

Checking for outliers using IQR

```
[14]: # outlier detection using IQR
for column in df_heart[choosen_features_nums]:
    for target in df_heart['target'].unique():
        q25 = df_heart[column][df_heart['target'] == target].quantile(0.25)
        q75 = df_heart[column][df_heart['target'] == target].quantile(0.75)
        iqr = q75 - q25
        print(target, '-', column.upper())
        print('Percentiles: 25th = %.3f, 75th = %.3f, IQR = %.3f' % (q25, q75, iqr))

        # Calculate the outlier cutoff
        cut_off = iqr * 1.5
        lower, upper = q25 - cut_off, q75 + cut_off

        # Identify outliers
        df_heart2 = pd.DataFrame(df_heart[df_heart['target'] == target][column])

        count = len(df_heart2[df_heart2[column] < lower].index)
        count += len(df_heart2[df_heart2[column] > upper].index)
        print('Identified outliers: ', count)

        # replacing outliers with NaN (Will be later replaced with feature mean)
        for index in df_heart2[df_heart2[column] < lower].index:
            df_heart.loc[index, column] = np.nan
        for index in df_heart2[df_heart2[column] > upper].index:
            df_heart.loc[index, column] = np.nan
```

1 - OLDPEAK

Percentiles: 25th = -0.023, 75th = 0.906, IQR = 0.929

Identified outliers: 5

0 - OLDPEAK

Percentiles: 25th = 0.623, 75th = 2.555, IQR = 1.932

Identified outliers: 2

1 - THALACH

Percentiles: 25th = 148.052, 75th = 172.048, IQR = 23.996

Identified outliers: 1

0 - THALACH

Percentiles: 25th = 124.972, 75th = 156.158, IQR = 31.186

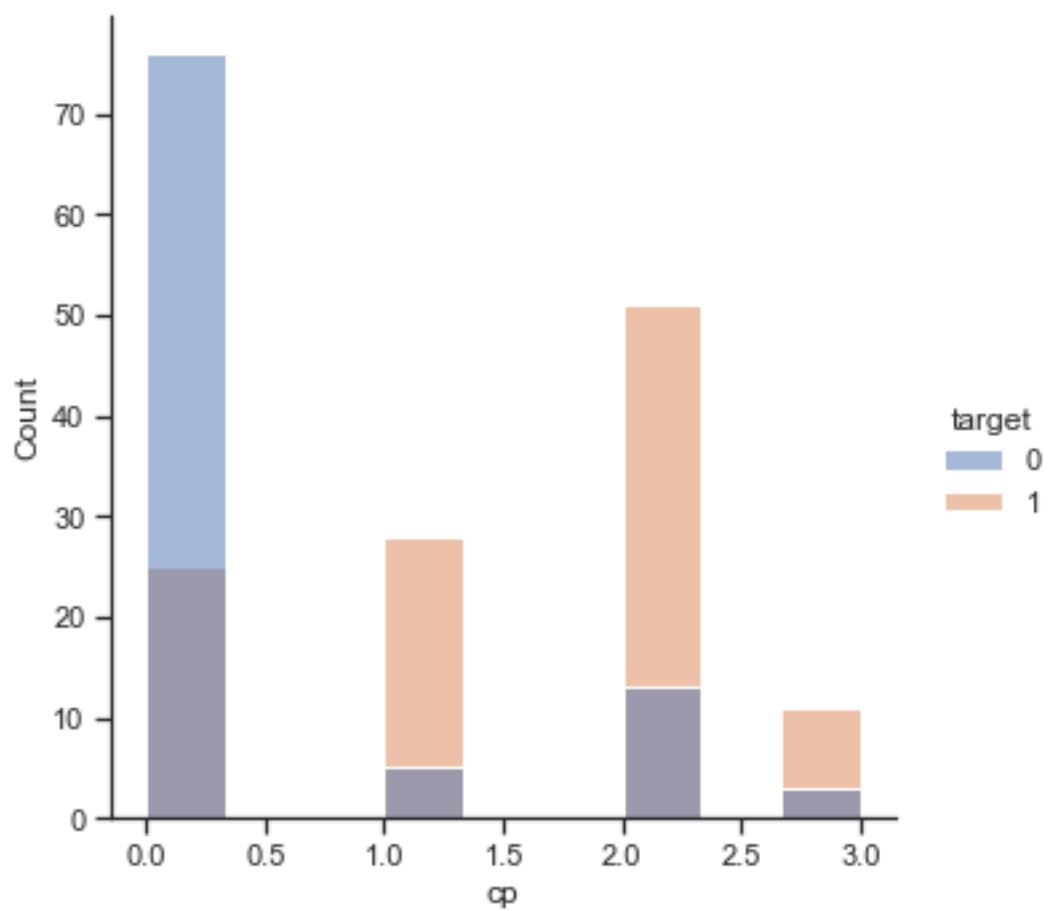
Identified outliers: 0

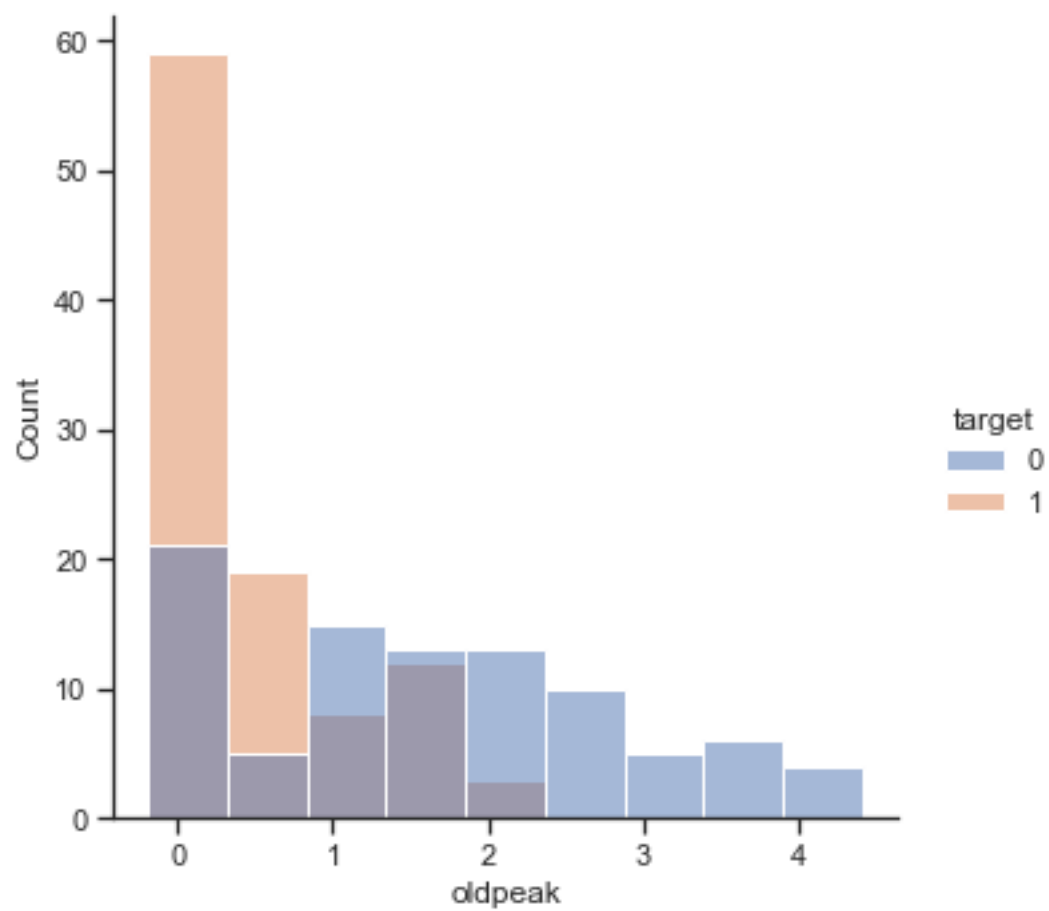
We observe that the number of outliers found corresponds to the box-plot. These outliers can be handled by replacing with feature mean.

