

Image denoising Using Magnetic Resonance Guided Positron Emission Tomography

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Abstract—With the growing interest in conducting multi- centre and multi-modality studies on neurological disorders, post-reconstruction PET image enhancement methods that take advantage of available anatomical information are becoming more important. In this work, a novel method for denoising PET images using the subject's registered T1-weighted MR image is proposed. The proposed method combines the non-local means approach with the twicing strategy from the image denoising literature to restore a reconstructed PET image. Preliminary analysis shows promising improvements in peak signal to noise ratio (PSNR) and contrast recovery coefficients (CRC) of the lesions when denoising simulated images reconstructed using the MLEM algorithm.

Index Terms—PET-MR, denoising, restoration, non-local means, twicing.

1. INTRODUCTION

PET is a powerful tool in both research and diagnosis of various brain disorders. However, reconstructed PET images usually suffer from low signal to noise ratio and low spatial resolution. Therefore, many methods have been devised to address these issues within or after reconstruction. In this work, we are interested in post-reconstruction methods. Post-reconstruction methods have two clear advantages. First, they can be applied to reconstructed images for which the raw measurement data is no longer available. Second, in multicentre datasets, performing minimal set of correction steps (Eg. attenuation/scatter corrections) at the scanner site can reduce the between-centre variability by applying the same post-reconstruction methods for partial volume correction and noise reduction to the whole dataset. Because of the low spatial resolution and high amount of noise in PET images, using anatomical information from MR images in reconstruction, denoising and partial volume correction of PET images has become increasingly popular. In this work, we propose a novel method which incorporates anatomical information from MR images for restoring PET images. Anatomical information has been previously utilized in a non-local means (NLM) method [2] for denoising PET images: authors in [3] have proposed using NLM for PET image denoising while preserving edges, by performing the average in the NLM only in ROIs derived from the subject's CT image. In this work however, instead of only using boundary information from the MR image, the weights in a rotationally invariant NLM method are obtained from the subject's registered T1-weighted MR image. The problem with using a different modality for weight computation is that the resulting image can become excessively blurred in regions where the anatomy in the MR image does not agree with the functional activity in the PET image. To overcome this problem and motivated by the well-known twicing

strategy, the PET-unique signals are denoised using weights obtained from the PET image and the result is added to the denoised image.

2. METHODOLOGY

A. Background

1) *Non-local means (NLM)*: NLM is a simple yet powerful denoising method[1]. It is based on the assumption that patches extracted from a natural image contain redundant information. In essence, in the NLM method a voxel \hat{x}_i in the denoised image \hat{x} is estimated as the weighted average of other voxels in the image with the weights obtained by a robust similarity measure based on the distance of the patches around voxels.

$$x_i = \frac{1}{\sum_{j \in \Lambda_i} w_j} \sum_{j \in \Lambda_i} x_j w_{ij} \quad (1)$$

where Λ_i is the neighbourhood around and including x_i , and P_k is the patch centred at location k in the image.

2) *Rotationally invariant nonlocal-means (RI-NLM3D)*: In order to increase the number of samples contributing to the average in the NLM method and thus reduce noise further, several rotationally invariant variations of NLM have been proposed. In this work, we use a very simple and computationally efficient rotationally invariant NLM method proposed by Manjon et al. for denoising MR images [5].

$$w_{i,j} = \frac{e^{-x_i - x_j} + \mu p_i - \mu p_j}{\sigma^2} \quad (2)$$

where μp_i is the mean value of the patch P_i . Note that the above similarity measure is sensitive to noise and hence usually a pre-filtered image is used to compute the weights, for subsequent application to the unfiltered image. Each step of the RINLM3D has 3 parameters: patch size, the σ parameter and neighbourhood size for the average.

