

Bi-exponential modeling of prostate diffusion weighted MR imaging acquired using high b values: clinical evaluations of advanced post-processing methods

Parisa Movahedi^{1,2}, Harri Merisaari^{1,3}, Ileana Montoya Perez^{1,2}, Jussi Toivonen^{1,2}, Pekka Taimen⁴, Peter J. Boström⁵, Janne Verho¹, Hannu J. Aronen^{1,6}, Tapio Pahikkala², and Ivan Jambor^{1,6}

¹Department of Diagnostic Radiology, University of Turku, Turku, Finland, ²Department of Future Technologies, University of Turku, Turku, Finland, ³Turku PET Center, University of Turku, Turku, Finland, ⁴Department of Pathology, Turku University Hospital, Turku, Finland, ⁵Department of Urology, Turku University Hospital, Turku, Finland, ⁶Medical Imaging Center of Southwest Finland, Turku University Hospital, Turku, Finland

Synopsis

The aim of this study was to evaluate various mathematical methods for enhanced parameter estimation of bi-exponential DWI (12 b values 0-2000 s/mm²) of prostate cancer. Least Squares (LSQ), Bayesian Shrinkage (BS) and Maximum Penalized Likelihood Estimation (MPLE) fitting methods were evaluated in the terms of Coefficients of Variation (CV), Contrast to Noise Ratio (CNR) and the Area under the curve (AUC) between tumor and non-tumor prostate tissue. BS and MPLE methods improved AUC and CNR values of bi-exponential model parameters and also decreased CV values in comparison with the commonly used LSQ fitting method.

Introduction

Diffusion weighted MR imaging (DWI) is a non-invasive imaging technique that has extensively being used for the detection and characterization of prostate cancer (PCa) during the last decade.^{1,2} Voxel-wise signal fitting utilizing bi-exponential model has potential of extracting more information from the DWI signal than the commonly used monoexponential model.³ Bi-exponential model could provide parameter estimates with more clinically relevant information about the underlying tissue. Unfortunately this approach usually results in parametric maps that are corrupted by noise due to low signal-to-noise ratio (SNR) of the DWI. Recent developments in Bayesian bi-exponential model fitting^{4,5} have claimed to overcome this obstacle by denoising the parametric maps of the bi-exponential model utilizing the neighboring voxel information.

Materials and Methods

The study was approved by institutional review board and each patient with historically confirmed prostate cancer (PCa) included in the study provided written informed consent. The DWI was performed using a 3T MR scanner with a single shot spin-echo echo planar imaging, monopolar diffusion gradient scheme, and the following parameters: TR/TE 3141/51 ms, FOV 250 × 250 mm², slice thickness 5 mm, diffusion gradient timing (Δ) 24.5 ms, diffusion gradient duration (δ) 12.6 ms, diffusion time ($\Delta - \delta/3$) 20.3 ms, 12 b values of [number of signal averages]: 0 [2], 100 [2], 300 [2], 500 [2], 700 [2], 900 [2], 1100 [2], 1300 [2], 1500 [2], 1700 [3], 1900 [4], 2000 [4] s/mm². Prostate cancer lesions were delineated using whole mount prostatectomy sections as the “gold standard”. Voxel-wise fitting of bi-exponential model (f, D_f, D_s), eq.1,⁶ was performed by Least Squares fitting (LSQ), Bayesian Shrinkage (BS)⁴ and Maximum Penalized Likelihood estimation (MPLE)⁵ methods on whole prostate area including the lesions.

$$\mathbf{Y} = c(fe^{-bD_f} + (1-f)e^{-bD_s}) \quad (1)$$

where N is the number of b-values, signal vector $\mathbf{Y} = (y_n)_{n=1}^N$ is the obtained signal model values estimated from the bi-exponential model Eq.(1), $\mathbf{b} = (b_n)_{n=1}^N$ is the b value set, c is the signal without diffusion weighting. D_s is the slow diffusion component, D_f is the fast decay component and f is the fraction between fast and slow diffusion components. While LSQ method only estimates each voxel parameters individually, both BS and MPLE utilize spatial prior characteristics of surrounding voxels as a prior knowledge when fitting the model parameters for individual voxels. MPLE hyper parameter set (kernel size, constraints and the weighting penalty) were optimized while BS and LSQ did not required any prior parameter settings. The goodness of the methods' fit were evaluated using parametric maps and Fitting errors. Furthermore Coefficients of Variation (CV), Contrast to Noise Ratio (CNR) and the Area under the curve (AUC) between each tumor and non-tumor prostate region was calculated and compared in-between the methods.

Results

In total, 50 patients were included in final analyses. The tumor visibility was improved in both BS and MPLE for parametric maps (Figure1). The mean values of all estimated parameter values were lowest for BS and highest for LSQ (Figure2). The CV was significantly lower in all parameters' estimation with BS method compared to other two methods ($p < 0.05$) (see Figure 3.A). The BS and MPLE provided better CNR with the model parameter values D_f and D_s , compared with the LSQ and BS has significantly lower CNR for f parameter (Figure 3.B).

For classification performance between cancer and non-cancer tissue types when voxels of all 50 cases were pooled, the BS and MPLE method provided better AUC values than LSQ. The best classification performance was with MPLE method in D_f and D_s while classification performance for f parameter was best with the Bayesian Shrinkage (Table 1). The AUC values calculated for each subject separately are shown in Table 2. Significant differences between methods were found in f and D_f parameters. Classification performance of f was best with BS. For D_f and D_s the classification performance was better with BS and MPLE than with LSQ, where in D_s the difference was statistically significant.

