# Generalized Tree-Like Self-Organizing Neural Networks with Dynamically Defined Neighborhood for Cluster Analysis

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Abstract. The paper presents a generalization of self-organizing neural networks of spanning-tree-like structures and with dynamically defined neighborhood (SONNs with DDN, for short) for complex cluster-analysis problems. Our approach works in a fully-unsupervised way, i.e., it operates on unlabelled data and it does not require to predefine the number of clusters in a given data set. The generalized SONNs with DDN, in the course of learning, are able to disconnect their neuron structures into sub-structures and to reconnect some of them again as well as to adjust the overall number of neurons in the system. These features enable them to detect data clusters of virtually any shape and density including both volumetric ones and thin, shell-like ones. Moreover, the neurons in particular sub-networks create multi-point prototypes of the corresponding clusters. The operation of our approach has been tested using several diversified synthetic data sets and two benchmark data sets yielding very good results.

**Keywords:** generalized self-organizing neural networks with dynamically defined neighborhood, multi-point prototypes of clusters, cluster analysis, unsupervised learning.

#### 1 Introduction

Data clustering or cluster analysis is an unsupervised process that aims at grouping unlabelled data records from a given data set into an unknown in advance number of cluster or groups. Elements of each cluster should be as "similar" as possible to each other and as "dissimilar" as possible from those of other clusters. Cluster analysis belongs to fundamental issues in data mining and machine learning with wide range of applications, cf., e.g., [15], [1].

This paper presents a technique for cluster analysis based on self-organizing neural networks (SONNs) of spanning-tree-like structures and with dynamically defined neighborhood (henceforward referred to as SONNs with DDN) outlined in Kohonen's work [10]. However, we propose an essential generalization of these networks by introducing original mechanisms that, in the course of learning:

L. Rutkowski et al. (Eds.): ICAISC 2014, Part II, LNAI 8468, pp. 713–725, 2014.

a) automatically adjust the number of neurons in the network, b) allow to disconnect the tree-like structure into sub-trees, and c) allow to reconnect some of the sub-trees preserving the no-loop spanning-tree properties. All these features enable them to detect data clusters of virtually any shape and density including both volumetric clusters and thin, piece-wise linear, shell, polygonal, etc. types of clusters. Similar, to above-listed, mechanisms have been proposed by us in [7], [8], [9] ([5], [6] present their earlier versions) to obtain - for clustering purposes - the so-called dynamic SONNs with one-dimensional neighborhood; they can be treated as a special case of the presently proposed generalized SONNs with DNN. First, their details are presented. Then, an illustration of their operation using several synthetic data sets containing data concentrations of various shapes and densities is given. Finally, their application to the clustering of two benchmark data sets is presented.

An idea of evolving topological structures of self-organizing neural networks has been addressed in the literature. However, some existing solutions do not directly deal with data clustering, e.g. in [11] and [14] evolving neuron trees are used to decrease the computational complexity of Winner-Takes-Most (WTM) learning algorithm. In [16] tree structures are used to visualize the obtained results for the purpose of comparison with conventional decision trees. Among data clustering techniques (to some extent alternative to our approach), evolving topological structures are proposed in [3], [13], and in [4], [2]. However, the approaches of [3], [13] do not enable to detect, in an automatic way, the number of clusters in data sets. In turn, the results of experiments presented in [4], [2] do not provide an information on how effective the proposed approaches are in terms of the automatic detection of the number of clusters in data sets.

# 2 Generalized SONNs with DDN for Clustering Analysis

First, we consider the conventional SONN with one-dimensional neighborhood. Assume that the network has n inputs  $x_1, x_2, \ldots, x_n$  and consists of m neurons arranged in a chain; their outputs are  $y_1, y_2, \ldots, y_m$ , where  $y_j = \sum_{i=1}^n w_{ji} x_i$ ,  $j = 1, 2, \ldots, m$  and  $w_{ji}$  are weights connecting the i-th input of the network with the output of the j-th neuron. Using vector notation  $(\mathbf{x} = (x_1, x_2, \ldots, x_n)^T, \mathbf{w}_j = (w_{j1}, w_{j2}, \ldots, w_{jn})^T)$ ,  $y_j = \mathbf{w}_j^T \mathbf{x}$ . The learning data consists of L input vectors  $\mathbf{x}_l$   $(l = 1, 2, \ldots, L)$ . The first stage of any Winner-Takes-Most (WTM) learning algorithm that can be applied to the considered network, consists in determining the neuron  $j\mathbf{x}$  winning in the competition of neurons when learning vector  $\mathbf{x}_l$  is presented to the network. Assuming the normalization of learning vectors, the winning neuron  $j\mathbf{x}$  is selected such that

