



Unsupervised segmentation using a self-organizing map and a noise model estimation in sonar imagery

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Abstract

This work deals with unsupervised sonar image segmentation. We present a new estimation and segmentation procedure on images provided by a high-resolution sonar. The sonar image is segmented into two kinds of regions: *shadow* (corresponding to a lack of acoustic reverberation behind each object lying on the seabed) and *reverberation* (due to the reflection of acoustic wave on the seabed and on the objects). The unsupervised contextual method we propose is defined as a two-step process. Firstly, the iterative conditional estimation is used for the estimation step in order to estimate the noise model parameters and to accurately obtain the proportion of each class in the maximum likelihood sense. Then, the learning of a Kohonen self-organizing map (SOM) is performed directly on the input image to approximate the discriminating functions, i.e. the contextual distribution function of the grey levels. Secondly, the previously estimated proportion, the contextual information and the Kohonen SOM, after learning, are then used in the *segmentation step* in order to classify each pixel on the input image. This technique has been successfully applied to real sonar images, and is compatible with an automatic processing of massive amounts of data. © 2000 Pattern Recognition Society. Published by Elsevier Science Ltd. All rights reserved.

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1. Introduction

In high-resolution sonar imagery, three kinds of regions can be visualized: *echo*, *shadow* and *sea-bottom reverberation*. The *echo* information is caused by the reflection of the acoustic wave from the object while the *shadow* zone corresponds to a lack of acoustic reverberation behind this object. The remaining information is called the *sea-bottom* reverberation area. On the pictures provided by a classification sonar, the echo features are generally less discriminant than the shadow shape for the classification sonar of object lying on the seafloor. For this reason, detection and classification of an object

located on the seafloor (as wrecks, rocks, man-made objects, and so on...) are generally based on the extraction and the identification of its associated cast shadow [1]. Thus, before any classification step, one must segment the sonar image between *shadow* areas and *reverberation* areas. In fact, the *sea-bottom reverberation* and the *echo* are considered as a single class.

Unfortunately, sonar images contain speckle noise [2] which affects any simple segmentation scheme such as a maximum likelihood (ML) segmentation. In this simple case, each pixel is classified only from its associated grey level intensity.

In order to face speckle noise and to obtain an accurate segmentation map, a solution consists in taking into account the contextual information, i.e. class of the neighborhood pixels. This can be done using Markov random field (MRF) models [3] and this is why Markovian assumption has been proposed in sonar imagery [4]. In this global bayesian method, pixels are classified

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using the whole information contained in the observed image simultaneously. Nevertheless, simple spatial MRF model have a limited ability to describe properties on large scale, and may not sufficient to ensure the regularization process of the set of labels when the sonar image contains high speckle noise. Such a model can be improved by using a large spatial neighborhood for each pixel [5], or a causal scale and spatial neighborhood [6] but this rapidly increases the complexity of segmentation algorithms and the parameter estimation procedure required to make this segmentation *unsupervised*. Besides, the segmentation and the estimation procedure with such a *priori* model requires a lot of computing time. Moreover, the uses of such a global method does not allow to take into account the noise correlation on the sonar image [7].

An alternate approach adopted here, uses a local method, i.e. takes into account the grey levels of the neighborhood pixels. In this scheme, each pixel is classified from information contained in its neighborhood. This method allowing to take into account the noise correlation is divided in two main steps: the model parameter estimation [8] and the segmentation algorithm which is fed with the previously estimated parameters.

In this paper, we adopt for the parameter estimation step an iterative method called iterative conditional estimation (ICE) [9] in order to estimate, in the ML sense, the noise model parameters and specially the proportion of each class (shadow and reverberation). Followed by the training of a competitive neural network as a Kohonen self-organizing map (SOM) [10] in order to approximate the discriminating function (i.e. the contextual distribution function of the grey level). For the segmentation step, we develop a contextual segmentation algorithm exploiting efficiently the previously estimated parameters, the input sonar image, and the topology of the resulting Kohonen SOM.

This paper is organized as follows. In Section 2, we detail the parameter estimation step based on the ICE procedure in Section 2.1, and the training step of the SOM in Section 2.2. Section 3 presents the segmentation step. Experimental results both on real scenes and synthetic sonar images are presented in subsection 3.3, where we compare the results obtained with the proposed scheme, a ML segmentation and a classical monoscale Markovian segmentation. Then a conclusion is drawn in Section 4.

2. Estimation step

2.1. Iterative conditional estimation

2.1.1. Introduction

We consider a couple of random fields $Z = (X, Y)$ with $Y = \{Y_s, s \in S\}$ the field of observations located on a lat-

tice S of N sites s , and $X = \{X_s, s \in S\}$ the label field. Each Y_s takes its value in $\Lambda_{obs} = \{0, \dots, 255\}$ and each X_s in $\{e_0 = \text{shadow}, e_1 = \text{reverberation}\}$. The distribution of (X, Y) is defined firstly by $P_X(x)$, the distribution of X assumed to be stationary and Gibbsian (i.e. Markovian) in this estimation step, and secondly by the site-wise likelihoods $P_{Y_s/X_s}(y_s/x_s)$. In this work, these likelihoods depend on the class label x_s . The observation Y is called the *incomplete data* whereas Z stands for the *complete data*.

