## Arrhythmia Insensitive Rapid Cardiac T1 Mapping Pulse Sequence: In Vivo Study

Michelle Fitts<sup>1,2</sup>, Elodie Breton<sup>3</sup>, Eugene G. Kholmovski<sup>2,4</sup>, Derek Dosdall<sup>2,5</sup>, Sathya Vijayakumar<sup>2,4</sup>, Kyung P. Hong<sup>1,2</sup>, Ravi Ranjan<sup>2,5</sup>, Nassir F. Marrouche<sup>2,5</sup>, Leon Axel<sup>6</sup>, and Daniel Kim<sup>2,4</sup>

<sup>1</sup>Bioengineering, University of Utah, Salt Lake City, Utah, United States, <sup>2</sup>CARMA Center, University of Utah, Salt Lake City, Utah, United States, <sup>3</sup>ICube, Strasbourg University, Strasbourg, Alsance, France, <sup>4</sup>UCAIR, Department of Radiology, University of Utah, Salt Lake City, Utah, United States, <sup>5</sup>Internal Medicine, University of Utah, Salt Lake City, Utah, United States, <sup>6</sup>Department of Radiology, New York University, New York, New York, United States

**Background:** The gold standard imaging modality for assessment of focal or patchy myocardial infarction/fibrosis is late gadolinium enhancement (LGE) MRI [1,2]. This imaging modality, however, cannot assess diffuse fibrosis because it requires a normal reference tissue to distinguish the fibrosis. LGE cardiac T<sub>1</sub> mapping is the only proven method for quantification of diffuse myocardial fibrosis [3]. The most widely used cardiac T<sub>1</sub> mapping pulse sequence is MOLLI [4], which is based on inversion-recovery (IR) magnetization T<sub>1</sub>. weighting and Look-Locker imaging. Unfortunately, MOLLI is sensitive to heart rate and rhythm and T<sub>2</sub> effects [5] and requires a long breath-hold duration. We present an arrhythmia-insensitive, rapid (AIR) cardiac T<sub>1</sub> mapping pulse sequence, which is also insensitive to T<sub>2</sub> effects, and compare its performance against the conventional IR-based MOLLI method.

**Methods**: We developed the AIR cardiac  $T_1$  mapping pulse sequence based on  $B_1$ -insensitive saturation recovery (SR) [6]  $T_1$ -weighting (insensitive to heart rate, rhythm) and two single-shot balanced steady-state free precession (b-SSFP) image acquisitions (proton density (PD) and  $T_1$ -weighted ( $T_1$ w)) with centric k-space ordering (rapid, insensitive to  $T_2$  blurring) [7,8]. Both MOLLI and AIR pulse sequences were implemented on two 3T whole-body MRI scanners (Tim Trio and Verio, Siemens Healthcare, Erlangen, Germany). Relevant



imaging parameters used for both MOLLI and AIR data acquisitions were: TR = 2.7 ms, TE = 1.1 ms, acquisition  $\overline{\text{matrix}} = 192$  (readout) x 144, slice thickness = 8 mm, flip angle = 35°, field of view (FOV) = 340 mm (readout) x 255 mm, GRAPPA parallel imaging factor R = 1.8, receiver bandwidth = 930 Hz/pixel, and temporal resolution = 217 ms. The MOLLI acquisition was performed in a breath-hold of 17 heart beats, using inversion time (TI) values as specified by Messroghli et al [3]. The AIR image acquisition was performed in a breath-hold of 2-3 heart beats, depending on the heart rate, with SR time delay (TD) = 600 ms. We elected to use TD = 600 ms to achieve a good balance between T<sub>1</sub> sensitivity and signal-to-noise ratio of T<sub>1</sub>w images [9]. T<sub>1</sub> was calculated pixel-wise by dividing the T<sub>1</sub>w image, I<sub>T1w</sub>, by PD image, I<sub>PD</sub>, to correct for the unknown equilibrium magnetization, cancel T<sub>2</sub> effects and compensate for receiver coil inhomogeneity, and then solving the Bloch equation governing T<sub>1</sub> relaxation that describes the ideal SR experiment (Eq.1). Customized software in MATLAB was used to manually segment the regions of interest (ROI) (left ventricular myocardium, blood pool). T<sub>1</sub> was calculated on a pixel-by-pixel basis and averaged within the ROI. For statistical analysis, T<sub>1</sub> measurements by MOLLI and AIR T<sub>1</sub> measurements were assessed using the Bland-Altman analysis. We compared the performances of AIR and MOLLI in ten human subjects and 17 large animals in sinus rhythm pre-contrast and 5, 10, and 15 minutes after contrast agent (Gd-BOPTA, 0.1 mmol/kg dose for numans). O.15 mmol/kg dose for animals) administration. Pulse sequence order was randomized to minimize bias due to their slightly different imaging times relative to the contrast agent administration time.

**Results**: Mean heart rates in humans, dogs, and goats were  $57 \pm 9$  beats-per-minute (bpm),  $86 \pm 15$  bpm, and  $107 \pm 15$  bpm, respectively. Compared with AIR T<sub>1</sub> maps, MOLLI T<sub>1</sub> maps yielded lower values and more spatial blurring (Fig.1). T<sub>1</sub> measurements made by MOLLI and AIR were strongly correlated (Pearson's correlation coefficient =0.99) but in poor agreement (Fig.2, mean

difference =161.77 ms, upper and lower 95% limits of agreements = 348.4ms and -24.4 ms, respectively). Averaging results over 10 humans, AIR T<sub>1</sub> measurement for left ventricular myocardium (1501  $\pm$  69 ms) agreed better than MOLLI T<sub>1</sub> measurement (1198 $\pm$ 46 ms) compared with a previous study which measured T<sub>1</sub> (1471 ms) of an excised heart using a rigorous IR pulse sequence with 35 TI values with 2 averages [10]. For interscan repeatability, the coefficient of repeatability (CR) was 49 ms (5% of mean) for MOLLI and 79 ms (7% of mean) for AIR.



**Conclusions**: Rapid and repeatable cardiac  $T_1$  mapping can be performed using our proposed AIR pulse sequence. This rapid cardiac  $T_1$  mapping pulse sequence may be clinically useful for assessment of myocardial fibrosis in patients. It may be particularly useful for imaging whole heart and for imaging patients with irregular heart rates or difficulty suspending respiration.

**References:** [1] Kim RJ, et al. Circulation 1999; 100(19):1992-2002. [2] Kim RJ, et al. N Engl J Med 2000; 343(20):1445-1453. [3] Mewton N, et al. J Am Coll Cardiol 2011; 57(8):891-903. [4] Messroghli DR, et al. MRM 2004; 52(1):141-146. [5] Gai N, MRM 2012; DOI: 10.1002/mrm.24251. [6] Kim D, et al. MRM 2009; 62(2):300-306. [7] Breton E, et al. SCMR 2011; O107. [8] Lattanzi R, et al. MRM 2011; 66(2):348-355. [9] Haacke et al. Magnetic resonance imaging. New York: Wiley-Liss; 1999: p. 637-667. [10] Stanisz GJ, et al. MRM 2005; 54(3):507-512.