

Dynamic generation of prototypes with self-organizing feature maps for classifier design

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Received 2 June 1999; accepted 1 October 1999

Abstract

We propose a new scheme for designing a nearest-prototype classifier using Kohonen's self-organizing feature map (SOFM). The net starts with the minimum number of prototypes which is equal to the number of classes. Then on the basis of the classification performance, new prototypes are generated dynamically. The algorithm merges similar prototypes and deletes less significant prototypes. If prototypes are deleted or new prototypes appear then they are fine tuned using Kohonen's SOFM algorithm with the winner-only update strategy. This adaptation continues until the system satisfies a termination condition. The classifier has been tested with several well-known data sets and the results obtained are quite satisfactory.

Keywords: Nearest-prototype classifier; Dynamic prototype generation; Self-organizing feature map; Split-merge technique

1. Introduction

Kohonen's self-organizing feature map (SOFM) has been successfully used in numerous fields of application such as speech recognition [1], robotics [2,3], industrial process control [5], image compression [4], etc. Designing of classifiers [6] and other pattern recognition systems based on SOFM [7] are some of the most successful areas of application. SOFM [8,9] has the interesting property of achieving a distribution of the weight vectors that approximates the distribution of the input data. This property of the SOFM can be exploited for designing nearest prototype classifiers. Here we propose a new approach for this. Although our training data are labeled, the SOFM is trained without using the class information. When the training is over, the weight vectors are converted into labeled prototypes of a classifier using the class information. The performance of the classifier is then evaluated. Based on the evaluation results a tuning

step consisting of deletion, merging, splitting and retraining of the net is performed. The evaluation and tuning are repeated until the number of prototypes stabilizes or the performance of the classifier reaches a satisfactory level. In case of highly overlapped class boundaries usually it is very difficult to estimate the adequate number of prototypes. Small number of prototypes suffer from large error rates, while at the other extreme a large number of prototypes make the system expensive. Here the tuning strategy is designed to strike a compromise between the classification performance and the number of prototypes for such data. These prototypes can be used to generate fuzzy rules for a fuzzy rule based pattern recognition system.

2. Self-organizing feature map

We view the self-organizing feature map as an algorithmic transformation $A_{\text{SOFM}}^D: R^p \rightarrow V(R^q)$ that is often advocated for visualization of metric-topological relationships and distributional density properties of feature vectors (signals) $X = \{\mathbf{x}_1, \dots, \mathbf{x}_N\}$ in R^p [9].

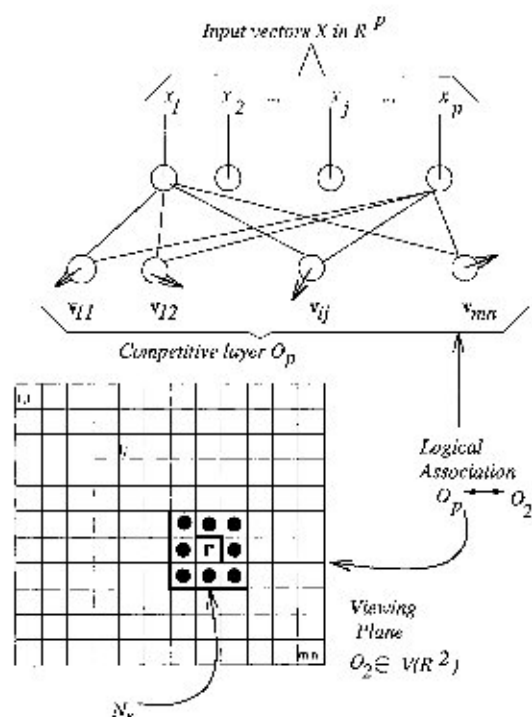


Fig. 1. The SOFM architecture.

SOFM is implemented through a neural-like network architecture as shown in Fig. 1 and it is believed to be similar in some ways to the biological neural network.

The visual display produced by A_{SOFM}^D helps to form hypotheses about topological structure present in X . Although, in this article we concentrate on $(m \times n)$ displays in R^2 , in principle X can be transformed onto a display lattice in R^q for any q . In practice, visual display can be made only for $q \leq 3$ and are usually made on a linear or planar configuration arranged as a rectangular or hexagonal lattice.

