

Comparison of Three Different Cardiac T2-Mapping Techniques at 1.5 Tesla



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Abstract

Background: T2-mapping techniques gain increasing acceptance to study myocardial edema in inflammatory and ischemic heart diseases.

Study Aim: To compare the performance of a breath-hold and two free-breathing T2-mapping techniques in a prospective design in healthy subjects.

Methods: Three different sequences for T2-mapping were tested on a clinical CMR scanner (Aera, Siemens, Germany) at 1.5T:

- Breath-hold 2D-acquisition technique (2D-BH)
- Free-breathing 2D-technique applying a diaphragmatic navigator (2D-FBNav), and

c) Free-breathing 3D-technique applying self-navigation for respiratory motion correction (3D-FBSN). T2-values were quantified in 6 segments per short-axis slice on 5 slices (2D-BH and 3D-FBSN) and 3 slices (2D-FBNav) covering the left ventricle. Analyses were also performed with larger segments (8- and 2-segment models). As a quality measure, the coefficient of variation (CV%=standard deviation of T2-value expressed as percentage of mean T2) was determined. Contours were drawn manually by 2 observers using commercial software (Gyrotools, Zurich, Switzerland). T2 differences between techniques, slices, and segments were evaluated by repeated-measures ANOVA and post-hoc Bonferroni-correction.

Results: With 2D-BH, diagnostic images were obtained in all 13 volunteers (=390 segments). With 2D-FBNav 3 slices/volunteer were acquired and quality was non-diagnostic in 5 slices yielding 204 segments for analysis. With 3D-FBSN 1 volunteer was not evaluable yielding 360 segments for analysis. Mean T2-values (=T2 averaged over all segments) were higher for the 3D-FBSN (54.4±5.4ms vs. 48.5±2.4ms and 45.9±4.4ms for 2D-BH and 2D-FBNav, respectively, p<0.002). The CV% of the 3D-FBSN technique was higher vs. both, 2D-BH and 2D-FBNav (12.1±5.4% vs. 7.0±1.1% and 8.3±3.9%, respectively, p<0.02).

Conclusion: At 1.5T, most reliable T2 results with a low coefficient of variation were obtained by the 2D-BH technique. The 2D-FBNav technique is considered as an alternative if breath-holding capacity is not sufficient. The 3D-FBSN technique is not at the same level of robustness as the breath-holding technique and not yet recommended for clinical use.

Keywords: Heart; MRI; T2-mapping; 1.5 Tesla; Techniques

Introduction

For the detection of inflammatory myocardial diseases [1] and the characterization of myocardial edema in the setting of ischemic heart disease, CMR techniques, and in particular, novel T2-mapping

techniques, gain increasing acceptance. While T2-weighted imaging strategies allow for the detection of myocardial edema, T2-mapping techniques are considered advantageous as they yield quantitative

measures of T2 and this should allow for more reliable inter-subject comparisons, e.g. for follow-up studies to evaluate treatment response. The quantitative T2 mapping strategy should also facilitate multicenter trials and inter-study comparisons. However, T2 mapping pulse sequences are demanding as they require 3 or more image acquisitions (with different T2-waiting times) to reconstruct the T2-map. Breath-hold approaches are proposed to generate T2 maps [2] but some compromises are unavoidable regarding spatial resolution and/or acquisition window duration in order to fit the acquisition into a single breath-hold.

Alternatively, free-breathing strategies could be superior with regard to spatial resolution and acquisition window duration, if breathing motion can be adequately corrected for [3,4]. For such breathing motion correction, navigators applied to the right hemidiaphragm were used [3] or self-navigation strategies [4]. The aim of this study was to evaluate the performance of 3 different T2-mapping techniques, i.e.

- a) A breath-hold 2D acquisition technique (2D-BH)
- b) A free-breathing 2D acquisition technique applying a diaphragmatic navigator approach (2D-FB_{Nav})
- c) A free-breathing 3D acquisition technique applying self-navigation for breathing motion correction (3D-FB_{SN}). As most clinical CMR studies are performed on 1.5T systems [5], both the 2D-FB_{Nav} and 3D-FB_{SN} pulses sequences developed for 3T [3,4] were transferred onto a 1.5T system for this comparative study.

