

# Introduction to Machine learning Project Phase 2 Team 8

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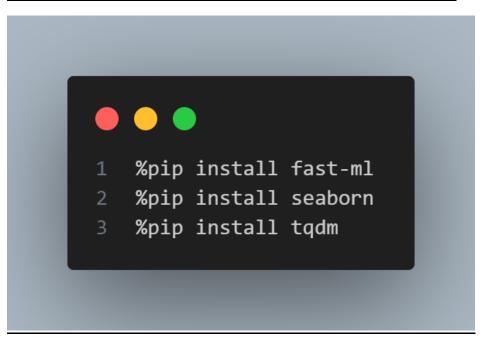
**Eng. Omar Elessawy** 

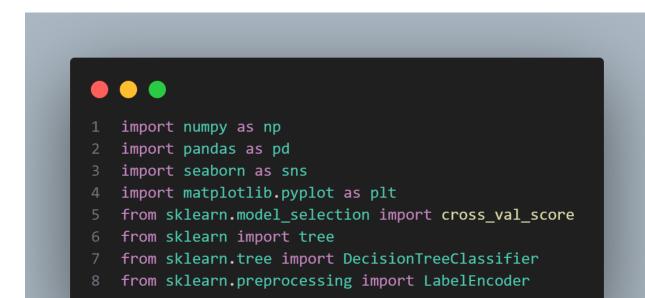
## **Decision Trees**

## Steps on how to run project

- 1) Open Decision tree.ipynb and then run the first cell to import all the required libraries
- 2) Project is ready to run!

## **Screenshots of the Code including the output**





```
train_dataset = pd.read_csv('updated_train.csv')
label_encoder = LabelEncoder()
train_dataset['Sex'] = label_encoder.fit_transform(train_dataset['Sex'])
```

```
# hyperparameter optemization
from sklearn.model_selection import GridSearchCV
from fast_ml.model_development import train_valid_test_split

X_train, y_train, X_valid, y_valid, X_test, y_test = train_valid_test_split(train_dataset, target = 'Survived',
train_size=0.7, valid_size=0.15, test_size=0.15)
```

```
1 #defining and Fitting decision tree model
            classifier = DecisionTreeClassifier()
           classifier.fit(X_train, y_train)
           y_pred = classifier.predict(X_test)
           from sklearn.tree import plot_tree, export_text
            plt.figure(figsize =(80,20))
            plot_tree(classifier, feature_names=X_train.columns, max_depth=2, filled=True);
                                                                                Sex <= 0.5
gini = 0.469
samples = 623
value = [389, 234]
                              Pclass <= 4.0
gini = 0.41
samples = 212
value = [61, 151]
                                                                                                                                    PCA3 <= 0.125
gini = 0.322
samples = 411
value = [328, 83]
      PCA1 <= 1.599
gini = 0.101
samples = 112
value = [6, 106]
                                                        PCA2 <= -0.286
gini = 0.495
samples = 100
value = [55, 45]
                                                                                                          Pclass <= 2.0
gini = 0.284
samples = 374
value = [310, 64]
                                                                                                                                                             PCA2 <= -0.246
gini = 0.5
                                                                                                                                                             samples = 37
value = [18, 19]
                                                                                                                               (...)
(...)
                          (...)
                                                  (...)
                                                                            (...)
                                                                                                     (...)
                                                                                                                                                        (...)
                                                                                                                                                                                 (...)
• • •
   param_grid = {
    'criterion': ['gini', 'entropy'],
    'max_depth': [None, 10, 20, 30, 40, 50, 60, 70, 80],
    'min_samples_split': [2, 5, 10, 15],
    'min_samples_leaf': [1, 2, 3, 4, 5, 6]
```

gs = GridSearchCV(classifier, param\_grid=param\_grid,scoring=["accuracy","f1","precision","recall","roc\_auc"],cv=10,n\_jobs=-1,refit="accuracy")
g\_res = gs.fit(X\_valid, y\_valid)
g\_res.best\_score\_

0.8137362637362637

```
1  # get the hyperparameters with the best score
2  g_res.best_params_

{'criterion': 'gini',
   'max_depth': None,
   'min_samples_leaf': 3,
   'min_samples_split': 10}
```

