

Practical Decision Tree

Decision Tree:

Decision tree is a one of the supervised machine learning algorithm. This algorithm can be used for regression and classification on problems. but mostly used classification problems.

The goal is to build up a classifire model that can predict the class or value of the target variable.

Graphical representation of all the possible solutions to a decision. Decisions are mainly based on some condition. Decision mode can easily be explained.

```
In [2]: #import Labrires
import numpy as np
import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn import tree
from sklearn.metrics import accuracy_score, confusion_matrix
import seaborn as sns
import matplotlib.pyplot as plt
```

```
In [3]: #import dataset
data='heart_1.csv'
df=pd.read_csv(data)
df.head()
```

```
Out[3]:
```

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

```
In [4]: df.shape
```

```
Out[4]: (303, 14)
```

```
In [5]: df[df['chol']>300].shape
```

```
Out[5]: (43, 14)
```

```
In [6]: f=df[df['thal']==2]
```

```
In [7]: f[f['target']==1].shape
```

```
Out[7]: (130, 14)
```

```
In [8]: df.shape
```

Out[8]: (303, 14)

In [9]: `df.isnull().sum()`

Out[9]:

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

In [10]: `df.dtypes`

Out[10]:

age	int64
sex	int64
cp	int64
trestbps	int64
chol	int64
fbs	int64
restecg	int64
thalach	int64
exang	int64
oldpeak	float64
slope	int64
ca	int64
thal	int64
target	int64

