

# HeartDisease prediction

## 0.1 1 : Indtroduction

```
[67]: import numpy as np
import pandas as pd
import matplotlib as plt
import seaborn as sns
import matplotlib.pyplot as plt
```

## 0.2 2 : Data Wrangling

```
[68]: data = pd.read_csv("SL_data.csv")
data.head()
```

```
[68]:
```

	age	sex	cp	trestbps	chol	fbs	restecg	thalach	exang	oldpeak	slope	\
0	63	1	3	145	233	1	0	150	0	2.3	0	
1	37	1	2	130	250	0	1	187	0	3.5	0	
2	41	0	1	130	204	0	0	172	0	1.4	2	
3	56	1	1	120	236	0	1	178	0	0.8	2	
4	57	0	0	120	354	0	1	163	1	0.6	2	

	ca	thal	target
0	0	1	1
1	0	2	1
2	0	2	1
3	0	2	1
4	0	2	1

```
[69]: print("(Rows, columns): " + str(data.shape))
data.columns
```

(Rows, columns): (303, 14)

```
[69]: Index(['age', 'sex', 'cp', 'trestbps', 'chol', 'fbs', 'restecg', 'thalach',
        'exang', 'oldpeak', 'slope', 'ca', 'thal', 'target'],
        dtype='object')
```

```
[70]: data.nunique(axis=0) # returns the number of unique values for each variable.
```

```
[70]: age          41
      sex          2
      cp          4
      trestbps    49
      chol       152
      fbs         2
      restecg     3
      thalach     91
      exang       2
      oldpeak     40
      slope       3
      ca          5
      thal        4
      target      2
      dtype: int64
```

1 *#summarizes the count, mean, standard deviation, min, and max for numeric variables.*

```
data.describe()
```

```
[71]: # Display the Missing Values

      print(data.isna().sum())
```

```
age          0
sex          0
cp           0
trestbps     0
chol         0
fbs          0
restecg      0
thalach      0
exang        0
oldpeak      0
slope        0
ca           0
thal         0
target       0
dtype: int64
```

```
[72]: data['target'].value_counts()
```

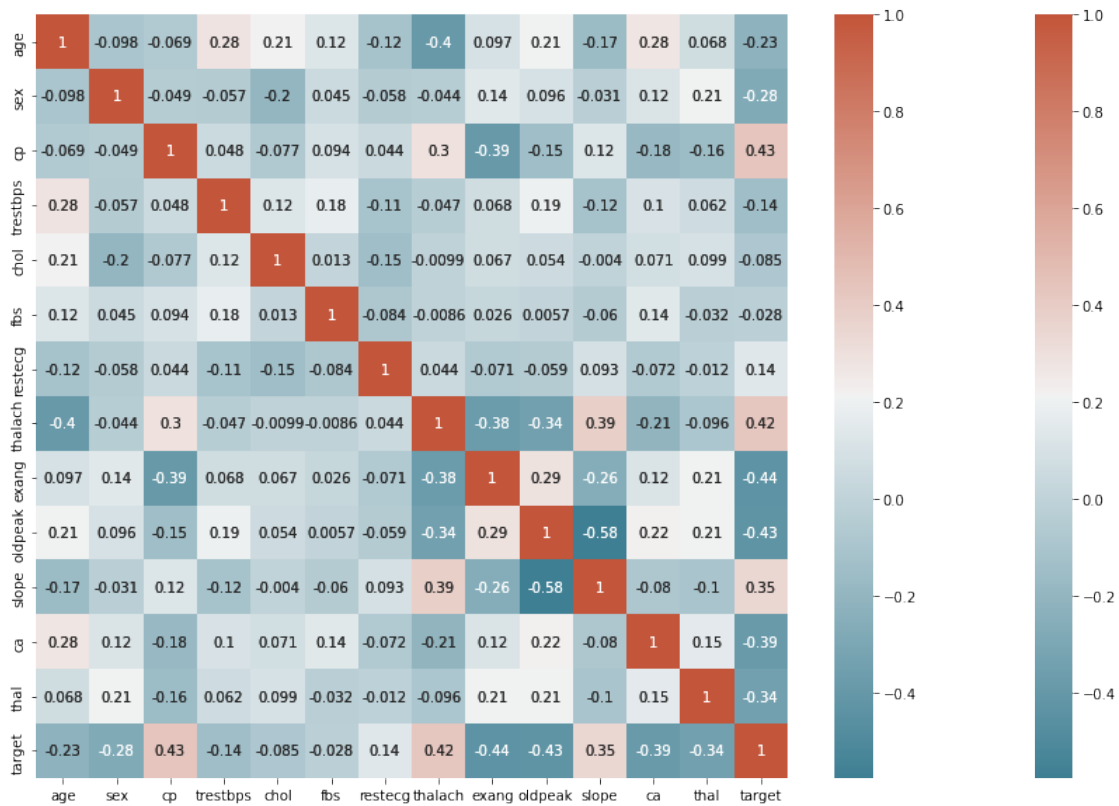
```
[72]: 1    165
      0    138
      Name: target, dtype: int64
```

