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## Assessment 3

[344 rows x 7 columns]

Perform the below Tasks to complete the Assignment: -

Clustering the data and performing classification algorithms

#### 1. Download the dataset: Dataset

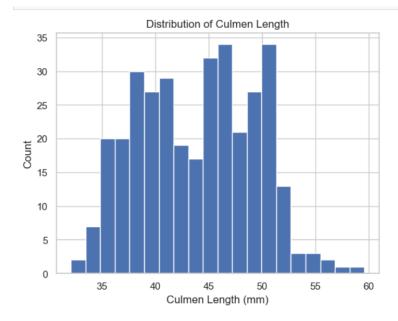
#### 2. Load the dataset into the tool

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler, LabelEncoder
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import classification_report, confusion_matrix
d = pd.read csv('C:\\Users\\wwwad\\Downloads\\penguins size.csv')
df=pd.DataFrame(d)
print(d)
   species
             island culmen_length_mm culmen_depth_mm flipper_length_mm
   Adelie Torgersen
  Adelie Torgersen
                                39.5
                                               17.4
                                                                186.0
1
   Adelie Torgersen
                                40.3
                                               18.0
                                                                195.0
   Adelie Torgersen
3
                                NaN
                                                NaN
                                                                 NaN
  Adelie Torgersen
                                36.7
                                                                193.0
339 Gentoo
            Biscoe
                                NaN
                                                NaN
                                                                 NaN
                                              14.3
15.7
340 Gentoo
            Biscoe
                                46.8
                                                                215.0
341
    Gentoo
             Biscoe
                                50.4
                                                                222.0
342 Gentoo
           Biscoe
                                45.2
                                              14.8
                                                                212.0
                                49.9
343 Gentoo
           Biscoe
                                               16.1
                                                                213.0
    body_mass_g
0
         3750.0
                 MALE
1
        3800.0 FEMALE
       3250.0 FEMALE
2
3
          NaN
                  NaN
4
       3450.0 FEMALE
          NaN
339
                  NaN
340
        4850.0 FEMALE
341
         5750.0
                 MALE
        5200.0 FEMALE
342
        5400.0
```

## 3. Perform Below Visualizations.

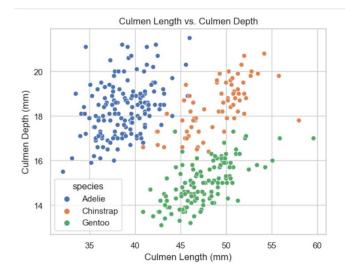
## Univariate Analysis

```
plt.hist(df['culmen_length_mm'], bins=20)
plt.xlabel('Culmen Length (mm)')
plt.ylabel('Count')
plt.title('Distribution of Culmen Length')
plt.show()
```

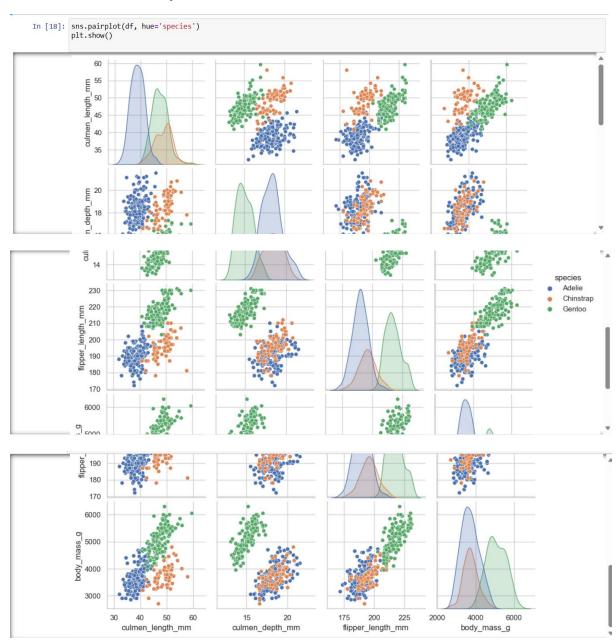


# • Bi- Variate Analysis

```
sns.scatterplot(x='culmen_length_mm', y='culmen_depth_mm', data=df, hue='species')
plt.xlabel('Culmen Length (mm)')
plt.ylabel('Culmen Depth (mm)')
plt.title('Culmen Length vs. Culmen Depth')
plt.show()
```



# • Multi-Variate Analysis



# 4. Perform descriptive statistics on the dataset.

1 [19]:	<pre>print(df.describe())</pre>					
		culmen_length_mm	culmen_depth_mm	flipper_length_mm	body_mass_g	
	count	342.000000	342.000000	342.000000	342.000000	
	mean	43.921930	17.151170	200.915205	4201.754386	
	std	5.459584	1.974793	14.061714	801.954536	
	min	32.100000	13.100000	172.000000	2700.000000	
	25%	39.225000	15.600000	190.000000	3550.000000	
	50%	44.450000	17.300000	197.000000	4050.000000	
	75%	48.500000	18.700000	213.000000	4750.000000	
	max	59,600000	21.500000	231.000000	6300.000000	

### 5. Check for Missing values and deal with them.