These are the optimal hyperparameters for the Decision tree classifier

```
1 y_pred = g_res.predict(X_test)
```

```
# precession score
from sklearn.metrics import precision_score
print("Precession Score:",precision_score(y_test, y_pred, average='macro'))
```

Precesion Score: 0.7868487043744776

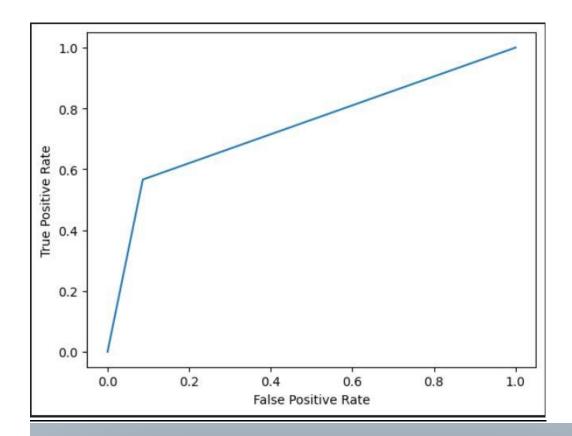
```
# Recall Score
from sklearn.metrics import recall_score
print("Recall Score:",recall_score(y_test, y_pred, average='macro'))
```

Recall Score: 0.7398089913813184

```
# F1-Score
from sklearn.metrics import f1_score
print("F1-Score:",f1_score(y_test, y_pred, average='macro'))
```

F1-Score: 0.749063670411985

```
# ROC/AUC Curves
%matplotlib inline
from sklearn.metrics import roc_curve
def plot_roc_curve(true_y, y_prob):
    fpr, tpr, thresholds = roc_curve(true_y, y_prob)
    plt.plot(fpr, tpr)
    plt.xlabel('False Positive Rate')
    plt.ylabel('True Positive Rate')
    plot_roc_curve(y_test,y_pred)
```



```
# ROC AUC Score
from sklearn.metrics import roc_auc_score
r_a_score = roc_auc_score(y_test, y_pred)
print("ROC-AUC-Score:", r_a_score)
```

