TEAM ID: PNT2022TMID48371

PROJECT TITLE: RETAIL STORE STOCK INVENTORY ANALYTICS

ASSIGNMENT DATE: 25.10.22

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1.Download the dataset

```
In [265]:
```

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

2.LOAD THE DATASET

```
In [266]:

df = pd.read_csv('abalone.csv')
```

```
In [267]:
```

```
df.head
```

```
Out[267]:
```

<box< th=""><th>d met</th><th>thod NDFr</th><th>ame.head o</th><th>of</th><th>Sex</th><th>Length</th><th>Diameter</th><th>Height</th><th>Whole weight</th><th>Shucked w</th></box<>	d met	thod NDFr	ame.head o	of	Sex	Length	Diameter	Height	Whole weight	Shucked w
eight	\									
0	M	0.455	0.365	0.095		0.514	0	0.2245		
1	M	0.350	0.265	0.090		0.225	5	0.0995		
2	F	0.530	0.420	0.135		0.677	0	0.2565		
3	M	0.440	0.365	0.125		0.516	0	0.2155		
4	I	0.330	0.255	0.080		0.205	0	0.0895		
4172	F	0.565	0.450	0.165		0.887	0	0.3700		
4173	M	0.590	0.440	0.135		0.966	0	0.4390		
4174	M	0.600	0.475	0.205		1.176	0	0.5255		
4175	F	0.625	0.485	0.150		1.094	5	0.5310		
4176	M	0.710	0.555	0.195		1.948	5	0.9455		

	Viscera	weight	Shell	weight	Rings
0		0.1010		0.1500	15
1		0.0485		0.0700	7
2		0.1415		0.2100	9
3		0.1140		0.1550	10
4		0.0395		0.0550	7
4172		0.2390		0.2490	11
4173		0.2145		0.2605	10
4174		0.2875		0.3080	9
4175		0.2610		0.2960	10
4176		0.3765		0.4950	12

[4177 rows x 9 columns] >

In [268]:

```
Age=1.5+df.Rings
```

```
df["Age"]=Age
df=df.rename(columns = {'whole weight':'whole_weight','Shucked weight':'Shucked_weight',
'Viscera weight':'Viscera_weight','Shell weight':'Shell_weight'})
df=df.drop(columns=["Rings"],axis=1)
df.head()
```

Out[268]:

	Sex	Length	Diameter	Height	Whole weight	Shucked_weight	Viscera_weight	Shell_weight	Age
0	М	0.455	0.365	0.095	0.5140	0.2245	0.1010	0.150	16.5
1	М	0.350	0.265	0.090	0.2255	0.0995	0.0485	0.070	8.5
2	F	0.530	0.420	0.135	0.6770	0.2565	0.1415	0.210	10.5
3	М	0.440	0.365	0.125	0.5160	0.2155	0.1140	0.155	11.5
4	ı	0.330	0.255	0.080	0.2050	0.0895	0.0395	0.055	8.5

In [269]:

df.tail()

Out[269]:

	Sex	Length	Diameter	Height	Whole weight	Shucked_weight	Viscera_weight	Shell_weight	Age
4172	F	0.565	0.450	0.165	0.8870	0.3700	0.2390	0.2490	12.5
4173	М	0.590	0.440	0.135	0.9660	0.4390	0.2145	0.2605	11.5
4174	М	0.600	0.475	0.205	1.1760	0.5255	0.2875	0.3080	10.5
4175	F	0.625	0.485	0.150	1.0945	0.5310	0.2610	0.2960	11.5
4176	М	0.710	0.555	0.195	1.9485	0.9455	0.3765	0.4950	13.5

3. Perform Below Visualizations

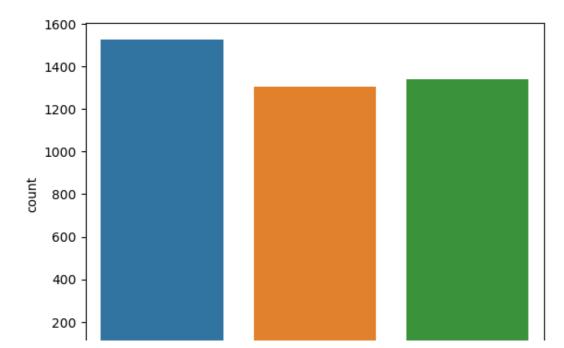
Univariate Analysis

```
In [270]:
```

```
sns.countplot(x='Sex',data=df)
```

Out[270]:

