

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
import tensorflow as tf
from tensorflow import keras
```

```
df=pd.read_csv(r"C:\Users\arjun\Downloads\
heart_failure_clinical_records_dataset.csv")
df.head()
```

	age	anaemia	creatinine_phosphokinase	diabetes
0	75.0	0	582	0
20				
1	55.0	0	7861	0
38				
2	65.0	0	146	0
20				
3	50.0	1	111	0
20				
4	65.0	1	160	1
20				

	high_blood_pressure	platelets	serum_creatinine	serum_sodium	sex
0	1	265000.00	1.9	130	1
1	0	263358.03	1.1	136	1
2	0	162000.00	1.3	129	1
3	0	210000.00	1.9	137	1
4	0	327000.00	2.7	116	0

	smoking	time	DEATH_EVENT
0	0	4	1
1	0	6	1
2	1	7	1
3	0	7	1
4	0	8	1

```
df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 299 entries, 0 to 298
Data columns (total 13 columns):
```

#	Column	Non-Null Count	Dtype
0	age	299 non-null	float64
1	anaemia	299 non-null	int64
2	creatinine_phosphokinase	299 non-null	int64
3	diabetes	299 non-null	int64
4	ejection_fraction	299 non-null	int64
5	high_blood_pressure	299 non-null	int64
6	platelets	299 non-null	float64
7	serum_creatinine	299 non-null	float64
8	serum_sodium	299 non-null	int64
9	sex	299 non-null	int64
10	smoking	299 non-null	int64
11	time	299 non-null	int64
12	DEATH_EVENT	299 non-null	int64

dtypes: float64(3), int64(10)
memory usage: 30.5 KB

df.describe()

	age	anaemia	creatinine_phosphokinase	diabetes \
count	299.000000	299.000000	299.000000	299.000000
mean	60.833893	0.431438	581.839465	0.418060
std	11.894809	0.496107	970.287881	0.494067
min	40.000000	0.000000	23.000000	0.000000
25%	51.000000	0.000000	116.500000	0.000000
50%	60.000000	0.000000	250.000000	0.000000
75%	70.000000	1.000000	582.000000	1.000000
max	95.000000	1.000000	7861.000000	1.000000

	ejection_fraction	high_blood_pressure	platelets \
count	299.000000	299.000000	299.000000
mean	38.083612	0.351171	263358.029264
std	11.834841	0.478136	97804.236869
min	14.000000	0.000000	25100.000000
25%	30.000000	0.000000	212500.000000
50%	38.000000	0.000000	262000.000000
75%	45.000000	1.000000	303500.000000
max	80.000000	1.000000	850000.000000

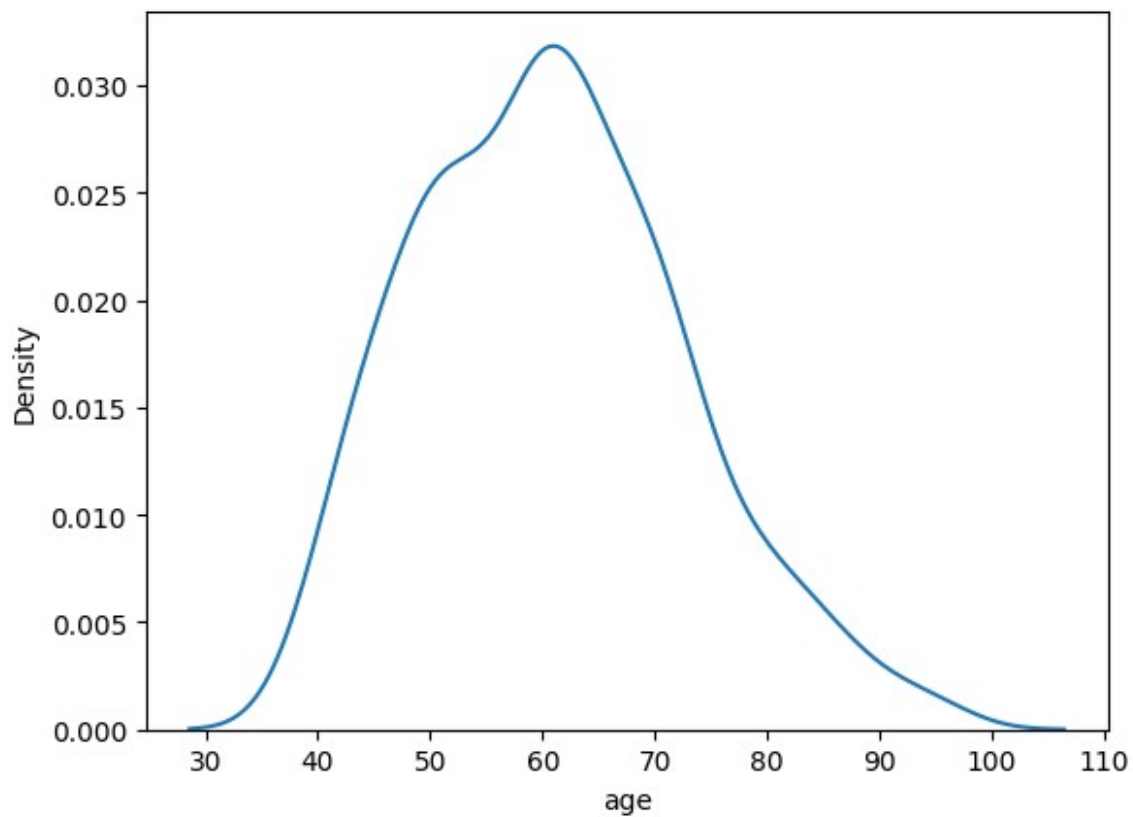
	serum_creatinine	serum_sodium	sex	smoking
time \				
count	299.000000	299.000000	299.000000	299.000000
299.000000				
mean	1.39388	136.625418	0.648829	0.32107
130.260870				
std	1.03451	4.412477	0.478136	0.46767
77.614208				
min	0.50000	113.000000	0.000000	0.00000
4.000000				

25%	0.90000	134.000000	0.000000	0.00000
73.000000				
50%	1.10000	137.000000	1.000000	0.00000
115.000000				
75%	1.40000	140.000000	1.000000	1.00000
203.000000				
max	9.40000	148.000000	1.000000	1.00000
285.000000				

	DEATH_EVENT
count	299.00000
mean	0.32107
std	0.46767
min	0.00000
25%	0.00000
50%	0.00000
75%	1.00000
max	1.00000

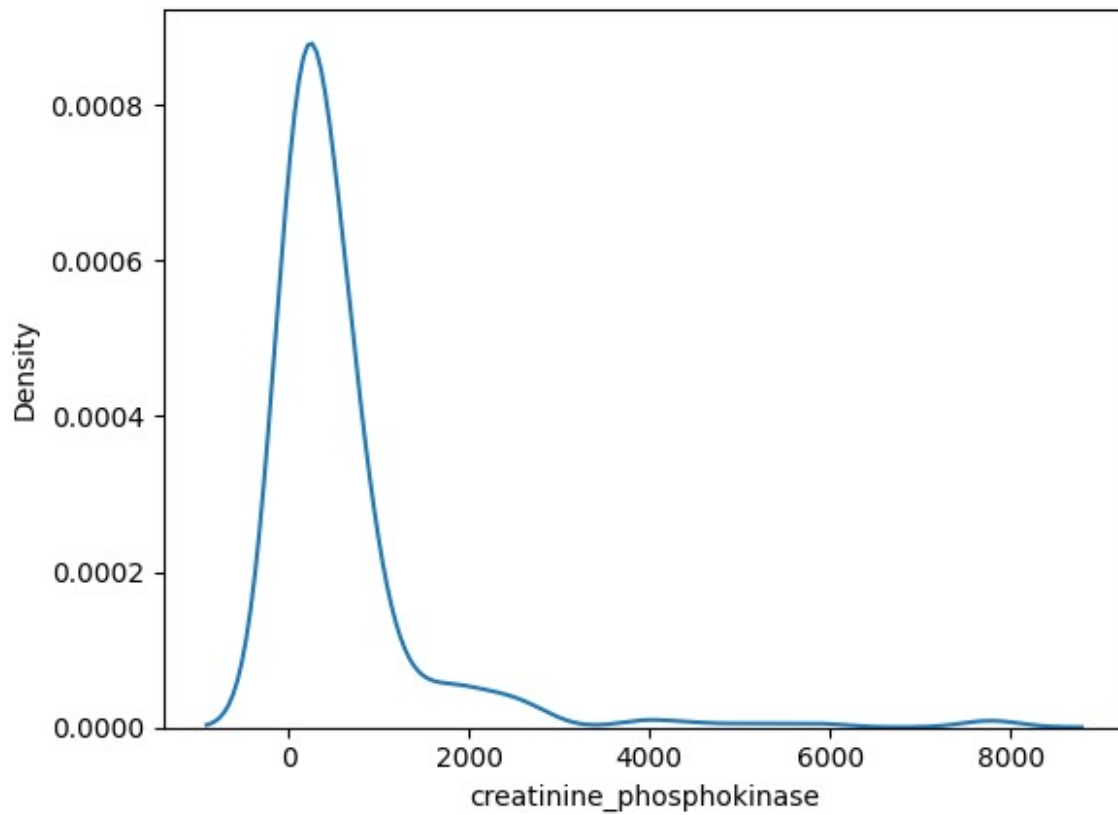
```
sns.kdeplot(df['age'])
```