$$d(\boldsymbol{x}_l, \boldsymbol{w}_{j\boldsymbol{x}}) = \min_{j=1,2,\dots,m} d(\boldsymbol{x}_l, \boldsymbol{w}_j), \tag{1}$$

where  $d(\mathbf{x}_l, \mathbf{w}_j)$  is a distance measure between  $\mathbf{x}_l$  and  $\mathbf{w}_j$ ; throughout this paper, the Euclidian distance measure will be applied

$$d_E(\boldsymbol{x}_l, \boldsymbol{w}_j) = \sqrt{\sum_{i=1}^n (x_{li} - w_{ji})^2}.$$
 (2)

The WTM learning rule can be formulated as follows:

$$\boldsymbol{w}_{j}(k+1) = \boldsymbol{w}_{j}(k) + \eta_{j}(k)N(j, j_{\boldsymbol{x}}, k)[\boldsymbol{x}(k) - \boldsymbol{w}_{j}(k)], \tag{3}$$

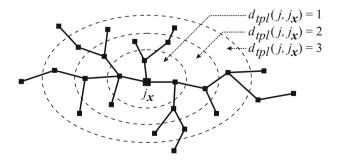
where k is the iteration number,  $\eta_j(k)$  is the learning coefficient, and  $N(j, j_{\boldsymbol{x}}, k)$  is the neighborhood function. At this point, we have to address the problem of a spanning-tree-like structure of the SONN with DDN. The neighborhood of a given neuron in such a topology is defined along the arcs emanating from that neuron as illustrated in Fig. 1. Therefore, the paths between any two neurons in such a structure are the pieces of SONN with one-dimensional neighborhood. The topological distance  $d_{tpl}(j, j_{\boldsymbol{x}})$  between the  $j_{\boldsymbol{x}}$ -th neuron and some other neurons is equal to 1 if those other neurons are direct neighbors of the  $j_{\boldsymbol{x}}$ -th one as shown in Fig. 1. The distance  $d_{tpl}(j, j_{\boldsymbol{x}}) = 2$  for the neurons that are second along all paths starting at the  $j_{\boldsymbol{x}}$ -th one (see Fig. 1), etc. The topological distance measure is the basis of the neighborhood function  $N(j, j_{\boldsymbol{x}}, k)$ . In this paper, the Gaussian-type neighborhood function is used:

$$N(j, j_{\boldsymbol{x}}, k) = e^{-\frac{d_{tpl}^2(j, j_{\boldsymbol{x}})}{2\lambda^2(k)}}$$

$$\tag{4}$$

where  $\lambda(k)$  is the radius of the neighborhood (the width of the Gaussian "bell"). As already mentioned, the generalization of the above-presented SONN with DDN consists in introducing mechanisms that allow the network:

- to automatically adjust the number of neurons in the network by removing low-active neurons from the network and adding new neurons in the areas of existing high-active neurons,
- II) to automatically disconnect the network, as well as to reconnect some of the sub-networks again preserving the no-loop spanning-tree properties.



**Fig. 1.** Examples of neighborhood of the jx-th neuron

These two features enable the generalized SONN with DDN to fit in the best way the structures that are "encoded" in data sets to display them to the user. In particular, the number of disconnected sub-networks is equal to the number of clusters detected in a given data set. Moreover, the neurons in a given sub-network create a multi-point prototype of the corresponding cluster. The mechanisms I and II are implemented by activating (under some conditions) - after each learning epoch - five successive operations (cf. [7]):

- 1) the removal of single, low-active neurons,
- 2) the disconnection of the network (sub-network) into two sub-networks,
- 3) the removal of small-size sub-networks,
- 4) the insertion of additional neurons into the neighborhood of high-active neurons in order to take over some of their activities,
- 5) the reconnection of two selected sub-networks.

