## Ávore de Decisão

### Mini-projeto 2 - Sistemas Inteligentes

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### Equipe 6

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# Setup

Importando dependências, conectando Google Drive, lendo datasets e criando os respectivos dataframes.

### **Dependências**

### In [122]:

```
import numpy as np
import pandas as pd

import matplotlib.pyplot as plt
import seaborn as sns

from sklearn.ensemble import RandomForestClassifier
from sklearn.tree import DecisionTreeClassifier, plot_tree
from sklearn.model_selection import train_test_split, GridSearchCV
from sklearn.metrics import classification_report, accuracy_score, precision_score, rec
all_score, f1_score, roc_auc_score, confusion_matrix
```

#### **Drive**

```
In [2]:
```

```
from google.colab import drive
drive.mount('/content/drive')

base_path = '/content/drive/MyDrive/'
```

Mounted at /content/drive

IRIS Dataset - <a href="https://www.kaggle.com/datasets/arshid/iris-flower-dataset">https://www.kaggle.com/datasets/arshid/iris-flower-dataset</a> <a href="https://www.kaggle.com/datasets/arshid/iris-flower-dataset">https://www.kaggle.com/datasets/arshid/iris-flower-dataset</a>)

```
In [3]:
```

```
iris_dataset_path = base_path + 'IRIS.csv'
```

Titanic Dataset - https://www.kaggle.com/c/titanic/data (https://www.kaggle.com/c/titanic/data)

### In [4]:

```
titanic_train_dataset_path = base_path + 'titanic/train.csv'
titanic_test_dataset_path = base_path + 'titanic/test.csv'
```

Lower Back Pain Symptoms Dataset - <a href="https://www.kaggle.com/datasets/sammy123/lower-back-pain-symptoms-dataset">https://www.kaggle.com/datasets/sammy123/lower-back-pain-symptoms-dataset</a>)

### In [204]:

```
pain_dataset_path = base_path + 'Dataset_spine.csv'
```

### Lendo datasets

### In [5]:

```
iris_df = pd.read_csv(iris_dataset_path)
```

### In [6]:

```
titanic_df = pd.read_csv(titanic_train_dataset_path)
titanic_test_df = pd.read_csv(titanic_test_dataset_path) # Não possui a coluna de class
ificação (Survived)
```

### In [230]:

```
pain_df = pd.read_csv(pain_dataset_path)
pain_df = pain_df.drop(pain_df.columns[[13]], axis=1) # Removendo a coluna 13
```

### **Funções**

### In [98]:

```
def plot_confusion_matrix(y_test, y_pred):
    cm = confusion_matrix(y_test, y_pred)
    fig, ax = plt.subplots(figsize=(2,2))
    sns.heatmap(cm, annot=True, fmt='d')
```

## Experimento: Iris Flower Dataset

### Análise do Dataset

```
In [ ]:
iris_df.info()
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 150 entries, 0 to 149
Data columns (total 5 columns):
    Column
                   Non-Null Count Dtype
    sepal_length 150 non-null
                                   float64
 0
 1
   sepal_width 150 non-null
                                   float64
     petal length 150 non-null
 2
                                   float64
 3
    petal_width
                   150 non-null
                                   float64
                                   object
     species
                   150 non-null
dtypes: float64(4), object(1)
memory usage: 6.0+ KB
In [ ]:
iris_df['species'].unique()
Out[ ]:
array(['Iris-setosa', 'Iris-versicolor', 'Iris-virginica'], dtype=object)
In [ ]:
iris_df['species'].value_counts()
Out[ ]:
Iris-setosa
                   50
Iris-versicolor
                   50
Iris-virginica
                   50
Name: species, dtype: int64
In [ ]:
#iris df.describe()
In [ ]:
#sns.pairplot(iris_df, hue ='species')
In [ ]:
#iris_df.hist(bins=50, figsize=(5,5))
#plt.show()
```

### In [ ]:

```
#sns.set_style("whitegrid")
#data = np.random.normal(size=(20, 6)) + np.arange(6) / 2
#sns.set(rc={'figure.figsize':(7,4)})
#sns.boxplot(data = iris_df.iloc[:,:-1])
```