```
<Axes: xlabel='age', ylabel='Density'>
```



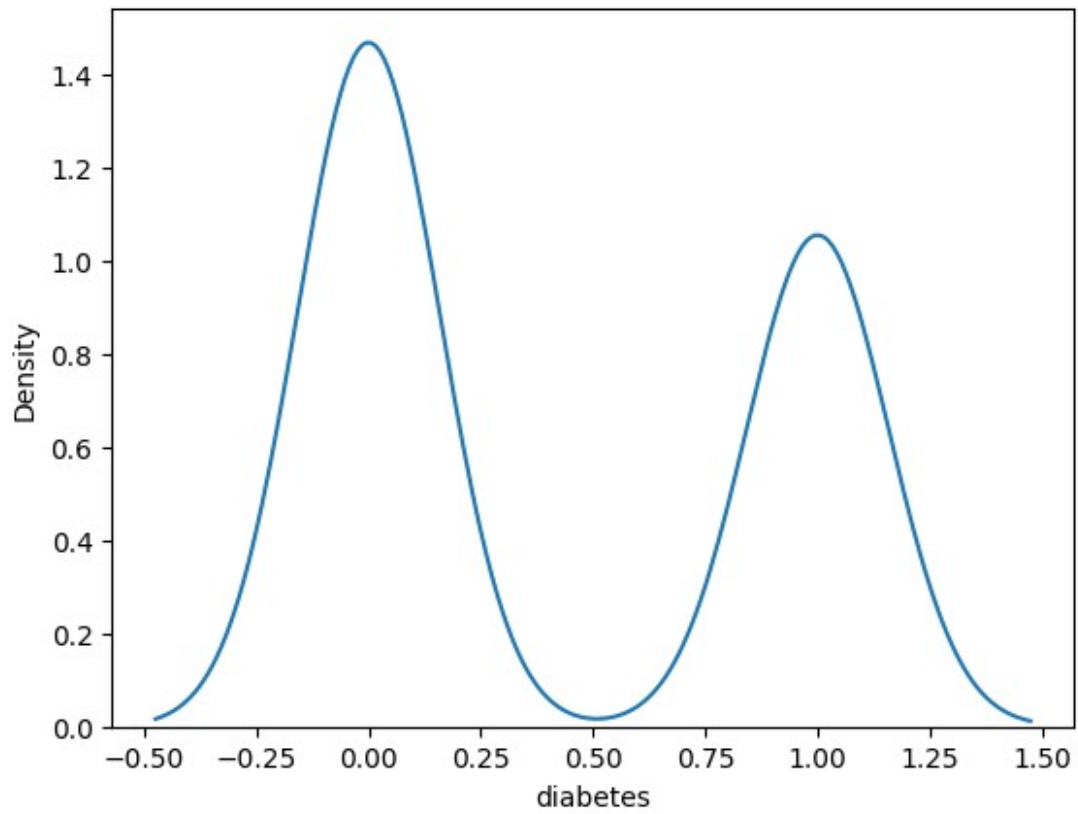
```
sns.kdeplot(df['creatinine_phosphokinase'])
```

```
<Axes: xlabel='creatinine_phosphokinase', ylabel='Density'>
```

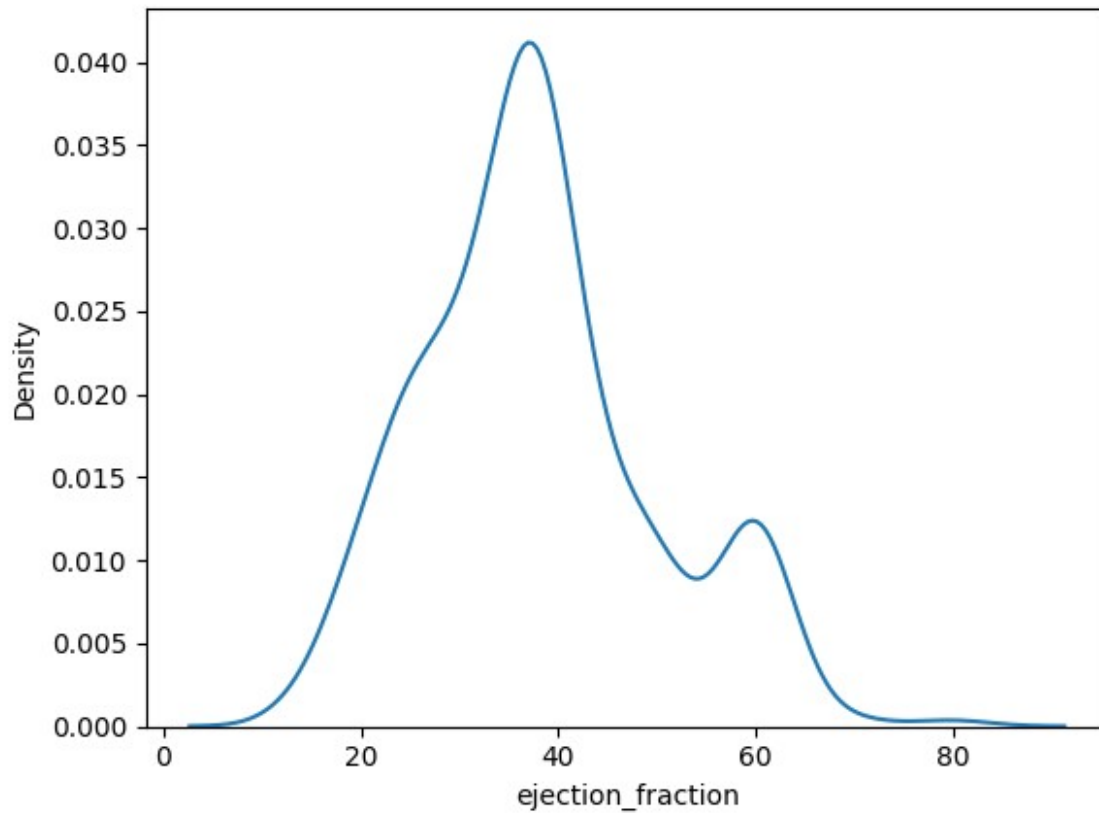


