NEURAL NETWORKS WITH SELF-ADAPTIVE TOPOLOGY

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ABSTRACT

Recently, it has been proposed that the biological networks change not only the synaptic strengths of connection but also partially their internal topologies, according to either the received external stimuli and the pre-existent connection layouts. Following this idea, in the paper a method is presented to dynamically adapt the topology of a neural network using only the information of the learning set. The method, although simply eliminates connections from an initial fully connected network, presents characteristics which can resemble some biological behaviour. Several experimental results obtained with recurrent and multilayered networks are reported in the paper to demonstrate the capabilities of the proposed method.

1. Introduction

It is well known that Artificial Neural Networks can solve a large variety of problems. In order to reach this goal, usually a formal model of the neuron is chosen and used to build a network with a given topology. Then this network is trained, adapting the synaptic weights of every neuron, by using a suitable learning algorithm. The network topologies which are extensively used, are chosen among a set of known fixed models that are characterized by the presence of dense connection matrices among all neurons (fully connected network) or among groups of neurons, as in the well known MultiLayer Perceptron (MLP) [1]. These topologies however are different from those found in biological neural systems, where the networks are often much less regular. Moreover, since a particular problem may be optimally solved by a network with a non-regular connection topology, the task of transforming an oversized regularly connected network into such optimal non-regular network is completely committed to the learning algorithm. Most of known learning algorithms, however, are not able to produce a reduced topology by zeroing some synaptic weights, instead they typically produce different structures spreading nonvanishing weights all over the network. The efficiency can result poor and several dimensions of a chosen configuration must be tried to yield an acceptable solution. In fact, if the network is too small, the input-output mapping cannot be learned with satisfactory accuracy. Conversely, if the network is too large, after a long training phase it will learn the given set correctly, but will badly generalize due to learning data overfitting [2]. Therefore, the goal when training a network is to find a topology large enough to learn the mapping and as small as possible to generalize correctly.

Two possible approach to produce networks with correct topologies have been proposed:
1) start with a small network and grow additional synapses and/or neurons until the desired behaviour is reached;
2) start with a large network and prune off synapses and/or neurons until the desired behaviour is retained.

The first approach, although it could be efficient since most of the job is performed on networks of small size, can be difficult to realize and control in order to reach network topologies which are "optimal" in some senses (see for example [3]). The second approach is much more studied and consists of finding a subset of network synaptic weights that, when set to zero, lead to the smallest increase of an error measure at the output. Several methods has been proposed in the last years in particular for the MLP [4,5,6,7,8]. However they can be slow, requiring periodic retraining of the network, and often require a "supervisor", a non-local algorithm which performs to pruning task. Moreover, some of them can require a difficult fine-tuning of the pruning coefficients in order to work correctly. Therefore it is highly desirable to find new methods which are able to dynamically adapt the network topology to the problem. In this paper, using some observations on the biological network plasticity, an idea is introduced to develop a procedure for varying the network topology by simply pruning off connections in an initial overdimensioned fully connected network.

2. The Proposed Method

The plasticity of brain, i. e. the capability of changing neural connections over time, is an extensively studied phenomenon. In this field the work of Hubel and Wiesel on measuring changes in the ocular dominance statistics of a large neuron sample of the visual cortex in mammalians (particularly cat and monkey) are a fundamental milestone. They found that monocular deprivation drives most neurons of the cortex to prefer the open eye, provided that the animal is still sufficiently young [9]. In particular it is now clear that the plasticity varies over the course of ontogenetic development and reaches a sharp maximum during the so-called critical period, during which a strong change of the brain structure is driven by external environmental stimuli. Recently [10], it has been proposed that, during the critical period, the network changes not only the synaptic strengths
of connection but also partially its internal topology. These changes are driven by either the received external stimuli and the pre-existent connection layout.

We have found that this idea can be applied also to simple artificial neural networks. By using only the external stimuli (i.e. input and output patterns) and the knowledge of the synaptic weights it is possible to adapt the network to the problem, changing concurrently the network synapses and the connection topology.