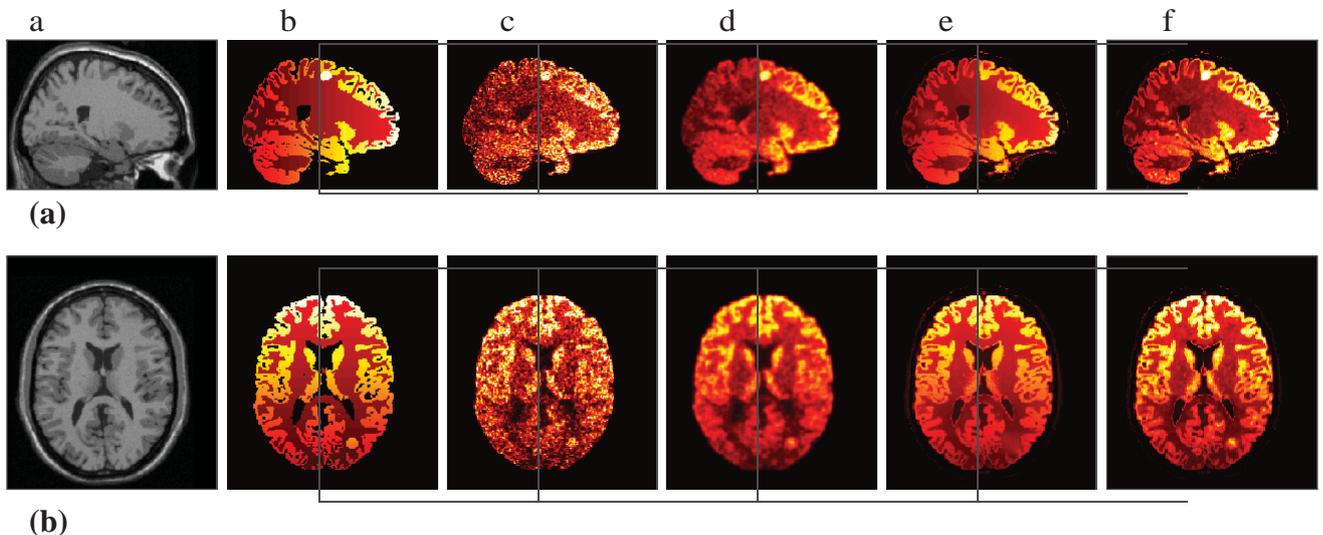


Figure 1: Denoising of a reconstructed image of a sample noisy realization. (a) Registered MR image. (b) The ground truth (c) Noisy reconstructed image after 50 iterations (where the best PSNR is obtained). (d) Post-smoothed image after 70 iterations (where the best PSNR was obtained), $\sigma = 1.2mm$ (CRC of lesions in: WM=0.38, GM=0.56). (e) Denoised image using weights from MR (step 1). (f) Final denoised image after step 2, (CRC of the lesions in: WM=0.58, GM=0.81).

A. Proposed method

We propose a two-step algorithm for restoring PET images. In the first step, the subject's registered MR image is used to compute the weights based on the rotationally invariant formulation in Eq. 2. These weights are then used to denoise the PET image.

The second step is motivated by the twicing method in the denoising literature [7]. In this step, first the residual image is obtained by subtracting the denoised image in the first step from the noisy image. Then, the noisy PET image is pre-filtered using a Gaussian filter and the resulting image is used to compute weights based on Eq. 2. The weights are then used to denoise the residual image. This way, signals specific to the PET image that have been lost in the first step (due to using weights computed from the MR image) are added to the denoised image.

I. SIMULATION STUDY

A simulation of ^{18}F FDG images was performed using the BrainWeb phantom (<http://brainweb.bic.mni.mcgill.ca/brainweb>). The corresponding segmented brain was resampled to HRRT image space (256x256x256).

Realistic concentrations in the grey matter (GM) and the white matter (WM), were estimated using the radioactivity distribution values from PET SORTEO (<http://sorteo.cermep.fr/home.php>). Two hot lesions, one in the GM and one in the WM were artificially added to the simulation. To further investigate the ability of the method to capture information specific to the PET image and absent in the MR image, the simulated PET radioactivity distribution was multiplied by a gradient 3D image in the coronal direction (Fig. 1.b). The noise-free 3D image was forward projected to generate noise-free sinogram data. Then Poisson noise was introduced to the sinogram data to

