## **Importing Libraries**

```
import numpy as np
import pandas as pd
from sklearn.metrics import confusion_matrix
from sklearn.model_selection import train_test_split
from sklearn.tree import DecisionTreeClassifier
from sklearn.metrics import accuracy_score
from sklearn.metrics import classification_report
from sklearn import tree
import matplotlib.pyplot as plt
import seaborn as sns
%matplotlib inline
```

# Import Data from Excel Sheet

```
In [2]: file_name = "Dataset.xls"
    sheet = 'Heart Disease 1'

df = pd.read_excel(io=file_name, sheet_name=sheet)
```

# **Data Processing**

## Select the given attributes

```
In [3]: #BMI, Smoking, PhysicalHealth, Race, Age
df = df[['BMI', 'Smoking', 'PhysicalHealth', 'Race', 'AgeCategory', 'SkinCancer']]
```

## Remove '0' values from PhysicalHealth

```
In [4]: df = df[df.PhysicalHealth != 0]
```

### Select first 100 rows

```
In [5]: df=df.head(100)
    df
```

Out[5]:		BMI	Smoking	PhysicalHealth	Race	AgeCategory	SkinCancer
	0	16.60	Yes	3	White	55-59	Yes
	2	26.58	Yes	20	White	65-69	No
	4	23.71	No	28	White	40-44	No
	5	28.87	Yes	6	Black	75-79	No
	6	21.63	No	15	White	70-74	Yes
	•••						
	218	33.01	Yes	14	Black	70-74	No
	219	25.84	Yes	20	White	55-59	No
	222	21.21	No	7	Black	80 or older	No
	224	21.30	No	2	White	60-64	Yes
	228	43.27	No	2	Black	50-54	No

100 rows × 6 columns

# Change the intervals to midpoint values

```
In [6]:
        for i in df['AgeCategory']:
            if i == '18-24':
                df['AgeCategory'] = df['AgeCategory'].replace(['18-24'], 21)
            elif i == '25-29':
                df['AgeCategory'] = df['AgeCategory'].replace(['25-29'], 27)
            elif i == '30-34':
                df['AgeCategory'] = df['AgeCategory'].replace(['30-34'], 32)
            elif i == '35-39':
                df['AgeCategory'] = df['AgeCategory'].replace(['35-39'], 37)
            elif i == '40-44':
                df['AgeCategory'] = df['AgeCategory'].replace(['40-44'], 42)
            elif i == '45-49':
                df['AgeCategory'] = df['AgeCategory'].replace(['45-49'], 47)
            elif i == '50-54':
                df['AgeCategory'] = df['AgeCategory'].replace(['50-54'], 52)
            elif i == '55-59':
                df['AgeCategory'] = df['AgeCategory'].replace(['55-59'], 57)
            elif i == '60-64':
                df['AgeCategory'] = df['AgeCategory'].replace(['60-64'], 62)
            elif i == '65-69':
                df['AgeCategory'] = df['AgeCategory'].replace(['65-69'], 67)
            elif i == '70-74':
                df['AgeCategory'] = df['AgeCategory'].replace(['70-74'], 72)
            elif i == '75-79':
                df['AgeCategory'] = df['AgeCategory'].replace(['75-79'], 77)
            elif i == '80 or older':
                df['AgeCategory'] = df['AgeCategory'].replace(['80 or older'], 87)
```

## Change string to Int/Float

```
In [7]: from sklearn.preprocessing import LabelEncoder
In [8]: le_race = LabelEncoder()
```

```
df['Race'] = le_race.fit_transform(df['Race'])
for i in df['Smoking']:
    if i == 'Yes':
        df['Smoking'] = df['Smoking'].replace(['Yes'], int(1))
    elif i == 'No':
        df['Smoking'] = df['Smoking'].replace(['No'], int(0))

for i in df['SkinCancer']:
    if i == 'Yes':
        df['SkinCancer'] = df['SkinCancer'].replace(['Yes'], int(1))
    elif i == 'No':
        df['SkinCancer'] = df['SkinCancer'].replace(['No'], int(0))

df
```