Methods

Volunteers: The study enrolled 13 healthy subjects (5 men, 8 women, age 24-47 years old). Exclusion criteria were presence of any cardiac pathology, a history of cardiac interventions, any past or current cardiac medication, and presence of a pacemaker/defibrillator. The study protocol was approved by the local Ethics Committee and all study subjects gave written informed consent before study participation.

T2 Mapping Pulse Sequences: In each subject, 3 different cardiac T2 mapping pulse sequences were acquired at 1.5T (Magnetom Area, Siemens Healthcare, Germany) in a single scanning session starting with a 2D breath-hold acquisition (2D-BH) [2], followed by a high spatial resolution free-breathing 2D acquisition (2D-FB_{Nav}) [3], and ending with a high spatial resolution free-breathing self-navigation 3D acquisition (3D-FB_{SN}) [4], Table 1. With the 2D-BH, 5 short-axis slices of the left ventricle (LV) were acquired distributed evenly along the long axis of the LV. With the 2D-FB_{Nav} sequence, only 3 LV short-axis slices were acquired (avoiding the most basal and most apical short-axis slice) to account for the longer acquisition duration for this sequence. Finally, with the 3D-FB_{SN} sequence the entire LV was covered.

Table 1: Magnetic resonance imaging parameters for the 3 different techniques

	2D-BH	2D-FB _{Nav}	3D-FB _{SN}
Repetition time TR (ms)	2.5	4.5	4.0
Echo time TE (ms)	1.06	2.1	1.32
Acquisition (steady-state free precession read-out)	cartesian	radial	radial
Acquisition matrix	192x156	256x256	128x128
Field-of-View (cm)	36 x 29	30 x 30	22 x 22
Slice thickness (mm)	8	8	1.72
Pixel Bandwidth (Hz)	1'185	590	910
Flip angle	35°	90°	70°
Number of heart beats waited	3	3	3
T2prep pulse intervals (ms)	0/25/50	0/30/60	0/30/60
k-line readouts/heart beat	76	21	30
Acquisition window (ms)	190	94.5	120

This table shows the MRI pulse sequence parameters for the three different techniques.

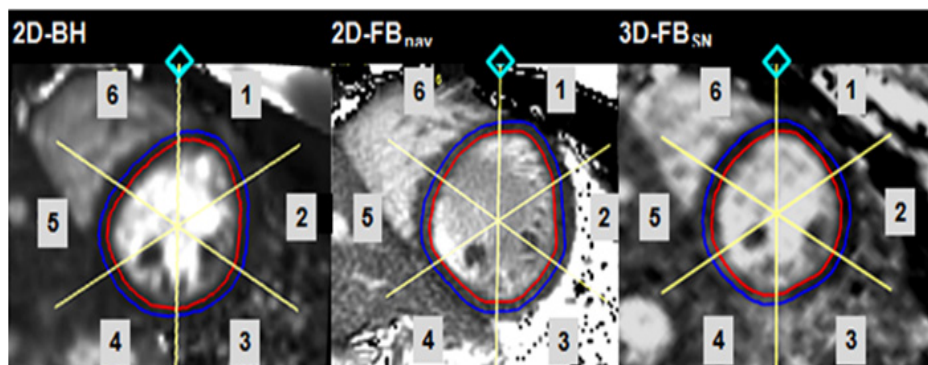


Figure 1: T2-mapping images in the 3 different techniques. Examples of a mid-ventricular short-axis T2 map obtained with 2D-BH (A), 2D-FB_{Nav} (B), and 3D-FB_{SN} (C). Reference point (cross) is the anterior insertion point of the right ventricle at the septum. Blue/red: epicardial/endocardial contours.