```
In [20]:
         print(df.isnull().sum())
         df.dropna(inplace=True)
         species
                                0
         island
                                0
         culmen length mm
                                2
         culmen depth mm
                                2
         flipper_length_mm
                                2
         body mass g
                                2
         sex
                               10
         dtype: int64
```

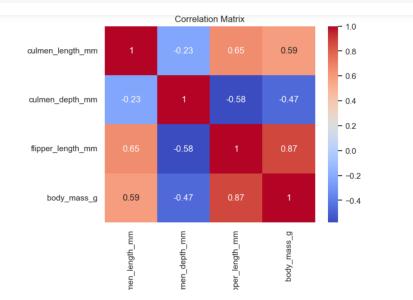
## 6. Find the outliers and replace them outliers

```
In [23]: from scipy import stats
z_scores = np.abs(stats.zscore(df.select_dtypes(include='number')))
threshold = 3
df_no_outliers = df[(z_scores < threshold).all(axis=1)]

print(f"Number of rows before removing outliers: {df.shape[0]}")
print(f"Number of rows after removing outliers: {df_no_outliers.shape[0]}")</pre>
Number of rows before removing outliers: 334
Number of rows after removing outliers: 334
```

# 7. Check the correlation of independent variables with the target

```
corr_matrix = df.corr()
sns.heatmap(corr_matrix, annot=True, cmap='coolwarm')
plt.title('Correlation Matrix')
plt.show()
```



### 8. Check for Categorical columns and perform encoding.

```
In [26]: categorical_columns = df.select_dtypes(include=['object']).columns.tolist()
         label_encoders = {}
         for col in categorical columns:
             le = LabelEncoder()
             df no outliers[col] = le.fit transform(df no outliers[col])
             label_encoders[col] = le
         for col, le in label encoders.items():
             print(f"Label encoding for {col}:")
             for label, code in zip(le.classes_, le.transform(le.classes_)):
                 print(f"{label}: {code}")
         Label encoding for island:
         Biscoe: 0
         Dream: 1
         Torgersen: 2
         Label encoding for sex:
         .: 0
         FEMALE: 1
         MALE: 2
```

# 9. Split the data into dependent and independent variables.

```
from sklearn.preprocessing import StandardScaler
scaler = standardScaler()
target_column = 'your_target_column_name_here'  # Replace with the actual target column name

columns_to_scale = [col for col in df_no_outliers.columns if col != target_column and df_no_outliers[col].dtype in ['int64', 'floif columns_to_scale:
    df_no_outliers[columns_to_scale] = scaler.fit_transform(df_no_outliers[columns_to_scale])

print("Mean of scaled features:")
    print(df_no_outliers[columns_to_scale].mean())
    print("Instandard deviation of scaled features:")
    print(df_no_outliers[columns_to_scale].std())
else:
    print("No numeric columns to scale.")
```

```
Mean of scaled features:
culmen length mm
                    -4.254747e-17
culmen_depth_mm
                    -1.276424e-16
flipper_length_mm
                     0.000000e+00
body_mass_g
                     4.254747e-17
dtype: float64
Standard deviation of scaled features:
culmen length mm
                     1.0015
culmen_depth_mm
                     1.0015
flipper_length_mm
                     1.0015
body mass g
                     1.0015
dtype: float64
```

#### 10. Scaling the data

```
scaler = StandardScaler()
target_column = 'species'

columns_to_scale = [col for col in X.columns if X[col].dtype in ['int64', 'float64']]
if columns_to_scale:
    X_scaled = scaler.fit_transform(X[columns_to_scale])
else:
    print("No numeric columns to scale.")

print("scaled Data:")
print(X_scaled[:5])

Scaled Data:
[[-0.89765322   0.78348666  -1.42952144  -0.57122888]
[-0.82429023   0.12189602  -1.07240838  -0.50901123]
[-0.67756427   0.42724555  -0.42960487  -1.19340546]
[-1.33783112   1.08883619  -0.5724501  -0.94453483]
[-0.86097173   1.75042684  -0.78671793  -0.6956642 ]]
```

#### 11. Split the data into training and testing

## 12.check the training and testing data shape.

```
[40]: # Split data into training and testing sets
X_train, X_test, y_train, y_test = train_test_split(X_scaled, y, test_size=0.2, random_state=42)

[41]: print(f"X_train shape: {X_train.shape}")
    print(f"X_test shape: {X_test.shape}")
    print(f"y_train shape: {y_train.shape}")
    print(f"y_test shape: {y_test.shape}")

X_train shape: (267, 4)
    X_test shape: (67, 4)
    y_train shape: (267,)
    y_test shape: (67,)
```