

IMMIGRATE: A Margin-based Feature Selection Method with Interaction Terms

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Abstract

Relief based algorithms have often been claimed to uncover feature interactions. However, it is still unclear whether and how interaction terms will be differentiated from marginal effects. In this paper, we propose IMMIGRATE algorithm by including and training weights for interaction terms. Besides applying the large margin principle, we focus on the robustness of the contributors of margin and consider local and global information simultaneously. Moreover, IMMIGRATE has been shown to enjoy attractive properties, such as robustness and combination with Boosting. We evaluate our proposed method on several tasks, which achieves state-of-the-art results significantly.

1. Introduction

Feature selection is one of the most fundamental problems in machine learning [Fukunaga, 2013]. Due to the simplicity and effectiveness, the Relief algorithm by Kira and Rendell [1992] has been proven to be one of the most successful feature selection algorithms. Following the large margin principle, it is interpreted to be an online learning algorithm that solves a convex optimization problem with a margin-based cost function. Compared with exhaustive or heuristic combinatorial searches, Relief decomposes a complex, global and nonlinear classification task into a simple and local one. Relief is the first algorithm that uses hypothesis-margin to calculate feature weights for the purpose of classification. See Gilad-Bachrach et al. [2004] for a formal definition of the hypothesis-margin. Besides, the same idea are shared

among early Relief-based algorithms, namely a margin is defined by the fixed 1-nearest-neighbor. Kononenko [1994] extended Relief to ReliefF, which uses multiple nearest neighbors to adjust feature weights. Gilad-Bachrach et al. [2004] proposed Simba, which updates the nearest neighbors every time when the feature weights are updated. Sun and Wu [2008] extended Relief from feature selection to feature extraction using local information. Some competitive feature selection methods have also been proposed based on the large hypothesis-margin principle [Crammer et al., 2003; Yang et al., 2008]. In particular, Sun and Li [2006] developed a Relief-based framework to include global information. Based on this framework, IM4E was proposed by Bei and Hong [2015] to balance margin-quantity maximization and margin-quality maximization. Besides Relief-based algorithms, the large margin method has been widely discussed in machine learning [Weinberger and Saul, 2009; Hariharan et al., 2010; Zhu et al., 2016].

However, although feature interactions are indirectly considered in the Relief-based algorithms by normalizing the feature weights, natural effects of association cannot be reflected by feature weights of Relief. For example, Relief and many of its extensions do not tell us whether the cause of a high feature weight is from its linear effect or its interaction with other features [Urbanowicz et al., 2018]. In addition, these methods cannot clearly reveal the influence of interaction terms on the generation capabilities of Relief-based classifiers and in particular, the degree of such influence, which is the motivation of our work.

To this end, *Iterative Max-MIn entropy marGin-maximization with inteRAction TErms* algorithm (IMMIGRATE, henceforth) is proposed in this paper. It has the following novelties. (1) Taking the stability of margin contributors' distribution into account, generalized quadratic form distance ($gd(\vec{x}_i, \vec{x}_j) = |\vec{x}_i - \vec{x}_j|^T \mathbf{W} |\vec{x}_i - \vec{x}_j|$, $\mathbf{W} \geq 0$, $\|\mathbf{W}\|_F^2 = 1$) based framework is proposed to capture interaction terms. (2) An iterative optimization method is designed for minimizing the cost function efficiently with a closed-form solution for matrix update. (3) A novel prediction method is proposed to apply the weight matrix by expected generalized distance. (4) A new classifier(Boosted IMMIGRATE) with stronger generation capability is formed by applying Boosting algorithm. Finally, to efficiently implement IMMIGRATE on high-dimensional dataset, IM4E-IMMIGRATE algorithm is designed for effective selection of interaction terms on them. Gene expression datasets are used to

demonstrate the effectiveness of the IM4E-IMMIGRATE algorithm. What’s more, the computation time of IMMIGRATE is comparable to other popular feature selection method with interaction terms. Experimental results show that IMMIGRATE achieves state-of-the-art results compared with most classifiers. Meanwhile, Boosted IMMIGRATE outperforms other Boosting classifiers significantly. Moreover, proposed IMMIGRATE and distance metric learning [Xing et al., 2003; Weinberger and Saul, 2009] share the similar distance metric form. However, our method is quite different. Besides different frameworks, in particular, we use generalized quadratic form distance instead of distance metric, where \mathbf{W} does not have to be positive defined. And the normalized \mathbf{W} represents weights for corresponding features or interactions, where more important terms have larger weights. Thus, IMMIGRATE can be regarded as generalized metric learning.

The rest of the paper is organized as follows. Section 2 explains the mathematical foundation of Relief algorithms. IMMIGRATE algorithm is proposed in Section 3. Section 4 summarizes and discusses the experiments on different datasets. The paper is concluded in Section 5.

2. Review: Relief Algorithm

The Relief algorithm [Kira and Rendell, 1992] provides a framework for calculating feature weights using a fixed number of instances. Feature weights are usually referred to as feature “scores” and range from -1 to 1. After calculating the feature weights, a certain threshold is set to select relevant features and discard irrelevant features. Relief can be viewed as a convex optimization problem with a cost function whose mathematical expression is shown in Eq. 2.1 if the threshold is set to be 0. Relief minimizes

$$C = \sum_{n=1}^M \left(\vec{w}^T |\vec{x}_n - NH(\vec{x}_n)| - \vec{w}^T |\vec{x}_n - NM(\vec{x}_n)| \right), \quad (2.1)$$

subject to : $\vec{w} \geq 0$ and $\|\vec{w}\|_2^2 = 1$,

where $NH(x)$ is the nearest “hit” (from the same class) of x ; $NM(x)$ is the nearest “miss” (from a different class) of x ; $|\vec{x}_n - NH(\vec{x}_n)|$ calculates the absolute element-wise differences; $\vec{w}^T |\vec{x}_n - NH(\vec{x}_n)|$ is the weighted Manhattan distance. Denote $\vec{u} = \sum_{n=1}^M (|\vec{x}_n - NH(\vec{x}_n)| - |\vec{x}_n - NM(\vec{x}_n)|)$. Minimizing

Algorithm 1 Original Relief Algorithm

State : Feature selection for binary classification

$N \leftarrow$ number of training instances

$A \leftarrow$ number of features(i.e. attributes)

$M \leftarrow$ number of randomly chosen instances out of M to update weight \vec{w}

initial all features weights to 0: $\vec{w} := 0$

for $i := 1$ **to** M **do**

 randomly select a “target” instance x_i

 find its $NH(x_i)$ and $NM(x_i)$ in Eq. 2.1

for $a := 1$ **to** A **do**

$w[a] := w[a] - (x_i[a] - NH(x_i)[a])^2/M + (x_i[a] - NM(x_i)[a])^2/M$

end for

end for

return the vector W of feature scores that estimate the quality of features

Eq. 2.1 can be solved using the Lagrange multiplier method and the Karush-Kuhn-Tucker condition by Kuhn and Tucker [2014], the solution is straightforward: $\vec{w} = (-\vec{u})^+ / \|(-\vec{u})^+\|_2$, where $(\vec{a})^+ = [\max(a_1, 0), \max(a_2, 0), \dots, \max(a_A, 0)]$. This solution to the original Relief algorithm is fundamental to understanding the Relief-based algorithms. The pseudo code of the Relief algorithm is listed in Algorithm 1.

3. IMMIGRATE Algorithm

The framework of IMMIGRATE algorithm is established in this section.

Let $\mathcal{D} = \{z_n | z_n = (\vec{x}_n, y_n) \in \mathcal{R}^{A+1}, \vec{x}_n \in \mathcal{R}^A, y_n \in \{-1, 1\}\}_{n=1}^N$, where N is the number of instances; A is the number of features; \vec{x}_n represents feature vector and y_n is class. Only binary classification is considered in this formulation. Since binary classification is the basic classification task in machine learning, it is often used to test the performance of feature selection algorithms. The definitions in our margin-based framework (e.g., hits and misses) make it easy to extend a binary classification formulation to a multiple classification problem. Our IMMIGRATE implementation in R is applicable for multiple classification tasks. Here, we define the following notations $\mathcal{H}_n = \{j \in \{1, 2, \dots, N\} | z_j \in \mathcal{D}, y_j = y_n \text{ and } j \neq n\}$,

$\mathcal{M}_n = \{j \in \{1, 2, \dots, N\} | z_j \in \mathcal{D}, y_j \neq y_n\}$, where \mathcal{H}_n and \mathcal{M}_n represent the index sets of hits and misses of the instance z_n , respectively.

