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### ASSINMENT 3

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
```

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

```
df.head()
```

	species	island	culmen_length_mm	culmen_depth_mm	flipper_length_mm	body_mass_g	sex
0	Adelie	Torgersen	39.1	18.7	181.0	3750.0	MALE
1	Adelie	Torgersen	39.5	17.4	186.0	3800.0	FEMALE
2	Adelie	Torgersen	40.3	18.0	195.0	3250.0	FEMALE
3	Adelie	Torgersen	NaN	NaN	NaN	NaN	NaN
4	Adelie	Torgersen	36.7	19.3	193.0	3450.0	FEMALE

```
df.shape
```



```
(344, 7)
```

```
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
```

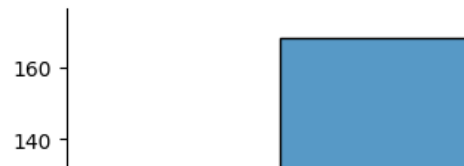
```
df.describe()
```

	culmen_length_mm	culmen_depth_mm	flipper_length_mm	body_mass_g	
<b>count</b>	342.000000	342.000000	342.000000	342.000000	
<b>mean</b>	43.921930	17.151170	200.915205	4201.754386	
<b>std</b>	5.459584	1.974793	14.061714	801.954536	
<b>min</b>	32.100000	13.100000	172.000000	2700.000000	
<b>25%</b>	39.225000	15.600000	190.000000	3550.000000	
<b>50%</b>	44.450000	17.300000	197.000000	4050.000000	
<b>75%</b>	48.500000	18.700000	213.000000	4750.000000	
<b>max</b>	59.600000	21.500000	231.000000	6300.000000	

## ▼ univariate analysis

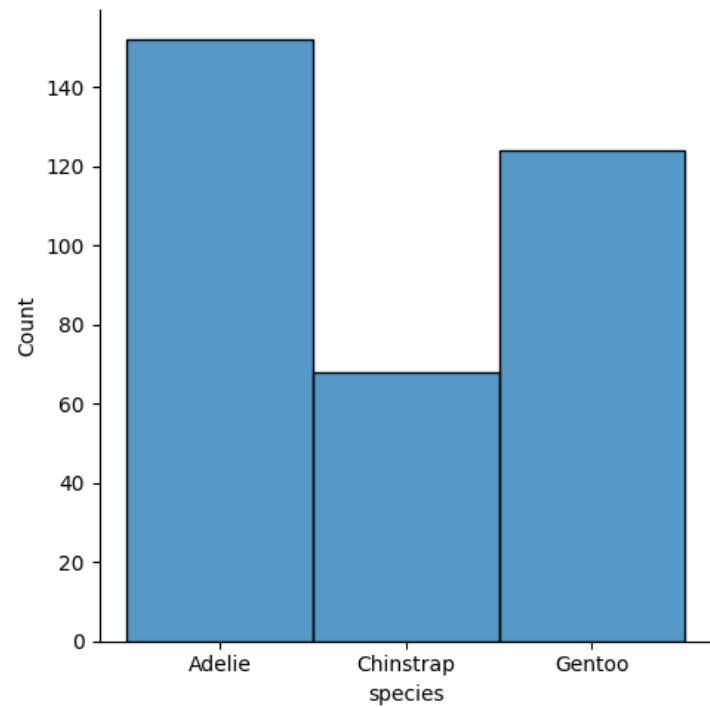
```
sns.displot(df.island)
```

