50. Regression algorithms o Decision Tree Regressor o Random Forest Regressor o Support Vector Regression

```
In [11]: import pandas as p
          from sklearn.metrics import r2_score
          from sklearn.model selection import train test split
          # Breast cancer dataset
          data = p.read_csv("breast-cancer.csv")
          data.info()
          data.drop(["id"],axis=1,inplace=True)
          M=data[data.diagnosis=="M"]
          B=data[data.diagnosis=="B"]
          data.diagnosis=[1 if i == "M" else 0 for i in data.diagnosis]
          x=data.drop(["diagnosis"],axis=1)
          y=data.diagnosis.values
          x_train,x_test,y_train,y_test=train_test_split(x,y,test_size=0.3,random_state=42)
          from sklearn.tree import DecisionTreeRegressor
          model=DecisionTreeRegressor()
          model.fit(x_train,y_train)
          y pred=model.predict(x test)
          print("\nAccuracy of the model using Decision tree regression algorithm is ",r2 score(y test,y pred))
          \textbf{from} \  \, \textbf{sklearn.ensemble} \  \, \textbf{import} \  \, \textbf{RandomForestRegressor}
          model1 = RandomForestRegressor()
          model1.fit(x_train,y_train)
          y_pred1 = model1.predict(x_test)
          print("\nAccuracy of the model using Random forest regression algorithm is ",r2_score(y_test,y_pred1))
          from sklearn.svm import SVR
          model2 = SVR(kernel='rbf')
          model2.fit(x_train,y_train)
          y pred2 = model2.predict(x test)
          print("\nAccuracy of the model using Support vector regression algorithm is ",r2_score(y_test,y_pred2))
```

```
RangeIndex: 569 entries, 0 to 568
Data columns (total 32 columns):
 # Column
                                                                    Non-Null Count Dtype
 0 id
                                                                  569 non-null int64
                                                                 569 non-null object
569 non-null float64
569 non-null float64
        diagnosis
radius_mean
 1
 3 texture_mean
4 perimeter_mean 569 non-null float64
5 area_mean 569 non-null float64
6 smoothness_mean 569 non-null float64
7 compactness_mean 569 non-null float64
8 concavity_mean 569 non-null float64
9 concave points_mean 569 non-null float64
10 symmetry_mean 569 non-null float64
11 fractal_dimension_mean 569 non-null float64
 11 fractal dimension mean 569 non-null float64
                                         569 non-null float64
 12 radius se
13 texture_se 569 non-null float64
14 perimeter_se 569 non-null float64
15 area_se 569 non-null float64
16 smoothness_se 569 non-null float64
17 compactness_se 569 non-null float64
18 concavity_se 569 non-null float64
19 concave points_se 569 non-null float64
20 symmetry_se 569 non-null float64
21 fractal_dimension_se 569 non-null float64
22 radius_worst 569 non-null float64
23 texture_worst 569 non-null float64
24 perimeter_worst 569 non-null float64
25 area_worst 569 non-null float64
26 smoothness_worst 569 non-null float64
27 compactness_worst 569 non-null float64
28 concavity_worst 569 non-null float64
29 concave points_worst 569 non-null float64
30 symmetry_worst 569 non-null float64
31 fractal_dimension_worst 569 non-null float64
31 fractal_dimension_worst 569 non-null float64
 13 texture_se
                                                                    569 non-null
                                                                                                           float64
 31 fractal dimension worst 569 non-null
                                                                                                           float64
dtypes: float64(30), int64(1), object(1)
memory usage: 142.4+ KB
Accuracy of the model using Decision tree regression algorithm is 0.6984126984126984
Accuracy of the model using Random forest regression algorithm is 0.852717328042328
```

Accuracy of the model using Support vector regression algorithm is 0.8114302960086689