In this step, we estimate the parameter vector Φ_y which defines $P_{Y/X}(y/x)$ by using the iterative method of estimation called iterative conditional estimation (ICE) [9]. This method requires to find an estimator, namely $\Phi_y(X, Y)$ for completely observed data. When X is unobservable, the iterative ICE procedure defines $\Phi_y^{[k+1]}$ as conditional expectations of $\hat{\Phi}_y$ given $Y = y$, computed according to the current value $\Phi_y^{[k]}$. This is the best approximation of Φ_y in terms of the mean square error [9]. By denoting E_k , the conditional expectation using $\Phi_y^{[k]}$, this iterative procedure is defined as follows:

- Initialize the noise model parameters to $\Phi_y^{[0]}$.
- $\Phi_y^{[k+1]}$ is computed from $\Phi_y^{[k]}$ and $Y = y$ by

$$\Phi_y^{[k+1]} = E_k[\hat{\Phi}_y/Y = y]. \quad (1)$$

The computation of this expectation is impossible in practice, but we can approach Eq. (1) owing to the law of large numbers by

$$\Phi_y^{[k+1]} = \frac{1}{n} [\hat{\Phi}_y(x_{(1)}, y) + \dots + \hat{\Phi}_y(x_{(n)}, y)], \quad (2)$$

where $x_{(i)}$, with $i = 1, \dots, n$ are realizations of X according to the posterior distribution $P_{X/Y, \Phi_y(x/y, \Phi_y^{[k]})}$. Finally, we can use the ICE procedure for our application because we get:

- An estimator $\hat{\Phi}_y(X, Y)$ of the complete data: we use a maximum likelihood (ML) estimator for the noise model parameter estimation (see Section 2.1.2.)
- An initial value $\Phi_y^{[0]}$ not too far from the optimal parameters (see Section 2.1.3).
- A way of simulating realizations of X according to the *posterior* distribution $P_{Y/X}(y/x)$ by using the Gibbs sampler [11]. For the *prior* model, we adopt an 8-connexity spatial neighborhood (see Fig. 1) in which $\beta_1, \beta_2, \beta_3, \beta_4$ represent the a priori potential associated to the horizontal, vertical, right and left diagonal binary cliques, respectively. In our application, we want to favour homogeneous regions. Then, we define potential functions associated to the two-site cliques of the form

$$\beta_{s,t} = \mu[1 - \delta(x_s, x_t)], \quad (3)$$

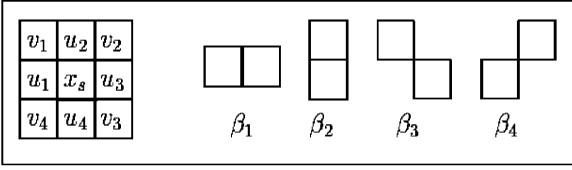


Fig. 1. Second-order neighborhood and two-site associated cliques.

where $\beta_{s,t} = \beta_1, \beta_2, \beta_3$ or β_4 according to the type of neighboring pair $\langle s, t \rangle$, μ is a predetermined positive constant and $\delta(\cdot)$ is the Kronecker function.

2.1.2. Estimation of the noise model parameters for the complete data

The Gaussian law $\mathcal{N}(\mu, \sigma^2)$, is an appropriate degradation model to describe the luminance y within *shadow* regions (essentially due to electronic noise) [12]. The most natural choice of the estimator $\hat{\Phi}_y(x = e_0, y)$ is the empirical mean and the empirical variance. If N_0 pixels are located in the shadow areas, we have

$$\hat{\mu}_{ML} = \frac{1}{N_0} \sum_{s \in S: x_s = e_0} y_s, \quad (4)$$

$$\sigma_{ML}^2 = \frac{1}{N_0 - 1} \sum_{s \in S: x_s = e_0} (y_s - \hat{\mu}_{ML})^2. \quad (5)$$

In order to take into account the speckle noise phenomenon [2] in the reverberation areas, we model the conditional density function of the reverberation class by a shifted Rayleigh law $\mathcal{R}(\min, \alpha^2)$ [12]

$$P(y_s/x_s = e_1) = \frac{y_s - \min}{\alpha^2} \exp \left[-\frac{(y_s - \min)^2}{\alpha^2} \right]. \quad (6)$$