Data Analysis: For the 2D-BH and 2D-FB_{Nav} acquisitions, on each slice, the epicardial and endocardial contours of the LV myocardium were manually drawn with a thickness of 3 mm, approximately 1.5 mm from each side of the middle line of the

myocardium (to avoid T2 values from the blood pool and epicardial fat). The LV short-axis slices were manually divided into 6 segments in a clockwise manner (similar to the AHA guidelines) [6] with the reference point chosen at the anterior insertion of the right ventricle

on the LV (Figure 1) yielding 30 and 18 segment models for the 2D-BH and 2D-FBNav acquisitions, respectively. For the analysis of the 3D-FBSN acquisition, 5 short-axis slices were selected out of the 3D volume based on anatomical landmarks (Figure 1) to match the short-axis slices of the 2D acquisitions (yielding 30 segments per

heart for this analysis). Analyses were performed using commercial software (Gyrotools GmbH, version 2.2.1, Zurich, Switzerland). To fully exploit the LV coverage by the 3D-FBSN acquisition, the heart was divided into larger segments yielding one 8-segment and three 2-segments models as shown in Figure 2.

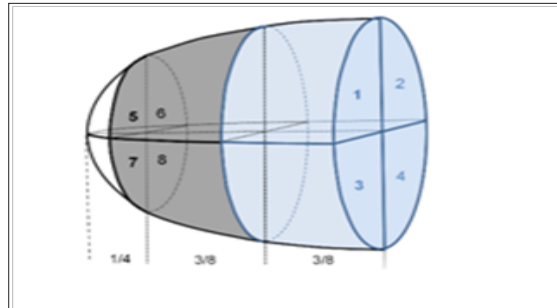


Figure 2: Heart division in the different segment-division models. The whole heart analysis yields one 8-segment model (antero-lateral (1;5), infero-lateral (2;6), infero-septal (4;8), and antero-septal (3;7) for the basal and midventricular level, respectively) as well as three 2-segment models dividing the heart into an anterior and inferior half (segments 1;3;5;7 and 2;4;6;8, respectively), a lateral and septal half (segments 1;2;5;6 and 3;4;7;8, respectively), and a basal and mid-ventricular half (segments 1;2;3;4 and 5;6;7;8, respectively). For these analyses the apex was excluded to minimize partial volume artifacts.

To compare these data with the 2D data sets, several segments and slices of the 2D data were combined to match these 8- and 2-segments models. For 2D-BH, e.g., the basal segments i.e. segments 1-4 on (Figure 2) were calculated by averaging slice 1, slice 2, and half of slice 3 and the mid segments i.e. segments 5-8 on (Figure 2) by averaging half of slice 3, and the entire slice 4 and slice 5. For 2D-FBNav the basal and mid segments were calculated by averaging basal slice 1 and half of slice 2 and by averaging half of slice 2 and slice 3, respectively. In analogy, segments of a slice were combined to match the 8-segment and 2-segment models. The pulse sequences used in this study are all well validated against known T2 values determined in phantoms [3,4,7].

As the true myocardial T2 in humans is difficult to obtain, we used the variability of T2 values measured in the volunteer group as an indicator of data quality. This variability is expected to be small for a robust measurement when performed in a homogenous group

of healthy volunteers. Measurement variability was expressed for each segment (of the 30-segment, 18-segment, 8-segment and the three 2-segment models obtained in the 13 volunteers) as the SD (for each segment) divided by the mean T2 value for this segment, i.e. by the coefficient of variation (CV% = SD expressed as percentage of the mean T2 value).

Statistics

Continuous data are presented as mean ± standard deviation. T2 values and CV% for the 3 pulse sequences were compared by 1-one way ANOVA for repeated measures with post-hoc Bonferroni correction. CV% for the 3 techniques was compared on the basis of the 18 segment model (3 slices with 6 segments each). CV% was also compared between the 18-segment, the 8-segment, and the three 2-segment models for each pulse sequence. A post-hoc corrected p value <0.05 was considered significant.

