Assignment -3

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Department	INFORMATION TECHNOLOGY

Problem Statement: Abalone Age Prediction

Description:

Predicting the age of abalone from physical measurements. The age of abalone is determined by cutting the shell through the cone, staining it, and counting the number of rings through a microscope -- a boring and time-consuming task. Other measurements, which are easier to obtain, are used to predict age. Further information, such as weather patterns and location (hence food availability) may be required to solve the problem.

Importing Modules

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

In [1]:

1. Dataset has been downloaded

#Name of the dataset: abalone.csv

2. Load the dataset into the tool

data=pd.read_csv("abalone.csv")
data.head()

Out[12]:

Sex Lenoth Diameter Height Whole weight Shucked weight Viscera weight Shell weight Rings

	Sex	Length	Diameter	Height	Whole weight	Shucked weight	Viscera weight	Shell weight	Rings
0	- M	0.455	0.365	0.095	0.5140	0.2245	0.1010	0.150	15
1	М	0.350	0.265	0.090	0.2255	0.0995	0.0485	0.070	7
2	F	0.530	0.420	0.135	0.6770	0.2565	0.1415	0.210	9
3	М	0.440	0.365	0.125	0.5160	0.2155	0.1140	0.155	10

	Sex	Length	Diameter	neight	whole weight	Shuckeu weight	visceta weight	onen weight	Kings
4		0.330	0.255	0.080	0.2050	0.0895	0.0395	0.055	7

Let's know the shape of the data

data.shape

Out[13]:

One additional task is that, we have to add the "Age" column using "Rings" data. We just have to add '1.5' to the ring data

Out[14]:

Length Diameter Height Whole_weight Shucked_weight Viscera_weight Shell_weight 0.5140 0.455 0.365 0.095 0.2245 0.1010 0.150 0.350 0.265 0.090 0.2255 0.0995 0.0485 0.070 0.530 0.420 0.135 0.6770 0.2565 0.1415 0.210 0.440 0.365 0.125 0.5160 0.2155 0.1140 0.155 0.255 0.080 0.2050 0.0895 0.0395 0.055

3. Perform Below Visualizations.

(i) Univariate Analysis

#

The term univariate analysis refers to the analysis of one variable. You can remember this because the prefix "uni" means "one." There are three common ways to perform univariate analysis on one variable: 1. Summary statistics – Measures the center and spread of values.

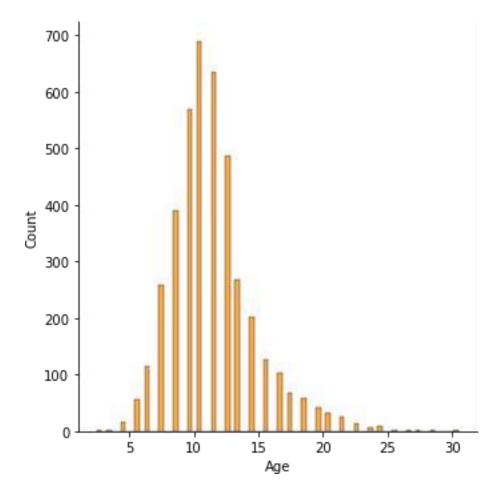
Histogram

sns.displot(data["Age"], color='darkorange')

In [16]:

Out[16]:

<seaborn.axisgrid.FacetGrid at 0x7fd3f837a430>

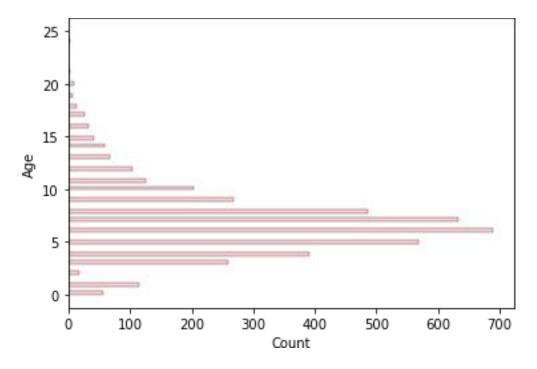


sns.histplot(y=data.Age,color='pink')