As shown in Fig. 1 input vectors $\mathbf{x} \in R^p$ are distributed by a fan-out layer to each of the $(m \times n)$ output nodes in the competitive layer. Each node in this layer has a weight vector (prototype) \mathbf{w}_{ij} attached to it. Let $O_p = \{\mathbf{w}_{ij}\} \subset R^p$ denote the set of $m \times n$ weight vectors. O_p is (logically) connected to a display grid $O_2 \subset V(R^2)$. (i, j) in the index set $\{1, 2, \dots, m\} \times \{1, 2, \dots, n\}$ is the logical address of the cell. There is a one-to-one correspondence between the $m \times n$ p -vectors \mathbf{w}_{ij} and the $m \times n$ cells $\{(i, j)\}$, i.e., $O_p \leftrightarrow O_2$. In the literature display cells are sometimes called nodes, or even neurons, and we shall use both of them.

The feature mapping algorithm starts with (usually) a random initialization of the weight vectors \mathbf{w}_{ij} . For notational clarity we suppress the double subscripts. Now let $\mathbf{x} \in R^p$ enter the network and let t denote the

current iteration number. Find $\mathbf{w}_{r, t-1}$, that best matches \mathbf{x} in the sense of minimum Euclidian distance in R^p . This vector has a (logical) "image" which is the cell in O_2 with subscript r . Next, a topological (spatial) neighborhood $N_r(t)$ centered at r is defined in O_2 , and its display cell neighbors are located. A 3×3 window, $N(r)$, centered at r corresponds to updating nine prototypes in R^p . Finally, $\mathbf{w}_{r, t-1}$ and the other weight vectors associated with cells in the spatial neighborhood $N_r(r)$ are updated using the rule

$$\mathbf{w}_{i, t} = \mathbf{w}_{i, t-1} + h_r(t)(\mathbf{x} - \mathbf{w}_{i, t-1}). \quad (1)$$

Here r is the index of the "winner" prototype

$$r = \underset{i}{\operatorname{arg\,min}} \{ \|\mathbf{x} - \mathbf{w}_{i, t-1}\| \} \quad (2)$$

and $\|\cdot\|$ is the Euclidian norm on R^p . The function $h_r(t)$ which expresses the strength of interaction between cells r and i in O_2 usually decreases with t , and for a fixed t it decreases as the distance (in O_2) from cell r to cell i increases. $h_r(t)$ is usually expressed as the product of a learning parameter α_i and a lateral feedback function $g_i(\operatorname{dist}(r, i))$. A common choice for g_i is $g_i(\operatorname{dist}(r, i)) = \exp^{-\operatorname{dist}(r, i)/\sigma_i}$. α_i and σ_i both decrease with time t . The topological neighborhood $N_r(r)$ also decreases with time. This scheme, when repeated long enough, usually preserves spatial order in the sense that weight vectors which are metrically close in R^p generally have, at termination of the learning procedure, visually close images in the viewing plane. We next provide a schematic description of the algorithm.

Algorithm A_{SOFM}^d (Kohonen):

Begin

Input X /** unlabeled data set $X = \{\mathbf{x}_i \in R^p : i = 1, 2, \dots, N\}$ **/

Input m, n /** the display grid size, a rectangle of size $m \times n$ is assumed **/

Input maxstep /** maximum number of updating steps **/

Input N_0 /** initial neighborhood size **/

Input α_0 /** the initial step size (learning coefficient) **/

Input σ_0 and σ_f /** parameters to control effective step size **/

/** Learning phase **/

Randomly generate initial weight vectors

$\{\mathbf{w}_{ij}, i = 1, 2, \dots, m; j = 1, 2, \dots, n, \mathbf{w}_{ij} \in R^p\}$

$t \leftarrow 0$

While($t < \text{maxstep}$)

Select randomly $\mathbf{x}(t)$ from X ;

find $r = \underset{i}{\operatorname{arg\,min}} \{ \|\mathbf{x}(t) - \mathbf{w}_i(t)\| \}$

