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Semi-automated Segmentation and Classification of Digital Breast Tomosynthesis Reconstructed Images

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Abstract

Digital breast tomosynthesis (DBT) is a limited-angle tomographic x-ray imaging technique that reduces the effect of tissue superposition observed in planar mammography. An integrated imaging platform that combines DBT with near infrared spectroscopy (NIRS) to provide co-registered anatomical and functional imaging is under development. Incorporation of anatomic priors can benefit NIRS reconstruction. In this work, we provide a segmentation and classification method to extract potential lesions, as well as adipose, fibroglandular, muscle and skin tissue in reconstructed DBT images that serve as anatomic priors during NIRS reconstruction. The method may also be adaptable for estimating tumor volume, breast glandular content, and for extracting lesion features for potential application to computer aided detection and diagnosis.

I. Introduction

Mammography provides a two-dimensional (2-D) projection image of the three-dimensional (3-D) breast that results in tissue superposition. This could lead to callbacks for additional imaging, missed cancers and false positives. Digital breast tomosynthesis [1] is a limited angle tomographic x-ray imaging technique that acquires multiple projections over an angular range of $\sim \pm 15^\circ$. The acquired projections are reconstructed to ~ 1 mm thick slices parallel to the x-ray imaging detector using analytical or iterative methods [1–5]. Breast

simulating phantom studies and clinical trials have shown the benefits of DBT [1, 5–7]. However, DBT provides anatomical images only. Near Infrared Spectroscopy (NIRS) allows non-invasive imaging of total hemoglobin, oxygen saturation, water content, optical scattering, and lipid concentration [8] through 10 cm of breast tissue. These properties are important physiological parameters that can characterize breast tissue and offer contrast in tumors [9, 10]. Increased total hemoglobin is likely in malignant versus benign lesions because of the leaky, more densely packed vasculature in breast cancers. Additionally, increased metabolism of malignant tumors could decrease hemoglobin oxygen saturation. However, NIRS exhibits poor spatial resolution due to excessive light scatter. Incorporating anatomic priors has shown to improve the spatial resolution and the quantitative accuracy of NIRS reconstruction [11–13]. Hence, we and others [14] are exploring the potential of a combined DBT-NIRS system.

II. Method

Segmentation and classification of reconstructed digital breast tomosynthesis images is non-trivial due to the severity of out-of-slice artifacts arising from angular under-sampling. In addition, DBT images contain more x-ray scatter than mammography due to the lack of an anti-scatter grid during acquisition of projections. X-ray scatter causes artifacts and quantitative inaccuracies in the reconstructed DBT images. Hence, a six-step process was developed and applied to anonymized images from a prior institutional review board (IRB)-approved, HIPAA-compliant clinical study [6]. Fig. 1 shows the imaging geometry (not drawn to scale) and the labeled axes that define the image coordinate system. Sequentially, the steps involved are: (1) angular-constrained bilateral filtering along the x-z plane; (2) unsharp masking along the x-y plane; (3) anisotropic diffusion filtering along the x-y plane; (4) background correction along the y-z plane; (5) fuzzy c-means based segmentation of the 3-D image stack; and (6) morphological operations for void filling of segmented areas. Rationale for each step is provided in their description below. All steps were implemented in MATLAB 7.7.0 (R2008) on an Intel Pentium workstation (2.67GHz dual quad-core processors, 24GB RAM, 64-bit Windows XP).

A. Angular-constrained bilateral filtering

The out-of-slice artifacts in DBT reconstructions appear as laterally shifted low-intensity copies of objects from slices other than in focus, with their intensity reducing as a function of distance from the slice in focus. If the reconstructed slices are stacked and viewed along the x-z plane, they appear as “X”-shaped artifacts. Fig. 2a shows an x-z plane from a clinical case that was reconstructed to 56 slices (z-axis) of 1 mm thickness. We chose a nonlinear edge-preserving bilateral filter [15–17] applied along the x-z plane to reduce these out-of-slice artifacts. Specifically, the bilateral filter proposed by Chen [16] was modified by incorporating a 2θ angular constraint corresponding to the acquisition angular range. Fig. 2b and 2c show the 23×23 Gaussian filter kernels before and after application of the $2\theta = 30^\circ$ constraint. Fig. 2d shows the corrected x-z plane corresponding to Fig. 2a that demonstrates a reduction in out-of-plane artifacts.