```
<seaborn.axisgrid.FacetGrid at 0x78c3542c7610>
```



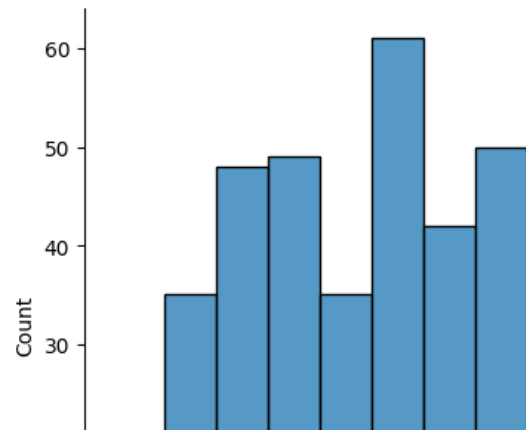
```
sns.displot(df.species)
```

```
<seaborn.axisgrid.FacetGrid at 0x78c3546e6260>
```



```
sns.displot(df.culmen_length_mm)
```

<seaborn.axisgrid.FacetGrid at 0x78c3542c7820>



```
sns.distplot(df.flipper_length_mm )
```

```
<ipython-input-13-ae65ebdd98e7>:1: UserWarning:
```

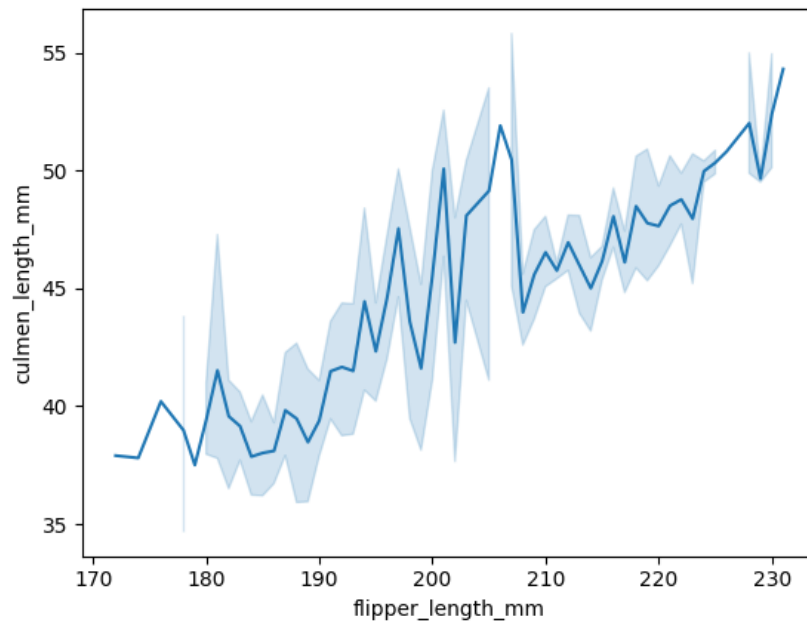
```
`distplot` is a deprecated function and will be removed in seaborn v0.14.0.
```

```
similar flexibility) or `histplot` (an axes-level function for histograms).
```

## ▼ bivariate analysis

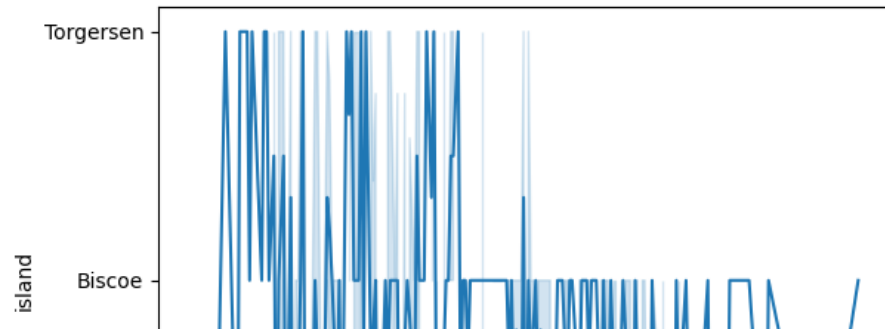
```
sns.lineplot(x=df.flipper_length_mm,y=df.culmen_length_mm)
```

