

CISC 889 Bioinformatics (Spring 2004)

Support Vector Machines I

The methodology

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Terminologies

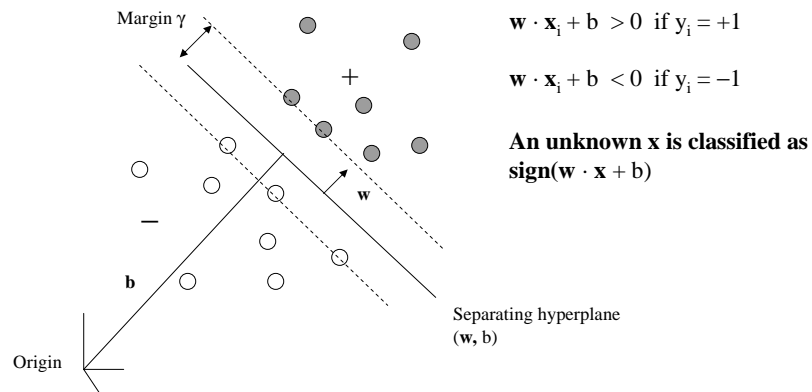
- An object \mathbf{x} is represented by a set of m attributes x^i , $1 \leq i \leq m$.
- A set of n training examples $S = \{ (\mathbf{x}_1, y_1), \dots, (\mathbf{x}_n, y_n) \}$, where y_i is the classification (or label) of instance \mathbf{x}_i .
 - For binary classification, $y_i = \{-1, +1\}$, and for k -class classification, $y_i = \{1, 2, \dots, k\}$.
 - Without loss of generality, we focus on binary classification.
- The task is to learn the mapping: $\mathbf{x}_i \rightarrow y_i$
- A machine is a learned function/mapping/hypothesis h :
$$\mathbf{x}_i \rightarrow h(\mathbf{x}_i, \alpha)$$
where α stands for parameters to be fixed during training.
- Performance is measured as

$$E = (1/2n) \sum_{i=1 \text{ to } n} |y_i - h(\mathbf{x}_i, \alpha)|$$

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Linear SVMs: find a hyperplane (specified by normal vector \mathbf{w} and perpendicular distance \mathbf{b} to the origin) that separates the positive and negative examples with the largest margin.



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Rosenblatt's Algorithm (1956)

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 $\eta$ ; // is the learning rate
 $\mathbf{w}_0 = \mathbf{0}$ ;  $b_0 = 0$ ;  $k = 0$ 
 $R = \max_{1 \leq i \leq n} \|\mathbf{x}_i\|$ 

error = 1; // flag for misclassification/mistake
while (error) { // as long as modification is made in the for-loop
    error = 0;

    for (i = 1 to n) {
        if ( $y_i (\langle \mathbf{w}_k, \mathbf{x}_i \rangle + b_k) \leq 0$ ) { // misclassification
             $\mathbf{w}_{k+1} = \mathbf{w}_k + \eta y_i \mathbf{x}_i$  // update the weight
             $b_{k+1} = b_k + \eta y_i R^2$  // update the bias
             $k = k + 1$ 
            error = 1;
        }
    }
}

return  $(\mathbf{w}_k, b_k)$  // hyperplane that separates the data, where k is the number of
// modifications.

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Questions w.r.t. Rosenblatt's algorithm

- Is the algorithm guaranteed to converge?
- How quickly does it converge?

Novikoff Theorem:

Let S be a training set of size n and $R = \max_{1 \leq i \leq n} \| \mathbf{x}_i \|$. If there exists a vector \mathbf{w}^* such that $\| \mathbf{w}^* \| = 1$ and

$$y_i (\mathbf{w}^* \cdot \mathbf{x}_i) \geq \gamma,$$

for $1 \leq i \leq n$, then the number of modifications before convergence is at most

$$(R/\gamma)^2.$$

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Proof:

1. $\mathbf{w}_t \cdot \mathbf{w}^* = \mathbf{w}_{t-1} \cdot \mathbf{w}^* + \eta y_i \mathbf{x}_i \cdot \mathbf{w}^* \geq \mathbf{w}_{t-1} \cdot \mathbf{w}^* + \eta \gamma$
 $\mathbf{w}_t \cdot \mathbf{w}^* \geq t \eta \gamma$
2. $\| \mathbf{w}_t \|^2 = \| \mathbf{w}_{t-1} \|^2 + 2 \eta y_i \mathbf{x}_i \cdot \mathbf{w}_{t-1} + \eta^2 \| \mathbf{x}_i \|^2$
 $\leq \| \mathbf{w}_{t-1} \|^2 + \eta^2 \| \mathbf{x}_i \|^2$
 $\leq \| \mathbf{w}_{t-1} \|^2 + \eta^2 R^2$
 $\| \mathbf{w}_t \|^2 \leq t \eta R^2$
3. $\sqrt{t \eta R} \| \mathbf{w}^* \| \geq \mathbf{w}_t \cdot \mathbf{w}^* \geq t \eta \gamma$
 $t \leq (R/\gamma)^2.$