## 1.1 3. Exploratory Data Analysis

### 3.1 : Correlation Matrix- correlations between all variables.

```
[73]: corr = data.corr()
plt.subplots(figsize=(15,10))
sns.heatmap(corr, xticklabels=corr.columns, yticklabels=corr.columns,
            annot=True, cmap=sns.diverging_palette(220, 20, as_cmap=True))
sns.heatmap(corr, xticklabels=corr.columns,
            yticklabels=corr.columns,
            annot=True,
            cmap=sns.diverging_palette(220, 20, as_cmap=True))
```

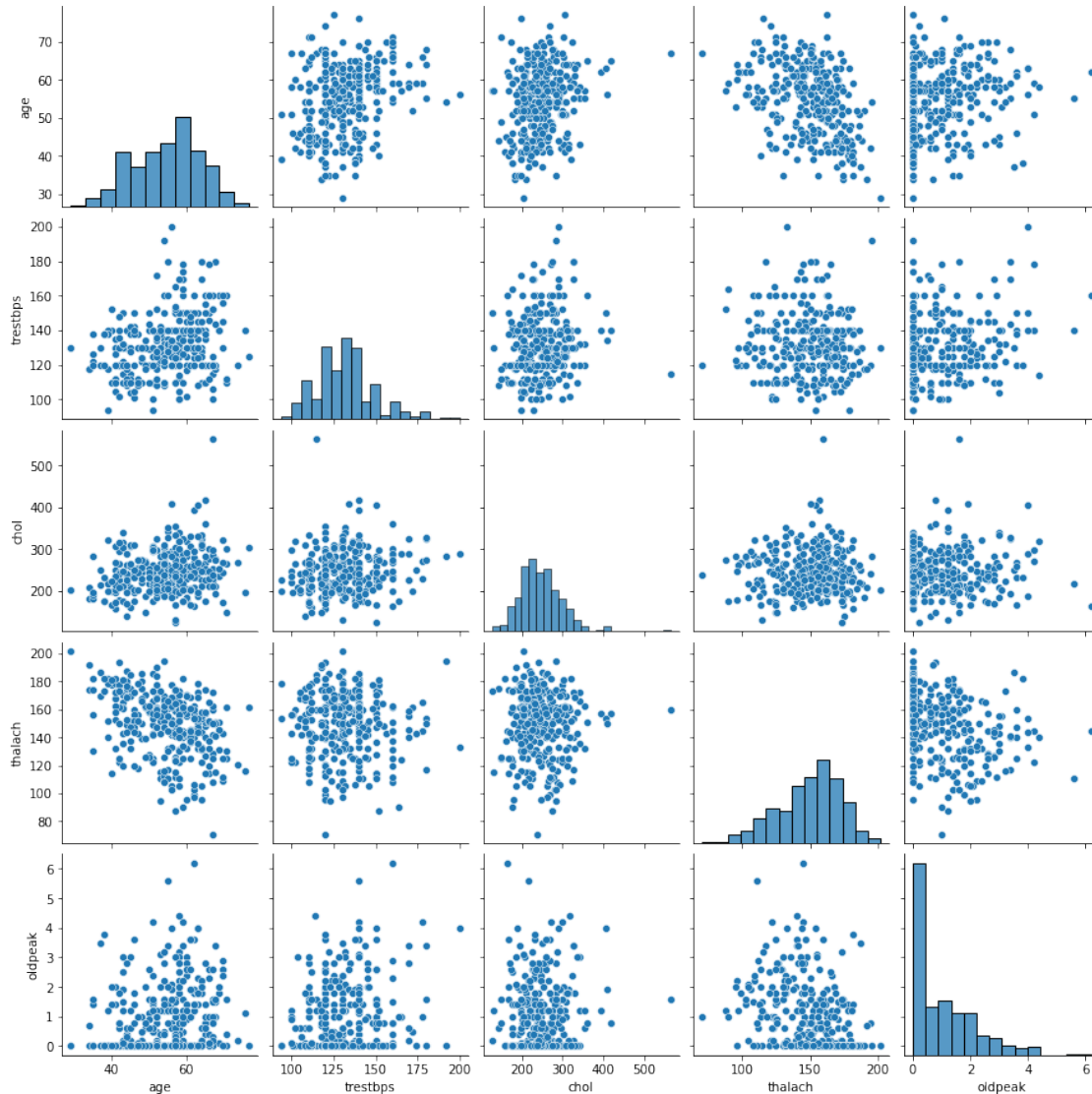
