# TP No 1 : Support Vector Machine (SVM)

## Useful links for this lab:

- \*\*\* http://scikit-learn.org/stable/modules/svm.html
- \*\* http://en.wikipedia.org/wiki/Support\_vector\_machine
- \*\* http://fr.wikipedia.org/wiki/Machine\_%C3%A0\_vecteurs\_de\_support

## - Introduction and Mathematical Foundations -

The SVMs were introduced by Vapnik [3], and are discussed in Chapter 12 of the book [1]. More mathematical details can be be found in Chapter 7 of the book [2]. The popularity of SVM methods, for binary classification in particular, comes from the fact that they rely on the application of algorithms of linear decision rules: we speak of hyperplanes (affine) separators. However, this search takes place in a higher-dimensional *feature space*, that is the image of the original input space transformed by a non-linear transformation  $\Phi$ .

The aim of this TP is to put into practice this type of classification techniques on actual and simulated data in the scikit-learn package (which implements the library in C LIBSVM) and to learn to control the parameters guaranteeing their flexibility (hyper-parameters, kernel).

#### **Definitions and Notations**

It is recalled that within the framework of the supervised binary classification we use the notations:

- $\mathcal{Y}$  The set of labels, commonly  $\mathcal{Y} = \{-1, 1\}$  in the case of binary classification,
- $\mathbf{x} = (x_1, \dots, x_p) \in \mathcal{X} \subset \mathbb{R}^p$  is an observation (or an example),
- $\mathcal{D}_n = \{(\mathbf{x}_i, y_i), i = 1, \dots n\}$  a learning set containing n examples and their labels,
- There is a probabilistic model that governs the generation of our observations according to random variables X and  $Y: \forall i \in \{1, ..., n\}, (\mathbf{x}_i, y_i) \overset{i.i.d}{\sim} (X, Y)$ .
- We try to build from the learning set  $\mathcal{D}_n$  a function  $\hat{f}: \mathcal{X} \mapsto \{-1,1\}$  which for an unknown point  $\mathbf{x}$  (*i.e.*, which is not present in the learning set) predicts its label:  $\hat{f}(\mathbf{x})$ . The rule considered here is called linear, in the sense that the space is separated by a hyperplane (affine): according to the position with respect thereto, then predicting +1 or -1.

## SVM and kernels for binary classification

SVM (nonlinear) techniques use an implicit function  $\Phi$  transforming the input space  $\mathcal{X} \subset \mathbb{R}^p$  in a Hilbert space  $(\mathcal{H}, \langle \cdot, \cdot \rangle)$  of higher dimension. Learning is then carried out using the model  $(\Phi(X), Y)$  in the space  $\mathcal{H}$ , of larger size of course, but in which it is hoped that the data is "more linearly separable". From a practical point of view, It should be noted that the calculation of projections  $\Phi(X)$  is not used in the method, only scalar products  $\langle \Phi(\mathbf{x}), \Phi(\mathbf{x}') \rangle$ ,  $(\mathbf{x}, \mathbf{x}') \in \mathcal{X}^2$ , are required. Now these are given by a kernel K, via the relation  $kernel\ trick$ :

$$K(\mathbf{x}, \mathbf{x}') = \langle \Phi(\mathbf{x}), \Phi(\mathbf{x}') \rangle.$$

The method therefore requires selecting a kernel (as well as other parameters). Among the possible choices, there are in particular the following kernels

- Linear Kernel:  $K(\mathbf{x}, \mathbf{x}') = \langle \mathbf{x}, \mathbf{x}' \rangle$  (corresponding to linear SVMs)
- Gaussian Kernel (Gaussian RBF)  $K(\mathbf{x}, \mathbf{x}') = \exp(-\gamma ||\mathbf{x} \mathbf{x}'||^2)$

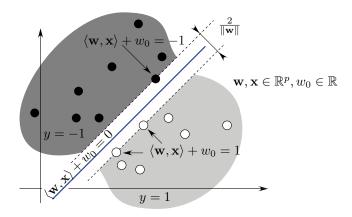


FIGURE 1 – Margin and separator hyperplane in separable classes (case of a linear kernel)