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



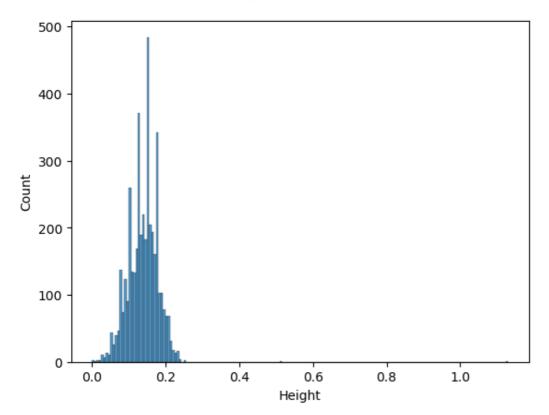


In [271]:

sns.histplot(df["Height"])

Out[271]:

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

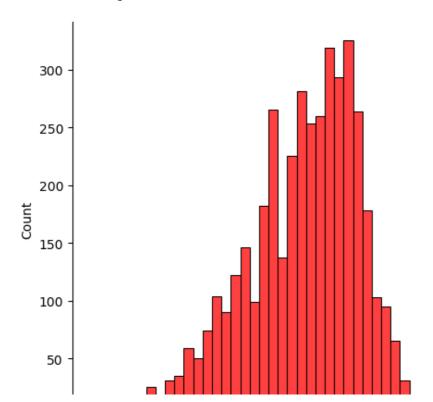


In [272]:

sns.displot(df["Length"],color='red')

Out[272]:

<seaborn.axisgrid.FacetGrid at 0x1af5e2f7820>

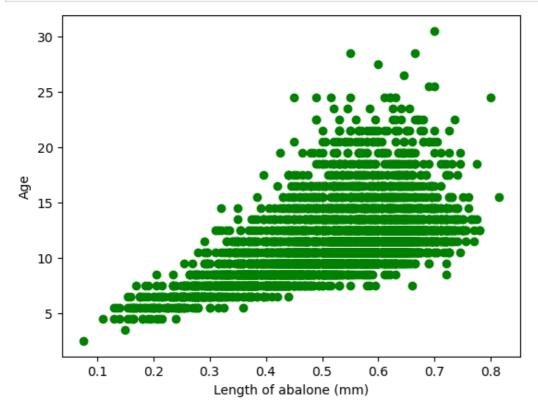


```
0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8
Length
```

Bi-Variate Analysis

```
In [273]:
```

```
plt.scatter(df['Length'], df['Age'], c='green')
plt.xlabel('Length of abalone (mm)')
plt.ylabel('Age')
plt.show()
```



In [274]:

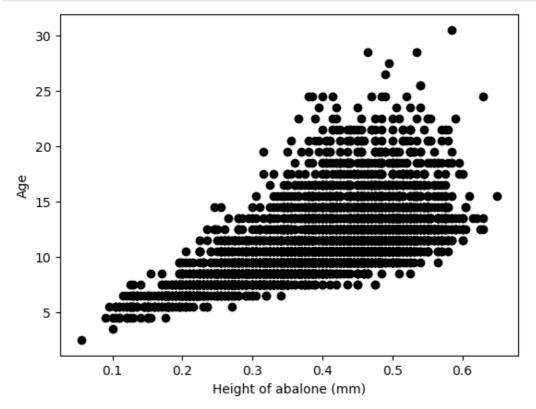
```
plt.scatter(df['Height'], df['Age'], c='purple')
plt.xlabel('Height of abalone (mm)')
plt.ylabel('Age')
plt.show()
```



```
0.0 0.2 0.4 0.6 0.8 1.0
Height of abalone (mm)
```

In [275]:

```
plt.scatter(df['Diameter'], df['Age'], c='black')
plt.xlabel('Height of abalone (mm)')
plt.ylabel('Age')
plt.show()
```



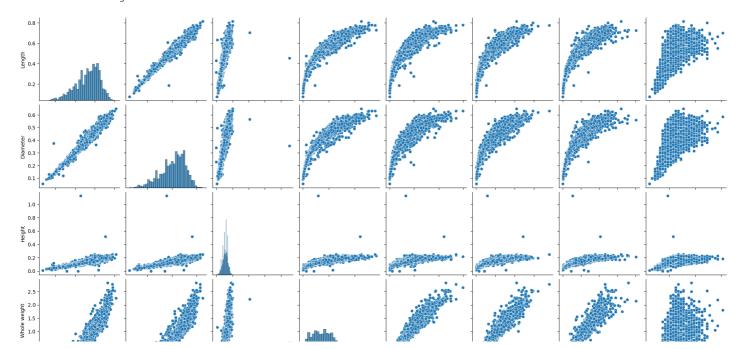
Multi-Variate Analysis

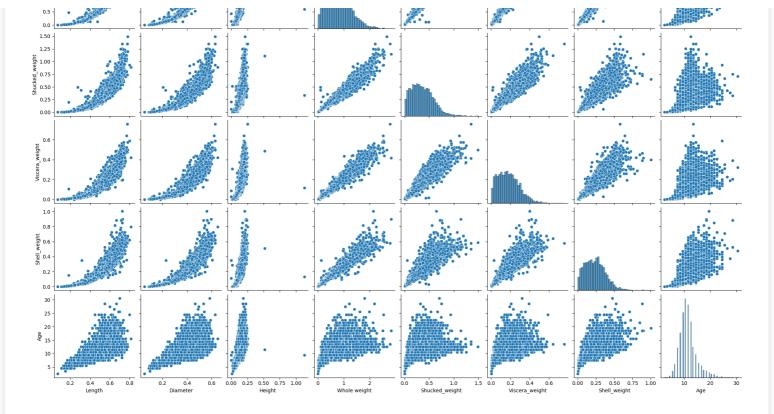
In [276]:

```
numerical_features = df.select_dtypes(include = [np.number]).columns
sns.pairplot(df[numerical_features])
```

