

# **AP EDA: Framingham**

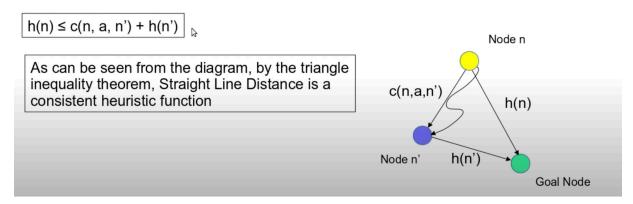
⊙ Course #	5100 - Intro Al
<b>≖</b> Task	Class Lecture
→ Parent Class	📦 *Intro to Al Syllabus, 🧠 🔥 🔥 Al FINAL PROJECT 🔥 🔥 🔥
	AP XGBoost, <a> Random Forest Exploratory Data Analysis (EDA)</a>
Ø URL	https://rajagopalvenkat.com/teaching/resources/Al/
☆ Progress	In progress
→ *Algorithms, Formulas & Equations DB*	*Artificial Intelligence, Artificial Intelligence Tutorial   Al Tutorial - GeeksforGeeks
<b>⊞</b> Week	10
■ Office Hours	Khoury Office Hours App for all office hours (online and in-person) for efficient queue management this semester.  Instructor - Raj  In-person, Meserve 303, Tuesday, 9:30 am - 11:30 am.  To schedule appointments outside of office hours, visit my appointments page.  If you decide to swing by on a whim and my office door is open, feel free to bug me
☑ Reviewed?	
Textbook File	https://artint.info/3e/html/ArtInt3e.html

#### Masters CS Northeastern DB\*

<b>⊼</b> Task	<b></b> ₩eek	• Class	Progress	■ Date	Due Date	<b>■</b> Rea
Syllabus	1	Anaconda Setup	Submitted!			
Homework	1	Problem Set 0: Background	Submitted!		@January 17, 2025	

# **Consistent Heuristic**

A heuristic h(n) is consistent (or monotonic) if, for every node n and every successor n' of n generated by any action a, the estimated cost of reaching the goal from n is no greater than the step cost of getting to n' plus the estimated cost of reaching the goal from n'



#### **Al Review**

# ▼ Exploratory Data Analysis (EDA)

"Framingham" heart disease dataset includes over 4,240 records,16 columns and 15 attributes. The goal of the dataset is to predict whether the patient has 10-year risk of future (CHD) coronary heart disease

#### **Heart Failure Attributes**

- 1. Age: age of the patient [years]
- 2. Sex: sex of the patient [M: Male, F: Female]
- 3. ChestPainType: chest pain type [TA: Typical Angina, ATA: Atypical Angina, NAP: Non-Anginal Pain, ASY: Asymptomatic]
- 4. RestingBP: resting blood pressure [mm Hg]
- 5. Cholesterol: serum cholesterol [mm/dl]
- 6. FastingBS: fasting blood sugar [1: if FastingBS > 120 mg/dl, 0: otherwise]
- RestingECG: resting electrocardiogram results [Normal: Normal, ST: having ST-T wave abnormality (T wave inversions and/or ST elevation or depression of > 0.05 mV), LVH: showing probable or definite left ventricular hypertrophy by Estes' criteria]
- 8. MaxHR: maximum heart rate achieved [Numeric value between 60 and 202]
- 9. ExerciseAngina: exercise-induced angina [Y: Yes, N: No]
- 10. Oldpeak: oldpeak = ST [Numeric value measured in depression]

- 11. ST\_Slope: the slope of the peak exercise ST segment [Up: upsloping, Flat: flat, Down: downsloping]
- 12. HeartDisease: output class [1: heart disease, 0: Normal]

sex
age
education
currentSmoker
cigsPerDay
BPMeds
prevalentStroke
prevalentHyp
diabetes
totChol
sysBP
diaBP
ВМІ
heartRate
glucose
TenYearCHD

```
male
            0
            0
age
education
             105
currentSmoker
                0
cigsPerDay
              29
BPMeds
              53
prevalentStroke 0
prevalentHyp
               0
diabetes
             0
            50
totChol
             0
sysBP
diaBP
             0
ВМІ
           19
heartRate
              1
            388
glucose
TenYearCHD
```

