## Phase 1

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
from sklearn.datasets import make_classification
from sklearn.model_selection import train_test_split
from sklearn.ensemble import
BaggingClassifier,RandomForestClassifier,AdaBoostClassifier
from sklearn.tree import DecisionTreeClassifier
from sklearn.metrics import accuracy_score,precision_score,
recall_score, fl_score, classification_report
from sklearn import tree
```

# Bagging

```
# Define 9 different colors
colors = ['b', 'g', 'r', 'c', 'm', 'y', 'k', 'purple', 'orange']
# Read data from CSV file
df = pd.read csv('/content/drive/My Drive/1/Dataset1.csv')
X = df.drop('Label', axis=1)
y = df['Label']
# Scatter plot of Feature 1 vs Feature 2
plt.scatter(df['Feature 1'], df['Feature 2'], c=y,
cmap=plt.cm.get cmap('rainbow', len(np.unique(y))))
plt.title('Scatter Plot of Feature1 vs Feature2')
plt.xlabel('Feature1')
plt.ylabel('Feature2')
plt.grid(True)
plt.colorbar(ticks=range(len(np.unique(y))), label='Label')
plt.show()
# Split the data into training and testing sets
X train, X test, y train, y test = train test split(X, y,
test size=0.2, random state=42)
# Number of estimators to evaluate
num estimators = [1, 2, 5, 10, 50, 100]
train accuracies = []
test accuracies = []
# Loop to test different numbers of estimators
for n in num estimators:
    # Create Bagging model with different numbers of estimators
```

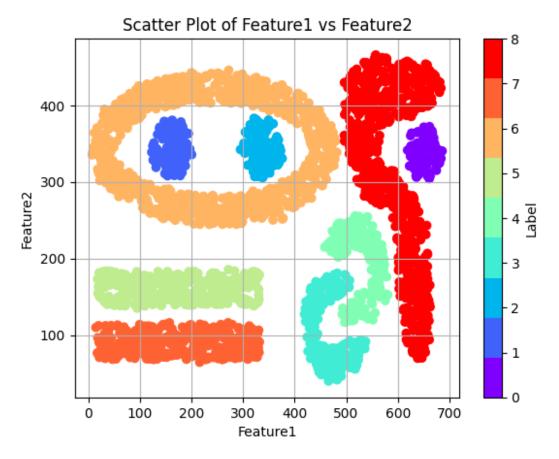
```
bagging clf =
BaggingClassifier(base estimator=DecisionTreeClassifier(max depth=4,
splitter='random'),
                                    n estimators=n,
                                    random state=42)
    # Train the model on training data
    bagging clf.fit(X train, y train)
    # Predict labels for training and testing data
    y_train_pred = bagging_clf.predict(X_train)
    y test pred = bagging clf.predict(X test)
    # Compute model accuracy
    train_accuracy = accuracy_score(y_train, y_train_pred)
    test accuracy = accuracy_score(y_test, y_test_pred)
    # Store accuracies in lists
    train accuracies.append(train accuracy)
    test accuracies.append(test accuracy)
    # Evaluate the model for training and testing data
    train_f1 = f1_score(y_train, y_train_pred, average='micro')
    test f1 = f1 score(y test, y test pred, average='micro')
    train precision = precision_score(y_train, y_train_pred,
average='micro')
    test precision = precision score(y test, y test pred,
average='micro')
    train recall = recall score(y train, y train pred,
average='micro')
    test recall = recall score(y test, y test pred, average='micro')
    print(f'n estimators Bagging = {n}:')
    print(f'Train Accuracy: {train accuracy:.2f}, Test Accuracy:
{test accuracy:.2f}')
    print(f'Train Precision: {train precision:.2f}, Test Precision:
{test precision:.2f}')
    print(f'Train Recall: {train recall:.2f}, Test Recall:
{test recall:.2f}')
    print(f'Train F1-score: {train f1:.2f}, Test F1-score:
{test f1:.2f}')
    print()
# Plot Bagging accuracy versus number of estimators
plt.figure(figsize=(10, 6))
plt.plot(num estimators, train accuracies, marker='o', label='Train
Accuracy')
plt.plot(num estimators, test accuracies, marker='o', label='Test
Accuracy')
```

```
plt.title('Bagging Accuracy vs. Number of Estimators')
plt.xlabel('Number of Estimators')
plt.ylabel('Accuracy')
plt.xticks(num estimators)
plt.legend()
plt.grid(True)
plt.tight layout()
plt.show()
# Display five random trees from the model
plt.figure(figsize=(20, 10))
for i in range(5):
    plt.subplot(2, 3, i + 1)
    # Plot the tree
    plot tree(bagging clf.estimators [i], filled=True) # Use
plot tree to plot the tree
    plt.title(f'Tree Bagging {i + 1}')
plt.tight layout()
plt.show()
# Plot decision boundaries
plt.figure(figsize=(15, 10))
# Create a mesh grid
x_{min}, x_{max} = X['Feature 1'].min() - 1, <math>X['Feature 1'].max() + 1
y_{min}, y_{max} = X['Feature 2'].min() - 1, <math>X['Feature 2'].max() + 1
xx, yy = np.meshgrid(np.arange(x min, x max, 0.1),
                     np.arange(y min, y max, 0.1))
for i in range(5):
    plt.subplot(2, 3, i + 1)
    # Predict the class for each point in the mesh grid
    Z = bagging clf.estimators [i].predict(np.c [xx.ravel(),
yy.ravel()])
    Z = Z.reshape(xx.shape)
    # Plot decision boundaries
    plt.contourf(xx, yy, Z, alpha=0.4, cmap=plt.cm.get cmap('rainbow',
len(np.unique(y))))
    # Plot original data points with predicted labels
    y pred tree = bagging clf.estimators [i].predict(X)
    for label in np.unique(y):
        plt.scatter(X[y_pred_tree == label]['Feature 1'],
X[y_pred_tree == label]['Feature 2'],
                    color=colors[label], label=f'Label {label}')
```

```
plt.title(f'Tree Bagging {i + 1}')
plt.legend()

plt.tight_layout()
plt.show()

<ipython-input-54-1fa8d6f22dc9>:11: MatplotlibDeprecationWarning: The
get_cmap function was deprecated in Matplotlib 3.7 and will be removed
two minor releases later. Use ``matplotlib.colormaps[name]`` or
``matplotlib.colormaps.get_cmap(obj)`` instead.
   plt.scatter(df['Feature 1'], df['Feature 2'], c=y,
cmap=plt.cm.get_cmap('rainbow', len(np.unique(y))))
```

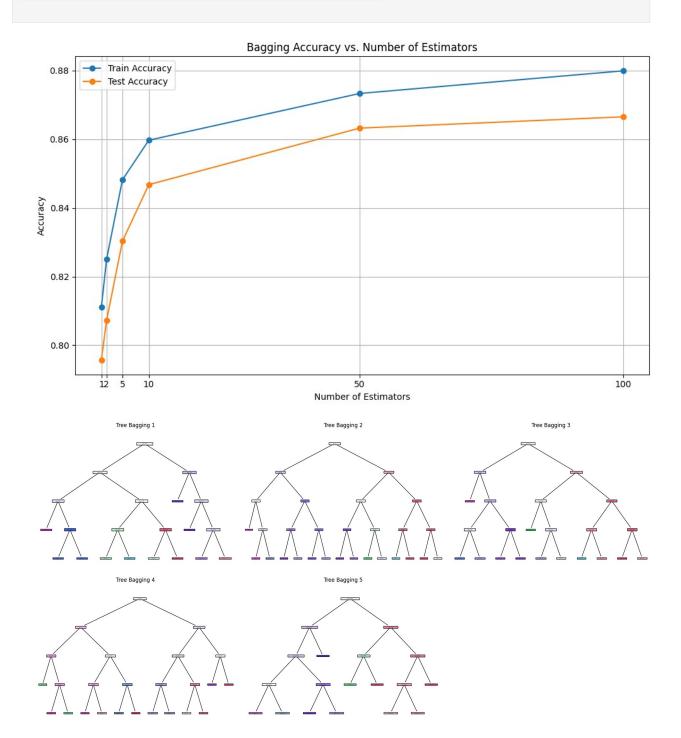


```
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/_base.py:166:
FutureWarning: `base_estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
   warnings.warn(
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/_base.py:166:
FutureWarning: `base_estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
   warnings.warn(
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/_base.py:166:
```

```
FutureWarning: `base estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
  warnings.warn(
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/ base.py:166:
FutureWarning: `base estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
  warnings.warn(
n estimators Bagging = 1:
Train Accuracy: 0.81, Test Accuracy: 0.80
Train Precision: 0.81, Test Precision: 0.80
Train Recall: 0.81, Test Recall: 0.80
Train F1-score: 0.81, Test F1-score: 0.80
n estimators Bagging = 2:
Train Accuracy: 0.83, Test Accuracy: 0.81
Train Precision: 0.83, Test Precision: 0.81
Train Recall: 0.83, Test Recall: 0.81
Train F1-score: 0.83, Test F1-score: 0.81
n estimators Bagging = 5:
Train Accuracy: 0.85, Test Accuracy: 0.83
Train Precision: 0.85, Test Precision: 0.83
Train Recall: 0.85, Test Recall: 0.83
Train F1-score: 0.85, Test F1-score: 0.83
n estimators Bagging = 10:
Train Accuracy: 0.86, Test Accuracy: 0.85
Train Precision: 0.86, Test Precision: 0.85
Train Recall: 0.86, Test Recall: 0.85
Train F1-score: 0.86, Test F1-score: 0.85
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/ base.py:166:
FutureWarning: `base estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
 warnings.warn(
n estimators Bagging = 50:
Train Accuracy: 0.87, Test Accuracy: 0.86
Train Precision: 0.87, Test Precision: 0.86
Train Recall: 0.87, Test Recall: 0.86
Train F1-score: 0.87, Test F1-score: 0.86
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/ base.py:166:
FutureWarning: `base estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
  warnings.warn(
```

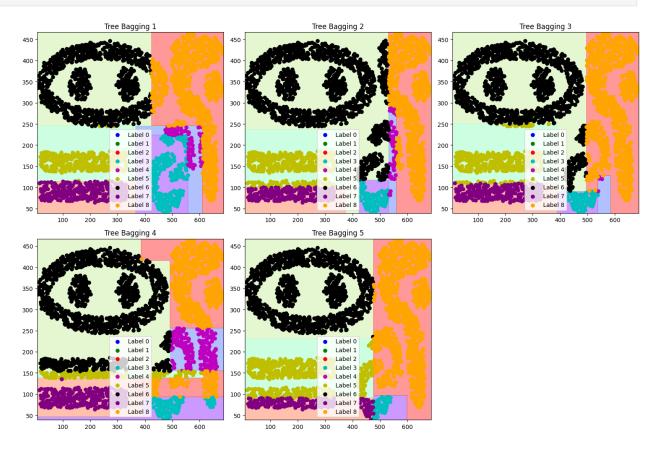
n\_estimators Bagging = 100:
Train Accuracy: 0.88, Test Accuracy: 0.87
Train Precision: 0.88, Test Precision: 0.87

Train Recall: 0.88, Test Recall: 0.87
Train F1-score: 0.88, Test F1-score: 0.87