Out[276]:

<seaborn.axisgrid.PairGrid at 0x1af61732d00>





In [277]:

plt.figure(figsize=(12,8));
sns.heatmap(df.corr(),cmap='PiYG',annot=True);



4. Perform descriptive statistics on the dataset

```
In [278]:

df.describe()
Out[278]:
```

	Length	Diameter	Height	Whole weight	Shucked_weight	Viscera_weight	Shell_weight	Age
count	4177.000000	4177.000000	4177.000000	4177.000000	4177.000000	4177.000000	4177.000000	4177.000000
mean std min	0.523992	0.407881	0.139516	0.828742	0.359367	0.180594	0.238831	11.433684
	0.120093	0.099240	0.041827	0.490389	0.221963	0.109614	0.139203	3.224169
	0.075000	0.055000	0.000000	0.002000	0.001000	0.000500	0.001500	2.500000
25%	0.450000	0.350000	0.115000	0.441500	0.186000	0.093500	0.130000	9.500000
50%	0.545000	0.425000	0.140000	0.799500	0.336000	0.171000	0.234000	10.500000
75%	0.615000	0.480000	0.165000	1.153000	0.502000	0.253000	0.329000	12.500000
max	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

```
In [279]:
```

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

Out[279]:

COTT

sex	U
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

```
In [280]:
```

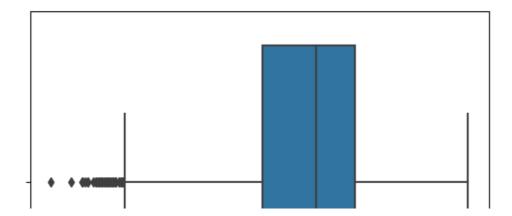
```
sns.boxplot(df['Length'])
```

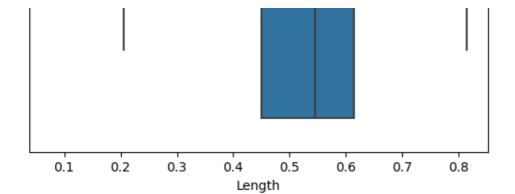
C:\Users\ELCOT\anaconda3\lib\site-packages\seaborn_decorators.py:36: FutureWarning: Pass the following variable as a keyword arg: x. From version 0.12, the only valid positional argument will be `data`, and passing other arguments without an explicit keyword will result in an error or misinterpretation.

warnings.warn(

Out[280]:

<AxesSubplot:xlabel='Length'>





In [281]:

```
q1 = df['Length'].quantile(0.25)
q2 = df['Length'].quantile(0.75)
iqr = q2-q1
q1, q2, iqr
```

Out[281]:

(0.45, 0.615, 0.1649999999999999)

In [282]:

```
upper limit = q2+(1.5*iqr)
lower limit = q1-(1.5*iqr)
lower limit, upper limit
```

Out[282]:

(0.2025000000000004, 0.8624999999999999)

In [283]:

```
new df = df.loc[(df['Length'] <= upper limit) & (df['Length'] >= lower limit)]
print('before removing outliers:', len(df))
print('after removing outliers:', len(new_df))
print('outliers:', len(df)-len(new df))
```

before removing outliers: 4177 after removing outliers: 4128 outliers: 49

In [284]:

```
new df = df.copy()
new_df.loc[(new_df['Length']>upper_limit), 'Length'] = upper_limit
new_df.loc[(new_df['Length']<lower_limit), 'Length'] = lower_limit</pre>
```

In [285]:

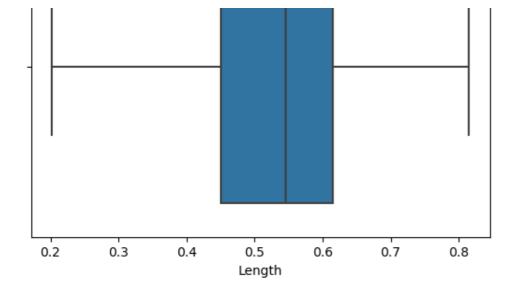
```
sns.boxplot(new df['Length'])
```

C:\Users\ELCOT\anaconda3\lib\site-packages\seaborn_decorators.py:36: FutureWarning: Pass the following variable as a keyword arg: x. From version 0.12, the only valid positional argument will be `data`, and passing other arguments without an explicit keyword will res ult in an error or misinterpretation.

