

**WORKSHOP PRESENTATION**

**Open Access**

# Fast, heart-rate independent, whole-heart, free-breathing, three-dimensional myocardial BOLD MRI at 3T with simultaneous $^{13}\text{N}$ -ammonia PET validation in canines

Hsin-Jung Yang<sup>1\*</sup>, Damini Dey<sup>1</sup>, Jane M Sykes<sup>2</sup>, John Butler<sup>2</sup>, Behzad Sharif<sup>1</sup>, Debiao Li<sup>1</sup>, Sotirios Tsaftaris<sup>4</sup>, Xiaoming Bi<sup>3</sup>, Piotr Slomka<sup>1</sup>, Frank S Prato<sup>2</sup>, Rohan Dharmakumar<sup>1</sup>

From 19th Annual SCMR Scientific Sessions  
Los Angeles, CA, USA. 27-30 January 2016

## Background

Myocardial BOLD MRI is a non-contrast approach for examining myocardial perfusion. Although recent developments have shown promising technical advancements, current myocardial BOLD MR methods are still limited by: (a) poor spatial coverage; (b) imaging confounders; and (c) imaging artifacts, particularly at 3T. To address these limitations, we developed a heart-rate independent, free-breathing 3D  $T_2$  mapping technique at 3T that utilizes near 100% imaging efficiency, which can be completed within 3 minutes. We tested our method in a canine model and validated our findings with simultaneously acquired  $^{13}\text{N}$ -ammonia PET perfusion data in a whole-body PET/MR system.

## Methods

Canines with and without LAD stenosis ( $n = 11$ ) were studied in a PET/MR system. The proposed sequence was prescribed at rest and under adenosine stress (140 mg/min/kg). Dynamic  $^{13}\text{N}$ -ammonia PET scans were acquired for validation purpose. PET images were analyzed using qPET software. In healthy dogs, mean myocardial  $T_2$  ( $T_{2\text{avg}}$ ) were measured at rest and stress and mean myocardial blood flow ( $Q_{\text{avg}}$ ) were derived from PET images in the corresponding slices. Myocardial BOLD Response (MBOLDR =  $T_{2\text{avg}}$  (stress): $T_{2\text{avg}}$  (rest)) and perfusion reserve (MPR =  $Q_{\text{avg}}$  (stress): $Q_{\text{avg}}$  (rest)) were computed and compared. In the stenosis study, the affected regions were identified from the PET

images and matched to the corresponding slices in BOLD data. Mean myocardial  $T_2$  and myocardial perfusion were measured at rest and stress in the affected and remote territories. MBOLDR and MPR from affected and remote regions were computed and compared against to each other.

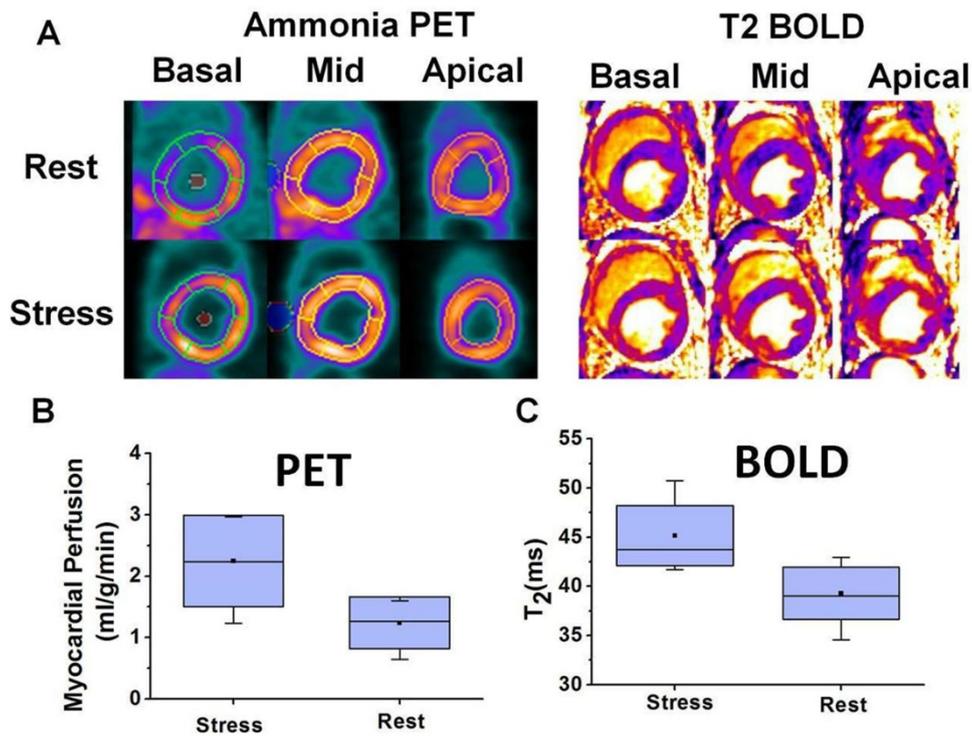
## Results

A representative set of BOLD and PET images acquired from a healthy dog under rest and stress are shown in Figure 1.  $T_{2\text{avg}}$  measured under stress were significantly higher than at rest ( $T_{2\text{avg}}$ :  $38.5 \pm 1.0$  ms (rest) vs.  $44.4 \pm 3.1$  ms (stress),  $p < 0.05$ ). As expected,  $Q_{\text{avg}}$  were significantly higher during adenosine stress relative to rest ( $Q_{\text{avg}}$ :  $0.8 \pm 0.1$  ml/mg/min (rest) vs.  $2.0 \pm 0.9$  ml/mg/min (stress);  $p < 0.05$ ). Linear regression of MBOLDR and MPR showed high correlation ( $R^2 = 0.67$ ,  $p < 0.05$ ). In Figure 2, a set of PET and BOLD images from a dog with LAD stenosis acquired during stress are presented (A and C). Perfusion defect was consistently observed in the LAD territory from both PET and BOLD images. Panel B shows MPR was significantly higher in the remote regions ( $2.8 \pm 1.7$ ) compare to the affected regions ( $1.4 \pm 1.0$ ),  $p < 0.05$ . Significant higher MBOLDR was also observed in panel D (Remote:  $1.09 \pm 0.04$ , Affected:  $1.00 \pm 0.03$ ,  $p < 0.05$ ).