3.1. Max-Min Entropy Principle

Given a distance metric $d(\vec{x}_i, \vec{x}_j)$ between two instances \vec{x}_i and \vec{x}_j , a hypothesis-margin [Gilad-Bachrach et al., 2004] is defined as $\rho_{n,h,m} = d(\vec{x}_n, \vec{x}_m) - d(\vec{x}_n, \vec{x}_h)$, where $\vec{x}_h, h \in \mathcal{H}_n$ and $\vec{x}_m, m \in \mathcal{M}_n$ represent the nearest hit and nearest miss for instance \vec{x}_n and $\mathcal{H}_n, \mathcal{M}_n$ are the index sets for hits and misses separately. Since the generalized distance metric is unknown due to the unknown feature weights, the nearest hit and nearest miss are undetermined under such a framework. Hence, the method proposed in Sun and Li [2006], Sun and Wu [2008], Sun et al. [2010] and Bei and Hong [2015] is adopted, where the margin is defined as follows.

$$\rho_n = \sum_{m \in \mathcal{M}_n} \beta_{n,m} d(\vec{x}_n, \vec{x}_m) - \sum_{h \in \mathcal{H}_n} \alpha_{n,h} d(\vec{x}_n, \vec{x}_h) \quad (3.2)$$

where $\alpha_{n,h} \geq 0$, $\beta_{n,m} \geq 0$, $\sum_{h \in \mathcal{H}_n} \alpha_{n,h} = 1$, $\sum_{m \in \mathcal{M}_n} \beta_{n,m} = 1$, for $\forall n \in \{1, \dots, N\}$. As in the above design, hidden random variable $\alpha_{n,h}$ represents the probability that \vec{x}_h is the nearest hit of instance \vec{x}_n , while hidden variable $\beta_{n,m}$ indicates the probability that \vec{x}_m is the nearest miss of instance \vec{x}_n . The derivations of $\alpha_{n,h}$ and $\beta_{n,m}$ are closely related to the generalized distance metric.

As proposed, the probabilities $\{\alpha_{n,h}\}$ and $\{\beta_{n,m}\}$ represent some distribution of hits and misses. The stability of margin contributors' distribution of \vec{x}_n can be defined using its hit probabilities $\{\alpha_{n,h}\}$ and miss probabilities $\{\beta_{n,m}\}$. Thus, the hit entropy and miss entropy are respectively defined as $E_{hit} = -\sum_{h \in \mathcal{H}_n} \alpha_{n,h} \log \alpha_{n,h}$ and $E_{miss} = -\sum_{m \in \mathcal{M}_n} \beta_{n,m} \log \beta_{n,m}$.

The following two scenarios help to explain the intuition of using the hit entropy and miss entropy. Scenario A: all neighbors are distributed evenly around the target instance; Scenario B: the neighbor distribution is highly uneven. In particular, one instance is quite close to the target and the rest are quite far away from the target. An easy experiment to test the stability of margin contributors' distribution is to discard one instance from the system and to check the change degree. In scenario A, if the nearest hit is discarded, the margin changes slightly since there are many other hits evenly distributed

around target. However, in scenario B, the disappearance of the nearest hit can largely reduce the margin since its hit probabilities are concentrated at a few hits. Thus, hit probabilities prefer large entropy, such as scenario A. Meanwhile. The miss probabilities prefer small entropy (e.g., scenario B) because the disappearance of the nearest miss can largely increase the margin. In such a max-min entropy framework, the hit entropy should be maximized and the miss entropy should be minimized. This max-min entropy principle [Bei and Hong, 2015] is an extension of the large margin principle and the hit entropy and miss entropy are optimized to be consistent with the large margin principle.

3.2. IMMIGRATE Algorithm

Here, we extend the margin in Eq. 3.2 by using generalized distance metric. To capture feature interactions, the IMMIGRATE algorithm is developed by choosing the distance metric in framework Eq. 3.2 as generalized quadratic form distance. The “IMMIGRATE” stands for *Iterative Max-Min entropy marGin-maximization with inteRAction TErms algorithm.*

The generalized quadratic form distance for instances z_i and z_j with a weight matrix \mathbf{W} is defined as

$$gd(\vec{x}_i, \vec{x}_j) = |\vec{x}_i - \vec{x}_j|^T \mathbf{W} |\vec{x}_i - \vec{x}_j|, \quad (3.3)$$

where $\mathbf{W} \geq 0$ and $\|\mathbf{W}\|_F^2 = 1$. The weight matrix is a natural extension of the weight vector since a weight vector can be represented as a diagonal matrix. The cost function Eq. 3.4 is designed to maximize the generalized

margin under max-min entropy principle.

$$\begin{aligned}
C = & \sum_{n=1}^N \left(\sum_{h \in \mathcal{H}_n} \alpha_{n,h} |\vec{x}_n - \vec{x}_h|^T \mathbf{W} |\vec{x}_n - \vec{x}_h| \right. \\
& \left. - \sum_{m \in \mathcal{M}_n} \beta_{n,m} |\vec{x}_n - \vec{x}_m|^T \mathbf{W} |\vec{x}_n - \vec{x}_m| \right) \\
& + \sigma \sum_{n=1}^N [E_{miss}(z_n) - E_{hit}(z_n)], \tag{3.4} \\
\text{subject to : } & \mathbf{W} \geq 0 \text{ and } \|\mathbf{W}\|_F^2 = 1, \\
& \sum_{h \in \mathcal{H}_n} \alpha_{n,h} = 1, \quad \sum_{m \in \mathcal{M}_n} \beta_{n,m} = 1 \quad \forall n, \\
& \alpha_{n,h} \geq 0 \quad \beta_{n,m} \geq 0 \quad \forall n,
\end{aligned}$$

where $E_{miss}(z_n) = -\sum_{m \in \mathcal{M}_n} \beta_{n,m} \log \beta_{n,m}$, $E_{hit}(z_n) = -\sum_{h \in \mathcal{H}_n} \alpha_{n,h} \log \alpha_{n,h}$ and σ, λ are both tune parameters. $\|\mathbf{W}\|_F^2$ is the Frobenius norm of \mathbf{W} .

$\|\mathbf{W}\|_F^2 = \sum_{i,j} w_{i,j}^2 = \sum_i \lambda_i^2$, with λ_i 's are eigenvalues of matrix \mathbf{W} .

Now an iterative optimization framework is proposed and three steps are included to iteratively minimize the cost function. The framework starts from a randomly generated weight matrix as the prior one and ends up until the change of cost reaches a preset limit, and then the posterior one is used as the final output.

Step 1: The optimization of cost function Eq. 3.4 starts from a randomly initialized \mathbf{W} (satisfying $\mathbf{W} \geq 0$ and $\|\mathbf{W}\|_F^2 = 1$). Then following two steps are iterated to minimize the cost function. Step 2: Fix \mathbf{W} , update $\{\alpha_{n,h}\}$ and $\{\beta_{n,m}\}$. Step 3: Fix $\{\alpha_{n,h}\}$ and $\{\beta_{n,m}\}$, update \mathbf{W} .