warnings.warn(

Out[285]:

<AxesSubplot:xlabel='Length'>



In [286]:

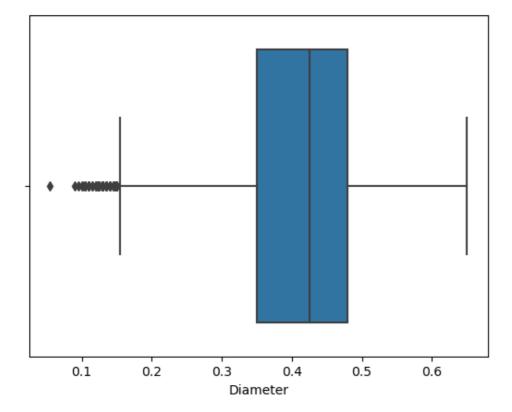
sns.boxplot(df['Diameter'])

C:\Users\ELCOT\anaconda3\lib\site-packages\seaborn_decorators.py:36: FutureWarning: Pass the following variable as a keyword arg: x. From version 0.12, the only valid positional argument will be `data`, and passing other arguments without an explicit keyword will result in an error or misinterpretation.

warnings.warn(

Out[286]:

<AxesSubplot:xlabel='Diameter'>



In [287]:

```
q1 = df['Diameter'].quantile(0.25)
q2 = df['Diameter'].quantile(0.75)
iqr = q2-q1
q1, q2, iqr
```

Out[287]:

(0.35, 0.48, 0.13)

In [288]:

```
upper_limit = q2 + (1.5*iqr)
lower_limit = q1 - (1.5*iqr)
lower_limit, upper_limit
Out[288]:
```

(0.1549999999999997, 0.675)

In [289]:

```
new_df = df.loc[(df['Diameter'] <= upper_limit) & (df['Diameter'] >= lower_limit)]
print('before removing outliers :', len(df))
print('after removing outliers :', len(new_df))
print('outliers :', len(df)-len(new_df))
```

before removing outliers : 4177 after removing outliers : 4118 outliers : 59

In [290]:

```
new_df = df.copy()
new_df.loc[(new_df['Diameter']>upper_limit), 'Diameter'] = upper_limit
new_df.loc[(new_df['Diameter']<lower_limit), 'Diameter'] = lower_limit</pre>
```

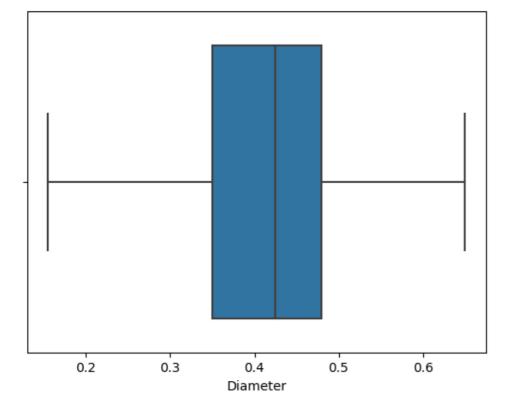
In [291]:

```
sns.boxplot(new_df['Diameter'])
```

C:\Users\ELCOT\anaconda3\lib\site-packages\seaborn_decorators.py:36: FutureWarning: Pass
the following variable as a keyword arg: x. From version 0.12, the only valid positional
argument will be `data`, and passing other arguments without an explicit keyword will res
ult in an error or misinterpretation.
 warnings.warn(

Out[291]:

<AxesSubplot:xlabel='Diameter'>



In [292]:

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

C:\Users\ELCOT\anaconda3\lib\site-packages\seaborn_decorators.py:36: FutureWarning: Pass the following variable as a keyword arg: x. From version 0.12, the only valid positional argument will be `data`, and passing other arguments without an explicit keyword will result in an experience of misinterprotection.

```
utt in an error or misinterbretation.
 warnings.warn(
Out[292]:
<AxesSubplot:xlabel='Height'>
   0.0
            0.2
                     0.4
                              0.6
                                      0.8
                                               1.0
                           Height
In [293]:
q1 = df['Height'].quantile(0.25)
q2 = df['Height'].quantile(0.75)
iqr = q2-q1
q1, q2, iqr
Out[293]:
(0.115, 0.165, 0.05)
In [294]:
upper limit = q2 + (1.5*iqr)
lower limit = q1 - (1.5*iqr)
lower_limit, upper_limit
Out[294]:
In [295]:
new df = df.loc[(df['Height'] <= upper limit) & (df['Height'] >= lower limit)]
print('before removing outliers :', len(df))
print('after removing outliers :', len(new df))
print('outliers :', len(df)-len(new_df))
before removing outliers : 4177
after removing outliers : 4148
outliers : 29
In [296]:
new df = df.copy()
```

new_df.loc[(new_df['Height']>upper_limit), 'Height'] = upper_limit
new_df.loc[(new_df['Height']<lower_limit), 'Height'] = lower_limit</pre>

In [297]:

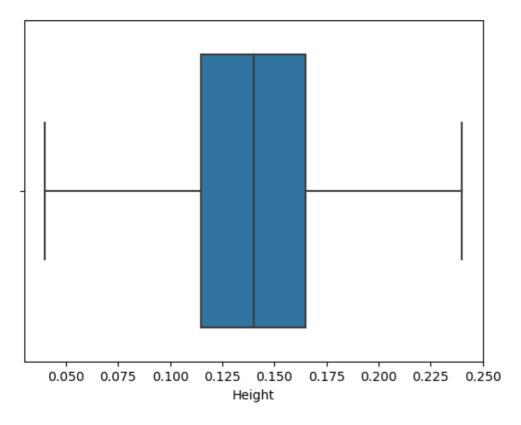
sns.boxplot(new_df['Height'])

C:\Users\ELCOT\anaconda3\lib\site-packages\seaborn_decorators.py:36: FutureWarning: Pass the following variable as a keyword arg: x. From version 0.12, the only valid positional argument will be `data`, and passing other arguments without an explicit keyword will result in an error or misinterpretation.

warnings.warn(

Out[297]:

<AxesSubplot:xlabel='Height'>



In [298]:

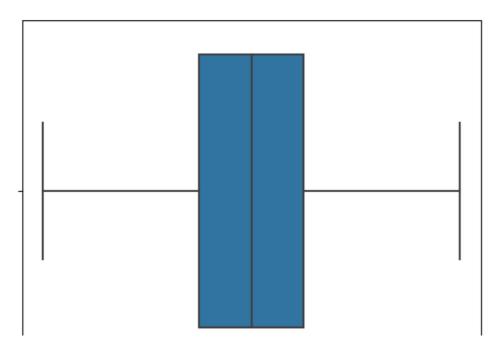
sns.boxplot(new_df['Height'])

C:\Users\ELCOT\anaconda3\lib\site-packages\seaborn_decorators.py:36: FutureWarning: Pass the following variable as a keyword arg: x. From version 0.12, the only valid positional argument will be `data`, and passing other arguments without an explicit keyword will result in an error or misinterpretation.

warnings.warn(

Out[298]:

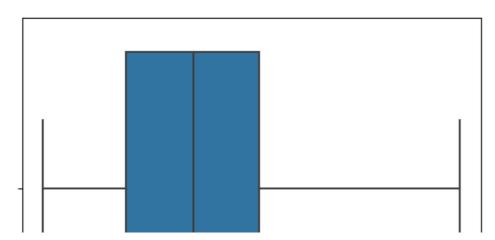
<AxesSubplot:xlabel='Height'>

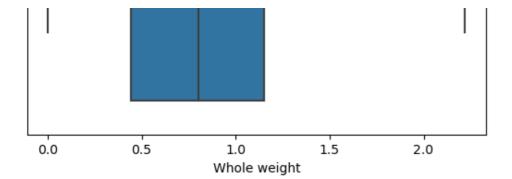


```
In [299]:
q1 = df['Whole weight'].quantile(0.25)
q2 = df['Whole weight'].quantile(0.75)
iqr = q2-q1
q1, q2, iqr
Out[299]:
(0.4415, 1.153, 0.7115)
In [300]:
upper limit = q2 + (1.5*iqr)
lower limit = q1 - (1.5*iqr)
lower limit, upper limit
Out[300]:
(-0.62575, 2.22025)
In [301]:
new df = df.loc[(df['Whole weight'] <= upper limit) & (df['Whole weight'] >= lower limit
print('before removing outliers :', len(df))
print('after removing outliers :', len(new df))
print('outliers :', len(df)-len(new_df))
before removing outliers : 4177
after removing outliers: 4147
outliers : 30
In [302]:
new df = df.copy()
new df.loc[(new df['Whole weight']>upper limit), 'Whole weight'] = upper limit
new df.loc[(new df['Whole weight'] < lower limit), 'Whole weight'] = lower limit</pre>
In [303]:
sns.boxplot(new df['Whole weight'])
C:\Users\ELCOT\anaconda3\lib\site-packages\seaborn\ decorators.py:36: FutureWarning: Pass
the following variable as a keyword arg: x. From version 0.12, the only valid positional
argument will be `data`, and passing other arguments without an explicit keyword will res
ult in an error or misinterpretation.
 warnings.warn(
```

Out[303]:

<AxesSubplot:xlabel='Whole weight'>





In [304]:

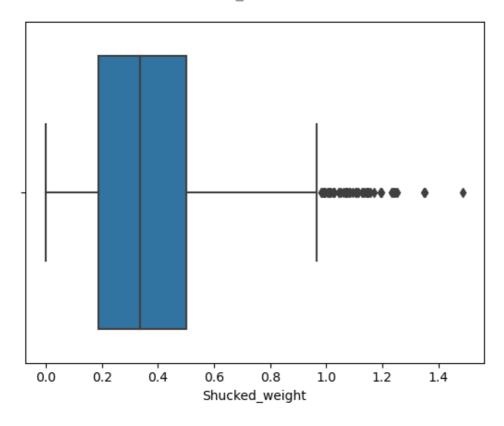
```
sns.boxplot(df['Shucked weight'])
```

C:\Users\ELCOT\anaconda3\lib\site-packages\seaborn_decorators.py:36: FutureWarning: Pass the following variable as a keyword arg: x. From version 0.12, the only valid positional argument will be `data`, and passing other arguments without an explicit keyword will result in an error or misinterpretation.

warnings.warn(

Out[304]:

<AxesSubplot:xlabel='Shucked_weight'>



In [305]:

```
q1 = df['Shucked_weight'].quantile(0.25)
q2 = df['Shucked_weight'].quantile(0.75)
iqr = q2-q1
q1, q2, iqr
```

Out[305]:

(0.186, 0.502, 0.316)

In [306]:

```
upper_limit = q2 + (1.5*iqr)
lower_limit = q1 - (1.5*iqr)
lower_limit, upper_limit
```

Out[306]:

(-0.288, 0.976)

In [307]:

```
new_df = df.loc[(df['Shucked_weight'] <= upper_limit) & (df['Shucked_weight'] >= lower_l
imit)]
print('before removing outliers :', len(df))
print('after removing outliers :', len(new_df))
print('outliers :', len(df)-len(new_df))
```

before removing outliers : 4177 after removing outliers : 4129 outliers : 48

In [308]:

```
new_df = df.copy()
new_df.loc[(new_df['Shucked_weight']>upper_limit), 'Shucked_weight'] = upper_limit
new_df.loc[(new_df['Shucked_weight']<lower_limit), 'Shucked_weight'] = lower_limit</pre>
```

In [309]:

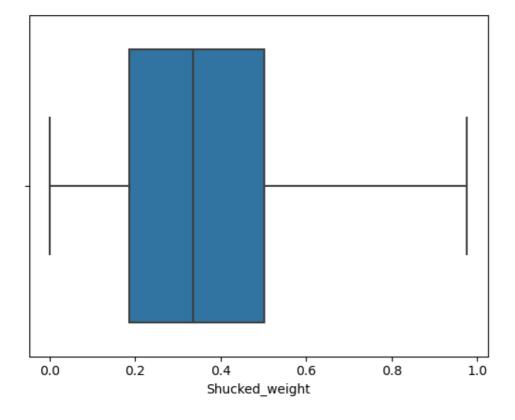
```
sns.boxplot(new_df['Shucked_weight'])
```

C:\Users\ELCOT\anaconda3\lib\site-packages\seaborn_decorators.py:36: FutureWarning: Pass the following variable as a keyword arg: x. From version 0.12, the only valid positional argument will be `data`, and passing other arguments without an explicit keyword will result in an error or misinterpretation.

warnings.warn(

Out[309]:

<AxesSubplot:xlabel='Shucked_weight'>



In [310]:

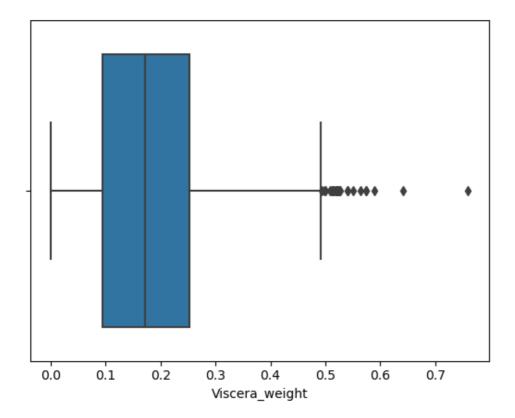
```
sns.boxplot(df['Viscera weight'])
```

C:\Users\ELCOT\anaconda3\lib\site-packages\seaborn_decorators.py:36: FutureWarning: Pass the following variable as a keyword arg: x. From version 0.12, the only valid positional argument will be `data`, and passing other arguments without an explicit keyword will result in an error or misinterpretation.

warnings.warn(

Out[310]:

```
<AxesSubplot:xlabel='Viscera_weight'>
```



7. Check for Categorical columns and perform encoding

```
In [311]:

df['Sex'].replace({'M':1,'F':0,'I':2},inplace=True)
df
```

Out[311]:

	Sex	Length	Diameter	Height	Whole weight	Shucked_weight	Viscera_weight	Shell_weight	Age
0	1	0.455	0.365	0.095	0.5140	0.2245	0.1010	0.1500	16.5
1	1	0.350	0.265	0.090	0.2255	0.0995	0.0485	0.0700	8.5
2	0	0.530	0.420	0.135	0.6770	0.2565	0.1415	0.2100	10.5
3	1	0.440	0.365	0.125	0.5160	0.2155	0.1140	0.1550	11.5
4	2	0.330	0.255	0.080	0.2050	0.0895	0.0395	0.0550	8.5
	•••								
4172	0	0.565	0.450	0.165	0.8870	0.3700	0.2390	0.2490	12.5
4173	1	0.590	0.440	0.135	0.9660	0.4390	0.2145	0.2605	11.5
4174	1	0.600	0.475	0.205	1.1760	0.5255	0.2875	0.3080	10.5
4175	0	0.625	0.485	0.150	1.0945	0.5310	0.2610	0.2960	11.5
4176	1	0.710	0.555	0.195	1.9485	0.9455	0.3765	0.4950	13.5

4177 rows × 9 columns

```
In [312]:
```

from sklearn.preprocessing import LabelEncoder,OneHotEncoder,StandardScaler

In [313]:

```
label_encoder = LabelEncoder()
df['Sex'] = label_encoder.fit_transform(df['Sex'])
df
```

Out[313]:

	Sex	Length	Diameter	Height	Whole weight	Shucked_weight	Viscera_weight	Shell_weight	Age
0	1	0.455	0.365	0.095	0.5140	0.2245	0.1010	0.1500	16.5
1	1	0.350	0.265	0.090	0.2255	0.0995	0.0485	0.0700	8.5
2	0	0.530	0.420	0.135	0.6770	0.2565	0.1415	0.2100	10.5
3	1	0.440	0.365	0.125	0.5160	0.2155	0.1140	0.1550	11.5
4	2	0.330	0.255	0.080	0.2050	0.0895	0.0395	0.0550	8.5
4172	0	0.565	0.450	0.165	0.8870	0.3700	0.2390	0.2490	12.5
4173	1	0.590	0.440	0.135	0.9660	0.4390	0.2145	0.2605	11.5
4174	1	0.600	0.475	0.205	1.1760	0.5255	0.2875	0.3080	10.5
4175	0	0.625	0.485	0.150	1.0945	0.5310	0.2610	0.2960	11.5
4176	1	0.710	0.555	0.195	1.9485	0.9455	0.3765	0.4950	13.5

4177 rows × 9 columns

```
In [314]:
```

```
enc = OneHotEncoder(drop='first')
enc_df = pd.DataFrame(enc.fit_transform(df[['Sex']]).toarray())
df = df.join(enc_df)
df.head()
```

Out[314]:

	Sex	Length	Diameter	Height	Whole weight	Shucked_weight	Viscera_weight	Shell_weight	Age	0	1
0	1	0.455	0.365	0.095	0.5140	0.2245	0.1010	0.150	16.5	1.0	0.0
1	1	0.350	0.265	0.090	0.2255	0.0995	0.0485	0.070	8.5	1.0	0.0
2	0	0.530	0.420	0.135	0.6770	0.2565	0.1415	0.210	10.5	0.0	0.0
3	1	0.440	0.365	0.125	0.5160	0.2155	0.1140	0.155	11.5	1.0	0.0
4	2	0.330	0.255	0.080	0.2050	0.0895	0.0395	0.055	8.5	0.0	1.0

8. Split the data into dependent and independent variables

```
In [315]:
```

```
x.info()
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 4177 entries, 0 to 4176
Data columns (total 11 columns):
 # Column Non-Null Count Dtype
 0 Length
                         4177 non-null float64
 1 Diameter
                         4177 non-null float64
 2 Height
                          4177 non-null float64
 Whole weight 4177 non-null float64
Shucked_weight 4177 non-null float64
Shucked_weight 4177 non-null float64
Viscera_weight 4177 non-null float64
Shell_weight 4177 non-null float64
Age 4177 non-null float64
                          4177 non-null uint8
      Sex F
      Sex_ I
                           4177 non-null uint8
 10 Sex_M
                          4177 non-null uint8
```

dtypes: float64(8), uint8(3)

memory usage: 273.4 KB

```
In [316]:
X = x.drop(['Age'], axis = 1)
In [317]:
X.head(2)
Out[317]:
   Length Diameter Height Whole weight Shucked_weight Viscera_weight Shell_weight Sex_F Sex_I Sex_M
   0.455
             0.365
                    0.095
                               0.5140
                                              0.2245
                                                            0.1010
                                                                          0.15
                                                                                  0
                                                                                        0
                                                                                               1
    0.350
             0.265
                    0.090
                               0.2255
                                              0.0995
                                                            0.0485
                                                                          0.07
                                                                                               1
                                                                                  0
                                                                                        0
In [318]:
y = x['Age']
In [319]:
y.head(2)
Out[319]:
0
    16.5
1
      8.5
Name: Age, dtype: float64
```