In [103]:

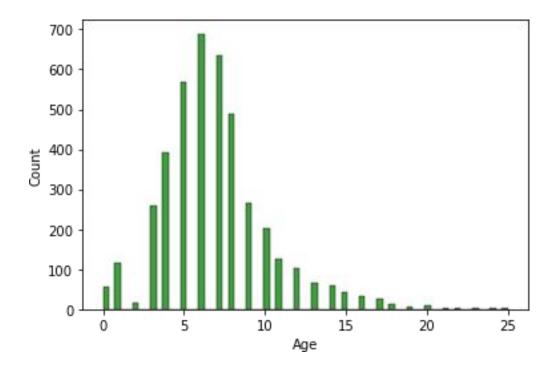
Out[103]:

<AxesSubplot:xlabel='Count', ylabel='Age'>



sns.histplot(x=data.Age,color='green')

<AxesSubplot:xlabel='Age', ylabel='Count'>



Boxplot

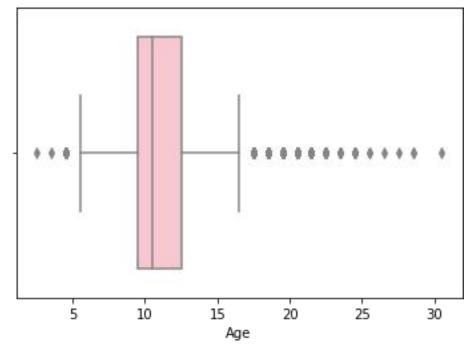
sns.boxplot(x=data.Age,color='pink')

In [106]:

Out[106]:

In [52]:

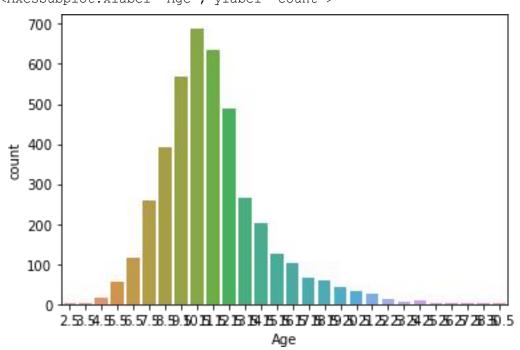
<AxesSubplot:xlabel='Age'>



Countplot

sns.countplot(x=data.Age)

<AxesSubplot:xlabel='Age', ylabel='count'>



In [51]:

Out[51]:

(ii) Bi-Variate Analysis

#

Image result for bivariate analysis in python It is a methodical statistical technique applied to a pair of variables (features/ attributes) of data to determine the empirical relationship between them. In order words, it is meant to determine any concurrent relations (usually over and above a simple correlation analysis).

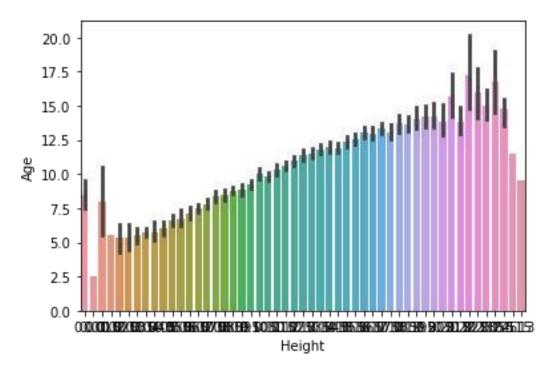


Barplot

sns.barplot(x=data.Height, y=data.Age)
<AxesSubplot:xlabel='Height', ylabel='Age'>

In [50]:

Out[50]:

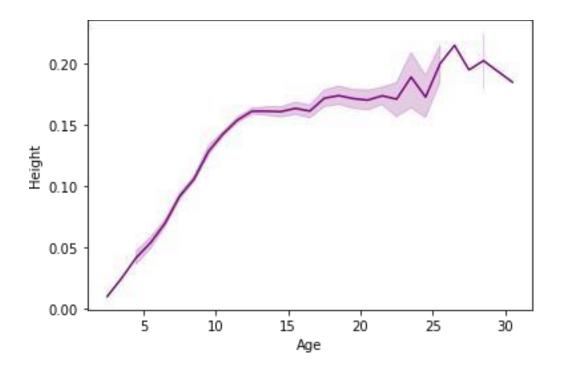


Linearplot

sns.lineplot(x=data.Age,y=data.Height, color='purple')
<AxesSubplot:xlabel='Age', ylabel='Height'>

