



# A road to classification in high dimensional space: the regularized optimal affine discriminant

Jianqing Fan,

*Princeton University, USA*

Yang Feng

*Columbia University, New York, USA*

and Xin Tong

*Princeton University, USA*

[Received November 2010. Final revision November 2011]

**Summary.** For high dimensional classification, it is well known that naively performing the Fisher discriminant rule leads to poor results due to diverging spectra and accumulation of noise. Therefore, researchers proposed independence rules to circumvent the diverging spectra, and sparse independence rules to mitigate the issue of accumulation of noise. However, in biological applications, often a group of correlated genes are responsible for clinical outcomes, and the use of the covariance information can significantly reduce misclassification rates. In theory the extent of such error rate reductions is unveiled by comparing the misclassification rates of the Fisher discriminant rule and the independence rule. To materialize the gain on the basis of finite samples, a regularized optimal affine discriminant (ROAD) is proposed. The ROAD selects an increasing number of features as the regularization relaxes. Further benefits can be achieved when a screening method is employed to narrow the feature pool before applying the ROAD method. An efficient constrained co-ordinate descent algorithm is also developed to solve the associated optimization problems. Sampling properties of oracle type are established. Simulation studies and real data analysis support our theoretical results and demonstrate the advantages of the new classification procedure under a variety of correlation structures. A delicate result on continuous piecewise linear solution paths for the ROAD optimization problem at the population level justifies the linear interpolation of the constrained co-ordinate descent algorithm.

**Keywords:** Fisher discriminant; High dimensional classification; Independence rule; Linear discriminant analysis; Regularized optimal affine discriminant

## 1. Introduction

Technological innovations have had a deep influence on society and on various areas of scientific research. High throughput data from microarray and proteomics technologies are frequently used in many contemporary statistical studies. In the case of microarray data, the dimensionality is frequently in thousands or beyond, whereas the sample size is typically of the order of tens. The large  $p$ –small  $n$  scenario poses challenges for the classification problems. We refer to Fan and Lv (2010) for an overview of statistical challenges that are associated with high dimensionality.

When the feature space dimension  $p$  is very high compared with the sample size  $n$ , the Fisher discriminant rule performs poorly owing to diverging spectra as demonstrated by Bickel and

*Address for correspondence:* Xin Tong, Department of Operations Research and Financial Engineering, Princeton University, Princeton, NJ 08540, USA.  
E-mail: xtong@princeton.edu

Levina (2004), who showed that the independence rule in which the covariance structure is ignored performs better than the naive Fisher rule (NFR) in the high dimensional setting. Fan and Fan (2008) demonstrated further that, even for the independence rules, a procedure using all the features can be as poor as random guessing owing to accumulation of noise in estimating population centroids in high dimensional feature space. As a result, Fan and Fan (2008) proposed the features annealed independence rule (FAIR) that selects a subset of important features for classification. Dudoit *et al.* (2002) reported that, for microarray data, ignoring correlations between genes leads to better classification results. Tibshirani *et al.* (2002) proposed the nearest shrunken centroid (NSC) which likewise employs the working independence structure. Similar problems are also studied in the machine learning community such as Domingos and Pazzani (1997) and Lewis (1998).

In microarray studies, correlation between different genes is an essential characteristic of the data and is usually not negligible. Other examples include proteomics, and metabolomics data where correlation between biomarkers is commonplace. More details can be found in Ackermann and Strimmer (2009). Intuitively, the independence assumption among genes leads to loss of critical information and hence is suboptimal. We believe that, in many cases, the crucial point is not whether to consider correlations, but how we can incorporate the covariance structure in the analysis with protection against diverging spectra and significant noise accumulation effect.

The set-up of the objective classification problem is now introduced. We assume in what follows that the variability of data under consideration can be described reasonably well by the means and variances. To be more precise, suppose that random variables representing two classes  $\mathcal{C}_1$  and  $\mathcal{C}_2$  follow  $p$ -variate normal distributions  $\mathbf{X}|Y = 1 \sim \mathcal{N}_p(\boldsymbol{\mu}_1, \boldsymbol{\Sigma})$  and  $\mathbf{X}|Y = 2 \sim \mathcal{N}_p(\boldsymbol{\mu}_2, \boldsymbol{\Sigma})$  respectively. Moreover, assume that  $\mathbb{P}(Y = 1) = \frac{1}{2}$ . This Gaussian discriminant analysis set-up is known for its good performance despite its rigid model structure. For any linear discriminant rule

$$\delta_{\mathbf{w}}(\mathbf{X}) = \mathbb{1}\{\mathbf{w}^T(\mathbf{X} - \boldsymbol{\mu}_a) > 0\}, \tag{1}$$

where  $\boldsymbol{\mu}_a = (\boldsymbol{\mu}_2 + \boldsymbol{\mu}_1)/2$ , and  $\mathbb{1}$  denotes the indicator function with value 1 corresponds to assigning  $\mathbf{X}$  to class  $\mathcal{C}_2$  and 0 class  $\mathcal{C}_1$ , the misclassification rate of the (pseudo)classifier  $\delta_{\mathbf{w}}$  is

$$W(\delta_{\mathbf{w}}) = \frac{1}{2} P_2\{\delta_{\mathbf{w}}(\mathbf{X}) = 0\} + \frac{1}{2} P_1\{\delta_{\mathbf{w}}(\mathbf{X}) = 1\} = 1 - \Phi\{\mathbf{w}^T \boldsymbol{\mu}_d / (\mathbf{w}^T \boldsymbol{\Sigma} \mathbf{w})^{1/2}\}, \tag{2}$$

where  $\boldsymbol{\mu}_d = (\boldsymbol{\mu}_2 - \boldsymbol{\mu}_1)/2$ , and  $P_i$  is the conditional distribution of  $\mathbf{X}$  given its class label  $i$ . We shall focus on such a linear classifier  $\delta_{\mathbf{w}}(\cdot)$ , and the mission is to find a good data projection direction  $\mathbf{w}$ . Note that the Fisher discriminant

$$\delta_F(\mathbf{X}) = \mathbb{1}\{(\boldsymbol{\Sigma}^{-1} \boldsymbol{\mu}_d)^T(\mathbf{X} - \boldsymbol{\mu}_a) > 0\} \tag{3}$$

is the *Bayes rule*. There are two fundamental difficulties in applying the Fisher discriminant whose missclassification rate is

$$1 - \Phi\{(\boldsymbol{\mu}_d^T \boldsymbol{\Sigma}^{-1} \boldsymbol{\mu}_d)^{1/2}\}. \tag{4}$$

The first difficulty arises from the noise accumulation effect in estimating the population centroids (Fan and Fan, 2008) when  $p$  is large. The second challenge is more severe: estimating the inverse of covariance matrix  $\boldsymbol{\Sigma}$  when  $p > n$  (Bickel and Levina, 2004). As a result, much previous research focused on the independence rules, which act as if  $\boldsymbol{\Sigma}$  is diagonal. However, correlation matters!

To illustrate this point, consider a case when  $p = 2$ . These two features can be selected from the original thousands of features, and we can estimate the correlation between two variables with reasonable accuracy. Let

$$\Sigma = \begin{pmatrix} 1 & \rho \\ \rho & 1 \end{pmatrix},$$

where  $\rho \in [0, 1)$  and  $\mu_d = (\mu_1, \mu_2)^T$ . Without loss of generality, assume that  $|\mu_1| \geq |\mu_2| > 0$ . The misclassification rate of the Fisher discriminant depends on

$$\Delta_p(\rho) = \mu_d^T \Sigma^{-1} \mu_d = \frac{1}{1 - \rho^2} (\mu_1^2 + \mu_2^2 - 2\rho\mu_1\mu_2). \tag{5}$$

Note that

$$\Delta'_p(\rho) > 0 \Leftrightarrow \mu_1\mu_2\rho^2 - (\mu_1^2 + \mu_2^2)\rho + \mu_1\mu_2 < 0.$$

Therefore, when  $\mu_1\mu_2 < 0$ ,  $\Delta'_p(\rho) > 0$  for all  $\rho \in [0, 1)$ . In contrast, when  $\mu_1\mu_2 > 0$ ,  $\Delta_p(\rho)$  decreases on  $\rho \in (0, \mu_2/\mu_1)$ , and increases on  $(\mu_2/\mu_1, 1)$ . When  $\rho \rightarrow 1$ ,  $\Delta_p \rightarrow \infty$  regardless of signs for  $\mu_1\mu_2$ , which in turn leads to vanishing classification error. However, if we use the independence rule (which is also called the naive Bayes rule), the optimal misclassification rate

$$1 - \Phi \left\{ \frac{\|\mu_d\|_2^2}{(\mu_d^T \Sigma \mu_d)^{1/2}} \right\} \tag{6}$$

depends on  $\Gamma(\rho) = \|\mu_d\|_2^4 / (\mu_d^T \Sigma \mu_d)$ , which is monotonically decreasing for  $\rho \in [0, 1)$ , with the limit  $(\mu_1^2 + \mu_2^2)^2 / (\mu_1 + \mu_2)^4$  that is smaller than 1 when  $\mu_1$  and  $\mu_2$  have the same sign. Hence, the optimal classification error by using the independence rule actually increases as correlation between features increases.

The above simple example shows that, by incorporating correlation information, the gain in terms of classification error can be substantial. Elaboration on this point in more realistic scenarios is provided in Section 2. Now it seems wise to use at least a part of the covariance structure to improve the performance of a classifier. So there is a need to estimate the covariance matrix  $\Sigma$ . Without structural assumptions on  $\Sigma$ , the pooled sample covariance  $\hat{\Sigma}$  is one natural estimate. But, for  $p > n$ , it is not considered as a good estimate of  $\Sigma$  in general. We are lucky here because our mission is not to construct a good estimate of the covariance matrix, but finding a good direction  $w$  that leads to a good classifier. To mimic the optimal data projection direction  $\Sigma^{-1} \mu_d$ , we do not adopt a direct plug-in approach, simply because it is unlikely that a product is a good estimate when at least one of its components is not. Instead, we find the data projection direction  $w$  by directly minimizing the classification error subject to a capacity constraint on  $w$ . From a broad spectrum of simulated and real data analysis, we are convinced that this approach leads to a robust and efficient sparse linear classifier.

Admittedly, our work is far from the first to use covariance for classification; support vector machines (Vapnik, 1995), for example, implicitly utilize covariance between covariates. Another notable work is ‘shrunk centroids regularized discriminant analysis’ (SCRDA) (Guo *et al.*, 2005), which calls for a version of regularized sample covariance matrix  $\hat{\Sigma}_{reg}$ , and soft thresholds on  $\hat{\Sigma}_{reg}^{-1} \hat{x}_i$ . Shao *et al.* (2011) consider a sparse linear discriminant analysis, assuming the sparsity on both the covariance matrix and the mean difference vector so that they can be regularized. They show that such a regularized estimator is asymptotically optimal under some conditions. However, to the best of our knowledge, this work is the first to select features by directly optimizing the misclassification rates, to use unregularized sample covariance information explicitly, and to establish the oracle inequality and risk approximation theory.

There is a huge literature on high dimensional classification. Examples include principal component analysis in Bair *et al.* (2006) and Zou *et al.* (2006), partial least squares in Nguyen and

Rocke (2002), Huang (2003) and Boulesteix (2004) and sliced inverse regression in Li (1991) and Antoniadis *et al.* (2003).

The rest of our paper is organized as follows. Section 2 provides some insights on the performances of naive Bayes (NB), Fisher discriminant and restricted Fisher discriminants. In Section 3, we propose the regularized optimal affine discriminant (ROAD) and variants of it. An efficient algorithm called constrained co-ordinate descent is constructed in Section 4. The main risk approximation results and continuous piecewise linear property of the solution path are established in Section 5. We conduct simulation and empirical studies in Section 6. A discussion is given in Section 7, and all proofs are relegated to Appendix A.

The data that are analysed in the paper and the programs that were used to analyse them can be obtained from

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## 2. Naive Bayes and Fisher discriminant

To compare the NB and Fisher discriminant at the population level, we assume without loss of generality that variables have been marginally standardized so that  $\Sigma$  is a correlation matrix. Recall that the NB discriminant has error rate (6) and the Fisher discriminant has error rate (4). Let  $\Gamma_p = \|\mu_d\|_2^4 / (\mu_d^T \Sigma \mu_d)$  and  $\Delta_p = \mu_d^T \Sigma^{-1} \mu_d$ . Denote by  $\{\lambda_i\}_{i=1}^p$  the eigenvalues and  $\{\xi_i\}_{i=1}^p$  eigenvectors of the matrix  $\Sigma$ . Decompose

$$\mu_d = a_1 \xi_1 + \dots + a_p \xi_p, \tag{7}$$

where  $\{a_i\}_{i=1}^p$  are the coefficients of  $\mu_d$  in this new orthonormal basis  $\{\xi_i\}_{i=1}^p$ . Using decomposition (7), we have

$$\begin{aligned} \Delta_p &= \sum_{j=1}^p a_j^2 / \lambda_j, \\ \Gamma_p &= \left( \sum_{j=1}^p a_j^2 \right)^2 / \sum_{j=1}^p \lambda_j a_j^2. \end{aligned} \tag{8}$$

The relative efficiency of the Fisher discriminant over the NB discriminant is characterized by  $\Delta_p / \Gamma_p$ . By the Cauchy–Schwartz inequality,

$$\Delta_p / \Gamma_p \geq 1.$$

The NB method performs as well as the Fisher discriminant only when  $\mu_d$  is an eigenvector of  $\Sigma$ .