### X Criando modelo

### In [ ]:

```
iris_x = iris_df.iloc[:, :-1]
iris_y = iris_df.iloc[:, -1]
iris_x_train, iris_x_test, iris_y_train, iris_y_test = train_test_split(iris_x, iris_y, test_size = 0.25, random_state = 0)
```

### In [ ]:

```
classifier = DecisionTreeClassifier()
classifier.fit(iris_x_train, iris_y_train)
```

### In [ ]:

```
plot_tree(classifier, feature_names=iris_df.columns[: -1])
plt.show()
```

```
petal_length <= 2.35
                                    gini = 0.665
                                  samples = 112
                                value = [37, 34, 41]
                                          petal length <= 4.95
                          gini = 0.0
                                               gini = 0.496
                        samples = 37
                                              samples = 75
                      value = [37, 0, 0]
                                            value = [0, 34, 41]
          petal width <= 1.65
                                                                            petal width <= 1.75
             gini = 0.153
                                                                                 \overline{gini} = 0.05
             samples = 36
                                                                                samples = 39
           value = [0, 33, 3]
                                                                              value = [0, 1, 38]
                                                                 petal width <= 1.65
                     sepal width <= 3.1
   gini = 0.0
                                                                                             gini = 0.0
                                                                      gini = 0.375
                         gini = 0.375
  samples = 32
                                                                                           samples = 35
                                                                     samples = 4
                        samples = 4
value = [0, 32, 0]
                                                                                         value = [0, 0, 35]
                       value = [0, 1, 3]
                                                                   value = [0, 1, 3]
              gini = 0.0
                                     gini = 0.0
                                                           gini = 0.0
                                                                                 gini = 0.0
             samples = 3
                                                          samples = 3
                                                                                samples = 1
                                   samples = 1
           value = [0, 0, 3]
                                  value = [0, 1, 0]
                                                        value = [0, 0, 3]
                                                                              value = [0, 1, 0]
```

### In [ ]:

```
iris_y_pred = classifier.predict(iris_x_test)
```

#### In [ ]:

```
report = classification_report(iris_y_test, iris_y_pred)
```

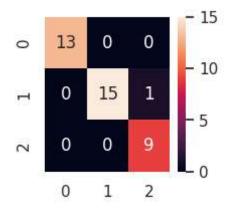
### In [ ]:

### print(report)

	precision	recall	f1-score	support	
Iris-setosa	1.00	1.00	1.00	13	
Iris-versicolor	1.00	0.94	0.97	16	
Iris-virginica	0.90	1.00	0.95	9	
accuracy			0.97	38	
macro avg	0.97	0.98	0.97	38	
weighted avg	0.98	0.97	0.97	38	

### In [ ]:

plot\_confusion\_matrix(iris\_y\_test, iris\_y\_pred)



# Projeto: Titanic Dataset

## Análise e limpeza do Dataset

### **Valores faltantes**

A primeira análise feita é verificar os valores faltantes. **Três colunas possuem valores nulos: Age, Cabin e Embarked.** Cada coluna terá um tratamento apropriado.

### In [9]:

titanic\_df.shape

### Out[9]:

(891, 12)

### In [10]:

```
#titanic_df.info()
titanic_df.isnull().sum()
```

### Out[10]:

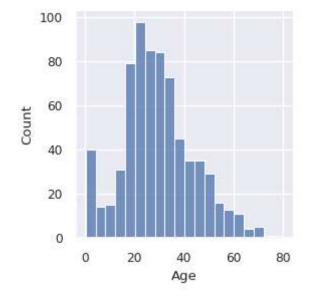
PassengerId	0
Survived	0
Pclass	0
Name	0
Sex	0
Age	177
SibSp	0
Parch	0
Ticket	0
Fare	0
Cabin	687
Embarked	2
dtype: int64	

### Age

- Como é uma coluna de valores numéricos, que a maioria dos valores não é nulo e que acreditamos ter grande relevância para a sobrevivência ou não, será utilizado um método de substituir os valores nulos por um outro valor.
- No plot vemos que se aproxima de uma distribuição assimétrica para a direita, então, então podemos substituir os valores faltantes pela mediana. [1] (https://vitalflux.com/pandas-impute-missing-values-mean-median-mode/#How to decide which imputation technique to use)
- Idealmente, deveríamos analisar a correlação entre a idade e outras features, observando a existência de uma correlação, então os valores nulos deveriam ser substituídos pela mediana do subconjunto (separado pelas outras features correlacionadas).