ROC-AUC-Score: 0.7398089913813184

```
# Extract the F1 scores and hyperparameters from the cv_results_

f1_scores = g_res.cv_results_['mean_test_f1']

params = g_res.cv_results_['params']

# Extract the values for each hyperparameter

max_depth_values = [param['max_depth'] for param in params]

# Create a 2D scatter plot to visualize the F1 scores

plt.figure(figsize=(12, 8))

plt.scatter(max_depth_values, f1_scores, c=f1_scores, cmap='viridis', marker='o')

plt.xlabel('Max Depth')

plt.ylabel('Mean Test F1 Score')

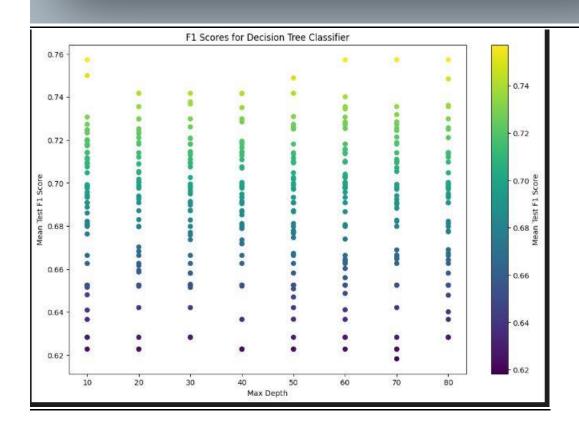
# Add a colorbar to the right of the plot

cbar = plt.colorbar()

cbar.set_label('Mean Test F1 Score')

plt.show()

plt.show()
```



```
# Extract the Accuracy scores and hyperparameters from the cv_results_
accuracy_scores = g_res.cv_results_['mean_test_accuracy']
params = g_res.cv_results_['params']

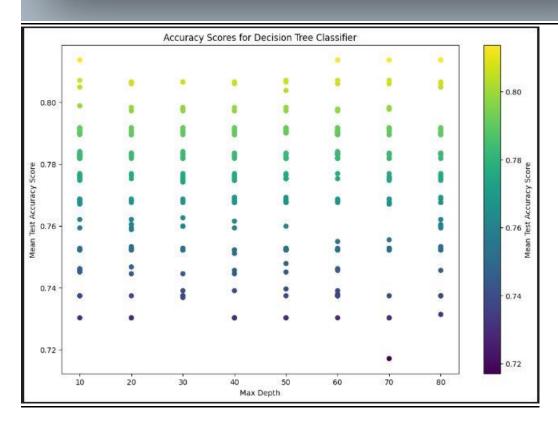
# Extract the values for each hyperparameter
max_depth_values = [param['max_depth'] for param in params]

# Create a 2D scatter plot to visualize the Accuracy scores
plt.figure(figsize=(12, 8))
plt.scatter(max_depth_values, accuracy_scores, c=accuracy_scores, cmap='viridis', marker='o')

plt.xlabel('Max Depth')
plt.ylabel('Mean Test Accuracy Score')

# Add a colorbar to the right of the plot
cbar = plt.colorbar()
cbar.set_label('Mean Test Accuracy Score')

plt.show()
```



```
# Extract the Recall scores and hyperparameters from the cv_results_
recall_scores = g_res.cv_results_['mean_test_recall']

params = g_res.cv_results_['params']

# Extract the values for each hyperparameter
max_depth_values = [param['max_depth'] for param in params]

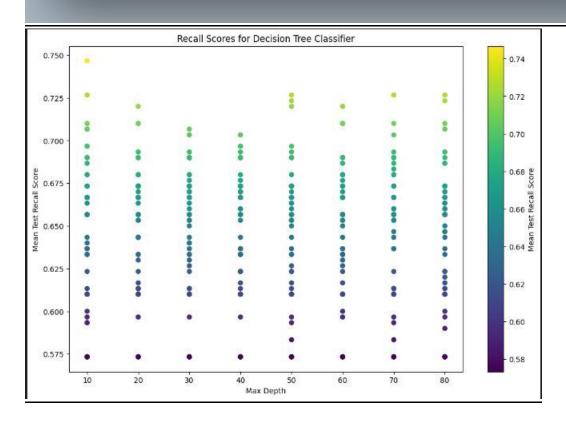
# Create a 2D scatter plot to visualize the Recall scores
plt.figure(figsize=(12, 8))
plt.scatter(max_depth_values, recall_scores, c=recall_scores, cmap='viridis', marker='o')

plt.xlabel('Max Depth')
plt.ylabel('Mean Test Recall Score')

# Add a colorbar to the right of the plot
cbar = plt.colorbar()
cbar.set_label('Mean Test Recall Score')

plt.show()

plt.show()
```



```
# Extract the Precision scores and hyperparameters from the cv_results_

precision_scores = g_res.cv_results_['mean_test_precision']

params = g_res.cv_results_['params']

# Extract the values for each hyperparameter

max_depth_values = [param['max_depth'] for param in params]

# Create a 2D scatter plot to visualize the Precision scores

plt.figure(figsize=(12, 8))

plt.scatter(max_depth_values, precision_scores, c=precision_scores, cmap='viridis', marker='o')

plt.xlabel('Max Depth')

plt.ylabel('Mean Test Precision Score')

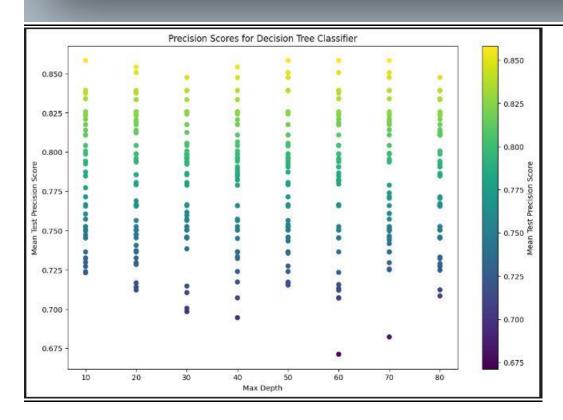
# Add a colorbar to the right of the plot

cbar = plt.colorbar()

cbar.set_label('Mean Test Precision Score')

plt.show()

plt.show()
```



#### **Multilayer Perceptron**

### Steps on how to run project

- 1) Open Multilayer-Perceptron.ipynb and then run the first cell to import all the required libraries
- 2) Project is ready to run!

