

AI&ML ASSIGNMENT-3

```
In [2]: import numpy as np
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt

from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
```

2. Loading the dataset

```
In [10]: df=pd.read_csv("penguins_size.csv")
df.head()
```

```
Out[10]:
```

	species	island	culmen_length_mm	culmen_depth_mm	flipper_length_mm	body_mass_g
0	Adelie	Torgersen	39.1	18.7	181.0	3750.0
1	Adelie	Torgersen	39.5	17.4	186.0	3800.0
2	Adelie	Torgersen	40.3	18.0	195.0	3250.0
3	Adelie	Torgersen	NaN	NaN	NaN	NaN
4	Adelie	Torgersen	36.7	19.3	193.0	3400.0



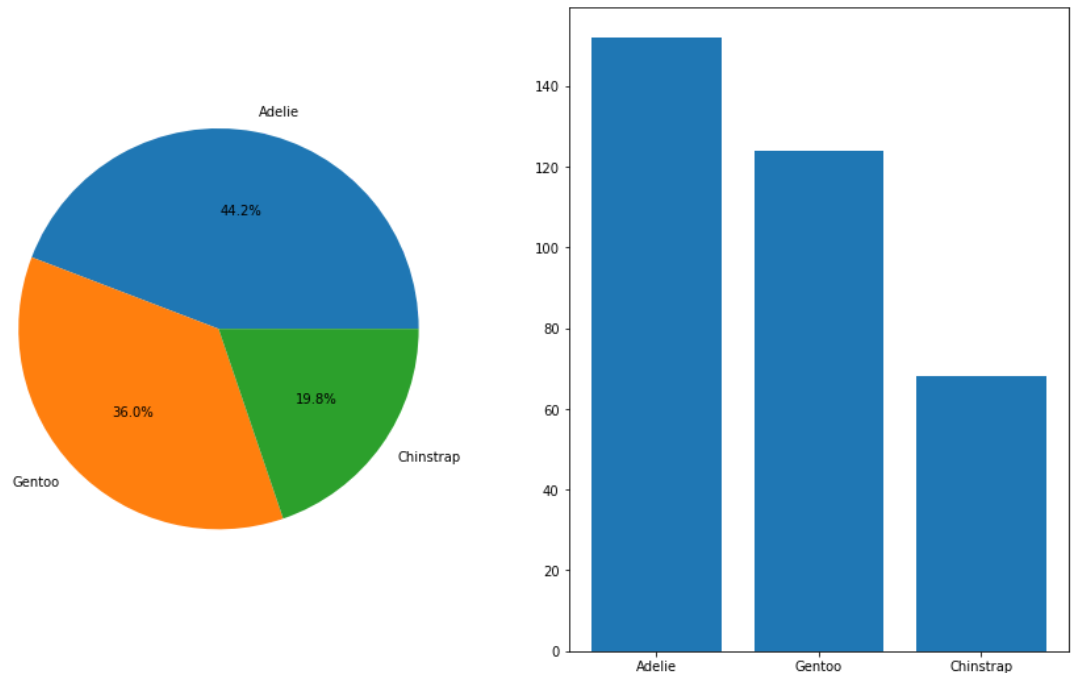
Univariate Analysis

```
In [12]: df['species'].value_counts()
```

```
Out[12]: Adelie      152
Gentoo      124
Chinstrap    68
Name: species, dtype: int64
```

```
In [254]: a=df['species'].value_counts().index
fig,ax = plt.subplots(1,2, figsize=(15,9))
ax[0].pie(df['species'].value_counts(),labels=a,autopct = "%1.1f%%")
ax[1].bar(a,df['species'].value_counts())
```