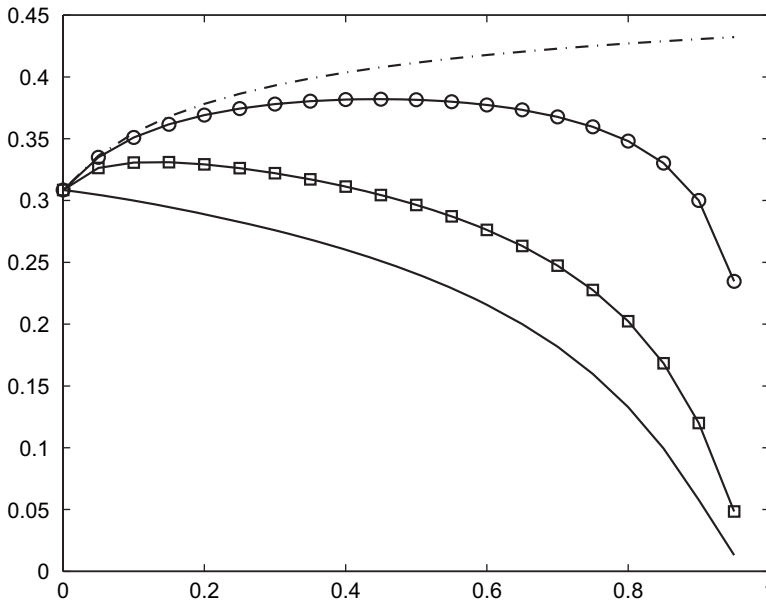
In general,  $\Delta_p / \Gamma_p$  can be much larger than 1. Since  $\Sigma$  is the correlation matrix,  $\sum_{j=1}^p \lambda_j = \text{tr}(\Sigma) = p$ . If  $\mu_d$  is equally loaded on  $\xi_j$ , then the ratio

$$\Delta_p / \Gamma_p = p^{-2} \sum_{j=1}^p \lambda_j \sum_{j=1}^p \lambda_j^{-1} = p^{-1} \sum_{j=1}^p \lambda_j^{-1}. \tag{9}$$

More generally, if  $\{a_j\}_{j=1}^p$  are realizations from a distribution with the second moment  $\sigma^2$ , then, by the law of large numbers,

$$\sum_{j=1}^p a_j^2 \lambda_j^{-1} \approx \sigma^2 \sum_{j=1}^p 1 / \lambda_j, \quad p^{-1} \sum_{j=1}^p a_j^2 \approx \sigma^2, \quad \sum_{j=1}^p \lambda_j a_j^2 \approx \sigma^2 \sum_{j=1}^p \lambda_j.$$

Hence, equation (9) holds approximately in this case. In other words, the right-hand side of equation (9) is approximately the relative efficiency of the Fisher discriminant over the NB



**Fig. 1.** Misclassification rates of the Fisher discriminant (—), NB (· · · ·) and restricted Fisher (10 features) (○) and restricted Fisher (20 features) discriminants (□) against  $\rho$

discriminant. Now suppose further that half of the eigenvalues of  $\Sigma$  are  $c$  and the other half are  $2 - c$ . Then, the right-hand side of equation (9) is  $\{c^{-1} + (2 - c)^{-1}\}/2$ . For example when the condition number is 10, this ratio is about 3. A high ratio translates into a large difference in error rates:  $1 - \Phi(\Gamma_p^{1/2})$  for the independence rule is much larger than  $1 - \Phi(3\Gamma_p^{1/2})$  for the Fisher discriminant. For example, when  $\Gamma_p^{1/2} = 0.5$ , we have 30.9% and 6.7% error rates respectively for the NB and Fisher discriminant.

To put the above arguments under a visual inspection, consider a case in which  $p = 1000$ ,  $\mu_d = (\mu_s^T, \mathbf{0}^T)^T$  with  $\mu_s = (0.1, 0.1, 0.1, 0.1, 0.1, 0.2, 0.2, 0.2, 0.2, 0.2)^T$  and  $\Sigma$  equals the equicorrelation matrix with pairwise correlation  $\rho$ . The vector  $\mu_d$  simulates the case in which 10 genes out of 1000 express mean differences. Fig. 1 depicts the theoretical error rates of the Fisher discriminant and the NB rule as functions of  $\rho$ .

It is not surprising that the Fisher discriminant rule performs significantly better than the NB discriminant as  $\rho$  deviates away from 0. The error rate of the NB rule actually increases with  $\rho$ , whereas the error rate of the Fisher discriminant tends to 0 as  $\rho$  approaches 1. This phenomenon is the same as what was shown analytically through the toy example in Section 1. To mimic the Fisher discriminant by a plug-in estimator, we need to estimate  $\Sigma^{-1}\mu_d$  with reasonable accuracy. This mission is difficult if not impossible. However, imitating a weaker oracle is more manageable. For example, when the samples are of reasonable size, we can select the 10 variables with differences in means by applying a two-sample  $t$ -test. Restricting to the best linear classifiers based on these  $s = 10$  variables, we have the optimal error rate

$$1 - \Phi\{(\mu_s^T \Sigma_s^{-1} \mu_s)^{1/2}\},$$

where the classification rule is  $\delta_{w^R}$  and  $w^R = ((\Sigma_s^{-1} \mu_s)^T, \mathbf{0}^T)^T$ . The performance of this oracle classifier is depicted by the sub-Fisher (10 features) discriminant in Fig. 1. It performs much better than the NB method. One can also employ the NB rule to the restricted feature space, but this method has exactly the same performance as the NB method in the whole space. Thus,

the restricted Fisher discriminant outperforms both the NB method with restricted features and the NB method using all features.

Mimicking the performance of the restricted Fisher discriminant is feasible. Instead of estimating a  $1000 \times 1000$  covariance matrix, we only need to gauge a  $10 \times 10$  submatrix. However, this restricted Fisher rule is not sufficiently powerful, as shown in Fig. 1. We can improve its performance by including the 10 most correlated variables in each of those selected features to account further for the correlation effect, giving rise to a 20-dimensional feature space. Since the variables are equally correlated in this example, we are free to choose any 10 variables among the other 990. The performance of such an enlarged restricted Fisher discriminant is represented by the sub-Fisher (20 features) discriminant in Fig. 1. It performs closely to the Fisher discriminant which uses the whole feature space, and it is feasible to implement with finite samples.

### 3. Regularized optimal affine discriminant

The misclassification rate of the Fisher discriminant is  $1 - \Phi(\Delta_p^{1/2})$ , where  $\Delta_p = \boldsymbol{\mu}_d^T \boldsymbol{\Sigma}^{-1} \boldsymbol{\mu}_d$ . However, for high dimensional data, it is impossible to achieve such a performance empirically. Among other reasons, the estimated covariance matrix  $\hat{\boldsymbol{\Sigma}}$  is ill conditioned or not invertible. One solution is to focus only on the  $s$  ( $\ll p$ ) most important features for classification. Ideally, the best  $s$  features should be those with the largest  $\Delta_s$  among all  $\binom{p}{s}$  possibilities, where  $\Delta_s$  is the counterpart of  $\Delta_p$  when only  $s$  variables are considered. Naive search for the best subset of size  $s$  is NP hard. Thus, we develop a regularized method to circumvent these two problems.

#### 3.1. Definition of regularized optimal affine discriminant

Recall that, by equation (2), minimizing the classification error  $W(\delta_w)$  is the same as maximizing  $\mathbf{w}^T \boldsymbol{\mu}_d / (\mathbf{w}^T \boldsymbol{\Sigma} \mathbf{w})^{1/2}$ , which is equivalent to minimizing  $\mathbf{w}^T \boldsymbol{\Sigma} \mathbf{w}$  subject to  $\mathbf{w}^T \boldsymbol{\mu}_d = 1$ . We would like to add a penalty function for capacity control. There are many ways to do regularization; for the literature on penalized methods, refer to the lasso (Tibshirani, 1996), smoothly clipped absolute deviation (Fan and Li, 2001), the elastic net (Zou and Hastie, 2005), the minimax concave penalty (Zhang, 2010) and related methods (Zou, 2006; Zou and Li, 2008). As our primary interest is classification error (the risk of the procedure), an  $L_1$ -constraint  $\|\mathbf{w}\|_1 \leq c$  is added for regularization, so the problem can be recast as

$$\mathbf{w}_c = \arg \min_{\|\mathbf{w}\|_1 \leq c, \mathbf{w}^T \boldsymbol{\mu}_d = 1} \mathbf{w}^T \boldsymbol{\Sigma} \mathbf{w}. \tag{10}$$

We name the classifier  $\delta_{w_c}(\cdot)$  the regularized optimal affine discriminant (ROAD). The existence of a feasible solution in equation (10) dictates that

$$c \geq 1 / \max_{1 \leq i \leq p} |\mu_{d,i}|. \tag{11}$$

When  $c$  is small, we obtain a sparse solution and achieve feature selection by using covariance information. When  $c \geq \|\boldsymbol{\Sigma}^{-1} \boldsymbol{\mu}_d\|_1 / (\boldsymbol{\mu}_d^T \boldsymbol{\Sigma}^{-1} \boldsymbol{\mu}_d)$ , the  $L_1$ -constraint is no longer binding and  $\delta_{w_c}$  reduces to the Fisher discriminant, which can be denoted by  $\delta_{w_\infty}$  ( $= \delta_F$ ). Therefore we have provided a family of linear discriminants, indexed by  $c$ , using from only one feature to all features. In some applications such as portfolio selection, the choice of  $c$  reflects the investor's tolerance upper bound on gross exposure. In other applications, when the user does not have a such a preference, the choice of  $c$  can be data driven. To accommodate both application

scenarios, we propose a co-ordinate descent algorithm (Section 4) to implement our ROAD proposal.

### 3.2. Variants of regularized optimal affine discriminant

At the sample level, NSCs (Tibshirani *et al.*, 2002) and the FAIR (Fan and Fan, 2008) both use shrunken versions of standardized mean differences to find the  $s$  features. In the same spirit, we consider the following diagonal regularized optimal affine discriminant (DROAD)  $\delta_{\mathbf{w}_c^I}$ , where

$$\mathbf{w}_c^I = \arg \min_{\|\mathbf{w}\|_1 \leq c, \mathbf{w}^T \boldsymbol{\mu}_d = 1} \mathbf{w}^T \text{diag}(\boldsymbol{\Sigma}) \mathbf{w}. \tag{12}$$

The DROAD will be compared with NSCs (Tibshirani *et al.*, 2002) and the FAIR (Fan and Fan, 2008) in the simulation studies, and all these independence-based rules will be compared with the ROAD and its two variants defined below.

A screening-based variant (to be proposed) of the ROAD aims at mimicking the performance of the sub-Fisher (10 features) discriminant in Fig. 1. A fast way to select features is independence screening, which uses the marginal information such as the two-sample  $t$ -test. We can also enlarge the selected feature subspace by incorporating the features which are most correlated with what have been chosen. This additional variant of the ROAD tracks the performance of sub-Fisher (20 features) discriminant in Fig. 1. We shall refer to the two variants of the ROAD as S-ROAD1 and S-ROAD2. More description of these procedures, along with their theoretical properties and numerical investigations, will be detailed in Sections 5 and 6.

A hint of the rationale behind including correlated features that do not show a difference in means between the two classes is revealed through the two-feature example in Section 1. Suppose that  $\mu_2 = 0$ . Then, by equation (5), the power of the discriminant using two features is  $1 - \Phi(\Delta_2^{1/2})$  where  $\Delta_2 = \mu_1^2 / (1 - \rho^2)$ , whereas with the first feature alone the misclassification rate is  $1 - \Phi(\Delta_1^{1/2})$  where  $\Delta_1 = \mu_1^2$ . Therefore when the correlation  $|\rho|$  is large, using two correlated features is far more powerful than employing only one feature, even though the second feature has no marginal discrimination power. More intuition is granted by this observation: at the population level, the best  $s$  features are not necessarily those with largest standardized mean differences. In other words, with the two-class Gaussian model in mind, when  $\boldsymbol{\Sigma}$  is the correlation matrix, the most powerful  $s$  features for classification are not necessarily the co-ordinates of  $\boldsymbol{\mu}_d$  with largest absolute values. This is illustrated by the next stylized example.

Let  $\mathbf{X}|Y=0 \sim \mathcal{N}(\boldsymbol{\mu}_1, \boldsymbol{\Sigma})$  and  $\mathbf{X}|Y=1 \sim \mathcal{N}(\boldsymbol{\mu}_2, \boldsymbol{\Sigma})$ , where  $\boldsymbol{\mu}_1 = (0, 0, 0)^T$ ,  $\boldsymbol{\mu}_2 = (4, 0.5, 1)^T$  and

$$\boldsymbol{\Sigma} = \begin{pmatrix} 1 & -0.25 & 0 \\ -0.25 & 1 & 0 \\ 0 & 0 & 1 \end{pmatrix}.$$

Suppose that the objective is to choose two out of three variables for classification. If we rank features by marginal information, e.g. by the absolute value of standardized mean differences, then we would choose the first and third features. However, denote  $\boldsymbol{\mu}_{d,ij}$  the mean difference vector for features  $i$  and  $j$ , and  $\boldsymbol{\Sigma}_{ij}$  the covariance matrix of features  $i$  and  $j$ ; then the classification power using features  $i$  and  $j$  depends on  $\Gamma_{ij} = \boldsymbol{\mu}_{d,ij}^T \boldsymbol{\Sigma}_{ij}^{-1} \boldsymbol{\mu}_{d,ij}$ . Simple calculation leads to

$$\Gamma_{12} = 18.4 > 17 = \Gamma_{13}.$$

Hence the most powerful two features for classification are not the first and third.

#### 4. Constrained co-ordinate descent

With a Lagrangian argument, we reformulate problem (10) as

$$\bar{\mathbf{w}}_\lambda = \arg \min_{\mathbf{w}^T \boldsymbol{\mu}_d = 1} \frac{1}{2} \mathbf{w}^T \boldsymbol{\Sigma} \mathbf{w} + \lambda \|\mathbf{w}\|_1. \tag{13}$$

In this section, we propose a constrained co-ordinate descent algorithm that is tailored to solving our minimization problem with linear constraints. Optimization (13) is a constrained quadratic programming problem and can be solved by existing software such as MOSEK. Although such software are well regarded in practice, they are slow for our application. The structure of problem (13) could be exploited to obtain a more efficient algorithm. In line with the algorithm LARS, we shall exploit the fact that the solution path has a piecewise linear property.

In the compressed sensing literature, it is common to replace an affine constraint by a quadratic penalty. We borrow this idea and consider the following approximation to equation (13):

$$\tilde{\mathbf{w}}_{\lambda, \gamma} = \arg \min \frac{1}{2} \mathbf{w}^T \boldsymbol{\Sigma} \mathbf{w} + \lambda \|\mathbf{w}\|_1 + \frac{1}{2} \gamma (\mathbf{w}^T \boldsymbol{\mu}_d - 1)^2. \tag{14}$$

In practice, we replace  $\boldsymbol{\Sigma}$  by the pooled sample covariance  $\hat{\boldsymbol{\Sigma}}$  and  $\boldsymbol{\mu}$  by the sample mean difference vector  $\hat{\boldsymbol{\mu}}_d$ . By theorem 6.7 in Ruszczyński (2006), we have

$$\tilde{\mathbf{w}}_{\lambda, \gamma} \rightarrow \bar{\mathbf{w}}_\lambda \quad \text{when } \gamma \rightarrow \infty.$$

Note that we do not have to enforce the affine constraint strictly, because it serves only to normalize our problem. In the optimization problem (14), when  $\lambda = 0$ , the solution  $\tilde{\mathbf{w}}_{0, \gamma}$  is always in the direction of  $\boldsymbol{\Sigma}^{-1} \boldsymbol{\mu}_d$ , the Fisher discriminant, regardless of the value of  $\gamma$ . In addition, this observation is confirmed in the data analysis (Section 6.2) by the insensitivity of choice for  $\gamma$ . Therefore we hold  $\gamma$  as a constant in practice.

