

POSTER PRESENTATION

Open Access

Integrated quantitative first-pass cardiac perfusion MRI protocol

Elodie Breton*, Daniel Kim, Sohae Chung, Leon Axel

From 2011 SCMR/Euro CMR Joint Scientific Sessions Nice, France. 3-6 February 2011

Introduction

The proposed contrast-enhanced first-pass perfusion cardiovascular MR (CMR) protocol integrates some recent technical MRI advances towards quantitative analysis of perfusion CMR images: fast multi-slice pulse sequence [1], robust saturation pulse [2], accurate dedicated AIF imaging [3], signal-to-concentration modeling [4], and higher SNR at 3T.

Purpose

To evaluate an integrated first-pass perfusion cardiovascular MR (CMR) protocol designed to determine absolute contrast-agent concentrations in blood and tissues.

Methods

A multi-slice saturation recovery (SR) pulse sequence with sequential SR time delays (TD) after a non-selective saturation pulse [2] was implemented at 3T (Fig. 1). The rationale for this acquisition scheme was to acquire a dedicated arterial input function (AIF) image with a short TD (50ms) in the aortic root and short-axis myocardial images with longer TD values (~150-400ms), to allow for the different amounts of T₁ shortening expected in blood and wall. First-pass perfusion CMR was performed in 7 volunteers (0.05mmol/kg, Gd-DTPA). A signal-to-concentration model was applied to calculate Gd-DTPA concentrations in blood and tissues [4,5]. A proton density-weighted (PDw) image was acquired in the first heartbeat, without the saturation pulse, in order to normalize the image signal, and obtain a theoretical signal-to-T₁ relationship based on Bloch equation in the center of k-space. Gd-DTPA concentrations were calculated assuming: fast water exchange condition [6], longitudinal relaxivity $r_1=3.8L.mmol^{-1}.s^{-1}$ [7], and baseline T_1 measured with a multi-point SR fit. TurboFLASH imaging parameters included: FOV= 350mm×315mm, slice thickness=8mm, matrix=160×144, in-plane resolution=2.2mm×2.2mm, TE/TR=1.2/2.4ms, flip angle 10°, temporal resolution=114ms, tSENSEx3, centric k-space trajectory, and receiver bandwidth= 1008Hz/pix. Total image acquisition time was 523ms for the acquisition of 4 slices, namely aortic root and SA base, mid, and apex levels, with respective TD values 50-164-278-393ms. Contours for the blood and left ventricle were drawn manually, and the myocardium was divided into 6 (base-mid) or 4 (apex) standard segments.

Results

Representative images at peak contrast in blood and myocardium are shown Fig. 2, along with representative AIF and myocardial segment time-responses. The peak

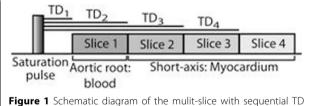


Figure 1 Schematic diagram of the mulit-slice with sequential 1D first-pass perfusion CMR pulse sequence.

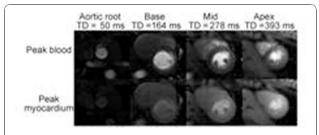
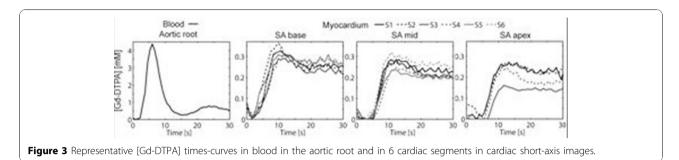


Figure 2 Representative images at peak blood and peak myocardium concentrations.







blood signal was not clipped in the short TD=50ms AIF images. Normalized signal in the myocardium increased along with TD; however similar [Gd-DTPA] were mea-

sured in all 3 short-axis images. First-pass perfusion peak Gd-DTPA concentrations were 3.95±0.080, 0.26 ±0.07mM in the blood and myocardium, respectively.

Conclusions

The proposed integrated first-pass perfusion CMR protocol at 3T produced AIF and myocardial wall Gd-DTPA concentrations consistent with previously published results. Future work includes evaluation of the integrated protocol in cardiac patients.

Published: 2 February 2011

References

- Nagel E, et al: Circulation 2003, 108:432-437.
- 2. Kim D, et al: Magn Reson Med 2008, 62:1368-1378.
- 3. Kim D, Axel LJ: Magn Reson Imaging 2006, 23:81-86.
- 4. Cernicau A, Axel L: Acad Radiol 2006, 13:686-693.
- 5. Breton E, et al: SCMR 2010.
- 6. Dobahue KM, et al: J Magn Reson Imaging 1997, 7:102-110.
- 7. Rohrer M, et al: invest Radiol 2005, 40:715-724.

doi:10.1186/1532-429X-13-S1-P60

Cite this article as: Breton *et al.*: **Integrated quantitative first-pass cardiac perfusion MRI protocol**. *Journal of Cardiovascular Magnetic Resonance* 2011 **13**(Suppl 1):P60.

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

