SLR MID EXAM

Data_set:

This dataset is a record of 7 common different fish species in fish market sales. With this dataset, a predictive model can be performed using machine friendly data and estimate the weight of fish can be predicted.

- 1. Species: Species name of fish
- 2. Weight: Weight of fish in gram
- 3. Length1: Vertical length in cm
- 4. Length2: Diagonal length in cm
- 5. Length3: Cross length in cm
- 6. Height: Height in cm
- 7. Width: Diagonal width in cm

Our dependent variable is 'Weight'. Independent variables are 'species', different lengths, 'height' and 'width'.

In [1]:

```
# Kindly change the below cells from markdown to code and execute it
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
import pandas as pd
%matplotlib inline
```

In [2]:

```
import pandas as pd
import csv
with open("data_set.csv","r")as file:
    reader=csv.reader(file)
df=pd.read_csv("data_set.csv")
df.head()
```

Out[2]:

	Species	Weight	Length1	Length2	Length3	Height	Width
0	Bream	242.0	23.2	25.4	30.0	11.5200	4.0200
1	Bream	290.0	24.0	26.3	31.2	12.4800	4.3056
2	Bream	340.0	23.9	26.5	31.1	12.3778	4.6961
3	Bream	363.0	26.3	29.0	33.5	12.7300	4.4555
4	Bream	430.0	26.5	29.0	34.0	12.4440	5.1340

1. Data Understanding (5 marks)

- a. Read the dataset (tab, csv, xls, txt, inbuilt dataset). What are the number of rows and no. of cols & types of variables (continuous, categorical etc.)? (1 MARK)
- b. Calculate five-point summary for numerical variables (1 MARK)
- c. Summarize observations for categorical variables no. of categories, % observations in each category. (1 mark)

d. Check for defects in the data such as missing values, null, outliers, etc. (2 marks)

```
In [3]:
```

```
# a. Read the dataset (tab, csv, xls, txt, inbuilt dataset).
# What are the number of rows and no. of cols & types of variables (continuous, categoric
al etc.)? (1 MARK)
print('number of rows:',df.shape[0])
print(' ')
print('number of columns:',df.shape[1])
print(' ')
print('Type of Variables:')
print(' ')
print(df.info())
number of rows: 159
number of columns: 7
Type of Variables:
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 159 entries, 0 to 158
Data columns (total 7 columns):
        159 non-null object
Species
           159 non-null float64
Weight
Length1
          159 non-null float64
```

None

In [4]:

Length2

Length3

Height

Width

```
# b. Calculate five-point summary for numerical variables (1 MARK)

df.describe()
```

Out[4]:

		Weight	Length1	Length2	Length3	Height	Width
C	ount	159.000000	159.000000	159.000000	159.000000	159.000000	159.000000
mean	nean	398.326415	26.247170	28.415723	31.227044	8.970994	4.417486
	std	357.978317	9.996441	10.716328	11.610246	4.286208	1.685804
min 25%	min	0.000000	7.500000	8.400000	8.800000	1.728400	1.047600
	25%	120.000000	19.050000	21.000000	23.150000	5.944800	3.385650
	50%	273.000000	25.200000	27.300000	29.400000	7.786000	4.248500
7	75%	650.000000	32.700000	35.500000	39.650000	12.365900	5.584500
ı	max	1650.000000	59.000000	63.400000	68.000000	18.957000	8.142000

159 non-null float64

159 non-null float64 159 non-null float64

159 non-null float64

dtypes: float64(6), object(1)

memory usage: 8.8+ KB

In [5]:

```
# c. Summarize observations for categorical variables - no. of categories, % observations
in each category. (1 mark)

df_cat=df['Species'].value_counts()
df_cat=pd.DataFrame(df_cat)
df_cat['%']=df_cat['Species'].apply(lambda x:round((x/159)*100,2))
df_cat
```

Out[5]:

	Species	%
Perch	56	35.22
Bream	35	22.01
Roach	20	12.58
Pike	17	10.69
Smelt	14	8.81
Parkki	11	6.92
Whitefish	6	3.77

In [6]:

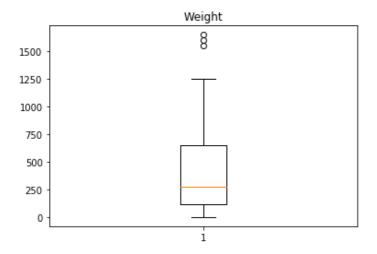
```
# d. Check for defects in the data such as missing values, null, outliers, etc. (2 marks)
print('Missing Values in the data :')
print(df.isnull().sum())
Missing Values in the data:
Species
           0
Weight
           0
Length1
           0
Length2
Length3
Height
Width
           0
dtype: int64
```