```
<Axes: xlabel='flipper_length_mm', ylabel='culmen_length_mm'>
```



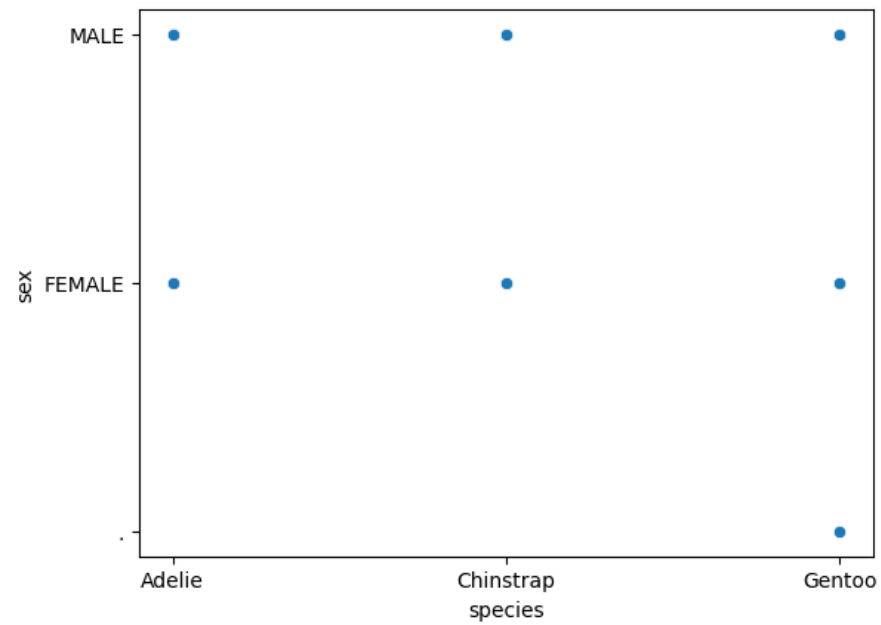
```
sns.lineplot(x=df.culmen_length_mm,y=df.island)
```

