### ML LAB 8

**Name: Soumyadeep Ganguly** 

**Reg No: 24MDT0082** 

```
In [4]: import pandas as pd
        import numpy as np
        import matplotlib.pyplot as plt
In [5]: df = pd.read csv('Book1.csv')
        df.head()
        df = df.drop(['furnishingstatus'], axis = 1)
        df.head()
Out[5]:
              price area bedrooms bathrooms stories parking
        0 13300000 7420
        1 12250000 8960
        2 12250000 9960
        3 12215000 7500
        4 11410000 7420
                                            1
                                                    2
In [6]: from sklearn.preprocessing import MinMaxScaler
        scaler = MinMaxScaler(feature_range=(0,1))
        scaled = scaler.fit transform(df)
        df_scaled = pd.DataFrame(scaled, columns = df.columns)
        df scaled.head()
```

```
Out[6]:
             price
                     area bedrooms bathrooms
                                               stories parking
       0 1.000000 0.356777
                               0.50
                                     1 0.880096 0.469597
                               0.50
                                     1.000000 1.000000 1.000000
       2 0.880096 0.542857
                               0.25
                                     3 0.876099 0.362637
                               0.50
                                     4 0.784173 0.356777
                               0.50
                                     0.000000 0.333333 0.666667
In [7]: from sklearn.model selection import train test split
       X = df scaled.drop(['price'], axis=1)
       y = df scaled['price']
       X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = 0.2, random_state= 42)
       from sklearn.tree import DecisionTreeRegressor
In [8]:
       DTR = DecisionTreeRegressor()
       DTR.fit(X train, y train)
Out[8]:
        ▼ DecisionTreeRegressor
       DecisionTreeRegressor()
In [9]: y pred = DTR.predict(X test)
       y_pred
```

Error is: 0.07675530700761357

## **Gradient Boosting Regressor**

```
Out[12]: array([0.17414174, 0.4752913, 0.17071984, 0.23444211, 0.19341953,
                0.20503384, 0.15387844, 0.18093334, 0.23547499, 0.20453393,
                0.15984301, 0.15009756, 0.12134007, 0.15984301, 0.40508986,
                 0.18893 , 0.11118068, 0.18085147, 0.21885721, 0.42330002,
                 0.11107113, 0.23444211, 0.18048463, 0.14834259, 0.42330002,
                 0.21979456, 0.16655274, 0.19840865, 0.42433569, 0.18048463,
                0.12134007, 0.23172445, 0.21271041, 0.12516125, 0.23494203,
                0.1991435 , 0.22773238, 0.23222437, 0.17213165, 0.18093334,
                0.15285543, 0.18093334, 0.15200926, 0.14644912, 0.277752 ,
                0.23494203, 0.25775041, 0.42330002, 0.1991435, 0.17370852])
In [13]: GBR Error = mean squared error(y pred, y test)
         print(f'Error is: {GBR Error}')
        Error is: 0.026793767598257433
In [14]: from sklearn.model selection import GridSearchCV
         param grid = {
           'n estimators':[ i for i in range(50, 501, 50)],
          'learning rate': list(np.linspace(0,1, 10)),
           'max depth': list(np.random.randint(1,10, 10))
         base model = GradientBoostingRegressor(random state=0)
In [15]: gs = GridSearchCV(estimator = base model, param grid = param grid, cv=5, n jobs = -1)
         gs.fit(X train, y train)
Out[15]:
                           GridSearchCV
                         best estimator :
                   GradientBoostingRegressor
                ▶ GradientBoostingRegressor
        print(f'Best parameters for "Gradient Boosting Regressor": {gs.best_params_}')
         print(f'Best Accuracy: {gs.best score }')
```

```
Best parameters for "Gradient Boosting Regressor": {'learning_rate': np.float64(0.11111111111111), 'max_depth': np.int32(1), 'n_estimators': 200}
Best Accuracy: 0.24953014819241837
```

