Support Vector Machines (SVM) in bioinformatics

## Day 1: Introduction to SVM

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## **3** days outline

- Day 1: Introduction to SVM
- Day 2: Applications in bioinformatics
- Day 3: Advanced topics and current research

## **Today's outline**

- 1. SVM: A brief overview (FAQ)
- 2. Simplest SVM: linear classifier for separable data
- 3. More useful SVM: linear classifiers for general data
- 4. Even more useful SVM: non-linear classifiers for general data
- 5. Remarks

## Part 1

## SVM: a brief overview (FAQ)

## What is a SVM?

- a family of learning algorithm for classification of objects into two classes (works also for regression)
- Input: a training set

 $\mathcal{S} = \{(x_1, y_1), \dots, (x_N, y_N)\}$ 

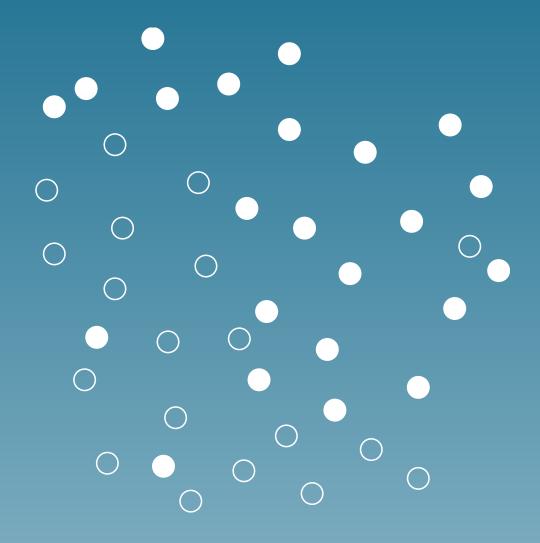
of objects  $x_i \in \mathcal{X}$  and their known classes  $y_i \in \{-1, +1\}$ .

• Output: a classifier  $f : \mathcal{X} \to \{-1, +1\}$  which predicts the class f(x) for any (new) object  $x \in \mathcal{X}$ .

#### Examples of classification tasks (more tomorrow)

- Optical character recognition: x is an image, y a character.
- Text classification: x is a text, y is a category (topic, spam / non spam...)
- Medical diagnosis: x is a set of features (age, sex, blood type, genome...), y indicates the risk.
- Protein secondary structure prediction: x is a string, y is a secondary structure

## Pattern recognition example



## Are there other methods for classification?

- Bayesian classifier (based on maximum a posteriori probability)
- Fisher linear discriminant
- Neural networks
- Expert systems (rule-based)
- Decision tree



## Why is it gaining popularity

- Good performance in real-world applications
- Computational efficiency (no local minimum, sparse representation...)
- Robust in high dimension (e.g., images, microarray data, texts)
- Sound theoretical foundations

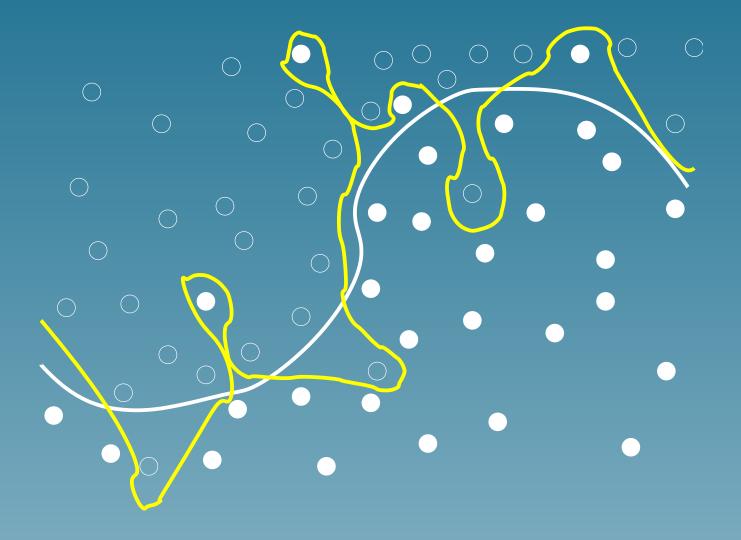
## Why is it so efficient?