```
<ipython-input-54-1fa8d6f22dc9>:108: MatplotlibDeprecationWarning: The
get cmap function was deprecated in Matplotlib 3.7 and will be removed
two minor releases later. Use ``matplotlib.colormaps[name]`` or
``matplotlib.colormaps.get_cmap(obj)`` instead.
  plt.contourf(xx, yy, Z, alpha=0.4, cmap=plt.cm.get cmap('rainbow',
len(np.unique(y))))
/usr/local/lib/python3.10/dist-packages/sklearn/base.py:432:
UserWarning: X has feature names, but DecisionTreeClassifier was
fitted without feature names
  warnings.warn(
<ipython-input-54-1fa8d6f22dc9>:108: MatplotlibDeprecationWarning: The
get cmap function was deprecated in Matplotlib 3.7 and will be removed
two minor releases later. Use ``matplotlib.colormaps[name]`` or
``matplotlib.colormaps.get cmap(obj)`` instead.
  plt.contourf(xx, yy, Z, alpha=0.4, cmap=plt.cm.get_cmap('rainbow',
len(np.unique(y))))
/usr/local/lib/python3.10/dist-packages/sklearn/base.py:432:
UserWarning: X has feature names, but DecisionTreeClassifier was
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  warnings.warn(
<ipython-input-54-1fa8d6f22dc9>:108: MatplotlibDeprecationWarning: The
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  plt.contourf(xx, yy, Z, alpha=0.4, cmap=plt.cm.get cmap('rainbow',
len(np.unique(y))))
/usr/local/lib/python3.10/dist-packages/sklearn/base.py:432:
UserWarning: X has feature names, but DecisionTreeClassifier was
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<ipython-input-54-1fa8d6f22dc9>:108: MatplotlibDeprecationWarning: The
get_cmap function was deprecated in Matplotlib 3.7 and will be removed
two minor releases later. Use ``matplotlib.colormaps[name]`` or
 `matplotlib.colormaps.get cmap(obj)`` instead.
  plt.contourf(xx, yy, Z, alpha=0.4, cmap=plt.cm.get cmap('rainbow',
len(np.unique(y))))
/usr/local/lib/python3.10/dist-packages/sklearn/base.py:432:
UserWarning: X has feature names, but DecisionTreeClassifier was
fitted without feature names
  warnings.warn(
<ipython-input-54-1fa8d6f22dc9>:108: MatplotlibDeprecationWarning: The
get cmap function was deprecated in Matplotlib 3.7 and will be removed
two minor releases later. Use ``matplotlib.colormaps[name]`` or
 `matplotlib.colormaps.get cmap(obj)`` instead.
  plt.contourf(xx, yy, Z, alpha=0.4, cmap=plt.cm.get cmap('rainbow',
len(np.unique(v))))
/usr/local/lib/python3.10/dist-packages/sklearn/base.py:432:
UserWarning: X has feature names, but DecisionTreeClassifier was
```

# fitted without feature names warnings.warn(



#### RandomForest

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from sklearn.tree import plot_tree
from sklearn.ensemble import RandomForestClassifier
from sklearn.model_selection import train_test_split
from sklearn.metrics import accuracy_score, precision_score,
recall_score, fl_score

# Define 9 different colors
colors = ['b', 'g', 'r', 'c', 'm', 'y', 'k', 'purple', 'orange']

# Read data from CSV file
df = pd.read_csv('/content/drive/My Drive/1/Dataset1.csv')

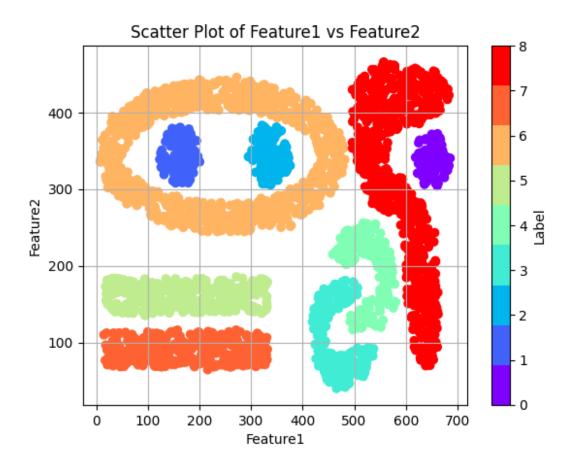
X = df.drop('Label', axis=1)
y = df['Label']

# Scatter plot of Feature 1 vs Feature 2
```

```
plt.scatter(df['Feature 1'], df['Feature 2'], c=y,
cmap=plt.cm.get cmap('rainbow', len(np.unique(y))))
plt.title('Scatter Plot of Feature1 vs Feature2')
plt.xlabel('Feature1')
plt.ylabel('Feature2')
plt.grid(True)
plt.colorbar(ticks=range(len(np.unique(y))), label='Label')
plt.show()
# Split the data into training and testing sets
X train, X test, y train, y test = train test split(X, y,
test size=0.2, random state=42)
# Number of estimators to evaluate
num estimators = [1, 2, 5, 10, 50, 100]
train accuracies = []
test accuracies = []
# Loop to test different numbers of estimators
for n in num estimators:
    # Create RandomForest model with different numbers of estimators
    rf clf = RandomForestClassifier(n estimators=n, max depth=4,
random state=42)
    # Train the model on training data
    rf clf.fit(X train, y train)
    # Predict labels for training and testing data
    y_train_pred = rf_clf.predict(X_train)
    y test pred = rf clf.predict(X test)
    # Compute model accuracy
    train accuracy = accuracy score(y train, y train pred)
    test accuracy = accuracy score(y test, y test pred)
    # Store accuracies in lists
    train accuracies.append(train accuracy)
    test accuracies.append(test accuracy)
    # Evaluate the model for training and testing data
    train f1 = f1 score(y train, y train pred, average='micro')
    test f1 = f1 score(y test, y test pred, average='micro')
    train precision = precision score(y train, y train pred,
average='micro')
    test precision = precision score(y test, y test pred,
average='micro')
    train recall = recall score(y train, y train pred,
```

```
average='micro')
    test recall = recall score(y test, y test pred, average='micro')
    print(f'n estimators RandomForest = {n}:')
    print(f'Train Accuracy: {train accuracy:.2f}, Test Accuracy:
{test accuracy:.2f}')
    print(f'Train Precision: {train precision:.2f}, Test Precision:
{test precision:.2f}')
    print(f'Train Recall: {train recall:.2f}, Test Recall:
{test recall:.2f}')
    print(f'Train F1-score: {train f1:.2f}, Test F1-score:
{test f1:.2f}')
    print()
# Plot RandomForest accuracy versus number of estimators
plt.figure(figsize=(10, 6))
plt.plot(num_estimators, train accuracies, marker='o', label='Train
Accuracy')
plt.plot(num estimators, test accuracies, marker='o', label='Test
Accuracy')
plt.title('RandomForest Accuracy vs. Number of Estimators')
plt.xlabel('Number of Estimators')
plt.ylabel('Accuracy')
plt.xticks(num estimators)
plt.legend()
plt.grid(True)
plt.tight layout()
plt.show()
# Display five random trees from the model
plt.figure(figsize=(20, 10))
for i in range(5):
    plt.subplot(2, 3, i + 1)
    # Plot the tree
    plot tree(rf clf.estimators [i], filled=True) # Use plot tree to
plot the tree
    plt.title(f'Tree RandomForest {i + 1}')
plt.tight layout()
plt.show()
# Plot decision boundaries
plt.figure(figsize=(15, 10))
# Create a mesh arid
x \min, x \max = X['Feature 1'].\min() - 1, X['Feature 1'].\max() + 1
y = min, y = X['Feature 2'].min() - 1, X['Feature 2'].max() + 1
xx, yy = np.meshgrid(np.arange(x min, x max, 0.1),
                     np.arange(y min, y max, 0.1))
```

```
for i in range(5):
    plt.subplot(2, 3, i + 1)
    # Predict the class for each point in the mesh grid
    Z = rf clf.estimators [i].predict(np.c [xx.ravel(), yy.ravel()])
    Z = Z.reshape(xx.shape)
    # Plot decision boundaries
    plt.contourf(xx, yy, Z, alpha=0.4, cmap=plt.cm.get cmap('rainbow',
len(np.unique(y))))
    # Plot original data points with predicted labels
    y pred tree = rf clf.estimators [i].predict(X)
    for label in np.unique(y):
        plt.scatter(X[y pred tree == label]['Feature 1'],
X[y pred tree == label]['Feature 2'],
                    color=colors[label], label=f'Label {label}')
    plt.title(f'Tree RandomForest {i + 1}')
    plt.legend()
plt.tight layout()
plt.show()
<ipython-input-55-27a5bddb699e>:19: MatplotlibDeprecationWarning: The
get_cmap function was deprecated in Matplotlib 3.7 and will be removed
two minor releases later. Use ``matplotlib.colormaps[name]`` or
``matplotlib.colormaps.get_cmap(obj)`` instead.
  plt.scatter(df['Feature 1'], df['Feature 2'], c=y,
cmap=plt.cm.get cmap('rainbow', len(np.unique(y))))
```



n estimators RandomForest = 1: Train Accuracy: 0.88, Test Accuracy: 0.85 Train Precision: 0.88, Test Precision: 0.85 Train Recall: 0.88, Test Recall: 0.85 Train F1-score: 0.88, Test F1-score: 0.85 n estimators RandomForest = 2: Train Accuracy: 0.90, Test Accuracy: 0.88 Train Precision: 0.90, Test Precision: 0.88 Train Recall: 0.90, Test Recall: 0.88 Train F1-score: 0.90, Test F1-score: 0.88 n estimators RandomForest = 5: Train Accuracy: 0.92, Test Accuracy: 0.90 Train Precision: 0.92, Test Precision: 0.90 Train Recall: 0.92, Test Recall: 0.90 Train F1-score: 0.92, Test F1-score: 0.90 n estimators RandomForest = 10: Train Accuracy: 0.92, Test Accuracy: 0.90 Train Precision: 0.92, Test Precision: 0.90 Train Recall: 0.92, Test Recall: 0.90 Train F1-score: 0.92, Test F1-score: 0.90

n estimators RandomForest = 50:

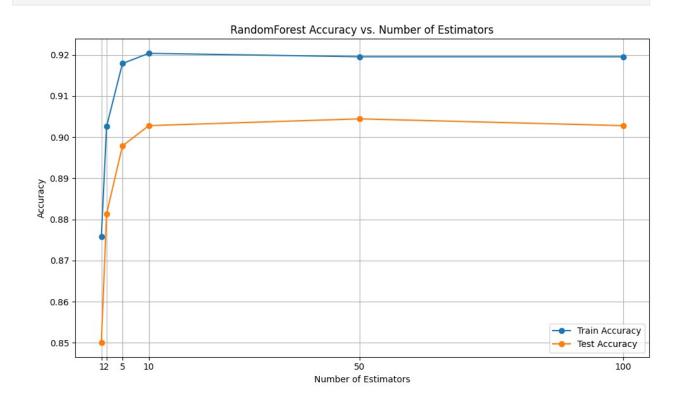
Train Accuracy: 0.92, Test Accuracy: 0.90 Train Precision: 0.92, Test Precision: 0.90

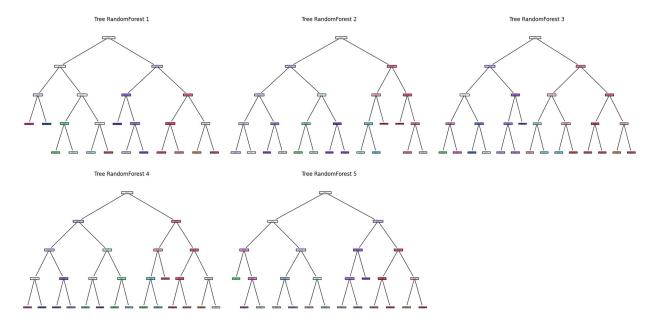
Train Recall: 0.92, Test Recall: 0.90 Train F1-score: 0.92, Test F1-score: 0.90

n estimators RandomForest = 100:

Train Accuracy: 0.92, Test Accuracy: 0.90 Train Precision: 0.92, Test Precision: 0.90

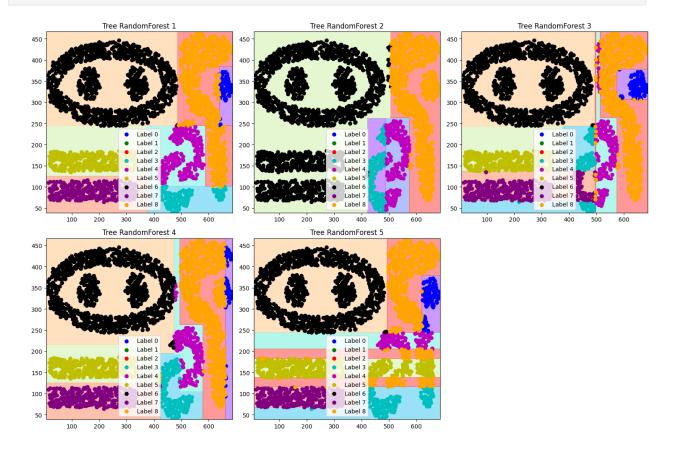
Train Recall: 0.92, Test Recall: 0.90 Train F1-score: 0.92, Test F1-score: 0.90





```
<ipython-input-55-27a5bddb699e>:115: MatplotlibDeprecationWarning: The
get cmap function was deprecated in Matplotlib 3.7 and will be removed
two minor releases later. Use ``matplotlib.colormaps[name]`` or
``matplotlib.colormaps.get_cmap(obj)`` instead.
  plt.contourf(xx, yy, Z, alpha=0.4, cmap=plt.cm.get cmap('rainbow',
len(np.unique(y))))
/usr/local/lib/python3.10/dist-packages/sklearn/base.py:432:
UserWarning: X has feature names, but DecisionTreeClassifier was
fitted without feature names
  warnings.warn(
<ipython-input-55-27a5bddb699e>:115: MatplotlibDeprecationWarning: The
get cmap function was deprecated in Matplotlib 3.7 and will be removed
two minor releases later. Use ``matplotlib.colormaps[name]`` or
``matplotlib.colormaps.get_cmap(obj)`` instead.
  plt.contourf(xx, yy, Z, alpha=0.4, cmap=plt.cm.get_cmap('rainbow',
len(np.unique(y))))
/usr/local/lib/python3.10/dist-packages/sklearn/base.py:432:
UserWarning: X has feature names, but DecisionTreeClassifier was
fitted without feature names
  warnings.warn(
<ipython-input-55-27a5bddb699e>:115: MatplotlibDeprecationWarning: The
get_cmap function was deprecated in Matplotlib 3.7 and will be removed
two minor releases later. Use ``matplotlib.colormaps[name]`` or
``matplotlib.colormaps.get cmap(obj)`` instead.
  plt.contourf(xx, yy, Z, alpha=0.4, cmap=plt.cm.get cmap('rainbow',
len(np.unique(y))))
/usr/local/lib/python3.10/dist-packages/sklearn/base.py:432:
UserWarning: X has feature names, but DecisionTreeClassifier was
fitted without feature names
  warnings.warn(
```

```
<ipython-input-55-27a5bddb699e>:115: MatplotlibDeprecationWarning: The
get cmap function was deprecated in Matplotlib 3.7 and will be removed
two minor releases later. Use ``matplotlib.colormaps[name]`` or
``matplotlib.colormaps.get cmap(obj)`` instead.
  plt.contourf(xx, yy, Z, alpha=0.4, cmap=plt.cm.get cmap('rainbow',
len(np.unique(y))))
/usr/local/lib/python3.10/dist-packages/sklearn/base.py:432:
UserWarning: X has feature names, but DecisionTreeClassifier was
fitted without feature names
  warnings.warn(
<ipython-input-55-27a5bddb699e>:115: MatplotlibDeprecationWarning: The
get_cmap function was deprecated in Matplotlib 3.7 and will be removed
two minor releases later. Use ``matplotlib.colormaps[name]`` or
``matplotlib.colormaps.get cmap(obj)`` instead.
  plt.contourf(xx, yy, Z, alpha=0.4, cmap=plt.cm.get_cmap('rainbow',
len(np.unique(y))))
/usr/local/lib/python3.10/dist-packages/sklearn/base.py:432:
UserWarning: X has feature names, but DecisionTreeClassifier was
fitted without feature names
  warnings.warn(
```



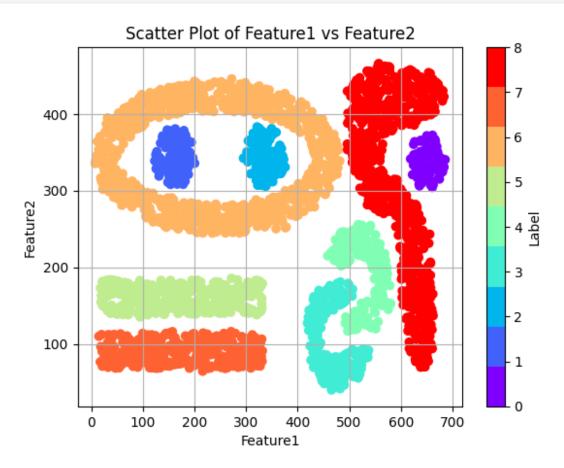
#### Adaboost

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from sklearn.tree import DecisionTreeClassifier, plot tree
from sklearn.ensemble import AdaBoostClassifier
from sklearn.model selection import train test split
from sklearn.metrics import accuracy score, precision score,
recall score, f1 score
# Define 9 different colors
colors = ['b', 'g', 'r', 'c', 'm', 'y', 'k', 'purple', 'orange']
# Read data from CSV file
df = pd.read csv('/content/drive/My Drive/1/Dataset1.csv')
X = df.drop('Label', axis=1)
y = df['Label']
# Scatter plot of Feature 1 vs Feature 2
plt.scatter(df['Feature 1'], df['Feature 2'], c=y,
cmap=plt.cm.get_cmap('rainbow', len(np.unique(y))))
plt.title('Scatter Plot of Feature1 vs Feature2')
plt.xlabel('Feature1')
plt.ylabel('Feature2')
plt.grid(True)
plt.colorbar(ticks=range(len(np.unique(y))), label='Label')
plt.show()
# Split the data into training and testing sets
X_train, X_test, y_train, y_test = train_test_split(X, y,
test_size=0.2, random_state=42)
# Number of estimators to evaluate
num estimators = [1, 2, 5, 10, 50, 100]
train accuracies = []
test accuracies = []
# Loop to test different numbers of estimators
for n in num estimators:
    # Create AdaBoost model with different numbers of estimators
    ada clf =
AdaBoostClassifier(base estimator=DecisionTreeClassifier(max depth=4,
splitter='random'),
                                 n estimators=n, random state=42)
    # Train the model on training data
    ada clf.fit(X train, y train)
```