The maximum value of the log-likelihood function is used to determine a ML estimator of the complete data. If \hat{y}_{\min} is the minimum grey level in the reverberation areas and N_1 the number of pixels located within this region, we obtain for $\hat{\Phi}_y(x = e_1, y)$ the following results [12]:

$$\sigma_{ML}^2 = \frac{1}{2N_1} \sum_{s \in S: x_s = e_1} (y_s - \hat{\min}_{ML})^2, \quad (7)$$

$$\hat{\min}_{ML} \simeq \hat{y}_{\min} - 1. \quad (8)$$

In the two cases, the proportion π_k of the k th class is given by the empirical frequency

$$\hat{\pi}_k = \frac{N_k}{N_0 + N_1} \quad \text{with } k \in \{0, 1\}. \quad (9)$$

2.1.3. Initialisation

The initial parameter values have a significant impact on the rapidity of the convergence of the ICE procedure

and the quality of the final estimates. In our application, we use the following method: the initial parameters of the noise model $\Phi_y^{[0]}$ are determined by applying a small non-overlapping sliding window over the image and calculating the sample mean, variance and minimum grey-level estimates. Each estimation calculated over the sliding window gives a “sample” \mathbf{x}_i , a three component vector. These samples $\{\mathbf{x}_1, \dots, \mathbf{x}_M\}$ are then clustered into two classes $\{e_0, e_1\}$ using the *K-means clustering procedure* [13]. This algorithm uses a similarity measure based on the Euclidean distance between samples. A criterion is based on the minimization of the related cost function defined by

$$J = \sum_{i=1}^K \sum_{\mathbf{x}_l \in C_i} |\mathbf{x}_l - c_i|^2, \quad (10)$$

where the second sum is over all samples in the i th cluster and c_i is the *center* of this cluster. It is easy to show that for a given set of samples and class assignments, J is minimized by choosing c_i to be the *sample mean* of the i th cluster. Moreover, when c_i is a sample mean, J is minimized by assigning \mathbf{x}_j to the class of the cluster with the nearest mean. A number of other criteria are given in Ref. [13]. The complete algorithm is outlined below:

- (1) Choose K initial clusters $c_1^{[1]}, \dots, c_K^{[1]}$. These could be arbitrarily chosen, but are usually defined by

$$c_i^{[1]} = \mathbf{x}_i \quad (1 \leq i \leq K). \quad (11)$$

- (2) At the k th step, assign the sample \mathbf{x}_l ($1 \leq l \leq M$) to cluster i if

$$\|\mathbf{x}_l - c_i^{[k]}\| < \|\mathbf{x}_l - c_j^{[k]}\| \quad (\forall j \neq i). \quad (12)$$

In fact, we reassign every sample to the cluster with the nearest mean. In the case of equality, we assign \mathbf{x}_l arbitrary to i or j .

- (3) Let $c_i^{[k]}$ denote the i th cluster after Step 2. Determine new clusters by

$$c_i^{[k+1]} = \frac{1}{N_i} \sum_{\mathbf{x}_l \in c_i^{[k]}} \mathbf{x}_l, \quad (13)$$

where N_i represents the number of samples in $c_i^{[k]}$. Thus, the new cluster position is the mean of the samples in the previous one.

- (4) Repeat until convergence is achieved, say $c_i^{[k+1]} = c_i^{[k]}, \quad \forall i$.

Although it is possible to find pathological cases where convergence never occurs [13], the algorithm does converge in all tested examples. The rapidity of convergence depends on the number K , the choice of initial cluster centers and the order in which the samples are considered. In our application $K = 2$. Fig. 2(a) represents a sonar image and the result of the *K-means* clustering algorithm is reported in Fig. 2(b).

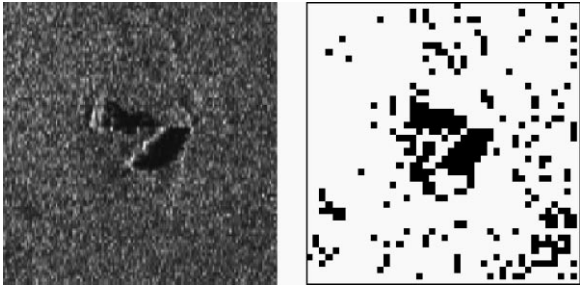


Fig. 2. K-means clustering procedure: (a) sonar picture involving an object and a rock shadow (b) segmentation according to the ML criterion with parameter estimation given by the K-means algorithm.

On the one hand, a small size window increases the accuracy of the segmentation and then the precision of the distribution mixture estimation. On the other hand, it decreases the number of pixels with which \mathbf{x}_i 's are computed and may increase the misclassification error. In our application, good results are obtained with a 6×6 pixels window. The ML estimation is then used over the K-means segmentation in order to find $\Phi_y^{[0]}$.

2.1.4. Parameter estimation procedure for the incomplete data

We can use the following algorithm to estimate the noise model parameters. Let us recall that this method takes into account the diversity of the laws in the distribution mixture estimation.

• Parameter initialization:

The K-means algorithm is used. Let us denote by $\Phi_y^{[0]}$ the obtained result.