```
<Axes: xlabel='culmen_length_mm', ylabel='island'>
```



```
sns.scatterplot(x=df.species, y=df.sex)
```

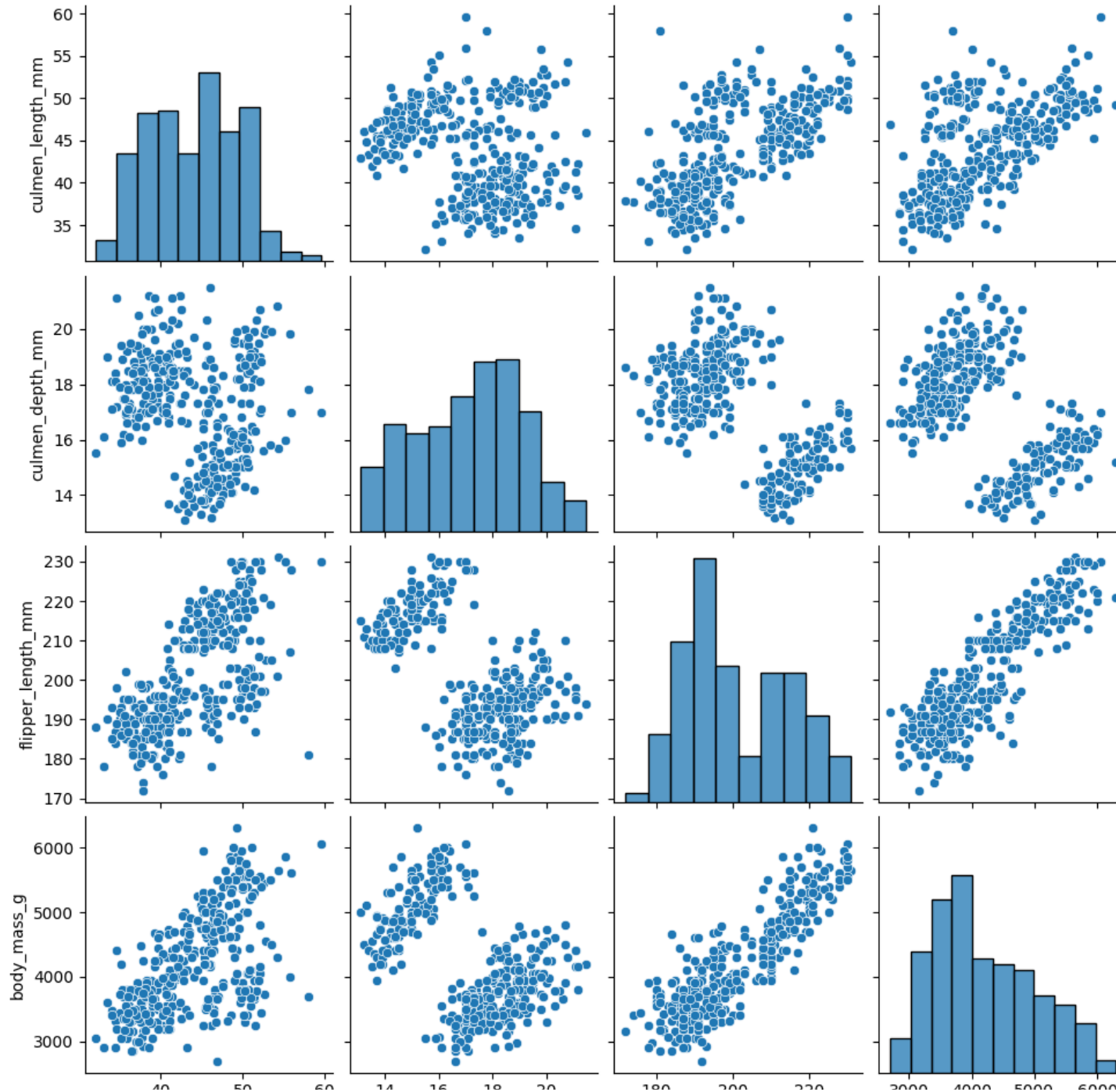
```
<Axes: xlabel='species', ylabel='sex'>
```



## ▼ Multivariate analysis

```
sns.pairplot(df)
```

&lt;seaborn.axisgrid.PairGrid at 0x78c34f890370&gt;

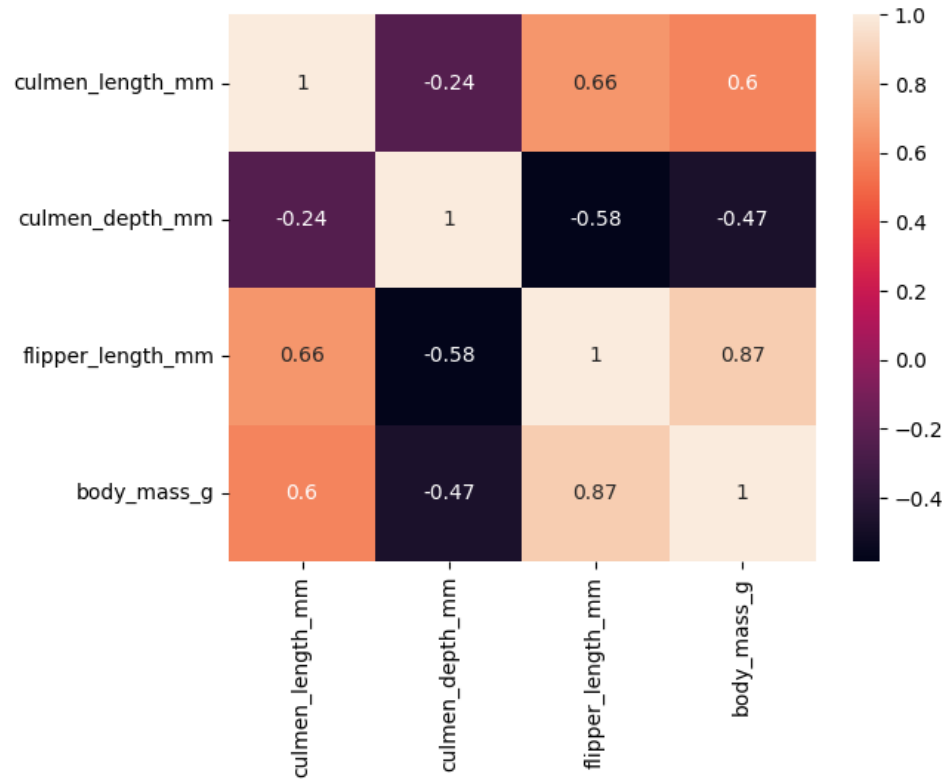