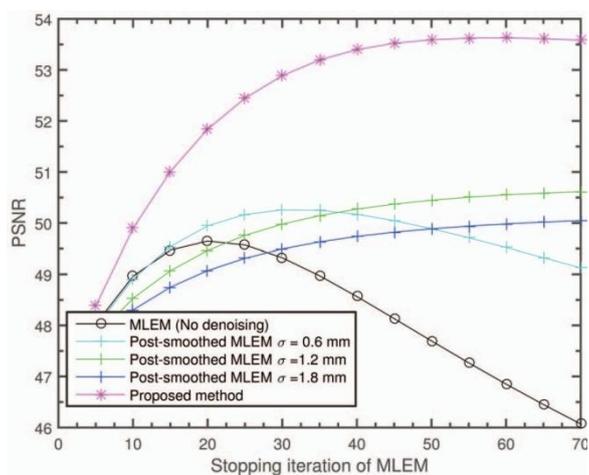
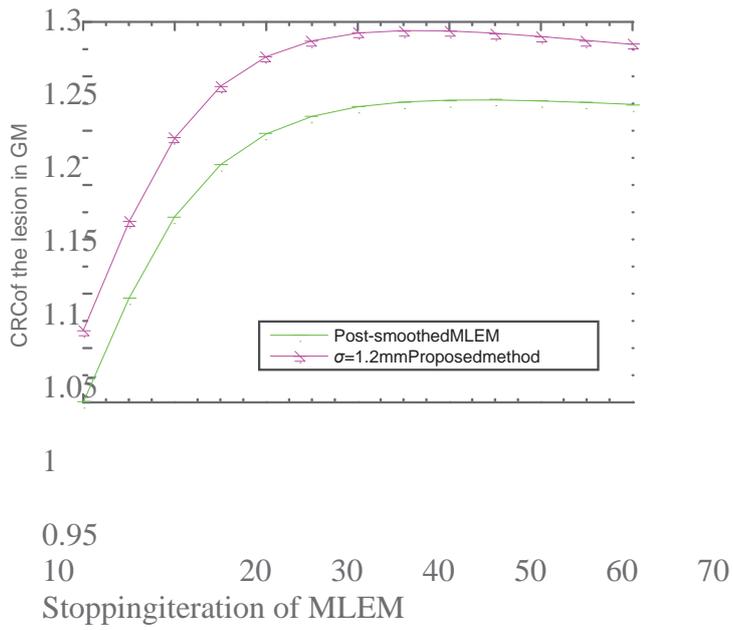


Figure 2: Mean PSNR values across 5 realizations for different methods as a function of stopping iteration of the MLEM reconstruction

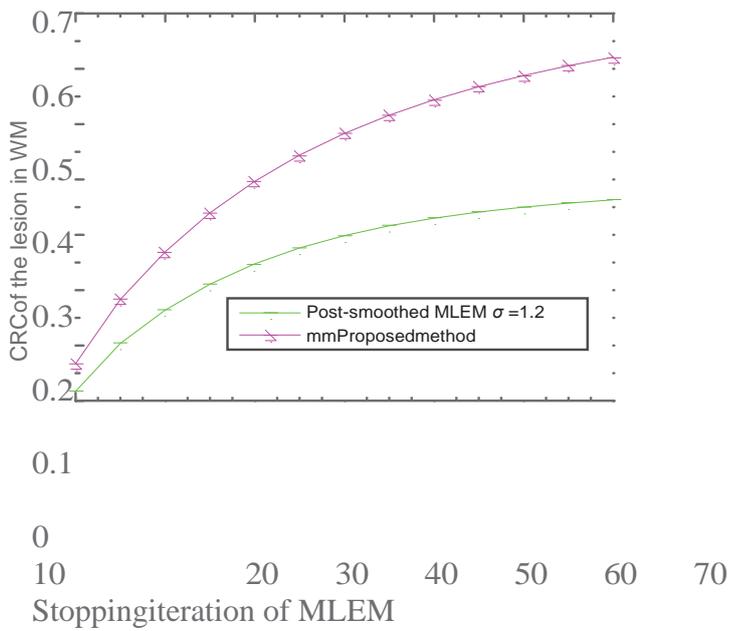
generate 10 noisy realizations. The resulting sinogram data were then reconstructed using the MLEM algorithm.