Note:

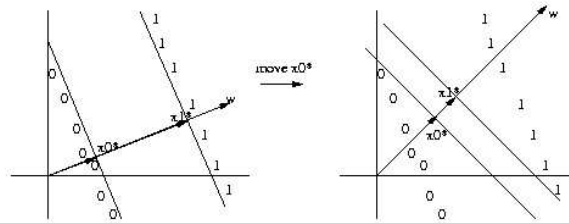
- Without loss of generality, the separating plane is assumed to pass the origin, i.e., no bias b is necessary.
- The learning rate η seems to have no bearing on this upper bound. (why?)
- What if the training data is not linearly separable, i.e., \mathbf{w}^* does not exist?

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Larger margin is preferred:

- converge more quickly
- generalize better



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Dual form

- The final hypothesis w is a linear combination of the training points:

$$w = \sum_{i=1 \text{ to } n} \alpha_i y_i x_i$$

where α_i are positive values proportional to the number of times misclassification of x_i has caused the weight to be updated.

- Vector α can be considered as alternative representation of the hypothesis; α_i can be regarded as an indication of the information content of the example x_i .
- The decision function can be rewritten as

$$\begin{aligned} h(x) &= \text{sign}(w \cdot x + b) \\ &= \text{sign}\left(\left(\sum_{j=1 \text{ to } n} \alpha_j y_j x_j\right) \cdot x + b\right) \\ &= \text{sign}\left(\sum_{j=1 \text{ to } n} \alpha_j y_j (x_j \cdot x) + b\right) \end{aligned}$$

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Rosenblatt's Algorithm in dual form

$\alpha = 0$; $b = 0$

$R = \max_{1 \leq i \leq n} \|x_i\|$

error = 1; // flag for misclassification

while (error) { // as long as modification is made in the for-loop

error = 0;

for (i = 1 to n) {

if ($y_i (\sum_{j=1}^{n} \alpha_j y_j (x_j \cdot x_i) + b) \leq 0$) { // misclassification

$\alpha_i = \alpha_i + 1$ // update the weight

$b = b + y_i R^2$ // update the bias

error = 1;

}

}

return (α, b) // hyperplane that separates the data, where k is the number of
// modifications.

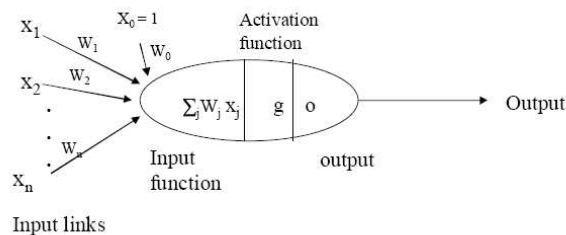
Notes:

- The training examples enter the algorithm as dot products ($x_j \cdot x_i$).
- α_i is a measure of information content; x_i with non-zero information content ($\alpha_i > 0$) are called support vectors, as they are located on the boundaries.

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Relationship to linear perceptrons

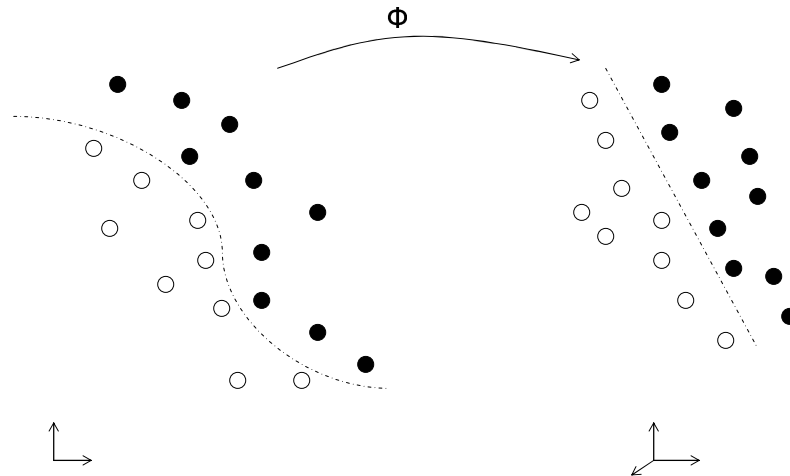


- Linear SVMs are almost identical to linear perceptrons
- They differ from each other when are generalize to handle non linear cases.