B. Unsharp masking

We applied an unsharp masking technique along the x-y plane to provide edge enhancement. The images were low pass filtered with a 5×5 Gaussian kernel, followed by subtraction of the low pass filtered image from the original image. Addition of the resultant image with a weight of 2 provided edge enhancement. This above process could also increase the high frequency noise in the images. Hence, we used anisotropic diffusion filtering for noise regularization.

C. Anisotropic diffusion filtering

We applied an edge-preserving, iterative, anisotropic diffusion filter described by Perona and Malik [18] for noise regularization along the x-y plane. After multiple attempts, we found that a diffusion constant value of 30 that controls the sensitivity to gradients, with 5 iterations, yielded desired results. This filtering step is necessary to minimize the effect of noise while preserving the edges for segmentation. Fig 3 shows a slice (x-y plane) of the clinical case prior to and after application of each of the aforementioned steps. Visually, the filtered images showed reduced noise and do not appear to distort lesion features. This is particularly important if the techniques proposed in here were to be adapted for lesion feature extraction in computer aided detection and diagnosis.

D. Background correction

Inclusion of x-ray scatter in DBT reconstructed images results in quantitative inaccuracies and cupping artifacts, wherein the slices (x-y plane) at the periphery of the breast have, on average, increased intensity relative to, those in the center of the breast. This effect coupled with substantial out-of-slice artifacts could be problematic for segmentation. If they are not accounted for and corrected, lesion contrast could be sufficient even in the top and bottom most slices (x-y plane) which are composed of skin, that could be segmented as part of the lesion. Hence, a background correction method is necessary.

The user provides the input (single mouse click) on the location of the lesion from which the volume of interest (VOI) encapsulating the lesion was extracted. Within the VOI, normalizing the signal intensity in each slice (x-y plane) by its average computed over all slices corrected for fluctuations in mean signal intensity across slices. Fig. 4 shows three slices (x-y plane) extracted from the volume of interest prior to and after normalization.

Chan [19] described a background correction technique that was used with a method for computer-aided classification of mammograms. We applied a similar correction along the y-z plane. Fig. 5 shows a slice extracted from the VOI where the lesion is prominent, before and after application of the background correction.

E. Fuzzy C-Means (FCM) segmentation

Fuzzy C-Means (FCM) clustering algorithm is a widely used automated and unsupervised technique for medical image segmentation [20]. Several investigators have applied this technique for lesion segmentation in digital mammography and breast MRI [19, 21, 22]. In our implementation, the FCM clustering algorithm classifies pixels with similar gray levels into a cluster. After initial arbitrary assignment of the center of a cluster, it is iteratively updated by minimizing an objective function that is dependent on the Euclidean distance of gray levels to the center of the cluster. We applied this FCM clustering method to segment the 3-D VOI into two clusters representing the background tissue and the lesion. This clustering generated a rough 2-D binary mask for each slice that needed further refinement. Fig. 6(a) shows the results of FCM clustering applied to a representative slice (x-y plane) of the VOI shown in Fig. 5(b).

F. Additional morphological operations

Masks generated from FCM clustering may result in isolated pixels that correspond to pixel gray levels similar to the lesion. Hence, additional morphological operations are required. The morphological operations performed were area filtering to extract the largest connected label to represent the lesion; 3-D void filling; removal of salt and pepper noise by morphological binary opening followed by binary closing; and, residual removal and mask smoothing by binary erosion followed by binary dilation. Fig. 6b through Fig. 6d shows the result after each step.

In addition to FCM-based segmentation of the lesion, application of the FCM clustering algorithm on each anisotropic diffusion filtered slice (x-y plane, similar to that shown in Fig. 3d) yielded segmentation for adipose, fibroglandular, skin, and muscle tissues. It was necessary to segment the lesion from the background independently of other tissues, as the gray level of the lesion is similar to fibroglandular tissue. Fig. 7a shows a representative DBT reconstructed slice prior to segmentation. Fig. 7b shows the segmented slice color-coded to represent skin (cyan), adipose (red), fibroglandular (yellow), muscle (magenta), and lesion (blue).

