

TÍTULO

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1 import numpy as np
2 from scipy import stats
3
4 # names of variables
5 labels = ['age', 'leptin', 'bmi', 'adiponectin', 'glucose',
6           'resistin', 'insulin', 'MCP1', 'HOMA']
7
8 # loads data
9 data = np.loadtxt (open (r'../..//data.csv', 'rb'), delimiter = ',')
10
11 # rewrites data as all the rows of data w/out nan cells
12 data = data [~np.isnan (data).any (axis=1)]
13
14 # separates parameters into matrix x
15 x = np.array ([list (data [x][: -1]) for x in range (len (data))])
16
17 # and class (1, 2) into vector y
18 y = np.array ([int (data [x][ -1]) for x in range (len (data))])
19
20 # removes outliers
21 data_no = data [(np.abs (stats.zscore (data)) < 3).all (axis = 1)]
22 # ↑ = No Outliers
23
24 x_no = np.array ([list (data_no [x][: -1]) for x in range (len (data_no))])
25 y_no = np.array ([int (data_no [x][ -1]) for x in range (len (data_no))])
```

Listing 1: Importaciones iniciales y preparacion de datos.

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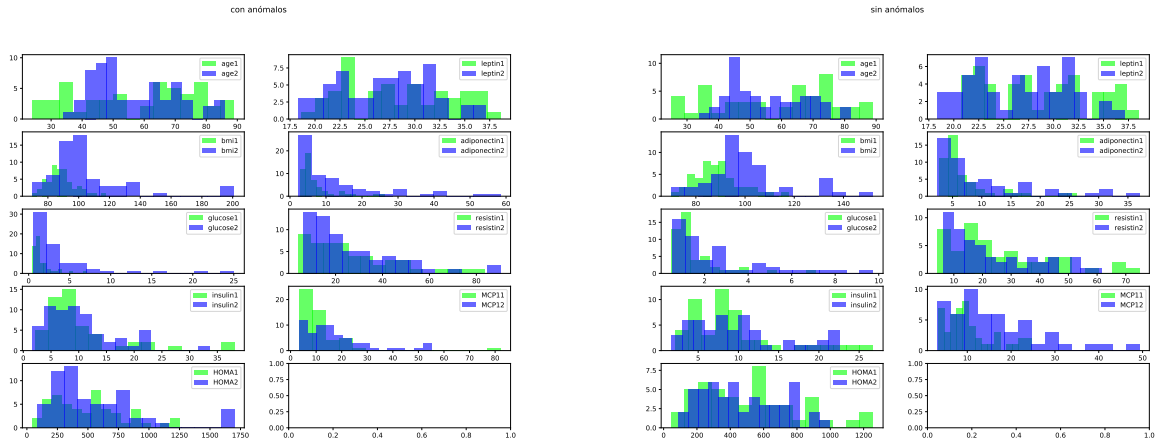


Fig. 1: Histogramas para datos con y sin anomalías.