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Non-linear mapping to a feature space



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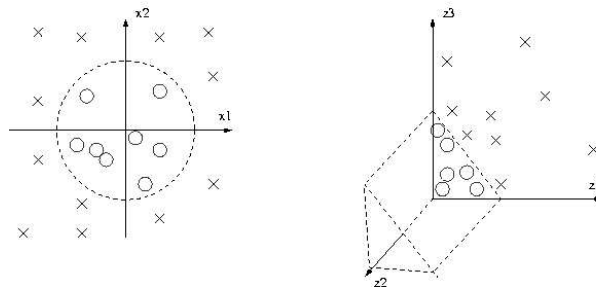
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Nonlinear SVMs

Input Space

Feature Space

$$\mathbf{x} \longrightarrow \Phi(\mathbf{x}) = \begin{pmatrix} x_1^2 \\ \sqrt{2}x_1x_2 \\ x_2^2 \end{pmatrix}$$



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Kernel function for mapping

- For input $\mathbf{X} = (x_1, x_2)$,
Define map $\Phi(\mathbf{X}) = (x_1 x_1, \sqrt{2} x_1 x_2, x_2 x_2)$.

$$K(\mathbf{X}, \mathbf{Y}) = \Phi(\mathbf{X}) \cdot \Phi(\mathbf{Y})$$

$$= (x_1 x_1, \sqrt{2} x_1 x_2, x_2 x_2) \cdot (y_1 y_1, \sqrt{2} y_1 y_2, y_2 y_2)$$
- Define Kernel function as $K(\mathbf{X}, \mathbf{Y}) = (\mathbf{X} \cdot \mathbf{Y})^2$.

$$= (x_1 x_1 y_1 y_1 + 2 x_1 x_2 y_1 y_2 + x_2 x_2 y_2 y_2)$$
- It has $K(\mathbf{X}, \mathbf{Y}) = \Phi(\mathbf{X}) \cdot \Phi(\mathbf{Y})$.

$$= (x_1 y_1 + x_2 y_2)(x_1 y_1 + x_2 y_2)$$
- We can compute the scalar product in feature space without computing Φ .**

$$= ((x_1, x_2) \cdot (y_1, y_2))^2$$

$$= (\mathbf{X} \cdot \mathbf{Y})^2$$

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Mercer's condition

Since kernel functions play an important role, it is important to know if a kernel gives dot products (in some higher dimension space).

For a kernel $K(x, y)$, if for any $g(x)$ such that $\int g(x)^2 dx$ is finite, we have

$$\int \int K(x, y) g(x) g(y) dx dy \geq 0,$$

then there exist a mapping Φ such that

$$K(x, y) = \Phi(x) \cdot \Phi(y)$$

Notes:

- Mercer's condition does not tell how to actually find Φ .
- Mercer's condition may be hard to check since it must hold for every $g(x)$.

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More kernel functions

some commonly used generic kernel functions

- Polynomial kernel: $K(\mathbf{x}, \mathbf{y}) = (1 + \mathbf{x} \cdot \mathbf{y})^p$
- Radial (or Gaussian) kernel: $K(\mathbf{x}, \mathbf{y}) = \exp(-\|\mathbf{x} - \mathbf{y}\|^2 / 2\sigma^2)$

Questions: By introducing extra dimensions (sometimes infinite), we can find a linearly separating hyperplane. But how can we be sure such a mapping to a higher dimension space will generalize well to unseen data? Because the mapping introduces flexibility for fitting the training examples, how to avoid overfitting?

Answer: Use the maximum margin hyperplane. (Vapnik theory)

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$$\mathbf{w} \cdot \mathbf{x}_+ + b = +1$$

$$\mathbf{w} \cdot \mathbf{x}_- + b = -1$$

$$\begin{aligned}\gamma &= \frac{1}{2} [(\mathbf{x}_+ \cdot \mathbf{w} / \|\mathbf{w}\|_2) - (\mathbf{x}_- \cdot \mathbf{w} / \|\mathbf{w}\|_2)] \\ &= 1 / \|\mathbf{w}\|_2\end{aligned}$$

Therefore, maximizing the geometric margin γ is equivalent to minimizing $\|\mathbf{w}\|_2$, under linear constraints.

$$\text{Min}_{\mathbf{w}, b} \langle \mathbf{w} \cdot \mathbf{w} \rangle$$

$$\text{subject to } y_i \langle \mathbf{w} \cdot \mathbf{x}_i \rangle + b \geq 1 \text{ for } i = 1, \dots, n$$

Lagrangian Theory

Quadratic programming optimization problem

... guaranteed to converge to the global minimum because of its being a convex

Note: advantages over the artificial neural nets

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Advanced Issues

- Soft margin
 - Allow misclassification, but with penalties
- Multiclass classification
 - Indirect: combine multiple binary classifiers into a single multiclass classifier
 - Direct: generalize binary classification methods
- SVM Regression
- Support vector clustering by Ben-Hur et al (2001)

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References and resources

- Cristianini & Shawe-Taylor, “*An introduction to Support Vector Machines*”, Cambridge University Press, 2000.
- Chris Burges, A tutorial
- www.kernel-machines.org
- SVMLight

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