```
[73]: <AxesSubplot:>
```



### 1.1.1 3.2 : A pairplot with only our continuous features.

```
[74]: subData = data[['age', 'trestbps', 'chol', 'thalach', 'oldpeak']]
      sns.pairplot(subData)
```

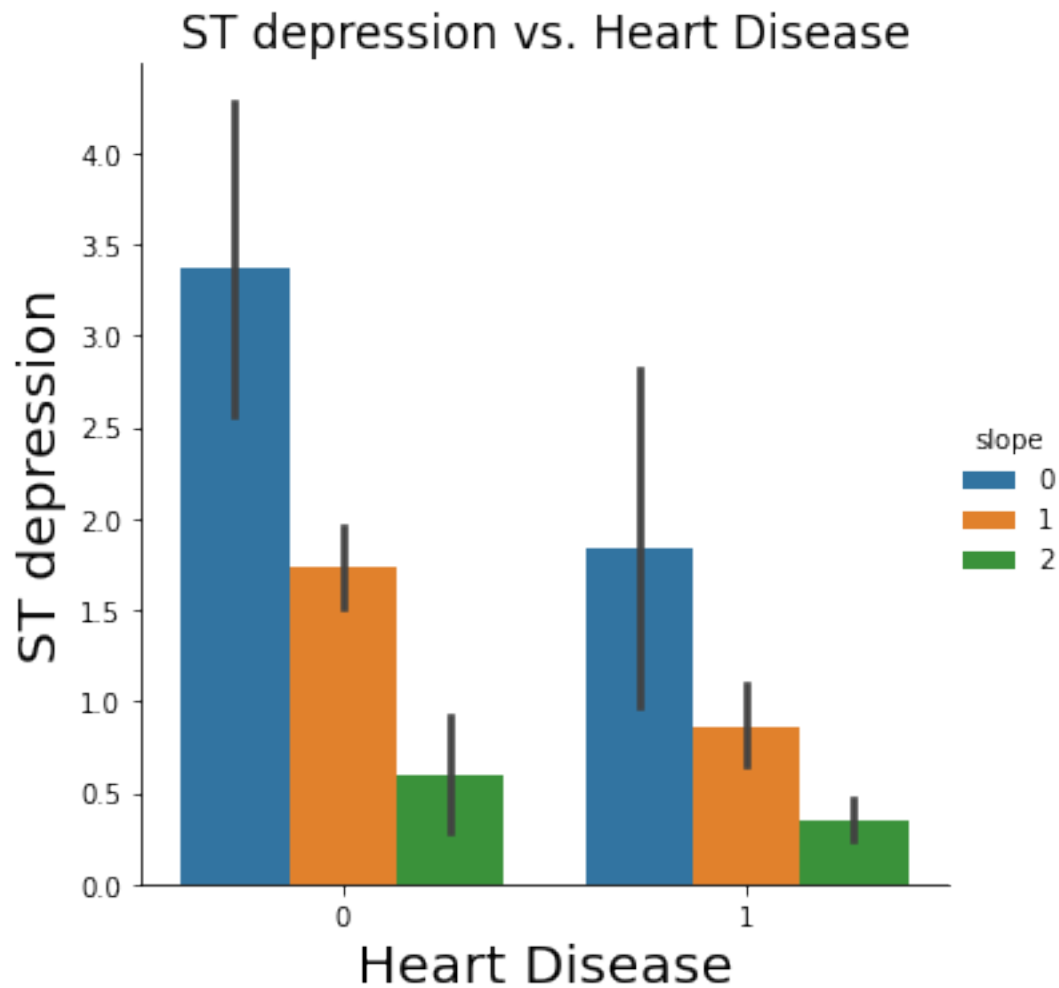
```
[74]: <seaborn.axisgrid.PairGrid at 0x7f4ab045bf10>
```



```
[75]: sns.catplot(x="target", y="oldpeak", hue="slope", kind="bar", data=data);

plt.title('ST depression vs. Heart Disease',size = 17)
plt.xlabel('Heart Disease',size=20)
plt.ylabel('ST depression',size=20)
```