Framingham \*number of missing values

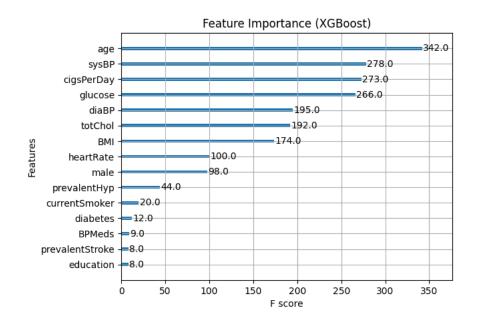
### ▼ XGBoost v1 Results @March 11, 2025 11:55 AM

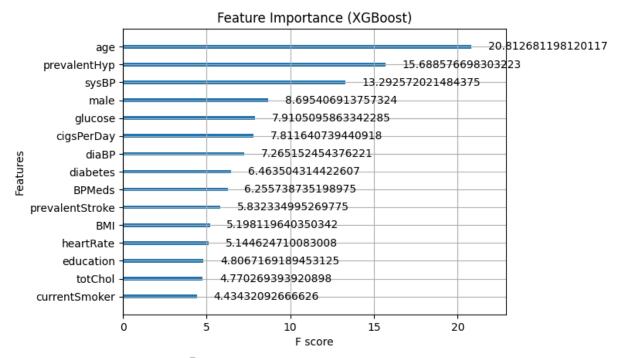
```
# 4. Train default XGBoost classifier model
#Set up basic parameters
params = {
  "objective": "binary:logistic", # Binary classification
  "eval_metric": "auc", # AUC is good for classification
  "seed": 42
}
# Train the model
model = xgb.train(params, dtrain, num_boost_round=100)
# Save model
model.save_model("xgboost_model.json")
# Define XGBoost classifier
xgb_clf = XGBClassifier(use_label_encoder=False, eval_metric="auc")
# Define hyperparameter grid
param_grid = {
  "n_estimators": [30, 40, 50, 75, 100, 150, 200, 300], # Number of trees
  "max_depth": [3, 5, 7, 8, 9, 10, 11, 12], # Tree depth
  "learning_rate": [0.01, 0.1, 0.2], # Step size shrinkage
  "subsample": [0.8, 1.0], # Percentage of samples used per tree
  "colsample_bytree": [0.8, 1.0] # Percentage of features used per tree
}
# Run GridSearchCV
grid_search = GridSearchCV(xgb_clf, param_grid, cv=3, scoring="roc_auc", verbose=2, n_jobs=-1)
grid_search.fit(X_train, y_train)
# Best parameters
print("Best parameters:", grid_search.best_params_)
# Save best model
best_model = grid_search.best_estimator_
best_model.save_model("best_xgboost_model.json")
#Fitting 3 folds for each of 768 candidates, totalling 2304 fits
Best parameters: {'colsample_bytree': 0.8, 'learning_rate': 0.01, 'max_depth': 3, 'n_estimators': 300, 'subsample': 0.8}
Accuracy: 0.8573113207547169
AUC Score: 0.7036613400616765
Confusion Matrix:
[[725 0]
[121 2]]
```

Plot

# Plot feature importance using built-in XGBoost method
xgb.plot\_importance(best\_model, importance\_type="weight") # "gain", "cover", or "weight"
plt.title("Feature Importance (XGBoost)")
plt.show()

- "weight": Number of times a feature appears in trees.
- "gain": Improvement in performance when a feature is used.
- "cover": Coverage of samples a feature influences.