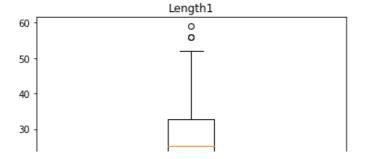
• There are no missing values in the data

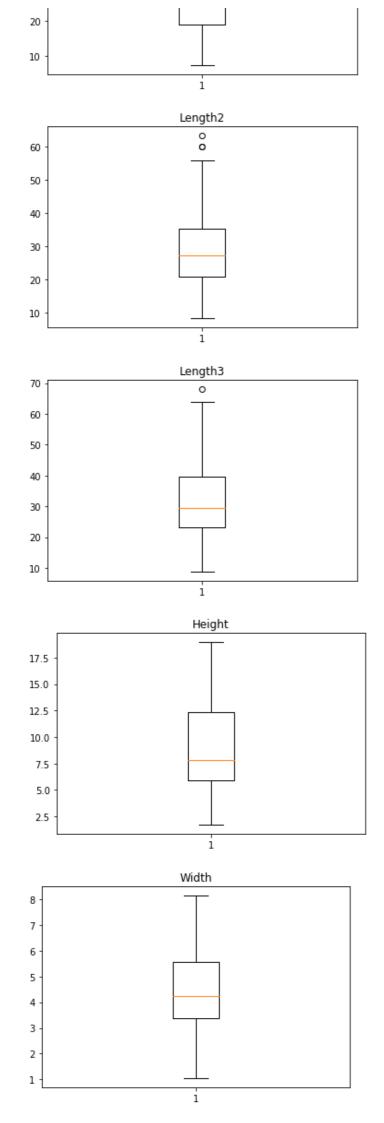
In [7]:

```
df_num=df.select_dtypes(np.number)
print('Outliers in the data :')
for features in df_num.columns:
    plt.boxplot(df_num[features])
    plt.title(features)
    plt.show()
```

Outliers in the data :







 We could see Outliers in weight,length1,length2 and length3 columns in the data with very less number of outliers

2. Data Preparation (10 marks)

- a. Fix the defects found above and do appropriate treatment if any. (3 marks)
- b. Visualize the data using relevant plots. Find out the variables which are highly correlated with target variable? (3 marks)
- c. Do you want to exclude some variables from the model based on this analysis? What other actions will you take? (2 marks)
- d. Split dataset into train and test (70:30). Are both train and test representative of the overall data? How would you ascertain this statistically? (2 marks)

```
In [8]:
# a. Fix the defects found above and do appropriate treatment if any. (3 marks)
# Since we could not see any null value/missing values in the data we dont need to handle
it.
# We just have to remove the outliers in the data using IQR method
```

```
q1=df['Weight'].quantile(0.25)
q3=df['Weight'].quantile(0.75)
iqr=q3-q1
df= df[(df['Weight']>q1-(1.5*iqr)) & (df['Weight']<q3+(1.5*iqr))]</pre>
```

```
In [9]:
```

```
q1=df['Length1'].quantile(0.25)
q3=df['Length1'].quantile(0.75)
iqr=q3-q1
df= df[(df['Length1']>q1-(1.5*iqr)) & (df['Length1']<q3+(1.5*iqr))]</pre>
```

```
In [10]:
```

```
q1=df['Length2'].quantile(0.25)
q3=df['Length2'].quantile(0.75)
iqr=q3-q1
df= df[(df['Length2']>q1-(1.5*iqr)) & (df['Length2']<q3+(1.5*iqr))]</pre>
```