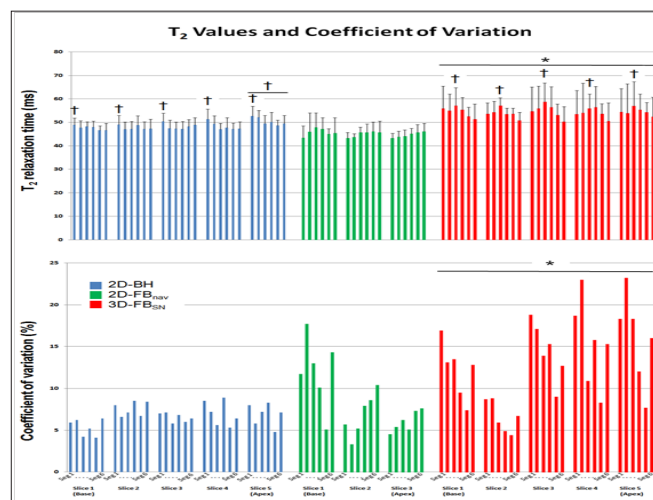


Figure 3: Segmental T₂ (A) and CV% (B) values for the three T₂-mapping sequences (30 segments for 2D-BH and 3D-FB_{SN}, 18 segments for 2D-FB_{Nav}). Average T₂ as well as CV% were both higher with 3D FB_{SN} vs 2D-BH and 2D-FB_{Nav} (*significant vs 2D-BH and 2D-FB_{nav}; † significant difference between segments or slices of the same pulse sequence; for details, see results section).

Results

Comparison of 2D-BH, 2D-FBNav, and 3D-FBSN

In all volunteers the 2D-BH T2 mapping sequences yielded complete data sets of diagnostic quality. With the 2D-FBNav sequence (with 3 slices per subject) 1 slice was of non-diagnostic quality in 3 subjects each, and in 2 additional subjects, slice 3 was not acquired (as the irregular breathing pattern did not allow finishing the acquisition in time). For the 3D-FBSN pulse sequence, data of one subject were not correctly acquired (due to inaccurate tracking of the heart silhouette). The T2 values, averaged over all segments, for 2D-BH and for 2D-FBNav were 48.5±2.4ms and 45.9±4.4ms, respectively, which were lower than 54.4±5.4ms measured by 3D-FBSN (p<0.002 for both, no difference for 2D-BH vs. 2D-FBNav, Figure 3A). The CV% of the 3D-FBSN technique was higher vs both, the 2D-BH and 2D-FBNav acquisition (CV% 12.1±5.4% vs 7.0±1.1% and vs 8.3±3.9%, respectively, p<0.02 for both, Figure 3B).

Distribution of T2 Values in Normal Myocardium for the three T2 Mapping Techniques

With 2D-BH, T2 of 50.4±2.9ms in slice 5 was slightly higher than in the other 4 slices (47.7±2.0ms, 47.8±3.1ms, 47.7±2.0ms, 48.3±2.9ms, respectively, p<0.001). Also, the T2 of the anterior segment was higher than the T2 of the other segments (50.5±3.2ms

vs segments 2, 3, 4, 5, and 6 with 48.8±2.6ms, 47.9±2.6ms, 48.4±3.1ms, 47.6±2.0ms, and 48.0±2.7ms, respectively, p<0.005). Inspection of Figure 3A indicates that slice 5 contributes most to the elevated T2 in the anterior segment. For 2D-FBNav, no differences in T2 were found between slices or segments. For 3D-FBSN, no differences in T2 between slices were found. However, T2 in the infero-lateral segment 3 was higher in comparison to the “opposite” antero-septal segment 6 (57.2±6.7ms vs 51.1±6.4ms, p<0.003; Figure 3A).

Influence of Size of Myocardial Segments on Robustness of T2 Measurements

For the models with larger segment sizes, T2 did not differ for the 3 techniques (Figure 4A). For 3D-FBSN, CV% showed a trend to diminish with increasing sizes of segments (CV% for 18-, 8-segment, and 2-segment models: 12.1±5.4%, 8.6±2.1%, 7.2±0.6%, and 7.6±1.9%, and 7.1±0.1%, respectively, ns, Figure 4B). Similarly, for 2D-FBnav, there was a trend of CV% to decrease with increasing segment size (CV% for 18-, 8-, and 2-segment models: 8.3±3.9%, 5.5±0.6%, 6.0±2.0%, 5.7±0.7%, and 6.1±0.1%, respectively, ns, Figure 4B). For the 2D-BH data, the models with 8 and 2 segments yielded slightly lower CV% vs. the 18-segment model (5.6±0.6%, 5.0±0.1%, 5.1±0.5%, 5.2±0.4% vs 7.0±1.1%, respectively p<0.001, Figure 4B).