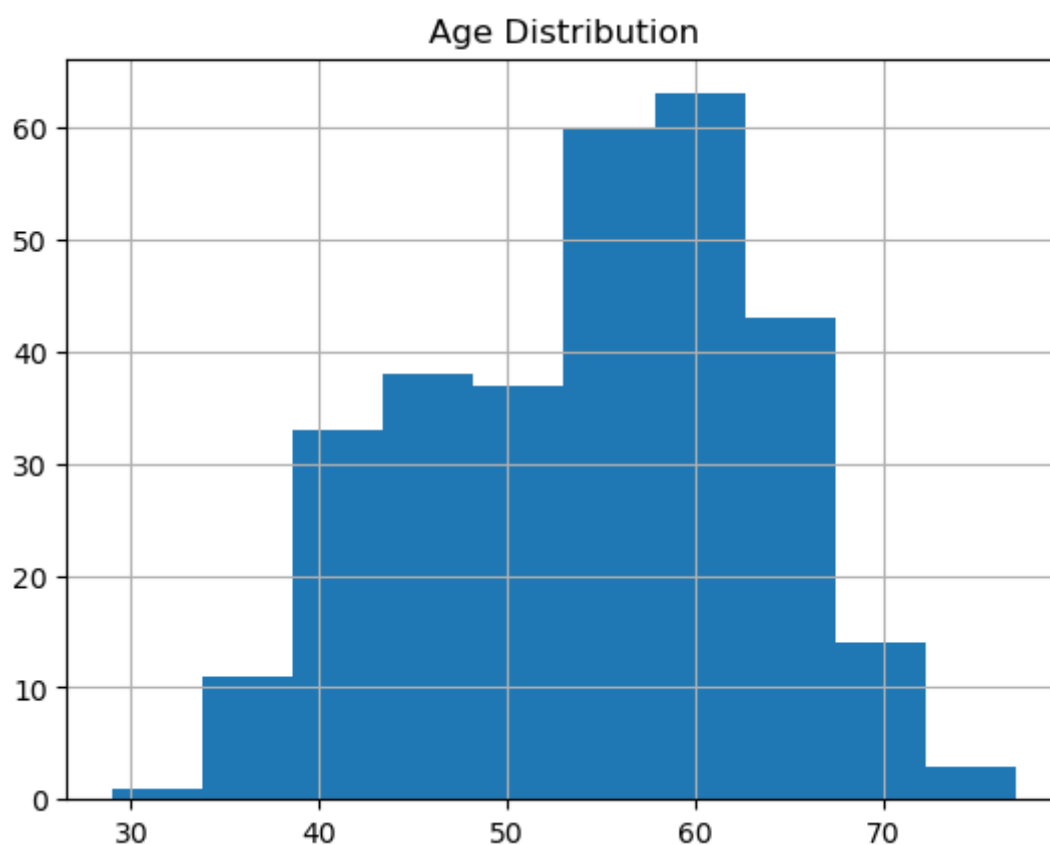
dtype: object

In [11]: `df.info()`

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 303 entries, 0 to 302
Data columns (total 14 columns):
 #   Column      Non-Null Count  Dtype  
---  -
 0   age         303 non-null   int64  
 1   sex         303 non-null   int64  
 2   cp          303 non-null   int64  
 3   trestbps    303 non-null   int64  
 4   chol        303 non-null   int64  
 5   fbs         303 non-null   int64  
 6   restecg     303 non-null   int64  
 7   thalach     303 non-null   int64  
 8   exang       303 non-null   int64  
 9   oldpeak     303 non-null   float64 
10   slope       303 non-null   int64  
11   ca          303 non-null   int64  
12   thal        303 non-null   int64  
13   target      303 non-null   int64  
dtypes: float64(1), int64(13)
memory usage: 33.3 KB
```

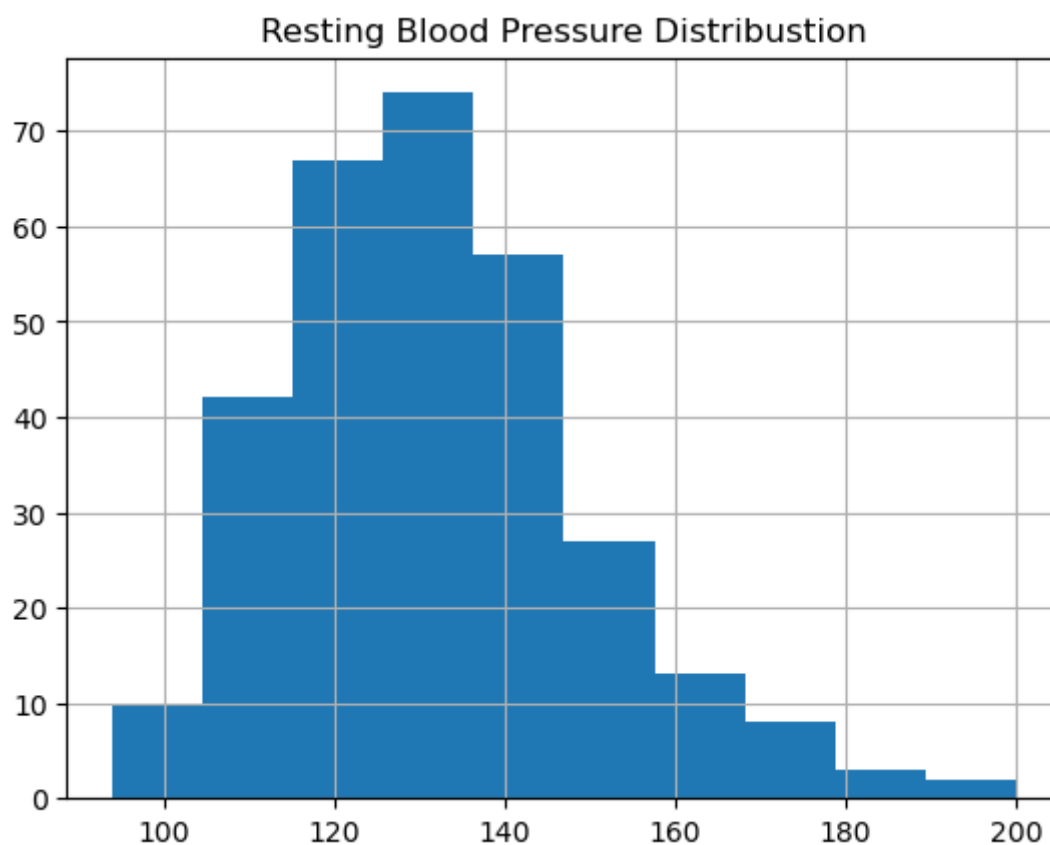
In [12]: `df['age'].hist(grid=True,bins=10);`
`plt.title('Age Distribution')`

Out[12]: Text(0.5, 1.0, 'Age Distribution')



```
In [13]: df['trestbps'].hist()  
plt.title('Resting Blood Pressure Distribution')
```

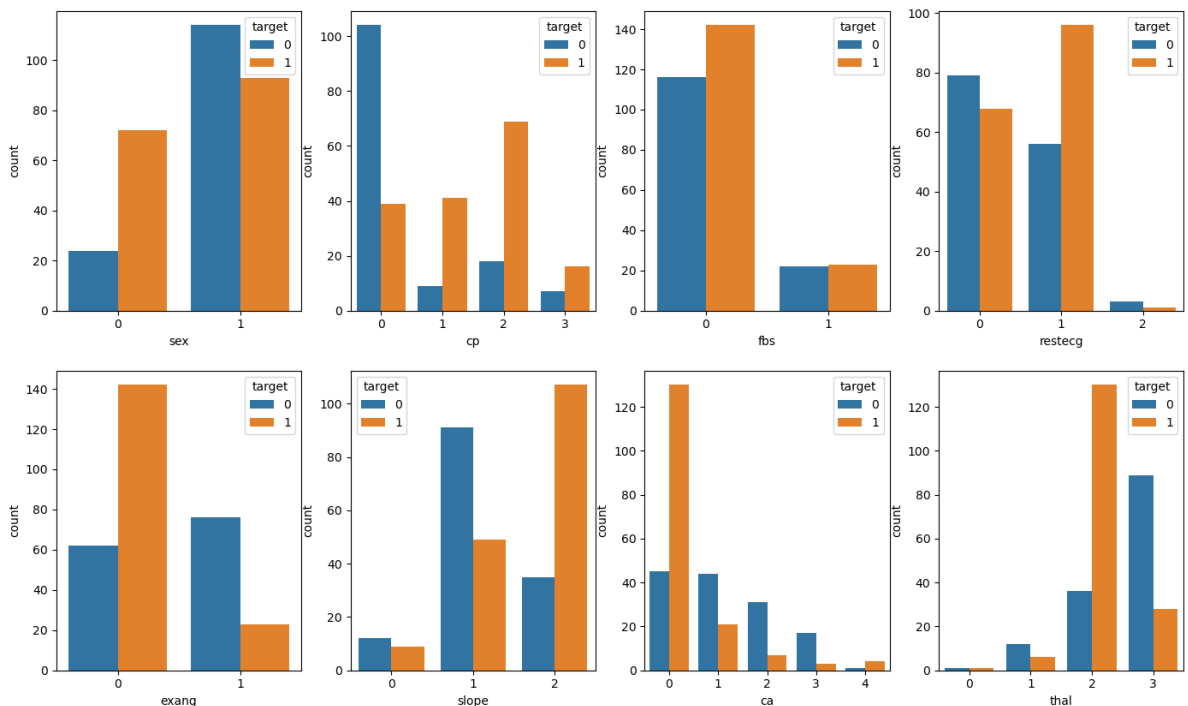
Out[13]: Text(0.5, 1.0, 'Resting Blood Pressure Distribution')



In the above graph, we are having a normal distribution