#### **Code Screenshots and Details**

```
%pip install fast-ml
%pip install scikit-learn
%pip install numpy
%pip install pandas
%pip install seaborn
```

First, we install all the required libraries for initializing and training our multilayer perceptron model.

```
import pandas as pd
from sklearn.preprocessing import LabelEncoder

train_dataset = pd.read_csv('../updated_train.csv')
label_encoder = LabelEncoder()
train_dataset['Sex'] = label_encoder.fit_transform(train_dataset['Sex'])

X = train_dataset.drop("Survived", axis=1)
y = train_dataset["Survived"]
```

Then we import our preprocessed dataset and encode the "Sex" Column as Multilayer perceptron only accepts numerical features. Afterwards, we split the data into input features (X) and the output (y)

```
from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.85,test_size=0.15)
```

Then we split the input and output features into training and testing sets, but this time we won't use the fast-ml library, as there is no need to create a validation set, which we will see as progress on, will be generated by the multilayer perceptron model via the validation\_factor parameter.

```
def generate_hidden_layer_sizes(num_sizes, min_nodes, max_nodes):
    return [(np.random.randint(min_nodes, max_nodes),) for _ in
range(num_sizes)]
```

This is a method we will use to generate a list of different hidden\_layer\_sizes (the number of hidden nodes per layer) that will be supplied to GridSearchCV to determine the optimal hyperparameter as per the supplied parameter grid.

```
import numpy as np
num_hidden_sizes = 10

min_nodes_per_layer = 50
max_nodes_per_layer = 250

hidden_layer_sizes = generate_hidden_layer_sizes(num_hidden_sizes,
min_nodes_per_layer, max_nodes_per_layer)
```

Now, we will call the generate\_hidden\_layer\_sizes method to generate different hidden layer sizes as described above

```
from sklearn.neural_network import MLPClassifier
from sklearn.model_selection import GridSearchCV
param_grid = {
    "activation": ["identity", "logistic", "tanh", "relu",],
    "solver": ["lbfgs", "sgd", "adam",],
    "learning_rate": ["constant", "invscaling", "adaptive", ],
    "early_stopping": [True, False,],
    "validation_fraction": [0.15,],
    "hidden_layer_sizes": hidden_layer_sizes
mlp = GridSearchCV(
    MLPClassifier(),
    param_grid=param_grid,
    scoring=["accuracy","f1","precision","recall","roc_auc"],
    n_jobs=-1,
    cv=2,
    refit="f1",
    verbose=True
mlp.fit(X_train,y_train)
mlp.best_score_
```

Finally, we will call the GridSearchCV object to supply it with a multilayer perceptron object, and parameter grid that act our hyperparameters with almost each and every possible value of each hyperparameter, and only two folds for simplicity, and the refit is going to be based primarily on F1-score

