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A Novel Approach to Simultaneous Image Segmentation and Bias Correction

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Abstract-

This paper presents a variational level set approach to joint segmentation and bias of correction images with intensity homogeneity. Our method is based on an observation that intensities in a relatively small local region are separable, despite of the inseparability of the intensities in the whole image caused by the intensity in homogeneity. We first define a weighted K-means clustering objective function for image intensities in a neighbourhood around each point, with the cluster centres having a multiplicative factor that estimates the bias within the neighbourhood. The objective function is then integrated over the domain and incorporated into entire variational level set formulation. The energy minimization is performed via a level set evolution process. Our method is able to estimate bias of quite general profiles. Moreover, it is robust to initialization, and therefore allows automatic applications. The proposed method has been used for images of various modalities with promising results.

Index Terms—Bias field; computer vision; energy minimization; image segmentation; variational approach

1.Introduction

A major problem for automatic segmentation of magnetic resonance (MR) images is the intensity inhomogeneity due to the bias field, which is caused by limitations in imaging devices and subject-induced susceptibility effect.

The bias can cause serious misclassifications when intensity-based

segmentation algorithms are used. Essentially, the misclassification is due to an overlap of the intensity range of different tissues introduced by the bias field, so that the voxels in different tissues are not separable based on their intensities. Intensity inhomogeneities also often occur in images of other modalities, such as X-ray and computed tomography images.

Bias correction has been extensively studied in the past two decades [1,2]. Methods of bias correction can be categorized into two classes: prospective methods and retrospective methods. Prospective methods aim to avoid inhomogeneities in the intensity acquisition process. These methods, capable of correcting intensity inhomogeneity induced by the imaging device, are not able to remove subject-induced intensity inhomogeneity. In contrast, retrospective methods only rely on the information the acquired images. Therefore, they can also remove intensity inhomogeneities regardless of their sources.

Early retrospective methods include those based on filtering [3], surface fitting [4, 5], and histogram [6]. Segmentation based methods [1, 7,8,9] are more attractive, as they unify segmentation and bias correction within a single framework. These methods consist of the interaction between segmentation and bias correction, which benefit each other to yield final segmentation and bias correction.

The generally accepted assumption on the bias field is that it is slowly varying. It is necessary and beneficial to preserve the slowly varying property of the computed bias field in segmentation based methods. In Wells *et al.*'s method [1], the directly computed bias field is



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not smooth, which would lead to poor bias correction and segmentation results.

A moving-average low pass filter is empirically used to force the bias field to be smooth. In [10], Pham and Prince proposed an energy minimization method for adaptive segmentation and estimation of the bias field. In their method, the smoothness of the bias field is ensured by adding a smoothing constraint term in their objective function, which leads to a highly expensive procedure to solve a space-varying difference equation.

Such an expensive smoothing procedure is avoided in some well-known parametric methods (e.g. [8,7]) by modelling the bias field as a polynomial, which is smooth by nature. However, due to limited approximation capability of polynomials, these methods are not able to approximate bias fields of general profiles, such as those in 7T MR images (see Fig. 5 for example).

In this paper, we propose a variational level set approach to bias correction and segmentation for images corrupted with intensity inhomogeneities.

A unique feature of our method is that the computed bias field is intrinsically ensured to be smooth by the data term in our variational formulation, without any additional effort to maintain the smoothness of the bias field, and it can approximate bias fields of more general profiles, such as those in 7T MR images. Moreover, our method is not sensitive to initialization, thereby allowing automatic applications.

2. Method

2.1 Model of Images with Intensity Inhomogeneity

Our method is based on the model commonly used to describe images with intensity Inhomogeneity:

$$I = b \cdot J + n (1) \tag{1}$$

Where I is the measured image intensity, \tilde{J} is the true signal to be restored, \tilde{b} is the bias field,

and n is noise. The superscript tilde in $^{\sim} J$ and $^{\sim} b$ is used to distinguish the unknown true signal $^{\sim} J$ and the bias field $^{\sim} b$ from their estimates, which will be denoted by J and b, respectively.

