2211CS020319 AIML-DELTA

## BDA HOLIDAY ASSIGNMENT -1

## In [23]: !pip install imbalanced-learn Requirement already satisfied: imbalanced-learn in c:\users\chait\anaconda3\lib\site-packages (0.12.4) Requirement already satisfied: scikit-learn>=1.0.2 in c:\users\chait\anaconda3\lib\site-packages (from imba lanced-learn) (1.0.2) Requirement already satisfied: threadpoolctl>=2.0.0 in c:\users\chait\anaconda3\lib\site-packages (from imb alanced-learn) (3.5.0) Requirement already satisfied: scipy>=1.5.0 in c:\users\chait\anaconda3\lib\site-packages (from imbalancedlearn) (1.9.1) Requirement already satisfied: joblib>=1.1.1 in c:\users\chait\anaconda3\lib\site-packages (from imbalanced -learn) (1.4.2) Requirement already satisfied: numpy>=1.17.3 in c:\users\chait\anaconda3\lib\site-packages (from imbalanced -learn) (1.24.4) WARNING: Ignoring invalid distribution -rotobuf (c:\users\chait\anaconda3\lib\site-packages) WARNING: Ignoring invalid distribution -rotobuf (c:\users\chait\anaconda3\lib\site-packages)

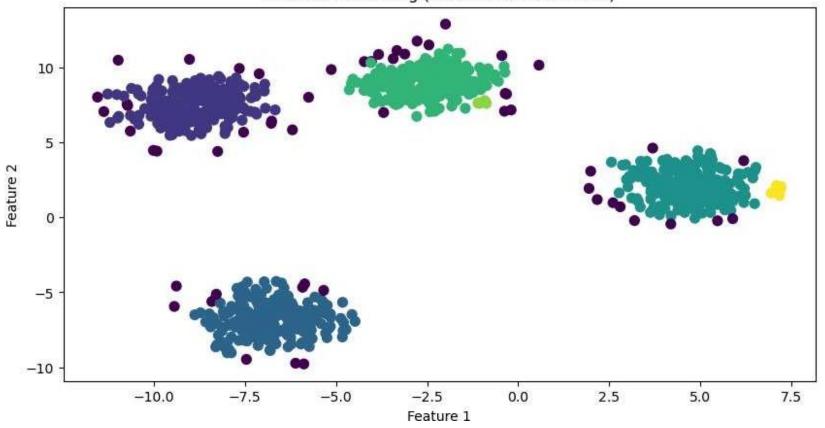
```
In [4]: #Question 1: Handling Imbalanced Datasets
        from sklearn.datasets import make classification
        from sklearn.model selection import train test split
        from sklearn.ensemble import RandomForestClassifier
        from sklearn.metrics import classification report
        from imblearn.over sampling import SMOTE
        from sklearn.utils.class weight import compute class weight
        # Generate a synthetic imbalanced dataset
        X, y = make classification(n samples=1000, n features=20, n classes=2,
                                   weights=[0.9, 0.1], random state=42)
        # Split the dataset into training and test sets
        X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=42)
        # Apply SMOTE for oversampling the minority class
        smote = SMOTE(random state=42)
        X train smote, y train smote = smote.fit resample(X train, y train)
        # Compute class weights to adjust for imbalance
        class weights = compute class weight('balanced', classes=[0, 1], y=y train)
        class weights dict = {0: class weights[0], 1: class weights[1]}
        # Train a RandomForestClassifier using the resampled data
        model = RandomForestClassifier(class weight=class weights dict, random state=42)
        model.fit(X train smote, y train smote)
        # Make predictions on the test set
        y pred = model.predict(X test)
        # Print classification report
        print(classification report(y test, y pred))
```

	precision	recall	f1-score	support
0 1	<b>0.</b> 96 <b>0.</b> 59	0.95 0.63	0.96 0.61	270 30
accuracy	0.78	0.79	0.92 0.78	300 300
macro avg weighted avg	0.92	0.92	0.78	300