Discussion

To the best of our knowledge, bi-exponential parametric map enhancement using BS and MPLE methods, have not yet been evaluated for their use in prostate cancer studies. Our results showed that spatially constraint bi-exponential methods offer possibilities for improved quantification of prostate DWI. BS method should be used with caution because while it globally optimizes the model parameters, the information might get lost if the tumor to prostate ratio is very small.

Conclusion

In conclusion, our results suggest that the presented advanced Bayesian based methods improves image contrast (CNR) and classification performance (AUC) in voxel-wise level of bi-exponential model which may lead to better detection and characterization of prostate cancer.

Acknowledgements

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Figures

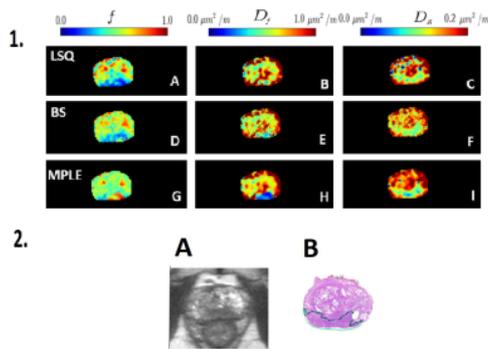


Figure 1: 1. Representative voxel-wise fitting results of a prostate with tumor area from clinical dataset. Parametric maps of estimated parameters are from LSQ (A-C), BS (D-F) and MPLE (G-I). 2. Corresponding T2-weighted image, with tumor in the peripheral zone (A) and the histological section with tumor delineation (B).

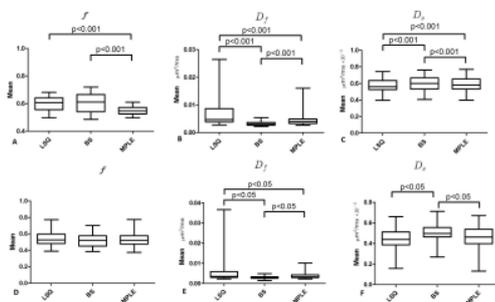


Figure 2: Box plots of the mean parameter values for both non-cancer prostate tissue (A-C) and tumor area (D-F) of 50 patients with histologically confirmed prostate cancer (clinical dataset).

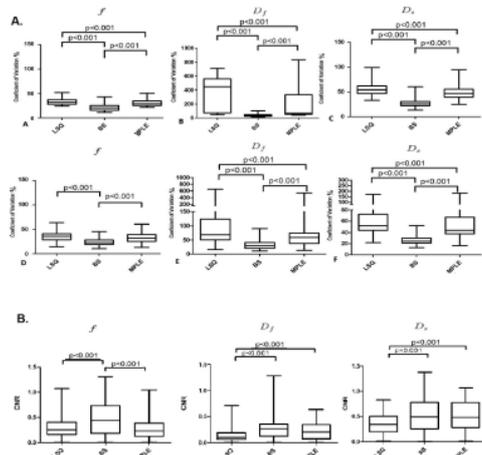


Figure 3: A. Box plots for Coefficient of Variation (CV) calculated for each parameter of bi-exponential model with LSQ, BS and MPLE for both non-tumor (A-C) and tumor (D-F) ROIs of 50 patients with histologically confirmed prostate cancer. B. Mean Contrast-to-Noise Ratio (CNR, between tumor and prostate) of all 50 cases from clinical DWI dataset, for each estimated parameter, f , D_f , D_s of bi-exponential model for BS, MPLE and LSQ.

AUC of all pooled voxels	f^*	D_f^{**}	D_s^*
LSQ	0.61 (0.612-0.626)	0.52 (0.512-0.527)	0.62 (0.617-0.629)
BS	0.70 (0.697-0.709)	0.53 (0.518-0.533)	0.66 (0.650-0.663)
MPLE	0.57 (0.529-0.601)	0.59 (0.585-0.600)	0.70 (0.697-0.708)

Table 1: AUC= Area under the curve values for classification of voxels between cancer and non-cancer tissue type inside prostate region for bi-exponential parameters. 95% confidence intervals are shown in brackets. Range of the AUC values over subjects is shown in brackets. * Difference between all methods is statistically significant ($p < 0.001$). ** Difference between MPLE and other methods is statistically significant ($p < 0.001$).

Median AUC of all cases	f	D_f	D_s
LSQ	0.60 (0.50-0.88)	0.57 (0.51-0.75)	0.62** (0.51-0.81)
BS	0.66* (0.51-0.95)	0.58 (0.50-0.83)	0.66 (0.51-0.90)
MPLE	0.59 (0.50-0.84)	0.60 (0.50-0.85)	0.68 (0.50-0.92)

Table 2: AUC= Medians of area under the curve values of 50 subjects for bi-exponential parameters. Each AUC was calculated for classification of voxels between cancer and non-cancer tissue type inside prostate region. Range of the AUC values over subjects is shown in brackets. * In f , BS difference to other methods is statistically significant ($p < 0.05$). ** In D_s , LSQ difference to other methods is statistically significant ($p < 0.05$).