

**WORKSHOP PRESENTATION**

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# Motion correction for free breathing quantitative myocardial $t_2$ mapping: impact on reproducibility and spatial variability

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## Background

Quantitative myocardial  $T_2$  mapping is a promising technique for *in-vivo* assessment of inflammation and edema [1]. Free breathing  $T_2$  mapping sequences increase the flexibility in the choice of the number of  $T_2$ prep echoes times ( $TE_{T_2P}$ ), but should be combined with respiratory motion correction technique [2]. In this study, we sought to evaluate the performance of the Adaptive Registration of varying Contrast-weighted images for improved Tissue Characterization (ARCTIC) algorithm [3] for in-plane motion correction in  $T_2$  mapping data and its impact on in-vivo reproducibility and spatial variability of myocardial  $T_2$  estimates.

## Methods

Seven healthy adult subjects ( $30 \pm 17$  y, 3 male) were imaged using a 1.5 T Phillips scanner.  $T_2$  mapping was performed using either 1) a " $T_{2P}4TE$ " sequence (4  $T_2$ prep echo times=[0, 25, 50,  $\infty$ ]), or 2) a " $T_{2P}20TE$ " sequence (20  $T_2$ prep echo times=[0, 25, 30, 35, ..., 95, 100,  $\infty$ ,  $\infty$ ,  $\infty$ ]) [4].  $TE_{T_2P}=\infty$  was simulated by acquiring an image immediately after a saturation pulse [4]. Each subject was imaged using eight  $T_2$  mapping scans in the following order: 1) breath-held  $T_{2P}4TE$  (BH), 2) free breathing  $T_{2P}4TE$  without respiratory navigator (FB), 3) free breathing  $T_{2P}4TE$  with respiratory navigator (FB+NAV), and 4) free breathing  $T_{2P}20TE$  with respiratory navigator (5 repetitions). The same 2D short axis slice was acquired with all scans using single-shot ECG-triggered acquisitions with balanced SSFP imaging read-out ( $TR/TE/\alpha=2.7\text{ms}/1.35\text{ms}/85^\circ$ ,  $FOV=240 \times 240\text{mm}^2$ ,

resolution= $2.5 \times 2.5 \times 8\text{mm}^3$ , 10 linear ramp-up pulses, SENSE rate=2, 51 phase encoding lines, linear ordering). Accuracy of in-plane motion correction was evaluated in the first three scans by measurements of the DICE similarity coefficients (DSC) (1: ideal registration, 0: none) and the myocardial boundary error (MBE) with and without using ARCTIC.  $T_2$  mapping reproducibility and spatial variability with and without using ARCTIC was evaluated over the entire myocardium using the 5 repetitions of the  $T_{2P}20TE$  sequence and 1) a subset of 4  $T_2$ prep echo times=[0ms, 25ms, 50ms,  $\infty$ ] (referred to as 4TE) and 2) all 20  $T_2$ prep echo times (referred to as 20TE).