!pip install --upgrade threadpoolctl

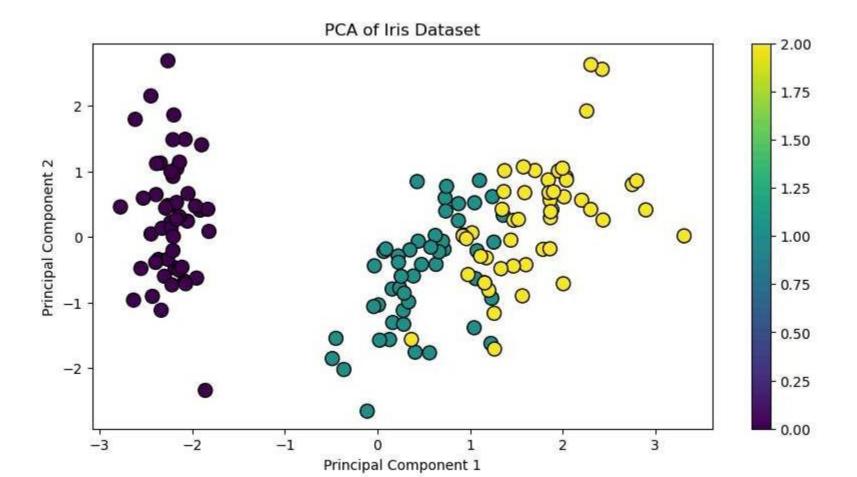
```
In [10]: #Question 2: Optimal Clusters for K-means
         from sklearn.datasets import make blobs
         from sklearn.cluster import DBSCAN
         from sklearn.metrics import silhouette score
         import matplotlib.pyplot as plt
         # Generate synthetic data
         X, _ = make_blobs(n_samples=1000, n_features=2, centers=4, cluster_std=1.0, random_state=42)
         # Apply DBSCAN
         dbscan = DBSCAN(eps=0.5, min_samples=5).fit(X)
         labels = dbscan.labels_
         # Calculate the silhouette score
         score = silhouette score(X, labels)
         # Plot the clusters
         plt.figure(figsize=(10, 5))
         plt.scatter(X[:, 0], X[:, 1], c=labels, cmap='viridis', s=50)
         plt.xlabel('Feature 1')
         plt.ylabel('Feature 2')
         plt.title(f'DBSCAN Clustering (Silhouette Score: {score:.2f})')
         plt.show()
```

DBSCAN Clustering (Silhouette Score: 0.50)



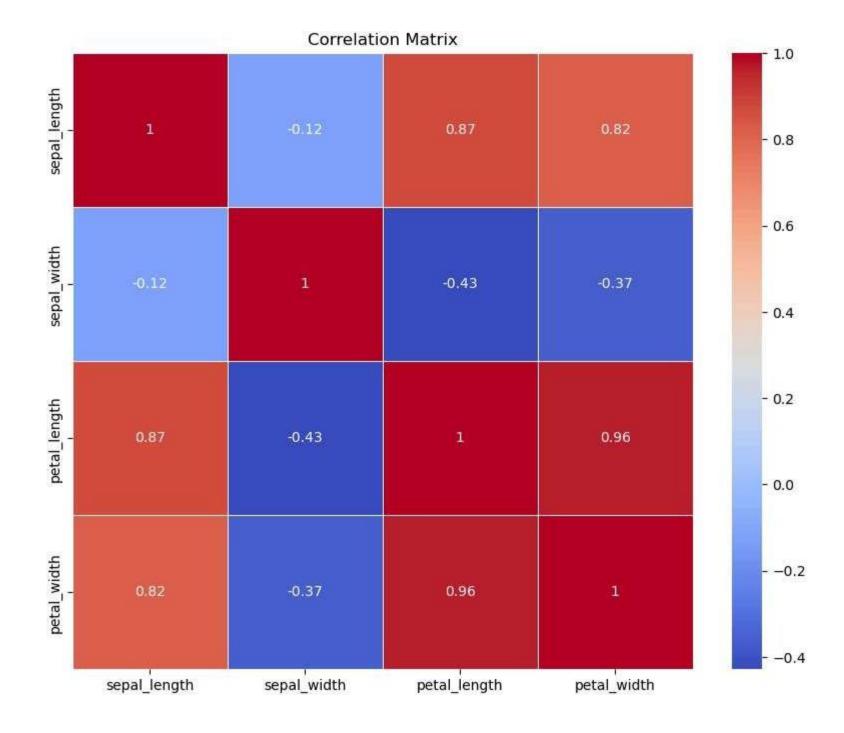
```
In [11]: #Question 3:Dimensionality reduction
         import numpy as np
         from sklearn.datasets import load iris
         from sklearn.preprocessing import StandardScaler
         from sklearn.decomposition import PCA
         import matplotlib.pyplot as plt
         # Load the dataset
         data = load iris()
         X = data.data
         y = data.target
         # Standardize the data
         scaler = StandardScaler()
         X scaled = scaler.fit_transform(X)
         # Apply PCA
         pca = PCA(n components=2)
         X pca = pca.fit transform(X scaled)
         # Explained variance
         explained_variance = pca.explained_variance_ratio_
         print(f"Explained variance by component: {explained variance}")
         # Plot the transformed data
         plt.figure(figsize=(10, 5))
         plt.scatter(X_pca[:, 0], X_pca[:, 1], c=y, cmap='viridis', edgecolor='k', s=100)
         plt.xlabel('Principal Component 1')
         plt.ylabel('Principal Component 2')
         plt.title('PCA of Iris Dataset')
         plt.colorbar()
         plt.show()
```