/** r and i stands for two-dimensional indices that uniquely identify a weight vector in O_p **/

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 $w_i(t+1) \leftarrow w_i(t) + \alpha_i g_i(\text{dist}(r, i)) [\mathbf{x}(t) - w_i(t)] \quad \forall i \in N_i(r)$ 
 $w_i(t+1) \leftarrow w_i(t) \quad \forall i \notin N_i(r)$ 
  /** dist( $i, j$ ) is the Euclidian distance between the
  centers of nodes  $r$  and  $i$  on the display lattice,  $g_i(d)$ 
  is the lateral feedback function, usually
   $g_i(d) = e^{-d/\sigma_i}$  **/
   $t \leftarrow t + 1$ 
   $\alpha_i \leftarrow \alpha_0(1 - t/\text{maxstep})$ 
   $N_i \leftarrow N_0 - t(N_0 - 1)/\text{maxstep}$ 
   $\sigma_i \leftarrow \sigma_0 - t(\sigma_0 - \sigma_f)/\text{maxstep}$ 
  /** there are other ways to readjust  $\alpha_i, N_i$  and  $\sigma_i$ ,
  and many choices
  for  $g_i$  **/
End While
/** Display phase **/
For each  $\mathbf{x} \in X$  find
   $r = \underset{i}{\text{arg min}} \{ \|\mathbf{x} - w_i\| \}$ , and mark the associated
  cell  $r$  in  $O_2$ .
End.
```

3. Labeling of SOFM prototypes

In this investigation we use a 1-D SOFM, but the algorithm can be extended to 2-D SOFM also. First we train a one-dimensional SOFM using the training data, of course, without using the class information of the input data. Initially the number of nodes in the SOFM is the same as the number of classes c . This is motivated by the fact that the smallest number of prototypes that may be required is equal to the number of classes. At the end of the training the weight vector distribution of the SOFM will reflect the distribution of the input data. These unlabeled prototypes are then labeled using class information. For each of N input feature vectors we identify the prototype closest to it, i.e., the winner node. Since no class information is used during the training, it is only natural that some prototypes may become the winner for data from more than one classes. For each prototype \mathbf{v}_i we compute a score D_{ij} , which is the number of data points from class j to which \mathbf{v}_i is the closest prototype. Due to strong interaction among the neighboring nodes of the SOFM during the training some prototypes may be so placed that for no input data they are the closest prototypes; i.e., D_{ij} is 0 for all j . Naturally we reject such prototypes. For the remaining prototypes the class label C_i of the prototype \mathbf{v}_i is determined as

$$C_i = \underset{j}{\text{arg max}} D_{ij}. \quad (3)$$

The scheme will assign a label to each of the c prototypes, but such a set of prototypes may not classify the data satisfactorily. For example, from (3) it is clear that $\sum_{j \neq C_i} D_{ij}$ data points will be wrongly classified by the

prototype \mathbf{v}_i . Hence we need further refinement of the initial set of prototypes $V_0 = \{\mathbf{v}_{10}, \mathbf{v}_{20}, \dots, \mathbf{v}_{c0}\} \subset R^p$, which we do next.

3.1. Refinement of prototypes

The prototypes generated by the SOFM algorithm represent the overall distribution of the input data. A set of prototypes useful for classification job must be capable of dealing with class specific characteristics (such as class boundaries) of the data. We present a strategy of modifying the initial set of prototypes V_0 leading to the enhancement of performance of the classifier. This process of modification is repeated till the number of prototypes and their performance stabilize within an acceptable level.

On m th iteration the prototype set V_{m-1} from previous iteration is used to generate the new set of prototypes V_m . The labeled prototypes V_{m-1} are used to classify a set of training data and their performance is monitored. Let W_i be the number of training data to which prototype \mathbf{v}_i is the closest one. Let $S_i = \max_j \{D_{ij}\} = D_{iC_i}$. Thus when \mathbf{v}_i is labeled as a prototype for class C_i , S_i training data points will be correctly classified by \mathbf{v}_i and $F_i = \sum_{j \neq C_i} D_{ij}$ data points will be incorrectly classified. Thus,

