

SPECT IMAGE RECONSTRUCTION USING COMPOUND PRIOR MODELS

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We propose a new iterative method for Maximum a Posteriori (MAP) reconstruction of SPECT (Single Photon Emission Computed Tomography) images. The method uses Compound Gauss Markov Random Fields (CGMRF) as prior model and is stochastic for the line process and deterministic for the reconstruction. Synthetic and real images are used to compare the new method with existing ones.

Keywords: SPECT imaging; compound Gauss Markov random fields; Bayesian reconstruction; simulated annealing; deterministic image reconstruction.

1. Introduction

SPECT images are observation data acquired by a gamma-camera following an orbit around the patient's body, at regularly spaced angles. A reconstructed image is the discrete representation of a slice or cross-section of the isotope distribution within the patient, transversal to the gamma-camera rotation axis.

Bayesian reconstruction methods have been extensively used to reconstruct medical images since they can improve the reconstructions with respect to the classical nonstatistical methods, such as FBP (Filtered Back Projection)¹⁵ and ART (Algebraic Reconstruction Techniques).²⁵

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In the Bayesian paradigm, the reconstructed image \hat{X} is usually selected as

$$\hat{X} = \arg \max_X P(X|Y) = \arg \max_X P(Y|X)P(X), \quad (1)$$

where $P(X)$ is a prior distribution incorporating information about the expected structure in the image X , and $P(Y|X)$ models the degradation process of projections Y from the pixel intensities of the emission source (patient).

When the energy function associated with the prior distribution is convex, pure descent algorithms assure that the global minimum of $-\ln P(X|Y)$ can be found.³ However, when the prior model explicitly favors the presence of discontinuities, the energy of the posterior distribution of the reconstruction given the data usually presents many local minima. In this case stochastic algorithms such as simulated annealing,^{11,23} deterministic methods^{2,5,6} and mixed-annealing algorithms,^{1,22} where stochastic steps alternate with deterministic ones, have been proposed. Unfortunately all these methods assume that the degradation process is Gaussian.

For SPECT images, a problem where the degradation process is Poisson, several reconstruction methods have been proposed that fit within the Bayesian framework (see, for instance Refs. 3, 10, 12, 13, 16, 17, 21, 27, 29 and 30). General strategies to find the MAP have been proposed.^{4,7-9} However, not much work has been reported on the use of Compound Gauss Markov Random Fields (CGMRF) to reconstruct such images (see however Refs. 18, 19 and 24).

In this work we propose a Bayesian technique for SPECT image reconstruction that uses Compound Gauss Markov Random Fields (CGMRF) as prior model and is stochastic for the line process and deterministic for the image process when computing their respective MAP estimates.

The rest of the paper is organized as follows. In Sec. 2 we define the observation and image models. Then, in Sec. 3 we propose a new method for finding the MAP estimate. The application of this method to synthetic and real images is described in Sec. 4. Finally, Sec. 5 concludes the paper.

2. Observation and Image Models

The observation model for emission tomography can be specified as a product of independent Poisson distributions²⁹

$$P(Y|X) = \prod_{s=1}^M \frac{(\sum_{i=1}^N A_{s,i}x_i)^{y_s} \exp\{-\sum_{t=1}^N A_{s,t}x_t\}}{y_s!}, \quad (2)$$

where M is the number of detectors, N is the number of pixels and A is the system matrix, $A_{s,i}$ is the probability that an emitted photon from pixel i reaches detector s .

The prior model we use is a CGMRF model. This model provides us with a means to control changes in the image using a hidden random field. A CGMRF model has two levels, an upper level which is the image to be restored and the

lower or hidden level, called the line process, which is an underlying random field governing the transition between submodels. The use of a line process, was introduced by Geman and Geman¹¹ for the discrete case. Extensions to the continuous case were presented by Jengs and Woods¹⁴ (see also Ref. 18 for medical image reconstruction).