For both steps of the method, the patch size is set to

33 and the σ value is set to 0.05. The neighbourhood is a patch of size $15 \times 15 \times 15$ for the first step and a patch of size 999 for the second step. Fig. 2 shows the mean PSNR value of the denoised image across 5 realizations as a function of stopping iteration of the MLEM algorithm for the proposed method and also for Gaussian post-smoothing with different σ values. The results indicate the superiority of the proposed method to Gaussian smoothing. The PSNR of the proposed method reaches a maximum at 50 iterations and then plateaus. Fig. 1 shows the output of different steps of the proposed method on a reconstructed PET image stopped after 50 iterations as well as Gaussian post-smoothing at its best performance (highest PSNR). Note how the lesions



(a)



(b)

Figure 3: The mean contrast recovery coefficients of the two lesions embedded in grey matter (GM) and white matter (WM) across 5 realizations.

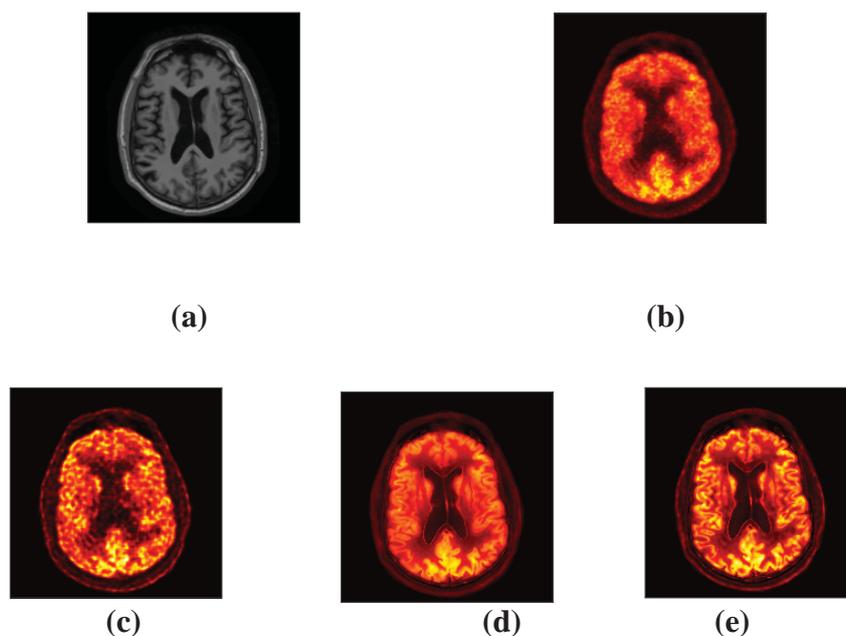


Figure 4: Denoising result of a baseline FDG image of an MCI patient from the ADNI dataset. (a) Registered MR image. (b) Noisy reconstructed image. (c) PVE corrected image. (d) denoised. (e) PVE corrected+denoised.

which disappeared in step one (Fig. 1(e)) become visible after adding the denoised residual (Fig. 1(f)).

In order to evaluate the ability of the proposed method in regions where functional anatomy in PET does not agree with the anatomical information in MR, the contrast recovery coefficients (CRC) of the two lesions were obtained and compared to those for Gaussian post smoothing. Fig. 3 compares the mean CRC of the lesions across multiple realizations for the proposed method and the Gaussian post-smoothing with the σ parameter value leading to maximum PSNR (i.e. $\sigma = 1.2\text{mm}$). The results indicate that the proposed method provides a higher contrast in these regions for all iterations of MLEM.

APPLICATION ON REAL DATA

An FDG image of a subject with mild cognitive impairment (MCI) with four 5-minute co-registered frames was obtained from the Alzheimer's Disease Neuroimaging Initiative (ADNI, <http://www.adni-info.org/>) database. The co-registered frames were then averaged together and the resulting static image was co-registered to the subject's baseline T1-weighted MR image. Then, the proposed method was applied to the resulting image with the same parameters used for the simulation data. In addition, in order to correct for PVE, 10 steps of the Richardson-Lucy deconvolution algorithm [4], [6] with FWHM equal to 6.5mm were applied to the image prior to denoising. The result of the denoising with and without PVE correction is shown in Fig. 4.

3. CONCLUSION

In this study a novel MR-guided PET image denoising method was proposed. Simulation results show that the proposed method is able to significantly improve the peak signal to noise ratio of PET images reconstructed using the MLEM algorithm. Furthermore, we show that the proposed method can be applied after the conventional PVE correction method to restore the PET images by both reducing the

noiseandrecoveringtheboundaries.

4. REFERENCES

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