

# Forming Neural Networks Design through Evolution

Eva Volná

University of Ostrava, 30ht dubna st. 22, 701 03 Ostrava, Czech Republic

**Abstract.** Neuroevolution techniques have been successful in many sequential decision tasks such as robot control and game playing. This paper aims at evolution in artificial neural networks (e.g. neuroevolution). Here is presented a neuroevolution system evolving populations of neurons that are combined to form the fully connected multilayer feedforward network with fixed architecture. In this paper, the transfer function has been shown to be an important part of architecture of the artificial neural network and have significant impact on an artificial neural network's performance. In order to test the efficiency of described method, we applied it to the alphabet coding problem.

## 1 Introduction to Neuroevolution

Evolutionary algorithms refer to a class of population-based stochastic search algorithms that are developed from ideas and principles of natural evolution. They include [1] evolution strategies, evolutionary programming, and genetic algorithms. Evolutionary algorithms are particularly useful for dealing with large complex problems which generate many local optima. They are less likely to be trapped in local minima than traditional gradient-based search algorithms. They do not depend on gradient information and thus are quite suitable for problems where such information is unavailable or very costly to obtain or estimate. They can even deal with problems, where no explicit and/or exact objective function is available. These features make them much more robust than many other search algorithms. Fogel [2] and Bäck et al. [3] give a good introduction to various evolutionary algorithms for optimization. One important feature of all these algorithms is their population-based search strategy. Individuals in a population compete and exchange information with each other in order to perform certain tasks. A general framework of evolutionary algorithms can be described as follows:

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generate the initial population G(0) at random, set i=0
REPEAT
  ..Evaluate each individual in the population;
  ..Select parents from G(i) based on their fitness
    in G(i);
  ..Apply search operators to parents and produce
    offspring which form G(i+1);
  ..i = i+ 1;
UNTIL 'termination criterion' is satisfied
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Neuroevolution represents a combination of neural networks and evolutionary algorithms, where neural networks are the phenotype being evaluated. The genotype is a compact representation that can be translated into an artificial neural network. Evolution has been introduced into artificial neural networks at roughly three different levels: connection weights, architectures, and learning rules. The evolution of connection weights provides a global approach to connection weights training, especially when gradient information of the error function is difficult or costly to obtain. Due to the simplicity and generality of the evolution and the fact that gradient-based training algorithms often have to be run multiple times in order to avoid being trapped in a poor local optimum, the evolutionary approach is quite competitive. The evolution of architectures enables artificial neural networks to adapt their topologies to different tasks without human intervention and thus provides an approach to automatic artificial neural network design. Simultaneous evolution of artificial neural network architectures and connection weights generally produces better results. The evolution of learning rules in artificial neural networks can be used to allow an artificial neural network to adapt its learning rule to its environment. In a sense, the evolution provides artificial neural network with the ability of learning to learn. Global search procedures such as evolutionary algorithms are usually computationally expensive. It would be better not to employ evolutionary algorithms at all three levels of evolution in neural networks. It is, however, beneficial to introduce global search at some levels of evolution, especially when there is little prior knowledge available at that level and the performance of the artificial neural network is required to be high, because the trial-and-error or heuristic methods are very ineffective in such circumstances. With the increasing power of parallel computers, the evolution of large artificial neural networks becomes feasible. Not only can such evolution discover possible new artificial neural network architectures and learning rules, but it also offers a way to model the creative process as a result of artificial neural network's adaptation to a dynamic environment.

## 2 Overview of the Evolution of Node Transfer Functions

The discussion on the evolution of architectures so far only deals with the topological structure of architecture. The transfer function of each node in the architecture has been usually assumed that is fixed and predefined by human experts, at least for all the nodes in the same layer. Little work has been only done on the evolution of node transfer function up to now. Mani proposed a modified backpropagation, which performs gradient descent search in the weight space as well as the transfer function space [4], but connectivity of artificial neural networks was fixed. Lovel and Tsoi investigated the performance of Neocognitrons with various S-cell and C-cell transfer functions, but did not adopt any adaptive procedure to search for an optimal transfer function automatically [5]. Stork et al. [6] were, to our best knowledge, the first to apply evolutionary algorithms to the evolution of both topological structures and node transfer functions even though only simple artificial neural networks with seven nodes were considered. The transfer function was specified in the structural genes in their genotypic representation. It was much more complex than the usual sigmoid function because authors in [6] tried to model biological neurons. White and Ligomenides [7]