The CGMRF model to be used here is introduced from a simpler one, the Conditional Auto-Regression (CAR) model.²⁶ This prior model is defined by

$$P(X) \propto \exp \left\{ -\frac{1}{2} \alpha X^t (I - \phi C) X \right\} \tag{3}$$

where α is a scaling parameter, for an 8-point neighborhood system, the (i, j) th element of matrix C is given by

$$C_{i,j} = \begin{cases} 2(0.5\sqrt{2} + 1)^{-1} & \text{if } d(i, j) = 1 \\ 2(\sqrt{2} + 1)^{-1} & \text{if } d(i, j) = \sqrt{2} \\ 0 & \text{otherwise} \end{cases}$$

$d(i, j)$ is the Euclidean distance between pixels i and j (see Ref. 28) and $|\phi| < 1/8$. Note that $\sum_j C_{i,j} = 8$ and since (see Ref. 26)

$$E[x_i | x_j, j \neq i] = \phi \sum_j C_{i,j} x_j, \tag{4}$$

when ϕ is just less than $1/8$, Eq. (4) means that the expectation of a pixel x_i given the rest of the image is similar to the mean of its eight neighbors weighted according to their distance to the pixel x_i .

If we assume a “toroidal edge correction”, we introduce the line process by rewriting Eq. (3) as

$$\begin{aligned} & -\log P(X, L) \\ & \propto \frac{\alpha\phi}{2} \left(\sum_i C_{i,i:1} (x_i - x_{i:1})^2 (1 - l_{[i,i:1]}) + \sum_i C_{i,i:2} (x_i - x_{i:2})^2 (1 - l_{[i,i:2]}) \right) \\ & \quad + \frac{\alpha\phi}{2} \left(\sum_i C_{i,i:5} (x_i - x_{i:5})^2 (1 - l_{[i,i:5]}) + \sum_i C_{i,i:6} (x_i - x_{i:6})^2 (1 - l_{[i,i:6]}) \right) \\ & \quad + \frac{\alpha}{2} \sum_i \beta [l_{[i,i:1]} + l_{[i,i:2]} + l_{[i,i:5]} + l_{[i,i:6]}] + \frac{\alpha}{2} \sum_i (1 - 8\phi) x_i^2, \end{aligned} \tag{5}$$

where $i : 1, i : 2, \dots, i : 8$ are the eight pixels around pixel i (see Fig. 1). If all the elements $l([i, j])$ take always the value zero we have a CAR model. However, if $l([i, j])$ takes the value zero if pixels i and j are not separated by an active line and one otherwise we have a CGMRF model. The parameter β is a scalar weight, which adjusts the introduction of the active line elements. For β very large the prior model becomes Gaussian. The line process acts as inhibitor or activator of the relation between two neighboring pixels depending on whether or not there exists an edge.

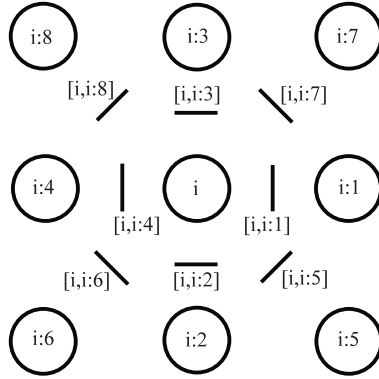


Fig. 1. Image and line sites.

3. MAP Estimation

Let us now proceed to find \hat{X} , \hat{L} , the MAP estimates of X and L , that is

$$\hat{X}, \hat{L} = \arg \max_{X, L} P(X, L|Y). \tag{6}$$

The method we propose for estimating the original image and the line process is stochastic for the line process and deterministic for the reconstruction, as we will describe now.

In order to estimate the line process we simulate the corresponding conditional *a posteriori* density function. Let us denote by $P_T(l_{[i,j]}|L_{[i,j]}, X, Y)$ the conditional *a posteriori* density function for the line process element $l_{[i,j]}$, given X, Y and the rest of L , ($L_{[i,j]} = (l_{[s,t]} : \forall [s,t] \neq [i,j])$). To simulate this density function, we have

$$P_T(l_{[i,j]} = 0|L_{[i,j]}, X, Y) \propto \exp \left[-\frac{1}{T} \frac{\alpha \phi C_{i,j}}{2} (x_i - x_j)^2 \right], \tag{7}$$

$$P_T(l_{[i,j]} = 1|L_{[i,j]}, X, Y) \propto \exp \left[-\frac{1}{T} \frac{\alpha \beta}{2} \right], \tag{8}$$

where T is the temperature.

