| Assignment Date | |
|---------------------|---------------------------|
| Student Name | ANJURU SAI SUNITH KARTHIK |
| Student Roll Number | 111519104006 |
| 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.

In [1]:

Importing Modules

0.440

0.365

0.125

0.5160

0.2155

```
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

```
In[12]:
data=pd.read csv("abalone.csv")
data.head()
                                                                                                                                                                     Out[12]:
            Length
                                Height
                                         Whole weight
                                                       Shucked weight
                                                                       Viscera weight
                                                                                      Shell weight
             0.455
                        0.365
                                 0.095
                                              0.5140
                                                              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
```

0.1140

0.155

| | Sex | Length | Diameter | Height | whole weight | Shucked weight | viscera weight | Such 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

Height Whole weight Shucked weight Viscera weight 0.455 0.365 0.095 0.5140 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 0.365 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

#

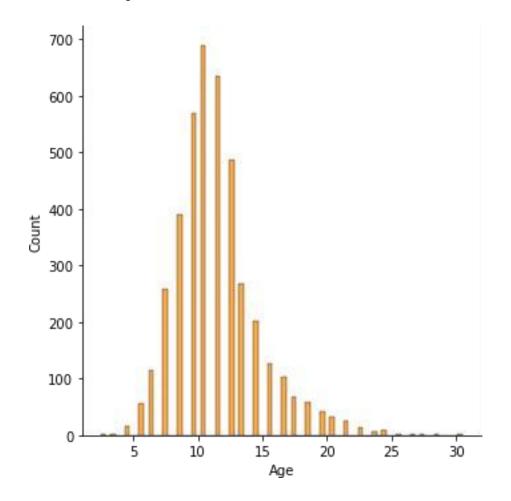
The term univariate analysis refers to the analysis of one variable. You can remember this because the prefix "uni" means "one." There are three common ways to perform univariate analysis on one variable: 1. Summary statistics – Measures the center and spread of values.

Histogram

sns.displot(data["Age"], color='darkorange')
<seaborn.axisgrid.FacetGrid at 0x7fd3f837a430>