We solve problem (14) by co-ordinate descent. Non-gradient algorithms seem to be less popular for convex optimization. For instance, the popular textbook *Convex Optimization* by Boyd and Vandenberghe (2004) does not even have a section on these methods. The co-ordinate descent method is an algorithm in which the  $p$  search directions are just unit vectors  $e_1, \dots, e_p$ , where  $e_i$  denotes the  $i$ th element in the standard basis of  $\mathbb{R}^p$ . These unit vectors are used as search directions in each search cycle until some convergence criterion has been met.

What makes the co-ordinate descent algorithm particularly attractive for problem (14) is that there is an explicit formula for each co-ordinate update. For a given  $\gamma$ , fix  $\tau$  and  $K$ ; then do the optimization on a grid (of log-scale) of  $\lambda$ -values:  $\tau \lambda_{\max} = \lambda_K < \lambda_{K-1} < \dots < \lambda_1 = \lambda_{\max}$ . The  $\lambda_{\max}$  is the minimum  $\lambda$ -value such that no variables enter the model; this is analogous to the minimum requirement on  $c$  in inequality (11). In our implementation, we take  $\tau = 0.001$  and  $K = 100$ . The problem is solved backwards from  $\lambda_{\max}$ . When  $\lambda = \lambda_{i+1}$ , we use the solution from  $\lambda = \lambda_i$  as the initial value. This kind of ‘warm start’ is very effective in improving computational efficiency.

Consider a co-ordinate descent step to solve problem (14). Without loss of generality, suppose that  $\tilde{w}_j$  for all  $j \geq 2$  are given, and we need to optimize with respect to  $w_1$ . The objective function now becomes

$$g(w_1) = \frac{1}{2} (w_1^T \quad \tilde{\mathbf{w}}_2^T) \begin{pmatrix} \boldsymbol{\Sigma}_{11} & \boldsymbol{\Sigma}_{12} \\ \boldsymbol{\Sigma}_{21} & \boldsymbol{\Sigma}_{22} \end{pmatrix} \begin{pmatrix} w_1 \\ \tilde{\mathbf{w}}_2 \end{pmatrix} + \lambda |w_1| + \lambda |\tilde{\mathbf{w}}_2| + \frac{1}{2} \gamma (\mathbf{w}^T \boldsymbol{\mu}_d - 1)^2.$$

When  $w_1 \neq 0$ , we have

$$\begin{aligned} g'(w_1) &= \boldsymbol{\Sigma}_{11} w_1 + \boldsymbol{\Sigma}_{12} \tilde{\mathbf{w}}_2 + \lambda \operatorname{sgn}(w_1) + \gamma (\mathbf{w}^T \boldsymbol{\mu}_d - 1) \boldsymbol{\mu}_{d1} \\ &= (\boldsymbol{\Sigma}_{11} + \gamma \boldsymbol{\mu}_{d1}^2) w_1 + (\boldsymbol{\Sigma}_{12} + \gamma \boldsymbol{\mu}_{d1} \boldsymbol{\mu}_{d2}^T) \tilde{\mathbf{w}}_2 + \lambda \operatorname{sgn}(w_1) - \gamma \boldsymbol{\mu}_{d1}. \end{aligned}$$



By simple calculation (Donoho and Johnstone, 1994), the co-ordinatewise update has the form

$$\tilde{w}_1 = \frac{S\{\gamma\mu_{d1} - (\Sigma_{12} + \gamma\mu_{d1}\mu_{d2}^T)\tilde{w}_2, \lambda\}}{\Sigma_{11} + \gamma\mu_{d1}^2},$$

where  $S(z, \lambda) = \text{sgn}(z)(|z| - \lambda)^+$  is the soft thresholding operator.

Now, we consider the convergence property of the co-ordinate descent algorithm. Here, although the objective function is not strictly convex, it is strictly convex in each of the co-ordinates.

To show that  $g(w_1)$  is strictly convex in  $w_1$ , we decompose it as

$$g(w_1) = g_1(w_1) + g_2(w_1),$$

where  $g_2(w_1) = \lambda|w_1|$  and

$$g_1(w_1) = \frac{1}{2}(w_1^T \quad \tilde{w}_2^T) \begin{pmatrix} \Sigma_{11} & \Sigma_{12} \\ \Sigma_{21} & \Sigma_{22} \end{pmatrix} \begin{pmatrix} w_1 \\ \tilde{w}_2 \end{pmatrix} + \lambda|\tilde{w}_2|_1 + \frac{1}{2}\gamma(w^T\mu_d - 1)^2.$$

$g_1(w_1)$  is a quadratic function of  $w_1$  and  $g_1''(w_1) = \Sigma_{11} + \gamma\mu_{d1}^2 > 0$  for all  $w_1 \in \mathbb{R}$ . Therefore, the function  $g_1(\cdot)$  is strictly convex on  $\mathbb{R}$ . Also, it is clear that  $g_2$  is convex on  $\mathbb{R}$ . Therefore  $g = g_1 + g_2$  is a strictly convex function on  $\mathbb{R}$ .

Combining the co-ordinatewise strict convexity with the fact that the non-differentiable part of the objective function is separable, theorem 5.1 of Tseng (2001) guarantees that co-ordinate descent algorithms converge to co-ordinatewise minima. Moreover, since all directional derivatives exist, every co-ordinatewise minimum is also a local minimum. A similar study on the convergence of the co-ordinate descent algorithm can be found in Breheny and Huang (2011).

In each co-ordinate update, the computational complexity is  $\mathcal{O}(p)$ . A complete cycle through all  $p$  variables costs  $\mathcal{O}(p^2)$  operations. From our experience, constrained co-ordinate descent converges quickly after a few cycles if a warm start is used for the initial solution. Let  $C$  denote the average number of cycles until convergence for each  $\lambda$ . Then our constrained co-ordinate descent algorithm enjoys computational complexity  $\mathcal{O}(CKp^2)$ . The DROAD can be similarly implemented by replacing the covariance matrix with its diagonal.

### 5. Asymptotic property

#### 5.1. Risk approximation

Let  $\hat{w}_c$  be a sample version of  $w_c$  in problem (10):

$$\hat{w}_c \in \arg \min_{\|w\|_1 \leq c, w^T\hat{\mu}_d = 1} w^T\hat{\Sigma}w. \tag{15}$$

The fact that  $\hat{\Sigma}$  is only positive semidefinite leads to potential non-uniqueness of  $\hat{w}_c$ . Now, we have three different classifiers:  $\delta_{w_\infty} = \mathbb{I}\{w_\infty^T(X - \mu_a) > 0\}$ ,  $\delta_{w_c} = \mathbb{I}\{w_c^T(X - \mu_a) > 0\}$  and  $\hat{\delta}_{w_c} = \mathbb{I}\{\hat{w}_c^T(X - \hat{\mu}_a) > 0\}$ . The first two are oracle classifiers, requiring knowledge of unknown parameters  $\mu_1, \mu_2$  and  $\Sigma$ , whereas the third is the feasible ROAD classifier based on the sample. Their classification errors are given by equation (2). Explicitly, the error rates are respectively  $W(\delta_{w_\infty})$  (see expression (4)),  $W(\delta_{w_c})$  and  $W(\hat{\delta}_{w_c})$ . By equation (2), an obvious estimator of the misclassification rate of  $\hat{\delta}_{w_c}$  is

$$W_n(\hat{\delta}_{w_c}) = 1 - \Phi \left\{ \frac{\hat{w}_c^T \hat{\mu}_d}{(\hat{w}_c^T \hat{\Sigma} \hat{w}_c)^{1/2}} \right\}. \tag{16}$$

Two questions arise naturally.

- (a) How close is  $W(\hat{\delta}_{\mathbf{w}_c})$ , the misclassification error of  $\hat{\delta}_{\mathbf{w}_c}$ , to that of its oracle  $W(\delta_{\mathbf{w}_c})$ ?
- (b) Does  $W_n(\hat{\delta}_{\mathbf{w}_c})$  estimate  $W(\delta_{\mathbf{w}_c})$  well?

Theorem 1 addresses these two questions. We introduce an intermediate optimization problem for convenience:

$$\mathbf{w}_c^{(1)} = \arg \min_{\|\mathbf{w}\|_1 \leq c, \mathbf{w}^T \hat{\boldsymbol{\mu}}_d = 1} \mathbf{w}^T \boldsymbol{\Sigma} \mathbf{w}.$$

*Theorem 1.* Let  $s_c = \|\mathbf{w}_c\|_0$ ,  $s_c^{(1)} = \|\mathbf{w}_c^{(1)}\|_0$  and  $\hat{s}_c = \|\hat{\mathbf{w}}_c\|_0$ . Assume that  $\lambda_{\min}(\boldsymbol{\Sigma}) \geq \sigma_0^2 > 0$ ,  $\|\hat{\boldsymbol{\Sigma}} - \boldsymbol{\Sigma}\|_\infty = O_p(a_n)$  and  $\|\hat{\boldsymbol{\mu}}_d - \boldsymbol{\mu}_d\|_\infty = O_p(a_n)$  for a given sequence  $a_n \rightarrow 0$ . Then, we have

$$W(\hat{\delta}_{\mathbf{w}_c}) - W(\delta_{\mathbf{w}_c}) = O_p(d_n)$$

and

$$W_n(\hat{\delta}_{\mathbf{w}_c}) - W(\hat{\delta}_{\mathbf{w}_c}) = O_p(b_n),$$

where  $b_n = (c^2 \vee s_c \vee s_c^{(1)})a_n$  and  $d_n = b_n \vee (\hat{s}_c a_n)$ .

*Remark 1.* In theorem 1,  $\|\cdot\|_\infty$  is the elementwise super norm. When  $\hat{\boldsymbol{\Sigma}}$  is the sample covariance, under some mild moment conditions,  $\|\hat{\boldsymbol{\Sigma}} - \boldsymbol{\Sigma}\|_\infty = O_p(\sqrt{\{\log(p)/n\}})$ ; hence we can take  $a_n = \sqrt{\{\log(p)/n\}}$ . The first result in theorem 1 shows the difference between the misclassification rate of  $\hat{\delta}_{\mathbf{w}_c}$  and its oracle version  $\delta_{\mathbf{w}_c}$ ; the second result indicates the error in estimating the true misclassification rate of the ROAD.

*Remark 2.* In view of equation (2), one intends to choose a  $\mathbf{w}$  that makes  $\mathbf{w}^T \boldsymbol{\Sigma} \mathbf{w}$  small and  $\mathbf{w}^T \boldsymbol{\mu}_d$  large. A compromise between these dual objectives leads to the utility function

$$U(\mathbf{w}) = -\mathbf{w}^T \boldsymbol{\Sigma} \mathbf{w} + \xi \boldsymbol{\mu}_d^T \mathbf{w}$$

as a proxy of the objective function (2) for a fixed  $\xi$ . For any  $\xi > 0$ , the optimal choice  $\mathbf{w}^* \in \arg \min U(\mathbf{w})$  leads to the Fisher discriminant rule. Consider also the regularized versions

$$\begin{aligned} \mathbf{w}_c^* &= \arg \min_{\|\mathbf{w}\|_1 \leq c} U(\mathbf{w}), \\ \hat{\mathbf{w}}_c^* &= \arg \min_{\|\mathbf{w}\|_1 \leq c} \hat{U}(\mathbf{w}), \end{aligned}$$

where  $\hat{U}(\mathbf{w})$  is the utility function with  $\boldsymbol{\Sigma}$  and  $\boldsymbol{\mu}_d$  estimated by  $\hat{\boldsymbol{\Sigma}}$  and  $\hat{\boldsymbol{\mu}}_d$ . Then, it is easy to see the following utility approximation: for any  $\|\mathbf{w}\|_1 \leq c$

$$|U(\mathbf{w}) - \hat{U}(\mathbf{w})| \leq \|\hat{\boldsymbol{\Sigma}} - \boldsymbol{\Sigma}\|_\infty c^2 + \xi c \|\hat{\boldsymbol{\mu}}_d - \boldsymbol{\mu}_d\|_\infty$$

and

$$|U(\hat{\mathbf{w}}_c^*) - U(\mathbf{w}_c^*)| \leq 2(\|\hat{\boldsymbol{\Sigma}} - \boldsymbol{\Sigma}\|_\infty c^2 + \xi c \|\hat{\boldsymbol{\mu}}_d - \boldsymbol{\mu}_d\|_\infty).$$

*Remark 3.* The most prominent technical challenge of our original problem (10) is due to different dualities of penalization problems. For the population version (10), it can be reduced, by the Lagrange multiplier method, to the utility  $U(\mathbf{w})$  optimization problem in remark 2 with a given  $\xi > 0$ , whereas, for the sample version (15), it can be reduced to the utility  $\hat{U}(\mathbf{w})$  optimization problem with a different  $\hat{\xi}$ . Therefore, the problem is not the same as the utility optimization problem in remark 2:  $\hat{\xi}$  is difficult to bound. In fact, it is much more difficult and yields more complicated results.

We now show how different the data projection direction in the regularized oracle can be from that in the Fisher discriminant. To gain better insight, we reformulate the  $L_1$ -constraint

problem as the following penalized version:

$$\mathbf{w}^\lambda = \arg \min_{\mathbf{w}: \mu_d^T \mathbf{w} = 1} \mathbf{w}^T \Sigma \mathbf{w} + \lambda \|\mathbf{w}\|_1. \tag{17}$$

The following characterizes its convergence to the Fisher discriminant weight  $\mathbf{w}_\infty$  as  $\lambda \rightarrow 0$ .

*Theorem 2.* Let  $s$  be the size of the set  $\{k : (\Sigma^{-1} \mu_d)_k \neq 0\}$ . Then, we have

$$\|\mathbf{w}^\lambda - \mathbf{w}_\infty\|_2 \leq \frac{\lambda \sqrt{s}}{\lambda_{\min}(\Sigma)},$$

where  $\mathbf{w}_\infty = \Sigma^{-1} \mu_d / (\mu_d^T \Sigma^{-1} \mu_d)$  is the normalized Fisher discriminant, optimizing problem (17) with  $\lambda = 0$ .