The operations 1, 3 and 4 are the components of the mechanism I, whereas the operations 2 and 5 govern the mechanism II. Based on experimental investigations, the following conditions for activating particular operations have been formulated.

Operation 1: The neuron no.  $j_r$  is removed from the network if its activity measured by the number of its wins  $win_{jr}$  - is below an assumed level  $win_{min}$ , i.e.,  $win_{jr} < win_{min}$ .  $win_{min}$  is experimentally selected parameter (usually,  $win_{min} \in \{2, 3, ..., 7\}$ ). The removal of the  $j_r$ -th neuron is followed by reconfiguration of the network topology as shown in Fig. 2. If the  $j_r$ -th neuron has only two neighbors (Fig. 2a), they are now topologically connected. In the case of three or more neighbors of the  $j_r$ -th unit (Fig. 2b), one of them, say  $j_1$ , which is nearest to the  $j_r$ -th one in terms of their weight-vector distance, is selected. Then, the remaining neighbors are topologically connected to the  $j_1$ -th unit.

Operation 2: The structure of the network is disconnected into two subnetworks by removing the topological connection between two neighboring neurons  $j_1$  and  $j_2$  (see Fig. 3) after fulfilling the following condition:  $d_{E,j_1j_2} > \alpha_{dsc}d_{E,avr}$  where  $d_{E,j_1j_2} = d_E(\mathbf{x}_{j_1},\mathbf{x}_{j_2})$  ( $d_E$  is defined in (2)),  $d_{E,avr} = \frac{1}{P}\sum_{p=1}^{P}d_{E,p}$  is the average distance between two neighboring neurons for all pairs of such neurons in the network ( $d_{E,p}$  is the  $d_E$  distance for the p-th pair of neighboring neurons,  $p = 1, 2, \ldots, P$ ), and  $\alpha_{dsc}$  is experimentally selected parameter governing the disconnection operation (usually,  $\alpha_{dsc} \in [2, 4]$ ).

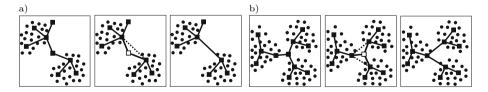


Fig. 2. Removal of single, low-active neuron connected with two (a) and three (in general, more than two) (b) neighboring neurons (illustrations of the exemplary network structure before, during, and after the operation, respectively)



Fig. 3. Disconnection of the network into two sub-networks (illustration of the exemplary network structure before, during, and after the operation, respectively)

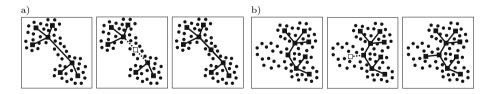
Operation 3: A sub-network that contains  $m_s$  neurons is removed from the system if  $m_s < m_{s,min}$ , where  $m_{s,min}$  is experimentally selected parameter (usually,  $m_{s,min} \in \{3,4\}$ ).

The operation of the insertion of additional neurons into the neighborhood of high-active neurons in order to take over some of their activities covers 2 cases denoted by 4a and 4b, respectively.

Operation 4a: A new neuron, labelled as (new), is inserted between two neighboring and high-active neurons  $j_1$  and  $j_2$  (see Fig. 4a) if they fulfil the following conditions:  $win_{j_1} > win_{max}$  and  $win_{j_2} > win_{max}$ , where  $win_{j_1}$  and  $win_{j_2}$  are the numbers of wins of particular neurons and  $win_{max}$  is experimentally selected parameter (usually  $win_{max} \in \{4, 5, ..., 9\}$ ). The weight vector  $\boldsymbol{w}_{(new)}$  of the new neuron is calculated as follows:  $\boldsymbol{w}_{(new)} = \frac{\boldsymbol{w}_{j_1} + \boldsymbol{w}_{j_2}}{2}$ .

