von Knobelsdorff-Brenkenhoff *et al. Journal of Cardiovascular Magnetic Resonance* 2013, **15**(Suppl 1):P37 http://www.jcmr-online.com/content/15/S1/P37

POSTER PRESENTATION



Open Access

Segment-based myocardial T1 and T2 mapping at 3T: feasibility and normal values

Florian von Knobelsdorff-Brenkenhoff^{1,2*}, Matthias A Dieringer^{1,2}, Marcel Prothmann^{1,2}, Andreas Greiser³, Thoralf Niendorf^{1,4}, Jeanette Schulz-Menger^{1,2}

From 16th Annual SCMR Scientific Sessions San Francisco, CA, USA. 31 January - 3 February 2013

Background

Myocardial T1 and T2 mapping using cardiovascular magnetic resonance imaging (CMR) is promising to improve disease detection and monitoring. We applied T1 and T2 mapping at 3T to study the technical feasibility and provide reference values in healthy volunteers.

Methods

Sixty healthy volunteers (30 males / 30 females, 20 in each age group: 20-39 years, 40-59 years, 60-80 years) underwent T1 and T2 mapping of the left ventricle in 3 short axis slices. For T2 maps, 3 single shot steady state free precession (SSFP) images with different T2 preparation times (0, 24, 55ms) were acquired (TE 1.0 ms, TR 2.4 ms, voxel 1.9 x 1.9x6 mm³). For T1 maps, Modified Look-Locker Inversion Recovery (MOLLI) technique with 11 single shot SSFP images was used before and after injection of gadolinium contrast (pre-contrast: TE 1.0 ms, TR 2.6 ms, voxel size 1.4-1.7 x 1.4-1.7 x 6 mm³). T1 and T2 relaxation times were quantified for each slice and each myocardial segment.

Results

With T2 maps, 97.7% of all segments were diagnostic and 2.3% were excluded (susceptibility artifact, Figure 1a). With T1 maps (pre-/post-contrast), 91.6% / 93.9% were

diagnostic, while 8.4% / 6.1% were excluded (7.7% / 3.2% susceptibility artifact (Figure 1b); 0.7% / 2.2% incorrect motion correction; 0% / 0.7% mistriggering). Mean T2 times and 95% tolerance interval were: base: 44.1ms (39.3-49-5 ms); middle: 45.1 ms (39.9-50.1 ms); apex: 46.9 ms (40.8-53.8 ms). Mean T1 times and 95% tolerance interval pre- and post-contrast were: Base: 1157.1 ms (1074.5-1246.0) and 427.3 ms (363.2-502.7 ms). Middle: 1158.7 ms (1074.0-1250.1 ms) and 411.2 ms (349.9-483.2 ms). Apex: 1180.6 ms (1073.9-1297.9 ms) and 399.7 ms (323.0-494.6 ms). The segmental results are depicted in figure 2. Inter- and intra-observer analysis of T2 (r=0.95; r=0.95) and T1 (r=0.91; r=0.93) demonstrated excellent agreement (p<0.0001).

Conclusions

T2 and T1 mapping at 3T was technically feasible, reference values for each myocardial segment are now provided, and observer dependency was low. However, 3T-related susceptibility artifacts and the relatively wide tolerance interval of T2 and T1 times must be considered during interpretation.

Funding

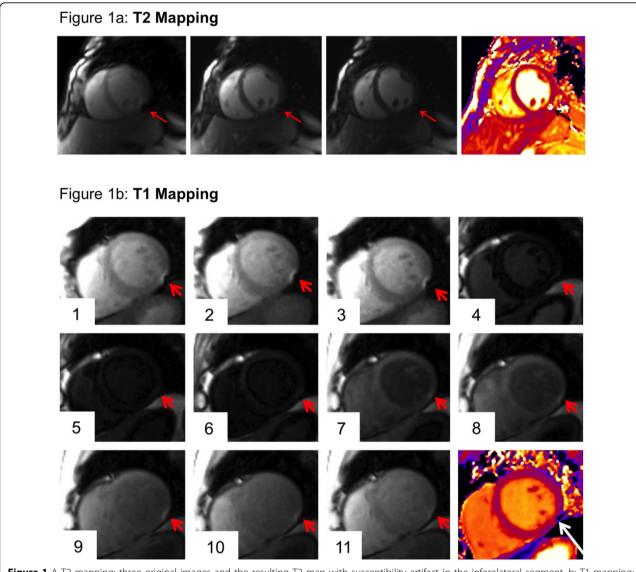
This project is supported by the Else Kröner-Fresenius Stiftung (Bad Homburg, Germany).

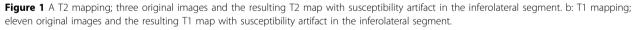
¹Berlin Ultrahigh Field Facility, Berlin, Germany

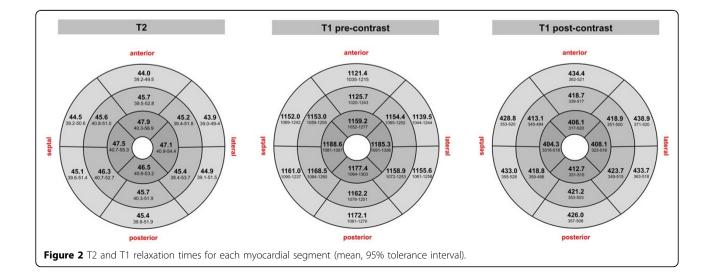
Full list of author information is available at the end of the article



© 2013 von Knobelsdorff-Brenkenhoff et al; licensee BioMed Central Ltd. This is an Open Access article distributed under the terms of the Creative Commons Attribution License (http://creativecommons.org/licenses/by/2.0), which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.







Author details

¹Berlin Ultrahigh Field Facility, Berlin, Germany. ²Working Group Cardiovascular MRI, Experimental and Clinical Research Center (Charite, MDC) and HELIOS Clinics, Berlin, Germany. ³Siemens Healthcare, Erlangen, Germany. ⁴Experimental and Clinical Research Center (Charite, MDC), Berlin, Germany.

Published: 30 January 2013

doi:10.1186/1532-429X-15-S1-P37

Cite this article as: von Knobelsdorff-Brenkenhoff *et al.*: **Segment-based myocardial T1 and T2 mapping at 3T: feasibility and normal values.** *Journal of Cardiovascular Magnetic Resonance* 2013 **15**(Suppl 1):P37.

Submit your next manuscript to BioMed Central and take full advantage of:

- Convenient online submission
- Thorough peer review
- No space constraints or color figure charges
- Immediate publication on acceptance
- Inclusion in PubMed, CAS, Scopus and Google Scholar
- Research which is freely available for redistribution

BioMed Central

Submit your manuscript at www.biomedcentral.com/submit