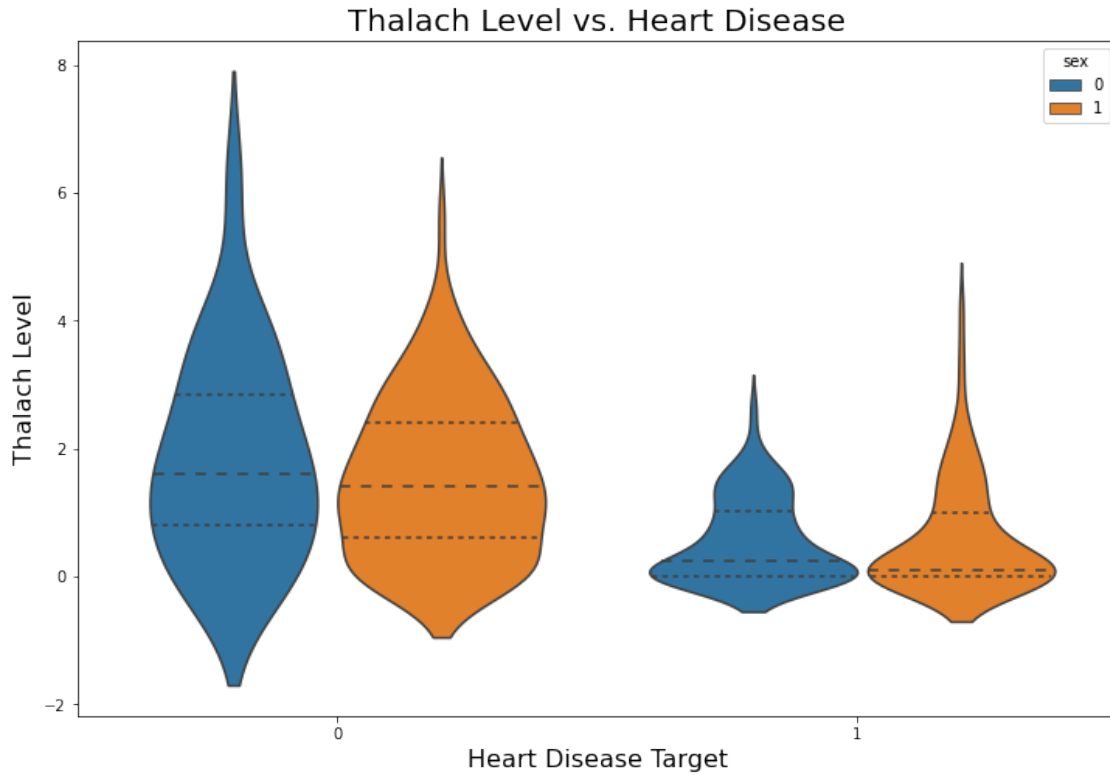
[75]: Text(26.426458333333343, 0.5, 'ST depression')



