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Variable Selection for SVM via Smoothing Spline ANOVA

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Abstract: It is well-known that the support vector machine paradigm is equivalent to solving a regularization problem in a reproducing kernel Hilbert space. The squared norm penalty in the standard support vector machine controls the smoothness of the classification function. We propose, under the framework of smoothing spline ANOVA models, a new type of regularization to conduct simultaneous classification and variable selection in the SVM. The penalty functional used is the sum of functional component norms, which automatically applies soft-thresholding operations to functional components hence yields sparse solutions. We suggest an efficient algorithm to solve the proposed optimization problem by iteratively solving the quadratic programming and linear programming. Numerical studies, on both simulated data and real datasets, show that the modified support vector machine gives very competitive performances compared to other popular classification algorithms, in terms of both classification accuracy and variable selection.

Key words and phrases: support vector machine, L_1 penalty, smoothing spline ANOVA, classification, sparsity.

1. Introduction

In classification problems, we are given a training data set of n examples from two or more populations. For each example $i, i = 1, ..., n$, in the training set, we observe an input vector $\mathbf{x}_i \in \mathbb{R}^d$, and a label y_i indicating one of the classes to which the example belongs. The binary classification problem is considered in this paper, and two classes are the positive class (labeled as $+1$) and the negative class (labeled as -1). Support vector machine (SVM) classifiers developed by Boser, Guyon, and Vapnik (1992) and Vapnik (1995) have gained popularity due to its promising performances in real-world applications such as text categorization, image recognition, gene expression array data analysis, and etc. However, the standard SVM decision rule utilizes all the input variables, which is not desirable when some variables are not relevant or have too much noise. Hastie, Tibshirani, and Friedman (2001) demonstrated that the standard SVM can suffer from the existence of irrelevant variables. An appropriate variable selection is therefore needed to obtain a compact classifier with improved accuracy.

Several methods have been proposed for conducting variable selection in the SVM. In the linear SVM setting, Bradley and Mangasarian (1998) suggested the 1-norm SVM which imposes the absolute value penalty on the coefficients hence produces a sparse directional vector for the separating plane. Recently Zhu, Rosset, Hastie, and Tibshirani (2003) studied the solution property of the 1-norm SVM and suggested an algorithm to find the whole solution path over a range of tuning parameters. Fung and Mangasarian (2004) developed a fast Newton algorithm to solve the dual problem of the 1-norm SVM. Another class of methods are kernel scaling methods proposed by Weston, Mukherjee, Chapelle, Pontil, Poggio, and Vapnik (2000) and Grandvalet and Canu (2002). A special issue on variable and feature selection published by *Journal of Machine Learning Research* in 2003 introduced other approaches like Bi, Bennett, Embrechts, Breneman, and Song (2003) and Rakotomamonjy (2003). Recently Bach, Lanckriet and Jordan (2004) considered the block 1-norm regularization for learning a sparse conic combination of kernels. Their formulation can also be used for variable selection in nonparametric setting.

Different from all the methods above, we formulate the SVM as a regularization problem in the reproducing kernel Hilbert space (RKHS) with a novel form of the penalty functional. The optimization problem consists of two parts: the data fit part represented by the hinge loss function functional, and the regularization penalty part defined as the sum of function component norms. In the Gaussian regression context, this penalty was proposed and studied by Lin and Zhang (2002) and named as the component smoothing and selection operator (COSSO). Following their terminology, we will refer to our method as the COSSO SVM. We will show that the COSSO SVM inherits the desired properties of the SVM, and more importantly, it conducts variable selection and classification simultaneously. For the linear classification, the COSSO SVM reduces to the 1-norm SVM.

This paper is organized as follows. Section 2 gives an overview on the SVM regularization problem and the smoothing spline ANOVA framework for multivariate function estimation. Section 3 proposes the COSSO SVM method and studies its solution properties such as the existence and finite representation of the optimal classifier. In Section 4, we give an iterative algorithm to solve the COSSO SVM, and show that only quadratic and linear programming problems are needed for implementation. We also discuss the issue of parameter tuning in this section. Simulation results and real

examples are presented in Section 5. The final discussion is given in Section 6. The proofs of the theorems are in the appendix.

2. SVM and Smoothing Functional ANOVA

2.1. Regularization Problem

In supervised learning problems, our task is to learn a classification rule $f : \mathbb{R}^d \to \{+1, -1\}$ from the training set, so that we can assign a class label to any new subject observed in the future. In the statistical framework, the training data $(\mathbf{x}_i, y_i), i = 1, ..., n$, are generally assumed to be independent realizations of the random pair $(\mathbf{X}, Y)$ which has a joint distribution $P(\mathbf{X}, Y)$. Define $p(\mathbf{x}) = \text{Prob}(Y = +1|\mathbf{X} = \mathbf{x})$. When the 0-1 loss is used, the optimal rule minimizing the expected loss is $\text{sign}[p(\mathbf{x}) - 1/2]$. This is known as the Bayes rule.

The linear SVM is a large margin classifier which separates two classes by maximizing the margin between them. When a linear separation is not plausible, the nonlinear SVM maps the data into a high dimensional feature space and then implements the linear classification in the feature space. It is well-known that the nonlinear SVM can be cast as a regularization problem in a reproducing kernel Hilbert space (RKHS). Let $\mathcal{H}$ be the RKHS associated with some reproducing kernel, and $||\cdot||$ be the functional norm of any element in $\mathcal{H}$. The standard SVM amounts to solving the regularization problem

$$\min_{f \in \mathcal{H}} \quad \frac{1}{n} \sum_{i=1}^{n} [1 - y_i f(\mathbf{x}_i)]_+ + \lambda ||f||^2. \tag{2.1}$$

Here $[\tau]_+ = \tau$, for $\tau > 0$; $= 0$, otherwise. The hinge loss function $[1 - yf]_+$ is a convex upper bound of the misclassification rate. λ is the tuning parameter that can be adaptively chosen by the data. Once the solution $\hat{f}$ is obtained, the classification rule is $\text{sign}[\hat{f}(\mathbf{x})]$. See Wahba (1999) and Evgeniou, Pontil, and Poggio (1999) for more details. Lin (2002) showed that, if the reproducing kernel Hilbert space is rich enough (for example, associated with the Gaussian kernel or the spline kernel), the solution to (2.1) approaches the Bayes rule $\text{sign}[p(\mathbf{x}) - 1/2]$ when $n \to \infty$. This result provides a theoretical justification for the superior performances of the nonlinear SVM.

