Support Vector Machines

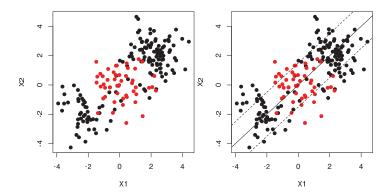
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Predictive Modeling

Support Vector Machines

- Support vector classifier is a natural approach for classification in the two-class setting, if the boundary between the two classes is linear
- In practice, we are faced with non-linear class boundaries



Support Vector Machines

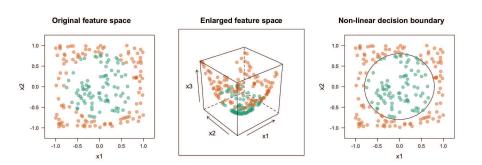
- **Support vector machine** (SVM) is an extension of the support vector classifier that results from enlarging the feature space
- Main idea: we may want to enlarge our feature space in order to accommodate a non-linear boundary between the classes
- Rather than fitting a support vector classifier using p features

$$X_1, X_2, \ldots, X_p$$

We could instead fit a support vector classifier using 2p features

$$X_1, X_1^2, X_2, X_2^2, \dots, X_p, X_p^2$$

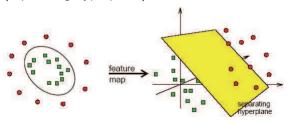
Enlarged Feature Space



We may enlarge the feature space by adding a third feature, say $X_3=X_1^2+X_2^2$

Enlarged Feature Space

- Why does enlarging the feature space lead to a non-linear decision boundary?
- In the enlarged feature space, the resulting decision boundary is in fact linear (seperating hyperplane)



• In the original feature space, the decision boundary is of the form q(x) = 0, where q is a quadratic polynomial, and its solutions are generally **non-linear**

Optimization Problem

Maximize M

$$\max_{\beta_0,\beta_1,\dots,\beta_p,\varepsilon_1,\varepsilon_2,\dots,\varepsilon_n} M \tag{1}$$

subject to

$$y_i \left(\beta_0 + \sum_{j=1}^p \beta_{j1} x_{j1} + \sum_{j=1}^p \beta_{j2} x_{j2}^2 \right) \ge M (1 - \varepsilon_i)$$
 (2)

for all i = 1, 2, ..., n

Optimization Problem

and subject to

$$\sum_{j=1}^{p} \sum_{k=1}^{2} \beta_{jk}^{2} = 1 \tag{3}$$

and to

$$\varepsilon_i \ge 0; \qquad \sum_{i=1}^n \varepsilon_i \le C$$
 (4)

where C is a nonnegative tuning parameter.

This optimization problem may become intractable for high dimensional feature space extension \rightarrow Kernel Trick

Kernel Trick: Support Vector Classifiers Rewritten

The linear support vector classifier can be represented as

$$f(x^*) = \beta_0 + \sum_{i=1}^n \alpha_i \langle x^*, x_i \rangle$$
 (5)

where x_i denotes training observations and x^* denotes a new observation we want to classify

• The inner product of two observations $x_i, x_{i'}$ is given by

$$\langle x_i, x_{i'} \rangle = \sum_{j=1}^r x_{ij} \cdot x_{i'j}$$
 (6)

Kernel Trick: Support Vector Classifiers Rewritten

- To estimate the parameters $\alpha_1, \ldots, \alpha_n$ and β_0 , all we need are the $\binom{n}{2}$ inner products $\langle x_i, x_{i'} \rangle$ between all pairs of training observations where $\binom{n}{2}$ means n(n-1)/2, and gives the number of pairs among a set of n items
- However, it turns out that α_i is **nonzero** only for the support vectors in the solution. If a training observation is not a support vector, then its α_i equals zero
- \bullet If ${\cal S}$ is the collection of indices of these support vectors, we can rewrite any support vector classifier function as

$$f(x^*) = \beta_0 + \sum_{i \in \mathcal{S}} \alpha_i \langle x^*, x_i \rangle \tag{7}$$

Kernel Trick: Kernel Functions

- Support vector machine (SVM) is an extension of the support vector classifier that results from enlarging the feature space in a specific way, using kernels
- In representing the linear classifier

$$f(x^*) = \beta_0 + \sum_{i \in \mathcal{S}} \alpha_i \langle x^*, x_i \rangle \tag{8}$$

and in computing its coefficients, all we need are inner products

• Every time the inner $\langle x, x_i \rangle$ appears in the representation for the support vector classifier, we replace it with a generalization of the inner product of the form

$$K(x_i, x_{i'}) \tag{9}$$

Kernel Trick: Kernel Functions

This results in the support vector machine

$$f(x^*) = \beta_0 + \sum_{i \in \mathcal{S}} \alpha_i K(x_i, x_{i'})$$
(10)

where

$$K(x_i, x_{i'}) \tag{11}$$

is called kernel function

 Kernel trick: Using such a generalized kernel instead of the standard linear kernel in the support vector classifier algorithm amounts to fitting a support vector classifier in a higher-dimensional space involving extended features!