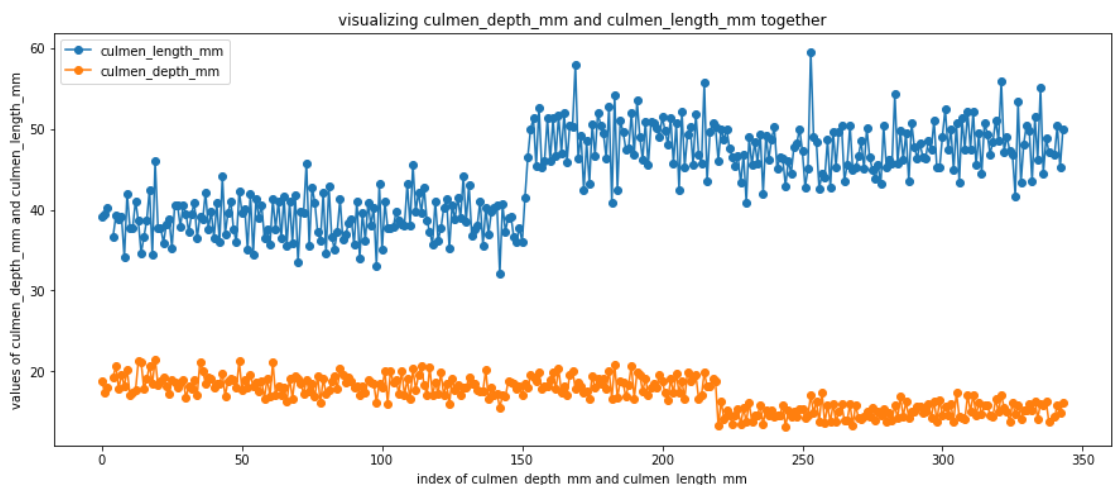
Out[254]: <BarContainer object of 3 artists>



Bi- Variate Analysis

```
In [262]: plt.figure(figsize=(15,6))
plt.plot(df['culmen_length_mm'],'o-')
plt.plot(df['culmen_depth_mm'],'o-')
plt.title("visualizing culmen_depth_mm and culmen_length_mm together")
plt.xlabel("index of culmen_depth_mm and culmen_length_mm")
plt.ylabel("values of culmen_depth_mm and culmen_length_mm")
plt.legend(['culmen_length_mm','culmen_depth_mm'])
```

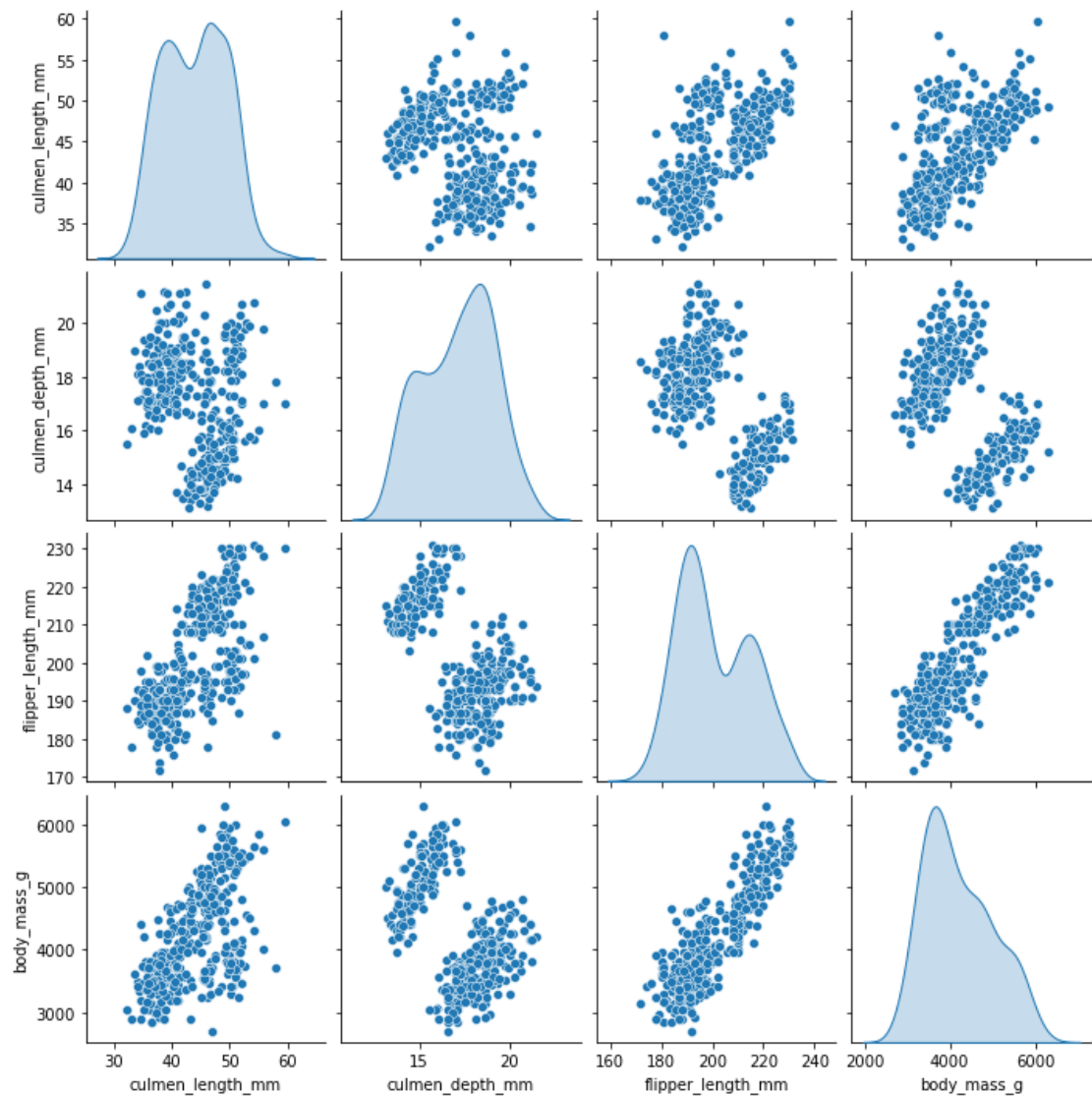
Out[262]: <matplotlib.legend.Legend at 0x1c1d4aad850>