```
sns.kdeplot(df['diabetes'])
```

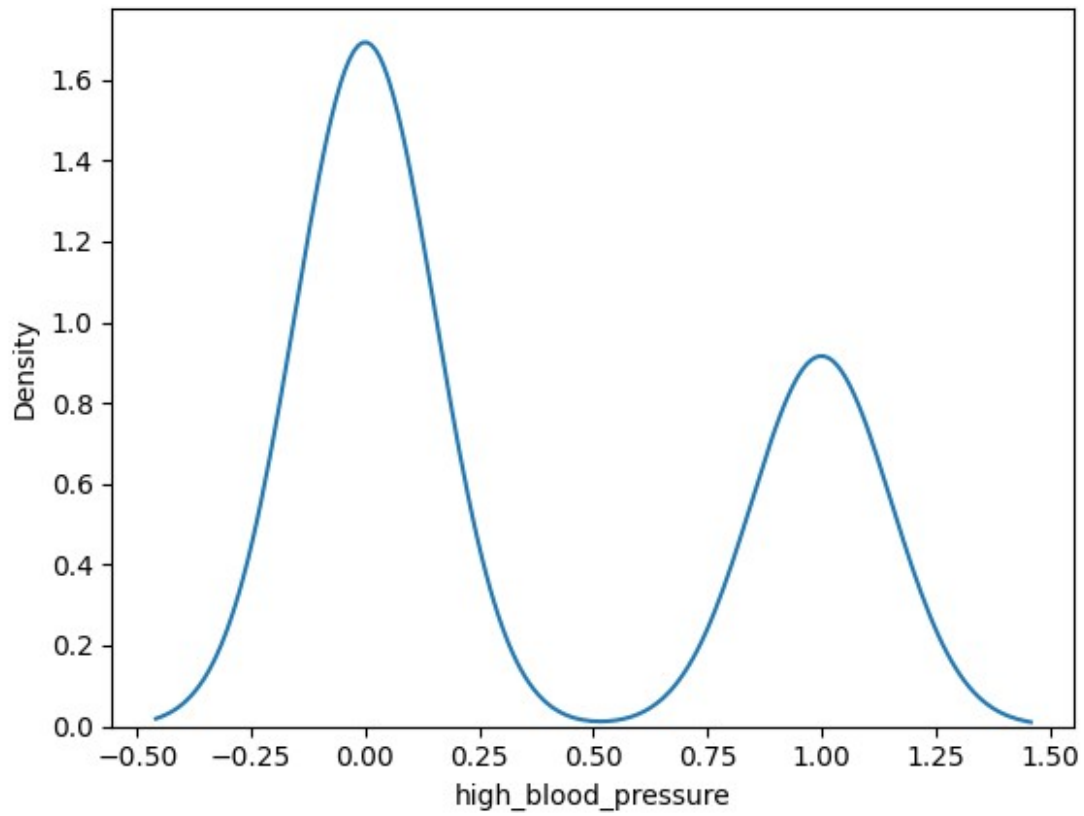
```
<Axes: xlabel='diabetes', ylabel='Density'>
```



```
sns.kdeplot(df['ejection_fraction'])  
<Axes: xlabel='ejection_fraction', ylabel='Density'>
```

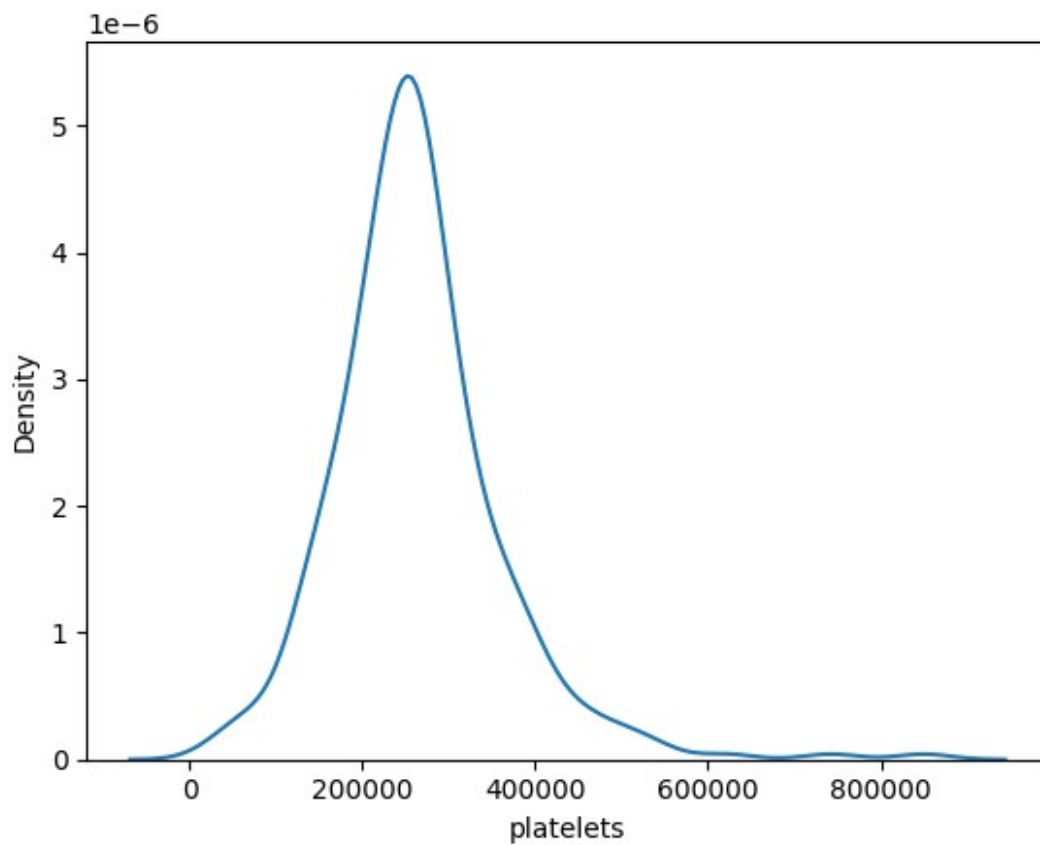


```
sns.kdeplot(df['high_blood_pressure'])  
<Axes: xlabel='high_blood_pressure', ylabel='Density'>
```

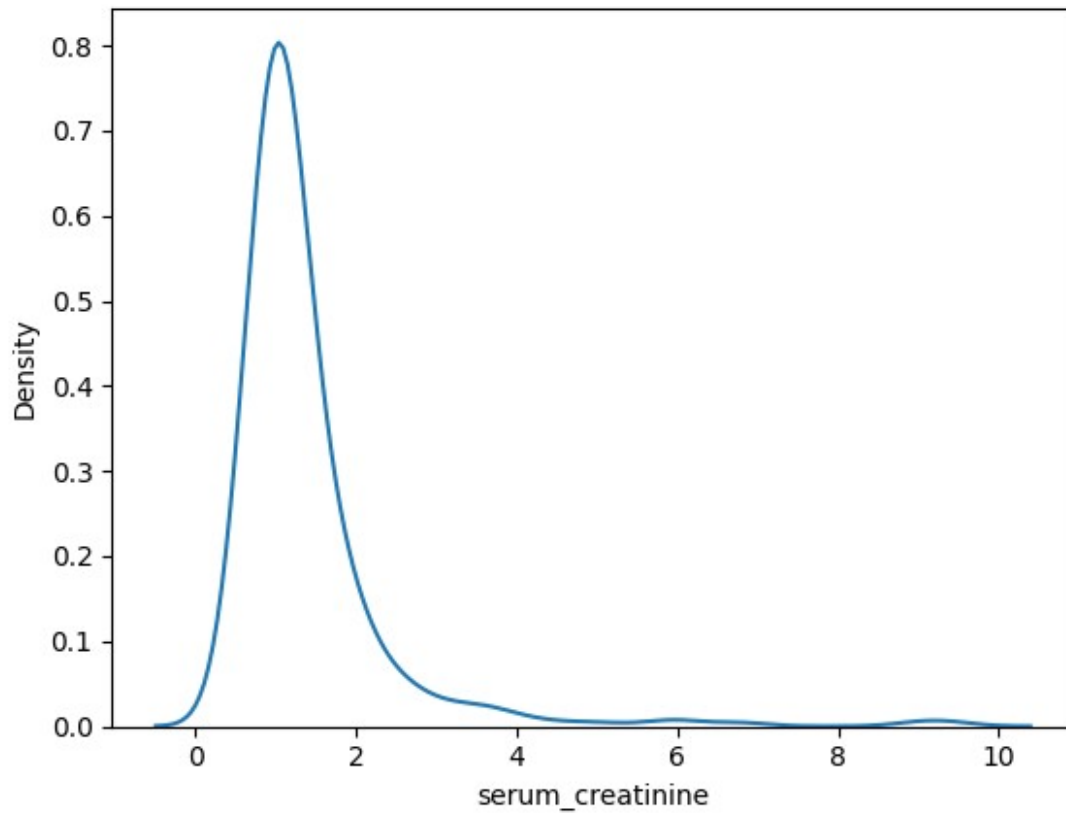


