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Impact of motion correction on reproducibility and spatial variability of quantitative myocardial T₂ mapping

Sébastien Roujol¹, Tamer A. Basha^{1,2}, Sebastian Weingärtner^{1,3}, Mehmet Akçakaya¹, Sophie Berg¹, Warren J. Manning^{1,4} and Reza Nezafat^{1*}

Abstract

Background: To evaluate and quantify the impact of a novel image-based motion correction technique in myocardial T₂ mapping in terms of measurement reproducibility and spatial variability.

Methods: Twelve healthy adult subjects were imaged using breath-hold (BH), free breathing (FB), and free breathing with respiratory navigator gating (FB + NAV) myocardial T₂ mapping sequences. Fifty patients referred for clinical CMR were imaged using the FB + NAV sequence. All sequences used a T₂ prepared (T₂prep) steady-state free precession acquisition. In-plane myocardial motion was corrected using an adaptive registration of varying contrast-weighted images for improved tissue characterization (ARCTIC). DICE similarity coefficient (DSC) and myocardial boundary errors (MBE) were measured to quantify the motion estimation accuracy in healthy subjects. T₂ mapping reproducibility and spatial variability were evaluated in healthy subjects using 5 repetitions of the FB + NAV sequence with either 4 or 20 T₂prep echo times (TE). Subjective T₂ map quality was assessed in patients by an experienced reader using a 4-point scale (1-non diagnostic, 4-excellent).

Results: ARCTIC led to increased DSC in BH data (0.85 ± 0.08 vs. 0.90 ± 0.02 , $p = 0.007$), FB data (0.78 ± 0.13 vs. 0.90 ± 0.21 , $p < 0.001$), and FB + NAV data (0.86 ± 0.05 vs. 0.90 ± 0.02 , $p = 0.002$), and reduced MBE in BH data (0.90 ± 0.40 vs. 0.64 ± 0.19 mm, $p = 0.005$), FB data (1.21 ± 0.65 vs. 0.63 ± 0.10 mm, $p < 0.001$), and FB + NAV data (0.81 ± 0.21 vs. 0.63 ± 0.08 mm, $p < 0.001$). Improved reproducibility (4TE: 5.3 ± 2.5 ms vs. 4.0 ± 1.5 ms, $p = 0.016$; 20TE: 3.9 ± 2.3 ms vs. 2.2 ± 0.5 ms, $p = 0.002$), reduced spatial variability (4TE: 12.8 ± 3.5 ms vs. 10.3 ± 2.5 ms, $p < 0.001$; 20TE: 9.7 ± 3.5 ms vs. 7.5 ± 1.4 ms) and improved subjective score of T₂ map quality (3.43 ± 0.79 vs. 3.69 ± 0.55 , $p < 0.001$) were obtained using ARCTIC.

Conclusions: The ARCTIC technique substantially reduces spatial mis-alignment among T₂-weighted images and improves the reproducibility and spatial variability of in-vivo T₂ mapping.

Keywords: Motion correction, Image registration, Quantitative myocardial tissue characterization, Myocardial T₂ mapping

* Correspondence: rnezafat@bidmc.harvard.edu

¹Department of Medicine (Cardiovascular Division), Beth Israel Deaconess Medical Center, 330 Brookline Ave, Boston, MA 02215, USA

Full list of author information is available at the end of the article

Background

The T_2 relaxation time is dependent on the amount of free water [1] and can be exploited as a potential marker of inflammation and edema [2–7]. In cardiac MR (CMR), T_2 changes are generally assessed using a dark blood T_2 -weighted acquisition [8]. Elevated signal intensity in T_2 -weighted images have been reported in presence of several cardiomyopathies such as myocarditis [2, 3], Tako-Tsubo [4], and acute myocardial infarction [5–7]. However, this technique only provides qualitative measurements and image interpretation can be limited by several factors including regional signal variations induced by phased array coil, elevated signal induced by sub-endocardial stagnant blood, and signal loss caused by through-plane motion [9, 10].

Quantitative myocardial T_2 mapping [11, 12] is an alternative technique, which shows promise for reducing uncertainties in interpretations of dark blood T_2 -weighted images. In this technique, several T_2 -weighted images are acquired, each with a different T_2 contrast. The signal intensity obtained from the T_2 -weighted images is then fit to a physical model of T_2 signal decay on a per-pixel basis, leading to the creation of a T_2 map. The acquisition of each T_2 -weighted image was initially performed using either spin echo/fast spin echo acquisitions [11–14] with varying echo times (TE) which results in very long scan time. Recently, T_2 -prepared (T_2 prep) [15] steady-state free precession (SSFP) acquisitions have been proposed and provide higher imaging efficiency [16]. These sequences can be acquired within a breath-hold [16, 17] or under free breathing conditions with respiratory motion correction techniques [16, 18, 19].

Despite the promise of this technique, its in-vivo reproducibility and precision have not been fully characterized. These two factors play a major role for clinical acceptance of any quantitative myocardial tissue characterization technique [20, 21]. The presence of motion among T_2 -weighted images is one of the main challenges in T_2 mapping and is expected to have important impact on the technique precision and reproducibility.

Breath-hold acquisitions can be used to reduce the impact of respiratory motion. However, some motion can still be detected in 40–60 % of patients due to their limited breath-holding capabilities, as reported by several T_1 mapping studies using breath-held acquisitions of ~11–17 heart beats [22–25]. The breath-hold approach imposes severe time limitations on the number of acquired T_2 -weighted images (typically ~3–4) since a rest time of ~4–6 heart beats is required between each acquisition to allow for full longitudinal magnetization recovery. Therefore, the use of a free breathing acquisition is attractive as it enables the acquisition of a larger number of T_2 -weighted images which may be beneficial to improve precision and reproducibility. On the other hand,

free breathing acquisitions require the use of respiratory navigators to account for through plane motion and image registration algorithms to correct for residual in-plane motion [18].

We recently developed a technique for Adaptive Registration of varying Contrast-weighted images for improved Tissue Characterization (ARCTIC) which we have evaluated for myocardial T_1 mapping [23]. In this study, we sought to investigate the performance of ARCTIC for T_2 mapping and its impact on in-vivo reproducibility and spatial variability of myocardial T_2 estimates.

Methods

All subjects were scanned using a 1.5 T Philips Achieva (Philips Healthcare, Best, The Netherlands) scanner with a 32-channel cardiac phased array receiver coil. This study was health insurance portability and accountability act (HIPAA) compliant and the imaging protocol was approved by our institutional review board (Committee on Clinical Investigations (CCI)) at the Beth Israel Deaconess Medical Center. Written informed consent was obtained from each participant.

