

POSTER PRESENTATION

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# Self-navigated three-dimensional cardiac $T_2$ mapping at 3T

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## Background

Cardiac  $T_2$  mapping using a variable  $T_2$  preparation module ( $T_2$ Prep) has recently gained attention for its ability to quantify the extent of edema (Giri, JCMR 2009). Due to time constraints, the  $T_2$  maps are commonly acquired as one or several two-dimensional slices, while the underlying pathology has a three-dimensional (3D) structure. The next logical step would therefore be to exploit recent hardware and software advances to directly acquire 3D  $T_2$  maps. To this end, we tested the feasibility of using a self-navigated 3D radial acquisition with a variable  $T_2$ Prep for 3D  $T_2$  mapping at 3T.

## Methods

Approval was obtained from the institutional review board. A 3D self-navigated undersampled balanced steady-state free precession (bSSFP) sequence (TR/TE=2.6/1.33ms, matrix 128<sup>3</sup>, flip angle 70 ) with a spiral phyllotaxis radial 3D trajectory (Piccini, MRM 2011) was implemented on a 3T clinical system (Skyra, Siemens AG). This self-navigated pulse sequence allows free breathing acquisitions with 100% scan efficiency, while ECG triggering every 2 heartbeats and TE<sub>T<sub>2</sub>Prep</sub>=60/30/0ms allow for a total acquisition time of ~18min with an isotropic spatial resolution of (1.7mm)<sup>3</sup>. The datasets were registered using 3D affine registration (Studholme, Med Image Anal 1996). Through Bloch equation simulations, the heart-rate-dependent  $T_1$  -relaxation-related offset in the  $T_2$ -fitting equation was ascertained. Subsequently, the validity and accuracy of the  $T_2$  fitting was tested in a phantom whose "true"  $T_2$  values were previously determined. The in vivo robustness of the  $T_2$

determination was then tested in 9 healthy adult subjects. Finally, the sequence was applied for the detection of edema in a 75-year-old male infarct patient after revascularization of his proximal left circumflex.

## Results

The Bloch equation simulations of the pulse sequence demonstrated that the input  $T_2$  value could be accurately fitted from the magnetization  $M$  with the equation [ $M=M_0e^{-TE/T_2} + 0.08M_0$ ], while the fitted  $T_2$  had only a ~3% variation over the common range of heart rates (Fig.1A). The phantom  $T_2$  maps demonstrated high homogeneity and fitting accuracy with the 3D sequence matching the 'true' value to within 1% (Fig.1B). The volunteer study (Fig.2A-C) suggested good agreement with previously reported  $T_2$  values at  $T_2=39.3\pm 3.9$ ms (Van Heeswijk, JACC Imaging 2012, in press). A region of significantly elevated  $T_2$  (60.4 $\pm$ 9.1 vs. 41.0 $\pm$ 4.5ms) was identified in the patient in the infero-lateral myocardium of the left ventricle (Fig.2D,E), consistent with the findings on X-ray coronary angiography.