Linear Kernel

If we choose in

$$f(x^*) = \beta_0 + \sum_{i \in \mathcal{S}} \alpha_i K(x^*, x_{i'})$$
(12)

for $K(x_i, x_{i'})$ the linear kernel function

$$K(x_i, x_{i'}) = \sum_{j=1}^{p} x_{ij} x_{i'j}$$
 (13)

this would just give us back the support vector classifier

 Linear kernel essentially quantifies the similarity of a pair of observations using Pearson (standard) correlation

Polynomial Kernel Function

Now we choose in

$$f(x^*) = \beta_0 + \sum_{i \in \mathcal{S}} \alpha_i K(x^*, x_{i'})$$
(14)

for $K(x_i, x_{i'})$ the **polynomial kernel function** of degree d

$$K(x_i, x_{i'}) = \left(1 + \sum_{j=1}^{p} x_{ij} x_{i'j}\right)^d$$
 (15)

where d is a positive polynomial integer

- Using such a kernel with d>1, instead of the standard linear kernel, in the support vector classifier algorithm leads to a much **more** flexible decision boundary
- Please see example 3.1 of the chapter Support Vector Machines

Radial Kernel

Now, we choose in

$$f(x^*) = \beta_0 + \sum_{i \in \mathcal{S}} \alpha_i K(x^*, x_{i'})$$
(16)

for $K(x_i, x_{i'})$ the radial kernel

$$K(x_i, x_{i'}) = \exp\left(-\gamma \sum_{j=1}^{p} (x_{ij} - x_{i'j})^2\right)$$
 (17)

where γ is a positive constant

Radial Kernel

• If a given test observation $x^* = (x_1^* \dots, x_p^*)$ is **far** from training observation x_i wrt Euclidean distance, then

$$\sum_{j=1}^{p} (x_{j}^{*} - x_{i'j})^{2}$$

will be large

Then

$$K(x_i, x_{i'}) = \exp\left(-\gamma \sum_{j=1}^{p} (x_{ij} - x_{i'j})^2\right)$$
 (18)

will be tiny

Radial Kernel

• Recall that the predicted class label for the test observation x^* is based on the sign of

$$f(x^*) = \beta_0 + \sum_{i \in \mathcal{S}} \alpha_i K(x^*, x_i)$$
(19)

- In other words, training observations x_i that are far from x^* will play essentially **no** role in the predicted class label for x^*
- This means that the radial kernel has very local behavior, in the sense that only nearby training observations have an effect on the class label of a test observation
- Please see example 3.2 of the chapter Support Vector Machines

Advantages of Kernel Functions

- What is the advantage of using a kernel rather than simply enlarging the feature space using functions of the original features?
- One advantage is computational, and it amounts to the fact that using kernels, for training one need only compute

$$K(x_i, x_{i'})$$

for all $\binom{n}{2}$ distinct training observation pairs i, i'

Advantages of Kernel Functions

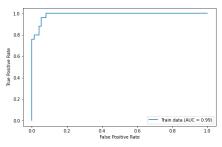
Training a SVM by computing

$$K(x_i,x_{i'})$$

for all $\binom{n}{2}$ distinct training observation pairs i, i' can be done without explicitly working in the enlarged feature space

- This is important because in many applications of SVMs, the enlarged feature space is so large that computations are intractable
- For some kernels, such as the radial kernel, the feature space is implicit and infinite-dimensional, so we could never do the computations there anyway.

 The ROC curve is a popular graphic for simultaneously displaying the two types of errors - false positive and true positive rates - for all possible thresholds.



• The name **ROC** is historic, and comes from communications theory. It is an acronym for **receiver operating characteristics**.

- SVMs output class labels for each observation
- However, it is also possible to obtain fitted values for each observation, which are the numerical scores (distances to separating hyperplane) used to obtain the class labels
- For an SVM with a non-linear kernel, the equation that yields the fitted value is given in

$$f(x^*) = \beta_0 + \sum_{i \in \mathcal{S}}^n \alpha_i K(x^*, x_i)$$
 (20)

- Relationship between the **fitted value** and the **class prediction** for a given observation is simple:
 - ▶ If the fitted value $f(x^*)$ exceeds a given threshold, then the observation is assigned to one class
 - ▶ If the fitted value $f(x^*)$ is **less** than this threshold, then it is assigned to the other class
- Threshold usually is zero, but we may choose a different value for the threshold

- A ROC curve is constructed by:
 - Choosing threshold value
 - **2** Computing fitted values $f(x^*)$ of observations
 - Classifying observations with respect to the chosen threshold
 - Computing the related true and false positive rates
 - For each threshold, display the corresponding true and false positive rate as ROC curve
- See example 4.1 of the Support Vector Machine chapter

SVMs with More than Two Classes

- So far, our discussion has been limited to the case of binary classification: that is, classification in the two-class setting
- How can we extend SVMs to the more general case where we have some arbitrary number of classes?
- The two most popular are the one-versus-one and one-versus-all approaches

One-versus-One Classification

- A **one-versus-one** or **all-pairs** approach constructs $\binom{K}{2}$ SVMs, each of which compares a pair of classes
- **Example:** one such SVM might compare the kth class, coded as +1, to the k'th class, coded as -1
- We classify a test observation using each of the $\binom{K}{2}$ classifiers, and we tally the number of times that the test observation is assigned to each of the K classes
- The final classification is performed by assigning the test observation to the class to which it was most frequently assigned in these $\binom{K}{2}$ pairwise classifications

One-versus-All Classification

- We fit K SVMs, each time comparing **one** of all the K classes to the **remaining** K-1 classes
- Let $\beta_{0k}, \alpha_{1k}, \ldots, \alpha_{pk}$ denote the parameters that result from fitting an SVM comparing the kth class (coded as +1) to the others (coded as -1)
- Let x^* denote a test observation. We assign the observation to the class for which

$$f(x^*) = \beta_{0k} + \sum_{i \in \mathcal{S}} \alpha_{ik} K(x^*, x_i)$$
(21)

is largest

• This amounts to a high level of confidence that the test observation belongs to the kth class rather than to any of the other classes.

Application to Gene Expression Data

- Khan data set consists of a number of tissue samples corresponding to four distinct types of small round blue cell tumors
- For each tissue sample, 2308 gene expression measurements are available
- **Goal:** By means of support vector machines to predict cancer subtype using gene expression measurements
- Please check example 5.3 of the Support Vector Machine chapter