## liver\_patient.csv

```
In [17]: df2 = pd.read csv("liver patient.csv")
         df2 = df2.drop(['Age', 'Gender'], axis = 1)
         df2.head()
Out[17]:
             Total_Bilirubin Direct_Bilirubin Alkaline_Phosphotase Alamine_Aminotransferase Aspartate_Aminotransferase Total_Protiens Albumin
                       0.7
                                       0.1
                                                            187
                                                                                      16
                                                                                                                 18
                                                                                                                                6.8
                                                                                                                                         3.3
          0
                      10.9
                                       5.5
                                                            699
                                                                                      64
                                                                                                                 100
                                                                                                                                7.5
                                                                                                                                         3.2
          1
          2
                       7.3
                                       4.1
                                                            490
                                                                                      60
                                                                                                                  68
                                                                                                                                7.0
                                                                                                                                         3.3
          3
                                                                                                                                6.8
                                                                                                                                         3.4
                       1.0
                                       0.4
                                                            182
                                                                                      14
                                                                                                                  20
                       3.9
                                       2.0
                                                            195
                                                                                      27
                                                                                                                                7.3
                                                                                                                                         2.4
          4
                                                                                                                  59
         scaler2 = MinMaxScaler(feature range=(0,1))
         scaled2 = scaler2.fit transform(df2)
         df2 scaled = pd.DataFrame(scaled2, columns = df2.columns)
         df2 scaled.head()
```

```
Out[18]:
             Total Bilirubin Direct Bilirubin Alkaline Phosphotase Alamine Aminotransferase Aspartate Aminotransferase Total Protiens Albumin
          0
                  0.004021
                                 0.000000
                                                      0.060576
                                                                                0.003015
                                                                                                           0.001626
                                                                                                                         0.594203 0.521739
                  0.140751
                                                      0.310699
                                                                                0.027136
                                                                                                           0.018296
                                                                                                                         0.695652  0.500000
          1
                                 0.275510
          2
                  0.092493
                                 0.204082
                                                                                0.025126
                                                      0.208598
                                                                                                           0.011791
                                                                                                                         0.623188 0.521739
          3
                  0.008043
                                 0.015306
                                                      0.058134
                                                                                0.002010
                                                                                                           0.002033
                                                                                                                         0.594203 0.543478
          4
                  0.046917
                                 0.096939
                                                      0.064485
                                                                                0.008543
                                                                                                           0.009961
                                                                                                                         0.666667 0.326087
         X2 = df2 scaled.drop(['liver disease'], axis = 1)
         y2 = df2 scaled['liver disease']
         X train2, X test2, y train2, y test2 = train test split(X2, y2, test size = 0.2, random state=0)
In [20]:
         from sklearn.tree import DecisionTreeClassifier
         from sklearn.metrics import accuracy score
         DTC = DecisionTreeClassifier()
         DTC.fit(X train2, y train2)
Out[20]:
          ▼ DecisionTreeClassifier
         DecisionTreeClassifier()
In [21]:
         y pred2 = DTC.predict(X test2)
In [22]: DTC score = accuracy score(y test2, y pred2)
         print(f'Error is: {DTC score}')
        Error is: 0.5811965811965812
In [23]: from sklearn.ensemble import GradientBoostingClassifier
         GBC = GradientBoostingClassifier(learning rate=0.01, n estimators=100, max depth = 3, random state=0)
         GBC.fit(X train2, y train2)
```

#### Q2

```
In [24]:
         param grid2 = {
          'n estimators':[ i for i in range(50, 501, 50)],
           'learning rate': list(np.linspace(0,1, 10)),
           'max depth': list(np.random.randint(1,10, 10))
         base model2 = GradientBoostingClassifier(random state=0)
         gs2 = GridSearchCV(estimator = base model2, param grid = param grid2,cv=5, n jobs = -1)
         gs2.fit(X train2, y train2)
Out[24]: ▶
                           GridSearchCV
                          best estimator :
                   GradientBoostingClassifier
                ▶ GradientBoostingClassifier
In [25]: print(f'Best parameters for "Gradient Boosting Classifier": {gs2.best params }')
         print(f'Best Accuracy: {gs2.best score }')
        Best parameters for "Gradient Boosting Classifier": {'learning rate': np.float64(0.111111111111111), 'max depth': np.int32(1),
        'n estimators': 50}
        Best Accuracy: 0.7424388011896592
```

# Reguarization techniques: Ridge and lasso regression.

Now we will try to look at ridge and lasso regression which are again regularization techniques 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