A simple procedure to prune off connection from a fully connected network is presented here. No new connections are created, although a similar approach could be followed to derive a procedure to add synapses to the network. In order to verify if a particular connection is necessary or not to solve a particular problem, a measure of the importance of the connection is necessary. According to the biological behaviour, this measure must be proportional either to the external stimuli and to pre-existent synaptic strength. A very simple formula can be derived by defining the following quantity:

- the synaptic activity of the connection of the i-th neuron coming from the j-th neuron and relative to the p-th training pattern

$$a_{ij}(p) = \left[ w_{ij} \cdot \text{Act}_j(p) \right]^2$$  \hspace{1cm} (1)

where $w_{ij}$ is the synaptic weight of the connection between the i-th and j-th neurons, and $\text{Act}_j(p)$ is the activation of the j-th neuron corresponding to the p-th input pattern. This activity can be averaged on the whole training set

$$\bar{a}_{ij} = \frac{1}{N_p} \sum_{p=1}^{N_p} a_{ij}(p)$$  \hspace{1cm} (2)

where $N_p$ is the number of the input training patterns. Since each neuron is driven by several different synapses, an average total synaptic activity relative to the i-th neuron can be defined

$$A_i = \sum_{j=1}^{N_j} \bar{a}_{ij}$$  \hspace{1cm} (3)

where $N_j$ is the total number of inputs to the i-th neuron. A measure of importance of the synapses can be obtained by comparing its averaged activity with respect to the the total average activity relative to the neuron which the synapse belongs to. Therefore the Percentage Average Synaptic Activity (PASA) can be defined as follows

$$\text{PASA}_{ij} = \frac{100 \bar{a}_{ij}}{A_i}$$  \hspace{1cm} (4)

During the learning phase the previous quantities vary until they reach stable values. The idea, used for changing the network topology, consists in dynamically eliminating all the connections which present a PASA lower than a threshold value varying according to a predefined curve.

In the forward step of the learning phase, in each neuron the synaptic activity of every connection relative to the current training pattern, is computed and accumulated. At the end of the training set all the PASA's can be computed. These values are compared with a threshold $\text{Th}(p)$ which is a function of the epochs in a way depending on the particular problem. The connections with a PASA lower than the threshold value are pruned off.

It can be noted that this algorithm is local in the sense that each neuron requires only local quantities to compute the PASA's. Moreover, the computational cost is low, since:

- the calculation of the synaptic activity per training pattern requires one multiplication/accumulation operation per synapsis (the multiplication can be avoided by using the absolute value instead of the square value in (1));
- the calculation of the PASA's of each connection relative to the i-th neuron at the end of one training set requires $(N_j-1)$ sums, one division and one multiplication per synapsis. The comparison between the PASA's and the threshold can be easily arranged in order to further reduce the computational burden.

3. Experimental Results on Recurrent Networks

A neural network is considered here in which there are input, hidden and output neurons and where all possible connections among these units are allowed, specifically including recursive connections and self-feedback connections. The algorithm for training the recursive network is derived from [11]. The proposed method has been tested on several simple applications, in particular the experimental results relative to the binary 2x2 bits multiplication problems are reported here.

The initial fully connected network used in this case was composed of 5 neurons with a total of 50 parameters. The PASA's have been further averaged on 100 epochs during the learning phase. Several trials were carried out using different initial sets of weights and offsets and various threshold curves have been tested for pruning off. Good results, in terms of convergence rate and pruning capability, have been obtained by using the curve in Fig. 1 which closely resembles the critical period of the biological neural networks.

