

Optimization Method Based Extreme Learning Machine For Classification

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Abstract

Extreme learning machine (ELM) as an emergent technology has shown its good performance in regression applications as well as in large dataset (and/or multi-label) classification applications. The ELM theory shows that the hidden nodes of the “generalized” single-hidden layer feedforward networks (SLFNs), which needn't be neuron alike, can be randomly generated and the universal approximation capability of such SLFNs can be guaranteed. This paper further studies ELM for classification in the aspect of the standard optimization method and extends ELM to a specific type of “generalized” SLFNs - support vector network. This paper shows that: 1) under the ELM learning framework, SVM's maximal margin property and the minimal norm of weights theory of feedforward neural networks are actually consistent; 2) from the standard optimization method point of view ELM for classification and SVM are equivalent but ELM has less optimization constraints due to its special separability feature; 3) as analyzed in theory and further verified by the simulation results, ELM for classification tends to achieve better generalization performance than traditional SVM. ELM for classification is less sensitive to user specified parameters and can be implemented easily.

Key words: Extreme learning machine, support vector machine, support vector network, ELM kernel, ELM feature space, equivalence between ELM and SVM, maximal margin, minimal norm of weights, primal and dual ELM networks.

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1 Introduction

There are three main approaches in the training of feedforward networks: 1) gradient-descent based (e.g. backpropagation (BP) method [1] for multi-layer feedforward “neural” networks); 2) least square based (e.g. extreme learning machines (ELMs) [2–6] for the “generalized” single-hidden layer feedforward networks (SLFNs)); 3) standard optimization method based (e.g. support vector machines (SVMs) [7] for a specific type of single-hidden layer feedforward networks, the so-called support vector network).

Support vector machines (SVMs) [7] have been extensively used in widespread applications. The main learning feature of SVM is that the standard optimization method is used to find the solution of maximizing the separating margin of two different classes while minimizing the training errors.

Extreme learning machines (ELMs) were originally developed for the single-hidden layer feedforward neural networks [2–4] and then extended to the “generalized” single-hidden layer feedforward networks (SLFNs) which may not be neuron alike [5,6]. ELM proposes to apply random computational nodes in the hidden layer, which may be independent of the training data. ELM [2,3] and its variants [4–6,8–12] mainly focus on the function approximation applications. Different from traditional learning algorithms for neural networks ELM not only tends to reach the smallest training error but also the smallest norm of output weights. Bartlett’s theory [13] shows that for feedforward neural networks reaching smaller training error the smaller the norm of weights is, the better generalization performance the networks tend to have.

Intuitively speaking, SVMs and ELMs do not seem to have much direct relevance although one of the main reasons behind the extension of ELM to “generalized” SLFNs is to possibly apply ELM learning approach in SVM in some way [5,6]. It is also not clear whether there is any close relationship between SVM’s maximal separating margin of two different classes and Bartlett’s minimal size of weights theory on feedforward neural networks [13]. However, Liu, et al [14] and Fréney and Verleysen [15] have made a significant contribution showing that the ELM learning approach can be applied to SVMs directly by simply replacing SVM kernels with (random) ELM kernels and better generalization can be achieved². Random ELM kernels can be applied in SVM, which validates the correctness of ELM’s theory on the “generalized” SLFNs [5,6]. Their proposed SVM with ELM kernel and the conventional SVM have the same optimization constraints. Their work is exciting and has resolved a challenging issue whether ELM can be used for SVM in real applications, however, the reason why ELM for SVM can achieve better generalization performance

² Liu, et al [14] originally applies ELM kernel in PSVM [16] and later Fréney and Verleysen [15] shows that ELM kernel can also be applied in the traditional SVM.

than the conventional SVM has not been answered clearly.

Different from the earlier work on ELM for SVM [14,15] which effectively and simply apply ELM kernels in SVM, this paper studies ELM for classification in the aspect of the standard optimization method and further shows that 1) SVM's maximal separating margin property and the ELM's minimal norm of output weights property are actually consistent and with ELM framework SVM's maximal separating margin property and Barlett's theory on feedforward neural networks remain consistent; 2) in theory ELM can be linearly extended to SVMs (but with less optimization constraints) instead of only replacing SVM kernels with ELM kernels, and thus the learning and implementation of SVMs can be made much simpler and more efficient indeed; 3) since according to the ELM theories [2–6] all the training data are linearly separable by a hyperplane passing through the origin with probability one in the ELM feature space, ELM for classification tends to achieve better generalization performance than SVM.

2 Brief of Support Vector Machine

In order to understand how Cortes and Vapnik's SVM [7] was proposed it may be useful to mention Rosenblatt's feature space on perceptrons [17] first³. As shown in Cortes and Vapnik's SVM [7], SVM can be seen as a specific type of SLFNs, the so-called support vector networks.

2.1 Rosenblatt's feature space

In 1962 Rosenblatt [17] investigated perceptrons (multi-layer feedforward neural networks). Due to lack of learning algorithms to adjust all the weights of perceptrons, Rosenblatt suggested a learning mechanism where only the weights of the connections from the last hidden layer to the output layer were adjusted. After all the rest weights fixed the input data are actually transformed into a feature space Z of the last hidden layer. In this feature space a linear decision function is constructed:

$$f(\mathbf{x}) = \text{sign} \left(\sum_{i=1}^L \alpha_i z_i(\mathbf{x}) \right) \quad (1)$$

where $z_i(\mathbf{x})$ is the output of the i -th neuron in the last hidden layer of a perceptron.

³ One may refer to [7] for the details of the relationship between SVM and Rosenblatt's feature space.

In order to find an alternative solution of $z_i(\mathbf{x})$, in 1995 Cortes and Vapnik [7] proposed the support vector machine which maps the data from the input space to some high dimensional feature space Z through some non-linear mapping chosen *a priori*. Standard optimization methods are used to find the separating hyperplane which maximizes the separating margins of two different classes in the feature space.

