Assignment Date		
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Student Roll Number	111519104013	
Maximum Marks	2 Marks	

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
```

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	Length	Diameter	Height	Whole weight	Shucked weight	Viscera weight	Shell weight	Rings
0	М	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	Height	Whole weight	Shucked weight	Viscera weight	Shell weight	Rings
4	I	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

In [14]: Age=1.5+data.Rings data["Age"]=Age data=data.rename(columns = {'Whole weight':'Whole weight','Shucked weight': 'Shucked weight', 'Viscera weight': 'Viscera weight', 'Shell weight': 'Shell weight'}) data=data.drop(columns=["Rings"],axis=1) data.head() Out[14]: Height Whole_weight Viscera_weight Shucked_weight Shell_weight 0.365 0.095 0.5140 0.2245 0.1010 0.150 0.090 0.265 0.070 0.6770 0.2565 0.1415 0.420 0.135 0.210

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.

#

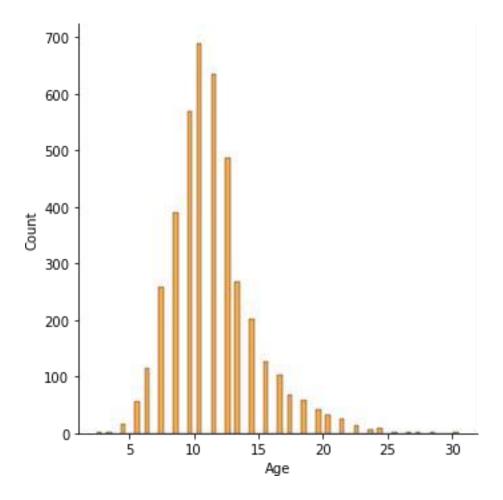
Histogram

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

Out[16]:

In[16]:

<seaborn.axisgrid.FacetGrid at 0x7fd3f837a430>

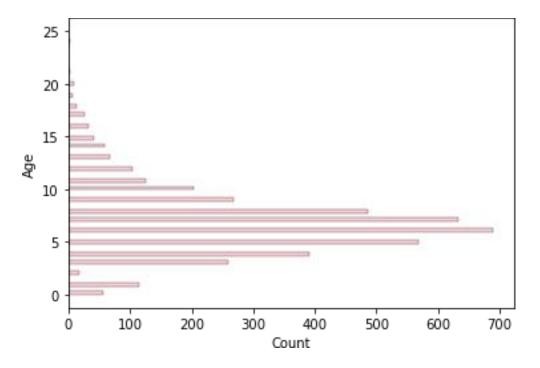


In[103]:

Out[103]:

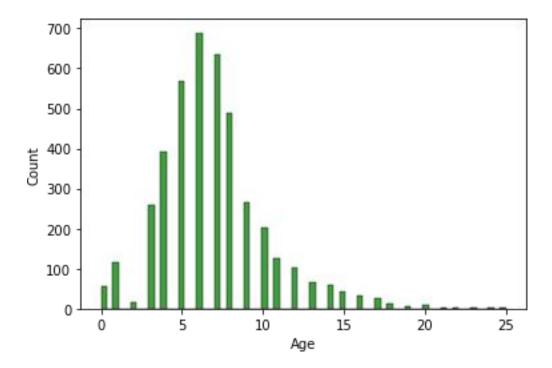
sns.histplot(y=data.Age,color='pink')

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



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

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



Boxplot

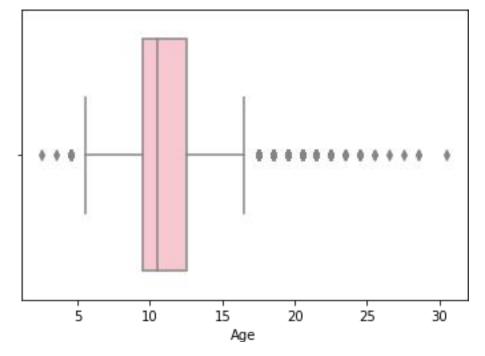
sns.boxplot(x=data.Age,color='pink')

In[106]:

Out[106]:

In [52]:

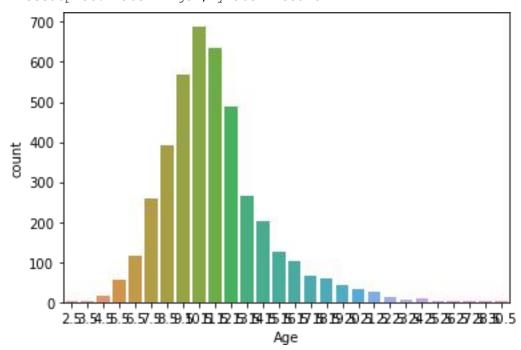
<AxesSubplot:xlabel='Age'>