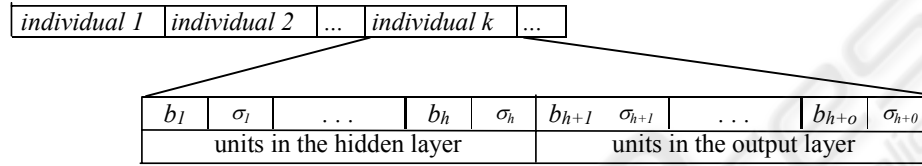
adopted a simpler approach to the evolution of both topological structures and node transfer functions. For each individual (i.e. artificial neural network) in the initial population, 80% nodes in the artificial neural network used the sigmoid transfer function and 20% nodes used the Gaussian transfer function. The evolution was used to decide the optimal mixture between these two transfer functions automatically. The sigmoid and Gaussian transfer function themselves were not evolvable. No parameters of the two functions were evolved. Liu and Yao [1] used evolutionary programming to evolve artificial neural networks with both sigmoidal and Gaussian nodes. Rather than fixing the total number of nodes and evolve mixture of different nodes, their algorithm allowed growth and shrinking of the whole artificial neural network by adding or deleting a node (either sigmoidal or Gaussian). The type of node added or deleted was determined at random. Hwang et al. [8] went one step further. They evolved topology of artificial neural network, node transfer function, as well as connection weights for projection neural networks. Sebald and Chellapilla [9] used the evolution of node transfer function as an example to show the importance of evolving representations. Representation and search are the two key issues in problem solving. Co-evolving solutions and their representations may be an effective way to tackle some difficult problems where little human expertise is available. In principle, the difference in transfer functions could be as large as that in the function type, e.g. that between a hard limiting threshold function and Gaussian function, or as small as that in one of parameters of the same type of function, e.g. the slope parameter of the sigmoid function. The decision on how to encode transfer functions in chromosomes depends on how much prior knowledge and computation time is available. This suggests some kind of indirect encoding method, which lets developmental rules to specify function parameters if the function type can be obtained through evolution, so that more compact chromosomal encoding and faster evolution can be achieved. One point worth mentioning here is the evolution of both connectivity and transfer functions at the same time [6] since they constitute a complete architecture. Encoding connectivity and transfer functions into the same chromosome makes it easier to explore nonlinear relations between them. Many techniques used in encoding and evolving connectivity could equally be used here.

### **3 Evolution Design of Neural Networks With Fixed Topology**

In the paper, the transfer function has been shown to be an important part of architecture of the artificial neural network, one has significant impact on artificial neural network's performance. Here is presented a neuroevolution system evolving populations of neurons that are combined to form the fully connected multilayer feedforward network with fixed architecture. Neuroevolution evolves transfer functions of each unit in hidden and output layers of the network. The system maintains diversity in the population, because a dominant neural phenotype is likely to end up in the same network more than once. As several different types of neurons are usually necessary to solve a problem, networks with too many copies of the same neuron are likely to fail. The dominant phenotype then loses fitness and becomes less dominant. The system works well because it makes sure neurons get the credit they deserve, unlike some other neuroevolution techniques, where bad neurons can share

in a good network or good neurons can be brought down by their network. It also works by decomposing the task, breaking the search into smaller, more manageable parts.

In the following is described a method of automatic searching the node transfer function architecture in multilayer feedforward network: First, we must propose neural network architecture before the main calculation. We get the number of input ( $m$ ) and output ( $o$ ) units from the training set. Next, we have to define the number of hidden units ( $h$ ) that is very confounding issue, because it is generally more difficult to optimize large networks than small ones. Thereafter the process of evolutionary algorithms is applied. Chromosomes are generated for every individual from the initial population as follows, see Fig. 1:



**Fig. 1.** Population of individuals and their chromosomes.

Symbols  $b_i$  ( $i = 1, \dots, h+o$ ) refers to varies types of activation functions [10]:

- $b_i = 1$ , if the activation function is a *binary sigmoid function*:

$$f(x) = \frac{1}{1 + \exp(-\sigma x)}. \quad (1)$$

where  $\sigma$  is the steepness parameter, which value is set in the initial population randomly (e.g.  $\sigma_i \in (0; 7>)$ ).