Given an estimate of the line process, L , and the observation, Y , we note that $-\ln P(X|L, Y)$ is convex in X and so a simple gradient method can be used to minimize it. A different more efficient procedure is followed in estimating the image X , as described next. Starting from the probability distribution $P(X|L, Y)$ we have

$$-\frac{\partial}{\partial x_i} \ln P(X|L, Y) = 1 - \sum_{s=1}^M \frac{y_s A_{s,i}}{\sum_{t=1}^N A_{s,t} x_t},$$

$$+ \alpha \phi \sum_{j \in \mathcal{N}_i} C_{i,j} (x_i - x_j) (1 - l_{[i,j]}) + \alpha (1 - 8 \phi) x_i,$$

and so at the minimum of $-\ln P(X|L, Y)$ we have

$$\begin{aligned}
 & 1 + \alpha x_i \left\{ \phi \sum_{j \in \mathcal{N}_i} C_{i,j} (1 - l_{[i,j]}) + (1 - 8\phi) \right\} \\
 &= \sum_{s=1}^M \frac{y_s A_{s,i}}{\sum_{t=1}^N A_{s,t} x_t} + \alpha \phi \sum_{j \in \mathcal{N}_i} C_{i,j} x_j (1 - l_{[i,j]}).
 \end{aligned}$$

Now, taking into account that $\phi \sum_{j \in \mathcal{N}_i} C_{i,j} + (1 - 8\phi) = 1$ and noting that when $l_{[i,j]} = 1$, $x_i(1 - l_{[i,j]}) = 0$ and $x_j(1 - l_{[i,j]}) = 0$, at the minimum of $-\ln P(X|L, Y)$ we have

$$1 + \alpha x_i = \sum_{s=1}^M \frac{y_s A_{s,i}}{\sum_{t=1}^N A_{s,t} x_t} + \alpha \phi \sum_{j \in \overline{\mathcal{N}}_i} x_j C_{i,j} + \alpha \phi x_i \sum_{j \in \underline{\mathcal{N}}_i} C_{i,j}, \quad (9)$$

where $\overline{\mathcal{N}}_i$ and $\underline{\mathcal{N}}_i$ denote, respectively, the neighboring pixels of i without or with an active line between i and j . We have removed their dependency on the current line process estimate for simplicity.

Then, multiplying both sides of Eq. (9) by $x_i/(1 + \alpha x_i)$ we obtain that the image X maximizing $P(X|L, Y)$ satisfies for all i

$$x_i = \mu_i \left(\phi \sum_{j \in \overline{\mathcal{N}}_i} x_j C_{i,j} + \phi x_i \sum_{j \in \underline{\mathcal{N}}_i} C_{i,j} \right) + (1 - \mu_i) x_i \sum_{s=1}^M \frac{y_s A_{s,i}}{\sum_{t=1}^N A_{s,t} x_t}, \quad (10)$$

with $\mu_i = \alpha x_i / (\alpha x_i + 1)$.

This equation is used to define an iterative process. Using the current estimate of the original image on the right-hand side of Eq. (10) we obtain a new estimate of it.

Furthermore the proposed iterative method is a modification of Newton's method²⁰ since from Eq. (10) we have

$$x_i^{\text{new}} = x_i^{\text{old}} + \frac{x_i^{\text{old}}}{1 + \alpha x_i^{\text{old}}} \frac{\partial}{\partial x_i} \ln P(X|L, Y), \quad (11)$$

and so it can be written as

$$X^k = X^{k-1} - M^{k-1} \nabla [-\ln P(X^{k-1}|L, Y)]^t, \quad (12)$$

where k denotes iteration and M^{k-1} is a diagonal matrix with entries $1/(1 + \alpha x_i^{k-1})$. We have used this iterative scheme in our test examples and observed experimentally that it converges. We also note that the proposed method does not need to calculate either the Hessian matrix or the optimal step of the gradient. Furthermore, if we start the iteration with a non-negative image estimate, the iterative process in Eq. (10) is guaranteed to provide always non-negative images and that when $\alpha = 0$ we obtain the classical E-M algorithm for Poisson distributions.

Let us now summarize and describe how to use Eqs. (7), (8) and (10) to find the MAP estimate of L and X :

Let $k = 1, 2, \dots$, be the sequence of iterations in which the sites (lines or pixels) are visited for updating.

- (1) Set $k = 0$ and assign an initial configuration, X^{-1} , L^{-1} , and an initial temperature $T = 1$.
- (2) The evolution $\hat{L}^{k-1} \rightarrow \hat{L}^k$ of the whole line process is simulated by the probability functions defined in Eqs. (7) and (8).
- (3) The evolution $\hat{X}^{k-1} \rightarrow \hat{X}^k$ of the whole image is obtained using \hat{X}^{k-1} on the right-hand side of Eq. (10) to obtain \hat{X}^k on the left-hand side of the same equation.
- (4) Set $k = k + 1$. Decrease the temperature T according to an annealing scheme.¹⁴ Go to step 2 to fit a stopping criterion.

