

Getting Started

In this lab, we'll learn how to use boosting algorithms to make classifications on the Pima Indians Dataset. You will find the data stored in the file 'pima-indians-diabetes.csv'. Our goal is to use boosting algorithms to determine whether a person has diabetes. Let's get started!

We'll begin by importing everything we need for this lab. Run cell below:

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
%matplotlib inline
from sklearn.model_selection import train_test_split, cross_val_score
from sklearn.ensemble import AdaBoostClassifier, GradientBoostingClassifier
from sklearn.metrics import accuracy_score, f1_score, confusion_matrix, classificati
```

Now, use Pandas to import the data stored in 'pima-indians-diabetes.csv' and store it in a DataFrame. Print the first five rows to inspect the data we've imported and ensure everything loaded correctly.

```
# Import the data
df = pd.read_csv('pima-indians-diabetes.csv')

# Print the first five rows
df.head()

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```

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| | Pregnancies | Glucose | BloodPressure | SkinThickness | Insulin | BMI |
|---|-------------|---------|---------------|---------------|---------|------|
| 0 | 6 | 148 | 72 | 35 | 0 | 33.6 |

| | Pregnancies | Glucose | BloodPressure | SkinThickness | Insulin | вмі |
|---|-------------|---------|---------------|---------------|---------|------|
| 1 | 1 | 85 | 66 | 29 | 0 | 26.6 |
| 2 | 8 | 183 | 64 | 0 | 0 | 23.3 |
| 3 | 1 | 89 | 66 | 23 | 94 | 28.1 |
| 4 | 0 | 137 | 40 | 35 | 168 | 43.1 |
| | | | | | | |

Cleaning, exploration, and preprocessing

The target we're trying to predict is the 'Outcome' column. A 1 denotes a patient with diabetes.

By now, you're quite familiar with exploring and preprocessing a dataset.

In the following cells:

- Check for missing values and deal with them as you see fit (if any exist)
- Count the number of patients with and without diabetes in this dataset
- Store the target column in a separate variable and remove it from the dataset
- Split the dataset into training and test sets, with a test_size of 0.25 and a random state of 42

```
# Check for missing values
df.isna().sum()
```

| Pregnancies | 0 | |
|--------------------------|---|--|
| Glucose | 0 | |
| BloodPressure | 0 | |
| SkinThickness | 0 | |
| Insulin | 0 | |
| BMI | 0 | |
| DiabetesPedigreeFunction | | |
| Age | 0 | |
| Outcome | 0 | |
| dtype: int64 | | |

Number of patients with and without diabetes
df['Outcome'].value_counts()

```
0 500
1 268
Name: Outcome, dtype: int64

target = df['Outcome']
df = df.drop('Outcome', axis=1)

# Split the data into training and test sets
X_train, X_test, y_train, y_test = train_test_split(df, target, test_size=0.25, rand)
```

Train the models

Now that we've explored the dataset, we're ready to fit some models!

In the cell below:

- Instantiate an AdaBoostClassifier (set the random_state for 42)
- Instantiate a GradientBoostingClassifer (set the random_state for 42)

```
# Instantiate an AdaBoostClassifier
adaboost_clf = AdaBoostClassifier(random_state=42)
# Instantiate an GradientBoostingClassifier
gbt_clf = GradientBoostingClassifier(random_state=42)
```

Now, fit the training data to both the classifiers:

learning_rate=0.1, loss='deviance', max_depth=3,

GradientBoostingClassifier(criterion='friedman_mse', init=None,

```
max_features=None, max_leaf_nodes=None,
min_impurity_decrease=0.0, min_impurity_split=None,
min_samples_leaf=1, min_samples_split=2,
min_weight_fraction_leaf=0.0, n_estimators=100,
n_iter_no_change=None, presort='auto',
random_state=42, subsample=1.0, tol=0.0001,
validation_fraction=0.1, verbose=0,
warm start=False)
```

Now, let's use these models to predict labels on both the training and test sets:

```
# AdaBoost model predictions
adaboost_train_preds = adaboost_clf.predict(X_train)
adaboost_test_preds = adaboost_clf.predict(X_test)

# GradientBoosting model predictions
gbt_clf_train_preds = gbt_clf.predict(X_train)
gbt_clf_test_preds = gbt_clf.predict(X_test)
```

