

Normalization of T2W-MRI Prostate Images using Rician *a priori*

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ABSTRACT

Prostate cancer is reported to be the second most frequently diagnosed cancer of men in the world. In practise, diagnosis can be affected by multiple factors which reduces the chance to detect the potential lesions. In the last decades, new imaging techniques mainly based on MRI are developed in conjunction with Computer-Aided Diagnosis (CAD) systems to help radiologists for such diagnosis. CAD systems are usually designed as a sequential process consisting of four stages: pre-processing, segmentation, registration and classification. As a pre-processing, image normalization is a critical and important step of the chain in order to design a robust classifier and overcome the inter-patients intensity variations. However, little attention has been dedicated to the normalization of T2W-MRI prostate images. In this paper, we propose a method based on a Rician *a priori* in order to normalize T2W-MRI prostate images. A comparison with the state-of-the-art methods is also provided. The normalization of the data is assessed by comparing the alignment of the patient Probability Density Function (PDF)s in both qualitative and quantitative manners. In both evaluation, the normalization using Rician *a priori* outperforms the other state-of-the-art methods.

Keywords: Prostate cancer, T2W-MRI, normalization, pre-processing, computer-aided diagnosis

1. INTRODUCTION

Prostate Cancer (CaP) has been reported the second most frequently diagnosed cancer of men accounting for 13.6%.¹ In United States, aside from skin cancer, CaP was considered to be the most commonly diagnosed cancer among men, implying that approximately 1 in 6 men will be diagnosed with CaP during their lifetime. The *American cancer society* also reported an estimated 233,000 new cases of prostate cancer in 2014.² To address these dramatic issues, more systematic screenings are organized through Prostate-Specific Antigen (PSA) test with further Transrectal Ultrasound biopsy if necessary. However, these tests are unreliable or invasive and that is why further investigations using Magnetic Resonance Imaging (MRI)-Computer-Aided Diagnosis (CAD) are motivated. In the past decades, several CAD systems have been proposed in order to assist the radiologists with their diagnosis. These systems are usually designed as a sequential process consisting of four stages: pre-processing, segmentation, registration and classification. As a pre-processing steps, image normalization is an important step of the chain. Normalization is a highly crucial step to overcome the inter-patient intensity variations occurring, enforce the repeatability, and achieve a robust classification.³ However, little attention has been dedicated to the problem of normalization of T2W-MRI prostate images.³

Artan *et al.*^{4,5} and Ozer *et al.*^{6,7} proposed to normalize the T2W-MRI images by computing the standard score (i.e., *z-score*) of the Peripheral Zone (PZ) pixels such as:

$$I_s(x) = \frac{I_r(x) - \mu_{PZ}}{\sigma_{PZ}}, \forall x \in PZ, \quad (1)$$

where, $I_s(x)$ and $I_r(x)$ are the standardized and the raw signal intensity, respectively, and μ_{PZ} and σ_{PZ} are the mean and standard deviation of the PZ signal intensity. This transformation enforces the image Probability

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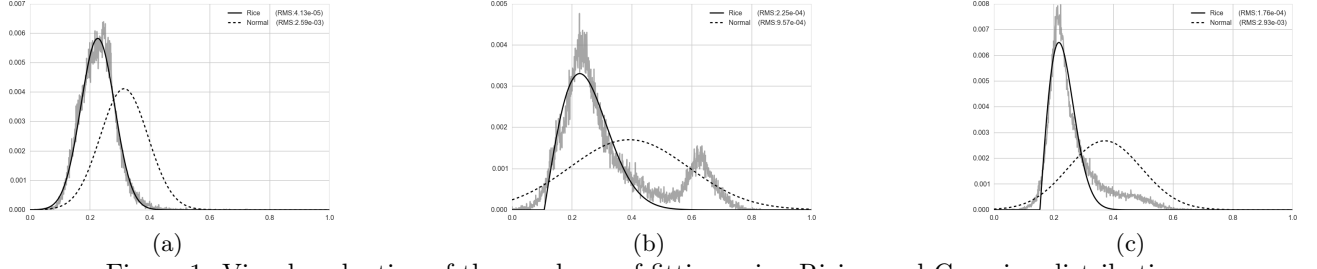


Figure 1: Visual evaluation of the goodness of fitting using Rician and Gaussian distribution.