Operation 4b: A new neuron (new) is inserted in the neighborhood of high-active neuron  $j_1$  surrounded by low-active neighbors (see Fig. 4b) if the following conditions are fulfilled:  $win_{j_1} > win_{max}$  and  $win_j < win_{max}$  for j such that  $d_{tpl}(j, j_1) = 1$ , where  $win_{j_1}$  and  $win_{max}$  are as in Operation 4a and  $win_j$  is the number of wins of the j-th neuron. The weight vector  $\mathbf{w}_{(new)} = [w_{(new)1}, w_{(new)2}, \dots, w_{(new)n}]^T$  is calculated as follows:  $w_{(new)i} = w_{j_1i}(1 + \xi_i)$ ,  $i = 1, 2, \dots, n$ , where  $\xi_i$  is a random number from the interval [-0.01, 0.01]. Therefore, particular components of high-active neuron  $j_1$ , after experimentally selected random modification in the range of [-1%, 1%], give the weight vector  $\mathbf{w}_{(new)}$  of the new neuron. It is a starting point for the new neuron in its further evolution as the learning progresses.

Operation 5: Two sub-networks  $S_1$  and  $S_2$  are reconnected by introducing topological connection between neurons  $j_1$  and  $j_2$  ( $j_1 \in S_1$ ,  $j_2 \in S_2$ ) - see Fig. 5 - after fulfilling condition:  $d_{E,j_1j_2} < \alpha_{con} \frac{d_{E,avr_{S_1}} + d_{E,avr_{S_2}}}{2}$ .  $d_{E,j_1j_2}$  is the same



**Fig. 4.** Insertion of additional neuron between two high-active neighbouring neurons (a) and into the neighborhood of a single high-active neuron (b) (illustrations of the exemplary network structure before, during, and after the operation, respectively)

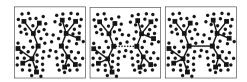


Fig. 5. Reconnection of two sub-networks (illustration of the exemplary network structure before, during, and after the operation, respectively)

as in Operation 2.  $d_{E,avr_{S_1}}$  and  $d_{E,avr_{S_2}}$  are calculated for sub-networks  $S_1$  and  $S_2$ , respectively, in the same way as  $d_{E,avr}$  is calculated in Operation 2 for the considered network.  $\alpha_{con}$  is experimentally selected parameter that controls the reconnection process (usually,  $\alpha_{con} \in [3,5]$ ).

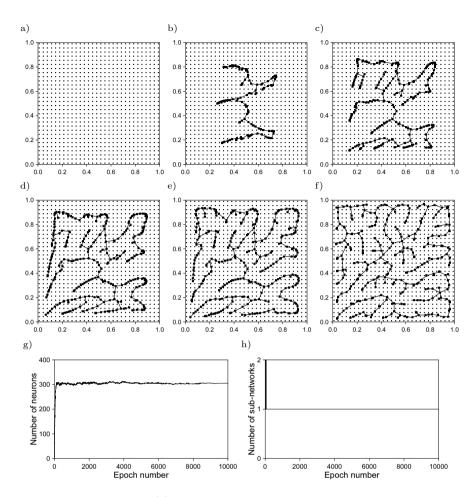
The conditions that govern Operations 1 through 5 are checked after each learning epoch. The condition that is fulfilled activates the appropriate operation.

In the experiments presented below, the following values of control parameters are selected:  $win_{min} = 2$ ,  $win_{max} = 4$ ,  $m_{s,min} = 3$ ,  $\alpha_{dsc} = 3$ , and  $\alpha_{con} = 4$ . Moreover, the learning process is carried out through 10000 epochs, the learning coefficient  $\eta_j(k) = \eta(k)$  of (3) linearly decreases over the learning horizon from  $7 \cdot 10^{-4}$  to  $10^{-6}$ , the neighborhood radius  $\lambda(k) = \lambda$  of (4) is equal to 2, and the initial number of neurons in the network (at the start of the learning process) is equal to 2.

### 3 Cluster Analysis in Two-Dimensional Synthetic Data Sets

Fig. 6 shows the performance of the generalized SONN with DDN applied to the set of uniformly distributed data (i.e., without any clusters in them). It is a hard-to-pass test for very many clustering techniques, especially those generating a predefined number of clusters - regardless of whether any clusters exist in data or not. Our approach perfectly passes this test. After initial jump to 2, the number of sub-networks (clusters) stabilizes on 1, i.e., the system detects one big cluster in data (see Fig. 6h) and generates a multi-point prototype for it (see neurons of Fig. 6f). Fig. 6g shows the adjustment of the number of neurons in the network in the course of learning.