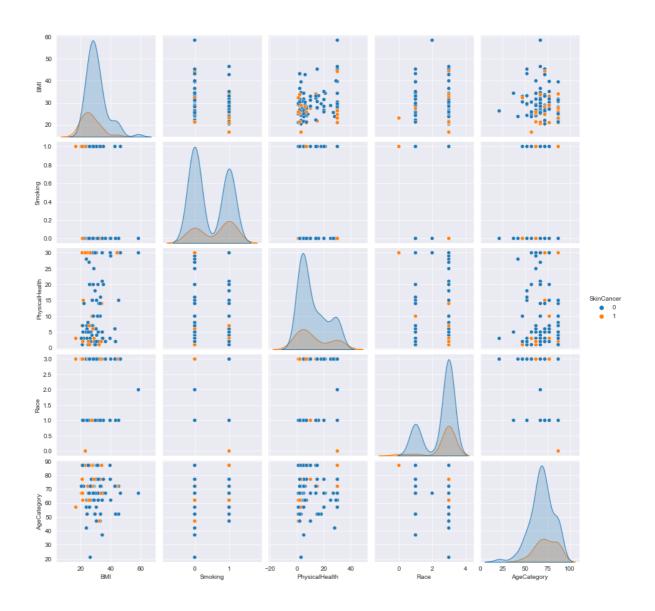
Out[8]:

	ВМІ	Smoking	PhysicalHealth	Race	AgeCategory	SkinCancer
0	16.60	1	3	3	57	1
2	26.58	1	20	3	67	0
4	23.71	0	28	3	42	0
5	28.87	1	6	1	77	0
6	21.63	0	15	3	72	1
•••						
218	33.01	1	14	1	72	0
219	25.84	1	20	3	57	0
222	21.21	0	7	1	87	0
224	21.30	0	2	3	62	1
228	43.27	0	2	1	52	0

100 rows × 6 columns

# **Exploratory Data Analysis**

```
In [9]: sns.pairplot(data=df, hue = 'SkinCancer')
Out[9]: <seaborn.axisgrid.PairGrid at 0x2242de58d60>
```



# **Model Fitting & Test**

# Split Input and Target Columns and Split Train-Test Data

```
In [10]: target = df['SkinCancer']
  inputs = df.drop(['SkinCancer'], axis=1)

x = inputs.values
y = target.values

X_train, X_test, y_train, y_test = train_test_split(x, y, test_size=0.3, random_startest_split(x, y, y, test_size=0.3, random_startest_split(x, y, y, test_size=0.3, random_startest_split(x, y, y, test_size
```

### **Decision Tree Classifier**

```
In [11]: clf_tree = DecisionTreeClassifier(criterion='entropy', random_state = 100, max_dept
clf_tree.fit(X_train, y_train)
```

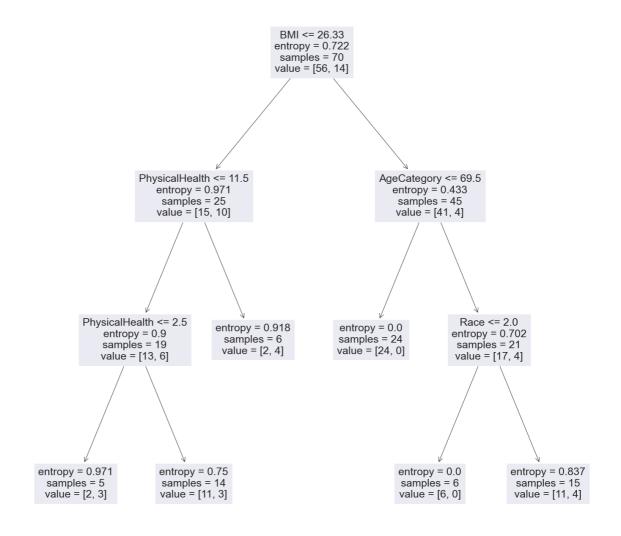
```
Out[11]:

DecisionTreeClassifier

DecisionTreeClassifier(criterion='entropy', max_depth=3, min_samples_leaf =5, random_state=100)
```