Multi-Variate Analysis

```
In [266]: sns.pairplot(df,diag_kind='kde')
```

```
Out[266]: <seaborn.axisgrid.PairGrid at 0x1c1d5231a30>
```

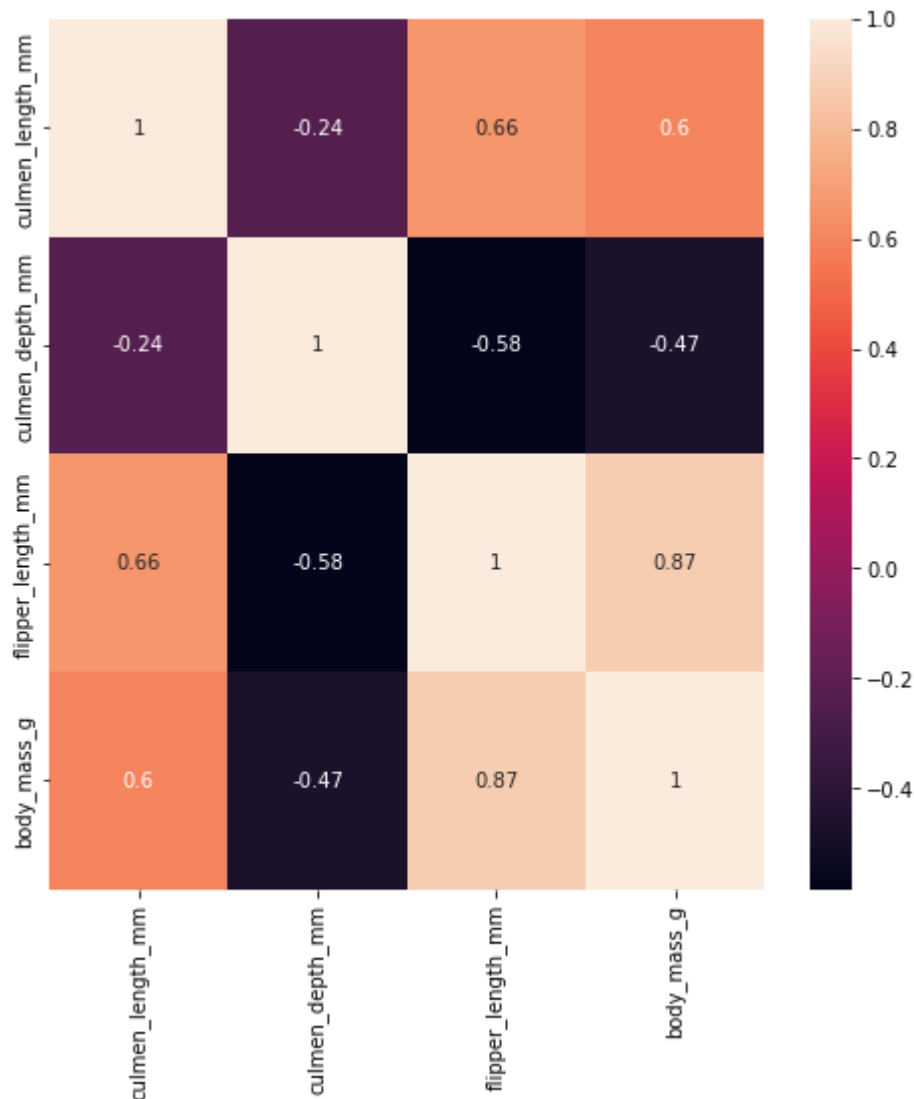


```
In [269]: sns.heatmap(df.corr(),annot=True)
```

