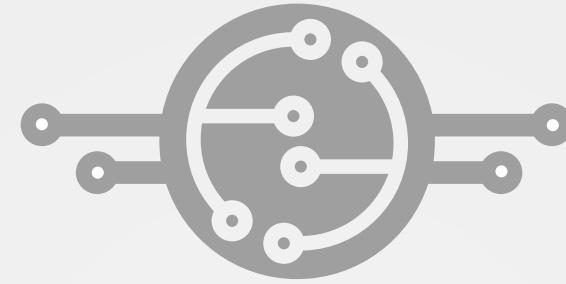


March 2024



Artificial intelligence

ML Challenge Thomas More
2023-2024

Team 6 - German Novykov, Pidnebesnyi Alex, Mark Vaikul

The challenge

Heart Disease predictions AI

In this challenge we have to design an AI model that will predict heart disease based on symptoms. Ideal use case for this system might be for improving doctors diagnosis.

We were given a training dataset and test dataset. We used three different models to train and then perform predictions. Our result was 91% accuracy on Kaggle.





Models used

Decision Tree

Tree-like graph of decisions and their possible consequences, including chance event outcomes, resource costs, and utility, to arrive at a prediction or decision.

Random forest

Combines the output of multiple decision trees to reach a single result

Linear Regression

Models the relationship between a dependent variable and one or more independent variables by fitting a linear equation to observed data

Machine learning steps

Processing Data

Transform categorical data (gender, location of pain) into numerical. We used OrdinalEncoder for that. For each unique string this encoder creates a number.

Training

Split the testing data into two datasets - training and testing. We use train split to evaluate our models performance and in effort to not overfit the model.

Testing

Let the model predict the heart disease and save it in the file





Processing Data

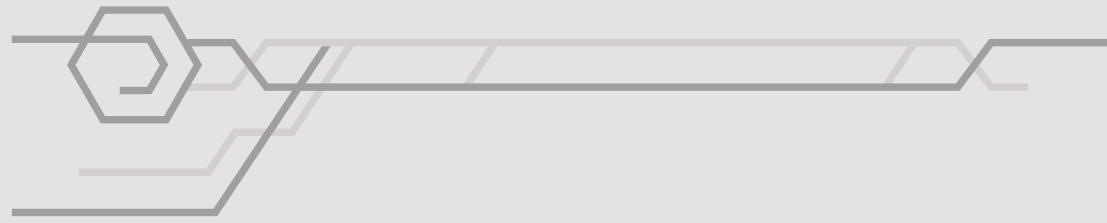
Changing faulty data

```
# Filter values for Cholesterol and RestingBP
chol_median = train_data.loc[train_data['Cholesterol'] != 0, 'Cholesterol'].median()
train_data['Cholesterol'].replace(0, chol_median, inplace=True)

resting_median = train_data.loc[train_data['RestingBP'] != 0, 'RestingBP'].median()
train_data['RestingBP'].replace(0, resting_median, inplace=True)
```

Faulty data decrease purity of the model. 0 values for Cholesterol and RestingBP may indicate wrongly inserted data





Processing Data

Adding interaction terms

```
train_data['Age_Chol_Interact'] = train_data['Age'] * train_data['Cholesterol']
```

✓ 0.0s

Combining two columns together can
increase model accuracy

Processing Data

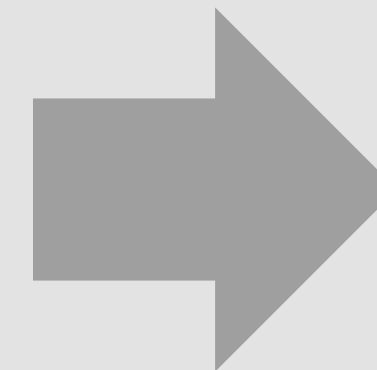
Encoding categorical features

```
categorical_columns = train_data.select_dtypes(include=['object']).columns

ordinal_encoder = ce.OrdinalEncoder(cols=categorical_columns)
features_encoded = ordinal_encoder.fit_transform(train_data.drop(['id', 'HeartDisease'], axis=1))
```

Model cannot recognise strings and objects to train on. OrdinalEncoder transforms every string into unique number.

ChestPainType
ASY
ASY
ASY
ATA
ASY



ChestPainType
1
1
1
2
1
...
2
1
3
4
1



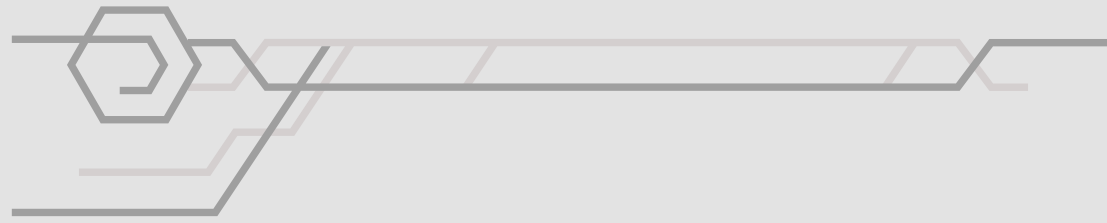
Processing Data

Scaling

```
standard_scaler = StandardScaler()  
features_scaled = standard_scaler.fit_transform(features_encoded)
```

Transforms features to have a mean of 0 and standard deviation of 1. This makes models to converge faster and perform better by treating all features equally in scale.

```
[[ 0.13874196 -0.51601569 -0.80799877 ... 0.1749016 -0.93481405  
 -0.48252039]  
 [ 1.4265348 -0.51601569 -0.80799877 ... 0.5565051 -0.93481405  
 1.71778046]  
 [ 0.24605803 -0.51601569 -0.80799877 ... -1.82851674 0.11623782  
 0.05494922]  
 ...  
 [-0.39783839 1.93792558 1.27527517 ... -0.87450801 -0.93481405  
 0.3984865 ]  
 [ 1.21190266 -0.51601569 2.31691214 ... 0.46110422 -0.93481405  
 1.48727803]  
 [ 1.21190266 -0.51601569 -0.80799877 ... 0.1749016 -0.93481405  
 0.64090192]]
```

Processing Data

Polynomial Features

```
poly_features_transformer = PolynomialFeatures(degree=2, include_bias=False)
features_poly = poly_features_transformer.fit_transform(features_scaled)
```