- Polynomial Kernel  $K(\mathbf{x}, \mathbf{x}') = (\alpha + \beta \langle \mathbf{x}, \mathbf{x}' \rangle)^{\delta}$  for a  $\delta > 0$
- Laplacian Kernel (Laplace RBF)  $K(\mathbf{x}, \mathbf{x}') = \exp(-\gamma ||\mathbf{x} \mathbf{x}'||)$  for a  $\gamma > 0$
- Hyperbolic tangent kernel (Sigmoid)  $K(\mathbf{x}, \mathbf{x}') = \tanh(\alpha + \beta \langle \mathbf{x}, \mathbf{x}' \rangle)$  for a pair  $(\alpha, \beta) \in \mathbb{R}^2$

A SVM classifier is of the form

$$\hat{f}_{\mathbf{w},w_0}(\mathbf{x}) = \operatorname{sign}\left(\langle \mathbf{w}, \Phi(\mathbf{x}) \rangle + w_0\right),\tag{1}$$

or  $\mathbf{w} \in \mathcal{H}$  and  $w_0 \in \mathbb{R}$  are parameters adjusted during phase learning from a sample of i.i.d. examples,  $\mathcal{D}_n = \{(\mathbf{x}_i, y_i), i = 1, \dots n\}$ . Note: in the case where the function  $\Phi$  is the identity on  $\mathbb{R}^p$ , we simply find that the decision rule is

$$\hat{f}_{\mathbf{w},w_0}(\mathbf{x}) = \operatorname{sign}\left(\sum_{i=1}^p w_i x_i + w_0\right).$$

The boundary associated with the decision rule (1) is the set  $\{\mathbf{x} : \langle \mathbf{w}, \Phi(\mathbf{x}) \rangle + w_0 = 0\}$ . It corresponds to a hyperplane in space  $\mathcal{H}$ , but is much more complex in  $\mathcal{X}$  (depending on the shape of the selected kernel).

In  $\mathcal{H}$ , the hyperplane is obtained by maximizing the margin separating the two classes. This requires to solve an optimization problem under linear constraints:

$$\begin{cases}
(\mathbf{w}^*, w_0^*, \xi^* \in \mathbb{R}^n) \in \underset{\mathbf{w} \in \mathcal{H}, w_0 \in \mathbb{R}, \xi \in \mathbb{R}^n}{\operatorname{arg \, min}} \left( \frac{1}{2} ||\mathbf{w}||^2 + C \sum_{i=1}^n \xi_i \right) \\
\text{s.t.} \quad \xi_i \ge 0, \qquad \forall i \in \{1, \dots, n\}, \\
y_i \left( \langle \mathbf{w}, \Phi(\mathbf{x}_i) + w_0 \right) \ge 1 - \xi_i, \qquad \forall i \in \{1, \dots, n\}.
\end{cases}$$
(2)

It can be shown that the solution  $\mathbf{w}$  can be expressed in the following way:

$$\mathbf{w}^* = \sum_{i=1}^n \alpha_i^* y_i \Phi(\mathbf{x}_i).$$

The indices i for which  $\alpha_i^* \neq 0$  are those for whom equality is carried out in the constraint, the corresponding points  $\mathbf{x}_i$  are called *support vectors* (vectours supports en français) of the decision. The coefficients  $\alpha_i^*$  denote the solutions of the dual problem which is a quadratic program (QP) under affine constraints:

$$\begin{cases}
\alpha^* \in \arg\max_{\alpha \in \mathbb{R}^n} \left( \sum_{i=1}^n \alpha_i - \frac{1}{2} \sum_{1 \le i, j \le n} \alpha_i \alpha_j y_i y_j K(\mathbf{x}_i, \mathbf{x}_j) \right) \\
\text{s.t.} \quad 0 \le \alpha_i \le C, \quad \forall i \in \{1, \dots, n\}, \\
\sum_{i=1}^n \alpha_i y_i = 0.
\end{cases} \tag{3}$$

The parameter C controls the complexity of the classifier insofar as it determines the cost of a misclassification: the higher the C value, the more the rule obtained is complex (the number of points for which one wants to minimize the classification error crop). This approach is called C-classification and is used with the object sklearn.svm.SVC in the module scikit-learn.