T_2 mapping acquisition scheme

T_2 mapping was performed using our recently reported T_2 mapping sequence [26] in which multiple T_2 -weighted images are acquired using an electrocardiogram (ECG)-triggered T_2 prep steady-state free precession (SSFP) acquisition with different T_2 prep echo times (TE_{T_2P}). A rest cycle of 6 s was used between the acquisitions of two successive T_2 -weighted images to ensure full re-growth of the longitudinal magnetization. The $TE_{T_2P} = 0$ image was acquired using 90° pulse followed immediately by a -90° pulse and a crusher gradient to ensure consistency with all other images in term of longitudinal signal reduction induced by imperfect 90° and -90° flip angles used in the T_2 prep. Finally, to model the signal re-growth induced by the SSFP imaging pulses, an infinitely long T_2 prep echo time ($TE_{T_2P} = \infty$) was simulated by acquiring an image immediately after a saturation pulse. In this study, this sequence has been evaluated with 4 T_2 prep echo times ($T_{2P}4TE$: 0, 25, 50, ∞) and 20 T_2 prep echo times ($T_{2P}20TE$: 0, 25, 30, 35, ..., 95, 100, ∞ , ∞ , ∞). For free breathing acquisitions, a respiratory navigator positioned immediately prior to the T_2 prep was used for end expiratory gating (window size = 5 mm). No T_2 prep or imaging pulses were applied if the navigator signal was outside the gating window to enable the acquisition of undisturbed signal in the next heartbeat.

In-plane motion correction

The ARCTIC approach was used to compensate for in-plane motion between T_2 -weighted images [23]. In this

approach, all images are registered individually to a common reference image, which was chosen as the first image of the series ($TE_{T2P} = 0$). The motion was then estimated in a two-step process. Affine motion descriptors are first estimated over a region of interest surrounding the heart. This global transformation is then provided as input of a more sophisticated local non-rigid motion estimation step using an extended formulation of the optical flow problem which enables the simultaneous estimation of both motion field and intensity variations on a per-pixel basis [27]. An additional term is used to constrain the motion estimates based on prior automatic tracking of specific feature points in the images [28–30]. In this algorithm, both motion field and intensity variation map are solved using an iterative scheme. A multi-resolution approach was used for the local non-rigid motion estimation step where the optical flow is initially estimated from first sub-resolution images and then refined using the full resolution images [31]. For each resolution level, the iterative scheme used 100 iterations and was repeated fifty times. These parameters were empirically optimized in this study. Since optical flow algorithms are well suitable for parallelization on graphic processing unit (GPU) [32–34], a GPU implementation of the method was used based on the compute unified device architecture (CUDA). More details about the algorithm can be found in [23].

T₂ map reconstruction

T₂ maps were reconstructed offline using a 3-parameter curve fitting model.

$$S(A, B, T_2, T_n) = Ae^{-t_n/T_2} + B. \quad (1)$$

where t_n is the T₂prep echo time of n^{th} T₂-weighted image, and A, B, and T₂ are the model parameters. A, B, and T₂ are estimated independently for each pixel using a Levenberg-Marquard optimizer with the online library provided in [35].

In-vivo study in healthy subjects

Twelve healthy adult subjects (32 ± 16 years, 6 male) without any history of cardiovascular disease underwent CMR examination. Each subject was imaged using eight T₂ mapping sequences in the following order:

1. Breath-held T_{2p}4TE
2. Free breathing T_{2p}4TE *without* respiratory navigator
3. Free breathing T_{2p}4TE *with* respiratory navigator
4. Free breathing T_{2p}20TE *with* respiratory navigator (5 repetitions).

All sequences were acquired in the short axis view using a single-shot ECG-triggered acquisition with SSFP

imaging readout and the following parameters: field of view = 240×240 mm², in-plane resolution = 2.5×2.5 mm², slice thickness = 8 mm, TR/TE = 2.7 ms/1.35 ms, flip angle = 85°, 10 linear ramp-up pulses, SENSE rate = 2, acquisition window = 138 ms, number of phase encoding lines = 51, linear k-space ordering. All T₂ scans were acquired in the same short axis orientation at the mid-diastolic cardiac phase using one single mid-ventricular slice.

Accuracy of motion correction was evaluated in the first three scans (T_{2p}4TE) by quantifying the motion between the T₂-weighted images without (uncorrected) and with in-plane motion correction using ARCTIC (motion corrected). Endocardial and epicardial contours were manually drawn in all T₂-weighted images of all T₂ mapping scans. The two contours were used to create a binary representation of the myocardium for each T₂-weighted image. The DICE similarity coefficient (DSC) [36] was then calculated between the myocardial binary mask of the reference image (M_{ref}) and the myocardial binary mask of each k^{th} T₂-weighted image (M_k) as follows:

$$\text{DSC} = \frac{2 \times \text{area}(M_{\text{ref}} \cap M_k)}{\text{area}(M_{\text{ref}}) + \text{area}(M_k)} \quad (2)$$

The myocardial boundary error (MBE), which provides a local alignment measure is also reported. MBE was measured as the average distance between the myocardial boundary of each T₂-weighted image (boundary of M_k) and the myocardial boundary of the reference image (boundary of M_{ref}) as follows:

$$\text{MBE}(M_k, M_{\text{ref}}) = \frac{1}{N} \sum_{i=1}^N \left\| P_{M_k}^i - P_{M_{\text{ref}}}^{\text{Closest}-i} \right\|_2 \quad (3)$$

Where $P_{M_k}^i$ is the i^{th} point along the boundary of M_k , $P_{M_{\text{ref}}}^{\text{Closest}-i}$ is the closest point of $P_{M_k}^i$ located on the boundary of M_{ref} . Since $TE_{T2P} = \infty$ images are very low signal-to-noise ratio (SNR) images, which makes the detection of the myocardial borders very difficult, no DSC/MBE were measured in those images. The statistical significant difference between DSCs (and MBEs) obtained with and without motion correction was evaluated using Wilcoxon signed rank tests. Statistical significance was considered at $p < 0.05$.