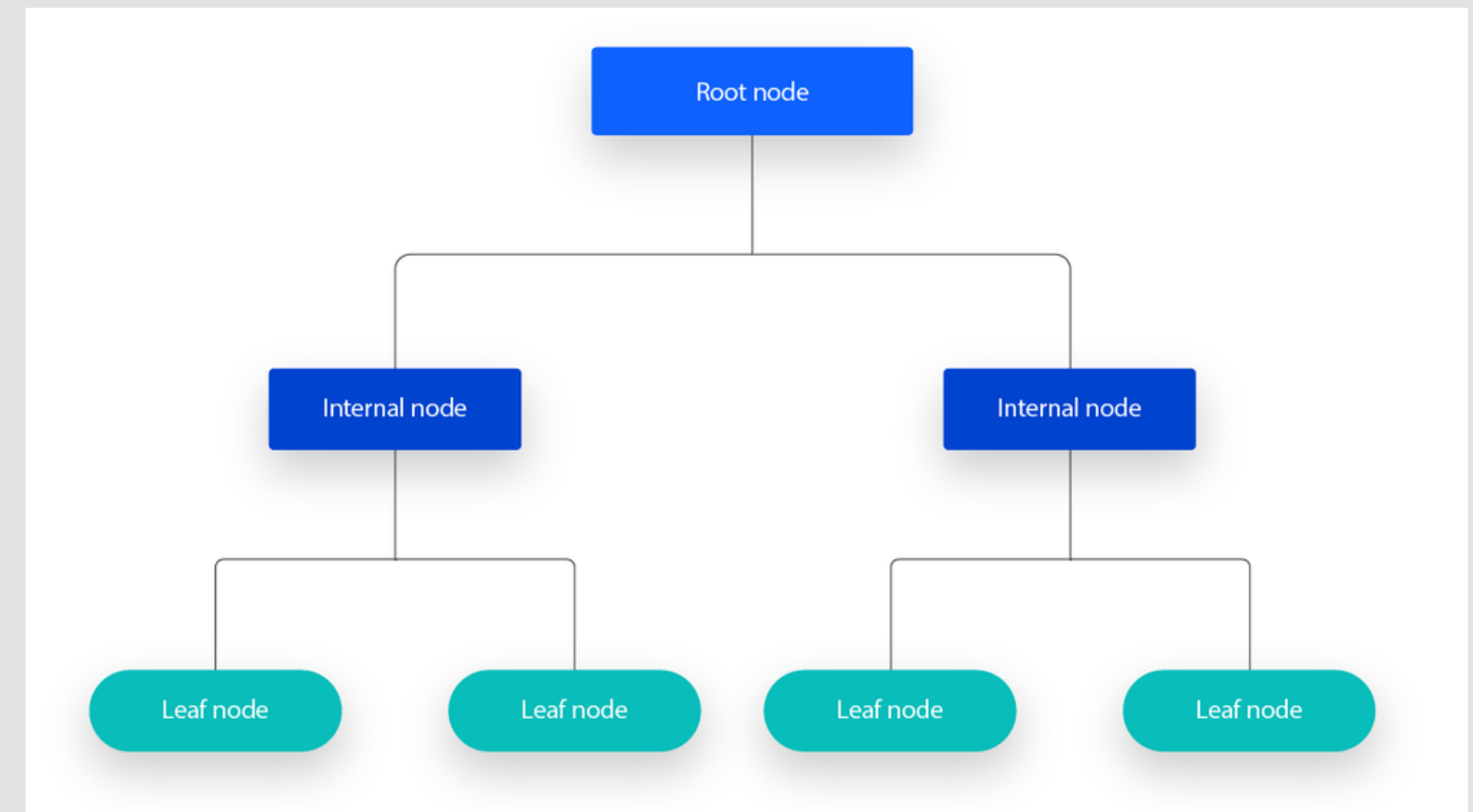
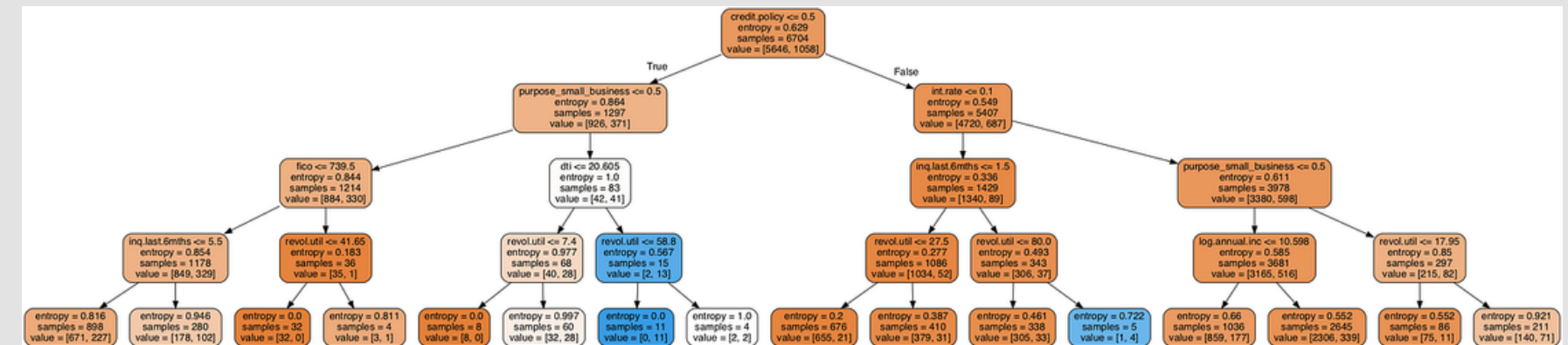
For each feature makes a copy with values raised to a degree.

```
[[ 0.13874196 -0.51601569 -0.80799877 ... 0.8738773 0.45106684
  0.23282593]
 [ 1.4265348 -0.51601569 -0.80799877 ... 0.8738773 -1.6058053
  2.95076971]
 [ 0.24605803 -0.51601569 -0.80799877 ... 0.01351123 0.00638718
  0.00301942]
 ...
 [-0.39783839 1.93792558 1.27527517 ... 0.8738773 -0.37251078
  0.15879149]
 [ 1.21190266 -0.51601569 2.31691214 ... 0.8738773 -1.39032839
  2.21199594]
 [ 1.21190266 -0.51601569 -0.80799877 ... 0.8738773 -0.59912411
  0.41075526]]
```

Decision Tree

Decision Tree

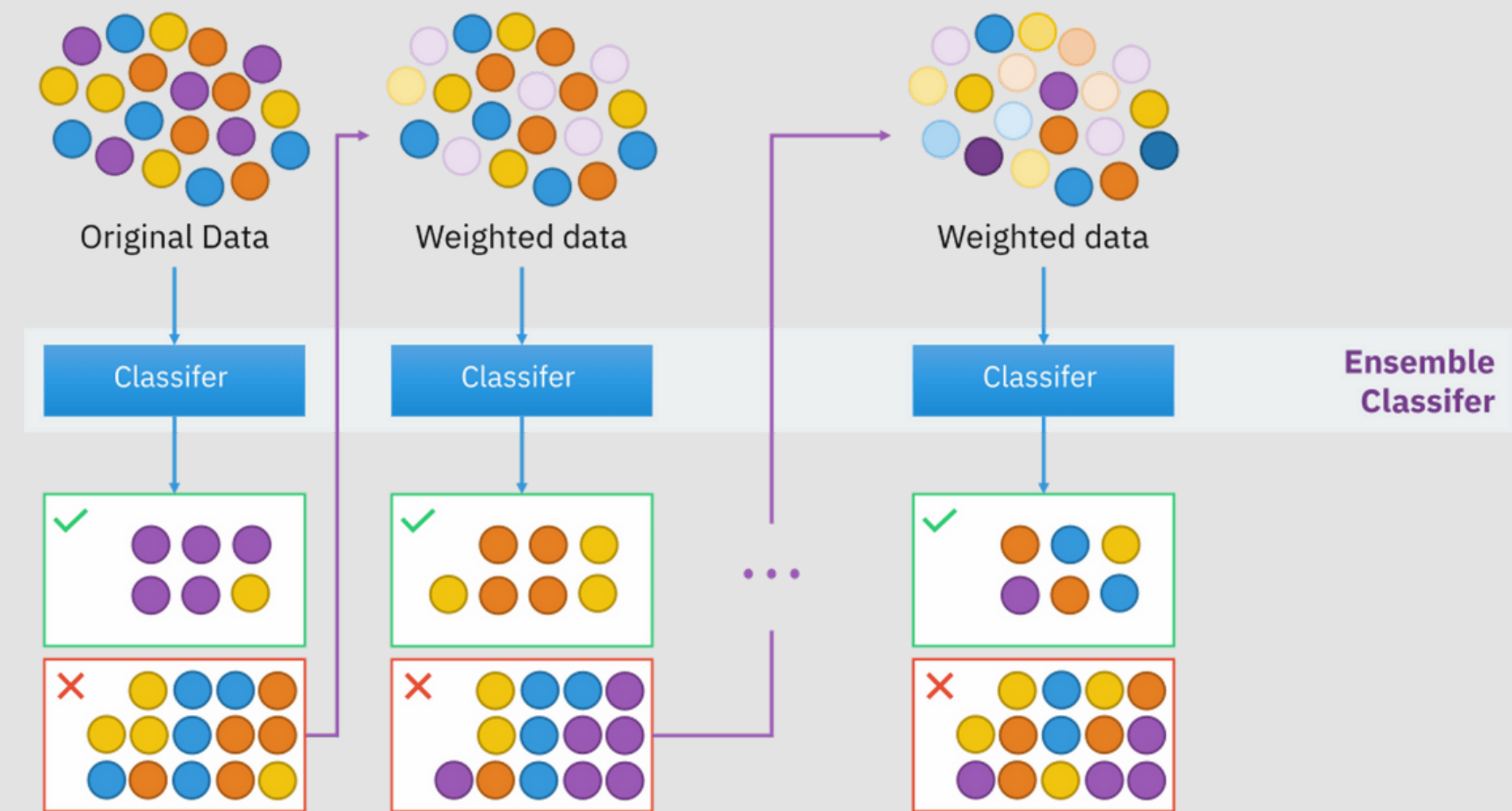
Tree structure, which consists of a root node, branches, internal nodes and leaf nodes. Starts with a root node, which does not have any incoming branches. The outgoing branches from the root node then feed into the internal nodes, also known as decision nodes

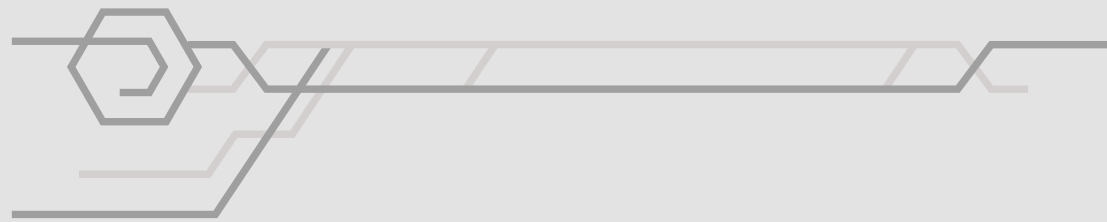


Adaboost

Adaptive Boost

Adaboost works by weighting incorrectly classified instances more heavily so that the subsequent weak learners focus more on the difficult cases.





Decision Tree

Performance

No hyperparameter tuning

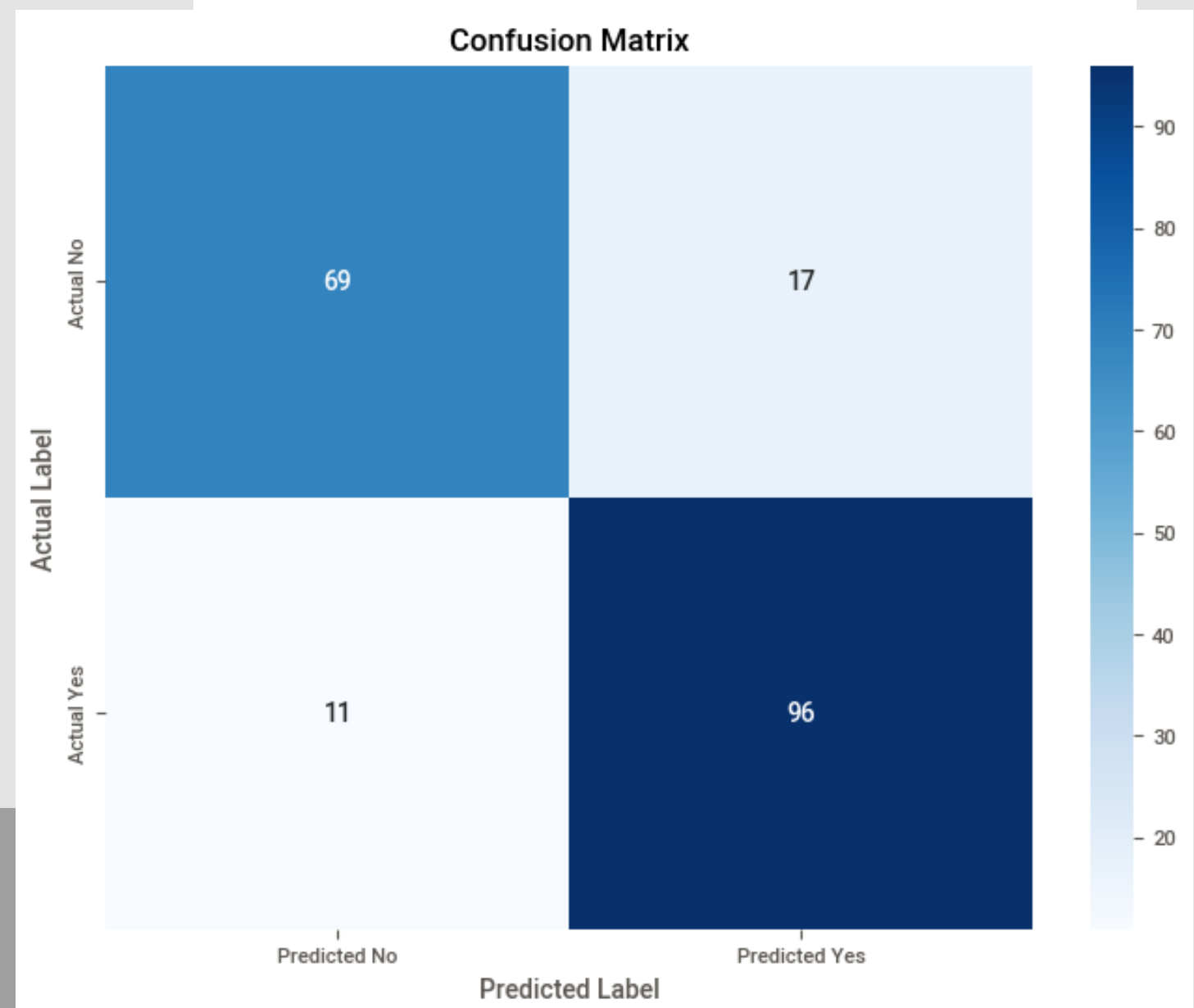
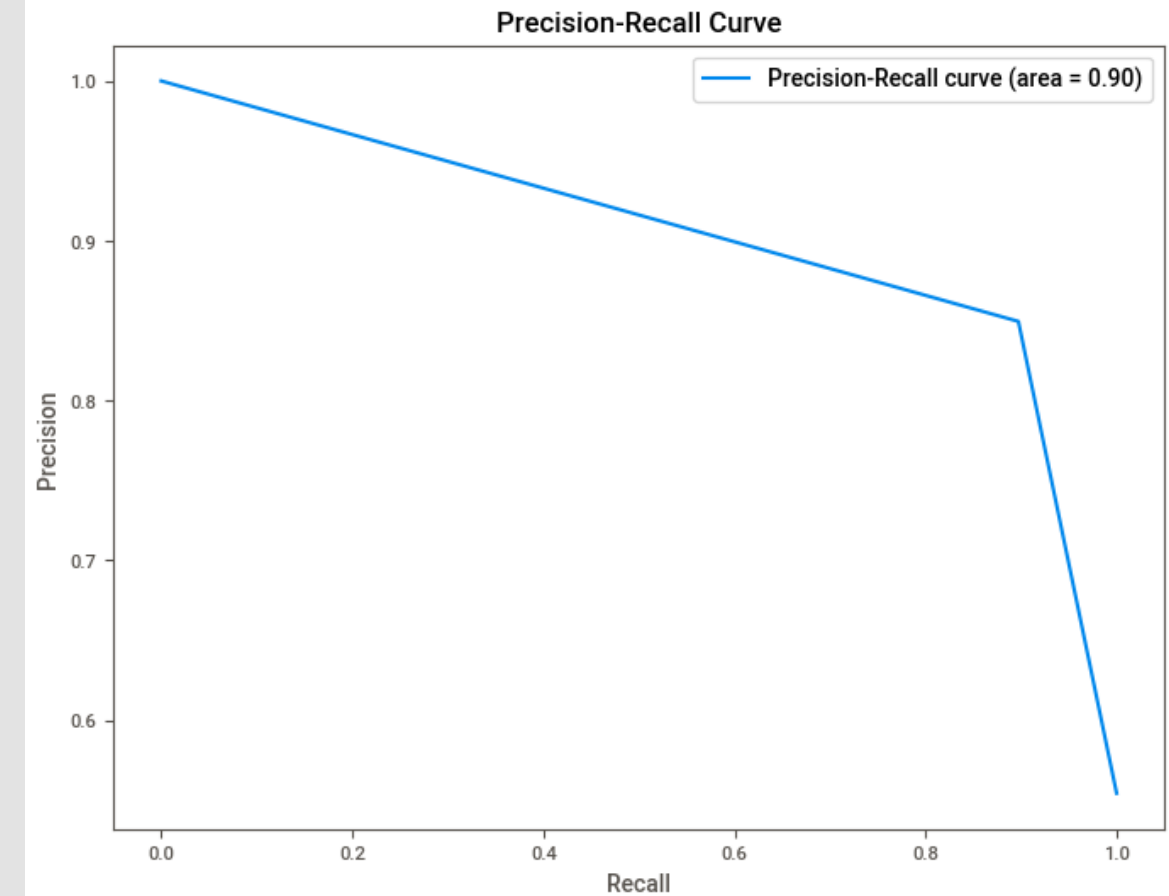
Decision Tree Accuracy: 0.7616580310880829

Hyperparameter tuning with GridSearch

Decision Tree Accuracy: 0.8549222797927462

With Adaboost

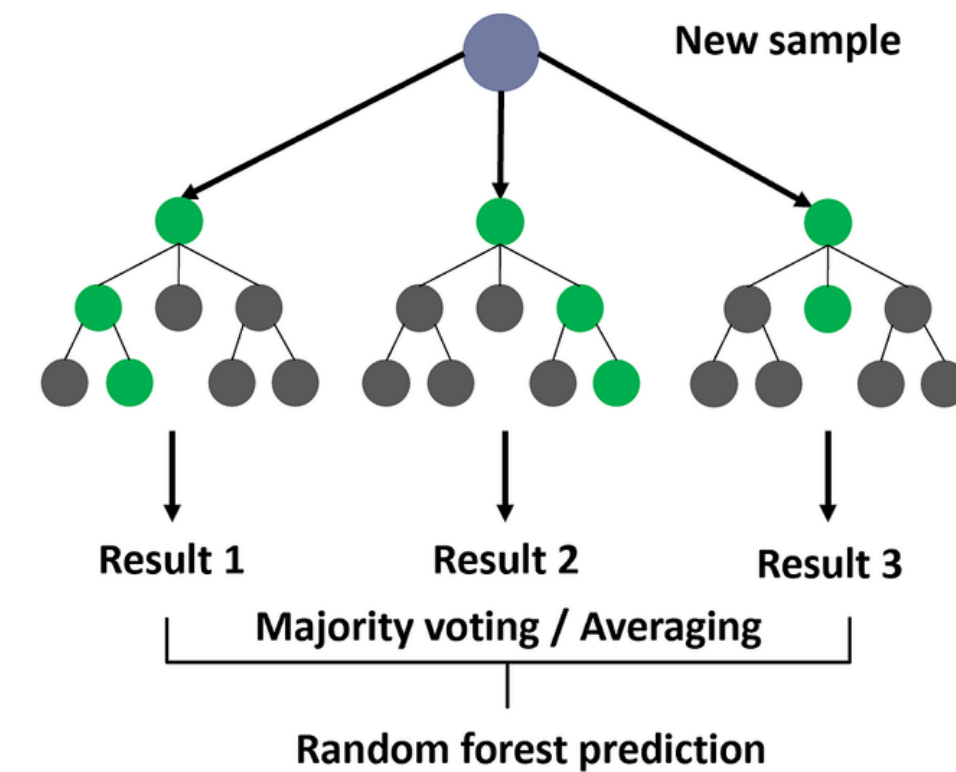
AdaBoost Decision Tree Accuracy: 0.8549222797927462

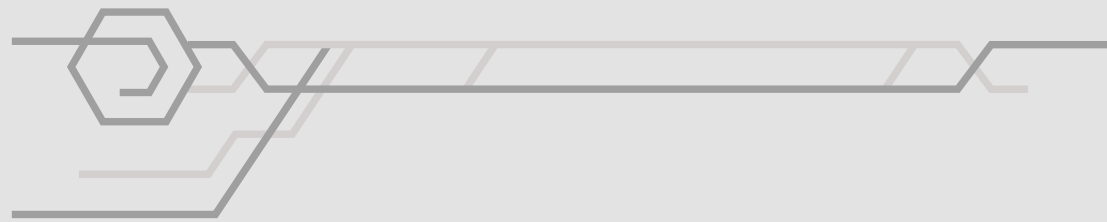


Random Forest

Random forest

The random forest algorithm is an extension of the bagging method as it utilizes both bagging and feature randomness to create an uncorrelated forest of decision trees, thus limiting the risk of overfitting



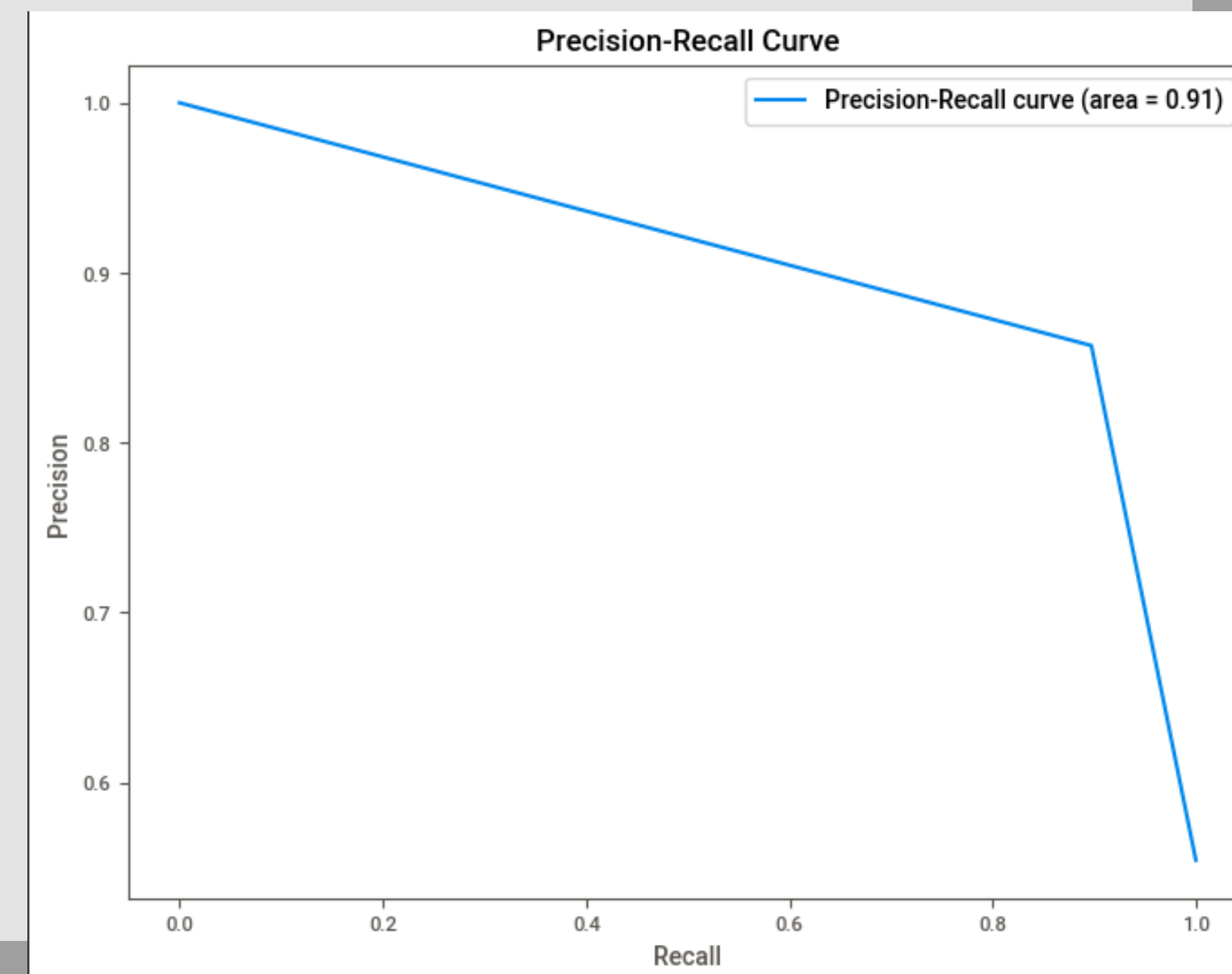
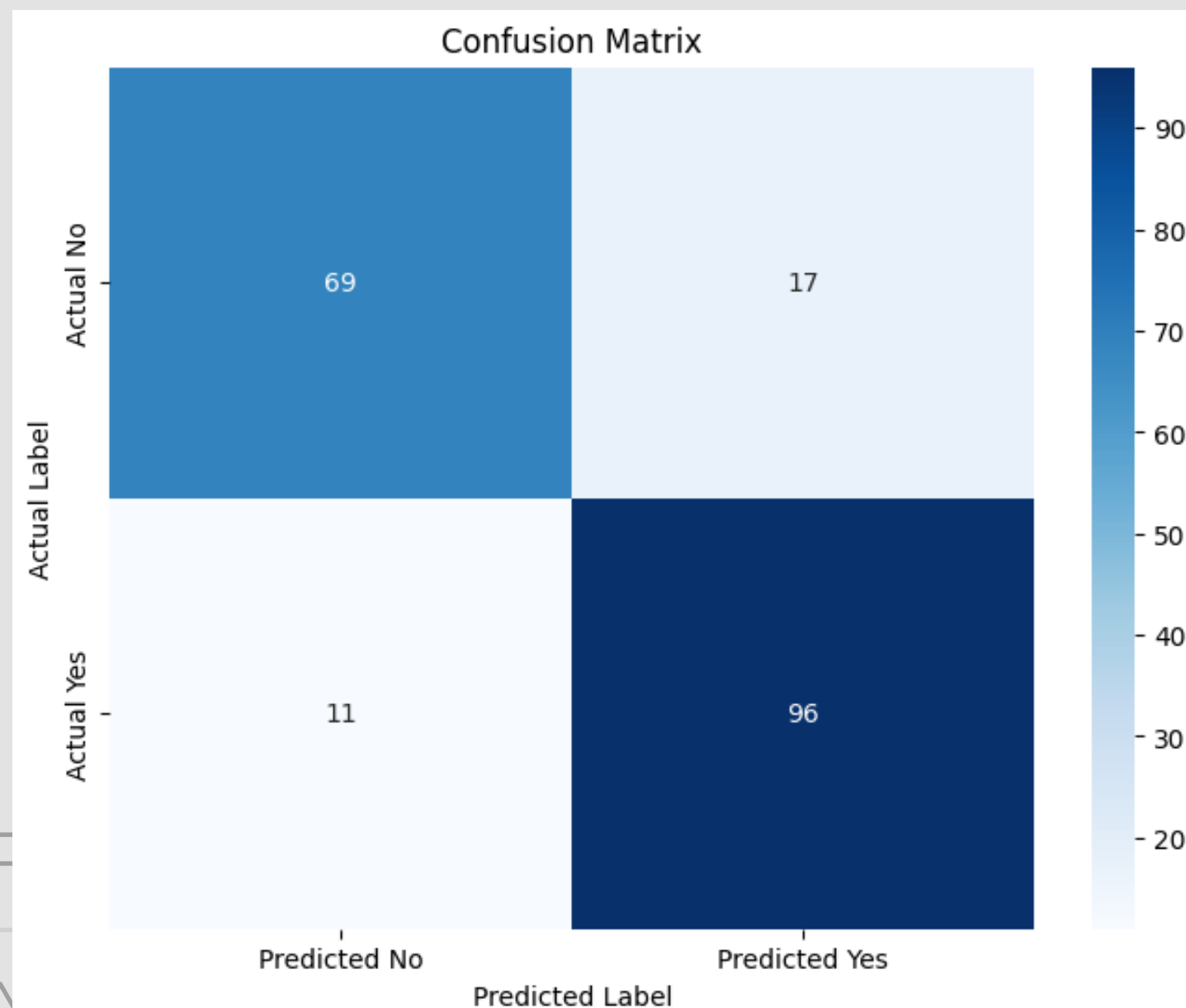


Random Forest

Performance

Random Forest accuracy

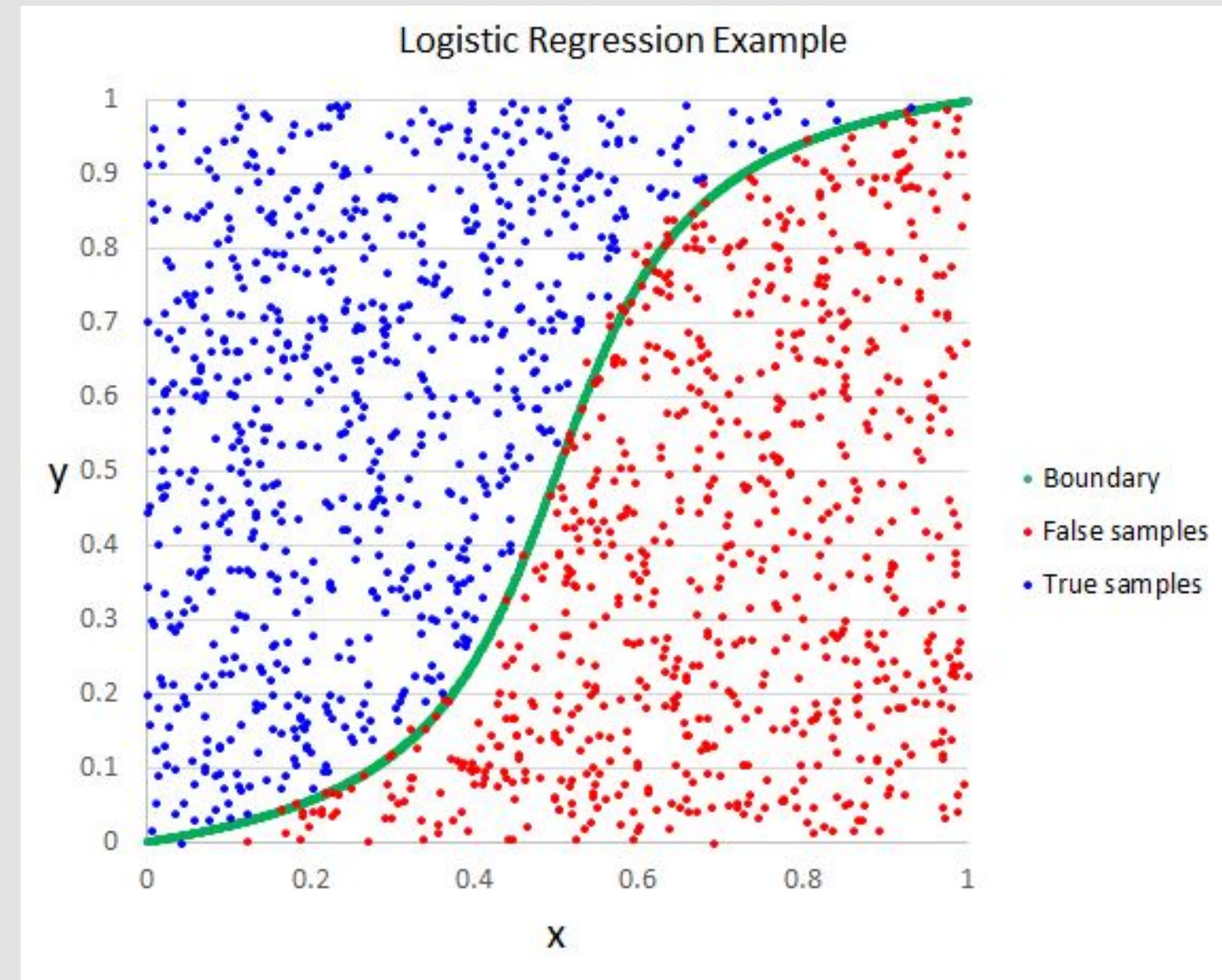
Random Forest Accuracy: 0.8653



Logistic Regression

Logistic Regression

Logistic regression is a Machine Learning classification algorithm that is used to predict the probability of certain classes based on some dependent variables.

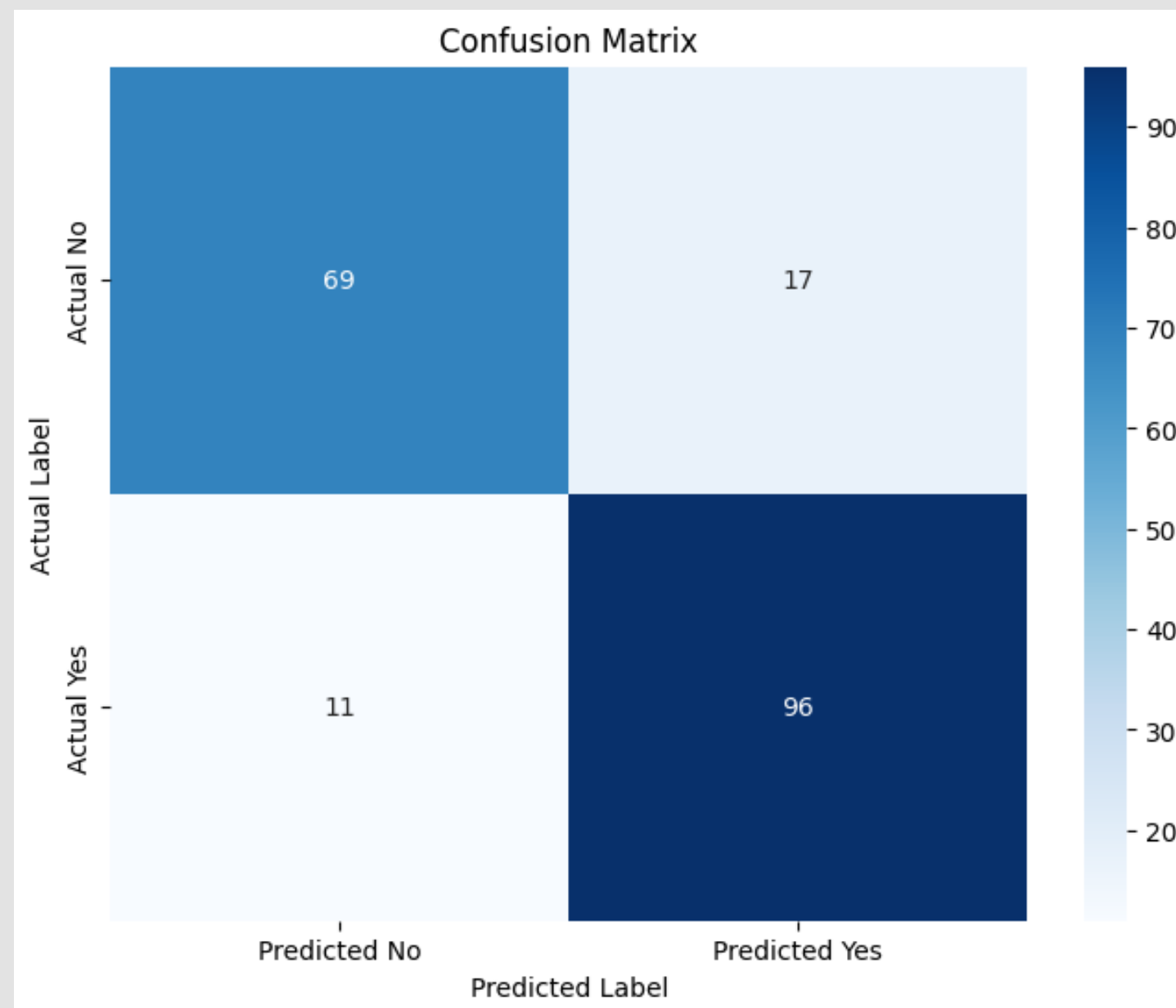




Logistic Regression

Performance

Logistic Regression Accuracy: 0.8549



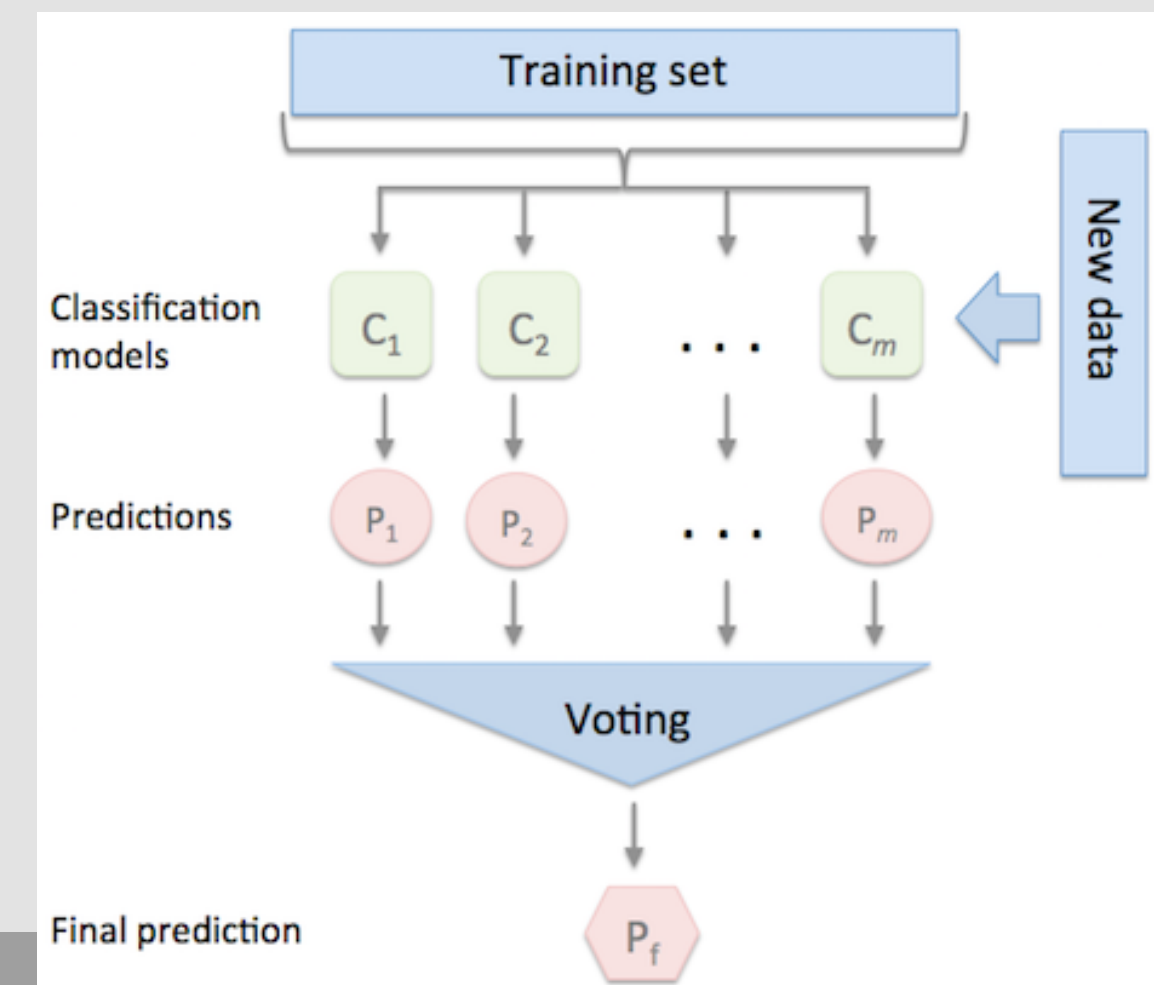
Combining the models

Voting Classifier

```
# Stacking Ensemble Setup
stacking_ensemble_classifier = VotingClassifier(estimators=base_classifiers, voting="hard", n_jobs=-1)

# Train Stacking Ensemble
stacking_ensemble_classifier.fit(features_train, target_train)
```

The next step is combining three models into one by using **Voting Classifier**. Every model makes a prediction and by counting, the prediction with most votes wins. This improves consistency, as if one model fails to work with data, other models will override the faulty prediction.



Stacking Ensemble Model Accuracy: 0.844559585492228



ChatGPT prompts

Code generated

Prompt:



You

in my dataset i have anomalyous data, like when colesterol equals to zero, which is impossible, how can i get rid of the outliers?

Answer:

Removing Anomalies Based on Domain Knowledge

You can remove rows where cholesterol equals zero (or any other impossible values) by using a conditional statement that filters your DataFrame. Here's how you can do it:

python

Copy code

```
# Assuming 'cholesterol' is the column name in your DataFrame
# Remove rows where cholesterol equals zero
hearts_cleaned = hearts[hearts['cholesterol'] != 0]

# Continue with your preprocessing on hearts_cleaned
```



ChatGPT prompts

Code generated

Answer:

Prompt:

 You



train_heart.csv
Spreadsheet


make a best pandas sklearn random forest model that you can do for this data (you can manipulate the data too in any way)

Same code that we
already have, no
improvements

Analysis



python

 Copy code

```
from sklearn.model_selection import train_test_split
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import accuracy_score, classification_report

# Splitting the data into features and target variable
X = df_encoded.drop('HeartDisease', axis=1)
y = df_encoded['HeartDisease']

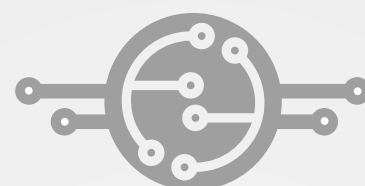
# Splitting the data into training and test sets
X_train, X_test, y_train, y_test = train_test_split(X, y,

# Initialize the Random Forest model
rf = RandomForestClassifier(n_estimators=100, random_state=

# Train the model
rf.fit(X_train, y_train)

# Predictions
y_pred = rf.predict(X_test)

# Evaluation
accuracy = accuracy_score(y_test, y_pred)
report = classification_report(y_test, y_pred)
```

THANKYOU