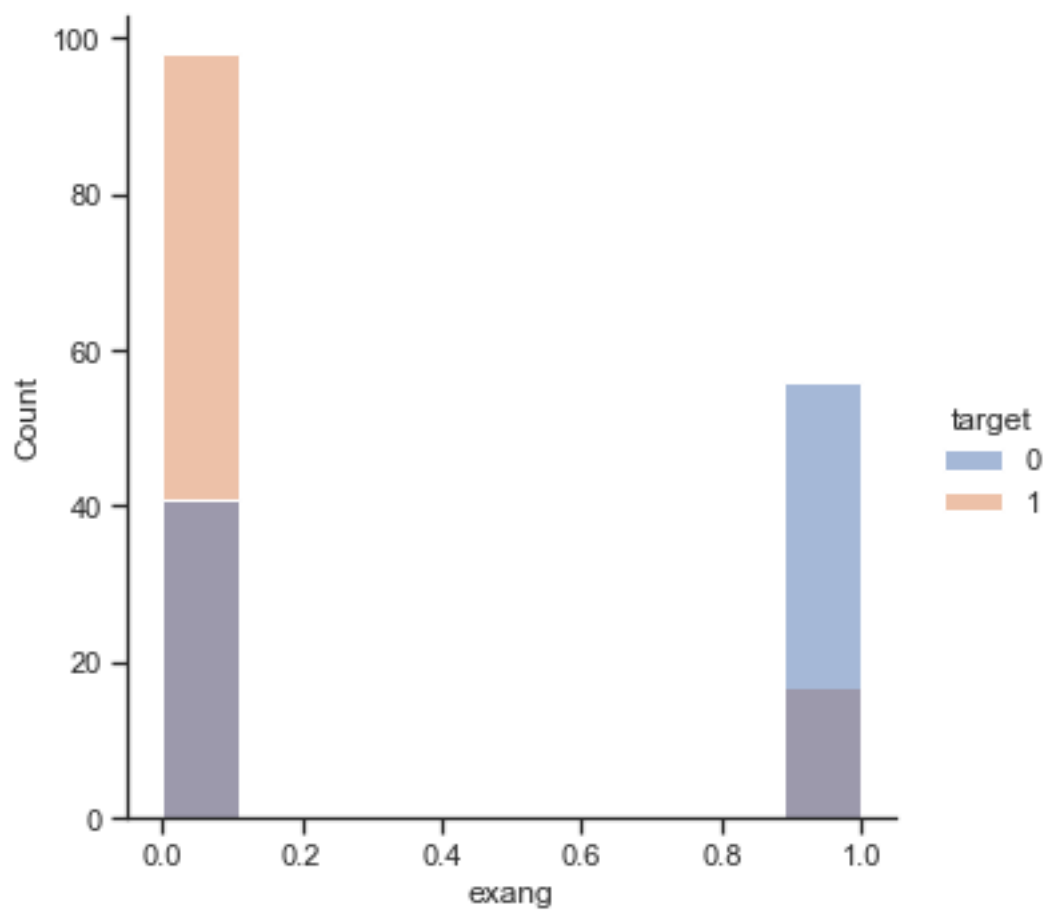
[CM4]

