adaboost

September 7, 2018

0.1 Modelo de predición de cancer de mama (Wisconsin uci)

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In [1]: import pandas as pd
       from sklearn.cross_validation import train_test_split
       from sklearn.preprocessing import StandardScaler
       from sklearn.tree import DecisionTreeClassifier
       from sklearn.ensemble import AdaBoostClassifier
       from sklearn.pipeline import Pipeline
/home/rafa/anaconda3/lib/python3.6/site-packages/sklearn/cross_validation.py:41: DeprecationWarn
  "This module will be removed in 0.20.", DeprecationWarning)
In [2]: # Importamos los datos desde el repositorio oficial
       df = pd.read_csv('https://archive.ics.uci.edu/ml/machine-learning-databases/breast-cance
       header=None)
In [3]: # Vemos la estructura del dataframe
        # Columna O --> ID
        # Columna 1 --> Clase
        # Columnas 2.. -> Datos
       df[:2]
Out [3]:
                               3
                                              5
       0 842302 M 17.99
                           10.38 122.8 1001.0 0.11840 0.27760
                                                                   0.3001 0.14710
        1 842517 M 20.57 17.77 132.9 1326.0 0.08474 0.07864 0.0869
                      22
                             23
                                            25
                                                    26
                                                            27
                                                                    28
                                    24
                                                                            29
       0
                   25.38 17.33 184.6 2019.0 0.1622 0.6656 0.7119 0.2654
                   24.99 23.41 158.8 1956.0 0.1238 0.1866 0.2416 0.1860
               30
                       31
       0 0.4601
                  0.11890
        1 0.2750 0.08902
        [2 rows x 32 columns]
In [4]: # Separamos el dataset en (Conjunto de datos, Etiquetas)
       X = df.loc[:, 2:].values
       y = df.loc[:, 1].values
```

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In [5]: # Separamos el conjunto total en dos subconjuntos (Train 80% / Test 20%)
        X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.20)
In [6]: weak_clf = DecisionTreeClassifier(criterion='entropy', max_depth=None, random_state=0)
In [7]: # Creamos un flujo de trabajo (pipeline):
        # Paso 1 -> Escalar los datos
        # Paso 2 -> Analisis de componentes principales (Reduce la dimensionalidad)
        # Paso 3 -> Construimos un modelo utilizando Adaboost
       tree = DecisionTreeClassifier(random_state=0)
        adaboost = AdaBoostClassifier(base_estimator=tree,
                                      n_estimators=100,
                                      learning_rate=0.01,
                                      random_state=0)
       pipe_lr = Pipeline([('scl', StandardScaler()),
                            ('clf', adaboost)])
        # Ajustamos el modelo a los datos de entrenamiento
        pipe_lr.fit(X_train, y_train)
        #Calculamos la precisión del modelo utilizando los datos de test
        print('Test Accuracy: %.3f' % pipe_lr.score(X_test, y_test))
Test Accuracy: 0.904
```