4. Experimental Results

Let us examine how the proposed method works on both synthetic and real images.

Figure 2 shows the results for the 128×128 synthetic image depicted in Fig. 2(a). The sinogram, shown in Fig. 2(b), was simulated assuming a system with 128 detectors equipped with a parallel hole collimator that described a semicircular orbit with 128 angles. The simulation assumed Poisson noise and no attenuation during the capturing process.

Figure 2(e) shows the reconstruction of this sinogram with the proposed method. The used parameters were chosen experimentally to be $\alpha = 19^{-1}$ and $\beta = 15.4$. The stopping criterion was $\|X^k - X^{k-1}\|^2 / \|X^k\|^2 < 10^{-9}$. The corresponding line process [Fig. 2(f)] shows that the method locates all the regions in the image and the general form of the objects is clearly distinguished.

In order to compare the results, other reconstruction methods using a Conditional Autoregressive (CAR) prior and a Generalized Gauss Markov Random Field (GGMRF)³ prior were applied to the sinogram in Fig. 2(b). The parameters needed for these reconstruction methods were $\alpha = 280^{-1}$ for the CAR prior and, for the GGMRF, $\sigma^{1.1} = 18$ for a shape parameter $p = 1.1$. Results are shown in Fig. 2(c) for the CAR prior and Fig. 2(d) for the GGMRF prior.

We can observe that the CAR prior penalizes in excess the edges while both the GGMRF and the proposed CGMRF priors better preserve them. It is important to note that the shape parameter in the GGMRF prior plays an important role in the edge preservation obtaining better results when it is close to 1. We fixed it to 1.1 since, experimentally, we observed that best results were obtained in that case. The CGMRF model provides much sharper edges and zones inside the regions are more uniform and similar to the original image.

We note that the CAR prior is a special case of both the GGMRF and CGMRF priors. The GGMRF prior generates the CAR prior when the shape parameter is set equal to 2 and, as noted previously, the CGMRF is equivalent to the CAR prior when β is very large. Note, however, that, in the presence of edges, the variance of the prior distribution, the inverse of the parameter α , is greater for the CAR