Density Function (PDF) to have a zero mean and a unit standard deviation. However, this normalization is not appropriate if the PDF do not follow a Gaussian distribution as illustrated in Fig. 1

Lv *et al.*⁸ used the method proposed by Nyul *et al.*⁹ For a given patient, a warping function is inferred by matching some specific landmarks (i.e., median and different percentiles) of the current PDF to the same landmarks learned during a training phase from several patients. The mapping between each landmark is performed using a linear mapping. Viswanath *et al.*¹⁰ used a variant of the previous method by segmenting first the image using region growing with a pre-defined homogeneity criterion and keeping only the largest region to build the PDF. Nevertheless, the warping functions inferred by these methods can suffer from abrupt changes around the landmarks position, leading to a disrupt PDF in the normalized image.

In this paper, we evaluate and compare different normalization approaches in the context of T2W-MRI prostate images normalization. Our contribution is threefold: (i) a normalization approach based on a Rician *a priori*; (ii) a normalization approach based on a method used in registration of functional data, without any assumption regarding the PDF of the data; (iii) a novel evaluation metric to asses quantitatively the alignment of the PDFs independently of the assumed distribution. These methods will be compared qualitatively and quantitatively, with both *z-score* normalization and piecewise-linear normalization.

2. METHODOLOGY

2.1 Normalization using Rician *a priori*

As stated in Sect. 1, proper normalization of the MRI data during pre-processing is a key problem that has been addressed using parametric and non-parametric strategies. We believe that normalizing MRI data using a parametric model based on a Rician distribution would improve the results for the parametric case. Expecting this improvement by changing the data model from the widely used Gaussian distribution to Rician distribution is reasonable. Indeed, Bernstein *et al.*¹¹ state that MRI data theoretically follows a Rayleigh distribution for a low Signal-to-Noise Ratio (SNR) scenarios while it appears closer to a Gaussian distribution when the SNR increases. Figure 1 shows the intensity spectrum for some MRI prostate data as well as the fitted Gaussian and Rician distributions. A qualitative assessment of the underlying distribution is performed by overlying the fitted distribution, while quantitative results of the fitting are given in terms of Root Mean Square. It can be highlighted that the Rician model better fits the data than the Gaussian model.

The normalization is carried out as: (i) Fit a Rician model to each prostate PDF using Least Squares minimization; (ii) Compute the mean (see Eq. (7)) and variance (see Eq. (3)) of the Rician model; (iii) Normalize the entire data using the *z-score* similarly as in Eq. (1).

$$\sigma \sqrt{\pi/2} L_{1/2}(-\nu^2/2\sigma^2) \quad (2)$$

$$2\sigma^2 + \nu^2 - \frac{\pi\sigma^2}{2} L_{1/2}^2\left(\frac{-\nu^2}{2\sigma^2}\right) \quad (3)$$

2.2 Normalization using generative models in functional data analysis

Srivastava *et al.* proposed a function registration framework which do not rely on any model regarding the functions themselves.

2.2.1 Square-Root Slope Function transform

The function registration is performed in a subspace using the Square-Root Slope Function (SRSF). The SRSF of a function f is defined as:

$$q(t) = \text{sign}(\dot{f}(t))\sqrt{|\dot{f}(t)|} . \quad (4)$$

The function f can be warped by a warping function γ , denoted $f \circ \gamma$ and can be expressed as in Eq. 5 using the SRSF.

$$\tilde{q}(t) = (q(t) \circ \gamma)\sqrt{\dot{\gamma}} . \quad (5)$$

Using these properties, a cost function can be defined and used inside the framework to align the different functions.

2.2.2 Cost function

The cost function used in the registration framework is expressed as in Eq. 6

$$D_y(f_1, f_2) = \inf_{\gamma \in \Gamma} \|q_1 - (q_2 \circ \gamma)\sqrt{\dot{\gamma}}\| . \quad (6)$$

2.2.3 Alignment

The alignment framework consists in the following steps:

- Compute u_f as in Eq. 7;
- For each function f_i , iteratively
 - Compute γ_i^* as in Eq. 8;
 - Compute \tilde{q}_i as in Eq. 5;
 - Update μ_f using the function \tilde{q}_i .

$$\mu_f = \arg \min_{f \in \mathcal{F}} \sum_{i=1}^n D_y(f, f_i)^2 . \quad (7)$$

$$\gamma_i^* = \arg \min_{f \in \mathcal{F}} \sum_{i=1}^n D_y(\mu_f, f_i)^2 . \quad (8)$$

3. EXPERIMENTS

Add the explanations regarding the piecewise-linear setup and percentiles. Add the implementation using the protocolclass.