In [49]:

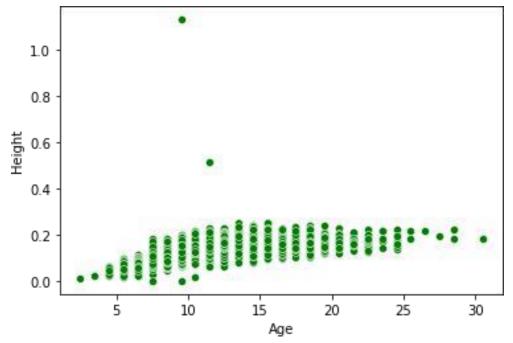
Out[49]:



Scatterplot

sns.scatterplot(x=data.Age,y=data.Height,color='green')

 $\verb| <AxesSubplot:xlabel='Age', ylabel='Height'> \\$

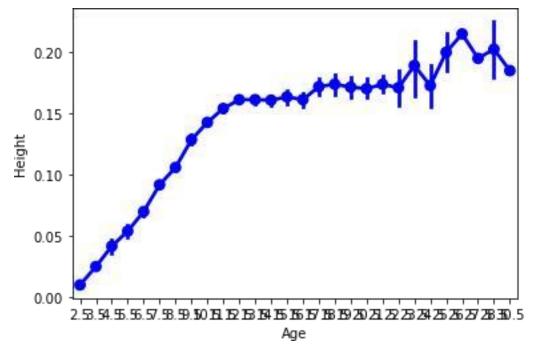


In [42]:

Out[42]:

Pointplot

sns.pointplot(x=data.Age, y=data.Height, color="blue")
<AxesSubplot:xlabel='Age', ylabel='Height'>



Regplot

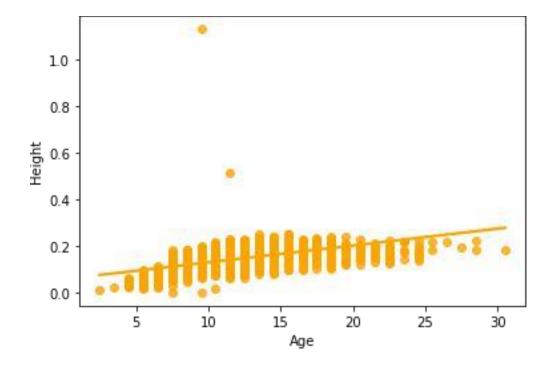
sns.regplot(x=data.Age,y=data.Height,color='orange')
<AxesSubplot:xlabel='Age', ylabel='Height'>

In [45]:

Out[45]:

In [48]:

Out[48]:



(iii) Multi-Variate Analysis

#

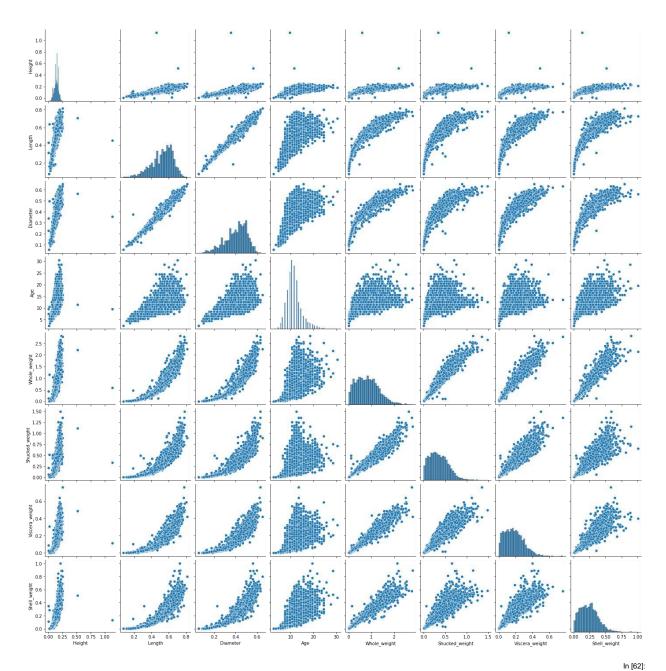
Multivariate analysis is based in observation and analysis of more than one statistical outcome variable at a time. In design and analysis, the technique is used to perform trade studies across multiple dimensions while taking into account the effects of all variables on the responses of interest.

