02 eda

July 20, 2025

1 Exploratory Data Analysis: Heart Failure Clinical Dataset

```
[1]: import pandas as pd
     import matplotlib.pyplot as plt
     import seaborn as sns
     sns.set(style="whitegrid")
     # Load dataset
     df = pd.read_csv('E:/hospital_analytics_project/data/raw/
      ⇔heart_failure_clinical_records_dataset.csv')
     df.head()
[1]:
                        creatinine_phosphokinase
                                                   diabetes
                                                             ejection_fraction
         age
              anaemia
     0 75.0
                                              582
                                                                             20
     1 55.0
                    0
                                             7861
                                                          0
                                                                             38
     2 65.0
                    0
                                              146
                                                          0
                                                                             20
     3 50.0
                     1
                                                          0
                                                                             20
                                              111
     4 65.0
                     1
                                              160
                                                          1
                                                                             20
        high_blood_pressure
                              platelets
                                         serum_creatinine
                                                            serum_sodium
                                                                           sex
     0
                              265000.00
                                                       1.9
                                                                      130
     1
                              263358.03
                                                       1.1
                                                                      136
                                                                             1
     2
                              162000.00
                                                       1.3
                                                                      129
                                                                             1
                              210000.00
     3
                                                       1.9
                                                                      137
                                                                             1
     4
                              327000.00
                                                       2.7
                                                                      116
                                                                             0
        smoking time
                        DEATH_EVENT
     0
                                  1
              0
     1
                    6
                                  1
     2
              1
                    7
                                  1
     3
              0
                    7
                                  1
              0
                    8
[2]: print("Shape:", df.shape)
     print("\nMissing Values:")
     print(df.isnull().sum())
```

```
print("\nData Types:")
print(df.dtypes)
print("\nTarget Distribution:")
print(df['DEATH_EVENT'].value_counts(normalize=True))
Shape: (299, 13)
Missing Values:
age
                             0
                             0
anaemia
{\tt creatinine\_phosphokinase}
                             0
diabetes
                             0
ejection_fraction
high_blood_pressure
                             0
platelets
                             0
serum_creatinine
                             0
serum_sodium
                             0
sex
                             0
smoking
                             0
                             0
time
                             0
DEATH_EVENT
dtype: int64
Data Types:
                             float64
age
anaemia
                               int64
creatinine_phosphokinase
                               int64
diabetes
                               int64
ejection_fraction
                               int64
high_blood_pressure
                               int64
                             float64
platelets
serum_creatinine
                             float64
serum_sodium
                               int64
                               int64
sex
smoking
                               int64
                               int64
time
DEATH_EVENT
                               int64
dtype: object
Target Distribution:
DEATH_EVENT
```

0

1

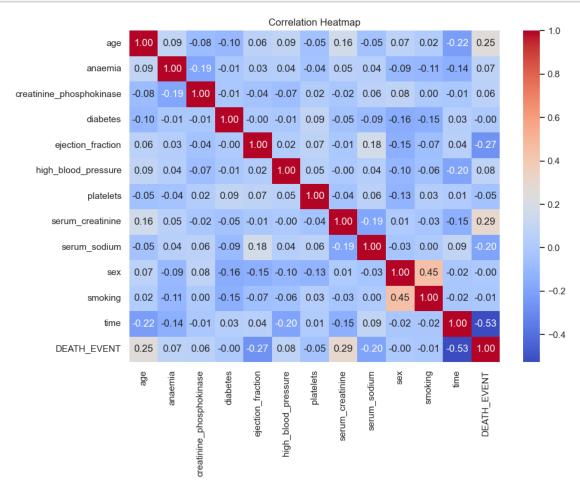
0.67893

0.32107

Name: proportion, dtype: float64

2

```
[3]: plt.figure(figsize=(10, 8))
    sns.heatmap(df.corr(numeric_only=True), annot=True, cmap='coolwarm', fmt=".2f")
    plt.title("Correlation Heatmap")
    plt.tight_layout()
    plt.show()
```



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

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

sns.boxplot(x="DEATH_EVENT", y="age", data=df, ax=axs[0], palette='Set2')
C:\Users\MSI\AppData\Local\Temp\ipykernel_7828\1517280060.py:6: FutureWarning:

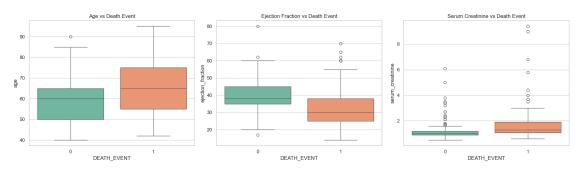
Passing `palette` without assigning `hue` is deprecated and will be removed in v0.14.0. Assign the `x` variable to `hue` and set `legend=False` for the same effect.

sns.boxplot(x="DEATH_EVENT", y="ejection_fraction", data=df, ax=axs[1],
palette='Set2')

C:\Users\MSI\AppData\Local\Temp\ipykernel_7828\1517280060.py:9: FutureWarning:

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

sns.boxplot(x="DEATH_EVENT", y="serum_creatinine", data=df, ax=axs[2],
palette='Set2')



1.1 Feature Engineering

```
[5]: df['is_elderly'] = (df['age'] > 65).astype(int)
    df['low_ejection'] = (df['ejection_fraction'] < 30).astype(int)
    df['high_creatinine'] = (df['serum_creatinine'] > 1.5).astype(int)

# Simple risk score based on 3 conditions
```

```
df['combined_risk_score'] = df['is_elderly'] + df['low_ejection'] +__

df['high_creatinine']
    df[['age', 'ejection_fraction', 'serum_creatinine', 'is_elderly', "
     [5]:
      age ejection_fraction serum_creatinine is_elderly low_ejection \
    0 75.0
                        20
                                      1.9
                                                  1
                                                              1
    1 55.0
                        38
                                      1.1
                                                  0
                                                              0
    2 65.0
                        20
                                      1.3
                                                  0
                                                              1
    3 50.0
                        20
                                      1.9
                                                  0
                                                              1
    4 65.0
                        20
                                      2.7
                                                  0
                                                              1
      high_creatinine combined_risk_score
    0
                  1
                                    3
    1
                  0
                                    0
    2
                  0
                                    1
                                    2
    3
                  1
```