The impact of in-plane motion correction on the reproducibility and spatial variability of T₂ mapping was evaluated using the five T_{2p}20TE scans. For each scan, T₂ maps were reconstructed without (uncorrected) and with prior in-plane motion correction using ARCTIC (motion corrected). The endocardial and epicardial border of the myocardium and the insertion point were manually drawn on each T₂ map. A six myocardial segment model [37] was automatically created for each single slice (1:anterior, 2:anterospetal, 3:inferospetal,

4:inferior, 5:inferolateral, 6:anterolateral). Segment-based analysis of reproducibility and spatial variability of T_2 estimates was then performed. Spatial variability was defined as the standard deviation of T_2 estimates over a given segment. Reproducibility was defined as the standard deviation over the 5 scans of the spatial average T_2 values in one given segment. Both reproducibility and spatial variability are reported in average over all segments for each subject, and in average over all subjects for each segment. To investigate the motion influence in T_2 mapping sequences using a limited number of T_2 prep echo times, this overall analysis was repeated using a subset of the T_2 -weighted images from each scan (4 T_2 prep echo times of 0, 25, 50, ∞). The statistical significant difference between uncorrected and ARCTIC motion corrected T_2 reproducibility (and spatial variability) measured for each subject (in average over all myocardial segments) was evaluated using Wilcoxon signed rank tests.

In-vivo study in patients

Fifty patients referred for clinical CMR (56 ± 14 y, 29 male) were imaged using the free breathing $T_{2p}4TE$ T_2 mapping sequence with respiratory navigator. All sequences were acquired in the short axis view using a single-shot ECG-triggered acquisition with SSFP imaging readout and the following parameters: field of view = 360×360 mm², in-plane resolution = 2×2 mm², slice thickness = 8 mm, slice number = 3, TR/TE = 2.9 ms/1.45 ms, flip angle = 85°, 10 linear ramp-up pulses, SENSE rate = 2, acquisition window = 270 ms, number of phase encoding lines = 93, linear k-space ordering. T_2 maps were reconstructed without and with ARCTIC motion correction.

A subjective qualitative analysis was performed by an experienced cardiologist. The initial motion level in uncorrected data was assessed for each slice as “no motion”, “small motion”, or “large motion” by visual inspection of all uncorrected T_2 -weighted images. Subjective assessment of uncorrected and motion correction T_2 maps (150 T_2 maps) followed. Each pair of uncorrected and motion correction T_2 maps were shown simultaneously to the reader side by side in a random order. The reader was blinded to the reconstruction approach (uncorrected vs. motion corrected). Each map was assessed in term of overall quality (1-non diagnostic/large artifacts/no confidence in interpreting T_2 values in more than half of the myocardial segments, 2-fair/moderate artifacts/confidence in interpreting T_2 values in more than half of the myocardial segments, 3-good/small motion artifacts/no confidence in interpreting T_2 values in at most one myocardial segment, 4-excellent/no motion artifact/confidence in interpreting T_2 values in all myocardial segments). Furthermore, for each pair of T_2 maps, the reader was asked

to evaluate if any of the two T_2 map had “1-inferior”, “2-similar”, or “3-superior” quality. Wilcoxon signed rank test was used to test the null hypothesis that the difference of overall T_2 map quality scores between uncorrected and motion corrected T_2 maps was zero. Statistical significance was considered at $p < 0.05$.

Results

All scans were successful. The nominal scan time (assuming 100% gating efficiency) corresponded to 13 heart beats for the $T_{2p}4TE$ sequence and to 99 heart beats for the $T_{2p}20TE$ sequence. The employed ARCTIC motion correction and reconstruction of one T_2 map with 20 T_2 prep echo times was 20s.

Figure 1 shows an example of the remaining in-plane motion between T_2 -weighted images acquired in one healthy subject using the $T_{2p}4TE$ sequence under breath-hold, free breathing, and free breathing with respiratory navigator gating. Motion artifacts can be observed in the reconstructed T_2 maps (see white arrows). In-plane motion correction improves the spatial alignment of T_2 -weighted images and results in visually improved T_2 map quality (Figure 1).

Figure 2 shows quantitative metrics of motion accuracy (DSC and MBE) obtained in healthy subjects using the three aforementioned acquisition sequences. Increased DSC and reduced MBE were observed in each of the three acquisition sequences. In the remaining part of this paragraph, DSC and MBE are reported as (uncorrected data vs. motion corrected data using ARCTIC). On average for all subjects, the DSC increased in breath-hold data (0.85 ± 0.08 vs. 0.90 ± 0.02 , $p = 0.007$), free breathing data (0.78 ± 0.13 vs. 0.90 ± 0.21 , $p < 0.001$), and free breathing data with respiratory navigator gating (0.86 ± 0.05 vs. 0.90 ± 0.02 , $p = 0.002$). The MBE decreased in breath-hold data (0.90 ± 0.40 vs. 0.64 ± 0.19 mm, $p = 0.005$), free breathing data (1.21 ± 0.65 vs. 0.63 ± 0.10 mm, $p < 0.001$), and free breathing data with respiratory navigator gating (0.81 ± 0.21 vs. 0.63 ± 0.08 mm, $p < 0.001$).

Figure 3 shows an example of multiple T_2 maps obtained in one healthy subject using the $T_{2p}20TE$ sequence acquired under free breathing conditions with respiratory navigator gating. T_2 maps are shown when reconstructed from only 4 T_2 prep echo times and from all 20 T_2 prep echo times. The level of artifacts in uncorrected T_2 maps appears higher than in motion corrected T_2 maps (see white arrows). As expected, motion artifact patterns have high spatial variability in uncorrected T_2 maps. Furthermore, the spatial variability of the myocardial T_2 estimates appears well reduced when using all 20 T_2 prep echo times compared to only 4 T_2 prep echo times.