The generally accepted assumption on the bias field is that it is smooth (or slowly varying). Ideally, the intensity \tilde{J} in each tissue should take a specific value $\tilde{c}i$ of the physical property being measured (e.g. the proton density for MR images). In general, we assume that the true image \tilde{J} and the bias field \tilde{b} have the following **Properties:**

(P1) The bias field b is slowly varying in the entire image domain. **(P2)** The true image intensities J are approximately a constant within each class of tissue, i.e.

 $\tilde{J}(\mathbf{x}) \approx \tilde{c}i$ for $\mathbf{x} \in \tilde{\Omega}i$, with $\tilde{C}\Omega i Ni = 1$ being a partition of Ω .

2.2 Energy Formulation

Our ultimate goal is to separate the image domain Ω into N disjoint regions Ω i, $i=1, \cdots, N$, based on the measured image I. However, due to the intensity inhomogeneity caused by the bias field, the measured intensities are not separable by using traditional intensity based classification methods. In this section, we will propose a new method for joint segmentation and bias correction. Our method is based on an observation that the intensities in a relatively small region are separable, which can be verified by the above assumptions (P1) and (P2) as explained in the following.

We consider a circular neighbourhood with a relatively small radius ρ cantered at each point \mathbf{x} in the image domain Ω , defined by $O\mathbf{x}$ $[\mathbf{y}:|\mathbf{y}-\mathbf{x}|\leq\rho]$. The partition $[\Omega_i]N$ i=1 induces a partition of the neighbourhood $O\mathbf{x}$, i.e., $\{O\mathbf{x}\cap \Omega_i\}N$ i=1. For a smooth function b, the values $b(\mathbf{y})$ for all \mathbf{y} in the circular neighbourhood $O\mathbf{x}$ can be well approximated by $b(\mathbf{x})$, which is at the centre of $O\mathbf{x}$. Therefore, the intensities b

 $(\mathbf{y})\mathcal{F}(\mathbf{y})$ in each subregion $O\mathbf{x} \cap {}^{\sim}\Omega i$ are approximately the constant $b(\mathbf{x}){}^{\sim}ci$. Thus, we have the following approximation



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$$b(\mathbf{y})J(\mathbf{y}) \approx b(\mathbf{x}) ci \text{ for } \mathbf{y} \in O\mathbf{x} \cap \Omega i$$
(2)

The constants(\mathbf{x}) ci can be considered as the approximations of the *cluster centres* (or means) of the clusters $\{I(\mathbf{y}): \mathbf{y} \in O\mathbf{x} \cap {}^{\sim}\Omega i\}$ within the neighbourhood Ox. Therefore, the intensities in the neighbourhood Ox are around N distinct cluster centres $\tilde{m}i \approx \tilde{b} (\mathbf{x})\tilde{c}i$. The multiplicative components(x) and ~ci of the cluster centres ~mi $\approx b(\mathbf{x}) ci$ can be estimated as the following. Consider the task of classifying the intensities I(y) in the neighbourhood Ox into N classes. In view of the reparability of the intensities within the neighbourhood Ox, this task can be performed by using the standard K-means clustering method. In this paper, we introduce a K-means clustering method based on the minimization of the following

Weighted objective function

$$E\mathbf{x} = N_{i} = 1 - O\mathbf{x} \cap \Omega i \ K(\mathbf{x} - \mathbf{y}) / I(\mathbf{y}) - b(\mathbf{x}) ci / 2d\mathbf{y}$$
(3)

Where $b(\mathbf{x})ci$ are the cluster centers to be optimized, and $K(\mathbf{x} - \mathbf{y})$ is a non-negative weighting function such that $K(\mathbf{x}-\mathbf{y}) = 0$ for $|\mathbf{x}-\mathbf{y}| > \rho$ and $O\mathbf{x} \ K(\mathbf{x}-\mathbf{y})d\mathbf{y} = 1$.

