Extreme Learning Machine with Enhanced Variation of Activation Functions

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Abstract: The main aim of this paper is to stress the fact that the sufficient variability of activation functions (AF) is important for an Extreme Learning Machine (ELM) approximation accuracy and applicability. A slight modification of the standard ELM procedure is proposed, which allows increasing the variance of each AF, without losing too much from the simplicity of random selection of parameters. The proposed modification does not increase the computational complexity of an ELM training significantly. Enhancing the variation of AFs results in reduced output weights norm, better numerical conditioning of the output weights calculation, smaller errors for the same number of the hidden neurons. The proposed approach works efficiently together with the Tikhonov regularization of ELM.

1 INTRODUCTION

Extreme Learning Machine (ELM) is widely accepted assignment of the learning algorithm for a single-hidden- layer, feedforward neural network. The main concepts behind the ELM are that: (i) the weights and biases of the hidden nodes are generated randomly and are not adjusted and (ii) the output weights are determined analytically. ELM may be applied for modeling (regression) as well as classification problems, and various modifications of the basic algorithm are possible. The literature concerning ELMs is numerous, but some recent review papers extensively describe the recent trends and modifications of the standard algorithm (Huang, Huang, Song, and You, 2015), (Liu, Lin, Fang, and Xu, 2015), (Lin, Liu, Fang, and Xu, 2015). Very short learning times and the simplicity of the algorithm are the most attractive features ELMs. On the other hand, several publications report that ELM may: create illcondition numerical problems (Chen, Zhu, and Wang, 2013), introduce overfitting, require too big number of neurons (Liu et al., 2015), that the influence of number of neurons and parameters of random weights selection on the resulting performance is unclear and requires analysis (Parviainen and Riihimäki, 2013). The necessity of improving the numerical properties of ELM was noticed in several recent publications (Akusok, Bjork, Miche, and Lendasse, 2015).

In this contribution a standard ELM applied for regression problems with batch data processing is considered. Numerical properties of the standard algorithm are commented and insufficient variation of the AFs is recognized as the reason of huge output weights and large modeling errors. The modification of the input weights and biases selection procedure is proposed to improve the ELM accuracy and applicability.

2 STANDARD ELM MODELING

2.1 Components of ELM

2.1.1 Training Data

The training data for a *n*-input ELM compose a batch of *N* samples:

$$\{(x_i, t_i), x_i \in \mathbb{R}^n, t_i \in \mathbb{R}, i = 1, \dots, N\}, (1)$$

where x_i denote the inputs and t_i the desired outputs, that form the target (column) vector:

$$T = \begin{bmatrix} t_1 & \cdots & t_N \end{bmatrix}^T. \tag{2}$$

It is commonly accepted that the inputs are normalized to the interval [0,1] each.

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2.1.2 Hidden Neurons

The single, hidden layer of M neurons transforms the input data into a different representation, called the feature space. The most popular are "projection-based neurons". Each *n*-dimensional input is projected by the input layer weights $w_k^T = [w_{k,1} \cdots w_{k,n}]$, $k = 1, \dots M$ and the bias b_k into the *k*-th neuron input and next a nonlinear transformation h_k , called activation function (AF), is applied to obtain the neuron output. The matrix form of the hidden layer performance on the batch of N samples is represented by a $N \times M$ matrix:

$$H = \begin{bmatrix} h_1(w_1^T x_1 + b_1) & \cdots & h_M(w_M^T x_1 + b_M) \\ \vdots & \ddots & \vdots \\ h_1(w_1^T x_N + b_1) & \cdots & h_M(w_M^T x_N + b_M) \end{bmatrix}.$$
 (3)

Although it is not obligatory that the hidden layer must contain only one kind of neurons, it is usually the case. Any piecewise differentiable function may be used as activation function: sigmoid, hyperbolic tangent, threshold are among the most popular.

Another type of neurons used in ELM is "distancebased" neurons, such as Radial Basis Functions (RBF) or multi-quadratic functions. Each neuron uses the distance from the centroid (represented by w_k^T) as the input to the nonlinear transformation (Guang-Bin Huang and Chee-Kheong Siew, 2004). The formal representation of ELM also generates the matrix similar to (3):

$$H = \begin{bmatrix} h_1(||w_1 - x_1||, b_1) & \cdots & h_M(||w_M - x_1||, b_M) \\ \vdots & \ddots & \vdots \\ h_1(||w_1 - x_N||, b_1) & \cdots & h_M(||w_M - x_N||, b_M) \end{bmatrix}. (4)$$

The acceptable number of neurons may be found using validation data, Leave-One-Out validation procedure, random adding or removing the neurons (Akusok et al., 2015) or ranking the neurons (Miche et al., 2010), (Feng, Lan, Zhang, and Qian, 2015), (Miche, van Heeswijk, Bas, Simula, and Lendasse, 2011).