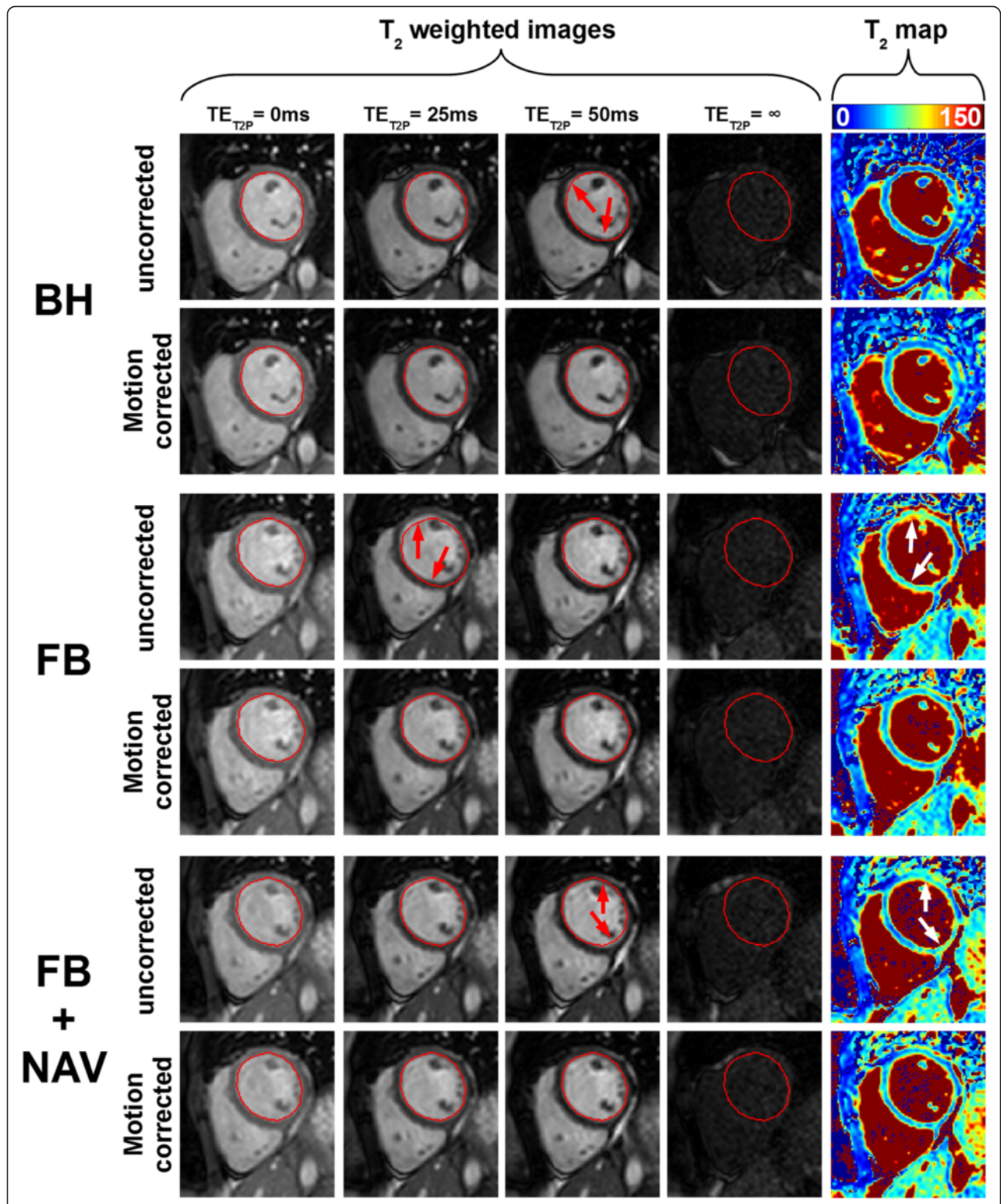
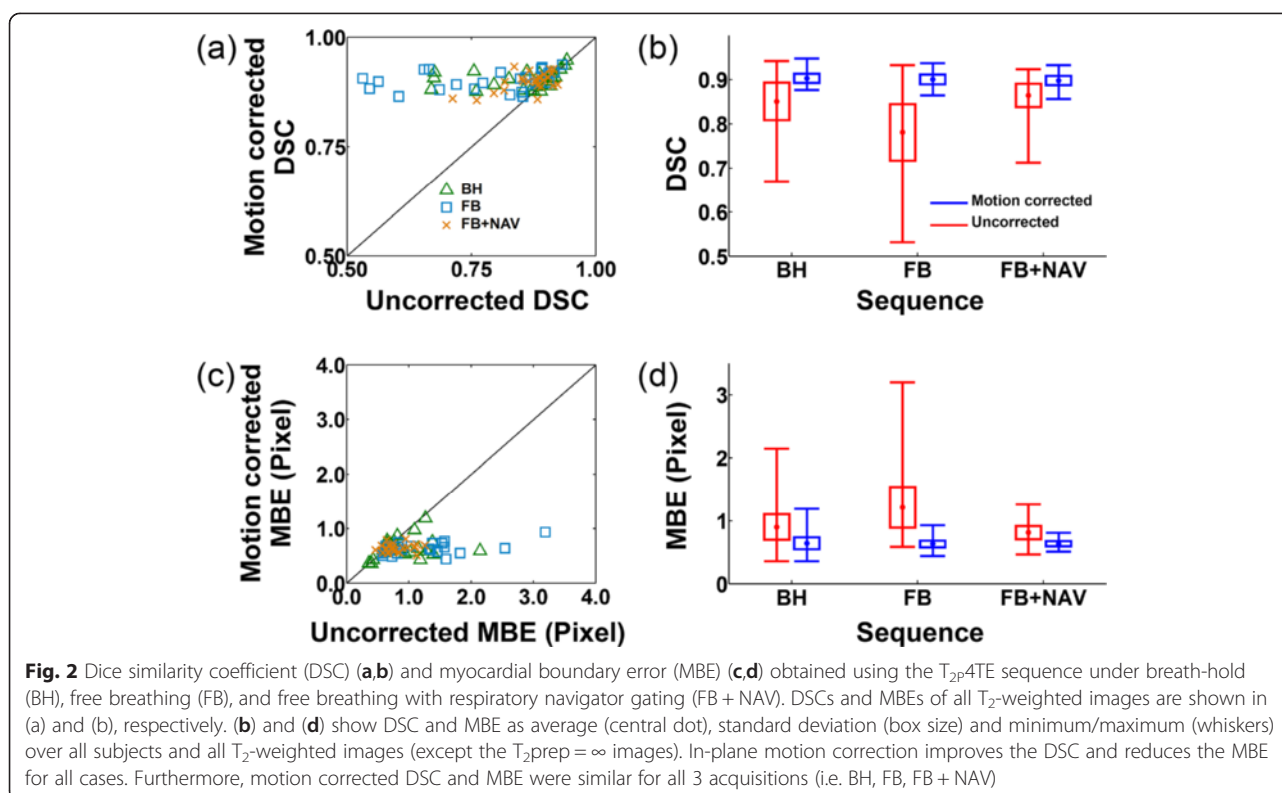


Fig. 1 T₂ scans from one subject acquired using the T_{2P}4TE sequence under breath-hold (BH), free breathing (FB), and free breathing with respiratory navigator gating (FB + NAV). Data are shown without (uncorrected) and with (motion corrected) in-plane motion correction. The endocardial contour of the LV myocardium, drawn on the reference image (1st image) of each scan, is reported in all subsequent T₂-weighted images to facilitate visual motion assessment. Misalignments observed among uncorrected images (red arrows) were substantially reduced after in-plane motion correction using ARCTIC. Furthermore, artifacts in uncorrected T₂ maps (white arrows) were reduced in motion corrected T₂ maps



