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Low-dose 2D X-ray angiography enhancement using 2-axis PCA for the preservation of blood-vessel region and noise minimization



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ABSTRACT

Enhancing 2D angiography while maintaining a low radiation dose has become an important research topic. However, it is difficult to enhance images while preserving vessel-structure details because X-ray noise and contrast blood vessels in 2D angiography have similar intensity distributions, which can lead to ambiguous images of vessel structures. In this paper, we propose a novel and fast vessel-enhancement method for 2D angiography. We apply filtering in the principal component analysis domain for vessel regions and background regions separately, using assumptions based on energy compaction. First, we identify an approximate vessel region using a Hessian-based method. Vessel and non-vessel regions are then represented sparsely by calculating their optimal bases separately. This is achieved by identifying periodic motion in the vessel region caused by the flow of the contrast medium through the blood vessels when viewed on the time axis. Finally, we obtain noise-free images by removing noise in the new coordinate domain for the optimal bases. Our method was validated for an X-ray system, using 10 low-dose sets for training and 20 low-dose sets for testing. The results were compared with those for a high-dose dataset with respect to noise-free images. The average enhancement rate was $93.11 \pm 0.71\%$. The average processing time for enhancing video comprising 50–70 frames was 0.80 ± 0.35 s, which is much faster than the previously proposed technique. Our method is applicable to 2D angiography procedures such as catheterization, which requires rapid and natural vessel enhancement.

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1. Introduction

Two-dimensional X-ray angiography is the most commonly used imaging modality for artery diseases monitoring and facilitating catheter-based procedures, such as FFR, IVUS, OCT, stent or peacemaker implantation. Briefly, it provides online two-dimensional projections of considered vessel to the clinician, which manually injects the contrast agent into the vessel since blood itself does not absorb enough amounts of X-rays to be visible. As a consequence, patients and medical staff can be exposed to X-ray radiation for prolonged periods. To

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http://dx.doi.org/10.1016/j.cmpb.2015.09.011 0169-2607/© 2015 Elsevier Ireland Ltd. All rights reserved. minimize this harmful exposure, very low levels of X-ray dosage are used. However, this leads to considerable degradation of image quality through quantum noise [1,2]. Therefore, the ability to enhance images while maintaining a low radiation dose has become an important research topic.

There have been a number of studies on vessel enhancement methods. One popular technique is based on the second-order Hessian matrix [3-6]. And there are approaches with other filters that use information from neighboring pixels, such as the directional filter-bank-based method [7,8], the method using morphological connected-set filters [9-11], and the method using curvilinear filters [12]. To use more information, several nonlocal filters have been proposed for both the spatial domain and the frequency domain. Buades et al. [13] proposed a weighted nonlocal filtering in spatial domain. In this method, noises are removed by a pointwise estimate of the image where each pixel is obtained as a weighted average of pixels centered at regions that are similar to the region centered at the estimated pixel. Because all pixels of image can be used for averages, the estimates are non-local. One popular technique that uses a nonlocal filter in the frequency domain is block-matching and 3D filtering (BM3D) [14]. BM3D filter uses a procedure termed grouping and collaborative filtering. Grouping procedure is the step to find mutually similar 2D image patches and stack them in 3D blocks. And collaborative filtering procedure is the step to estimate all grouped 3D blocks by filtering them jointly by using hard-threshold or wiener filter in transform-domain. Video block-matching and 3D filtering (VBM3D) [15] applies BM3D successfully to a video system. In this method, groups are 3D blocks of mutually similar 2D patches extracted from a set of consecutive frames of the video sequence. BM3D and V-BM3D schemes are applied in medical imaging applications [16–18].

Using the assumptions related to sparse representation in BM3D, it is possible for noise-free images to be sparsely represented in the transform domain. Even though the temporal Fourier transform is effective in making the signal sparse when the image undergoes periodic motion, other transforms might be more efficient for other types of motion. Unlike the Fourier transform, the principal component analysis (PCA) transform is data dependent. It is well known that the PCA transform gives optimal energy compaction, with most of the energy accounted for by only a few coefficients [19]. Here, energy relates to image information such as the brightness magnitude and the variation of brightness between neighborhoods. Our method is based on these assumptions. The 2D angiography used in clinical practice involves the flow through the blood vessels of a contrast medium, which is a substance used in medical imaging to enhance the X-ray images of fluids within the body. The resulting images will contain more motion, both general and periodic, than those for other video applications. This fact enables sparser and less noisy images to be produced by using the PCA transform and removing low-energy coefficients. In addition, we are obtaining the PCA transform matrix of each vessel region and background region from difference of motion properties, which will preserve important information about the blood-vessel structures.

In this paper, we present a novel vessel enhancement method based on removing separately the noise in the extracted vessel region and in the background region within the PCA domain. Conventional approaches enhance images by combining neighboring pixels in a 3D kernel which includes temporal information of those neighboring pixels. However, our method uses only temporal information without the convolution's operation, which decreases the computation time by removing the convolution calculations required in a spatial domain, enhances images non-artificially and reduces the number of parameters to be optimized. Furthermore, we enhance the vessel region and the background region separately in PCA domain. And, it removes noise while preserving the details of vessel structures.

The remainder of the paper is organized as follows. Section 2 describes the proposed three-step method for 2Dangiography vessel enhancement. Section 3 presents the results of the application of the proposed method to clinical datasets in comparison with other previous methods. Finally, we discuss results/correctness analysis, novelty and future work in Section 4 and summarize our method in Section 5.

2. 2D angiography enhancement method

The proposed method comprises three main steps, as shown in Fig. 1. First, we extract an approximation of the whole vessel structure from the 2D angiography and partition the whole image into a vessel region and a background region. Second, all pixels are transformed from the spatial domain to the time domain, with all pixels transformed to a new coordinate system having a basis in each region obtained by PCA. Finally, frames are enhanced using a hard threshold in the PCA domain, and each region is converted back from the PCA domain to the spatial domain by an inverse transformation.

2.1. Extraction of approximate vessel region

In this step, we extract an approximate vessel region that will include most of the flowing vessels for each frame of the input 2D angiography. This process is a necessary step in distinguishing between intensity variation caused by noise and that caused by the contrast medium. A blood vessel emphasized by the contrast medium is the most important structure in 2D angiography. The structure should be preserved and distinguished from noise.

To extract the vessel region, we construct a digital subtraction angiography (DSA) by subtracting the average of images obtained after injecting contrast medium into the blood vessels from the average of images before the injection. Because the overall noise distribution approximates a Poisson distribution in X-ray imaging, the expectation of noise asymptotically approaches zero. Averaging the images therefore has a denoising effect. We find candidate vessels via the Hessian-based method of multiscale vessel enhancement [3]. We then remove wrongly extracted noise and connect broken vessels by applying a morphological operator. In addition, we make the edges of the vessel region robust by including an area beyond the calculated extent of the blood vessels (see Fig. 2). Download English Version:

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