



Joint myocardial T1 and T2 mapping

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Joint myocardial T₁ and T₂ mapping

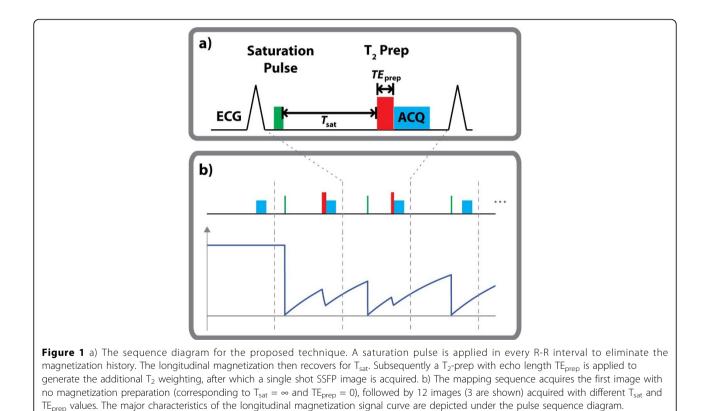
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Background

Recent studies suggest that quantitative myocardial T_1 mapping allows assessment of focal and diffuse fibrosis in the myocardium [1]. Quantitative T_2 mapping has also been proposed to overcome challenges associated with T_2 weighted imaging [2]. These maps are traditionally acquired with different sequences, necessitating

image registration to evaluate them jointly. A sequence that can jointly estimate T_1 and T_2 maps has been proposed [3], but it requires multiple relaxation cycles, which necessitates a lengthy free-breathing acquisition. In [4], an alternative joint estimation sequence was proposed based on the inversion-recovery SSFP curve. In this study, we sought to develop a saturation-recovery

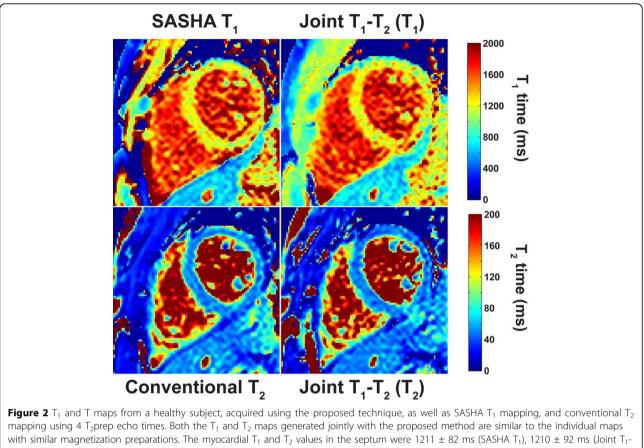


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with similar magnetization preparations. The myocardial T_1 and T_2 values in the septum were 1211 ± 82 ms (SASHA T_1), 1210 ± 92 ms (Joint T_1 - T_2), 49.0 ± 5.8 ms (conventional T_2) and 47.3 ± 6.5 ms (Joint T_1 - T_2) for each technique. The methods generated with the proposed method were acquired in the same time as each individual map, and are jointly registered by design.

based heart-rate independent sequence that can be acquired in a breath-hold and that allows for simultaneous estimation of quantitative T_1 and T_2 maps.

Methods

The sequence diagram is depicted in Figure 1. At every heartbeat, a saturation pulse is applied to eliminate the magnetization history. The longitudinal magnetization then recovers for T_{sat} based on the T_1 value. Subsequently a T_2 -prep pulse [5] with echo length TE_{prep} is applied to generate the additional T_2 weighting, after which a single shot SSFP image is acquired. The process is repeated for 13 heartbeats with various (T_{sat}^{k} , TE_{prep}^{k}) corresponding to heartbeat k, to sample different T_1 - T_2 weighted images. The first heartbeat is acquired with no magnetization preparation.

The T_1 and T_2 maps were estimated jointly by voxelwise least squares fitting to a 4-parameter signal model, A (1- $\exp(-T_{sat}{}^k/T_1)$) $\exp(-TE_{prep}{}^k/T_2)$ + B. Phantom imaging of 14 vials with different T_1/T_2 values were performed and compared to inversion-recovery and CPMG spin-echo references, respectively. Breath-held in-vivo imaging was performed on 5 healthy adult subjects, and the maps were compared to SASHA T_1 maps [6] and to T_2 maps [7].

Results

Phantom imaging resulted in T_1 and T_2 values not significantly different than the references (P = 0.481 and 0.479 respectively). Example in-vivo T_1 and T_2 maps are depicted in Figure 2, comparing various techniques. The T_1 and T_2 values were in good agreement (1211 ± 82 ms vs. 1210 ± 92 ms for T_1 ; 49.0 ± 5.8 ms and 47.3 ± 6.5 ms for T_2).

Conclusions

The proposed sequence allows for the simultaneous estimation of accurate and jointly registered quantitative T_1 and T_2 maps with similar accuracy and precision to saturation-based T_1 mapping and to T_2 mapping of same duration.

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References

- 1. Mewton: JACC 2011.
- 2. Giri: JCMR 2009.
- 3. Sinclair: JMRI 2009.
- 4. Santini: MRM 2014.
- 5. Brittain: MRM 1995.
- 6. Chow: MRM 2013.
- 7. Akçakaya: MRM 2014.

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