```
# Predict labels for training and testing data
    y train pred = ada clf.predict(X train)
    y test pred = ada clf.predict(X test)
    # Compute model accuracy
    train_accuracy = accuracy_score(y_train, y_train_pred)
    test_accuracy = accuracy_score(y_test, y_test_pred)
    # Store accuracies in lists
    train accuracies.append(train accuracy)
    test accuracies.append(test accuracy)
    # Evaluate the model for training and testing data
    train f1 = f1 score(y train, y train pred, average='micro')
    test f1 = f1 score(y test, y test pred, average='micro')
    train precision = precision score(y train, y train pred,
average='micro')
    test precision = precision score(y test, y test pred,
average='micro')
    train recall = recall score(y train, y train pred,
average='micro')
    test recall = recall score(y test, y test pred, average='micro')
    print(f'n estimators AdaBoost = {n}:')
    print(f'Train Accuracy: {train accuracy:.2f}, Test Accuracy:
{test accuracy:.2f}')
    print(f'Train Precision: {train precision: 2f}, Test Precision:
{test precision:.2f}')
    print(f'Train Recall: {train recall:.2f}, Test Recall:
{test recall:.2f}')
    print(f'Train F1-score: {train f1:.2f}, Test F1-score:
{test f1:.2f}')
    print()
# Plot AdaBoost accuracy versus number of estimators
plt.figure(figsize=(10, 6))
plt.plot(num estimators, train accuracies, marker='o', label='Train
Accuracy')
plt.plot(num estimators, test accuracies, marker='o', label='Test
Accuracy')
plt.title('AdaBoost Accuracy vs. Number of Estimators')
plt.xlabel('Number of Estimators')
plt.ylabel('Accuracy')
plt.xticks(num estimators)
plt.legend()
plt.grid(True)
plt.tight layout()
plt.show()
```

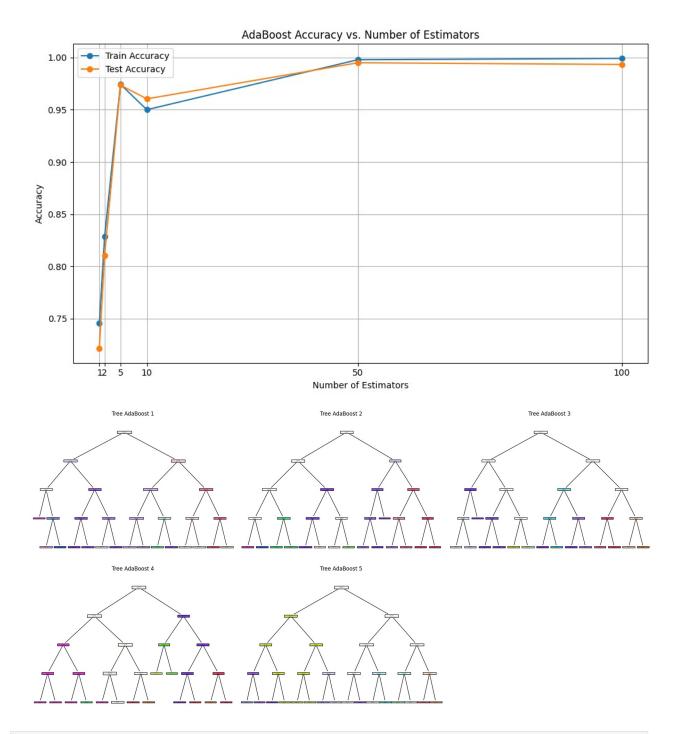
```
# Display five random trees from the model
plt.figure(figsize=(20, 10))
for i in range(5):
    plt.subplot(2, 3, i + 1)
    # Plot the tree
    plot tree(ada clf.estimators [i], filled=True) # Use plot tree to
plot the tree
    plt.title(f'Tree AdaBoost {i + 1}')
plt.tight layout()
plt.show()
# Plot decision boundaries
plt.figure(figsize=(15, 10))
# Create a mesh arid
x \min, x \max = X['Feature 1'].\min() - 1, X['Feature 1'].\max() + 1
y = min, y = X['Feature 2'].min() - 1, <math>X['Feature 2'].max() + 1
xx, yy = np.meshgrid(np.arange(x min, x max, 0.1),
                     np.arange(y min, y max, 0.1))
for i in range(5):
    plt.subplot(2, 3, i + 1)
    # Predict the class for each point in the mesh grid
    Z = ada clf.estimators [i].predict(np.c [xx.ravel(), yy.ravel()])
    Z = Z.reshape(xx.shape)
    # Plot decision boundaries
    plt.contourf(xx, yy, Z, alpha=0.4, cmap=plt.cm.get cmap('rainbow',
len(np.unique(y))))
    # Plot original data points with predicted labels
    y pred tree = ada clf.estimators [i].predict(X)
    for label in np.unique(y):
        plt.scatter(X[y_pred_tree == label]['Feature 1'],
X[y pred tree == label]['Feature 2'],
                    color=colors[label], label=f'Label {label}')
    plt.title(f'Tree AdaBoost {i + 1}')
    plt.legend()
plt.tight layout()
plt.show()
<ipython-input-56-9d42626d182c>:19: MatplotlibDeprecationWarning: The
get cmap function was deprecated in Matplotlib 3.7 and will be removed
two minor releases later. Use ``matplotlib.colormaps[name]`` or
 `matplotlib.colormaps.get cmap(obj)`` instead.
```

plt.scatter(df['Feature 1'], df['Feature 2'], c=y,
cmap=plt.cm.get cmap('rainbow', len(np.unique(y))))



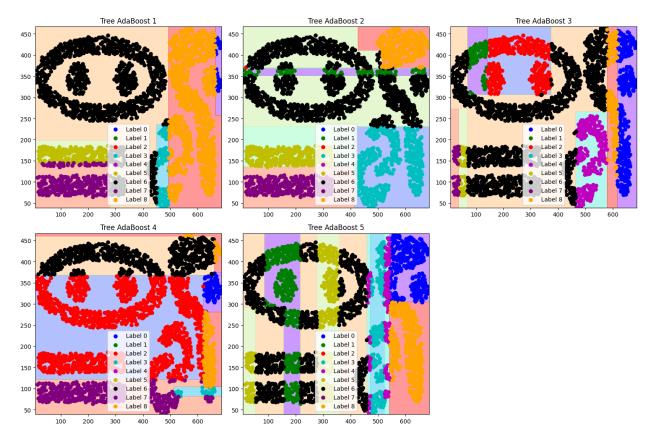
```
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/_base.py:166:
FutureWarning: `base estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
  warnings.warn(
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/_base.py:166:
FutureWarning: `base_estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
  warnings.warn(
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/ base.py:166:
FutureWarning: `base_estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
  warnings.warn(
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/ base.py:166:
FutureWarning: `base_estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
  warnings.warn(
n estimators AdaBoost = 1:
Train Accuracy: 0.75, Test Accuracy: 0.72
```

```
Train Precision: 0.75, Test Precision: 0.72
Train Recall: 0.75, Test Recall: 0.72
Train F1-score: 0.75, Test F1-score: 0.72
n estimators AdaBoost = 2:
Train Accuracy: 0.83, Test Accuracy: 0.81
Train Precision: 0.83, Test Precision: 0.81
Train Recall: 0.83, Test Recall: 0.81
Train F1-score: 0.83, Test F1-score: 0.81
n estimators AdaBoost = 5:
Train Accuracy: 0.97, Test Accuracy: 0.97
Train Precision: 0.97, Test Precision: 0.97
Train Recall: 0.97, Test Recall: 0.97
Train F1-score: 0.97, Test F1-score: 0.97
n estimators AdaBoost = 10:
Train Accuracy: 0.95, Test Accuracy: 0.96
Train Precision: 0.95, Test Precision: 0.96
Train Recall: 0.95, Test Recall: 0.96
Train F1-score: 0.95, Test F1-score: 0.96
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/ base.py:166:
FutureWarning: `base_estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
  warnings.warn(
n estimators AdaBoost = 50:
Train Accuracy: 1.00, Test Accuracy: 1.00
Train Precision: 1.00, Test Precision: 1.00
Train Recall: 1.00, Test Recall: 1.00
Train F1-score: 1.00, Test F1-score: 1.00
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/ base.py:166:
FutureWarning: `base estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
 warnings.warn(
n estimators AdaBoost = 100:
Train Accuracy: 1.00, Test Accuracy: 0.99
Train Precision: 1.00, Test Precision: 0.99
Train Recall: 1.00, Test Recall: 0.99
Train F1-score: 1.00, Test F1-score: 0.99
```



<ipython-input-56-9d42626d182c>:116: MatplotlibDeprecationWarning: The
get\_cmap function was deprecated in Matplotlib 3.7 and will be removed
two minor releases later. Use ``matplotlib.colormaps[name]`` or
``matplotlib.colormaps.get\_cmap(obj)`` instead.
 plt.contourf(xx, yy, Z, alpha=0.4, cmap=plt.cm.get\_cmap('rainbow',
len(np.unique(y))))
/usr/local/lib/python3.10/dist-packages/sklearn/base.py:432:
UserWarning: X has feature names, but DecisionTreeClassifier was