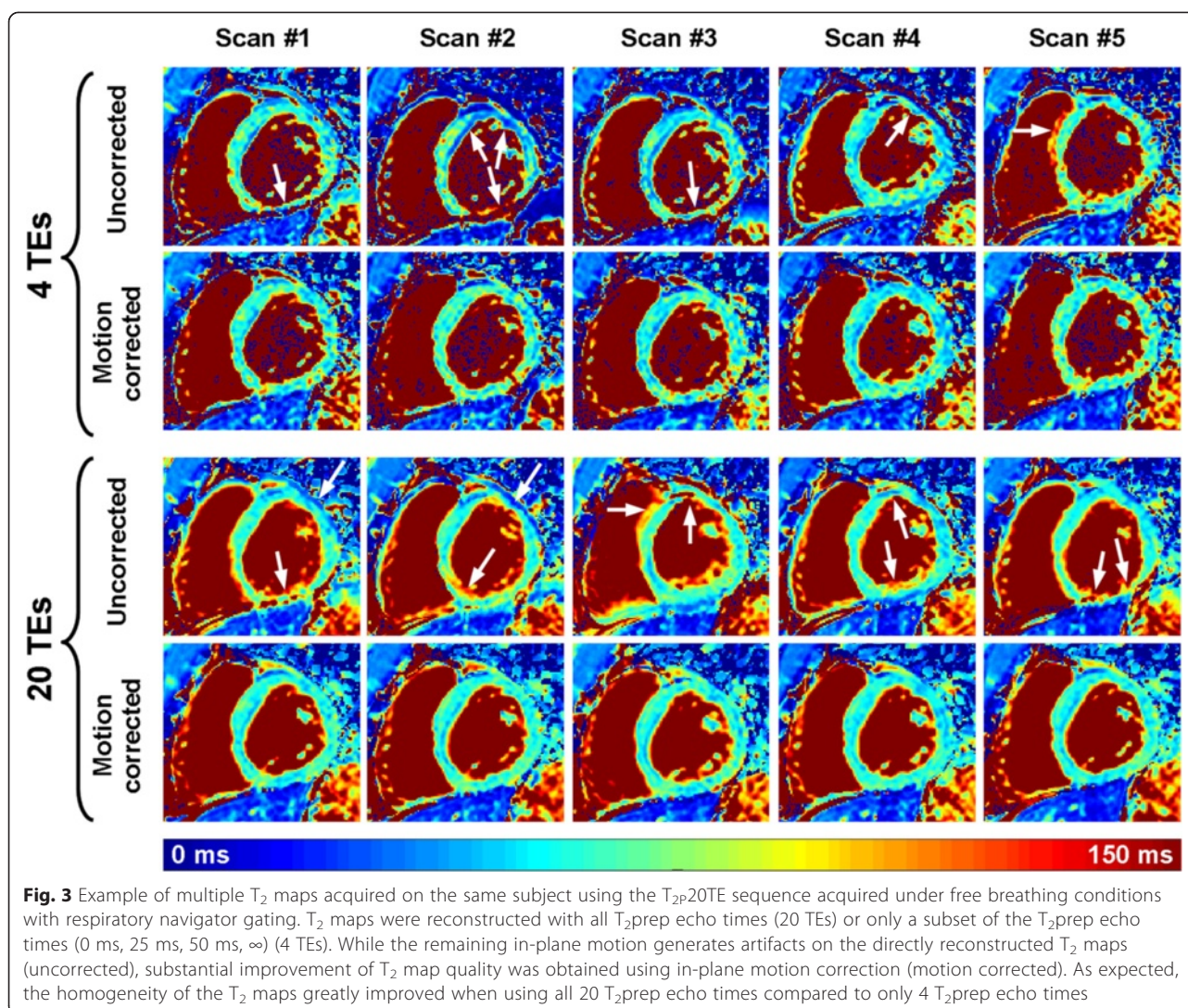
Figures 4 and 5 summarize the reproducibility and spatial variability of T_2 measurements obtained in healthy subjects using the $T_{2p}20TE$ sequence. Results are shown for uncorrected and motion corrected T_2 maps reconstructed using either 4 T_{2prep} echo times or 20 T_{2prep} echo times. Reproducibility and spatial variability are reported as uncorrected T_2 maps vs. motion corrected T_2 maps using ARCTIC. Improved reproducibility was observed over all subjects and myocardial segments in T_2 maps reconstructed from 4 T_{2prep} echo times (5.3 ± 2.5 ms vs. 4.0 ± 1.5 ms, $p = 0.016$) and 20 T_{2prep} echo times (3.9 ± 2.3 ms vs. 2.2 ± 0.5 ms, $p = 0.002$). Similarly, reduced spatial variability was observed over all subjects and myocardial segments in T_2 maps reconstructed from 4 T_{2prep} echo times (12.8 ± 3.5 ms vs. 10.3 ± 2.5 ms, $p < 0.001$) and 20 T_{2prep} echo times (9.7 ± 3.5 ms vs. 7.5 ± 1.4 ms, $p = 0.005$).

As expected, T_2 maps reconstructed using 20 T_{2prep} echo times had better reproducibility than those reconstructed using only 4 T_{2prep} echo times in both uncorrected data (3.9 ± 2.3 ms vs. 5.3 ± 2.5 ms, respectively, $p = 0.007$) and motion corrected (2.2 ± 0.5 ms vs. 4.0 ± 1.5 ms, respectively, $p < 0.001$). The spatial variability of myocardial T_2 estimates reconstructed using 20 T_{2prep} echo times was also lower than the one obtained with 4 T_{2prep} echo times in both uncorrected data (9.7 ± 3.5 ms vs. $12.8 \pm$

3.5 ms, respectively, $p < 0.001$) and motion corrected data (7.5 ± 1.4 ms vs. 10.3 ± 2.5 ms, respectively, $p < 0.001$).

Figure 6 shows example uncorrected and ARCTIC motion corrected T_2 maps obtained in four patients. Large regional variations and artifacts can be observed in uncorrected T_2 maps (see white arrows). The proposed ARCTIC motion correction substantially improved the T_2 map quality in all 4 patients.