### In [11]:

```
sns.set(font_scale=0.8)
sns.displot(titanic_df['Age'], height=3)
plt.show()
```



```
In [12]:
```

```
titanic_df['Age'] = titanic_df['Age'].fillna(titanic_df['Age'].median())
```

### Cabin

- A maioria dos valores são nulos (687 de 891), não são numéricos e possuem muitos valores únicos.
- Poderia ser feita uma tentativa de extrair informações relevantes dos dados que existem, visto que as strings consistem de uma letra seguida por um ou dois números, podemos deduzir que está relacionada a posição da cabine no navio, podendo então influenciar na sobrevivência de um passageiro. [2] <a href="https://www.kaggle.com/code/ccastleberry/titanic-cabin-features/notebook">(https://www.kaggle.com/code/ccastleberry/titanic-cabin-features/notebook)</a>
- Entretanto, optaremos por descartar a coluna devido a quantidade de valores faltantes.

### In [13]:

```
titanic_df['Cabin'].value_counts()
Out[13]:
B96 B98
                4
G6
                4
C23 C25 C27
                4
C22 C26
                3
F33
                3
E34
                1
C7
                1
C54
                1
E36
                1
C148
Name: Cabin, Length: 147, dtype: int64
In [14]:
titanic_df = titanic_df.drop(columns=['Cabin'])
```

### **Embarked**

• Como se tratam de valores categóricos, os dois valores faltantes serão subsituídos pela moda.

```
In [15]:
```

```
titanic_df['Embarked'].value_counts()

Out[15]:

S   644
C   168
Q   77
Name: Embarked, dtype: int64
```

### In [16]:

```
titanic_df['Embarked'] = titanic_df['Embarked'].fillna(titanic_df['Embarked'].mode()
[0])
```

### In [17]:

```
titanic_df['Embarked'].value_counts()

Out[17]:

S  646
C  168
O  77
```

### Colunas consideradas irrelevantes

Name: Embarked, dtype: int64

Algumas colunas possuem informações irrelevantes: identificadores que não apresentam relação direta com algo que possa definir a sobrevivência ou são valores aleatórios:

- Passengerld
- Ticket

### In [18]:

```
titanic_df = titanic_df.drop(columns=['PassengerId'])
titanic_df = titanic_df.drop(columns=['Ticket'])
```

### Substituição dos valores categóricos por numéricos

### Name

- Essa coluna é composta apenas de valores únicos, mas todos os indivíduos possuem títulos (Mr., Mrs., Miss., etc).
- É possível extrair a informação do título do passageiro e criar uma nova coluna.

### In [19]:

891

```
titanic_df['Name'].unique().size
Out[19]:
```

```
In [20]:
```

```
titles = titanic_df.Name.str.extract(' ([A-Za-z]+)\.', expand=False)
print(titles.size)
titles.value_counts()
```

#### 891

### Out[20]:

517 Mr Miss 182 125 Mrs 40 Master 7 Dr Rev 6 2 Mlle 2 Major 2 Col Countess 1 Capt 1 Ms 1 1 Sir 1 Lady Mme 1 1 Don Jonkheer 1

Name: Name, dtype: int64

- Como Mr, Miss, Mrs e Master são os únicos títulos que se repetem mais de dez vezes, vamos optar por substituir o restante por 'Other'. E outros títulos que equivalem a esses títulos também serão subsituídos.
- O seguinte mapeamento será feito:
  - Mr = 0, Mrs = 1, Miss = 2, Master = 3, Other = 4