```
fitted without feature names
  warnings.warn(
<ipython-input-56-9d42626d182c>:116: MatplotlibDeprecationWarning: The
get cmap function was deprecated in Matplotlib 3.7 and will be removed
two minor releases later. Use ``matplotlib.colormaps[name]`` or
 `matplotlib.colormaps.get_cmap(obj)`` instead.
  plt.contourf(xx, yy, Z, alpha=0.4, cmap=plt.cm.get cmap('rainbow',
len(np.unique(y))))
/usr/local/lib/python3.10/dist-packages/sklearn/base.py:432:
UserWarning: X has feature names, but DecisionTreeClassifier was
fitted without feature names
  warnings.warn(
<ipython-input-56-9d42626d182c>:116: MatplotlibDeprecationWarning: The
get cmap function was deprecated in Matplotlib 3.7 and will be removed
two minor releases later. Use ``matplotlib.colormaps[name]`` or
 `matplotlib.colormaps.get cmap(obj)`` instead.
  plt.contourf(xx, yy, Z, alpha=0.4, cmap=plt.cm.get_cmap('rainbow',
len(np.unique(v))))
/usr/local/lib/python3.10/dist-packages/sklearn/base.py:432:
UserWarning: X has feature names, but DecisionTreeClassifier was
fitted without feature names
  warnings.warn(
<ipython-input-56-9d42626d182c>:116: MatplotlibDeprecationWarning: The
get cmap function was deprecated in Matplotlib 3.7 and will be removed
two minor releases later. Use ``matplotlib.colormaps[name]`` or
``matplotlib.colormaps.get cmap(obj)`` instead.
  plt.contourf(xx, yy, Z, alpha=0.4, cmap=plt.cm.get_cmap('rainbow',
len(np.unique(y))))
/usr/local/lib/python3.10/dist-packages/sklearn/base.py:432:
UserWarning: X has feature names, but DecisionTreeClassifier was
fitted without feature names
  warnings.warn(
<ipython-input-56-9d42626d182c>:116: MatplotlibDeprecationWarning: The
get cmap function was deprecated in Matplotlib 3.7 and will be removed
two minor releases later. Use ``matplotlib.colormaps[name]`` or
``matplotlib.colormaps.get cmap(obj)`` instead.
  plt.contourf(xx, yy, Z, alpha=0.4, cmap=plt.cm.get cmap('rainbow',
len(np.unique(y))))
/usr/local/lib/python3.10/dist-packages/sklearn/base.py:432:
UserWarning: X has feature names, but DecisionTreeClassifier was
fitted without feature names
  warnings.warn(
```



# Stacked Learning

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from sklearn.tree import DecisionTreeClassifier
from sklearn.ensemble import RandomForestClassifier,
AdaBoostClassifier
from sklearn.neighbors import KNeighborsClassifier
from sklearn.model selection import train test split
from sklearn.metrics import accuracy score, precision score,
recall score, f1 score
# Load the dataset
df = pd.read csv('/content/drive/My Drive/1/Dataset1.csv')
X = df.drop('Label', axis=1)
y = df['Label']
# Split the data into training and testing sets
X train, X test, y train, y test = train test split(X, y,
test size=0.2, random state=42)
# Define base classifiers
base classifiers = {
```

```
'knn': KNeighborsClassifier(n neighbors=3),
    'decision tree': DecisionTreeClassifier(max depth=4,
random state=42),
    'random forest': RandomForestClassifier(n estimators=10,
max depth=4, random state=42),
    'adaboost':
AdaBoostClassifier(base estimator=DecisionTreeClassifier(max depth=4),
n estimators=10, random state=42)
# Train base classifiers
for name, clf in base classifiers.items():
    clf.fit(X train, y train)
    y train pred = clf.predict(X train)
    y test pred = clf.predict(X test)
    print(f'{name} - Train Accuracy: {accuracy score(y train,
y train pred):.2f}, Test Accuracy: {accuracy score(y test,
y test pred):.2f}')
# Create new features for the meta-classifier
meta features train = np.column stack([clf.predict(X train) for clf in
base classifiers.values()])
meta features test = np.column stack([clf.predict(X test) for clf in
base classifiers.values()])
# Define and train the meta-classifier
meta classifier = DecisionTreeClassifier(max depth=4, random state=42)
meta classifier.fit(meta features train, y train)
# Evaluate the stacked model
y train pred meta = meta classifier.predict(meta features train)
y test pred meta = meta classifier.predict(meta features test)
train accuracy = accuracy score(y train, y train pred meta)
test accuracy = accuracy score(y test, y test pred meta)
train precision = precision score(y train, y train pred meta,
average='micro')
test precision = precision score(y test, y test pred meta,
average='micro')
train recall = recall score(y train, y train pred meta,
average='micro')
test_recall = recall_score(y_test, y_test_pred_meta, average='micro')
train f1 = f1 score(y train, y train pred meta, average='micro')
test_f1 = f1_score(y_test, y_test_pred_meta, average='micro')
print(f'Stacked Model:')
print(f'Train Accuracy: {train accuracy:.2f}, Test Accuracy:
```

```
{test accuracy:.2f}')
print(f'Train Precision: {train precision:.2f}, Test Precision:
{test precision:.2f}')
print(f'Train Recall: {train recall:.2f}, Test Recall:
{test recall:.2f}')
print(f'Train F1-score: {train f1:.2f}, Test F1-score: {test f1:.2f}')
knn - Train Accuracy: 1.00, Test Accuracy: 1.00
decision tree - Train Accuracy: 0.91, Test Accuracy: 0.89
random forest - Train Accuracy: 0.92, Test Accuracy: 0.90
adaboost - Train Accuracy: 1.00, Test Accuracy: 0.99
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/ base.py:166:
FutureWarning: `base estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
 warnings.warn(
Stacked Model:
Train Accuracy: 0.90, Test Accuracy: 0.89
Train Precision: 0.90, Test Precision: 0.89
Train Recall: 0.90, Test Recall: 0.89
Train F1-score: 0.90, Test F1-score: 0.89
```