3.2.1. Fix \mathbf{W} , Update $\{\alpha_{n,h}\}$ and $\{\beta_{n,m}\}$

Fixing \mathbf{W} , $\alpha_{n,h}$ and $\beta_{n,m}$ can be obtained from Eq. 3.5, where $gd(\vec{x}_i, \vec{x}_j) = |\vec{x}_i - \vec{x}_j|^T \mathbf{W} |\vec{x}_i - \vec{x}_j|$ is the newly defined distance metric.

$$\begin{aligned}\alpha_{n,h} &= \exp\left(\frac{-gd(\vec{x}_n, \vec{x}_h)}{\sigma}\right) / \sum_{j \in \mathcal{H}_n} \exp\left(\frac{-gd(\vec{x}_n, \vec{x}_j)}{\sigma}\right), \\ \beta_{n,m} &= \exp\left(\frac{-gd(\vec{x}_n, \vec{x}_m)}{\sigma}\right) / \sum_{k \in \mathcal{M}_n} \exp\left(\frac{-gd(\vec{x}_n, \vec{x}_k)}{\sigma}\right),\end{aligned}\tag{3.5}$$

The derivative of the cost function with respect to $(\alpha_{n,h}, \beta_{n,m})$ is

$$\frac{\partial^2 C}{\partial(\alpha_{n,h}, \beta_{n,m})} = \begin{pmatrix} \sigma/\alpha_{n,h} & \partial^2 C / \partial\beta_{n,m} \alpha_{n,h} \\ \partial^2 C / \partial\beta_{n,m} \alpha_{n,h} & -\sigma/\beta_{n,m} \end{pmatrix},\tag{3.6}$$

$$\left| \frac{\partial^2 C}{\partial(\alpha_{n,h}, \beta_{n,m})} \right| = -\frac{\sigma^2}{(\alpha_{n,h} \beta_{n,m})} - \left(\frac{\partial^2 C}{\partial\beta_{n,m} \alpha_{n,h}} \right)^2 < 0.\tag{3.7}$$

Therefore, when fixing \mathbf{W} , a saddle point in $(\alpha_{n,h}, \beta_{n,m})$ space can be found.

3.2.2. Fix $\{\alpha_{n,h}\}$ and $\{\beta_{n,m}\}$, Update \mathbf{W}

Fixing $\alpha_{n,h}$ and $\beta_{n,m}$, the derivation of \mathbf{W} is a little computationally arduous. However, a closed form solution for \mathbf{W} is derived in Theorem 1.

Theorem 3.1. *Fixing $\{\alpha_{n,h}\}$ and $\{\beta_{n,m}\}$, the cost function Eq. 3.4 has a closed-form solution for updating \mathbf{W} .*

$$\Sigma = \sum_{n=1}^N \Sigma_{n,H} - \Sigma_{n,M}, \quad \Sigma \psi_i = \mu_i \psi_i,\tag{3.8}$$

where $\Sigma_{n,H} = \sum_{h \in \mathcal{H}_n} \alpha_{n,h} |\vec{x}_n - \vec{x}_h| |\vec{x}_n - \vec{x}_h|^T$, $\Sigma_{n,M} = \beta_{n,m} |\vec{x}_n - \vec{x}_m| |\vec{x}_n - \vec{x}_m|^T$, and $\|\psi_i\|_2^2 = 1$, $\mu_1 \leq \mu_2 \leq \dots \leq \mu_A$. ψ_i 's and μ_i 's are the eigenvectors and eigenvalues of Σ separately.

$$\mathbf{W} = \Phi \Phi^T,\tag{3.9}$$

where $\Phi = (\sqrt{\eta_1}\psi_1, \sqrt{\eta_2}\psi_2, \dots, \sqrt{\eta_A}\psi_A)$, $\sqrt{\eta_i} = \sqrt{(-\mu_i)^+ / \sqrt{\sum_{i=1}^A ((-\mu_i)^+)^2}}$.

The proof of Theorem 1 is in the supplementary material.

Under the iterative optimization framework, start with Step 1, the optimization algorithm iteratively executes Steps 2 and 3 until convergence. The hyperparameter σ is tuned by cross-validation. In application, when the change of objective cost function is less than the preset limit, the iterative optimization will stop. This procedure is shown computationally efficient for IMMIGRATE algorithm.

Algorithm 2 IMMIGRATE Algorithm

State : Feature selection for binary classification

$N \leftarrow$ number of training instances

$A \leftarrow$ number of features(i.e. attributes)

Input : a training dataset $\{z_n = (\vec{x}_n, y_n)\}_{n=1, \dots, N}$

Initialize: Let $t = 0$, randomly initialize $\mathbf{W}^{(0)}$ satisfying nonnegative $\mathbf{W}^{(0)} \geq 0$, and $\|\mathbf{W}^{(0)}\|_F^2 = 1$

1: **repeat**

2: Calculate $\{\alpha_{n,h}^{(t+1)}\}$ and $\{\beta_{n,m}^{(t+1)}\}$ using Eq. 3.5 with $gd(\vec{x}_i, \vec{x}_j) = |\vec{x}_i - \vec{x}_j|^T \mathbf{W}^{(t)} |\vec{x}_i - \vec{x}_j|$

3: Calculate $\mathbf{W}^{(t+1)}$ using Theorem 1 and Eq. 5.15.

4: **until** the change of C in Eq. 3.4 is small enough or the iteration indicator t reaches a preset limit

Return $\mathbf{W}^{(t)}$

3.2.3. Weight Pruning

The previous Relief-based algorithms remove weights that are lower than a preset threshold. The remaining features are used in K-nearest neighbor classifier to classify new samples. IMMIGRATE adopts a similar method and adds an optional step to prune small weights: Step 4 - Set elements in \mathbf{W} , which are smaller than a preset threshold (empirically, the preset threshold is chosen to be $1/A$, where A is the number of features), to 0, and normalize

the \mathbf{W} so that its Frobenius norm is 1. Here, we set that Step 2, 3, 4 are iterated to minimize the cost function after half of the maximum iteration number(preset limit).

3.2.4. New Prediction Approach

The prediction method is improved by the learned weight matrix \mathbf{W} as expected generalized distance.

$$y' = \arg \min_c \sum_{y_n=c} \alpha_n^c(\vec{x}') gd(\vec{x}', \vec{x}_n), \quad (3.10)$$

where c denotes the class, and

$$\alpha_n^c(\vec{x}') = \frac{\exp(-gd(\vec{x}', \vec{x}_n)/\sigma)}{\sum_{y_k=c} \exp(-gd(\vec{x}', \vec{x}_k)/\sigma)}, \quad (3.11)$$

For a new sample $z' = (\vec{x}', y')$, the learned weight matrix \mathbf{W} is used to select a class for z' using Eq. 3.10 and Eq. 3.11, where $gd(\vec{x}', \vec{x}_n) = |\vec{x}' - \vec{x}_n|^T \mathbf{W} |\vec{x}' - \vec{x}_n|$.

Here, the new samples are divided in the class of the shortest expected generalized distance. Compared with original Relief-based algorithm, the obtained weights for features and interaction terms are exploited by the expected generalized distance in Eq. 3.10 and Eq. 3.11.

Actually, this new prediction method is to maximize the margin contribution of \vec{x}' (Eq. 3.2): where $\mathcal{H}_{\vec{x}'}$ and $\mathcal{M}_{\vec{x}'}$ are the index sets of hits and misses of the new sample \vec{x}' separately. In Eq. 3.10, a class c whose corresponding index set of hits is $\mathcal{H}_{\vec{x}'}$ is chosen. For other classes except c , the corresponding samples are included in the misses. To minimize Eq. 3.10, a class is selected to minimize the second term and maximize the first term, in which the margin contribution of \vec{x}' is maximized.

3.3. Boosting IMMIGRATE

Boosting [Schapire, 1990] has been widely applied in practice by using a set of weak learners to create a strong learner. Meanwhile, Bagging [Breiman,

[1996] is also used to improve the performance of classifiers. There are three widely used algorithms, including Boosting based: Ada Boost [Freund et al. \[1996\]](#); [Freund and Mason \[1999\]](#), XgBoost [Chen and Guestrin \[2016\]](#) and Bagging based: Random Forest [Liaw et al. \[2002\]](#). They have been shown competitive in many applications in terms of prediction accuracy. To use IMMIGRATE as the base classifier in the AdaBoost algorithm [Freund et al. \[1996\]](#), the cost function of IMMIGRATE is changed from Eq. 3.4 to Eq. 3.12 with the same constraints, where $D(x)$ is the corresponding sample weight for sample x . In the ***BOOSTED IMMIGRATE Algorithm*** (**BIM**, Algorithm 3), the maximum iteration number is limited for each IMMIGRATE classifier to create relatively weak learners. Also, according to the experiments, different hyperparameter σ can provide different capabilities for IMMIGRATE classifiers. Thus, for chosen $\sigma_{max} > \sigma_{min}$ and T (number of classifiers for boosting), BIM algorithm uses σ_t , where σ_t starts from σ_{max} and gradually decreases to $\sigma_t \times (\sigma_{min}/\sigma_{max})^{1/T}$ each time until it is not greater than σ_{min} .