```
sns.kdeplot(df['platelets'])
```

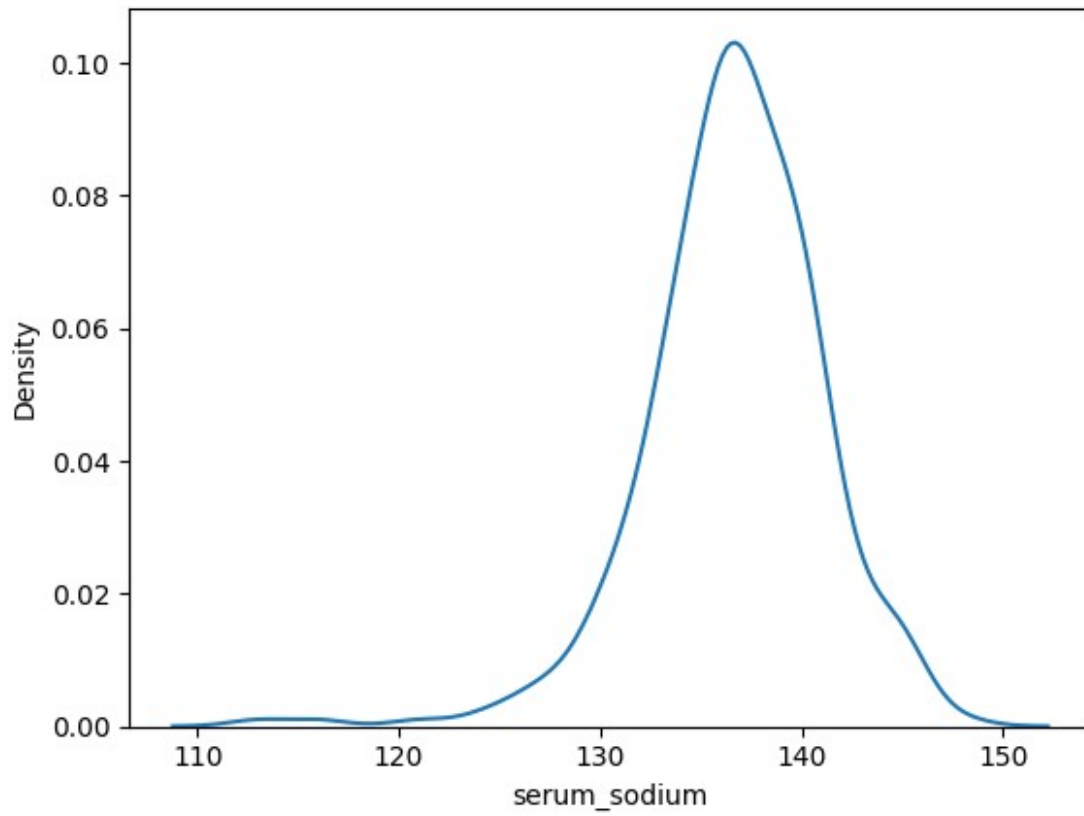
```
<Axes: xlabel='platelets', ylabel='Density'>
```



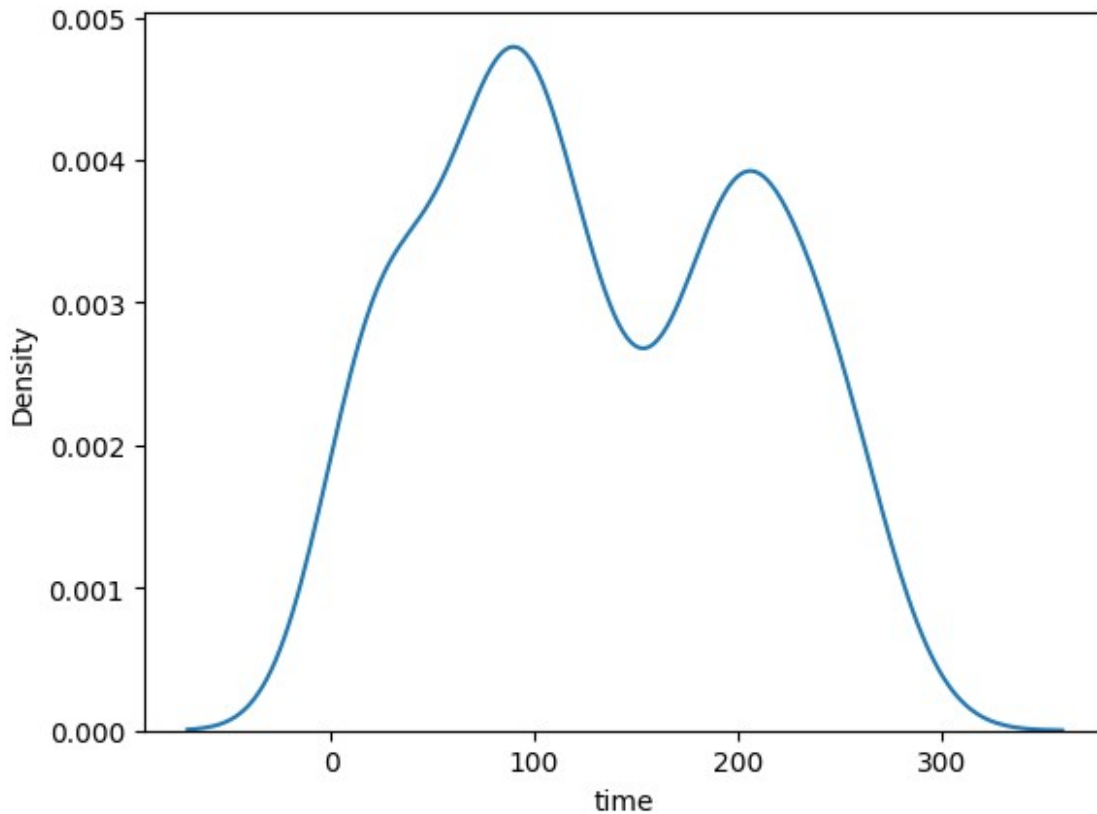
```
sns.kdeplot(df['serum_creatinine'])  
<Axes: xlabel='serum_creatinine', ylabel='Density'>
```

```
sns.kdeplot(df['serum_sodium'])  
<Axes: xlabel='serum_sodium', ylabel='Density'>
```



```
sns.kdeplot(df['time'])  
<Axes: xlabel='time', ylabel='Density'>
```



```
df['DEATH_EVENT'].value_counts()
```

```
DEATH_EVENT
```

```
0      203
```

```
1       96
```

```
Name: count, dtype: int64
```

```
"""from sklearn.preprocessing import RobustScaler,MinMaxScaler
```

```
# Initialize RobustScaler
```

```
scaler = MinMaxScaler()
```

```
# Apply scaling (convert Series to 2D before transforming)
```

```
df['serum_sodium'] = scaler.fit_transform(df[['serum_sodium']])
```

```
df['serum_creatinine'] =
```

```
scaler.fit_transform(df[['serum_creatinine']])
```

```
df['platelets'] = scaler.fit_transform(df[['platelets']])
```

```
df['creatinine_phosphokinase'] =
```

```
scaler.fit_transform(df[['creatinine_phosphokinase']])
```

```
df['time'] = scaler.fit_transform(df[['time']])
```

```
df['age'] = scaler.fit_transform(df[['age']])
```

```
df['ejection_fraction'] =
```

```
scaler.fit_transform(df[['ejection_fraction']])"""
```

```

"from sklearn.preprocessing import RobustScaler,MinMaxScaler\n\n#
Initialize RobustScaler\nscaler = MinMaxScaler()\n\n# Apply scaling
(convert Series to 2D before transforming)\ndf['serum_sodium'] =
scaler.fit_transform(df[['serum_sodium']])\ndf['serum_creatinine'] =
scaler.fit_transform(df[['serum_creatinine']])\ndf['platelets'] =
scaler.fit_transform(df[['platelets']])\
ndf['creatinine_phosphokinase'] =
scaler.fit_transform(df[['creatinine_phosphokinase']])\ndf['time'] =
scaler.fit_transform(df[['time']])\ndf['age'] =
scaler.fit_transform(df[['age']])\ndf['ejection_fraction'] =
scaler.fit_transform(df[['ejection_fraction']])"

"""import pandas as pd

def remove_outliers_all_columns(df):

    Removes outliers from all numerical columns in a DataFrame using
    the IQR method.

