Experiment_8_assessment

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- 1 Data Mining and machine Learning
- 2 Experiment 8
- 2.1 26 February
- 3 Name: Tufan Kundu
- 4 Reg no.: 24MDT0184
- 5 Decision Tree: Gradient boosting
- 6 Q1. Today we will try to see how gradient boosting can be implemented both manually and using the inbuilt classes.
- 6.0.1 importing th necessary libraries

```
[1]: import numpy as np
import pandas as pd
from sklearn.preprocessing import MinMaxScaler
from sklearn.model_selection import train_test_split
from sklearn.tree import DecisionTreeRegressor
from sklearn.metrics import mean_squared_error, r2_score
```

6.0.2 Loading the dataset

```
[2]: df = pd.read_csv(r"D:\study material\VIT_Data_Science\Winter_Sem\Data Mining_

→and Machine Learning Lab\Class_notes\ML_exp2\Book1.csv")

df
```

[2]:	price	area	bedrooms	bathrooms	stories	parking	furnishingstatus
0	13300000	7420	4	2	3	2	furnished
1	12250000	8960	4	4	4	3	furnished
2	12250000	9960	3	2	2	2	semi-furnished
3	12215000	7500	4	2	2	3	furnished
4	11410000	7420	4	1	2	2	furnished
		•••	•••		•••		

semi-furnished	0	2	1	3	5320	4550000	244
unfurnished	2	2	1	3	5360	4550000	245
semi-furnished	0	1	1	3	3520	4550000	246
unfurnished	3	4	1	4	8400	4550000	247
semi-furnished	0	1	2	2	4100	4543000	248

[249 rows x 7 columns]

6.0.3 Dropping the unnecessary column

```
[3]: df.drop('furnishingstatus',axis = 1, inplace = True) df
```

[3]:		price	area	bedrooms	bathrooms	stories	parking	
	0	13300000	7420	4	2	3	2	
	1	12250000	8960	4	4	4	3	
	2	12250000	9960	3	2	2	2	
	3	12215000	7500	4	2	2	3	
	4	11410000	7420	4	1	2	2	
		•••	•••	•••		•••		
	244	4550000	5320	3	1	2	0	
	245	4550000	5360	3	1	2	2	
	246	4550000	3520	3	1	1	0	
	247	4550000	8400	4	1	4	3	
	248	4543000	4100	2	2	1	0	

[249 rows x 6 columns]

6.0.4 Performing min-max scaling

```
[4]: scaler = MinMaxScaler()
X = scaler.fit_transform(df)
```

6.0.5 Setting the x and y(target) variable

6.0.6 Train test split

```
[6]: x_train,x_test,y_train,y_test = train_test_split(x,y,test_size=0.

→2,random_state=0)
```

6.0.7 Fitting the data in decision tree

6.1 Now we will see how we can implement the gradient boosting technique with inbuilt class.

MSE for GradientBoosting Regressor: 0.017811289322634986 R2 score for GradientBoosting Regressor: 0.27594382660242045

- 6.1.1 Error reduced significantly by using gradient boosting technique
- 6.2 Perform hyperparameter tuning using GridsearchCV by giving an parameter grid with the hyperparameters n estimators, learning rate and max depth. Each parameter should be having ateast 10 values in the parameter grid.

Best parameters:

```
{'learning_rate': 0.1, 'max_depth': 1, 'n_estimators': 700}
Best model :
```

GradientBoostingRegressor(max_depth=1, n_estimators=700)

y",mean_squared_error(y_test,y_pred))
print("r2 score:",r2 score(y test,y pred))

MSE(optimised after hyperparameter tuning): 0.018494705889616964 r2 score: 0.24816189709905545

- 6.3 Similarly you can import GradientBoostingClassifier from sklearn.ensemble. Use the liver patient dataset and fit a Decision tree and GradientBoostingClassfier
- 6.3.1 importing the libraries

```
[11]: from sklearn.tree import DecisionTreeClassifier from sklearn.ensemble import GradientBoostingClassifier from sklearn.metrics import accuracy_score
```