Figure 7 shows the subjective assessment of T_2 map quality obtained in 50 patients. Overall ($N = 150$ T_2 maps), ARCTIC motion corrected T_2 maps had higher quality score than uncorrected T_2 maps (3.69 ± 0.55 vs. 3.43 ± 0.79 , $p < 0.001$). In the relative comparison of T_2 map quality, uncorrected T_2 maps has superior, similar, and inferior quality than ARCTIC motion corrected T_2 maps in 4 maps (3%), 99 maps (66%), and 47 maps (31%), respectively. Furthermore, the motion level was assessed as “no motion” in 35 slices (23%), “small motion” in 69 slices (46%), and “large motion” in 46 slices (30%). In “no motion” data, all ARCTIC motion corrected and uncorrected T_2 maps received a subjective quality score of 4.0 and 97% of them had similar relative quality. In “small motion” data, ARCTIC motion corrected T_2 maps had higher subjective quality score (3.71 ± 0.49 vs. 3.61 ± 0.60 , $p = 0.015$) and superior (23%), similar (75%) and

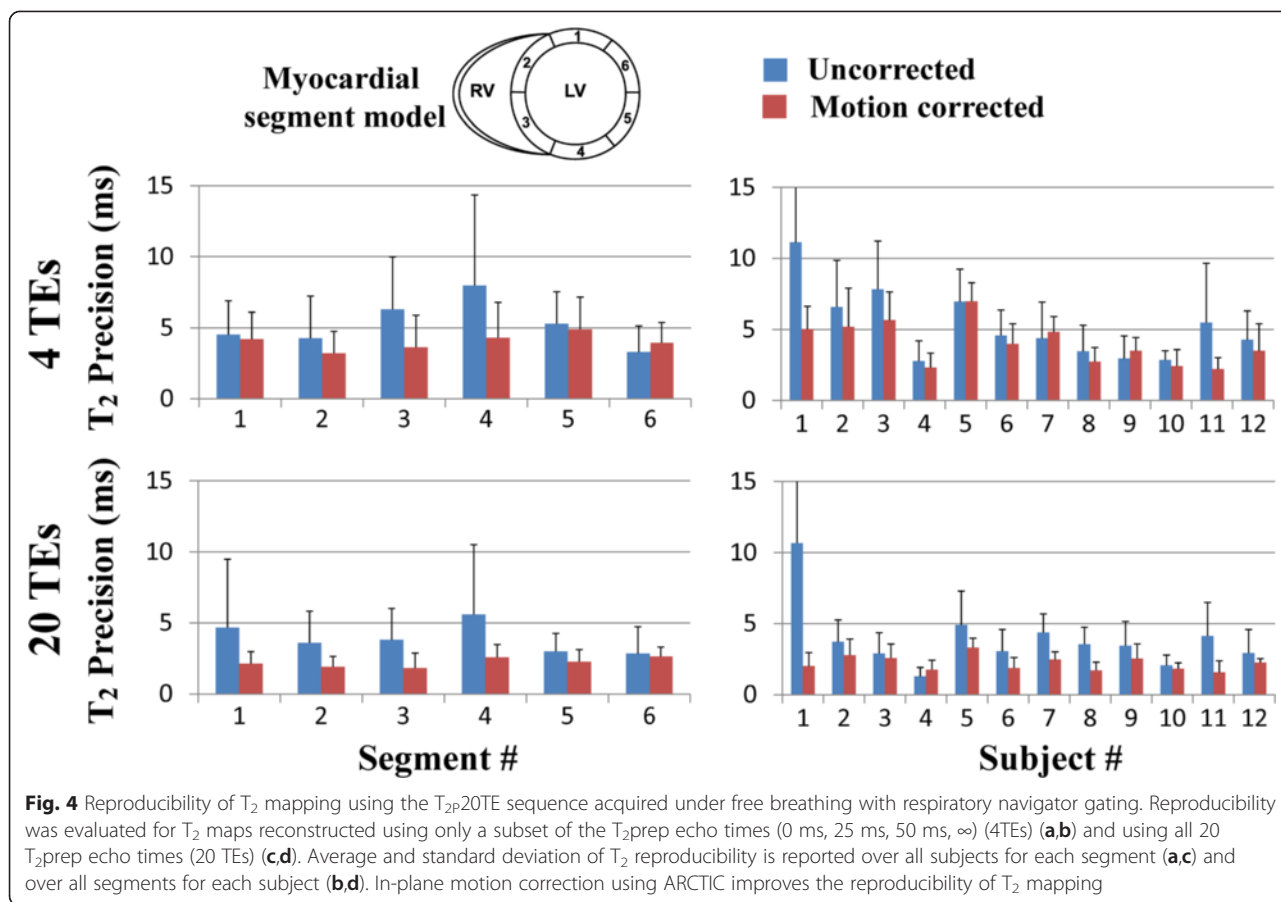


inferior (1%) relative quality than uncorrected T_2 maps. In “large motion” data, ARCTIC motion corrected T_2 maps had higher subjective quality score (3.41 ± 0.69 vs. 2.72 ± 0.83 , $p < 0.001$) and superior (65%), similar (28%) and inferior (6%) relative quality than uncorrected T_2 maps.

Discussion

In this study, we demonstrate the benefit of in-vivo in-plane ARCTIC motion correction in myocardial T_2 mapping. The method provides improved alignment of the myocardium in T_2 -weighted images acquired with breath-hold acquisitions and free breathing acquisitions with and without respiratory navigator gating. ARCTIC motion correction improves T_2 map quality which results in improved reproducibility and spatial variability of myocardial T_2 estimates. Finally, the CPU/GPU implementation of ARCTIC substantially reduces the computation time of the T_2 map reconstruction to 20s which is suitable for clinical applicability.

DSCs and MBEs found in this study are in good agreement with previous studies [23–25, 38]. As expected higher mis-alignments were observed using free breathing acquisitions without respiratory navigator gating. DSCs/MBEs improvement was obtained in all three types of acquisitions. This confirms the benefit of motion correction, even for data acquired with a breath-hold. This is likely because 40–60% of patients fail to sustain a stable breath-hold in these conditions [23–25]. Furthermore, similar DSCs/MBEs were obtained after motion correction using the three acquisition conditions (breath-hold and free breathing with and without respiratory navigator). It is important to note that through-plane motion cannot be compensated when using the free breathing acquisition without respiratory navigator gating. In this case, the efficacy of in-plane motion correction algorithms depends on the subject’s heart orientation in relation to his respiratory movement. The use of respiratory navigator appears thus



desirable to enable through plane motion compensation in free breathing acquisitions. The registration accuracy was not evaluated in the $TE_{T2P} = \infty$ images since the contrast is too low to identify the myocardium. Motion correction is expected to have slightly lower accuracy in those images due to the expected limited ability to compensate for complex motion.