▼ Logistic Regression Results @March 10, 2025

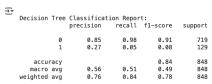
₹	Logistic Regr	ession Class		Report: f1-score	support
	0	0.85	0.99	0.92	719
	1	0.47	0.05	0.10	129
	accuracy			0.85	848
	macro avg	0.66	0.52	0.51	848
	weighted avg	0.79	0.85	0.79	848

Logistic Regression ROC-AUC: 0.6995935353796724

Logistic Regression ROC Curve (AUC = 0.6996)

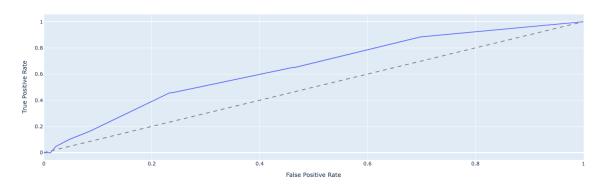


# ▼ Decision Tree Results @March 10, 2025



Decision Tree ROC-AUC: 0.6444135373203523

Decision Tree ROC Curve (AUC = 0.6444)



# **▼** Framingham Dataset

Info	Result
Total rows	4240
df.shape	(4240, 16)

### **▼** Heart Failure Attributes

### 1. Age: age of the patient [years]

- 2. Sex: sex of the patient [M: Male, F: Female]
- 3. ChestPainType: chest pain type [TA: Typical Angina, ATA: Atypical Angina, NAP: Non-Anginal Pain, ASY: Asymptomatic]
- 4. RestingBP: resting blood pressure [mm Hg]
- 5. Cholesterol: serum cholesterol [mm/dl]
- 6. FastingBS: fasting blood sugar [1: if FastingBS > 120 mg/dl, 0: otherwise]
- 7. RestingECG: resting electrocardiogram results [Normal: Normal, ST: having ST-T wave abnormality (T wave inversions and/or ST elevation or depression of > 0.05 mV), LVH: showing probable or definite left ventricular hypertrophy by Estes' criteria]
- 8. MaxHR: maximum heart rate achieved [Numeric value between 60 and 202]
- 9. ExerciseAngina: exercise-induced angina [Y: Yes, N: No]
- 10. Oldpeak: oldpeak = ST [Numeric value measured in depression]
- 11. ST\_Slope: the slope of the peak exercise ST segment [Up: upsloping, Flat: flat, Down: downsloping]
- 12. HeartDisease: output class [1: heart disease, 0: Normal]

### ▼ \*After converting all features to numeric

<class< th=""><th>'pandas.core.frame.DataFrame'&gt;</th><th></th><th></th></class<>	'pandas.core.frame.DataFrame'>		
Index:	4240	entries	
Data	columns	(total	
#	Column	Non-Null	
0	male	4240	Int64
1	age	4240	Int64
2	education	4240	Int64
3	currentSmoker	4240	Int64
4	cigsPerDay	4240	Int64
5	BPMeds	4240	Int64
6	prevalentStroke	4240	Int64
7	prevalentHyp	4240	Int64
8	diabetes	4240	Int64
9	totChol	4240	Int64
10	sysBP	4240	Float64
11	diaBP	4240	Float64
12	ВМІ	4240	Float64
13	heartRate	4240	Int64
14	glucose	4240	Int64
15	TenYearCHD	4240	Int64
dtypes:	Float64(3)		
memory	usage:	629.4	

df.info()	#	Column	Non-Null	Count	Dtype
	0	male	4240	non-null	int64
	1	age	4240	non-null	int64

	2	education	4135	non-null	float64
	3	currentSmoker	4240	non-null	int64
	4	cigsPerDay	4211	non-null	float64
	5	BPMeds	4187	non-null	float64
	6	prevalentStroke	4240	non-null	int64
	7	prevalentHyp	4240	non-null	int64
	8	diabetes	4240	non-null	int64
	9	totChol	4190	non-null	float64
	10	sysBP	4240	non-null	float64
	11	diaBP	4240	non-null	float64
	12	ВМІ	4221	non-null	float64
	13	heartRate	4239	non-null	float64
	14	glucose	3852	non-null	float64
	15	TenYearCHD	4240	non-null	int64
dtypes:	float64(9),	int64(7)			
memory	usage:	530.1	KB		