6.3.2 Loading the dataset

```
[12]: df = pd.read_csv(r"D:\study material\VIT_Data_Science\Winter_Sem\Data Mining

→ and Machine Learning Lab\Class_notes\ML_exp4\liver_patient.csv")

df
```

```
[12]:
                Gender Total Bilirubin Direct Bilirubin Alkaline Phosphotase
           Age
      0
            65
                Female
                                     0.7
                                                        0.1
                                                                                187
                                                        5.5
                                                                                699
      1
            62
                  Male
                                    10.9
      2
                  Male
                                     7.3
                                                        4.1
                                                                                490
            62
      3
            58
                  Male
                                     1.0
                                                        0.4
                                                                                182
      4
            72
                  Male
                                     3.9
                                                        2.0
                                                                                195
      578
            60
                  Male
                                     0.5
                                                        0.1
                                                                                500
```

```
579
      40
             Male
                                 0.6
                                                     0.1
                                                                              98
580
      52
             Male
                                 0.8
                                                     0.2
                                                                              245
581
             Male
                                 1.3
                                                     0.5
                                                                              184
      31
582
      38
                                 1.0
                                                     0.3
             Male
                                                                              216
     Alamine_Aminotransferase Aspartate_Aminotransferase Total_Protiens \
0
                              16
                                                             18
                                                                              6.8
1
                              64
                                                            100
                                                                             7.5
2
                              60
                                                             68
                                                                             7.0
3
                              14
                                                             20
                                                                             6.8
4
                              27
                                                             59
                                                                             7.3
                                                                             5.9
578
                              20
                                                             34
579
                              35
                                                             31
                                                                             6.0
580
                              48
                                                             49
                                                                             6.4
581
                              29
                                                             32
                                                                             6.8
582
                                                                             7.3
                              21
                                                             24
              Albumin_and_Globulin_Ratio
                                              liver_disease
0
          3.3
                                        0.90
          3.2
1
                                        0.74
                                                            1
2
          3.3
                                        0.89
                                                            1
3
          3.4
                                        1.00
                                                            1
4
          2.4
                                        0.40
                                                            1
. .
                                        0.37
                                                           0
578
          1.6
579
          3.2
                                        1.10
                                                            1
580
          3.2
                                        1.00
                                                            1
581
          3.4
                                        1.00
                                                            1
582
                                                            0
          4.4
                                        1.50
```

[583 rows x 11 columns]

6.3.3 Dropping unnecessary columns

```
[13]: df.drop(['Age','Gender'],axis =1 , inplace=True)
df
```

```
[13]:
           Total_Bilirubin Direct_Bilirubin Alkaline_Phosphotase
      0
                        0.7
                                            0.1
                                                                    187
      1
                       10.9
                                            5.5
                                                                   699
      2
                        7.3
                                            4.1
                                                                   490
      3
                        1.0
                                            0.4
                                                                   182
      4
                        3.9
                                            2.0
                                                                   195
      578
                        0.5
                                            0.1
                                                                   500
      579
                        0.6
                                            0.1
                                                                    98
```

```
0.8
                                                              245
580
                                      0.2
581
                  1.3
                                      0.5
                                                              184
582
                  1.0
                                      0.3
                                                              216
     Alamine_Aminotransferase Aspartate_Aminotransferase Total_Protiens \
0
                             16
                                                            18
                                                                            6.8
                             64
                                                           100
1
                                                                            7.5
2
                             60
                                                            68
                                                                            7.0
3
                             14
                                                            20
                                                                            6.8
4
                             27
                                                            59
                                                                            7.3
. .
578
                             20
                                                            34
                                                                            5.9
579
                             35
                                                            31
                                                                            6.0
                                                            49
                                                                            6.4
580
                             48
581
                             29
                                                            32
                                                                            6.8
582
                                                                            7.3
                             21
                                                            24
     Albumin Albumin_and_Globulin_Ratio liver_disease
         3.3
                                       0.90
0
         3.2
                                       0.74
1
                                                           1
2
         3.3
                                       0.89
                                                           1
3
         3.4
                                       1.00
                                                           1
4
         2.4
                                       0.40
                                                           1
578
         1.6
                                       0.37
                                                           0
         3.2
579
                                       1.10
                                                           1
580
         3.2
                                       1.00
                                                           1
581
         3.4
                                       1.00
                                                           1
         4.4
582
                                       1.50
                                                           0
[583 rows x 9 columns]
```