• ICE procedure. $\Phi_y^{[k+1]}$ is computed from $\Phi_y^{[k]}$ in the following way:

- Using the Gibbs sampler, n realizations $x_{(1)}, \dots, x_{(n)}$ are simulated according to the posterior distribution with parameter vector $\Phi_y^{[k]}$, and with $P_{Y_s/X_s}(y_s/x_s = e_0)$ a Gaussian law for shadow area, $P_{Y_s/X_s}(y_s/x_s = e_1)$ a shifted Rayleigh law for reverberation area.
- For each $x_{(i)}$ with $i = 1, \dots, n$, the parameter vector Φ_y is estimated with the ML estimator on each class:
- $\Phi_y^{[k+1]}$ is obtained from $\Phi_y(x_{(i)}, y)$ with $(1 \leq i \leq n)$ by using Eq. (2).

If the sequence $\Phi_y^{[k]}$ becomes steady, the ICE procedure is ended and one proceeds the segmentation using the estimated parameters. We can use all these estimated parameters in order to get a complete unsupervised Markovian segmentation (see Section 3.2) or only use the proportion of each class in the Kohonen SOM-based unsupervised segmentation described in Section 2.2.

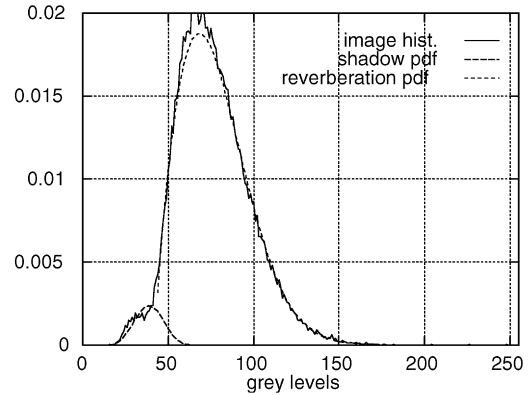


Fig. 3. Image histogram of the picture reported in Fig. 2 with the estimated Gaussian and Rayleigh laws.

Table 1

Estimated parameters on the pictures reported in Fig. 2(a). π stands for the proportion of the two classes within the sonar image. μ and σ^2 are the Gaussian parameters (*shadow area*). Min and α are the Rayleigh law parameters (*reverberation area*). $\Phi_y^{[0]}$ represents the initial parameter estimates and the final estimates are denoted $\hat{\Phi}_y$.

Initialisation of K-means procedure			
$\Phi_{y(\text{shadow})}^{[final]}$	0.04(π)	36(μ)	55(σ^2)
$\Phi_{y(\text{sea-bottom})}^{[final]}$	0.96(π)	39(min)	1061(α^2)
ICE procedure			
$\Phi_{y(\text{shadow})}^{[final]}$	0.03(π)	32(μ)	17(σ^2)
$\Phi_{y(\text{sea-bottom})}^{[final]}$	0.97(π)	39(min)	1591(α^2)

We calibrate the weight of the “stochastic” aspect of the ICE by choosing n . When n increases, the “stochastic” aspect of the algorithm decreases. The choice of a small value for n ($n = 1$ in our application) can increase its efficiency [14].

Fig. 3 represents the mixture of distributions of the sonar image reported in Fig. 2(a). The obtained results are given in Table 1.

The quality of the estimations is difficult to appreciate in absence of real values. We can roughly perform such evaluation by comparing the image histogram with the probability density mixture corresponding to the estimated parameters. Fig. 3(a) shows the resulting mixture solution in graphical form. The two dashed curves in the figures represent the individual components $P_{Y/X}(y/e_m)$ with $0 \leq m \leq K$. The histogram is quite close to the mixture densities based on the estimated parameters, and a segmentation with these estimates can be done as shown in the following section.

2.2. Self-organizing map

2.2.1. Introduction

Researches on neurobiology have shown that centers of diverse activities as thought, speech, vision, hearing, lie in specific areas of the cortex and these areas are ordered to preserve the topological relations between informations while performing a dimensionality reduction of the representation space. Such organization led Kohonen to develop the SOM algorithm [10]. This kind of competitive neural network is composed of one or two dimensional array of processing elements or neurons in the input space. All these neurons receive the same inputs from external world. Learning is accomplished by iterative application of unlabeled input data. As training process, the neurons evolve in the input space in order to approximate the distribution function of the input vectors. After this step, large-dimensional input vectors are, in a sense, projected down on the one or two-dimensional map in a way that maintains the natural order of the input data. This dimensional reduction could allow us to visualize and to use easily, on a one or two-dimensional array, important relationships among the data that might go unnoticed in a high-dimensional space.