### Q3

```
In [32]: from sklearn.linear model import RidgeCV, LassoCV
In [26]: df3 = df.copy()
         df3.head()
         scaler3 = MinMaxScaler()
In [28]: df3.head(2)
Out[28]:
               price area bedrooms bathrooms stories parking
         0 13300000 7420
         1 12250000 8960
                                                            3
In [29]: data = scaler3.fit transform(df)
         df3 = pd.DataFrame(data=data, columns=df3.columns)
         df3.head(2)
Out[29]:
               price
                        area bedrooms bathrooms
                                                    stories
                                                          parking
         0 1.000000 0.356777
                                         0.5
         1 0.880096 0.469597
                                         1.000000 1.000000 1.000000
In [30]: X = df3.iloc[:,1:].values
         y = df3.iloc[:,0].values
         X train, X test, y train, y test = train test split(X, y, test size=0.2, random state=0)
In [31]:
        alpha_values = np.logspace(-2,4,100)
         alpha values
```

```
Out[31]: array([1.0000000e-02, 1.14975700e-02, 1.32194115e-02, 1.51991108e-02,
                1.74752840e-02, 2.00923300e-02, 2.31012970e-02, 2.65608778e-02,
                 3.05385551e-02, 3.51119173e-02, 4.03701726e-02, 4.64158883e-02,
                 5.33669923e-02, 6.13590727e-02, 7.05480231e-02, 8.11130831e-02,
                 9.32603347e-02, 1.07226722e-01, 1.23284674e-01, 1.41747416e-01,
                1.62975083e-01, 1.87381742e-01, 2.15443469e-01, 2.47707636e-01,
                 2.84803587e-01, 3.27454916e-01, 3.76493581e-01, 4.32876128e-01,
                 4.97702356e-01, 5.72236766e-01, 6.57933225e-01, 7.56463328e-01,
                 8.69749003e-01, 1.00000000e+00, 1.14975700e+00, 1.32194115e+00,
                1.51991108e+00, 1.74752840e+00, 2.00923300e+00, 2.31012970e+00,
                 2.65608778e+00, 3.05385551e+00, 3.51119173e+00, 4.03701726e+00,
                4.64158883e+00, 5.33669923e+00, 6.13590727e+00, 7.05480231e+00,
                 8.11130831e+00, 9.32603347e+00, 1.07226722e+01, 1.23284674e+01,
                1.41747416e+01, 1.62975083e+01, 1.87381742e+01, 2.15443469e+01,
                 2.47707636e+01, 2.84803587e+01, 3.27454916e+01, 3.76493581e+01,
                4.32876128e+01, 4.97702356e+01, 5.72236766e+01, 6.57933225e+01,
                7.56463328e+01, 8.69749003e+01, 1.00000000e+02, 1.14975700e+02,
                1.32194115e+02, 1.51991108e+02, 1.74752840e+02, 2.00923300e+02,
                 2.31012970e+02, 2.65608778e+02, 3.05385551e+02, 3.51119173e+02,
                4.03701726e+02, 4.64158883e+02, 5.33669923e+02, 6.13590727e+02,
                7.05480231e+02, 8.11130831e+02, 9.32603347e+02, 1.07226722e+03,
                1.23284674e+03, 1.41747416e+03, 1.62975083e+03, 1.87381742e+03,
                 2.15443469e+03, 2.47707636e+03, 2.84803587e+03, 3.27454916e+03,
                 3.76493581e+03, 4.32876128e+03, 4.97702356e+03, 5.72236766e+03,
                 6.57933225e+03, 7.56463328e+03, 8.69749003e+03, 1.00000000e+04])
In [33]: ridge cv = RidgeCV(alphas=alpha values, store cv values=True)
         ridge cv.fit(X train,y train)
        e:\VIT Study Materials\SEM 2\Data Mining and ML\LAB\.venv\Lib\site-packages\sklearn\linear_model\_ridge.py:2375: FutureWarning:
        'store cv values' is deprecated in version 1.5 and will be removed in 1.7. Use 'store cv results' instead.
          warnings.warn(
```

```
Out[33]:
                                                RidgeCV
         RidgeCV(alphas=array([1.00000000e-02, 1.14975700e-02, 1.32194115e-02, 1.51991108e-02,
                1.74752840e-02, 2.00923300e-02, 2.31012970e-02, 2.65608778e-02,
                3.05385551e-02, 3.51119173e-02, 4.03701726e-02, 4.64158883e-02,
                5.33669923e-02, 6.13590727e-02, 7.05480231e-02, 8.11130831e-02,
                9.32603347e-02, 1.07226722e-01, 1.23284674e-01, 1.41747416e-01,
                1.62975083e-01, 1.87381742e-0...
                4.03701726e+02, 4.64158883e+02, 5.33669923e+02, 6.13590727e+02,
                7.05480231e+02, 8.11130831e+02, 9.32603347e+02, 1.07226722e+03,
                1.23284674e+03, 1.41747416e+03, 1.62975083e+03, 1.87381742e+03,
In [34]: ridge pred = ridge cv.predict(X test)
In [35]: ridge cv.alpha
Out[35]: np.float64(0.49770235643321115)
        mean squared error(y test, ridge pred)
         0.019544354076968272
Out[37]:
In [38]: ridge cv.coef
Out[38]: array([0.2916599, 0.0894537, 0.30197724, 0.13122731, 0.14923577])
In [39]: lasso cv = LassoCV(alphas=alpha values,cv=5,random state=0)
         lasso cv.fit(X train, y train)
```

```
Out[39]:
                                               LassoCV
        LassoCV(alphas=array([1.00000000e-02, 1.14975700e-02, 1.32194115e-02, 1.51991108e-02,
               1.74752840e-02, 2.00923300e-02, 2.31012970e-02, 2.65608778e-02,
               3.05385551e-02, 3.51119173e-02, 4.03701726e-02, 4.64158883e-02,
                5.33669923e-02, 6.13590727e-02, 7.05480231e-02, 8.11130831e-02,
               9.32603347e-02, 1.07226722e-01, 1.23284674e-01, 1.41747416e-01,
               1.62975083e-01, 1.87381742e-0...
               4.03701726e+02, 4.64158883e+02, 5.33669923e+02, 6.13590727e+02,
               7.05480231e+02, 8.11130831e+02, 9.32603347e+02, 1.07226722e+03,
               1.23284674e+03, 1.41747416e+03, 1.62975083e+03, 1.87381742e+03,
In [41]: lasso pred = lasso cv.predict(X test)
In [42]: lasso cv.alpha
```