The experiments are conducted on a subset of public multi-parametric MRI prostate dataset available at <http://visor.udg.edu/i2cvb/>.¹² This dataset was acquired from a cohort of patients with higher-than-normal level of PSA. The acquisition was performed using a 3T whole body MRI scanner (Siemens Magnetom Trio TIM, Erlangen, Germany) using sequences to obtain T2W-MRI. Aside of the MRI examinations, these patients also underwent a guided-biopsy. Finally, the dataset was composed of a total of 20 patients of which 18 patients had biopsy proven CaP and 2 patients were “healthy” with negative biopsies. In this study, our subset consists of 17 patients with CaP. The prostate organ as well as the prostate zones (i.e., PZ, Central Gland) and CaP were manually segmented by an experienced radiologist.

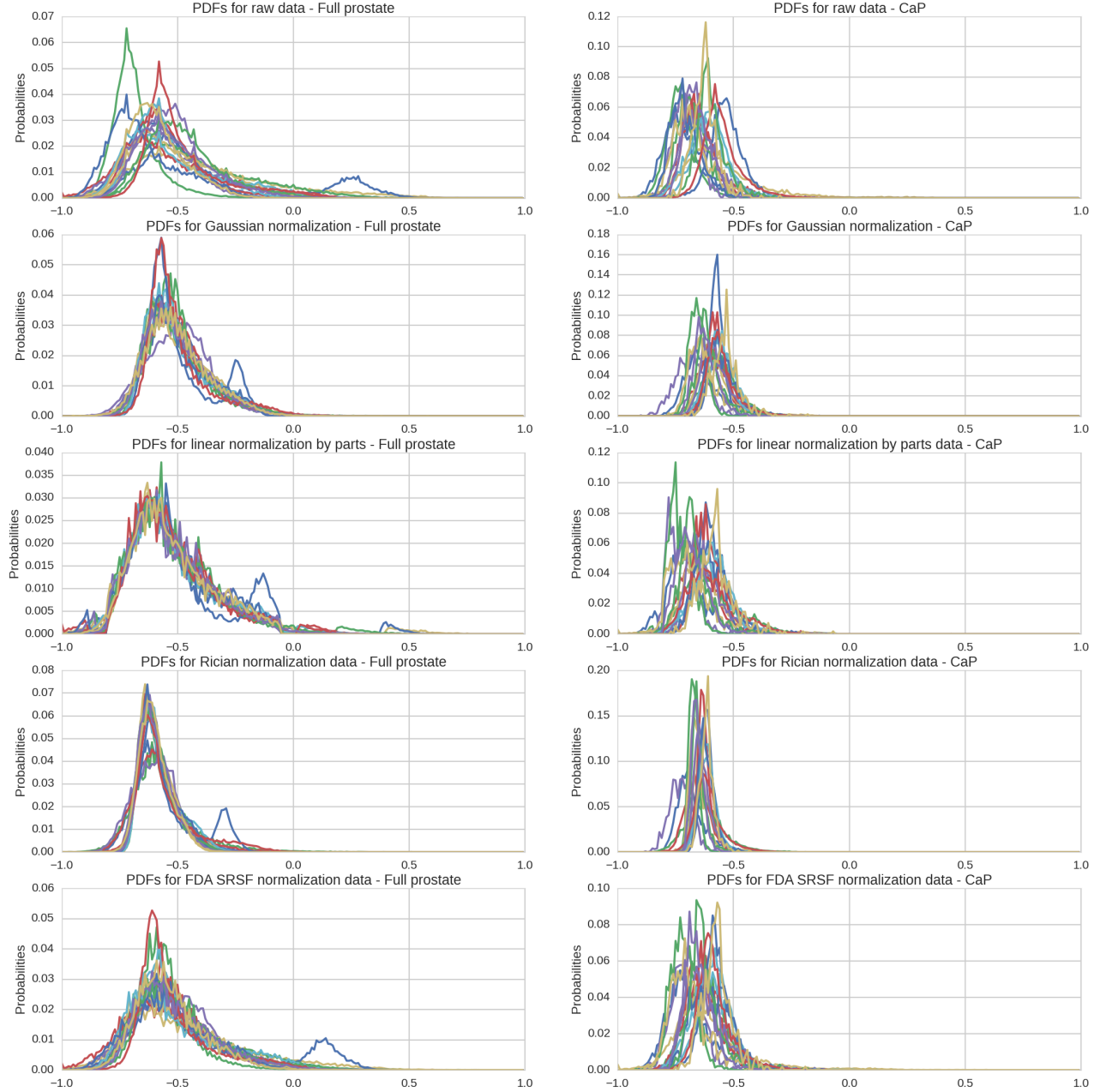


Figure 2: Qualitative evaluation by visual inspection of the alignment of the PDFs for the full prostate and the CaP.

4. RESULTS

4.1 Qualitative

Figure 2 depicts the alignment of the different PDFs using the different methods implemented. All the methods seem to address the problem of the PDF alignment of the full prostate data. However, the Rician normalization seems to outperform the other methods when focusing solely on the CaP data. The PDF computed in this specific area is more skewed from its original shape in the case of the linear normalization by parts than with the two other normalization strategies.

