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Mapping Cardiac Fiber Orientations from High-Resolution DTI to High-Frequency 3D Ultrasound

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Abstract

The orientation of cardiac fibers affects the anatomical, mechanical, and electrophysiological properties of the heart. Although echocardiography is the most common imaging modality in clinical cardiac examination, it can only provide the cardiac geometry or motion information without cardiac fiber orientations. If the patient’s cardiac fiber orientations can be mapped to his/her echocardiography images in clinical examinations, it may provide quantitative measures for diagnosis, personalized modeling, and image-guided cardiac therapies. Therefore, this project addresses the feasibility of mapping personalized cardiac fiber orientations to three-dimensional (3D) ultrasound image volumes. First, the geometry of the heart extracted from the MRI is translated to 3D ultrasound by rigid and deformable registration. Deformation fields between both geometries from MRI and ultrasound are obtained after registration. Three different deformable registration methods were utilized for the MRI-ultrasound registration. Finally, the cardiac fiber orientations imaged by DTI are mapped to ultrasound volumes based on the extracted deformation fields. Moreover, this study also demonstrated the ability to simulate electricity activations during the cardiac resynchronization therapy (CRT) process. The proposed method has been validated in two rat hearts and three canine hearts. After MRI/ultrasound image registration, the Dice similarity scores were more than 90% and the corresponding target errors were less than 0.25 mm. This proposed approach can provide cardiac fiber orientations to ultrasound images and can have a variety of potential applications in cardiac imaging.

Keywords

Cardiac fiber orientations; 3D ultrasound; Deformable image registration; Magnetic resonance imaging (MRI); diffusion tensor imaging (DTI); heart imaging
1. INTRODUCTION

Echocardiography is the most common and important imaging modality in cardiac clinical examinations because it is dynamic, noninvasive, fast, and inexpensive [1–3]. Unfortunately, the spatial resolution of ultrasound imaging is not as high as that of MRI or CT. Ultrasound imaging can only provide the geometry or motion information of the heart at the organ level and cannot supply the cardiac fiber orientations at the tissue level. However, these fiber orientations in the heart are important and affect the cardiac anatomy, mechanical properties, and the electrophysiology [4]. These properties then determine the heart pumping functions. Therefore, during the echocardiography imaging in routine clinical examinations, if the patient’s personalized fiber orientations can be mapped to his/her echocardiographic images, it may provide useful and comprehensive parameters such as the geometric shape deformation, stress distributions during a beating period, and the electric action spreading in the heart [4–6]. These parameters can improve or even change cardiac diagnosis or quantitative modeling [7]. Thus, it will be much more useful for long-term examinations of those patients who suffer from chronic cardiac diseases such as hypertrophy or even heart failure. Another important possible application is that these mapped cardiac fibers can help ultrasound-guided cardiac therapies such as the prediction of pacing acute effects during cardiac resynchronization therapy, surgical plans and guidance of cardiac ablation in vivo, as well as the evaluations after these therapies [5, 8]. Recently, significant progress has been made in fast diffusion tensor imaging (DTI) that can measure cardiac fiber orientations in vivo [9]. This progress makes it possible to map personalized cardiac fiber orientations in routine echocardiography examinations. Although cardiac fiber orientations are important for both clinical diagnosis and therapies, mapping cardiac fiber orientation to ultrasound images has not yet been investigated.

In this project, we address the feasibility of mapping the personalized cardiac fiber orientations to 3D ultrasound volumes. First, we apply high-frequency ultrasound imaging with a 30 MHz transducer to scan rat hearts ex vivo. Second, we utilized a 7.0 T MRI scanner to acquire both high-resolution T1 and DTI images from the same heart. Third, after MR imaging, the geometries of the rat hearts are manually segmented from both ultrasound and MR images. We next register the MRI volumes to the 3D ultrasound volumes using both rigid and deformable registration methods. Among these steps, the accuracy of the registration is a key factor [10–17]. Thus, three deformable registration methods [18–21], i.e. Demons, b-spline free form deformation (b-spline), and large deformable diffeomorphic metric mapping (LDDMM), were compared in this study. Finally, according to the cardiac geometry and fiber orientation relationships [5, 6, 22–24], the cardiac fiber orientations obtained by DTI are registered and mapped to the ultrasound volumes based on the extracted deformation fields between MRI and ultrasound.

2. METHODS

2.1 Experiment procedures

The hearts of male Sprague Dawley rates were excised and then quickly perfused by 1 × PBS to clean the residual blood in heart pools and vessels. The hearts were then fixed by 4% phosphate buffered paraformaldehyde (PFA) solution for 14 hours and then were embedded
in 2% agar phantoms for the following imaging procedures. First, the phantoms were settled on the imaging plat and imaged by the Vevo 2100 ultrasound system (FUJIFILM VisualSonics, Inc., Toronto, Canada) with a 30 MHz transducer. B-mode ultrasound images of the hearts in the short-axis view were acquired slice by slice from base to apex at a 0.2 mm thickness. The length of each pixel in the B-mode image was 0.03 mm. Second, the phantoms were placed in a Biospec 7 T MRI system (Bruker Corporation, Massachusetts, USA). A RF coil with an inner diameter of 30 mm was used to transmit/receive the signals. Before DTI imaging, T1 anatomical images were acquired at a voxel resolution of 0.078 × 0.078 × 0.156 mm³. Then, the cardiac fiber orientations were imaged in 60 directions by the spin echo sequence at the 0.234 mm isotropic resolution. The total MRI time was 25 hours for each heart. After the data acquisition, cardiac fiber orientations from DTI were mapped to ultrasound volumes based on the following procedures, as shown in Figure 1.

2.2 Mapping cardiac fiber orientation from DTI to ultrasound

Geometry and fiber orientation reconstruction—Both ultrasound images and T1 weighted MR images were manually segmented using the Analyze software (AnalyzeDirect Inc., Overland Park, KS). Then the binary geometric volumes of the hearts were reconstructed. Based on the segmented heart mask from T1 images, the cardiac fiber orientations were reconstructed from the DTI data.

Geometric registrations from MRI to ultrasound—Although both MRI and ultrasound geometries were for the same heart, their resolutions and imaging angles were different. Thus, the MRI volumes were registered to their corresponding ultrasound ones using a rigid registration method. During the ultrasound imaging procedure, the ultrasound probe pressed the heart and caused deformation. The fiber orientations also affected the ultrasound images and generated imaging artifacts [25, 26]. In order to correct the deformation, three deformable registration methods, i.e. Demons, b-spline, and LDDMM, were implemented to register the MRI and ultrasound images.

2.3 Cardiac fiber orientation mapping

After the geometric registration, the transformation field from MRI to ultrasound was extracted. The reconstructed cardiac fiber orientations were mapped to the ultrasound volumes. During this process, besides the rigid transformation, the cardiac fibers were also reoriented by the strategy of preservation of principle directions (PPD), which was previously used for DTI fiber reorientations [27, 28].

2.4 Evaluations

Quantitative evaluation of the method is conducted by comparing the processed image with the corresponding reference image [29–32]. Two evaluation methods Dice similarity score and target errors are utilized here. The Dice score is used as the performance assessment of the registration, as shown in Figure 2(a). It is computed as follows:

\[
Dice(R, S) = \frac{2 \text{Vol}u\text{m}(R \cap S)}{\text{Vol}u\text{m}(R) + \text{Vol}u\text{m}(S)}
\]  

(3)
where $R$ and $S$ represent the voxel set in the volumes of both registered and corresponding reference volumes, respectively.

The other evaluation method is the target registration error, which calculates the distance between corresponding markers in both images, shown as Figure 2(b). We used the papillary muscles in the hearts as the anatomic markers for the calculation of the target errors. The distance between the mass centers in the corresponding markers was calculated for registration evaluation.

3. RESULTS

3.1 Registrations on canine MRI volumes

We first used the canine cardiac dataset, which was shared by the Center of Cardiovascular Bioinformatics and Modeling (CCBM) at Johns Hopkins University [33], to test the three deformable registration methods, i.e. b-spline, Demons, and LDDMM. Three canine cardiac MRI volumes, which were segmented as binary volumes, were applied to this registration evaluation purpose. Before the deformable registrations, the volumes were registered using rigid-body registration. The corresponding volume Dice similarity scores are listed in Table 1. The results demonstrated that deformable registration improve the match between the two cardiac MRI geometries of the canine heart as compared to rigid registration.

3.2 Registration of ultrasound and MRI volumes of rat hearts

To test the feasibility, two rat hearts were imaged and analyzed in this experiment. First, both ultrasound and T1-weighted MR images were manual segmented as binary volumes using Analyze. Then the cardiac fiber orientations were reconstructed by using the segmented T1 binary volume as the myocardium mask. The processed results are shown in Figure 2.

Registration between ultrasound and MRI volumes—As shown in Figure 4(a–c), the segmented MRI volume and its corresponding segmented ultrasound volume of each rat heart were registered using a rigid-body registration followed by the deformable registration: b-spline, Demons, and LDDMM.

After three different registration approaches, their corresponding volume Dice similarity score are listed in Table 2. We also used the papillary muscles connected to the endocardium of the hearts as the anatomic markers for the registration evaluation. Their corresponding target registration errors are listed in Table 3. It can be seen that the Dice and target errors of rigid registration results are around 85% (Dice) and 0.5 mm (target error). After deformable registration, all Dice scores are higher than 90% and all target errors are less than 0.25 mm. This demonstrates that these deformable registration approaches improved the accuracy of the registration between MRI and ultrasound geometries.

Mapping cardiac fiber orientations to the ultrasound volume—After the geometric registration, based on the transformation field of the deformable registrations, the cardiac fiber orientations were mapped to the ultrasound volumes, as shown in Figure 4(d-f).
This mapping procedure was conducted using two steps: (1) relocation of the DTI point positions and (2) reorientation of the cardiac fibers based on PPD [27].