```
40      50      60      14      16      20      180      200      220      3000      4000      5000      6000
      culmen_length_mm      culmen_depth_mm      flipper_length_mm      body_mass_g
```

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

```
<ipython-input-19-8df7bcac526d>:1: FutureWarning: The default value of numeric_only in DataFrame.corr is deprecated. In a future version, it will default to False. Select only
sns.heatmap(df.corr(),annot=True)
```

```
<Axes: >
```





Double-click (or enter) to edit



## ▼ descriptive statistics

```
df.describe()
```



	culmen_length_mm	culmen_depth_mm	flipper_length_mm	body_mass_g	
<b>count</b>	342.000000	342.000000	342.000000	342.000000	
<b>mean</b>	43.921930	17.151170	200.915205	4201.754386	
<b>std</b>	5.459584	1.974793	14.061714	801.954536	
<b>min</b>	32.100000	13.100000	172.000000	2700.000000	
<b>25%</b>	39.225000	15.600000	190.000000	3550.000000	
<b>50%</b>	44.450000	17.300000	197.000000	4050.000000	

```
df.head()
```

	species	island	culmen_length_mm	culmen_depth_mm	flipper_length_mm	body_mass_g	sex	
<b>0</b>	Adelie	Torgersen	39.1	18.7	181.0	3750.0	MALE	
<b>1</b>	Adelie	Torgersen	39.5	17.4	186.0	3800.0	FEMALE	
<b>2</b>	Adelie	Torgersen	40.3	18.0	195.0	3250.0	FEMALE	
<b>3</b>	Adelie	Torgersen	NaN	NaN	NaN	NaN	NaN	
<b>4</b>	Adelie	Torgersen	36.7	19.3	193.0	3450.0	FEMALE	

```
df.isnull().any()
```

```
species      False
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
culmen_depth_mm    2
flipper_length_mm    2
body_mass_g        2
sex           10
dtype: int64
```

```
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)
df['sex'].fillna(df['sex'].mode()[0],inplace=True)
```

```
df.isnull().sum()
```

```
species      0
island        0
culmen_length_mm  0
culmen_depth_mm  0
flipper_length_mm  0
body_mass_g   0
sex           0
dtype: int64
```