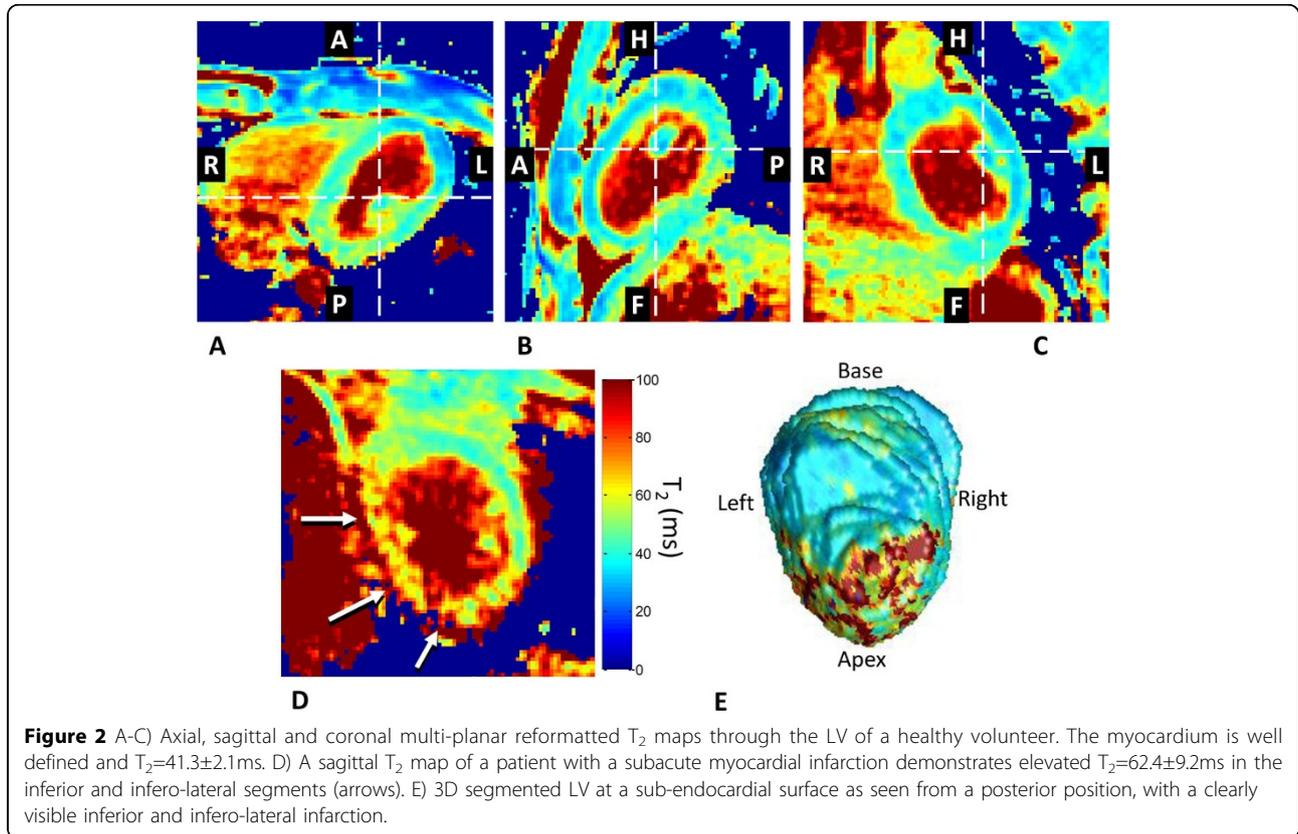
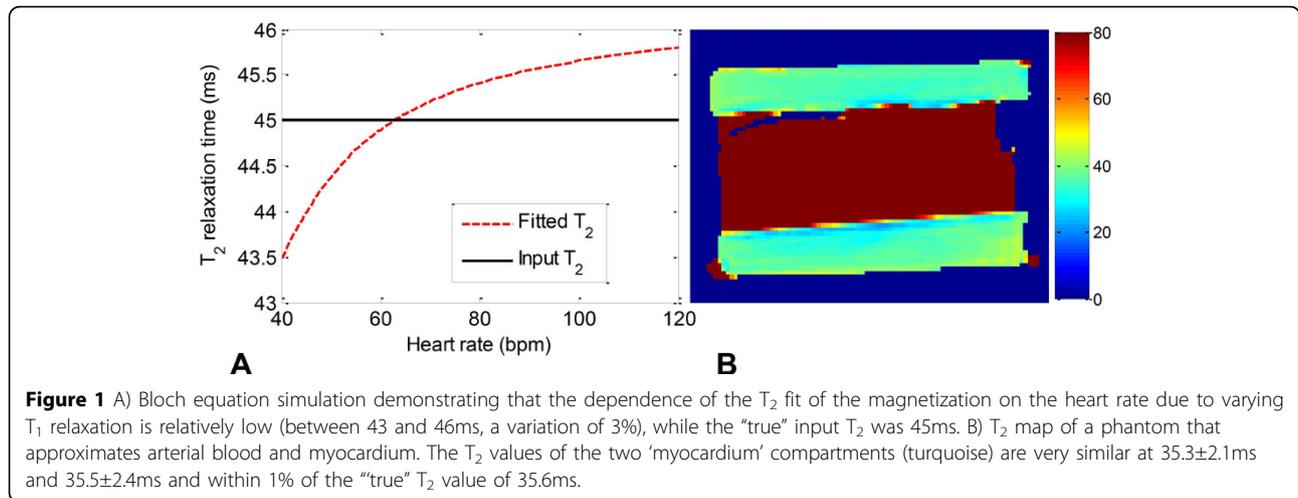
## Conclusions

The proposed technique provides an easy and time-efficient way to obtain accurate isotropic  $T_2$  maps of the whole heart. Accurate  $T_2$  values were obtained in the phantom, while those in volunteers are consistent with previously reported values. The preliminary patient study demonstrated elevated  $T_2$  in the infarcted region as expected.

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