

POSTER PRESENTATION

Open Access

Intra-MRI extraction of diagnostic electrocardiograms using dynamic feedback from carotidal magnetohydrodynamic voltages

Thomas S Gregory^{1*}, Kevin J Wu¹, Ehud J Schmidt², John Oshinski³, Zion T Tse¹

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

Background

During Cardiac Magnetic Resonance Imaging (CMR), blood plasma electrolytes ejected into the aorta during early systole interact with the static magnetic field of the MR scanner (B₀) to produce a Magnetohydrodynamic (MHD) Effect [1]. Electrocardiograms (ECGs) recorded in the presence of B₀ are overlaid with induced MHD voltages (V_{MHD}), leading to non-robustly synchronized imaging [2], and preventing reliable physiological monitoring inside the MRI [3]. Previous methods have sought to separate between V_{MHD} and the true ECG (ECG_{real}) through adaptive filtering [3], independent component analysis [4], and advanced computational models [5]. However, these methods are based on a static model, which has limited accuracy during varying-rate heart-beats. We aim to develop accurate ECG_{real} extraction, as well as real-time Stroke Volume (SV) estimation (proportional to the integral of MHD over systole) [6], with the advantage of physiological feedback through the real-time monitoring of common carotidal MHD, through which the previously static MHD template can be dynamically updated, providing an increased level of accuracy during variations in heart rate, and a continuous estimation of V_{MHD} and ECG_{real}, for the patient's entire duration inside the MRI.

Methods

12-lead ECGs were acquired in two (n=2) healthy volunteers during 20-second breath-holds in a 3T MRI (Figure 1ab) with the heart at isocenter. A secondary monitor was used to acquire a single anterior-posterior bipolar lead placed approximately on the left common

carotid artery (Figure 1c). ECGs were acquired inside (ECG $_{\rm real}$ + V $_{\rm MHD}$) and outside (ECG $_{\rm real}$) the MRI bore during an initial phase in which a static MHD template was extracted, based on lead subtraction. Carotidal MHD was extracted from the single bipolar lead and phase-compensated to match V $_{\rm MHD}$ obtained from the 12-lead ECG. Carotid MHD was subsequently used to adaptively train a Least Mean Squares filter (Figure 1d) to update the MHD template and produce: (1) clean 12-lead ECGs; and (2) an accurate SV estimate [6] (Figure 1e).

Results

The adaptive filtering method was shown to reduce $V_{\rm MHD}$ in the acquired 12-lead ECGs, with residual noise forming <5% of the R-wave amplitude. The method preserved the true S-T segment, while requiring only a short training phase for the 12-lead ECG (10-15 seconds). The Pearson's Correlation Coefficient between Aortic and Carotid MHD increased from 0.51 to 0.88 after the adaptive filtering routine was applied. Figure 1f shows the extracted 12-lead ECG acquired inside the MRI bore after the training phase.

Conclusions

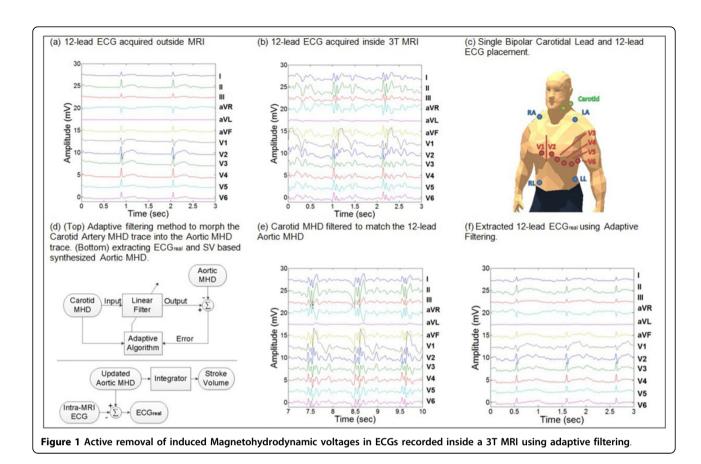
A method to extract true sinus rhythm beats from intra-MRI 12-lead ECGs was presented and shown to provide accurate dynamic measurements of induced $V_{\rm MHD}$ using Carotid artery MHD and ECG_{real} to allow for advanced physiological monitoring inside the MRI.

Authors' details

¹College of Engineering, University of Georgia, Athens, GA, USA. ²Radiology, Brigham and Women's Hospital, Boston, MA, USA. ³Radiology, Emory University Hospital, Atlanta, GA, USA.

¹College of Engineering, University of Georgia, Athens, GA, USA Full list of author information is available at the end of the article





Published: 27 January 2016

References

- 1. Gupta: IEEE Trans BioMed Eng 2008.
- 2. Gregory: MRM 2014.
- 3. Tse: MRM 2013.
- 4. Krug: Comp Card 2012.
- 5. Oster: Comp Method Biomed 2014.
- 6. Gregory: JCMR 2015.

doi:10.1186/1532-429X-18-S1-P214

Cite this article as: Gregory et al.: Intra-MRI extraction of diagnostic electrocardiograms using dynamic feedback from carotidal magnetohydrodynamic voltages. Journal of Cardiovascular Magnetic Resonance 2016 18(Suppl 1):P214.

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