The model of SOM used in our application is a one-dimensional array of n nodes. To each neuron N_i , a weight vector $w_i = (w_{i1}, w_{i2}, \dots, w_{ip})^T \in \mathbb{R}^p$ is associated. During learning procedure, an input vector $\mathbf{x} \in \mathbb{R}^p$ randomly selected among vectors of the training set, is connected to all neurons in parallel. The input \mathbf{x} is compared with all the neurons in the Euclidean distance sense *via* variable scalar weight w_{ij} . At the k th step, we assign the vector \mathbf{x} to the winning or leader neuron N_l if:

$$\|\mathbf{x} - \mathbf{w}_l^{[k]}\| = \min_i \|\mathbf{x} - \mathbf{w}_i^{[k]}\|. \quad (14)$$

All the neurons within a certain neighborhood around the *leader* participate in the weight-update process. Considering random initial values for $\mathbf{w}_i^{[0]}$ ($0 \leq i \leq n$), this learning process can be described by the following iterative procedure:

$$\mathbf{w}_i^{[k+1]} = \mathbf{w}_i^{[k]} + \mathcal{H}_{li}^{[k]}(\mathbf{x}^{[k]} - \mathbf{w}_i^{[k]}). \quad (15)$$

The lateral interactions among topographically close elements are modeled by the application of a *neighborhood function* or a smoothing Kernel defined over the winning neuron [10]. This Kernel can be written in terms of the Gaussian function

$$\mathcal{H}_{li}^{[k]} = \alpha^{[k]} \exp\left(-\frac{d^2(l, i)}{2(\sigma^{[k]})^2}\right), \quad (16)$$

where $d(l, i) = \|l - i\|$ is the distance between the node l and i in the array, $\alpha^{[k]}(t)$ is the learning-rate factor and $\sigma^{[k]}$ defines the width of the Kernel at the iteration k . For

the convergence, it is necessary that $H_{li}^{[k]} \rightarrow 0$ when $k \rightarrow T$, where T is the total number of steps of the process [10]. Therefore, for the first step, $\alpha^{[k]}$ should start with a value that is close to unity, thereafter decreasing monotonically [10]. To achieve this task, we use

$$\alpha^{[k]} = \alpha^{[0]}\left(1 - \frac{k}{T}\right). \quad (17)$$

Moreover, as learning proceeds, the size of the neighborhood should be diminished until it encompasses only a single unit. So, we applied for the width of the Kernel the monotonically decreasing function:

$$\sigma^{[k]} = \sigma^{[0]}\left(\frac{\sigma^{[T-1]}}{\sigma^{[0]}}\right)^{k/(T-1)} \quad (18)$$

The ordering of the map occurs during the first steps, while the remaining steps are only needed for the fine adjustment of the weight values.

2.2.2. Iterative learning step

The learning process is performed directly on the real image to be segmented. An input vector is filled with the grey levels of the pixels contained in a 3×3 pixels window sliding over the image (cf. Fig. 4). Therefore, each neuron has nine weights allowing to locate it in the input space. At each step, the location of the window in the image is randomly chosen and the weights are modified according to Eq. (15). Experiments have shown that this training strategy provides as good results as an ordered image scanning process while spending less processing time.

σ has a significant impact on the quality of the convergence. We have to start with a fairly large value to globally order the map. The initial value σ_0 of σ can be half the length of the network. During learning, σ has to decrease monotonically until it reaches a small value. Experiments have shown that $\sigma_{T-1} = 0.1$ is a good choice and provides the minimum quantization error

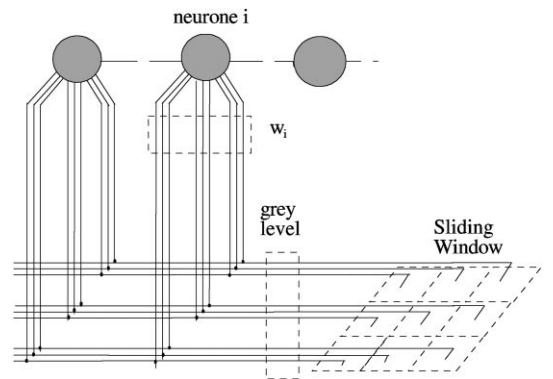


Fig. 4. Model of the SOM used for the segmentation. An 3×3 sliding window is used to feed the SOM.

defined by

$$E_{quant} = \frac{1}{N} \sum_{i=1}^N \|\mathbf{x}_i - \mathbf{w}_l\|, \quad (19)$$

where the summation is over all the N windows of the image and \mathbf{w}_l is the weight vector associated to the leader neuron of the input vector \mathbf{x}_i after learning step.

3. Segmentation step

3.1. SOM segmentation

The classification task consists in running the sliding window over the image. For each location of the window, the corresponding input vector \mathbf{x} is compared with all the neurons using Eq. (14). The *winning neuron*, the one which leads to the smallest distance, gives the class of the pixel located in the center of the window. However, before any classification task, we have to *calibrate* the map in order to associate the label *shadow* or *reverberation* to each neuron.