6.3.4 Min max scaling

6.3.5 Train test split

```
[16]: ##Fitting in decision tree classifier
model_dtc = DecisionTreeClassifier()
model_dtc.fit(x_train,y_train)
y_pred_dtc = model_dtc.predict(x_test)
```

Accuracy score when fitting the model via decision tree classifier:59.82905982905983 %
Accuracy score when fitting the model via gradient boosting classifier:65.8119658119658 %

6.4 Q2. Perform hyperparameter tuning on with the hyperparameters n estimators, learning rate and max depth. Each parameter should be having ateast 10 values in the parameter grid. Find the best combination of the parameters to get the better accuracy.

Best parameters:

```
{'learning_rate': 0.01, 'max_depth': 1, 'n_estimators': 400}
Best model:
GradientBoostingClassifier(learning_rate=0.01, max_depth=1, n_estimators=400)
```

Accuracy score after hyperparameter tuning:65.8119658119658 %

6.5 Reguarization techniques: Ridge and lasso regression.

Now we will try to look at ridge and lasso regression which are again regularization tech niques used to minimize the variance or reduce overfitting of data. The lasso regression also kind of helps to know the best features in the modeling. Because it will take some coefficients of the model which are not that relevant to zero

6.6 Q3. To perform ridge and lasso regression download the Book1.csv dataset to do house price prediction.

6.6.1 Reading the dataset

[37]:		price	area	bedrooms	bathrooms	stories	parking
	0	13300000	7420	4	2	3	2
	1	12250000	8960	4	4	4	3
	2	12250000	9960	3	2	2	2
	3	12215000	7500	4	2	2	3
	4	11410000	7420	4	1	2	2
		•••	•••	•••		•••	
	244	4550000	5320	3	1	2	0
	245	4550000	5360	3	1	2	2
	246	4550000	3520	3	1	1	0
	247	4550000	8400	4	1	4	3
	248	4543000	4100	2	2	1	0

[249 rows x 6 columns]

6.6.2 Min max scaling

```
[38]: X = scaler.fit_transform(df)
```

6.6.3 Setting x and y

6.6.4 Train test split

```
[40]: x_train,x_test,y_train,y_test = train_test_split(x,y,test_size=0.

-2,random_state=0)
```

```
[41]: from sklearn.linear_model import RidgeCV, LassoCV
      ## Defining alpha values for tuning
     alpha_values = np.logspace(-2,4,100)
     print()
     #-----
      # Ridge Regression with RidgeCV
      #-----
     ridge_cv = RidgeCV(alphas = alpha_values, store_cv_values = True)
     ridge_cv.fit(x_train,y_train)
     ridge_pred = ridge_cv.predict(x_test)
     print("Best Ridge Alpha:",ridge_cv.alpha_)
     print("Ridge Regression MSE:", mean_squared_error(y_test,ridge_pred))
     print("Ridge coefficients:",ridge_cv.coef_)
     print()
     print()
      #Lasso regression with Lassocv
     lasso_cv = LassoCV(alphas=alpha_values,cv = 5, random_state = 0)
     lasso_cv.fit(x_train,y_train)
     lasso_pred = lasso_cv.predict(x_test)
     print("Best Lasso Alpha:",lasso_cv.alpha_)
     print("Lasso Regression MSE:", mean_squared_error(y_test,lasso_pred))
     print("lasso coefficients:",lasso_cv.coef_)
```

```
Best Ridge Alpha: 0.49770235643321115
Ridge Regression MSE: 0.019544354076968272
Ridge coefficients: [0.2916599 0.0894537 0.30197724 0.13122731 0.14923577]
```