$$W_i = S_i + F_i$$

and

$$W_i = \sum_j D_{ij}.$$

Let $X = \{\mathbf{x}_1, \dots, \mathbf{x}_N\}$ be the set of training data and N_j be the number of training data from class j . The refinement stage uses $(c+1)$ parameters, a global retention parameter α and a set of class-wise retention parameters β_k (one for each class), to evaluate the performance of each prototype. α and β_k are computed dynamically (not fixed) for m th iteration using the following formula:

$$\alpha_m = \frac{1}{K_1 |V_{m-1}|},$$

$$\beta_{mk} = \frac{1}{K_2 |V_{mk}^*|},$$

where $V_{mk}^* = \{\mathbf{v}_i | \mathbf{v}_i \in V_{m-1}, C_i = k\}$, K_1 and K_2 are two user supplied constants whose choice is discussed in detail later. Thus, actually the algorithm uses 2 (not $c+1$) user supplied parameters.

Based on the result of the evaluation, different operations may be performed on the prototypes to generate a new set of prototypes. We use the following operations in our algorithm.

Merging of a prototype with respect to a class: Let a prototype \mathbf{v}_i represent D_{ik} training data from class k . To

merge v_i w.r.t. class k we identify the prototype v_i closest to v_i where $C_i = k$ (i.e., v_i also is a prototype for class k). Let us denote X_{ij} as the set of training data vectors from class j whose nearest prototype is v_i . When we merge v_i with v_i w.r.t. class k , v_i is updated according to the equation,

$$v_i = \frac{W_i v_i + \sum_{x \in X_k} x}{W_i + D_{ik}} \quad (4)$$

Note that we do not say here, when to merge. This will be discussed later.

Modifying a labeled prototype: A prototype v_i is modified according to the following equation:

$$v_i = \frac{\sum_{x \in X_{C_i}} x}{D_{iC_i}} \quad (5)$$

Splitting a prototype: A prototype v_i is split into r new prototypes for r different classes according to the following rule. For each of r new prototypes v_i of class C_i we compute

$$v_i = \frac{\sum_{x \in X_{C_i}} x}{D_{iC_i}} \quad (6)$$

The prototype v_i is deleted. So after the splitting the number of prototypes is increased by $r - 1$.

Deleting a prototype: The prototype v_i is deleted so that the number of prototypes is reduced by one.

Now we are in a position to schematize the evaluation and enhancement strategy for the prototypes as follows.

Repeat for all $v_i \in V_{m-1}$ until termination condition is satisfied.

If $W_i \neq D_{iC_i}$ and $W_i < \alpha N$ and there is at least another prototype for class C_i

then delete v_i . (Global deletion)

/ If a prototype is not a pure one (i.e., it represents data from more than one classes) and does not represent a reasonable number of points, it fails to qualify to become a prototype. However if there is no other prototype for class C_i the prototype is retained */*

Else if $W_i > \alpha N$ but $D_{ij} < \beta_{mj} N_j$ for all classes then merge v_i for the classes for which $D_{ij} > 0$ and delete v_i .

(Merge and delete)

/ The prototype represents a reasonable number of points, but not a reasonable number of points from any particular class so that it can qualify as a prototype for a particular class. But we cannot ignore the prototype completely. We logically first split v_i into s prototypes $v_{i1}, v_{i2}, \dots, v_{is}$, $s \leq c$, c is the total number of classes for which $D_{ij} > 0$, and then merge v_{ij} to its closest prototype from class j . v_i is then deleted. */*

Else if $W_i > \alpha N$ and $D_{iC_i} > \beta_{mC_i} N_{C_i}$ but $D_{ij} < \beta_{mj} N_j$ for all $j \neq C_i$

then merge v_i with respect to all the classes other than C_i for which $D_{ij} > 0$ using (4) and modify v_i using (5). (Merge and modify)