### In [21]:

```
titles = titles.replace(['Lady', 'Countess','Capt', 'Col', 'Don', 'Dr', 'Major', 'Rev',
    'Sir', 'Jonkheer', 'Dona'], 'Other')
titles = titles.replace('Mlle', 'Miss')
titles = titles.replace('Ms', 'Miss')
titles = titles.replace('Mme', 'Mrs')
titles = titles.map({"Mr": 0, "Mrs": 1, "Miss": 2, "Master": 3, "Other": 4})
titles.value_counts()
```

### Out[21]:

```
0 517
2 185
1 126
3 40
4 23
Name: Name, dtype: int64
```

### In [22]:

```
titanic_df = titanic_df.drop(columns=['Name'])
```

```
In [23]:
titanic_df['Title'] = titles
```

#### **Outras colunas**

```
In [26]:
```

```
titanic_df['Sex'] = titanic_df['Sex'].map({'male': 0, 'female': 1})
In [27]:
titanic_df['Embarked'] = titanic_df['Embarked'].map({'S': 0, 'C': 1, 'Q': 2})
```

### Combinar colunas

• Embarked (S = 0, C = 1, Q = 2)

• Podemos combinar as colunas Sibsp e Parch para ter uma coluna representando o tamanho da família.

```
In [28]:
```

```
#family = titanic_df['SibSp'] + titanic_df['Parch']
#titanic_df['Family'] = family
In [29]:
```

#titanic\_df = titanic\_df.drop(columns=['SibSp', 'Parch'])

```
Dataset após limpeza
```

### In [30]:

```
titanic_df.info()

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 891 entries, 0 to 890
Data columns (total 9 columns):
```

```
Column
             Non-Null Count Dtype
   -----
             -----
   Survived 891 non-null
                             int64
0
             891 non-null
1
   Pclass
                             int64
2
   Sex
             891 non-null
                             int64
3
   Age
             891 non-null
                             float64
4
             891 non-null
                             int64
   SibSp
5
   Parch
             891 non-null
                             int64
6
   Fare
             891 non-null
                             float64
7
   Embarked 891 non-null
                             int64
   Title
             891 non-null
                             int64
```

dtypes: float64(2), int64(7)

memory usage: 62.8 KB

### In [31]:

```
titanic_df.head()
```

### Out[31]:

	Survived	Pclass	Sex	Age	SibSp	Parch	Fare	Embarked	Title
0	0	3	0	22.0	1	0	7.2500	0	0
1	1	1	1	38.0	1	0	71.2833	1	1
2	1	3	1	26.0	0	0	7.9250	0	2
3	1	1	1	35.0	1	0	53.1000	0	1
4	0	3	0	35.0	0	0	8.0500	0	0

### \* Criando modelo

### In [32]:

```
titanic_x = titanic_df.drop(columns=['Survived'])
titanic_y = titanic_df['Survived']

titanic_x_train, titanic_x_test, titanic_y_train, titanic_y_test = train_test_split(titanic_x, titanic_y, test_size = 0.2)
```

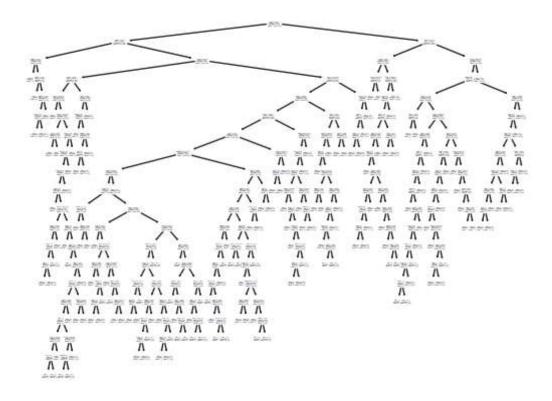
### 1. Classificador com parâmetros default

### In [ ]:

```
classifier1 = DecisionTreeClassifier()
classifier1.fit(titanic_x_train, titanic_y_train)
```

### In [256]:

plot\_tree(classifier1, feature\_names=titanic\_df.drop(columns=['Survived']).columns)
plt.show()