```
In [14]: fig, axes=plt.subplots(nrows=2,ncols=4,figsize=(17,10))
cat_feat=['sex','cp','fbs','restecg','exang','slope','ca','thal','target']

for idx, feature in enumerate(cat_feat):
    if feature != 'target':
        ax=axes[int(idx/4),idx%4]
        sns.countplot(x=feature,hue='target',data=df,ax=ax)
```



Let's get some insights from this chart

Chest pain: The heart disease rate is greater among the patients that feel any chest pain.

Restecg-Electrocardiograph results: The rate of heart disease diagnosis is higher for patients with a ST-T wave abnormality.

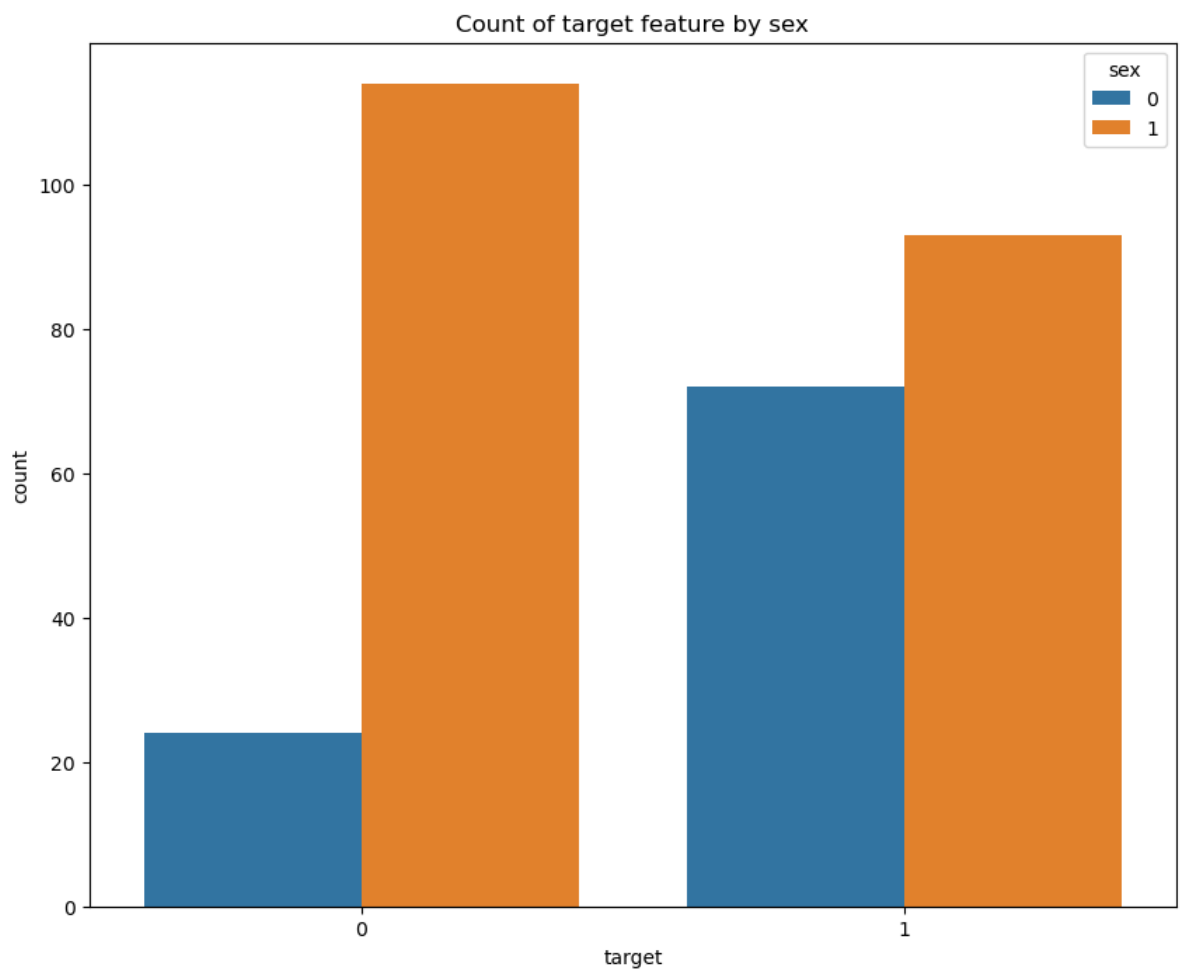
Slope: The ratio of patients diagnosed with heart disease is higher for slope=2.

ca: The diagnosed ratio decreases of ca between 1 and 3.

Thal: The diagnosed ratio is higher for thal=2.