2.2. Smoothing Spline ANOVA

The smoothing spline analysis of variance (SS-ANOVA) models provide a general framework for high dimensional function estimation, as shown in Wahba (1990) and

Gu (2002). In the SS-ANOVA, any function $f(\mathbf{x}) = f(x^{(1)}, ..., x^{(d)})$ on a product domain $\mathcal{X}$ has a functional ANOVA decomposition

$$f(\mathbf{x}) = b + \sum_{j=1}^{d} f_j(x^{(j)}) + \sum_{j<k} f_{jk}(x^{(j)}, x^{(k)}) + \text{all higher-order interactions}, \quad (2.2)$$

where b is constant, f_j's are the main effects, and f_{jk}'s are the two-factor interactions. Each main effect $f_j, j = 1, \cdots, d$, is estimated in a reproducing kernel Hilbert space, denoted by $\mathcal{H}_j = [1] \oplus \bar{\mathcal{H}}_j$. The entire model space $\mathcal{H}$ is then the tensor product space $\otimes_{j=1}^{d} \mathcal{H}_j$ which admits the following tensor sum decomposition

$$\otimes_{j=1}^{d} \mathcal{H}_j = [1] \oplus \sum_{j=1}^{d} \bar{\mathcal{H}}_j \oplus \sum_{j<k} [\bar{\mathcal{H}}_j \otimes \bar{\mathcal{H}}_k] \oplus \cdots . \quad (2.3)$$

The space $\otimes_{j=1}^{d} \mathcal{H}_j$ is also an RKHS, and its reproducing kernel is the sum of the reproducing kernels of those component spaces. Each functional component in (2.2) lies in a subspace in the orthogonal decomposition (2.3) of $\otimes_{j=1}^{d} \mathcal{H}_j$. The identifiability of the components is assured by side conditions through averaging operators. Without loss of generality, we assume $\mathcal{X} = [0, 1]^d$. A typical choice of $\mathcal{H}_j$ is the ℓ-th order Sobolev Hilbert space: $W_\ell[0, 1] = \{g : g, g', ..., g^{(\ell-1)}$ are absolutely continuous, $g^{(\ell)} \in \mathcal{L}_2[0, 1]\}$. In particular, the space $W_2[0, 1]$ is an RKHS when equipped with the norm

$$\|g\|^2 = [\int_0^1 g(t)dt]^2 + [\int_0^1 g'(t)dt]^2 + \int_0^1 [g''(t)]^2 dt, \quad \forall g \in W_2[0, 1].$$

It has the reproducing kernel $1 + R(s, t)$, where

$$R(s, t) = k_1(s)k_1(t) + k_2(s)k_2(t) - k_4(|s - t|), \quad (2.4)$$

$k_1(t) = t - \frac{1}{2}$, $k_2(t) = \frac{1}{2}\{k_1^2(t) - \frac{1}{12}\}$, and $k_4(t) = \frac{1}{24}\{k_1^4(t) - \frac{1}{2}k_1^2(t) + \frac{7}{240}\}$.

In the applications, usually only low order interaction terms in the decomposition (2.2) are retained for easy computation and interpretation. Correspondingly, the space $\otimes_{j=1}^{d} \mathcal{H}_j$ represented in (2.3) is truncated to some proper subspace $\mathcal{F}$. We write $\mathcal{F}$ as

$$\mathcal{F} = \{1\} \oplus_{\alpha=1}^{q} \mathcal{F}^\alpha, \quad (2.5)$$

where $\mathcal{F}^1, ..., \mathcal{F}^q$ are q orthogonal subspaces of $\otimes_{j=1}^{d} \mathcal{H}_j$. The space $\mathcal{F}$ is an RKHS with the induced norm $\|\cdot\|$.

3. The COSSO SVM

3.1. Formulation

We propose a new type of regularization for the SVM, in the framework of smoothing spline ANOVA, by solving

$$\min_{f\in\mathcal{F}} \frac{1}{n}\sum_{i=1}^{n}[1-y_i f(\mathbf{x}_i)]_+ + \tau^2 \sum_{\alpha=1}^{q}\|P^\alpha f\|, \tag{3.1}$$

where $P^\alpha f$ is the projection of f onto the subspace $\mathcal{F}^\alpha$. The parameter τ is a tuning parameter which should be properly chosen, and we will discuss the tuning issue in Section 4. The penalty $\sum_{\alpha=1}^{q}\|P^\alpha f\|$ is a sum of RKHS component norms, different from the squared RKHS norm penalty used in the standard SVM. Lin and Zhang (2002) proposed and studied this type of regularization in the penalized likelihood regression setting. Two special cases of (3.1) will be considered in this paper. Assume $\mathcal{F} = \{1\} \oplus_{j=1}^{d} \bar{\mathcal{H}}_j$, then we have the additive model:

$$\min_{f\in\mathcal{F}} \quad \frac{1}{n}\sum_{i=1}^{n}[1-y_i f(\mathbf{x}_i)]_+ + \tau^2 \sum_{j=1}^{d}\|f_j\|, \quad \text{where} \quad f(\mathbf{x}) = b + \sum_{j=1}^{d} f_j(x^{(j)}).$$

Each $\bar{\mathcal{H}}_j$ is the Sobolev space $W_2[0,1]$ associated with the reproducing kernel R given in (2.4). In this case, the selection of main effect components is equivalent to variable selection. The other important case is the two-way interaction model which includes all the main effects and the second-order interaction effects:

$$\min_{f\in\mathcal{F}} \quad \frac{1}{n}\sum_{i=1}^{n}[1-y_i f(\mathbf{x}_i)]_+ + \tau^2 \Big[\sum_{j=1}^{d}\|f_j\| + \sum_{j<k}\|f_{jk}\|\Big],$$

where $f(\mathbf{x}) = b + \sum_{j=1}^{d} f_j(x^{(j)}) + \sum_{j<k} f_{jk}(x^{(j)}, x^{(k)})$. We have $q = d(d+1)/2$. The choice of the additive or two-way interaction form mainly depends on the nature of data. When additive models are not adequate, two-way or higher interaction models should be considered.