### 1.1.2 3.3 : Violin & Box Plots

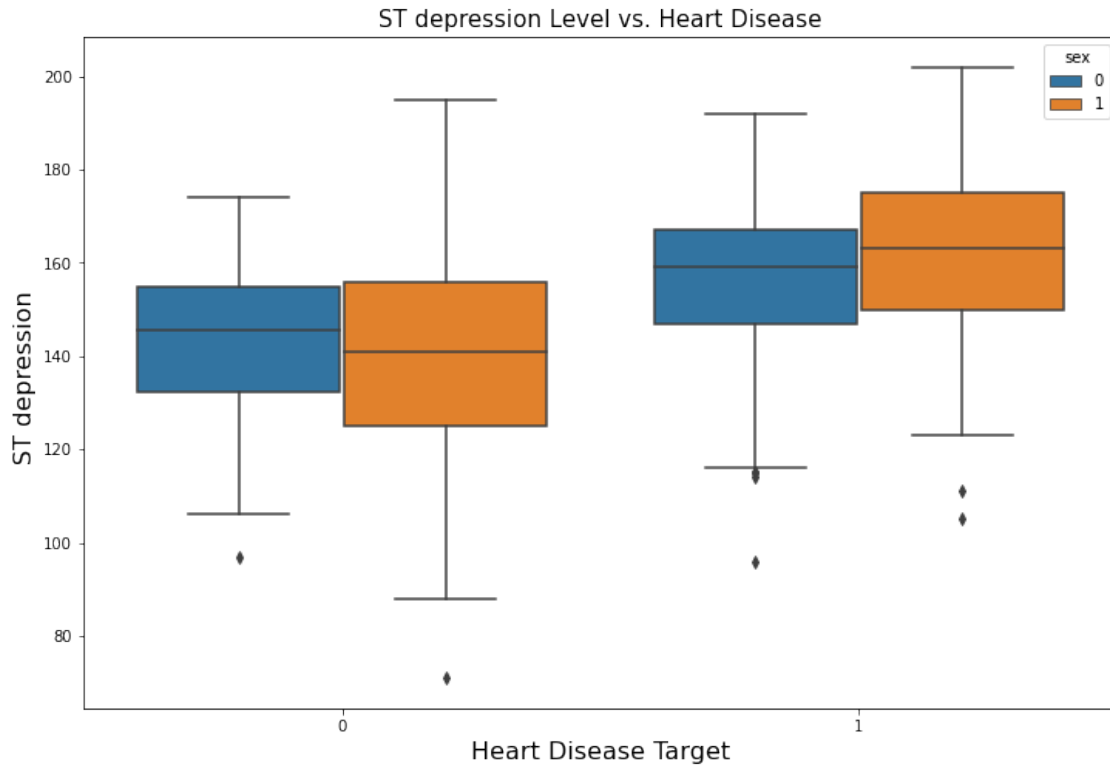
```
[76]: plt.figure(figsize=(12,8))
sns.violinplot(x= 'target', y= 'oldpeak',hue="sex", inner='quartile',data= data_1)
plt.title("Thalach Level vs. Heart Disease",fontsize=20)
plt.xlabel("Heart Disease Target", fontsize=16)
plt.ylabel("Thalach Level", fontsize=16)
```

[76]: Text(0, 0.5, 'Thalach Level')



```
[77]: plt.figure(figsize=(12,8))
sns.boxplot(x= 'target', y= 'thalach',hue="sex", data=data )
plt.title("ST depression Level vs. Heart Disease", fontsize = 15)
plt.xlabel("Heart Disease Target",fontsize=16)
plt.ylabel("ST depression ", fontsize=16)
```

```
[77]: Text(0, 0.5, 'ST depression ')
```



### 1.1.3 3.4 : Filtering data by positive Heart Disease patient

```
[78]: pos_data = data[data['target']==1]
pos_data.describe()
```

```
[78]:
```

	age	sex	cp	trestbps	chol	fbs	\
count	165.000000	165.000000	165.000000	165.000000	165.000000	165.000000	
mean	52.496970	0.563636	1.375758	129.303030	242.230303	0.139394	
std	9.550651	0.497444	0.952222	16.169613	53.552872	0.347412	
min	29.000000	0.000000	0.000000	94.000000	126.000000	0.000000	
25%	44.000000	0.000000	1.000000	120.000000	208.000000	0.000000	
50%	52.000000	1.000000	2.000000	130.000000	234.000000	0.000000	
75%	59.000000	1.000000	2.000000	140.000000	267.000000	0.000000	
max	76.000000	1.000000	3.000000	180.000000	564.000000	1.000000	

	restecg	thalach	exang	oldpeak	slope	ca	\
count	165.000000	165.000000	165.000000	165.000000	165.000000	165.000000	
mean	0.593939	158.466667	0.139394	0.583030	1.593939	0.363636	
std	0.504818	19.174276	0.347412	0.780683	0.593635	0.848894	
min	0.000000	96.000000	0.000000	0.000000	0.000000	0.000000	
25%	0.000000	149.000000	0.000000	0.000000	1.000000	0.000000	