# Phase 2

### **Preprocess**

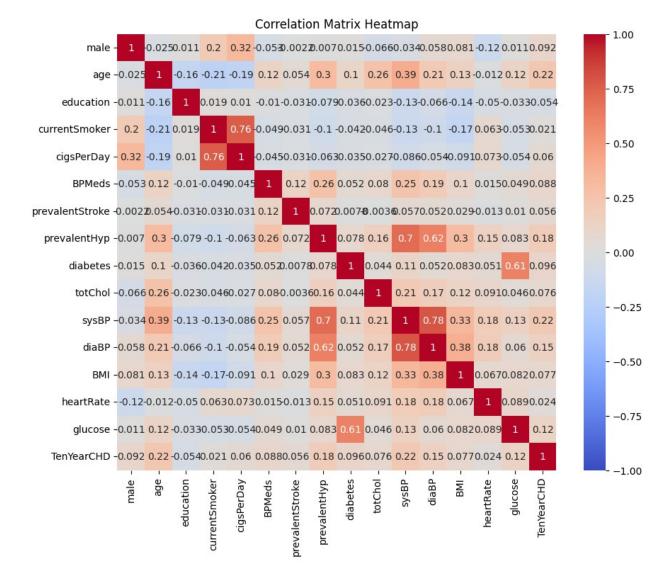
```
import pandas as pd
df = pd.read csv('/content/drive/My Drive/1/Dataset2.csv')
df.info()
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 4240 entries, 0 to 4239
Data columns (total 16 columns):
#
     Column
                       Non-Null Count
                                        Dtype
- - -
     _ _ _ _ _
                       _ _ _ _ _ _ _ _ _ _ _ _ _ _ _ _
 0
     male
                       4240 non-null
                                        int64
 1
     age
                       4240 non-null
                                        int64
 2
                                        float64
     education
                       4135 non-null
 3
     currentSmoker
                       4240 non-null
                                        int64
 4
                       4211 non-null
                                        float64
     cigsPerDay
 5
     BPMeds
                       4187 non-null
                                        float64
     prevalentStroke 4240 non-null
 6
                                        int64
                                        int64
 7
                       4240 non-null
     prevalentHyp
 8
     diabetes
                       4240 non-null
                                        int64
 9
     totChol
                       4190 non-null
                                        float64
                       4240 non-null
                                        float64
 10 svsBP
```

```
11 diaBP
                      4240 non-null
                                      float64
                                      float64
 12
    BMI
                      4221 non-null
13 heartRate
                      4239 non-null
                                      float64
 14
    alucose
                      3852 non-null
                                      float64
15 TenYearCHD
                     4240 non-null
                                      int64
dtypes: float64(9), int64(7)
memory usage: 530.1 KB
df['education'].unique()
array([ 4., 2., 1., 3., nan])
# Calculate the mode of the column
mode value = df['education'].mode()[0]
# Fill NaN values with the mode
df['education'].fillna(mode value, inplace=True)
df['cigsPerDay'].unique()
array([ 0., 20., 30., 23., 15., 9., 10., 5., 35., 43., 1., 40.,
3.,
        2., nan, 12., 4., 18., 25., 60., 14., 45., 8., 50., 13.,
11.,
       7., 6., 38., 29., 17., 16., 19., 70.])
# Calculate the median of the column
median value = df['cigsPerDay'].median()
# Fill NaN values with the median
df['cigsPerDay'].fillna(median value, inplace=True)
df['BPMeds'].unique()
array([ 0., 1., nan])
df = df.dropna(subset=['BPMeds'])
# Calculate the median of the column
median value = df['totChol'].median()
# Fill NaN values with the median
df['totChol'].fillna(median_value, inplace=True)
<ipython-input-66-6f836d55e12c>:5: SettingWithCopyWarning:
A value is trying to be set on a copy of a slice from a DataFrame
See the caveats in the documentation:
https://pandas.pydata.org/pandas-docs/stable/user guide/indexing.html#
returning-a-view-versus-a-copy
  df['totChol'].fillna(median value, inplace=True)
```

```
# Calculate the median of the column
mean value = df['BMI'].mean()
# Fill NaN values with the median
df['BMI'].fillna(mean value, inplace=True)
<ipython-input-67-a33f1e4e780a>:5: SettingWithCopyWarning:
A value is trying to be set on a copy of a slice from a DataFrame
See the caveats in the documentation:
https://pandas.pydata.org/pandas-docs/stable/user guide/indexing.html#
returning-a-view-versus-a-copy
  df['BMI'].fillna(mean value, inplace=True)
# Calculate the median of the column
median value = df['heartRate'].median()
# Fill NaN values with the median
df['heartRate'].fillna(median_value, inplace=True)
<ipython-input-68-25274b80e945>:5: SettingWithCopyWarning:
A value is trying to be set on a copy of a slice from a DataFrame
See the caveats in the documentation:
https://pandas.pydata.org/pandas-docs/stable/user guide/indexing.html#
returning-a-view-versus-a-copy
 df['heartRate'].fillna(median value, inplace=True)
# Calculate the median of the column
median_value = df['glucose'].median()
# Fill NaN values with the median
df['glucose'].fillna(median value, inplace=True)
<ipython-input-69-3502970513b1>:5: SettingWithCopyWarning:
A value is trying to be set on a copy of a slice from a DataFrame
See the caveats in the documentation:
https://pandas.pydata.org/pandas-docs/stable/user_guide/indexing.html#
returning-a-view-versus-a-copy
  df['glucose'].fillna(median value, inplace=True)
df.head()
{"summary":"{\n \"name\": \"df\",\n \"rows\": 4187,\n \"fields\":
      {\n \"column\": \"male\",\n \"properties\": {\n
\"dtype\": \"number\",\n \"std\": 0,\n
                                                  \"min\": 0,\n
                    \"num_unique_values\": 2,\n
\"max\": 1,\n
                                                       \"samples\":
                                    ],\n
[\n
            0,\n
                         1\n
                                                 \"semantic type\":
                                                },\n {√n
\"\",\n
              \"description\": \"\"\n
                                        }\n
\"column\": \"age\",\n \"properties\": {\n
                                                     \"dtype\":
```

```
\"number\",\n \"std\": 8,\n \"min\": 32,\n \"max\": 70,\n \"num_unique_values\": 39,\n \"samples\": [\n 34,\n 70\n ],\n \"semantic_type\": \"\",\n \"description\": \"\"\n }\n },\n {\n
\"min\": 1.0,\n \"max\": 4.0,\n \"num_unique_values\":
4,\n \"samples\": [\n 2.0,\n 3.0\n ],\n \"semantic_type\": \"\",\n \"description\": \"\"\n
0,\n \"min\": 0,\n \"max\": 1,\n
\"num_unique_values\": 2,\n \"samples\": [\n
                                                       1, n
\"description\": \"\"\n }\n {\n \"column\": \"cigsPerDay\",\n \"properties\": {\n \"dtype\": \"number\",\n \"std\": 11.901904311142495,\n \"min\":
                                             \"column\":
0.0,\n \"max\": 70.0,\n \"num_unique_values\": 33,\n \"samples\": [\n 19.0,\n 4.0\n ],\n \"semantic_type\": \"\",\n \"description\": \"\"\n }\
n },\n {\n \"column\": \"BPMeds\",\n \"properties\":
{\n \"dtype\": \"number\",\n
                                       \"std\":
0.16954428739626196,\n \"min\": 0.0,\n \"max\'\"num_unique_values\": 2,\n \"samples\": [\n
                                                \mbox{"max}": 1.0,\n
                                                      1.0,\n
0.0\n ],\n \"semantic_type\": \"\",\n
\"description\": \"\"\n }\n {\n \"column\":
\"prevalentStroke\",\n \"properties\": {\n \"dtype\":
\"semantic type\":
                                                     \"dtype\":
n \"dtype\": \"number\",\n \"std\": 21.98290660966593,\n \"min\": 83.5,\n \"max\": 295.0,\n
```

```
117.0,\
\"diaBP\",\n \"properties\": {\n \"dtype\": \"number\",\n
\"std\": 11.878463990074186,\n \"min\": 48.0,\n \"max\":
142.5,\n \"num_unique_values\": 146,\n \"samples\": [\n
86.5,\n 108.\overline{5}\n ],\n \"semantic_type\": \"\",\n \"description\": \"\"\n }\n \\"column\":
\"BMI\",\n \"properties\": {\n \"dtype\": \"number\",\n
\"std\": 4.067178635883306,\n \"min\": 15.54,\n \"max\":
56.8,\n \"num_unique_values\": 1360,\n \"samples\": [\n
19.66,\n 30.99\n ],\n \"semantic_type\": \"\",\n \"description\": \"\"n }\n {\n \"column\": \"heartRate\",\n \"properties\": {\n \"dtype\": \"number\",\n \"std\": 12.053463897069951,\n
\"min\": 44.0,\n \"max\": 143.0,\n
\"num_unique_values\": 73,\n \"samples\": [\n
                                                             85.0,\n
\"glucose\",\n\\"properties\": {\n\\"dtype\": \"number\",\n\\"std\": 22.932432152397404,\n\\"min\": 40.0,\n
\"max\": 394.0,\n \"num_unique_values\": 142,\n \"samples\": [\n 332.0,\n 74.0\n
\"semantic_type\": \"\",\n \"description\": \"\"\n }\
n },\n {\n \"column\": \"TenYearCHD\",\n \"properties\": {\n \"dtype\": \"number\",\n \"std\":
0,\n \"min\": 0,\n \"max\": 1,\n \"num_unique_values\": 2,\n \"samples\": [\n
                                                            1, n
0\n ],\n \"semantic type\": \"\",\n
n}","type":"dataframe","variable name":"df"}
# Calculate the correlation matrix
correlation matrix = df.corr()
import seaborn as sns
import matplotlib.pyplot as plt
plt.figure(figsize=(10, 8))
# Draw the heatmap
sns.heatmap(correlation matrix, annot=True, cmap='coolwarm', vmin=-1,
vmax=1, center=0)
# Add title and labels
plt.title('Correlation Matrix Heatmap')
plt.show()
```



```
columns_to_drop = set()

# Iterate through the correlation matrix
for i in range(len(correlation_matrix.columns)):
    for j in range(i):
        if abs(correlation_matrix.iloc[i, j]) > threshold:
            column_to_drop = correlation_matrix.columns[i]
            columns_to_drop.add(column_to_drop)

# Create a new DataFrame with only the columns that have low correlation
df_low_corr = df.drop(columns=columns_to_drop)

print("Columns to drop due to high correlation:")
print(columns_to_drop)
print("\nDataFrame with only low correlated columns:")
print(df_low_corr)
```

```
Columns to drop due to high correlation:
{'cigsPerDay', 'diaBP'}
DataFrame with only low correlated columns:
                                                     prevalentStroke \
      male
            age
                 education currentSmoker BPMeds
         1
             39
                        4.0
                                         0
                                                0.0
1
         0
             46
                        2.0
                                          0
                                                0.0
                                                                    0
2
                                                                    0
                                          1
         1
             48
                        1.0
                                                0.0
3
                                          1
                                                                    0
             61
                        3.0
                                                0.0
4
             46
                        3.0
                                          1
                                                0.0
                                                                    0
. . .
                                                                    0
4234
             51
                                         1
                                                0.0
         1
                        3.0
4236
         0
             44
                        1.0
                                          1
                                                0.0
                                                                    0
             52
                                                                    0
4237
         0
                        2.0
                                          0
                                                0.0
                                                                    0
4238
             40
                        3.0
                                          0
                                                0.0
         1
4239
             39
                        3.0
                                          1
                                                0.0
                                                                    0
      prevalentHyp diabetes totChol sysBP
                                                       heartRate
                                               BMI
glucose \
                                 195.0 106.0
                                               26.97
                                                            80.0
77.0
                                 250.0 121.0 28.73
                            0
                                                            95.0
1
76.0
2
                                 245.0 127.5
                                               25.34
                                                            75.0
70.0
                                 225.0
                                       150.0
                                                28.58
                                                            65.0
103.0
4
                                 285.0
                                       130.0
                                                23.10
                                                            85.0
85.0
. . .
4234
                                 207.0
                                       126.5
                                               19.71
                                                            65.0
68.0
4236
                                 210.0 126.5
                                                            86.0
                                               19.16
78.0
4237
                                 269.0 133.5 21.47
                                                            80.0
107.0
4238
                                 185.0 141.0 25.60
                                                            67.0
72.0
4239
                                 196.0 133.0 20.91
                                                            85.0
80.0
      TenYearCHD
0
               0
1
               0
2
               0
3
               1
4
               0
4234
```

```
4236
               0
4237
               0
4238
               0
4239
               0
[4187 rows x 14 columns]
df low corr
{"summary":"{\n \"name\": \"df_low_corr\",\n \"rows\": 4187,\n
0,\n \"min\": 0,\n \"max\": 1,\n \"num_unique_values\": 2,\n \"samples\": [\n
                                                            0, n
1\n ],\n \"semantic_type\": \"\",\n
\"num_unique_values\": 39,\n \"samples\": [\n
\"education\",\n \"properties\": {\n \"dtype\": \"number\",\n \"std\": 1.01787672260344,\n \"min\":
1.0,\n \"max\": 4.0,\n \"num_unique_values\": 4,\n
\"samples\": [\n 2.0,\n
                                         \overline{3.0}\n ],\n
\"semantic type\": \"\",\n \"description\": \"\"\n
n },\n {\n \"column\": \"currentSmoker\",\n
\"properties\": {\n \"dtype\": \"number\",\n \"std\":
0,\n \"min\": 0,\n \"max\": 1,\n \"num_unique_values\": 2,\n \"samples\": [\n
                                                            1, n
0\n ],\n \"semantic_type\": \"\",\n
\"std\": 0.16954428739626196,\n \"min\": 0.0,\n \"max\":
1.0,\n \"num_unique_values\": 2,\n \"samples\": [\n 1.0,\n 0.0\n ],\n \"semantic_type\": \"\",\n \"description\": \"\"\n }\n },\n {\n \"column\": \"prevalentStroke\",\n \"properties\": {\n \"dtype\":
\"number\",\n \"std\": 0,\n \"min\": 0,\n \"max\": 1,\n \"num_unique_values\": 2,\n \"semantic_type\": [\n 1,\n 0\n ],\n \"semantic_type\": \"\",\n \"description\": \"\"\n }\n },\n {\n
\"column\": \"prevalentHyp\",\n \"properties\": {\n
\"dtype\": \"number\",\n \"std\": 0,\n \"min\": 0,\n \"max\": 1,\n \"num_unique_values\": 2,\n \"samples\":
                   \"semantic_type\":
\"column\": \"diabetes\",\n \"properties\": {\n
                                                        \"dtype\":
\"number\",\n \"std\": 0,\n \"min\": 0,\n
\"max\": 1,\n \"num_unique_values\": 2,\n \"samples\":
```

```
\"column\": \"totChol\",\n \"properties\": {\n \"dtype\": \"number\",\n \"std\": 44.2237265788299,\n \"min\":
107.0,\n \"max\": 696.0,\n \"num_unique_values\": 248,\n \"samples\": [\n 311.0,\n 205.0\n ],\n
\"semantic_type\": \"\",\n \"description\": \"\"\n }\
    \"dtype\": \"number\",\n \"std\": 21.98290660966593,\n
\"min\": 83.5,\n \"max\": 295.0,\n
\"num_unique_values\": 233,\n \"samples\": [\n 11 n 191.5\n ],\n \"semantic_type\": \"\",\n \"description\": \"\"\n }\n {\n \"column\":
                                                       117.0,\
                                      \"dtype\": \"number\",\n
\"BMI\",\n \"properties\": {\n
\"std\": 4.067178635883306,\n \"min\": 15.54,\n \"max\":
56.8,\n \"num unique values\": 1360,\n \"samples\": [\n
19.66,\n 30.99\n ],\n \"semantic_type\": \"\",\n \"description\": \"\"n }\n }\n {\n \"column\": \"heartRate\",\n \"properties\": {\n \"dtype\": \"number\",\n \"std\": 12.053463897069951,\n
\"min\": 44.0,\n \"max\": 143.0,\n
\"num_unique_values\": 73,\n \"samples\": [\n
                                                       85.0,\n
\"max\": 394.0,\n \"num_unique_values\": 142,\n \"samples\": [\n 332.0,\n 74.0\n
\"semantic_type\": \"\",\n \"description\": \"\"\n }\
0\n ],\n \"semantic type\": \"\",\n
n}","type":"dataframe","variable name":"df low corr"}
```