In the context of linear classification, the COSSO SVM actually reduces to the 1-norm SVM suggested by Bradley and Mangasarian (1998). The argument is as follows. For the input space $\mathcal{X} = [0,1]^d$, the linear SVM has the separating hyperplane $f(x) = b + \sum_{j=1}^{d} w_j x^{(j)}$. Consider the linear function space $\mathcal{F} = \{1\} \oplus \{x^{(1)} - 1/2\} \oplus \ldots \oplus \{x^{(d)} - 1/2\}$ with the usual L_2 inner product on $\mathcal{F}$: $(f,g) = \int_{\mathcal{X}} fg$. For each

f_j, its RKHS norm penalty becomes $\|f_j\| = (12)^{-1/2}|w_j|$, which is equivalent to the absolute value penalty used in the 1-norm SVM.

Define the penalty functional $J(f) = \sum_{\alpha=1}^{q} \|P^\alpha f\|$. It is straightforward to show that $J(f)$ is convex in f and it defines a pseudo-norm in the space $\mathcal{F}$. The following theorem guarantees the existence of the solution to (3.1).

Theorem 3.1 *Let $\mathcal{F}$ be a reproducing kernel Hilbert space of functions over an input space $\mathcal{X}$. Assume that $\mathcal{F}$ can be decomposed as in (2.5). Then there exists a minimizer of (3.1) in $\mathcal{F}$.*

Though the model space $\mathcal{F}$ is infinite dimensional, the solution to (3.1) can be shown to lie in a finite dimensional subspace of $\mathcal{F}$. This is an important property also satisfied by the standard smoothing spline. We will demonstrate later that, the finite representation of the solution makes it feasible to implement the COSSO SVM in practice. Here is the representer theorem and its proof is similar to that of the smoothing spline (Kimeldorf and Wahba (1971)).

Theorem 3.2 *Let $\hat{f} = \hat{b} + \sum_{\alpha=1}^{q} \hat{f}_\alpha$ be the minimizer of (3.1) with $\hat{f}_\alpha \in \mathcal{F}^\alpha$. Then $\hat{f}_\alpha \in span\{R_\alpha(\mathbf{x}_i, \cdot), i = 1, ..., n\}$, where $R_\alpha(\cdot, \cdot)$ is the reproducing kernel of $\mathcal{F}^\alpha$.*

3.2. Equivalent Optimization Problem

In this section we will derive an equivalent formulation of (3.1) which naturally leads to an iterative algorithm. We introduce a new vector $\boldsymbol{\theta} = (\theta_1, ..., \theta_q)^{\mathrm{T}}$ and consider the optimization problem

$$\min_{f \in \mathcal{F}, \boldsymbol{\theta}} \frac{1}{n} \sum_{i=1}^{n} [1 - y_i f(\mathbf{x}_i)]_+ \quad + \quad \lambda_0 \sum_{\alpha=1}^{q} \theta_\alpha^{-1} \|P^\alpha f\|^2 + \lambda \sum_{\alpha=1}^{q} \theta_\alpha,$$
$$\text{subject to} \quad \theta_\alpha \geq 0, \quad \alpha = 1, ..., q. \tag{3.2}$$

Note there is only one tuning parameter τ in (3.1) while there are two parameters (λ_0, λ) in (3.2). In fact, $\lambda > 0$ is the real tuning parameter while $\lambda_0 > 0$ is a constant that can be fixed at any positive value. In Section 4, we will explain that the redundancy of λ_0 is intentional for computational convenience. We have the following theorem.

Theorem 3.3 *Set $\lambda = \tau^4/(4\lambda_0)$. (i) If $\hat{f}$ minimizes (3.1), setting $\hat{\theta}_\alpha = \lambda_0^{1/2} \lambda^{-1/2} \|P^\alpha \hat{f}\|$*

for $\alpha = 1, ..., q$, then the pair $(\hat{\boldsymbol{\theta}}, \hat{f})$ minimizes (3.2). (ii) On the other hand, if a pair $(\hat{\boldsymbol{\theta}}, \hat{f})$ minimizes (3.2), then $\hat{f}$ minimizes (3.1).

Theorem 3.3 states that, with proper choice of parameters, solving (3.1) and solving (3.2) always give the same optimal classifier $\hat{f}$. In practice, we choose to solve (3.2) since its objective function can be easily handled by standard quadratic programming and linear programming techniques.

The non-negative θ_α's can be regarded as scaling parameters and they are interpretable for the purpose of variable selection. Using the standard smoothing spline results, we can show that the solution to (3.2) has the following form

$$\hat{f}(\mathbf{x}) = b + \sum_{\alpha=1}^{q} \hat{\theta}_\alpha \sum_{i=1}^{n} \hat{c}_i R_\alpha(\mathbf{x}_i, \mathbf{x}) = b + \sum_{\alpha=1}^{q} \hat{\theta}_\alpha \hat{f}_\alpha(\mathbf{x}). \tag{3.3}$$

The expression in (3.3) suggests that whether $\hat{\theta}_\alpha = 0$ or not directly determines the appearance of the αth component of the classification function. For the additive model, if $\hat{\theta}_j = 0$, the minimizer is then taken to satisfy $\|\hat{f}_j\| = 0$, implying that the variable X_j is not selected. In this paper we use the convention 0/0=0.

4. Algorithm

Given any $\boldsymbol{\theta}$, we note that solving (3.2) is equivalent to solving the standard SVM problem (2.1) with the reproducing kernel $R_\theta = \sum_{\alpha=1}^{q} \theta_\alpha R_\alpha$, where R_α is the reproducing kernel of $\mathcal{F}^\alpha$. Lin, Wahba, Zhang, and Lee (2002) showed that the solution f is given by

$$f(\mathbf{x}) = b + \sum_{i=1}^{n} c_i R_\theta(\mathbf{x}_i, \mathbf{x}).$$

Define $\mathbf{c} = (c_1, ..., c_n)^{\mathrm{T}}$, $\mathbf{f} = (f(\mathbf{x}_1), ..., f(\mathbf{x}_n))^{\mathrm{T}}$, $\mathbf{y} = (y_1, ..., y_n)^{\mathrm{T}}$, and the diagonal matrix $Y = \mathrm{diag}[y_1, ..., y_n]$. Let $\mathbf{1}_n$ and $\mathbf{0}_n$ be the column vector of ones and zeros with length n, and I_n be the identity matrix of dimension n. With some abuse of notations, we also use R_α for the $n \times n$ matrix $\{R_\alpha(\mathbf{x}_i, \mathbf{x}_j)\}, i = 1, ..., n, j = 1, ..., n$, and use R_θ for the matrix $\sum_{\alpha=1}^{q} \theta_\alpha R_\alpha$. Then we have $\mathbf{f} = R_\theta \mathbf{c} + b\mathbf{1}_n$, and (3.2) becomes

$$\min_{\boldsymbol{\theta} > \mathbf{0}, b, \mathbf{c}} \frac{1}{n} \sum_{i=1}^{n} [1 - y_i f(\mathbf{x}_i)]_+ + \lambda_0 \sum_{\alpha=1}^{q} \theta_\alpha \mathbf{c}^{\mathrm{T}} R_\alpha \mathbf{c} + \lambda \sum_{\alpha=1}^{q} \theta_\alpha. \tag{4.1}$$

4.1. Quadratic and Linear Programming

It is possible to minimize the objective function in (4.1) with respect to $\boldsymbol{\theta}$ and $(b, \boldsymbol{c})$

simultaneously. We propose alternatively solving $(b, \mathbf{c})$ with $\boldsymbol{\theta}$ fixed and solving $\boldsymbol{\theta}$ with $(b, \mathbf{c})$ fixed, since both problems can be solved using standard software easily.

When $\boldsymbol{\theta}$ is fixed, we need to solve the SVM associated with the kernel R_θ:

$$\min_{b,\mathbf{c}} \quad \frac{1}{n}\sum_{i=1}^{n}[1 - y_i f(\mathbf{x}_i)]_+ + \lambda_0 \mathbf{c}^{\mathrm{T}} R_\theta \mathbf{c} \tag{4.2}$$

Typically the dual problem of (4.2) is solved using the quadratic programming problem (QP). Introducing the dual variables $\boldsymbol{a} = (a_1, ..., a_n)^{\mathrm{T}}$ and the matrix $H = \frac{1}{2n\lambda_0} Y R_\theta Y$, the dual problem is

$$\max L = -\frac{1}{2}\boldsymbol{a}^{\mathrm{T}} H \boldsymbol{a}, \quad \text{subject to} \quad \mathbf{y}^{\mathrm{T}}\boldsymbol{a} = 0, \quad \mathbf{0}_n \le \boldsymbol{a} \le \mathbf{1}_n.$$

From the dual solution, we compute $\mathbf{c} = \frac{1}{2n\lambda_0} Y \boldsymbol{a}$. Define $A = \mathrm{diag}[a_1, ..., a_n]$. The constant b is derived using the Karash-Kuhn-Tucker optimality condition as $b = [\mathbf{1}_n^{\mathrm{T}} A(I_n - A)(\mathbf{y} - R_\theta \mathbf{c})]/[\boldsymbol{a}^{\mathrm{T}}(\mathbf{1}_n - \boldsymbol{a})]$,

When $(b, \mathbf{c})$ is fixed, we need to solve the linear programming (LP) under linear inequality constraints. Let $\mathbf{g}_\alpha = R_\alpha \mathbf{c}$ and G be the matrix with the αth column as $\mathbf{g}_\alpha$, $\alpha = 1, ..., q$. The objective function in (4.1) becomes $\frac{1}{n}\sum_{i=1}^{n}\left[1 - y_i f(\mathbf{x}_i)\right]_+ + \lambda_0 \mathbf{c}^{\mathrm{T}} G\boldsymbol{\theta} + \lambda \boldsymbol{\theta}^{\mathrm{T}} \mathbf{1}_q$. Using slack variables $\mathbf{z} = (z_1, ..., z_n)^{\mathrm{T}}$, the hinge loss function can be replaced by $\mathbf{z}^{\mathrm{T}}\mathbf{1}_n$ plus two constraints $\mathbf{z} \ge \mathbf{0}_n$ and $\mathbf{z} \ge \mathbf{1} - Y(G\boldsymbol{\theta} + b\mathbf{1}_n)$. In addition, the term $\lambda \boldsymbol{\theta}^{\mathrm{T}}\mathbf{1}_q$ can be changed into the constraint $\boldsymbol{\theta}^{\mathrm{T}}\mathbf{1}_q \le M$. The parameter M is a tuning parameter which replaces λ, and there is one-to-one corresponding relationship between them. The final optimization problem is

$$\begin{aligned} \min_{\mathbf{z},\boldsymbol{\theta}} \quad & \frac{1}{n}\mathbf{z}^{\mathrm{T}}\mathbf{1}_n + \lambda_0 \mathbf{c}^{\mathrm{T}} G\boldsymbol{\theta} \\ \text{subject to} \quad & \mathbf{z} + \boldsymbol{\theta}^{\mathrm{T}}(YG) \ge (I - Yb)\mathbf{1}_n, \quad \mathbf{z} \ge \mathbf{0}_n, \quad \boldsymbol{\theta}^{\mathrm{T}}\mathbf{1}_q \le M, \quad \boldsymbol{\theta} \ge \mathbf{0}_q. \end{aligned} \tag{4.3}$$

This is a linear optimization problem with polyhedral constraints. Popular algorithms for solving the LP include the simplex method and the interior-point method. Many optimization packages are in wide use as well, such as CPLEX, MATLAB, GAMS, and MINOS. In our implementations, we used the MATLAB optimization toolbox. The following algorithm is proposed to solve the COSSO SVM:

1. Initialization: $\boldsymbol{\theta} = \mathbf{1}_q$;

2. With $\boldsymbol{\theta}$ fixed at current values, solve the dual problem of (4.2) for $(b, \mathbf{c})$;

3. With $(b, \mathbf{c})$ fixed at current values, solve (4.3) for $\boldsymbol{\theta}$;

4. With the new $\boldsymbol{\theta}$, go to step 2 until the convergence.

When R_θ is strictly positive definite, this algorithm is guaranteed to converge because the objective function in (4.1) is bounded below by zero and each iteration between step 2 and step 3 always results in improved updates. It is well-known that the LP is solvable in polynomial time, say, the algorithm given by Anstreicher (1999) has the computational complexity $O(n^3/\log(n)L)$ where L is the bit length of input variables. Since the complexity of the QP is $O(n^3)$, the overall computation for one iteration is cubic in time. Numerical studies show that one-step update was practically sufficient to give good approximate solutions.

4.2. Parameter Tuning

Smoothing parameters balance the tradeoff between the hinge loss fit and the penalty on the functional components. The choice of parameters is typically done by minimizing either an estimate of generalization error or other related performance measure. We consider minimizing the generalized comparative Kullback-Liebler (GCKL) distance proposed in Wahba (1999). Given a fitted classifier f_λ, the GCKL is defined as:

$$\begin{aligned} GCKL(\lambda) &= E_p\big[\ \frac{1}{n}\sum_{i=1}^{n}(1 - Y_i f_{\lambda i})_+\big] \\ &= \frac{1}{n}\sum_{i=1}^{n}\ \big[p_i(1 - f_{\lambda i})_+ + (1 - p_i)(1 + f_{\lambda i})_+\big]. \end{aligned} \tag{4.4}$$

Here f_λ is fixed and the expectation is taken over the true conditional probability $p(\mathbf{x}) = P(Y = +1|\mathbf{x} = \mathbf{x})$. The GCKL can be seen as an upper bound of the misclassification rate. Since the GCKL depends on the underlying distribution $P(\mathbf{X}, Y)$, it is only computable in simulations. For real data with unknown $p(\mathbf{x})$, the leaving-out-one cross validation proxy of GCKL $\frac{1}{n}\sum_{i=1}^{n}[1 - y_i f_\lambda^{[-i]}(\mathbf{x}_i)]_+$ can be used as a tuning criterion. Here $f_\lambda^{[-i]}$ is the solution with the ith data point deleted. In practice, we suggest using the five-fold cross validation (CV) estimate of the GCKL. The training set is randomly split into five subsets of approximately equal sizes. Then one subset is left out, and the COSSO SVM is fitted using the other four subsets and then the hinge loss is evaluated on the left-out subset. This procedure is repeated five times in this fashion with each subset being left out once. For the COSSO SVM, we need

to tune λ, or equivalently M in (4.3). The parameter λ_0 seems redundant in addition to M, however the proper choice of λ_0 in (4.2) helps to solve the SVM in step 2 more stably. In practice, we suggest tuning λ_0 once when step 2 is executed for the first time and fix it thereafter.

5. Numerical Examples

In this section we study the empirical performances of the COSSO SVM through simulated examples and some real data sets. The fitted classifier is evaluated in both its classification and variable selection performances. We simulate the examples for both the additive model and two-way interaction model. In each experiment setting, we generate 100 data sets for fitting and tuning, and an extra test set of $10,000$ points to compute the expected misclassification rate (EMR) of the classifier. We summarize the average EMR, model size, the frequency of each variable appearing in the COSSO SVM classifier over the 100 runs.

5.1. Example 1: additive model

Consider an additive model with $\mathcal{X} = [0,1]^{10}$. We generate $\mathbf{X}$ uniformly and the binary response Y with the conditional logit function

$$f(\mathbf{x}) = 3x^{(1)} + \pi \sin(\pi x^{(2)}) + 8(x^{(3)})^5 + \frac{2}{e-1}e^{x^{(4)}} - 6.$$

Therefore $X^{(5)}, ..., X^{(10)}$ are uninformative in this example. The tuning parameter M was tuned by the GCKL. The Bayes misclassification rate is 0.216, which is the optimal rate based on the true p. Consider two settings: $n = 250$ and $n = 500$. In Table 1 we report the average EMR and model size of the additive COSSO SVM in 100 runs. The values in the parentheses are the standard errors of the corresponding mean values. Table 2 shows the appearance frequency of each variable in the final model. When $n = 250$, the COSSO SVM always selects $X^{(1)}, X^{(2)}, X^{(3)}$, and selects $X^{(4)}$ in 98 runs. Sometimes some of uninformative variables are selected as well. When the sample size increases to 500, the COSSO SVM never misses any important variable, and it selects unimportant variables with a much lower frequency. The left plot in Figure 5.1 shows that the COSSO SVM gives the correct true model in 74 of 100 runs.

Table 1: The average EMR and model size of the additive COSSO SVM classifier.

n	EMR	Model Size
250	0.234 (0.009)	5.07
500	0.225 (0.006)	4.46

Table 2: The appearance frequency of the input variables in the COSSO SVM classifiers.

n	X_1	X_2	X_3	X_4	X_5	X_6	X_7	X_8	X_9	X_{10}
250	100	100	100	98	14	18	18	17	23	19
500	100	100	100	100	9	12	10	8	2	5

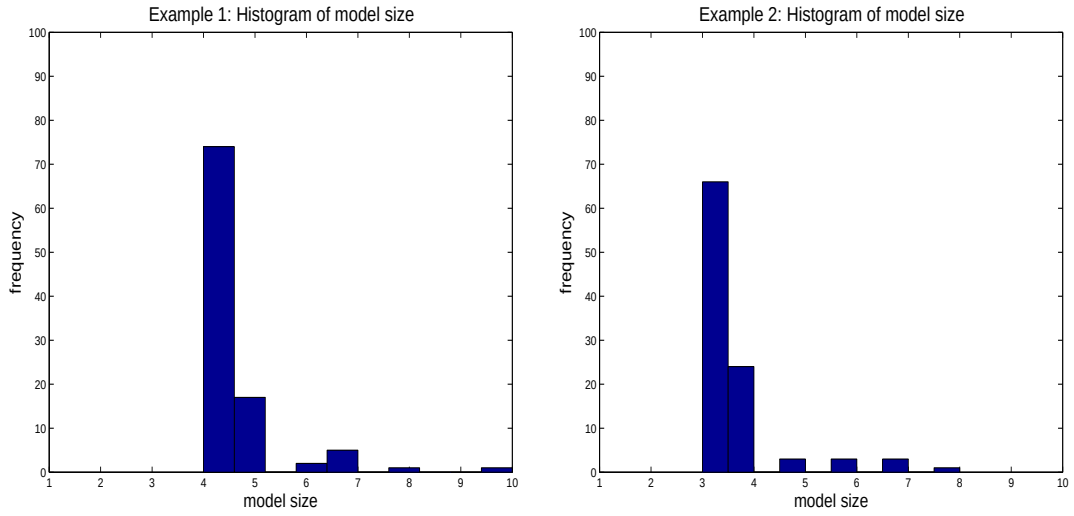

Table 5.1: The histogram of the model size given by the COSSO SVM in the 100 runs when $n = 500$. The plot on the left is for the additive model, and the plot on the right is for the two-way interaction model.

5.2. Example 2: two-way interaction model

In this example, we generate four input variables independently from Unif$[0, 1]$. The true logit function contains the important main effects $X^{(1)}, X^{(2)}$ and their interaction:

$$f(\mathbf{x}) = 4x^{(1)} + \pi \sin\left(\pi x^{(1)}\right) + 6x^{(2)} - 8\left(x^{(2)}\right)^3 + 3\cos\left(2\pi(x^{(1)} - x^{(2)})\right) - 5.$$

The Bayes error is 0.155. We fit the two-way interaction model in two settings $n = 300$ and $n = 500$. Table 3 shows the COSSO SVM never misses any important main or interaction term in the 100 runs under each setting. When n increases from 300 to 500, the COSSO SVM selects the correct model size more precisely as shown in Table 4. The distribution of the model size in 100 runs is depicted in Figure 5.1 (right plot), showing that the correct model is chosen by the COSSO SVM in 66 of 100 runs.

Table 3: The average EMR and model size of the two-way interaction COSSO SVM classifier.

n	EMR	Model Size
300	0.198 (0.016)	4.56 (1.73)
500	0.182 (0.010)	3.56 (1.03)

Table 4: The appearance frequency of the input variables in the COSSO SVM classifiers.

n	X_1	X_2	X_3	X_4	X_1, X_2	X_1, X_3	X_1, X_4	X_2, X_3	X_2, X_4	X_3, X_4
300	100	100	23	20	100	22	24	28	24	15
500	100	100	6	5	100	13	8	7	5	12

5.3. Real Data

Gestel, Suykens, Baesens, Viaene, Vantheienen, Dedene, Moor, and Vandewalle (2004) conducted a benchmark study comparing a number of commonly used machine learning techniques including the SVM; least squares SVM (LS-SVM); linear discriminant analysis (LDA); quadratic discriminant analysis (QDA); logistic regression (Logit); the decision tree algorithm C4.5; Holte's one-rule classifier (oneR); instance based learners (IB); and Naive Bayes method. They found out the least squares SVM (LS-SVM) with the radial basis function (RBF) kernel performs best among six types of LS-SVMs. Thus we only include the LS-SVM with RBF kernel and linear kernel for our comparison. They considered two instance based learners (IB1 and IB10) and two types of Naive Bayes methods, and we only report the better performance of the two. There are five binary classification datasets with continuous predictors in their study, and we test the performance of the COSSO SVM on these datasets. The results for the other algorithms are taken from their paper.

The datasets are the BUPA Liver Disorder data, the Johns Hopkins University Ionosphere data; the PIMA Indian Diabetes; the Sonar, Mines vs. Rocks data; and

the Wisconsin Breast Cancer data. The basic features of the datasets and the performances of different algorithms are summarized in Table 5. Following Gestel, Suykens, Baesens, Viaene, Vantheienen, Dedene, Moor, and Vandewalle (2004), for each dataset we randomly select 2/3 of the data for training and tuning, and test on the remaining 1/3 of the data. We do this randomization 10 times and report the average test set performance and sample standard deviation for the COSSO SVM. The best average test set performances are denoted in bold face for each dataset in Table 5. The additive COSSO SVM is fitted and its performances on these benchmark datasets are very competitive to other algorithms.

Table 5: Comparison of the test set performance of the COSSO SVM with those of SVM, LS-SVM, LDA, QDA, Logit, C4.5, oneR, IB, Naive Bayes, and the Majority Rule. The results of the other algorithms are taken from the paper Gestel et. al. (2004).

	BUPA	Ionosphere	Pima Indian	Sonar MR	Wisc. BC
n	345	351	768	208	683
d	6	33	8	60	9
COSSO SVM	**72.0** (5.0)	89.6 (2.6)	**77.3** (2.3)	**78.6** (2.6)	95.8 (1.2)
SVM (linear)	67.7 (2.6)	87.1 (3.4)	77.0 (2.4)	74.1 (4.2)	96.3 (1.0)
SVM (RBF)	70.4 (3.2)	95.4 (1.7)	**77.3** (2.2)	75.0 (6.6)	96.4 (1.0)
LS-SVM (linear)	65.6 (3.2)	87.9 (2.0)	76.8 (1.8)	72.6 (3.7)	95.8 (1.0)
LS-SVM (RBF)	70.2 (4.1)	**96.0** (2.1)	76.8 (1.7)	73.1 (4.2)	96.4 (1.0)
LDA	65.4 (3.2)	87.1 (2.3)	76.7 (2.0)	67.9 (4.9)	95.6 (1.1)
QDA	62.2 (3.6)	90.6 (2.2)	74.2 (3.3)	53.6 (7.4)	94.5 (0.6)
Logit	66.3 (3.1)	86.2 (3.5)	77.2 (1.8)	68.4 (5.2)	96.1 (1.0)
C4.5	63.1 (3.8)	90.6 (2.2)	73.5 (3.0)	72.1 (2.5)	94.7 (1.0)
oneR	56.3 (4.4)	83.6 (4.8)	71.3 (2.7)	62.6 (5.5)	91.8 (1.4)
IB	61.3 (6.2)	87.2 (2.8)	73.6 (2.4)	77.7 (4.4)	96.4 (1.2)
Naive Bayes	63.7 (4.5)	92.1 (2.5)	75.5 (1.7)	71.6 (3.5)	**97.1** (0.9)
Majority Rule	56.5 (3.1)	64.4 (2.9)	66.8 (2.1)	54.4 (4.7)	66.2 (2.4)

The number of interaction terms in the two-way interaction model is $d(d-1)/2$, which can be very large even for a moderate d. For example, in the Sonar, Mine, Rock data there are 60 variables and the full two-way ANOVA model has $1,770$ interaction terms. This can cause great difficulty in both model fitting and interpretation. Therefore, when d is large, additive models are often preferred in practical situations, and maybe sufficient. We also fitted the two-way interaction models for three data sets

having fewer than 10 variables (BUPA, Pima Indian, and Wisconsin BC), and they gave similar classification performances as additive models but take much longer to fit. For example, for the BUPA data, the average accuracy of the two-way interaction model is 73.0% and that of the additive model is 72.0%.

6. Discussion

The COSSO SVM is attractive in its compact mathematical formulation and nice solution properties. The novel regularization setup naturally combines smoothing and shrinkage-type operations on the ANOVA components of the classifier. In addition, the COSSO SVM includes the 1-norm SVM as a special case. Numerical studies demonstrate its desirable performances when compared with other classification schemes.

The proposed idea provides a general framework for variable selection in the SVM. We focus on the two-class classification in this paper, however, the idea can be used to multi-class classification problems. See the technical report of Lee, Kim, Lee, and Koo (2004). In addition, it is straightforward to generalize the COSSO SVM to nonstandard classification situations where: (i) different costs are used for different types of misclassification; (ii) the proportions of two classes in samples do not represent those in populations. Let the false positive and the false negative cost be c^+ and c^- respectively, the proportions of two classes in populations be π_0^+ and π_0^-, and those in samples be π^+ and π^-. Following Lin, Lee, and Wahba (2002), we define the weight w function on the label by $w(+1) = c^-\pi^-\pi_0^+$ and $w(-1) = c^+\pi^+\pi_0^-$. The non-standard COSSO SVM can be proposed as

$$\min_f \quad \frac{1}{n}\sum_{i=1}^{n} w(y_i)[1 - y_i f(\mathbf{x}_i)]_+ + \tau^2 \sum_{\alpha=1}^{q} \|P^\alpha f\|. \tag{6.1}$$

The algorithm suggested in Section 4 can be adapted to solve this problem.

For high dimension low sample size data, or, $d \gg n$, linear classifiers often give better performances than nonlinear ones in many applications (Hastie, Tibshirani, and Friedman 2001). This fact is related to the asymptotic results in Hall and Marron (2004): when $d \to \infty$ with n fixed, all the pairwise distances between any two points are asymptotically identical to each other, so all the points form an n-simplex. Linear classifiers are natural choices to discriminate two simplices. In those situations, the L_1-norm SVM (Bradley and Mangasarian 1998; Zhu, Hastie, Rosset, and Tibshirani 2003) may be sufficient for classification and variable selection.

Appendix 1
Proof of Solution Existence

Proof of Theorem 3.1. Denote the functional to be minimized in (3.1) by

$$A(f) = \sum_{i=1}^{n}[1 - y_i f(\mathbf{x}_i)]_+ + \tau^2 J(f).$$

Then $A(f)$ is convex and continuous. And we have the following relationship

$$\sum_{\alpha=1}^{q} \|P^\alpha f\|^2 \le J^2(f) \le q \sum_{\alpha=1}^{q} \|P^\alpha f\|^2. \tag{A1}$$

Without loss of generality, we assume $\tau = 1$.

Define $\mathcal{F}_1 = \oplus_{\alpha=1}^{q}\mathcal{F}^\alpha$. By (A1) we have that $J(f_1) \ge \|f_1\|$ for any $f_1 \in \mathcal{F}_1$. Let $R_{\mathcal{F}_1}$ be the reproducing kernel of $\mathcal{F}_1$ and $\langle\cdot,\cdot\rangle_{\mathcal{F}_1}$ be the inner product in $\mathcal{F}_1$. Denote $a = \max_{i=1}^{n} R_{\mathcal{F}_1}^{1/2}(\mathbf{x}_i, \mathbf{x}_i)$. By the reproducing property of the kernel, we have for any $f_1 \in \mathcal{F}_1$ and $i = 1, ..., n$,

$$\begin{aligned} |f_1(\mathbf{x})| &= |\langle f_1(\cdot), R_{\mathcal{F}_1}(\mathbf{x}_i, \cdot)\rangle_{\mathcal{F}_1}| \le \|f_1\|\langle R_{\mathcal{F}_1}(\mathbf{x}_i, \cdot), R_{\mathcal{F}_1}(\mathbf{x}_i, \cdot)\rangle_{\mathcal{F}_1}^{1/2} \\ &= \|f_1\| R_{\mathcal{F}_1}^{1/2}(\mathbf{x}_i, \mathbf{x}_i) \le a\|f_1\| \le aJ(f_1). \end{aligned} \tag{A2}$$