50%	1.000000	161.000000	0.000000	0.200000	2.000000	0.000000
75%	1.000000	172.000000	0.000000	1.000000	2.000000	0.000000
max	2.000000	202.000000	1.000000	4.200000	2.000000	4.000000

	thal	target
count	165.000000	165.0
mean	2.121212	1.0
std	0.465752	0.0
min	0.000000	1.0
25%	2.000000	1.0
50%	2.000000	1.0
75%	2.000000	1.0
max	3.000000	1.0

### 1.1.4 3.5 : Filtering data by Negative Heart Disease patient

```
[79]: neg_data = data[data['target']==0]
neg_data.describe()
```

```
[79]:
```

	age	sex	cp	trestbps	chol	fbs	\
count	138.000000	138.000000	138.000000	138.000000	138.000000	138.000000	
mean	56.601449	0.826087	0.478261	134.398551	251.086957	0.159420	
std	7.962082	0.380416	0.905920	18.729944	49.454614	0.367401	
min	35.000000	0.000000	0.000000	100.000000	131.000000	0.000000	
25%	52.000000	1.000000	0.000000	120.000000	217.250000	0.000000	
50%	58.000000	1.000000	0.000000	130.000000	249.000000	0.000000	
75%	62.000000	1.000000	0.000000	144.750000	283.000000	0.000000	
max	77.000000	1.000000	3.000000	200.000000	409.000000	1.000000	

	restecg	thalach	exang	oldpeak	slope	ca	\
count	138.000000	138.000000	138.000000	138.000000	138.000000	138.000000	
mean	0.449275	139.101449	0.550725	1.585507	1.166667	1.166667	
std	0.541321	22.598782	0.499232	1.300340	0.561324	1.043460	
min	0.000000	71.000000	0.000000	0.000000	0.000000	0.000000	
25%	0.000000	125.000000	0.000000	0.600000	1.000000	0.000000	
50%	0.000000	142.000000	1.000000	1.400000	1.000000	1.000000	
75%	1.000000	156.000000	1.000000	2.500000	1.750000	2.000000	
max	2.000000	195.000000	1.000000	6.200000	2.000000	4.000000	

	thal	target
count	138.000000	138.0
mean	2.543478	0.0
std	0.684762	0.0
min	0.000000	0.0
25%	2.000000	0.0
50%	3.000000	0.0



```
75%      3.000000      0.0
max      3.000000      0.0
```

```
[80]: print("(Positive Patients ST depression): " + str(pos_data['oldpeak'].mean()))
      print("(Negative Patients ST depression): " + str(neg_data['oldpeak'].mean()))
```

```
(Positive Patients ST depression): 0.5830303030303029
(Negative Patients ST depression): 1.5855072463768118
```

```
[81]: print("(Positive Patients thalach): " + str(pos_data['thalach'].mean()))
      print("(Negative Patients thalach): " + str(neg_data['thalach'].mean()))
```

```
(Positive Patients thalach): 158.46666666666667
(Negative Patients thalach): 139.1014492753623
```

## 1.2 4. Machine Learning and Predictive Analytics

### 1.2.1 4.1 : Preparing Data for Modeling

```
[82]: X = data.iloc[:, :-1].values
      y = data.iloc[:, -1].values
```

### 1.2.2 4.2 : Splitting data into the Training and Test sets

```
[83]: from sklearn.model_selection import train_test_split
      x_train, x_test, y_train, y_test = train_test_split(X,y,test_size = 0.2,
      ↪random_state = 1)
```

```
[84]: from sklearn.preprocessing import StandardScaler
      sc = StandardScaler()
      x_train = sc.fit_transform(x_train)
      x_test = sc.transform(x_test)
```

### 1.2.3 4.3 : Modeling and Training

#### Model 1: Logistic Regression

```
[85]: from sklearn.metrics import classification_report
      from sklearn.linear_model import LogisticRegression

      model1 = LogisticRegression(random_state=1) # get instance of model
      model1.fit(x_train, y_train) # Train/Fit model

      y_pred1 = model1.predict(x_test) # get y predictions
      print(classification_report(y_test, y_pred1)) # output accuracy
```

	precision	recall	f1-score	support
0	0.77	0.67	0.71	30
1	0.71	0.81	0.76	31
accuracy			0.74	61
macro avg	0.74	0.74	0.74	61
weighted avg	0.74	0.74	0.74	61

