

## **In vivo cardiac diffusion tensor imaging in free-breathing conditions**

Hongjiang Wei, Magalie Viallon, Benedicte Delattre, Vinay Pai, Han Wen, Hui Xue, Christoph Guetter, Marie-Pierre Jolly, Pierre Croisille, Yuemin Zhu

► **To cite this version:**

Hongjiang Wei, Magalie Viallon, Benedicte Delattre, Vinay Pai, Han Wen, et al.. In vivo cardiac diffusion tensor imaging in free-breathing conditions. *Journal of Cardiovascular Magnetic Resonance*, BioMed Central, 2013, 15 (Suppl 1), pp.P231. <inserm-00782655>

**HAL Id: inserm-00782655**

**<http://www.hal.inserm.fr/inserm-00782655>**

Submitted on 30 Jan 2013

**HAL** is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers.

L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.

POSTER PRESENTATION

Open Access

# In vivo cardiac diffusion tensor imaging in free-breathing conditions

Hongjiang Wei<sup>1\*</sup>, Magalie Viallon<sup>1,2</sup>, Benedicte M Delattre<sup>1</sup>, Vinay M Pai<sup>3</sup>, Han Wen<sup>3</sup>, Hui Xue<sup>4</sup>, Christoph Guetter<sup>4</sup>, Marie-Pierre Jolly<sup>4</sup>, Pierre Croisille<sup>1,5</sup>, Yuemin Zhu<sup>1</sup>

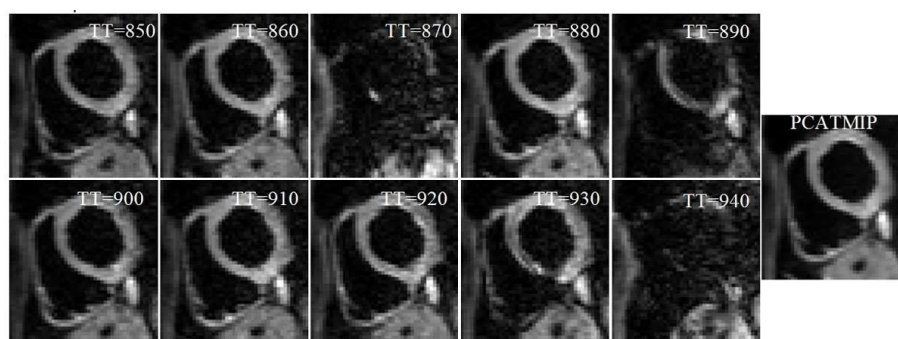
From 16th Annual SCMR Scientific Sessions  
San Francisco, CA, USA. 31 January - 3 February 2013

## Background

The fiber structure of the human heart contributes significantly to efficient ventricular function in the presence of disease. DTI provides a non-invasive approach for the three-dimensional depiction of the myocardial fiber architecture. The biggest problem for in vivo cardiac DTI is the signal loss caused by motion. Recently, to cope with human physiological motion problem, a robust method called PCATMIP was proposed (Rapacchi, Invest radiol 2011) that uses principal component analysis (PCA) filtering to improve the signal-to-noise ratio (SNR) and temporal maximum intensity projection (TMIP) approach to compensate the signal loss. While performing cardiac DTI during subject's breath-hold may be not realistic to apply in clinical routine, achieving acquisitions during subject free-breathing represents an ultimate objective. In this study, our objective was to obtain in vivo DTI parameters of the human heart with free-breathing.