2.2 Maximal separating margins in feature space

Given a set of training data (\mathbf{x}_i, t_i) , $i = 1, \dots, N$, where $\mathbf{x}_i \in \mathbf{R}^d$ and $t_i \in \{-1, 1\}$, these training data are usually not separable in the input space in most cases. If the training data cannot be separated by a linear decision function $\mathbf{w} \cdot \mathbf{x} + b = 0$ one can map the training data \mathbf{x}_i from the input space to a feature space Z through a mapping $\phi(\mathbf{x}) : \mathbf{x}_i \rightarrow \phi(\mathbf{x}_i)$. In this feature space, to separate the training data with a minimal training error we have:

$$\begin{aligned} \mathbf{w} \cdot \phi(\mathbf{x}_i) + b &\geq 1 - \xi_i & \text{if } t_i = 1 \\ \mathbf{w} \cdot \phi(\mathbf{x}_i) + b &\leq -1 + \xi_i & \text{if } t_i = -1 \end{aligned} \quad (2)$$

that is

$$t_i(\mathbf{w} \cdot \phi(\mathbf{x}_i) + b) \geq 1 - \xi_i, \quad i = 1, \dots, N \quad (3)$$

Vectors \mathbf{x}_i for which $t_i(\mathbf{w} \cdot \phi(\mathbf{x}_i) + b) = 1$ is termed support vectors. The hyperplane $\mathbf{w} \cdot \phi(\mathbf{x}) + b = 0$ is the unique one which separates the training data with a maximal margin in the feature space: it maximizes the distance $2/\|\mathbf{w}\|$ between two different classes in the feature space Z .

To maximize such separating margin and to minimize the training error is equivalent to:

$$\begin{aligned} \text{Minimize: } L_P &= \frac{1}{2}\|\mathbf{w}\|^2 + C \sum_{i=1}^N \xi_i \\ \text{Subject to: } &t_i(\mathbf{w} \cdot \phi(\mathbf{x}_i) + b) \geq 1 - \xi_i, \quad i = 1, \dots, N \\ &\xi_i \geq 0, \quad i = 1, \dots, N \end{aligned} \quad (4)$$

where C is a user specified parameter and provides a tradeoff between the distance of the separating margin and the training error.

Based on the Karush-Kuhn-Tucker theorem [18], to train such a SVM is equiv-

alent to solving the following dual optimization problem:

$$\begin{aligned}
& \text{minimize: } L_D = \frac{1}{2} \sum_{i=1}^N \sum_{j=1}^N t_i t_j \alpha_i \alpha_j \phi(\mathbf{x}_i) \cdot \phi(\mathbf{x}_j) - \sum_{i=1}^N \alpha_i \\
& \text{subject to: } \sum_{i=1}^N t_i \alpha_i = 0 \\
& \quad 0 \leq \alpha_i \leq C, i = 1, \dots, N
\end{aligned} \tag{5}$$

where each Lagrange multiplier α_i corresponds to a training example (\mathbf{x}_i, t_i) .

We also have

$$\mathbf{w} = \sum_{s=1}^{N_s} \alpha_s t_s \phi(\mathbf{x}_s) \tag{6}$$

where N_s is the number of support vectors \mathbf{x}_s .

2.3 SVM kernel functions

Kernel functions $K(\mathbf{u}, \mathbf{v}) = \phi(\mathbf{u}) \cdot \phi(\mathbf{v})$ are usually used in the implementation of SVM learning algorithm. In this case, we have:

$$\begin{aligned}
& \text{minimize: } L_D = \frac{1}{2} \sum_{i=1}^N \sum_{j=1}^N t_i t_j K(\mathbf{x}_i, \mathbf{x}_j) \alpha_i \alpha_j - \sum_{i=1}^N \alpha_i \\
& \text{subject to: } \sum_{i=1}^N t_i \alpha_i = 0 \\
& \quad 0 \leq \alpha_i \leq C, i = 1, \dots, N
\end{aligned} \tag{7}$$

The SVM kernel function $K(\mathbf{u}, \mathbf{v})$ needs to satisfy Mercer's condition [7]. The decision function of SVM is:

$$f(\mathbf{x}) = \text{sign} \left(\sum_{s=1}^{N_s} \alpha_s t_s K(\mathbf{x}, \mathbf{x}_s) + b \right) \tag{8}$$

3 Brief of Extreme Learning Machine

Extreme learning machine (ELM) [2–4] was originally proposed for the single-hidden layer feedforward neural networks and then extended to the generalized single-hidden layer feedforward networks where the hidden layer needn't be neuron alike [5,6]. In ELM *all* the hidden node parameters are randomly generated (even before ELM sees the training data) without tuning.

3.1 ELM feature space

The output of ELM is

$$f(\mathbf{x}) = \sum_{i=1}^L \beta_i G(\mathbf{a}_i, b_i, \mathbf{x}) = \boldsymbol{\beta} \cdot h(\mathbf{x}) \quad (9)$$

where β_i is the output weight from the i -th hidden node to the output node and $G(\mathbf{a}_i, b_i, \mathbf{x})$ is the output of the i -th hidden node. $h(\mathbf{x}) = [G(\mathbf{a}_1, b_1, \mathbf{x}), \dots, G(\mathbf{a}_L, b_L, \mathbf{x})]^T$ is the output vector of the hidden layer with respect to the input \mathbf{x} . $h(\mathbf{x})$ actually maps the data from the d -dimensional input space to the L -dimensional hidden layer feature space (*ELM feature space*) H . For the binary classification applications, the decision function of ELM is

$$f(\mathbf{x}) = \text{sign} \left(\sum_{i=1}^L \beta_i G(\mathbf{a}_i, b_i, \mathbf{x}) \right) = \text{sign} (\boldsymbol{\beta} \cdot h(\mathbf{x})) \quad (10)$$

3.2 Minimal norm of output weights

Different from traditional learning algorithms ELM not only tends to reach the smallest training error but also the smallest norm of output weights. According to Bartlett's theory [13], for feedforward neural networks reaching smaller training error, the smaller the norm of weights is, the better generalization performance the networks tends to have. We conjecture that this may be true to generalized SLFNs. ELM is to minimize the training error as well as the norm of the output weights [2,3]:

$$\begin{aligned} \text{Minimize: } & \sum_{i=1}^N \|\boldsymbol{\beta} \cdot h(\mathbf{x}_i) - t_i\| \\ \text{and} & \\ \text{Minimize: } & \|\boldsymbol{\beta}\| \end{aligned} \quad (11)$$

Seen from (10), to minimize the norm of the output weights $\|\boldsymbol{\beta}\|$ is actually to maximize the distance of the separating margins of the two different classes in the ELM feature space: $2/\|\boldsymbol{\beta}\|$ although the minimal norm least square method instead of the standard optimization method was used in the original implementation of ELM [2,3].

4 ELM for Classification

This section shows that with the standard optimization method ELM can be linearly extended to SVM (with less optimization constraints) and the implementation of SVM can be made much simpler.

Seen from the decision function of SVM (8): $f(\mathbf{x}) = \text{sign} \left(\sum_{s=1}^{N_s} \alpha_s t_s K(\mathbf{x}, \mathbf{x}_s) + b \right)$, apparently SVM can be considered as one of the “generalized” single-hidden layer feedforward networks (SLFNs) which has N_s hidden nodes. The output function of the s -th hidden node of this support vector network is $K(\mathbf{x}, \mathbf{x}_s)$, the output weight between the s -th hidden node to the output node (with bias b) is $\alpha_s t_s$.

Huang, et al [5,6] extends ELM from the SLFNs with neuron type of hidden nodes to the generalized SLFNs where the hidden nodes may not neuron alike. One of the aims of this extension is to possibly apply ELM learning approach in SVM. Since both ELM and SVM work for SLFNs and the hidden layers of both ELM and SVM are not tuned, the learning mechanism of ELM and SVM may be combined in some way: 1) similar to the standard ELM but different from SVM, the random kernels are used; 2) similar to SVM but different the least square solution of ELM, the standard optimization method is adopted to find the solution of ELM, resulting in support vectors as well.

4.1 Optimization method based solution to ELM

Different from the standard SVM [7], in theory [2–6] any set of distinct training data transformed from the input space to the ELM feature space with the ELM mapping $h(\mathbf{x})$ are linearly separable in the ELM feature space with probability one. However, it is most possible that some testing data may be within the classification margin if zero training error is strictly obtained. In this sense, one may wish to separate the training data with an acceptable minimal training error instead of the zero training error so that the testing error can thus be minimized accordingly:

$$\begin{aligned} \boldsymbol{\beta} \cdot h(\mathbf{x}_i) &\geq 1 - \xi_i & \text{if } t_i = 1 \\ \boldsymbol{\beta} \cdot h(\mathbf{x}_i) &\leq -1 + \xi_i & \text{if } t_i = -1 \end{aligned} \quad (12)$$

That is,

$$t_i \boldsymbol{\beta} \cdot h(\mathbf{x}_i) \geq 1 - \xi_i, \quad i = 1, \dots, N \quad (13)$$

Thus, from the standard optimization theory point of view, the objective (11) of ELM in minimizing both the training errors and the output weights can be

written as:

$$\begin{aligned}
& \text{Minimize: } L_P = \frac{1}{2} \|\boldsymbol{\beta}\|^2 + C \sum_{i=1}^N \xi_i \\
& \text{Subject to: } t_i \boldsymbol{\beta} \cdot h(\mathbf{x}_i) \geq 1 - \xi_i, \quad i = 1, \dots, N \\
& \quad \xi_i \geq 0, \quad i = 1, \dots, N
\end{aligned} \tag{14}$$

which is very similar to SVM's optimization problem (4) with two main differences:

- (1) Different from the conventional SVM the randomness is adopted in the ELM mapping $h(\mathbf{x})$, that is, all the parameters of $h(\mathbf{x})$ are chosen randomly.
- (2) The bias b is not required in the ELM's optimization constrains since in theory the separating hyperplane in the ELM feature space passes through the origin.

The Lagrange function of the primal ELM optimization (14) is:

$$L_{\text{ELM}}(\boldsymbol{\beta}, \boldsymbol{\xi}, \boldsymbol{\alpha}, \boldsymbol{\mu}) = \frac{1}{2} \boldsymbol{\beta} \cdot \boldsymbol{\beta} + C \sum_{i=1}^N \xi_i - \sum_{i=1}^N \alpha_i (t_i \boldsymbol{\beta} \cdot h(\mathbf{x}_i) - (1 - \xi_i)) - \sum_{i=1}^N \mu_i \xi_i \tag{15}$$

where α_i and μ_i are the Lagrange multipliers and are non-negative values.

In order to find the optimal solutions of (15) we should have:

$$\frac{\partial L_{\text{ELM}}(\boldsymbol{\beta}, \boldsymbol{\xi}, \boldsymbol{\alpha}, \boldsymbol{\mu})}{\partial \boldsymbol{\beta}} = 0 \implies \boldsymbol{\beta} = \sum_{i=1}^N \alpha_i t_i h(\mathbf{x}_i) \tag{16}$$

$$\frac{\partial L_{\text{ELM}}(\boldsymbol{\beta}, \boldsymbol{\xi}, \boldsymbol{\alpha}, \boldsymbol{\mu})}{\partial \boldsymbol{\xi}} = 0 \implies C = \alpha_i + \mu_i, \quad \forall i \tag{17}$$

Substitute (16) and (17) into (15) and to train ELM for classification is then equivalent to solving the following dual optimization problem:

$$\begin{aligned}
& \text{minimize: } L_D = \frac{1}{2} \sum_{i=1}^N \sum_{j=1}^N t_i t_j \alpha_i \alpha_j h(\mathbf{x}_i) \cdot h(\mathbf{x}_j) - \sum_{i=1}^N \alpha_i \\
& \text{subject to: } 0 \leq \alpha_i \leq C, i = 1, \dots, N
\end{aligned} \tag{18}$$

Different from the conventional dual SVM optimization problem (5), the above dual ELM optimization problem does not have the condition $\sum_{i=1}^N t_i \alpha_i = 0, \forall i$ due to the truth that in theory the separating hyperplane in the ELM feature space tends to pass through the origin.

4.2 ELM kernel

An ELM kernel function can be defined as

$$\begin{aligned} K_{\text{ELM}}(\mathbf{x}_i, \mathbf{x}_j) &= h(\mathbf{x}_i) \cdot h(\mathbf{x}_j) \\ &= [G(\mathbf{a}_1, b_1, \mathbf{x}_i), \dots, G(\mathbf{a}_L, b_L, \mathbf{x}_i)]^T \cdot [G(\mathbf{a}_1, b_1, \mathbf{x}_j), \dots, G(\mathbf{a}_L, b_L, \mathbf{x}_j)]^T \end{aligned} \quad (19)$$

where $G(\mathbf{a}, b, \mathbf{x})$ is a nonlinear piecewise continuous function satisfying ELM universal approximation capability theorems [4–6] and $\{(\mathbf{a}_i, b_i)\}_{i=1}^L$ are randomly generated according to any continuous probability distribution.

Thus, we have:

$$\begin{aligned} \text{minimize: } L_D &= \frac{1}{2} \sum_{i=1}^N \sum_{j=1}^N t_i t_j K_{\text{ELM}}(\mathbf{x}_i, \mathbf{x}_j) \alpha_i \alpha_j - \sum_{i=1}^N \alpha_i \\ \text{subject to: } &0 \leq \alpha_i \leq C, i = 1, \dots, N \end{aligned} \quad (20)$$

The decision function of ELM is

$$f(\mathbf{x}) = \text{sign} \left(\sum_{s=1}^{N_s} \alpha_s t_s K_{\text{ELM}}(\mathbf{x}, \mathbf{x}_s) \right) \quad (21)$$

4.3 ELM primal and dual networks

Seen from ELM output functions (10) and (21), after applying the optimization method to ELM, we have $\boldsymbol{\beta} = \sum_{s=1}^{N_s} \alpha_s t_s h(\mathbf{x}_s)$, where N_s is the number of the support vectors \mathbf{x}_s , and thus we have the primal ELM network as shown in Figure 1(a).

On the other hand, seen from the ELM network output function (21), after applying the standard optimization method to ELM, we can also obtain a dual network (the so-called support vector network) for ELM (cf. Figure 1(b)) which has N_s number of hidden nodes each with ELM kernel $K_{\text{ELM}}(\mathbf{x}, \mathbf{x}_s)$ while the output weight linking the s -th hidden layer to the output node is $\alpha_s t_s$.

4.4 Karush-Kuhn-Tucker Conditions of ELM

Based on the Karush-Kuhn-Tucker (KKT) theorem [18], the necessary conditions (often referred to as KKT conditions) of the primal ELM optimization (14) should be:

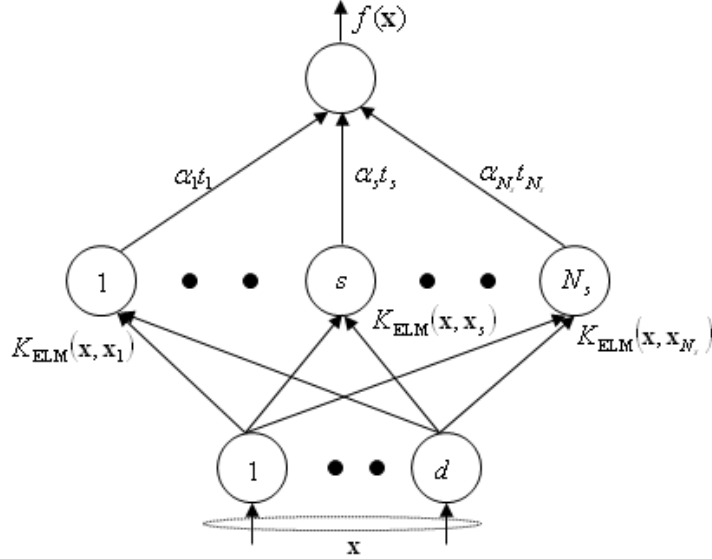
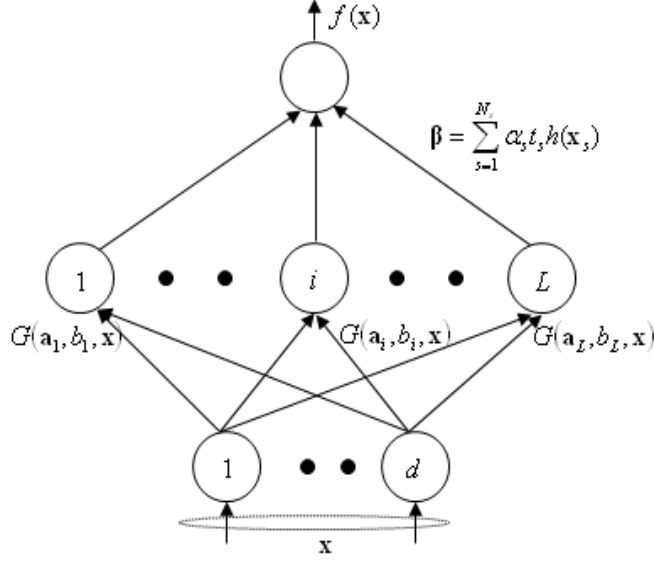


Figure 1. (a) Primal ELM Network: 1) L hidden nodes; 2) The output of the i -th hidden node is $G(\mathbf{a}_i, b_i, \mathbf{x}_i)$; 3) the output weight vector linking the hidden layer to the output node is $\boldsymbol{\beta} = \sum_{i=1}^N \alpha_i t_i K_{\text{ELM}}(\mathbf{x}, \mathbf{x}_i)$. (b) Dual ELM Network: 1) N_s hidden nodes where N_s is the number of support vectors; 2) The output of the s -th hidden node is $K_{\text{ELM}}(\mathbf{x}, \mathbf{x}_s)$: $K_{\text{ELM}}(\mathbf{x}, \mathbf{x}_s) = [G(\mathbf{a}_1, b_1, \mathbf{x}), \dots, G(\mathbf{a}_L, b_L, \mathbf{x})]^T \cdot [G(\mathbf{a}_1, b_1, \mathbf{x}_s), \dots, G(\mathbf{a}_L, b_L, \mathbf{x}_s)]^T$; 3) the output weight linking the s -th hidden layer to the output node is $\alpha_s t_s$.

Primal feasibility:

$$t_i \boldsymbol{\beta} \cdot h(\mathbf{x}_i) - (1 - \xi_i) \geq 0, \quad \forall i \quad (22)$$

$$\xi_i \geq 0, \quad \forall i \quad (23)$$

Dual feasibility:

$$\alpha_i \geq 0, \quad \forall i \quad (24)$$

$$\mu_i \geq 0, \quad \forall i \quad (25)$$

Complementary slackness:

$$\alpha_i (t_i \boldsymbol{\beta} \cdot h(\mathbf{x}_i) - (1 - \xi_i)) = 0, \quad \forall i \quad (26)$$

$$\mu_i \xi_i = 0, \quad \forall i \quad (27)$$

Furthermore,

(1) if $\alpha_i = 0$: from (17) and (27) we have $\xi_i = 0$. Thus, from (22) we have

$$t_i \boldsymbol{\beta} \cdot h(\mathbf{x}_i) \geq 1 \quad (28)$$

(2) if $0 < \alpha_i < C$: from (17) and (27) we have $\xi_i = 0$. Thus, from (26) we have

$$t_i \boldsymbol{\beta} \cdot h(\mathbf{x}_i) = 1 \quad (29)$$

(3) if $\alpha_i = C$: from (17) and (27) we have $\xi_i \geq 0$. Thus, from (26) we have

$$t_i \boldsymbol{\beta} \cdot h(\mathbf{x}_i) \leq 1 \quad (30)$$

Thus, the QP problem is solved when for all i :

$$\begin{aligned} \alpha_i = 0 &\implies t_i \boldsymbol{\beta} \cdot h(\mathbf{x}_i) \geq 1 \\ 0 < \alpha_i < C &\implies t_i \boldsymbol{\beta} \cdot h(\mathbf{x}_i) = 1 \\ \alpha_i = C &\implies t_i \boldsymbol{\beta} \cdot h(\mathbf{x}_i) \leq 1 \end{aligned} \quad (31)$$

5 Discussions

In essence it has different physical meanings of introducing the training error in the optimization constraints of both SVM and ELM: the reason why SVM permits the training errors is that some training data may not be linearly separable in the conventional SVM feature space. According to the ELM theory [2–6] all the training data are linearly separable in the ELM feature space, the reason why ELM permits the training errors is to eliminate the possible overfitting and thus to minimize the testing errors and to further improve the generalization performance. According to ELM theories [2–6] widespread nonlinear piecewise continuous functions can be used as ELM feature mapping $h(\mathbf{x})$. As pointed out by Fréney and Verleysen [15] the ELM kernel function $K_{\text{ELM}}(\mathbf{x}, \mathbf{x}_i)$ always satisfies the Mercer’s condition.

5.1 Differences between SVM's separating hyperplane and ELM's separating hyperplane

Liu, et al [14] and Fréney and Verleysen [15] provides an efficient extreme learning approach to SVM by simply replacing the SVM kernels with ELM kernels. Generally speaking, their work adopts the same optimization constraints as the conventional SVM.

The essence of SVM is: after the training data are mapped into the SVM feature space and there exists a hyperplane which can separate these data with a maximal margin. It is reasonable to think that such separating hyperplane in SVM may not necessarily pass through the origin in the SVM feature space and thus a bias b is required in the optimization constraint of SVM: $t_i(\mathbf{w} \cdot \phi(\mathbf{x}_i) + b) \geq 1 - \xi_i$. Liu, et al [14] and Fréney and Verleysen [15] proposed the SVM with ELM kernels with the same optimization constraint assumption as the conventional SVM. The corresponding Lagrange function of the primal SVM optimization (4) is:

$$L_{\text{SVM}}(\mathbf{w}, \boldsymbol{\xi}, \boldsymbol{\alpha}, \boldsymbol{\mu}, b) = \frac{1}{2} \mathbf{w} \cdot \mathbf{w} + C \sum_{i=1}^N \xi_i - \sum_{i=1}^N \alpha_i (t_i(\mathbf{w} \cdot \phi(\mathbf{x}_i) + b) - (1 - \xi_i)) - \sum_{i=1}^N \mu_i \xi_i \quad (32)$$

In order to find the optimal solutions of (32) one of the KKT conditions (necessary conditions) for the conventional SVM (as well as for SVM with ELM kernels [14,15]) is:

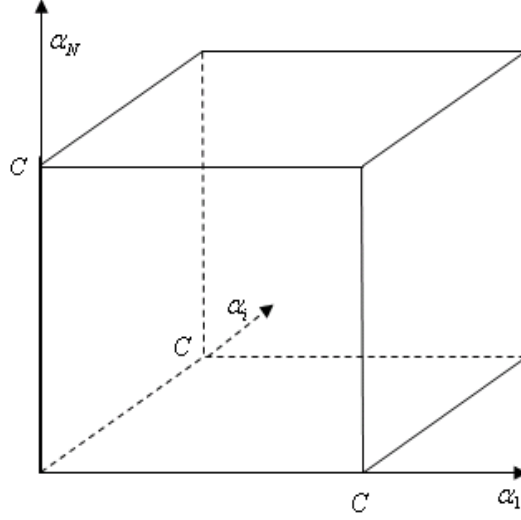
$$\frac{\partial L_{\text{SVM}}(\mathbf{w}, \boldsymbol{\xi}, \boldsymbol{\alpha}, \boldsymbol{\mu}, b)}{\partial b} = 0 \implies \sum_{i=1}^N \alpha_i t_i = 0 \quad (33)$$

This is the reason why SVM learning needs to satisfy both optimization conditions (5): $\sum_{i=1}^N \alpha_i t_i = 0$ and $0 \leq \alpha_i \leq C, \forall i$.

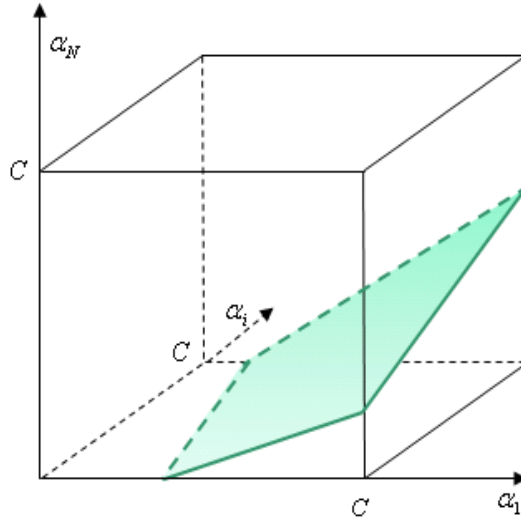
According to ELM theories [2–6] the separating hyperplane of ELM basically passes through the origin in the SVM feature space, there is no bias b in the optimization constraint of ELM: $t_i \boldsymbol{\beta} \cdot h(\mathbf{x}_i) \geq 1 - \xi_i$ and thus, different from SVM, ELM learning does not need to satisfy the condition: $\sum_{i=1}^N \alpha_i t_i = 0$.

SVM and ELM have similar optimization objective functions and SVM learning needs to satisfy both optimization conditions (5): $\sum_{i=1}^N \alpha_i t_i = 0$ and $0 \leq \alpha_i \leq C, \forall i$ while ELM learning only needs to satisfy the loose condition: $0 \leq \alpha_i \leq C, \forall i$, thus, obviously SVM tends to find a solution which is sub-optimal to ELM's solution.

This view can also become clear from a different observation aspect. ELM



(a)



(b)

Figure 2. (a) ELM finds the optimal solution in the entire cube $[0, C]^N$ of the ELM feature space without any other constraints. (b) SVM always searches for the optimal solution in the hyperplane $\sum_{i=1}^N \alpha_i t_i = 0$ (shaded area) within the cube $[0, C]^N$ of the SVM feature space.

finds the optimal solution in the entire cube $[0, C]^N$ of the ELM feature space without any other constraints (cf. Figure 2(a)). However, SVM always searches for the optimal solution in the hyperplane $\sum_{i=1}^N \alpha_i t_i = 0$ within the cube $[0, C]^N$ of the SVM feature space (cf. Figure 2(b)). In this sense, SVM's search area depends more on the target output vector $[t_1, \dots, t_N]^T$ instead of the combination of (\mathbf{x}_i, t_i) . In other words, given two training datasets $\{(\mathbf{x}_i^{(1)}, t_i^{(1)})\}_{i=1}^N$ and $\{(\mathbf{x}_i^{(2)}, t_i^{(2)})\}_{i=1}^N$ and $\{\mathbf{x}_i^{(1)}\}_{i=1}^N$ and $\{\mathbf{x}_i^{(2)}\}_{i=1}^N$ are totally irrelevant/independent, if $[t_1^{(1)}, \dots, t_N^{(1)}]^T$ is similar or close to $[t_1^{(2)}, \dots, t_N^{(2)}]^T$

(although this rarely happens in applications) SVM may have similar search areas of the cube $[0, C]^N$ for two different cases, which actually implies that the bias b may not be used in SVMs.⁴

5.2 Maximal separating margin and minimal norm of output weights

The aim of ELM is to minimize the norm of the output weights β : $\min \|\beta\|$. However, the distance of the separating boundaries of the two classes in the ELM feature $2/\|\beta\|$, thus to minimize the norm of the output weights β is actually to maximize the separating margin.

6 Performance Verification

In this section, the performance of ELM for classification is compared with SVM on some benchmark binary classification problems: 11 UCI datasets [20] and 2 Gene expression datasets (leukemia and colon [21]) (cf Table 1). The training and testing data of the first 7 datasets of Table 1 are reshuffled at each trial of simulation while the training and testing data of the last 6 datasets remain fixed for all trials of simulations. Different from the ELM learning approach to SVM proposed by Liu, et al [14], Frénay and Verleysen [15] simply replaces the conventional kernel with ELM kernel in the standard SVM method. We will compare our method with ELM learning approach to SVM proposed by Frénay and Verleysen [15].

All the simulations for SVM and ELM for classification algorithms are carried out in MATLAB 2007 environment running in a Pentium 4, 2.53 GHZ CPU. The simulations for SVM are carried out using Matlab-coded SVM package: SVM and Kernel Methods Matlab Toolbox [22]. Although it is fully written in Matlab, it can perform fast due to the optimization of QP solver.

6.1 User specified parameters

The popular Gaussian kernel function $K(\mathbf{u}, \mathbf{v}) = \exp(-\gamma\|\mathbf{u} - \mathbf{v}\|^2)$ is used in SVM. Sigmoid type of ELM kernels are used in ELM and ELM for SVM [15]:

⁴ Poggio, et al [19] studies the role of the bias b in SVMs and shows that the bias b may not be required in some cases. However, different from our work, Poggio, et al [19] does not show that from the cross-application point of view the bias b should not be used in SVMs. A detail comparison of SVMs with and without the bias b is beyond the scope of this paper and will be reported in a separate work.

Datasets	# Attributes	# Training data	# Testing data
Breast-cancer	10	300	383
liver-disorders	6	200	145
heart	13	70	200
ionosphere	34	100	251
Pimadata	8	400	368
Pwlinear	10	100	100
Sonar	60	100	158
Monks Problem 1	6	124	432
Monks Problem 2	6	169	432
Splice	60	1000	2175
A1a	123	1605	30956
leukemia	7129	38	34
Colon	2000	30	32

Table 1
Specification of tested binary classification problems

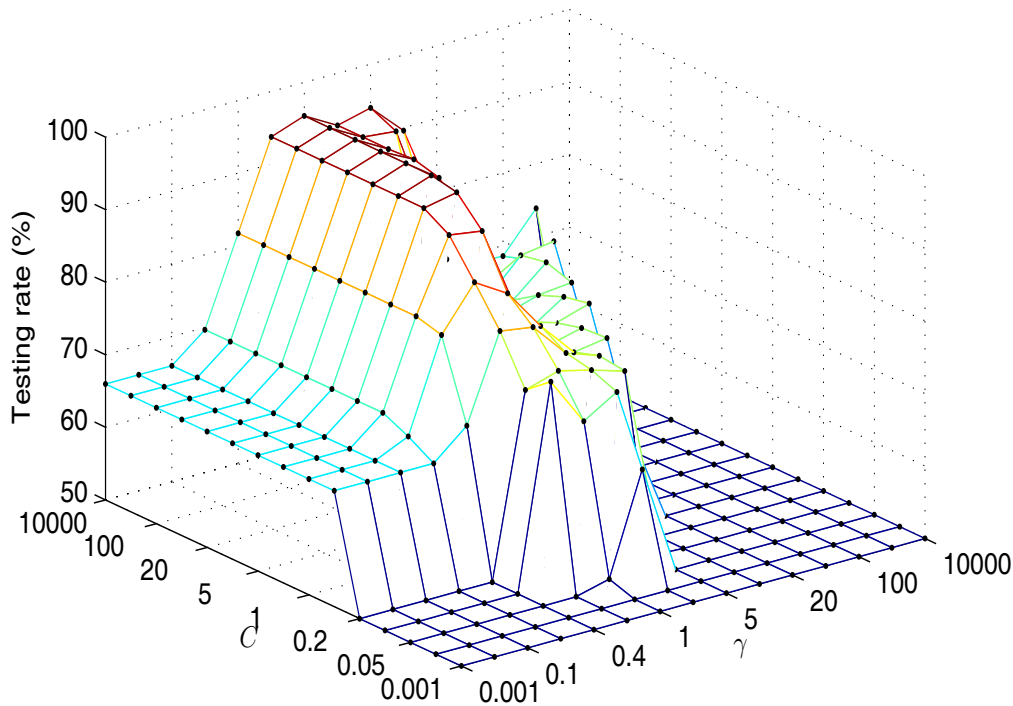


Figure 3. The performance of SVM is sensitive to the parameters (C, γ) : an example on Monks problem 1.

$K_{\text{ELM}}(\mathbf{x}, \mathbf{x}_s) = [G(\mathbf{a}_1, b_1, \mathbf{x}), \dots, G(\mathbf{a}_L, b_L, \mathbf{x})]^T \cdot [G(\mathbf{a}_1, b_1, \mathbf{x}_s), \dots, G(\mathbf{a}_L, b_L, \mathbf{x}_s)]^T$
where $G(\mathbf{a}, b, \mathbf{x}) = 1/(1 + \exp(-(\mathbf{a} \cdot \mathbf{x} + b)))$. In order to achieve good generalization performance, the cost parameter C and kernel parameter γ of SVM need to be chosen appropriately. Similar to Ghanty et al [23] we have tried a

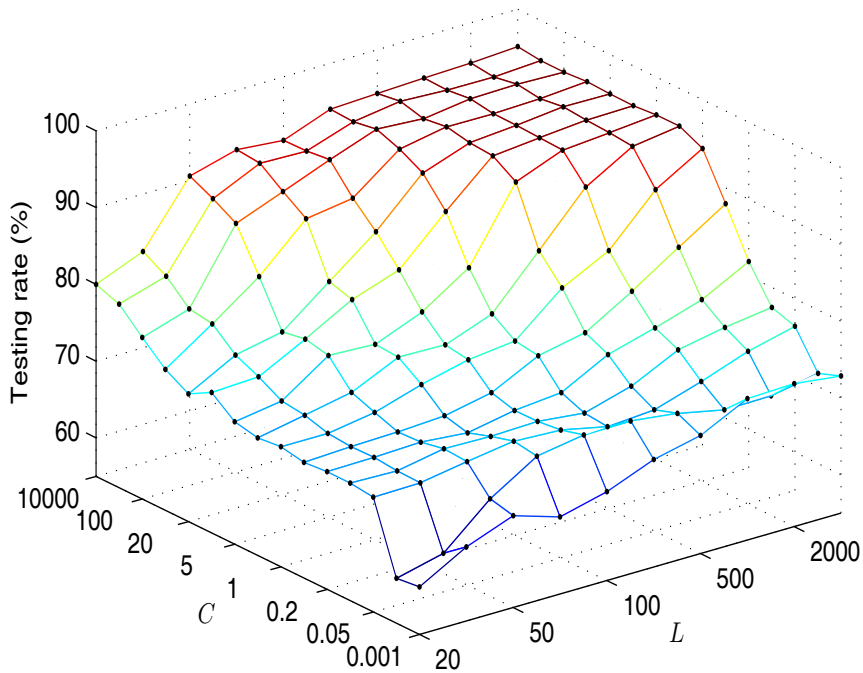


Figure 4. The performance of ELM is not very sensitive to the parameters (C, L) and a good testing accuracy can be achieved as long as L is large enough: an example on Monks problem 1.

wide range of C and γ . For each dataset we have used 15 different values of C and 15 different values of γ resulting in a total of 225 pairs of (C, γ) . The 15 different values of C are 0.001, 0.01, 0.05, 0.1, 0.2, 0.5, 1, 2, 5, 10, 20, 50, 100, 1000, 10000 and the 15 different values of γ are 0.0001, 0.001, 0.01, 0.1, 0.2, 0.4, 0.8, 1, 2, 5, 10, 20, 100, 1000 and 10000.

It is known that the generalization performance of SVM usually depends closely on the combination of (C, γ) (see Figure 3 for an example). The best generalization performance is usually achieved in a narrow range of such combinations. Thus, when SVM applied the best combination of (C, γ) needs to be chosen for each dataset.

For ELM for classification and SVM with ELM kernel [15]), there are two parameters: the cost parameter C and the number of hidden nodes L . For each problem we have also used 15 different values of C and 10 different values of L resulting in a total of 150 pairs of (C, L) . The 15 different values of C are 0.001, 0.01, 0.05, 0.1, 0.2, 0.5, 1, 2, 5, 10, 20, 50, 100, 1000, and 10000 and the 10 different values of L are 20, 30, 50, 80, 100, 200, 500, 1000, 2000, 3000. After extensive simulations, it is found that the performance of ELM and SVM with ELM kernel is actually not so sensitive to the combination of (C, L) . As mentioned in [15], the good performance of ELM for SVM can be

achieved as long as the number of hidden node L is large enough. It is true to ELM as well. As observed from Figure 4 (which is true to other datasets as well), the generalization performance of ELM tends to monotonically increase with the number of hidden nodes L . In all the simulations for 11 UCI datasets, L is set 1000. One only needs to adjust the cost parameter C although the performance of ELM is not very sensitive to C either. The insensitivity of ELM performance on the user specified parameters makes the implementation of ELM very effectively.

Twenty trials have been conducted for each problem, with training and testing data sets randomly generated for each trial. Simulation results including the average testing accuracy, the corresponding standard deviation (Dev), and the number of resulted support vectors are given in this section.

6.2 Performance comparison on UCI benchmark datasets

As observed from Table 2, SVM with ELM kernels and the conventional SVM achieve similar generalization performance, however, the conventional SVM face the trivial and time-consuming issue in finding the best combination of learning parameters while users can choose the learning parameters for SVM with ELM kernels easily. Among the three learning methods, ELM usually achieves the best generalization performance.

6.3 Performance comparison on DNA microarray datasets

The leukemia dataset was taken from a collection of leukemia patient samples [24]. The dataset consists of 72 samples: 25 samples of AML, and 47 samples of ALL. Each sample is measured over 7,129 genes. The colon microarray dataset consists of 22 normal and 40 tumor tissue samples. In this dataset, each sample contains 2,000 genes.

The minimum-redundancy-maximum-relevance (MRMR) feature selection method was developed by Ding and Peng, et al [25], and further studied in Ding and Peng [26]. MRMR considers both maximum relevance with the class labels and minimum redundancy among genes. That has six cases, such as MID, MIQ, FCD, FCQ, FDM, FSQ. The MRMR feature selection is among the most powerful methods to select a subset of features from a big pool, or just do dimension reduction. This method has been well cited and used for many different applications. Seen from Table 3, ELM generally achieves the best generalization performance in these two DNA microarray applications.

Datasets	SVM											ELM				
	Gaussian Kernel						ELM Kernel [15]					C	Training Time (s)	Testing Rate (%)	Testing Dev (%)	# SVs
	(C, γ)	Training Time (s)	Testing Rate (%)	Testing Dev (%)	# SVs	C	Training Time (s)	Testing Rate (%)	Testing Dev (%)	# SVs						
Breast- cancer	(5, 50)	0.1118	94.20	0.87	190	10^{-3}	0.1442	96.28	0.59	65	10^{-3}	0.1423	96.32	0.75	66	
Liver- disorders	(10, 2)	0.0972	68.24	4.58	158	10	0.1685	71.79	3.02	132	10	0.1734	72.34	2.55	131	
Heart	(10^4 , 5)	0.0382	76.00	3.85	41	50	0.0474	75.32	2.22	37	50	0.0344	76.25	2.70	36	
Ionosphere	(5, 5)	0.0218	90.58	1.22	30	0.1	0.0396	88.78	1.77	25	10^{-3}	0.0359	89.48	1.12	32	
Pimadata	(10^3 , 50)	0.2049	76.43	1.57	209	0.01	0.2942	76.54	1.89	218	0.01	0.2867	77.27	1.33	217	
Pwlinear	(10^4 , 10^3)	0.0357	84.35	3.28	54	10^{-3}	0.0560	84.10	2.88	49	10^{-3}	0.0486	86.00	1.92	51	
Sonar	(20, 1)	0.0412	83.33	3.55	70	10^4	0.0498	78.81	3.75	44	10^4	0.0467	81.53	3.78	50	
Monks Problem 1	(10, 1)	0.0424	95.37	0	70	10^4	0.0612	94.49	0.51	37	10^4	0.0821	95.19	0.41	40	
Monks Problem 2	(10^4 , 5)	0.0860	83.80	0	77	10^4	0.1077	85.52	0.59	75	10^4	0.0920	85.14	0.57	75	
Splice	(2, 20)	2.0683	84.05	0	704	0.01	3.2076	85.34	0.41	507	0.01	3.5912	85.50	0.54	509	
Ala	(2, 5)	5.6275	84.25	0	680	0.01	5.6017	84.13	10^{-3}	656	0.01	5.4542	84.36	0.79	666	

Table 2. Comparison between the conventional SVM, SVM with ELM kernel [15] and ELM for classification.

Datasets	SVM										ELM				
	Gaussian Kernel					ELM Kernel [15]					(C, L)	Training Time (s)	Testing Rate (%)	Testing Dev (%)	# SVs
	(C, γ)	Training Time (s)	Testing Rate (%)	Testing Dev (%)	# SVs	(C, L)	Training Time (s)	Testing Rate (%)	Testing Dev (%)	# SVs					
Before Gene Selection															
leukemia	$(10^3, 10^3)$	3.2674	82.35	0	29	(0.05, 3000)	0.9796	80.15	2.44	35	(0.01, 3000)	0.2878	81.18	2.37	35
Colon	$(10^3, 10^3)$	0.2391	81.25	0	24	(0.01, 3000)	0.2757	81.87	2.54	28	(0.01, 3000)	0.1023	82.50	2.86	28
After Gene Selection (60 Genes Obtained for Each Case)															
leukemia	(2, 20)	0.0640	100	0	12	(0.01, 3000)	0.0316	100	0	13	(10, 3000)	0.0199	100	0	11
Colon	(2, 50)	0.0166	87.50	0	26	(0.05, 20)	0.0160	82.34	4.22	25	(0.001, 500)	0.0161	89.06	2.10	28

Table 3. Comparison between the conventional SVM, SVM with ELM kernel [15] and ELM in gene classification applications.

7 Conclusions

This paper studies ELM for classification with the standard optimization method. It is found that ELM and SVM are actually consistent in some manner: 1) to minimize the norm of output weights in ELM for classification is actually to maximize the distance of the separating margin of two different classes in the ELM feature space, which is equivalent to SVM's maximal separating margin property; 2) both ELM and SVM are to maximize the separating margin as well as to minimize the training errors.

This paper shows that the separating hyperplane tends to pass through the origin of the ELM feature space, resulting in less optimization constraints and better generalization performance than SVM. It is also pointed out that the generalization performance of ELM is less sensitive to the learning parameters especially the number of hidden nodes. Thus, compared to SVM, users can use ELM easily and effectively by avoiding tedious and time-consuming parameter tuning. From the implementation point of view, such ELM implementation may straightforward be extended to other SVM variants and those extensions are worth studying further in the future.

Similar to the conventional SVM, one may easily apply ELM in multi-label classification applications by using one-against-one (OAO) or one-against-all (OAA) methods.

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