The weights and the biases of the hidden neurons are generated on random. The uniform distributions in [-1,1] for the weights and in [0, 1] for the biases is the most popular choice. If the data are normalized to have the zero mean and the unit variance the normal distribution may be used to generate the neuron parameters (Akusok et al., 2015).

2.1.3 Output Weights

The output F of an ELM is obtained by applying the

output weights β to the hidden neurons, therefore the outputs for all samples in the batch are:

$$F = H\beta. \tag{5}$$

The output weights β are found by minimizing the approximation error:

$$E = \|H\beta - T\|^2.$$
(6)

The optimal solution is:

$$\beta_{opt} = H^+ T, \tag{7}$$

where H^+ is the Moore–Penrose generalized inverse of matrix H.

The matrix *H* possesses *N* rows and *M* columns. If the number of training samples *N* is bigger than the number of hidden neurons M, $N \ge M$, and the matrix *H* has full column rank, then

$$H^{+} = (H^{T}H)^{-1}H^{T}.$$
 (8)

If the number of training samples N is smaller than the number of hidden neurons M, N < M, and the matrix H has full row-rank, then

$$H^{+} = H^{T} (HH^{T})^{-1}, (9)$$

but the last case is impractical in modeling.

2.2 Numerical Aspects of ELM

The calculation of the output weights is the most sensitive stage of the ELM algorithm and may be a reason for various numerical difficulties. Each of two ways may be selected to calculate β_{opt} : specialized algorithms for calculation of pseudoinverse (8), or simply, calculation of following matrices:

$$\Lambda = H^T H, \ \Omega = HT, \ \beta_{opt} = \Lambda^{-1} \Omega.$$
 (10)

The computational complexity of both approaches is similar, but the second one is characterized by smaller memory requirements (Akusok et al., 2015). For the both approaches working with moderate condition number of H^TH is crucial for the algorithm stability and is necessary to get moderate values of β_{opt} . Huge output weights may reinforce the round-off errors of arithmetical operations performed by the network and make the application impossible.

The well-known therapy for numerical problems in ELM caused by ill-conditioned matrix H is Tikhonov regularization of the least-square problem (Huang et al., 2015), (Akusok et al., 2015). Instead of minimizing (6), the weighted problem is considered, with the performance index:

$$E_{\mathcal{C}} = \|\beta\|^2 + C\|H\beta - T\|^2, \qquad (11)$$

where C > 0 is a design parameter. The optimal solution is:

$$\beta_{Copt} = \left(\frac{1}{c}I + H^T H\right)^{-1} H^T T.$$
(12)

Inevitably, this modification degrades the modeling accuracy. The proper choice of C depend on the problem structure and it is difficult to formulate general rules. The structure of the matrix H depends on the number of neurons, samples, and inputs, and on the shape of activations functions. The target vector has no influence on H.

To demonstrate that ill-conditioning of the matrix $H^{T}H$ may really cause problems, a simple example is demonstrated. Consider the sigmoid activation functions with input weights and biases taken randomly according to the uniform distribution in the interval [-1,1], samples selected according to the same distribution in the unit cube and calculate the *cond*($H^{T}H$). The mean of condition coefficients obtained in 20 experiments, for several numbers of inputs is presented in fig. 1.

The number of hidden neurons was always smaller than a half of a number of samples. The condition number of $H^T H$ increases rapidly with a number of neurons and samples. Performing any calculations with the numbers as big as 10^{14} in the matrix Λ^{-1} is not reasonable.

As it is visible, the problem of ill-conditioning escalates for small-dimensional modeling, paradoxically. The regularization is not able to solve this difficulty efficiently. To keep the condition number in reasonable constraints $\frac{1}{c} < 10^{-5}$ must be used, as it is presented in fig. 2. It is much bigger than typically applied values $\frac{1}{c} = 50 \times machine \ epsilon$.

