

Programming

Winter 2020/2021

Number 07, Submission Deadline: Jan. 12, 2021

1. The diabetes data set (7P)

Scikit-Learn provides the following diabetes data set that has been published by

Bradley Efron, Trevor Hastie, Iain Johnstone and Robert Tibshirani (2004) "Least Angle Regression," *Annals of Statistics* (with discussion), 407-499.

The authors describe the data set as follows:

Ten baseline variables, age, sex, body mass index, average blood pressure, and six blood serum measurements were obtained for each of $n = 442$ diabetes patients, as well as the response of interest, a quantitative measure of disease progression one year after baseline.

1. Inform yourself about Lasso regression. Briefly describe its key properties (2P)
2. Perform a regression analysis of the diabetes data set with Lasso. Split the dataset in a train (70%) and a test set (30%). (1P)
3. Describe and visualize your results (2P)
4. Compare your results with the test-dataset. (2P)

```
[22]: from sklearn.datasets import load_diabetes
import pandas as pd

diabetes = load_diabetes()
data = pd.DataFrame(diabetes.data, columns=diabetes.feature_names)

data.head()
```

```
[22]:      age      sex      bmi      bp      s1      s2      s3  \
0  0.038076  0.050680  0.061696  0.021872 -0.044223 -0.034821 -0.043401
1 -0.001882 -0.044642 -0.051474 -0.026328 -0.008449 -0.019163  0.074412
2  0.085299  0.050680  0.044451 -0.005671 -0.045599 -0.034194 -0.032356
3 -0.089063 -0.044642 -0.011595 -0.036656  0.012191  0.024991 -0.036038
```

```

4  0.005383 -0.044642 -0.036385  0.021872  0.003935  0.015596  0.008142

           s4           s5           s6
0 -0.002592  0.019908 -0.017646
1 -0.039493 -0.068330 -0.092204
2 -0.002592  0.002864 -0.025930
3  0.034309  0.022692 -0.009362
4 -0.002592 -0.031991 -0.046641

```

2. The breast cancer Wisconsin diagnostic data set (8P)

Another data set that Scikit-Learn provides is the *breast cancer Wisconsin diagnostic data set* that was first published by

W.N. Street, W.H. Wolberg and O.L. Mangasarian. Nuclear feature extraction for breast tumor diagnosis. IS&T/SPIE 1993 International Symposium on Electronic Imaging: Science and Technology, volume 1905, pages 861-870, San Jose, CA, 1993.

The data set comprises data of 569 patients and consists of features that are computed from a digitized image of a fine needle aspirate (FNA) of a breast mass. They describe characteristics of the cell nuclei present in the image.

```

[31]: from sklearn.datasets import load_breast_cancer

breast_cancer = load_breast_cancer()
data = pd.DataFrame(breast_cancer.data, columns=breast_cancer.
↳feature_names)

data.head()

```

```

[31]:   mean radius  mean texture  mean perimeter  mean area  mean smoothness  \
↳\
0      17.99         10.38         122.80         1001.0         0.11840
1      20.57         17.77         132.90         1326.0         0.08474
2      19.69         21.25         130.00         1203.0         0.10960
3      11.42         20.38          77.58          386.1         0.14250
4      20.29         14.34         135.10         1297.0         0.10030

      mean compactness  mean concavity  mean concave points  mean symmetry  \
0          0.27760         0.3001         0.14710         0.2419
1          0.07864         0.0869         0.07017         0.1812

```

2	0.15990	0.1974	0.12790	0.2069
3	0.28390	0.2414	0.10520	0.2597
4	0.13280	0.1980	0.10430	0.1809

	mean fractal dimension	...	worst radius	worst texture	worst _□
↪perimeter \					
0	0.07871	...	25.38	17.33	184.
↪60					
1	0.05667	...	24.99	23.41	158.
↪80					
2	0.05999	...	23.57	25.53	152.
↪50					
3	0.09744	...	14.91	26.50	98.
↪87					
4	0.05883	...	22.54	16.67	152.
↪20					

	worst area	worst smoothness	worst compactness	worst concavity \
0	2019.0	0.1622	0.6656	0.7119
1	1956.0	0.1238	0.1866	0.2416
2	1709.0	0.1444	0.4245	0.4504
3	567.7	0.2098	0.8663	0.6869
4	1575.0	0.1374	0.2050	0.4000

	worst concave points	worst symmetry	worst fractal dimension
0	0.2654	0.4601	0.11890
1	0.1860	0.2750	0.08902
2	0.2430	0.3613	0.08758
3	0.2575	0.6638	0.17300
4	0.1625	0.2364	0.07678

[5 rows x 30 columns]

1. Inform yourself about decision tree classification with Scikit-Learn. Briefly describe the classification algorithms that Scikit-Learn provides (2P)
2. Perform a classification analysis of the breast cancer data set. In doing so,
 1. Use cross validation in your analysis; justify your choice(s) of the number of partitions (1P)
 2. Run the analysis for all decision tree algorithms that Scikit-Learn provides (2P)
 3. Evaluate the classification quality of the algorithms with your (justified) choice

- of metric (2P)
4. Visualize your results. (1P)