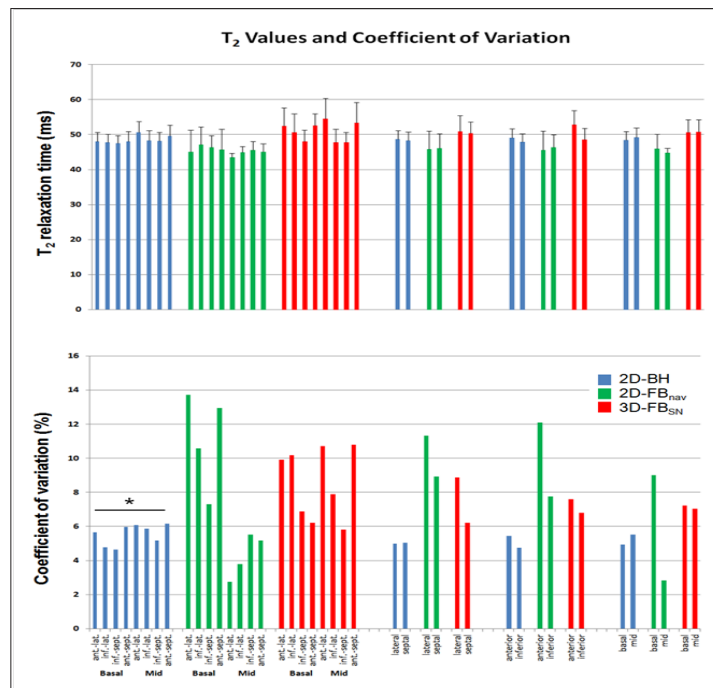


Figure 4: Segmental T₂ (A) and CV% (B) values for the three T₂-mapping sequences for models with 8 and 2 segments per heart. For models with larger segments, CV% tended to decrease with 2D-FB_{Nav}, 3D-FB_{SN}, which reached statistical significance for 2D-BH (*significant vs 18-segment model; for details, see results section).

Discussion

For both, the 2D-BH and the 2D-FBNav technique, very similar T2 values were obtained with 48.5±2.4ms and 45.9±4.4ms, respectively. Also, these T2 values were homogeneously distributed in the 18 segments per heart as shown in Figure 3A. For the 2D-FBNav technique, T2 values were not different over the 3 short-axis slices or in the different segments. For the 2D-BH technique, the

T2 values were also homogeneously distributed. Only in the most apical slice 5, the T2 was elevated by 2-3ms in comparison to the other slices. Similarly, with the 2D-BH sequence T2 in the anterior segment 1 was slightly higher than in the other segments. As shown in Figure 3A, inspection of individual segmental T2 data indicates, that the apical slice 5 is mostly contributing to the T2 elevation in anterior segment. A likely explanation for the slightly elevated T2

in the apical slice is a partial volume artifact in the apical region. With the 3D-FBSN technique, higher T₂ values were obtained in comparison with the other two techniques. This T₂ elevation was most prominent in the infero-lateral segment, where T₂ was higher than in the “opposite” antero-septal segment (Figure 3A).

This may be explained by the fact that the infero-lateral segment is more distant from the surface coil than the antero-septal segment causing a reduced signal-to-noise ratio (SNR) in this infero-lateral segment. A slight T₂ overestimation in regions with lower SNR has been shown for 3D-FBSN at 3T [4,8]. Regarding the robustness of the three acquisition strategies, the 2D-BH technique proved to be very reliable. In all subjects, the 2D-BH technique yielded analyzable data and the CV% was small with 7.0±1.1%. As shown in Figure 3B, the CV% is also acceptably small for 2D-FBNav with 8.3±3.9%. However, 5 out of 39 short-axis slices were not evaluable with 2D-FBNav, i.e. data were diagnostic in only 87%, and navigator gating is typically related to longer acquisitions. Nevertheless, in case of inadequate breath-holding capabilities, the 2D-FBNav sequence can be considered as an alternative for T₂ mapping.