```
df.head()
```

	species	island	culmen_length_mm	culmen_depth_mm	flipper_length_mm	body_mass_g	sex
0	Adelie	Torgersen	39.10	18.7	181.0	3750.0	MALE
1	Adelie	Torgersen	39.50	17.4	186.0	3800.0	FEMALE
2	Adelie	Torgersen	40.30	18.0	195.0	3250.0	FEMALE
3	Adelie	Torgersen	44.45	17.3	197.0	4050.0	MALE
4	Adelie	Torgersen	36.70	19.3	193.0	3450.0	FEMALE

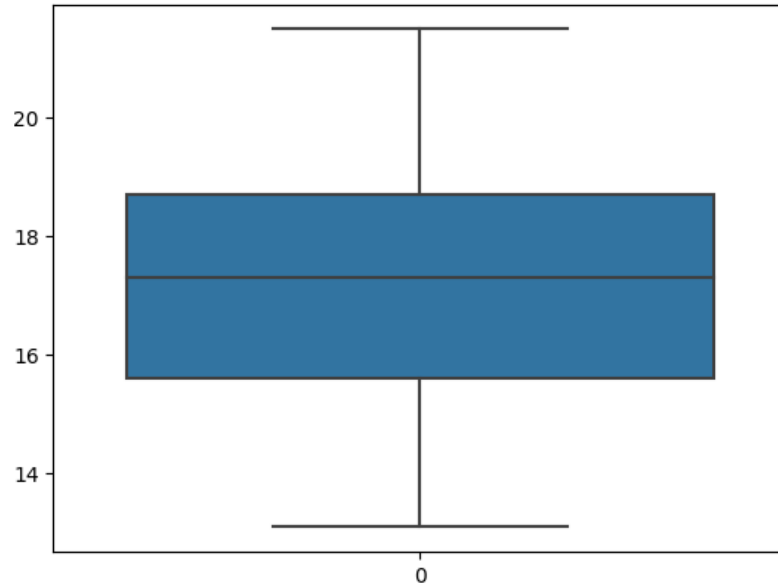
```
sns.boxplot(df['culmen_length_mm'])
```

&lt;Axes: &gt;

60

```
sns.boxplot(df['culmen_depth_mm'])
```

&lt;Axes: &gt;



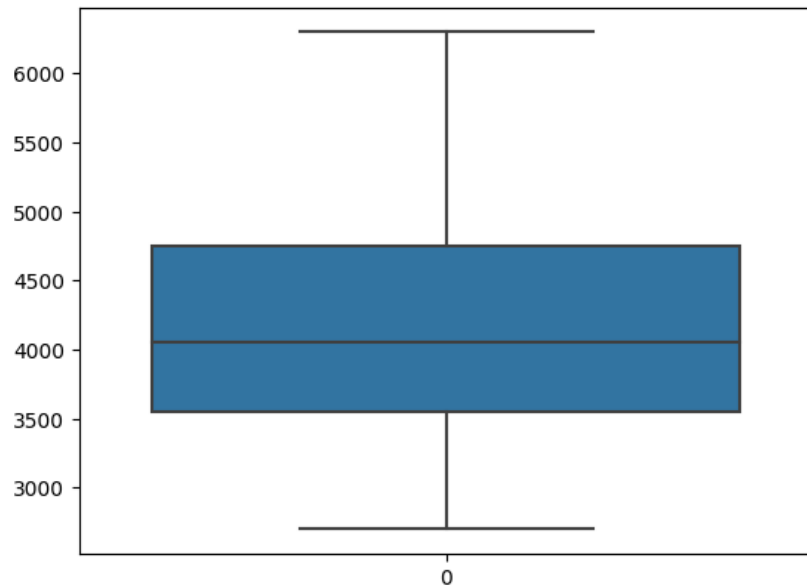
```
sns.boxplot(df['flipper_length_mm'])
```

&lt;Axes: &gt;



```
sns.boxplot(df['body_mass_g'])
```

&lt;Axes: &gt;



## ▼ 7 checking correlation

```
df.corr()
```

```
<ipython-input-44-2f6f6606aa2c>:1: FutureWarning: The default value of numeric_only in DataFrame.corr is deprecated. In a future version, it will default to False. Select only
```

## ▼ 8 check for categorical columns and perform encoding

```
from sklearn.preprocessing import LabelEncoder
le=LabelEncoder()
```

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

```
df.head()
```

	species	island	culmen_length_mm	culmen_depth_mm	flipper_length_mm	body_mass_g	sex
0	0	2	39.10	18.7	181.0	3750.0	2
1	0	2	39.50	17.4	186.0	3800.0	1
2	0	2	40.30	18.0	195.0	3250.0	1
3	0	2	44.45	17.3	197.0	4050.0	2
4	0	2	36.70	19.3	193.0	3450.0	1

## ▼ finding correlation between target column and all other columns after encoding the target column into numerical column