```
In [15]: plt.rcParams['figure.figsize']=(10,8)
sns.countplot(x='target', hue='sex',data=df);
plt.title('Count of target feature by sex')
```

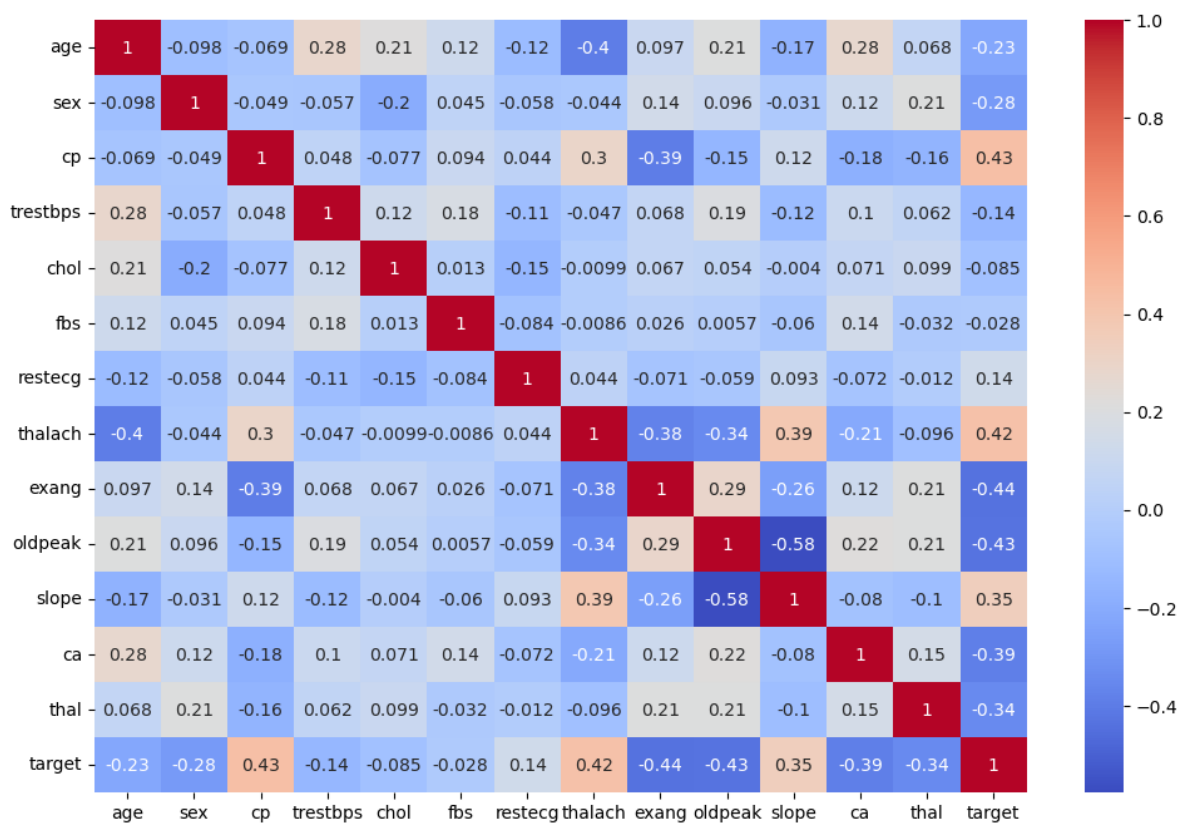
```
Out[15]: Text(0.5, 1.0, 'Count of target feature by sex')
```



The amount of healthy male people is greater than the amount of unhealthy for women the number of unhealthy women is higher