### In [257]:

titanic\_y\_pred = classifier1.predict(titanic\_x\_test)

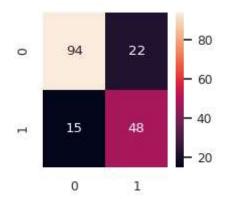
### In [258]:

titanic\_report = classification\_report(titanic\_y\_test, titanic\_y\_pred)
print(titanic\_report)

	precision	recall	f1-score	support
0	0.86	0.81	0.84	116
1	0.69	0.76	0.72	63
accuracy			0.79	179
macro avg	0.77	0.79	0.78	179
weighted avg	0.80	0.79	0.80	179

### In [259]:

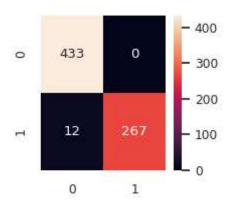
```
plot_confusion_matrix(titanic_y_test, titanic_y_pred)
```



### In [261]:

```
titanic_y_pred = classifier1.predict(titanic_x_train)
titanic_report = classification_report(titanic_y_train, titanic_y_pred)
print(titanic_report)
plot_confusion_matrix(titanic_y_train, titanic_y_pred)
```

	precision	recall	f1-score	support
0	0.97	1.00	0.99	433
1	1.00	0.96	0.98	279
accuracy			0.98	712
macro avg	0.99	0.98	0.98	712
weighted avg	0.98	0.98	0.98	712



### 2. Utilizando GridSearchCV

### In [262]:

```
parameters = {
    'criterion': ["gini", "entropy", "log_loss"],
    'splitter': ["best", "random"],
    'max_depth': [3, 10, 30, None],
    'max_features': ["sqrt", "log2", None],
    'min_samples_split': [2, 16, 32],
    'max_leaf_nodes': [5, 10, 40, None],
    'random_state': [13, 42, 84, None]
}
```

```
In [263]:
```

```
classifier = DecisionTreeClassifier()
```

### In [264]:

```
grid = GridSearchCV(estimator = classifier, param_grid = parameters)
grid.fit(titanic_x_train, titanic_y_train)
```

### Out[264]:

```
► GridSearchCV
► estimator: DecisionTreeClassifier
► DecisionTreeClassifier
```

### In [265]:

```
print(grid.best_score_)
print(grid.best_params_)
```

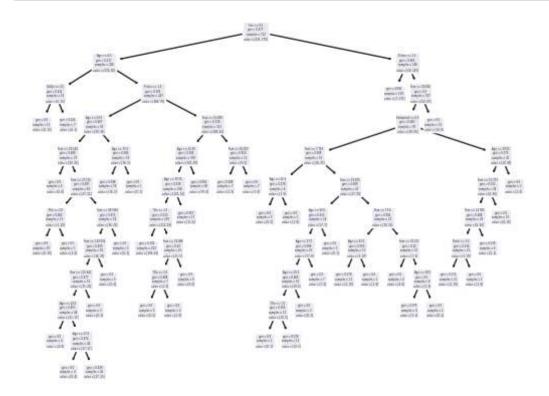
### 0.8343346794051021

```
{'criterion': 'gini', 'max_depth': 30, 'max_features': None, 'max_leaf_nod es': 40, 'min_samples_split': 2, 'random_state': None, 'splitter': 'best'}
```

### In [ ]:

### In [283]:

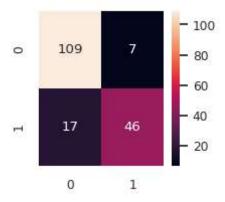
plot\_tree(classifier2, feature\_names=titanic\_df.drop(columns=['Survived']).columns)
plt.show()



### In [288]:

```
titanic_y_pred = classifier2.predict(titanic_x_test)
titanic_report = classification_report(titanic_y_test, titanic_y_pred)
print(titanic_report)
plot_confusion_matrix(titanic_y_test, titanic_y_pred)
```

	precision	recall	f1-score	support
0	0.87	0.94	0.90	116
1	0.87	0.73	0.79	63
accuracy			0.87	179
macro avg	0.87	0.83	0.85	179
weighted avg	0.87	0.87	0.86	<b>1</b> 79