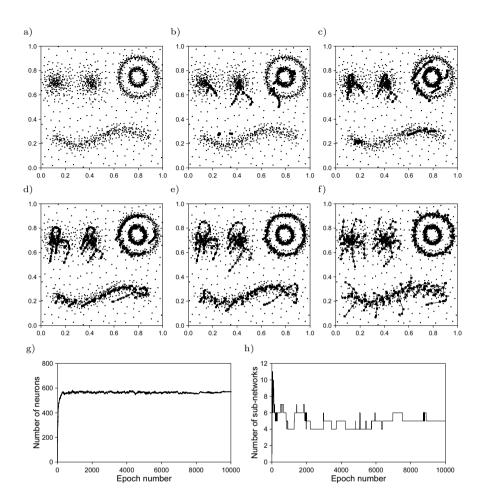
Figs. 7, 8, and 9 present further illustrations of the performance of our approach applied to various two-dimensional synthetic data sets. Fig. 7a presents data set used in [17]; it contains two overlapped Gaussian distributions, two concentric rings, and a sinusoidal curve with 10% noise added to data. Fig. 8a presents data set with various types of clusters in it. Both, thin piece-wise linear and two-ellipsoidal as well as volumetric of various shapes and densities clusters are considered. Finally, Fig. 9a presents a "classical" two-spiral data set. As the above-listed figures show, in all of these data sets our approach detects the correct numbers of clusters and generates multi-point prototypes for them.



**Fig. 6.** Synthetic data set (a) and the evolution of the generalized SONN with DDN in it in learning epochs: b) no. 20, c) no. 50, d) no. 100, e) no. 200, and f) no. 10 000 (end of learning), as well as plots of the number of neurons (g) and the number of sub-networks (clusters) (h) vs. epoch number

## 4 Cluster Analysis in Selected Benchmark Data Sets

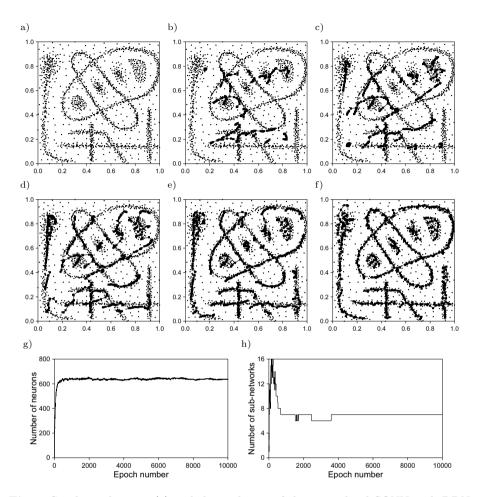
Our approach will now be tested using two multidimensional benchmark data sets such as *Breast Cancer Wisconsin* (*Diagnostic*) and *Congressional Voting Records* (*BCWD* and *CVR*, for short) [12]. *BCWD* data set has 569 records and 30 numerical attributes, whereas *CVR* data set - 435 records and 16 nominal attributes. It is essential to note that our approach does not utilize the knowledge on class assignments of particular records and on the number of classes (equal to 2



**Fig. 7.** Synthetic data set (a) and the evolution of the generalized SONN with DDN in it in learning epochs: b) no. 20, c) no. 50, d) no. 100, e) no. 200, and f) no. 10 000 (end of learning), as well as plots of the number of neurons (g) and the number of sub-networks (clusters) (h) vs. epoch number

in both sets). Our approach works in a fully-unsupervised way, i.e., it operates on unlabelled data and without any predefinition of the number of clusters (classes).