III. Conclusion

The method presented here demonstrates a strategy for segmenting DBT images that are challenging due to out-of-slice artifacts and inclusion of x-ray scatter. The only user input required is the identification of the location of a possible lesion. Segmentation and classification of tissue types is fully automated. To date, the method has been applied to seven clinical datasets. Visually, the lack of lesion distortion suggests that the proposed technique may be adaptable for initial image processing as part of a computer-aided detection or diagnosis technique. We are pursuing additional refinements including improved estimation of skin layer thickness and determining filter parameters that are best suited for the task. Quantitative validation such as verifying the estimated tumor volume and glandular fraction content, and suitability of the method to provide NIRS anatomic priors are subjects of ongoing investigations.

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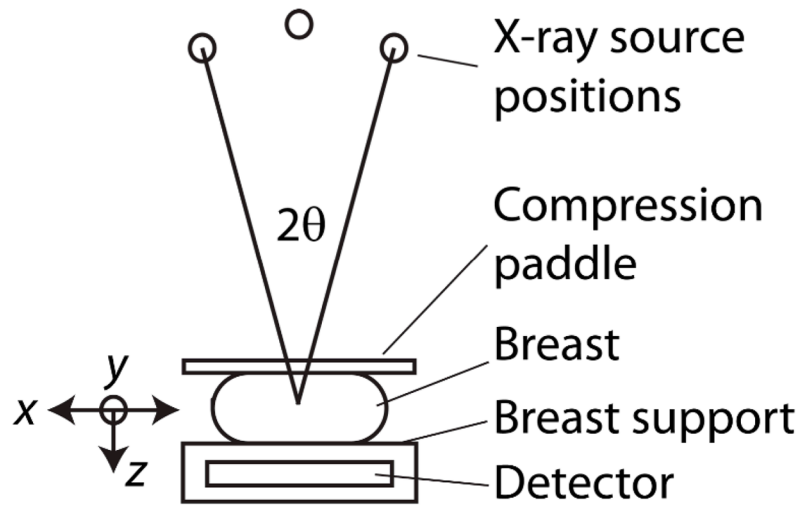


Fig. 1. Imaging geometry for DBT acquisition (not drawn to scale), over the angular range of 2θ . The labeled axes define the orientation in which the filters are applied.

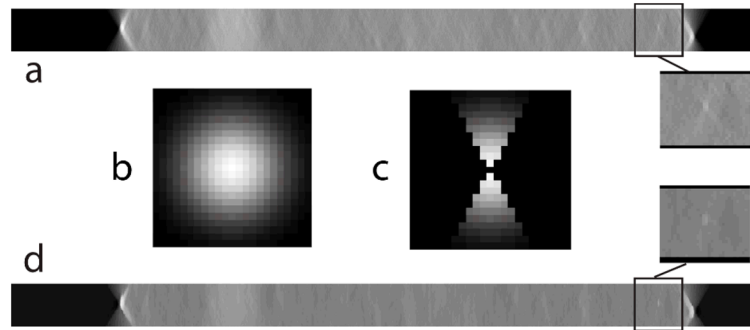


Fig. 2.

a. Shows an x-z plane image from the 3-D DBT image stack for a clinical case. b. 23×23 Gaussian kernel used in the bilateral filter before modification. c. Modified Gaussian kernel with an angular constraint of 2θ corresponding to the angular range during projection view acquisition. d. x-z plane after application of the angular-constrained bilateral filter that shows a reduction in out-of-plane artifacts. Zoomed ($1.5\times$) regions of interest before and after application of the filter are shown.