### 5.2. Screening-based regularized optimal affine discriminant

Following the idea of sure independence screening in Fan and Lv (2008), we prescreen all the features before applying the ROAD method. The advantage of this two-step procedure is that we have control on the total number of features that are used in the final classification rule. A popular method for independent feature selection is the two-sample  $t$ -test (Tibshirani *et al.*, 2002; Fan and Fan, 2008), which is a specific case of marginal screening in Fan and Lv (2008). The sure screening property of such a method was demonstrated in Fan and Fan (2008), which selects consistently the features with different means in the same settings as ours.

Once the features have been selected, we apply the ROAD method, producing the original screening-based ROAD S-ROAD1.

- (a) Employ a screening method to obtain  $k$  features.
- (b) Apply the ROAD to the  $k$  selected features.

In the first step, we use the  $t$ -statistics as the screening criterion and determine a data-driven threshold. This idea is motivated by a false discovery rate criterion for choosing the marginal screening threshold in Zhao and Li (2010). A random permutation  $\pi$  of  $\{1, \dots, n\}$  is used to decouple  $\mathbf{X}_i$  and  $Y_i$  so that the resulting data  $(\mathbf{X}_{\pi(i)}, Y_i)$  follow a null model, by which we mean that features have no prediction power for the class label. More specifically, the screening step is carried out as follows.

- Step 1:* calculate the  $t$ -statistic  $t_j$  for each feature  $j$ , where  $j = 1, \dots, p$ .
- Step 2:* for the permuted data pairs  $(\mathbf{X}_{\pi(i)}, Y_i)$ , recalculate the  $t$ -statistic  $t_j^*$ , for  $j = 1, \dots, p$ . (Intuitively, if  $j$  is the index of an important feature,  $|t_j|$  should be larger than most of  $|t_j^*|$ , because the random permutation is meant to eliminate the prediction power of features.)
- Step 3:* for  $q \in [0, 1]$ , let  $\omega_{(q)}$  be the  $q$ th quantile of  $\{|t_j^*|, j = 1, 2, \dots, p\}$ . Then, the selected set  $\mathcal{A}$  is defined as

$$\mathcal{A} = \{j \mid |t_j| \geq \omega_{(q)}\}.$$

The choice of threshold is made to retain the features whose  $t$ -statistics are significant in the two-sample  $t$ -test. Alternatively, if the user knows his  $k$  (owing to budget constraints, etc.), then he can just rank  $|t_j|$ s and choose the threshold accordingly.

S-ROAD1 tracks the performance of oracle procedures like the sub-Fisher (10 features) discriminant in Fig. 1. The feature space that is obtained in the first step can be expanded by including those features which are most correlated with what have already been selected. This additional variant, S-ROAD2, aims at achieving the performance of the sub-Fisher (20 features) type of procedure in Fig. 1.

To elaborate on the theoretical properties of S-ROAD1, assume with no loss of generality that the first  $k$  variables are selected in the screening step. Denote by  $\Sigma_k$  the upper left  $k \times k$  block of  $\Sigma$  and  $\mu_k$  the first  $k$  co-ordinates of  $\mu_d$ . Let

$$\beta_c = \arg \min_{\|\beta\|_1 \leq c, \beta^T \mu_k = 1} \beta^T \Sigma_k \beta.$$

The quantities  $\hat{\beta}_c$  and  $\beta_c^{(1)}$  are defined similarly to  $\hat{w}_c$  and  $w_c^{(1)}$  (which were defined immediately before theorem 1). Then denote by  $y_c = (\beta_c^T, \mathbf{0}^T)^T$ ,  $\hat{y}_c = (\hat{\beta}_c^T, \mathbf{0}^T)^T$  and  $y_c^{(1)} = (w_c^{(1)}, \mathbf{0}^T)^T$ . The next two theorems can be verified along lines similar to those for theorems 1 and 2. Hence, the proofs have been omitted.

*Theorem 3.* If  $\|\hat{\Sigma}_k - \Sigma_k\|_\infty = O_p[\sqrt{\{\log(k)/n\}}]$ ,  $\|\hat{\mu}_k - \mu_k\|_\infty = O_p[\sqrt{\{\log(k)/n\}}]$ , and  $\lambda_{\min}(\Sigma_k) \geq \delta_0 > 0$ , then we have

$$W(\hat{\delta}_{y_c}) - W(\delta_{y_c}) = O_p(e_n)$$

and

$$W_n(\hat{\delta}_{y_c}) - W(\delta_{y_c}) = O_p(e_n),$$

where  $e_n = (c^2 \vee k)\sqrt{\{\log(k)/n\}}$ .

This result is cleaner than theorem 1, as the rate does not involve  $s_c$  and  $\hat{s}_c$ : they are simply replaced by the upper bound  $k$ . Accurate bounds for  $s_c$  and  $\hat{s}_c$  are of interest for future exploration, but they are beyond the scope of this paper.

*Theorem 4.* Let  $y_k^\lambda = \arg \min_{y: \mu_d^T y = 1, y \in M_k} R(y) + \lambda \|y\|_1$  where  $M_k$  is the subspace in  $R^p$  with the last  $p - k$  components being 0, and  $y^0 = ((\Sigma_k^{-1} \mu_k)^T / (\mu_k^T \Sigma_k^{-1} \mu_k), \mathbf{0}^T)^T$ . Then we have

$$\|y_k^\lambda - y^0\|_2 \leq \frac{\lambda \sqrt{k}}{\lambda_{\min}(\Sigma_k)}.$$

### 5.3. Continuous piecewise linear solution path

We use the word ‘linear’ when referring to ‘affine’, in line with the *status quo* in the statistical community. Continuous piecewise linear paths are of much interest to statisticians, as the property reduces the computational complexity of solutions and justifies the linear interpolations of solutions at discrete points. Previous well-known investigations include Efron *et al.* (2004) and Rosset and Zhu (2007). Our set-up differs from others mainly in that, in addition to a complexity penalty, there is also an affine constraint. Our proof calls in point set topology, and is purely geometrical, in a spirit that is very different from the existing ones. In particular, we stress that the continuity property is intuitively correct, but it is far from a trivial consequence of the assumptions. We also believe that the claim holds true even if the  $(p - 1)$ -dimensional affine subspace constraint is replaced by more generic constraints, though the technicality of the proof must be more involved.

*Theorem 5.* Let  $\mu_d \in \mathbb{R}^p$  be a constant, and  $\Sigma$  be a positive definite matrix of dimension  $p \times p$ . Let

$$w_c = \arg \min_{\|w\|_1 \leq c, w^T \mu_d = 1} w^T \Sigma w;$$

then  $w_c$  is a continuous piecewise linear function in  $c$ .

*Proposition 1.*  $W(\delta_{w_c})$  is a Lipschitz function in  $c$ .

*Proof.* Recall that

$$W(\delta_{w_c}) = 1 - \Phi\{1/R(w_c)^{1/2}\}.$$

By theorem 5 and the fact that composition of Lipschitz functions is again Lipschitz, the conclusion holds.

## 6. Numerical investigation

In this section, several simulation and real data studies are conducted. We compare the ROAD and its variants S-ROAD1 (screening-based ROAD version 1), S-ROAD2 (screening-based ROAD version 2) and DROAD with the NSC method, SCRDA, the FAIR, the NB method and the naive Fisher rule (which uses the generalized inverse of the sample covariance matrix), as well as the oracle.

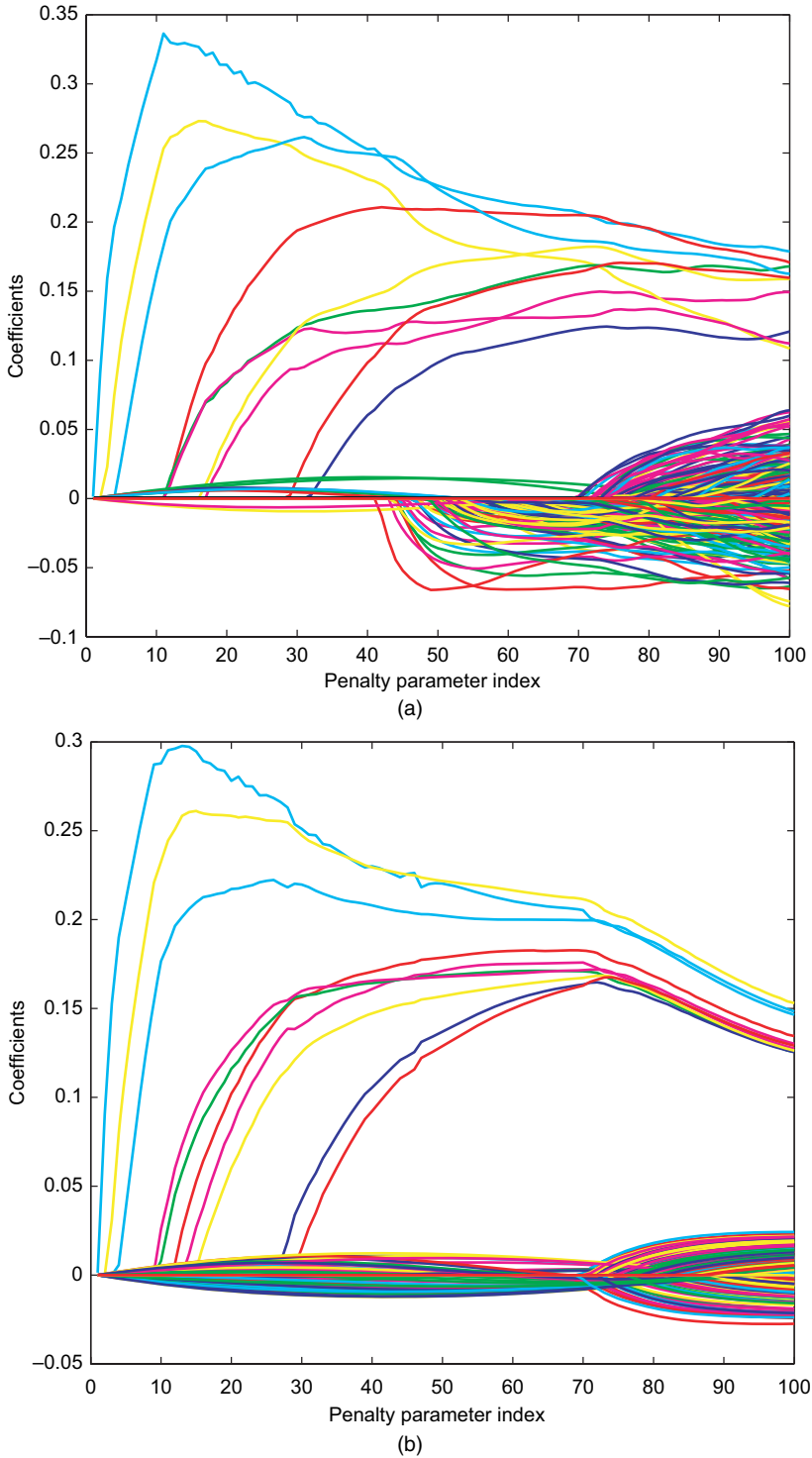
In all simulation studies, the number of variables is  $p = 1000$ , and the sample size of the training and testing data is  $n = 300$  for each class. Each simulation is repeated 100 times to test the stability of the method. Without loss of generality, the mean vector of the first class  $\mu_1$  is set to be  $\mathbf{0}$ . We use fivefold cross-validation to choose the penalty parameter  $\lambda$ .

### 6.1. Equal correlation setting, sparse fixed signal

In this subsection, we consider the setting where  $\Sigma_{i,i} = 1$  for all  $i = 1, \dots, p$  and  $\Sigma_{i,j} = \rho$  for all  $i, j = 1, \dots, p$  and  $i \neq j$ , and take  $\mu_2$  to be a sparse vector:  $\mu_2 = (\mathbf{1}_{10}^T, \mathbf{0}_{990}^T)^T$ , where  $\mathbf{1}_d$  is a length  $d$  vector with all entries 1, and  $\mathbf{0}_d$  is a length  $d$  vector with all entries 0, where the sparsity size is  $s_0 = 10$ . Also, we fix  $\gamma = 10$  in problem (14) for this simulation. Sensitivity of the performance due to the choice of  $\gamma$  will be investigated in the next subsection.

The solution paths for the ROAD and DROAD of one realization are rendered in Fig. 2. It is clear from Fig. 2 that, as the penalty parameter decreases (the index increases), both the ROAD and DROAD use more features. Also, the cut-off point for the DROAD, where the number of features starts to increase dramatically, tends to come later than that for the ROAD.