Unlike 2D-BH and 2D-FBNav, the 3D-FBSN technique is associated with significantly larger CV% of 12.1±5.4% and as a consequence, this self-navigated T₂ mapping technique in its current form loses too much SNR in the transition from 3T to 1.5T, and would require further refinement before it can be recommended for clinical application. One might argue, that

analyzing only 5 slices out of a 3D volume (as acquired with this 3D-FBSN sequence), would not adequately exploit the full 3D-FBSN performance. While the CV% of the 3D-FBSN technique tended to improve with larger segments (from 12.1±5.4% for 18 segments to 7.1±0.1% for 2 segments), it was not superior in comparison to the 2D-BH technique even when compared with small segments (CV% of 7.0±1.1% for the 18-segment model).

In recent publications, T₂ values of healthy volunteers were reported as shown in Figure 5 using a breath-hold T₂prep BSSFP technique [2,9-12] (corresponding to 2D-BH used in the current study) or a gradient-echo (echo-planar) multi-echo spin-echo technique (GraSE technique) either applied during a breath-hold [12-14] or combined with navigators for free breathing [12,15]. As shown in Figure 5, which summarizes these findings, normal T₂ values measured by the different techniques were similar, ranging from 52.2ms to 58.6ms [2,9-15], while the T₂ normal value for the current 2D-BH approach was 48.5ms. When comparing the variability of the T₂ measurements, the current 2D-BH technique with a CV% of 4.9% compares well with the CV% reported in the literature ranging from 3.7% to 6.8% [2,9-15], while the variability of the tested 2D-FBNav and 3D-FBSN are considerably higher with 9.6% and 9.9%, respectively. This translates in an upper limit of normal T₂ values for the 2D-BH technique of 53.3ms which is lower than reported previously ranging from 57.3ms to 60.1ms using the same T₂prep BSSFP technique [2,9-12].

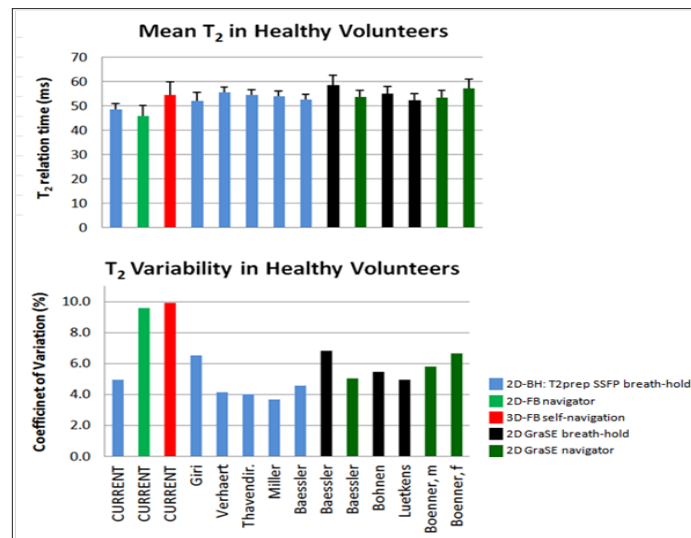


Figure 5: Mean T₂ values measured with 2D-BH, 2D-FB_{Nav} and 3D-FB_{SN} versus mean T₂ values from the literature (2,9-15). For 2D-BH, T₂ tends to be smaller in comparison to published T₂ (Figure 5a), while variability, i.e. CV%, is comparable to other 2D breath-hold techniques reported in the literature (Figure 5b). For the free-breathing 2D-FB_{Nav} and 3D-FB_{SN}, CV% is substantially higher versus published data. Gra SE: gradient-echo (echo-planar) multi-echo spin-echo technique; other abbreviations as mentioned in the text.