```
In [16]: plt.figure(figsize=(12,8))  
sns.heatmap(df.corr(),annot=True,cmap='coolwarm')
```

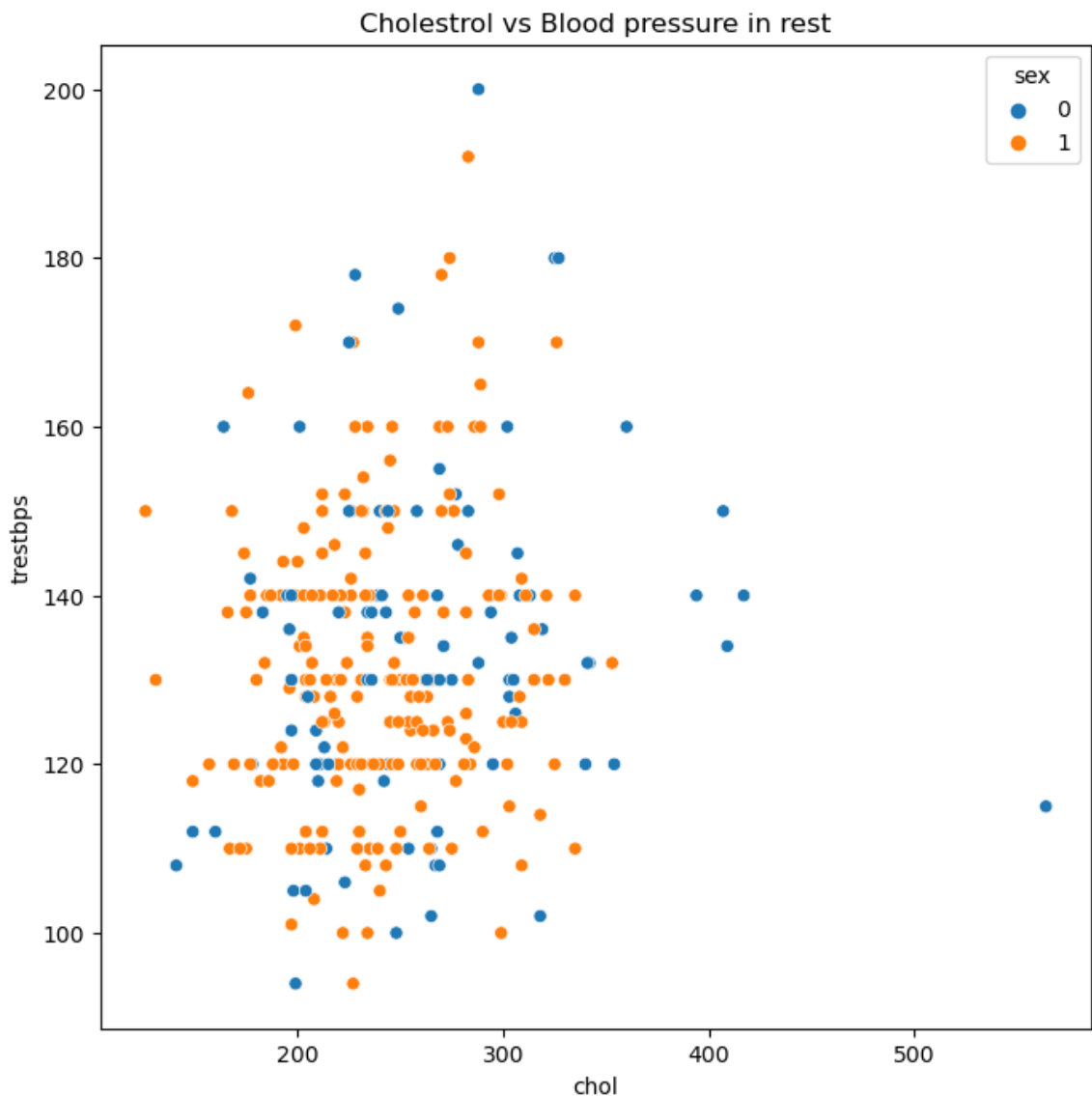
```
Out[16]: <AxesSubplot:>
```



Apparently there are no feature with a pretty strong correlation (above [0.7])

```
In [21]: plt.rcParams['figure.figsize']=(8,8)
sns.scatterplot(x='chol',y='trestbps',hue='sex',size=None,data=df)
plt.title('Cholestrol vs Blood pressure in rest')
```

```
Out[21]: Text(0.5, 1.0, 'Cholestrol vs Blood pressure in rest')
```



As can be seen there is a patient with high cholesterol. But there not a specific between those feel pain during exercise practice and those of not feel pain. we can use hue to filter by sex. it's also possible to filter using size='label_to_filter'

```
In [22]: X=df.drop(columns=['target'])#independant variable
y=df['target']#dependant variable
print(X.shape)
print(y.shape)
```

```
(303, 13)
(303,)
```

```
In [23]: X_train, X_test, y_train, y_test = train_test_split(X,y, random_state=0,test_size=0.3)
```