Let n^+ and n^- be respectively the number of sample points from $+1$ class and -1 class in the data. Define $\rho = \min\{2n^+/n, 2n^-/n\}$. Consider the set

$$D = \{f \in \mathcal{F} : f = b + f_1, \text{with } \ b \in \{1\}, f_1 \in \mathcal{F}_1, J(f) \le \rho, |b| \le 1 + a\}.$$

Then D is a closed, convex, and bounded set. By Theorem 4 of Tapia and Thompson (1978, page 162), there exists a minimizer of (3.1) in D. Let the minimizer be $\bar{f}$. Direct calculation gives us

$$\sum_{i=1}^{n}[1 - y_i f(\mathbf{x}_i)]_+ = \sum_{y_i=+1}[1 - f(\mathbf{x}_i)]_+ + \sum_{y_i=-1}[1 + f(\mathbf{x}_i)]_+,$$

hence $A(+1) = 2n^-/n$ for the function $f(\mathbf{x}) \equiv +1$ and $A(-1) = 2n^+/n$ for the function $f(\mathbf{x}) \equiv -1$. Since the constant functions $+1$ and -1 are both in D, we must have $A(\bar{f}) < \min\{A(+1), A(-1)\} = \rho$.

On the other hand, for any $f \notin D$, one of the following must happen:

(i) When $J(f) > \rho$, we have $A(f) \ge J(f) > \rho$.

(ii) When $J(f) \le \rho$, $f = b + f_1$, $f_1 \in \mathcal{F}$ and $b > 1 + a$, we use (A2) to get that, for any $i = 1, ..., n$, $b + f_1(\mathbf{x}_i) \ge b - a > 1$ and

$$\sum_{y_i=+1} [1 - b_1 - f_1(\mathbf{x}_i)]_+ + \sum_{y_i=-1} [1 + b_1 + f(\mathbf{x}_i)]_+ \ge \sum_{y_i=-1} [1 + b_1 + f(\mathbf{x}_i)]_+ > 2n^-.$$

We then have $A(f) > 2n^-/n$.

(iii) When $J(f) \le \rho$, $f = b + f_1$, $f_1 \in \mathcal{F}$ and $b < -1 - a$, we use (A2) to get that, for any $i = 1, ..., n$, $b + f_1(\mathbf{x}_i) \le b + a < -1$ and

$$\sum_{y_i=-1} [1 - b_1 - f_1(\mathbf{x}_i)]_+ + \sum_{y_i=-1} [1 + b_1 + f(\mathbf{x}_i)]_+ \ge \sum_{y_i=+1} [1 - b_1 - f(\mathbf{x}_i)]_+ > 2n^+.$$

We then have $A(f) > 2n^+/n$.

Hence for any $f \notin D$, we have $A(f) > A(\bar{f})$. Therefore $\bar{f}$ is a minimizer of (3.1) in $\mathcal{F}$.

Appendix 2

Proof of Representer Theorem

Proof of Theorem 3.2. For any $f \in \mathcal{F}$, we can write $f = b + \sum_{\alpha=1}^q f_\alpha$ with $f_\alpha \in \mathcal{F}^\alpha$. Let the projection of f_α onto $\text{span}\{R_\alpha(\mathbf{x}_i, \cdot), i = 1, ..., n\} \subset \mathcal{F}^\alpha$ be denoted by g_α, and its orthogonal complement by h_α. Then $f_\alpha = g_\alpha + h_\alpha$, and $\|f_\alpha\|^2 = \|g_\alpha\|^2 + \|h_\alpha\|^2$, $\alpha = 1, ..., q$. Since the reproducing kernel of $\mathcal{F}$ is $1 + \sum_{\alpha=1}^q R_\alpha$ is, we have

$$f(\mathbf{x}_i) = \langle 1 + \sum_{\alpha=1}^q R_\alpha(\mathbf{x}_i, \cdot), b + \sum_{\alpha=1}^q (g_\alpha + h_\alpha) \rangle = b + \sum_{\alpha=1}^q \langle R_\alpha(\mathbf{x}_i, \cdot), g_\alpha \rangle,$$

where $\langle \cdot, \cdot \rangle$ is the inner product in $\mathcal{F}$. Therefore (3.1) can be written as

$$\frac{1}{n} \sum_{i=1}^n \left[1 - y_i(b + \sum_{\alpha=1}^q \langle R_\alpha(\mathbf{x}_i, \cdot), g_\alpha \rangle)\right]_+ + \tau^2 \sum_{\alpha=1}^q (\|g_\alpha\|^2 + \|h_\alpha\|^2)^{1/2}.$$

Therefore any minimizer f satisfies $h_\alpha = 0$, $\alpha = 1, ..., q$. The theorem is proved.

Proof of Theorem 3.3. Denote the functional in (3.1) by $A(f)$, and the functional in (3.2) by $N(\theta, f)$. For any $\theta_\alpha \ge 0, f \in \mathcal{F}$, we have

$$\lambda_0 \theta_\alpha^{-1} \|P^\alpha f\|^2 + \lambda \theta_\alpha \ge 2\lambda_0^{1/2} \lambda^{1/2} \|P^\alpha f\| = \tau^2 \|P^\alpha f\|,$$

and the equality holds if and only if $\theta_\alpha = \lambda_0^{1/2} \lambda^{-1/2} \|P^\alpha f\|$. Therefore $N(\boldsymbol{\theta}, f) \ge A(f)$ for any $\theta_\alpha \ge 0$, $\alpha = 1, ..., q$, and $f \in \mathcal{F}$, and the equality holds if and only if $\theta_\alpha = \lambda_0^{1/2} \lambda^{-1/2} \|P^\alpha f\|$, $\alpha = 1, ..., q$.

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