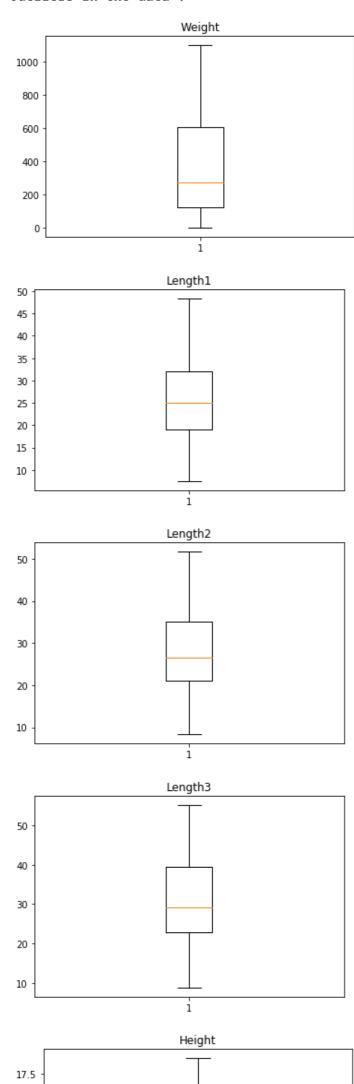
```
In [11]:
```

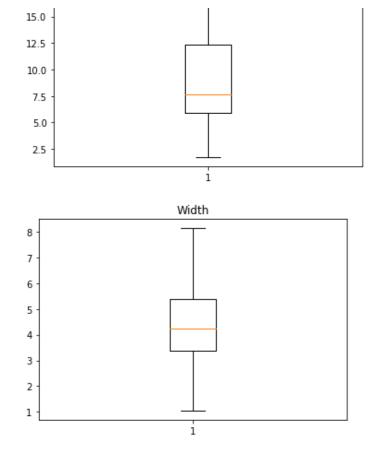
```
q1=df['Length3'].quantile(0.25)
q3=df['Length3'].quantile(0.75)
iqr=q3-q1
df= df[(df['Length3']>q1-(1.5*iqr)) & (df['Length3']<q3+(1.5*iqr))]</pre>
```

```
In [12]:
```

```
df.shape
Out[12]:
  (155, 7)
In [13]:
```

```
df_num=df.select_dtypes(np.number)
print('Outliers in the data :')
for features in df_num.columns:
    plt.boxplot(df_num[features])
    plt.title(features)
    plt.show()
```



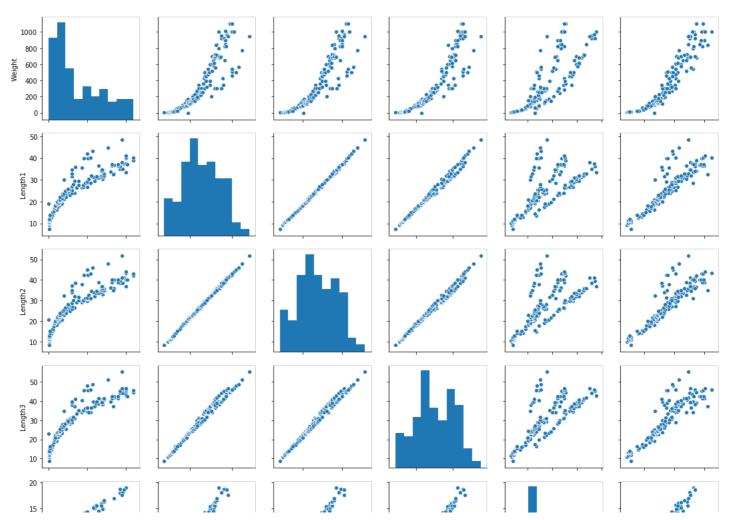


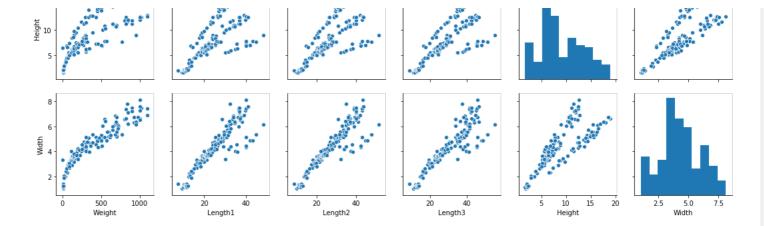
In [14]:

b. Visualize the data using relevant plots.
Find out the variables which are highly correlated with target variable? (3 marks)
sns.pairplot(df)

Out[14]:

<seaborn.axisgrid.PairGrid at 0x7f4de3e44978>





In [15]:

df.corr()

Out[15]:

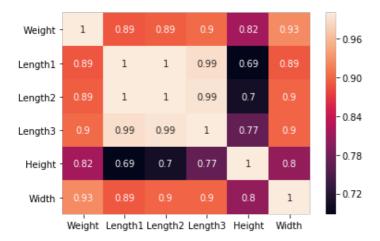
	Weight	Length1	Length2	Length3	Height	Width
Weight	1.000000	0.890173	0.894415	0.902576	0.815237	0.925664
Length1	0.890173	1.000000	0.999385	0.990205	0.688050	0.891260
Length2	0.894415	0.999385	1.000000	0.992765	0.704041	0.897855
Length3	0.902576	0.990205	0.992765	1.000000	0.768038	0.898315
Height	0.815237	0.688050	0.704041	0.768038	1.000000	0.803268
Width	0.925664	0.891260	0.897855	0.898315	0.803268	1.000000

In [17]:

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

Out[17]:

<matplotlib.axes. subplots.AxesSubplot at 0x7f4de05f8c50>



- From the above analysis, the below variables are highly correlated withe target variable
- Length1
- Length2
- Length3
- Height
- Width

In [21]:

sns.boxplot(df['Species'],df['Weight'])

Out[21]:

<matplotlib.axes. subplots.AxesSubplot at 0x7f4de0517358>

1000 - Bream Roach Whitefish Parkki Species Perch Pike Smelt

 From the above boxplot on categorical column we could clearly see the variation in the mean weight of different categories, which shows that the Species columns contributes to the weight of the fish.

In [22]:

```
# c. Do you want to exclude some variables from the model based on this analysis?
# What other actions will you take? (2 marks)
```

- From the above plots and correlation matrix it is clearly visible that length 1,length 2, length 3 are highly correlated and could tend to have **Multicollinearity** with a correlation of close to 1 and adding redundancy to the data.
- Hence removing the redundant columns of length1 and length2 will be appropriate to reduce the redundancy

Checking Multicoolinearity

```
In [33]:
```

```
df_samp=df_num.drop(columns=['Weight'])
```

In [34]:

```
from statsmodels.stats.outliers_influence import variance_inflation_factor
vif=pd.DataFrame()
vif['features']=df_samp.columns
vif['vif']=[variance_inflation_factor(df_samp.values,i) for i in range(df_samp.shape[1]
)]
vif
```

Out[34]:

	features	vif
0	Length1	11640.943653
1	Length2	15183.972890
2	Length3	3093.805839
3	Height	75.546906
4	Width	96.285071

• From VIF factor it is clearly visible that length 1,length2,length3 are having high multicollinearity with each other.

In [37]:

d. Split dataset into train and test (70:30). Are both train and test representative of

```
# How would you ascertain this statistically? (2 marks)
df dummy=pd.get dummies(data=df,columns=['Species'],drop first=True)
In [42]:
x=df dummy.drop(columns='Weight')
y=df dummy['Weight']
from sklearn.model selection import train test split
x train,x test,y train,y test=train test split(x,y,test size=0.3,random state=0)
In [43]:
x train.mean()
Out[43]:
                    25.294444
Length1
Length2
                    27.389815
Length3
                    30.183333
Height
                    8.805746
Width
                    4.288494
Species Parkki
                    0.083333
Species Perch
                    0.342593
Species Pike
                    0.101852
Species_Roach
Species_Smelt
                    0.129630
                    0.092593
Species Whitefish
                    0.027778
dtype: float64
In [45]:
x test.mean()
Out[45]:
                    25.925532
Length1
Length2
                    28.097872
                    30.842553
Length3
                     9.248279
Height
                     4.520638
Width
Species_Parkki
                    0.042553
Species_Perch
                    0.404255
Species Pike
                     0.042553
Species_Roach
                     0.127660
Species_Smelt
                     0.085106
Species_Whitefish
                     0.063830
```

- From the above mean analysis it is clear that the mean of all the training variables is almost close to the mean of all the test variables.
- Hence we can conclude that the splitted data is the correct representative of the original data.

3. Model Building (15 marks)

dtype: float64

the overall data?