Although the choice of the weighting function is flexible, it is preferable to use a weighting function $K(\mathbf{x}-\mathbf{y})$ such that larger weights are assigned to the data $I(\mathbf{y})$ for \mathbf{y} closer to the center \mathbf{x} of the neighborhood $O\mathbf{x}$. In this paper, the weighting function K is chosen as a truncated Gaussian kernel

$$K(\mathbf{u}) = 1$$
 a $e - |\mathbf{u}|/2/2\sigma^2$ for $|\mathbf{u}| \le \rho 0$ else
Where a is a constant such that $K(\mathbf{u}) = 1$. The above objective function $K(\mathbf{u}) = 1$.

Rewritten as

$$E\mathbf{x} = N \ \underline{i=1} \ \underline{\Omega}i \ K(\mathbf{x} - \mathbf{y})/I(\mathbf{y}) - b(\mathbf{x})ci/2d\mathbf{x}$$
(4)

Due to the fact that $K(\mathbf{x} - \mathbf{y}) = 0$ for $\mathbf{y} / \in O\mathbf{x}$.

As mentioned above, the measured intensities $I(\mathbf{y})$ within the neighbourhood $O\mathbf{x}$ are separable, and therefore could be classified into N clusters by minimizing the objective function $E\mathbf{x}$, which results in the optimal cluster centers mi and an optimal partition of $O\mathbf{x}$. However, we still cannot determine the components $b(\mathbf{x})$ and ci of the computed cluster centers mi. Moreover, our ultimate goal is to find an optimal set of a partition of the entire image domain Ω , the bias

field b, and the constants ci. The minimization of a single objective function $E\mathbf{x}$, which is defined for a point \mathbf{x} , does not achieve this goal. We need to minimize $E\mathbf{x}$ for all the points \mathbf{x} . This can be achieved by minimizing the integral of $E\mathbf{x}$ over Ω . Therefore, we define an energy $E _{-} E\mathbf{x}d\mathbf{x}$, i.e.

$$E ___N _i=1 _ \Omega i K(\mathbf{x} - \mathbf{y})/I(\mathbf{y}) - b(\mathbf{x})ci/2d\mathbf{y}_d\mathbf{x}$$
 (5)

Directly minimizing the energy with the partition $\{\Omega i\}N$ i=1 as a variable is not convenient. We will use one or multiple level set functions to represent a partition $\{\Omega i\}N$ i=1. The energy minimization can thus be performed by solving a level set evolution equation.

3. Level Set Formulation

We first consider the case of N=2. In this case, the image domain is partitioned into two regions $\{\Omega i\}2$ i=1. These two regions can be represented by the regions separated by the zero level contour of a function φ , i.e., $\Omega 1 = \{\varphi > 0\}$ and $\Omega 2 = \{\varphi < 0\}$. Using Heaviside function H, the energy E in Eq. (5) can be expressed as an energy in terms of φ , b, and \mathbf{c} as below

 $E(\varphi, b, \mathbf{c}) = __2 _i = 1 _K(\mathbf{x} - \mathbf{y})/I(\mathbf{y}) - b(\mathbf{x})ci/2Mi(\varphi(\mathbf{y}))d\mathbf{y}_d\mathbf{x}$ (6) where $M1(\varphi(\mathbf{x})) = H(\varphi(\mathbf{x}))$ and $M2(\varphi(\mathbf{x})) = 1 - H(\varphi(\mathbf{x}))$. In practice, we use a smoothed Heaviside function $H_(x) = 1$ 2 [1+ 2π arctan($x_$)] to approximate the original Heaviside function H, with $_=1$ as used in [11]. It is necessary to add a regularization term $R(\varphi)$ to the above energy in the following energy functional:

 $F(\varphi, b, c1, c2) = E(\varphi, b, \mathbf{c}) + R(\varphi)$ (7) where $R(\varphi)$ $v = \sqrt{VH(\varphi)}/d\mathbf{x} + \mu = (\sqrt{V\varphi} - 1)2d\mathbf{x}$. The first term in R serves to regularize the zero level contour of φ as in typical level set methods [11], while the second term regularizes the entire level set function φ by penalizing its deviation from Signed distance, as in the level set methods proposed by Li et al. [12,13]. The energy $E(\varphi, b, \phi)$ c) is the data term in our variational framework. Similarly, we can use multiple level set functions $\varphi 1, \cdots, \varphi n$ to represent regions $\{\Omega i\} N i=1$ with N = 2n as in [14]. For convenience, we use a vector valued function $\Phi = (\varphi 1, \cdots, \varphi n)$ to represent the functions $\varphi 1, \dots, \varphi n$. The energy for general multiphase formulation of our method can be defined as



