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Non-crossing large-margin probability estimation and its application to robust SVM via preconditioning

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Abstract

Many large-margin classifiers such as the Support Vector Machine (SVM) sidestep estimating conditional class probabilities and target the discovery of classification boundaries directly. However, estimation of conditional class probabilities can be useful in many applications. Wang, Shen, and Liu (2008) bridged the gap by providing an interval estimator of the conditional class probability via bracketing. The interval estimator was achieved by applying different weights to positive and negative classes and training the corresponding weighted large-margin classifiers. They propose to estimate the weighted large-margin classifiers individually. However, empirically the individually estimated classification boundaries may suffer from crossing each other even though, theoretically, they should not.

In this work, we propose a technique to ensure non-crossing of the estimated classification boundaries. Furthermore, we take advantage of the estimated conditional class probabilities to precondition our training data. The standard SVM is then applied to the preconditioned training data to achieve robustness. Simulations and real data are used to illustrate their finite sample performance.

Key Words and Phrases

Conditional class probability; large-margin classifier; preconditioning; SVM

1 Introduction

Binary classification or discrimination handles data with a binary response, where we are interested in studying the relationship between a *p*-vector predictor *x* and a binary response *y* ∈ {−1, +1}. In particular we seek a classification rule to predict the binary response for an arbitrary input predictor. A well-known traditional likelihood-based statistical approach is the logistic regression, which tries to estimate the conditional probability $p(x) = P(Y = 1 | X = 1)$ *x*) and make prediction $\hat{y} = 1$ if $\hat{p}(x) > 1/2$ and -1 otherwise for any *x*. See McCullagh and Nelder (1989) for a detailed introduction to logistic regression.

Recently many margin-based classifiers have been proposed in the literature. Examples include the support vector machine (SVM) (Vapnik, 1998; Cortes and Vapnik, 1995), Boosting (Freund and Schapire, 1997; Friedman, Hastie, and Tibshirani, 2000), *ψ*-learning (Shen, Tseng, Zhang, and Wong, 2003), etc. By sidestepping the conditional class probability estimation and targeting directly at classification, many margin-based classification methods may yield good classification performance. However, one main drawback of some margin-based classifiers such as the SVM is that they are not able to

To provide class probability estimation for large-margin classifiers, Wang et al. (2008) proposed an interval estimation method to estimate the conditional class probability. By noting that ${x : \hat{f}(x) = 0}$ estimates the Bayes boundary ${x : p(x) - 1/2 = 0}$ for a marginbased classifier sign($\hat{f}(x)$), they propose to run different weighted margin-based classification individually to estimate Bayes boundaries $\{x : p(x) - p = 0\}$ for a prespecified grid of *p*. Each weighted classifier can estimate whether $p(x) > p$ or not without estimating $p(x)$ itself. In this way, an interval estimator is obtained by bracketing the conditional class probability $p(x)$. Their approach is nonparametric in the sense that it does not make any parametric assumption on $p(x)$.

Theoretically Bayes boundaries $\{x : p(x) - p = 0\}$ for different *p* should not cross each other. To bracket the conditional class probability, we estimate them using the classification boundaries of the weighted margin-based classifiers based on a finite training data set. Thus the individually estimated boundaries may unavoidably cross each other, especially when the training sample size is small. This crossing phenomenon was also pointed out by Wang et al. (2008). Their probability bracketing method cannot avoid crossing. In this work, we address this crossing issue and provide a technique to enforce non-crossing of the estimated classification boundaries. With this technique, we can ensure non-crossing of the estimated boundaries, at least over the convex hull of the training data points, where the training data convex hull represents the smallest convex hull in the input space which contains all the training data points.

As mentioned earlier, the estimated conditional class probability can provide us with some confidence of our prediction for any x . Our method yields an estimate of $p(x_i)$ for each training dataset point (x_i, y_i) . For a clean training data with a low noise level, we generally expect a training observation with class probability estimate bigger (smaller) than 0.5 to have class label $y_i = +1$ ($y_i = -1$, respectively), especially so for those with probabilities close to 1 or 0. A training point with large (small) probability estimate and class label $y_i =$ -1 ($y_i = +1$, respectively) is typically an outlier. It is known that such outliers may pull the classification boundaries towards themselves and as a result can reduce the classification performance (Wu and Liu, 2007). In order to improve classification performance in such cases, we propose to precondition the training data set based on the estimated conditional class probabilities; i.e., we change the class labels of potential outliers to the corresponding correct class labels. After preconditioning the training set, any standard margin-based classifier may be applied to obtain a robust classification rule.

In the literature there are some other robust margin-based classifiers such as *ψ*-learning (Shen et al., 2003 and Liu and Shen, 2006) and the robust truncated-hinge-loss SVM (RSVM) (Wu and Liu, 2007). These robust methods typically involve non-convex optimization, which is more expensive to solve than convex minimization. Our preconditioning method serves as another technique to achieve robustness. Compared to *ψ*learning and the RSVM, our preconditioned SVM can avoid non-convex optimization.

The rest of the article is organized as follows. Some background on the SVM and weighted SVM is given in Section 2. Section 3 presents our technique which provides non-crossing probability estimation. In Section 4, we propose to robustify margin-based classifiers via preconditioning. Simulation and real data analysis are reported in Sections 5 and 6.

2 Standard and weighted SVMs

For binary classification, a training data set $\{(x_i, y_i) : i = 1, 2, \dots, n\}$ is typically given where *n* is the sample size, $x_i \in R^p$ is the *p*-dimensional input vector, and $y_i \in \{-1, +1\}$ is a binary response. The goal is to construct a function $f(\cdot)$ whose sign will be used as the classification rule, namely $\hat{y} = sign(f(x))$ for any new input *x*.

The SVM was first invented from the idea of searching for the optimal separating hyperplane using an elegant margin interpretation (Vapnik, 1998; Cristianini and Shawe-Taylor, 2000). The idea can be traced back to the method of portraits (Vapnik and Lerner, 1963). It is now known that the SVM is equivalent to solving the following optimization

$$
\min_{f(\cdot)} \left\{ \sum_{i=0}^{n} H(\mathbf{y}_i f(\mathbf{x}_i)) + \lambda J(f) \right\},\tag{1}
$$

where $H(\eta) = \max(0, 1 - \eta)$ is the hinge loss function and $J(f)$ is the roughness penalty of $f(\cdot)$ (Wahba, 1998). Here $y_i f(x_i)$ is the functional margin. A positive functional margin value indicates correct classification and the corresponding $0 - 1$ loss can be expressed as $I(y_i f(x_i))$ > 0 . Besides the SVM, many other classifiers can be fit into the regularization framework. In fact the hinge loss can be replaced by a general large-margin loss $V(u)$ satisfying the condition that $V(u)$ is non-increasing in u , thereby penalizing small margin values more than large margin values. Some common loss functions include the exponential loss $V(u)$ = exp($-u$) (Friedman et al., 2000), the logistic loss $V(u) = \log(1 + \exp(-u))$ (Lin et al., 2000; Zhu and Hastie, 2005), the ψ -loss $V(u) = 1 - sign(u)$ if $u \ge 1$ or $u < 0$, 2(1 – *u*) otherwise (Shen et al., 2003), etc. Figure 1 displays some common large margin loss functions.

Lin (2002) showed that the theoretical minimizer of the hinge loss is $sign(p(x) - 0.5)$. Consequently, the SVM only targets at the Bayes classification boundary asymptotically. As a result, the SVM boundary $\{x : \hat{f}(x) = 0\}$ estimates the Bayes boundary $\{x : p(x) = 0.5\}$, where $\hat{f}(\cdot)$ is the solution to (1). Lin et al. (2002) proposed a weighted hinge loss by giving different weights for the positive and negative classes and solves a weighted version of (1), namely,

$$
\min \left\{ (1 - \pi) \sum_{i: y_i = 1} H(f(x_i)) + \pi \sum_{i: y_i = -1} H(-f(x_i)) + \lambda J(f) \right\}
$$
\n(2)

for some $\pi \in [0, 1]$. It reduces to (1) when $\pi = 1/2$. The weighted SVM (2) was originally proposed to handle the problem of unequal representation of categories in the training sample or unequal costs for positive and negative misclassifications with π and $1 - \pi$ being the known costs for negative and positive classes, respectively. Interestingly, it turns out that the classification boundary of the optimizer $\hat{f}(\cdot)$ of (2) estimates the Bayes boundary {*x* : *p*(*x*) $=$ π .

Motivated by the properties of a weighted loss function, Wang et al. (2008) proposed to solve (2) for many different π 's and, consequently, to estimate the conditional probability $p(x)$ via bracketing the probability. For simplicity, they proposed to solve (2) individually over a grid $0 = \pi_1 < \pi_2 < \cdots < \pi_K = 1$. The corresponding algorithm for estimating the conditional class probability can be summarized as follows.

Wang et al.'s Algorithm (2008):

- **1.** Initialize $\pi_k = (k-1)/(K-1)$ for $k = 1, 2, ..., K$.
- **2.** Train a weighted large-margin classifier for π_k by solving (2) and denote the solution as $\hat{f}_{\pi k}(\mathbf{x})$ for $k = 1, 2, ..., K$.
- **3.** For each *x*, compute $k^* = k^*(x) \triangleq \max\{k: \widehat{f}_{\pi_k}(x) \geq 0\}$ and

 $k_* = k_*(x) \triangleq \min\{k: \widehat{f}_{\pi_k}(x) \leq 0\}$. The estimated conditional class probability is given

by
$$
\widehat{p}(x) = \frac{1}{2}(\pi_{k^*} + \pi_{k_*}).
$$

When the estimated classification boundaries $\{x : \hat{f}_{\pi k}(x) = 0\}$ do not cross each other, $k^*(x)$ and $k_*(x)$ should satisfy $|k^*(x) - k_*(x)| \le 1$ for any x . Unfortunately individually optimized boundaries for different *π^k* 's may cross each other, especially for small training samples. As a result, one may have contradictory probability estimation using multiple individually optimized classification functions. To solve the crossing problem, we propose a technique to enforce non-crossing constraints in this paper. Furthermore, we propose to robustify the SVM via preconditioning by taking advantage of the estimated probabilities.

3 Non-crossing class probability estimation

As discussed in Section 2, the weighted SVM with π and $1 - \pi$ being weights for negative and positive classes targets at the Bayes boundary $\{x : p(x) = \pi\}$. Note that $\{x : p(x) \geq \pi_i\} \subset$ ${x : p(x) \geq \pi_l}$ for $\forall \pi_j > \pi_l$. As a result, classification boundaries with different π 's should not cross each other theoretically. However, the estimated classification boundaries may cross each other if each weighted SVM is calculated separately.

In this section, we present a new technique to ensure that the estimated classification boundaries do not cross each other. We illustrate the non-crossing technique first in the linear case in Section 3.1. Its extension to the nonlinear case via the kernel trick is then discussed in Section 3.2. As mentioned in Section 2, we aim to solve (2) for many different π_k 's ($0 = \pi_1 < \pi_2 < \cdots < \pi_K = 1$) to obtain an estimator for the conditional class probability. Denote by k_0 the index such that π_{k_0} is the closest to 1/2 among all different π_k 's, namely,

$$
|\pi_{k_0} - 1/2| = \min_{1 \le k \le K} |\pi_k - 1/2|
$$

3.1 Linear boundaries

By making the linearity assumption on the classification boundaries, we assume that $f_k(x) =$ $x^Tw_k + b_k$ corresponds to π_k for $k = 1, 2, \dots, K$. For the linear function $f_k(x)$, we choose its roughness penalty to be the 2-norm penalty, namely $J(f_k) = \boldsymbol{w}_k^T \boldsymbol{w}_k$.

Our non-crossing algorithm begins with estimating the classification boundary corresponding to π_{k_0} by solving the following optimization problem

$$
\min\left\{ (1 - \pi_{k_0}) \sum_{i: y_i = 1} [1 - (\boldsymbol{x}_i^T \boldsymbol{w} + b)]_+ + \pi_{k_0} \sum_{i: y_i = -1} [1 + (\boldsymbol{x}_i^T \boldsymbol{w} + b)]_+ + \lambda \sum_{j=1}^p w_j^2 \right\},\tag{3}
$$

where the regularization parameter can be tuned either using a separate independent tuning data set or cross-validation. Denote the corresponding estimator by \hat{w}_{k_0} and \hat{b}_{k_0} .

For any $k > k_0$, we require that the classification boundary corresponding to π_k does not cross that of π_{k-1} within the convex hull of the data points. In particular, the convex hull $H_{convex}(x_1, x_2, \dots, x_n)$ represents the smallest convex set in the input data space which

contains all the training points x_1, x_2, \dots, x_n . Since the region of the training data is a set where the testing data points are likely to fall in, our technique aims to avoid crossing of different boundaries in $H_{convex}(x_1, x_2, \dots, x_n)$. To achieve this goal, we solve the following constrained optimization problem

$$
\min \left\{ (1 - \pi_k) \sum_{i: y_i = 1} [1 - x_i^T \mathbf{w} + b] \mathbf{1}_+ + \pi_k \sum_{i: y_i = -1} [1 + (x_i^T \mathbf{w} + b)]_+ + \lambda \sum_{j=1}^p w_j^2 \right\}, \text{ s.t. } \mathbf{x}^T \mathbf{w} + b \le 0 \text{ for any } \mathbf{x} \in \{\mathbf{x} : \mathbf{x}^T \widehat{\mathbf{w}}_{k-1} + \widehat{\mathbf{b}}_{k-1} + \widehat{\mathbf
$$

(4)

where \hat{w}_{k-1} and \hat{b}_{k-1} represent the solution corresponding to π_{k-1} .

A a remark, we note that a direct way for calculation of the constraint set in (4) is to find the set of intersections of all pairwise line segments between two training data points with the surface. However, the computation can be greatly reduced by the fact that the restriction is satisfied as long as $x^T w + b \le 0$ holds for any x in the intersecion between the surface $x \in$ ${x : x^T \hat{\mathbf{w}}_{k-1} + \hat{b} = 0}$ and the edges of the convex hull $H_{convex}(x_1, x_2, \dots, x_n)$. Furthermore, only a small subset of all pairwise line segments are edges of the convex hull. Thus, we need to find all edges of $H_{convex}(x_1, x_2, \dots, x_n)$ and enumerate the intersection between these edges and the surface $\mathbf{x} \in \{ \mathbf{x} : \mathbf{x}^T \hat{\mathbf{w}}_{k-1} + \hat{b} = 0 \}.$

Notice that any edge of $H_{convex}(x_1, x_2, \dots, x_n)$ can be represented as a segment between two training data points. If $p \le n$, we can first derive all facets of the convex hull and each edge should fall in one or more of the facets. A facet of the convex hull with *p* vertice lies in a hyperplane, such that all training data points fall either on the hyperplane or on the same side of that hyperplane. If $p \ge n$, all data points are vertices and any pairwise line segment between two training points is an edge of $H_{convex}(x_1, x_2, \dots, x_n)$.

Once the points in the intersection in (4) are enumerated, the constrained weighted SVM given in (4) can be implemented using quadratic programming (QP).

Similarly we can estimate the classification boundary corresponding to π_k for $k < k_0$ by solving

$$
\min \left\{ (1 - \pi_k) \sum_{i: y_i = 1} [1 - x_i^T \mathbf{w} + b]_+ + \pi_k \sum_{i: y_i = -1} [1 + (x_i^T \mathbf{w} + b)]_+ + \lambda \sum_{j=1}^p w_j^2 \right\}, \text{ s.t. } \mathbf{x}^T \mathbf{w} + b \ge 0 \text{ for any } \mathbf{x} \in \{\mathbf{x} : \mathbf{x}^T \widehat{\mathbf{w}}_{k+1} + \widehat{b}_{k+1} \}
$$

$$
^{(5)}
$$

Note that we begin with the center π_{k_0} and move to the two ends sequentially to estimate the classification boundaries. By using constraints (4) and (5), it is guaranteed that our estimated classification boundaries do not cross each other within the convex hull of our data. Then Step 3 of Wang et al. (2008)'s algorithm can be applied to estimate the conditional class probability. The main difference of our new algorithm is to enforce non-crossing constraints on the corresponding classification boundaries.

3.2 Nonlinear boundaries

Nonlinear classification can be achieved by using the kernel trick. Denote by $K(\cdot, \cdot)$ a reproducing kernel function. Then by the Representer Theorem, the classification function

can be expressed as $f(x) = \sum_{i=1}^{n} \theta_i K(x_i, x) + \beta$. In this case the roughness penalty is given by (Wahba, 1998). Letting *F* denote the reproducing kernel Hilbert space (RKHS), the weighted SVM (2) becomes

$$
\min_{f \in F} \left\{ (1 - \pi) \sum_{i: y_i = 1} H(f(x_i)) + \pi \sum_{i: y_i = -1} H(-f(x_i)) + \lambda J(f) \right\}.
$$
\n(6)

Once we plug the expressions of $f(x)$ and $J(f)$ into (6), the optimization reduces to a QP problem with respect to $\theta = (\theta_1, \theta_2, \dots, \theta_n)^T$ and β . To further simplify the problem, let $\tilde{x_i}$ = $(K(x_i, x_1), K(x_i, x_2), \dots, K(x_i, x_n))^T$. Then $f(x_i) = \tilde{x}_i^T \theta + \beta$. Therefore the non-crossing technique for the linear case can be applied to the kernel case as well.

Denote by *K* a $n \times n$ matrix with (i, j) -element being $K(x_i, x_j)$. We solve

$$
\min_{\theta,\beta} \quad \left\{ (1-\pi_{k_0}) \sum_{i:y_i=1} [1-(\tilde{x}_i^T \theta + \beta)]_+ + \pi_{k_0} \sum_{i:y_i=-1} [1+(\tilde{x}_i^T \theta + \beta)]_+ + \lambda \theta^T \mathbf{K} \theta \right\}
$$

and denote the corresponding solutions as $\hat{\theta}_{k_0}$ and $\hat{\beta}_{k_0}$. Then, for $k > k_0$, we sequentially solve

$$
\min_{\theta,\beta} \left\{ (1 - \pi_k) \sum_{i:y_i=1} [1 - (\tilde{x}_i^T \theta + \beta)]_+ + \pi_k \sum_{i:y_i=-1} [1 + (\tilde{x}_i^T \theta + \beta)]_+ + \lambda \sum_{j=1}^p w_j^2 \right\},\
$$

s.t. $\tilde{x}^T \theta$

$$
+ \beta \le 0 \text{ for any } \tilde{x} \in \{\tilde{x} : \tilde{x}^T \widehat{\theta}_{k-1} + \widehat{\beta}_{k-1} = 0\} \cap H_{conver}(\tilde{x}_1, \tilde{x}_2, \dots, \tilde{x}_n)
$$
(7)

and denote the corresponding solutions as $\pmb{\theta}_k$ and $\hat{\beta_k}$; for $k < k_0$, we sequentially solve

$$
\min_{\theta,\beta} \left\{ (1 - \pi_k) \sum_{i:y_i=1} [1 - (\tilde{x}_i^T \theta + \beta)]_+ + \pi_k \sum_{i:y_i=-1} [1 + (\tilde{x}_i^T \theta + \beta)]_+ + \lambda \sum_{j=1}^p w_j^2 \right\},
$$

s.t. $\tilde{x}^T \theta$
 $+ \beta \ge 0$ for any $\tilde{x} \in {\tilde{x}} : \tilde{x}^T \widehat{\theta}_{k+1} + \widehat{\beta}_{k+1} = 0$ \cap $H_{conver}(\tilde{x}_1, \tilde{x}_2, \dots, \tilde{x}_n)$ (8)

Note that calculation of the set $\{\hat{x}: \hat{x}^T \hat{\theta}_{k+1} + \hat{\beta}_{k+1} = 0\} \cap H_{convex}(\hat{x}_1, \hat{x}_2, \dots, \hat{x}_n)$ is similar to that of the linear case. The difference here is that we consider the *n*-dimensional kernel space instead of the *p*-dimensional linear space.

The estimated classification boundary for π_k is given by $\{x:\sum_{i=1} \theta_{ki} K(x_i, x) + \beta_k = 0\}$, where θ_{ki} is the *i*th element of θ_k . The constraints (7) and (8) guarantee the desired non-crossing. Once the classification boundaries with different weights are obtained, Step 3 of Wang et al.'s algorithms can be applied to estimate the conditional class probability as in the linear case.

4 Robust learning via preconditioning

In Section 3, we present a technique to estimate classification functions with different weights and without crossing to estimate class probabilities. In this section, we propose to make use of the estimated class probabilities to precondition the training data. This preconditioning operation may help to remove potential outliers and consequently yield more robust classifiers.

In the literature, there is some existing work suggesting that the SVM may not be robust to outliers or mislabeled data points (Shen et al. 2003). One main reason is that the hinge loss used in the SVM is unbounded. Consequently, potential outliers or mislabeled data points may pull the classification boundary towards themselves and thus reduce the classification performance.

For a training data set with a low noise level, we expect that most data points fall in their corresponding class sets. A training point is likely to be an outlier if it is far from its own class. Specifically, for a non-outlier data point x_i with $y_i = 1$ ($y_i = -1$), its estimated conditional class probability $\hat{p}(x_i)$ should be likely to be larger (smaller, respectively) than 0.5. This gives us a potential method to identify outliers. If there are some data points x_i with $y_i = 1$ and $\hat{p}(\mathbf{x}_i)$ close to zero or some data points \mathbf{x}_i with $y_i = -1$ and $\hat{p}(\mathbf{x}_i)$ close to one, then these points (x_i, y_i) 's are likely to be outliers. The outliers can be mislabeled data due to error or a result of sampling biases.

Once we have identified outliers, we can correct them by flipping their binary response from y_i to $-y_i$ and train the standard SVM with the modified training data set. In this way, we are able to remove the effect of these outliers from deteriorating the performance of the SVM. We call this technique preconditioning. In the literature, preconditioning has been applied in the regression problems. In particular, Paul, Bair, Hastie and Tibshirani (2008) proposed to first estimate the regression function to yield a preconditioned response variable. Then they apply standard variable selection techniques to the preconditioned response variable. They argued that the preconditioning step helps to get better variable selection performance. Our preconditioning method is similar to that of Paul et al. (2008) in the sense that we generate a more "clean" binary response variable first and then apply the standard large-margin classifiers to achieve robustness.

We now give more details of our technique of preconditioning. We first choose a prespecified preconditioning level *π̃*, which can be any number between 0 and 0.5. For our training data $\{(x_i, y_i), i = 1, 2, \dots, n\}$, we estimate the conditional class probability $\hat{p}(x_i)$ for each data point using our non-crossing technique. For any x_i , we precondition the binary response by defining

$$
\widetilde{y}_i = \begin{cases}\n-1 & \text{if } \widehat{p}(\mathbf{x}_i) \in [0, \pi) \\
y_i & \text{if } \widehat{p}(\mathbf{x}_i) \in [\pi, 1 - \pi] \\
1 & \text{if } \widehat{p}(\mathbf{x}_i) \in (1 - \pi, 1].\n\end{cases}
$$

Here $\pi \in [0, 0.5)$ controls the level of preconditioning. A large π indicates an aggressive preconditioning while a small *π̃* means that preconditioning is used conservatively. When *π̃* $= 0$, no preconditioning is applied and our preconditioned SVM reduces to the standard SVM.

Once preconditioning has been applied, we train the standard SVM (1) over the preconditioned training data set $\{(\mathbf{x}_i, \tilde{y}_i), i = 1, 2, \dots, n\}$ to get our final classification rule.

Letting $\tilde{f}_{\overline{n}}(\cdot)$ be the solution of (1) using the preconditioned data with the optimal tuning parameter λ , then our final classification rule is sign($\tilde{f}_{\pi}^*(x)$).

Compared to other robust large-margin classifiers such as *ψ*-learning (Shen et al, 2003) and the RSVM (Wu and Liu, 2007), our preconditioning method is another technique to robustify the SVM without nonconvex optimization. In Section 5, we perform a simulation study to compare the performance of our preconditioned SVM with that of the RSVM.

5 Simulation

In this section we use several simulated examples to illustrate the performance of our proposed methods. Examples 1, 2, 3 and 4 are linear while examples 5 and 6 are nonlinear. For the linear case, we use the 2-norm penalty. Example 1 compares the estimated conditional class probability using Wang et al. (2008) with that of our proposed noncrossing version. Examples 2-6 are used to show improvement of the preconditioned SVM over the standard SVM. In all simulated examples, we use an independent tuning set $\{(x_{i, tune}, y_{i, tune}), i = 1, 2, \cdots, n_{tune}\}\$ to select the regularization parameter λ in (2) by minimizing

$$
(1 - \pi_k) \sum_{i: y_{i, \text{tune}} = 1} H(\widehat{f_\lambda}(x_{i, \text{tune}})) + \pi_k \sum_{i: y_{i, \text{tune}} = -1} H(-\widehat{f_\lambda}(x_{i, \text{tune}}))
$$

over a grid of *λ*, where $\hat{f}_\lambda(\cdot)$ denotes the solution of the weighted SVM with the tuning parameter *λ*.

5.1 Linear learning

Example 1 (Class probability estimation comparison)—In this example, the two dimensional predictor $\mathbf{x} = (x_1, x_2)^T$ is uniformly distributed over the square $\{(x_1, x_2)^T : |x_1| +$ $|x_2| \leq 2$. Conditional on $X = x$, *Y* takes +1 with probability $p(x) = \exp(4(x_1 + x_2))$ / $(1+\exp(4(x_1+x_2)))$ and -1 with probability $1-\exp(4(x_1+x_2))/(1+\exp(4(x_1+x_2)))$, namely, (X, Y, Y) *Y*) comes from a logistic regression model. We set the sample size $n = 100$, and $\pi_k = 0.01(k)$ -1) for $k = 1, 2, \dots, 101$. Independent tuning and test data sets are generated to tune the regularization parameter *λ* and evaluate performance, respectively. Both Wang et al.'s method and our non-crossing version for conditional class probability estimation are applied. As the true conditional class probability is known in this simulation, we report the empirical generalized Kullback-Leibler (eGLK) loss over both the training and test data sets. The eGLK loss of a probability estimator $\hat{p}(x)$ for a set $\{(x_i, y_i), i = 1, 2, \dots, n\}$ with true conditional class probability $p(x)$, is defined as

$$
\text{eGLK}(p,\widehat{p}) = \frac{1}{n} \sum_{i=1}^{n} \left(p(x_i) \log \frac{p(x)}{\widehat{p}(x)} + (1 - p(x_i)) \log \frac{1 - p(x)}{1 - \widehat{p}(x)} \right).
$$

Average eGLK of the training set and test set over 100 repetitions are reported in Table 1 with standard errors in parentheses. We can see slight improvement of our non-crossing version, however the improvement is not statistically significant. Nevertheless, the noncrossing version leads to better statistical interpretation.

For a typical sample, we compare Wang et al.'s method with our new non-crossing version by plotting the estimated classification boundaries for different *π^k* 's in Figure 2. The left panel corresponds to Wang et al.'s method while the right panel is for our new non-crossing version. It is clear from the plot that the estimated classification boundaries do cross each

other for some π_k 's if they are estimated individually. However, by enforcing the noncrossing constraints, our new version leads to non-crossing classification boundaries which give better interpretation. This figure demonstrates the usefulness of the non-crossing constraints.

Example 2—In this example, the data $\{(x_i, y_i), i = 1, 2, \dots, n\}$ are generated as *i.i.d.* copies of (*X, Y*) with *X* being uniformly distributed over the square $\{(x_1, x_2) : |x_1| + |x_2| \le 2\}$ and *Y* $=$ sign($x_1 + x_2$) with probability 0:9 and $Y = -\text{sign}(x_1 + x_2)$ with probability 0.1. We set the sample size $n = 100$. An independent tuning set of size *n* is used to select the tuning parameter. In this probability estimation step, we set $\pi_k = 0.05(k-1)$ for $k = 1, 2, \dots, 21$. Over 100 repetitions, average testing errors over an independent test set of size 100*n* and corresponding standard errors are reported in Table 2 for the standard SVM, our preconditioned SVMs with different preconditioning levels π , and the RSVM. Using the standard SVM as the baseline, the difference, ratio and relative improvement are reported as well. For example, when $\pi = 0.1$, the difference and ratio improvements are given by 0.1289 $-0.1294 = -0.0005$, $0.1289/0.1294 = 0.9961$ and $(0.1294 - 0.1289)/(0.1294 - 0.1000) =$ 0.0172 (up to rounding error), respectively. From Table 2, we can see that our preconditioned SVM improves over the standard SVM at all different levels of preconditioning and the improvement is maximized at $\pi = 0.3$. The RSVM leads to a slightly larger improvement using nonconvex optimization.

Example 3 (moderate dimension: $d = 8$ **)—In this example, we increase the** dimensionality of the feature space to 8. The eight-dimensional predictor $\mathbf{x} = (x_1, x_2, \dots, x_n)$

 $(x_8)^T$ is generated from the uniform distribution over $\{x:\sum_{j=1}|x_j|\leq 8\}$. Conditional on $X=x$, $Y = sign(\sum_{j=1}^{8} x_j)$ with probability 0.9 and $Y = -sign(\sum_{j=1}^{8} x_j)$ with probability 0.1. The sample size is chosen to be $n = 200$. Simulation results based on 100 replications are reported in Table 3. The message is the similar to the previous example. In particular, our preconditioned SVM does better than the standard SVM but worse than the RSVM.

Example 4 (high dimension: $d = 50$ **)—In this example, we increase the dimensionality** of the feature space to 50. Predictors are independent of each other and are marginally

distributed as the standard normal distribution. Conditional on $X = x$, $Y = sign(\sum_{j=1}^{50} x_j)$ with

probability 0.9 and $Y = -\text{sign}(\sum_{j=1}^{50} x_j)$ with probability 0.1. The sample size is chosen to be $n = 200$. Simulation results are reported in Table 4. The message is very similar to the previous examples. However, the advantages of the robust methods are much smaller than before, possibly due to the difficulty of high dimension.

5.2 Kernel learning

Example 5—In this example we consider a nonlinear Bayes boundary. The two dimensional predictor $\mathbf{x} = (x_1, x_2)^T$ is uniformly distributed over the square $\{(x_1, x_2)^T : \max(\mathbf{x})\}$ x_1 , $|x_2|$ \leq 2}. Conditional on $X = x$, the binary response *Y* takes sign sign $(x_1^2 + x_2^2 - 8/\pi)$ with probability 0:9 and $-\text{sign}(x_1^2 + x_2^2 - 8/\pi)$ with probability 0.1. Sample size *n* is set to be 100. We implement the nonlinear SVM using the kernel trick with the Gaussian kernel

 $K(x, x') = \exp(-||x - x'||_2^2)$, where $||x||_2 = \sqrt{x_1^2 + x_2^2}$. Results over 100 repetitions are reported in Table 5. Interestingly, the RSVM performs slightly worse than the SVM for this example, while our preconditioned SVM improves over the standard SVM.

Example 6—In this example, two-dimensional predictors are generated uniformly from the

unit disc $\{(x_1, x_2): x_1^2 + x_2^2 < 1\}$. Conditional on $X = x$, $Y = sign(x_1)$ with probability 0.9 and $Y =$ $-\text{sign}(x_1)$ with probability 0.1. The sample size is chosen to be *n* = 100. A similar example was previously used by Shen et al. (2003).

Simulation results are reported in Table 6. In this example, both the RSVM and our preconditioned SVM give improvement over the standard SVM. Compared to the RSVM, our preconditioned SVM delivers more accurate classification accuracy.

It is worthwhile to point out that although both the RSVM and preconditioned SVM can yield more robust classifiers, it is interesting to see from our simulated examples that they perform differently for linear versus nonlinear examples. In particular, the RSVM generally works slightly better than the preconditioned SVM in linear examples, and the preconditioned SVM works better than the RSVM in kernel examples. Further investigation is needed to examine whether this pattern holds in general.

As a remark, we note that the value of *π̃* greatly affects of the performance of our preconditioned SVM. The larger π is, the more change we perform on the training data. In practice, one may use a data adaptive procedure to select π to optimize the performance for the preconditioned SVM. For example, one may treat π as a tuning parameter and select the value which gives the most accurate classification performance.

6 Liver disorder data

In this section we apply our preconditioned SVM to the dataset Liver-disorder. This dataset is available at the UCI Machine Learning Repository. More information can be found at the UCI webpage [http://archive.ics.uci.edu/ml/index.html.](http://archive.ics.uci.edu/ml/index.html) It has a total of 345 observations with six predictor variables and a binary response.

To analyze the data, we first standardize each predictor variable to have mean zero and standard deviation one. Then we randomly split the dataset into training, tuning, and testing sets, each of size 115. As in Wu and Liu (2007), in order to study robustness of our preconditioned SVM for both the training and tuning sets, we contaminate the binary output by randomly choosing a fixed percent of their observations and changing the class output to the other class. Contamination with different percentages, *perc* = 0%, 5%, and 10%, are considered and applied to both the training and tuning sets, but not to the testing set.

To examine the performance of our proposed method, we apply both the standard linear SVM and our preconditioned linear SVM with different preconditioning levels *π̃*'s. Average testing errors over 10 random splitting are reported in Table 7 for different methods with corresponding standard errors in parentheses. For each different *perc*, the entry with boldface corresponds to the best performance and each entry with an underline indicates that it improves over the standard SVM. Table 7 shows that a relatively high level of preconditioning improves over the standard SVM either for the case without contamination (*perc* = 0%) or with contamination (*perc* = 5% or 10%). This clearly demonstrates that our preconditioned SVM is more robust than the standard SVM.

7 Discussion

The SVM only estimates the classification boundary without producing the class probability estimation. To get class probability estimation for the SVM, Wang et al. (2008) makes use of multiple weighted SVM boundaries to yield nonparametric class probability estimation.

However, since each weighted SVM is calculated separately, different weighted classification boundaries may cross each other.

In this work, we first propose a technique to estimate different classification boundaries without crossing using weighted SVMs with constraints. Once the estimated conditional class probabilities are obtained, we make use of that to precondition our training data set. The operation of preconditioning may help to remove potential outliers. To get robust classifiers, we propose to apply the standard SVM on the preconditioned training data set. Our simulated and real examples clearly demonstrate that our preconditioned SVM improves over the standard SVM.

To estimate multiple classification boundaries, we only make use of the 2-norm penalty. However, when the dimensionality is large with many noise variables in the predictor, it is necessary to consider other penalties that are able to achieve variable selection, such as the LASSO penalty (Tibshirani, 1996; Zhu et al, 2004) and SCAD penalty (Fan and Li, 2001). It will be interesting to consider simultaneous classification and variable selection for the SVM as in Zhang, Liu, Wu, and Zhu (2008).

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Figure 1. Plot of several common binary large margin loss functions.

Figure 2.

Estimated classification boundaries for different π_k in a typical sample of Example 1 with π $= 0.05, 0.06, 0.07, 0.08, 0.09, 0.10, 0.20, 0.30, 0.40, 0.50, 0.60, 0.70, 0.80, 0.90, 0.96, 0.97,$ 0.98, 0.99: Wang et al.'s method is in the left panel and our new non-crossing version is in the right panel.

Table 1

Simulation results for Example 1

Table 2

Simulation results of Example 2 Simulation results of Example 2 **π̃**

 $\overline{1}$

Table 3

Simulation results of Example 3 Simulation results of Example 3 **π̃**

Table 4

Simulation results of Example 4 Simulation results of Example 4 **π̃**

Table 5

Simulation results of Example 5 Simulation results of Example 5 **π̃**

Table 6

Simulation results of Example 6 Simulation results of Example 6

Table 7

Numerical results of the Liver Disease data Numerical results of the Liver Disease data