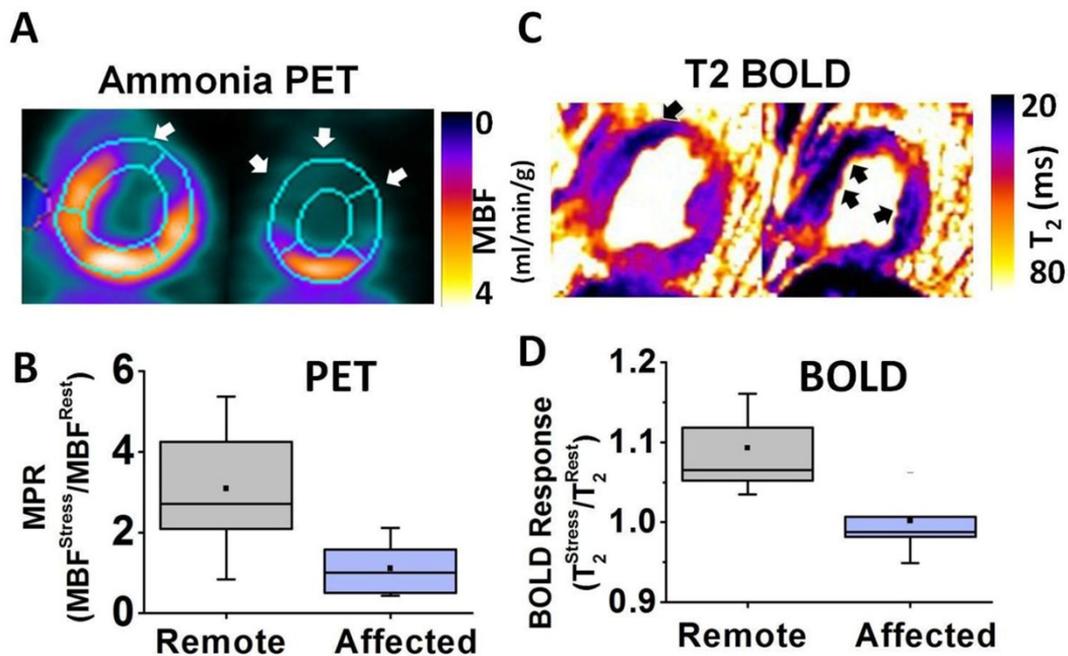
## Conclusions

The proposed BOLD CMR approach permits rapid whole LV assessment of BOLD changes between rest and adenosine stress. The BOLD responses were very closely correlated with PET perfusion, suggesting that the proposed BOLD CMR method is a viable approach

<sup>1</sup>Cedars Sinai Medical Center, Los Angeles, CA, USA  
Full list of author information is available at the end of the article



**Figure 1** PET MBF versus Myocardial  $T_2$  in Healthy Canines. Representative short-axis PET images of myocardial blood flow and myocardial  $T_2$  maps at rest and stress are shown in panel A. Both BOLD and PET images demonstrate significant signal elevation during stress compared to rest. Rest and stress Mean myocardial blood flow and Myocardial  $T_2$  from matched slices are compared in panel B and C, respectively. An average of 2.5 fold increase of myocardial blood flow during stress is observed (panel B) along with a 15%  $T_2$  elevation presented in panel C.



**Figure 2** PET MPR versus Myocardial BOLD Response in Canines with LAD Stenosis. Representative short-axis PET images showing myocardial blood flow (panel A) and myocardial  $T_2$  maps (panel C) acquired during adenosine infusion are shown. Hypoperfused territories are highlighted with arrows in both PET and BOLD images. Note the close correspondence between myocardial perfusion defects identified in the PET and BOLD images. MPR and BOLD Response between remote and affected territories are compared in panels B and D, respectively. MPR in the remote territory was significantly higher than MPR in the affected territories. Similar observations were made with myocardial BOLD response (panel D).

for imaging myocardial perfusion. The method remains to be validated in patients.

#### Authors' details

<sup>1</sup>Cedars Sinai Medical Center, Los Angeles, CA, USA. <sup>2</sup>Lawson Health Research Institute, London, ON, Canada. <sup>3</sup>Siemens USA, Los Angeles, CA, USA. <sup>4</sup>Institute for Advanced Studies Lucca, Lucca, Italy.

Published: 27 January 2016

doi:10.1186/1532-429X-18-S1-W2

**Cite this article as:** Yang *et al.*: Fast, heart-rate independent, whole-heart, free-breathing, three-dimensional myocardial BOLD MRI at 3T with simultaneous <sup>13</sup>N-ammonia PET validation in canines. *Journal of Cardiovascular Magnetic Resonance* 2016 **18**(Suppl 1):W2.

**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](http://www.biomedcentral.com/submit)