```
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
```

## ▼ split the data into dependent and independent variable

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

	island	culmen_length_mm	culmen_depth_mm	flipper_length_mm	body_mass_g	sex
0	2	39.10	18.7	181.0	3750.0	2
1	2	39.50	17.4	186.0	3800.0	1
2	2	40.30	18.0	195.0	3250.0	1
3	2	44.45	17.3	197.0	4050.0	2
4	2	36.70	19.3	193.0	3450.0	1

```
y=df['species']
y
```

```
0    0
1    0
2    0
3    0
4    0
..
339  2
340  2
341  2
342  2
343  2
Name: species, Length: 344, dtype: int64
```

## ▼ scaling a 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_mm	body_mass_g	sex	
0	1.0	0.254545	0.666667	0.152542	0.291667	1.0	
1	1.0	0.269091	0.511905	0.237288	0.305556	0.5	
2	1.0	0.298182	0.583333	0.389831	0.152778	0.5	
3	1.0	0.449091	0.500000	0.423729	0.375000	1.0	
4	1.0	0.167273	0.738095	0.355932	0.208333	0.5	

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=0.3,random_state=0)
```

x\_train

	island	culmen_length_mm	culmen_depth_mm	flipper_length_mm	body_mass_g	sex	
219	0.5	0.658182	0.666667	0.440678	0.298611	0.5	
271	0.0	0.596364	0.119048	0.813559	0.722222	1.0	
266	0.0	0.487273	0.095238	0.644068	0.416667	0.5	
335	0.0	0.836364	0.345238	0.983051	0.875000	1.0	
217	0.5	0.636364	0.607143	0.355932	0.298611	1.0	
...	...	...	...	...	...	...	
323	0.0	0.618182	0.226190	0.949153	0.777778	1.0	
192	0.5	0.614545	0.761905	0.644068	0.347222	1.0	
117	1.0	0.189091	0.880952	0.457627	0.298611	1.0	
47	0.5	0.196364	0.690476	0.118644	0.076389	1.0	
172	0.5	0.374545	0.500000	0.152542	0.250000	0.5	

240 rows × 6 columns

y\_train

```
219    1
271    2
266    2
335    2
```

```

217 1
    ..
323 2
192 1
117 0
47  0
172 1
Name: species, Length: 240, dtype: int64

```

x\_test

	island	culmen_length_mm	culmen_depth_mm	flipper_length_mm	body_mass_g	sex
141	0.5	0.309091	0.488095	0.254237	0.215278	1.0
6	1.0	0.247273	0.559524	0.152542	0.256944	0.5
60	0.0	0.130909	0.452381	0.220339	0.125000	0.5
249	0.0	0.650909	0.261905	0.813559	0.791667	1.0
54	0.0	0.087273	0.595238	0.254237	0.055556	0.5
...	...	...	...	...	...	...
81	1.0	0.392727	0.535714	0.406780	0.555556	1.0
1	1.0	0.269091	0.511905	0.237288	0.305556	0.5
120	1.0	0.149091	0.488095	0.254237	0.125000	0.5
8	1.0	0.072727	0.595238	0.355932	0.215278	1.0
313	0.0	0.632727	0.357143	0.881356	0.819444	1.0

104 rows × 6 columns

y\_test

```

141 0
6    0
60   0
249 2
54   0
    ..
81   0
1    0
120 0
8    0
313 2
Name: species, Length: 104, dtype: int64

```

x\_train.shape



```
(240, 6)
```

```
x_test.shape
```

```
(104, 6)
```

```
y_train.shape
```

```
(240,)
```

```
y_test.shape
```

```
(104,)
```

---

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