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).

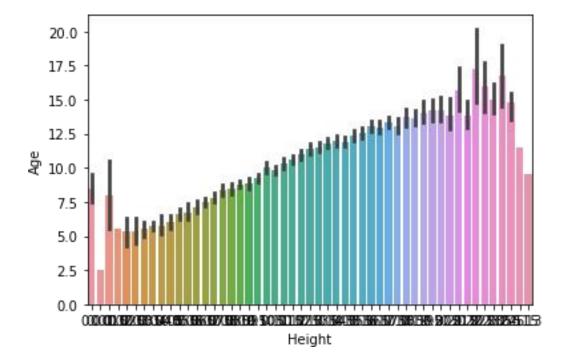


Barplot

sns.barplot(x=data.Height,y=data.Age)

Out[50]:

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

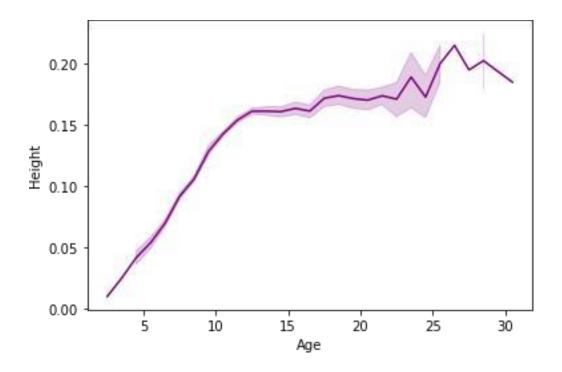


Linearplot

sns.lineplot(x=data.Age,y=data.Height, color='purple')

Out[49]:

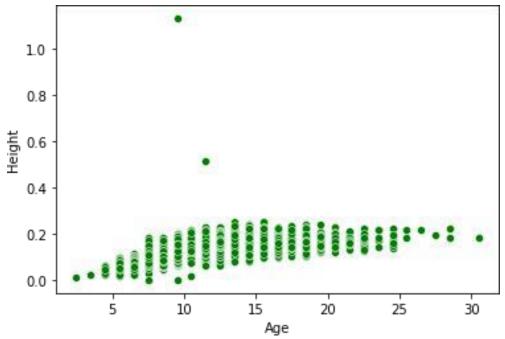
<AxesSubplot:xlabel='Age', ylabel='Height'>



Scatterplot

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

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



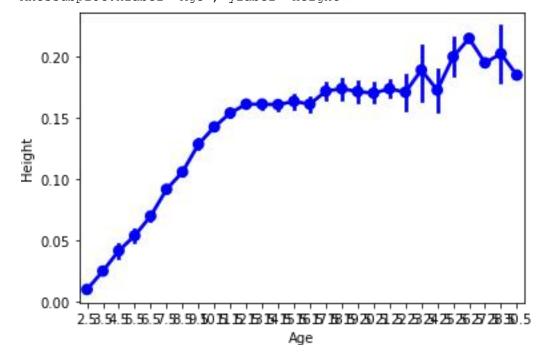
In [42]:

Out[42]:

Pointplot

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

In [45]: Out[45]:

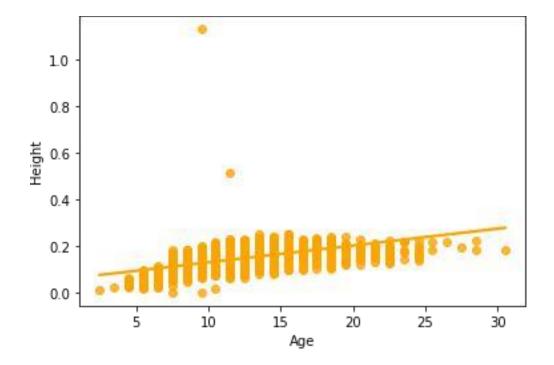


Regplot

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

In [48]:

Out[48]:



(iii) Multi-Variate Analysis

#

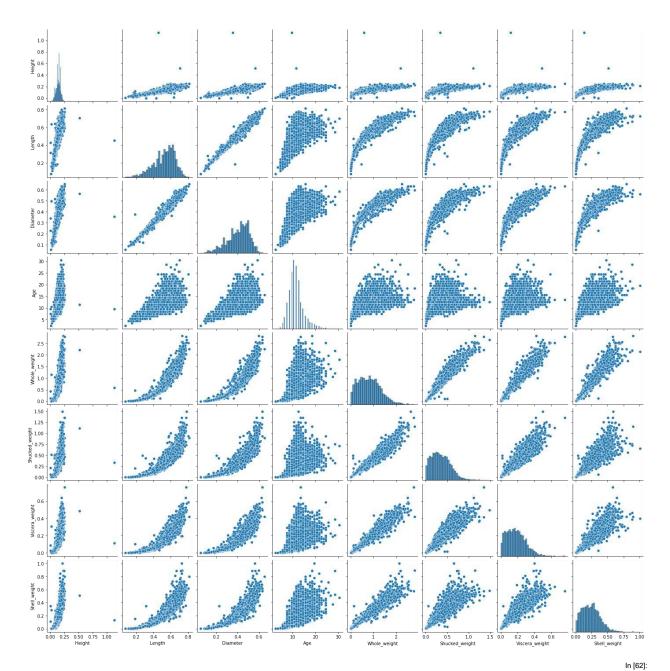
Multivariate analysis is based in observation and analysis of more than one statistical outcome variable at a time. In design and analysis, the technique is used to perform trade studies across multiple dimensions while taking into account the effects of all variables on the responses of interest.

#

Pairplot

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

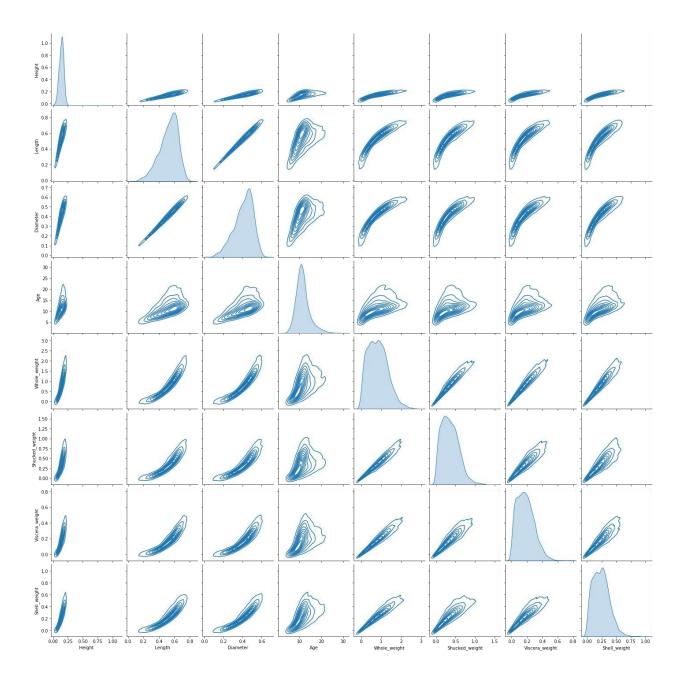
<seaborn.axisgrid.PairGrid at 0x7fd3d93e1040>



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

6. Find the outliers and replace them outliers

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

dtype: int64

 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
 9.5

In [65]:

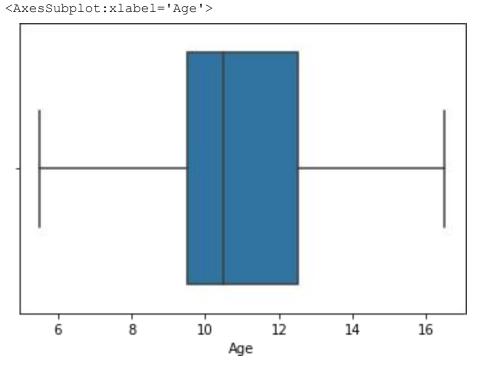
In [64]:

Out[64]:

Out[65]:

```
0.615
           0.48
               0.165
                      1.1530
                               0.502
                                       0.2530
                                              0.329
                                                   12.5
                                                                                        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
                      0.1400
Height
                      0.7995
Whole weight
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]:
```

Shell_weight



Height

Whole_weight

Shucked_weight

Viscera_weight

7. Check for Categorical columns and perform encoding

data.head()
Out[68]:

In [68]:

```
Height
                                   Whole_weight
                                                   Shucked_weight
                                                                      Viscera_weight
Length
           Diameter
                                                                                         Shell_weight
 0.455
                         0.095
                                         0.5140
                                                            0.2245
                                                                              0.1010
              0.365
                                                                                               0.150
                                                                                               0.070
                         0.090
                                                            0.0995
              0.420
                         0.135
                                                            0.2565
                                                                                               0.210
                                                                                               0.155
                                                            0.0895
                                                                                               0.055
```

 $\textbf{from} \ \text{sklearn.preprocessing} \ \textbf{import} \ \texttt{LabelEncoder}$

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

data.head()

Whole_weight Shucked_weight Viscera_weight 0.455 0.365 0.095 0.5140 0.2245 0.1010 0.150 0.350 0.2255 0.265 0.090 0.0995 0.0485 0.070 0.440 0.125 0.2155 0.1140 0.155 0.255 0.055

8. Split the data into dependent and independent variables

```
In [84]:
y = data["Sex"]
y.head()
                                                                                                Out[84]:
0
      2
      2
1
      0
3
      2
Name: Sex, dtype: int64
                                                                                                 In [85]:
x=data.drop(columns=["Sex"],axis=1)
x.head()
                                                                                                Out[85]:
```

In [83]:

Out[83]:

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

In[88]:
X_Train.shape, X_Test.shape

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

Y_Train.shape, Y_Test.shape

Out[89]:
((3341,), (836,))
In[90]:
```

In [86]:

Out[86]:

In [87]:

X_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	
X_T	est.h	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_T	rain.	head(()						
314 352		1 1							
883 362	} !7	2 2							
	ne: Se		ype:	int64					
Y_Test.head()									

Out[93]:

11. Build the Model

Name: Sex, dtype: int64

668 2 1580 1

3784 2 463 1 2615 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]:
   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
0.73 0.75
0.48 0.42
            0
                             0.49
                                        0.46
                                                     249
```

0.74

0.44

0.42

291

296

accurac	:y		0.55	836
macro av	rg 0.55	0.55	0.55	836
weighted av	rg 0.55	0.55	0.55	836