$$\begin{aligned}
C = & \sum_{n=1}^N D(\vec{x}_n) \left(\sum_{h \in \mathcal{H}_n} \alpha_{n,h} |\vec{x}_n - \vec{x}_h|^T \mathbf{W} |\vec{x}_n - \vec{x}_h| \right. \\
& \left. - \sum_{m \in \mathcal{M}_n} \beta_{n,m} |\vec{x}_n - \vec{x}_m|^T \mathbf{W} |\vec{x}_n - \vec{x}_m| \right) \\
& + \sigma \sum_{n=1}^N D(\vec{x}_n) [E_{miss}(z_n) - E_{hit}(z_n)], \tag{3.12}
\end{aligned}$$

$$\begin{aligned}
\text{subject to : } & \sum_{n=1}^N D(\vec{x}_n) = 1, \quad D(\vec{x}_n) \geq 0, \quad \forall n, \\
& \mathbf{W} \geq 0 \text{ and } \|\mathbf{W}\|_F^2 = 1, \quad \alpha_{n,h} \geq 0 \quad \beta_{n,m} \geq 0 \quad \forall n, \\
& \sum_{h \in \mathcal{H}_n} \alpha_{n,h} = 1, \quad \sum_{m \in \mathcal{M}_n} \beta_{n,m} = 1 \quad \forall n,
\end{aligned}$$

Algorithm 3 BIM Algorithm

State : Boosted IMMIGRATE for binary classification

$N \leftarrow$ number of training instances

$A \leftarrow$ number of features(i.e. attributes)

$T \leftarrow$ number of classifiers for BIM

Input : a training dataset $\{z_n = (\vec{x}_n, y_n)\}_{n=1, \dots, N}$

Initialize : for each \vec{x}_n , set $D_1(\vec{x}_n) = 1/N$

- 1: **for** $t := 1$ **to** T **do**
- 2: Limit Max Iteration of IMMIGRATE less than preset
- 3: Train weak IMMIGRATE classifier $h_t(x)$ using a chosen σ_t and weights $D_t(x)$ by Eq. 3.12
- 4: Compute the error rate ϵ_t as

$$\epsilon_t = \sum_{i=1}^N D_t(x_i) I[y_i \neq h_t(x_i)]$$

- 5: **if** $\epsilon_t \geq 1/2$ or $\epsilon_t = 0$ **then**
- 6: Discard h_t , $T = T - 1$ and Continue
- 7: **end if**
- 8: Set $\alpha_t = 0.5 \times \log((1 - \epsilon_t)/\epsilon_t)$
- 9: Update $D(x_i)$: For each x_i ,
 $D_{t+1}(x_i) = D_t(x_i) \exp(\alpha_t I[y_i \neq h_t(x_i)])$
- 10: Normalize $D_{t+1}(x_i)$, so that $\sum_{i=1}^N D_{t+1}(x_i) = 1$
- 11: **end for**

Output:

$$h_{final}(x) = \arg \max_{y \in \{0,1\}} \sum_{t: h_t(x)=y} \alpha_t$$

3.4. Apply IMMIGRATE to high-dimensional data

When applied to high-dimensional, IMMIGRATE can incur high computational complexity as it considers the interactions every feature pair. To reduce computational costs, we designed a pipeline IM4E-IMMIGRATE. Firstly, IM4E is applied to learn a weight vector and based to the learned weight vector, we choose features whose corresponding weights are above a preset

threshold (empirically, the preset threshold is chosen to be $2/A$, where A is the number of features). The selected weights are used to initialize the diagonal elements in the weight matrix of IMMIGRATE. Finally, IMMIGRATE is executed. IM4E-IMMIGRATE is a sub-optimal, however, effective and efficient solution to apply IMMIGRATE to high dimensional data. It can also be boosted to produce a stronger algorithm.

4. Experiments

IMMIGRATE has been compiled in R and the corresponding R package can be found on [CRAN](#) and [Github](#).

4.1. Results on Synthetic Dataset

In this experiment, the robustness of IMMIGRATE algorithm is tested using synthesized dataset where two features are purposely designed for interaction term selection. A feature selection algorithm is called robust if the results obtained from original dataset are consistent with the ones from datasets with noises. We generate dataset1 with 200 samples as follows. 100 samples with class 0 and 100 samples with class 1 are randomly generated from Gaussian distributions with mean $[4,2]$, variance $\text{diag}[1,1]$ and with mean $[6,0]$, variance $\text{diag}[1,1]$ separately. Noises with class 0 and class 1 are randomly generated from a Gaussian distribution with mean $[8,-2]$, variance $\text{diag}[8,8]$ and with mean $[2,4]$, variance $\text{diag}[8,8]$, respectively. The scatter plot of dataset with 10% noise is shown in Fig. 1. The noises are designed to disturb the detection of the interaction term. The level of noises starts from 0.05, and gradually increases 0.05 each time until greater than 0.5. IMMIGRATE and LFE are run on the synthesized dataset and the weights corresponding to the interaction term between features 1 and 2 are collected. Interaction weights learned by IMMIGRATE and LFE are plotted in Fig. 2, which clearly shows that IMMIGRATE is more robust than LFE.

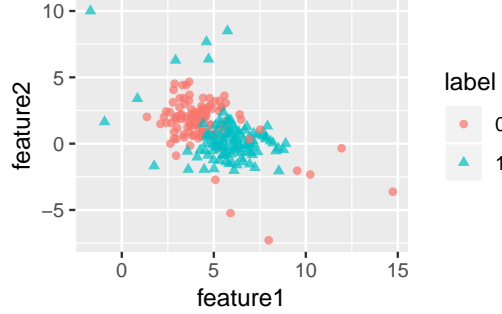


Figure 1: Synthesized Dataset1 with 10% noise.

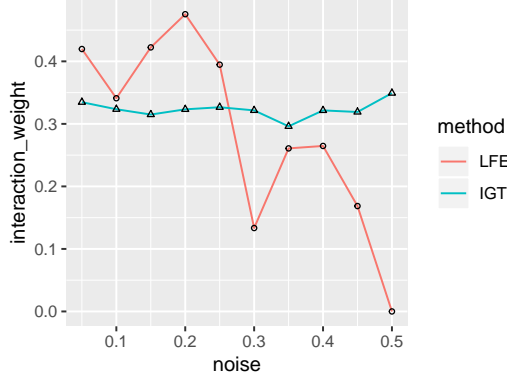


Figure 2: IMMIGRATE is more robust than LFE when learning the weights of interaction from noisy datasets

4.2. Results on Real Datasets

IMMIGRATE is also compared with several existing popular methods using real datasets from UCI database. Cross validation is used to test the performance. The following existing algorithms are used in this experiment: Support Vector Machine [Soentpiet et al., 1999] with Sigmoid Kernel (SV1), with Radial basis function Kernel (SV2), LASSO [Tibshirani, 1996] (LAS), Decision Tree [Freund and Mason, 1999] (DT), Naive Bayes Classifier [John and Langley, 1995] (NBC), Radial basis function Network [Haykin, 1994] (RBF), 1-Nearest Neighbor [Aha et al., 1991] (1NN), 3-Nearest Neighbor (3NN), Large Margin Nearest Neighbor [Weinberger and Saul, 2009] (LMN), Relief [Kira and Rendell, 1992] (REL), ReliefF [Kononenko, 1994; Robnik-Šikonja and Kononenko, 2003] (RFF), Simba [Gilad-Bachrach et al., 2004]

Table 1: Summarizes the accuracies on five high-dimensional gene expression datasets.

