# SPLEX TME 1

# Exploratory Analysis with Significance Tests. Multiple Hypothesis Testing. Handling Missing Data. AdaBoost

### Data

We explore two data sets downloadable from the Machine Learning Repository (http://archive.ics.uci.edu/ml/index.php)

- Breast Cancer Wisconsin (Diagnostic) Data Set (https://archive.ics.uci.edu/ml/datasets/Breast+Cancer+Wisconsin+(Diagnostic))
- Mice Protein Expression Data Set (https://archive.ics.uci.edu/ml/datasets/Mice+Protein+ Expression)

## **Exploratory Analysis**

We will perform an exploratory analysis with Python. You can use the Spyder environment which is already installed on the machines.

You will need to load the following packages:

```
import numpy as np
import pandas as pd
import scipy.stats as stats
import matplotlib.pyplot as plt
import statsmodels.sandbox.stats.multicomp as sm
```

- 1. Load the Breast Cancer data set using pd.read\_table() (for the Mice Data Set you will need pd.ExcelFile())
- 2. Some data sets have missing data. You can impute them by replacing the missing values by median values with fillna(data, inplace=True) (you will need to impute data in the Mice Data Set)
- 3. Both the Mice data set and the Breast Cancer are binary classification tasks (M and B are two classes in the Breast Cancer, and Ts65Dn and Control for the Mice)
- 4. Find the correlation coefficients between variables with stats.pearsonr(). Are there a lot of variables which are strongly correlated? What is the meaning of the sign of the correlation coefficient?
- 5. Run the Wilcoxon test (if you have two classes or Kruskal-Wallis test if you have more than two classes) stats.wilcoxon() to find variables which are significant to discriminate two classes.
- 6. We perform a multiple hypothesis testing (since we have a lot of variables), and we need to adjust the p-values. You can adjust the p-values with sm.multipletests()

Consider different adjustment methods:

http://jpktd.blogspot.fr/2013/04/multiple-testing-p-value-corrections-in.html What is the most and the least stringent methods of adjustment? Can you explain why?

- 7. Compare the distributions of variables in two classes with stats.ttest\_ind()
- 8. Boxplot the distributions of the variables of the observations from class 1 and from class 2 using boxplot(). Are the plotted distributions coherent with the results obtained by stats.ttest\_ind()?

9. What are your conclusions? What variables are the most significant? What significance threshold (significance level) should be chosen? What multiple hypothesis adjustment method would you use? Why?

# AdaBoost

- 1. Test the AdaBoost classifier
  - http://scikit-learn.org/stable/modules/generated/sklearn.ensemble.AdaBoostClassifier.html
- 2. Test various number of weak lerners
- 3. Run the AdaBoost on a 2-dimensional problem and analyse the class boundary http://scikit-learn.org/stable/auto\_examples/ensemble/plot\_adaboost\_twoclass.html# sphx-glr-auto-examples-ensemble-plot-adaboost-twoclass-py
- 4. Is AdaBoost a linear or a non-linear classifier?