Best Lasso Alpha: 0.01 Lasso Regression MSE: 0.01991781625293486 lasso coefficients: [0. 0. 0.13504156 0.07345709 0.07836195]

6.7 Stacking

```
[33]: import warnings
      warnings.filterwarnings("ignore")
[35]: # Import necessary libraries
      import numpy as np
      import pandas as pd
      import matplotlib.pyplot as plt
      from sklearn import preprocessing
      from sklearn.model selection import train test split
      from sklearn.metrics import accuracy score
      from sklearn.tree import DecisionTreeClassifier
      from sklearn.ensemble import BaggingClassifier, RandomForestClassifier,
       →AdaBoostClassifier, StackingClassifier
      from sklearn.linear_model import LogisticRegression
      # Load dataset
      data = pd.read_csv(r"D:\study material\VIT_Data_Science\Winter_Sem\Data_Mining_

→and Machine Learning Lab\Class_notes\ML_exp4\liver_patient.csv")
      # Extract target variable (Y) and drop unnecessary columns
      Y = data.liver_disease
      data.drop(['Age', 'Gender', 'liver_disease'], axis=1, inplace=True)
      # Normalize data using MinMaxScaler
      scaler = preprocessing.MinMaxScaler()
      X_scaled = scaler.fit_transform(data)
      X = pd.DataFrame(X_scaled[:, 0:8]) # Retaining first 8 normalized features
      # Split dataset into training and testing sets (90% train, 10% test)
      X_train, X_test, y_train, y_test = train_test_split(X, Y, test_size=0.10, __
       →random_state=0)
      # Define base classifiers
      DT = DecisionTreeClassifier()
      BC = BaggingClassifier(n_estimators=10, random_state=0) # Bagging Classifier
      PC = BaggingClassifier(n_estimators=10, bootstrap=True, random_state=0) #__
       → Pasting Classifier
      RFC = RandomForestClassifier(n_estimators=10, max_features="sqrt", __
       ⇒random_state=0) # Random Forest
      ABC = AdaBoostClassifier(estimator=DecisionTreeClassifier(max_depth=1),__
       on_estimators=500, random_state=0) # AdaBoost
      # Train the base models
      DT.fit(X_train, y_train)
      BC.fit(X_train, y_train)
```

```
PC.fit(X_train, y_train)
RFC.fit(X_train, y_train)
ABC.fit(X_train, y_train)
# Make predictions
pred_DT = DT.predict(X_test)
pred_BC = BC.predict(X_test)
pred_PC = PC.predict(X_test)
pred RFC = RFC.predict(X test)
pred_ABC = ABC.predict(X_test)
# Print accuracy of individual models
print(f"Decision Tree Accuracy:{accuracy_score(y_test, pred_DT)*100} %")
print(f"Bagging Accuracy:{accuracy_score(y_test, pred BC)*100} %")
print(f"Pasting Accuracy:{accuracy_score(y_test, pred PC)*100} %")
print(f"Random Forest Accuracy:{accuracy_score(y_test, pred_RFC)*100} %")
print(f"AdaBoost Accuracy:{accuracy_score(y_test, pred ABC)*100} %")
# Define Stacking Classifier with Logistic Regression as the final estimator
estimators = [('dt', DT), ('bc', BC), ('pc', PC), ('rfc', RFC), ('abc', ABC)]
stk = StackingClassifier(estimators=estimators,__
 →final_estimator=LogisticRegression(), passthrough=True)
# Train the stacking classifier
stk.fit(X_train, y_train)
# Make predictions with stacking classifier
pred_stk = stk.predict(X_test)
# Print accuracy of Stacking Classifier
print(f"Stacking Accuracy:{accuracy_score(y_test, pred_stk)*100} %")
Decision Tree Accuracy:66.10169491525424 %
```

Decision Tree Accuracy:66.10169491525424 % Bagging Accuracy:72.88135593220339 % Pasting Accuracy:72.88135593220339 % Random Forest Accuracy:74.57627118644068 % AdaBoost Accuracy:77.96610169491525 % Stacking Accuracy:72.88135593220339 %