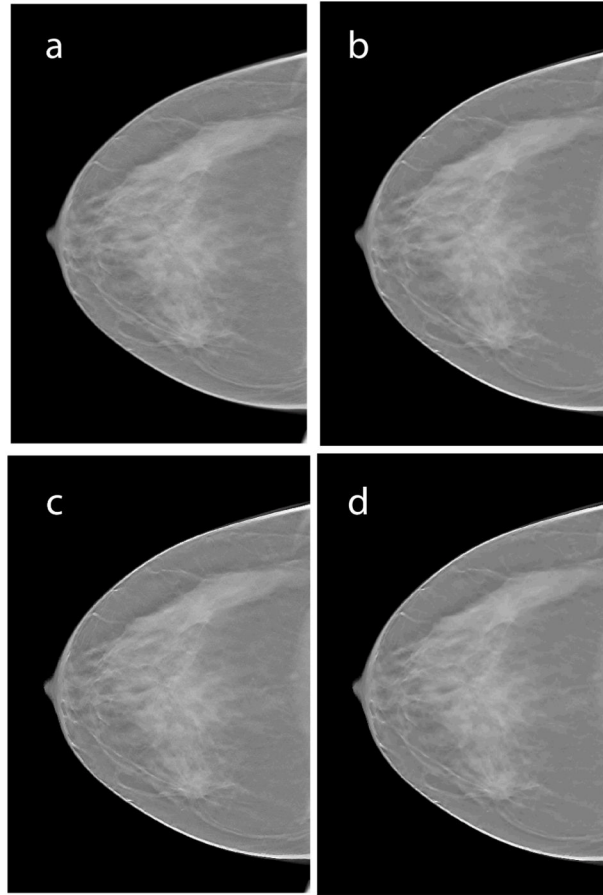


Fig. 3. Shows a single slice (x-y plane) of the reconstructed clinical case where the lesion is prominent prior to and after application of each of the following steps: a. reconstructed slice prior to filtering; b. after angle-constrained bilateral filtering; c. after unsharp masking; d. after anisotropic diffusion filtering. Visually the filtered images showed reduced noise and did not distort lesion features.

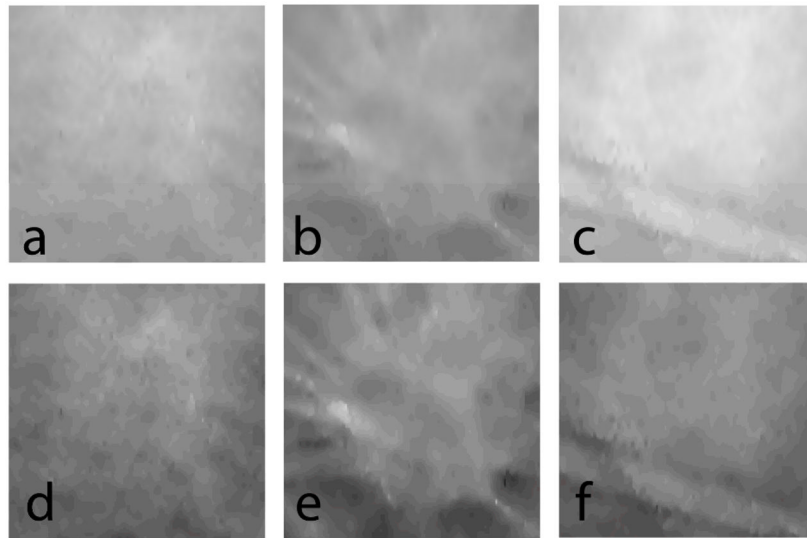


Fig. 4. Three slices (x - y plane) extracted from the volume of interest (VOI) encapsulating the lesion: a. top slice that is in contact with the compression paddle; b. slice that best represents the lesion; and c. bottom slice that is contact with the breast support. (a–c) are prior to and (d–f) are after VOI normalization.

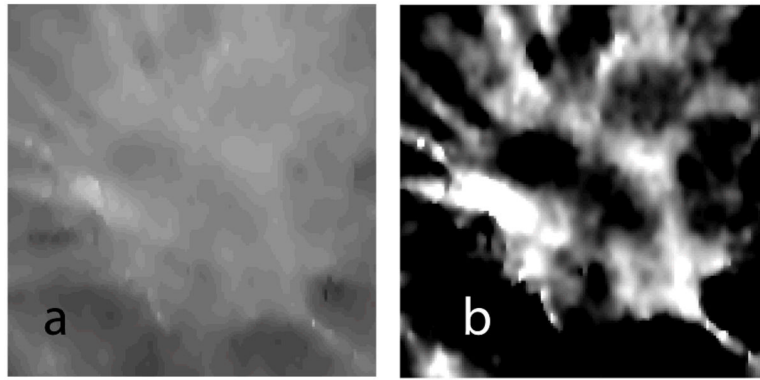


Fig. 5. A slice (x-y plane) containing the lesion extracted from the VOI before (a), and after (b) background correction.

