Einleitung

Der vorliegende Report dient zur Beschreibung des Projekts im Rahmen der Data Exploration Vorlesung. Die Abgabe umfasst ein GitHub Repositpory mit dem erstellten Code und dem Report in Form eines Jupyter Notebooks. Ziel dieses Projekts ist es anhand des, im Folgenden beschriebenen Datensatzes, eine exlorative Datenanalyse zu betreiben und ein Machine Learning Modell zu entwickeln, das zuverlässige Ergebnisse liefert.

Requirements

Um alle benötigten Requirements zu installieren, wird der folgende Befehl verwendet.

In []: ! pip install -r requirements.txt

```
Requirement already satisfied: appnope==0.1.4 in ./venv/lib/python3.9/site
-packages (from -r requirements.txt (line 1)) (0.1.4)
Requirement already satisfied: asttokens==2.4.1 in ./venv/lib/python3.9/si
te-packages (from -r requirements.txt (line 2)) (2.4.1)
Requirement already satisfied: comm==0.2.1 in ./venv/lib/python3.9/site-pa
ckages (from -r requirements.txt (line 3)) (0.2.1)
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Requirement already satisfied: cycler==0.12.1 in ./venv/lib/python3.9/site
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-packages (from -r requirements.txt (line 6)) (1.8.1)
Requirement already satisfied: decorator==5.1.1 in ./venv/lib/python3.9/si
te-packages (from -r requirements.txt (line 7)) (5.1.1)
Requirement already satisfied: exceptiongroup==1.2.0 in ./venv/lib/python
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Requirement already satisfied: executing==2.0.1 in ./venv/lib/python3.9/si
te-packages (from -r requirements.txt (line 9)) (2.0.1)
Requirement already satisfied: fonttools==4.49.0 in ./venv/lib/python3.9/s
ite-packages (from -r requirements.txt (line 10)) (4.49.0)
Requirement already satisfied: importlib-metadata==7.0.1 in ./venv/lib/pyt
hon3.9/site-packages (from -r requirements.txt (line 11)) (7.0.1)
Requirement already satisfied: importlib-resources==6.1.1 in ./venv/lib/py
thon3.9/site-packages (from -r requirements.txt (line 12)) (6.1.1)
Requirement already satisfied: ipykernel==6.29.2 in ./venv/lib/python3.9/s
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e-packages (from -r requirements.txt (line 14)) (8.18.1)
Requirement already satisfied: jedi==0.19.1 in ./venv/lib/python3.9/site-p
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Requirement already satisfied: joblib==1.3.2 in ./venv/lib/python3.9/site-
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Requirement already satisfied: jupyter_client==8.6.0 in ./venv/lib/python
3.9/site-packages (from -r requirements.txt (line 17)) (8.6.0)
Requirement already satisfied: jupyter_core==5.7.1 in ./venv/lib/python3.
9/site-packages (from -r requirements.txt (line 18)) (5.7.1)
Requirement already satisfied: kiwisolver==1.4.5 in ./venv/lib/python3.9/s
ite-packages (from -r requirements.txt (line 19)) (1.4.5)
Requirement already satisfied: matplotlib==3.8.3 in ./venv/lib/python3.9/s
ite-packages (from -r requirements.txt (line 20)) (3.8.3)
Requirement already satisfied: matplotlib-inline==0.1.6 in ./venv/lib/pyth
on3.9/site-packages (from -r requirements.txt (line 21)) (0.1.6)
Requirement already satisfied: nest-asyncio==1.6.0 in ./venv/lib/python3.
9/site-packages (from -r requirements.txt (line 22)) (1.6.0)
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Requirement already satisfied: packaging==23.2 in ./venv/lib/python3.9/sit
e-packages (from -r requirements.txt (line 24)) (23.2)
Requirement already satisfied: pandas==2.2.0 in ./venv/lib/python3.9/site-
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Requirement already satisfied: pexpect==4.9.0 in ./venv/lib/python3.9/site
-packages (from -r requirements.txt (line 27)) (4.9.0)
Requirement already satisfied: pillow==10.2.0 in ./venv/lib/python3.9/site
-packages (from -r requirements.txt (line 28)) (10.2.0)
Requirement already satisfied: platformdirs==4.2.0 in ./venv/lib/python3.
9/site-packages (from -r requirements.txt (line 29)) (4.2.0)
Requirement already satisfied: prompt-toolkit==3.0.43 in ./venv/lib/python
3.9/site-packages (from -r requirements.txt (line 30)) (3.0.43)
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Requirement already satisfied: psutil==5.9.8 in ./venv/lib/python3.9/site-
       packages (from -r requirements.txt (line 31)) (5.9.8)
       Requirement already satisfied: ptyprocess==0.7.0 in ./venv/lib/python3.9/s
       ite-packages (from -r requirements.txt (line 32)) (0.7.0)
       Requirement already satisfied: pure-eval==0.2.2 in ./venv/lib/python3.9/si
       te-packages (from -r requirements.txt (line 33)) (0.2.2)
       Requirement already satisfied: Pygments==2.17.2 in ./venv/lib/python3.9/si
       te-packages (from -r requirements.txt (line 34)) (2.17.2)
       Requirement already satisfied: pyparsing==3.1.1 in ./venv/lib/python3.9/si
       te-packages (from -r requirements.txt (line 35)) (3.1.1)
       Requirement already satisfied: python-dateutil==2.8.2 in ./venv/lib/python
       3.9/site-packages (from -r requirements.txt (line 36)) (2.8.2)
       Requirement already satisfied: pytz==2024.1 in ./venv/lib/python3.9/site-p
       ackages (from -r requirements.txt (line 37)) (2024.1)
       Requirement already satisfied: pyzmq==25.1.2 in ./venv/lib/python3.9/site-
       packages (from -r requirements.txt (line 38)) (25.1.2)
       Requirement already satisfied: scikit-learn==1.4.1.post1 in ./venv/lib/pyt
       hon3.9/site-packages (from -r requirements.txt (line 39)) (1.4.1.post1)
       Requirement already satisfied: scipy==1.12.0 in ./venv/lib/python3.9/site-
       packages (from -r requirements.txt (line 40)) (1.12.0)
       Requirement already satisfied: seaborn==0.13.2 in ./venv/lib/python3.9/sit
       e-packages (from -r requirements.txt (line 41)) (0.13.2)
       Requirement already satisfied: six==1.16.0 in ./venv/lib/python3.9/site-pa
       ckages (from -r requirements.txt (line 42)) (1.16.0)
       Requirement already satisfied: stack-data==0.6.3 in ./venv/lib/python3.9/s
       ite-packages (from -r requirements.txt (line 43)) (0.6.3)
       Requirement already satisfied: threadpoolctl==3.3.0 in ./venv/lib/python3.
       9/site-packages (from -r requirements.txt (line 44)) (3.3.0)
       Requirement already satisfied: tornado==6.4 in ./venv/lib/python3.9/site-p
       ackages (from -r requirements.txt (line 45)) (6.4)
       Requirement already satisfied: traitlets==5.14.1 in ./venv/lib/python3.9/s
       ite-packages (from -r requirements.txt (line 46)) (5.14.1)
       Requirement already satisfied: typing_extensions==4.9.0 in ./venv/lib/pyth
       on3.9/site-packages (from -r requirements.txt (line 47)) (4.9.0)
       Requirement already satisfied: tzdata==2024.1 in ./venv/lib/python3.9/site
       -packages (from -r requirements.txt (line 48)) (2024.1)
       Requirement already satisfied: wcwidth==0.2.13 in ./venv/lib/python3.9/sit
       e-packages (from -r requirements.txt (line 49)) (0.2.13)
       Requirement already satisfied: zipp==3.17.0 in ./venv/lib/python3.9/site-p
       ackages (from -r requirements.txt (line 50)) (3.17.0)
In []: # importing all required libraries
```

```
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
import numpy as np

# machine learning libraries
from sklearn.ensemble import RandomForestClassifier
from sklearn.linear_model import LogisticRegression
from sklearn.metrics import accuracy_score, confusion_matrix, classificat
from sklearn.model_selection import GridSearchCV
from sklearn.metrics import roc_auc_score
from sklearn.metrics import roc_curve
from sklearn.svm import SVC
from sklearn.model_selection import train_test_split

%matplotlib inline
```

Data Quality Check & Data Characterization

Die verwendeten Daten

Bei den verwendeten Daten handelt es sich um einen Kaggle Datensatz (https://www.kaggle.com/datasets/fedesoriano/heart-failure-prediction/data; letzter Abruf: 04.04.2024). Der Datensatz enthält Informationen von 918 Patienten und umfasst zwölf verschiedene Merkmale, darunter de- mografische Angaben wie Alter und Geschlecht, klinische Messungen wie Ruheblutdruck und maximale Herzfrequenz, sowie Informationen zu Symptomen wie Brustschmerzen und zuvor dia- gnostizierten Herzkrankheiten. Die Daten widerspiegeln auch medizinischen Tests wie Ruhe- Elektrokardiogrammen und Belastungsuntersuchungen.

```
# defining path in which the data is stored
In [ ]:
         data = "data/heart.csv"
In [ ]: # reading the data
         df = pd.read csv(data)
         df
Out[]:
                                                    Cholesterol FastingBS
                         ChestPainType RestingBP
                                                                            RestingECG
               Age Sex
            0
                40
                                               140
                                                            289
                                                                         0
                                                                                 Normal
                      Μ
                                    ATA
            1
                49
                      F
                                    NAP
                                                160
                                                            180
                                                                         0
                                                                                 Normal
                37
                      Μ
                                    ATA
                                                130
                                                            283
                                                                                     ST
            3
                48
                      F
                                                138
                                                            214
                                                                         0
                                    ASY
                                                                                 Normal
            4
                54
                                    NAP
                                                150
                                                            195
                                                                         0
                                                                                 Normal
                      Μ
         913
                45
                                     TA
                                                110
                                                            264
                                                                         0
                                                                                 Normal
         914
                68
                      М
                                    ASY
                                               144
                                                            193
                                                                          1
                                                                                 Normal
                                    ASY
         915
                57
                                               130
                                                            131
                                                                         0
                                                                                 Normal
                      Μ
                                                                                    LVH
         916
                57
                      F
                                    ATA
                                                130
                                                            236
                                                                         0
         917
                38
                                    NAP
                                               138
                                                            175
                                                                         0
                                                                                 Normal
```

918 rows × 12 columns

Beschreibung der Attribute:

- Age: Alter des Patienten [Jahre]
- Sex: Geschlecht des Patienten [M: Männlich, F: Weiblich]

 ChestPainType: Brustschmerztyp [TA: Typische Angina, ATA: Atypische Angina, NAP: Nicht-Anginaler Schmerz, ASY: Asymptomatisch]

- RestingBP: Ruheblutdruck [mm Hg]
- Cholesterol: Serumcholesterin [mm/dl]
- FastingBS: Nüchternblutzucker [1: Wenn Nüchternblutzucker > 120 mg/dl, 0: Ansonsten]
- RestingECG: Ruheelektrokardiogrammergebnisse [Normal: Normal, ST: Mit ST-T-Wellen-Abnormalitäten (T-Wellen-Inversionen und/oder ST-Hebungen oder Senkungen von > 0,05 mV), LVH: Zeigt wahrscheinliche oder definitive linksventrikuläre Hypertrophie nach Estes-Kriterien]
- MaxHR: Maximale erreichte Herzfrequenz [Numerischer Wert zwischen 60 und 202]
- ExerciseAngina: Belastungsinduzierte Angina [J: Ja, N: Nein]
- Oldpeak: ST-Depression = ST [Numerischer Wert gemessen in Depression]
- ST_Slope: Die Steigung des Spitzen-Übungs-ST-Segments [Up: Aufsteigend, Flat: Flach, Down: Absteigend]
- HeartDisease: Ausgabeklasse [1: Herzkrankheit, 0: Normal]

```
In []: # using the pandas method "describe()"" to get a describtion of the datas
# ".T" transposes the dataframe (rows and columns are switched)
df.describe().T
```

Out[]:		count	mean	std	min	25%	50%	75%	max
	Age	918.0	53.510893	9.432617	28.0	47.00	54.0	60.0	77.0
	RestingBP	918.0	132.396514	18.514154	0.0	120.00	130.0	140.0	200.0
	Cholesterol	918.0	198.799564	109.384145	0.0	173.25	223.0	267.0	603.0
	FastingBS	918.0	0.233115	0.423046	0.0	0.00	0.0	0.0	1.0
	MaxHR	918.0	136.809368	25.460334	60.0	120.00	138.0	156.0	202.0
	Oldpeak	918.0	0.887364	1.066570	-2.6	0.00	0.6	1.5	6.2
	HeartDisease	918.0	0.553377	0.497414	0.0	0.00	1.0	1.0	1.0

Bereits nachdem man sich die Beschreibung des Datensatzes anschaut, kann man feststellen, dass die SPalten "Cholesterol" und "RestingBP" unerwartete minimal Werte aufweisen (ruhe Puls und Cholisterinspiegel können keine Werte von 0 annehmen).

```
In []: # count null values
  null_values_count = (df['RestingBP'] == 0).sum()
  print("Anzahl der Nullwerte in der Spalte 'RestingBP':", null_values_coun
```

Anzahl der Nullwerte in der Spalte 'RestingBP': 1

```
In [ ]: # delete the only patient with the null value in RestingBP
df = df[df['RestingBP'] != 0]
```

Da es nur bei einem Patienten eine vermutliche Fehlmessung gab, wird dieser Patient aus dem Datensatz gelöscht.

```
In [ ]: # count null values
        null_values_count = (df['Cholesterol'] == 0).sum()
        print("Anzahl der Nullwerte in der Spalte 'Cholesterol':", null_values_co
       Anzahl der Nullwerte in der Spalte 'Cholesterol': 171
        Leider weisen dennoch 171 Patienten bei Cholesterol den Wert 0 auf. Dies war bei der
        initialen explorativen Datenanalyse nicht auf den ersten Blick ersichtlich. Da das
        löschen von 171 Einträgen problematisch ist, wird in den fehldenen Stellen der
        durschnittliche Cholesterol Wert des Datensatzes eingesetzt. Somit sollen
        erheblichere Verfälschungen im Machine Learning Model im nachhinein vermieden
        werden.
In [ ]: # we don't wont the 0 values, when calculating the mean value
        df_cleaned = df[df['Cholesterol'] != 0]
        # calculate mean value
        average chol = round(df cleaned['Cholesterol'].mean())
        print("Durchschnittlicher Cholesterinspiegel nach Entfernen von Nullwerte
       Durchschnittlicher Cholesterinspiegel nach Entfernen von Nullwerten (ohne
       Nachkommastellen): 245
In []: # replace 0 values with the mean value
        df.loc[df['Cholesterol'] == 0, 'Cholesterol'] = average_chol
In [ ]: # check if the anomaly still exists
        df["Cholesterol"].min()
Out[]: 85
In []: # checking for missung values in the dataframe
        missing_values = df.isnull().sum()
        missing_values
Out[]: Age
                            0
                            0
         Sex
         ChestPainType
                            0
         RestingBP
                            0
         Cholesterol
                            0
         FastingBS
         RestingECG
                            0
         MaxHR
                            0
         ExerciseAngina
                            0
         0ldpeak
         ST_Slope
                            0
         HeartDisease
                            0
         dtype: int64
In [ ]: # checking for duplicated rows in the dataframe
        duplicates = df.duplicated().sum()
        duplicates
```