## Bagging

```
# Define 9 different colors
colors = ['b', 'g', 'r', 'c', 'm', 'y', 'k', 'purple', 'orange']

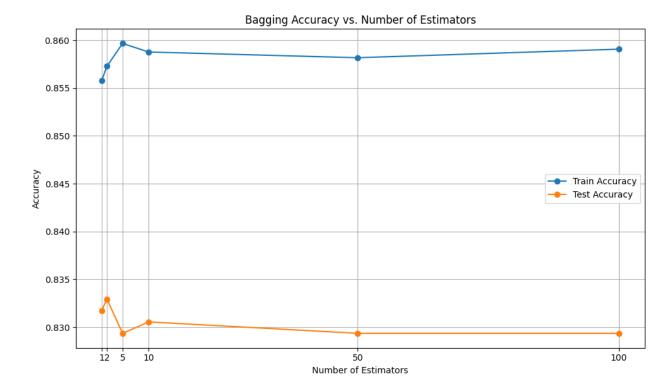
X = df_low_corr.drop('TenYearCHD', axis=1)
y = df_low_corr['TenYearCHD']

# Split the data into training and testing sets
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
```

```
# Number of estimators to evaluate
num estimators = [1, 2, 5, 10, 50, 100]
train accuracies = []
test accuracies = []
# Loop to test different numbers of estimators
for n in num estimators:
    # Create Bagging model with different numbers of estimators
    bagging clf =
BaggingClassifier(base estimator=DecisionTreeClassifier(max depth=4,
splitter='random'),
                                    n estimators=n,
                                    random state=42)
    # Train the model on training data
    bagging clf.fit(X train, y train)
    # Predict labels for training and testing data
    y_train_pred = bagging_clf.predict(X_train)
    y test pred = bagging clf.predict(X test)
    # Compute model accuracy
    train_accuracy = accuracy_score(y_train, y_train_pred)
    test accuracy = accuracy score(y test, y test pred)
    # Store accuracies in lists
    train accuracies.append(train accuracy)
    test accuracies.append(test accuracy)
    # Evaluate the model for training and testing data
    train_f1 = f1_score(y_train, y_train_pred, average='micro')
    test f1 = f1 score(y test, y test pred, average='micro')
    train precision = precision score(y train, y train pred,
average='micro')
    test precision = precision score(y test, y test pred,
average='micro')
    train recall = recall score(y train, y train pred,
average='micro')
    test recall = recall score(y test, y test pred, average='micro')
    print(f'n estimators Bagging = {n}:')
    print(f'Train Accuracy: {train accuracy:.2f}, Test Accuracy:
{test accuracy:.2f}')
    print(f'Train Precision: {train precision:.2f}, Test Precision:
{test precision:.2f}')
    print(f'Train Recall: {train_recall:.2f}, Test Recall:
{test recall:.2f}')
```

```
print(f'Train F1-score: {train f1:.2f}, Test F1-score:
{test f1:.2f}')
    print()
# Plot Bagging accuracy versus number of estimators
plt.figure(figsize=(10, 6))
plt.plot(num estimators, train accuracies, marker='o', label='Train
Accuracy')
plt.plot(num estimators, test accuracies, marker='o', label='Test
Accuracy')
plt.title('Bagging Accuracy vs. Number of Estimators')
plt.xlabel('Number of Estimators')
plt.ylabel('Accuracy')
plt.xticks(num estimators)
plt.legend()
plt.grid(True)
plt.tight layout()
plt.show()
n estimators Bagging = 1:
Train Accuracy: 0.86, Test Accuracy: 0.83
Train Precision: 0.86, Test Precision: 0.83
Train Recall: 0.86, Test Recall: 0.83
Train F1-score: 0.86, Test F1-score: 0.83
n estimators Bagging = 2:
Train Accuracy: 0.86, Test Accuracy: 0.83
Train Precision: 0.86, Test Precision: 0.83
Train Recall: 0.86, Test Recall: 0.83
Train F1-score: 0.86, Test F1-score: 0.83
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/ base.py:166:
FutureWarning: `base_estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
  warnings.warn(
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/ base.py:166:
FutureWarning: `base_estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
  warnings.warn(
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/ base.py:166:
FutureWarning: `base_estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
  warnings.warn(
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/_base.py:166:
FutureWarning: `base estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
  warnings.warn(
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/ base.py:166:
FutureWarning: `base_estimator` was renamed to `estimator` in version
```

```
1.2 and will be removed in 1.4.
 warnings.warn(
n estimators Bagging = 5:
Train Accuracy: 0.86, Test Accuracy: 0.83
Train Precision: 0.86, Test Precision: 0.83
Train Recall: 0.86, Test Recall: 0.83
Train F1-score: 0.86, Test F1-score: 0.83
n estimators Bagging = 10:
Train Accuracy: 0.86, Test Accuracy: 0.83
Train Precision: 0.86, Test Precision: 0.83
Train Recall: 0.86, Test Recall: 0.83
Train F1-score: 0.86, Test F1-score: 0.83
n estimators Bagging = 50:
Train Accuracy: 0.86, Test Accuracy: 0.83
Train Precision: 0.86, Test Precision: 0.83
Train Recall: 0.86, Test Recall: 0.83
Train F1-score: 0.86, Test F1-score: 0.83
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/ base.py:166:
FutureWarning: `base_estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
  warnings.warn(
n estimators Bagging = 100:
Train Accuracy: 0.86, Test Accuracy: 0.83
Train Precision: 0.86, Test Precision: 0.83
Train Recall: 0.86, Test Recall: 0.83
Train F1-score: 0.86, Test F1-score: 0.83
```

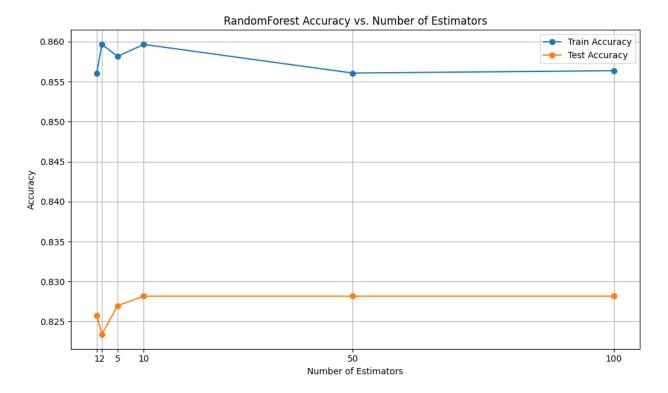


#### RandomForest

```
# Define 9 different colors
colors = ['b', 'g', 'r', 'c', 'm', 'y', 'k', 'purple', 'orange']
X = df_low_corr.drop('TenYearCHD', axis=1)
y = df low corr['TenYearCHD']
# Split the data into training and testing sets
X_train, X_test, y_train, y_test = train_test_split(X, y,
test_size=0.2, random_state=42)
# Number of estimators to evaluate
num estimators = [1, 2, 5, 10, 50, 100]
train accuracies = []
test accuracies = []
# Loop to test different numbers of estimators
for n in num estimators:
    # Create RandomForest model with different numbers of estimators
    rf clf = RandomForestClassifier(n estimators=n, max depth=4,
random state=42)
```

```
# Train the model on training data
    rf clf.fit(X train, y train)
    # Predict labels for training and testing data
    y train pred = rf clf.predict(X train)
    y test pred = rf clf.predict(X test)
    # Compute model accuracy
    train accuracy = accuracy score(y train, y train pred)
    test accuracy = accuracy score(y test, y test pred)
    # Store accuracies in lists
    train accuracies.append(train accuracy)
    test accuracies.append(test accuracy)
    # Evaluate the model for training and testing data
    train_f1 = f1_score(y_train, y_train_pred, average='micro')
    test_f1 = f1_score(y_test, y_test_pred, average='micro')
    train precision = precision score(y train, y train pred,
average='micro')
    test precision = precision score(y test, y test pred,
average='micro')
    train recall = recall score(y train, y train pred,
average='micro')
    test recall = recall score(y test, y test pred, average='micro')
    print(f'n estimators RandomForest = {n}:')
    print(f'Train Accuracy: {train accuracy:.2f}, Test Accuracy:
{test accuracy:.2f}')
    print(f'Train Precision: {train precision:.2f}, Test Precision:
{test precision:.2f}')
    print(f'Train Recall: {train recall:.2f}, Test Recall:
{test recall:.2f}')
    print(f'Train F1-score: {train f1:.2f}, Test F1-score:
{test_f1:.2f}')
    print()
# Plot RandomForest accuracy versus number of estimators
plt.figure(figsize=(10, 6))
plt.plot(num estimators, train accuracies, marker='o', label='Train
Accuracy')
plt.plot(num estimators, test accuracies, marker='o', label='Test
Accuracy')
plt.title('RandomForest Accuracy vs. Number of Estimators')
plt.xlabel('Number of Estimators')
plt.ylabel('Accuracy')
plt.xticks(num estimators)
plt.legend()
```

```
plt.grid(True)
plt.tight layout()
plt.show()
n estimators RandomForest = 1:
Train Accuracy: 0.86, Test Accuracy: 0.83
Train Precision: 0.86, Test Precision: 0.83
Train Recall: 0.86, Test Recall: 0.83
Train F1-score: 0.86, Test F1-score: 0.83
n estimators RandomForest = 2:
Train Accuracy: 0.86, Test Accuracy: 0.82
Train Precision: 0.86, Test Precision: 0.82
Train Recall: 0.86, Test Recall: 0.82
Train F1-score: 0.86, Test F1-score: 0.82
n estimators RandomForest = 5:
Train Accuracy: 0.86, Test Accuracy: 0.83
Train Precision: 0.86, Test Precision: 0.83
Train Recall: 0.86, Test Recall: 0.83
Train F1-score: 0.86, Test F1-score: 0.83
n estimators RandomForest = 10:
Train Accuracy: 0.86, Test Accuracy: 0.83
Train Precision: 0.86, Test Precision: 0.83
Train Recall: 0.86, Test Recall: 0.83
Train F1-score: 0.86, Test F1-score: 0.83
n estimators RandomForest = 50:
Train Accuracy: 0.86, Test Accuracy: 0.83
Train Precision: 0.86, Test Precision: 0.83
Train Recall: 0.86, Test Recall: 0.83
Train F1-score: 0.86, Test F1-score: 0.83
n estimators RandomForest = 100:
Train Accuracy: 0.86, Test Accuracy: 0.83
Train Precision: 0.86, Test Precision: 0.83
Train Recall: 0.86, Test Recall: 0.83
Train F1-score: 0.86, Test F1-score: 0.83
```