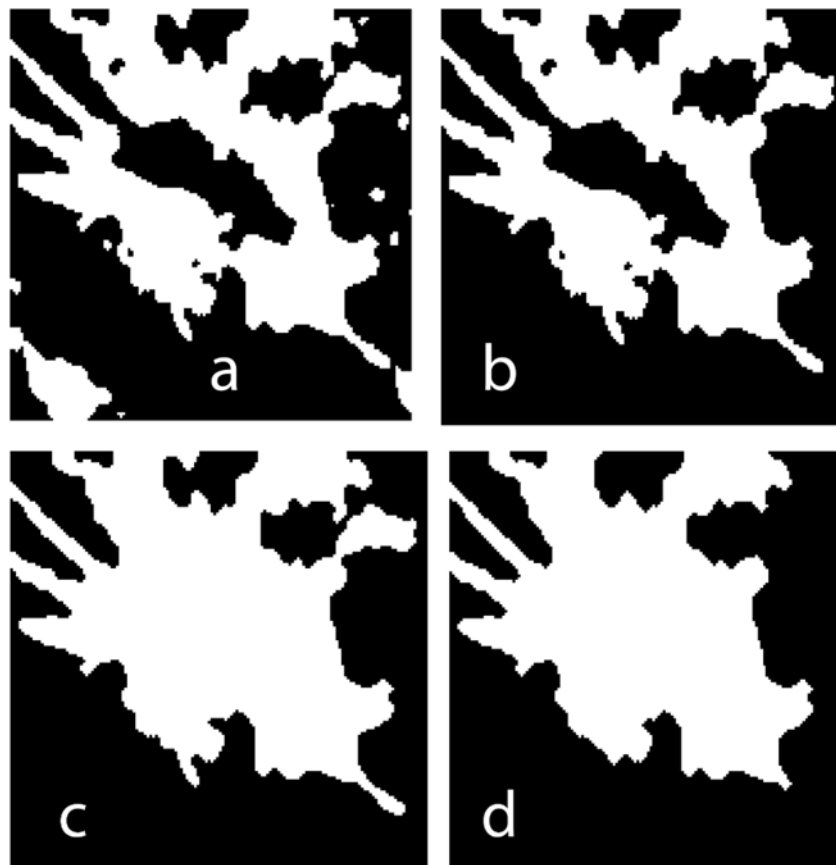


Fig. 6. A slice (x-y plane) containing the lesion extracted from the VOI (a) after FCM-clustering based segmentation, (b) after area filtering to remove isolated segments, (c) after 3-D void filling, and, (d) final smooth mask.

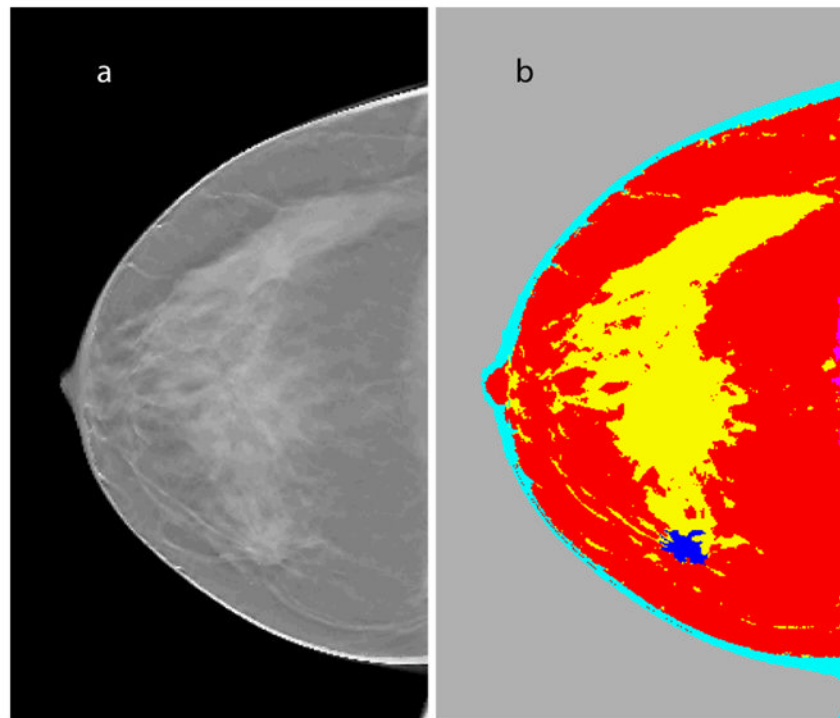


Fig. 7.
a. Representative slice (x-y plane) from DBT reconstruction that prominently shows the lesion; b. segmented slice color-coded to represent skin (cyan), adipose (red), fibroglandular (yellow), muscle (magenta), and lesion (blue).