3 ENHANCED VARIATION OF ACTIVATION FUNCTIONS

The ELM is not able to learn the features from the data, as a fully trained neural network does. The randomly chosen weights happen to specify a linear mapping to the space of neuron arguments. Therefore the nonlinear mapping of the data into a feature space should be able to extract the features sufficient for predicting the target variable of a regression task. It is not achieved at all times if the typical rules of neuron construction (described in section 2) are applied.

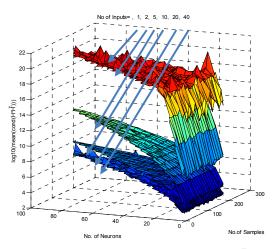


Figure 1: Logarithm of the condition number of $H^T H$ as a function of batch size and the number of hidden neurons for different number of inputs.

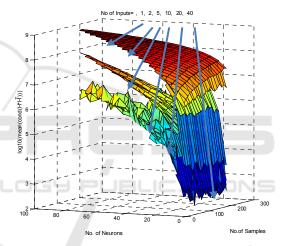


Figure 2: Logarithm of the condition number of $H^T H$ as a function of batch size and the number of hidden neurons for different number of inputs and the regularisation parameter $C = 10^5$.

For instance consider a case with 4 neurons, 2 inputs, 100 samples, weights and biases selection according to the uniform distribution in [-1,1]. The mean condition number of $H^T H$ in 2000 experiments is ~ 10⁶ and the exemplary plots of activation functions (from the last experiment) are presented in fig. 3.

It may be noticed that the activation functions are almost linear and that the range of theirs outputs is limited.

The importance of sufficient AFs variability was noticed previously (Parviainen and Riihimäki, 2013), (Kabziński, 2015). The challenge is to correct the random mechanism of weights and biases creation to increase variability without losing too much from the simplicity of random selection.

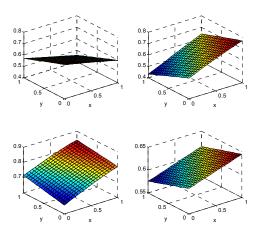


Figure 3: Plots of exemplary sigmoid activation functions obtained from a standard ELM.

For sigmoid AFs the following procedure may be proposed:

The first step to enlarge the variation of a sigmoid activation function is to increase the range of uniform random selection of input weights. It must be suitably fixed to guarantee that the sigmoid operation neither remains linear nor too strongly saturates in the input domain. Therefore the weights are selected from uniform random distribution in the interval [-q, q]. The values 5 < q < 10 seems suitable.

The selected weights for the *k*-th neuron are divided into positive and negative.

Next, the biases may be selected to ensure that the range of a sigmoid function is sufficiently large. The minimal value of the sigmoid function 200

$$h_k(x) = \frac{1}{1 + exp(-(w_k^T x + b_k))}$$
(13)

in the unit cube is achieved at the vertex selected according to the following rules:

$$w_{k,i} > 0 \Rightarrow x_i = 0, w_{k,i} < 0 \Rightarrow x_i = 1 \ i = 1, ..., n,$$

(14)

and equals

$$h_{k,min} = \frac{1}{1 + exp(-(\sum_{i:w_{k,i} < 0} w_{k,i} + b_k))}, \quad (15)$$

while the maximal value is achieved at the point given by:

$$w_{k,i} > 0 \Rightarrow x_i = 1, w_{k,i} < 0 \Rightarrow x_i = 0 \ i = 1, ..., n$$

(16)

and equals

$$h_{k,max} = \frac{1}{1 + exp(-(\sum_{i:w_{k,i} > 0} w_{k,i} + b_k))}.$$
 (17)

Assuming that the minimal value of sigmoid AF should be smaller than the given r_1 , and the maximal value should be bigger than r_2 , $0 < r_1 < r_2 < 1$ yields the range for the bias selection. It follows from the inequalities

$$h_{k,min} < r_1, \ h_{k,max} > r_2,$$
 (18)

that the bias b_k should be selected on random, uniformly from $b_k \in [\bar{a}, \tilde{a}]$, where

$$\bar{a} = -\sum_{i:w_{k,i}>0} w_{k,i} - ln\left(\frac{1}{r_2} - 1\right),$$
(19)

$$\tilde{a} = -ln\left(\frac{1}{r_1} - 1\right) - \sum_{i:w_{k,i} < 0} w_{k,i}.$$
 (20)

Of course, this approach does not guarantee that the range of the AF will be $[r_1, r_2]$, as the parameters are still random variables, but at least has chance to be. A similar strategy may be applied for other "projection based" AFs.