#

Pairplot

sns.pairplot(data=data[["Height","Length","Diameter","Age","Whole_weight","Sh ucked_weight","Viscera_weight","Shell_weight"]])

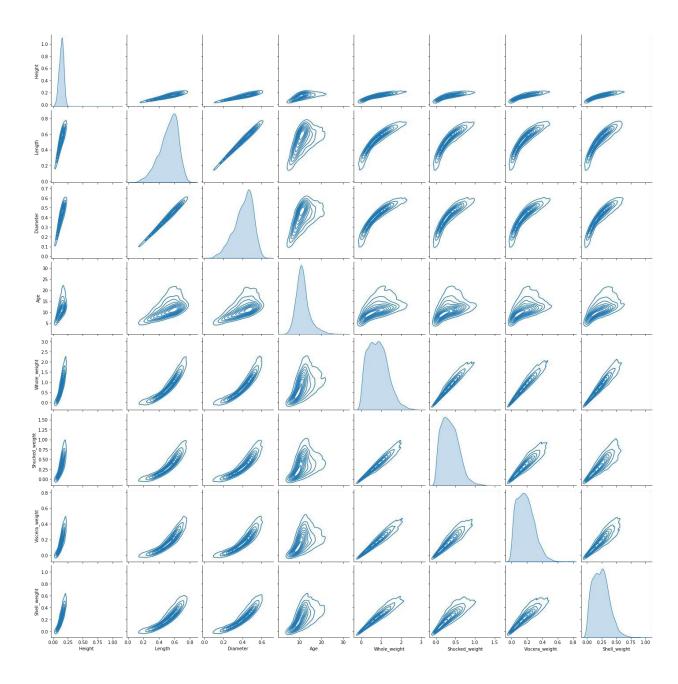
<seaborn.axisgrid.PairGrid at 0x7fd3d93e1040>



sns.pairplot(data=data[["Height","Length","Diameter","Age","Whole_weight","Sh
ucked_weight","Viscera_weight","Shell_weight"]],kind="kde")

Out[62]:

<seaborn.axisgrid.PairGrid at 0x7fd39840c790>



4. Perform descriptive statistics on the dataset

| Count | 4177 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 | 4177.000000 |

	Sex	Length	Diameter	Height	Whole_weight	Shucked_weight	Viscera_weight	Shell_weight	Age
top	М	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN
freq	1528	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN
mean	NaN	0.523992	0.407881	0.139516	0.828742	0.359367	0.180594	0.238831	11.433684
std	NaN	0.120093	0.099240	0.041827	0.490389	0.221963	0.109614	0.139203	3.224169
min	NaN	0.075000	0.055000	0.000000	0.002000	0.001000	0.000500	0.001500	2.500000
25%	NaN	0.450000	0.350000	0.115000	0.441500	0.186000	0.093500	0.130000	9.500000
50%	NaN	0.545000	0.425000	0.140000	0.799500	0.336000	0.171000	0.234000	10.500000
75%	NaN	0.615000	0.480000	0.165000	1.153000	0.502000	0.253000	0.329000	12.500000
max	NaN	0.815000	0.650000	1.130000	2.825500	1.488000	0.760000	1.005000	30.500000

5. Check for Missing values and deal with them

data.isnull().sum()

Sex 0

Length 0

Diameter 0

Height 0

Whole_weight 0

Shucked_weight 0

Viscera_weight 0

Shell_weight 0

Age 0

dtype: int64

6. Find the outliers and replace them outliers

outliers=data.quantile(q=(0.25,0.75))
outliers

	Length	Diameter	Height	Whole_weight	Shucked_weight	Viscera_weight	Shell_weight	Age
0.25	0.450	0.35	0.115	0.4415	0.186	0.0935	0.130	0.5