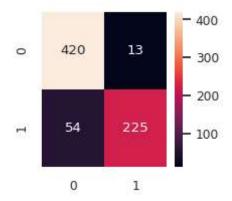


### Checando overfitting:

### In [287]:

```
titanic_y_pred = classifier2.predict(titanic_x_train)
titanic_report = classification_report(titanic_y_train, titanic_y_pred)
print(titanic_report)
plot_confusion_matrix(titanic_y_train, titanic_y_pred)
```

	precision	recall	f1-score	support
0	0.89	0.97	0.93	433
1	0.95	0.81	0.87	279
accuracy			0.91	712
macro avg	0.92	0.89	0.90	712
weighted avg	0.91	0.91	0.90	712



### 3. Tentando obter uma árvore menor com score similar

### In [272]:

```
parameters = {
    'criterion': ["gini", "entropy", "log_loss"],
    'splitter': ["best", "random"],
    'max_depth': [2, 4, 6],
    'max_features': ["sqrt", "log2", None],
    'min_samples_split': [2, 4, 8],
    'max_leaf_nodes': [4, 7, 10],
    'random_state': [13, 42, 84, None]
}
```

### In [273]:

```
classifier = DecisionTreeClassifier()
```

### In [ ]:

```
grid = GridSearchCV(estimator = classifier, param_grid = parameters)
grid.fit(titanic_x_train, titanic_y_train)
```

### In [275]:

```
print(grid.best_score_)
print(grid.best_params_)
```

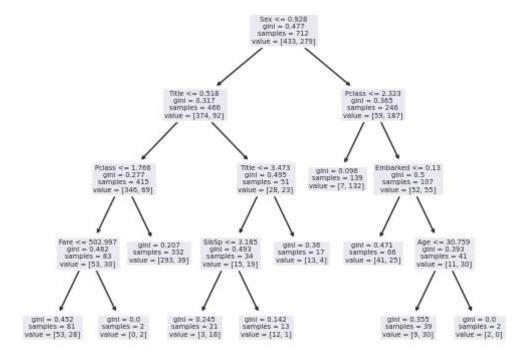
#### 0.8329065300896286

```
{'criterion': 'gini', 'max_depth': 4, 'max_features': None, 'max_leaf_node
s': 10, 'min_samples_split': 2, 'random_state': 13, 'splitter': 'random'}
```

### In [ ]:

### In [277]:

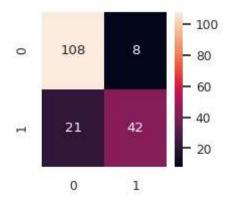
plot\_tree(classifier3, feature\_names=titanic\_df.drop(columns=['Survived']).columns)
plt.show()



### In [294]:

```
titanic_y_pred = classifier3.predict(titanic_x_test)
titanic_report = classification_report(titanic_y_test, titanic_y_pred)
print(titanic_report)
plot_confusion_matrix(titanic_y_test, titanic_y_pred)
```

	precision	recall	f1-score	support
0	0.84	0.93	0.88	116
1	0.84	0.67	0.74	63
accuracy			0.84	179
macro avg	0.84	0.80	0.81	179
weighted avg	0.84	0.84	0.83	179

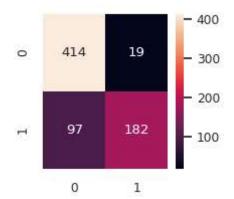


### Verificando overfitting

### In [281]:

```
titanic_y_pred = classifier3.predict(titanic_x_train)
titanic_report = classification_report(titanic_y_train, titanic_y_pred)
print(titanic_report)
plot_confusion_matrix(titanic_y_train, titanic_y_pred)
```

support	f1-score	recall	precision	
433	0.88	0.96	0.81	0
279	0.76	0.65	0.91	1
712	0.84			accuracy
712	0.82	0.80	0.86	macro avg
712	0.83	0.84	0.85	weighted avg



## **W** Utilizando Random Forests

Aplicando o RandomForestClassifier com a configuração default no problema Titanic.