```
In [25]: print(X_train.shape)
print(X_test.shape)
```

```
(212, 13)
(91, 13)
```

```
In [26]: clf=tree.DecisionTreeClassifier()
clf.fit(X_train,y_train)
#predict
y_train_pred=clf.predict(X_train)
y_test_pred=clf.predict(X_test)
```

```
In [27]: y_train_pred
```

```
Out[27]: array([1, 1, 0, 1, 1, 0, 0, 1, 0, 0, 1, 0, 1, 0, 0, 0, 1, 1, 1, 0, 0, 0,
0, 1, 0, 1, 1, 1, 0, 1, 1, 1, 0, 1, 1, 1, 1, 1, 1, 0, 0, 1, 1,
0, 1, 1, 0, 0, 0, 0, 1, 0, 1, 1, 0, 1, 1, 1, 1, 1, 1, 0, 0, 1, 0,
0, 0, 1, 1, 1, 0, 1, 0, 1, 1, 0, 1, 1, 1, 1, 1, 0, 0, 1, 0, 1, 0,
0, 1, 0, 0, 1, 1, 1, 0, 0, 1, 1, 0, 1, 0, 1, 0, 1, 1, 0, 1, 1, 1,
0, 1, 0, 0, 0, 1, 1, 1, 1, 0, 1, 0, 1, 0, 1, 1, 1, 0, 1, 0, 1, 0,
0, 0, 1, 1, 0, 1, 0, 1, 1, 0, 0, 1, 0, 1, 1, 0, 1, 1, 1, 1, 1, 0,
0, 1, 1, 1, 0, 0, 1, 1, 1, 1, 1, 0, 1, 1, 0, 1, 1, 0, 1, 1, 0, 0,
1, 1, 1, 0, 0, 0, 1, 0, 0, 1, 0, 0, 1, 0, 1, 1, 0, 0, 0, 0, 1, 0,
1, 1, 1, 0, 0, 0, 0, 1, 0, 0, 0, 1, 1, 0], dtype=int64)
```

```
In [28]: y_test_pred
```

```
Out[28]: array([0, 0, 1, 0, 0, 0, 0, 0, 0, 1, 1, 0, 1, 1, 1, 0, 1, 0, 0, 1, 0,
                0, 0, 1, 1, 0, 0, 0, 1, 0, 0, 1, 1, 1, 1, 0, 1, 0, 0, 1, 0, 1, 0,
                0, 0, 0, 1, 0, 1, 1, 1, 1, 0, 1, 0, 1, 1, 1, 1, 1, 1, 0, 1, 1, 1,
                1, 0, 1, 1, 0, 1, 1, 0, 0, 0, 0, 0, 1, 1, 0, 0, 0, 1, 0, 1, 0, 1,
                0, 1, 0], dtype=int64)
```

```
In [30]: confusion_matrix(y_train_pred,y_train)
```

```
Out[30]: array([[ 94,  0],
                [ 0, 118]], dtype=int64)
```

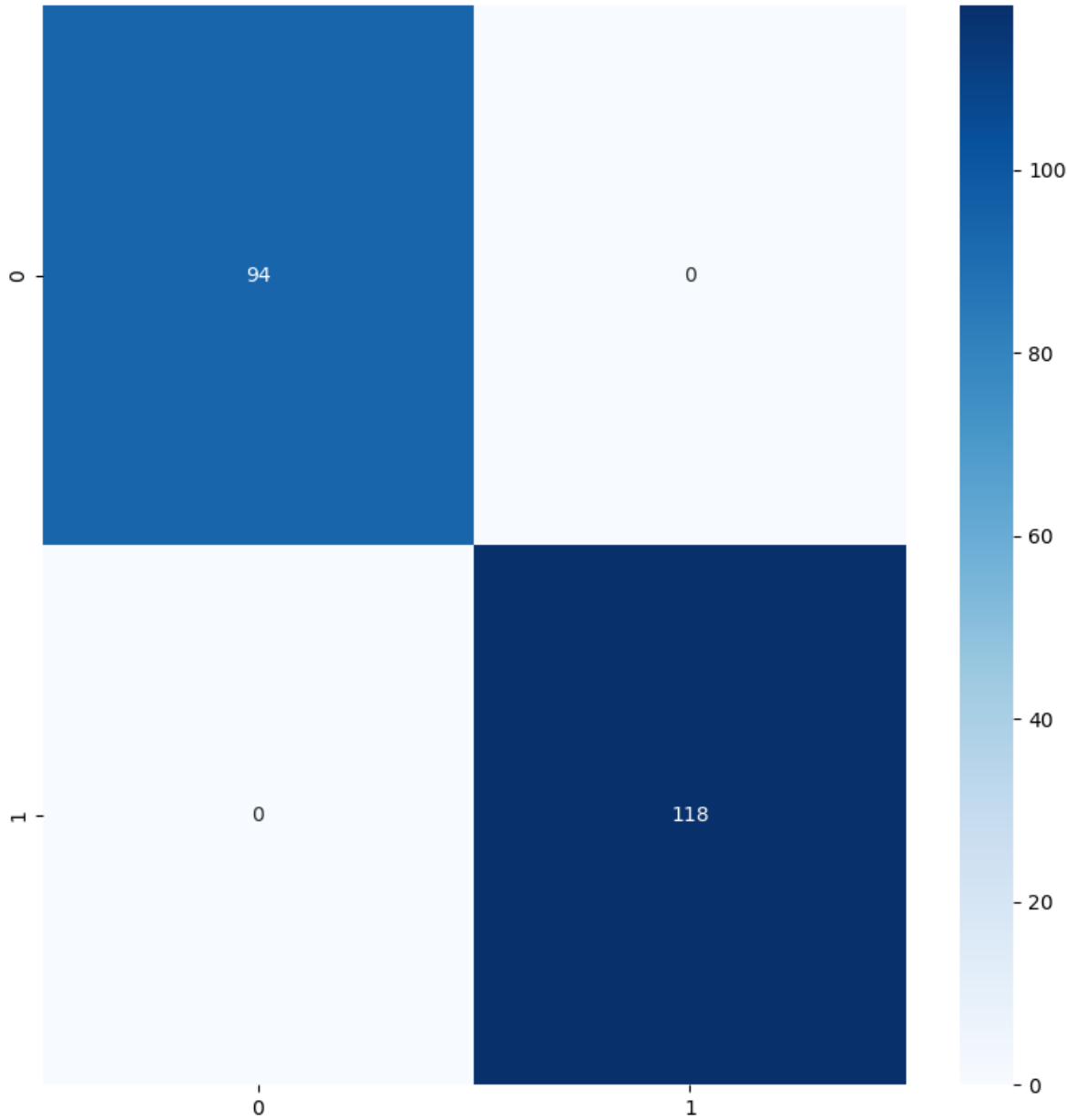
```
In [29]: #helper function
def plot_confusionmatrix(y_train_pred, y_train, dom):
    print(f'{dom} Confusion matrix')
    cf=confusion_matrix(y_train_pred,y_train)
    sns.heatmap(cf,annot=True,cmap='Blues',fmt='g')
    plt.tight_layout()
    plt.show()
```