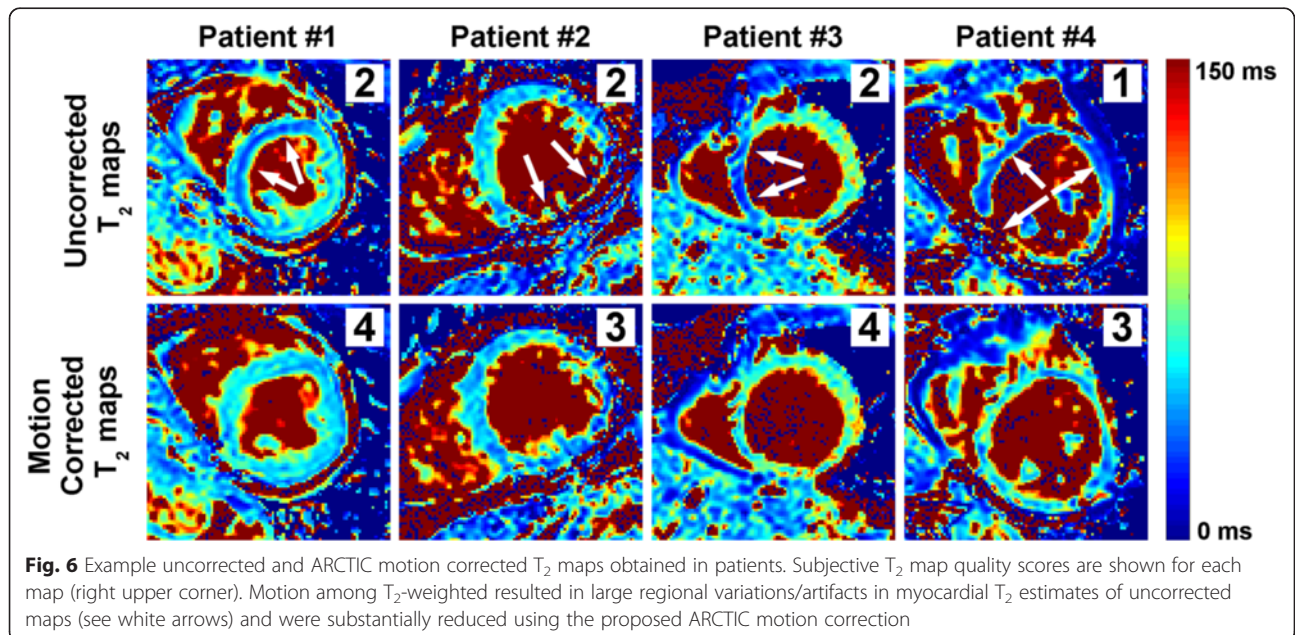
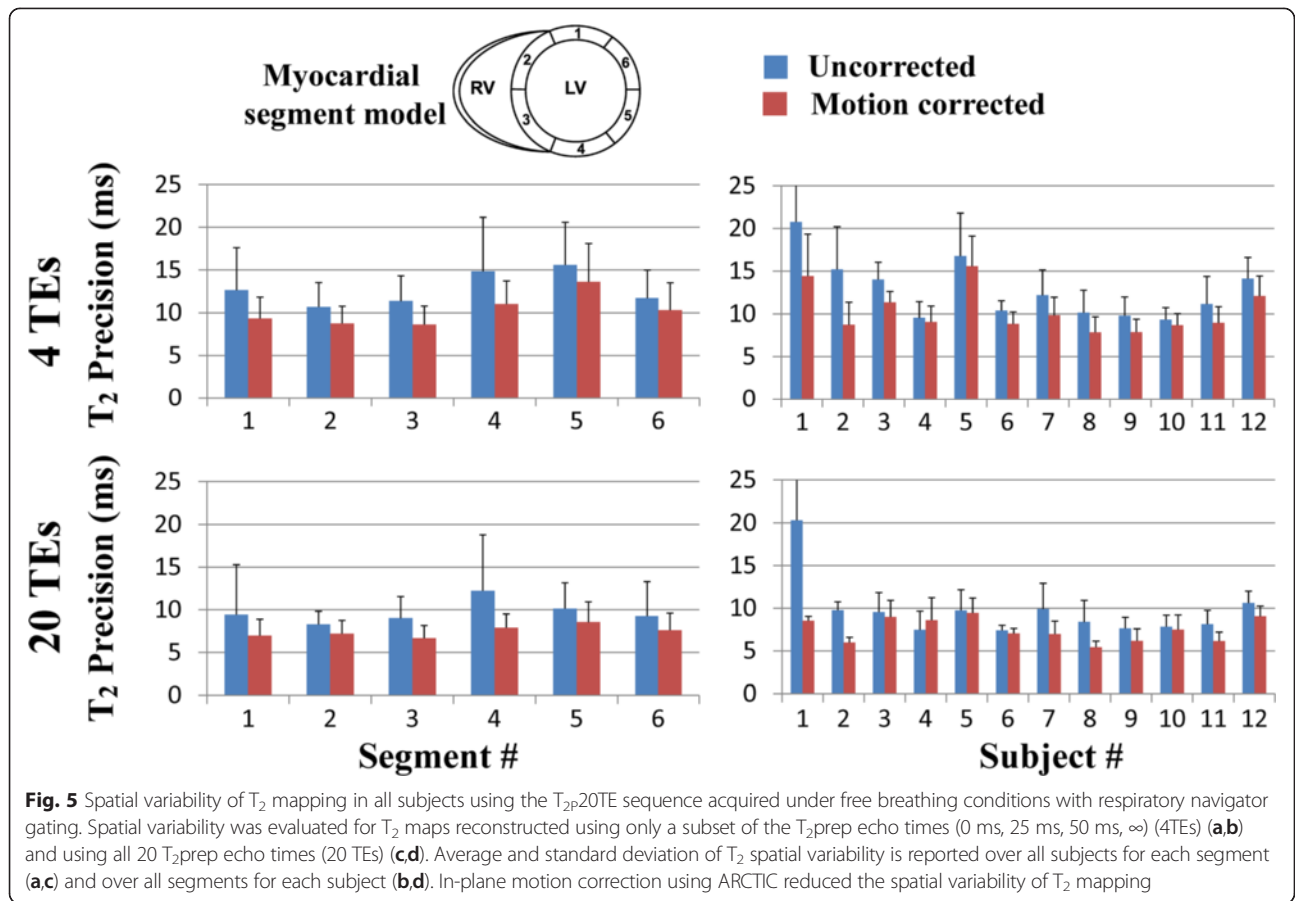
The ARCTIC approach successfully corrected the encountered motion in all subjects. In this study, the heart motion patterns were mainly influenced by the breathing activity of the subjects and to lesser extent to their RR-interval variations. However, the motion pattern can be more complex in patients imaged during arrhythmic events. The performance of the method in such conditions was not investigated and should be addressed in future work.