Data	SV1	SV2	LAS	DT	NBC	1NN	3NN	SOD	RF	XGB	IM4	EGT	B4G
GLI	85.1	86.0	85.2	83.8	83.0	88.7	87.7	88.7	87.6	86.3	87.5	89.1	89.9
COL	73.7	82.0	80.6	69.2	71.1	72.1	77.9	78.1	82.6	79.5	84.3	78.6	82.5
ELO	72.9	90.2	74.6	77.3	76.3	85.6	91.3	86.9	79.2	77.9	88.9	88.6	88.4
BRE	76.0	88.7	91.4	76.4	69.4	83.0	73.6	82.6	86.3	87.3	88.1	90.2	91.5
PRO	71.3	69.9	87.9	86.4	68.0	83.2	82.7	83.2	91.8	90.5	88.0	89.5	89.7
W,T,L ¹	0,0, 5	1,0, 4	0,1, 4	0,0, 5	0,0, 5	0,0, 5	1,0, 4	0,0, 5	1,1, 3	1,0, 4	1,1, 3	-, -, -	-, -, -

¹ The last row shows the number of times each method W,T,L (win,tie,loss) compared with Boosted IM4E-IMMIGRATE(**B4G**) by paired t -test.

(SIM), Linear Discriminant Analysis [Fisher, 1936] (LDA). In addition, several methods designed for detecting interaction terms are included: LFE [Sun and Wu, 2008], Stepwise conditional likelihood variable selection for Discriminant Analysis [Li and Liu, 2018] (SOD), hierNet [Bien et al., 2013] (HIN). What’s more, for the comparison among Boosting methods, three most widely used and competitive ones are used: Adaptive Boosting Freund et al. [1996]; Freund and Mason [1999] (ADB), Random Forest Liaw et al. [2002] (RF), and XgBoost Chen and Guestrin [2016] (XGB). In the discussion of the experimental results, IM4E is abbreviated as IM4, IMMIGRATE as IGT, BIM as BIM and Boosted IM4E-IMMIGRATE as B4G.

These methods are set in the same way suggested in original papers: LMNN uses 3-NN classifier; Relief and Simba use Euclidean distance and 1-NN classifier; ReliefF use Manhattan distance and k-NN classifier (k=1,3,5 is decided by internal cross-validation); in SODA, gam (=0,0.5,1) is determined by internal cross-validation and logistic regression is used for prediction. The IM4E algorithm owns two hyperparameters λ and σ , where we fix λ as 1 since it has no actual contribution and tune σ as suggested in Bei and Hong [2015]. The IMMIGRATE algorithm has one hyperparameter σ . When tuning σ , σ starts from $\sigma_0 = 4$ and gradually decreases to half each time until it is not larger than 0.2. For large-scale datasets, we choose σ which gives us the best results. The BIM algorithm uses σ_t , $\sigma_{max} = 4$, $\sigma_{min} = 0.2$ are chosen, the number of classifiers: $T = 100$, σ_t starts from σ_{max} and gradually decreases to $\sigma_t \times (\sigma_{min}/\sigma_{max})^{1/T}$ each time until it is not greater than σ_{min} . The preset threshold in IM4E-IMMIGRATE is $1/A$, where A is the number

of features.

4.2.1. Results on Gene Expression Datasets

Gene expression datasets typically have thousands of features. Comparison on five publicly available gene expression datasets are carried out: GLI[Freije et al., 2004], Colon[Alon et al., 1999](COL), Myeloma[Tian et al., 2003](ELO), Breast[Van’t Veer et al., 2002](BRE), Prostate[Singh et al., 2002](PRO). All five datasets have more than ten thousand features. Feature selection methods are widely tested in these high-dimensional datasets.

10-fold cross-validation is performed for ten times, namely 100 trials are carried out. The average accuracy is reported on the corresponding datasets in Table 1. The last row “(W,T,L)” indicates the number of times each algorithm W,T,L (win,tie,loss) when compared with Boosted IM4E-IMMIGRATE (B4G) by the paired Student’s t -test with the significance level of $\alpha = 0.05$. Algorithm A is significantly better than (i.e. win) another algorithm B on a dataset C if the p -value of the paired Student’s t -test with corresponding null hypothesis is less than $\alpha = 0.05$.

As shown in Table 1, although Boosted IM4E-IMMIGRATE (B4G) is not always the best, it outperforms other methods in most cases. In particular, when comparing IM4E-IMMIGRATE(EGT) with other methods, it also outperforms in most cases.

4.2.2. Results on UCI Datasets

UCI datasets from Frank and Asuncion [2010] are used to compare the performance of a cohort of algorithms. The used datasets include Breast Cancer Wisconsin (Prognostic) (BCW), Cryotherapy (CRY), Wholesale customers (CUS), Ecoli (ECO), Glass Identification (GLA), Haberman’s Survival (HMS), Immunotherapy (IMM), Ionosphere (ION), Lymphograph (LYM), MONK’s Problems (MON), Parkinsons (PAR), Pima-Indians-Diabetes (PID), Connectionist Bench (Sonar, Mines vs. Rocks) (SMR), Statlog (Heart) (STA), Urban Land Cover (URB), User Knowledge Modeling (USE) and Wine (WIN). For datasets with more than two classes, the largest two classes are used in this experiment. In addition, we use three large-scale data with respect to the sample size: Waveform Database Generator (WAV*, 3304 samples),

Crowdsourced Mapping (CRO*, 9903 samples) and Electrical Grid Stability Simulated (ELE*, 10000 samples).

10-fold cross-validation is also performed for ten times. Tables 2 and 3 show the average accuracy on the corresponding datasets. The last row “(W,T,L)” indicates the number of times each algorithm W,T,L (win,tie,loss) when compared with IMMIGRATE(IGT) in Table 2 and BIM in Table 3 separately by using the paired Student’s t -test with the significance level of $\alpha = 0.05$.

Although IMMIGRATE or BIM is not always the best, it outperforms other methods significantly in most cases in terms of Cross Validation classification accuracy. Based on Table 2, IMMIGRATE and BIM achieve state-of-the-art performance as base classifier and booster version separately.

Moreover, in view of generalized metric learning, compared with LMNN, better generalized metric for classification is obtained via IMMIGRATE in most cases. To show the results of feature weights, the heat maps of four datasets: GLA, LYM, SMR and STA are supplemented.

5. Conclusion & Discussion

In this paper, a novel feature selection algorithm IMMIGRATE is proposed for detecting and weighting interaction terms, including the extended version of Boosted IMMIGRATE(BIM) and IM4E-IMMIGRATE. Large margin and max-min entropy principle are used to present a generalized quadratic form distance based framework for feature learning. Non-linear margin-based cost function is proposed. To minimize the cost function, an iterative optimization framework is designed for implementing the IMMIGRATE algorithm and the close-form of matrix update is derived in Theorem 1. IMMIGRATE outperforms most methods in tasks and achieve state-of-the-art results. And BIM outperforms other Boosting algorithms. Its robustness is clearly demonstrated on synthetic dataset where we know the ground truth. In conclusion, compared with other Relief-based algorithms, IMMIGRATE mainly has the following advantages: (1) both local and global information are considered; (2) interaction terms are used; (3) robust and less prone to noise; (4) state-of-the-art generalization capabilities; (5) easily boosted. What’s more, the computation time of IMMIGRATE is comparable to other feature selection

Table 2: Summarizes the accuracies on UCI datasets.

