1. penguins_size.csv is downloaded

▼ 2. Load the dataset into the tool.

```
import numpy as np
import pandas as pd

df = pd.read_csv('/content/penguins_size.csv')
df.head()
```

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

▼ 3.1. Perform Univariate Analysis

from matplotlib import rcParams
import seaborn as sns

sns.distplot(df.body_mass_g)

<ipython-input-4-176964dae727>:1: UserWarning:

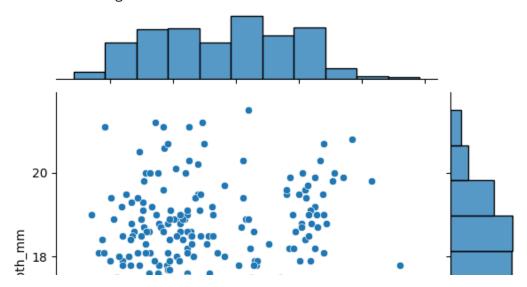
`distplot` is a deprecated function and will be removed in sea

Please adapt your code to use either `displot` (a figure-level similar flexibility) or `histolot` (an axes-level function for

▼ 3.2. Perform Bivariate Analysis

sns.jointplot(x='culmen_length_mm',y='culmen_depth_mm',data=df)

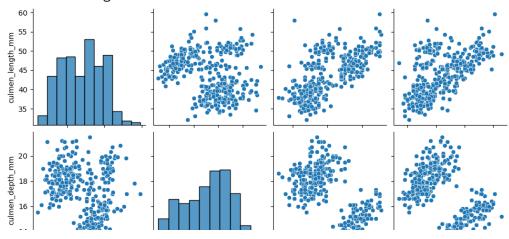
<seaborn.axisgrid.JointGrid at 0x7c313325c6a0>



▼ 3.3. Perform Multi-Variate Analysis

sns.pairplot(df)

<seaborn.axisgrid.PairGrid at 0x7c31298f71f0>



▼ 4. Perform descriptive statistics on the dataset.

df.describe()

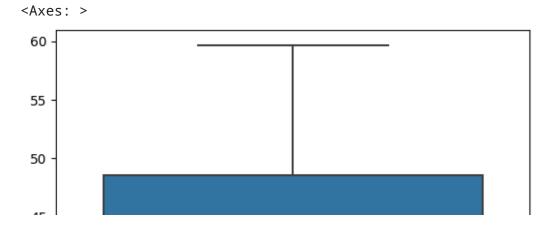
▼ 5. Check for Missing values and deal with them.

count

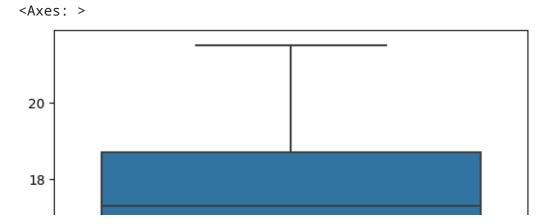
```
df.isnull().any() #Checking is there any null values in our dataset
                            False
     species
     island
                            False
     culmen_length_mm
                             True
     culmen_depth_mm
                             True
     flipper_length_mm
                             True
     body_mass_g
                             True
     sex
                             True
     dtype: bool
df.isnull().sum()
     species
                             0
     island
                             0
     culmen_length_mm
                             2
                             2
     culmen_depth_mm
                             2
     flipper length mm
                             2
     body_mass_g
                            10
     sex
     dtype: int64
# Code to replace null values in numerical columns with MEDIAN
df['culmen_length_mm'].fillna(df['culmen_length_mm'].median(),inplace=True)
df['culmen_depth_mm'].fillna(df['culmen_depth_mm'].median(),inplace=True)
df['flipper_length_mm'].fillna(df['flipper_length_mm'].median(),inplace=True)
df['body_mass_g'].fillna(df['body_mass_g'].median(),inplace=True)
# Code to replace null values in categorical column with MODE
df['sex'].fillna(df['sex'].mode().iloc[0],inplace=True)
# Now all null values are replaced with median and mode and dealt properly.
df.isnull().any()
     species
                            False
     island
                            False
     culmen_length_mm
                            False
     culmen depth mm
                            False
     flipper_length_mm
                            False
                            False
     body_mass_g
                            False
     sex
     dtype: bool
```

▼ 6. Find the outliers and replace the outliers

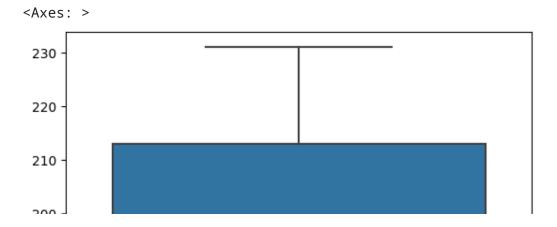
sns.boxplot(df.culmen_length_mm)



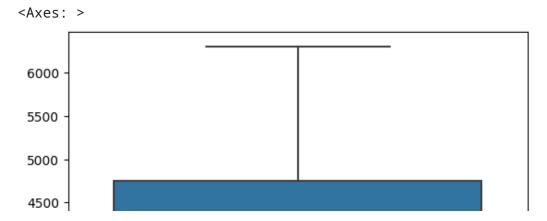
sns.boxplot(df.culmen_depth_mm)



sns.boxplot(df.flipper_length_mm)



sns.boxplot(df.body_mass_g)



Hence there are no outliers in the dataset.

▼ 7. Check for Categorical columns and perform encoding.

```
from sklearn.preprocessing import LabelEncoder
le = LabelEncoder()
df['sex'] = le.fit_transform(df['sex'])
df['species'] = le.fit_transform(df['species'])
df['island'] = le.fit_transform(df['island'])
df.head()
```

	species	island	culmen_length_mm	culmen_depth_mm	flipper_
0	0	2	39.10	18.7	
1	0	2	39.50	17.4	
2	0	2	40.30	18.0	
3	0	2	44.45	17.3	
4	0	2	36.70	19.3	

8. Check the correlation of independent variables with the target (TARGET IS SPECIES and remaining are independent)

df.corr().species.sort_values(ascending=False)

```
      species
      1.000000

      flipper_length_mm
      0.850819

      body_mass_g
      0.747547

      culmen_length_mm
      0.728706

      sex
      -0.003823

      island
      -0.635659

      culmen_depth_mm
      -0.741282

      Name: species, dtype: float64
```

▼ 9. Split the data into dependent and independent variables

```
X=df.drop(columns=['species'],axis=1)
X.head()
```

	island	culmen_length_mm	culmen_depth_mm	flipper_length_m
0	2	39.10	18.7	181.
1	2	39.50	17.4	186.
2	2	40.30	18.0	195.

Y=df['species']
Y.head()

Name: species, dtype: int64

▼ 10. Scaling the data

from sklearn.preprocessing import MinMaxScaler
scale = MinMaxScaler()
X_scaled = pd.DataFrame(scale.fit_transform(X),columns=X.columns)
X_scaled.head()

	island	culmen_length_mm	culmen_depth_mm	flipper_length_m
0	1.0	0.254545	0.666667	0.15254
1	1.0	0.269091	0.511905	0.23728
2	1.0	0.298182	0.583333	0.38983
3	1.0	0.449091	0.500000	0.42372
4	1.0	0.167273	0.738095	0.35593

▼ 11. Split the data into training and testing

from sklearn.model_selection import train_test_split
X_train,X_test,Y_train,Y_test = train_test_split(X_scaled,Y,test_size=0.2,random_state=0)

▼ 12. Check the training and testing data shape.

X_train.shape (275, 6)

The End!!!

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