## Results

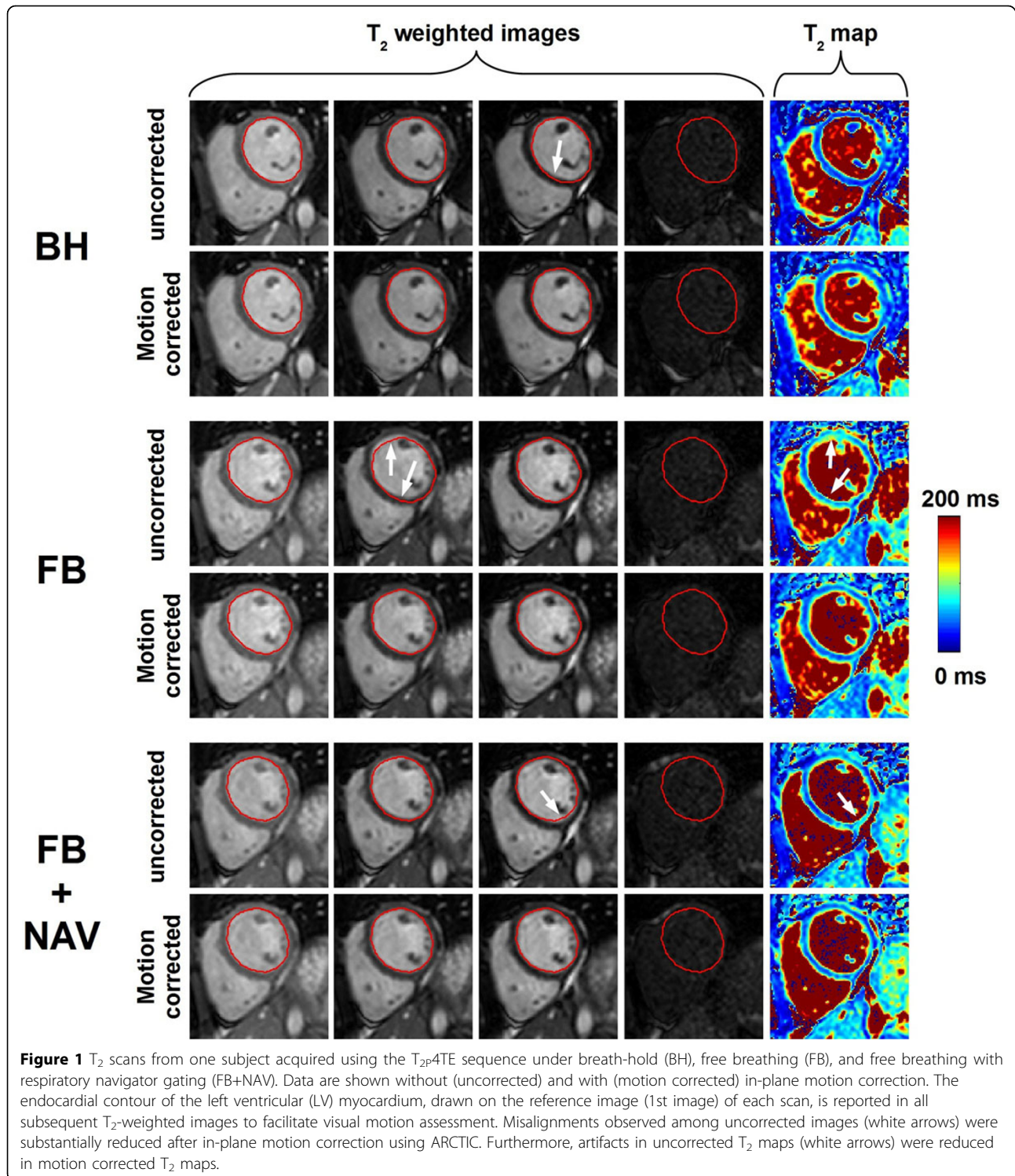
ARCTIC increased DSC in BH data ( $0.90 \pm 0.02$  vs.  $0.87 \pm 0.05$ ,  $p=0.09$ ), FB data ( $0.91 \pm 0.02$  vs.  $0.79 \pm 0.15$ ,  $p=0.009$ ), and FB+NAV data ( $0.90 \pm 0.02$  vs.  $0.86 \pm 0.08$ ,  $p=0.039$ ), and reduced MBE in BH data ( $0.63 \pm 0.09$  vs.  $0.74 \pm 0.12$ ,  $p=0.049$ ), FB data ( $0.60 \pm 0.12$  vs.  $1.16 \pm 0.71$ ,  $p=0.007$ ), and FB+NAV data ( $0.61 \pm 0.13$  vs.  $0.83 \pm 0.28$ ,  $p=0.025$ ). ARCTIC improved the reproducibility (4TE:  $5.0 \pm 2.3\text{ms}$  vs.  $5.9 \pm 3.1\text{ms}$ ,  $p=0.011$ ; 20TE:  $2.4 \pm 1.0\text{ms}$  vs.  $4.3 \pm 3.9\text{ms}$ ,  $p=0.002$ ) and reduced spatial variability (4TE:  $11.1 \pm 3.6\text{ms}$  vs.  $13.7 \pm 4.3\text{ms}$ ,  $p<0.001$ ; 20TE:  $7.9 \pm 1.8\text{ms}$  vs.  $10.6 \pm 5.3\text{ms}$ ,  $p=0.001$ ) of in-vivo  $T_2$  mapping.