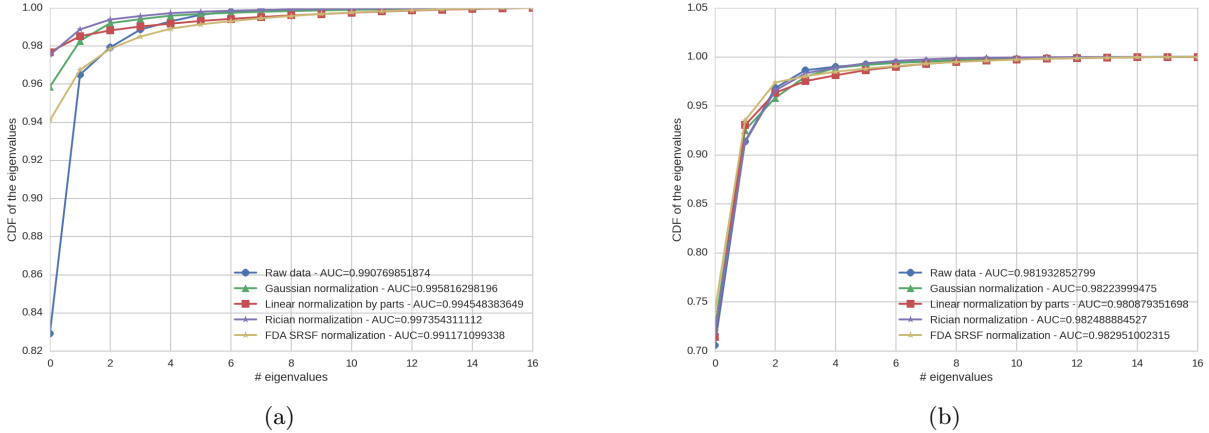


Figure 3: Spectral evaluation using PCA decomposition: (a) evaluation considering the full prostate, (b) evaluation considering only the CaP.

4.2 Quantitative

A spectral evaluation is performed by decomposing the set of normalized PDFs using PCA under the assumption that they are linearly dependent. Intuitively, the eigenvalues of the PCA decomposition are correlated with the alignment of the different PDFs. Thus, in the case of a perfect alignment of the PDFs, the first eigenvalue is much greater than the remaining since that the first eigenvector encodes all the information. In the contrary, in the case of a misalignment of the PDFs, more eigenvectors are needed to encode the information synonymous with larger eigenvalues. Thus, we propose to use the cumulative sum of the normalized eigenvalues as well as the Area Under this Curve (AUC), as depicted in Fig. 3. Rician normalization outperforms the other methods with an AUC of 0.9974 and 0.9824 considering the full prostate and CaP, respectively.

5. CONCLUSION AND FUTURE WORK

In this paper, we propose to normalize the T2W-MRI prostate images based on a Rician *a priori*. An extensive comparison was conducted showing that the Rician normalization outperforms the Gaussian and linear normalization by parts for T2W-MRI prostate images normalization.

Note. This work has not been submitted for publication or presentation elsewhere

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