Out[]: 0

```
In [ ]: # determining unique values of categorial columns in the dataframe
        categorical_columns = ['Sex', 'ChestPainType', 'RestingECG', 'ExerciseAng
        for col in categorical_columns:
            unique_values = df[col].unique()
            print(f"Eindeutige Werte für {col}:")
            print(unique_values)
       Eindeutige Werte für Sex:
       ['M' 'F']
       Eindeutige Werte für ChestPainType:
       ['ATA' 'NAP' 'ASY' 'TA']
       Eindeutige Werte für RestingECG:
       ['Normal' 'ST' 'LVH']
       Eindeutige Werte für ExerciseAngina:
       ['N' 'Y']
       Eindeutige Werte für ST_Slope:
       ['Up' 'Flat' 'Down']
In [ ]: # get dataframe info
        df.info()
       <class 'pandas.core.frame.DataFrame'>
       Index: 917 entries, 0 to 917
       Data columns (total 12 columns):
        #
                           Non-Null Count Dtype
            Column
                            917 non-null
        0
           Age
                                            int64
        1
            Sex
                            917 non-null
                                            object
        2
           ChestPainType
                          917 non-null
                                            object
        3
           RestingBP
                            917 non-null
                                            int64
          Cholesterol
        4
                            917 non-null
                                           int64
        5
          FastingBS
                            917 non-null
                                            int64
                                          object
        6
           RestingECG
                            917 non-null
        7
                            917 non-null
           MaxHR
                                            int64
        8
            ExerciseAngina 917 non-null
                                            object
        9
            0ldpeak
                            917 non-null
                                            float64
        10 ST_Slope
                            917 non-null
                                            object
        11 HeartDisease
                            917 non-null
                                            int64
       dtypes: float64(1), int64(6), object(5)
       memory usage: 93.1+ KB
In []: # getting the highest values of each column (categorial columns may be ig
        df.max()
Out[]: Age
                            77
        Sex
                            Μ
        {\tt ChestPainType}
                           TΑ
        RestingBP
                           200
        Cholesterol
                           603
        FastingBS
                            1
                           ST
        RestingECG
        MaxHR
                           202
                            Υ
        ExerciseAngina
        0ldpeak
                           6.2
        ST_Slope
                           Up
        HeartDisease
                            1
        dtype: object
In [ ]:
       # same goes for this but for minimal values
        df.min()
```

```
28
Out[]: Age
         Sex
                              F
                            ASY
         ChestPainType
        RestingBP
                             80
        Cholesterol
                             85
        FastingBS
                              0
        RestingECG
                            LVH
        MaxHR
                             60
        ExerciseAngina
                              Ν
        0ldpeak
                           -2.6
        ST Slope
                           Down
        HeartDisease
                              0
        dtype: object
In [ ]: # check how many unique elements the dataset contains in each column
        df.nunique()
Out[]: Age
                            50
                             2
         Sex
        ChestPainType
                             4
        RestingBP
                            66
         Cholesterol
                           221
         FastingBS
                             2
        RestingECG
                             3
        MaxHR
                           119
        ExerciseAngina
                             2
                            53
        0ldpeak
        ST Slope
                             3
                             2
        HeartDisease
        dtype: int64
```

Die Analyse zur Datenqualität liefert auf den ersten Blick, bis auf die 2 Anomalien, kaum Mängel, da es keine fehlenden Einträge oder duplizierte Zeilen gibt. Auch die Spalten mit den kategorischen Werten liefern saubere und "aufgeräumte" Werte. Der Datensatz ist im Allgemeinen sehr gut gepflegt.

Exploratory Data Analysis

Im folgenden werden die Daten analysiert und statistische Verteilungen und Merkmale, sowie Anhängigkeiten zwischen verschiedenen Attributen werden grafisch aufgezeigt.

```
In []: # visualize disease distribution in the dataset

colors_red_green = ["#9aff9a", "#ff3030"]

sns.countplot(x='HeartDisease', data=df, palette=colors_red_green)

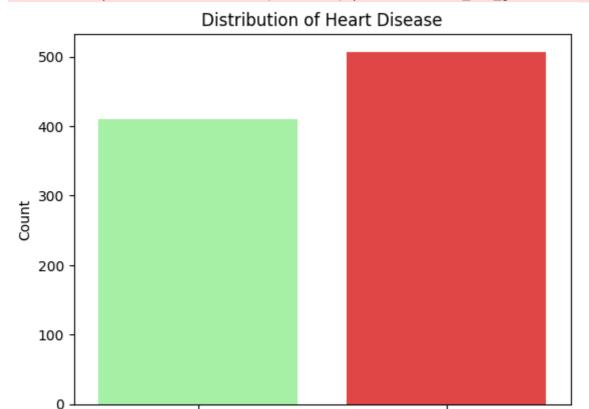
plt.xlabel('Heart Disease')
plt.ylabel('Count')
plt.title('Distribution of Heart Disease')

plt.show()
```

/var/folders/3l/_xvv3581559_krvl1r82px5w0000gn/T/ipykernel_96307/182032164 9.py:5: FutureWarning:

Passing `palette` without assigning `hue` is deprecated and will be remove d in v0.14.0. Assign the `x` variable to `hue` and set `legend=False` for the same effect.

sns.countplot(x='HeartDisease', data=df, palette=colors_red_green)



```
In [ ]: heart_disease_distribution = df['HeartDisease'].value_counts()
heart_disease_distribution
```

Heart Disease

1

0

Out[]: HeartDisease 1 507 0 410

Name: count, dtype: int64

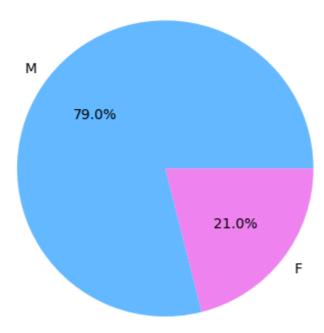
Die erste Visualisierung zeigt die Verteilung zwischen gesunden und kranken Patienten. Es ist eine leichte Inbalance der Werte vorhanden. Nach Absprache mit dem Dozenten kann diese Aufgrund ihrer leichten Ausprägung in diesem Fall ignoriert werden. Der Datensatz enthält 98 mehr betroffene als gesunde Patienten.

```
In []: distribution = df["Sex"].value_counts()
    colors = ['#63b8ff', '#ee82ee']

plt.title("Distribution of Sex")

plt.pie(distribution, labels=distribution.index, colors=colors, autopct='
    plt.show()
```

Distribution of Sex



Dieses Kuchendiagramm zeigt die Verteilung der Geschlächter in den Daten. 79% der Patienten sind männlich und 21% sind weiblich.

```
In [ ]: fig = plt.figure(figsize=(12, 6))
        gs = fig.add_gridspec(1, 3, width_ratios=[2, 1, 1])
        # total age distribution
        ax1 = fig.add_subplot(gs[0])
        sns.boxplot(x=df["Age"], ax=ax1, color='#5c5c5c')
        ax1.set_title('total age distribution')
        # female age distribution
        ax2 = fig.add_subplot(gs[1])
        sns.boxplot(x='Sex', y='Age', data=df[df['Sex'] == 'F'], ax=ax2, palette=
        ax2.set_title('female age distribution')
        # male age distribution
        ax3 = fig.add_subplot(gs[2])
        sns.boxplot(x='Sex', y='Age', data=df[df['Sex'] == 'M'], ax=ax3, palette=
        ax3.set_title('male age distribution')
        plt.tight_layout()
        plt.show()
```

/var/folders/3l/_xvv3581559_krvl1r82px5w0000gn/T/ipykernel_96307/267784640 5.py:11: FutureWarning:

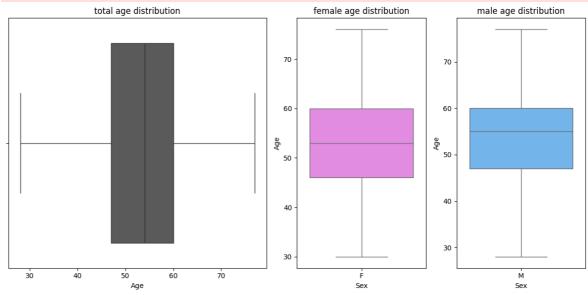
Passing `palette` without assigning `hue` is deprecated and will be remove d in v0.14.0. Assign the `x` variable to `hue` and set `legend=False` for the same effect.

sns.boxplot(x='Sex', y='Age', data=df[df['Sex'] == 'F'], ax=ax2, palette
=['#ee82ee'])

/var/folders/3l/_xvv3581559_krvl1r82px5w0000gn/T/ipykernel_96307/267784640
5.py:16: FutureWarning:

Passing `palette` without assigning `hue` is deprecated and will be remove d in v0.14.0. Assign the `x` variable to `hue` and set `legend=False` for the same effect.

sns.boxplot(x='Sex', y='Age', data=df[df['Sex'] == 'M'], ax=ax3, palette
=['#63b8ff'])



Diese Boxplots zeigen die gesamte Altersverteilung sowie die Verteilung pro Geschlecht. Die Altersspanne liegt im Durschschnitt zwischen 48 und 60 Jahren.

```
In []: counts = df["Age"].value_counts()
    print("Counts for Age:")
    print(counts)
```

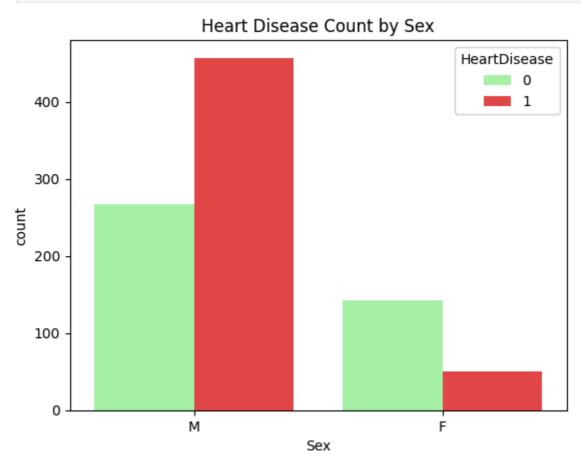
```
Counts for Age:
Age
54
      51
58
      42
55
      40
56
      38
57
       38
52
      36
51
      35
59
       35
62
      35
53
      33
      32
60
48
      31
61
      31
63
      30
50
      25
46
      24
41
      24
43
      24
64
      22
65
      21
49
      21
47
      19
44
       19
42
       18
45
      18
38
       16
67
       15
39
      15
66
      13
69
       13
40
      13
35
      11
37
       11
68
       10
34
        7
74
        7
70
        7
36
        6
        5
71
32
        5
72
        4
29
        3
        3
75
        2
33
        2
77
        2
76
        2
31
30
        1
28
        1
73
        1
```

Name: count, dtype: int64

Es gibt auch Patienten die "sehr" jung oder alt sind. Der jüngste Patient ist 28 und der älteste ist 77. Es kommen allerdings wenige Personen in diesem Datensatz vor, die an diese Altersgrenzen stoßen.

Im folgenden werden die verschiedenen kategorischen Attribute je nach Haufigkeit der Erkrankungen dargestellt.

```
In []: sns.countplot(x='Sex', hue='HeartDisease', data=df, palette=['#9aff9a', '
    plt.xlabel("Sex")
    plt.title("Heart Disease Count by Sex")
    plt.show()
```



In diesem Datensatz gibt es innerhalt der männlichen Patientengruppe deutlich mehr Herzerkrankte, während es bei der weiblichen Gruppe weniger Betroffene gibt. Man beachte, dass der Datensatz mehr männliche Einträge enthält als weibliche.

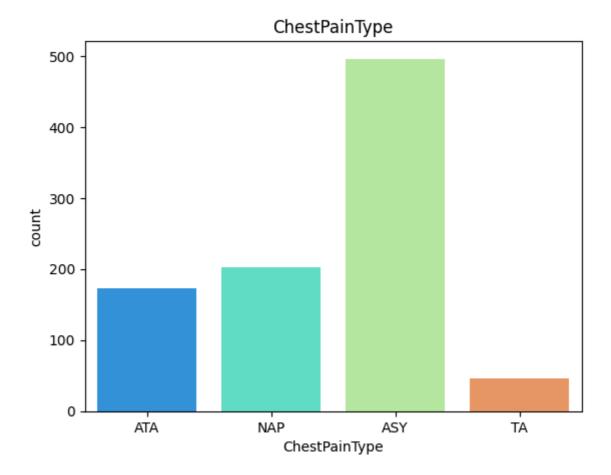
```
In []: sns.countplot(x=df['ChestPainType'], palette="rainbow")
    plt.title('ChestPainType')

/var/folders/3l/_xvv3581559_krvl1r82px5w0000gn/T/ipykernel_96307/344348670
9.py:1: FutureWarning:

Passing `palette` without assigning `hue` is deprecated and will be remove d in v0.14.0. Assign the `x` variable to `hue` and set `legend=False` for the same effect.

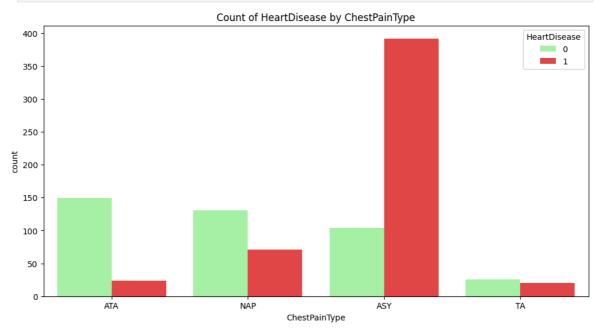
sns.countplot(x=df['ChestPainType'], palette="rainbow")

Out[]: Text(0.5, 1.0, 'ChestPainType')
```



Die häufigste Ausprägung bei den Brustschmerzen sind die asymptomatischen Brustschmerzen. Die wenigsten Fälle beschreiben typical angina chest pain.

```
In []: plt.figure(figsize=(12, 6))
    sns.countplot(x='ChestPainType', hue='HeartDisease', data=df, palette=col
    plt.title('Count of HeartDisease by ChestPainType')
    plt.show()
```



Interessanterweise zeigen Patienten mit asystomatischen Brustschmerzen am häufigsten eine Herzkrankheit auf. Bei der Gruppe TA gibt es in etwas gleich viele

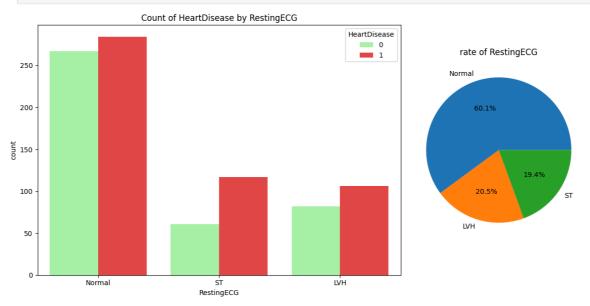
Gesunde wie Erkrankte. In den anderen beiden Gruppen überwiegt die Anzahl der gesunden Patienten.

```
In []: fig = plt.figure(figsize=(12, 6))
    gs = fig.add_gridspec(1, 2, width_ratios=[2, 1])

ax1 = fig.add_subplot(gs[0])
    sns.countplot( x='RestingECG', hue='HeartDisease', data=df, palette=colors
    ax1.set_title('Count of HeartDisease by RestingECG')

ax2 = fig.add_subplot(gs[1])
    types = df['RestingECG'].value_counts()
    ax2.pie(types, labels=types.index, autopct='%1.1f%%')
    ax2.set_title('rate of RestingECG')

plt.tight_layout()
    plt.show()
```



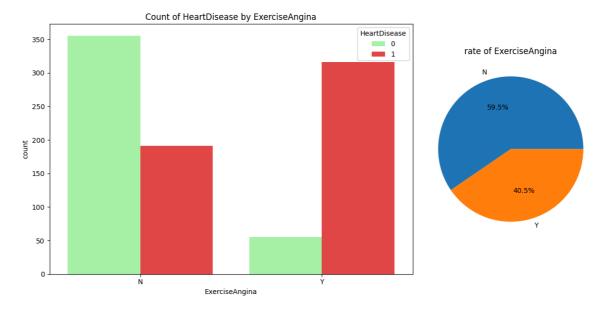
Die Ruheelektrokardiogrammergebnisse zeigen überwiegend normale Werte (60%). Die Gruppen LVH und ST (Beschreibung siehe oben) sind mit jeweils annähernd 20% seltener vertreten. Pro Gruppe gibt es allerdings stets mehr erkrankte als gesunde Patienten.

```
In []: fig = plt.figure(figsize=(12, 6))
    gs = fig.add_gridspec(1, 2, width_ratios=[2, 1])

ax1 = fig.add_subplot(gs[0])
    sns.countplot( x='ExerciseAngina', hue='HeartDisease', data=df, palette=co
    ax1.set_title('Count of HeartDisease by ExerciseAngina')

ax2 = fig.add_subplot(gs[1])
    types = df['ExerciseAngina'].value_counts()
    ax2.pie(types, labels=types.index, autopct='%1.1f%%')
    ax2.set_title('rate of ExerciseAngina')

plt.tight_layout()
    plt.show()
```