Data	SV1	SV2	LAS	DT	NBC	RBF	1NN	3NN	LMN	REL	RFF	SIM	LFE	LDA	SOD	hIN	IM4	IGT
BCW	61.4	66.6	71.4	70.5	62.4	56.9	68.2	72.2	69.5	66.4	67.1	67.7	67.1	73.9	65.2	71.8	66.4	74.5
CRY	72.9	90.6	87.4	85.3	84.4	89.7	89.1	85.4	87.8	73.8	77.2	79.7	86.0	88.6	86.0	87.9	86.2	89.8
CUS	86.5	88.9	89.6	89.6	89.5	86.8	86.5	88.7	88.8	82.1	84.7	84.3	86.4	90.3	90.8	90.3	87.5	90.1
ECO	92.9	96.9	98.6	98.6	97.8	94.6	96.0	97.8	97.8	89.0	90.7	91.2	93.1	99.0	97.9	98.7	97.5	98.2
GLA	64.2	76.7	72.3	79.4	69.5	73.0	81.1	78.1	79.4	64.1	63.5	67.1	81.2	72.0	75.3	75.0	78.0	87.5
HMS	63.8	64.5	67.7	72.5	67.2	66.8	66.0	69.3	71.2	65.3	66.0	65.7	64.9	69.0	67.4	69.4	66.6	69.2
IMM	74.3	70.6	74.4	84.1	77.9	67.3	69.4	77.9	76.7	69.9	71.8	69.0	75.0	75.2	72.3	70.2	80.7	83.8
ION	80.5	93.5	83.6	87.4	89.4	79.9	86.7	84.1	84.5	85.8	86.2	84.2	91.0	83.3	90.3	92.6	88.3	92.9
LYM	83.6	81.5	85.2	75.2	83.6	71.1	77.2	82.8	86.6	64.9	71.0	70.4	79.6	85.2	79.3	84.8	83.3	87.2
MON	74.4	91.7	75.0	86.4	74.0	68.2	75.1	84.4	84.9	61.4	61.8	65.0	64.8	74.4	91.9	97.2	75.6	99.5
PAR	72.7	72.5	77.1	84.8	74.1	71.5	94.6	91.4	91.8	87.3	90.3	84.6	94.0	85.6	88.2	89.5	83.2	93.8
PID	65.6	73.1	74.7	74.3	71.2	70.3	70.3	73.5	74.0	64.8	68.0	67.0	67.8	74.5	75.7	74.1	72.1	74.7
SMR	73.5	83.9	73.6	72.3	70.3	67.1	86.9	84.7	86.1	69.5	78.3	81.0	84.3	73.1	70.5	83.0	76.4	86.5
STA	69.8	71.6	70.8	68.9	71.0	69.5	67.8	70.8	71.3	59.7	64.0	63.0	66.7	71.3	71.8	69.2	70.8	75.9
URB	85.2	87.9	88.1	82.6	85.8	75.3	87.2	87.5	87.9	81.9	83.2	73.0	87.9	73.0	87.9	88.3	87.4	89.9
USE	95.7	95.2	97.2	93.2	90.6	84.9	90.5	91.5	92.0	54.5	63.7	69.5	85.8	96.9	96.2	96.5	94.1	96.4
WIN	98.3	99.3	98.6	93.1	97.3	97.2	96.4	96.6	96.5	87.2	95.0	95.0	93.8	99.7	92.9	98.9	98.2	99.0
CRO*	75.4	97.5	89.9	91.0	88.8	75.4	98.4	98.5	98.6	98.5	98.7	95.1	98.6	89.1	95.2	95.5	81.9	98.2
ELE*	72.3	95.7	79.9	80.0	82.5	70.8	81.1	83.9	89.7	64.6	75.4	76.2	79.8	79.9	93.7	93.6	83.2	93.7
WAV*	90.0	91.9	92.2	86.2	91.4	84.0	86.5	88.3	88.8	77.6	80.0	83.6	84.7	91.8	92.0	92.1	91.1	92.4
W,T,L ¹	0,0,	202 ,2,	161 ,4,	151 ,3,	160 ,1,	190 ,0,	201 ,2,	170 ,2,	181 ,3,	160 ,1,	190 ,1,	190 ,1,	190 ,2,	181 ,4,	153 ,4,	131 ,7,	121 ,0,	19 -, -

¹ The last row shows the number of times each method W,T,L (win,tie,loss) compared with IMMIGRATE (**IGT**) by paired *t*-test.

Table 3: Summarizes the accuracies on UCI datasets.

Data	ADB	RF	XGB	BIM
BCW	78.2	78.6	78.6	78.3
CRY	90.4	92.9	89.9	91.5
CUS	90.8	91.1	91.4	91.0
ECO	98.0	98.9	98.2	98.6
GLA	85.0	87.0	87.9	86.8
HMS	65.8	72.1	70.0	72.0
IMM	77.2	84.2	81.7	86.1
ION	92.1	93.5	92.5	93.1
LYM	84.8	87.0	87.4	88.1
MON	98.4	95.8	99.1	99.7
PAR	90.5	91.0	91.9	93.2
PID	73.5	76.0	75.1	76.2
SMR	81.4	82.8	83.3	86.6
STA	69.0	71.3	69.5	74.1
URB	87.9	88.6	88.8	91.4
USE	96.0	95.3	94.9	96.1
WIN	97.5	99.1	98.2	99.1
CRO*	97.3	97.4	98.5	98.6
ELE*	91.1	92.3	95.2	94.1
WAV*	89.5	91.2	90.8	93.3
W,T,L ¹	0,3, <u>17</u>	1,8, <u>11</u>	2,4, <u>14</u>	-, -, -

¹ The last row shows the number of times each method W,T,L (win,tie,loss) compared with Boosted IMMIGRATE(**BIM**) by paired *t*-test.

methods with interaction terms, such as SODA, hierNet. The well compiled codes for IMMIGRATE and BIM are matrix-based, where parallel computing can be implemented to accelerate.

Several points are required to be further explored in this work. (1) For large-scale datasets, the computation is not quite efficient, which is a common problem for metric learning. Training with well selected prototype [Garcia et al., 2012] might work efficiently. (2) Since the margin-based framework is not restricted to binary classification scenarios, multiple class classification can be achieved easily based on this work. (3) IMMIGRATE only considers pair-wise interactions between features. Interactions among multiple features also play an important role in many cases. Our work provides a basis for exploring new algorithms for detecting interactions among multiple features. (4) In section 3.2.3, small weights are removed to obtain sparse solution. Imposing l_0 or l_1 penalties might also work well to get sparse matrix. l_1 -regularization has been shown by Ng [2004] to have advantages on finding sparse solutions, which is quite valuable for high-dimensional data. (5) Strategy of selecting σ is remaining open to solve. Based on the experimental results, to achieve our best results, different σ 's are chosen for different datasets.

Supplementary Material

Heat Maps

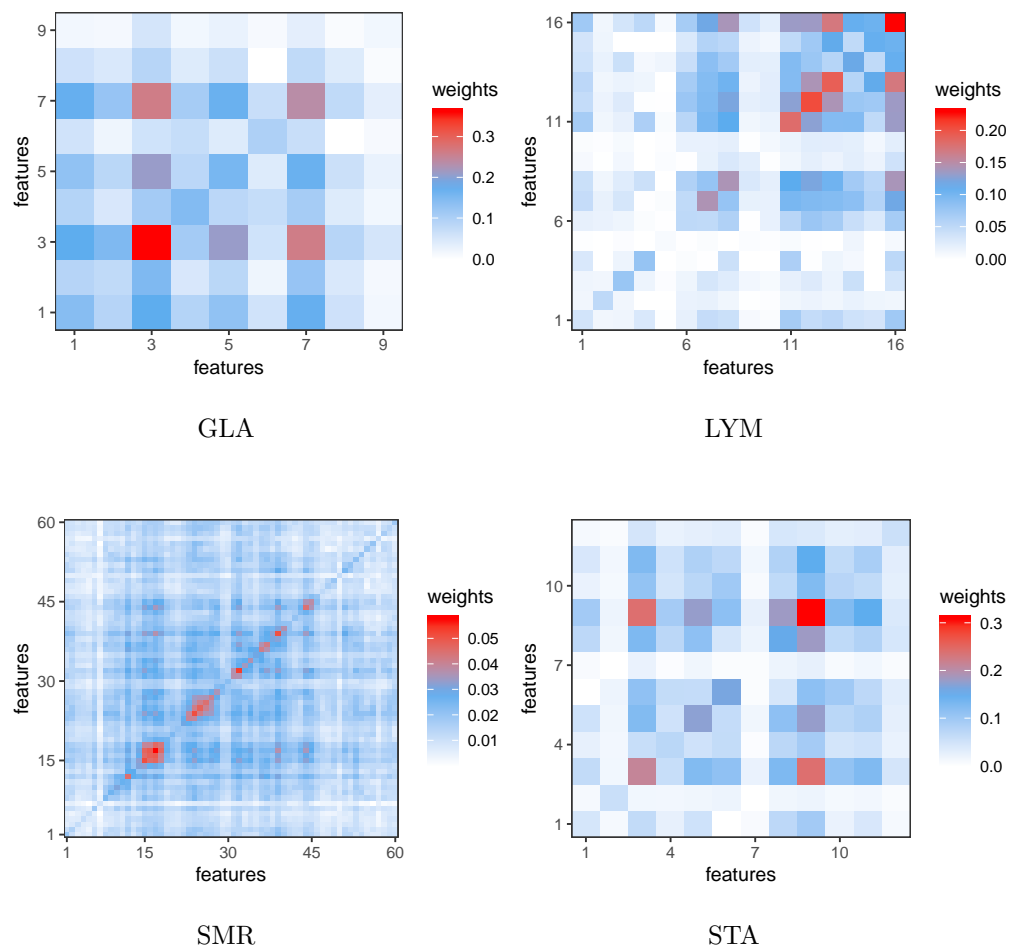


Figure: Heat Maps of Feature Weights Learned by IMMIGRATE.
The color bar shows the value of corresponding colors.