The reproducibility and spatial variability of T₂ mapping was improved using ARCTIC. The use of 20 T₂-weighted images improved the reproducibility and the spatial variability of T₂ mapping (over the use of only 4 T₂-weighted images) by a factor of 2 and 1.4, respectively. Therefore, the choice of the number of T₂prep echo times depends on the desired trade-off between acquisition time and T₂ map quality. Further studies are warranted to determine the clinically relevant threshold

providing satisfactory T₂ map quality in a reasonable amount of time.

Reproducibility and spatial variability of T₂ estimates were found similar in all myocardial segments when using 20 TEs. However, slight differences seemed to be observed when using 4TEs only, especially in the myocardial segment #4 (inferior wall). Several factors could have contributed to this observation including 1) increased sensitivity to cardiac motion and partial voluming in the free wall due to reduced wall thickness, 2) increased field inhomogeneity in myocardial segments located at the heart/lung interface. Future studies are warranted to study the impact of each of these factors.

In this study, the data were acquired using our recently developed T₂ mapping sequence. The ARCTIC approach is expected to provide similar motion correction performance using other T₂ mapping sequences. Nevertheless, the impact of motion correction on the reproducibility and spatial variability of other T₂ mapping sequences may be different and is beyond the scope of this study. Furthermore, all data were acquired in 2D. 3D myocardial T₂ mapping may represent a valuable approach for true 3D assessment of pathological tissues [39, 40]. The extension of the ARCTIC approach to 3D is straightforward and is expected to provide similar



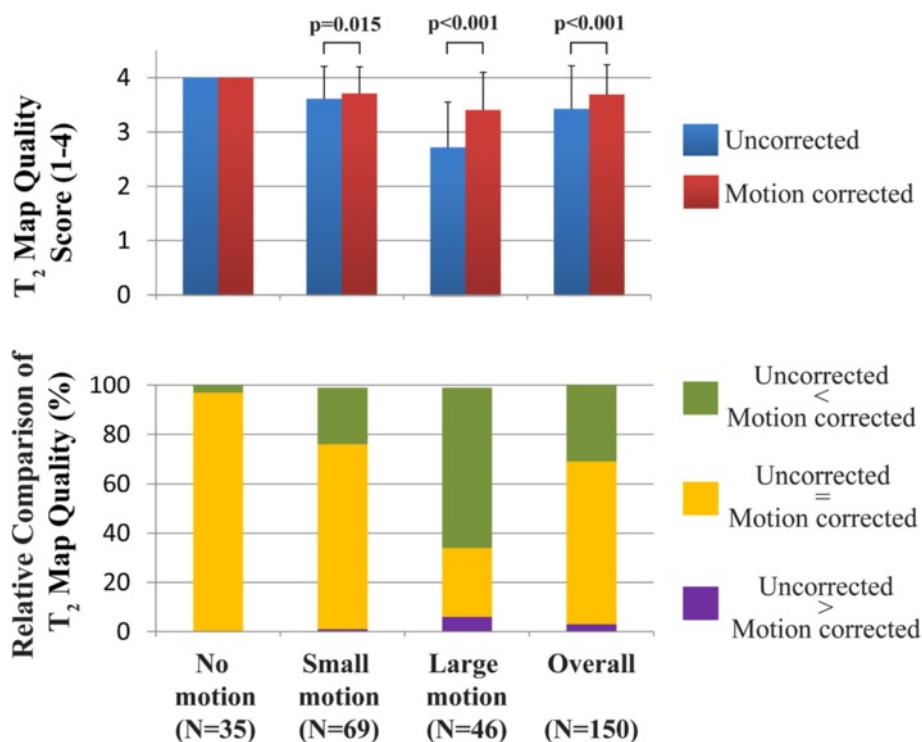


Fig. 7 Subjective assessment of T_2 map quality in patients. In-plane motion correction using ARCTIC increased T_2 map quality scores (3.69 ± 0.55 vs. 3.43 ± 0.79 , $p < 0.001$)

improvement of the reproducibility and spatial variability of 3D T_2 mapping.

There are several limitations in this study. In the in-vivo analysis of reproducibility and spatial variability, the 4TEs data were extracted from the $T_{2P}20TE$ sequence and were thus not acquired using the $T_{2P}4TE$ sequence. However, since the $T_{2P}20TE$ sequence was acquiring with respiratory gating, the potential bias in reproducibility and spatial variability obtained in the 4TEs data should have been kept to the minimum. Finally, the study was only performed in healthy adult subjects with limited sample size. Further studies are warranted to confirm the benefit of the ARCTIC approach to improve the reproducibility and spatial variability of myocardial T_2 mapping in patients.

Conclusions

The ARCTIC technique substantially reduces spatial mis-alignment among T_2 -weighted images. This method improves the reproducibility and reduces the spatial variability of in-vivo T_2 mapping. Furthermore, the in-vivo reproducibility and spatial variability of T_2 mapping is improved using a higher number of T_2 prep echo times combined with ARCTIC motion correction.

Abbreviations

T_2 prep: T_2 prepared; ARCTIC: Adaptive registration of varying contrast-weighted images for improved tissue characterization; BH: Breath-hold; FB: Free breathing; FBNAV: Free breathing conditions with respiratory navigator gating; DSC: DICE similarity coefficient; MBE: Myocardial boundary errors (MBE); TE: Echo times; SSFP: Steady-state free precession; HIPAA: Health insurance portability and accountability act; ECG: Electrocardiogram; TET2P: T_2 prep echo times; TR: Repetition time; FOV: Field of view; SENSE: Sensitivity encoding; SNR: Signal-to-noise ratio; GPU: Graphic processing unit; CUDA: Compute unified device architecture.

Competing interests

SR, WJM and RN have a pending patent for methods for correcting motion for tissue characterization sequences. TB, MA, WJM, and RN have a pending patent for system and method for assessing T_2 relaxation times with improved accuracy.

Authors' contributions

SR participated in the study design and coordination, developed the ARCTIC approach, carried out the motion correction/reconstruction of the data, performed the data analysis and drafted the manuscript. TB developed the prospective T_2 mapping sequence and participated in data acquisition. SW developed the T_2 mapping reconstruction code. MA participated in the data acquisition. SB was in charge of subject recruitment. WJM helped in revising the manuscript. RN conceived the study, participated in the study design and interpretation of the data. All authors read and approved the final manuscript.

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Author details

¹Department of Medicine (Cardiovascular Division), Beth Israel Deaconess Medical Center, 330 Brookline Ave, Boston, MA 02215, USA. ²Biomedical Engineering Department, Cairo University, Giza, Egypt. ³Computer Assisted Clinical Medicine, University Medical Center Mannheim, Heidelberg University, Mannheim, Germany. ⁴Radiology, Beth Israel Deaconess Medical Center and Harvard Medical School, Boston, MA, USA.

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