Assuming that the input vector $\mathbf{x}_0 = (0, \dots, 0)^t$ should represent a window setting on a perfect *shadow* area, it is very useful to define the distance graph representing the Euclidean distance in the nine-dimensional space between the point \mathbf{x}_0 and all the neurons. Such a graph is given in Fig. 5 and Fig. 6 respectively before and after learning for a hundred-neuron network.

Both these figures show that the maximal distance between two successive cells is widely smaller after learning than before. We can deduce that, after learning, neurons that are topologically close in the array are close

in the input space too. As a matter of fact, neurons that are physical neighbors should respond to a similar input vectors. The calibration of the map uses this topological property and the portion π_0 estimated in Section 2.1 of the pixels labelled as *shadow* in the image. This process can be summarized as follows:

- (1) Initially, we affect the class *reverberation* to all neurons.
- (2) We seek the most evident prototype of the shadow class. This neuron is the winning unit according to Eq. (14) when inputting the vectors \mathbf{x}_0 . Then, we affect it to the *shadow* class.
- (3) We affect the *shadow* class to pixels for which the leader neuron belongs to the *shadow* class. We can deduce the intermediate *shadow* portion π_{int} provided by the resulting image.
- (4) If π_{int} is smaller than π_0 , we have to select an additional prototype of a shadow class among neurons of the *reverberation* class. According to the topological preserving properties of the map, this additional neuron should be a direct neighbor of an already *shadow* labelled neuron. Among both the possible neighbors, we take the one which has the smallest Euclidean distance with the point \mathbf{x}_0 . Go to 3.
- (5) If π_{int} is larger than π_0 , we stop the process.

Experiments have shown that E_{quant} is a monotonically decreasing function of the number of steps and reaches an asymptotic value for large value of T . One hundred times the number of network units seems to be a reasonable compromise solution between speed and quality of learning. In our application, 100 neurons have been chosen for the network.

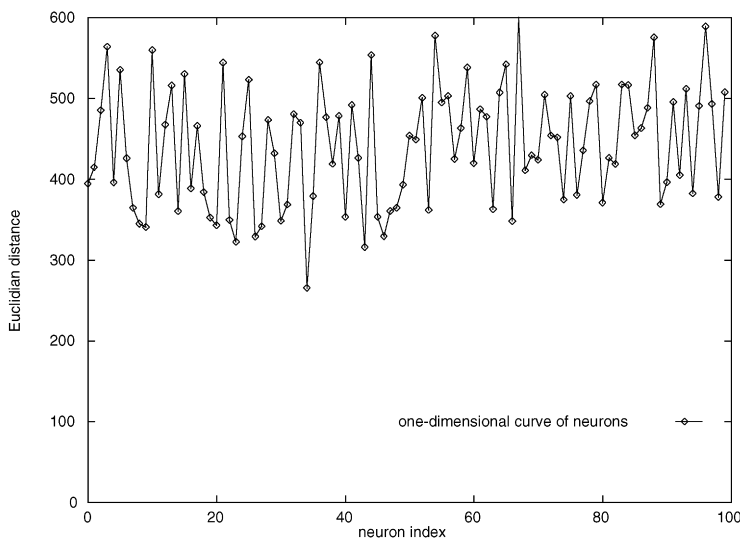


Fig. 5. The distance graph between the 100 neurons of the SOM before learning.

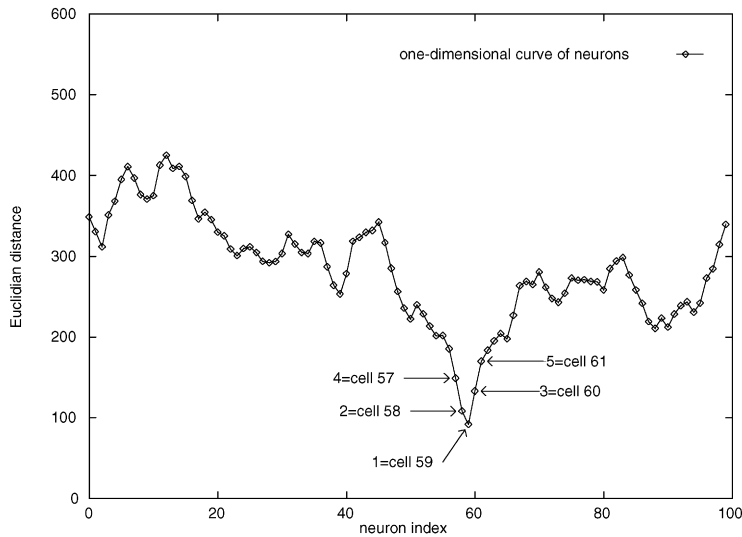


Fig. 6. The distance graph between neurons obtained after learning.