- Still a research subject
- Always try to classify objects with large confidence, which prevent from overfitting
- No strong hypothesis on the data generation process (contrary to Bayesian approaches)

## What is overfitting?

- There is always a trade-off between good classification of the training set, and good classification of future objects (generalization performance)
- Overfitting means fitting too much the training data, which degrades the generalization performance
- Very important in large dimensions, or with complex non-linear classifiers.

## **Overfitting example**



### What is Vapnik's Statistical Learning Theory

- The mathematical foundation of SVM
- Gives conditions for a learning algorithm to generalize well
- The "capacity" of the set of classifiers which can be learned must be controlled

#### Why is it relevant for bioinformatics?

- Classification problems are very common (structure, function, localization prediction; analysis of microarray data; ...)
- Small training sets in high dimension is common
- Extensions of SVM to non-vector objects (strings, graphs...) is natural



# Simplest SVM: Linear SVM for separable training sets

#### Framework

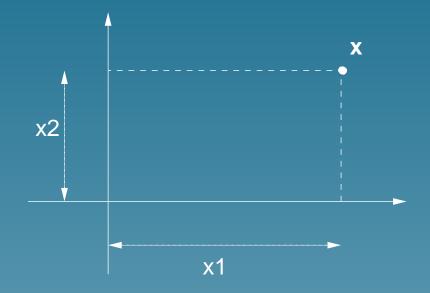
• We suppose that the object are finite-dimensional real vectors:  $\mathcal{X} = \mathbb{R}^n$  and an object is:

$$\vec{x} = (x_1, \dots, x_m).$$

•  $x_i$  can for example be a feature of a more general object

 Example: a protein sequence can be converted to a 20-dimensional vector by taking the amino-acid composition

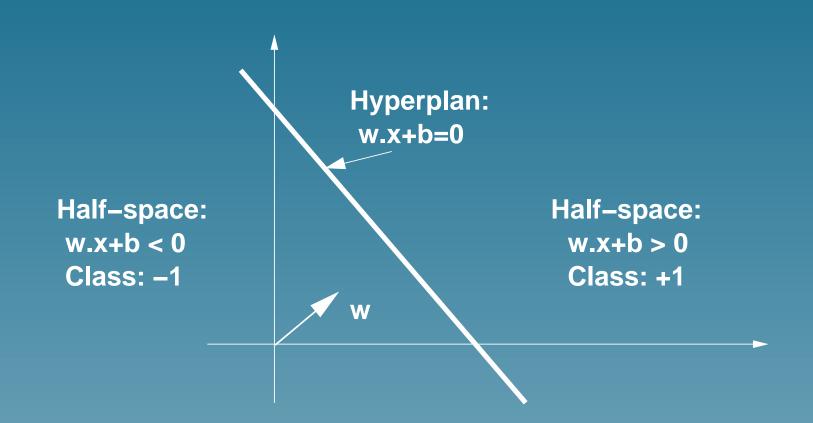
## Vectors and inner product



inner product:

$$\vec{x}.\vec{x'} = x_1 x'_1 + x_2 x'_2 \quad (+ \ldots + x_m x'_m) \tag{1}$$
$$= ||\vec{x}||.||\vec{x'}||.\cos(\vec{x},\vec{x'}) \tag{2}$$

