

# Clinical Assessment of B1+ Inhomogeneity Effects on Quantitative Prostate MRI at 3.0 T

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**Target Audience:** Clinicians and physicists interested in quantitative multi-parametric MRI for prostate cancer.

**Purpose:** Dynamic contrast-enhanced MRI (DCE-MRI) has an important role in multi-parametric MRI for prostate cancer detection and characterization. A primary step in the pharmacokinetic modeling analysis includes conversion of signal intensity curves into contrast agent concentration curves, which requires knowledge of the pre-contrast  $T_1$  ( $T_{10}$ ) values. Variable flip angle (VFA) imaging is commonly used for  $T_{10}$  measurements since it can provide fast volumetric  $T_1$  mapping [1], however, it heavily depends on the set of flip angles used, and therefore is sensitive to any flip angle variation. Transmit RF ( $B_1^+$ ) field inhomogeneity creates the flip angle variation, which tends to be 30 - 50% across the abdomen at 3T [2]. In this work, we measure  $B_1^+$  variation in the pelvis using the reference region VFA method [3] and evaluate the impact on the estimated  $T_1$  values in prostate and pelvic muscle with and without compensating for  $B_1^+$  variation in a total of 108 prostate patients at 3T.

**Methods:** Experiments were performed on two 3.0T systems (TIM Trio and Skyra, Siemens) in a total of 108 men undergoing clinically indicated prostate MRI, ranging in age between 48 and 87 years (age =  $65.4 \pm 7.8$  years and mass =  $82.0 \pm 13.5$  kg). Local IRB and informed consent was obtained. All MRI scans were feet first. A body coil was used for RF transmission, and the automatic pre-scan was used to calibrate RF transmission. A circular polarization (CP) mode was used for Trio (n = 18), and an elliptical polarization (EP) mode was used for Skyra (n = 90).

A 3D SPGR sequence with a dual-echo bipolar readout was used and TEs were chosen to be in- and out-of-phase (TE1 = 1.23ms and TE2 = 2.46ms) at 3T. A 2-point Dixon fat-water separation algorithm was used to generate fat- and water-only images [4], and B1+ maps were further calculated using RR-VFA [3] in prostate and in-house software (Matlab R2013, Osirix 5.9, Pixmeo SARL).

Maps of relative flip angle (rFA) as a percentage were calculated by dividing the actual flip angle by the nominal flip angle, and  $T_1$  maps were calculated with and without FA correction. Regions of Interest (ROI) were manually drawn on the prostate, left and right muscle in the pelvis, and we computed rFA and  $T_1$  before and after FA correction, averaged over these ROIs (Fig 1). Paired t-tests and f-tests were performed to compare the means and standard deviations of rFA and  $T_1$ .

**Results and Discussion:** Qualitative assessment (Fig 1) showed that  $T_1$  estimation was improved following FA correction.

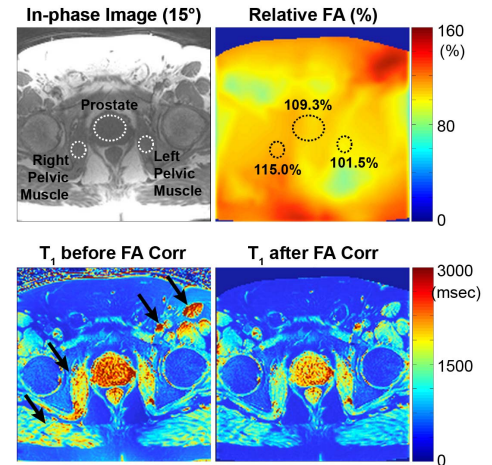
Statistical results in different tissues from 108 prostate patients are shown in Table 1. The average rFA was  $104.5 \pm 8.0\%$  in the prostate, ranging from 70.2% to 121.6%, which indicated that the FA correction is crucial to achieve more accurate quantitative results. The difference of mean  $T_1$  between the left and right pelvic muscle was 143ms before FA correction and was reduced to 27ms after the FA correction; the standard deviations of all  $T_1$  estimates were reduced by half after FA correction, resulting in a more uniform  $T_1$  value for each tissue ( $p = 9.77 \times 10^{-9}$ ). The reduction in these values gives more confidence in  $T_1$  correction results.

The average estimated  $T_1$  value in prostate is reduced from 2422ms to 2209ms with FA correction, which may be a more accurate estimation in prostate  $T_1$ .

The rFA and  $T_1$  values were also compared between two RF transmission modes: EP mode - Skyra and CP mode - Trio (Fig 2). From the box plots, a significant difference of  $B_1^+$  characteristics can be clearly seen. Skyra has relatively higher rFA while Trio has lower rFA, which leads to inconsistency between scanners in estimating  $T_1$  values. The average value of prostate  $T_1$  between scanners differed by 601ms, and the difference was significantly reduced to 65ms after FA correction. Paired t-tests were applied to data between different scanners, and the p value decreased from  $1.22 \times 10^{-5}$  to  $2.78 \times 10^{-1}$  after correction, showing a more uniform prostate  $T_1$  value after FA correction. Since an important application for quantitative imaging is to compare results from different scanners during follow-up studies, FA correction is necessary for accurate  $T_1$  estimation.

**Conclusion:** We have measured  $B_1^+$  inhomogeneity and the  $T_1$  value in 108 prostate MRI exams and have shown improved  $T_1$  measurements in both prostate and pelvic muscles. The FA correction provides a more uniform  $T_1$  value for each tissue and significantly reduces the inconsistency between different scanners. Since the RR-VFA method does not require additional scan time, it is recommended to use RR-VFA to correct  $T_1$  mapping for 3T prostate MRI.

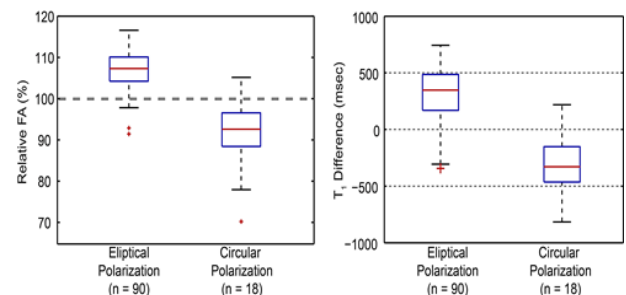
**References:** [1] Deoni et al., MRM 2003. [2] Azian et al., JMRI 2010. [3] Sung, et al., MRM 2013. [4] Ma et al. MRM. 2004.



**Fig 1:** A representative example of inaccuracies of  $T_1$  measurements due to  $B_1^+$  inhomogeneity.

**Table 1:** relative FA values and  $T_1$  values obtained from prostate and both pelvic muscles

|                     | rFA (%)         | $T_1$ before FA corr (msec) | $T_1$ after FA corr (msec) |
|---------------------|-----------------|-----------------------------|----------------------------|
| Prostate            | $104.5 \pm 8.1$ | $2421.9 \pm 411.3$          | $2209.3 \pm 232.6$         |
| Left Pelvic Muscle  | $106.3 \pm 6.7$ | $1701.1 \pm 206.6$          | $1506.2 \pm 103.7$         |
| Right Pelvic Muscle | $110.7 \pm 7.2$ | $1834.5 \pm 267.2$          | $1479.2 \pm 105.2$         |



**Fig 2:** Comparison of relative FA (a) and difference of prostate  $T_1$  (b) between two types of RF transmission.