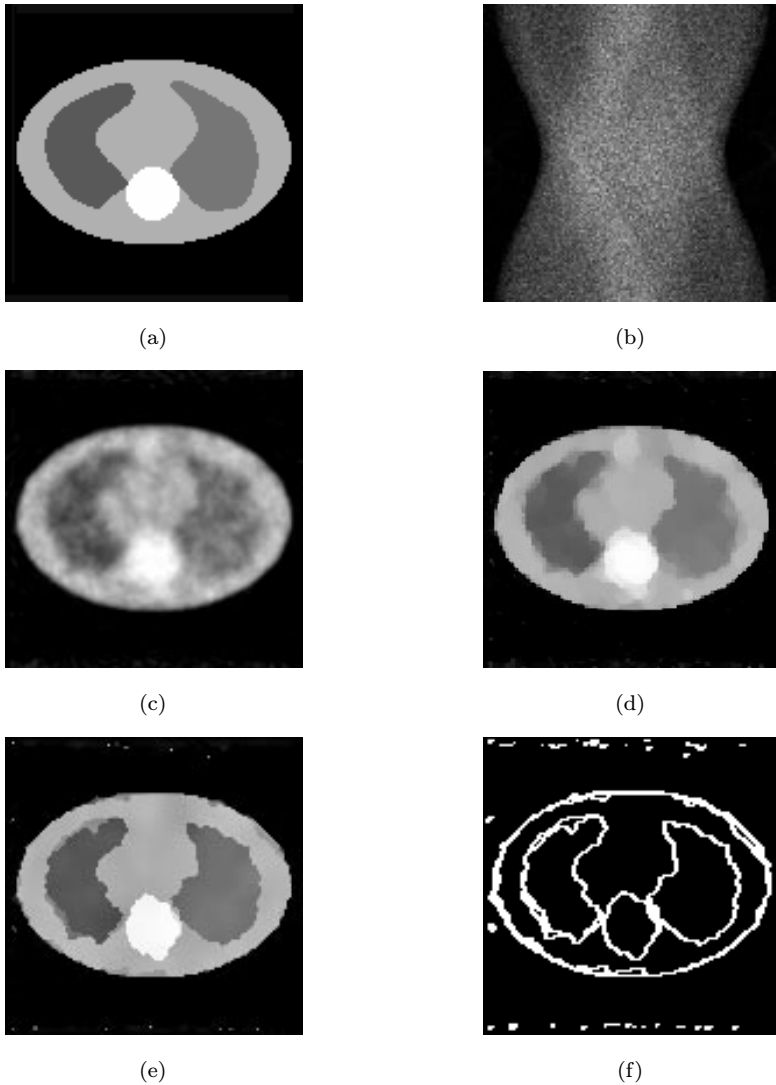


Fig. 2. (a) Original image. (b) Simulated sinogram. Reconstruction using a (c) CAR prior, (d) GGMRF prior and (e) CGMRF prior. (f) Line process corresponding to the reconstruction in (e).

prior than for the CGMRF one. Since the CAR prior does not take into account the presence of edges, the variance of the model increases due to the difference between the pixel in both sides of an edge. However, the CGMRF prior takes into account that pixels across the edges are not related and, hence, the variance of the model, and so the variance within regions, decreases.

Another example with a synthetic image is shown in Fig. 3. The simulation process from the original image [Fig. 3(a)] to the corresponding sinogram [Fig. 3(b)] was

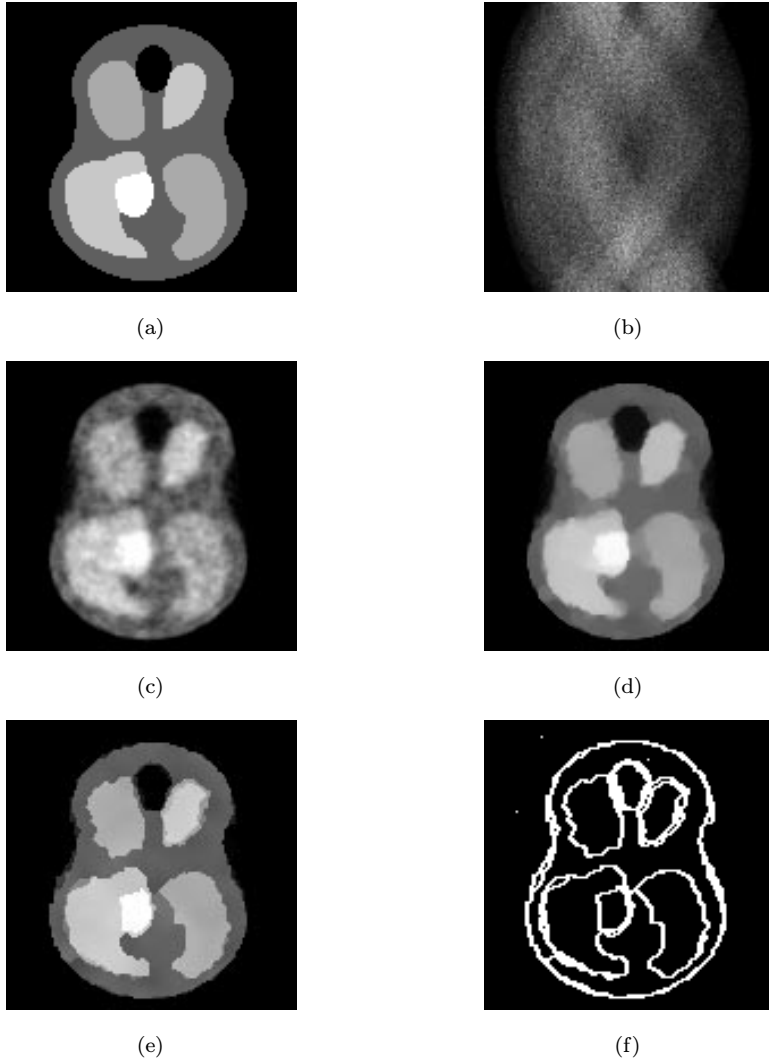


Fig. 3. (a) Original image. (b) Simulated sinogram. Reconstruction using a (c) CAR prior, (d) GGMRF prior and (e) CGMRF prior. (f) Line process corresponding to the reconstruction in (e).

performed as in the previous example. Reconstruction with the proposed method with $\alpha = 24.5^{-1}$ and $\beta = 17.5$ is shown in Fig. 3(e) and, its corresponding line process, in Fig. 3(f). Results for the CAR prior ($\alpha = 563$) and the GGMRF prior ($\sigma^{1.1} = 21.2$ for a shape parameter $p = 1.1$), are depicted in Figs. 3(c) and 3(d), respectively. As in the previous test, the CAR prior oversmooths the edges, furthermore, it produces artifacts in the interior areas. Sharper edges and smooth areas are obtained for the CGMRF prior which, also, recognized all the regions quite accurately.