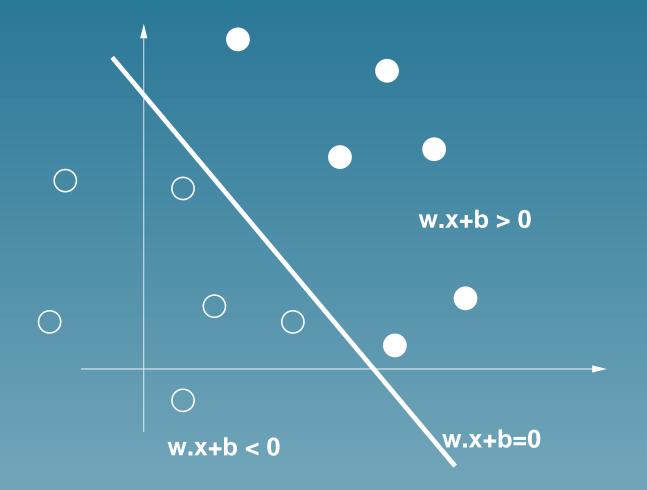
## Linear classifier



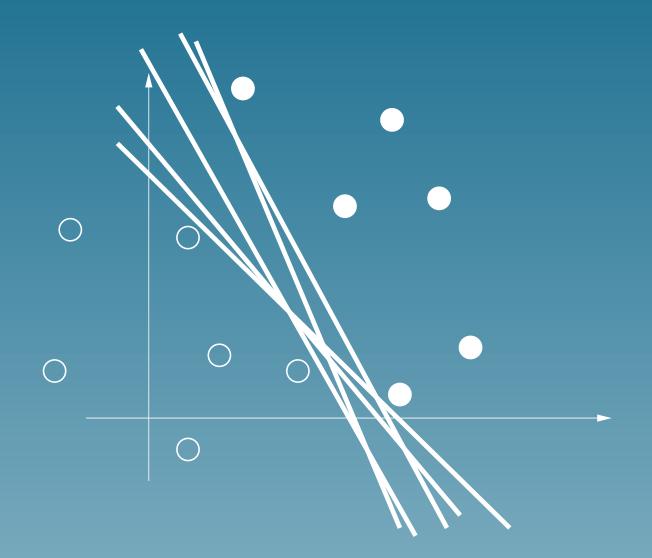
Classification is base on the sign the decision function:

$$f_{\vec{w},b}(\vec{x}) = \vec{w}.\vec{x} + b$$

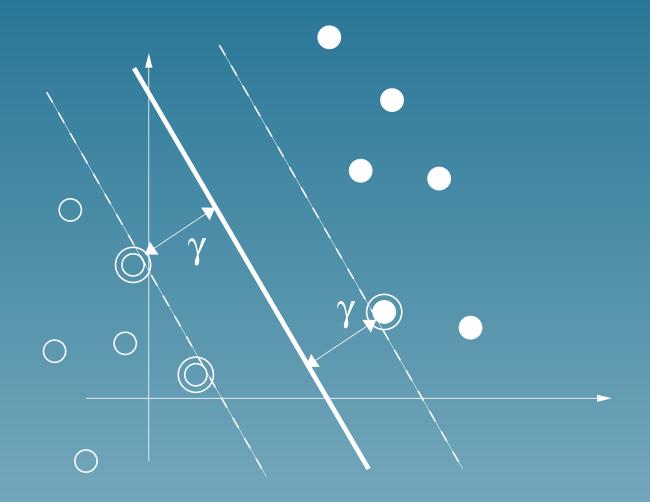
## Linearly separable training set



## Which one is the best?

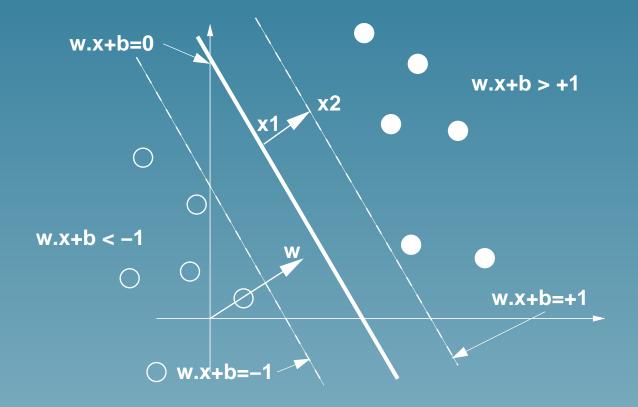


## Vapnik's answer : LARGEST MARGIN



## How to find the optimal hyperplane?

For a given linear classifier  $f_{\vec{w},b}$  consider the tube defined by the values -1 and +1 of the decision function:



## The width of the tube is $1/||\vec{w}||$

Indeed, the points  $\vec{x_1}$  and  $\vec{x_2}$  satisfy:

 $\begin{cases} \vec{w}.\vec{x_1} + b = 0, \\ \vec{w}.\vec{x_2} + b = 1. \end{cases}$ 

By subtracting we get  $\vec{w}.(\vec{x}_2 - \vec{x}_1) = 1$ , and therefore:

$$\gamma = ||\vec{x}_2 - \vec{x}_1|| = \frac{1}{||\vec{w}||}.$$

# All training points should be on the right side of the tube

For positive examples  $(y_i = 1)$  this means:

 $\vec{w}.\vec{x}_i + b \ge 1$ 

For negative examples  $(y_i = -1)$  this means:

 $\vec{w}.\vec{x}_i + b \le -1$ 

Both cases are summarized as follows:

 $\forall i = 1, \dots, N, \qquad \overline{y_i (\vec{w}.\vec{x_i} + b)} \ge 1$ 

## Finding the optimal hyperplane

The optimal hyperplane is defined by the pair  $(\vec{w}, b)$  which solves the following problem:

Minimize:

 $||\vec{w}||^2$ 

under the constraints:

 $\forall i = 1, ..., N, \qquad y_i (\vec{w}.\vec{x}_i + b) - 1 \ge 0.$ 

This is a classical quadratic program.

## How to find the minimum of a convex function?

If  $h(u_1, \ldots, u_n)$  is a convex and differentiable function of n variable, then  $\vec{u}^*$  is a minimum if and only if:

$$\nabla h(u^*) = \begin{pmatrix} \frac{\partial h}{\partial u_1}(\vec{u}^*) \\ \vdots \\ \frac{\partial h}{\partial u_1}(\vec{u}^*) \end{pmatrix} = \begin{pmatrix} 0 \\ \vdots \\ 0 \end{pmatrix}$$

$$h(u)$$

$$u^*$$

# How to find the minimum of a convex function with linear constraints?

Suppose that we want the minimum of h(u) under the constraints:

$$g_i(\vec{u}) \ge 0, \quad i = 1, \dots, N,$$

where each function  $g_i(\vec{u})$  is affine.

We introduce one variable  $\alpha_i$  for each constraint and consider the Lagrangian:

$$L(\vec{u}, \vec{\alpha}) = h(\vec{u}) - \sum_{i=1}^{N} \alpha_i g_i(\vec{u}).$$

## Lagrangian method (ctd.)

For each  $\vec{\alpha}$  we can look for  $\vec{u}_{\alpha}$  which minimizes  $L(\vec{u}, \vec{\alpha})$  (with no constraint), and note the dual function:

$$L(\vec{\alpha}) = \min_{\vec{u}} L(\vec{u}, \vec{\alpha}).$$

The dual variable  $\vec{\alpha}^*$  which maximizes  $L(\vec{\alpha})$  gives the solution of the primal minimization problem with constraint:

$$\vec{u}^* = \vec{u}_{\alpha^*}.$$

## **Application to optimal hyperplane**

In order to minimize:

$$\frac{1}{2}||\vec{w}||^2$$

under the constraints:

 $\forall i = \overline{1, \dots, N}, \quad y_i \left( \vec{w} \cdot \vec{x}_i + b \right) - 1 \ge 0.$ 

we introduce one dual variable  $\alpha_i$  for each constraint, i.e., for each training point. The Lagrangian is:

$$L(\vec{w}, b, \vec{\alpha}) = \frac{1}{2} ||\vec{w}||^2 - \sum_{i=1}^{N} \alpha_i \left( y_i \left( \vec{w} \cdot \vec{x}_i + b \right) - 1 \right).$$

#### Solving the dual problem

#### The dual problem is to find $\alpha^*$ maximize

$$L(\vec{\alpha}) = \sum_{i=1}^{N} \alpha_i - \frac{1}{2} \sum_{i,j=1}^{N} \alpha_i \alpha_j y_i y_j \vec{x}_i \cdot \vec{x}_j,$$

under the (simple) constraints  $\alpha_i \ge 0$  (for i = 1, ..., N), and

$$\sum_{i=1}^{N} \alpha_i y_i = 0.$$

 $\vec{\alpha}^*$  can be easily found using classical optimization softwares.