#### root

- |-- male: string (nullable = true)
- -- age: string (nullable = true)
- -- education: string (nullable = true)
- |-- currentSmoker: string (nullable = true)
- -- cigsPerDay: string (nullable = true)
- |-- BPMeds: string (nullable = true)
- |-- prevalentStroke: string (nullable = true)
- |-- prevalentHyp: string (nullable = true)
- -- diabetes: string (nullable = true)
- -- totChol: string (nullable = true)
- -- sysBP: string (nullable = true)
- -- diaBP: string (nullable = true)
- |-- BMI: string (nullable = true)
- -- heartRate: string (nullable = true)
- |-- glucose: string (nullable = true)
- -- TenYearCHD: string (nullable = true)

import pandas as pd import numpy as np import matplotlib.pylab as plt import seaborn as sns plt.style.use('ggplot') pd.set\_option('max\_columns', 200)

```
→ <class 'pandas.core.frame.DataFrame'>
    RangeIndex: 4240 entries, 0 to 4239
    Data columns (total 16 columns):
         Column
                         Non-Null Count
                                          Dtype
     #
     0
                          4240 non-null
         male
                                          int64
                          4240 non-null
                                          int64
     1
         age
     2
         education
                          4135 non-null
                                          float64
     3
         currentSmoker
                          4240 non-null
                                          int64
                                          float64
         cigsPerDay
                          4211 non-null
         BPMeds
                          4187 non-null
                                          float64
         prevalentStroke 4240 non-null
                                          int64
         prevalentHyp
                          4240 non-null
                                          int64
     8
         diabetes
                          4240 non-null
                                          int64
                          4190 non-null
                                          float64
         totChol
     10
         sysBP
                          4240 non-null
                                          float64
         diaBP
                          4240 non-null
                                          float64
     11
                                          float64
     12
         BMI
                          4221 non-null
     13
         heartRate
                          4239 non-null
                                          float64
     14
                          3852 non-null
                                          float64
         glucose
     15 TenYearCHD
                          4240 non-null
                                          int64
    dtypes: float64(9), int64(7)
```

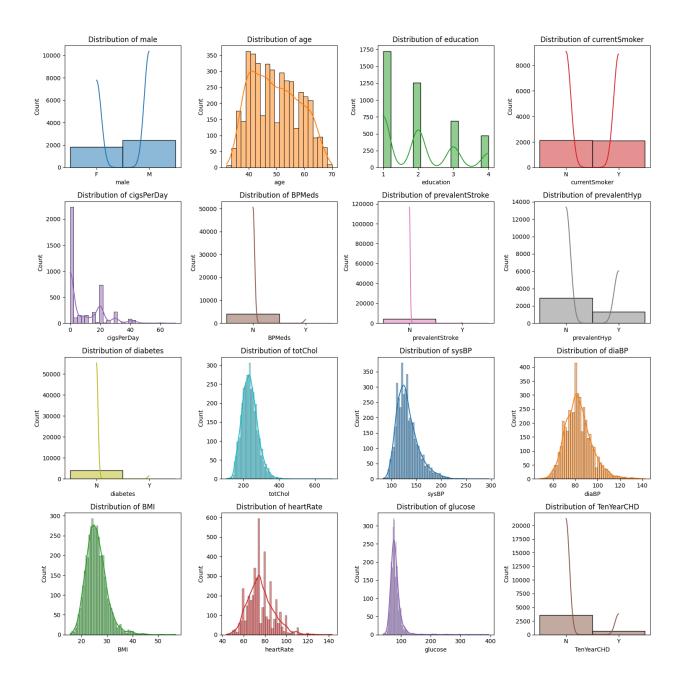
```
'male', 'age', 'education', 'currentSmoker', 'cigsPerDay', 'BPMeds',
   'prevalentStroke', 'prevalentHyp', 'diabetes', 'totChol', 'sysBP',
   'diaBP', 'BMI', 'heartRate', 'glucose', 'TenYearCHD'
```