In[16]:

Out[16]:

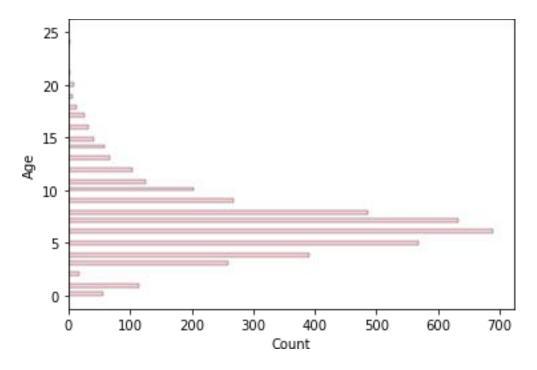


In[103]:

Out[103]:

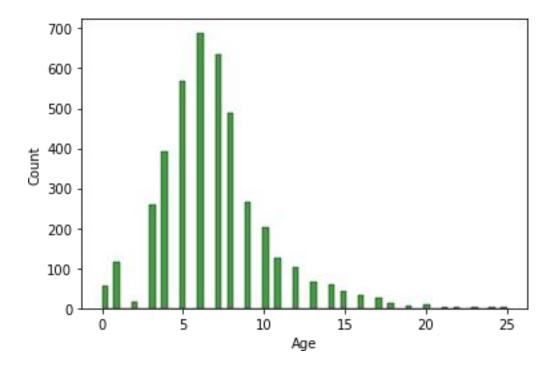
sns.histplot(y=data.Age,color='pink')

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



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

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



Boxplot

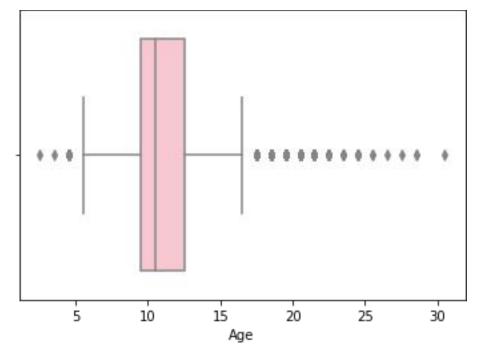
sns.boxplot(x=data.Age,color='pink')

In[106]:

Out[106]:

In [52]:

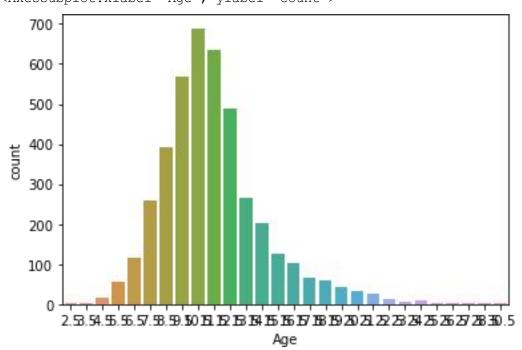
<AxesSubplot:xlabel='Age'>



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).

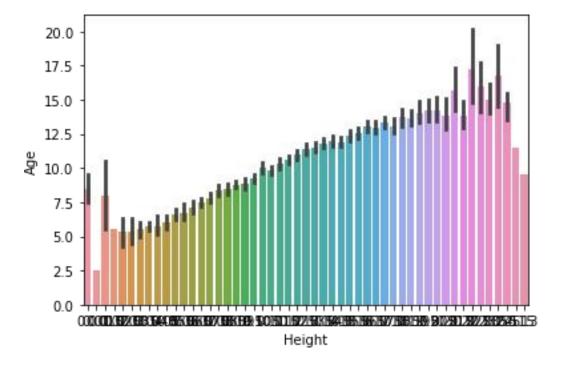


Barplot

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

Out[50]:

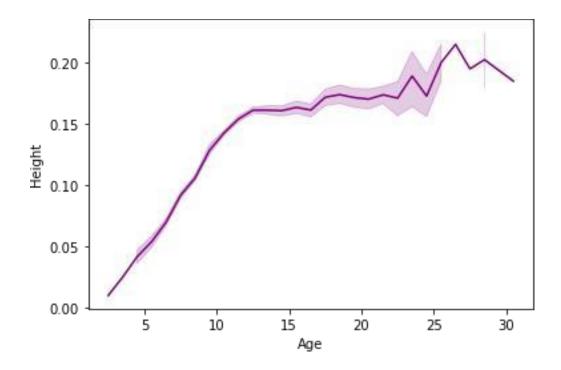
<AxesSubplot:xlabel='Height', ylabel='Age'>



Linearplot

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

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



Scatterplot

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

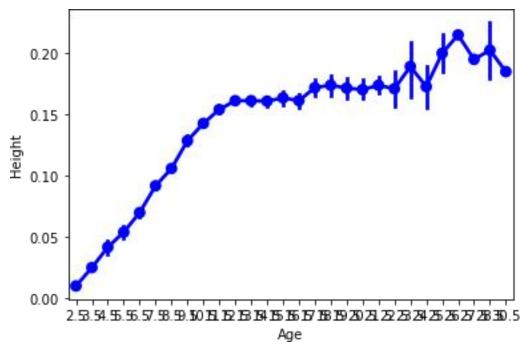
1.0 - 0.8 - 46 0.6 - 0.4 - 0.2 - 0.0 - 5 10 15 20 25 30

In [42]:

Out[42]:

Pointplot

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



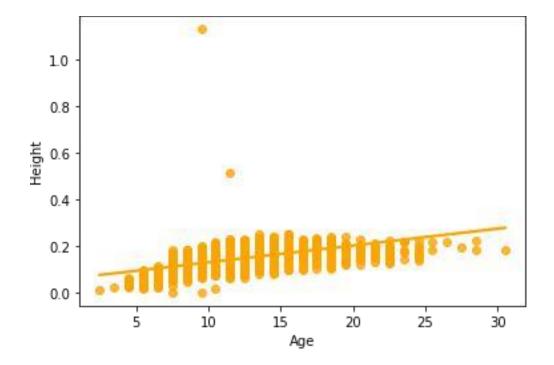
Regplot

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.

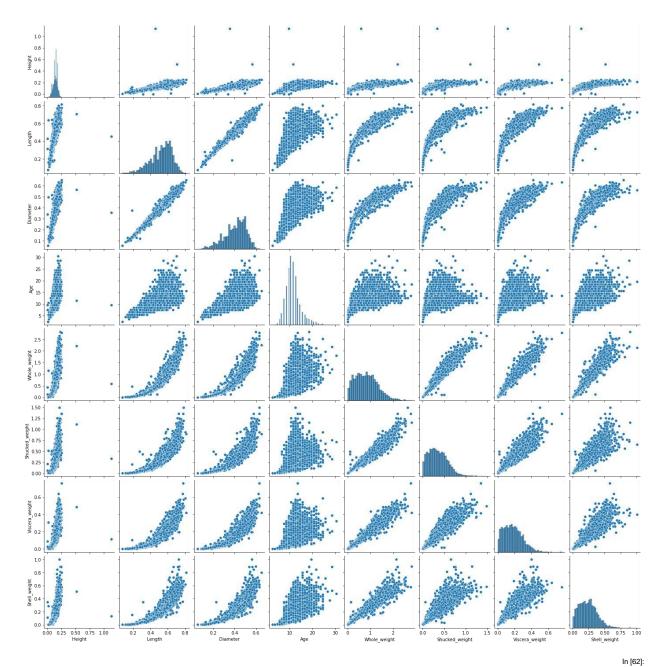
#

Pairplot

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

Out[57]:

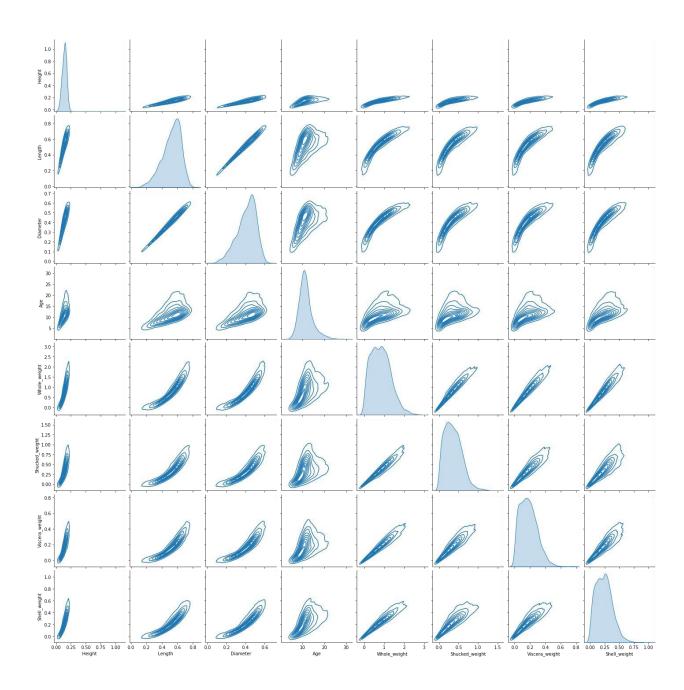
<seaborn.axisgrid.PairGrid at 0x7fd3d93e1040>



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

<seaborn.axisgrid.PairGrid at 0x7fd39840c790>

Out[62]:



4. Perform descriptive statistics on the dataset

| In[63]:
| 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 | 417

| | 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

Out[65]:

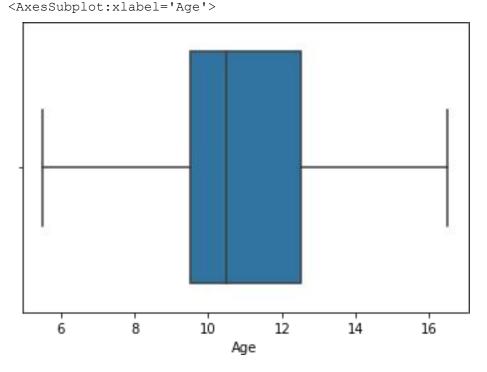
Length Diameter Height Whole_weight Shucked_weight Viscera_weight Age

In [64]:

Out[64]:

0.25 0.450 0.35 0.115 0.4415 0.186 0.0935 0.130 9.5