## **Recovering the optimal hyperplane**

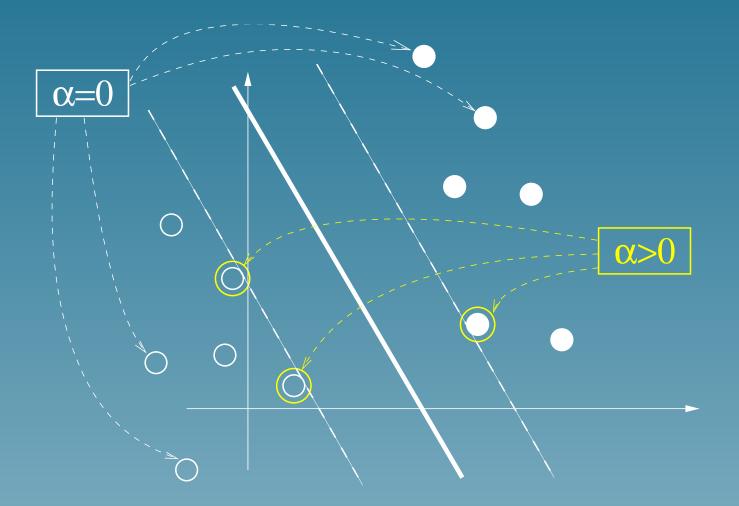
Once  $\vec{\alpha}^*$  is found, we recover  $(\vec{w}^*, b^*)$  corresponding to the optimal hyperplane.  $w^*$  is given by:

$$\vec{w}^* = \sum_{i=1}^N \alpha_i \vec{x}_i$$

and the decision function is therefore:

$$f^{*}(\vec{x}) = \vec{w}^{*}.\vec{x} + b^{*}$$
$$= \sum_{i=1}^{N} \alpha_{i}\vec{x}_{i}.\vec{x} + b^{*}.$$

## Interpretation : support vectors



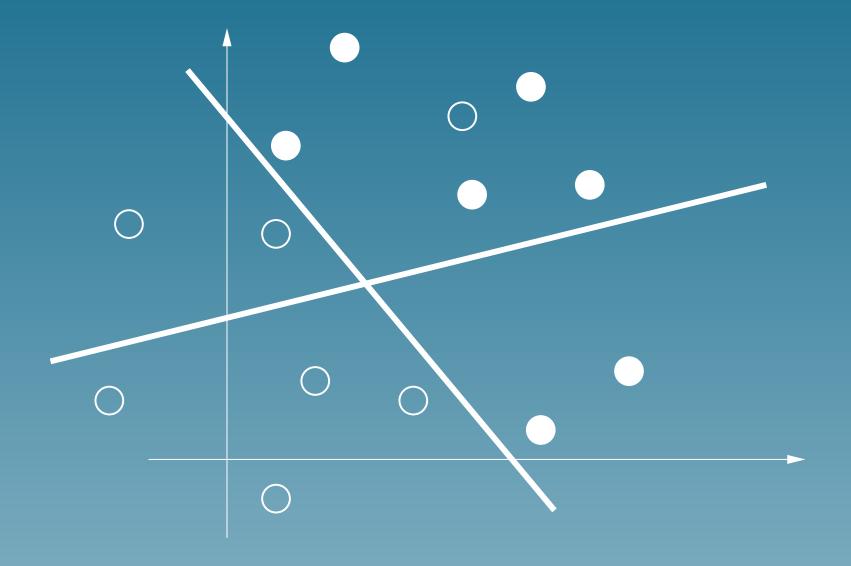
## Simplest SVM: conclusion

- Finds the optimal hyperplane, which corresponds to the largest margin
- Can be solved easily using a dual formulation
- The solution is sparse: the number of support vectors can be very small compared to the size of the training set
- Only support vectors are important for prediction of future points.
   All other points can be forgotten.



## More useful SVM: Linear SVM for general training sets

## In general, training sets are not linearly separable



## What goes wrong?

#### The dual problem, maximize

$$L(\vec{\alpha}) = \sum_{i=1}^{N} \alpha_i - \frac{1}{2} \sum_{i,j=1}^{N} \alpha_i \alpha_j y_i y_j \vec{x}_i \cdot \vec{x}_j$$

under the constraints  $\alpha_i \geq 0$  (for  $i = 1, \ldots, N$ ), and

$$\sum_{i=1}^{N} \alpha_i y_i = 0,$$

has no solution: the larger some  $\alpha_i$ , the larger the function to maximize.