Out[42]: np.float64(0.01)

lasso cv.coef In [43]:

, 0. Out[43]: array([0. , 0.13504156, 0.07345709, 0.07836195])

In [45]: mean squared error(y test, lasso pred)

Out[45]: 0.01991781625293486

## **Stacking**

In [48]: from sklearn.ensemble import BaggingClassifier, RandomForestClassifier, AdaBoostClassifier, StackingClassifier from sklearn.linear model import LogisticRegression

```
In [55]: data = pd.read csv('liver patient.csv')
         v = data.liver disease
         data.drop(['Age', 'Gender', 'liver disease'], axis=1, inplace=True)
In [56]: MM = MinMaxScaler()
         X1 = MM.fit transform(data)
         X = pd.DataFrame(X1[:, 0:8])
         X train, X test, y train, y test = train test split(X, y, test size=0.10, random state=0)
In [57]: DT = DecisionTreeClassifier()
         BC = BaggingClassifier(n estimators=10, random state=0)
         PC = BaggingClassifier(n estimators=10, bootstrap=True, random state=0)
         RFC = RandomForestClassifier(n estimators=10, max features="sqrt", random state=0)
         ABC = AdaBoostClassifier(estimator=DecisionTreeClassifier(max depth=1), n estimators=500, random state=0)
         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)
         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("Decision Tree Accuracy:",accuracy score(y test,pred DT))
In [58]:
         print("Bagging Accuracy:",accuracy score(y test,pred BC))
         print("Pasting Accuracy:",accuracy score(y test,pred PC))
         print("Random Forest Accuracy:",accuracy score(y test,pred RFC))
         print("AdaBoost Accuracy:",accuracy score(y test,pred ABC))
        Decision Tree Accuracy: 0.6440677966101694
        Bagging Accuracy: 0.7288135593220338
        Pasting Accuracy: 0.7288135593220338
        Random Forest Accuracy: 0.7457627118644068
        AdaBoost Accuracy: 0.7457627118644068
```

```
In [59]: estimators=[('dt',DT),('bc',BC),('pc',PC),('rfc',RFC),('abc',ABC)]
    stk=StackingClassifier(estimators=estimators, final_estimator=LogisticRegression(), passthrough=True)
    stk.fit(X_train,y_train)
    pred_stk=stk.predict(X_test)
    print("Stacking Accuracy:",accuracy_score(y_test,pred_stk))

Stacking Accuracy: 0.7288135593220338
In []:
```