()
       sns.heatmap(df.corr(numeric_only=True), annot=True, cmap="coolwarm")
       plt.show()
       sns.pairplot(df.select_dtypes(include='number').iloc[:, :4])
plt.show()
                                         SeniorCitizen
                                                                                  800
        5000
                                                                                  600
        4000
        3000
                                                                                  400
        2000
                                                                                  200
        1000
             0.0
                                            MonthlyCharges
        1200
        1000
        800
        600
        400 -
        200
                20
                                                          80
                                                                       100
                                                                                     120
                                                                               - 1.0
        SeniorCitizen
                                                                               - 0.8
                                                                              - 0.6
       tenure
                                                           0.25
                                                                               - 0.4
       MonthlyCharges
                                                                               - 0.2
                                       0.25
              SeniorCitizen
                                      tenure
                                                     MonthlyCharges
```



```
from sklearn.preprocessing import MinMaxScaler

scaler = MinMaxScaler()
num_df = df.select_dtypes(include='number')
scaled = scaler.fit_transform(num_df)
scaled_df = pd.DataFrame(scaled, columns=num_df.columns)
df[num_df.columns] = scaled_df
print("Data scaled")

Data scaled
```

```
[14]: # Outlier Removat (IQR)
num_df = df.select_dtypes(include='number')
Q1 = num_df.quantile(0.25)
Q3 = num_df.quantile(0.75)
IQR = Q3 - Q1
df = df[~((num_df < (Q1 - 1.5 * IQR)) | (num_df > (Q3 + 1.5 * IQR))).any(axis=1)]
print("Outliers removed. Shape: ", df.shape)
Outliers removed. Shape: (5901, 21)
```

```
[36]: # Random Forest
           rf = RandomForestClassifier()
           rf.fit(X, y)
           print("Top 5 Features (RandomForest):\n", pd.Series(rf.feature_importances_, index=X.columns).sort_values(ascending=False).head())
           Top 5 Features (RandomForest):
            MonthlyCharges
                                               0.044491
            tenure
                                              0.041623
                                             0.027985
           Contract_Two year
           InternetService_Fiber optic
                                              0.027399
           PaymentMethod_Electronic check 0.024558
           dtype: float64
   [42]: from sklearn.impute import SimpleImputer
          import pandas as pd
          from sklearn.ensemble import AdaBoostClassifier
          # Create an imputer to handle missing values
         imputer = SimpleImputer(strategy='mean') # You can choose other strategies like 'median', 'most_frequent', etc.
          # Fit and transform the data to handle missing values
         X_imputed = imputer.fit_transform(X)
          # Now we can use X_imputed in the AdaBoost model
         ada = AdaBoostClassifier()
         ada.fit(X_imputed, y)
         print("Top 5 Features (AdaBoost):\n", pd.Series(ada.feature_importances_, index=X.columns).sort_values(ascending=False).head())
         C:\Users\ASUS\anaconda3\Lib\site-packages\sklearn\ensemble\_weight_boosting.py:519: FutureWarning: The SAMME.R algorithm (the default) is deprecated and will be removed in 1.6. Use the SAMME algorithm to circumvent this warning.
           warnings.warn(
          Top 5 Features (AdaBoost):
           Contract_Two year
                                               0.08
          InternetService_Fiber optic
                                             0.04
          TechSupport_No internet service 0.04
          Contract One year
                                             0.04
          OnlineSecurity_Yes
                                            0.02
          dtype: float64
     [44]: # XGBoost
            xgb = XGBClassifier(use_label_encoder=False, eval_metric='mlogloss')
            print("Top 5 Features (XGBoost):\n", pd.Series(xgb.feature_importances_, index=X.columns).sort_values(ascending=False).head())
            C:\Users\ASUS\anaconda3\Lib\site-packages\xgboost\training.py:183: UserWarning: [22:40:55] WARNING: C:\actions-runner\_work\xgboo
            st\xgboost\src\learner.cc:738:
            Parameters: { "use_label_encoder" } are not used.
             bst.update(dtrain, iteration=i, fobj=obj)
            Top 5 Features (XGBoost):
                                            0.346791
             Contract Two year
                                           0.167848
            Contract One year
            InternetService_Fiber optic
                                         0.092792
            OnlineSecurity_Yes
                                           0.031399
            InternetService No
                                          0.023542
            dtype: float32
[20]: # Clustering (DBSCAN)
      from sklearn.cluster import DBSCAN
      import pandas as pd
      from sklearn.impute import SimpleImputer
      # Handle missing values using SimpleImputer
      imputer = SimpleImputer(strategy='mean') # You can choose 'mean', 'median', 'most_frequent', or 'constant'
      X_imputed = imputer.fit_transform(X)
      # Now apply DBSCAN on the imputed data
      dbscan = DBSCAN(eps=0.5, min_samples=5)
      clusters = dbscan.fit_predict(X_imputed)
      df["Cluster"] = clusters
      print("DBSCAN Clustering Done. Cluster counts:\n", pd.Series(clusters).value_counts())
      DBSCAN Clustering Done. Cluster counts:
```

-1 5901

Name: count, dtype: int64

```
# Train/Test + Evaluation
from sklearn.model_selection import train_test_split
from sklearn.metrics import classification_report, f1_score

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)

model = RandomForestClassifier()
model.fit(X_train, y_train)
y_pred = model.predict(X_test)

print("\nClassification Report:\n", classification_report(y_test, y_pred))
print(f"Macro F1 Score: {f1_score(y_test, y_pred, average='macro'):.4f)")
```

Classification Report:

	precision	recall	f1-score	support
0	0.82	0.95	0.88	916
1	0.62	0.26	0.37	265
accuracy			0.80	1181
macro avg	0.72	0.61	0.63	1181
weighted avg	0.77	0.80	0.77	1181

Macro F1 Score: 0.6260