Another way to control complexity (*i.e.*, the number of support vectors), called  $\nu$ -classification, is to consider, in place of the dual problem described above, the following problem:

$$\begin{cases}
\alpha^* \in \underset{\alpha \in \mathbb{R}^n}{\min} \left( \frac{1}{2} \sum_{1 \le i, j \le n} \alpha_i \alpha_j y_i y_j K(\mathbf{x}_i, \mathbf{x}_j) \right) \\
\text{s.c.} \quad 0 \le \alpha_i \le 1, \quad \forall i \in \{1, \dots, n\}, \\
\sum_{i=1}^n \alpha_i y_i = 0, \\
\sum_{i=1}^n \alpha_i \ge \nu.
\end{cases} \tag{4}$$

where  $\nu \in [0,1]$  is a parameter approximating the percentage of support vectors among the training data. This approach is used with the object sklearn.svm.NuSVC in the module scikit-learn.

#### Extensions to multi-class case

In case the output variable Y has more than two modalities, there are several ways to extend the methods of the binary case directly.

"One Vs One". In the case where it is desired to predict a label that has  $K \geq 3$  classes, we can consider all the pairs of labels (k,l) possible,  $1 \leq k < l \leq K$  (there are K(K-1)/2) and learning a classifier  $C_{k,l}(X)$  for each of them. The prediction then corresponds to the label that won the most "duels".

"One Vs. All". For each class k, we learn a classifier to discriminate between populations Y = k and  $Y \neq k$ . From the a posteriori probability estimates, we assign the most probable estimate.

- Implementation -

We will use the object sklearn.svm.SVC:

```
from sklearn.svm import SVC
```

The file svm\_script.py provides a complete example of classification using the function SVC. Use this example to familiarize yourself with the syntax and then reproduce a similar experience with the data iris as suggested below.

1) Based on the documentation at the following link:

```
http://scikit-learn.org/stable/modules/svm.html
```

Write a code that will classify class 1 against class 2 of the dataset iris using the first two variables and a linear kernel. Leaving half of the data aside, evaluate the model's generalization performance. The dataset iris is obtained using the following lines of code:

```
from sklearn import datasets
iris = datasets.load_iris()
X = iris.data
y = iris.target
X = X[y != 0, :2]
y = y[y != 0]
```

- Compare the result with the one obtained by using a polynomial kernel-based SVM.
- 3) Show that the primal problem solved by the SVM can be rewritten as follows:

$$\underset{\mathbf{w} \in \mathcal{H}, w_0 \in \mathbb{R}}{\arg \min} \left( \frac{1}{2} \|\mathbf{w}\|^2 + C \sum_{i=1}^n \left[ 1 - y_i \left( \langle \mathbf{w}, \Phi(\mathbf{x}_i) \rangle + w_0 \right) \right]_+ \right)$$

4) Explain the sentence: "An SVM minimizes the classification error by using a convex majorant of the function which is 1 when the margin is negative and 0 otherwise." The function  $x \to [1-x]_+ = \max(0, 1-x)$  is called *Hinge* (charnière in French).

## **SVM GUI**

- Start the script svm\_gui.py available at the link: http://scikit-learn.org/stable/auto\_examples/applications/svm\_gui.html
  - This application allows a real-time evaluation of the impact that the choice of the kernel and the regularization parameter C have.
- Generate a very unbalanced data set with much more points in one class than in the other (at least 90% vs 10%).
- Using a linear kernel and decreasing the parameter C what do you observe?
  This phenomenon can be corrected in practice by weighting more errors on the lesser class (parameter class\_weight de SVC) or by a re-calibration technique (used with SVC(..., probability=True)).

## Classification of faces

The following example is a face classification problem. The database to be used is available at the following address: http://perso.telecom-paristech.fr/~gramfort/lfw.zip.

By modifying the sample code given in the last part of the file svm\_script.py:

- 5) Show the influence of the regularization parameter. For example, the prediction error can be displayed as a function of C on a logarithmic scale between 1e5 and 1e-5.
- 6) By adding nuisance variables, thus increasing the number of variables to the number of learning points fixed, show that performance drops.
- 7) Explain why the features are centered and reduced 1.148-150.
- 8) What is the effect of choosing a non-linear RBF kernel on prediction? You will be able to improve the prediction with a reduction of dimension based on the object sklearn.decomposition.RandomizedPCA.

## To go further: Computation of the duality gap

The following example:

```
http://scikit-learn.org/stable/auto_examples/svm/plot_separating_hyperplane.html# example-svm-plot-separating-hyperplane-py
```

explains how to access the parameters estimated by the model (vector of coefficients  $\mathbf{w}$  in the attribute coef,  $w_0$  attribute intercept, list of support vectors, coefficients of the dual problem).