    Parameters:
    df (pd.DataFrame): The input DataFrame.

    Returns:
    pd.DataFrame: DataFrame with outliers removed from all numerical
    columns.

    df_filtered = df.copy() # Make a copy to avoid modifying the
    original DataFrame

    for column in
df_filtered.select_dtypes(include=['number']).columns:
        Q1 = df_filtered[column].quantile(0.25)
        Q3 = df_filtered[column].quantile(0.75)
        IQR = Q3 - Q1

        lower_bound = Q1 - 1.5 * IQR
        upper_bound = Q3 + 1.5 * IQR

        df_filtered = df_filtered[(df_filtered[column] >= lower_bound)
& (df_filtered[column] <= upper_bound)]

    return df_filtered

df = remove_outliers_all_columns(df) """

```

```

"import pandas as pd\n\ndef remove_outliers_all_columns(df):\n    \n    Removes outliers from all numerical columns in a DataFrame using the\n    IQR method.\n\n    Parameters:\n    df (pd.DataFrame): The input\n    DataFrame.\n\n    Returns:\n    pd.DataFrame: DataFrame with outliers\n    removed from all numerical columns.\n    \n    df_filtered = df.copy()\n    # Make a copy to avoid modifying the original DataFrame\n    \n    for\n    column in df_filtered.select_dtypes(include=['number']).columns:\n    Q1 = df_filtered[column].quantile(0.25)\n    Q3 =\n    df_filtered[column].quantile(0.75)\n    IQR = Q3 - Q1\n\n    lower_bound = Q1 - 1.5 * IQR\n    upper_bound = Q3 + 1.5 * IQR\n\n    df_filtered = df_filtered[(df_filtered[column] >= lower_bound) &\n    (df_filtered[column] <= upper_bound)]\n    \n    return df_filtered\n\n\ndef = remove_outliers_all_columns(df) "

```

```

import seaborn as sns
import matplotlib.pyplot as plt

```

```

# Define number of columns to plot
num_columns = df.shape[1] # Total columns in DataFrame

```

```

# Set up subplots
fig, axes = plt.subplots(nrows=num_columns, ncols=1, figsize=(8, 4 *
num_columns))

```

```

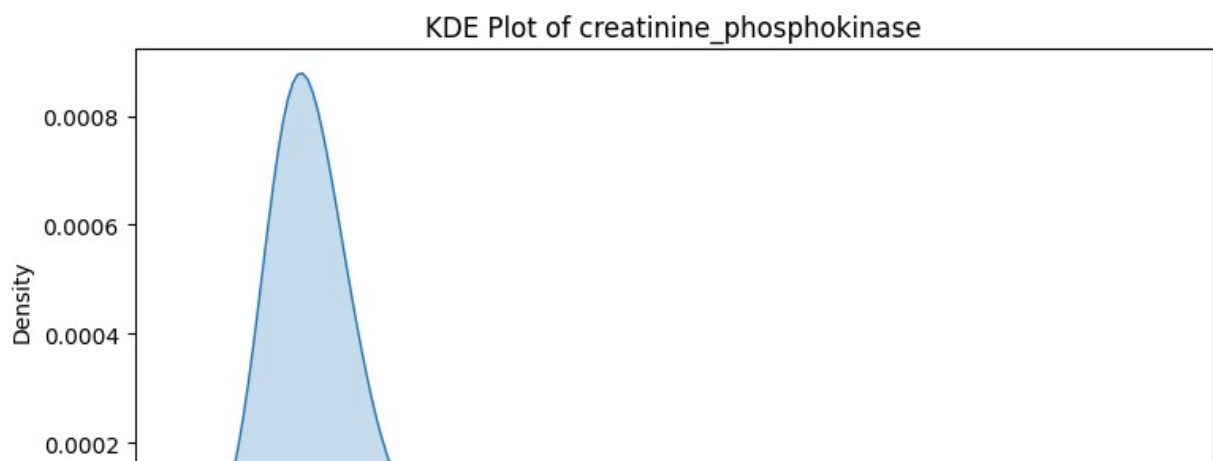
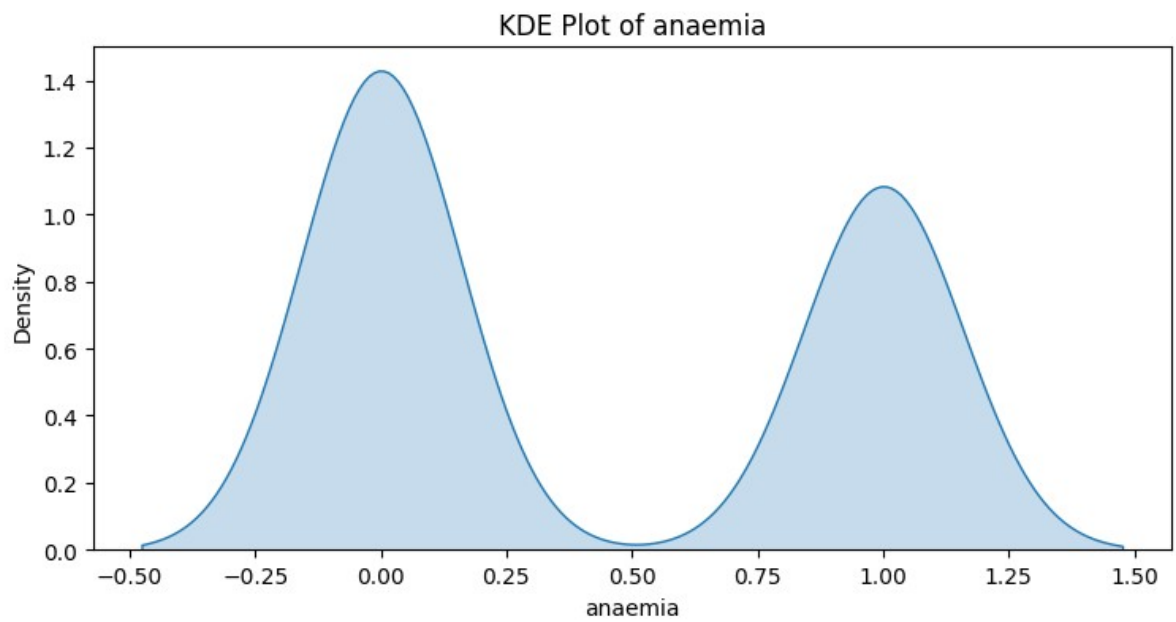
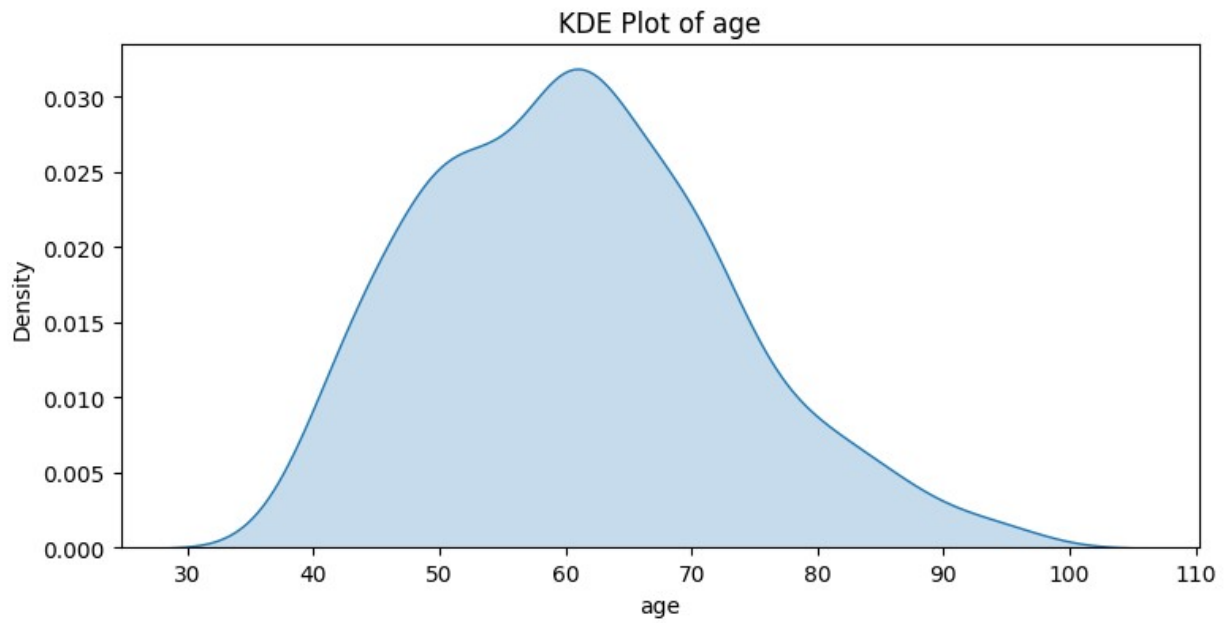
# Loop through each column and plot KDE
for i, column in enumerate(df.columns):
    sns.kdeplot(df[column], ax=axes[i], fill=True)
    axes[i].set_title(f'KDE Plot of {column}')

```

