

A boosting approach for prostate cancer detection using multi-parametric MRI

Guillaume Lemaître^{a,c} and Joan Massich^a and Robert Martí^c and Jordi Freixenet^c and Joan C. Vilanova^d and Paul M. Walker^b and Désiré D. Sidibé^a and Fabrice Mériaudeau^a

^aLE2I-UMR CNRS 6306, Université de Bourgogne, 12 rue de la Fonderie, 71200 Le Creusot, France;

^bLE2I-UMR CNRS 6306, Université de Bourgogne, Avenue Alain Savary, 21000 Dijon, France;

^cViCOROB, Universitat de Girona, Campus Montilivi, Edifici P4, 17071 Girona, Spain;

^dDepartment of Magnetic Resonance, Clinica Girona, Lorenzana 36, 17002 Girona, Spain

ABSTRACT

This document shows the desired format and appearance of a manuscript prepared for the Proceedings of the SPIE. It contains general formatting instructions and hints about how to use LaTeX. The LaTeX source file that produced this document, `article.tex` (Version 3.3), provides a template, used in conjunction with `spie.cls` (Version 3.3).

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1. INTRODUCTION

On a worldwide scale, prostate cancer (PCa) has been reported as the second most frequently diagnosed men cancers accounting for 13.6%.¹ Statistically, the estimated number of new diagnosed cases was 899,000 with no less than 258,100 estimated deaths.¹ In United States, aside from skin cancer, PCa was declared to be the most commonly diagnosed cancer among men, implying that around one in seven men will be diagnosed with PCa during their lifetime.²

Since its introduction in mid-1980s, prostate-specific antigen (PSA) is widely used for PCa screening³ and has shown to improve early detection of PCa.⁴ However, several trials conducted in Europe and United States conclude that PSA screening suffers from low specificity.⁵⁻⁷ Thus, current research focus on developing new screening methods to improve PCa detection. In this perspective, Magnetic resonance imaging (MRI) techniques have recently shown promising results for PCa detection. Furthermore, three different modalities are currently investigated: (i) T₂ Weighted (T₂-W) MRI, (ii) Dynamic Contrast-Enhanced (DCE) MRI and (iii) Diffusion Weighted (DW) MRI.

Several researches have been carried out in order to investigate the contributions of machine learning classifiers for PCa detection using the three aforementioned 3T multi-parametric MRI such as Support Vector Machines (SVM),⁸⁻¹² probabilistic boosting tree¹³ or probabilistic neural network.¹³ However, these studies use different datasets and evaluation statistics to report their results leading to an impossibility to give rise to a fair comparison.

In this research, we investigate the performance of gradient boosting for PCa detection using 3T multi-parametric MRI. Two different features extraction strategies have been chosen in order to feed the classifier: (i) voxel-based and (ii) 3D texton-based. An evaluation of both strategies as well as the contribution of each modality is provided. Furthermore, the dataset used for this experimentation is part of our future benchmarking platform I2CVB available at <http://visor.udg.edu/i2cvb/> and are available for future comparisons.

Further author information: (Send correspondence to G.L.)

G.L.: E-mail: guillaume.lemaitre@udg.edu

2. MATERIAL AND METHODS

2.1 Data

The multi-parametric MRI 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 T₂-W MRI, DCE MRI and DW MRI. Aside of the MRI examination, 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 PCa and 2 patients were “healthy” with negative biopsies. The prostate organ as well as the prostate zones (i.e., peripheral zone (PZ) and central gland (CG)) and PCa were manually segmented by an experienced radiologist. Therefore, 13 patients had a PCa in the PZ, 3 patients had PCa in the CG, 2 patients had invasive PCa in both PZ and CG and finally 3 patients were considered as “healthy”.

The Apparent Diffusion Coefficient (ADC) maps were computed using the scanner software and the DW MRI. The DCE MRI sequence consist in a kinetic study composed of 40 samples over time. These DCE MRI sequences and ADC maps were resampled using the spatial information of the T₂-W MRI sequence with dimensions of $448 \times 360 \times 64$ and voxel spacing of $0.68 \times 0.68 \times 1.25$ mm³. Linear interpolation was used to compute missing data during the up-sampling. The resampling was implemented in C++ using the Insight Segmentation and Registration Toolkit.¹⁴

2.2 Classification framework

2.2.1 Feature extraction strategies

Table 1. Overview of voxel features extracted in our classification framework.