C:\Users\HP\AppData\Local\Temp\ipykernel_15880\4277794465.py:1: FutureWarning: The default value of numeric_only in DataFrame.corr is deprecated. In a future version, it will default to False. Select only valid columns or specify the value of numeric_only to silence this warning.

```
sns.heatmap(df.corr(),annot=True)
```

Out[269]: <AxesSubplot:>



4. DESCRIPTIVE STATISTICS ON OUR DATASET

In [273]: `df.info()`

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 344 entries, 0 to 343
Data columns (total 7 columns):
#   Column                Non-Null Count  Dtype
---  -
0   species               344 non-null    object
1   island                344 non-null    object
2   culmen_length_mm       342 non-null    float64
3   culmen_depth_mm       342 non-null    float64
4   flipper_length_mm     342 non-null    float64
5   body_mass_g           342 non-null    float64
6   sex                   334 non-null    object
dtypes: float64(4), object(3)
memory usage: 18.9+ KB
```

In [274]: `df.dtypes`

```
Out[274]: species                object
island                  object
culmen_length_mm        float64
culmen_depth_mm         float64
flipper_length_mm       float64
body_mass_g             float64
sex                     object
dtype: object
```

In [275]: `df.describe()`

```
Out[275]:
```

	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. CHECKING FOR MISSING VALUES AND IMPUTING THEM WITH STATISTICAL METHODS

(MEAN FOR NUMERIC DATA AND MODE FOR CATEGORICAL DATA)

```
In [277]: df.isnull().sum() # columns except species and island are having nu  
#except sex columns remaining columns are of type float
```

```
Out[277]: species          0  
island          0  
culmen_length_mm    2  
culmen_depth_mm     2  
flipper_length_mm   2  
body_mass_g         2  
sex               10  
dtype: int64
```