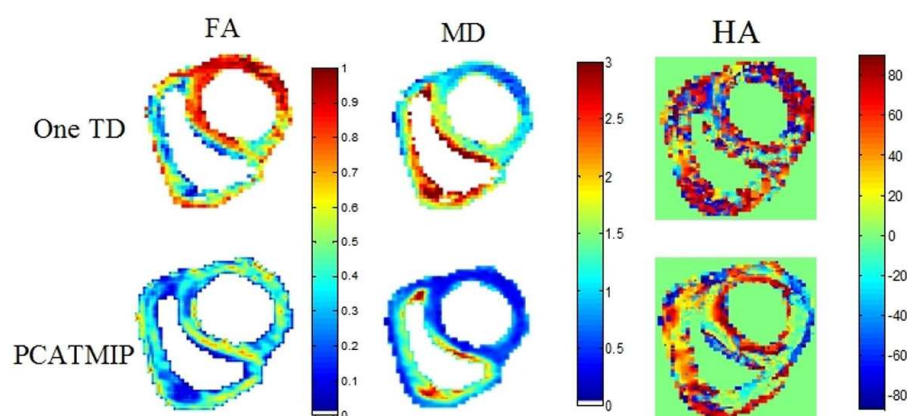
## Methods

To cope with intensity fluctuations arising due to motion, our strategy was to acquire multiple diffusion weighted (DW) images at different time points during the diastole in each consecutive cardiac cycle; after each time frame was acquired, the trigger delay was increased by 10ms. At each trigger delay, we obtained 12 direction DW images and  $b=0$ . We acquired 10 DT-MRI slices across the whole heart. The total scan time is about 20 minutes at an average heart rate of 60bpm. The MRI parameters are: TE/TR=51/100ms, spatial resolution=2.6x2.6x6mm<sup>3</sup>, acceleration rate=2 (GRAPPA), partial Fourier=6/8, base resolution matrix=90x160, and  $b=200$ s/mm<sup>2</sup>. Free-breathing DWI scans were then registered using a non-rigid registration algorithm that preserves high accuracy and consistency of the data. Finally, PCATMIP algorithm was applied to the registered images to obtain motion-reduced ones.



**Figure 1** Free-breathing cardiac DW images from 10 repetitions acquired at different time points and processed using PCATMIP.

<sup>1</sup>Creatis; CNRS (UMR 5220), Inserm (U1044), Insa Lyon, Villeurbanne, France  
Full list of author information is available at the end of the article



**Figure 2** FA, MD, and HA maps comparison between one trigger delay and PCATMIP, the chosen reference trigger delay was the one that exhibited the highest homogeneity and signal intensity in the myocardium.

## Results

Lost signal was recovered and the intensity of the DW images was substantially enhanced (Fig. 1). After processing by PCATMIP, both FA ( $0.43 \pm 0.05$ ) and MD ( $0.75 \pm 0.12 \times 10^{-3} \text{ mm}^2/\text{s}$ ) are smaller than those obtained from 1TD acquisition ( $0.56 \pm 0.05$  and  $1.37 \pm 0.16 \times 10^{-3} \text{ mm}^2/\text{s}$ , respectively) measured over the LV. Coherent helix angle (HA) variations across the ventricle's wall were observed (Fig. 2). The myocardial fiber orientation of the LV showed a circularly symmetric pattern, which reflects the rotation characteristic of cardiac fibers.

## Conclusions

This study demonstrates the feasibility of in vivo cardiac DTI in healthy volunteers. The PCATMIP can be used to minimize the motion-induced signal loss. The proposed acquisition and processing scheme allow to obtain the in vivo DTI parameters while the subject was freely breathing, which opens interesting perspectives for in vivo cardiac DTI clinical applications.

## Funding

This work was supported by the French ANR 2009 (under ANR-09-BLAN-0372-01).

## Author details

<sup>1</sup>Creatis; CNRS (UMR 5220), Inserm (U1044); Insa Lyon, Villeurbanne, France.

<sup>2</sup>Department of Radiology, University Hospitals of Geneva, Geneva, Switzerland.

<sup>3</sup>Imaging Physics Lab, BBC/NHLBI/NIH, Bethesda, MD, USA.

<sup>4</sup>Siemens Corporate Research, Princeton, NJ, USA. <sup>5</sup>Jean-Monnet University, Saint-Etienne, France.

Published: 30 January 2013

doi:10.1186/1532-429X-15-S1-P231

Cite this article as: Wei et al.: In vivo cardiac diffusion tensor imaging in free-breathing conditions. *Journal of Cardiovascular Magnetic Resonance* 2013 15(Suppl 1):P231.

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

Submit your manuscript at  
www.biomedcentral.com/submit

 BioMed Central