Then we will print our best achieved score, which is approximately 75%

#### 0.7466949109284875

```
y_pred = mlp.predict(X_test)
```

In this line of code, we just create a variable that will be used for calculating precision, recall, f1-score, and roc/auc measures

```
from sklearn.metrics import recall_score
print(f"Multilayer perceptron recall score:{recall_score(y_test, y_pred)}")

✓ 0.0s
Multilayer perceptron recall score:0.6862745098039216
```

```
from sklearn.metrics import f1_score
print(f"Multilayer perceptron f1 score:{f1_score(y_test, y_pred)}")

✓ 0.0s

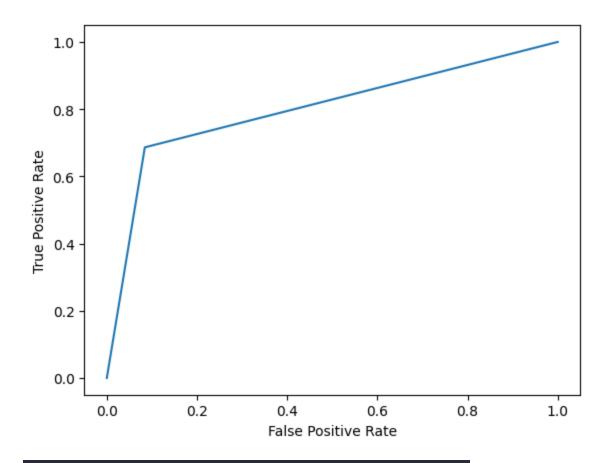
Multilayer perceptron f1 score:0.7526881720430109
```

```
import matplotlib.pyplot as plt
from sklearn.metrics import roc_curve

def plot_roc_curve(true_y, y_prob):
    fpr, tpr, thresholds = roc_curve(true_y, y_prob)
    plt.plot(fpr, tpr)
    plt.xlabel('False Positive Rate')
    plt.ylabel('True Positive Rate')
```

Above is a function to plot roc/auc curve

```
from sklearn.metrics import roc_auc_score
plot_roc_curve(y_test, y_pred)
plt.show()
print(f'Multilayer perceptron AUC score: {roc_auc_score(y_test, y_pred)}')
```

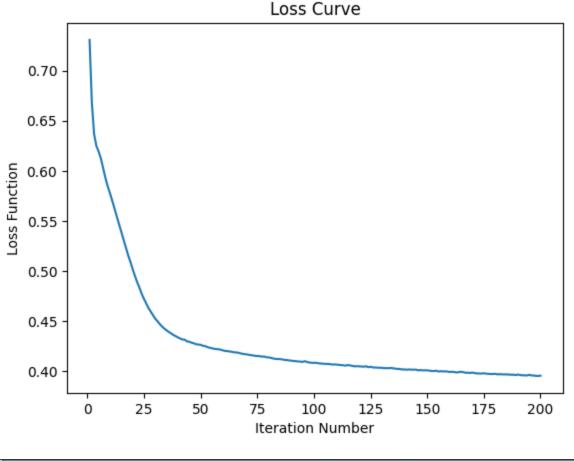


Multilayer perceptron AUC score: 0.8009685802031656

```
import matplotlib.pyplot as plt
iterations = [i for i in range(1,mlp.best_estimator_.n_iter_+1)]
loss_curve = mlp.best_estimator_.loss_curve_

plt.plot(iterations,loss_curve)
plt.xlabel("Iteration Number")
plt.ylabel("Loss Function")
plt.title("Loss Curve")
```

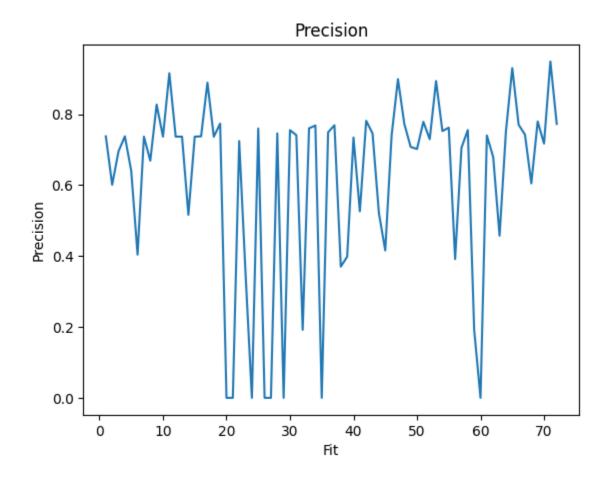
This is a function to visualize the loss curve of our best estimator(model)



These are the properties of the best estimator as per the parameters in parameter grid, and are the **optimal hyperparameters** 

Now we will visualize the fit against accuracy metric (precision, recall, f1-score, and roc/auc)

```
fits = [fit for fit in range(1,73)]
accuracy = mlp.cv_results_["mean_test_precision"][::10]
plt.plot(fits,accuracy)
plt.xlabel("Fit")
plt.ylabel("Precision")
plt.title("Precision")
```



The code is similar for all other accuracy change metrics, note that many of the fits have been removed for better visualization of the accuracy change