- Based on this example, write a code that will calculate the value of the primal and dual objective function values and verify that the values are close.
- How does the difference between the two values vary when the optimization tolerance is varied (parameter tol of SVC)?

```
from time import time
import pylab as pl
from sklearn.cross_validation import train_test_split
from sklearn.datasets import fetch_lfw_people
from sklearn.svm import SVC
# Download the data (if not already on disk); load it as numpy arrays
lfw_people = fetch_lfw_people(min_faces_per_person=70, resize=0.4,
                       color=True, funneled=False, slice_=None,
                       download_if_missing=True)
# data_home='.'
# introspect the images arrays to find the shapes (for plotting)
images = lfw_people.images / 255.
n_samples, h, w, n_colors = images.shape
# the label to predict is the id of the person
target_names = lfw_people.target_names.tolist()
# Pick a pair to classify such as
names = ['Tony Blair', 'Colin Powell']
# names = ['Donald Rumsfeld', 'Colin Powell']
idx0 = (lfw_people.target == target_names.index(names[0]))
idx1 = (lfw_people.target == target_names.index(names[1]))
images = np.r_[images[idx0], images[idx1]]
n_samples = images.shape[0]
y = np.r_[np.zeros(np.sum(idx0)), np.ones(np.sum(idx1))].astype(np.int)
# Extract features
# features using only illuminations
X = (np.mean(images, axis=3)).reshape(n_samples, -1)
# # or compute features using colors (3 times more features)
# X = images.copy().reshape(n_samples, -1)
# Scale features
X -= np.mean(X, axis=0)
X /= np.std(X, axis=0)
# Split data into a half training and half test set
# X_train, X_test, y_train, y_test, images_train, images_test = \
  train_test_split(X, y, images, test_size=0.5, random_state=0)
# X_train, X_test, y_train, y_test = \
  train_test_split(X, y, test_size=0.5, random_state=0)
indices = np.random.permutation(X.shape[0])
train_idx, test_idx = indices[:X.shape[0] / 2], indices[X.shape[0] / 2:]
X_train, X_test = X[train_idx, :], X[test_idx, :]
y_train, y_test = y[train_idx], y[test_idx]
images_train, images_test = images[
```

```
train_idx, :, :, :], images[test_idx, :, :, :]
# Quantitative evaluation of the model quality on the test set
print "Fitting the classifier to the training set"
t0 = time()
clf = SVC(kernel='linear', C=1.0)
clf = clf.fit(X_train, y_train)
print "Predicting the people names on the testing set"
t0 = time()
y_pred = clf.predict(X_test)
print "done in %0.3fs" % (time() - t0)
print "Chance level : %s" % max(np.mean(y), 1. - np.mean(y))
print "Accuracy : %s" % clf.score(X_test, y_test)
# Look at the coefficients
pl.figure()
pl.imshow(np.reshape(clf.coef_, (h, w)))
# Qualitative evaluation of the predictions using matplotlib
def plot_gallery(images, titles, n_row=3, n_col=4):
   """Helper function to plot a gallery of portraits"""
   pl.figure(figsize=(1.8 * n_col, 2.4 * n_row))
   pl.subplots_adjust(bottom=0, left=.01, right=.99, top=.90,
                  hspace=.35)
   for i in range(n_row * n_col):
      pl.subplot(n_row, n_col, i + 1)
      pl.imshow(images[i])
      pl.title(titles[i], size=12)
      pl.xticks(())
      pl.yticks(())
def title(y_pred, y_test, names):
  pred_name = names[int(y_pred)].rsplit(' ', 1)[-1]
   true_name = names[int(y_test)].rsplit(' ', 1)[-1]
   return 'predicted: %s\ntrue: %s' % (pred_name, true_name)
prediction_titles = [title(y_pred[i], y_test[i], names)
                for i in range(y_pred.shape[0])]
plot_gallery(images_test, prediction_titles)
pl.show()
```

# Références

[1] T. Hastie, R. Tibshirani, and J. Friedman. *The elements of statistical learning*. Springer Series in Statistics. Springer, New York, second edition, 2009. http://www-stat.stanford.edu/~tibs/ElemStatLearn/. 1

- [2] B. Schölkopf and A. J. Smola. Learning with kernels: Support vector machines, regularization, optimization, and beyond. MIT press, 2002. 1
- [3]Vladimir N<br/> Vapnik. Statistical learning theory. Wiley, 1998.  ${\color{black}1}$