![Fig. 1 Threshold function Th(p) used in the binary multiplication experiment.](image)

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Two sets of experiments were carried out after the proposed algorithm had reduced the original network by pruning off several connections. In the first the obtained topology was tested by performing a new learning phase on pruned networks loaded with small random set of weights and offsets. The performance of these networks were at least as good as those of the original fully connected network and often they worked better. On the contrary, in the second set of experiments several networks were built from the initial fully connected network by pruning off at random positions the same number of connections eliminated by the proposed algorithm. These networks, during the learning phase, gave
very poor results, often they did not converge at all, with respect to the equivalent network obtained by the proposed algorithm.

4. Experimental Results on Multilayered Networks

The proposed method has been tested also with the well-known Multilayer Perceptron (MLP) model [1]. The results of several simple experiments are reported here. In all cases, the PASA's are averaged on 2 learning epochs, although this was seen to be not critical. The threshold function used in these experiments is the gaussian function:

\[ \text{Th}(p) = \frac{1}{\sigma} \exp \left( -\frac{(p - m)^2}{2\sigma^2} \right) \]

where "v", "m", and \( \sigma \) are constant values, and the factor 100 has been removed. Moreover, a simple procedure has been added to the pruning algorithm which allows in a layer to prune off an entire neuron if it is not connected to any neuron of the upper layer.

In the first experiment, networks with 2 inputs, 4 hidden neurons and 1 output have been used to classify 100 points equally distributed inside and outside a unit circle on the X-Y plane. The high output classifies points inside the circle, while the null output classifies points outside it. 50 networks with different, randomly selected initial weights have been generated and used in the experiment. Each network was trained over 300 epoches using the 100 points as learning set. Fig. 2 shows the original network (a) and three common topologies obtained after pruning (b,c,d).

The various simulations have shown that the best results are obtained when the gaussian threshold function is centred on a value positioned after few tens of epoches but before the complete convergence has been reached. The values of "m" and \( \sigma \) parameters are therefore not critical providing that the network is not disturbed during the early and late phases of learning. Typical values in this example range from 10 to 60 for "m" and from 4 to 8 for \( \sigma \). The "v" value depends also on the size of the network, i.e. on the number of synapses per neuron, and it is not particularly critical. Notice that selecting and \( \sigma \) small allows to greatly reduce the computational burden since most of the learning is performed on a reduced network.

In the second experiment, networks with 2 inputs, 8 hidden neurons and 1 output have been used to classify 100 points equally distributed inside and outside a unit circle on the X-Y plane. A regular grid of 500 points was used as learning set, while 2000 points, randomly selected inside the region, were used as test set. Fig. 3 shows the original network (a) and three common topologies obtained after pruning (b,c,d).

Table 1 MSE and maximum output error obtained by the networks of Fig. 3 with the training and test set. The number of wrong classifications is also reported.

<table>
<thead>
<tr>
<th></th>
<th>training set</th>
<th>test set (2000 points)</th>
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<tr>
<td></td>
<td>MSE MaxErr</td>
<td>MSE MaxErr failed</td>
</tr>
<tr>
<td>Pruned 62.5 %</td>
<td>0.0602 0.4293</td>
<td>0.0662 0.99 174</td>
</tr>
<tr>
<td>Pruned 0%</td>
<td>0.0543 0.506</td>
<td>0.0655 0.99 169</td>
</tr>
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5. Conclusions

A simple generalization experiment has been also performed. Table 1 reports the MSE and maximum error values obtained by the 2 networks after 150 training epoches, using the test set as input. Although in this case the redundancy of the complete network is little, the pruned network shows similar or better performance on the test set with respect to the complete network, using a number of synapses reduced to the 37.5% of the original.

References

Fig. 3 (a) Complete network; (b) network obtained by the proposed method with \( v=0.1, m=30, \sigma=4 \).

Fig. 4 Plot of the threshold function used in fig. 3b vs the number of epochs.

Fig. 5 Plot of the MSE vs the number of learning epochs obtained (a) by the network of fig. 3a and (b) by the network of fig. 3b.

Fig. 6 Plot of the maximum output error vs the number of learning epochs obtained (a) by the network of fig. 3a and (b) by the network of fig. 3b.