<class 'pandas.core.frame.DataFrame'>

51. Build decision tree-based model for Breast Cancer Wisconsin (diagnostic) dataset.[Classifier]

```
In [16]: import numpy as n
         import pandas as p
         import seaborn as s
         import matplotlib.pyplot as m
         from sklearn.metrics import r2_score
         from sklearn.tree import DecisionTreeClassifier
         from sklearn.model selection import train test split
         from sklearn.metrics import accuracy score,confusion matrix
         data = p.read csv("breast-cancer.csv")
         data.info()
         data.drop(["id"],axis=1,inplace=True)
         M=data[data.diagnosis=="M"]
         B=data[data.diagnosis=="B"]
         m.title("Malignant vs Benign Tumor")
         m.xlabel("Radius Mean"); m.ylabel("Texture Mean")
         m.scatter(M.radius mean,M.texture mean,color='red',label='Malignant',alpha=0.3)
         m.scatter(B.radius mean, B.texture mean, color='lime', label='Bengin', alpha=0.4)
         m.legend(); m.show()
         data.diagnosis=[1 if i == "M" else 0 for i in data.diagnosis]
         x=data.drop(["diagnosis"],axis=1)
         y=data.diagnosis.values
         x_train,x_test,y_train,y_test=train_test_split(x,y,test_size=0.3,random_state=42)
         dt=DecisionTreeClassifier()
         dt.fit(x train,y train)
         y_pred=dt.predict(x_test)
         #Confusion Matrix
```

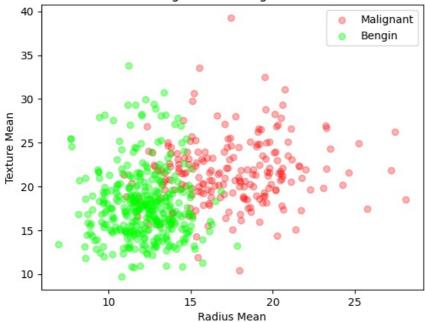
```
cm = confusion_matrix(y_test,y_pred)
s.heatmap(cm,annot=True,fmt='d',cmap="Blues")
m.title("Confusion matrics ")
m.show()
print("Accuracy of the classifier model is",accuracy_score(y_test,y_pred))
```

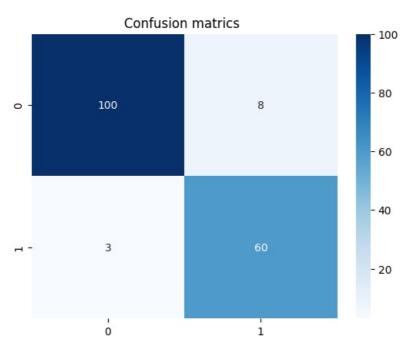
<class 'pandas.core.frame.DataFrame'> RangeIndex: 569 entries, 0 to 568 Data columns (total 32 columns):

#	Column	Non-Null Count	Dtype
0	id	569 non-null	int64
1	diagnosis	569 non-null	object
2	radius mean	569 non-null	float64
3	texture mean	569 non-null	float64
4	perimeter mean	569 non-null	float64
5	area_mean	569 non-null	float64
6	smoothness_mean	569 non-null	float64
7	compactness_mean	569 non-null	float64
8	concavity_mean	569 non-null	float64
9	concave points_mean	569 non-null	float64
10	symmetry_mean	569 non-null	float64
11	<pre>fractal_dimension_mean</pre>	569 non-null	float64
12	radius_se	569 non-null	float64
13	texture_se	569 non-null	float64
14	perimeter_se	569 non-null	float64
15	area_se	569 non-null	float64
16	smoothness_se	569 non-null	float64
17	compactness_se	569 non-null	float64
18	concavity_se	569 non-null	float64
19	concave points_se	569 non-null	float64
20	symmetry_se	569 non-null	float64
21	<pre>fractal_dimension_se</pre>	569 non-null	float64
22	radius_worst	569 non-null	float64
23	texture_worst	569 non-null	float64
24	perimeter_worst	569 non-null	float64
25	area_worst	569 non-null	float64
26	smoothness_worst	569 non-null	float64
27	compactness_worst	569 non-null	float64
28	concavity_worst	569 non-null	float64
29	concave points_worst	569 non-null	float64
30	symmetry_worst	569 non-null	float64
31	<pre>fractal_dimension_worst</pre>		float64
dtypes: float64(30), int64(1), object(1)			
memory usage: 142 4+ KB			

memory usage: 142.4+ KB

Malignant vs Benign Tumor





Accuracy of the classifier model is 0.935672514619883

In []:

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