# df\_visualize

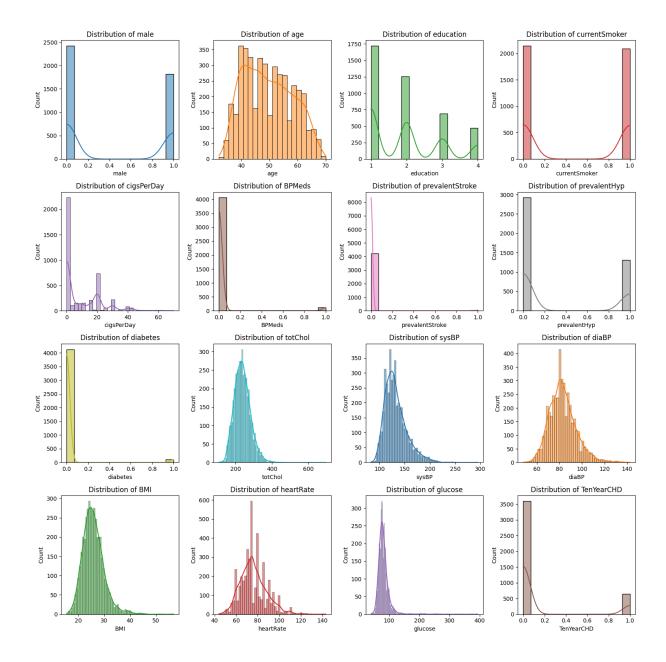
[5]

df.info()

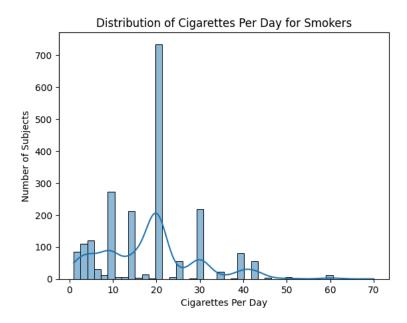
memory usage: 530.1 KB



# ▼ df original



▼ Smokers: 2095



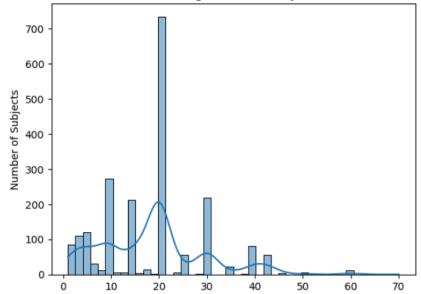
```
df_smoker = df_visualize[df_visualize['currentSmoker'] == 'Y']
df_smoker.info()

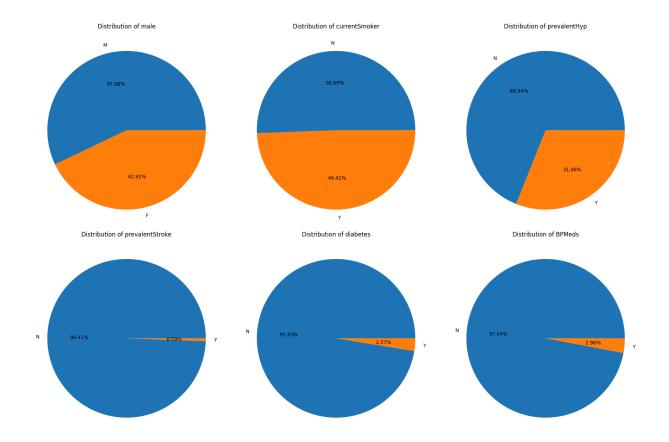
# Plot the distribution of cigsPerDay for those records
sns.histplot(df_smoker['cigsPerDay'], kde=True)
plt.title("Distribution of Cigarettes Per Day for Smokers")
plt.xlabel("Cigarettes Per Day")
plt.ylabel("Number of Subjects")
plt.show()
```

<class 'pandas.core.frame.DataFrame'>
Index: 2095 entries, 2 to 4239
Data columns (total 16 columns):