- a. Fit a base model and observe the overall R- Squared, RMSE and MAPE values of the model. Please comment on whether it is good or not. (3 marks)
- b. Check for multi-collinearity and treat the same. (2 marks)
- c. How would you improve the model? Write clearly the changes that you will make before refitting the model. Fit the final model. (6 marks)
- d. Write down a business interpretation/explanation of the model which variables are affecting the target the most and explain the relationship. Feel free to use charts or graphs to explain. (2 marks)
- e. What changes from the base model had the most effect on model performance? (2 marks)

```
In [46]:
# a. Fit a base model and observe the overall R- Squared, RMSE and MAPE values of the mod
el.
# Please comment on whether it is good or not. (3 marks)
from sklearn.linear model import LinearRegression
In [47]:
base model=LinearRegression()
In [48]:
base model.fit(x train, y train)
Out[48]:
LinearRegression()
In [49]:
y pred=base model.predict(x test)
In [50]:
from sklearn.metrics import r2_score, mean_squared_error
print('R2 score of the base model is',r2 score(y test,y pred))
R2 score of the base model is 0.9290256329220582
In [51]:
mse=mean squared error(y test, y pred)
rmse=np.sqrt(mse)
print('RMSE score of the base model is',rmse)
RMSE score of the base model is 81.25878675299707
In [100]:
y pred train=base model.predict(x train)
In [102]:
mse=mean_squared_error(y_train,y_pred_train)
rmse=np.sqrt(mse)
print('RMSE score of the base model is', rmse)
RMSE score of the base model is 71.37047100437793
In [104]:
mape=np.mean(np.abs((y test-y pred)/y test))*100
print('MAPE score of the base model is', mape)
MAPE score of the base model is inf
 • From the above analysis the r2 score is close to 92% which is good enough

    the rmse score is 81.25 which is the variance error and 71.37 is the bias error which not shows much of

   overfitting in the model
In [54]:
# b. Check for multi-collinearity and treat the same. (2 marks)
```

In [55]:

df samp=df num.drop(columns=['Weight'])

```
In [56]:
from statsmodels.stats.outliers_influence import variance_inflation_factor
vif=pd.DataFrame()
vif['features'] = df samp.columns
vif['vif']=[variance_inflation_factor(df_samp.values,i) for i in range(df_samp.shape[1]
) ]
vif
Out[56]:
  features
                 vif
0 Length1 11640.943653
1 Length2 15183.972890
2 Length3
          3093.805839
   Height
            75.546906
    Width
            96.285071
In [59]:
df samp=df num.drop(columns=['Weight','Length1','Length2'])
In [60]:
from statsmodels.stats.outliers influence import variance inflation factor
vif=pd.DataFrame()
vif['features'] = df samp.columns
vif['vif']=[variance inflation factor(df samp.values,i) for i in range(df samp.shape[1]
vif
Out[60]:
  features
               vif
0 Length3 41.909181
   Height 15.220064
    Width 48.769166
In [61]:
df_samp=df_num.drop(columns=['Weight','Length1','Length2','Length3'])
In [62]:
from statsmodels.stats.outliers influence import variance inflation factor
vif=pd.DataFrame()
vif['features'] = df samp.columns
vif['vif']=[variance inflation factor(df samp.values,i) for i in range(df samp.shape[1]
) ]
vif
```

Out[62]:

 features
 vif

 0
 Height
 14.869484

 1
 Width
 14.869484

 Removed all the multicollinearity columns and still multicollinearity can be seen hence we can go with Lasso to reduce the effect of Multicollinearity.

c. How would you improve the model?

Write clearly the changes that you will make before refitting the model. Fit the final model. (6 marks)

 We can use the feature selection technique to find the significant features in the model and refit the model to improve the performance of the model

```
In [72]:
from sklearn.feature selection import RFE
In [80]:
lr=LinearRegression()
rfe feat=RFE(estimator=lr,n features to select=5)
In [81]:
rfe sel feat=rfe feat.fit(x,y)
In [84]:
sel feat=rfe sel feat.get support()
In [87]:
sel feat=list(x.columns[sel feat])
In [88]:
print('the selected features are : ',sel feat)
the selected features are : ['Width', 'Species Perch', 'Species Roach', 'Species Smelt',
'Species Whitefish']

    the feature selection technique removed the insignificant features and the multicollinear features from the

   model and now the model can be built with the selected features
In [105]:
final model=LinearRegression()
In [106]:
final model.fit(x train, y train)
Out[106]:
LinearRegression()
In [107]:
y pred final=final model.predict(x test)
In [108]:
print('the R2 score of the final model is:',r2 score(y test,y pred final))
the R2 score of the final model is: 0.9290256329220582
In [109]:
mse=mean_squared_error(y_test,y_pred_final)
rmse=np.sqrt(mse)
print('RMSE score of the base model is',rmse)
RMSE score of the base model is 81.25878675299707
```

In [95]: # d. Write down a business interpretation/explanation of the model -

which variables are affecting the target the most and explain the relationship.
Feel free to use charts or graphs to explain. (2 marks)

import statsmodels.api as sm
xc=sm.add_constant(x)
model=sm.OLS(y,x).fit()

model.summary()