#### Adaboost

```
# Define 9 different colors
colors = ['b', 'g', 'r', 'c', 'm', 'y', 'k', 'purple', 'orange']
X = df_low_corr.drop('TenYearCHD', axis=1)
y = df low corr['TenYearCHD']
# Split the data into training and testing sets
X_train, X_test, y_train, y_test = train_test_split(X, y,
test_size=0.2, random_state=42)
# Number of estimators to evaluate
num estimators = [1, 2, 5, 10, 50, 100]
train accuracies = []
test accuracies = []
# Loop to test different numbers of estimators
for n in num estimators:
    # Create AdaBoost model with different numbers of estimators
    ada clf =
AdaBoostClassifier(base estimator=DecisionTreeClassifier(max depth=4,
splitter='random'),
```

```
n estimators=n, random state=42)
    # Train the model on training data
    ada clf.fit(X train, y train)
    # Predict labels for training and testing data
    y train pred = ada clf.predict(X train)
    y test pred = ada clf.predict(X test)
    # Compute model accuracy
    train_accuracy = accuracy_score(y_train, y_train_pred)
    test accuracy = accuracy score(y test, y test pred)
    # Store accuracies in lists
    train accuracies.append(train accuracy)
    test accuracies.append(test accuracy)
    # Evaluate the model for training and testing data
    train f1 = f1 score(y train, y train pred, average='micro')
    test f1 = f1 score(y test, y test pred, average='micro')
    train precision = precision score(y train, y train pred,
average='micro')
    test precision = precision_score(y_test, y_test_pred,
average='micro')
    train recall = recall score(y train, y train pred,
average='micro')
    test_recall = recall_score(y_test, y_test_pred, average='micro')
    print(f'n estimators AdaBoost = {n}:')
    print(f'Train Accuracy: {train accuracy:.2f}, Test Accuracy:
{test accuracy:.2f}')
    print(f'Train Precision: {train precision:.2f}, Test Precision:
{test precision:.2f}')
    print(f'Train Recall: {train_recall:.2f}, Test Recall:
{test recall:.2f}')
    print(f'Train F1-score: {train f1:.2f}, Test F1-score:
{test f1:.2f}')
    print()
# Plot AdaBoost accuracy versus number of estimators
plt.figure(figsize=(10, 6))
plt.plot(num estimators, train accuracies, marker='o', label='Train
Accuracy')
plt.plot(num estimators, test accuracies, marker='o', label='Test
Accuracy')
plt.title('AdaBoost Accuracy vs. Number of Estimators')
plt.xlabel('Number of Estimators')
plt.ylabel('Accuracy')
```

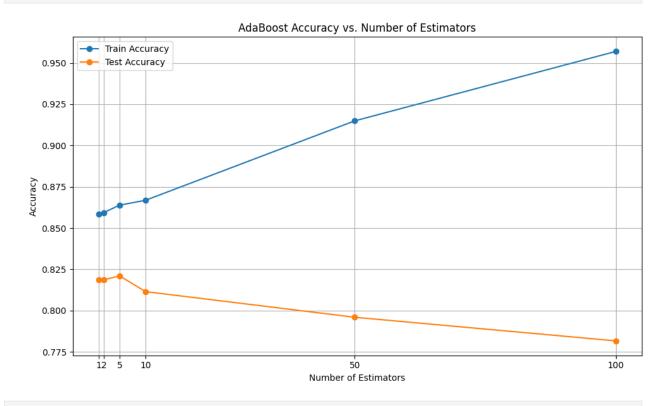
```
plt.xticks(num estimators)
plt.legend()
plt.grid(True)
plt.tight layout()
plt.show()
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/ base.py:166:
FutureWarning: `base_estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
  warnings.warn(
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/ base.py:166:
FutureWarning: `base estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
  warnings.warn(
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/ base.py:166:
FutureWarning: `base_estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
  warnings.warn(
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/_base.py:166:
FutureWarning: `base estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
  warnings.warn(
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/ base.py:166:
FutureWarning: `base estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
  warnings.warn(
n estimators AdaBoost = 1:
Train Accuracy: 0.86, Test Accuracy: 0.82
Train Precision: 0.86, Test Precision: 0.82
Train Recall: 0.86, Test Recall: 0.82
Train F1-score: 0.86, Test F1-score: 0.82
n estimators AdaBoost = 2:
Train Accuracy: 0.86, Test Accuracy: 0.82
Train Precision: 0.86, Test Precision: 0.82
Train Recall: 0.86, Test Recall: 0.82
Train F1-score: 0.86, Test F1-score: 0.82
n estimators AdaBoost = 5:
Train Accuracy: 0.86, Test Accuracy: 0.82
Train Precision: 0.86, Test Precision: 0.82
Train Recall: 0.86, Test Recall: 0.82
Train F1-score: 0.86, Test F1-score: 0.82
n estimators AdaBoost = 10:
Train Accuracy: 0.87, Test Accuracy: 0.81
Train Precision: 0.87, Test Precision: 0.81
Train Recall: 0.87, Test Recall: 0.81
```

```
Train F1-score: 0.87, Test F1-score: 0.81

n_estimators AdaBoost = 50:
Train Accuracy: 0.91, Test Accuracy: 0.80
Train Precision: 0.91, Test Precision: 0.80
Train Recall: 0.91, Test Recall: 0.80
Train F1-score: 0.91, Test F1-score: 0.80

/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/_base.py:166:
FutureWarning: base_estimator was renamed to estimator in version
1.2 and will be removed in 1.4.
    warnings.warn(

n_estimators AdaBoost = 100:
Train Accuracy: 0.96, Test Accuracy: 0.78
Train Precision: 0.96, Test Precision: 0.78
Train Recall: 0.96, Test Recall: 0.78
Train F1-score: 0.96, Test F1-score: 0.78
```



## Stacked Learner

import numpy as np
import pandas as pd

```
import matplotlib.pyplot as plt
from sklearn.tree import DecisionTreeClassifier
from sklearn.ensemble import RandomForestClassifier,
AdaBoostClassifier
from sklearn.neighbors import KNeighborsClassifier
from sklearn.model selection import train test split
from sklearn.metrics import accuracy score, precision score,
recall score, f1 score
X = df low corr.drop('TenYearCHD', axis=1)
y = df low corr['TenYearCHD']
# Split the data into training and testing sets
X_train, X_test, y_train, y_test = train_test_split(X, y,
test size=0.2, random state=42)
# Define base classifiers
base classifiers = {
    'knn': KNeighborsClassifier(n neighbors=3),
    'decision tree': DecisionTreeClassifier(max depth=4,
random state=42),
    'random forest': RandomForestClassifier(n estimators=10,
max depth=4, random state=42),
    'adaboost':
AdaBoostClassifier(base estimator=DecisionTreeClassifier(max depth=4),
n estimators=10, random state=42)
# Train base classifiers
for name, clf in base classifiers.items():
    clf.fit(X train, y train)
    y train pred = clf.predict(X train)
    y_test_pred = clf.predict(X_test)
    print(f'{name} - Train Accuracy: {accuracy score(y train,
y train pred):.2f}, Test Accuracy: {accuracy score(y test,
y test pred):.2f}')
# Create new features for the meta-classifier
meta features train = np.column stack([clf.predict(X train) for clf in
base classifiers.values()])
meta features test = np.column stack([clf.predict(X test) for clf in
base classifiers.values()])
# Define and train the meta-classifier
meta classifier = DecisionTreeClassifier(max depth=4, random state=42)
meta classifier.fit(meta features train, y train)
# Evaluate the stacked model
y train pred meta = meta classifier.predict(meta features train)
y test pred meta = meta classifier.predict(meta features test)
```

```
train accuracy = accuracy score(y train, y train pred meta)
test accuracy = accuracy score(y test, y test pred meta)
train precision = precision score(y train, y train pred meta,
average='micro')
test precision = precision score(y test, y test pred meta,
average='micro')
train_recall = recall_score(y_train, y_train_pred_meta,
average='micro')
test recall = recall score(y test, y test pred meta, average='micro')
train_f1 = f1_score(y_train, y_train_pred_meta, average='micro')
test_f1 = f1_score(y_test, y_test_pred_meta, average='micro')
print(f'Stacked Model:')
print(f'Train Accuracy: {train accuracy:.2f}, Test Accuracy:
{test accuracy:.2f}')
print(f'Train Precision: {train precision:.2f}, Test Precision:
{test precision:.2f}')
print(f'Train Recall: {train recall:.2f}, Test Recall:
{test recall:.2f}')
print(f'Train F1-score: {train f1:.2f}, Test F1-score: {test f1:.2f}')
knn - Train Accuracy: 0.89, Test Accuracy: 0.79
decision tree - Train Accuracy: 0.86, Test Accuracy: 0.82
random forest - Train Accuracy: 0.86, Test Accuracy: 0.83
adaboost - Train Accuracy: 0.88, Test Accuracy: 0.80
/usr/local/lib/python3.10/dist-packages/sklearn/ensemble/ base.py:166:
FutureWarning: `base_estimator` was renamed to `estimator` in version
1.2 and will be removed in 1.4.
 warnings.warn(
Stacked Model:
Train Accuracy: 0.89, Test Accuracy: 0.78
Train Precision: 0.89, Test Precision: 0.78
Train Recall: 0.89, Test Recall: 0.78
Train F1-score: 0.89, Test F1-score: 0.78
```