The simulation results for the pairwise correlations ranging from 0 to 0.9 are shown in Tables 1 and 2. The results for the naive Fisher rule are not included in these (and the subsequent) tables because the test classification error is always around 50%, i.e. it is about the same as a random guess. Also in the tables are the screening-based versions of the the ROAD. S-ROAD1 refers to the version where we first apply the two-sample  $t$ -test to select any features with the corresponding  $t$ -test statistic with absolute value larger than the maximum absolute  $t$ -test statistic value calculated on the permuted data. S-ROAD2 does the same except, for each variable in S-ROAD1's prescreened set, it adds an additional variable which is most correlated with that variable. Fig. 3, which is a graphical summary of Table 1, presents the median test errors for various methods. We can see from Table 1 and Fig. 3 that the oracle classification error decreases as  $\rho$  increases. This is due to a similar reason to the two-dimensional showcase in Section 1. When  $\rho \rightarrow 1$ , all the variables contribute in the same way to boost the classification power. The ROAD performs reasonably close to the oracle, whereas working independence-based methods such as the DROAD, NSC, FAIR and NB methods fail when  $\rho$  is large. The huge discrepancy shows the advantage of employing the correlation structure. Since SCRDA also employs the correlation structure, it does not fail when  $\rho$  is large. However, the ROAD still outperforms SCRDA in all the correlation settings. S-ROAD1 and S-ROAD2 both have misclassification rates which are similar to that of the ROAD. It is worth emphasizing that the merits of the



**Fig. 2.** Solution path for (a) the ROAD and (b) the DROAD: equal correlation setting ( $\rho = 0.5$ ) and sparse signal ( $s_0$ ) as in Section 6.1

**Table 1.** Equal correlation setting and fixed signal: median of the percentage for testing classification error and standard deviations (in parentheses)†

$\rho$	Results for the following methods:								
	ROAD	S-ROAD1	S-ROAD2	DROAD	SCRDA	NSC	FAIR	NB	Oracle
0	6.0 (1.2)	6.0 (1.1)	6.0 (1.2)	5.7 (1.1)	6.3 (1.0)	5.9 (1.0)	5.7 (1.0)	11.2 (1.4)	5.5 (1.1)
0.1	6.3 (2.5)	12.2 (5.0)	8.8 (2.4)	11.6 (5.1)	10.3 (1.4)	11.1 (3.0)	12.4 (1.4)	26.8 (10.1)	5.0 (0.9)
0.2	5.3 (1.0)	16.0 (6.3)	8.7 (2.5)	16.1 (7.5)	8.5 (1.2)	14.5 (4.3)	17.3 (1.7)	34.8 (11.6)	4.0 (0.8)
0.3	4.2 (0.9)	19.1 (7.9)	7.8 (2.6)	19.1 (9.4)	6.6 (1.1)	17.1 (5.5)	20.8 (1.7)	39.3 (12.3)	3.2 (0.7)
0.4	3.2 (0.8)	22.8 (9.4)	6.5 (2.6)	22.2 (9.9)	4.8 (1.0)	20.5 (6.1)	23.2 (1.8)	41.6 (11.3)	2.0 (0.6)
0.5	2.0 (0.6)	25.8 (11.0)	4.8 (1.4)	25.2 (10.2)	2.9 (0.7)	23.2 (6.0)	25.3 (1.6)	43.5 (11.1)	1.3 (0.5)
0.6	1.0 (0.4)	18.3 (12.4)	3.3 (1.3)	28.1 (10.3)	1.5 (0.5)	25.8 (5.7)	26.8 (1.8)	44.4 (12.1)	0.7 (0.3)
0.7	0.3 (0.2)	15.5 (13.6)	1.7 (1.0)	29.1 (10.1)	0.5 (0.3)	27.0 (8.2)	28.2 (2.0)	45.2 (12.3)	0.2 (0.2)
0.8	0.0 (0.1)	5.0 (14.0)	0.3 (0.4)	29.5 (9.9)	0.0 (0.1)	28.3 (8.7)	29.2 (2.0)	46.2 (10.3)	0.0 (0.1)
0.9	0.0 (0.0)	0.6 (14.8)	0.0 (0.1)	30.3 (7.6)	0.0 (0.2)	29.9 (8.0)	30.2 (1.9)	46.8 (8.8)	0.0 (0.0)

†Signals all equal to 1.  $s_0 = 10$ .

**Table 2.** Equal correlation setting and fixed signal: median of number of non-zero coefficients and standard deviations (in parentheses)†

$\rho$	Results for the following methods:						
	ROAD	S-ROAD1	S-ROAD2	DROAD	SCRDA	NSC	FAIR
0	16.00 (24.16)	10.00 (1.31)	17.00 (4.31)	29.50 (58.54)	10.00 (13.25)	10.00 (44.86)	11.00 (1.62)
0.1	117.50 (30.50)	11.00 (3.32)	21.00 (4.15)	14.00 (122.02)	1000.00 (345.48)	35.50 (117.32)	10.00 (0.27)
0.2	130.50 (33.33)	11.00 (6.99)	22.00 (8.98)	15.50 (111.42)	1000.00 (0.00)	95.00 (120.17)	10.00 (0.69)
0.3	136.50 (36.16)	11.00 (11.56)	22.00 (10.38)	17.50 (106.16)	1000.00 (0.00)	103.50 (117.52)	9.00 (1.19)
0.4	135.00 (34.43)	10.00 (14.21)	22.00 (17.07)	10.00 (98.10)	1000.00 (0.00)	70.00 (131.65)	8.00 (1.33)
0.5	138.50 (38.17)	9.00 (21.71)	22.00 (21.56)	10.00 (105.33)	1000.00 (0.00)	65.00 (137.97)	7.00 (1.30)
0.6	148.00 (49.74)	10.50 (27.92)	22.00 (31.88)	10.00 (110.23)	1000.00 (0.00)	38.00 (141.91)	6.00 (1.30)
0.7	170.50 (52.29)	11.00 (37.37)	22.00 (41.76)	1.00 (118.43)	1000.00 (0.00)	27.50 (140.10)	5.00 (1.20)
0.8	203.00 (27.72)	12.00 (50.36)	24.00 (59.23)	1.00 (143.83)	1000.00 (10.92)	15.00 (157.98)	5.00 (1.29)
0.9	151.50 (8.02)	14.00 (55.32)	28.00 (50.45)	1.00 (153.27)	1000.00 (56.30)	14.00 (225.38)	3.00 (1.08)

†Signals all equal to 1.  $s_0 = 10$ .

screening-based ROADS mainly lie in the computation cost, which is reduced significantly by the prescreening step.

The ROAD is a very robust estimator. It performs well even when all the variables are independent, in which case there could be a large amount of noise for fitting the covariance matrix. Table 1 indicates that the ROAD has almost the same performance as the DROAD, NSC and FAIR methods under the independence assumption, i.e.  $\rho = 0$ . As  $\rho$  increases, the edge of the ROAD becomes more substantial. In general, the ROAD is recommended on the grounds that, even with a pairwise correlation of about 0.1 (which is quite common in microarray data as well as financial data), the gain is substantial.

Another interesting observation is that the DROAD performs similarly to the NSC and FAIR methods in terms of classification error. An intuitive explanation is that they are all ‘sparse’ independence rules. NSCs use soft thresholding on the standardized sample mean

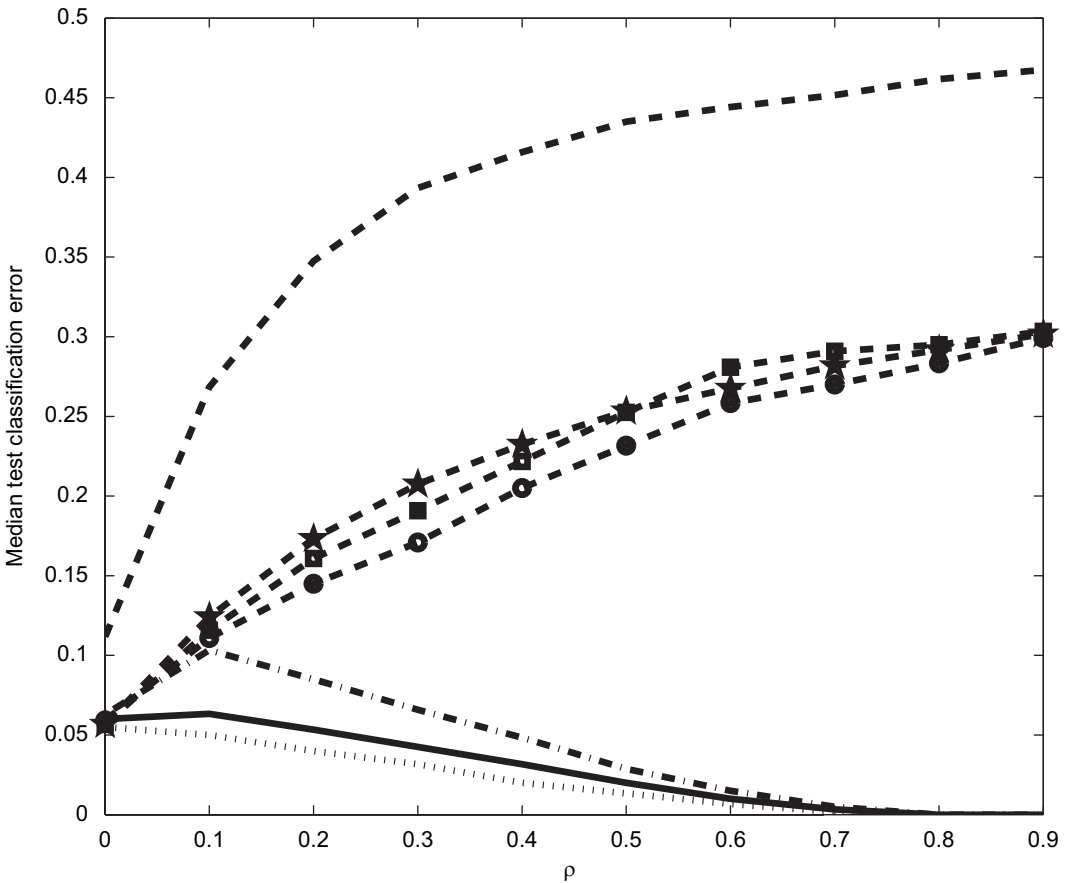


Fig. 3. Median classification error as a function of  $\rho$  in the equicorrelation matrix (sparse  $\mu_d$  as in Section 6.1): —, ROAD; ■, DROAD; - · - ·, SCRDA; ●, NSC; ★, FAIR; - - -, NB; ·····, oracle

difference, and its equivalent lasso derivation can be found in Wang and Zhu (2007). The FAIR method selects features with large marginal  $t$ -statistics in absolute values, whereas the DROAD is another  $L_1$  penalized independence rule, whose implementation is different from NSCs.

Table 2 summarizes the number of features that are selected by different classifiers. Note that the ROAD mimics Fisher discriminant co-ordinate  $\Sigma^{-1}\mu_d$ , which has  $p = 1000$  non-zero entries under our simulated model. Therefore, the large number of features selected by the ROAD is not out of expectation.

### 6.2. Effect of $\gamma$

Under the settings of the previous subsection, we look into the variation of the ROAD's performance as  $\gamma$  changes. In Table 3, the number of active variables varies; however, the median classification error remains about the same for a broad range of  $\gamma$ -values. The reason is that the cross-validation step chooses the 'best'  $\lambda$  according to a specific  $\gamma$ . Therefore, the final performance remains almost unchanged. Since our primary concern is the classification error, we fix  $\gamma = 10$  for simplicity in the subsequent simulations and in the real data analysis.



**Table 3.** Equal correlation setting, signals all equal to 1 and  $s_0 = 10$ : results for the ROAD and various  $\gamma$

$\gamma$	Result for the following values of $\rho$ :		
	$\rho=0$	$\rho=0.5$	$\rho=0.9$
<i>Median classification error (%)</i>			
0.01	5.8 (1.2)	2.7 (0.6)	0.2 (0.2)
0.1	6.0 (1.2)	2.0 (0.6)	0.2 (0.1)
1	6.0 (1.3)	2.0 (0.6)	0.0 (0.1)
10	6.0 (1.2)	2.0 (0.6)	0.0 (0.0)
100	6.2 (1.2)	2.3 (0.6)	0.0 (0.1)
<i>Median number of non-zeros</i>			
0.01	14.0 (19.2)	129.5 (42.5)	657.0 (179.6)
0.1	14.0 (19.6)	137.0 (37.6)	773.5 (103.2)
1	16.5 (22.9)	139.0 (37.9)	514.0 (39.7)
10	16.0 (24.2)	138.5 (38.2)	151.5 (8.0)
100	22.0 (16.1)	114.5 (9.4)	94.0 (9.6)

**6.3. Block diagonal correlation setting, sparse fixed signal**

In this subsection, we follow the same set-up as in Section 6.1 except that the covariance matrix  $\Sigma$  is taken to be block diagonal. The first block is a  $20 \times 20$  equicorrelated matrix and the second block is a  $(p - 20) \times (p - 20)$  equicorrelated matrix, both with pairwise correlation  $\rho$ . In other words,  $\Sigma_{i,i} = 1$  for all  $i = 1, \dots, p$ ,  $\Sigma_{i,j} = \rho$  for all  $i, j = 1, \dots, 20$  and  $i \neq j$ ,  $\Sigma_{i,j} = \rho$  for all  $i, j = 21, \dots, p$  and  $i \neq j$ , and the rest of the elements are 0s. As before, we examine the performances of various estimators when  $\rho$  varies. The percentages for testing error and the number of selected features in the estimators are shown in Tables 4 and 5 respectively.

In this block diagonal setting, we have observed similar results to those in Section 6.1: the ROAD and S-ROAD2 perform significantly better than the other methods. One interesting phenomenon is that S-ROAD1 does not perform well when  $\rho$  is large. The reason is that the

**Table 4.** Block diagonal correlation setting and sparse fixed signal: median of the percentage for testing classification error and standard deviations (in parentheses)†

$\rho$	Results for the following methods:								
	ROAD	S-ROAD1	S-ROAD2	DROAD	SCRDA	NSC	FAIR	NB	Oracle
0.1	6.0 (1.2)	6.0 (1.1)	6.0 (1.2)	5.7 (1.1)	6.0 (0.1)	5.5 (0.3)	5.7 (1.0)	11.2 (1.4)	5.5 (1.1)
0.1	10.8 (3.6)	13.0 (4.8)	10.3 (3.0)	12.8 (4.4)	13.0 (0.3)	12.5 (0.8)	12.7 (1.5)	25.7 (7.6)	8.8 (1.2)
0.2	10.7 (4.1)	18.0 (5.7)	9.7 (3.6)	17.7 (5.9)	14.2 (1.1)	17.2 (0.4)	17.7 (1.6)	34.4 (7.9)	8.8 (1.2)
0.3	9.5 (3.8)	23.2 (5.5)	8.8 (4.0)	23.2 (5.6)	12.7 (0.9)	20.0 (0.8)	20.4 (1.6)	38.3 (7.5)	7.7 (1.0)
0.4	8.0 (3.3)	29.7 (4.2)	7.5 (4.2)	29.3 (4.1)	11.0 (1.2)	23.8 (1.3)	23.2 (1.8)	41.0 (6.9)	6.6 (1.1)
0.5	6.2 (2.6)	30.1 (3.9)	5.7 (0.9)	30.0 (3.1)	8.7 (0.4)	26.2 (1.7)	25.1 (1.7)	42.2 (6.6)	5.0 (1.0)
0.6	4.2 (0.9)	30.3 (4.2)	4.0 (0.8)	30.3 (2.2)	6.4 (0.1)	26.5 (1.2)	26.8 (1.8)	43.6 (7.0)	3.5 (0.7)
0.7	2.3 (0.7)	30.0 (6.4)	2.2 (0.7)	30.6 (2.1)	2.5 (0.7)	28.1 (3.2)	28.2 (2.0)	44.2 (6.5)	1.8 (0.6)
0.8	0.8 (0.4)	29.8 (9.8)	0.7 (0.4)	30.6 (2.1)	0.6 (0.4)	29.2 (1.6)	29.2 (2.0)	44.8 (5.7)	0.7 (0.3)
0.9	0.0 (0.1)	29.8 (12.8)	0.0 (0.1)	30.6 (1.9)	0.2 (0.2)	29.2 (1.2)	30.2 (1.9)	45.2 (4.9)	0.0 (0.1)

†Signals all equal to 1.  $s_0 = 10$ .