/home/deploy/anaconda3/lib/python3.7/site-packages/numpy/core/fromnumeric.py:2389: Future Warning: Method .ptp is deprecated and will be removed in a future version. Use numpy.ptp instead.

return ptp(axis=axis, out=out, **kwargs)

Out[95]:

OLS Regression Results

OLO Hegression He	Juito					
Dep. Variable:	W	eight	R-squa	red (und	centered):	0.970
Model:		OLS Ad	j. R-squa	red (un	centered):	0.968
Method:	Least Squ	ares		F	-statistic:	427.4
Date:	Fri, 18 Sep	2020	ı	Prob (F	-statistic):	5.13e-104
Time:	10:1	13:23		Log-L	ikelihood:	-905.49
No. Observations:		155			AIC:	1833.
Df Residuals:		144			BIC:	1866.
Df Model:		11				
Covariance Type:	nonro	bust				
	coef	std err	t	P>iti	[0.025	0.975]
Length1	62.4023	34.681	1.799	0.074	-6.147	130.952
Length2	103.2937	41.019	2.518	0.013	22.216	184.372
Length3	-147.6639	18.715	-7.890	0.000	-184.655	-110.673
Height	22.5274	12.551	1.795	0.075	-2.280	47.335
Width	117.7837	20.385	5.778	0.000	77.491	158.077
Species_Parkki	-331.2196	29.444	-11.249	0.000	-389.417	-273.022
Species_Perch	-474.3285	41.545	-11.417	0.000	-556.446	-392.211
Species_Pike	-357.5681	102.967	-3.473	0.001	-561.090	-154.046
Species_Roach	-331.5962	37.179	-8.919	0.000	-405.083	-258.110
Species_Smelt	-205.4095	36.447	-5.636	0.000	-277.449	-133.370
Species_Whitefish	-304.1895	51.376	-5.921	0.000	-405.738	-202.641
Omnibus: 9	.252 D urt	oin-Watso	n: 0.8	27		
Prob(Omnibus): 0	.010 Jarque	e-Bera (JE	3): 10.5	99		
Skew: 0	.432	Prob(JE	3): 0.004	99		
Kurtosis: 3	.946	Cond. N	lo. 90	05.		

Warnings:

[1] Standard Errors assume that the covariance matrix of the errors is correctly specified.

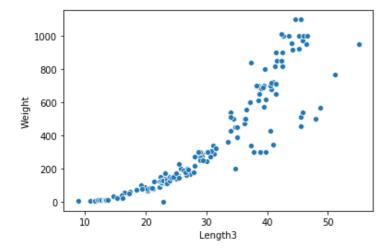
- the weight of the fish increases by 62.4 grams from the average weight for a unit change of 1 cm in length.
- the width can have a high high correlation as weight increases by 117.7 grams for a unit change of 1 cm in length

In [97]:

```
sns.scatterplot(x['Length3'],y)
```

Out[97]:

<matplotlib.axes._subplots.AxesSubplot at 0x7f4d7dee89e8>

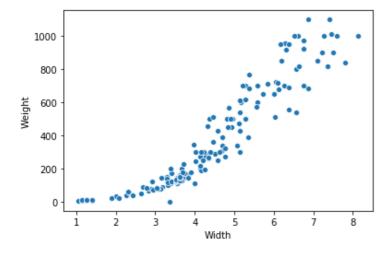


In [98]:

```
sns.scatterplot(x['Width'],y)
```

Out[98]:

<matplotlib.axes._subplots.AxesSubplot at 0x7f4d7deadb38>

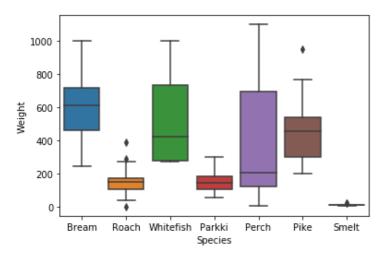


In [99]:

```
sns.boxplot(df['Species'],df['Weight'])
```

Out[99]:

 ${\tt <matplotlib.axes._subplots.AxesSubplot}$ at ${\tt 0x7f4d7de747b8}{\tt >}$



• From the above boxplot on categorical column we could clearly see the variation in the mean weight of different categories, which shows that the **Species** columns contributes to the weight of the fish.

e. What changes from the base model had the most effect on model performance? (2 marks)

- choosing the significant predictors from the model and significant features from the model is increasing the performance of the model.
- removing the multicollinearity columns from the model will give a generalized and reliable model to predict the weigth of the fish.

In []: