

Cardiac Diffusion Tensor MRI

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Introduction

Cardiac motion occurs at many scales. Fast and slow. Big and small. MRI has developed over several decades to freeze the motion of the heart or capture its dynamics all of which has led to advanced diagnostic methods for evaluating heart function. Central to our ability to freeze cardiac motion has been increasingly advanced gradient hardware [1]. Initially, this permitted capturing the dynamics of the beating heart in a breath-hold [2]. More recently, the fast and ultrafast gradient hardware permits encoding the diffusive motion of water molecules in a beating heart!

What can we measure?

Imaging the self-diffusion of water in the beating heart – termed cardiac diffusion weighted imaging (cDWI)¹ – presents numerous challenges, but also makes available new mechanisms for generating image contrast. cDWI is poised to provide molecular-

level, quantitative insight to microstructural organization and changes in organization (i.e. remodeling) that occur with disease. For example, sensitivity to changes in the apparent diffusion coefficient (ADC, mm²/s) provides information about how freely water molecules diffuse. Another measure, termed fraction anisotropy (FA), characterizes the directional dependence of the diffusion and is zero for tissues that have isotropic microstructure and can be as high as one for highly anisotropic tissues [3].

How do cDWI parameters change?

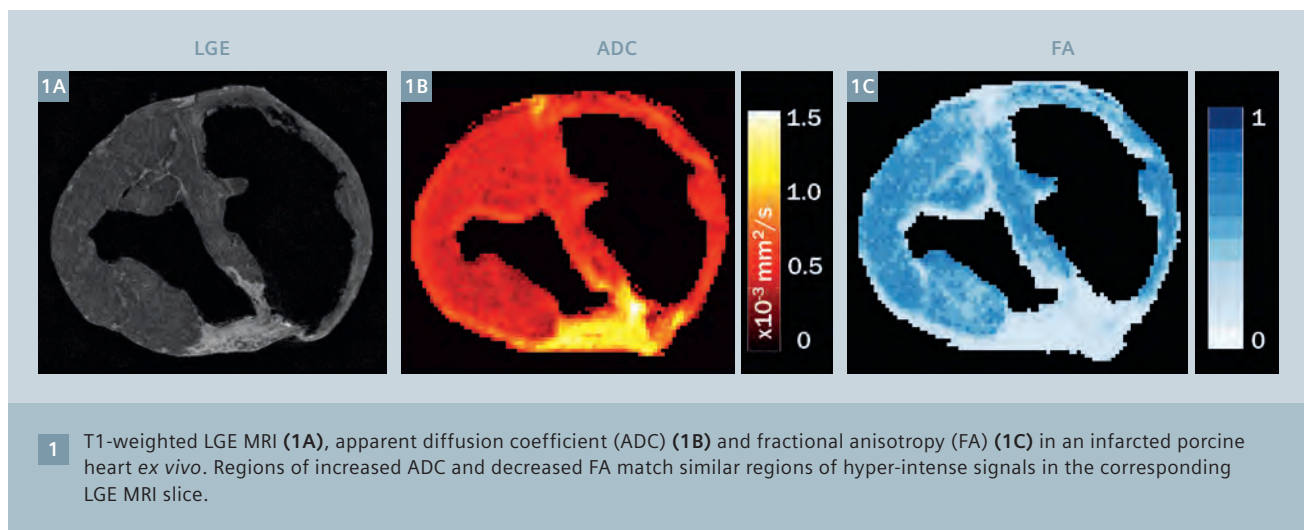
Much of what we know about *in vivo* changes in ADC comes from the neuro literature wherein it is widely appreciated that ADC decreases in acute strokes, pseudo-normalizes within weeks, and subsequently increases above baseline [4]. Currently, less is known about *in vivo* changes to ADC in areas of acute myocardial infarction, but similar findings have been reported [5]. ADC is known to increase significantly in diffuse fibrosis [6] and more so in chronic infarcts [7]. In fact, because

ex vivo cardiac DT-MRI has been practical already for several years we can easily observe the diffusive-level changes in chronic infarcts. Our detailed *ex vivo* work in chronic swine infarcts clearly demonstrates that the ADC is significantly elevated and that the FA is significantly decreased within the infarct, both of which show excellent correlation to the region identified using post-contrast T1-weighted MRI (Fig. 1). As *in vivo* cardiac diffusion weighted imaging becomes more widely available these findings are sure to be reported.

cDWI for myocardial infarcts?

Myocardial infarctions (MI) are the most common form of heart disease and a leading cause of death worldwide [8]. Non-invasive imaging methods for identifying and characterizing affected regions are critical to the diagnosis of MI and the management of post-MI ventricular remodeling [9, 10]. LGE MRI has demonstrated significant clinical value as the gold standard in detecting the location and extent of MI [11]. However, the injection of a GBCA is contraindicated in

¹ The product is still under development and not commercially available yet. Its future availability cannot be ensured.



patients with poor renal function, which poses a significant limitation because this group comprises ~40% of patients with cardiovascular disease and also exhibits increased cardiovascular mortality rates [12]. There does not currently exist a clinically acceptable imaging method for detecting and evaluating MI in this large and high-risk patient group. cDWI may be able to fill this role.

How is the heart microstructurally organized?

The heart is comprised of billions of individual myocytes coordinated into a highly organized network best described as a continuously branching syncytium. In the healthy heart, the organized myocytes are grossly characterized by a transmural change in orientation. The myocytes are further organized into bundles termed sheetlets [13] whose dynamics give rise to ventricular wall thickening [14]. Shortening along the myocyte's long-axis and shearing between the sheetlets are both critical to ventricular performance.

How does cDWI estimate directional information?

cDWI can be used to estimate the local diffusion tensor which comprises information about the magnitudes of diffusion (e.g. ADC and FA) and also the directionality of diffusion. We know from careful *ex vivo* work that the primary eigenvector of the diffusion tensor corresponds to the long axis of the myocytes [15] and that the tertiary eigenvector is generally related to the surface normal of the local sheetlets [16]. Consequently, *in vivo* measurement of the diffusion tensor provides direct insight to changes in microstructural orientations in health and disease [17].

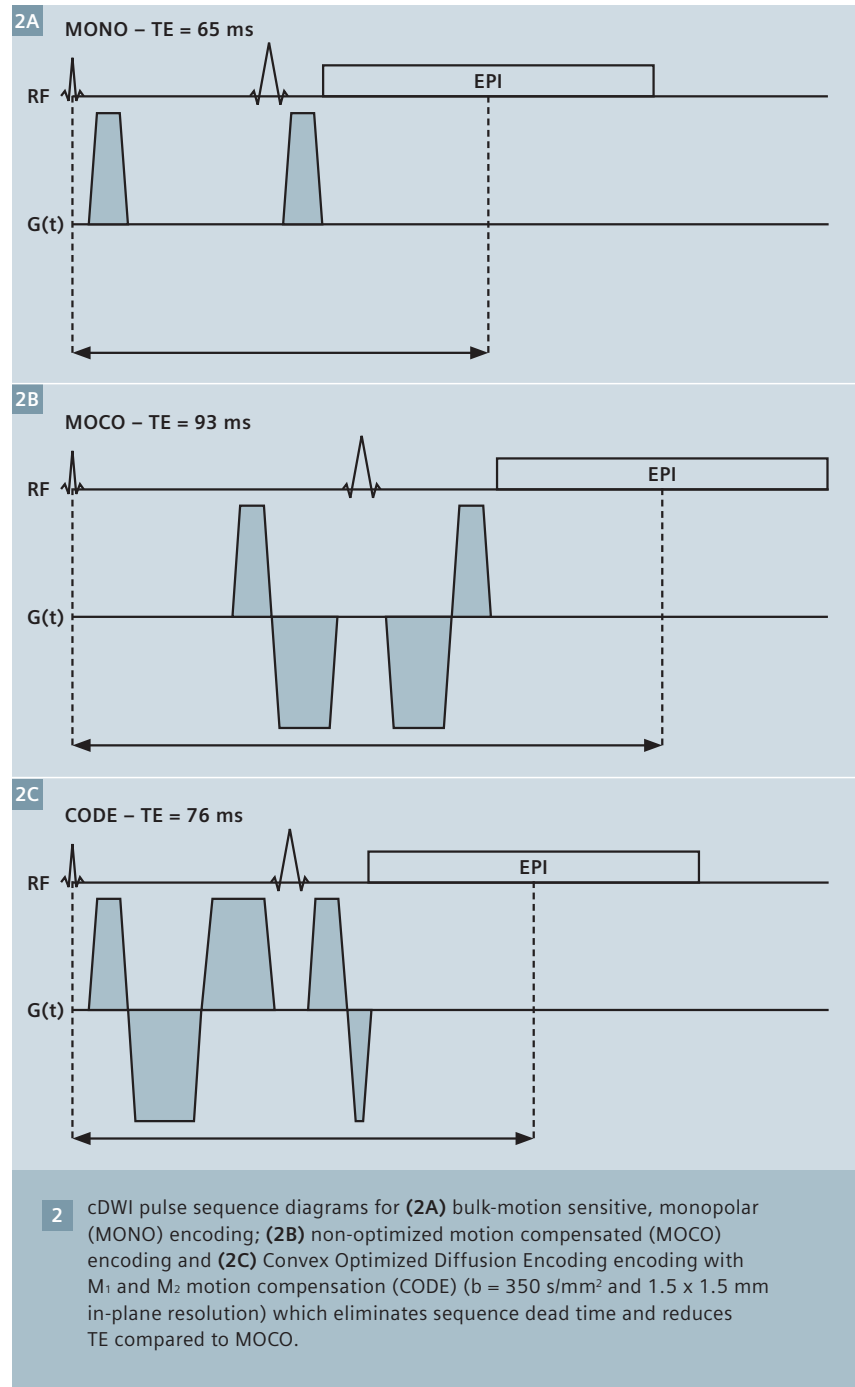
What about the bulk motion sensitivity of DWI?

cDWI measures diffusive motion on the microscopic scale using diffusion encoding gradients with large amplitudes (40 to 80 mT/m) and significant durations (5-10 ms is typical). Consequently, even subtle cardiac bulk motion can significantly corrupt the cDWI measurements unless steps are taken to carefully mitigate bulk motion artifacts. Several approaches have been proposed to address this issue.

For example, diffusion encoding with Stimulated Echo Acquisition Mode (STEAM) [18] limits bulk-motion sensitivity by using very short diffusion encoding gradients distributed across two heartbeats. This approach has been widely adopted and is quite promising, but necessarily has a longer scan time due to two heart-beat encoding, modest SNR because of the stimulated echo, and is confounded by cardiac strain sensitivity and heart rate variability [19].

cDWI with SE-EPI as an alternative to STEAM?

An alternative is single-shot spin-echo (SE) echo planar imaging (SE-EPI) which is insensitive to strain and has better SNR performance and shorter scan times than STEAM [20]. However, SE-EPI cDWI with traditional monopolar encoding is highly sensitive to cardiac bulk motion and requires customized acquisition strategies and post



processing to achieve acceptable image quality [21].

How can bulk motion sensitivity be mitigated?

One approach to mitigating bulk motion artifacts is to use diffusion encoding gradient waveforms that are insensitive to bulk motion, but still sensitive to diffusive motion. Recent work has shown that motion compensated (MOCO) diffusion gradient waveforms with nulled first (M_1) and second (M_2) moments (time-weighted gradient waveform integrals) that are unaffected by bulk linear velocities and accelerations, can also encode diffusion in the presence of cardiac motion [22-24]. When applied to SE-EPI cDWI, MOCO gradients combine bulk motion robustness with strain insensitivity and short scan times. However, moment nulling for the diffusion encoding gradients (i.e. MOCO) necessarily increases the total diffusion encoding duration as compared to monopolar waveforms (Fig. 2). Consequently, conferring bulk motion insensitivity unavoidably

increases the echo time (TE), which reduces SNR. This is exacerbated in high-resolution imaging with long EPI readout intervals, which introduces lengthy dead times within the sequence before the refocusing pulse (Fig. 2). This generally limits cDWI to ~ 2.5 mm in-plane resolution, which is insufficient for evaluation of myocardial infarcts per cardiac MRI guidelines [25]. Herein we describe a technique that we have found to largely overcome this limitation.

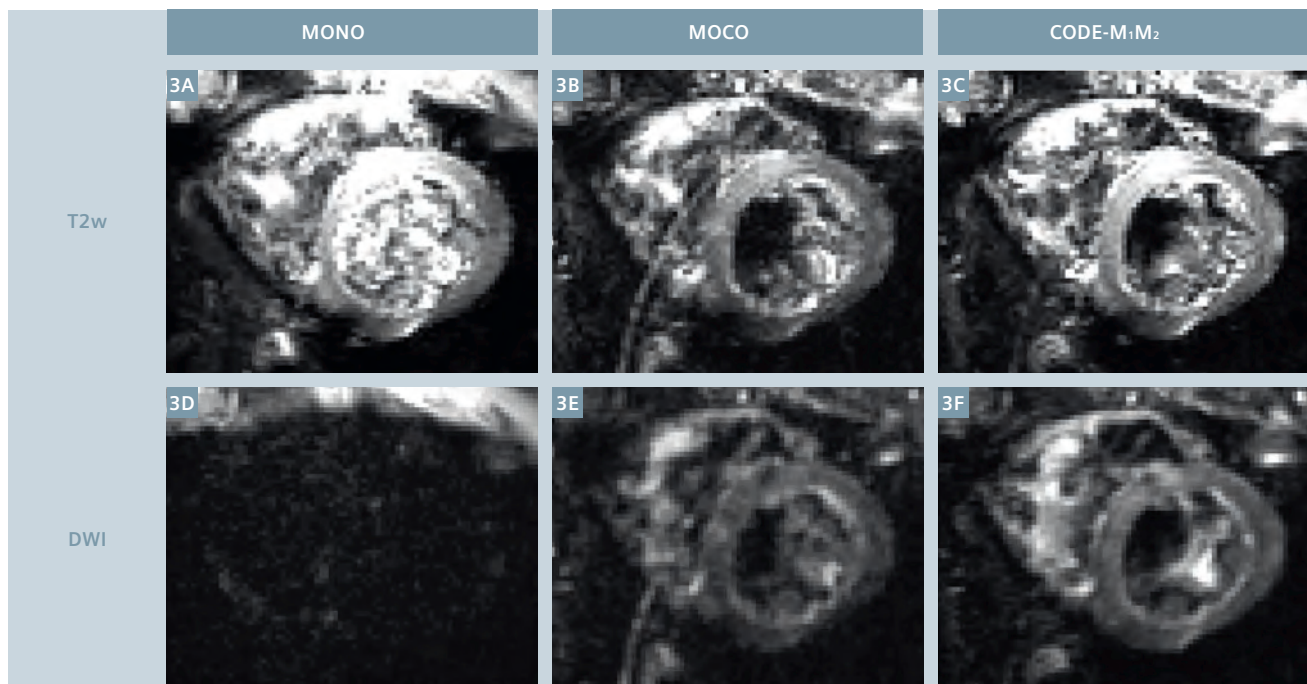
Methods

How are the diffusion encoding gradients designed?

Traditional diffusion encoding gradient waveform design uses an ad hoc approach that involves combining trapezoidal gradient waveforms with closed-form definitions that meet hardware constraints (G_{Max} and SR_{Max}) and ensure the desired properties (M_0 , M_1 , M_2 , b-value), but are agnostic to changes in sequence timing. This tends to lead to inefficiencies in pulse sequence timing that can extend TEs (Fig. 2).

How did we optimize diffusion encoding gradients design?

A general approach to gradient design has been proposed [26] which uses numerical optimization to select the most time-efficient waveform that adheres to all hardware (G_{Max} and SR_{Max}), gradient waveform (M_0 , M_1 , M_2 , b-value, etc.) and pulse sequence (field-of-view, bandwidth, resolution, etc.) constraints. This approach was recently used to design the fastest possible velocity encoding gradients for measuring through plane blood flow [27]. We have applied this concept to more efficiently encode diffusion in the beating heart using convex optimized diffusion encoding (CODE) gradients [28]. This approach calculates the diffusion encoding gradient waveform that minimizes TE for any spatial resolution (Fig. 2). By optimizing the diffusion encoding gradient waveform we can remove all diffusion encoding dead time and achieve M_1 and M_2 moment compensated diffusion encoding in shorter TEs than any previous method. This development permits bulk-motion



3 cDWI images with $b = 0$ (3A–C) and $b = 350$ s/mm² (3D–F) are shown for a healthy volunteer at a systolic cardiac phase with MONO, MOCO, and CODE. MONO cDWI had the highest SNR with $b = 0$ but were severely motion corrupted with diffusion encoding. MOCO and CODE were both robust to cardiac motion with $b = 350$ s/mm² but CODE had higher SNR due to its shorter TE.

insensitive cDWI with 1.5 mm in-plane spatial resolution that matches clinical LGE MRI and enables cDWI to be a practical GBCA-free method for the clinical evaluation of MI.

How are CODE cDWI images acquired?

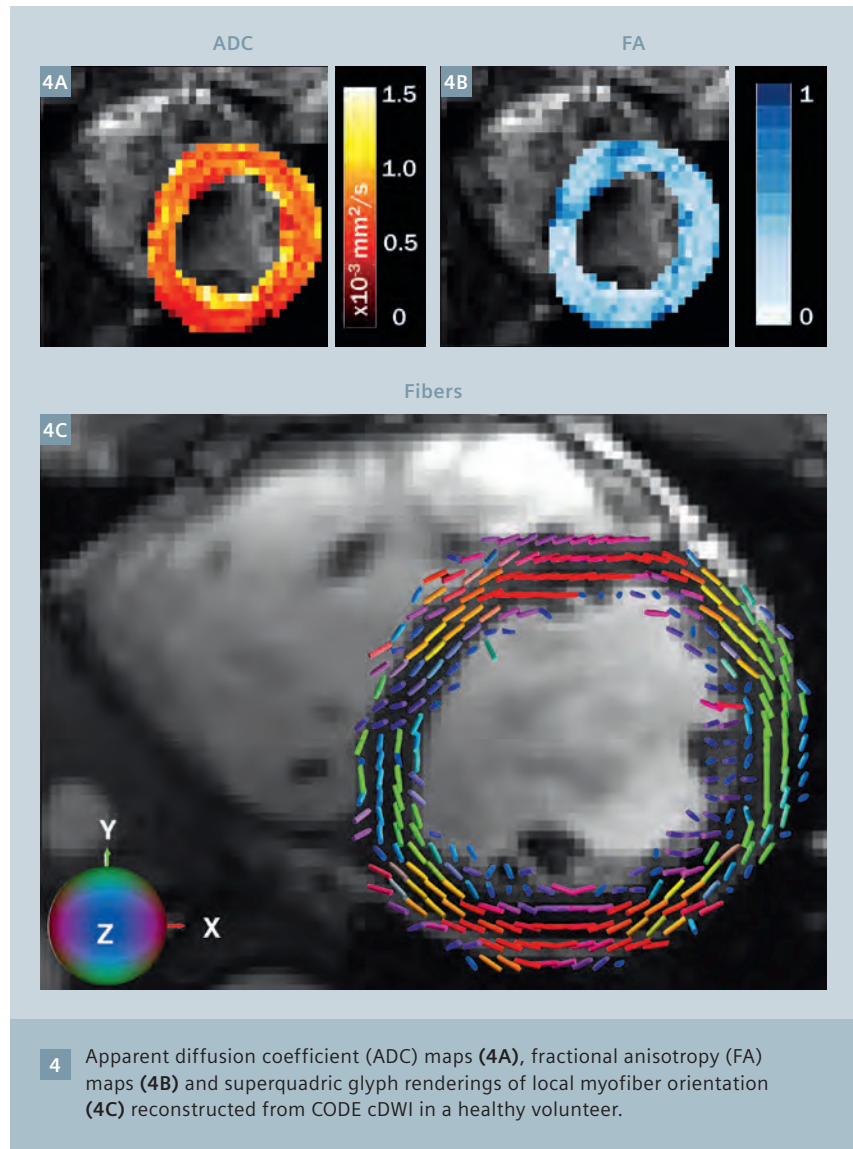
High resolution (1.5 x 1.5 x 5.0 mm) cDWI images ($b = 350 \text{ s/mm}^2$) were acquired in healthy volunteers on a 3T scanner (MAGNETOM Prisma) using three encoding schemes: monopolar (TE = 65 ms), non-optimized MOCO (TE = 93 ms) and CODE (TE = 76 ms) encoding. Diffusion was encoded along three orthogonal directions and three averages were acquired to improve image SNR. Imaging was performed under free breathing with navigator triggering (scan time: ~1 minute per technique). An additional CODE cDTI dataset was acquired with coarser resolution (2.5 x 2.5 x 5.0 mm) for fiber mapping (TE = 59 ms). This acquisition included six diffusion encoding directions and ten signal averages (scan time: ~10 minutes).

How does CODE cDWI compare with other approaches?

Monopolar cDWI resulted in dramatic bulk motion signal dropouts (Fig. 3) that were eliminated with MOCO and CODE. CODE also improved the SNR compared to conventional MOCO (Fig. 3). ADC, FA, and myocyte orientation maps were reconstructed from the CODE cDTI data and demonstrate the high SNR and high quality of the parametric and orientation maps (Fig. 4).

Discussion

We have developed, implemented, and demonstrated a novel approach to cDWI using convex optimized diffusion encoding (CODE) gradients. This approach is robust to cardiac bulk motion and enables sensitivity to diffusion in the beating heart. With the ability to routinely measure diffusion in the beating heart a number of quantitative imaging metrics become available. While the apparent diffusion coefficient (ADC) and the fractional anisotropy (FA) are the principal measures reported in the neuro literature, there are many additional metrics that



may prove specifically useful in the heart [29]. In particular, measures of local myocyte coherence and/or disarray, and changes to myocyte and sheetlet orientations may provide exceptional insight about myocardial function and dysfunction.

Future work will see broader application of CODE cDWI in both healthy volunteers and patients with a variety of clinical indications. Moving forward cDWI may lead directly to insights about microstructural organization in the healthy heart and may become a routine method for evaluating changes to cardiac microstructure without the need for GBCA. Central to the pursuit of this goal is a partnership between

physicists and physicians that enables novel insights arising from the latest MRI technologies.

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