3.3 Predicting cardiac electricity interventions based on the mapped ultrasound geometry and fiber orientations

Finally, we applied this cardiac fiber orientation mapping method to predict the 3D electrophysiological properties, as shown in Figure 5. Biomechanical simulations and predictions based on images is an important issue in clinical diagnosis and therapies [34]. In order to predict the cardiac electricity interventions, both reconstructed 3D ultrasound mesh geometry (Figure 5(a)) and the mapped cardiac fiber orientations (Figure 5(b)) were utilized for this purpose. Figure 5(c) illustrated the cardiac resynchronization therapy (CRT) conducted by a pacemaker, where the blue pipe was the internally placed electrode. One end of the electrode was connected to the pacemaker and the other one was attached to the myocardium of the heart apex.

Then, the cardiac electricity distributions around the 3D geometry were simulated. The action potentials at three different time points (15 ms, 30 ms and 45 ms) after the impulse of the pacemaker are shown in Figure 5(d–f). The geometric mesh was generated by iso2mesh [35] and the cardiac electricity simulations were simulated by Chaste [36].

4. CONCLUSIONS

In this project, we proposed a cardiac fiber imaging method and test the feasibility of mapping personalized cardiac fiber orientations from MR DTI data to 3D ultrasound volumes, which included data reconstruction, rigid and deformable registration, and fiber reorientation. We also compared the registration results based on different deformable registration methods. Furthermore, we utilized these results to simulate the electrophysiological properties around the heart during CRT process. By providing fiber orientation information on ultrasound images, this technique has the potential to provide measurable information for the examination of the chronic cardiac diseases and for image-guided cardiac therapies.

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References


Figure 1.
Flowchart of mapping cardiac fiber orientation from DTI to ultrasound
Figure 2.
Illustrations of both Dice and target error evaluations. (a) Dice similarity between the registered volumes (yellow and red volumes). (b) Target registration error, the distance (white double arrow) between the mass points (red and blue dots) of two registered targets.
Figure 3.
Processed results of the rat hearts. (a) Ultrasound image. (b) T1-weighted MR image. (c) DTI image. (d) The segmented ultrasound image. (e) The segmented MR image. (f) The reconstructed cardiac fiber orientations by tensor decomposition from DTI data.
Figure 4.
Mapping cardiac fiber orientations to the ultrasound volume based on the deformation field of both ultrasound and MRI volumes. (a) The reconstructed ultrasound volume. (b) The reconstructed MRI volume. (c) Deformed MRI volume after rigid and deformable registration. (d) The cardiac fiber orientations from DTI. (e) The relocated and reoriented fiber orientations based on the deformation fields from MRI to ultrasound. (f) The mapping results of both fiber orientations and the ultrasound volume.
Figure 5.
Predicting cardiac electricity distributions during cardiac resynchronization therapy based on the mapped ultrasound geometry and fiber orientations. (a) The reconstructed mesh volume of ultrasound. (b) The mapped cardiac fiber orientations. (c) The cardiac resynchronization therapy (CRT) conducted by an internal electrode (red line) from a pacemaker. (d–f) The simulation results of the action potentials at three different time points (15 ms, 30 ms and 45 ms) after the impulse of the pacemaker. The colors indicate the action potentials from −90 mV to 40 mV.
### Table 1
Registrations of canine cardiac MRI volumes

<table>
<thead>
<tr>
<th>Registration methods</th>
<th>Volume Dice similarity score (%)</th>
<th>Volume 1 → 2</th>
<th>Volume 2 → 3</th>
<th>Volume 3 → 1</th>
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<tr>
<td>Rigid</td>
<td>57.5</td>
<td>55.7</td>
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<td>72.2</td>
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<td>B-spline</td>
<td>89.1</td>
<td>88.9</td>
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<tr>
<td>Demons</td>
<td>96.6</td>
<td>94.9</td>
<td></td>
<td>97.2</td>
</tr>
<tr>
<td>LDDMM</td>
<td>96.2</td>
<td>95.3</td>
<td></td>
<td>96.8</td>
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</table>
Table 2

Volume Dice similarity Scores for the registration between MRI to ultrasound of the same heart

<table>
<thead>
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<th>Volume Dice similarity Score (%)</th>
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<th></th>
<th></th>
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<td>Rigid</td>
<td>86.5</td>
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<tr>
<td>Rat 1</td>
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<td>95.0</td>
<td>97.8</td>
<td>97.3</td>
</tr>
<tr>
<td>Rigid</td>
<td>86.5</td>
<td>93.7</td>
<td>97.6</td>
<td>98.4</td>
</tr>
<tr>
<td>Rat 2</td>
<td>84.1</td>
<td>95.0</td>
<td>97.8</td>
<td>97.3</td>
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</table>
Table 3
Target registration errors of the registration between MRI to ultrasound of the same rat heart

<table>
<thead>
<tr>
<th>Data sets</th>
<th>Target Errors (mm)</th>
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<th></th>
<th></th>
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</thead>
<tbody>
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<td></td>
<td>Rigid</td>
<td>B-spline</td>
<td>Demons</td>
<td>LDDMM</td>
</tr>
<tr>
<td>Rat 1</td>
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<td>0.19</td>
<td>0.19</td>
<td>0.08</td>
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<tr>
<td>Rat 2</td>
<td>0.42</td>
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