**Table 5.** Block diagonal correlation setting and fixed signal: median of the number of non-zero coefficients and standard deviations (in parentheses)†

$\rho$	Results for the following methods:						
	ROAD	S-ROAD1	S-ROAD2	DROAD	SCRDA	NSC	FAIR
0	16.00 (24.16)	10.00 (1.31)	17.00 (4.31)	29.50 (58.54)	10.00 (1.15)	10.00 (1.73)	11.00 (1.62)
0.1	48.50 (35.99)	10.00 (2.73)	20.00 (3.77)	14.00 (26.73)	33.00 (17.79)	65.00 (38.84)	18.00 (2.67)
0.2	48.00 (31.48)	10.00 (4.59)	20.00 (5.84)	10.00 (18.23)	38.00 (117.54)	10.00 (16.17)	18.00 (2.77)
0.3	47.50 (42.75)	9.00 (5.28)	20.00 (6.03)	10.00 (11.80)	208.00 (103.94)	10.00 (13.58)	18.00 (3.91)
0.4	40.50 (32.42)	1.00 (4.82)	20.00 (10.08)	1.00 (9.25)	27.00 (90.95)	33.00 (14.22)	17.00 (5.43)
0.5	40.50 (33.23)	1.00 (4.88)	20.00 (10.10)	1.00 (8.51)	24.00 (76.79)	10.00 (1.15)	7.00 (5.98)
0.6	39.50 (30.03)	1.00 (3.74)	20.00 (14.53)	1.00 (5.92)	127.50 (6.36)	6.50 (2.12)	6.00 (5.98)
0.7	40.00 (41.35)	1.00 (4.71)	20.00 (8.07)	1.00 (2.49)	94.50 (2.12)	9.50 (0.71)	5.00 (5.52)
0.8	55.00 (58.67)	1.00 (6.20)	20.00 (18.32)	1.00 (0.93)	58.00 (2.83)	6.00 (5.66)	5.00 (4.84)
0.9	120.00 (30.66)	1.00 (21.29)	20.00 (30.46)	1.00 (0.35)	20.00 (0.00)	8.00 (2.83)	3.00 (3.81)

†Signals all equal to 1.  $s_0 = 10$ .

current true model has 20 important features and, by looking only at the marginal contribution, S-ROAD1 misses some important variables, as shown in Table 4. Indeed, because those features have no expressed mean differences, it does not fully take advantage of highly correlated features. In contrast, S-ROAD2 can pick up all the important variables, takes advantage of correlation structure and leads to a sparser model than the ROAD. In view of the results from this simulation setting and the previous setting, we recommend S-ROAD2 over S-ROAD1.

6.4. Block diagonal negative correlation setting, sparse fixed signal

In this subsection, we again follow a similar set-up to that in Section 6.1. Here, the covariance matrix  $\Sigma$  is taken to be block diagonal with each block size equal to 10. Each block is an equicorrelated matrix with pairwise correlation  $\rho = -0.1$ . In other words,  $\Sigma = \text{diag}(\Sigma_0, \dots, \Sigma_0)$ , where  $\Sigma_0$  is a  $10 \times 10$  equicorrelated matrix with correlation  $-0.1$ . Here,  $\mu_2 = 0.5(\mathbf{1}_5^T, \mathbf{0}_5^T, \mathbf{1}_5^T, \mathbf{0}_{985}^T)^T$  and the sparsity size is  $s_0 = 10$ . As before, we examine the performances of various estimators when  $\rho$  varies. The percentages for testing error and the number of selected features in the estimators are shown in Table 6.

**Table 6.** Block diagonal negative correlation setting, sparse fixed signal: median error and number of non-zero coefficients with standard deviations in parentheses

Method	Error (%)	Non-zero coefficients
ROAD	7.3 (3.4)	168.00 (47.59)
S-ROAD1	16.0 (5.2)	10.00 (2.40)
S-ROAD2	12.7 (3.4)	20.00 (3.58)
DROAD	17.8 (8.0)	15.50 (15.32)
SCRDA	18.5 (1.1)	24.00 (0.58)
NSC	20.8 (0.6)	41.00 (17.90)
FAIR	24.8 (2.1)	59.00 (4.27)
NB	33.5 (2.1)	—
Oracle	3.2 (0.7)	—

**Table 7.** Random-correlation setting, double-exponential signal: median error and number of non-zero coefficients with standard deviations in parentheses

Method	Error (%)	Non-zero coefficients
ROAD	2.0 (0.6)	83.00 (39.54)
S-ROAD1	11.0 (5.2)	4.00 (8.13)
S-ROAD2	5.8 (3.9)	9.00 (10.69)
DROAD	17.0 (2.2)	1.00 (3.89)
SCRDA	5.2 (1.1)	1000.00 (0.00)
NSC	16.2 (1.3)	4.00 (0.58)
FAIR	17.0 (1.6)	1.00 (0.17)
NB	46.2 (2.4)	—
Oracle	1.3 (0.5)	—

6.5. Random-correlation setting, double-exponential signal

To evaluate the stability of the ROAD, we take a random matrix  $\Sigma$  as the correlation structure and use a signal  $\mu$  whose non-zero entries come from a double-exponential distribution. A random-covariance matrix  $\Sigma$  is generated as follows.

- (a) For a given integer  $m$  (here we take  $m = 10$ ), generate a  $p \times m$  matrix  $\Omega$  where  $\Omega_{i,j} \sim \text{Unif}(-1, 1)$ . Then the matrix  $\Omega\Omega^T$  is positive semidefinite.
- (b) Denote  $c_\Omega = \min_i \{(\Omega\Omega^T)_{ii}\}$ . Let  $\Xi = \Omega\Omega^T + c_\Omega \mathbf{I}$ , where  $\mathbf{I}$  is the identity matrix. It is clear that  $\Xi$  is positive definite.
- (c) Normalize the matrix  $\Xi$  to obtain  $\Sigma$  whose diagonal elements are 1.

For the signal, we take  $\mu$  to be a sparse vector with sparsity size  $s = 10$ , and the non-zero elements are generated from the double-exponential distribution with density function

$$f(x) = \exp(-2|x|).$$

Table 7 summarizes the results. It shows that, even under the random-correlation setting and random signals, our ROAD procedure still outperforms other competing classification rules such as SCRDA, NSCs and the FAIR in terms of the classification error.

6.6. Real data

Though the ROAD seems to perform best in a broad spectrum of idealized experiments, it must be tested against reality. We now evaluate the performance of our newly proposed estimator on three popular gene expression data sets: ‘leukaemia’ (Golub *et al.*, 1999), ‘lung cancer’ (Gordon *et al.*, 2002) and the ‘neuroblastoma data set’ (Oberthuer *et al.*, 2006). The first two data sets come with predetermined, separate training and test sets of data vectors. The leukaemia data set contains  $p = 7129$  genes for  $n_1 = 27$  acute lymphoblastic leukaemia and  $n_2 = 11$  acute myeloid leukaemia vectors in the training set. The test set includes 20 acute lymphoblastic leukaemia and 14 acute myeloid leukaemia vectors. The lung cancer data set contains  $p = 12533$  genes for  $n_1 = 16$  adenocarcinoma and  $n_2 = 16$  mesothelioma training vectors, along with 134 adenocarcinoma and 15 mesothelioma test vectors. The neuroblastoma data set, which was obtained via the ‘MicroArray quality control phase-II’ project, consists of gene expression profiles for  $p = 10707$  genes from 251 patients of the German neuroblastoma trials NB90–NB2004, diagnosed between 1989 and 2004. We analysed the gene expression data with

**Table 8.** Classification error and number of selected genes by various methods for the leukaemia data†

<i>Method</i>	<i>Training error</i>	<i>Testing error</i>	<i>Number of genes selected</i>
ROAD	0	1	40
S-ROAD1	0	3	49
S-ROAD2	0	1	66
SCRDA	1	2	264
FAIR	1	1	11
NSC	1	3	24
NB	0	5	7129

†The training and testing samples are of sizes 38 and 34 respectively.

**Table 9.** Classification error and number of selected genes by various methods for the lung cancer data†

<i>Method</i>	<i>Training error</i>	<i>Testing error</i>	<i>Number of genes selected</i>
ROAD	1	1	52
S-ROAD1	1	4	56
S-ROAD2	1	1	54
SCRDA	0	3	2410
FAIR	0	7	31
NSC	0	10	38
NB	6	36	12533

†The training and testing samples are of sizes 32 and 149 respectively.

**Table 10.** Classification error and number of selected genes by various methods for the neuroblastoma data†

<i>Method</i>	<i>Training error</i>	<i>Testing error</i>	<i>Number of genes selected</i>
ROAD	3	33	33
S-ROAD1	22	47	1
S-ROAD2	14	37	9
SCRDA	16	37	1
FAIR	15	44	18
NSC	16	35	41
NB	14	32	10707

†The training and testing samples are of sizes 83 and 163 respectively.

3-year event-free survival, which indicates whether a patient survived 3 years after the diagnosis of neuroblastoma. There are 239 subjects with 3-year event-free survival information available (49 positive and 190 negative). We randomly select 83 subjects (19 positive and 64 negative, which are about a third of the total subjects) as the training set and the rest as the test set. The readers can find more details about the data sets in Golub *et al.* (1999), Gordon *et al.* (2002) and Oberthuer *et al.* (2006).

Following Dudoit *et al.* (2002) and Fan and Fan (2008), we standardized each sample to zero mean and unit variance. The classification results for the ROAD, S-ROAD1, S-ROAD2, SCRDA, FAIR, NSC and NB methods are shown in Tables 8–10. For the leukaemia and lung cancer data, the ROAD performs the best in terms of classification error. For the neuroblastoma data, the NB method performs best; however, it makes use of all 10707 genes, which is not very desirable. In contrast, the ROAD has a competitive performance in terms of classification error and it selects only 33 genes. Although SCRDA has a close performance, the number of selected variables varies considerably for the three data set (264, 2410 and 1). Overall, the ROAD is a robust classification tool for high dimensional data.

## 7. Discussion

With a simple two-class Gaussian model, we explored the bright side of using correlation structure for high dimensional classification. Targeting directly on the classification error, the ROAD employs an unregularized pooled sample covariance matrix and sample mean difference vector without suffering from the curse of dimensionality and accumulation of noise. The sparsity of chosen features is evident in simulations and real data analysis; however, we have not discovered intuitively good conditions on  $\Sigma$  and  $\mu_d$ , such that a certain desirable sparsity pattern of  $\hat{w}_c$  follows. We resolve a part of the problem by introducing screening-based variants of the ROAD, but the precise control of the sparsity size is worth further investigation. Furthermore, we can explore the conditions for model selection consistency.

The MATLAB language software for implementing the ROAD can be downloaded from the MATLAB central file exchange, <http://www.mathworks.com/matlabcentral/fileexchange/33160>.

In this paper, we have restricted ourselves to linear rules. They can be easily extended to non-linear discriminants via transformations such as low order polynomials or spline basis functions. One may also use the popular ‘kernel tricks’ in the machine learning community. See, for example, Hastie *et al.* (2009) for more details. After the features have been transformed, we can apply the ROAD method. One essential technical challenge of the current paper is rooted in a stochastic linear constraint. The precise role of this constraint has not been completely pinned down. In what follows, a preliminary proposal is provided for extending the ROAD to multiclass settings.

### 7.1. Extension to multiclass settings

In this section, we outline an extension of the ROAD to multiclass classification problems. Suppose that there are  $K$  classes and, for  $j = 1, \dots, K$ , the  $j$ th class has mean  $\mu_j$  and common covariance  $\Sigma$ . Denote the overall mean of features by  $\mu_a = K^{-1} \sum_{j=1}^K \mu_j$ . Fisher’s reduced rank approach to multiclass classification is a minimum distance classifier in some lower dimensional projection space. The first step is to find  $s \leq K - 1$  discriminant co-ordinates ( $w_1^*, \dots, w_s^*$ ) that separate the population centroids  $\{\mu_j\}_{j=1}^K$  the most in the projected space  $S = \text{span}(w_1^*, \dots, w_s^*)$ . Then the population centroids  $\mu_j$  and new observation  $\mathbf{X}$  are both projected onto  $S$ . The observation  $\mathbf{X}$  will be assigned to the class whose projected centroid is closest to the projection of

$\mathbf{X}$  onto  $\mathcal{S}$ . Note that it is usually not necessary to compute all  $K - 1$  discriminant co-ordinates whose span is that of all  $K$  population centroids; the process can stop as long as the projected population centroids are well spread out in  $\mathcal{S}$ .

We adopt the above procedure for multiclass classification. However, the large  $p$ -small  $n$  scenario demands regularization in selecting discriminant co-ordinates. Indeed, in Fisher’s proposal the first discriminant co-ordinate  $\mathbf{w}_1^*$  is the solution of

$$\max_{\mathbf{w}} \frac{\mathbf{w}^T \mathbf{B} \mathbf{w}}{\mathbf{w}^T \Sigma \mathbf{w}}, \tag{18}$$

where  $\mathbf{B} = \Psi^T \Psi$ , and the  $j$ th column of  $\Psi^T$  is  $\mu_j - \mu_a$ . A multiple of  $\mathbf{B}$  is the between-class variance matrix. The second discriminant co-ordinate  $\mathbf{w}_2^*$  is the maximizer of  $\mathbf{w}^T \mathbf{B} \mathbf{w} / (\mathbf{w}^T \Sigma \mathbf{w})$  with constraint  $\mathbf{w}_1^{*T} \Sigma \mathbf{w} = 0$ , and the subsequent discriminant co-ordinates are determined analogously.

Since solving problem (18) is the same as looking for the eigenvector of  $\Sigma^{-1/2} \mathbf{B} \Sigma^{-1/2}$  corresponding to the largest eigenvalue, diverging spectrum and noise accumulation must be considered when we work on the sample. To address these issues, we regularize  $\mathbf{w}$  as in the binary case,

$$\min_{\|\mathbf{w}\|_1 \leq c, \mathbf{w}^T \mathbf{B} \mathbf{w} = 1} \mathbf{w}^T \Sigma \mathbf{w}, \tag{19}$$

whose solution is the first regularized discriminant co-ordinate  $\bar{\mathbf{w}}_1^*$ . Here, equation (19) is related to the null space method in Krzanowski *et al.* (1995). The second regularized discriminant co-ordinate is obtained by solving problem (19) with additional constraint  $\bar{\mathbf{w}}_1^{*T} \Sigma \mathbf{w} = 0$ . Other regularized discriminant co-ordinates can be found similarly. With these  $s$  ( $\leq K - 1$ ) regularized discriminant co-ordinates, the classifier is now based on the minimum distance to the projected centroids in the  $s$ -dimensional space that is spanned by  $\{\bar{\mathbf{w}}_j^*\}_{j=1}^s$ .