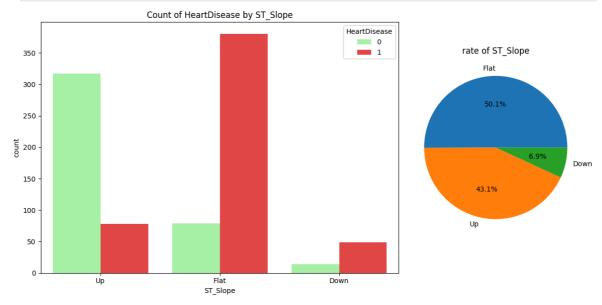
Die Mehrheit im Datensatz besitzt keine Belastungsinduzierte Brustschmerzen. Diejenigen Patienten die derartige Brustschmerzen aufweisen, haben jedoch signigfikant öfter eine Herzerkranung als die andere Gruppe.

```
In []: fig = plt.figure(figsize=(12, 6))
    gs = fig.add_gridspec(1, 2, width_ratios=[2, 1])

ax1 = fig.add_subplot(gs[0])
    sns.countplot( x='ST_Slope', hue='HeartDisease', data=df, palette=colors_r
    ax1.set_title('Count of HeartDisease by ST_Slope')

ax2 = fig.add_subplot(gs[1])
    types = df['ST_Slope'].value_counts()
    ax2.pie(types, labels=types.index, autopct='%1.1f%%')
    ax2.set_title('rate of ST_Slope')

plt.tight_layout()
    plt.show()
```



ST_Slope beschreibt die Steigung des peak exercise ST Segments. Wie man der Visualisierung entnehmen kann gibt es überwiegend flache und und steigende ST

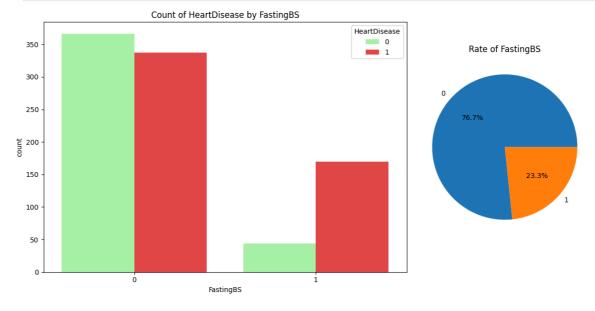
Segmente. Bis auf den steigenden Segmenten gibt es in jeder Gruppe deutlich mehr Herzerkrankte zu geben. Wie es scheint, sind Patienten mit einer steigenden Kurve wahrscheinlicher gesund.

```
In []: fig = plt.figure(figsize=(12, 6))
    gs = fig.add_gridspec(1, 2, width_ratios=[2, 1])

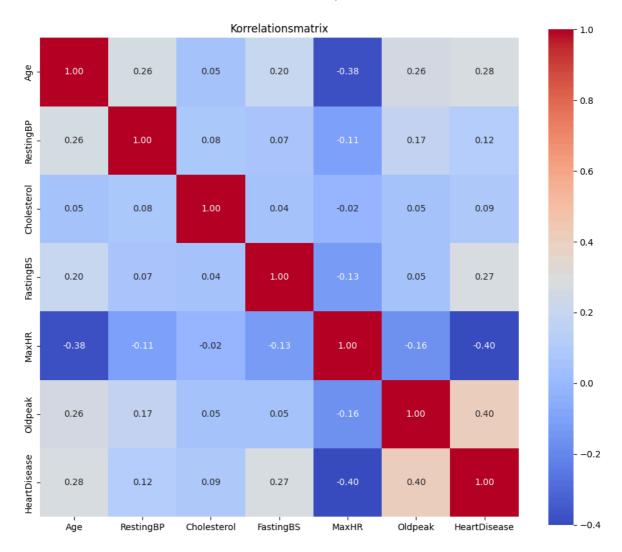
# Countplot für HeartDisease nach FastingBS
    ax1 = fig.add_subplot(gs[0])
    sns.countplot(x='FastingBS', hue='HeartDisease', data=df, palette=colors_ax1.set_title('Count of HeartDisease by FastingBS')

# Pie-Chart für die Verteilung von FastingBS
    ax2 = fig.add_subplot(gs[1])
    types = df['FastingBS'].value_counts()
    ax2.pie(types, labels=types.index, autopct='%1.1f%%')
    ax2.set_title('Rate of FastingBS')

plt.tight_layout()
    plt.show()
```



Das Attribut FastingBS beschreibt den nüchternen Blutzuckerspiegel eines Patienten, während Werte von 1 einen Blutzuckerspiegel von > 120 mg/dl kennzeichnen. Werte darunter sind mit 0 beschrieben. Der Großteil der Patienten fällt unter die Gruppe 0. In dieser Gruppe gibt es annähernd gleich viele Patienten mit sowie ohne Krankheit. In der Gruppe mit dem höheren Blutzuckerspiegel haben weitaus mehr Patienten eine Herzkrankheit.



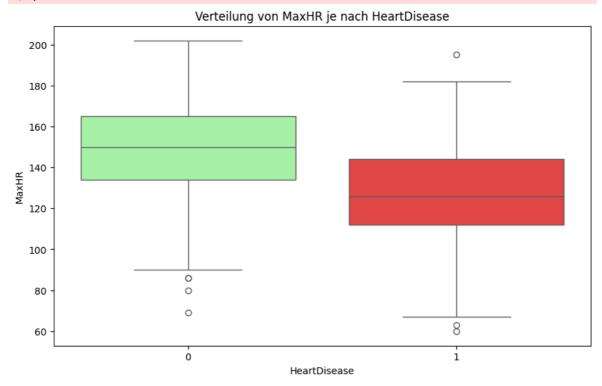
Um die Korrelationen zwischen den einzelnen Attributen zu ermittlen, wird diese Korrelationsmatrix erstellt. Die stärkste Korrelation weisen die Attribute Oldpeak und Heartdisease auf. Das lässt darauf schließen, dass sich je nach Gruppe innerhalb des Attributs Oldpeak eine genauere Aussage über den Gesundheitszustands eines Patienten fallen lässt. Weitere, jedoch schwächere Korrelationen (>= 0.20) herrschen zwischen den Attributen Age und Heartdisease, Age und Oldpeak, RestingPB und Age, FastingBS und Age, MaxHR und Cholesterol. Auffällig ist die negative Korrelation zwsichen MaxHR und Heartdisease. Der Wert -0,40 besagt, dass ein Patient mit Herzerkrankung einen tendenziell niedrigeren Maximalen Puls hat. Dies erschien auf den ersten Blick merkwürdig, da die Annahme herrschte, Herzerkrankte menschen hätten einen höheren Puls.

```
In []: # Erstellen des Boxplots
plt.figure(figsize=(10, 6))
sns.boxplot(x='HeartDisease', y='MaxHR', data=df, palette=colors_red_gree
plt.title('Verteilung von MaxHR je nach HeartDisease')
plt.xlabel('HeartDisease')
plt.ylabel('MaxHR')
plt.show()
```

/var/folders/3l/_xvv3581559_krvl1r82px5w0000gn/T/ipykernel_96307/208614592 5.py:3: FutureWarning:

Passing `palette` without assigning `hue` is deprecated and will be remove d in v0.14.0. Assign the `x` variable to `hue` and set `legend=False` for the same effect.

sns.boxplot(x='HeartDisease', y='MaxHR', data=df, palette=colors_red_gre
en)



Doch dieser Boxplot bestätigt den Wert in der Korrelationsmatrix.

Outlier Detection

In den nächsten werden die Outlier im Datensatz analysiert.

```
In []: # Wähle nur numerische Spalten aus
numeric_cols = df.select_dtypes(include=[np.number]).columns

# Definiere eine Funktion zur Identifizierung von Ausreißern
def detect_outliers(data):
    """
    Detect outliers in the given DataFrame.

Parameters:
    - data (DataFrame): The DataFrame containing the data.

Returns:
    - outliers (list): A list of indices corresponding to the outliers in
    """
    outliers = []
    for col in data.columns:
        q1 = data[col].quantile(0.25)
        q3 = data[col].quantile(0.75)
        iqr = q3 - q1
```

```
lower_bound = q1 - 1.5 * iqr
    upper_bound = q3 + 1.5 * iqr
    outlier_indices = data[(data[col] < lower_bound) | (data[col] > u
    outliers.extend(outlier_indices)
    return outliers

# Finde Ausreißer
outliers_indices = detect_outliers(df[numeric_cols])

# Entferne doppelte Indizes
outliers_indices = list(set(outliers_indices))

# Drucke die Ausreißer
print("Indices of outliers:", outliers_indices)

# Drucke die Ausreißer-Datensätze
print("Outlier rows:")
print(df.iloc[outliers_indices])
```

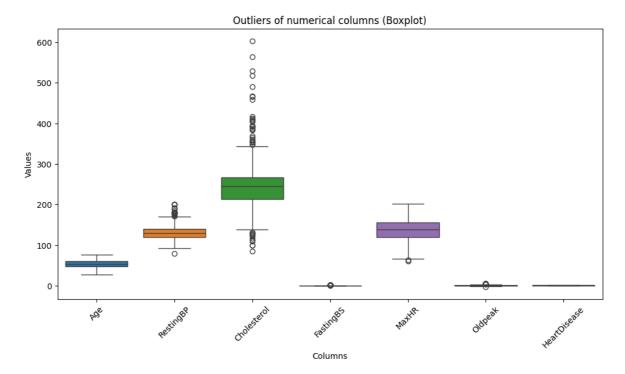
```
Indices of outliers: [515, 516, 518, 521, 522, 531, 532, 536, 537, 538, 2
8, 541, 30, 544, 546, 547, 36, 549, 550, 38, 553, 554, 556, 557, 559, 563,
52, 564, 58, 571, 826, 573, 574, 575, 577, 579, 580, 69, 582, 68, 584, 58
5, 76, 589, 78, 592, 593, 594, 595, 84, 86, 599, 604, 605, 606, 607, 97, 9
8, 610, 612, 613, 102, 103, 616, 105, 108, 109, 621, 624, 112, 117, 120, 6
32, 123, 639, 128, 132, 644, 650, 658, 659, 660, 149, 666, 667, 155, 160,
673, 672, 675, 165, 166, 679, 682, 686, 182, 185, 187, 189, 190, 702, 701,
718, 208, 210, 725, 728, 732, 734, 224, 738, 227, 744, 238, 752, 241, 242,
759, 247, 250, 256, 771, 774, 263, 775, 780, 782, 784, 785, 274, 275, 278,
790, 791, 793, 795, 284, 796, 799, 802, 803, 294, 295, 296, 297, 298, 299,
300, 809, 302, 303, 304, 305, 306, 308, 309, 820, 311, 312, 313, 314, 315,
316, 317, 318, 319, 320, 321, 322, 323, 324, 325, 838, 327, 328, 329, 330,
331, 842, 333, 334, 335, 843, 337, 338, 339, 340, 341, 342, 855, 343, 344,
850, 347, 349, 350, 869, 871, 872, 365, 880, 370, 372, 887, 888, 377, 378,
900, 389, 901, 390, 393, 394, 395, 396, 397, 398, 399, 400, 401, 402, 915,
403, 404, 405, 406, 407, 408, 409, 411, 410, 412, 413, 414, 415, 420, 422,
423, 424, 425, 430, 437, 441, 442, 443, 444, 448, 454, 457, 458, 460, 914,
469, 472, 473, 908, 475, 476, 477, 478, 480, 481, 482, 485, 486, 911, 491,
496, 498, 500, 503, 504, 505, 508, 511]
Outlier rows:
```

	Age	Sex	ChestPainType	RestingBP	Cholesterol	FastingBS	RestingECG
\							
516	68	М	NAP	150	195	1	Normal
517	65	М	ASY	150	235	0	Normal
519	63	М	ASY	96	305	0	ST
522	50	М	ASY	144	349	0	LVH
523	59	М	ASY	124	160	0	Normal
504	62	М	ASY	158	210	1	Normal
505	55	М	NAP	136	245	1	ST
506	75	М	ASY	136	225	0	Normal
509	58	М	ASY	110	198	0	Normal
512	35	М	NAP	123	161	0	ST

	MaxHR	ExerciseAngina	0ldpeak	ST_Slope	HeartDisease
516	132	N	0.0	Flat	1
517	120	Υ	1.5	Flat	1
519	121	Υ	1.0	Up	1
522	120	Υ	1.0	Up	1
523	117	Υ	1.0	Flat	1
504	112	Υ	3.0	Down	1
505	131	Υ	1.2	Flat	1
506	112	Υ	3.0	Flat	1
509	110	N	0.0	Flat	1
512	153	N	-0.1	Up	0

[275 rows x 12 columns]

```
In []: plt.figure(figsize=(12, 6))
    sns.boxplot(data=df[numeric_cols])
    plt.xticks(rotation=45) # Rotiere die Beschriftungen der x-Achse für bes
    plt.title('Outliers of numerical columns (Boxplot)')
    plt.xlabel('Columns')
    plt.ylabel('Values')
    plt.show()
```



Die Outlier Detection liefert bis gute Ergebnisse. Dadurch, dass die Anomalien bei der Analyse der Datenqulität behoben worden, gibt es keine weiteren erheblichen Einschränkungen in den Daten. Während Serum Cholesterol Werte von > 600 äußert gefährlich erscheinen, sind diese in der Realität dennoch möglich.

Machine Learning

Der vorliegende Datensatz liefert ein binäres Klassifikationsproblem. Um einen ersten Ansatz für die Auswahl eines endgültigen Classifiers zu ermitteln, wurden im Folgenden 3 verschiedene Classifier getestet.

Diese wären:

- Randomforest
- Logistic Regression
- Support Vector Machine

Die Auswahl dieser drei Classifie beruht auf ihrer Effektivität bei binären Klassifikationsproblemen. Jeder Algorithmus bietet spezifische Vorzüge und kann unterschiedliche Aspekte des Problems abdecken.

Metriken

Für die Bewertung des Models werden insbesondere die folgenden Metriken verwendet:

- Recall
- F1-Score
- ROC-AUC-Score

Der F1-Score ist ein harmonisches Mittelmaß zwischen Präzision und Recall und ermöglicht eine ausgewogene Bewertung von False Positives und False Negatives.

Der Recall bewertet die Fähigkeit des Modells, positive Instanzen korrekt zu identifizieren, was besonders wichtig ist, um sicherzustellen, dass keine relevanten Fälle übersehen werden.

Der ROC-AUC-Score bewertet die Fähigkeit des Modells, zwischen den Klassen zu unterscheiden, indem er die Fläche unter der ROC-Kurve misst, wobei ein höherer Wert auf eine bessere Leistung hinweist.

Diese Metriken in Kombination bieten eine umfassende Bewertung des Modells für das binäre Klassifikationsproblem.

Feature Engineering

Da die Kategorischen Attribute nicht vom Classifier erkannt werden, wird hier ein One-Hot-Encoding angewandt. One-Hot-Encoding ist eine Methode zur Umwandlung von kategorischen Variablen in ein binäres Format, das von maschinellen Lernalgorithmen besser verstanden werden kann.

```
In []: # featrue engineering
df_encoded = pd.get_dummies(df, columns=["Sex", "ChestPainType", "Resting")
```

Im nächsten Schritt werden die target und feature Variablen festgelegt.

```
In []: # preparation for train/test split
    target = df_encoded["HeartDisease"]
    features = df_encoded.drop("HeartDisease", axis=1)
```

Train-/Testsplit

Um einen Bias im Machine Learning Model zu vermeiden, splittet man den Datensatz auf in Trainings- und Testdaten. Der Trainingsdatensatz wird verwendet, um das tatsächliche Modell zu erstellen, das der Algorithmus verwenden wird, wenn er neuen Daten ausgesetzt ist.