The method was also tested on real data obtained by a Siemens Orbiter 66601 detector with the collimator being a parallel hole. The gamma-camera, equipped with 64 detectors, described a circular orbit, at 5.625 degrees steps (64 angles). Therefore, the size of the real image is 64×64 . The data corresponds to the inferior part of the liver (the right lobe and the center posterior hepatic zone, crossed there by blood vessels). The data provided by the detector system and its FBP reconstruction are shown in Figs. 4(a) and 4(b), respectively. Looking at the figure, it is

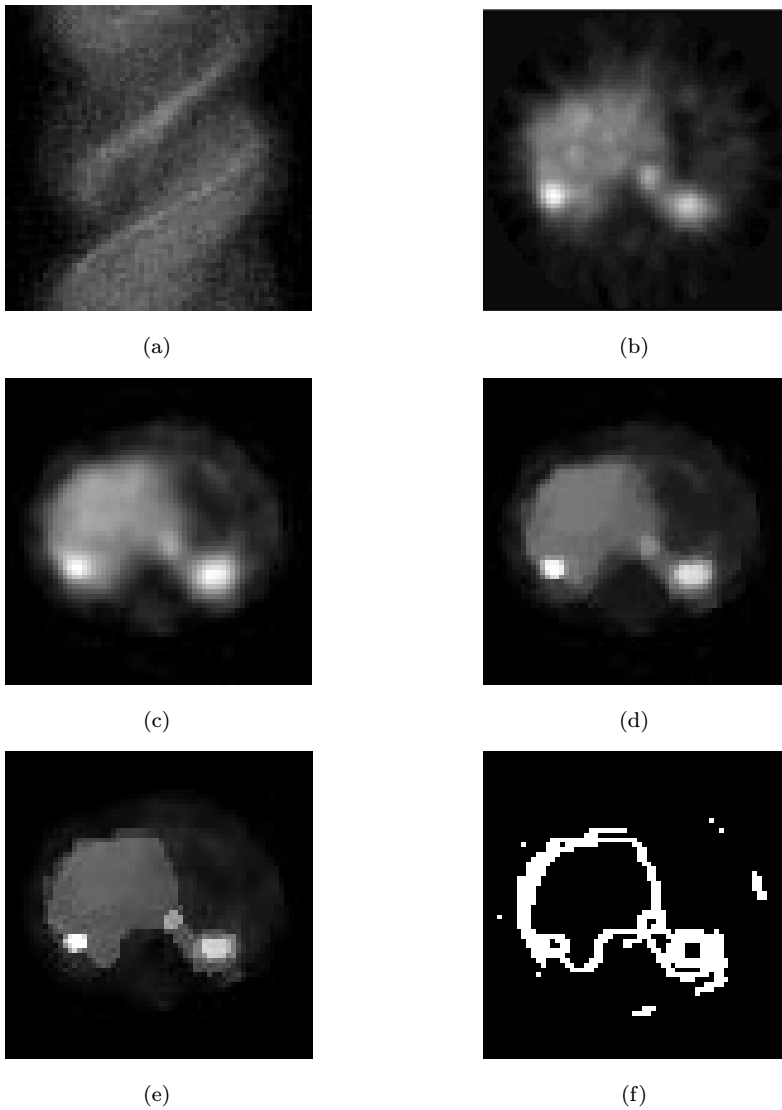
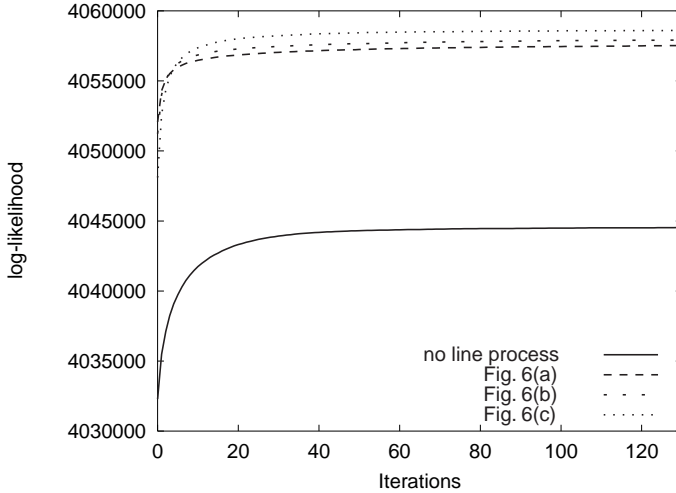
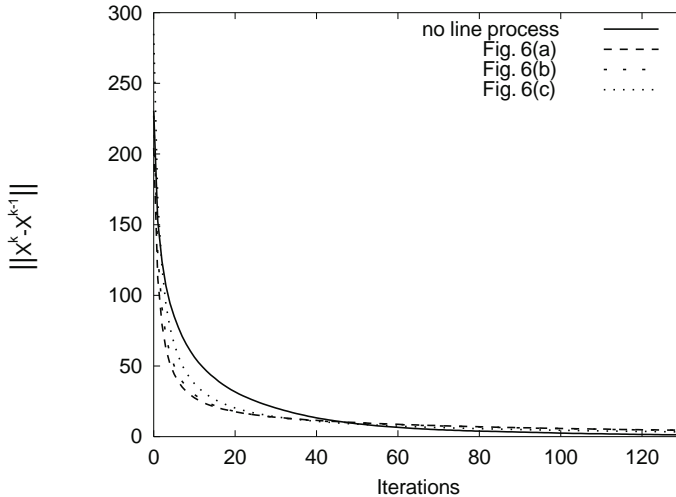