The implementation and theoretical properties for a multiclass ROAD are interesting topics for future research.

### Acknowledgements

The authors thank the Joint Editor, the Associate Editor and two referees, whose comments have greatly improved the scope and presentation of the paper. The financial support from National Science Foundation grant DMS-0704337 and National Institutes of Health grant R01-GM072611 is greatly acknowledged.

### Appendix A: Proofs

#### A.1. Proof of theorem 1

We now show the first part of theorem 1. Let  $f_0(\mathbf{w}) = \mathbf{w}^T \mu_d / (\mathbf{w}^T \Sigma \mathbf{w})^{1/2}$ ,  $f_1(\mathbf{w}) = \mathbf{w}^T \hat{\mu}_d / (\mathbf{w}^T \Sigma \mathbf{w})^{1/2}$  and  $f_2(\mathbf{w}) = \mathbf{w}^T \hat{\mu}_d / (\mathbf{w}^T \hat{\Sigma} \mathbf{w})^{1/2}$ . Then, it follows easily that

$$|f_0(\mathbf{w}_c) - f_2(\hat{\mathbf{w}}_c)| \leq \Lambda_1 + \Lambda_2,$$

where  $\Lambda_1 = |f_0(\mathbf{w}_c) - f_1(\mathbf{w}_c^{(1)})|$  and  $\Lambda_2 = |f_1(\mathbf{w}_c^{(1)}) - f_2(\hat{\mathbf{w}}_c)|$ . We now bound both terms separately in the following two steps.

##### A.1.1. Step 1 (bound $\Lambda_1$ )

For any  $\mathbf{w}$ , we have

$$\begin{aligned}
 |f_0(\mathbf{w}) - f_1(\mathbf{w})| &\leq \left| \frac{\mathbf{w}^T \boldsymbol{\mu}_d}{(\mathbf{w}^T \boldsymbol{\Sigma} \mathbf{w})^{1/2}} - \frac{\mathbf{w}^T \hat{\boldsymbol{\mu}}_d}{(\mathbf{w}^T \hat{\boldsymbol{\Sigma}} \mathbf{w})^{1/2}} \right| \\
 &\leq \frac{\|\mathbf{w}\|_1 \|\hat{\boldsymbol{\mu}}_d - \boldsymbol{\mu}_d\|_\infty}{\|\mathbf{w}\|_2 \lambda_{\min}^{1/2}(\boldsymbol{\Sigma})} \\
 &\leq \sqrt{\|\mathbf{w}\|_0} \frac{\|\hat{\boldsymbol{\mu}}_d - \boldsymbol{\mu}_d\|_\infty}{\sigma_0} \\
 &= \sqrt{\|\mathbf{w}\|_0} O_p(a_n).
 \end{aligned} \tag{20}$$

Since  $\mathbf{w}_c^{(1)}$  maximizes  $f_1(\cdot)$ , it follows that

$$\begin{aligned}
 f_0(\mathbf{w}_c) - f_1(\mathbf{w}_c^{(1)}) &= f_0(\mathbf{w}_c) - f_1(\mathbf{w}_c) + \{f_1(\mathbf{w}_c) - f_1(\mathbf{w}_c^{(1)})\} \\
 &\leq f_0(\mathbf{w}_c) - f_1(\mathbf{w}_c),
 \end{aligned} \tag{21}$$

and similarly noticing that  $w_c$  maximizes  $f_0(\cdot)$ , we have

$$\begin{aligned}
 f_1(\mathbf{w}_c^{(1)}) - f_0(\mathbf{w}_c) &= f_1(\mathbf{w}_c^{(1)}) - f_0(\mathbf{w}_c^{(1)}) + \{f_0(\mathbf{w}_c^{(1)}) - f_0(\mathbf{w}_c)\} \\
 &\leq f_1(\mathbf{w}_c^{(1)}) - f_0(\mathbf{w}_c^{(1)}).
 \end{aligned} \tag{22}$$

Combining the results of inequalities (21) and (22) and using equation (20), we conclude that

$$\Lambda_1 = |f_0(\mathbf{w}_c) - f_1(\mathbf{w}_c^{(1)})| = O_p\{(s_c \vee s_c^{(1)})a_n\}.$$

By the Lipschitz property of  $\Phi$ ,

$$|\Phi\{f_1(\mathbf{w}_c^{(1)})\} - \Phi\{f_0(\mathbf{w}_c)\}| = O_p\{(s_c \vee s_c^{(1)})a_n\}.$$

### A.1.2. Step 2 (bound $\Lambda_2$ )

Note that  $\mathbf{w}_c^{(1)}$  and  $\hat{\mathbf{w}}_c$  both are in the set  $\{\mathbf{w}: \mathbf{w}^T \boldsymbol{\mu}_d = 1, \|\mathbf{w}\|_1 \leq 1\}$ . Therefore, by definition of minimizers, we have

$$\begin{aligned}
 \mathbf{w}_c^{(1)T} \boldsymbol{\Sigma} \mathbf{w}_c^{(1)} - \hat{\mathbf{w}}_c^T \boldsymbol{\Sigma} \hat{\mathbf{w}}_c &\leq 0, \\
 \hat{\mathbf{w}}_c^T \hat{\boldsymbol{\Sigma}} \hat{\mathbf{w}}_c - \mathbf{w}_c^{(1)T} \hat{\boldsymbol{\Sigma}} \mathbf{w}_c^{(1)} &\leq 0.
 \end{aligned}$$

Consequently,

$$\begin{aligned}
 \mathbf{w}_c^{(1)T} \boldsymbol{\Sigma} \mathbf{w}_c^{(1)} - \hat{\mathbf{w}}_c^T \hat{\boldsymbol{\Sigma}} \hat{\mathbf{w}}_c &= (\mathbf{w}_c^{(1)T} \boldsymbol{\Sigma} \mathbf{w}_c^{(1)} - \hat{\mathbf{w}}_c^T \boldsymbol{\Sigma} \hat{\mathbf{w}}_c) + \hat{\mathbf{w}}_c^T \boldsymbol{\Sigma} \hat{\mathbf{w}}_c - \hat{\mathbf{w}}_c^T \hat{\boldsymbol{\Sigma}} \hat{\mathbf{w}}_c \\
 &\leq \hat{\mathbf{w}}_c^T (\boldsymbol{\Sigma} - \hat{\boldsymbol{\Sigma}}) \hat{\mathbf{w}}_c \\
 &\leq \|\boldsymbol{\Sigma} - \hat{\boldsymbol{\Sigma}}\|_\infty \|\hat{\mathbf{w}}_c\|_1^2 \\
 &\leq c^2 \|\boldsymbol{\Sigma} - \hat{\boldsymbol{\Sigma}}\|_\infty \\
 &= O_p(a_n c^2).
 \end{aligned} \tag{23}$$

By the same argument, we also have

$$\begin{aligned}
 \hat{\mathbf{w}}_c^T \hat{\boldsymbol{\Sigma}} \hat{\mathbf{w}}_c - \mathbf{w}_c^{(1)T} \boldsymbol{\Sigma} \mathbf{w}_c^{(1)} &= (\hat{\mathbf{w}}_c^T \hat{\boldsymbol{\Sigma}} \hat{\mathbf{w}}_c - \mathbf{w}_c^{(1)T} \hat{\boldsymbol{\Sigma}} \mathbf{w}_c^{(1)}) + \mathbf{w}_c^{(1)T} \hat{\boldsymbol{\Sigma}} \mathbf{w}_c^{(1)} - \mathbf{w}_c^{(1)T} \boldsymbol{\Sigma} \mathbf{w}_c^{(1)} \\
 &\leq \mathbf{w}_c^{(1)T} (\hat{\boldsymbol{\Sigma}} - \boldsymbol{\Sigma}) \mathbf{w}_c^{(1)} \\
 &\leq c^2 \|\boldsymbol{\Sigma} - \hat{\boldsymbol{\Sigma}}\|_\infty \\
 &= O_p(a_n c^2).
 \end{aligned} \tag{24}$$

Combination of expressions (23) and (24) leads to

$$|\hat{\mathbf{w}}_c^T \hat{\boldsymbol{\Sigma}} \hat{\mathbf{w}}_c - \mathbf{w}_c^{(1)T} \boldsymbol{\Sigma} \mathbf{w}_c^{(1)}| = O_p(a_n c^2).$$

Let  $g(x) = \Phi(x^{-1/2})$ . The function  $g$  is Lipschitz on  $(0, \infty)$ , as  $g'(x)$  is bounded on  $(0, \infty)$ . Hence,  $|\Phi\{f_2(\hat{\mathbf{w}}_c)\} - \Phi\{f_0(\mathbf{w}_c^{(1)})\}| = O_p(a_n c^2)$ . Thus,

$$\begin{aligned} |W_n(\hat{\delta}_{\mathbf{w}_c}, \boldsymbol{\theta}) - W(\delta_{\mathbf{w}_c}, \boldsymbol{\theta})| &\leq |\Phi\{f_2(\hat{\mathbf{w}}_c)\} - \Phi\{f_0(\mathbf{w}_c^{(1)})\}| + |\Phi\{f_1(\hat{\mathbf{w}}_c^{(1)})\} - \Phi\{f_0(\mathbf{w}_c)\}| \\ &= O_p\{(s_c \vee s_c^{(1)})a_n\} + O_p(a_n c^2) \\ &= O_p(b_n). \end{aligned}$$

We now prove the second result of the theorem 1. Since  $|\hat{\mathbf{w}}_c^T \boldsymbol{\Sigma} \hat{\mathbf{w}}_c - \hat{\mathbf{w}}_c^T \hat{\boldsymbol{\Sigma}} \hat{\mathbf{w}}_c| = O_p(a_n c^2)$ , we have

$$|\Phi\{f_1(\hat{\mathbf{w}}_c)\} - \Phi\{f_2(\hat{\mathbf{w}}_c)\}| = O_p(a_n c^2). \tag{25}$$

By equations (20) and (25), and the first part of theorem 1, we have

$$\begin{aligned} |W(\hat{\delta}_{\mathbf{w}_c}, \boldsymbol{\theta}) - W(\delta_{\mathbf{w}_c}, \boldsymbol{\theta})| &= |\Phi\{f_0(\hat{\mathbf{w}}_c)\} - \Phi\{f_0(\mathbf{w}_c)\}| \\ &\leq |\Phi\{f_0(\hat{\mathbf{w}}_c)\} - \Phi\{f_1(\hat{\mathbf{w}}_c)\}| + |\Phi\{f_1(\hat{\mathbf{w}}_c)\} - \Phi\{f_2(\hat{\mathbf{w}}_c)\}| + |\Phi\{f_2(\hat{\mathbf{w}}_c)\} - \Phi\{f_0(\mathbf{w}_c)\}| \\ &= O_p(\hat{s}_c a_n) + O_p(a_n c^2) + O_p(b_n) \\ &= O_p(d_n). \end{aligned}$$

This completes the proof of theorem 1.

### A.2. Proof of theorem 2

Let  $\mathbf{w}^\lambda = \mathbf{w}_\infty + \gamma^\lambda$ . Then, from the definition of  $\mathbf{w}^\lambda$ , we have

$$\begin{aligned} \gamma^\lambda &= \underset{\boldsymbol{\mu}_d^T \mathbf{w}_\infty + \boldsymbol{\mu}_d^T \gamma = 1}{\operatorname{arg\,min}} R(\mathbf{w}_\infty + \gamma) + \lambda \|\mathbf{w}_\infty + \gamma\|_1 \\ &= \underset{\boldsymbol{\mu}_d^T \gamma = 0}{\operatorname{arg\,min}} f(\gamma), \end{aligned} \tag{26}$$

where  $f(\gamma) = R(\gamma) + \lambda \sum_{k \in K^c} |\gamma_k| + \lambda \sum_{k \in K} (|\mathbf{w}_\infty^k + \gamma_k| - |\mathbf{w}_\infty^k|)$ . In the last statement, we used the fact that

$$\mathbf{w}_\infty^T \boldsymbol{\Sigma} \gamma = \boldsymbol{\mu}_d^T \gamma / (\boldsymbol{\mu}_d^T \boldsymbol{\Sigma}^{-1} \boldsymbol{\mu}_d) = 0.$$

We write  $\gamma$  for  $\gamma^\lambda$  for short in what follows.

By equation (26), we have  $f(\gamma) \leq f(\mathbf{0}) = 0$ . This implies that

$$R(\gamma) \leq \lambda \sum_{k \in K} (|\mathbf{w}_\infty^k| - |\mathbf{w}_\infty^k + \gamma_k|) \leq \lambda \sum_{k \in K} |\gamma_k| \leq \lambda \sqrt{s} \|\gamma\|_2.$$

However,  $R(\gamma) \geq \lambda_{\min}(\boldsymbol{\Sigma}) \|\gamma\|_2^2$ . Bringing the upper and lower bound of  $R(\gamma)$  together, we conclude that

$$\|\gamma\|_2 \leq \frac{\lambda \sqrt{s}}{\lambda_{\min}(\boldsymbol{\Sigma})}.$$

The proof is now complete.

### A.3. Proof of theorem 5

By the positive definiteness of  $\boldsymbol{\Sigma}$ ,  $\boldsymbol{\Sigma}^{-1}$  and  $\boldsymbol{\Sigma}^{-1/2}$  are also positive definite. Let  $\mathbf{v} = \boldsymbol{\Sigma}^{1/2} \mathbf{w}$ ; then the transformation  $\mathbf{v} \mapsto \mathbf{w}$  is linear. Define

$$\mathbf{v}_c = \underset{\|\boldsymbol{\Sigma}^{-1/2} \mathbf{v}\|_1 \leq c, \mathbf{v}^T \bar{\boldsymbol{\mu}}_d = 1}{\operatorname{arg\,min}} \mathbf{v}^T \mathbf{v},$$

where  $\bar{\boldsymbol{\mu}}_d = \boldsymbol{\Sigma}^{-1/2} \boldsymbol{\mu}_d$ . It is enough to show that  $\mathbf{v}_c$  is piecewise linear in  $c$ .