```
In [4]: x=['culmen_length_mm', 'culmen_depth_mm','flipper_length_mm', 'body  
for column in x:  
    df[column].fillna(df[column].mean(),inplace =True)  
df['sex'].fillna(df['sex'].mode()[0],inplace =True)
```

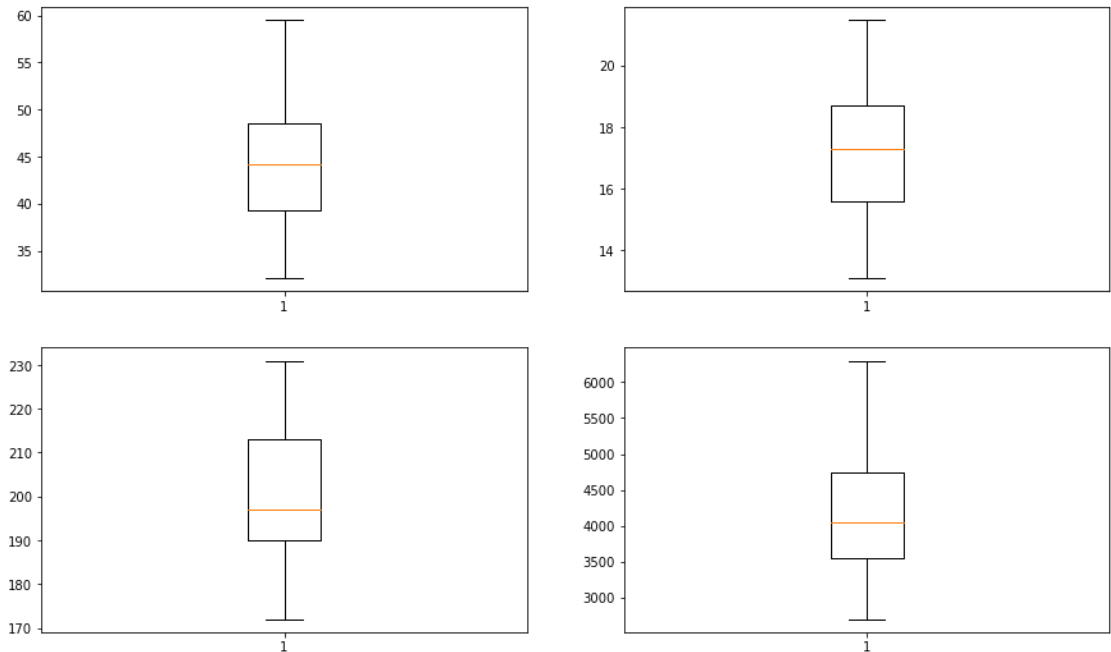
```
In [279]: #again checking for null values  
df.isnull().sum()
```

```
Out[279]: species          0  
island          0  
culmen_length_mm    0  
culmen_depth_mm     0  
flipper_length_mm   0  
body_mass_g         0  
sex                 0  
dtype: int64
```

6. FINDING THE OUTLIERS AND REPLACING THEM

```
In [280]: fig,ax = plt.subplots(2,2, figsize=(15,9))
ax[0][0].boxplot(df['culmen_length_mm'])
ax[0][1].boxplot(df['culmen_depth_mm'])
ax[1][0].boxplot(df['flipper_length_mm'])
ax[1][1].boxplot(df['body_mass_g']) #no outliers in following column
```

```
Out[280]: {'whiskers': [<matplotlib.lines.Line2D at 0x1c1d75d5eb0>,
<matplotlib.lines.Line2D at 0x1c1d75e2280>],
'caps': [<matplotlib.lines.Line2D at 0x1c1d75e2610>,
<matplotlib.lines.Line2D at 0x1c1d75e29a0>],
'boxes': [<matplotlib.lines.Line2D at 0x1c1d75d5b20>],
'medians': [<matplotlib.lines.Line2D at 0x1c1d75e2d30>],
'fliers': [<matplotlib.lines.Line2D at 0x1c1d75ef100>],
'means': []}
```



```
In [282]: def funct(col):
    print(col + "\n")
    q1 = df[col].quantile(0.25)
    q3 = df[col].quantile(0.75)
    print(f" First quartile of {col} is q1= {q1} \n Second quartile")
    iqr=q3-q1
    print(f" IQR OF {col} is {iqr}")
    upper_limit = q3+1.5*iqr
    lower_limit =q1-1.5*iqr
    print(f" Upper limit of {col} is: {upper_limit} \n Lower limit")
    print()
```