Explained variance by component: [0.72962445 0.22850762]



```
#Question 4: Correlations in a Dataset
In [12]:
         import pandas as pd
         import seaborn as sns
         import matplotlib.pyplot as plt
         # Load a sample dataset
         df = sns.load dataset('iris')
         # Calculate correlation matrix
         correlation matrix = df.corr()
         # Print the correlation matrix
         print(correlation_matrix)
         # Visualize the correlation matrix using a heatmap
         plt.figure(figsize=(10, 8))
         sns.heatmap(correlation matrix, annot=True, cmap='coolwarm', linewidths=0.5)
         plt.title('Correlation Matrix')
         plt.show()
```

```
sepal_length sepal_width petal_length petal_width
sepal length
                 1.000000
                                           0.871754
                            -0.117570
                                                        0.817941
sepal_width
                -0.117570
                            1.000000
                                          -0.428440
                                                       -0.366126
petal_length
                 0.871754
                                           1.000000
                                                       0.962865
                             -0.428440
petal width
                 0.817941
                            -0.366126
                                           0.962865
                                                       1.000000
```



Mean Imputation:
A B

```
In [15]: #Question 6: Detect and Remove Duplicates
         #import pandas as pd
         # Sample dataset with duplicates
         data = \{'A': [1, 2, 2, 4, 5],
                 'B': [5, 6, 6, 8, 9],
                 'C': [1, 2, 2, 4, 5]}
         df = pd.DataFrame(data)
         # Detect duplicates
         duplicates = df[df.duplicated()]
         print("Detected Duplicates:\n", duplicates)
         # Remove duplicates
         df cleaned = df.drop duplicates()
         print("DataFrame after Removing Duplicates:\n", df_cleaned)
         Detected Duplicates:
             А В С
         2 2 6 2
```

DataFrame after Removing Duplicates:

A B C
0 1 5 1
1 2 6 2
3 4 8 4
4 5 9 5

```
In [16]: #Question 7: Random Forest Regression for Housing Prices
         import pandas as pd
         from sklearn.model selection import train test split
         from sklearn.ensemble import RandomForestRegressor
         from sklearn.metrics import mean_squared_error, r2_score
         from sklearn.datasets import load boston
         # Load the dataset
         boston = load_boston()
         df = pd.DataFrame(boston.data, columns=boston.feature_names)
         df['PRICE'] = boston.target
         # Handle missing values if any (for demonstration, Boston dataset has no missing values)
         # df.fillna(df.mean(), inplace=True)
         # Split the data into training and test sets
         X = df.drop('PRICE', axis=1)
         y = df['PRICE']
         X train, X test, y train, y test = train_test_split(X, y, test_size=0.2, random_state=42)
         # Train the Random Forest Regressor
         model = RandomForestRegressor(n_estimators=100, random_state=42)
         model.fit(X train, y train)
         # Make predictions on the test set
         y_pred = model.predict(X_test)
         # Evaluate the model
         mse = mean_squared_error(y_test, y_pred)
         r2 = r2 \ score(y \ test, y \ pred)
         print(f"Mean Squared Error: {mse}")
         print(f"R^2 Score: {r2}")
         # Feature importance
         importances = model.feature_importances_
         feature importances = pd.DataFrame({'Feature': X.columns, 'Importance': importances})
         feature importances = feature importances.sort values(by='Importance', ascending=False)
         print("Feature Importances:\n", feature importances)
```

C:\Users\chait\anaconda3\lib\site-packages\sklearn\utils\deprecation.py:87: FutureWarning: Function load bo ston is deprecated; `load boston` is deprecated in 1.0 and will be removed in 1.2. The Boston housing prices dataset has an ethical problem. You can refer to the documentation of this function for further details. The scikit-learn maintainers therefore strongly discourage the use of this dataset unless the purpose of the code is to study and educate about ethical issues in data science and machine learning. In this special case, you can fetch the dataset from the original source:: import pandas as pd import numpy as np data url = "http://lib.stat.cmu.edu/datasets/boston" raw df = pd.read csv(data url, sep="\s+", skiprows=22, header=None) data = np.hstack([raw df.values[::2, :], raw df.values[1::2, :2]]) target = raw\_df.values[1::2, 2] Alternative datasets include the California housing dataset (i.e. :func:`~sklearn.datasets.fetch california housing`) and the Ames housing dataset. You can load the datasets as follows:: from sklearn.datasets import fetch california housing housing = fetch california\_housing() for the California housing dataset and:: from sklearn.datasets import fetch openml housing = fetch\_openml(name="house\_prices", as\_frame=True) for the Ames housing dataset.

warnings.warn(msg, category=FutureWarning)

Mean Squared Error: 7.901513892156864

R^2 Score: 0.8922527442109116

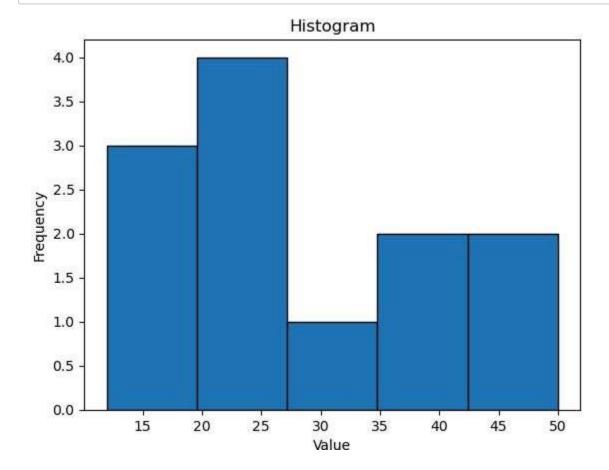
Feature Importances:

	car c impor	
	Feature	Importance
5	RM	0.503845
12	LSTAT	0.309509
7	DIS	0.060549
0	CRIM	0.038062
10	PTRATIO	0.016313
9	TAX	0.015661
4	NOX	0.015544
6	AGE	0.013840
11	В	0.012154
2	INDUS	0.007953
8	RAD	0.003811
1	ZN	0.001756
3	CHAS	0.001004

```
In [17]: #Question 8: Histogram, Bar Chart, and Pie Chart
    import matplotlib.pyplot as plt
    import pandas as pd

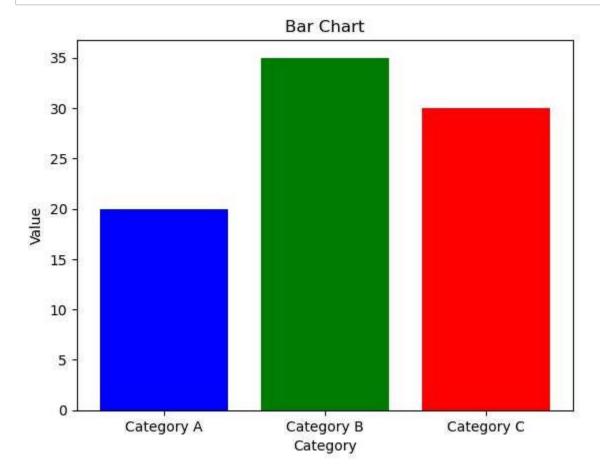
# Sample data
    data = [12, 15, 17, 20, 22, 25, 27, 30, 35, 40, 45, 50]

# Plot histogram
    plt.hist(data, bins=5, edgecolor='black')
    plt.xlabel('Value')
    plt.ylabel('Frequency')
    plt.title('Histogram')
    plt.show()
```



```
In [18]: # Sample data
    categories = ['Category A', 'Category B', 'Category C']
    values = [20, 35, 30]

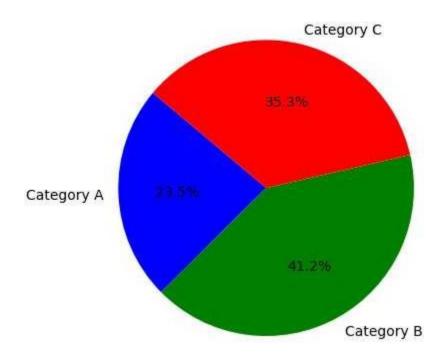
# Plot bar chart
    plt.bar(categories, values, color=['blue', 'green', 'red'])
    plt.xlabel('Category')
    plt.ylabel('Value')
    plt.title('Bar Chart')
    plt.show()
```



```
In [19]: # Sample data
    categories = ['Category A', 'Category B', 'Category C']
    values = [20, 35, 30]

# Plot pie chart
    plt.pie(values, labels=categories, autopct='%1.1f%%', colors=['blue', 'green', 'red'], startangle=140)
    plt.title('Pie Chart')
    plt.show()
```

## Pie Chart



```
In [20]: #Question 9: Linear and Logistic Regression
         import pandas as pd
         from sklearn.model selection import train test split
         from sklearn.linear model import LinearRegression
         from sklearn.metrics import mean squared error, r2 score
         # Sample dataset
         data = {'Area': [650, 785, 1200, 1600, 2100],
                 'Price': [80000, 105000, 180000, 230000, 300000]}
         df = pd.DataFrame(data)
         # Features and target variable
         X = df[['Area']]
         v = df['Price']
         # Split the data
         X train, X test, y train, y test = train test split(X, y, test size=0.2, random state=42)
         # Train the model
         model = LinearRegression()
         model.fit(X train, y train)
         # Make predictions
         y pred = model.predict(X test)
         # Evaluate the model
         mse = mean squared error(y test, y pred)
         r2 = r2_score(y_test, y_pred)
         print(f"Mean Squared Error: {mse}")
         print(f"R^2 Score: {r2}")
```

```
R^2 Score: nan
C:\Users\chait\anaconda3\lib\site-packages\sklearn\metrics\_regression.py:796: UndefinedMetricWarning: R^2
score is not well-defined with less than two samples.
   warnings.warn(msg, UndefinedMetricWarning)
```

Mean Squared Error: 4306026.030872749

```
Value Lag_1 Lag_2
Date
2022-01-01
            10 10.0 10.0
2022-01-02
            12 10.0 10.0
2022-01-03
            14 12.0 10.0
2022-01-04
            13 14.0 12.0
2022-01-05
            15 13.0 14.0
2022-01-06
            18 15.0 13.0
2022-01-07
            20 18.0 15.0
2022-01-08
            19 20.0 18.0
2022-01-09
            21 19.0 20.0
            23 21.0 19.0
2022-01-10
```

```
In [ ]:
```