The segmental normal values of the 2D-BH approach for the 17-segment model are given in Table 2. In our study the ROI's for T₂ determination were drawn strictly avoiding signals from blood pool or epicardial fat, which was achieved by restricting the ROI to a width of 3 mm. This may have contributed to the low variability of data even when T₂ values were assessed in small segments allowing T₂ quantification in 30 segments per heart. The fact, that sequence performances were assessed in healthy volunteers

only, but not in patients, is certainly a limitation as breath-hold capability may differ in patients. Nonetheless, the results allow the notion that for free-breathing strategies, 2D-FBNav performed better than 3D-FBSN. The patient examples as shown in Figures 6-8 illustrate the clinical usefulness of the normal values and the bull's eyes representation (based on the normal values given in Table 2) to establish diagnoses in patients and to monitor disease evolution.

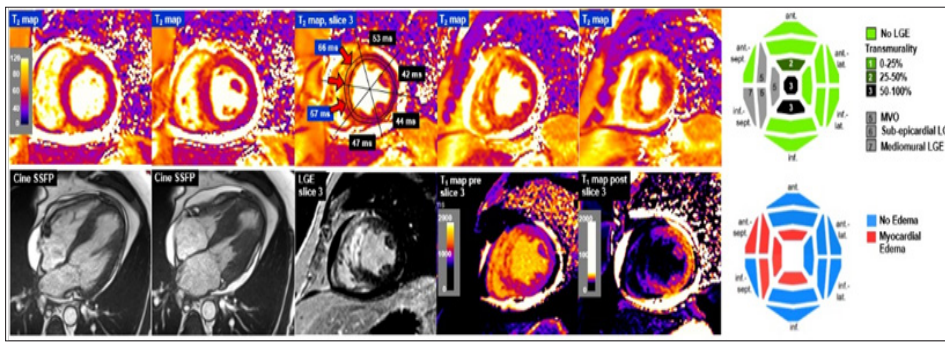


Figure 6: Example: Acute ST elevation myocardial infarction. 50-years old diabetic patient with anterior STEMI, investigated by CMR 5 days after percutaneous coronary intervention of the left anterior descending coronary artery. Cine acquisitions demonstrate severe hypokinesia of the anterior and septal walls with global LV ejection fraction of 45%, mild pericardial effusion, and increased LV mass of 103g/m² (normal <78g/m²) probably due to myocardial edema (LVEDV: 83ml/m²). The bull's eyes show distribution of necrosis (with microvascular obstruction) and of myocardial edema quantified by T₂ mapping. Distribution of late gadolinium enhancement (LGE) can also be quantitated by T₁ mapping before and after contrast medium administration. The T₂ mapping bull's eye shown in Figures 6-8 uses the threshold values for elevated T₂ as presented in Table 2.

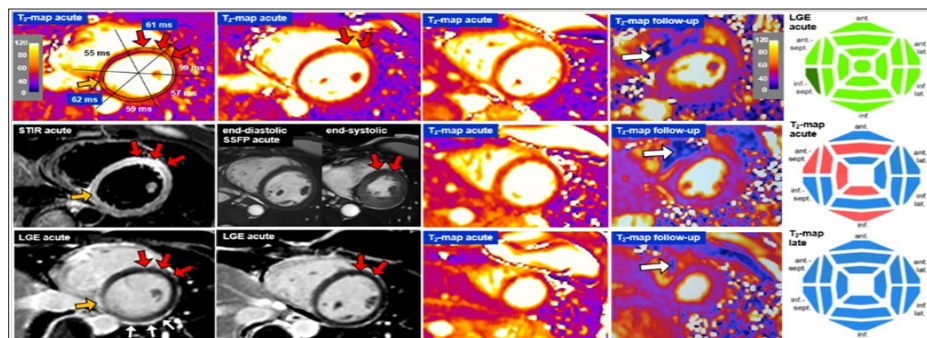


Figure 7: Example: Acute myocarditis. 44-year old female patient with complete AV block and chest discomfort, no elevation of troponines. The CMR examination reveals extensive myocardial edema by elevated T₂ values, see also bulls eye. A non-quantitative STIR image demonstrates increased signal in concordance with the T₂ mapping images (red arrows). Minimal necrosis in the basal infero-septal segment (orange arrow) is detected by late gadolinium enhancement (LGE), while the edematous hypokinetic anterior wall (red arrows on SSFP images) is free of necrosis. Enhancement of the pericardium (white arrows) is compatible with mild pericarditis. Endomyocardial biopsies confirm lymphocytic infiltrates, i.e. myocarditis. Follow-up CMR 4 months later confirms complete resolution of myocardial edema in all LV segments. White arrows on the follow-up T₂ maps indicate artifacts due to an electrode in the RV of an MR-conditional ICD.