```

# Adjust layout for better spacing
plt.tight_layout()
plt.show()

```



ANN MODEL

```
import numpy as np
import pandas as pd
import tensorflow as tf
from tensorflow.keras.models import Sequential
from tensorflow.keras.layers import Dense
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import MinMaxScaler
from imblearn.over_sampling import SMOTE
from collections import Counter

# 2 Define Columns to Scale
columns_to_scale = ['age', 'creatinine_phosphokinase',
                    'ejection_fraction',
                    'platelets', 'serum_creatinine',
                    'serum_sodium', 'time'] # Example columns to scale

# 3 Separate Features and Target
X = df.drop(columns=['DEATH_EVENT']) # Drop target column
y = df['DEATH_EVENT'] # Target variable

# 4 Train-Test Split (Stratified for Balance)
X_train, X_test, y_train, y_test = train_test_split(X, y,
                                                    test_size=0.2, random_state=42, stratify=y)

# 5 Apply SMOTE for Balancing the Dataset (Before Scaling)
smote = SMOTE(sampling_strategy=0.5, random_state=42)
X_train_resampled, y_train_resampled = smote.fit_resample(X_train,
                                                          y_train)

# 6 Apply MinMaxScaler to Selected Columns
scaler = MinMaxScaler()
X_train_resampled[columns_to_scale] =
scaler.fit_transform(X_train_resampled[columns_to_scale])
X_test[columns_to_scale] = scaler.transform(X_test[columns_to_scale])

# 7 Define ANN Model
model = Sequential([
    Dense(16, activation='relu',
input_shape=(X_train_resampled.shape[1],)), # Input layer
    Dense(8, activation='relu'), # Hidden layer
    Dense(1, activation='sigmoid') # Output layer for binary
classification
])

# 8 Compile Model
model.compile(optimizer='adam', loss='binary_crossentropy',
metrics=['accuracy'])
```

9 Train Model

```
history = model.fit(X_train_resampled, y_train_resampled, epochs=50,  
batch_size=32, validation_data=(X_test, y_test))
```

Evaluate Model

```
loss, accuracy = model.evaluate(X_test, y_test)  
print(f"Test Accuracy: {accuracy:.4f}")
```

Epoch 1/50

```
8/8 [=====] - 3s 85ms/step - loss: 0.6668 -  
accuracy: 0.6667 - val_loss: 0.6389 - val_accuracy: 0.6833
```

Epoch 2/50

```
8/8 [=====] - 0s 25ms/step - loss: 0.6619 -  
accuracy: 0.6667 - val_loss: 0.6341 - val_accuracy: 0.6833
```

Epoch 3/50

```
8/8 [=====] - 0s 24ms/step - loss: 0.6573 -  
accuracy: 0.6667 - val_loss: 0.6300 - val_accuracy: 0.6833
```

Epoch 4/50

```
8/8 [=====] - 0s 24ms/step - loss: 0.6532 -  
accuracy: 0.6667 - val_loss: 0.6260 - val_accuracy: 0.6833
```

Epoch 5/50

```
8/8 [=====] - 0s 24ms/step - loss: 0.6491 -  
accuracy: 0.6667 - val_loss: 0.6230 - val_accuracy: 0.6833
```

Epoch 6/50

```
8/8 [=====] - 0s 24ms/step - loss: 0.6450 -  
accuracy: 0.6667 - val_loss: 0.6197 - val_accuracy: 0.6833
```

Epoch 7/50

```
8/8 [=====] - 0s 22ms/step - loss: 0.6413 -  
accuracy: 0.6667 - val_loss: 0.6170 - val_accuracy: 0.6833
```

Epoch 8/50

```
8/8 [=====] - 0s 24ms/step - loss: 0.6373 -  
accuracy: 0.6667 - val_loss: 0.6135 - val_accuracy: 0.6833
```

Epoch 9/50

```
8/8 [=====] - 0s 22ms/step - loss: 0.6339 -  
accuracy: 0.6667 - val_loss: 0.6108 - val_accuracy: 0.6833
```

Epoch 10/50

```
8/8 [=====] - 0s 23ms/step - loss: 0.6305 -  
accuracy: 0.6667 - val_loss: 0.6080 - val_accuracy: 0.6833
```

Epoch 11/50

```
8/8 [=====] - 0s 25ms/step - loss: 0.6271 -  
accuracy: 0.6667 - val_loss: 0.6052 - val_accuracy: 0.6833
```

Epoch 12/50

```
8/8 [=====] - 0s 25ms/step - loss: 0.6241 -  
accuracy: 0.6667 - val_loss: 0.6025 - val_accuracy: 0.6833
```

Epoch 13/50

```
8/8 [=====] - 0s 23ms/step - loss: 0.6208 -  
accuracy: 0.6667 - val_loss: 0.6001 - val_accuracy: 0.6833
```

Epoch 14/50