Das Testset ist der letzte Datensatz, der verwendet wird. Die Genauigkeit bei der Vorhersage des Testsets entspricht der Genauigkeit des ML-Algorithmus.

Für den train/test Split wird ein Verhältnis von 80/20 gewählt.

```
In [ ]: # train/test split (80%/20%)
features_train, features_test, target_train, target_test = train_test_spl
```

Classifier

Die folgende Methoden werden verwendet, um die drei gewählten Classifier zur fitten, die Scores anzuzeigen und jeweils die Confusion Matrix auszuegebn.

```
In [ ]: def model(classifier):
            Train the classifier on the training data and evaluate its performance
            Parameters:

    classifier: The classifier model to be trained and evaluated.

            Returns:
            None
            0.00
            classifier.fit(features_train, target_train)
            prediction = classifier.predict(features_test)
            print("Accuracy: {:.2%}".format(accuracy_score(target_test, predictio))
            print("ROC_AUC Score: {:.2%}".format(roc_auc_score(target_test, predi
        def model evaluation(classifier):
            Evaluate the classifier using various performance metrics and visuali
            Parameters:
            - classifier: The trained classifier model.
            Returns:
            None
            .....
            # disply confusion Matrix
            cm = confusion_matrix(target_test, classifier.predict(features_test))
            names = ['True Neg', 'False Pos', 'False Neg', 'True Pos']
            counts = [value for value in cm.flatten()]
            percentages = ['{:.2%}'.format(value) for value in cm.flatten() / np.
            labels = [f'{v1}\n{v2}\n{v3}'  for v1, v2, v3 in zip(names, counts, pe
            labels = np.asarray(labels).reshape(2, 2)
            sns.heatmap(cm, annot=labels, cmap=colors, fmt='')
            # show classification Report
            print(classification_report(target_test, classifier.predict(features_
```

ML: Random Forrest

Der Random Forest Classifier ist ein Algorithmus für die Klassifizierung, der auf der Kombination mehrerer Entscheidungsbäume basiert. Er eignet sich gut für die Vorhersage von Herzkrankheiten aufgrund seiner Fähigkeit, mit verschiedenen Datentypen umzugehen und robuste Ergebnisse zu liefern. Er heißt "Random" Forest, da beim Algorithmus zwei zufällige Prozesse ablaufen. Zum einen das Bootstrapping zum anderen die Feature Auswahl beim erstellen der Entscheidungsbäume. Der Algorithmus baut also eine Vielzahl an Bäumen, die auf zufälligen Daten des Datensatzes basieren.

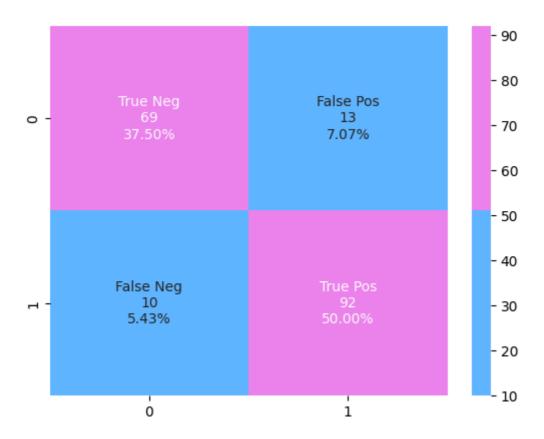
```
In [ ]: # defining RFC
forest = RandomForestClassifier()
```

```
# get scores
model(forest)
```

Accuracy: 87.50% ROC_AUC Score: 87.17%

In []: # get evalutation
model_evaluation(forest)

	precision	recall	f1-score	support
0 1	0.87 0.88	0.84 0.90	0.86 0.89	82 102
accuracy macro avg	0.87	0.87	0.88 0.87	184 184
weighted avg	0.87	0.88	0.87	184



ML: Logistic Regression

Die logistische Regression ist ein Algorithmus zur Klassifizierung, der die Wahrscheinlichkeit für das Eintreten eines Ereignisses basierend auf einer oder mehreren unabhängigen Variablen schätzt. Dabei nutzt sie die logistische Funktion, um die Vorhersage zwischen 0 und 1 zu skalieren. Sie eignet sich gut für binäre Klassifizierungsaufgaben wie die Vorhersage von Herzkrankheiten.

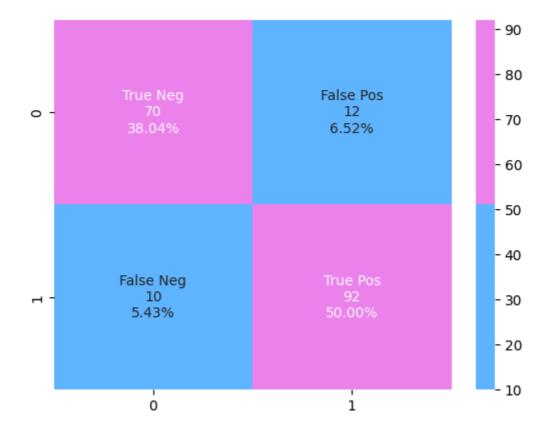
```
In []: # defining LRC
    classifier_lr = LogisticRegression(max_iter=10000)
```

```
# get scores
model(classifier_lr)
```

Accuracy: 88.04% ROC_AUC Score: 87.78%

In []: # get scores
model_evaluation(classifier_lr)

	precision	recall	f1-score	support
0 1	0.88 0.88	0.85 0.90	0.86 0.89	82 102
accuracy macro avg weighted avg	0.88 0.88	0.88 0.88	0.88 0.88 0.88	184 184 184



ML: Support Vector Machine

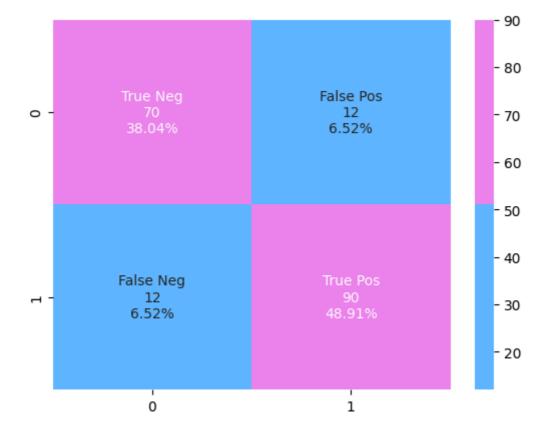
Die Support Vector Machine ist ebenfalls ein Klassifizierungsalgorithmus, der darauf abzielt, eine Trennung zwischen den verschiedenen Klassen zu finden, indem er die beste Entscheidungsgrenze (Hyperplane) zwischen den Datenpunkten sucht. Sie funktioniert, indem sie den Abstand zwischen den Datenpunkten maximiert und gleichzeitig eine minimale Fehlerrate aufweist. SVM eignet sich gut für datengetriebene Anwendungen mit komplexen Entscheidungsgrenzen und kann auch mit nicht-linearen Daten umgehen, indem sie den sogenannten Kernel-Trick anwendet. In Bezug auf Herzkrankheiten eignet sich die SVM, wenn die Daten gut separierbar sind und klare Entscheidungsgrenzen zwischen den Klassen existieren.

```
In []: # defining SVM
svc = SVC(kernel = 'linear', C = 0.1)
# get scores
model(svc)
```

Accuracy: 86.96% ROC_AUC Score: 86.80%

In []: # get evaluation
model_evaluation(svc)

support	f1-score	recall	precision	
82 102	0.85 0.88	0.85 0.88	0.85 0.88	0 1
184 184 184	0.87 0.87 0.87	0.87 0.87	0.87 0.87	accuracy macro avg weighted avg

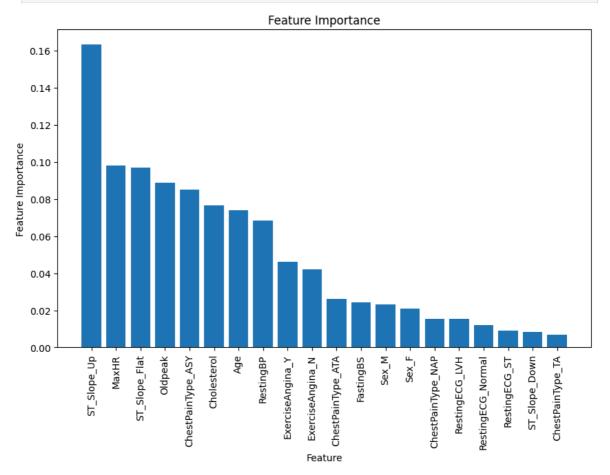


Feature Importance

```
In []: # plot for feature importances
    importances = forest.feature_importances_
    indices = np.argsort(importances)[::-1]
    feature_names = features_train.columns

plt.figure(figsize=(10, 6))
    plt.title("Feature Importance")
    plt.bar(range(features_train.shape[1]), importances[indices], align="cent plt.xticks(range(features_train.shape[1]), feature_names[indices], rotati plt.xlabel("Feature")
```

plt.ylabel("Feature Importance")
plt.show()



Gridsearch Prameter Tuning

Das Parameter-Tuning wird nur für den Random Forest Classifier (RFC) durchgeführt, da dieser bereits ohne Tuning die beste Leistung erzielt hat. Der RFC ist bekannt für seine Vielseitigkeit und Robustheit, insbesondere bei binären Klassifikationsproblemen wie im vorliegenden Fall. Durch das Feintuning seiner Hyperparameter kann die Vorhersagegenauigkeit weiter optimiert und potenzielles Overfitting reduziert werden. Dies ermöglicht eine präzisere Identifizierung von Herzkrankheiten, was in medizinischen Anwendungen von entscheidender Bedeutung ist.

Das Parameter-Tuning wird mithilfe von Grid Search durchgeführt, einem Ansatz zur systematischen Suche nach den besten Hyperparameter-Kombinationen für ein Machine Learning-Modell. Grid Search durchläuft vordefinierte Kombinationen von Hyperparametern und bewertet die Leistung des Modells anhand einer bestimmten Metrik für jede Kombination. In unserem Fall optimieren wir den Receiver Operating Characteristic Area Under Curve (ROC AUC) Score. Der ROC AUC Score ist eine Metrik, die die Fähigkeit eines Modells bewertet, zwischen den Klassen zu unterscheiden und die Trade-offs zwischen True Positive Rate und False Positive Rate darstellt. Für das binäre Klassifikationsproblem mit Herzkrankheiten ist es

wichtig, dass unser Modell eine hohe Unterscheidungskraft zwischen kranken und gesunden Patienten aufweist, weshalb wir den ROC AUC Score optimieren.

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In [ ]: # defining the grid search parameters
        param_grid = {
            'n_estimators': [100, 200, 300],
            'max_depth': [None, 10, 20, 30],
            'min_samples_split': [2, 5, 10],
            'min_samples_leaf': [1, 2, 4]
In [ ]: # defining the hyperparameter space to identify the optimal combination o
        grid_search = GridSearchCV(estimator=forest,
                                    param_grid=param_grid,
                                    scoring= "roc_auc",
                                    refit="roc_auc",
                                    cv=5,
                                    n_{jobs=-1}
                                    verbose=4)
In [ ]: # searching for the best combination
        grid_search.fit(features_train, target_train)
```

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Fitting 5 folds for each of 108 candidates, totalling 540 fits
[CV 3/5] END max_depth=None, min_samples_leaf=1, min_samples_split=2, n_es
timators=100;, score=0.941 total time=
                                         0.2s
[CV 5/5] END max_depth=None, min_samples_leaf=1, min_samples_split=2, n_es
timators=100;, score=0.870 total time=
                                         0.2s
[CV 2/5] END max depth=None, min samples leaf=1, min samples split=2, n es
timators=100;, score=0.970 total time=
                                         0.3s
[CV 1/5] END max_depth=None, min_samples_leaf=1, min_samples_split=2, n_es
timators=100;, score=0.916 total time=
                                         0.3s
[CV 4/5] END max_depth=None, min_samples_leaf=1, min_samples_split=2, n_es
timators=100;, score=0.911 total time=
                                         0.2s
[CV 3/5] END max_depth=None, min_samples_leaf=1, min_samples_split=2, n_es
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                                         0.4s
[CV 1/5] END max_depth=None, min_samples_leaf=1, min_samples_split=2, n_es
timators=200;, score=0.918 total time=
                                         0.4s
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                                         0.4s
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                                         0.2s
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                                         0.4s
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                                         0.1s
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timators=300;, score=0.917 total time=
                                         0.5s
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timators=300;, score=0.944 total time=
                                         0.5s
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                                         0.2s
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[CV 3/5] END max_depth=None, min_samples_leaf=1, min_samples_split=5, n_es
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timators=300;, score=0.978 total time= 0.6s [CV 5/5] END max_depth=None, min_samples_leaf=2, min_samples_split=5, n_es timators=100;, score=0.869 total time= 0.1s [CV 4/5] END max_depth=None, min_samples_leaf=2, min_samples_split=5, n_es timators=100;, score=0.917 total time= 0.2s [CV 3/5] END max depth=None, min samples leaf=2, min samples split=2, n es timators=300;, score=0.944 total time= 0.6s [CV 4/5] END max_depth=None, min_samples_leaf=2, min_samples_split=2, n_es timators=300;, score=0.912 total time= 0.6s [CV 5/5] END max_depth=None, min_samples_leaf=2, min_samples_split=2, n_es timators=300;, score=0.872 total time= 0.6s [CV 1/5] END max_depth=None, min_samples_leaf=2, min_samples_split=5, n_es timators=200;, score=0.926 total time= 0.3s [CV 2/5] END max_depth=None, min_samples_leaf=2, min_samples_split=5, n_es timators=200;, score=0.978 total time= 0.3s [CV 3/5] END max_depth=None, min_samples_leaf=2, min_samples_split=5, n_es timators=200;, score=0.947 total time= 0.3s [CV 5/5] END max_depth=None, min_samples_leaf=2, min_samples_split=5, n_es timators=200;, score=0.872 total time= 0.3s [CV 4/5] END max_depth=None, min_samples_leaf=2, min_samples_split=5, n_es timators=200;, score=0.914 total time= 0.3s [CV 1/5] END max_depth=None, min_samples_leaf=2, min_samples_split=10, n_e stimators=100;, score=0.927 total time= 0.2s [CV 1/5] END max_depth=None, min_samples_leaf=2, min_samples_split=5, n_es timators=300;, score=0.925 total time= 0.4s [CV 3/5] END max_depth=None, min_samples_leaf=2, min_samples_split=10, n_e stimators=100;, score=0.948 total time= 0.1s [CV 2/5] END max_depth=None, min_samples_leaf=2, min_samples_split=10, n_e stimators=100;, score=0.978 total time= 0.2s [CV 2/5] END max depth=None, min samples leaf=2, min samples split=5, n es timators=300;, score=0.977 total time= 0.4s [CV 4/5] END max_depth=None, min_samples_leaf=2, min_samples_split=10, n_e stimators=100;, score=0.912 total time= 0.1s [CV 3/5] END max_depth=None, min_samples_leaf=2, min_samples_split=5, n_es timators=300;, score=0.944 total time= 0.5s [CV 5/5] END max_depth=None, min_samples_leaf=2, min_samples_split=10, n_e stimators=100;, score=0.876 total time= 0.1s [CV 4/5] END max_depth=None, min_samples_leaf=2, min_samples_split=5, n_es timators=300;, score=0.914 total time= 0.4s [CV 5/5] END max_depth=None, min_samples_leaf=2, min_samples_split=5, n_es timators=300;, score=0.878 total time= 0.4s [CV 1/5] END max_depth=None, min_samples_leaf=2, min_samples_split=10, n_e stimators=200;, score=0.925 total time= 0.3s [CV 2/5] END max_depth=None, min_samples_leaf=2, min_samples_split=10, n_e stimators=200;, score=0.979 total time= 0.2s [CV 3/5] END max_depth=None, min_samples_leaf=2, min_samples_split=10, n_e stimators=200;, score=0.945 total time= 0.3s [CV 5/5] END max_depth=None, min_samples_leaf=2, min_samples_split=10, n_e stimators=200;, score=0.868 total time= 0.2s [CV 4/5] END max_depth=None, min_samples_leaf=2, min_samples_split=10, n_e stimators=200;, score=0.915 total time= 0.3s [CV 2/5] END max_depth=None, min_samples_leaf=4, min_samples_split=2, n_es timators=100;, score=0.978 total time= 0.1s [CV 1/5] END max_depth=None, min_samples_leaf=4, min_samples_split=2, n_es timators=100;, score=0.925 total time= 0.1s [CV 1/5] END max_depth=None, min_samples_leaf=2, min_samples_split=10, n_e stimators=300;, score=0.926 total time= 0.4s [CV 3/5] END max_depth=None, min_samples_leaf=4, min_samples_split=2, n_es timators=100;, score=0.941 total time= 0.1s [CV 2/5] END max_depth=None, min_samples_leaf=2, min_samples_split=10, n_e

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stimators=100;, score=0.918 total time=
                                          0.2s
[CV 2/5] END max_depth=None, min_samples_leaf=4, min_samples_split=5, n_es
timators=300;, score=0.979 total time=
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stimators=100;, score=0.873 total time=
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timators=300;, score=0.909 total time=
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stimators=200;, score=0.875 total time=
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mators=100;, score=0.921 total time=
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mators=100;, score=0.974 total time=
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stimators=300;, score=0.942 total time=
                                        0.4s
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                                       0.3s
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mators=200;, score=0.921 total time=
                                      0.3s
[CV 3/5] END max_depth=10, min_samples_leaf=1, min_samples_split=2, n_esti
mators=200;, score=0.943 total time=
                                      0.3s
[CV 5/5] END max_depth=10, min_samples_leaf=1, min_samples_split=2, n_esti
mators=200;, score=0.874 total time=
                                       0.3s
[CV 4/5] END max_depth=10, min_samples_leaf=1, min_samples_split=2, n_esti
mators=200;, score=0.917 total time=
                                       0.4s
[CV 2/5] END max_depth=10, min_samples_leaf=1, min_samples_split=5, n_esti
mators=100;, score=0.976 total time=
                                      0.1s
[CV 1/5] END max_depth=10, min_samples_leaf=1, min_samples_split=5, n_esti
mators=100;, score=0.922 total time=
                                      0.1s
[CV 3/5] END max_depth=10, min_samples_leaf=1, min_samples_split=5, n_esti
mators=100;, score=0.941 total time=
                                       0.1s
[CV 1/5] END max_depth=10, min_samples_leaf=1, min_samples_split=2, n_esti
mators=300;, score=0.920 total time=
                                       0.4s
[CV 2/5] END max_depth=10, min_samples_leaf=1, min_samples_split=2, n_esti
```