4 EFFECTIVENESS OF ELM WITH ENHANCED VARIATION OF ACTIVATION FUNCTIONS

To demonstrate the effectiveness of the proposed method, consider a two-dimensional function

$$z = \sin(2\pi(x_1 + x_2), x_1, x_2 \in [0, 1].$$
(21)

200 samples selected on random constitute the training set, and 100 samples the testing sets. The surface (21) was modeled by the standard ELM (S-ELM), ELM with the enhanced variation of AFs (EV-ELM) with parameters $r_1 = 0.1, r_2 = 0.9$, standard ELM with regularization (R-ELM) and ELM with regularization and the enhanced variation of AFs (EV-R-ELM) with parameters $C = 10^5, r_1 = 0.1, r_2 = 0.9$.

In fig. 4,5 the modeling errors and the conditioning of S-ELM and EV-ELM are compared. Application of EV-ELM allows smaller modeling errors and far better numerical properties of the obtained model. The output weights are $\sim 10^7$ times smaller in EV-ELM then in S-ELM.

The reason for this improvement is visible in fig. 6 where the plots of 4 AFs (selected from 60) are presented. Fig. 6 and 3 demonstrate that in EV-ELM AFs are nonlinear and cover wider subinterval in [0,1], however not all of them vary from r_1 to r_2 .

The advantage of EV-ELM is even more obvious if it is applied together with the regularization. Fig. 7, 8 demonstrate the comparison between R-ELM and

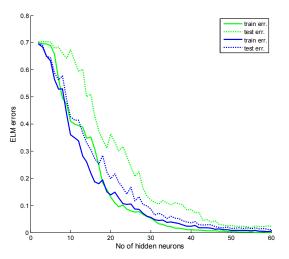


Figure 4: Testing (dotted) and training (solid) errors for S-ELM (brighter - green) and EV-ELM (darker – blue).

EV-R-ELM. The regularization alone is able to keep the output weights on a moderate level, but the modeling becomes unacceptably inaccurate. The EV-R-ELM improves the modeling accuracy and the conditioning together. Similar experiments were conducted for different modeling problems: with the training output corrupted by noise, with multiple inputs, with numerous extrema, and always the conclusions were similar.

5 CONCLUSIONS

The main aim of this paper was to stress the fact that the sufficient variability of AFs is important for an ELM model accuracy and applicability. A slight modification of the standard ELM procedure is proposed, which allows enhancing the variance of each AF, without losing too much from the simplicity of random selection of parameters.

First, it is proposed to change the parameter of the random distribution of the input weights, next to modify the random distribution for the basses, taken already established weights into account. The proposed modification does not increase the computational complexity of an ELM training significantly – only two simple calculations for each bias are added. Enhancing the variation of AFs results in reduced output weights norm, better numerical conditioning of the output weights calculation, smaller errors for the same number of the hidden neurons. The proposed approach works efficiently together with the Tikhonov regularization applied in ELM.

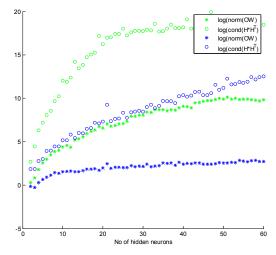


Figure 5: Logarithm of the norm of the output weights (stars) and $log(cond(HH^T)$ (circles) for S-ELM (brighter - green) and EV-ELM (darker – blue).

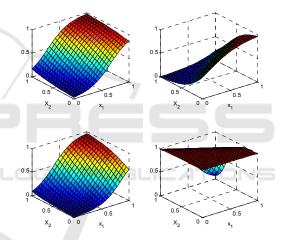


Figure 6: Selected AFs (4 from 60) from the EV-ELM model.

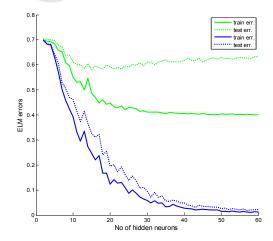


Figure 7: Testing (dotted) and training (solid) errors for R-ELM (brighter - green) and EV-R-ELM (darker – blue).

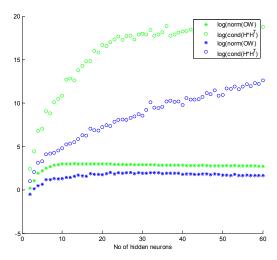


Figure 8: Logarithm of the norm of the output weights (stars) and $log(cond(HH^T)$ (circles) for R-ELM (brighter - green) and EV-R-ELM (darker – blue).

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