Fig. 4. (a) Observed sinogram. Reconstruction using (b) FBP, (c) CAR prior, (d) GGMRF prior and (e) CGMRF prior. (f) Line process corresponding to the reconstruction in (e).

clear that the CAR, GGMRF and CGMRF prior methods (depicted in Figs. 4(c)–4(e), respectively) provide better results than FBP [see Fig. 4(c)] although, the CAR prior oversmooths the edges. The CGMRF prior method again provides better results than the CAR and the GGMRF ones since areas with different levels of vascularization can be better distinguished. More concretely, the vessel in the center



(a)



(b)

Fig. 5. Convergence results for the synthetic image in Fig. 2(a). (a) Log-likelihood value of the iterative process described in Eq. (10) for the line process configurations shown in Fig. 6, (b) norm of the difference between two consecutive iterations of the iterative process described in Eq. (10) for the line process configurations shown in Fig. 6.

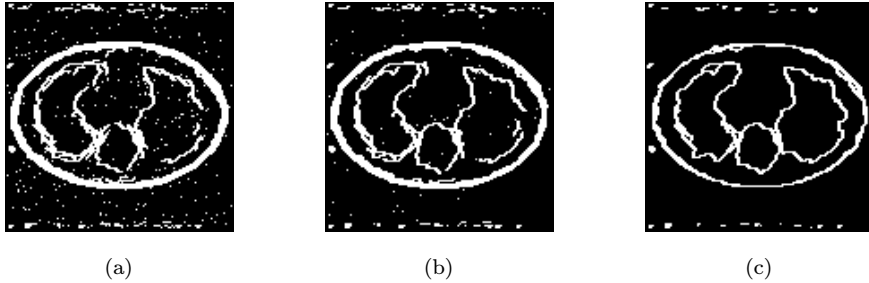


Fig. 6. Fixed line process configurations used in Fig. 5.

part of the image, barely distinguished in the GGMRF reconstruction is clear and sharp in the CGMRF one. Also, the shade around the central objects, due to the refraction on other organs near to the liver, is better eliminated by this method.

In order to test the convergence of the method we run our method on different fixed line process configurations [see Fig. 6]. We obtained in all our tests, a continuous increment in the log-likelihood [see Fig. 5] and, also, a continuous decrease of the norm of the difference between two consecutive iterations [Fig. 5(b)].

5. Conclusions

In this paper we have presented a new method that can be used to reconstruct SPECT images. This method uses a prior model with a line process and the MAP estimation is performed by simulated annealing for the line process and a deterministic iterative scheme for the image. This iterative scheme has been shown to converge experimentally and to provide always non-negative solutions when the iteration starts with a non-negative image. The experimental results demonstrate the improved performance of the proposed algorithm over other competing algorithms.

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