#### Forcing a solution

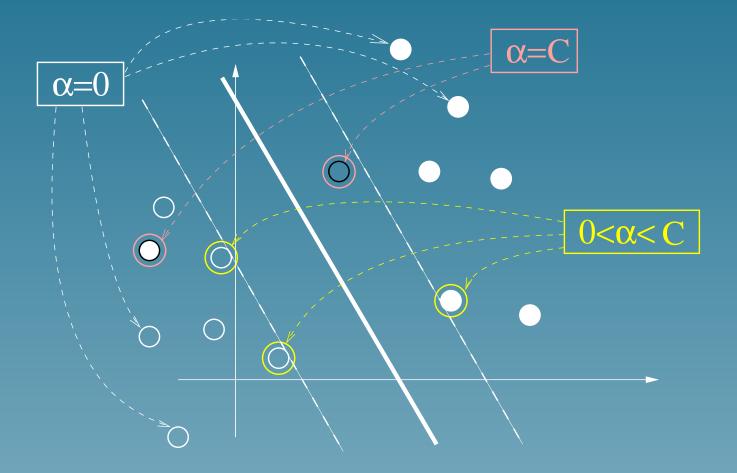
One solution is to limit the range of  $\vec{\alpha}$ , to be sure that one solution exists. For example, maximize

$$L(\vec{\alpha}) = \sum_{i=1}^{N} \alpha_i - \frac{1}{2} \sum_{i,j=1}^{N} \alpha_i \alpha_j y_i y_j \vec{x}_i \cdot \vec{x}_j$$

under the constraints:

$$\begin{cases} 0 \le \alpha_i \le C, & \text{for } i = 1, \dots, N \\ \sum_{i=1}^N \alpha_i y_i = 0. \end{cases}$$

# Interpretation



#### Remarks

• This formulation finds a trade-off between:

minimizing the training error
maximizing the margin

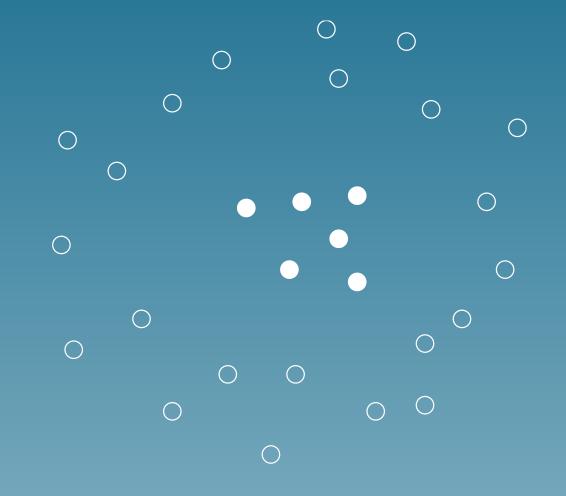
Other formulations are possible to adapt SVM to general training sets.

 All properties of the separable case are conserved (support vectors, sparseness, computation efficiency...)

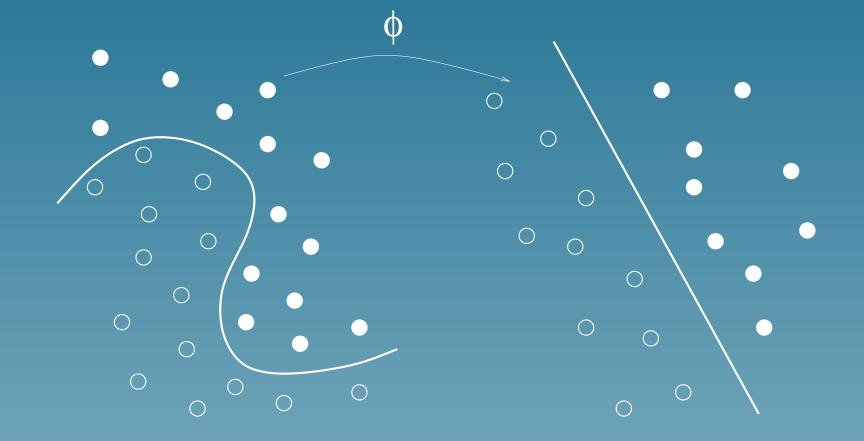


# General SVM: Non-linear classifiers for general training sets

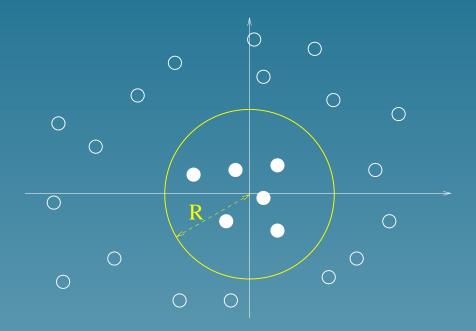
#### Sometimes linear classifiers are not interesting