```
In [283]: z=['culmen_length_mm', 'culmen_depth_mm', 'flipper_length_mm', 'body',
for col in z:
    funct(col)
```

culmen_length_mm

First quartile of culmen_length_mm is q1= 39.275
 Second quartile of culmen_length_mm is q3= 48.5
 IQR OF culmen_length_mm is 9.225000000000001
 Upper limit of culmen_length_mm is: 62.337500000000006
 Lower limit of culmen_length_mm is: 25.437499999999996

culmen_depth_mm

First quartile of culmen_depth_mm is q1= 15.6
 Second quartile of culmen_depth_mm is q3= 18.7
 IQR OF culmen_depth_mm is 3.0999999999999996
 Upper limit of culmen_depth_mm is: 23.349999999999998
 Lower limit of culmen_depth_mm is: 10.95

flipper_length_mm

First quartile of flipper_length_mm is q1= 190.0
 Second quartile of flipper_length_mm is q3= 213.0
 IQR OF flipper_length_mm is 23.0
 Upper limit of flipper_length_mm is: 247.5
 Lower limit of flipper_length_mm is: 155.5

body_mass_g

First quartile of body_mass_g is q1= 3550.0
 Second quartile of body_mass_g is q3= 4750.0
 IQR OF body_mass_g is 1200.0
 Upper limit of body_mass_g is: 6550.0
 Lower limit of body_mass_g is: 1750.0

7. CHECKING THE CORRELATION OF INDEPENDENT VARIABLES WITH TARGET

```
In [18]: df.corr().species.sort_values(ascending=False)
```

```
Out[18]: species            1.000000
flipper_length_mm    0.854307
body_mass_g          0.750491
culmen_length_mm     0.731369
sex                  0.002262
island              -0.635659
culmen_depth_mm     -0.744076
Name: species, dtype: float64
```

8. CHECKING FOR CATEGORICAL COLUMNS AND PERFORMING


```
In [14]: df.species.value_counts()
```

```
Out[14]: Adelie      152
          Gentoo     124
          Chinstrap   68
          Name: species, dtype: int64
```

```
In [15]: df.island.value_counts()
```

```
Out[15]: Biscoe      168
          Dream      124
          Torgersen   52
          Name: island, dtype: int64
```

```
In [16]: df['sex']=df['sex'].replace(".", "MALE")
          df.sex.value_counts()
```

```
Out[16]: MALE      169
          FEMALE    165
          Name: sex, dtype: int64
```

```
In [17]: from sklearn.preprocessing import LabelEncoder
          le = LabelEncoder()

          df.species= le.fit_transform(df.species)
          df.island= le.fit_transform(df.island)
          df.sex = le.fit_transform(df.sex)
```

```
In [290]: df.head()
```

```
Out[290]:
```

	species	island	culmen_length_mm	culmen_depth_mm	flipper_length_mm	body_mass_g
0	0	2	39.10000	18.70000	181.000000	3750.00000
1	0	2	39.50000	17.40000	186.000000	3800.00000
2	0	2	40.30000	18.00000	195.000000	3250.00000
3	0	2	43.92193	17.15117	200.915205	4201.75438
4	0	2	36.70000	19.30000	193.000000	3450.00000

9. SPLIT THE DATA INTO DEPENDENT AND INDEPENDENT VARIABLES.

```
In [291]: X=df.drop(columns=['sex'],axis=1) #dependent variables
          y=df.sex #independent variables
```

10. SCALING THE DATA

```
In [292]: from sklearn.preprocessing import MinMaxScaler
          scale =MinMaxScaler()
```

```
In [293]: X_scaled= pd.DataFrame(scale.fit_transform(X),columns =X.columns)
X_scaled.head()
```

```
Out[293]:
```

	species	island	culmen_length_mm	culmen_depth_mm	flipper_length_mm	body_mass_g
0	0.0	1.0	0.254545	0.666667	0.152542	0.29166
1	0.0	1.0	0.269091	0.511905	0.237288	0.30555
2	0.0	1.0	0.298182	0.583333	0.389831	0.15277
3	0.0	1.0	0.429888	0.482282	0.490088	0.41715
4	0.0	1.0	0.167273	0.738095	0.355932	0.20833

11. SPLIT THE DATA INTO TRAINING AND TESTING

```
In [294]: from sklearn.model_selection import train_test_split
X_train,X_test,y_train,y_test = train_test_split(X_scaled,y,test_si
```

12.CHECK THE TRAINING AND TESTING DATA SHAPE.

```
In [295]: X_train.shape
```

```
Out[295]: (275, 6)
```

```
In [296]: X_test.shape
```

```
Out[296]: (69, 6)
```

```
In [297]: y_train.shape
```

```
Out[297]: (275,)
```

```
In [298]: y_test.shape
```

```
Out[298]: (69,)
```