/ The prototype represents points from more than one classes, however, the points from one class only are well represented by the prototype. According to our labeling scheme the prototype is labeled with the most represented class. Thus we merge v_i with respect to the classes other than C_i using (4) and then modify v_i by (5). */*

Else if $W_i > \alpha N$ and $D_{ij} > \beta_{mj} N_j$ for more than one class then merge v_i w.r.t. classes for which $D_{ij} < \beta_{mj} N_j$ by (4) and split v_i into new prototypes for the classes for which $D_{ij} > \beta_{mj} N_j$ by (5). Add these new prototypes to the new set of prototypes V_m .

(Merge and split)

/ The prototype represents points reasonably well from more than one classes. So we merge the prototype with respect to the classes whose data are not represented reasonably well and split the prototype into one for each class whose data are reasonably well represented by v_i . */*

Let V_m be the union of the unaltered prototypes of V_{m-1} and the modified as well as the new prototypes.

Run the SOFM algorithm on V_m with winner-only update (i.e., no neighbor is updated) strategy using the same training data as input.

/ At this stage we want only to fine tune the prototypes. If the neighbors also are updated the prototypes again might migrate to represent points from more than one class. */*

Termination conditions

The algorithm may terminate under any one of the following three conditions.

- (i) Satisfactory recognition score defined in terms of percentage of correct classifications (ϵ).
- (ii) Stability of prototypes.
- (iii) A maximum number of iterations (I_{max}) reached.

Proper use of condition (i) requires some knowledge of the data. However, even if we do not have the same, we can always set a high (conservative) percentage for (ϵ), say 95%.

Condition (ii) can be checked by a parameter δ_p using the following condition:

$$\frac{||V_{m-1}| - |V_m||}{|V_{m-1}|} < \delta_p \quad (7)$$

where $|V_{m-1}|$ is the number of prototypes in V_{m-1} .

Thus the algorithm terminates when between two successive iterations the number of prototypes do not change significantly.

Condition (iii) is used to protect against infinite looping of the algorithm for some data with highly overlapped structures for which the chosen values of ϵ and δ_p may not be reachable.

4. Results

Several data sets have been used to judge the performance of the algorithm. But we report here results for five data sets: *Iris*, *Glass*, *Breast Cancer*, *Vowel* and *Norm4*. *Iris* data [10] have 150 points in four dimensions that are from three classes each with 50 points. *Glass* data [11] consist of 214 samples with nine attributes from six classes. *Breast cancer* [12] data have 569 points in 30 dimensions from two classes. The vowel [15] data set consists of 871 samples of discrete phonetically balanced speech samples for the Telugu vowels in consonant–vowel nucleus–consonant (CNC) form. These samples are generated from three male informants (in the age group of 25–30 yr) on an AKAI-type recorder. The spectrographic analysis was done on a Kay Sonograph Model 7029-A. The data have three features as the first three formant frequencies. The data set *Norm4* [13] is a sample of 800 points consisting of 200 points each from the four components of a mixture of 4 class 4-variate normals. All our reported results are obtained on the entire data sets.

Table 1 summarizes the classification performances. We used the values $K_1 = 3$, $K_2 = 6$, $\delta_p = 0.2$, $\epsilon = 95\%$ and $J_{\max} = 10$.

It is well known that classes 2 and 3 of *Iris* have some overlap and the typical re-substitution error with a nearest-prototype classifier defined by three prototypes obtained by some clustering algorithm is 15–16 (i.e., about 10% error with three prototypes). Our algorithm terminated with seven prototypes in three iterations. The performance of the proposed system with seven prototypes is quite good resulting only in 2.66% error.