mators=300;, score=0.974 total time= 0.4s [CV 4/5] END max_depth=10, min_samples_leaf=1, min_samples_split=5, n_esti mators=100;, score=0.918 total time= 0.1s [CV 5/5] END max_depth=10, min_samples_leaf=1, min_samples_split=5, n_esti mators=100;, score=0.868 total time= 0.1s [CV 3/5] END max depth=10, min samples leaf=1, min samples split=2, n esti mators=300;, score=0.944 total time= 0.4s [CV 5/5] END max_depth=10, min_samples_leaf=1, min_samples_split=2, n_esti mators=300;, score=0.870 total time= 0.4s [CV 4/5] END max_depth=10, min_samples_leaf=1, min_samples_split=2, n_esti mators=300;, score=0.913 total time= 0.4s [CV 1/5] END max_depth=10, min_samples_leaf=1, min_samples_split=5, n_esti mators=200;, score=0.923 total time= 0.2s [CV 3/5] END max_depth=10, min_samples_leaf=1, min_samples_split=5, n_esti mators=200;, score=0.943 total time= 0.2s [CV 2/5] END max_depth=10, min_samples_leaf=1, min_samples_split=5, n_esti mators=200;, score=0.975 total time= 0.3s [CV 4/5] END max_depth=10, min_samples_leaf=1, min_samples_split=5, n_esti mators=200;, score=0.910 total time= 0.3s [CV 5/5] END max_depth=10, min_samples_leaf=1, min_samples_split=5, n_esti mators=200;, score=0.869 total time= 0.3s [CV 1/5] END max_depth=10, min_samples_leaf=1, min_samples_split=10, n_est imators=100;, score=0.923 total time= 0.1s [CV 2/5] END max_depth=10, min_samples_leaf=1, min_samples_split=10, n_est imators=100;, score=0.977 total time= 0.1s [CV 3/5] END max_depth=10, min_samples_leaf=1, min_samples_split=10, n_est imators=100;, score=0.944 total time= 0.1s [CV 2/5] END max_depth=10, min_samples_leaf=1, min_samples_split=5, n_esti mators=300;, score=0.975 total time= 0.4s [CV 1/5] END max depth=10, min samples leaf=1, min samples split=5, n esti mators=300;, score=0.919 total time= 0.4s [CV 3/5] END max_depth=10, min_samples_leaf=1, min_samples_split=5, n_esti mators=300;, score=0.943 total time= 0.4s [CV 4/5] END max_depth=10, min_samples_leaf=1, min_samples_split=10, n_est imators=100;, score=0.915 total time= 0.1s [CV 5/5] END max_depth=10, min_samples_leaf=1, min_samples_split=10, n_est imators=100;, score=0.871 total time= 0.2s [CV 4/5] END max_depth=10, min_samples_leaf=1, min_samples_split=5, n_esti mators=300;, score=0.912 total time= 0.4s [CV 5/5] END max_depth=10, min_samples_leaf=1, min_samples_split=5, n_esti mators=300;, score=0.873 total time= 0.4s [CV 1/5] END max_depth=10, min_samples_leaf=1, min_samples_split=10, n_est imators=200;, score=0.926 total time= 0.3s [CV 2/5] END max_depth=10, min_samples_leaf=1, min_samples_split=10, n_est imators=200;, score=0.977 total time= 0.4s [CV 3/5] END max_depth=10, min_samples_leaf=1, min_samples_split=10, n_est imators=200;, score=0.947 total time= 0.4s [CV 4/5] END max_depth=10, min_samples_leaf=1, min_samples_split=10, n_est imators=200;, score=0.915 total time= 0.4s [CV 5/5] END max_depth=10, min_samples_leaf=1, min_samples_split=10, n_est imators=200;, score=0.871 total time= 0.4s [CV 1/5] END max_depth=10, min_samples_leaf=2, min_samples_split=2, n_esti mators=100;, score=0.924 total time= 0.2s [CV 2/5] END max_depth=10, min_samples_leaf=2, min_samples_split=2, n_esti mators=100;, score=0.973 total time= 0.2s [CV 3/5] END max_depth=10, min_samples_leaf=2, min_samples_split=2, n_esti mators=100;, score=0.942 total time= 0.2s [CV 4/5] END max_depth=10, min_samples_leaf=2, min_samples_split=2, n_esti mators=100;, score=0.910 total time= 0.1s [CV 3/5] END max_depth=10, min_samples_leaf=1, min_samples_split=10, n_est

imators=300;, score=0.944 total time= 0.5s [CV 5/5] END max_depth=10, min_samples_leaf=2, min_samples_split=2, n_esti mators=100;, score=0.868 total time= 0.2s [CV 4/5] END max_depth=10, min_samples_leaf=1, min_samples_split=10, n_est imators=300;, score=0.907 total time= 0.5s [CV 1/5] END max depth=10, min samples leaf=1, min samples split=10, n est imators=300;, score=0.924 total time= 0.7s [CV 2/5] END max_depth=10, min_samples_leaf=1, min_samples_split=10, n_est imators=300;, score=0.979 total time= 0.7s [CV 5/5] END max_depth=10, min_samples_leaf=1, min_samples_split=10, n_est imators=300;, score=0.869 total time= 0.5s [CV 1/5] END max_depth=10, min_samples_leaf=2, min_samples_split=2, n_esti mators=200;, score=0.927 total time= 0.4s [CV 2/5] END max_depth=10, min_samples_leaf=2, min_samples_split=2, n_esti mators=200;, score=0.973 total time= 0.3s [CV 3/5] END max_depth=10, min_samples_leaf=2, min_samples_split=2, n_esti mators=200;, score=0.944 total time= 0.3s [CV 4/5] END max_depth=10, min_samples_leaf=2, min_samples_split=2, n_esti mators=200;, score=0.915 total time= 0.3s [CV 5/5] END max_depth=10, min_samples_leaf=2, min_samples_split=2, n_esti mators=200;, score=0.871 total time= 0.3s [CV 2/5] END max_depth=10, min_samples_leaf=2, min_samples_split=5, n_esti mators=100;, score=0.977 total time= 0.1s [CV 1/5] END max_depth=10, min_samples_leaf=2, min_samples_split=5, n_esti mators=100;, score=0.917 total time= 0.2s [CV 3/5] END max_depth=10, min_samples_leaf=2, min_samples_split=5, n_esti mators=100;, score=0.942 total time= 0.1s [CV 1/5] END max_depth=10, min_samples_leaf=2, min_samples_split=2, n_esti mators=300;, score=0.924 total time= 0.4s [CV 2/5] END max depth=10, min samples leaf=2, min samples split=2, n esti mators=300;, score=0.977 total time= 0.5s [CV 3/5] END max_depth=10, min_samples_leaf=2, min_samples_split=2, n_esti mators=300;, score=0.945 total time= 0.4s [CV 4/5] END max_depth=10, min_samples_leaf=2, min_samples_split=5, n_esti mators=100;, score=0.916 total time= 0.2s [CV 5/5] END max_depth=10, min_samples_leaf=2, min_samples_split=5, n_esti mators=100;, score=0.872 total time= 0.2s [CV 4/5] END max_depth=10, min_samples_leaf=2, min_samples_split=2, n_esti mators=300;, score=0.912 total time= 0.5s [CV 5/5] END max_depth=10, min_samples_leaf=2, min_samples_split=2, n_esti mators=300;, score=0.877 total time= 0.6s [CV 1/5] END max_depth=10, min_samples_leaf=2, min_samples_split=5, n_esti mators=200;, score=0.923 total time= 0.4s [CV 2/5] END max_depth=10, min_samples_leaf=2, min_samples_split=5, n_esti mators=200;, score=0.975 total time= 0.4s [CV 3/5] END max_depth=10, min_samples_leaf=2, min_samples_split=5, n_esti mators=200;, score=0.943 total time= 0.3s [CV 4/5] END max_depth=10, min_samples_leaf=2, min_samples_split=5, n_esti mators=200;, score=0.913 total time= 0.4s [CV 5/5] END max_depth=10, min_samples_leaf=2, min_samples_split=5, n_esti mators=200;, score=0.869 total time= 0.3s [CV 1/5] END max_depth=10, min_samples_leaf=2, min_samples_split=10, n_est imators=100;, score=0.918 total time= 0.1s [CV 2/5] END max_depth=10, min_samples_leaf=2, min_samples_split=10, n_est imators=100;, score=0.976 total time= 0.1s [CV 3/5] END max_depth=10, min_samples_leaf=2, min_samples_split=10, n_est imators=100;, score=0.944 total time= 0.1s [CV 1/5] END max_depth=10, min_samples_leaf=2, min_samples_split=5, n_esti mators=300;, score=0.927 total time= 0.4s [CV 2/5] END max_depth=10, min_samples_leaf=2, min_samples_split=5, n_esti

mators=300;, score=0.977 total time= 0.4s [CV 4/5] END max_depth=10, min_samples_leaf=2, min_samples_split=10, n_est imators=100;, score=0.918 total time= 0.1s [CV 5/5] END max_depth=10, min_samples_leaf=2, min_samples_split=10, n_est imators=100;, score=0.874 total time= 0.1s [CV 3/5] END max depth=10, min samples leaf=2, min samples split=5, n esti mators=300;, score=0.942 total time= 0.4s [CV 5/5] END max_depth=10, min_samples_leaf=2, min_samples_split=5, n_esti mators=300;, score=0.868 total time= 0.4s [CV 4/5] END max_depth=10, min_samples_leaf=2, min_samples_split=5, n_esti mators=300;, score=0.915 total time= 0.5s [CV 1/5] END max_depth=10, min_samples_leaf=2, min_samples_split=10, n_est imators=200;, score=0.926 total time= 0.3s [CV 2/5] END max_depth=10, min_samples_leaf=2, min_samples_split=10, n_est imators=200;, score=0.980 total time= 0.3s [CV 3/5] END max_depth=10, min_samples_leaf=2, min_samples_split=10, n_est imators=200;, score=0.944 total time= 0.3s [CV 4/5] END max_depth=10, min_samples_leaf=2, min_samples_split=10, n_est imators=200;, score=0.911 total time= 0.3s [CV 5/5] END max_depth=10, min_samples_leaf=2, min_samples_split=10, n_est imators=200;, score=0.874 total time= 0.3s [CV 1/5] END max_depth=10, min_samples_leaf=4, min_samples_split=2, n_esti mators=100;, score=0.922 total time= 0.1s [CV 2/5] END max_depth=10, min_samples_leaf=4, min_samples_split=2, n_esti mators=100;, score=0.979 total time= 0.1s [CV 3/5] END max_depth=10, min_samples_leaf=4, min_samples_split=2, n_esti mators=100;, score=0.940 total time= 0.1s [CV 4/5] END max_depth=10, min_samples_leaf=4, min_samples_split=2, n_esti mators=100;, score=0.912 total time= 0.1s [CV 1/5] END max depth=10, min samples leaf=2, min samples split=10, n est imators=300;, score=0.925 total time= 0.4s [CV 2/5] END max_depth=10, min_samples_leaf=2, min_samples_split=10, n_est imators=300;, score=0.978 total time= 0.4s [CV 5/5] END max_depth=10, min_samples_leaf=4, min_samples_split=2, n_esti mators=100;, score=0.875 total time= 0.2s [CV 3/5] END max_depth=10, min_samples_leaf=2, min_samples_split=10, n_est imators=300;, score=0.941 total time= 0.4s [CV 4/5] END max_depth=10, min_samples_leaf=2, min_samples_split=10, n_est imators=300;, score=0.911 total time= 0.4s [CV 5/5] END max_depth=10, min_samples_leaf=2, min_samples_split=10, n_est imators=300;, score=0.869 total time= 0.4s [CV 4/5] END max_depth=10, min_samples_leaf=4, min_samples_split=2, n_esti mators=200;, score=0.912 total time= 0.2s [CV 3/5] END max_depth=10, min_samples_leaf=4, min_samples_split=2, n_esti mators=200;, score=0.943 total time= 0.3s [CV 1/5] END max_depth=10, min_samples_leaf=4, min_samples_split=2, n_esti mators=200;, score=0.930 total time= 0.3s [CV 2/5] END max_depth=10, min_samples_leaf=4, min_samples_split=2, n_esti mators=200;, score=0.977 total time= 0.4s [CV 5/5] END max_depth=10, min_samples_leaf=4, min_samples_split=2, n_esti mators=200;, score=0.870 total time= 0.3s [CV 1/5] END max_depth=10, min_samples_leaf=4, min_samples_split=5, n_esti mators=100;, score=0.926 total time= 0.1s [CV 2/5] END max_depth=10, min_samples_leaf=4, min_samples_split=5, n_esti mators=100;, score=0.974 total time= 0.1s [CV 1/5] END max_depth=10, min_samples_leaf=4, min_samples_split=2, n_esti mators=300;, score=0.926 total time= 0.4s [CV 3/5] END max_depth=10, min_samples_leaf=4, min_samples_split=5, n_esti mators=100;, score=0.949 total time= 0.1s [CV 4/5] END max_depth=10, min_samples_leaf=4, min_samples_split=5, n_esti