9. Scale the independent variables

```
In [320]:
scale = StandardScaler()
scaledX = scale.fit transform(x)
print(scaledX)
[-0.57455813 -0.43214879 -1.06442415 ... -0.67483383 -0.68801788]
  1.316677161
[-1.44898585 -1.439929
                      -1.18397831 ... -0.67483383 -0.68801788
  1.31667716]
[ 0.05003309  0.12213032  -0.10799087  ...  1.48184628  -0.68801788
 -0.75948762]
              0.67640943 1.56576738 ... -0.67483383 -0.68801788
[ 0.6329849
  1.31667716]
[0.84118198 \quad 0.77718745 \quad 0.25067161 \dots \quad 1.48184628 \quad -0.68801788
 -0.75948762]
1.31667716]]
```

10. Split the data into training and testing

```
In [321]:
X.shape, y.shape
Out[321]:
((4177, 10), (4177,))
In [322]:
from sklearn.model_selection import train_test_split
x_train, x_test, y_train, y_test = train_test_split(X,y, test_size=0.2, random_state=42)
```

```
In [323]:
print(' x tain.shape : ', x train.shape)
print(' y_tain.shape : ',y_train.shape)
print(' x test.shape : ', x test.shape)
print(' y test.shape : ', y test.shape)
 x tain.shape : (3341, 10)
 y_tain.shape : (3341,)
 x test.shape: (836, 10)
 y test.shape : (836,)
10. Build the Model, 11. Train the Model, 12. Test the Model
In [324]:
from sklearn.linear model import LinearRegression
lr = LinearRegression()
lr.fit(x train, y train)
lr_pred = lr.predict(x_test)
In [325]:
from sklearn.ensemble import RandomForestRegressor
from sklearn.metrics import mean squared error, make scorer
from sklearn.model selection import RandomizedSearchCV
rf = RandomForestRegressor()
param = {
    'max depth': [3,6,9,12,15],
    'n estimators':[10,50,100,150,200]
rf search = RandomizedSearchCV(rf, param distributions=param, n iter=5, scoring=make scorer
(mean squared error), n jobs=-1, cv=5, verbose=3)
rf search.fit(x train, y train)
Fitting 5 folds for each of 5 candidates, totalling 25 fits
Out[325]:
RandomizedSearchCV(cv=5, estimator=RandomForestRegressor(), n iter=5, n jobs=-1,
                   param_distributions={'max_depth': [3, 6, 9, 12, 15],
                                        'n_estimators': [10, 50, 100, 150,
                                                         200]},
                   scoring=make scorer(mean squared error), verbose=3)
In [326]:
means = rf search.cv results ['mean test score']
params = rf search.cv results ['params']
for mean, param in zip(means, params):
    print("%f with: %r" % (mean, param))
```

print('Best parameters with the minimum Mean Square Error are:' ,param)

Best parameters with the minimum Mean Square Error are: {'n estimators': 100, 'max depth'

if mean == min(means):

: 9}

In [327]:

rf.fit(x train, y train)

rf pred = rf.predict(x test)

4.664623 with: {'n_estimators': 200, 'max_depth': 6} 4.618707 with: {'n_estimators': 100, 'max_depth': 15} 4.644619 with: {'n_estimators': 200, 'max_depth': 15} 5.677870 with: {'n_estimators': 150, 'max_depth': 3} 4.581780 with: {'n_estimators': 100, 'max_depth': 9}

rf = RandomForestRegressor(n estimators=50, max depth=6)

14. Measure the performance using Metrics

```
In [328]:
```

```
from sklearn import metrics
print('Linear Regression :')
print('-----')
print('MAE:', metrics.mean_absolute_error(y_test, lr_pred))
print('MSE:', metrics.mean_squared_error(y_test, lr_pred))
print('RMSE:', np.sqrt(metrics.mean_squared_error(y_test, lr_pred)))
print('R2 Score:', metrics.r2_score(y_test, lr_pred))
print('\n\n')
```

MAE: 1.5944508821770336 MSE: 4.892375672262822 RMSE: 2.211871531591024 R2 Score: 0.5480572061259404

Linear Regression:

In [329]:

```
from sklearn import metrics
print('Random Forest Contains:')
print('-----')
print('MAE:', metrics.mean_absolute_error(y_test, rf_pred))
print('MSE:', metrics.mean_squared_error(y_test, rf_pred))
print('RMSE:', np.sqrt(metrics.mean_squared_error(y_test, rf_pred)))
print('R2 Score:', metrics.r2_score(y_test, rf_pred))
```

Random Forest Contains:
----MAE: 1.5580369509719958
MSE: 5.025592967383406
RMSE: 2.241783434541215
R2 Score: 0.535750997326301