#	Column	Non-Null Count	Dtype	
0	male	2095 non-null	object	
1	age	2095 non-null	int64	
2	education	2046 non-null	float64	
3	currentSmoker	2095 non-null	object	
4	cigsPerDay	2066 non-null	float64	
5	BPMeds	2072 non-null	float64	
6	prevalentStroke	2095 non-null	object	
7	prevalentHyp	2095 non-null	object	
8	diabetes	2095 non-null	object	
9	totChol	2064 non-null	float64	
10	sysBP	2095 non-null	float64	
11	diaBP	2095 non-null	float64	
12	BMI	2088 non-null	float64	
13	heartRate	2094 non-null	float64	
14	glucose	1890 non-null	float64	
15	TenYearCHD	2095 non-null	object	
dtypes: float64(9), int64(1), object(6)				
memo	ry usage: 278.2+	KB		

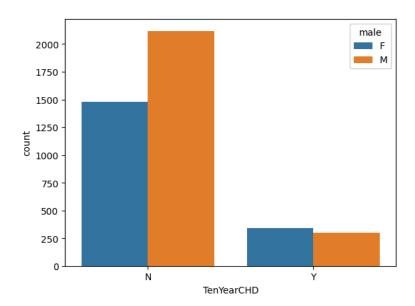
## Distribution of Cigarettes Per Day for Smokers

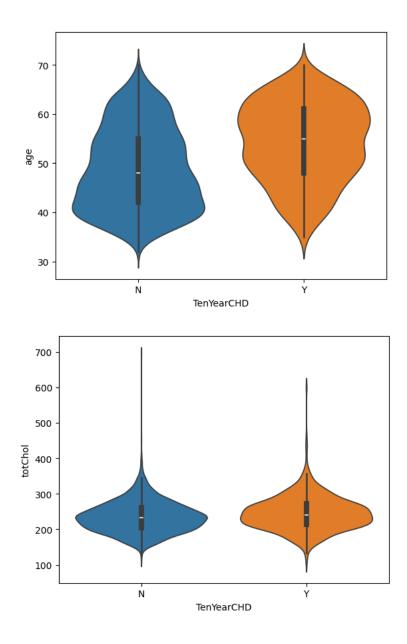




### **Violin Plot**

sns.violinplot(x=df\_visualize["age"])
sns.countplot(x=df\_visualize["TenYearCHD"],hue=df\_visualize["male"])





# **Logistic Regression**

# 1. Data Encoding

from sklearn.preprocessing import LabelEncoder, StandardScaler from sklearn.model\_selection import train\_test\_split

# Encode all columns with LabelEncoder (this converts categorical values to numeric) df = df.apply(LabelEncoder().fit\_transform)

# Split data into features (X) and target (y)

```
X = df.drop(columns=['TenYearCHD']) # All columns except the target
y = df['TenYearCHD'] # Target column
```

#### 2. Logistic Regression Model

```
# Split the data into train and test sets (stratified to preserve target class proportions)

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42, stratify=y)

# Standardize the features for logistic regression (helps the model converge)

scaler = StandardScaler()

X_train_scaled = scaler.fit_transform(X_train)

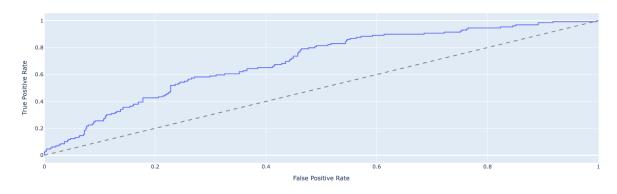
X_test_scaled = scaler.transform(X_test)
```