Let  $\Omega_c = \{\mathbf{v} : \|\boldsymbol{\Sigma}^{-1/2} \mathbf{v}\|_1 \leq c\}$  and  $S = \{\mathbf{v} : \mathbf{v}^T \bar{\boldsymbol{\mu}}_d = 1\}$ . When  $c$  is small, the solution set is  $\emptyset$ ; when  $c$  is large, the constraint  $\Omega_c$  is inactive. Denote by ‘ $a$ ’ the smallest ‘ $c$ ’ such that  $\Omega_c \cap S \neq \emptyset$ , and by ‘ $b$ ’ the smallest such that  $\mathbf{v}_c$  are the same for all  $c \geq b$ . Hence we are interested in  $c \in [a, b]$ , when changes in  $c$  actually affect the solution.

Let  $P$  be the projection of the origin  $O$  onto the hyperplane  $S$  in the  $p$ -dimensional space. Let

$$\mathcal{F}_c = \{S_{1,c}^0, \dots, S_{j_0,c}^0; S_{1,c}^1, \dots, S_{j_1,c}^1; \dots; S_{1,c}^{p-1}, \dots, S_{j_{p-1},c}^{p-1}\},$$

where  $S_{j,c}^i$  denotes an  $i$ -dimensional face of  $\Omega_c$ , i.e.  $S_{j,c}^0$  represents a vertex,  $S_{j,c}^1$  an edge and  $S_{j,c}^{p-1}$  a facet. It is clear that  $\mathcal{F}_c$  is a finite set.



Define a mapping  $\varphi : [a, b] \rightarrow \mathbb{Z} \times \mathbb{Z}$ , where  $\varphi(c) = (i, j)$  such that

- (a)  $\mathbf{v}_c \in S_{j,c}^i$  and
- (b)  $i$  is minimal.

By definition, this mapping is single valued.

For any  $c_0 \in (a, b]$ , denote  $D_{c_0} = \{(i, j) | \forall \varepsilon > 0, \exists c \in [c_0 - \varepsilon, c_0) \text{ subject to } \varphi(c) = (i, j)\}$ . The set  $D_{c_0}$  is non-empty because the collection  $\{(i, j) \in \mathbb{Z} \times \mathbb{Z} | S_{j,c}^i \in \mathcal{F}_c\}$  is finite. Then the theorem follows from compactness of  $[a, b]$  and lemma 2, remark 4 and lemma 3.

*Lemma 1.*  $\forall c_0 \in (a, b], \exists \varepsilon > 0$  such that  $\forall (i, j) \in D_{c_0}$  and  $\forall c \in (c_0 - \varepsilon, c_0)$ ,  $P_{j,c}^i \in S_{j,c}^{i_0} \cap S$ , where  $P_{j,c}^i$  is the projection of  $P$  onto  $S \cap \widehat{S}_{j,c}^i$ , and  $\widehat{S}_{j,c}^i$  denotes the  $i$ -dimensional affine space in which  $S_{j,c}^i$  embeds, and  $S_{j,c}^{i_0}$  is the interior of  $S_{j,c}^i$ , where the topology is the natural subspace topology restricted to  $\widehat{S}_{j,c}^i$ .

*Proof.* Fix  $c_0 \in (a, b]$ . For any  $(i, j) \in D_{c_0}$  and  $\bar{\varepsilon} > 0$ , by the definition of  $D_{c_0}$ , there exists  $c' \in [c_0 - \bar{\varepsilon}, c_0)$  such that  $\varphi(c') = (i, j)$ . The minimality of  $i$  in the definition for  $\varphi$  implies that  $\mathbf{v}_{c'} = P_{j,c'}^i \in S_{j,c'}^{i_0}$ , which is in the interior of  $S_{j,c'}^i$ . Therefore,  $P_{j,c'}^i \in S_{j,c'}^{i_0} \cap S$ . By arbitrariness of  $\bar{\varepsilon}$ ,  $\exists (c_n) \nearrow c_0$  such that  $P_{j,c_n}^i \in S_{j,c_n}^{i_0} \cap S$  for all  $n$ .

It can also be shown that  $\{c | P_{j,c}^i \in S_{j,c}^{i_0} \cap S\}$  is connected: let  $P_{j,c_1}^i \in S_{j,c_1}^{i_0} \cap S$  and  $P_{j,c_2}^i \in S_{j,c_2}^{i_0} \cap S$ ,  $c_1 < c_2$ . For any  $c_3 \in (c_1, c_2)$ ,  $P_{j,c_3}^i$  is on the line segment with end points  $P_{j,c_1}^i$  and  $P_{j,c_2}^i$  because  $\widehat{S}_{j,c}^i$  are parallel affine subspaces in  $\mathbb{R}^p$ . Let  $S_{j,\text{cone}}^i := \cup_{c \geq 0} S_{j,c}^{i_0}$ ; then it is a cone. Since  $P_{j,c_1}^i \in S_{j,c_1}^{i_0}$  and  $P_{j,c_2}^i \in S_{j,c_2}^{i_0}$ , we have  $P_{j,c_3}^i \in S_{j,c_3}^{i_0}$ . Then,  $P_{j,c_3}^i \in S_{j,c_3}^{i_0} \cap S \cap \widehat{S}_{j,c_3}^i = S_{j,c_3}^{i_0} \cap S$ . Hence,  $\exists \varepsilon_{ij} > 0$  such that, for all  $c \in [c_0 - \varepsilon_{ij}, c_0)$ ,  $P_{j,c}^i \in S_{j,c}^{i_0}$ . Take  $\varepsilon = \min_{(i,j) \in D_{c_0}} \varepsilon_{ij}$ ; the claim follows.

*Lemma 2.*  $\forall c_0 \in (a, b], D_{c_0}$  is a singleton, and  $\exists \varepsilon' > 0$  such that  $\mathbf{v}_c$  is linear in  $c$  on  $(c_0 - \varepsilon', c_0)$ .

*Proof.* Fix  $c_0 \in (a, b]$ . We claim that, for some  $(i, j) \in D_{c_0}$ , there are positive  $\varepsilon' (\leq \varepsilon$  that validates lemma 1) such that, for any  $c \in (c_0 - \varepsilon', c_0)$ ,  $\mathbf{v}_c = P_{j,c}^i$ . Assume that the claim is not correct; then pick any  $(i, j) \in D_{c_0}$ ; there is a sequence  $\{c_k\}$  ( $c_k \neq c_{k'}$  if  $k \neq k'$ ) converging to  $c_0$  from the left subject to  $\mathbf{v}_{c_k} \neq P_{j,c_k}^i$ . Without loss of generality, take  $\{c_k\} \subset (c_0 - \varepsilon, c_0)$ . Lemma 1 implies that  $P_{j,c_k}^i \in S_{j,c_k}^{i_0} \cap S$ . If  $\mathbf{v}_{c_k} \in S_{j,c_k}^{i'}$ , we would have  $\mathbf{v}_{c_k} = P_{j,c_k}^{i'}$ . Hence  $\mathbf{v}_{c_k} \notin S_{j,c_k}^i$ . By finiteness of the index pairs in  $\mathcal{F}_c$ , there exists  $(i', j') \neq (i, j)$  such that  $\varphi(c) = (i', j')$  for  $c \in \{c_k\}$ , where  $\{c_k\}$  is some subsequence of  $\{c_k\}$ . This implies that  $(i', j') \in D_{c_0}$ , which together with lemma 1 imply that  $\mathbf{v}_c = P_{j',c}^{i'}$  for  $c \in \{c_k\}$ . Therefore

$$\|P_{j',c}^{i'} - P\|_2 < \|P_{j,c}^i - P\|_2$$

for  $c \in \{c_k\}$ .

However, because  $(i, j) \in D_{c_0}$ , there are infinitely many  $c' \in (c_0 - \varepsilon, c_0)$  such that  $\|P_{j,c'}^i - P\|_2 \geq \|P_{j,c'}^{i'} - P\|_2$ . Therefore,

$$g(c) = \|P - P_{j,c}^i\|_2^2 - \|P - P_{j,c}^{i'}\|_2^2$$

changes signs infinitely many times on  $(c_0 - \varepsilon, c_0)$ . This leads to a contradiction because  $P_{j,c}^i$  and  $P_{j,c}^{i'}$  are both linear functions of  $c$ . Hence, the conclusion holds.

To show that  $D_{c_0}$  is a singleton, suppose that it has two distinct elements  $(i, j)$  and  $(i', j')$ . We have shown that  $\mathbf{v}_c = P_{j,c}^i$  and  $\mathbf{v}_c = P_{j',c}^{i'}$  for all  $c$  in a left neighbourhood of  $c_0$  (not including  $c_0$ ). Also we have  $P_{j,c}^i \in S_{j,c}^{i_0}$  and  $P_{j',c}^{i'} \in S_{j',c}^{i'_0}$  by lemma 1. This can be true only when  $S_{j,c}^{i_0} \subset S_{j',c}^{i'_0}$  (or vice versa), but then  $i < i'$ , contradicting minimality in the definition of  $D_{c_0}$ .

*Remark 4.* Similarly,  $\forall c_0 \in [a, b), \exists \varepsilon' > 0$  such that  $\mathbf{v}_c$  is linear in  $c$  on  $(c_0, c_0 + \varepsilon')$ .

*Lemma 3.*  $\mathbf{v}_c$  is a continuous function of  $c$  on  $[a, b]$ .

*Proof.* The continuity follows from two parts.

- (a)  $\forall c_0 \in [a, b), \exists \varepsilon > 0$  such that  $\mathbf{v}_c$  is continuous on  $[c_0, c_0 + \varepsilon)$ . Indeed, let

$$h(c) = \min_{\|\Sigma^{-1/2}\mathbf{v}\|_1 \leq c, \mathbf{v}^T \boldsymbol{\mu}_d = 1} \mathbf{v}^T \mathbf{v}.$$

We know that the mapping  $c \mapsto \mathbf{v}_c (= P_{j,c}^i)$  is linear and hence continuous on  $(c_0, c_0 + \varepsilon)$  for some small  $\varepsilon > 0$ . It only remains to show that the mapping is right continuous at  $c_0$ . Note here that  $h(c) = \|P_{j,c}^i\|_2^2$  for  $c \in (c_0, c_0 + \varepsilon)$ . Let  $L = \lim_{c \downarrow c_0} P_{j,c}^i$ . It is clear that  $L \in S_{j,c_0}^i$ . Because  $L \in \Omega_{c_0} \cap S$ ,  $h(c_0) \leq \|L\|_2^2$ . This inequality must take the equals sign because  $h(\cdot)$  is monotone decreasing, and  $h(c) = \|P_{j,c}^i\|_2^2 \rightarrow \|L\|_2^2$  as  $c$  approaches  $c_0$  from the right. Because  $\mathbf{v}_{c_0}$  is unique,  $\mathbf{v}_{c_0} = L = \lim_{c \downarrow c_0} P_{j,c}^i = \lim_{c \downarrow c_0} \mathbf{v}_c$ .

- (b)  $\forall c_0 \in (a, b], \exists \varepsilon > 0$  such that  $v_c$  is continuous on  $(c_0 - \varepsilon, c_0]$ . Again, it remains to show that there is no jump at  $c_0$ . Let  $(i_{c_0}, j_{c_0}) = \varphi(c_0)$ . Clearly  $P_{j_{c_0}, c_0}^{i_{c_0}} \in S_{j_{c_0}, c_0}^{i_{c_0}}$ . Introduce a notion of parallelism of affine subspaces in  $\mathbb{R}^p$ . We denote  $\widehat{S}_{j_{c_0}, c}^{i_{c_0}} \parallel S$ , if, only by translation,  $\widehat{S}_{j_{c_0}, c}^{i_{c_0}}$  becomes a subset of  $S$  (or vice versa in other situations); use the notation  $\widehat{S}_{j_{c_0}, c}^{i_{c_0}} \not\parallel S$  otherwise.

If  $\widehat{S}_{j_{c_0}, c}^{i_{c_0}} \parallel S$ , for  $c$  in some left neighbourhood of  $c_0$ ,  $P_{j_{c_0}, c}^{i_{c_0}}$  exists and  $P_{j_{c_0}, c}^{i_{c_0}} \in S_{j_{c_0}, c}^{i_{c_0}}$ . Note that  $P_{j_{c_0}, c}^{i_{c_0}} \in \Omega_c \cap S$ , and  $\|P_{j_{c_0}, c}^{i_{c_0}}\|_2 \rightarrow \|P_{j_{c_0}, c_0}^{i_{c_0}}\|_2$  as  $c$  approaches  $c_0$  from the left. Since  $h(\cdot)$  is monotone decreasing, obviously  $h(c) \rightarrow \|P_{j_{c_0}, c_0}^{i_{c_0}}\|_2 = h(c_0)$ . This shows the left continuity of  $h$  at  $c_0$ . Suppose that  $D_{c_0} = \{(i, j)\}$ ; then we know, on a left neighbourhood of  $c_0$  (not including  $c_0$ ),  $v_c = P_{j, c}^i$ . Let  $E = \lim_{c \uparrow c_0} P_{j, c}^i$ ; then  $E \in \Omega_{c_0} \cap S$ . Note that  $\|P_{j_{c_0}, c}^{i_{c_0}}\|_2 \geq \|P_{j, c}^i\|_2$  for all  $c$  in  $c_0$ 's left neighbourhood, so we have  $\|P_{j_{c_0}, c}^{i_{c_0}}\|_2 \geq \|E\|_2$ . In contrast,  $\|P_{j_{c_0}, c_0}^{i_{c_0}}\|_2 \leq \|E\|_2$  by the definition of  $P_{j_{c_0}, c_0}^{i_{c_0}}$ . Also, consider the uniqueness of distance minimizing point in  $\Omega_{c_0} \cap S$  to origin  $O$ ,  $E = P_{j_{c_0}, c_0}^{i_{c_0}}$ , and hence  $v_c$  has left continuity at  $c_0$ .

If  $\widehat{S}_{j_{c_0}, c}^{i_{c_0}} \not\parallel S$ ,  $\exists Q \in \Omega_{c_0 - \varepsilon/2} \cap S$  such that  $Q \neq P_{j_{c_0}, c_0}^{i_{c_0}}$ . When  $c$  goes from  $c_0 - \varepsilon/2$  to  $c_0$ , there is a point  $Q_c \in \Omega_c \cap S$  moving on the line segment from  $Q$  to  $P_{j_{c_0}, c_0}^{i_{c_0}}$ . Therefore,  $h(\cdot)$  is left continuous at  $c_0$ . Replace  $P_{j_{c_0}, c}^{i_{c_0}}$  by  $Q_c$  in the previous paragraph; the left continuity of  $v_c$  at  $c_0$  follows from the same argument.

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