Extraction strategy	Name	Size	Short description
Voxel-based	V _{T₂-W}	1	Intensity of a voxel in the T ₂ -W MRI
	V _{ADC}	1	Intensity of a voxel in the ADC map
	V _{DCE}	40	Intensities of a voxel along the whole serie in the DCE MRI
	V _{PZ}	1	Boolean value of a voxel membership to the PZ
	V _{CG}	1	Boolean value of a voxel membership to the CG
3D texton-based	T _{T₂-W}	243	Intensities vector for a window of $9 \times 9 \times 3$ voxels in the T ₂ -W MRI
	T _{ADC}	243	Intensities vector for a window of $9 \times 9 \times 3$ voxels in the ADC map
	T _{DCE}	9720	Intensities vector for a window of $9 \times 9 \times 3$ along the whole serie in the DCE MRI
	T _{PZ}	243	Boolean vector of voxels memberships to the PZ for a window of $9 \times 9 \times 3$
	T _{CG}	243	Boolean vector of voxels memberships to the CG for a window of $9 \times 9 \times 3$

A summary of the features extracted as well as the strategies chosen are summarized in Table 1. Two main strategies are applied to extract features. In the voxel-based approach, at each voxel location, the intensities for the different MRI modalities are extracted as well as the membership of this voxel to belong to the PZ or CG. The 3D texton-based approach extend this extraction for a 3D window of size $9 \times 9 \times 3$ around the central voxel. In both case, the vectors V(.) and T(.) extracted are scaled using min-max normalization.

Then, the different concatenation of the vectors V(.) and T(.) are summarized in Table 2. Different combinations are further tested in order to observe the contribution of each data feature.

Table 2. Overview of the different concatenations tested for the classification.

Name	V _{T2-W}	V _{ADC}	V _{DCE}	V _{PZ}	V _{CG}	T _{T2-W}	T _{ADC}	T _{DCE}	T _{PZ}	T _{CG}
V ₁	✗	✗	✓	✗	✗	✗	✗	✗	✗	✗
V ₂	✗	✓	✗	✗	✗	✗	✗	✗	✗	✗
V ₃	✗	✓	✓	✗	✗	✗	✗	✗	✗	✗
V ₄	✓	✗	✗	✗	✗	✗	✗	✗	✗	✗
V ₅	✓	✗	✓	✗	✗	✗	✗	✗	✗	✗
V ₆	✓	✓	✗	✗	✗	✗	✗	✗	✗	✗
V ₇	✓	✓	✓	✓	✓	✗	✗	✗	✗	✗
T ₁	✗	✗	✗	✗	✗	✗	✗	✓	✗	✗
T ₂	✗	✗	✗	✗	✗	✗	✓	✗	✗	✗
T ₃	✗	✗	✗	✗	✗	✗	✓	✓	✗	✗
T ₄	✗	✗	✗	✗	✗	✓	✗	✗	✗	✗
T ₅	✗	✗	✗	✗	✗	✓	✗	✓	✗	✗
T ₆	✗	✗	✗	✗	✗	✓	✓	✗	✗	✗
T ₇	✗	✗	✗	✗	✗	✓	✓	✓	✓	✓

2.2.2 Gradient boosting

In this research, we use a gradient boosting classifier¹⁵ originally proposed by Friedman^{16,17} in order to implement the computer-aided detection and diagnosis (CAD) for PCa detection. Gradient boosting is in fact a reformulation of the well-known AdaBoost¹⁸ in which boosting real-valued weak learners

2.2.3 Validation model

k-cross validation

3. RESULTS

Include the two figures with one for voxel-based and the other one for texton-based and discuss briefly.

4. DISCUSSION

Discuss the results.

- Single modality is not working as good as multi-parametric
- Which single modality is better.
- What is the increase of the zone information.
- Voxel-based vs texton-based

5. CONCLUSION

5.1 Future works

- Check the difference with other features usually extracted.
- Check the results difference with a registration of the three modalities.

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