3.2. Markovian segmentation

The segmentation of sonar images in two classes can be viewed as a statistical labelling problem according to a global Bayesian formulation in which the posterior distribution $P_{X/Y}(x/y) \propto \exp [-U(x, y)]$ has to be maximized [3]. In our case, the corresponding *posterior* energy $U(x, y)$ to be minimized is

$$U(x, y) = \underbrace{\sum_{s \in S} \psi_s(x_s, y_s)}_{U_1(x, y)} + \underbrace{\sum_{\langle s, t \rangle} \beta_{s, t} [1 - \delta(x_s, x_t)]}_{U_2(x, y)} \quad (20)$$

where U_1 denotes the adequacy between observations and labels ($\Psi_s(x_s, y_s) = \ln [P_{X_s/Y_s}(x_s/y_s)]$) and U_2 expresses the energy of the priori model. In order to minimize this energy function, we use a deterministic relaxation technique called ICM algorithm [3–5,15].

3.3. Results on real scenes

We compare the segmentation performance of the proposed SOM-based algorithm described in Section 3.1 with a ML segmentation and a classical Markovian segmentation using a deterministic relaxation technique such as the ICM [3]. All the segmentation results exploit the parameter estimation step presented in Section 2.1. This estimation step is used to estimate both the noise model parameters for the ML segmentation, the Mar-

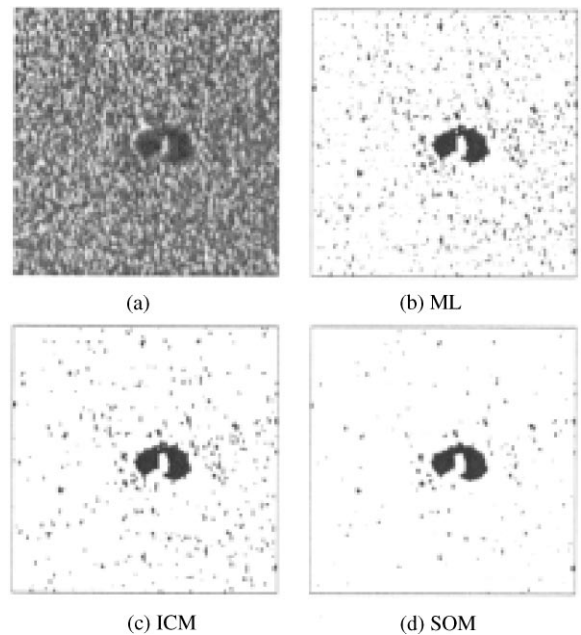


Fig. 7. (a) A real sonar image involving a sandy sea floor with the cast shadow of a tyre. Two-class segmentation results obtained with this image using: (b) ML segmentation, (c) Markovian segmentation with a deterministic relaxation technique as ICM, (d) the SOM based segmentation result (see Table 2 for the estimated parameters). The SOM segmentation exhibits a good robustness against the speckle noise (which induces false small shadow areas compared to the others approaches).

kovian segmentation and the proportion of the *shadow* class for the SOM segmentation.

Figs. 6–9 show the segmentation results obtained with the different methods. Example of the noise model

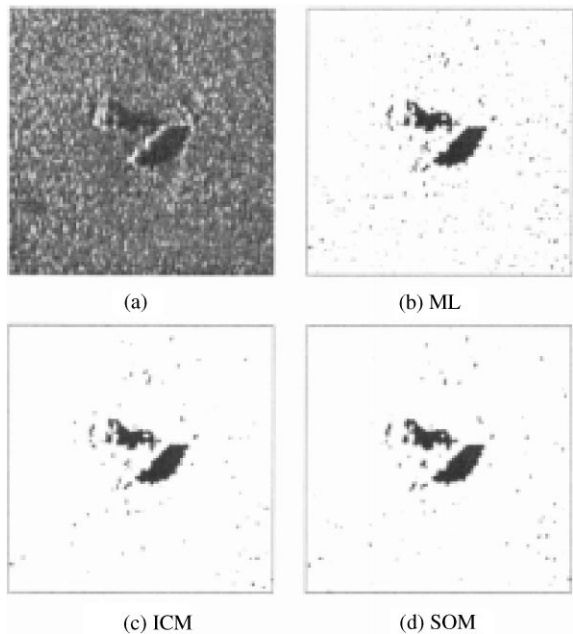


Fig. 8. A real sonar image involving an object and a rock shadows. Two-class segmentation results obtained with (a), (b) ML segmentation, (c) Markovian segmentation with ICM technique, (d) SOM based segmentation method (see Table 3 for the estimated parameters). The ML and ICM do not permit to totally eliminate the speckle noise effect (creating *shadow* mislabelled isolated pixels).

Table 2
Estimated parameters on the picture reported in Fig. 6

ICE procedure			
$\Phi_{y(shadow)}^{[final]}$	0.02 _(π)	36 _(μ)	85 _(σ^2)
$\Phi_{y(sea-bottom)}^{[final]}$	0.98 _(π)	46 _(min)	1878 _(σ^2)

Table 3
Estimated parameters on the picture reported in Fig. 7

ICE procedure			
$\Phi_{y(shadow)}^{[final]}$	0.03 _(π)	25 _(μ)	32 _(σ^2)
$\Phi_{y(sea-bottom)}^{[final]}$	0.97 _(π)	35 _(min)	1430 _(σ^2)

parameters Φ_y obtained with our scheme are given in Tables 2–4.