mators=100;, score=0.912 total time= 0.1s [CV 3/5] END max_depth=10, min_samples_leaf=4, min_samples_split=2, n_esti mators=300;, score=0.944 total time= 0.5s [CV 2/5] END max_depth=10, min_samples_leaf=4, min_samples_split=2, n_esti mators=300;, score=0.980 total time= 0.5s [CV 4/5] END max depth=10, min samples leaf=4, min samples split=2, n esti mators=300;, score=0.908 total time= 0.4s [CV 5/5] END max_depth=10, min_samples_leaf=4, min_samples_split=5, n_esti mators=100;, score=0.872 total time= 0.2s [CV 5/5] END max_depth=10, min_samples_leaf=4, min_samples_split=2, n_esti mators=300;, score=0.874 total time= 0.4s [CV 1/5] END max_depth=10, min_samples_leaf=4, min_samples_split=5, n_esti mators=200;, score=0.922 total time= 0.3s [CV 3/5] END max_depth=10, min_samples_leaf=4, min_samples_split=5, n_esti mators=200;, score=0.942 total time= 0.2s [CV 2/5] END max_depth=10, min_samples_leaf=4, min_samples_split=5, n_esti mators=200;, score=0.980 total time= 0.3s [CV 4/5] END max_depth=10, min_samples_leaf=4, min_samples_split=5, n_esti mators=200;, score=0.911 total time= 0.2s [CV 5/5] END max_depth=10, min_samples_leaf=4, min_samples_split=5, n_esti mators=200;, score=0.869 total time= 0.2s [CV 2/5] END max_depth=10, min_samples_leaf=4, min_samples_split=10, n_est imators=100;, score=0.976 total time= 0.1s [CV 1/5] END max_depth=10, min_samples_leaf=4, min_samples_split=10, n_est imators=100;, score=0.924 total time= 0.1s [CV 2/5] END max_depth=10, min_samples_leaf=4, min_samples_split=5, n_esti mators=300;, score=0.979 total time= 0.3s [CV 3/5] END max_depth=10, min_samples_leaf=4, min_samples_split=10, n_est imators=100;, score=0.946 total time= 0.1s [CV 1/5] END max depth=10, min samples leaf=4, min samples split=5, n esti mators=300;, score=0.926 total time= 0.4s [CV 4/5] END max_depth=10, min_samples_leaf=4, min_samples_split=10, n_est imators=100;, score=0.908 total time= 0.1s [CV 3/5] END max_depth=10, min_samples_leaf=4, min_samples_split=5, n_esti mators=300;, score=0.948 total time= 0.4s [CV 5/5] END max_depth=10, min_samples_leaf=4, min_samples_split=10, n_est imators=100;, score=0.881 total time= 0.2s [CV 4/5] END max_depth=10, min_samples_leaf=4, min_samples_split=5, n_esti mators=300;, score=0.911 total time= 0.4s [CV 5/5] END max_depth=10, min_samples_leaf=4, min_samples_split=5, n_esti mators=300;, score=0.872 total time= 0.4s [CV 1/5] END max_depth=10, min_samples_leaf=4, min_samples_split=10, n_est imators=200;, score=0.925 total time= 0.3s [CV 3/5] END max_depth=10, min_samples_leaf=4, min_samples_split=10, n_est imators=200;, score=0.943 total time= 0.3s [CV 2/5] END max_depth=10, min_samples_leaf=4, min_samples_split=10, n_est imators=200;, score=0.978 total time= 0.3s [CV 4/5] END max_depth=10, min_samples_leaf=4, min_samples_split=10, n_est imators=200;, score=0.911 total time= 0.3s [CV 5/5] END max_depth=10, min_samples_leaf=4, min_samples_split=10, n_est imators=200;, score=0.871 total time= 0.3s [CV 2/5] END max_depth=20, min_samples_leaf=1, min_samples_split=2, n_esti mators=100;, score=0.973 total time= 0.1s [CV 3/5] END max_depth=20, min_samples_leaf=1, min_samples_split=2, n_esti mators=100;, score=0.941 total time= 0.1s [CV 1/5] END max_depth=10, min_samples_leaf=4, min_samples_split=10, n_est imators=300;, score=0.925 total time= 0.4s [CV 2/5] END max_depth=10, min_samples_leaf=4, min_samples_split=10, n_est imators=300;, score=0.978 total time= 0.3s [CV 1/5] END max_depth=20, min_samples_leaf=1, min_samples_split=2, n_esti

mators=100;, score=0.921 total time= 0.2s [CV 3/5] END max_depth=10, min_samples_leaf=4, min_samples_split=10, n_est imators=300;, score=0.946 total time= 0.3s [CV 4/5] END max_depth=20, min_samples_leaf=1, min_samples_split=2, n_esti mators=100;, score=0.916 total time= 0.1s [CV 4/5] END max depth=10, min samples leaf=4, min samples split=10, n est imators=300;, score=0.914 total time= 0.4s [CV 5/5] END max_depth=20, min_samples_leaf=1, min_samples_split=2, n_esti mators=100;, score=0.879 total time= 0.1s [CV 5/5] END max_depth=10, min_samples_leaf=4, min_samples_split=10, n_est imators=300;, score=0.874 total time= 0.4s [CV 4/5] END max_depth=20, min_samples_leaf=1, min_samples_split=2, n_esti mators=200;, score=0.922 total time= 0.3s [CV 1/5] END max_depth=20, min_samples_leaf=1, min_samples_split=2, n_esti mators=200;, score=0.922 total time= 0.3s [CV 2/5] END max_depth=20, min_samples_leaf=1, min_samples_split=2, n_esti mators=200;, score=0.970 total time= 0.3s [CV 3/5] END max_depth=20, min_samples_leaf=1, min_samples_split=2, n_esti mators=200;, score=0.942 total time= 0.3s [CV 5/5] END max_depth=20, min_samples_leaf=1, min_samples_split=2, n_esti mators=200;, score=0.863 total time= 0.4s [CV 1/5] END max_depth=20, min_samples_leaf=1, min_samples_split=5, n_esti mators=100;, score=0.916 total time= 0.1s [CV 2/5] END max_depth=20, min_samples_leaf=1, min_samples_split=5, n_esti mators=100;, score=0.977 total time= 0.1s [CV 1/5] END max_depth=20, min_samples_leaf=1, min_samples_split=2, n_esti mators=300;, score=0.920 total time= 0.4s [CV 2/5] END max_depth=20, min_samples_leaf=1, min_samples_split=2, n_esti mators=300;, score=0.978 total time= 0.4s [CV 3/5] END max depth=20, min samples leaf=1, min samples split=5, n esti mators=100;, score=0.947 total time= 0.1s [CV 4/5] END max_depth=20, min_samples_leaf=1, min_samples_split=5, n_esti mators=100;, score=0.909 total time= 0.1s [CV 3/5] END max_depth=20, min_samples_leaf=1, min_samples_split=2, n_esti mators=300;, score=0.945 total time= 0.5s [CV 5/5] END max_depth=20, min_samples_leaf=1, min_samples_split=5, n_esti mators=100;, score=0.877 total time= 0.1s [CV 5/5] END max_depth=20, min_samples_leaf=1, min_samples_split=2, n_esti mators=300;, score=0.873 total time= 0.3s [CV 4/5] END max_depth=20, min_samples_leaf=1, min_samples_split=2, n_esti mators=300;, score=0.917 total time= 0.4s [CV 1/5] END max_depth=20, min_samples_leaf=1, min_samples_split=5, n_esti mators=200;, score=0.920 total time= 0.3s [CV 2/5] END max_depth=20, min_samples_leaf=1, min_samples_split=5, n_esti mators=200;, score=0.974 total time= 0.3s [CV 4/5] END max_depth=20, min_samples_leaf=1, min_samples_split=5, n_esti mators=200;, score=0.915 total time= 0.2s [CV 3/5] END max_depth=20, min_samples_leaf=1, min_samples_split=5, n_esti mators=200;, score=0.943 total time= 0.3s [CV 5/5] END max_depth=20, min_samples_leaf=1, min_samples_split=5, n_esti mators=200;, score=0.869 total time= 0.2s [CV 2/5] END max_depth=20, min_samples_leaf=1, min_samples_split=10, n_est imators=100;, score=0.978 total time= 0.2s [CV 1/5] END max_depth=20, min_samples_leaf=1, min_samples_split=10, n_est imators=100;, score=0.926 total time= 0.2s [CV 1/5] END max_depth=20, min_samples_leaf=1, min_samples_split=5, n_esti mators=300;, score=0.921 total time= 0.4s [CV 3/5] END max_depth=20, min_samples_leaf=1, min_samples_split=10, n_est imators=100;, score=0.944 total time= 0.2s [CV 4/5] END max_depth=20, min_samples_leaf=1, min_samples_split=10, n_est

imators=100;, score=0.912 total time= 0.1s [CV 2/5] END max_depth=20, min_samples_leaf=1, min_samples_split=5, n_esti mators=300;, score=0.979 total time= 0.5s [CV 5/5] END max_depth=20, min_samples_leaf=1, min_samples_split=10, n_est imators=100;, score=0.867 total time= 0.2s [CV 3/5] END max depth=20, min samples leaf=1, min samples split=5, n esti mators=300;, score=0.942 total time= 0.5s [CV 4/5] END max_depth=20, min_samples_leaf=1, min_samples_split=5, n_esti mators=300;, score=0.913 total time= 0.4s [CV 5/5] END max_depth=20, min_samples_leaf=1, min_samples_split=5, n_esti mators=300;, score=0.873 total time= 0.5s [CV 1/5] END max_depth=20, min_samples_leaf=1, min_samples_split=10, n_est imators=200;, score=0.927 total time= 0.3s [CV 2/5] END max_depth=20, min_samples_leaf=1, min_samples_split=10, n_est imators=200;, score=0.975 total time= 0.3s [CV 3/5] END max_depth=20, min_samples_leaf=1, min_samples_split=10, n_est imators=200;, score=0.941 total time= 0.3s [CV 5/5] END max_depth=20, min_samples_leaf=1, min_samples_split=10, n_est imators=200;, score=0.870 total time= 0.2s [CV 4/5] END max_depth=20, min_samples_leaf=1, min_samples_split=10, n_est imators=200;, score=0.915 total time= 0.3s [CV 1/5] END max_depth=20, min_samples_leaf=2, min_samples_split=2, n_esti mators=100;, score=0.928 total time= 0.1s [CV 2/5] END max_depth=20, min_samples_leaf=2, min_samples_split=2, n_esti mators=100;, score=0.974 total time= 0.1s [CV 1/5] END max_depth=20, min_samples_leaf=1, min_samples_split=10, n_est imators=300;, score=0.923 total time= 0.4s [CV 3/5] END max_depth=20, min_samples_leaf=2, min_samples_split=2, n_esti mators=100;, score=0.944 total time= 0.1s [CV 3/5] END max depth=20, min samples leaf=1, min samples split=10, n est imators=300;, score=0.941 total time= 0.3s [CV 2/5] END max_depth=20, min_samples_leaf=1, min_samples_split=10, n_est imators=300;, score=0.977 total time= 0.4s [CV 4/5] END max_depth=20, min_samples_leaf=2, min_samples_split=2, n_esti mators=100;, score=0.916 total time= 0.1s [CV 4/5] END max_depth=20, min_samples_leaf=1, min_samples_split=10, n_est imators=300;, score=0.914 total time= 0.3s [CV 5/5] END max_depth=20, min_samples_leaf=2, min_samples_split=2, n_esti mators=100;, score=0.868 total time= 0.1s [CV 5/5] END max_depth=20, min_samples_leaf=1, min_samples_split=10, n_est imators=300;, score=0.869 total time= 0.3s [CV 2/5] END max_depth=20, min_samples_leaf=2, min_samples_split=2, n_esti mators=200;, score=0.974 total time= 0.2s [CV 3/5] END max_depth=20, min_samples_leaf=2, min_samples_split=2, n_esti mators=200;, score=0.946 total time= 0.3s [CV 4/5] END max_depth=20, min_samples_leaf=2, min_samples_split=2, n_esti mators=200;, score=0.914 total time= 0.3s [CV 1/5] END max_depth=20, min_samples_leaf=2, min_samples_split=2, n_esti mators=200;, score=0.926 total time= 0.3s [CV 5/5] END max_depth=20, min_samples_leaf=2, min_samples_split=2, n_esti mators=200;, score=0.871 total time= 0.2s [CV 2/5] END max_depth=20, min_samples_leaf=2, min_samples_split=5, n_esti mators=100;, score=0.975 total time= 0.2s [CV 1/5] END max_depth=20, min_samples_leaf=2, min_samples_split=5, n_esti mators=100;, score=0.919 total time= 0.2s [CV 3/5] END max_depth=20, min_samples_leaf=2, min_samples_split=5, n_esti mators=100;, score=0.943 total time= 0.2s [CV 1/5] END max_depth=20, min_samples_leaf=2, min_samples_split=2, n_esti mators=300;, score=0.923 total time= 0.4s [CV 3/5] END max_depth=20, min_samples_leaf=2, min_samples_split=2, n_esti

mators=300;, score=0.945 total time= 0.4s [CV 2/5] END max_depth=20, min_samples_leaf=2, min_samples_split=2, n_esti mators=300;, score=0.975 total time= 0.5s [CV 5/5] END max_depth=20, min_samples_leaf=2, min_samples_split=5, n_esti mators=100;, score=0.870 total time= 0.2s [CV 4/5] END max depth=20, min samples leaf=2, min samples split=5, n esti mators=100;, score=0.915 total time= 0.2s [CV 4/5] END max_depth=20, min_samples_leaf=2, min_samples_split=2, n_esti mators=300;, score=0.910 total time= 0.5s [CV 5/5] END max_depth=20, min_samples_leaf=2, min_samples_split=2, n_esti mators=300;, score=0.870 total time= 0.5s [CV 1/5] END max_depth=20, min_samples_leaf=2, min_samples_split=5, n_esti mators=200;, score=0.924 total time= 0.3s [CV 3/5] END max_depth=20, min_samples_leaf=2, min_samples_split=5, n_esti mators=200;, score=0.945 total time= 0.3s [CV 4/5] END max_depth=20, min_samples_leaf=2, min_samples_split=5, n_esti mators=200;, score=0.914 total time= 0.3s [CV 5/5] END max_depth=20, min_samples_leaf=2, min_samples_split=5, n_esti mators=200;, score=0.874 total time= 0.3s [CV 2/5] END max_depth=20, min_samples_leaf=2, min_samples_split=5, n_esti mators=200;, score=0.977 total time= 0.4s [CV 1/5] END max_depth=20, min_samples_leaf=2, min_samples_split=10, n_est imators=100;, score=0.924 total time= 0.1s [CV 1/5] END max_depth=20, min_samples_leaf=2, min_samples_split=5, n_esti mators=300;, score=0.920 total time= 0.4s [CV 2/5] END max_depth=20, min_samples_leaf=2, min_samples_split=10, n_est imators=100;, score=0.978 total time= 0.1s [CV 3/5] END max_depth=20, min_samples_leaf=2, min_samples_split=10, n_est imators=100;, score=0.937 total time= 0.1s [CV 2/5] END max depth=20, min samples leaf=2, min samples split=5, n esti mators=300;, score=0.976 total time= 0.4s [CV 4/5] END max_depth=20, min_samples_leaf=2, min_samples_split=10, n_est imators=100;, score=0.908 total time= 0.1s [CV 4/5] END max_depth=20, min_samples_leaf=2, min_samples_split=5, n_esti mators=300;, score=0.911 total time= 0.3s [CV 5/5] END max_depth=20, min_samples_leaf=2, min_samples_split=10, n_est imators=100;, score=0.873 total time= 0.1s [CV 3/5] END max_depth=20, min_samples_leaf=2, min_samples_split=5, n_esti mators=300;, score=0.944 total time= 0.4s [CV 5/5] END max_depth=20, min_samples_leaf=2, min_samples_split=5, n_esti mators=300;, score=0.869 total time= 0.4s [CV 1/5] END max_depth=20, min_samples_leaf=2, min_samples_split=10, n_est imators=200;, score=0.928 total time= 0.3s [CV 2/5] END max_depth=20, min_samples_leaf=2, min_samples_split=10, n_est imators=200;, score=0.980 total time= 0.3s [CV 3/5] END max_depth=20, min_samples_leaf=2, min_samples_split=10, n_est imators=200;, score=0.948 total time= 0.3s [CV 4/5] END max_depth=20, min_samples_leaf=2, min_samples_split=10, n_est imators=200;, score=0.911 total time= 0.3s [CV 5/5] END max_depth=20, min_samples_leaf=2, min_samples_split=10, n_est imators=200;, score=0.873 total time= 0.3s [CV 1/5] END max_depth=20, min_samples_leaf=4, min_samples_split=2, n_esti mators=100;, score=0.929 total time= 0.1s [CV 2/5] END max_depth=20, min_samples_leaf=4, min_samples_split=2, n_esti mators=100;, score=0.979 total time= 0.1s [CV 1/5] END max_depth=20, min_samples_leaf=2, min_samples_split=10, n_est imators=300;, score=0.924 total time= 0.4s [CV 3/5] END max_depth=20, min_samples_leaf=4, min_samples_split=2, n_esti mators=100;, score=0.945 total time= 0.1s [CV 3/5] END max_depth=20, min_samples_leaf=2, min_samples_split=10, n_est