#### **Plot Tree**

```
In [12]: fig, ax = plt.subplots(figsize=(25,25))
    tree.plot_tree(clf_tree, ax=ax, feature_names=['BMI', 'Smoking', 'PhysicalHealth',
    plt.show()
```



### **Show Predicted Values**

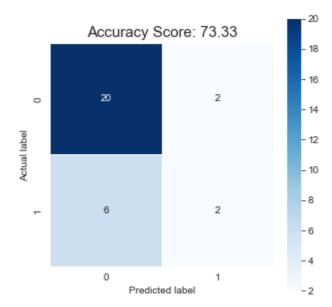
### **Show Results**

```
In [14]:
        print("Confusion Matrix:\n",confusion_matrix(y_test, y_pred))
         print ("Accuracy : ",accuracy_score(y_test,y_pred)*100)
         print("Report :\n",classification_report(y_test, y_pred))
        Confusion Matrix:
         [[20 2]
         [62]]
        Report :
                      precision
                                   recall f1-score
                                                    support
                   0
                          0.77
                                    0.91
                                             0.83
                                                        22
                                    0.25
                   1
                          0.50
                                             0.33
                                                         8
                                             0.73
                                                        30
            accuracy
                                    0.58
                                             0.58
                                                        30
           macro avg
                          0.63
        weighted avg
                          0.70
                                    0.73
                                             0.70
                                                        30
```

## **Confusion Matrix - Using Seaborn**

```
In [15]: cm = confusion_matrix(y_test, y_pred)
   plt.figure(figsize=(5,5))
   sns.heatmap(data=cm, linewidths=.5, annot=True, square = True, cmap = 'Blues')
   plt.ylabel('Actual label')
   plt.xlabel('Predicted label')
   all_sample_title = 'Accuracy Score: {:.2f}'.format(accuracy_score(y_test,y_pred)*10
   plt.title(all_sample_title, size = 15)
```

Out[15]: Text(0.5, 1.0, 'Accuracy Score: 73.33')



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Types of Data:

BMI -> Numerical (Interval)

Smoking > Categorical (Nomed)

Physical Health > Numerical

Race -> Categorical

Age Category > Numerical

Skin Cancer > Categorial

Calculating Entropy

For BMI, we can make data intervals:

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$$\alpha < 20 \rightarrow 1$$

Entropy (BMI) 
$$= -\frac{1}{100} \log_2(\frac{1}{100}) - \frac{22}{100} \log_2(\frac{22}{100})$$
  
 $-\frac{39}{100} \log_2(\frac{39}{100}) - \frac{25}{100} \log_2(\frac{25}{100})$   
 $-\frac{5}{100} \log_2(\frac{5}{100}) - \frac{9}{100} \log_2(\frac{8}{100})$   
 $= 2.08441$ 

# Smoking:

... Entropy (Smoking) = 
$$-\frac{53}{100}\log_2(\frac{53}{100}) - \frac{47}{100}\log_2(\frac{47}{100})$$
  
= 0.9974

# Physical Health:

:. Entropy (Physical Health) 
$$= \frac{-31}{106} \log_2 \left( \frac{31}{100} \right) - \frac{31}{100} \log_2 \left( \frac{31}{100} \right)$$

$$= \frac{11}{100} \log_2 \left( \frac{11}{100} \right) - \frac{27}{100} \log_2 \left( \frac{27}{100} \right)$$

$$= 1.90789$$

Race:

Entropy (Race) = 
$$-\frac{1}{100}\log_2\left(\frac{1}{100}\right) - \frac{20}{100}\log_2\left(\frac{20}{100}\right)$$
  
 $-\frac{1}{100}\log_2\left(\frac{1}{100}\right) - \frac{78}{100}\log_2\left(\frac{78}{100}\right)$   
= 0.87685

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Age Category:

$$x < 30 \rightarrow 1$$

$$-\frac{1}{100}\log_2\left(\frac{1}{100}\right) - \frac{5}{100}\log_2\left(\frac{5}{100}\right)$$

$$-\frac{44}{100}\log_2\left(\frac{44}{100}\right) - \frac{50}{100}\left(\frac{50}{100}\right)$$

noil gardal

= 1,30368

Skin Cancer:

FF 5 05 2 4213

00 = 00 > 4-54