```
8/8 [=====] - 0s 23ms/step - loss: 0.6178 -
```

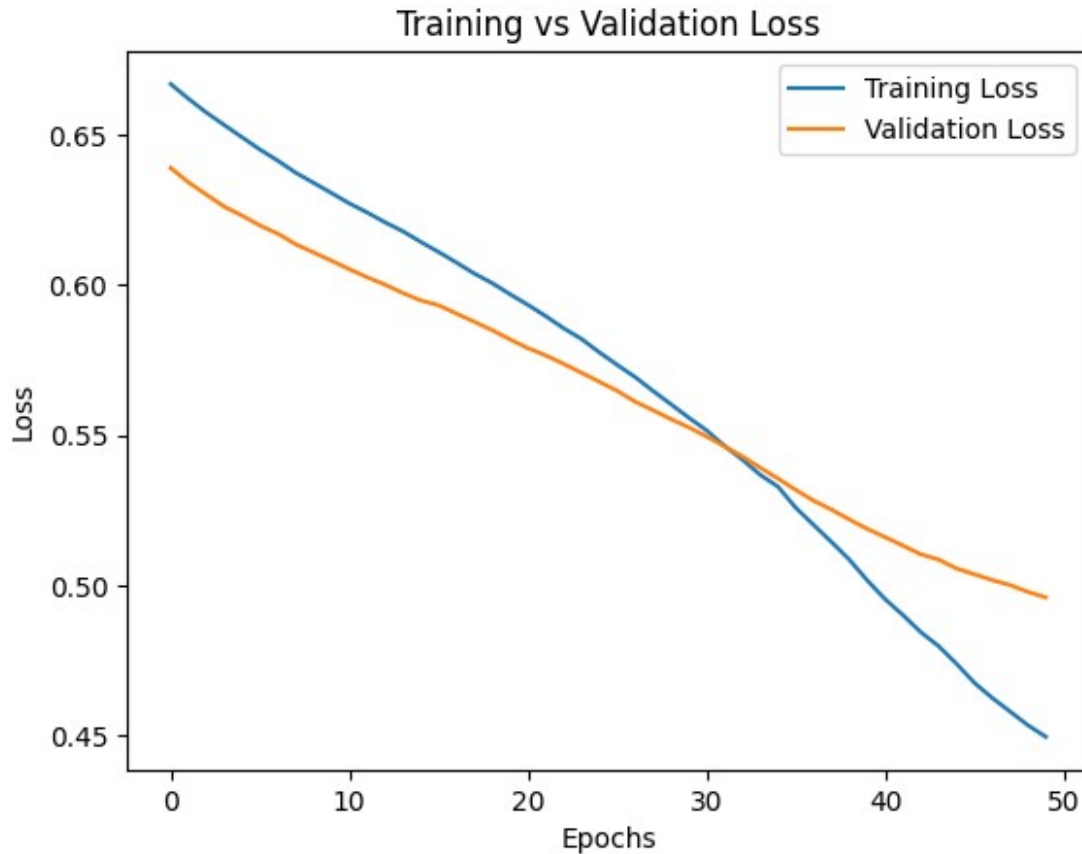

accuracy: 0.6667 - val_loss: 0.5973 - val_accuracy: 0.6833
Epoch 15/50
8/8 [=====] - 0s 25ms/step - loss: 0.6143 -
accuracy: 0.6667 - val_loss: 0.5948 - val_accuracy: 0.6833
Epoch 16/50
8/8 [=====] - 0s 24ms/step - loss: 0.6109 -
accuracy: 0.6667 - val_loss: 0.5932 - val_accuracy: 0.6833
Epoch 17/50
8/8 [=====] - 0s 23ms/step - loss: 0.6074 -
accuracy: 0.6667 - val_loss: 0.5904 - val_accuracy: 0.6833
Epoch 18/50
8/8 [=====] - 0s 25ms/step - loss: 0.6037 -
accuracy: 0.6667 - val_loss: 0.5877 - val_accuracy: 0.6833
Epoch 19/50
8/8 [=====] - 0s 25ms/step - loss: 0.6006 -
accuracy: 0.6667 - val_loss: 0.5849 - val_accuracy: 0.6833
Epoch 20/50
8/8 [=====] - 0s 23ms/step - loss: 0.5969 -
accuracy: 0.6667 - val_loss: 0.5818 - val_accuracy: 0.6833
Epoch 21/50
8/8 [=====] - 0s 23ms/step - loss: 0.5933 -
accuracy: 0.6667 - val_loss: 0.5789 - val_accuracy: 0.6833
Epoch 22/50
8/8 [=====] - 0s 24ms/step - loss: 0.5896 -
accuracy: 0.6667 - val_loss: 0.5765 - val_accuracy: 0.6833
Epoch 23/50
8/8 [=====] - 0s 27ms/step - loss: 0.5855 -
accuracy: 0.6667 - val_loss: 0.5737 - val_accuracy: 0.6833
Epoch 24/50
8/8 [=====] - 0s 25ms/step - loss: 0.5820 -
accuracy: 0.6667 - val_loss: 0.5707 - val_accuracy: 0.6833
Epoch 25/50
8/8 [=====] - 0s 25ms/step - loss: 0.5775 -
accuracy: 0.6667 - val_loss: 0.5677 - val_accuracy: 0.6833
Epoch 26/50
8/8 [=====] - 0s 26ms/step - loss: 0.5733 -
accuracy: 0.6667 - val_loss: 0.5647 - val_accuracy: 0.6833
Epoch 27/50
8/8 [=====] - 0s 25ms/step - loss: 0.5693 -
accuracy: 0.6667 - val_loss: 0.5611 - val_accuracy: 0.6833
Epoch 28/50
8/8 [=====] - 0s 25ms/step - loss: 0.5647 -
accuracy: 0.6667 - val_loss: 0.5583 - val_accuracy: 0.6833
Epoch 29/50
8/8 [=====] - 0s 23ms/step - loss: 0.5603 -
accuracy: 0.6667 - val_loss: 0.5553 - val_accuracy: 0.6833
Epoch 30/50
8/8 [=====] - 0s 25ms/step - loss: 0.5558 -
accuracy: 0.6667 - val_loss: 0.5526 - val_accuracy: 0.6833

Epoch 31/50
8/8 [=====] - 0s 25ms/step - loss: 0.5516 - accuracy: 0.6667 - val_loss: 0.5495 - val_accuracy: 0.6833
Epoch 32/50
8/8 [=====] - 0s 25ms/step - loss: 0.5464 - accuracy: 0.6667 - val_loss: 0.5463 - val_accuracy: 0.6833
Epoch 33/50
8/8 [=====] - 0s 25ms/step - loss: 0.5419 - accuracy: 0.6667 - val_loss: 0.5429 - val_accuracy: 0.6833
Epoch 34/50
8/8 [=====] - 0s 25ms/step - loss: 0.5368 - accuracy: 0.6790 - val_loss: 0.5392 - val_accuracy: 0.6833
Epoch 35/50
8/8 [=====] - 0s 26ms/step - loss: 0.5328 - accuracy: 0.6790 - val_loss: 0.5355 - val_accuracy: 0.7000
Epoch 36/50
8/8 [=====] - 0s 24ms/step - loss: 0.5258 - accuracy: 0.7037 - val_loss: 0.5317 - val_accuracy: 0.7167
Epoch 37/50
8/8 [=====] - 0s 24ms/step - loss: 0.5201 - accuracy: 0.7160 - val_loss: 0.5282 - val_accuracy: 0.7167
Epoch 38/50
8/8 [=====] - 0s 25ms/step - loss: 0.5144 - accuracy: 0.7160 - val_loss: 0.5252 - val_accuracy: 0.7167
Epoch 39/50
8/8 [=====] - 0s 24ms/step - loss: 0.5086 - accuracy: 0.7202 - val_loss: 0.5219 - val_accuracy: 0.7000
Epoch 40/50
8/8 [=====] - 0s 26ms/step - loss: 0.5017 - accuracy: 0.7407 - val_loss: 0.5188 - val_accuracy: 0.7000
Epoch 41/50
8/8 [=====] - 0s 24ms/step - loss: 0.4954 - accuracy: 0.7613 - val_loss: 0.5161 - val_accuracy: 0.7667
Epoch 42/50
8/8 [=====] - 0s 24ms/step - loss: 0.4901 - accuracy: 0.7860 - val_loss: 0.5132 - val_accuracy: 0.7667
Epoch 43/50
8/8 [=====] - 0s 24ms/step - loss: 0.4844 - accuracy: 0.7901 - val_loss: 0.5103 - val_accuracy: 0.7667
Epoch 44/50
8/8 [=====] - 0s 25ms/step - loss: 0.4798 - accuracy: 0.7819 - val_loss: 0.5086 - val_accuracy: 0.7667
Epoch 45/50
8/8 [=====] - 0s 26ms/step - loss: 0.4739 - accuracy: 0.7819 - val_loss: 0.5056 - val_accuracy: 0.7667
Epoch 46/50
8/8 [=====] - 0s 24ms/step - loss: 0.4676 - accuracy: 0.7860 - val_loss: 0.5037 - val_accuracy: 0.7667
Epoch 47/50

```
8/8 [=====] - 0s 24ms/step - loss: 0.4625 -  
accuracy: 0.7984 - val_loss: 0.5018 - val_accuracy: 0.7833  
Epoch 48/50  
8/8 [=====] - 0s 26ms/step - loss: 0.4580 -  
accuracy: 0.8107 - val_loss: 0.5001 - val_accuracy: 0.7833  
Epoch 49/50  
8/8 [=====] - 0s 24ms/step - loss: 0.4535 -  
accuracy: 0.8148 - val_loss: 0.4978 - val_accuracy: 0.7833  
Epoch 50/50  
8/8 [=====] - 0s 24ms/step - loss: 0.4496 -  
accuracy: 0.8148 - val_loss: 0.4960 - val_accuracy: 0.7833  
2/2 [=====] - 0s 6ms/step - loss: 0.4960 -  
accuracy: 0.7833  
Test Accuracy: 0.7833
```

```
import matplotlib.pyplot as plt
```

```
# Assuming history is from model.fit()  
plt.plot(history.history['loss'], label='Training Loss')  
plt.plot(history.history['val_loss'], label='Validation Loss')  
plt.xlabel('Epochs')  
plt.ylabel('Loss')  
plt.legend()  
plt.title("Training vs Validation Loss")  
plt.show()
```



```

train_loss, train_acc = model.evaluate(X_train_resampled,
y_train_resampled)
test_loss, test_acc = model.evaluate(X_test, y_test)

print(f"Training Accuracy: {train_acc:.4f}")
print(f"Testing Accuracy: {test_acc:.4f}")

8/8 [=====] - 0s 11ms/step - loss: 0.4462 -
accuracy: 0.8148
2/2 [=====] - 0s 13ms/step - loss: 0.4960 -
accuracy: 0.7833
Training Accuracy: 0.8148
Testing Accuracy: 0.7833