# Solution: non-linear mapping to a feature space



#### Example



Let  $\Phi(\vec{x}) = (x_1^2, x_2^2)'$ ,  $\vec{w} = (1, 1)'$  and b = 1. Then the decision function is:

$$f(\vec{x}) = x_1^2 + x_2^2 - R^2 = \vec{w} \cdot \Phi(\vec{x}) + b,$$

#### Kernel (simple but important)

For a given mapping  $\Phi$  from the space of objects  $\mathcal{X}$  to some feature space, the kernel of two objects x and x' is the inner product of their images in the features space:

$$\forall x, x' \in \mathcal{X}, \quad K(x, x') = \vec{\Phi}(x).\vec{\Phi}(x').$$

Example: if  $\vec{\Phi}(\vec{x}) = (x_1^2, x_2^2)'$ , then  $K(\vec{x}, \vec{x}') = \vec{\Phi}(\vec{x}) \cdot \vec{\Phi}(\vec{x}') = (x_1)^2 (x_1')^2 + (x_2)^2 (x_2')^2.$ 

#### Training a SVM in the feature space

Replace each  $\vec{x}.\vec{x}'$  in the SVM algorithm by K(x,x')

The dual problem is to maximize

$$L(\vec{\alpha}) = \sum_{i=1}^{N} \alpha_i - \frac{1}{2} \sum_{i,j=1}^{N} \alpha_i \alpha_j y_i y_j K(x_i, x_j),$$

under the constraints:

$$\begin{cases} 0 \le \alpha_i \le C, & \text{for } i = 1, \dots, N \\ \sum_{i=1}^N \alpha_i y_i = 0. \end{cases}$$

## **Predicting with a SVM in the feature space**

The decision function becomes:

$$f(x) = \vec{w}^* \cdot \vec{\Phi}(x) + b^*$$

$$= \sum_{i=1}^N \alpha_i K(x_i, x) + b^*.$$
(4)

#### The kernel trick

- The explicit computation of  $\vec{\Phi}(x)$  is not necessary. The kernel K(x, x') is enough. SVM work implicitly in the feature space.
- It is sometimes possible to easily compute kernels which correspond to complex large-dimensional feature spaces.

#### Kernel example

For any vector  $\vec{x} = (x_1, x_2)'$ , consider the mapping:

$$\Phi(\vec{x}) = \left(x_1^2, x_2^2, \sqrt{2}x_1x_2, \sqrt{2}x_1, \sqrt{2}x_2, 1\right)'.$$

The associated kernel is:

$$K(\vec{x}, \vec{x}') = \Phi(\vec{x}) \cdot \Phi(\vec{x}')$$
  
=  $(x_1 x'_1 + x_2 x'_2 + 1)^2$   
=  $(\vec{x} \cdot \vec{x}' + 1)^2$ 

#### **Classical kernels for vectors**

• Polynomial:

$$K(x, x') = (x \cdot x' + 1)^d$$

• Gaussian radial basis function

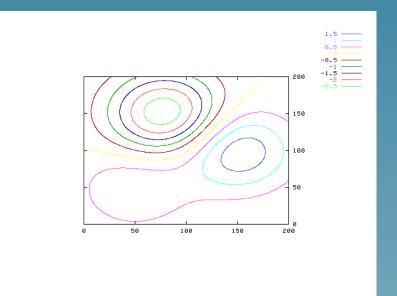
$$K(x, x') = \exp\left(\frac{||x - x'||^2}{2\sigma^2}\right)$$

Sigmoid

$$K(x, x') = \tanh(\kappa x \cdot x' + \theta)$$

## Example: classification with a Gaussian kernel

$$f(\vec{x}) = \sum_{i=1}^{N} \alpha_i \exp\left(\frac{||\vec{x} - \vec{x}_i||^2}{2\sigma^2}\right)$$





# Conclusion (day 1)

## Conclusion

- SVM is a simple but extremely powerful learning algorithm for binary classification
- The freedom to choose the kernel offers wonderful opportunities (see day 3: one can design kernels for non-vector objects such as strings, graphs...)
- More information : http://www.kernel-machines.org
- Lecture notes (draft) on my homepage