### In [ ]:

```
classifier4 = RandomForestClassifier()
classifier4.fit(titanic_x_train, titanic_y_train)
```

### In [129]:

```
titanic_y_pred = classifier4.predict(titanic_x_test)
```

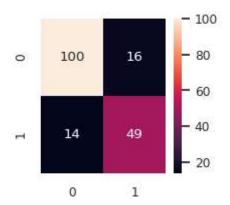
### In [130]:

```
titanic_report = classification_report(titanic_y_test, titanic_y_pred)
print(titanic_report)
```

	precision	recall	f1-score	support
0	0.88	0.86	0.87	116
1	0.75	0.78	0.77	63
accuracy			0.83	179
macro avg	0.82	0.82	0.82	179
weighted avg	0.83	0.83	0.83	179

### In [131]:

```
plot_confusion_matrix(titanic_y_test, titanic_y_pred)
```



## 🐣 Utilizando AD no problema de Detecção de Lombalgia

### In [290]:

```
pain_x = pain_df.drop(columns=['Class_att'])
pain_y = pain_df['Class_att']

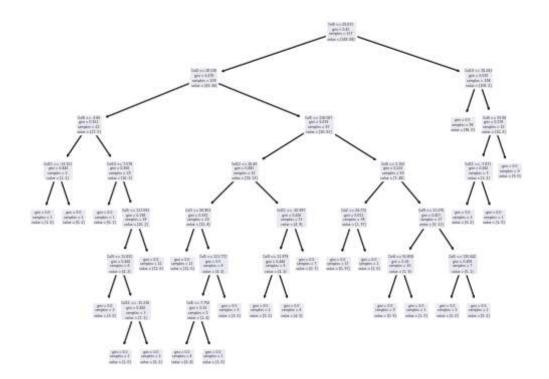
pain_x_train, pain_x_test, pain_y_train, pain_y_test = train_test_split(pain_x, pain_y, test_size = 0.3)
```

### In [ ]:

```
classifier5 = DecisionTreeClassifier()
classifier5.fit(pain_x_train, pain_y_train)
```

### In [292]:

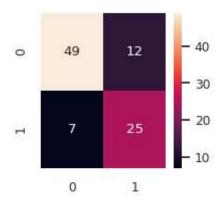
```
plot_tree(classifier5, feature_names=pain_df.drop(columns=['Class_att']).columns)
plt.show()
```



### In [293]:

```
pain_y_pred = classifier5.predict(pain_x_test)
pain_report = classification_report(pain_y_test, pain_y_pred)
print(pain_report)
plot_confusion_matrix(pain_y_test, pain_y_pred)
```

	precision	recall	f1-score	support
Abnormal Normal	0.88 0.68	0.80 0.78	0.84 0.72	61 32
accuracy macro avg weighted avg	0.78 0.81	0.79 0.80	0.80 0.78 0.80	93 93 93





### **Titanic**

### Árvore com configuração padrão:

- Acurácia na base de teste: 0.79
- Acurácia na base de treino: 0.98 (overfitting!)

### Árvore com configuração 1:

```
{'criterion': 'gini', 'max_depth': 30, 'max_features': None, 'max_leaf_nodes': 4
0, 'min_samples_split': 2, 'random_state': None, 'splitter': 'best'}
```

- Maior score no grid: 0.83
- Acurácia na base de teste: 0.87
- · Acurácia na base de treino: 0.91

### Árvore com configuração 2:

```
{'criterion': 'gini', 'max_depth': 4, 'max_features': None, 'max_leaf_nodes': 1
0, 'min_samples_split': 2, 'random_state': 13, 'splitter': 'random'}
```

- Maior score no grid: 0.83
- · Acurácia na base de teste: 0.84
- · Acurácia na base de treino: 0.84

### Random Forest com configuração padrão

Acurácia: 0.83

### Detecção de Lombalgia

- Acurácia com MLP: 0.82
- Acurácia com AD (configuração padrão): 0.80