```
Height
                    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
          Diameter
                                                                                    Shell_weight
 0.455
                        0.095
                                       0.5140
                                                         0.2245
                                                                                          0.150
             0.365
 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
                                                                                          0.155
                                       0.5160
                                                         0.2155
                                                                          0.1140
                        0.125
                                                         0.0895
                                                                                          0.055
 0.330
                                       0.2050
```

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]:

| | 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()

Height Whole weight Shucked weight Viscera weight Length Diameter Shell_weight -0.574558 -0.607685 -0.432149 -1.064424 -0.641898 -0.726212 -0.638217 1.555152 -1.448986 -1.170910 -1.439929 -1.183978 -1.230277 -1.205221 -1.212987 -0.884841 0.050033 0.122130 -0.107991 -0.309469 -0.463500 -0.356690 -0.207139 -0.274842 -0.648238 -0.699476 -0.432149 -0.347099 -0.637819 -0.607600 -0.602294 0.030157 -1.215968 -1.423087 -1.272086 -1.287337 -1.615544 -1.540707 -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]:

| 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_I | 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 |
| ΥТ | 'rain | head(| `\ | | | | | |
| 314 | | 1 | . / | | | | | |
| 352 883 | 21 | 1 2 | | | | | | |
| 362 | | 2 | | | | | | |
| 210 | 16 | 2 | | | | | | |

In [93]:

Out[93]:

668 2 1580 1 3784 2 463 1

Name: Sex, dtype: int64

Y_Test.head()

2

2615

Name: Sex, dtype: int64

11. Build the Model

```
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]:
   0 1 2
 0 122 29 98
 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

| 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 |