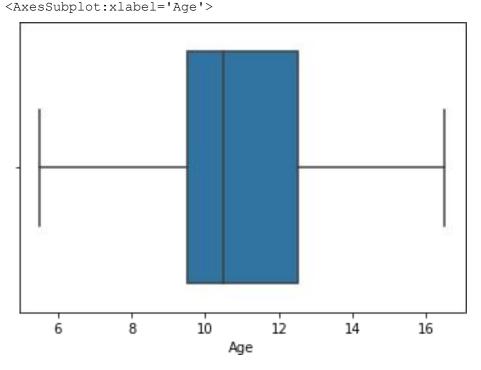
In [64]:

Out[64]:

In [65]:

Out[65]:

```
Length
                    Whole_weight
                           Shucked_weight
                                    Viscera_weight
                                             Shell_weight
                                                                                         In [66]:
a = data.Age.quantile(0.25)
b = data.Age.quantile(0.75)
c = b - a
lower limit = a - 1.5 * c
data.median(numeric only=True)
                                                                                         Out[66]:
Length
                      0.5450
Diameter
                      0.4250
Height
                      0.1400
Whole weight
                      0.7995
Shucked weight
                      0.3360
Viscera weight
                      0.1710
Shell weight
                      0.2340
                     10.5000
dtype: float64
                                                                                         In [67]:
data['Age'] = np.where(data['Age'] < lower limit, 7, data['Age'])</pre>
sns.boxplot(x=data.Age, showfliers = False)
                                                                                         Out[67]:
```



7. Check for Categorical columns and perform encoding

data.head()

Out[68]:

In [68]:

```
Height
                                  Whole_weight
                                                 Shucked_weight Viscera_weight
Length
                                                                                      Shell_weight
                                        0.5140
              0.365
                        0.095
                                                          0.2245
                                                                                            0.150
 0.350
                                        0.2255
              0.265
                        0.090
                                                          0.0995
                                                                            0.0485
                                                                                            0.070
 0.530
              0.420
                        0.135
                                        0.6770
                                                          0.2565
                                                                            0.1415
                                                                                            0.210
                                                          0.2155
              0.365
                        0.125
                                        0.5160
                                                                                            0.155
```

from sklearn.preprocessing import LabelEncoder

```
lab = LabelEncoder()
data.Sex = lab.fit_transform(data.Sex)
```

data.head()

	Sex	Length	Diameter	Height	Whole_weight	Shucked_weight	Viscera_weight	Shell_weight	Age
0	2	0.455	0.365	0.095	0.5140	0.2245	0.1010	0.150	12
1	2	0.350	0.265	0.090	0.2255	0.0995	0.0485	0.070	4
2	0	0.530	0.420	0.135	0.6770	0.2565	0.1415	0.210	6
3	2	0.440	0.365	0.125	0.5160	0.2155	0.1140	0.155	7
4	1	0.330	0.255	0.080	0.2050	0.0895	0.0395	0.055	4

8. Split the data into dependent and independent variables

Out[83]:

```
y = data["Sex"]
y.head()

Out[84]:

0      2
1      2
2      0
3      2
4      1
Name: Sex, dtype: int64

x=data.drop(columns=["Sex"],axis=1)
x.head()

Out[85]:

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```

	Length	Diameter	Height	Whole_weight	Shucked_weight	Viscera_weight	Shell_weight	Age
0	0.455	0.365	0.095	0.5140	0.2245	0.1010	0.150	12
1	0.350	0.265	0.090	0.2255	0.0995	0.0485	0.070	4
2	0.530	0.420	0.135	0.6770	0.2565	0.1415	0.210	6
3	0.440	0.365	0.125	0.5160	0.2155	0.1140	0.155	7
4	0.330	0.255	0.080	0.2050	0.0895	0.0395	0.055	4

9. Scale the independent variables

from sklearn.preprocessing import scale
X_Scaled = pd.DataFrame(scale(x), columns=x.columns)
X_Scaled.head()

Whole_weight Shucked_weight Length Viscera_weight -0.574558 -1.064424 -0.607685 -0.432149 -0.641898 -0.726212 -0.638217 1.555152 -1.448986 -1.439929 -1.183978 -1.230277 -1.170910 -1.205221 -1.212987 -0.884841 0.050033 0.122130 -0.107991 -0.309469 -0.463500 -0.356690 -0.207139 -0.274842 -0.699476 -0.432149 -0.347099 -0.637819 -0.648238 -0.607600 -0.602294 0.030157 -1.423087 -1.215968 -1.287337 -1.615544 -1.540707 -1.272086 -1.320757 -0.884841

10. 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)

X_Train.shape, X_Test.shape

((3341, 8), (836, 8))

Y_Train.shape, Y_Test.shape

Out[89]:
((3341,), (836,))
```