# Dummy Input (Without Target Column)
dummy_input = np.array([[75, 0, 582, 0, 20, 1, 265000.00, 1.9, 130, 1,
0, 4]]) # 12 Features

# Convert to DataFrame with Proper Column Names
dummy_input_df = pd.DataFrame(dummy_input, columns=X.columns) #
X.columns ensures correct order

# Scale Only Selected Columns

```

```

dummy_input_df[columns_to_scale] =
scaler.transform(dummy_input_df[columns_to_scale])

# Convert to NumPy for Prediction
dummy_input_scaled = dummy_input_df.to_numpy()

# Make a Prediction
prediction = model.predict(dummy_input_scaled)

# Convert Probability to Class (0 or 1)
predicted_class = (prediction > 0.5).astype(int)

print(f"Predicted Probability: {prediction[0][0]:.4f}")
print(f"Predicted Death Event: {predicted_class[0][0]}")

```

WARNING:tensorflow:5 out of the last 7 calls to <function Model.make_predict_function.<locals>.predict_function at 0x0000021382204040> triggered tf.function retracing. Tracing is expensive and the excessive number of tracings could be due to (1) creating @tf.function repeatedly in a loop, (2) passing tensors with different shapes, (3) passing Python objects instead of tensors. For (1), please define your @tf.function outside of the loop. For (2), @tf.function has reduce_retracing=True option that can avoid unnecessary retracing. For (3), please refer to https://www.tensorflow.org/guide/function#controlling_retracing and https://www.tensorflow.org/api_docs/python/tf/function for more details.

1/1 [=====] - 0s 179ms/step

Predicted Probability: 0.6417

Predicted Death Event: 1

```

import numpy as np
import pandas as pd

```

```

# Define column names

```

```

columns = ["age", "anaemia", "creatinine_phosphokinase", "diabetes",
           "ejection_fraction",
           "high_blood_pressure", "platelets", "serum_creatinine",
           "serum_sodium",
           "sex", "smoking", "time"]

```

```

# Case 1: High Risk of Death (Expected DEATH_EVENT = 1)

```

```

high_risk_patient = np.array([[85, 1, 1000, 1, 15, 1, 150000, 2.5,
125, 1, 1, 3]])

```

```

# Case 2: Low Risk of Death (Expected DEATH_EVENT = 0)

```

```

low_risk_patient = np.array([[40, 0, 250, 0, 50, 0, 300000, 1.0, 140,
0, 0, 30]])

```

```

# Convert to DataFrame for scaling
high_risk_df = pd.DataFrame(high_risk_patient, columns=columns)
low_risk_df = pd.DataFrame(low_risk_patient, columns=columns)

# Scale specific columns (same scaler used for training)
columns_to_scale = ["age", "creatinine_phosphokinase",
                    "ejection_fraction",
                    "platelets", "serum_creatinine", "serum_sodium",
                    "time"]

high_risk_df[columns_to_scale] =
scaler.transform(high_risk_df[columns_to_scale])
low_risk_df[columns_to_scale] =
scaler.transform(low_risk_df[columns_to_scale])

# Make predictions
high_risk_pred = model.predict(high_risk_df)
low_risk_pred = model.predict(low_risk_df)

print(f"High Risk Patient - Predicted Probability: {high_risk_pred[0]
[0]:.4f}, Predicted Death Event: {int(high_risk_pred[0][0] > 0.5)}")
print(f"Low Risk Patient - Predicted Probability: {low_risk_pred[0]
[0]:.4f}, Predicted Death Event: {int(low_risk_pred[0][0] > 0.5)}")

WARNING:tensorflow:6 out of the last 8 calls to <function
Model.make_predict_function.<locals>.predict_function at
0x0000021382204040> triggered tf.function retracing. Tracing is
expensive and the excessive number of tracings could be due to (1)
creating @tf.function repeatedly in a loop, (2) passing tensors with
different shapes, (3) passing Python objects instead of tensors. For
(1), please define your @tf.function outside of the loop. For (2),
@tf.function has reduce_retracing=True option that can avoid
unnecessary retracing. For (3), please refer to
https://www.tensorflow.org/guide/function#controlling\_retracing and
https://www.tensorflow.org/api\_docs/python/tf/function for more
details.
1/1 [=====] - 0s 176ms/step
1/1 [=====] - 0s 58ms/step
High Risk Patient - Predicted Probability: 0.7465, Predicted Death
Event: 1
Low Risk Patient - Predicted Probability: 0.2658, Predicted Death
Event: 0

import numpy as np
import pandas as pd
# Define column names
columns = ["age", "anaemia", "creatinine_phosphokinase", "diabetes",
          "ejection_fraction",
          "high_blood_pressure", "platelets", "serum_creatinine",
          "serum_sodium",

```

```

        "sex", "smoking", "time"]
# Case 1: High Risk of Death (Expected DEATH_EVENT = 1)
high_risk_patient = np.array([[85, 1, 1000, 1, 15, 1, 150000, 2.5,
125, 1, 1, 3]])
# Case 2: Low Risk of Death (Expected DEATH_EVENT = 0)
low_risk_patient = np.array([[40, 0, 250, 0, 50, 0, 300000, 1.0, 140,
0, 0, 30]])
# Case 3: Moderate-High Risk of Death (Expected DEATH_EVENT = 1)
moderate_high_risk = np.array([[72, 1, 850, 1, 25, 1, 200000, 1.9,
130, 1, 0, 6]])
# Case 4: Moderate-Low Risk of Death (Expected DEATH_EVENT = 0)
moderate_low_risk = np.array([[55, 0, 350, 0, 45, 0, 280000, 1.2, 138,
0, 1, 25]])
# Convert to DataFrame for scaling
high_risk_df = pd.DataFrame(high_risk_patient, columns=columns)
low_risk_df = pd.DataFrame(low_risk_patient, columns=columns)
moderate_high_df = pd.DataFrame(moderate_high_risk, columns=columns)
moderate_low_df = pd.DataFrame(moderate_low_risk, columns=columns)
# Scale specific columns (same scaler used for training)
columns_to_scale = ["age", "creatinine_phosphokinase",
"ejection_fraction",
                    "platelets", "serum_creatinine", "serum_sodium",
"time"]
high_risk_df[columns_to_scale] =
scaler.transform(high_risk_df[columns_to_scale])
low_risk_df[columns_to_scale] =
scaler.transform(low_risk_df[columns_to_scale])
moderate_high_df[columns_to_scale] =
scaler.transform(moderate_high_df[columns_to_scale])
moderate_low_df[columns_to_scale] =
scaler.transform(moderate_low_df[columns_to_scale])
# Make predictions
high_risk_pred = model.predict(high_risk_df)
low_risk_pred = model.predict(low_risk_df)
moderate_high_pred = model.predict(moderate_high_df)
moderate_low_pred = model.predict(moderate_low_df)
print(f"High Risk Patient - Predicted Probability: {high_risk_pred[0]
[0]:.4f}, Predicted Death Event: {int(high_risk_pred[0][0] > 0.5)}")
print(f"Low Risk Patient - Predicted Probability: {low_risk_pred[0]
[0]:.4f}, Predicted Death Event: {int(low_risk_pred[0][0] > 0.5)}")
print(f"Moderate-High Risk Patient - Predicted Probability:
{moderate_high_pred[0][0]:.4f}, Predicted Death Event:
{int(moderate_high_pred[0][0] > 0.5)}")
print(f"Moderate-Low Risk Patient - Predicted Probability:
{moderate_low_pred[0][0]:.4f}, Predicted Death Event:
{int(moderate_low_pred[0][0] > 0.5)}")

1/1 [=====] - 0s 60ms/step
1/1 [=====] - 0s 62ms/step
1/1 [=====] - 0s 90ms/step

```

```
1/1 [=====] - 0s 50ms/step
High Risk Patient - Predicted Probability: 0.7465, Predicted Death
Event: 1
Low Risk Patient - Predicted Probability: 0.2658, Predicted Death
Event: 0
Moderate-High Risk Patient - Predicted Probability: 0.7400, Predicted
Death Event: 1
Moderate-Low Risk Patient - Predicted Probability: 0.2834, Predicted
Death Event: 0
```