- $b_i = 2$ , if the activation function is a *binary step function with threshold  $\theta$* .

$$f(x) = \begin{cases} 1 & \text{if } x \geq \theta \\ 0 & \text{if } x < \theta \end{cases}. \quad (2)$$

the steepness parameter  $\sigma_i$  is not define here thus we assigned value 0 to it.

- $b_i = 3$  if the activation function is a *Gaussian function*:

$$f(x) = \exp(-x^2). \quad (3)$$

the steepness parameter  $\sigma_i$  is not define here thus we assigned value 0 to it.

–  $b_i = 4$ , if the activation function is a *saturated linear function*:

$$f(x) = \begin{cases} 0 & \text{if } x < 0 \\ x & \text{if } 0 \leq x \leq 1. \\ 1 & \text{if } x > 1 \end{cases} \quad (4)$$

the steepness parameter  $\sigma_i$  is not define here thus we assigned value 0 to it.

Next, we calculate an error value ( $E$ ) between the desired and the real output after defined partial training with genetic algorithms. Adaptation of each individual starts with randomly generated weight values that are the same for each neural network in the given population. On the basis of it is calculated a fitness function for every individual as follows:

$$Fitness_i = E_{max} - E_i. \quad (5)$$

for  $i = 1, \dots, N$ ,

where  $E_i$  is error for the  $i$ -th network after a partial adaptation;

$E_{max}$  is a maximal error for the given task,

$E_{max} = o \times pattern$

( $o$  is number of output units and  $pattern$  is number of patterns);

$N$  is the number of individuals in the population.

All of the calculated fitness function values of the two consecutive generations are sorted descending and the neural network representation attached to the first half creates the new generation. For each fitness function is calculated the probability of reproduction its existing individual by standard method [11]. One-point *crossover* was used to generate two offspring. If the input condition of *mutation* is fulfilled (e.g. if a randomly number is generated, that is equal to the defined constant), one of the individual is randomly chosen. There is randomly replaced one place in its genetic representation by a random value from the set of permitted values. Our calculation finishes, when the population is composed only from the same individuals.

## 4 Experiments

In order to test the efficiency of described method, we applied it to the alphabet coding problem that exists in cryptography. Neural networks can be also used in encryption or decryption algorithms, where parameters of adapted neural networks are included to cipher keys [12]. Cipher keys must have several heavy attributes. The best one is the singularity of encryption and cryptanalysis [13]. Encryption is a process in which we transform the open text (e.g. news) to cipher text according to rules. Cryptanalysis of the news is the inverse process, in which the receiver of the cipher transforms it to the original text. The open text is composed from alphabet characters, digits and punctuation marks. The cipher text has usually the same composition as the open text.

We worked with multilayer neural networks, which topologies were based on the training set (see Table 1). The chain of chars of the plain text in a training set is equivalent to a binary value that is 96 less than its ASCII code. The cipher text is then a random chain of bits.

**Table 1.** The set of patterns (the training set).

THE PLAIN TEXT			THE CIPHER TEXT	THE PLAIN TEXT			THE CIPHER TEXT
Char	ASCII code (DEC)	The chain of bits	The chain of bits	Char	ASCII code (DEC)	The chain of bits	The chain of bits
a	97	00001	000010	n	110	01110	011100
b	98	00010	100110	o	111	01111	101000
c	99	00011	001011	p	112	10000	001010
d	100	00100	011010	q	113	10001	010011
e	101	00101	100000	r	114	10010	010111
f	102	00110	001110	s	115	10011	100111
g	103	00111	100101	t	116	10100	001111
h	104	01000	010010	u	117	10101	010100
i	105	01001	001000	v	118	10110	001100
j	106	01010	011110	w	119	10111	100100
k	107	01011	001001	x	120	11000	011011
l	108	01100	010110	y	121	11001	010001
m	109	01101	011000	z	122	11010	001101

The initial population contains 30 three-layer feedforward neural networks. Each network architecture is 5 - 5 - 6 (e.g. five units in the input layer, five units in the hidden layer, and six units in the output layer), because the alphabet coding problem is not linearly separable and therefore we cannot use neural network without hidden units. The nets are fully connected. We use the genetic algorithm with the following parameters: probability of mutation is 0.01 and probability of crossover is 0.5. Adaptation of each neural network in given population starts with randomly generated weight values that are the same for each neural network in the population. We also used genetic algorithms with the same parameters for the partial neural network adaptation, where number of generations for a partial adaptation was 500. Its chromosome representation is described in [14].