Breast cancer data have been used in Ref. [12] to train a linear programming-based diagnostic system by a

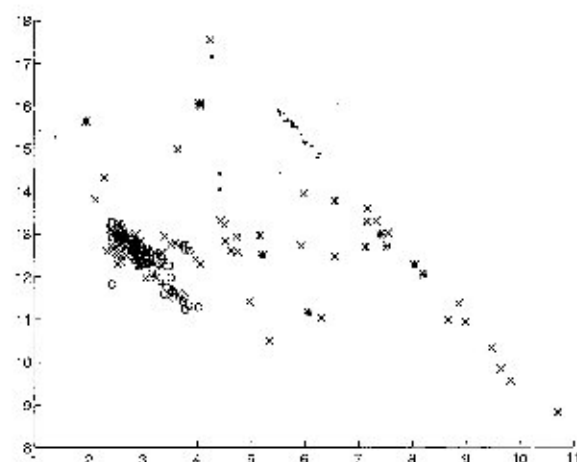


Fig. 2. Scatterplot of the glass data along two most significant principal components.

variant of multisurface method (MSM) called MSM-tree and about 97.5% accuracy was obtained. *Breast cancer* data of a similar kind have also been used in a recent study [14] with 74.0% accuracy with 100 rules. Our classifier could achieve as low as 11.07% error with only five prototypes and it is quite good.

Glass data shows a high percentage of error; this is possibly unavoidable, because a scatterplot (Fig. 2) of the two principal components shows that the data for class 3 are almost randomly distributed among the data points from other classes. In fact the points from class 3 (represented by +) are not visible in the scatterplot. In Ref. [14] the recognition score for the glass data is 64.4%, i.e., about 35% error. Our classifier could realize more than 78% accuracy with 30 prototypes generated in seven iterations of the algorithm.

Although the vowel data set has three features, we used only the first two features. Bayes classifier for this data set [16] gives an overall recognition score of 79.2%. Fig. 3, the scatterplot of vowel data depicts that there are substantial overlap among different classes and hence some misclassification is unavoidable. The proposed classifier could achieve nearly 79% correct classification with 15 prototypes.

The performance on *Norm4* [13] with only four prototypes, i.e., one prototype/class is excellent too. In this case the SOFM based classifier could achieve up to 96% accuracy with only four prototypes.

5. Conclusions

We have proposed a simple but powerful approach of finding a set of reliable prototypes for designing

Table 1
Performance of the classifier for different data sets

Data set	Size	No. of prototypes		No. of iterations	% of error
		Initial	Final		
<i>Iris</i>	150	3	7	4	2.66%
<i>Glass</i>	214	6	30	7	21.29%
<i>Breast cancer</i>	569	2	5	6	11.07%
<i>Vowel</i>	871	6	15	5	21.01%
<i>Norm4</i>	800	4	4	1	3.75%

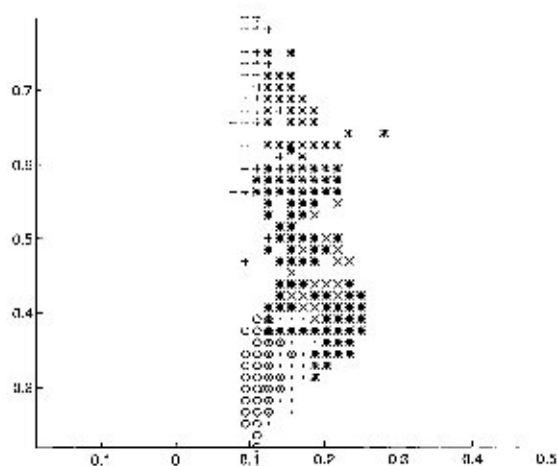


Fig. 3. Scatterplot of the vowel data.

nearest-prototype classifiers. The algorithm first finds a set of representative prototypes from the training data using SOFM disregarding the class information. These prototypes are labeled following a “most-likely class” heuristic. Subsequent stages of tuning cycles fine-tune the prototype set to realize a better class discrimination. Depending on the performance of the classifier prototypes are deleted, merged, modified or split. The retention parameters try to strike a compromise between error rate and the number of the prototypes. Global retention parameter $\alpha(K_1)$ prevents uncontrolled increase in the number of prototypes while the class-wise retention parameters $\beta_k(K_2)$ try to generate prototypes as pure as possible resulting in an increase in the number of prototypes. Proper choice of K_1 and K_2 is needed for balancing the opposing tendencies generated by α and β_k s. It is found that for data with well separated classes (such as Norm4 used here) the process is comparatively insensitive to the change of values of K_1 and K_2 , but for data with highly overlapped classes (like Glass and Vowel) the performance of the system varies considerably with the change of value of K_1 and K_2 , especially K_2 . The process, to some extent, depends on δ_p also. Further investigation is required to provide a guideline for selection of these parameters. Since, the proposed scheme can generate a small number of good prototypes for 1-NP classifier, they can be used to extract fuzzy rules for classifier design also. This is currently under investigation.

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