imators=300;, score=0.946 total time= 0.4s [CV 4/5] END max_depth=20, min_samples_leaf=4, min_samples_split=2, n_esti mators=100;, score=0.912 total time= 0.2s [CV 5/5] END max_depth=20, min_samples_leaf=4, min_samples_split=2, n_esti mators=100;, score=0.870 total time= 0.2s [CV 2/5] END max depth=20, min samples leaf=2, min samples split=10, n est imators=300;, score=0.978 total time= 0.6s [CV 4/5] END max_depth=20, min_samples_leaf=2, min_samples_split=10, n_est imators=300;, score=0.911 total time= 0.4s [CV 5/5] END max_depth=20, min_samples_leaf=2, min_samples_split=10, n_est imators=300;, score=0.870 total time= 0.4s [CV 1/5] END max_depth=20, min_samples_leaf=4, min_samples_split=2, n_esti mators=200;, score=0.929 total time= 0.3s [CV 2/5] END max_depth=20, min_samples_leaf=4, min_samples_split=2, n_esti mators=200;, score=0.979 total time= 0.3s [CV 3/5] END max_depth=20, min_samples_leaf=4, min_samples_split=2, n_esti mators=200;, score=0.945 total time= 0.3s [CV 4/5] END max_depth=20, min_samples_leaf=4, min_samples_split=2, n_esti mators=200;, score=0.908 total time= 0.3s [CV 5/5] END max_depth=20, min_samples_leaf=4, min_samples_split=2, n_esti mators=200;, score=0.869 total time= 0.2s [CV 1/5] END max_depth=20, min_samples_leaf=4, min_samples_split=5, n_esti mators=100;, score=0.922 total time= 0.1s [CV 3/5] END max_depth=20, min_samples_leaf=4, min_samples_split=5, n_esti mators=100;, score=0.944 total time= 0.1s [CV 2/5] END max_depth=20, min_samples_leaf=4, min_samples_split=5, n_esti mators=100;, score=0.980 total time= 0.1s [CV 1/5] END max_depth=20, min_samples_leaf=4, min_samples_split=2, n_esti mators=300;, score=0.928 total time= 0.4s [CV 2/5] END max depth=20, min samples leaf=4, min samples split=2, n esti mators=300;, score=0.977 total time= 0.4s [CV 4/5] END max depth=20, min samples leaf=4, min samples split=5, n esti mators=100;, score=0.908 total time= 0.1s [CV 3/5] END max_depth=20, min_samples_leaf=4, min_samples_split=2, n_esti mators=300;, score=0.942 total time= 0.4s [CV 5/5] END max_depth=20, min_samples_leaf=4, min_samples_split=2, n_esti mators=300;, score=0.871 total time= 0.3s [CV 5/5] END max_depth=20, min_samples_leaf=4, min_samples_split=5, n_esti mators=100;, score=0.872 total time= 0.1s [CV 4/5] END max_depth=20, min_samples_leaf=4, min_samples_split=2, n_esti mators=300;, score=0.913 total time= 0.4s [CV 1/5] END max_depth=20, min_samples_leaf=4, min_samples_split=5, n_esti mators=200;, score=0.925 total time= 0.3s [CV 2/5] END max_depth=20, min_samples_leaf=4, min_samples_split=5, n_esti mators=200;, score=0.978 total time= 0.3s [CV 3/5] END max_depth=20, min_samples_leaf=4, min_samples_split=5, n_esti mators=200;, score=0.942 total time= 0.2s [CV 5/5] END max_depth=20, min_samples_leaf=4, min_samples_split=5, n_esti mators=200;, score=0.875 total time= 0.3s [CV 4/5] END max_depth=20, min_samples_leaf=4, min_samples_split=5, n_esti mators=200;, score=0.912 total time= 0.3s [CV 1/5] END max_depth=20, min_samples_leaf=4, min_samples_split=10, n_est imators=100;, score=0.924 total time= 0.1s [CV 2/5] END max_depth=20, min_samples_leaf=4, min_samples_split=10, n_est imators=100;, score=0.979 total time= 0.1s [CV 3/5] END max_depth=20, min_samples_leaf=4, min_samples_split=5, n_esti mators=300;, score=0.944 total time= 0.3s [CV 1/5] END max_depth=20, min_samples_leaf=4, min_samples_split=5, n_esti mators=300;, score=0.926 total time= 0.4s [CV 3/5] END max_depth=20, min_samples_leaf=4, min_samples_split=10, n_est

imators=100;, score=0.947 total time= 0.1s [CV 2/5] END max_depth=20, min_samples_leaf=4, min_samples_split=5, n_esti mators=300;, score=0.979 total time= 0.4s [CV 4/5] END max_depth=20, min_samples_leaf=4, min_samples_split=10, n_est imators=100;, score=0.908 total time= 0.1s [CV 4/5] END max depth=20, min samples leaf=4, min samples split=5, n esti mators=300;, score=0.910 total time= 0.3s [CV 5/5] END max_depth=20, min_samples_leaf=4, min_samples_split=10, n_est imators=100;, score=0.875 total time= 0.1s [CV 5/5] END max_depth=20, min_samples_leaf=4, min_samples_split=5, n_esti mators=300;, score=0.877 total time= 0.4s [CV 2/5] END max_depth=20, min_samples_leaf=4, min_samples_split=10, n_est imators=200;, score=0.979 total time= 0.2s [CV 3/5] END max_depth=20, min_samples_leaf=4, min_samples_split=10, n_est imators=200;, score=0.944 total time= 0.2s [CV 4/5] END max_depth=20, min_samples_leaf=4, min_samples_split=10, n_est imators=200;, score=0.908 total time= 0.2s [CV 5/5] END max_depth=20, min_samples_leaf=4, min_samples_split=10, n_est imators=200;, score=0.873 total time= 0.2s [CV 1/5] END max_depth=30, min_samples_leaf=1, min_samples_split=2, n_esti mators=100;, score=0.915 total time= 0.1s [CV 1/5] END max_depth=20, min_samples_leaf=4, min_samples_split=10, n_est imators=200;, score=0.923 total time= 0.4s [CV 1/5] END max_depth=20, min_samples_leaf=4, min_samples_split=10, n_est imators=300;, score=0.927 total time= 0.3s [CV 2/5] END max_depth=20, min_samples_leaf=4, min_samples_split=10, n_est imators=300;, score=0.978 total time= 0.3s [CV 2/5] END max_depth=30, min_samples_leaf=1, min_samples_split=2, n_esti mators=100;, score=0.970 total time= 0.2s [CV 3/5] END max depth=20, min samples leaf=4, min samples split=10, n est imators=300;, score=0.943 total time= 0.3s [CV 4/5] END max_depth=30, min_samples_leaf=1, min_samples_split=2, n_esti mators=100;, score=0.915 total time= 0.1s [CV 3/5] END max_depth=30, min_samples_leaf=1, min_samples_split=2, n_esti mators=100;, score=0.939 total time= 0.1s [CV 5/5] END max_depth=30, min_samples_leaf=1, min_samples_split=2, n_esti mators=100;, score=0.870 total time= 0.2s [CV 5/5] END max_depth=20, min_samples_leaf=4, min_samples_split=10, n_est imators=300;, score=0.872 total time= 0.4s [CV 4/5] END max_depth=20, min_samples_leaf=4, min_samples_split=10, n_est imators=300;, score=0.911 total time= 0.5s [CV 2/5] END max_depth=30, min_samples_leaf=1, min_samples_split=2, n_esti mators=200;, score=0.973 total time= 0.3s [CV 1/5] END max_depth=30, min_samples_leaf=1, min_samples_split=2, n_esti mators=200;, score=0.920 total time= 0.3s [CV 3/5] END max_depth=30, min_samples_leaf=1, min_samples_split=2, n_esti mators=200;, score=0.944 total time= 0.3s [CV 4/5] END max_depth=30, min_samples_leaf=1, min_samples_split=2, n_esti mators=200;, score=0.919 total time= 0.3s [CV 5/5] END max_depth=30, min_samples_leaf=1, min_samples_split=2, n_esti mators=200;, score=0.868 total time= 0.3s [CV 1/5] END max_depth=30, min_samples_leaf=1, min_samples_split=5, n_esti mators=100;, score=0.926 total time= 0.1s [CV 2/5] END max_depth=30, min_samples_leaf=1, min_samples_split=5, n_esti mators=100;, score=0.972 total time= 0.1s [CV 3/5] END max_depth=30, min_samples_leaf=1, min_samples_split=5, n_esti mators=100;, score=0.944 total time= 0.2s [CV 1/5] END max_depth=30, min_samples_leaf=1, min_samples_split=2, n_esti mators=300;, score=0.919 total time= 0.4s [CV 2/5] END max_depth=30, min_samples_leaf=1, min_samples_split=2, n_esti

mators=300;, score=0.977 total time= 0.4s [CV 3/5] END max_depth=30, min_samples_leaf=1, min_samples_split=2, n_esti mators=300;, score=0.938 total time= 0.4s [CV 4/5] END max_depth=30, min_samples_leaf=1, min_samples_split=5, n_esti mators=100;, score=0.909 total time= 0.1s [CV 5/5] END max depth=30, min samples leaf=1, min samples split=2, n esti mators=300;, score=0.875 total time= 0.4s [CV 4/5] END max_depth=30, min_samples_leaf=1, min_samples_split=2, n_esti mators=300;, score=0.916 total time= 0.4s [CV 5/5] END max_depth=30, min_samples_leaf=1, min_samples_split=5, n_esti mators=100;, score=0.870 total time= 0.2s [CV 1/5] END max_depth=30, min_samples_leaf=1, min_samples_split=5, n_esti mators=200;, score=0.920 total time= 0.2s [CV 2/5] END max_depth=30, min_samples_leaf=1, min_samples_split=5, n_esti mators=200;, score=0.976 total time= 0.3s [CV 3/5] END max_depth=30, min_samples_leaf=1, min_samples_split=5, n_esti mators=200;, score=0.944 total time= 0.3s [CV 4/5] END max_depth=30, min_samples_leaf=1, min_samples_split=5, n_esti mators=200;, score=0.910 total time= 0.2s [CV 5/5] END max_depth=30, min_samples_leaf=1, min_samples_split=5, n_esti mators=200;, score=0.877 total time= 0.2s [CV 2/5] END max_depth=30, min_samples_leaf=1, min_samples_split=10, n_est imators=100;, score=0.978 total time= 0.2s [CV 1/5] END max_depth=30, min_samples_leaf=1, min_samples_split=10, n_est imators=100;, score=0.922 total time= 0.2s [CV 3/5] END max_depth=30, min_samples_leaf=1, min_samples_split=10, n_est imators=100;, score=0.942 total time= 0.1s [CV 3/5] END max_depth=30, min_samples_leaf=1, min_samples_split=5, n_esti mators=300;, score=0.942 total time= 0.3s [CV 1/5] END max depth=30, min samples leaf=1, min samples split=5, n esti mators=300;, score=0.921 total time= 0.4s [CV 2/5] END max_depth=30, min_samples_leaf=1, min_samples_split=5, n_esti mators=300;, score=0.977 total time= 0.4s [CV 4/5] END max_depth=30, min_samples_leaf=1, min_samples_split=10, n_est imators=100;, score=0.915 total time= 0.1s [CV 5/5] END max_depth=30, min_samples_leaf=1, min_samples_split=10, n_est imators=100;, score=0.868 total time= 0.1s [CV 4/5] END max_depth=30, min_samples_leaf=1, min_samples_split=5, n_esti mators=300;, score=0.917 total time= 0.4s [CV 5/5] END max_depth=30, min_samples_leaf=1, min_samples_split=5, n_esti mators=300;, score=0.870 total time= 0.4s [CV 3/5] END max_depth=30, min_samples_leaf=1, min_samples_split=10, n_est imators=200;, score=0.946 total time= 0.2s [CV 1/5] END max_depth=30, min_samples_leaf=1, min_samples_split=10, n_est imators=200;, score=0.920 total time= 0.3s [CV 2/5] END max_depth=30, min_samples_leaf=1, min_samples_split=10, n_est imators=200;, score=0.977 total time= 0.3s [CV 4/5] END max_depth=30, min_samples_leaf=1, min_samples_split=10, n_est imators=200;, score=0.912 total time= 0.2s [CV 5/5] END max_depth=30, min_samples_leaf=1, min_samples_split=10, n_est imators=200;, score=0.870 total time= 0.2s [CV 1/5] END max_depth=30, min_samples_leaf=2, min_samples_split=2, n_esti mators=100;, score=0.926 total time= 0.1s [CV 2/5] END max_depth=30, min_samples_leaf=2, min_samples_split=2, n_esti mators=100;, score=0.974 total time= 0.1s [CV 1/5] END max_depth=30, min_samples_leaf=1, min_samples_split=10, n_est imators=300;, score=0.922 total time= 0.3s [CV 3/5] END max_depth=30, min_samples_leaf=2, min_samples_split=2, n_esti mators=100;, score=0.947 total time= 0.1s [CV 2/5] END max_depth=30, min_samples_leaf=1, min_samples_split=10, n_est