Proof of Theorem 3.1

$$\begin{aligned}
\min_{\vec{w}^T} C &= \sum_{n=1}^N \left(\sum_{h \in \mathcal{H}_n} \alpha_{n,h} |\vec{x}_n - \vec{x}_h|^T \mathbf{W} |\vec{x}_n - \vec{x}_h| - \sum_{m \in \mathcal{M}_n} \beta_{n,m} |\vec{x}_n - \vec{x}_m|^T \mathbf{W} |\vec{x}_n - \vec{x}_m| \right), \\
&+ \sigma \sum_{n=1}^N [E_{miss}(z_n) - E_{hit}(z_n)], \\
&\text{subject to : } \mathbf{W} \geq 0 \text{ and } \|\mathbf{W}\|_F^2 = 1, \\
&\sum_{h \in \mathcal{H}_n} \alpha_{n,h} = 1, \quad \sum_{m \in \mathcal{M}_n} \beta_{n,m} = 1 \quad \forall n,
\end{aligned} \tag{5.13}$$

where $\|\mathbf{W}\|_F^2$ is the Frobenius norm of \mathbf{W} . $\|\mathbf{W}\|_F^2 = \sum_{i,j} w_{i,j}^2 = \sum_i \lambda_i^2$, with λ_i s are eigenvalues of matrix \mathbf{W} .

Theorem 1. *Fixing $\{\alpha_{n,h}\}$ and $\{\beta_{n,m}\}$, the cost function eq.3.4 has a closed-form solution for updating \mathbf{W} .*

$$\Sigma = \sum_{n=1}^N \Sigma_{n,H} - \Sigma_{n,M}, \quad \Sigma \psi_i = \mu_i \psi_i, \tag{5.14}$$

where $\Sigma_{n,H} = \sum_{h \in \mathcal{H}_n} \alpha_{n,h} |\vec{x}_n - \vec{x}_h| |\vec{x}_n - \vec{x}_h|^T$, $\Sigma_{n,M} = \sum_{m \in \mathcal{M}_n} \beta_{n,m} |\vec{x}_n - \vec{x}_m| |\vec{x}_n - \vec{x}_m|^T$, and $\|\psi_i\|_2^2 = 1$, $\mu_1 \leq \mu_2 \leq \dots \leq \mu_A$. ψ_i 's and μ_i 's are the eigenvectors and eigenvalues of Σ separately.

$$\mathbf{W} = \Phi \Phi^T, \tag{5.15}$$

where $\Phi = (\sqrt{\eta_1} \psi_1, \sqrt{\eta_2} \psi_2, \dots, \sqrt{\eta_A} \psi_A)$, $\sqrt{\eta_i} = \sqrt{(-\mu_i)^+ / \sum_{i=1}^A ((-\mu_i)^+)^2}$.

Proof of Thm. 3.1

1. Since \mathbf{W} is distance metric matrix, it is symmetric and positive-semidefinite. Eigenvalue decomposition of \mathbf{W} is

$$\begin{aligned}
\mathbf{W} &= P \Lambda P^T = P \Lambda^{1/2} \Lambda^{1/2} P^T, \\
&= [\sqrt{\lambda_1} p_1, \dots, \sqrt{\lambda_A} p_A] [\sqrt{\lambda_1} p_1, \dots, \sqrt{\lambda_A} p_A]^T,
\end{aligned} \tag{5.16}$$

where \mathbf{P} is orthogonal matrix. Thus, $\langle p_i, p_j \rangle = 0$.

Let $\Phi = [\phi_1, \dots, \phi_A] = [\sqrt{\lambda_1}p_1, \dots, \sqrt{\lambda_A}p_A]$, where $\langle \phi_i, \phi_j \rangle = 0$ and $\lambda_1 > \lambda_2 > \dots > \lambda_A$.

2. The constraint $\|\mathbf{W}\|_F^2 = 1$ can be simplified since \mathbf{W} can be decomposed to be some orthogonal vectors.

$$\|\mathbf{W}\|_F^2 = \sum_{i,j} w_{i,j}^2 = \sum_i (\phi_i^T \phi_i)^2 = 1 \quad (5.17)$$

3. Let us rearrange the Eq. 5.13.

$$\begin{aligned} \sum_{h \in \mathcal{H}_n} \alpha_{n,h} |\vec{x}_n - \vec{x}_h|^T \mathbf{W} |\vec{x}_n - \vec{x}_h| &= \text{tr}(\mathbf{W} \sum_{h \in \mathcal{H}_n} \alpha_{n,h} |\vec{x}_n - \vec{x}_h| |\vec{x}_n - \vec{x}_h|^T), \\ \text{tr}(\mathbf{W} \Sigma_{n,H}) &= \text{tr}(\Sigma_{n,H} \sum_{i=1}^A \phi_i \phi_i^T) = \sum_{i=1}^A \phi_i^T \Sigma_{n,H} \phi_i, \end{aligned} \quad (5.18)$$

Then, Eq. 5.13 can be simplified as follows.

$$\begin{aligned} \min_{\vec{w}^T} C &= \sum_{i=1}^A \phi_i^T \Sigma \phi_i, \\ \text{subject to } : \|\mathbf{W}\|_F^2 &= \sum_i (\phi_i^T \phi_i)^2 = 1, \langle \phi_i, \phi_j \rangle = 0, \end{aligned} \quad (5.19)$$

where $\Sigma = \sum_{n=1}^N \Sigma_{n,H} - \Sigma_{n,M}$ and $\Sigma_{n,H} = \sum_{h \in \mathcal{H}_n} \alpha_{n,h} |\vec{x}_n - \vec{x}_h| |\vec{x}_n - \vec{x}_h|^T$, $\Sigma_{n,M} = \beta_{n,m} |\vec{x}_n - \vec{x}_m| |\vec{x}_n - \vec{x}_m|^T$.

The orthogonal condition will be ignored by us when deriving the closed form solution because as we can notice at the last step, this condition has already been satisfied.

4. The Lagrangian of Eq. 5.19 is easy to obtain.

$$L = \sum_{i=1}^A \phi_i^T \Sigma \phi_i + \lambda \left(\sum_{i=1}^A (\phi_i^T \phi_i)^2 - 1 \right), \quad (5.20)$$

Derive L with respect to ϕ_i ,

$$\partial L / \partial \phi_i = 2\Sigma\phi_i + 4\lambda\phi_i^T\phi_i\phi_i = 0, \quad (5.21)$$

Denote $\phi_i / \|\phi_i\|_2 := \psi_i$. From Eq. 5.21,

$$\Sigma \psi_i = \mu_i \psi_i, \quad (5.22)$$

where $\mu_i = -2\lambda\|\phi_i\|_2^2$. ψ_i and μ_i are the eigenvector and eigenvalue of Σ separately.