```
In [31]: print(f'Train Score {accuracy_score(y_train_pred,y_train)}')
print(f'Test Score {accuracy_score(y_test_pred,y_test)}')
```

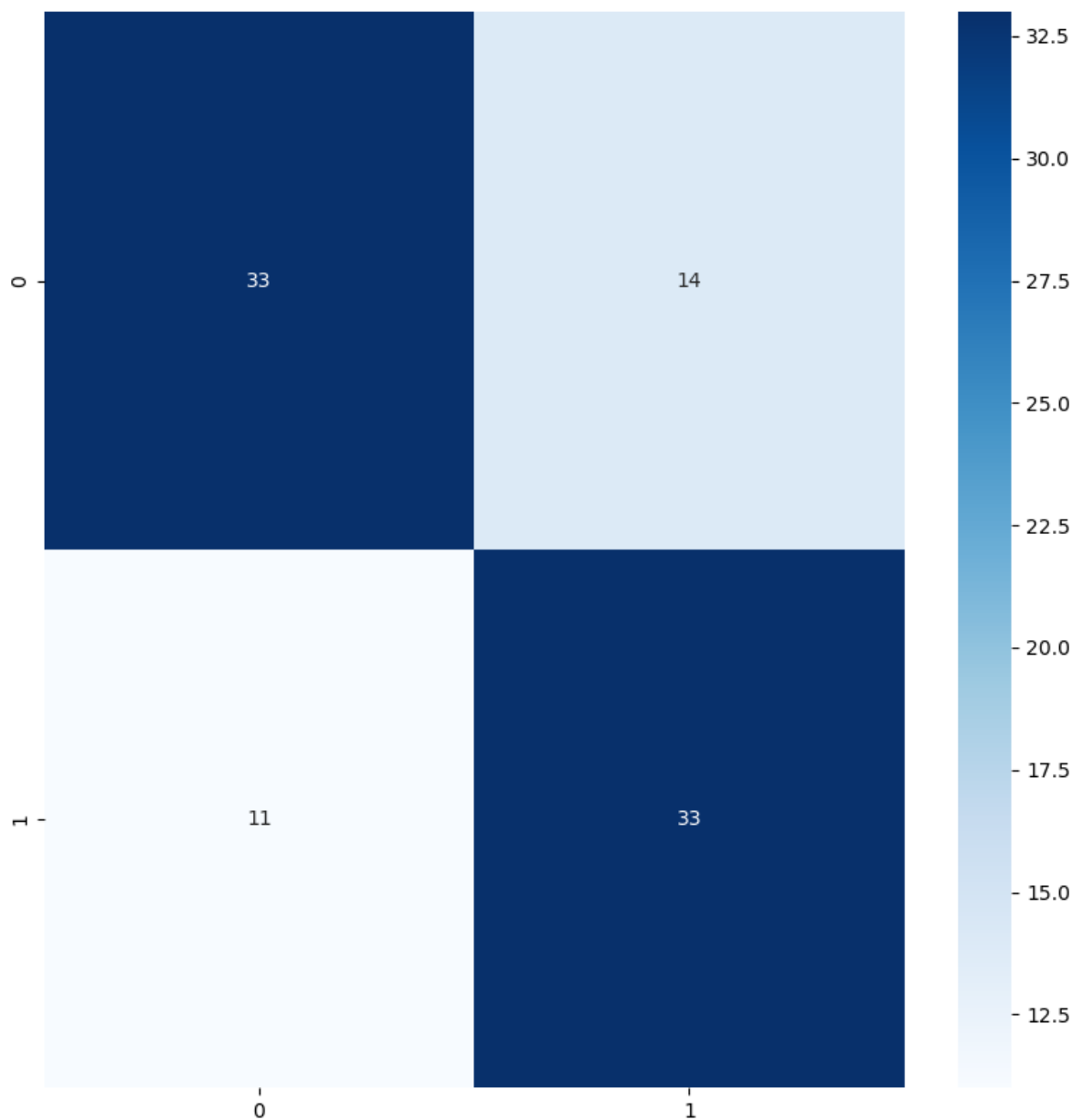
Train Score 1.0
Test Score 0.7252747252747253