History of the error function is shown in the figure 3. There are shown average values of error functions in the given population. Other numerical simulations give very similar results. The “*binary sigmoid function*” represents an average value after adaptation with the binary sigmoid activation function consecutively with all



steepness parameters  $\sigma = \{1,2,3,4,5,6,7\}$ . The “*binary step function*” represents an adaptation with the binary step activation function (with the threshold  $\theta$ ), the “*saturated linear function*” represents an adaptation with the saturated linear activation function, and “*Gaussian activation*” represents an adaptation with the Gaussian activation function. Each of these mentioned representations is associated with all units in given neural network architecture. Opposite of this, the “*best individual*” represents an adaptation of the best individual in population, which chromosome is the following, see Fig. 2:

$b_1$	$\alpha$	$b_2$	$\sigma_2$	$b_3$	$\sigma_3$	$b_4$	$\alpha_4$	$b_5$	$\sigma_5$	$b_6$	$\sigma_6$	$b_7$	$\sigma_7$	$b_8$	$\sigma_8$	$b_9$	$\sigma_9$	$b_{10}$	$\sigma_{10}$	$b_{11}$	$\sigma_{11}$
1	5	1	7	1	1	3	0	1	5	3	0	1	7	3	0	1	5	1	6	1	2
units in the hidden layer												units in the output layer									

Fig. 2. The “best individual” chromosome in the last population.

## 5 Conclusions

All networks solve the alphabet coding task in our experiment, but artificial neural network with evolving transfer functions of each unit works well, because several different types of neurons are usually necessary to solve a problem. We can see that the proposed technique is really efficient for the presented purpose, see the Fig. 3. Networks with too many copies of the same neuron work usually worse.

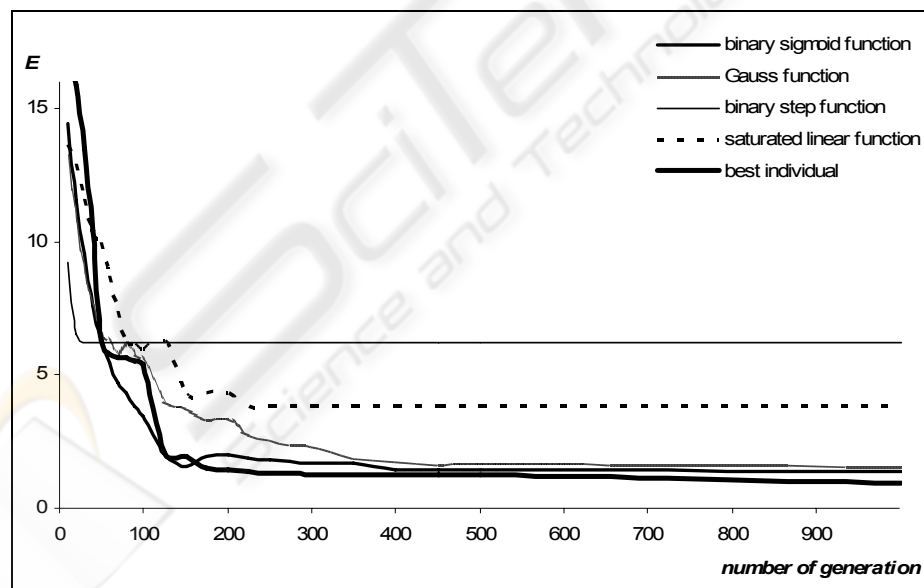


Fig. 3. The error function history.

Here, the transfer function is shown to be an important part of architecture of the artificial neural network and have significant impact on artificial neural network's

performance. Transfer functions of different units can be different and decided automatically by an evolutionary process, instead of assigned by human experts. In general, nodes within a group, like layer, in an artificial neural network tend to have the same type of transfer function with possible difference in some parameters, while different groups of nodes might have different types of transfer function.

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