#### 3. Training, Predicting & Evaluating Logistic Regression

```
from sklearn.linear_model import LogisticRegression
from sklearn.metrics import classification_report, roc_curve, auc
import plotly.express as px
# Initialize and train logistic regression with increased iterations if needed
log_model = LogisticRegression(max_iter=1000, random_state=42)
log_model.fit(X_train_scaled, y_train)
# Make predictions on the test set
y_pred_log = log_model.predict(X_test_scaled)
# Print the classification report
print("Logistic Regression Classification Report:")
print(classification_report(y_test, y_pred_log))
# ROC-AUC Calculation for Logistic Regression
fpr, tpr, _ = roc_curve(y_test, log_model.predict_proba(X_test_scaled)[:, 1])
roc_auc = auc(fpr, tpr)
print("Logistic Regression ROC-AUC:", roc_auc)
# Plot ROC Curve using Plotly Express
roc_data = pd.DataFrame({'False Positive Rate': fpr, 'True Positive Rate': tpr})
fig = px.line(roc_data, x='False Positive Rate', y='True Positive Rate',
        title=f'Logistic Regression ROC Curve (AUC = {roc_auc:.4f})',
        labels={'False Positive Rate': 'False Positive Rate', 'True Positive Rate': 'True Positive Rate'})
# Add the reference line for a random classifier
fig.add_shape(type='line', x0=0, y0=0, x1=1, y1=1, line=dict(dash='dash', color='grey'))
fig.show()
Logistic Regression Classification Report:
        precision recall f1-score support
```

```
0.85 0.99 0.92
                              719
     0
         0.47
                0.05 0.10
                             129
                              848
 accuracy
                       0.85
                    0.52
                          0.51
 macro avg
             0.66
                                 848
weighted avg
              0.79
                    0.85
                            0.79
                                   848
Logistic Regression ROC-AUC: 0.6995935353796724
```

Logistic Regression ROC Curve (AUC = 0.6996)



#### **Decision Tree**

```
from sklearn.tree import DecisionTreeClassifier
from sklearn.metrics import classification_report, roc_curve, auc

# Initialize and train the Decision Tree Classifier (limiting max_depth to control overfitting)
dt_model = DecisionTreeClassifier(max_depth=4, random_state=42)
dt_model.fit(X_train, y_train)

# Make predictions on the test set
y_pred_dt = dt_model.predict(X_test)

# Print the classification report
print("\nDecision Tree Classification Report:")
print(classification_report(y_test, y_pred_dt))

# ROC-AUC Calculation for Decision Tree (if predict_proba is available)
if hasattr(dt_model, "predict_proba"):
    fpr_dt, tpr_dt, _ = roc_curve(y_test, dt_model.predict_proba(X_test)[:, 1])
    roc_auc_dt = auc(fpr_dt, tpr_dt)
    print("Decision Tree ROC-AUC:", roc_auc_dt)
```

```
# Optional: Plot ROC Curve using Plotly Express for Decision Tree

roc_data_dt = pd.DataFrame({'False Positive Rate': fpr_dt, 'True Positive Rate': tpr_dt})

fig_dt = px.line(roc_data_dt, x='False Positive Rate', y='True Positive Rate',

title=f'Decision Tree ROC Curve (AUC = {roc_auc_dt:.4f})',

labels={'False Positive Rate': 'False Positive Rate', 'True Positive Rate': 'True Positive Rate'})

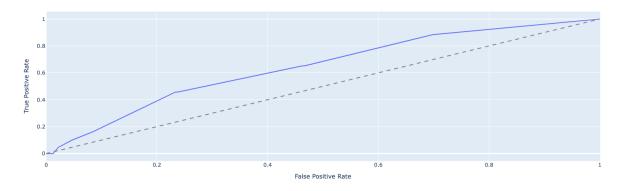
fig_dt.add_shape(type='line', x0=0, y0=0, x1=1, y1=1, line=dict(dash='dash', color='grey'))

fig_dt.show()
```

```
| Decision Tree | Classification | Reports | R
```

Decision Tree ROC-AUC: 0.6444135373203523

Decision Tree ROC Curve (AUC = 0.6444)



```
Decision Tree Classification Report:
       precision recall f1-score support
          0.85
               0.98
                       0.91
                              719
         0.27
                0.05 0.08
                              129
                       0.84
                              848
  accuracy
 macro avg
             0.56
                    0.51 0.49
                                  848
weighted avg
             0.76 0.84
Decision Tree ROC-AUC: 0.6444135373203523
```