

# Machine Learning for Medicine TP 4

## Feature Selection Model Selection

The goal of the TME is to learn various techniques of feature selection.

**Data** (both data sets are provided)

- Molecular classification of leukemia data set of *Golub et al. 1999* contains gene expressions of 72 patients and 3562 genes.
- Breast cancer data set

You will need to load at least the following packages:

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from sklearn.linear_model import ElasticNet
from sklearn.svm import LinearSVC
from sklearn.feature_selection import SelectFromModel
from sklearn import linear_model
```

### Analysis

Repeat the same analyses for the two data sets.

To read the data:

- For the *Golub et al. 1999* data

```
X = pd.read_csv('data/Golub_X', sep=' ') # Observations
y = pd.read_csv('data/Golub_y', sep=' ') # Classes
```

- For the Breast cancer data

```
X = pd.read_csv('data/Breast.txt', sep=' ')
y = X.as_matrix()[:, 30] # Classes
X = X.as_matrix()[:, 0:29] # Observations
```

We will use the `sklearn` Python library only.

1. A simple heuristic approach is to delete features whose variance is less than a threshold. Try it (with two different arbitrary thresholds) but do not expect this method to return an optimal performance (although it can be quite efficient on some data sets).

[http://scikit-learn.org/stable/modules/feature\\_selection.html](http://scikit-learn.org/stable/modules/feature_selection.html)

2. Univariate feature selection with statistical tests to get rid of features which are not statistically significant with respect to the vector of class. Try the `SelectFdr` function that computes p-values for an estimated false discovery rate.

[http://scikit-learn.org/stable/modules/generated/sklearn.feature\\_selection.SelectFdr.html#sklearn.feature\\_selection.SelectFdr](http://scikit-learn.org/stable/modules/generated/sklearn.feature_selection.SelectFdr.html#sklearn.feature_selection.SelectFdr)

3.  $L_1$ -based feature selection is designed to find an optimal solution. The sparsity parameter is important (since it controls the number of non-zero parameters: if too many parameters are kept, no really feature selection; if too few parameters are chosen, it is possible that the accuracy is very poor).

- (a) Logistic regression penalized by the  $L_1$  penalty term

```
linear_model.Lasso(alpha=alpha)
```

- (b) A support vector machine penalized by the  $L_1$  penalty term

```
LinearSVC(C=C, penalty="l1", dual=False)
```

- (c) Explore the Elastic Net which is a compromise between the  $L_1$  and  $L_2$  penalty terms.

```
ElasticNet(alpha=alpha, l1_ratio=0.7)
```

4. How many features do you keep using these different methods? It is quite normal that each method selects a different number of features.
5. What method leads to the best performance (on the given data sets) ?

## References

- *The original Lasso paper:*

Tibshirani, R. (1996). Regression shrinkage and selection via the lasso. J. Royal. Statist. Soc B., Vol. 58, No. 1, pages 267–288

<http://statweb.stanford.edu/~tibs/lasso/lasso.pdf>

- T. Hastie, R. Tibshirani, and M. Wainwright. *Statistical Learning with Sparsity. The Lasso and Generalizations*. (a good book)

[https://web.stanford.edu/~hastie/StatLearnSparsity\\_files/SLS\\_corrected\\_1.4.16.pdf](https://web.stanford.edu/~hastie/StatLearnSparsity_files/SLS_corrected_1.4.16.pdf)