In [86]:

Out[86]:

In [87]:

X_7	Train.	.head(()					
	Length	Diameter	Height	Whole_weight	Shucked_weight	Viscera_weight	Shell_weight	Age
3141	-2.864726	-2.750043	-1.423087	-1.622870	-1.553902	-1.583867	-1.644065	-1.799838
3521	-2.573250	-2.598876	-2.020857	-1.606554	-1.551650	-1.565619	-1.626104	-1.494839
883	1.132658	1.230689	0.728888	1.145672	1.041436	0.286552	1.538726	1.555152
3627	1.590691	1.180300	1.446213	2.164373	2.661269	2.330326	1.377072	0.030157
2106	0.591345	0.474853	0.370226	0.432887	0.255175	0.272866	0.906479	1.250153
Х	Test.	nead()						
_	Length	Diameter	Height	Whole_weight	Shucked_weight	Viscera_weight	Shell_weight	Age
668	0.216591	0.172519	0.370226	0.181016	-0.368878	0.569396	0.690940	0.945154
1580	-0.199803	-0.079426	-0.466653	-0.433875	-0.443224	-0.343004	-0.325685	-0.579842
3784	0.799543	0.726798	0.370226	0.870348	0.755318	1.764639	0.565209	0.335156
463	-2.531611	-2.447709	-2.020857	-1.579022	-1.522362	-1.538247	-1.572219	-1.799838
2615	1.007740	0.928354	0.848442	1.390405	1.415417	1.778325	0.996287	0.640155
Y_7	Train.	.head(()					
314 352		1 1						
883 362	3	2 2						
210 Nar		2 ex, dt	ype:	int64				
Y_7	Test.	nead()						
668	3	2						

11. Build the Model

Name: Sex, dtype: int64

1 2

```
In [94]:
from sklearn.ensemble import RandomForestClassifier
model = RandomForestClassifier(n estimators=10,criterion='entropy')
                                                                                 In [95]:
model.fit(X Train, Y Train)
                                                                                 Out[95]:
RandomForestClassifier(criterion='entropy', n_estimators=10)
                                                                                 In [96]:
y predict = model.predict(X Test)
                                                                                 In [97]:
y predict train = model.predict(X Train)
12. Train the Model
                                                                                 In [98]:
from sklearn.metrics import
accuracy_score,confusion_matrix,classification_report
                                                                                 In [99]:
print('Training accuracy: ',accuracy score(Y Train,y predict train))
Training accuracy: 0.9787488775815624
13.Test the Model
                                                                                 In [100]:
print('Testing accuracy: ',accuracy score(Y Test,y predict))
Testing accuracy: 0.5526315789473685
14. Measure the performance using Metrics
                                                                                 In [101]:
pd.crosstab(Y Test, y predict)
                                                                                Out[101]:
col_0 0 1 2
 0 122 29 98
   37 217 37
 2 120 53 123
                                                                                 In [102]:
print(classification_report(Y_Test,y_predict))
               precision recall f1-score support
                   0.44 0.49 0.46
0.73 0.75 0.74
0.48 0.42 0.44
                                                     249
```

1

291

296

accuracy			0.55	836
macro avg	0.55	0.55	0.55	836
weighted avg	0.55	0.55	0.55	836