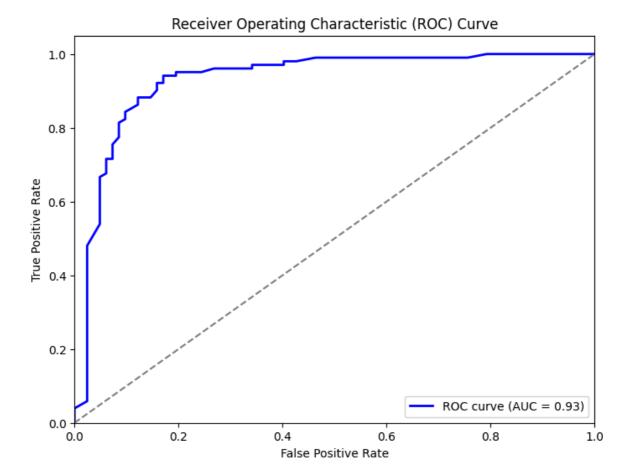
imators=300;, score=0.977 total time= 0.4s [CV 4/5] END max_depth=30, min_samples_leaf=2, min_samples_split=2, n_esti mators=100;, score=0.910 total time= 0.1s [CV 5/5] END max_depth=30, min_samples_leaf=2, min_samples_split=2, n_esti mators=100;, score=0.870 total time= 0.1s [CV 3/5] END max depth=30, min samples leaf=1, min samples split=10, n est imators=300;, score=0.945 total time= 0.4s [CV 4/5] END max_depth=30, min_samples_leaf=1, min_samples_split=10, n_est imators=300;, score=0.909 total time= 0.4s [CV 5/5] END max_depth=30, min_samples_leaf=1, min_samples_split=10, n_est imators=300;, score=0.872 total time= 0.4s [CV 1/5] END max_depth=30, min_samples_leaf=2, min_samples_split=2, n_esti mators=200;, score=0.923 total time= 0.3s [CV 2/5] END max_depth=30, min_samples_leaf=2, min_samples_split=2, n_esti mators=200;, score=0.975 total time= 0.3s [CV 4/5] END max_depth=30, min_samples_leaf=2, min_samples_split=2, n_esti mators=200;, score=0.909 total time= 0.2s [CV 3/5] END max_depth=30, min_samples_leaf=2, min_samples_split=2, n_esti mators=200;, score=0.947 total time= 0.3s [CV 5/5] END max_depth=30, min_samples_leaf=2, min_samples_split=2, n_esti mators=200;, score=0.868 total time= 0.2s [CV 1/5] END max_depth=30, min_samples_leaf=2, min_samples_split=5, n_esti mators=100;, score=0.920 total time= 0.1s [CV 2/5] END max_depth=30, min_samples_leaf=2, min_samples_split=5, n_esti mators=100;, score=0.975 total time= 0.1s [CV 3/5] END max_depth=30, min_samples_leaf=2, min_samples_split=5, n_esti mators=100;, score=0.942 total time= 0.1s [CV 1/5] END max_depth=30, min_samples_leaf=2, min_samples_split=2, n_esti mators=300;, score=0.925 total time= 0.4s [CV 4/5] END max depth=30, min samples leaf=2, min samples split=5, n esti mators=100;, score=0.913 total time= 0.1s [CV 5/5] END max_depth=30, min_samples_leaf=2, min_samples_split=5, n_esti mators=100;, score=0.875 total time= 0.1s [CV 2/5] END max_depth=30, min_samples_leaf=2, min_samples_split=2, n_esti mators=300;, score=0.974 total time= 0.4s [CV 3/5] END max_depth=30, min_samples_leaf=2, min_samples_split=2, n_esti mators=300;, score=0.946 total time= 0.4s [CV 4/5] END max_depth=30, min_samples_leaf=2, min_samples_split=2, n_esti mators=300;, score=0.915 total time= 0.4s [CV 1/5] END max_depth=30, min_samples_leaf=2, min_samples_split=5, n_esti mators=200;, score=0.927 total time= 0.2s [CV 5/5] END max_depth=30, min_samples_leaf=2, min_samples_split=2, n_esti mators=300;, score=0.873 total time= 0.4s [CV 2/5] END max_depth=30, min_samples_leaf=2, min_samples_split=5, n_esti mators=200;, score=0.975 total time= 0.2s [CV 3/5] END max_depth=30, min_samples_leaf=2, min_samples_split=5, n_esti mators=200;, score=0.943 total time= 0.2s [CV 4/5] END max_depth=30, min_samples_leaf=2, min_samples_split=5, n_esti mators=200;, score=0.916 total time= 0.2s [CV 5/5] END max_depth=30, min_samples_leaf=2, min_samples_split=5, n_esti mators=200;, score=0.872 total time= 0.2s [CV 1/5] END max_depth=30, min_samples_leaf=2, min_samples_split=10, n_est imators=100;, score=0.926 total time= 0.1s [CV 2/5] END max_depth=30, min_samples_leaf=2, min_samples_split=10, n_est imators=100;, score=0.979 total time= 0.1s [CV 3/5] END max_depth=30, min_samples_leaf=2, min_samples_split=10, n_est imators=100;, score=0.944 total time= 0.1s [CV 1/5] END max_depth=30, min_samples_leaf=2, min_samples_split=5, n_esti mators=300;, score=0.923 total time= 0.4s [CV 2/5] END max_depth=30, min_samples_leaf=2, min_samples_split=5, n_esti

mators=300;, score=0.976 total time= 0.4s [CV 4/5] END max_depth=30, min_samples_leaf=2, min_samples_split=10, n_est imators=100;, score=0.908 total time= 0.1s [CV 4/5] END max_depth=30, min_samples_leaf=2, min_samples_split=5, n_esti mators=300;, score=0.915 total time= 0.3s [CV 5/5] END max depth=30, min samples leaf=2, min samples split=10, n est imators=100;, score=0.871 total time= 0.1s [CV 3/5] END max_depth=30, min_samples_leaf=2, min_samples_split=5, n_esti mators=300;, score=0.944 total time= 0.4s [CV 5/5] END max_depth=30, min_samples_leaf=2, min_samples_split=5, n_esti mators=300;, score=0.871 total time= 0.4s [CV 2/5] END max_depth=30, min_samples_leaf=2, min_samples_split=10, n_est imators=200;, score=0.979 total time= 0.2s [CV 1/5] END max_depth=30, min_samples_leaf=2, min_samples_split=10, n_est imators=200;, score=0.923 total time= 0.3s [CV 4/5] END max_depth=30, min_samples_leaf=2, min_samples_split=10, n_est imators=200;, score=0.914 total time= 0.2s [CV 3/5] END max_depth=30, min_samples_leaf=2, min_samples_split=10, n_est imators=200;, score=0.943 total time= 0.3s [CV 5/5] END max_depth=30, min_samples_leaf=2, min_samples_split=10, n_est imators=200;, score=0.870 total time= 0.2s [CV 1/5] END max_depth=30, min_samples_leaf=4, min_samples_split=2, n_esti mators=100;, score=0.920 total time= 0.1s [CV 2/5] END max_depth=30, min_samples_leaf=4, min_samples_split=2, n_esti mators=100;, score=0.979 total time= 0.1s [CV 3/5] END max_depth=30, min_samples_leaf=4, min_samples_split=2, n_esti mators=100;, score=0.942 total time= 0.1s [CV 1/5] END max_depth=30, min_samples_leaf=2, min_samples_split=10, n_est imators=300;, score=0.927 total time= 0.4s [CV 3/5] END max depth=30, min samples leaf=2, min samples split=10, n est imators=300;, score=0.946 total time= 0.3s [CV 2/5] END max_depth=30, min_samples_leaf=2, min_samples_split=10, n_est imators=300;, score=0.978 total time= 0.4s [CV 4/5] END max_depth=30, min_samples_leaf=4, min_samples_split=2, n_esti mators=100;, score=0.910 total time= 0.1s [CV 5/5] END max_depth=30, min_samples_leaf=4, min_samples_split=2, n_esti mators=100;, score=0.872 total time= 0.1s [CV 4/5] END max_depth=30, min_samples_leaf=2, min_samples_split=10, n_est imators=300;, score=0.910 total time= 0.4s [CV 5/5] END max_depth=30, min_samples_leaf=2, min_samples_split=10, n_est imators=300;, score=0.873 total time= 0.5s [CV 1/5] END max_depth=30, min_samples_leaf=4, min_samples_split=2, n_esti mators=200;, score=0.925 total time= 0.3s [CV 2/5] END max_depth=30, min_samples_leaf=4, min_samples_split=2, n_esti mators=200;, score=0.980 total time= 0.4s [CV 3/5] END max_depth=30, min_samples_leaf=4, min_samples_split=2, n_esti mators=200;, score=0.944 total time= 0.4s [CV 5/5] END max_depth=30, min_samples_leaf=4, min_samples_split=2, n_esti mators=200;, score=0.873 total time= 0.4s [CV 4/5] END max_depth=30, min_samples_leaf=4, min_samples_split=2, n_esti mators=200;, score=0.912 total time= 0.4s [CV 2/5] END max_depth=30, min_samples_leaf=4, min_samples_split=5, n_esti mators=100;, score=0.977 total time= 0.1s [CV 4/5] END max_depth=30, min_samples_leaf=4, min_samples_split=5, n_esti mators=100;, score=0.906 total time= 0.1s [CV 1/5] END max_depth=30, min_samples_leaf=4, min_samples_split=5, n_esti mators=100;, score=0.928 total time= 0.3s [CV 3/5] END max_depth=30, min_samples_leaf=4, min_samples_split=5, n_esti mators=100;, score=0.944 total time= 0.2s

[CV 4/5] END max_depth=30, min_samples_leaf=4, min_samples_split=2, n_esti

mators=300;, score=0.911 total time= 0.4s [CV 1/5] END max_depth=30, min_samples_leaf=4, min_samples_split=2, n_esti mators=300;, score=0.929 total time= 0.7s [CV 2/5] END max_depth=30, min_samples_leaf=4, min_samples_split=2, n_esti mators=300;, score=0.980 total time= 0.6s [CV 5/5] END max depth=30, min samples leaf=4, min samples split=5, n esti mators=100;, score=0.871 total time= 0.2s [CV 3/5] END max_depth=30, min_samples_leaf=4, min_samples_split=2, n_esti mators=300;, score=0.943 total time= 0.6s [CV 5/5] END max_depth=30, min_samples_leaf=4, min_samples_split=2, n_esti mators=300;, score=0.874 total time= 0.6s [CV 1/5] END max depth=30, min samples leaf=4, min samples split=5, n esti mators=200;, score=0.924 total time= 0.3s [CV 3/5] END max_depth=30, min_samples_leaf=4, min_samples_split=5, n_esti mators=200;, score=0.945 total time= 0.4s [CV 2/5] END max_depth=30, min_samples_leaf=4, min_samples_split=5, n_esti mators=200;, score=0.977 total time= 0.4s [CV 5/5] END max_depth=30, min_samples_leaf=4, min_samples_split=5, n_esti mators=200;, score=0.873 total time= 0.4s [CV 4/5] END max_depth=30, min_samples_leaf=4, min_samples_split=5, n_esti mators=200;, score=0.908 total time= 0.4s [CV 1/5] END max_depth=30, min_samples_leaf=4, min_samples_split=10, n_est imators=100;, score=0.920 total time= 0.2s [CV 3/5] END max_depth=30, min_samples_leaf=4, min_samples_split=10, n_est imators=100;, score=0.943 total time= 0.1s [CV 2/5] END max_depth=30, min_samples_leaf=4, min_samples_split=10, n_est imators=100;, score=0.979 total time= 0.2s [CV 1/5] END max_depth=30, min_samples_leaf=4, min_samples_split=5, n_esti mators=300;, score=0.926 total time= 0.5s [CV 4/5] END max depth=30, min samples leaf=4, min samples split=10, n est imators=100;, score=0.914 total time= 0.1s [CV 2/5] END max depth=30, min samples leaf=4, min samples split=5, n esti mators=300;, score=0.979 total time= 0.4s [CV 5/5] END max_depth=30, min_samples_leaf=4, min_samples_split=10, n_est imators=100;, score=0.868 total time= 0.1s [CV 3/5] END max_depth=30, min_samples_leaf=4, min_samples_split=5, n_esti mators=300;, score=0.944 total time= 0.5s [CV 4/5] END max_depth=30, min_samples_leaf=4, min_samples_split=5, n_esti mators=300;, score=0.908 total time= 0.4s [CV 5/5] END max_depth=30, min_samples_leaf=4, min_samples_split=5, n_esti mators=300;, score=0.873 total time= 0.5s [CV 2/5] END max_depth=30, min_samples_leaf=4, min_samples_split=10, n_est imators=200;, score=0.980 total time= 0.3s [CV 1/5] END max_depth=30, min_samples_leaf=4, min_samples_split=10, n_est imators=200;, score=0.929 total time= 0.3s [CV 4/5] END max_depth=30, min_samples_leaf=4, min_samples_split=10, n_est imators=200;, score=0.909 total time= 0.3s [CV 3/5] END max_depth=30, min_samples_leaf=4, min_samples_split=10, n_est imators=200;, score=0.943 total time= 0.3s [CV 5/5] END max_depth=30, min_samples_leaf=4, min_samples_split=10, n_est imators=200;, score=0.874 total time= 0.4s [CV 1/5] END max_depth=30, min_samples_leaf=4, min_samples_split=10, n_est imators=300;, score=0.931 total time= 0.4s [CV 2/5] END max_depth=30, min_samples_leaf=4, min_samples_split=10, n_est imators=300;, score=0.979 total time= 0.4s [CV 5/5] END max_depth=30, min_samples_leaf=4, min_samples_split=10, n_est imators=300;, score=0.873 total time= 0.4s [CV 3/5] END max_depth=30, min_samples_leaf=4, min_samples_split=10, n_est imators=300;, score=0.943 total time= 0.4s

```
[CV 4/5] END max_depth=30, min_samples_leaf=4, min_samples_split=10, n_est
       imators=300;, score=0.906 total time= 0.4s
Out[]:
                    GridSearchCV
         ▶ estimator: RandomForestClassifier
             RandomForestClassifier
In []:
       # get best combination and result
        print("Beste Hyperparameter-Kombinationen: ", grid_search.best_params_)
        print("Beste Performance: ", grid_search.best_score_)
       Beste Hyperparameter-Kombinationen: {'max_depth': None, 'min_samples_lea
       f': 2, 'min_samples_split': 10, 'n_estimators': 100}
       Beste Performance: 0.928276497165386
In [ ]: target_pred_proba = forest.predict_proba(features_test)[:, 1]
        # compute ROC curve and ROC-AUC score
        fpr, tpr, thresholds = roc_curve(target_test, target_pred_proba)
        roc_auc = roc_auc_score(target_test, target_pred_proba)
        # plot ROC curve
        plt.figure(figsize=(8, 6))
        plt.plot(fpr, tpr, color='blue', lw=2, label='ROC curve (AUC = %0.2f)' %
        plt.plot([0, 1], [0, 1], color='gray', linestyle='--')
        plt.xlim([0.0, 1.0])
        plt.ylim([0.0, 1.05])
        plt.xlabel('False Positive Rate')
        plt.ylabel('True Positive Rate')
        plt.title('Receiver Operating Characteristic (ROC) Curve')
        plt.legend(loc="lower right")
        plt.show()
```



Evaluation und Ergebnisdarstellung

Basierend auf der Evaluation der drei Klassifikationsmodelle – Random Forest Classifier, Logistische Regression und Support Vector Machine – erzielte der RFC vor dem Parameter-Tuning die beste Leistung mit einer Genauigkeit von 88.59% und einem ROC AUC Score von 88.27%. Der precision, recall und f1-score für beide Klassen (Herzkrankheit und Normal) zeigen eine ausgeglichene Leistung des Modells. Nach dem Parameter-Tuning wurden die Hyperparameter des RFC optimiert, wodurch eine verbesserte Leistung mit einer ROC AUC Score von 92.78% erzielt wurde. Dies unterstreicht die Wirksamkeit des gewählten Ansatzes und die Fähigkeit des Modells, zwischen Herzkrankheit und Normalzustand zu unterscheiden.

Vorhersage-Demo

```
In []: selected_data_point = features_train.iloc[0:1, :]
    selected_target = target_train.iloc[0]

# make the prediction for the chosen data point
    prediction = forest.predict(selected_data_point)

# disply the chosen data point, the true class and the predicted class
    print("Ausgewählter Datenpunkt:")
    print(selected_data_point)
```

print("\nWahre Klasse des ausgewählten Datenpunkts:", selected_target)
print("\nVorhersage für den ausgewählten Datenpunkt:", prediction)

```
Ausgewählter Datenpunkt:
```

```
Age RestingBP Cholesterol FastingBS MaxHR Oldpeak Sex_F Sex_M \
161 49 128 212 0 96 0.0 False True ChestPainType_ASY ChestPainType_ATA ChestPainType_NAP \
161 True False False
```

```
ChestPainType_TA RestingECG_LVH RestingECG_Normal RestingECG_ST \
161 False False True False
```

Wahre Klasse des ausgewählten Datenpunkts: 1

Vorhersage für den ausgewählten Datenpunkt: [1]

Wie man anhand des Beispiels sieht, erkennt das Model den gegeben Datenpunkt korrekt als Herzerkrankung an.