```
In [32]: plot_confusionmatrix(y_train_pred,y_train,dom='Train')
          plot_confusionmatrix(y_test_pred,y_test,dom='Test')
```

Train Confusion matrix



Test Confusion matrix



```
In [48]: c_parameter_name='max_depth'
c_parameter_value=[1,2,3,4,5,6,7,8,9,10,11,12,13,14,15]
df=pd.DataFrame(columns=[c_parameter_name,'accuracy'])

for input_parameter in c_parameter_value:
    model=tree.DecisionTreeClassifier(max_depth=input_parameter,splitter='best')
    model.fit(X_train,y_train)
    y_pred=model.predict(X_test)
    acc_score=accuracy_score(y_test,y_pred)*100
    df=df.append({c_parameter_name:input_parameter, 'accuracy':acc_score}, ignore_
print(df)
```

	max_depth	accuracy
0	1.0	76.923077
1	2.0	73.626374
2	3.0	81.318681
3	4.0	76.923077
4	5.0	73.626374
5	6.0	72.527473
6	7.0	73.626374
7	8.0	71.428571
8	9.0	72.527473
9	10.0	72.527473
10	11.0	71.428571
11	12.0	71.428571
12	13.0	71.428571
13	14.0	75.824176
14	15.0	75.824176

```
C:\Users\Admin\AppData\Local\Temp\ipykernel_11352\3743572166.py:10: FutureWarning:  
The frame.append method is deprecated and will be removed from pandas in a future  
version. Use pandas.concat instead.  
df=df.append({'c_parameter_name':input_parameter, 'accuracy':acc_score}, ignore_in  
dex=True)  
C:\Users\Admin\AppData\Local\Temp\ipykernel_11352\3743572166.py:10: FutureWarning:  
The frame.append method is deprecated and will be removed from pandas in a future  
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```

```
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version. Use pandas.concat instead.
    df=df.append({c_parameter_name:input_parameter, 'accuracy':acc_score}, ignore_in
dex=True)
```

```
In [54]: from sklearn.tree import DecisionTreeClassifier
for input_parameter in c_parameter_value:
    model= DecisionTreeClassifier(max_depth=input_parameter, splitter='best')
    model.fit(X_train,y_train)
    y_pred1=model.predict(X_test)
    acc_score=accuracy_score(y_test,y_pred)*100
    df=df.append({c_parameter_name:input_parameter, 'accuracy':acc_score},ignore_in
```

```
C:\Users\Admin\AppData\Local\Temp\ipykernel_11352\2617876694.py:7: FutureWarning:
The frame.append method is deprecated and will be removed from pandas in a future
version. Use pandas.concat instead.
    df=df.append({c_parameter_name:input_parameter, 'accuracy':acc_score},ignore_ind
ex=True)
C:\Users\Admin\AppData\Local\Temp\ipykernel_11352\2617876694.py:7: FutureWarning:
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version. Use pandas.concat instead.
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ex=True)
C:\Users\Admin\AppData\Local\Temp\ipykernel_11352\2617876694.py:7: FutureWarning:
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```

```
ex=True)
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The frame.append method is deprecated and will be removed from pandas in a future
version. Use pandas.concat instead.
    df=df.append({c_parameter_name:input_parameter, 'accuracy':acc_score},ignore_ind
ex=True)
```

```
In [47]: from sklearn.metrics import classification_report
print(classification_report(y_test_pred,y_test))
```

	precision	recall	f1-score	support
0	0.75	0.70	0.73	47
1	0.70	0.75	0.73	44
accuracy			0.73	91
macro avg	0.73	0.73	0.73	91
weighted avg	0.73	0.73	0.73	91

```
In [ ]:
```