### Model 2: K-NN (K-Nearest Neighbors)

```
[86]: from sklearn.metrics import classification_report
      from sklearn.neighbors import KNeighborsClassifier

      model2 = KNeighborsClassifier() # get instance of model
      model2.fit(x_train, y_train) # Train/Fit model

      y_pred2 = model2.predict(x_test) # get y predictions
      print(classification_report(y_test, y_pred2)) # output accuracy
```

	precision	recall	f1-score	support
0	0.78	0.70	0.74	30
1	0.74	0.81	0.77	31
accuracy			0.75	61
macro avg	0.76	0.75	0.75	61
weighted avg	0.76	0.75	0.75	61

### Model 3: SVM

```
[87]: from sklearn.metrics import classification_report
      from sklearn.svm import SVC

      model3 = SVC(random_state=1) # get instance of model
      model3.fit(x_train, y_train) # Train/Fit model

      y_pred3 = model3.predict(x_test) # get y predictions
      print(classification_report(y_test, y_pred3)) # output accuracy
```

	precision	recall	f1-score	support
0	0.80	0.67	0.73	30
1	0.72	0.84	0.78	31
accuracy			0.75	61

macro avg	0.76	0.75	0.75	61
weighted avg	0.76	0.75	0.75	61

#### Model 4: Naives Bayes Classifier

```
[88]: from sklearn.metrics import classification_report
      from sklearn.naive_bayes import GaussianNB

      model4 = GaussianNB() # get instance of model
      model4.fit(x_train, y_train) # Train/Fit model

      y_pred4 = model4.predict(x_test) # get y predictions
      print(classification_report(y_test, y_pred4)) # output accuracy
```

	precision	recall	f1-score	support
0	0.79	0.73	0.76	30
1	0.76	0.81	0.78	31
accuracy			0.77	61
macro avg	0.77	0.77	0.77	61
weighted avg	0.77	0.77	0.77	61

#### Model 5: Decision Trees

```
[89]: from sklearn.metrics import classification_report
      from sklearn.tree import DecisionTreeClassifier

      model5 = DecisionTreeClassifier(random_state=1) # get instance of model
      model5.fit(x_train, y_train) # Train/Fit model

      y_pred5 = model5.predict(x_test) # get y predictions
      print(classification_report(y_test, y_pred5)) # output accuracy
```

	precision	recall	f1-score	support
0	0.68	0.70	0.69	30
1	0.70	0.68	0.69	31
accuracy			0.69	61
macro avg	0.69	0.69	0.69	61
weighted avg	0.69	0.69	0.69	61

#### Model 6: Random Forest

```
[90]: from sklearn.metrics import classification_report
      from sklearn.ensemble import RandomForestClassifier

      model6 = RandomForestClassifier(random_state=1) # get instance of model
      model6.fit(x_train, y_train) # Train/Fit model

      y_pred6 = model6.predict(x_test) # get y predictions
      print(classification_report(y_test, y_pred6)) # output accuracy
```

	precision	recall	f1-score	support
0	0.88	0.70	0.78	30
1	0.76	0.90	0.82	31
accuracy			0.80	61
macro avg	0.82	0.80	0.80	61
weighted avg	0.81	0.80	0.80	61

#### 4.4 : Confusion Matrix

```
[91]: from sklearn.metrics import confusion_matrix, accuracy_score
      cm = confusion_matrix(y_test, y_pred6)
      print(cm)
      accuracy_score(y_test, y_pred6)
```

```
[[21  9]
 [ 3 28]]
```

```
[91]: 0.8032786885245902
```

##### 4.4.1 : Confusion Matrix: Interpretation

1 : True Positive = 21

2 : True negative = 28

3 : 9 is False positive and its false

4 : 3 is False Negative and it is false

1.2.4 Accuracy =  $(TP + TN) / (TP + TN + FP + FN)$ .

Accuracy =  $(21 + 28) / (21 + 28 + 9 + 3) = 0.80 = 80\%$  accuracy

### 1.3 5: Which Feature is more Important

```
[92]: # get importance
importance = model6.feature_importances_

# summarize feature importance
for i,v in enumerate(importance):
    print('Feature: %0d, Score: %.5f' % (i,v))
```