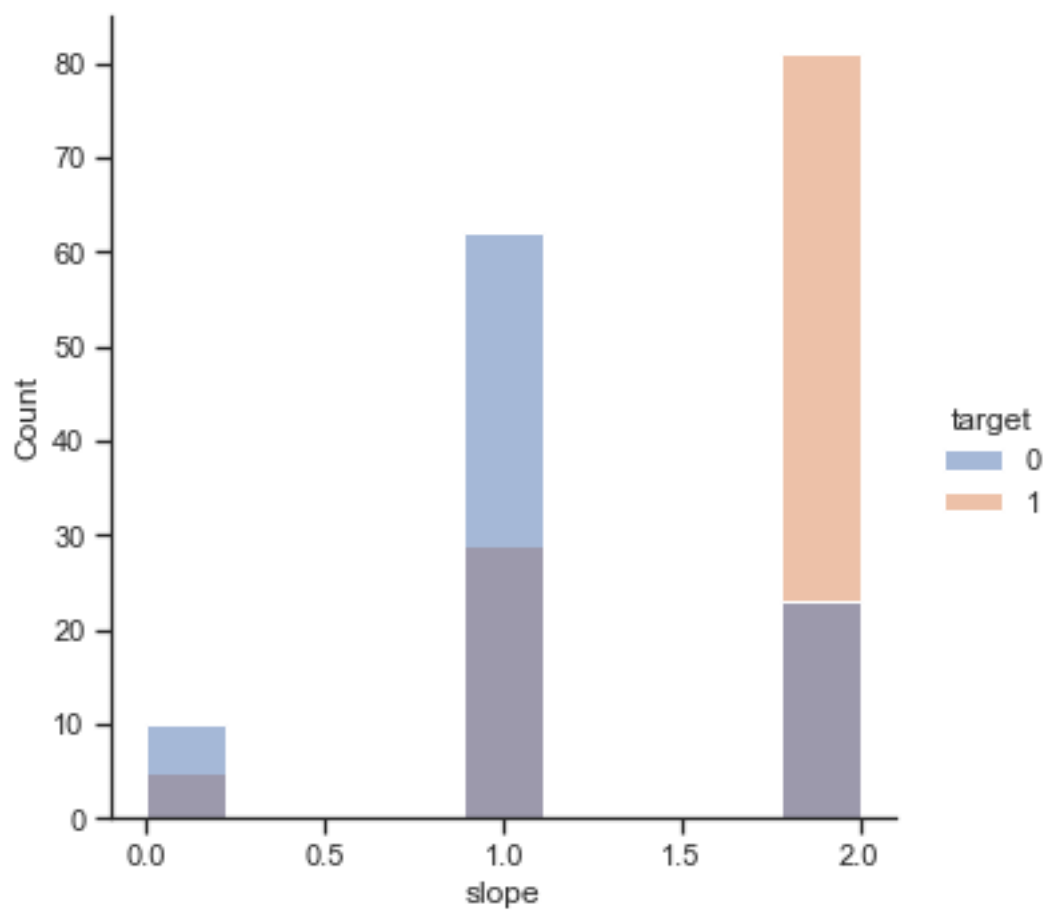
Histogram plot of the features

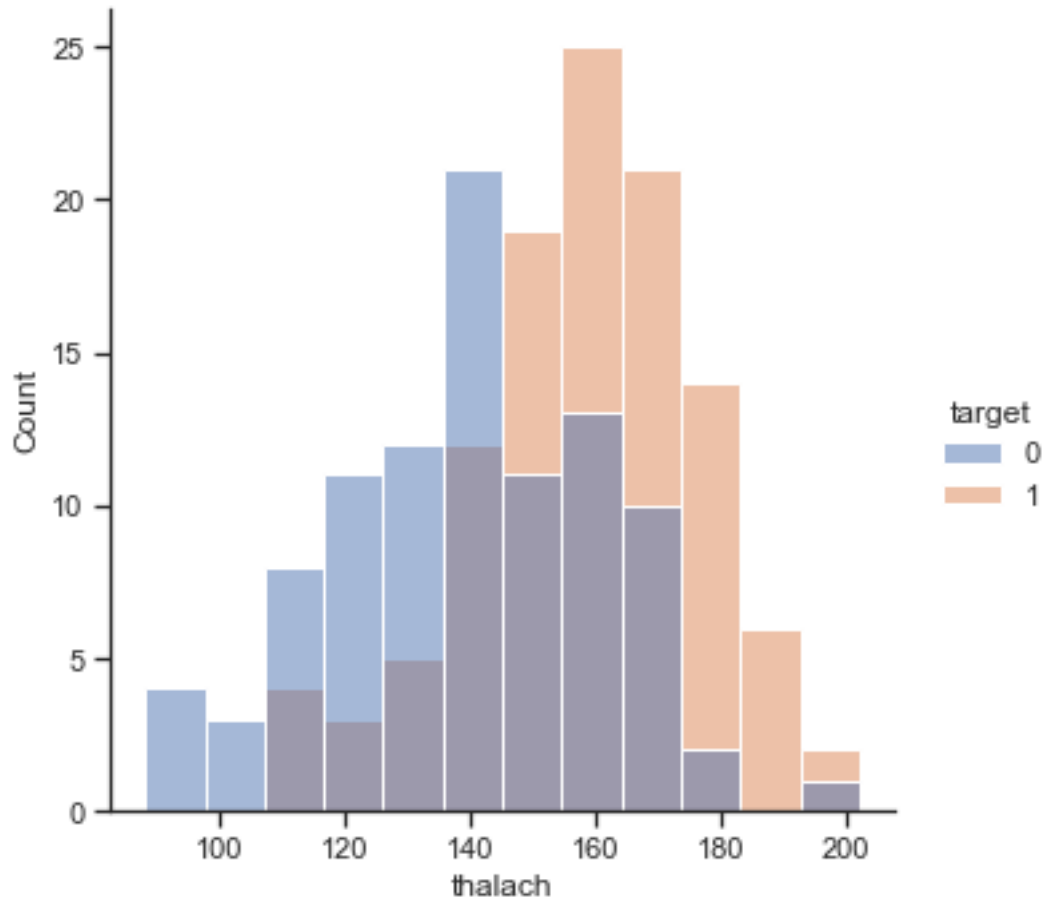
```
[15]: # plot histogram
for column in df_heart[choosen_features]:
    sns.displot(df_heart, x=column, hue="target")
```











From the above histograms, we can see the number of present (1) and absent (0) heart disease cases in each features.

[CM5]

Data Cleaning

Checking for null / NaN values (missing data)

```
[16]: # checking for any null / NaN values
df_heart.isnull().values.any()
```

[16]: True

```
[17]: # checking for any null / NaN values
df_heart.isna().sum()
```

```
[17]: age      0
      sex      0
      cp      0
      trestbps  7
      chol    10
      fbs      0
```