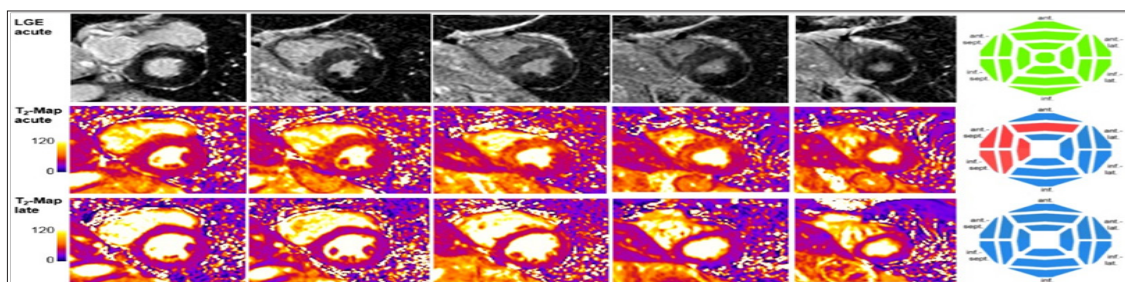
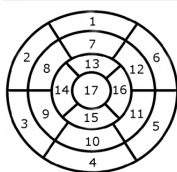


Figure 8: Example: After an emotional stress situation at work, a 60-year old patient presents with chest pain, slightly elevated troponine, and septal hypokinesia on echocardiography. Invasive coronary angiography revealed normal coronary arteries. In the acute phase, T2-mapping on CMR demonstrates extensive myocardial edema (see also bull's eye plot) associated with septal hypokinesia, normal global EF of 62%, and concentric remodeling (LVEDV 112ml, LV mass 130g), while LGE imaging excluded necrosis of the LV myocardium. A follow-up CMR study showed complete resolution of myocardial edema on T2-mapping (see bull's eye), a reduction in LV mass to 114 g, and slight increase in LVEDV and LVEF to 133ml and 72%, respectively. Considering the patient's history and the normalization of all CMR pathologies, the most likely diagnoses are takotsubo cardiomyopathy or a severe spasm of the left anterior descending coronary artery.

Table 2: Upper limit of normal for T2 values of the 2D-BH technique - 17-Segment Model.

S.NO	Segment	Upper limit of Normal T ₂ (ms)
1	Basal Anterior	55.3
2	Basal Antero-Septal	53.2
3	Basal infero-septal	51.2
4	Basal Inferior	54.3
5	Basal Infero-Lateral	52.2
6	Basal antero-lateral	52.4
7	Mid Anterior	57.4
8	Mid Antero-Septal	55.3
9	Mid Infero-Septal	54.1
10	Mid Inferior	53.6
11	Mid Infero-Lateral	52.9
12	Mid Antero-Lateral	54.2
13	Apical Anterior	59.9
14	Apical Septal	53.1
15	Apical Inferior	57.1
16	Apical Lateral	54.9
17	Apex	--



Upper limit of normal is upper 95%-confidence limit. The normal values of the 16 segments were calculated as follows: Values of slice 3 (=mid-ventricular slice) yielded values for segments 7 to 12. Averaging slice 1 and 2 (basal slices) yielded values for segments 1 to 6 and averaging slice 4 and 5 yielded values for segments 13 to 16. For the apical slice, segment 14 was the mean of the 2 septal segments of slice 4 and 5, segment 16 was the mean of the 2 lateral segments of slice 4 and 5.

Conclusion

Most reliable T2 results with a low coefficient of variation were obtained by the 2D-BH technique. The 2D-FBNav technique is considered as an alternative if breath-holding capacity is not sufficient. The 3D-FBSN technique is not at the same level of robustness as the breath-holding techniques and further refinement is needed before it can be considered an alternative for clinical application.

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