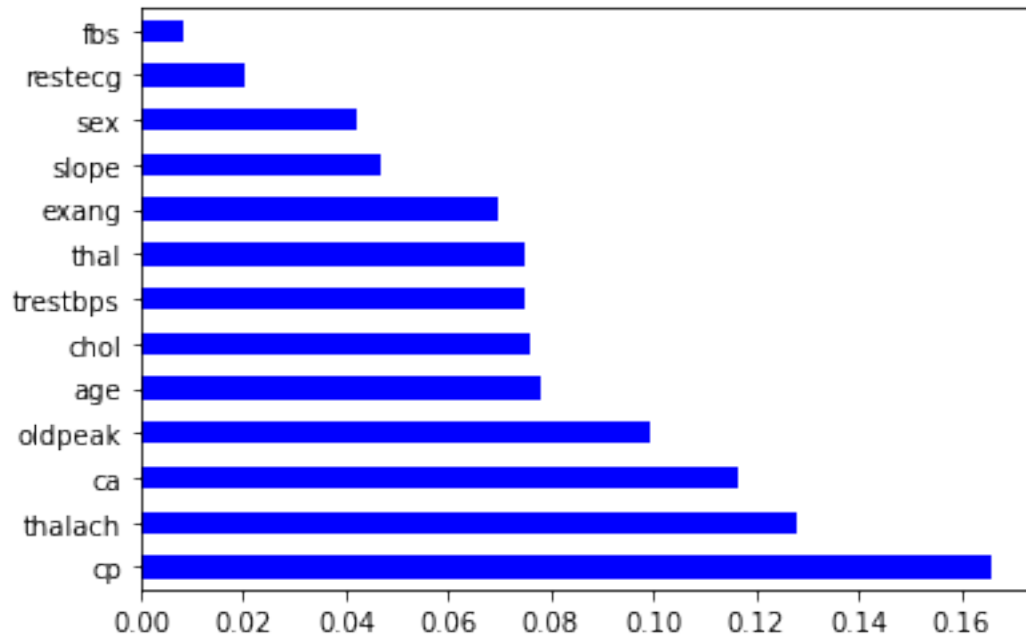
```
Feature: 0, Score: 0.07814
Feature: 1, Score: 0.04206
Feature: 2, Score: 0.16580
Feature: 3, Score: 0.07477
Feature: 4, Score: 0.07587
Feature: 5, Score: 0.00828
Feature: 6, Score: 0.02014
Feature: 7, Score: 0.12772
Feature: 8, Score: 0.06950
Feature: 9, Score: 0.09957
Feature: 10, Score: 0.04677
Feature: 11, Score: 0.11667
Feature: 12, Score: 0.07473
```

1.3.1 Since Feature : 7 has highest score (0.16580)

1.3.2 Feature :7 is more Important

```
[93]: index= data.columns[:-1]
importance = pd.Series(model6.feature_importances_, index=index)
importance.nlargest(13).plot(kind='barh', colormap='winter')
```

```
[93]: <AxesSubplot:>
```



from the graph above, we can conclude that following Features are more Important

1 : chest pain type (cp)

2 : maximum heart rate achieved (thalach)

3 : number of major vessels (ca)

4 : ST depression (oldpeak)

## 1.4 6 : Predictions

### 1.4.1 6.1 : Test set results: Predictions

```
[94]: y_pred = model6.predict(x_test)
      print(np.concatenate((y_pred.reshape(len(y_pred),1), y_test.
      ↪ reshape(len(y_test),1)),1))
```

```
[[0 0]
 [1 1]
 [0 0]
 [0 0]
```

[0 0]  
[0 0]  
[0 0]  
[1 1]  
[0 0]  
[1 1]  
[1 1]  
[0 0]  
[1 0]  
[0 0]  
[0 0]  
[1 0]  
[1 1]  
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```
[0 0]
[1 1]
[1 1]
[1 1]
[1 1]
[0 0]
[1 0]
[0 0]
[1 1]]
```

**1 :** First value is predicted value, Second value is actual value.

**2 :** If the predicted value = actual value then prediction correct

## **1.5 7 : Conclusions**

**7.1 :** Random Forest algorithm accuracy = 80%

**7.2 :** A ccuracy above 70% is considered good

**7.3 :** Above 80% acccuracy chances of Overfitting

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