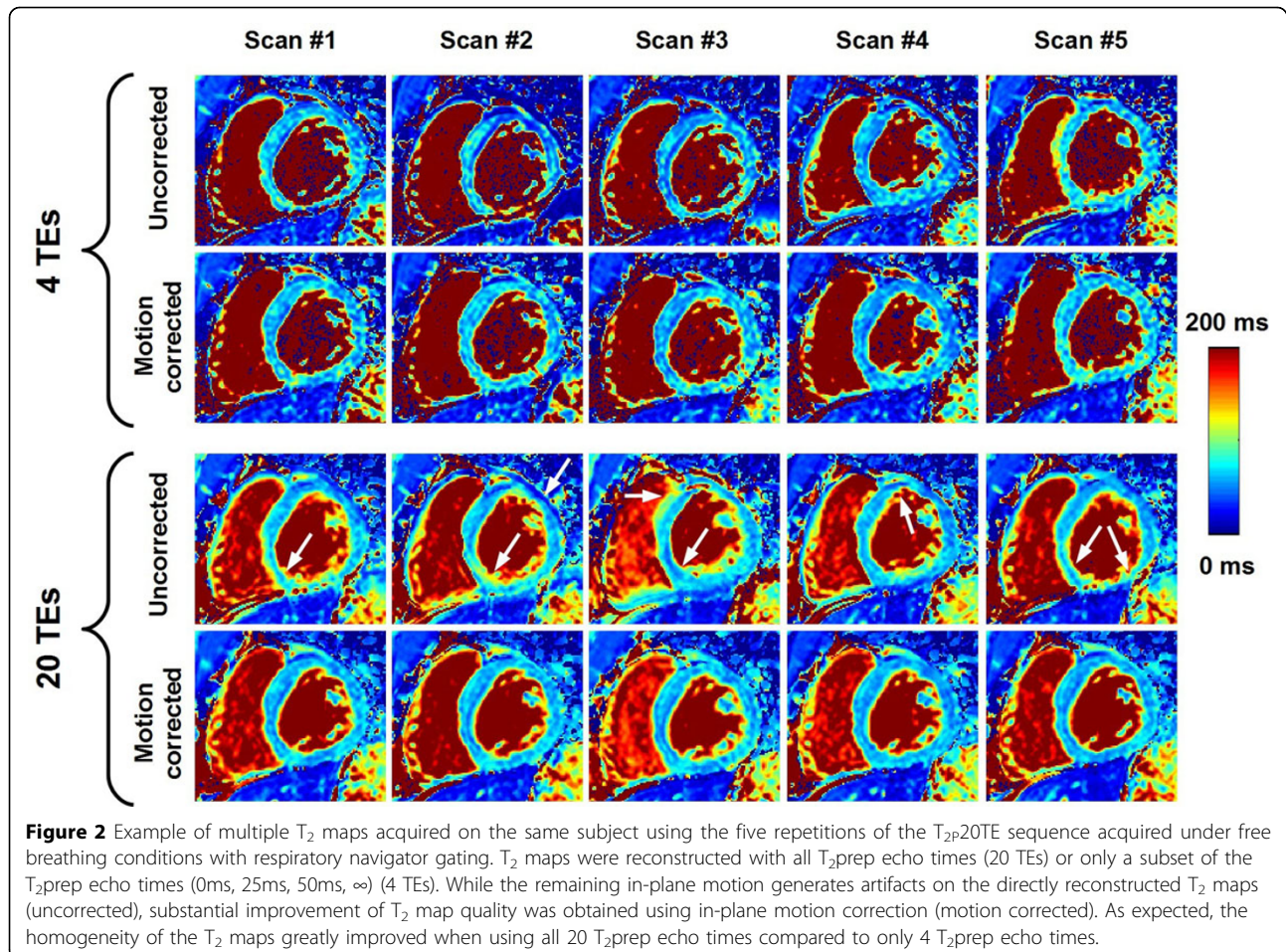
## Conclusions

The ARCTIC technique substantially reduces spatial mis-alignment among  $T_2$ -weighted images and improves both the reproducibility and the spatial variability of in-vivo  $T_2$  mapping.

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