Creado por:

Isabel Maniega

Decission Tree Classification

```
In [2]: import pandas as pd
         from sklearn.tree import DecisionTreeClassifier
         from sklearn.model_selection import train_test_split
         from sklearn import metrics
 In [3]: # Dataset:
         # https://www.kaggle.com/datasets/uciml/pima-indians-diabetes-database?resource=download
 In [5]: col_names = ["pregnant", "glucose", "bp", "skin", "insulin", "bmi", "pedigree", "age", "label"]
         pima = pd.read csv("diabetes.csv", header=None, names=col names)
 In [6]:
         pima.head()
 Out[6]:
             pregnant glucose
                                     bp
                                               skin insulin bmi
                                                                          pedigree age
                                                                                         label
                                                   Insulin BMI DiabetesPedigreeFunction Age Outcome
         0 Pregnancies
                     Glucose BloodPressure SkinThickness
         1
                   6
                         148
                                     72
                                                35
                                                       0 33.6
                                                                             0.627
                                                                                   50
                                                                                            1
         2
                   1
                         85
                                     66
                                                29
                                                       0 26.6
                                                                             0.351
                                                                                   31
                                                                                           0
         3
                         183
                                     64
                                                       0 23.3
                                                                             0.672
                                                                                   32
                                                                                           1
         4
                   1
                         89
                                     66
                                                23
                                                       94 28.1
                                                                                           0
                                                                             0.167
                                                                                   21
         pima.drop(index=pima.index[0], axis=0, inplace=True)
 In [8]:
         pima.head()
 Out[8]:
           pregnant glucose bp skin insulin bmi pedigree age label
         1
                                       0 33.6
                 6
                       148 72
                                35
                                                0.627
                                                      50
                                                             1
         2
                       85 66
                                                0.351
                                29
                                       0 26.6
                                                      31
                                                             0
                 8
                       183 64
                                       0 23.3
         3
                                0
                                                0.672
                                                      32
                                                             1
         4
                 1
                       89 66
                                23
                                      94 28.1
                                                0.167
                                                      21
                                                             0
         5
                       137 40
                                35
                                     168 43.1
                                                2.288
                                                      33
                                                             1
 In [9]: feature_cols = ["pregnant", "glucose", "bp", "insulin", "bmi", "pedigree", "age"]
         X = pima[feature cols]
         y = pima.label
In [10]: X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=1)
In [11]: # Creamos el clasificador
         clf = DecisionTreeClassifier()
         clf = clf.fit(X_train, y_train)
         y_pred = clf.predict(X_test)
         y_pred
'1',
                '1', '0', '0', '0', '1', '0', '0', '1', '0',
                                                            '1', '0',
                                                       '1',
                                                            '1',
                                                                 '0',
                                                                      '1',
                '0', '1', '0', '0', '0', '1',
                                             '0', '1',
                                                  '0',
                                                       '0',
                                                            '0',
                                                                           '1',
                '1', '0', '0', '0', '0', '0', '0',
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                    '0', '1', '0', '1', '0', '0', '1',
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                     '0',
                         '0',
                                   'Θ',
                                        '0',
                                             '0',
                                                  'Θ',
                                   '1',
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                    '0', '0', '0',
                                             'Θ',
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                    '0', '0', '0', '0',
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                                                  '1',
                                                       ' 0 '
                                                                 '0'
                     '0', '0', '0', '0',
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                              '0',
                                   '0',
                         '0',
                    ' 0 '
                                        '1',
                                             ' 0 '
                                                  '0',
                                                       ' 0 '
                                                            '0'
                                                                 ' 0 '
                                                                      ' 0 '
                                                  '1',
                                                       '0',
                '0', '1', '0', '0', '1', '1', '1',
                                                            '1', '1',
                                                                      '0',
                '0', '0', '1', '0', '0', '0', '0', '1', '0'], dtype=object)
In [12]: y_test
Out[12]: 286
         102
         582
                0
         353
                0
         727
         242
                0
         600
                0
         651
                0
         12
                1
         215
                1
         Name: label, Length: 231, dtype: object
In [13]: print("Accuracy:", metrics.accuracy_score(y_test, y_pred))
         Accuracy: 0.6796536796536796
In [14]: # Otra opción de clasificador
         clf = DecisionTreeClassifier(criterion="entropy", max_depth=3)
         clf = clf.fit(X_train, y_train)
         y_pred = clf.predict(X_test)
         print("Accuracy:", metrics.accuracy_score(y_test, y_pred))
         Accuracy: 0.7705627705627706
```

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