Figs. 10 and 11 as well as Tables 1 and 2 present the performance of our approach applied to both data sets. First, Figs. 10b and 11b show that our approach detects the correct number of clusters in both data sets. Second, since the number of classes and class assignments are known in both original data



**Fig. 8.** Synthetic data set (a) and the evolution of the generalized SONN with DDN in it in learning epochs: b) no. 20, c) no. 50, d) no. 100, e) no. 200, and f) no. 10 000 (end of learning), as well as plots of the number of neurons (g) and the number of sub-networks (clusters) (h) vs. epoch number

sets, a direct verification of the obtained results is also possible (see Tables 1 and 2). The percentages of correct decisions, equal to 90.51% (BCWD data set) and 94.71% (CVR data set), regarding the class assignments are very high (especially, that they have been achieved by the unsupervised-learning systems operating on benchmark data sets).

Malignant

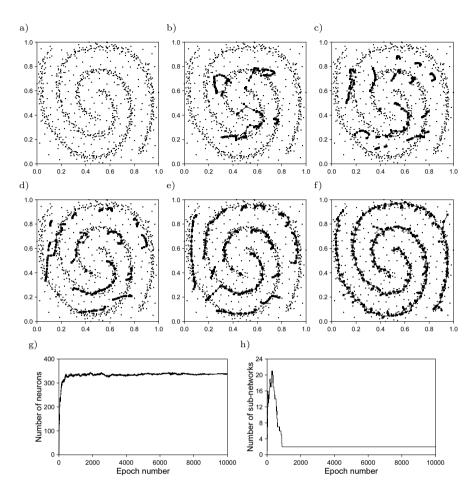
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ALL

212

357

569



**Fig. 9.** Synthetic data set (a) and the evolution of the generalized SONN with DDN in it in learning epochs: b) no. 20, c) no. 50, d) no. 100, e) no. 200, and f) no. 10 000 (end of learning), as well as plots of the number of neurons (g) and the number of sub-networks (clusters) (h) vs. epoch number

Number Number of decisions for Number Number Percentage Class label of records sub-network labelled: of correct of correct of wrong MalignantBenigndecisions decisions decisions

166

8

174

**Table 1.** Clustering results for BCWD data set

46

349

395

166

349

515

46

8

**54** 

78.30%

97.76%

90.51%

Class label	Number of records	Number of decisions for sub-network labelled:		Number of correct	Number of wrong	Percentage of correct
		Republican	Democrat	decisions	decisions	decisions
Republican	168	158	10	158	10	94.05%
Democrat	267	13	254	254	13	95.13%
		-			- 0	001-070

Table 2. Clustering results for CVR data set

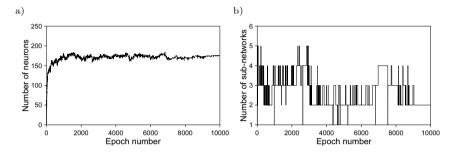


Fig. 10. Plots of the number of neurons (a) and the number of sub-networks (clusters) (b) vs. epoch number (BCWD data set)

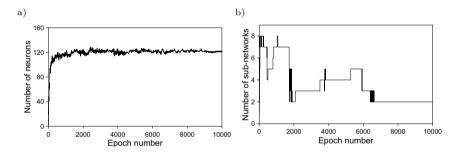


Fig. 11. Plots of the number of neurons (a) and the number of sub-networks (clusters) (b) vs. epoch number (CVR data set)

#### 5 Conclusions

The generalized SONNs with DDN that can be effectively applied in complex, multidimensional cluster-analysis problems have been presented in this paper. Our approach works in a fully-unsupervised way, i.e., it operates on unlabelled data and it does not require to predefine the number of clusters in a given data set. The proposed networks, in the course of learning, are able to disconnect their neuron structures into sub-structures and to reconnect some of them again as well as to adjust the overall number of neurons in the system. These features enable them to detect data clusters of virtually any shape and density

including both volumetric ones and thin, shell-like ones. Moreover, the neurons in particular sub-networks create multi-point prototypes of the corresponding clusters. The operation of our approach has been illustrated by means of several diversified synthetic data sets and then our approach has been tested using two benchmark data sets (*Breast Cancer Wisconsin (Diagnostic)* and *Congressional Voting Records*) yielding very good results.

Acknowledgments. The numerical experiments reported in this paper have been performed using computational equipment purchased in the framework of the EU Operational Programme Innovative Economy (POIG.02.02.00-26-023/09-00) and the EU Operational Programme Development of Eastern Poland (POPW.01.03.00-26-016/09-00).

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