Now, complete the following function and use it to calculate the accuracy and f1-score for each model:

```
def display_acc_and_f1_score(true, preds, model_name):
    acc = accuracy_score(true, preds)
    f1 = f1_score(true, preds)
    print("Model: {}".format(model_name))
    print("Accuracy: {}".format(acc))
    print("F1-Score: {}".format(f1))

print("Training Metrics")
display_acc_and_f1_score(y_train, adaboost_train_preds, model_name='AdaBoost')
print("")
display_acc_and_f1_score(y_train, gbt_clf_train_preds, model_name='Gradient Boosted
print("")
print("Testing Metrics")
display_acc_and_f1_score(y_test, adaboost_test_preds, model_name='AdaBoost')
print("")
display_acc_and_f1_score(y_test, adaboost_test_preds, model_name='Gradient Boosted Train_preds, model_name='Gradient Boosted Train_pre
```

Training Metrics Model: AdaBoost

Accuracy: 0.83506944444444444 F1-Score: 0.7493403693931399 Model: Gradient Boosted Trees Accuracy: 0.940972222222222 F1-Score: 0.9105263157894736

Testing Metrics Model: AdaBoost

Accuracy: 0.72395833333333334 F1-Score: 0.618705035971223

Model: Gradient Boosted Trees

Accuracy: 0.75

Let's go one step further and create a confusion matrix and classification report for each. Do so in the cell below:

```
adaboost_confusion_matrix = confusion_matrix(y_test, adaboost_test_preds)
adaboost_confusion_matrix
```

```
array([[96, 27],
[26, 43]])
```

```
gbt_confusion_matrix = confusion_matrix(y_test, gbt_clf_test_preds)
gbt_confusion_matrix
```

```
array([[96, 27],
[21, 48]])
```

adaboost_classification_report = classification_report(y_test, adaboost_test_preds)
print(adaboost_classification_report)

| | precision | recall | f1-score | support |
|--------------|-----------|--------|----------|---------|
| 0 | 0.79 | 0.78 | 0.78 | 123 |
| 1 | 0.61 | 0.62 | 0.62 | 69 |
| accuracy | | | 0.72 | 192 |
| macro avg | 0.70 | 0.70 | 0.70 | 192 |
| weighted avg | 0.72 | 0.72 | 0.72 | 192 |

```
gbt_classification_report = classification_report(y_test, gbt_clf_test_preds)
print(gbt classification report)
```

| | precision | recall | f1-score | support |
|--------------|-----------|--------|----------|---------|
| | | | | |
| 0 | 0.82 | 0.78 | 0.80 | 123 |
| 1 | 0.64 | 0.70 | 0.67 | 69 |
| | | | | |
| accuracy | | | 0.75 | 192 |
| macro avg | 0.73 | 0.74 | 0.73 | 192 |
| weighted avg | 0.76 | 0.75 | 0.75 | 192 |

Question: How did the models perform? Interpret the evaluation metrics above to answer this question.

Write your answer below this line:

As a final performance check, let's calculate the 5-fold cross-validated score for each model!

Recall that to compute the cross-validation score, we need to pass in:

- A classifier
- All training data
- All labels
- The number of folds we want in our cross-validation score

Since we're computing cross-validation score, we'll want to pass in the entire dataset, as well as all of the labels.

In the cells below, compute the mean cross validation score for each model.

```
print('Mean Adaboost Cross-Val Score (k=5):')
print(cross_val_score(adaboost_clf, df, target, cv=5).mean())
# Expected Output: 0.7631270690094218

Mean Adaboost Cross-Val Score (k=5):
0.7631270690094218
```

```
print('Mean GBT Cross-Val Score (k=5):')
print(cross_val_score(gbt_clf, df, target, cv=5).mean())
# Expected Output: 0.7591715474068416

Mean GBT Cross-Val Score (k=5):
0.7591715474068416
```

These models didn't do poorly, but we could probably do a bit better by tuning some of the important parameters such as the *Learning Rate*.

Summary

In this lab, we learned how to use scikit-learn's implementations of popular boosting algorithms such as AdaBoost and Gradient Boosted Trees to make classification predictions on a real-world dataset!

Releases

No releases published

Packages

No packages published

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Languages

• Jupyter Notebook 100.0%