5. Let $\phi_i = \sqrt{\eta_i}\psi_i$, $\eta_i \geq 0$. Thus, $C = \sum_{i=1}^A \sqrt{\eta_i}\psi_i^T \Sigma \sqrt{\eta_i}\psi_i = \sum_{i=1}^A \eta_i \mu_i \psi_i^T \psi_i = \sum_{i=1}^A \eta_i \mu_i$, and $\|\mathbf{W}\|_F^2 = \sum_i (\sqrt{\eta_i}\psi_i^T \sqrt{\eta_i}\psi_i)^2 = \sum_i (\eta_i)^2 = 1$,

Then, Eq. 5.19 can be simplified to be

$$\min_{\vec{w}^T} C = \sum_{i=1}^A \eta_i \mu_i, \text{ subject to } \sum_{i=1}^A (\eta_i)^2 = 1, \eta_i \geq 0 \quad (5.23)$$

6. It is excited to notice Eq. 5.23 is exactly the same as the original Relief Algorithm.

$$\vec{\eta} = (-\vec{\mu})^+ / \|(-\vec{\mu})^+\|_2, \quad (5.24)$$

where $(\vec{a})^+ = [\max(a_1, 0), \max(a_2, 0), \dots, \max(a_I, 0)]$. And $\phi_i = \sqrt{\eta_i}\psi_i$

Using $\Phi = [\phi_1, \dots, \phi_A] = [\sqrt{\lambda_1}p_1, \dots, \sqrt{\lambda_A}p_A]$,

$$\mathbf{W} = \Phi\Phi^T \quad (5.25)$$

The orthogonal condition is achieved here because $\|\mathbf{W}\|_F^2 = \sum_i (\phi_i^T \phi_i)^2 = 1$

Aha, D. W., Kibler, D., and Albert, M. K. (1991). Instance-based learning algorithms. *Machine learning*, 6(1):37–66.

Alon, U., Barkai, N., Notterman, D. A., Gish, K., Ybarra, S., Mack, D., and Levine, A. J. (1999). Broad patterns of gene expression revealed by clustering analysis of tumor and normal colon tissues probed by oligonucleotide arrays. *Proceedings of the National Academy of Sciences*, 96(12):6745–6750.

- Bei, Y. and Hong, P. (2015). Maximizing margin quality and quantity. In *Machine Learning for Signal Processing (MLSP), 2015 IEEE 25th International Workshop on*, pages 1–6. IEEE.
- Bien, J., Taylor, J., and Tibshirani, R. (2013). A lasso for hierarchical interactions. *Annals of statistics*, 41(3):1111.
- Breiman, L. (1996). Bagging predictors. *Machine learning*, 24(2):123–140.
- Chen, T. and Guestrin, C. (2016). Xgboost: A scalable tree boosting system. In *Proceedings of the 22nd acm sigkdd international conference on knowledge discovery and data mining*, pages 785–794. ACM.
- Crammer, K., Gilad-Bachrach, R., Navot, A., and Tishby, N. (2003). Margin analysis of the lvq algorithm. In *Advances in neural information processing systems*, pages 479–486.
- Fisher, R. A. (1936). The use of multiple measurements in taxonomic problems. *Annals of eugenics*, 7(2):179–188.
- Frank, A. and Asuncion, A. (2010). Uci machine learning repository [<http://archive.ics.uci.edu/ml>]. irvine, ca: University of california. *School of information and computer science*, 213:2–2.
- Freije, W. A., Castro-Vargas, F. E., Fang, Z., Horvath, S., Cloughesy, T., Liao, L. M., Mischel, P. S., and Nelson, S. F. (2004). Gene expression profiling of gliomas strongly predicts survival. *Cancer research*, 64(18):6503–6510.
- Freund, Y. and Mason, L. (1999). The alternating decision tree learning algorithm. In *icml*, volume 99, pages 124–133.
- Freund, Y., Schapire, R. E., et al. (1996). Experiments with a new boosting algorithm. In *Icml*, volume 96, pages 148–156. Citeseer.
- Fukunaga, K. (2013). *Introduction to statistical pattern recognition*. Elsevier.
- Garcia, S., Derrac, J., Cano, J., and Herrera, F. (2012). Prototype selection for nearest neighbor classification: Taxonomy and empirical study. *IEEE transactions on pattern analysis and machine intelligence*, 34(3):417–435.

- Gilad-Bachrach, R., Navot, A., and Tishby, N. (2004). Margin based feature selection-theory and algorithms. In *Proceedings of the twenty-first international conference on Machine learning*, page 43. ACM.
- Hariharan, B., Zelnik-Manor, L., Varma, M., and Vishwanathan, S. (2010). Large scale max-margin multi-label classification with priors. In *Proceedings of the 27th International Conference on Machine Learning (ICML-10)*, pages 423–430. Citeseer.
- Haykin, S. (1994). *Neural networks: a comprehensive foundation*. Prentice Hall PTR.
- John, G. H. and Langley, P. (1995). Estimating continuous distributions in bayesian classifiers. In *Proceedings of the Eleventh conference on Uncertainty in artificial intelligence*, pages 338–345. Morgan Kaufmann Publishers Inc.
- Kira, K. and Rendell, L. A. (1992). A practical approach to feature selection. In *Machine Learning Proceedings 1992*, pages 249–256. Elsevier.
- Kononenko, I. (1994). Estimating attributes: analysis and extensions of relief. In *European conference on machine learning*, pages 171–182. Springer.
- Kuhn, H. W. and Tucker, A. W. (2014). Nonlinear programming. In *Traces and emergence of nonlinear programming*, pages 247–258. Springer.
- Li, Y. and Liu, J. S. (2018). Robust variable and interaction selection for logistic regression and general index models. *Journal of the American Statistical Association*, pages 1–16.
- Liaw, A., Wiener, M., et al. (2002). Classification and regression by random-forest. *R news*, 2(3):18–22.
- Ng, A. Y. (2004). Feature selection, l_1 vs. l_2 regularization, and rotational invariance. In *Proceedings of the twenty-first international conference on Machine learning*, page 78. ACM.
- Robnik-Šikonja, M. and Kononenko, I. (2003). Theoretical and empirical analysis of relieff and rrelieff. *Machine learning*, 53(1-2):23–69.
- Schapire, R. E. (1990). The strength of weak learnability. *Machine learning*, 5(2):197–227.

- Singh, D., Febbo, P. G., Ross, K., Jackson, D. G., Manola, J., Ladd, C., Tamayo, P., Renshaw, A. A., D’Amico, A. V., Richie, J. P., et al. (2002). Gene expression correlates of clinical prostate cancer behavior. *Cancer cell*, 1(2):203–209.
- Soentpiet, R. et al. (1999). *Advances in kernel methods: support vector learning*. MIT press.
- Sun, Y. and Li, J. (2006). Iterative relief for feature weighting. In *Proceedings of the 23rd international conference on Machine learning*, pages 913–920. ACM.
- Sun, Y., Todorovic, S., and Goodison, S. (2010). Local-learning-based feature selection for high-dimensional data analysis. *IEEE transactions on pattern analysis and machine intelligence*, 32(9):1610–1626.
- Sun, Y. and Wu, D. (2008). A relief based feature extraction algorithm. In *Proceedings of the 2008 SIAM International Conference on Data Mining*, pages 188–195. SIAM.
- Tian, E., Zhan, F., Walker, R., Rasmussen, E., Ma, Y., Barlogie, B., and Shaughnessy Jr, J. D. (2003). The role of the wnt-signaling antagonist dkk1 in the development of osteolytic lesions in multiple myeloma. *New England Journal of Medicine*, 349(26):2483–2494.
- Tibshirani, R. (1996). Regression shrinkage and selection via the lasso. *Journal of the Royal Statistical Society. Series B (Methodological)*, pages 267–288.
- Urbanowicz, R. J., Meeker, M., La Cava, W., Olson, R. S., and Moore, J. H. (2018). Relief-based feature selection: introduction and review. *Journal of biomedical informatics*.
- Van’t Veer, L. J., Dai, H., Van De Vijver, M. J., He, Y. D., Hart, A. A., Mao, M., Peterse, H. L., Van Der Kooy, K., Marton, M. J., Witteveen, A. T., et al. (2002). Gene expression profiling predicts clinical outcome of breast cancer. *nature*, 415(6871):530.
- Weinberger, K. Q. and Saul, L. K. (2009). Distance metric learning for large margin nearest neighbor classification. *Journal of Machine Learning Research*, 10(Feb):207–244.

- Xing, E. P., Jordan, M. I., Russell, S. J., and Ng, A. Y. (2003). Distance metric learning with application to clustering with side-information. In *Advances in neural information processing systems*, pages 521–528.
- Yang, M., Wang, F., and Yang, P. (2008). A novel feature selection algorithm based on hypothesis-margin. *JCP*, 3(12):27–34.
- Zhu, J., Song, J., and Chen, B. (2016). Max-margin nonparametric latent feature models for link prediction. *arXiv preprint arXiv:1602.07428*.