Experiments indicate that the SOM segmentation requires less computation than the Markovian segmenta-

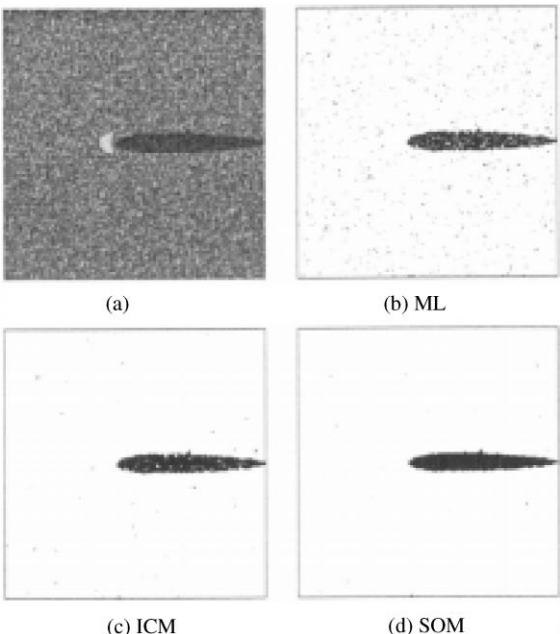


Fig. 9. (a) A synthetic sonar image of a sphere lying on the seabed. Segmentation results obtained with: (b) ML segmentation, (c) Markovian ICM technique, (d) SOM-based segmentation method.

Table 4
Estimated parameters on the picture reported in Fig. 8

ICE procedure			
$\Phi_{y(shadow)}^{[final]}$	0.03 _(π)	34 _(μ)	39 _(σ^2)
$\Phi_{y(reverberation)}^{[final]}$	0.97 _(π)	42 _(min)	1412 _(σ^2)

tion (30'' for the SOM estimation-segmentation, whereas roughly 100'' are required for unsupervised scale causal Markovian modelization [16] on IBM 43P-200 MHz workstation). Besides, the ICM algorithm do not permit to decrease the number of false alarm (wrong detections) due to the speckle noise effect. The SOM segmentation performs better, exhibits a good robustness versus speckle noise (false alarms have been eliminated), and allows us to preserve the shadow shapes of little rocks.

Manufactured objects or rock shadows are better segmented with our method than with the others (cf. Figs. 7–9) and their shape are close to the result we expected. The cast shadow of a manufactured object (a cylinder) reported in Fig. 10 has a geometric shape (contrary to the cast shadow of the rock) that will be discriminant for the classification step.

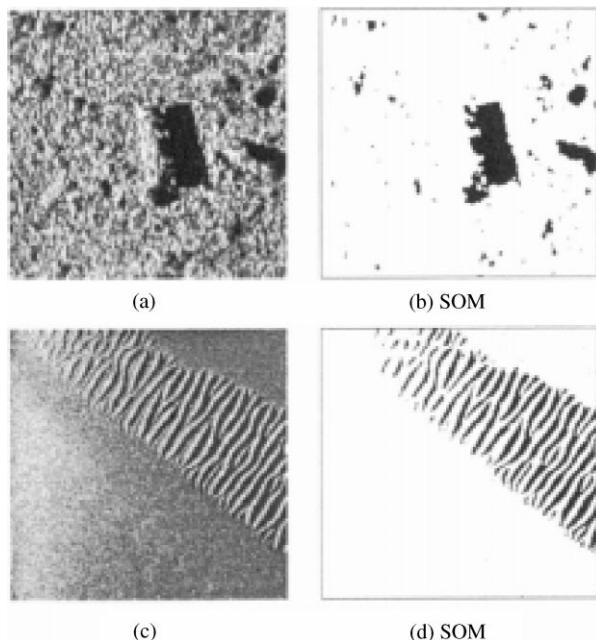


Fig. 10. Real sonar images of a cylindrical object (a) and of ridges of sand (c). Their corresponding SOM-based segmentation results are depicted, respectively, in (b) and (d).

4. Conclusion

We have described an unsupervised segmentation procedure based on a *parameter estimation step* (which offers an appropriate estimation of the noise model) and a segmentation step well adapted for sonar image segmentation problem. The estimation step takes into account the diversity of the laws in the distribution mixture of a sonar image and can be used with the Kohonen SOM-based segmentation in order to solve the difficult problem of unsupervised sonar image segmentation. This scheme is computationally simple and appears as an interesting alternative to existing complex hierarchical Markovian methods. This method has been validated on several real sonar images demonstrating the efficiency and robustness of this scheme.

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