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$$F(\Phi, b, \mathbf{c}) = (N_i = 1 - K(\mathbf{x} - \mathbf{y})/I(\mathbf{y}) - b(\mathbf{x})ci/2Mi(\Phi(\mathbf{y}))d\mathbf{y})d\mathbf{x} + n_i = 1 R(\varphi i)$$
 (8)

Where Mi (Φ) are functions of Φ which are designed such that N i=1Mi (Φ) = 1. The definition of Mi in the four-phase case is given in [14]. For N=3 and two level set functions $\varphi 1$ and $\varphi 2$, we can define $M1(\varphi 1, \varphi 2) = H(\varphi 1)H(\varphi 2)$, $M2(\varphi 1, \varphi 2) = H(\varphi 1)(1-H(\varphi 2))$, and $M3(\varphi 1, \varphi 2) = 1-H(\varphi 1)$ to obtain a three-phase formulation. We only describe the energy minimization for the two-phase case in this paper (the multi-phase case can be solved with the similar procedure). For fixed \mathbf{c} and \mathbf{b} , the minimization of \mathbf{F} (φ , \mathbf{c} , and \mathbf{b}) consists in solving the level set evolution equation as the gradient descent equation

$$\frac{\partial \varphi - \partial F}{\partial t = \partial \varphi} \tag{9}$$

4 Experimental Results and Validation

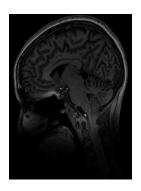
We use the parameters $\sigma = 4$, $\mu = 1$, and ν = 0.001×2552 for all the images in this paper. Our method is robust to the initialization of the constants $\mathbf{c} = (c1, \dots, cN)$, the bias field b, and functions. the level set For automatic applications, the constants $c1, \dots, cN$ can be initialized as N equally spaced numbers between the minimum and maximum intensities of the original image, and the bias field b is initialized as b = 1. The level set functions can be automatically generated or manually initialized by the users. The number of phases N depends on the number of tissue types in the images, which is usually known in practice.

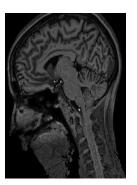
We first demonstrate our method in the two-phase case (i.e. N=2). For example, Fig. 1 shows the result of our method for an X-ray image. Intensity inhomogeneity is obvious in this image. We use this example to show the desirable capability of our method in joint segmentation and bias correction. The bias corrected image is given by the quotient L/b. It is worth noting that our method allows for flexible initialization





Fig. 1. Applications of our method to an X-ray image. (a) Original image and initial contour (dashed black line); (d) Bias corrected image.





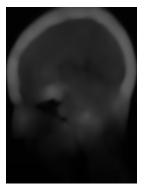


Fig. 5. Application of our method to a 7T

MR image. (a) Original image; (b) Bias corrected image; (c) Computed bias field.

Our method has been tested on 7T MR images. At 7T, significant gains in image resolution can be obtained due to the increase in signal-to-noise ratio. However, susceptibility-induced gradients scale with main field, while the imaging gradients are currently limited to essentially the same strengths as used at lower field strengths (i.e., 3T). Such effects are most pronounced at air/tissue interfaces, as can be seen in Fig. 5(a) at the base of the frontal lobe. This appears as a localized and stronger bias, which is challenging to traditional methods for



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bias correction. This result shows the ability of our method to capture and correct such bias, as shown in Fig. 5(b) and 5(c).

5 Conclusions

We have presented a unified framework of bias correction and segmentation. A unique advantage of our method is that the smoothness of the computed bias field is intrinsically ensured by the data term in our variational formulation. Our method is able to capture bias of quite general profiles, and can be used for images of various modalities.

Moreover, it is robust to initialization, thereby allowing automatic applications. Comparisons with two well-known bias correction methods demonstrate the advantages of the proposed method.

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