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Heterogeneous growth-induced prestrain in the heart

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Abstract

Even when entirely unloaded, biological structures are not stress-free, as shown by Y.C. Fung's seminal opening angle experiment on arteries and the left ventricle. As a result of this prestrain, subject-specific geometries extracted from medical imaging do not represent an unloaded reference configuration necessary for mechanical analysis, even if the structure is externally unloaded. Here we propose a new computational method to create physiological residual stress fields in subject-specific left ventricular geometries using the continuum theory of fictitious configurations combined with a fixed-point iteration. We also reproduced the opening angle experiment on four swine models, to characterize the range of normal opening angle values. The proposed method generates residual stress fields which can reliably reproduce the range of opening angles between 8.7 ± 1.8 and 16.6 ± 13.7 as measured experimentally. We demonstrate that including the effects of prestrain reduces the left ventricular stiffness by up to 40%, thus facilitating the ventricular filling, which has a significant impact on cardiac function. This method can improve the fidelity of subject-specific models to improve our understanding of cardiac diseases and to optimize treatment options.

Keywords

Residual stress; Prestrain; Opening angle; Patient-specific modeling; Finite element method; Finite strain

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Conflict of interest

None.

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1. Introduction

Biological tissues inherently contain some prestrain, or equivalently residual stress (Vaishnav and Vossoughi, 1987; Fung, 1993; Humphrey, 2002). This has been evidenced in the opening angle experiments, performed originally on the artery in (Chuong and Fung, 1986), and then on the left ventricle in (Omens and Fung, 1990). The exact origin of the prestrain is still an open question: it may be induced by inhomogeneous tissue growth, or the constant exchange of matter within the tissues and associated remodeling, cell death, and renewal, *etc.* (Fung, 1993; Humphrey, 2002). Moreover, prestrain potentially exists across multiple spatial scales, including the sub-cellular (Kumar, Maxwell, et al., 2006; Deguchi, Ohashi and Sato, 2006), cellular, and structural levels (Fung, 1993; Humphrey, 2002; Guo, Lanir and Kassab, 2007).

It is important to distinguish between these different types of prestrain. Indeed, sub-cellular prestrain surely affects the response of individual cells, but is confined to the cells and cannot be characterized on the macroscopic level (Kumar, Maxwell, et al., 2006; Deguchi, Ohashi and Sato, 2006; Webster, Ng and Fletcher, 2014). Conversely, structural, or global, residual stress (imagine a stress-free straight beam glued into a circular shape for which a single cut would release all the residual stress at once) are most probably not seen *in vivo*. In support of this, while original experiments showed that the first cut released most of the residual stress (Liu and Fung, 1989; Han and Fung, 1996), later experiments revealed that subsequent cuts always release additional residual stress (Greenwald, Moore, et al., 1997). Hence, we investigate the prestrain generated on the cellular scale, a scale where living tissues are composed of many different components in frictional contact and constant relative motion. Cellular prestrain is analogous to the one present in engineering materials (e.g., between grains in metals) or between components in composites, which is usually induced by the process or differences in coefficients of thermal expansion. It is present everywhere in living systems and it can have a tremendous impact on their macroscopic behavior.

To address cardiovascular diseases, one approach is the use of personalized models for quantitative diagnostic, prognostic, and treatment optimization (Lee, Genet, et al., 2014b). The current bottlenecks in personalized cardiac modeling include the assessment of myofiber architecture (Toussaint, Stoeck, et al., 2013; Harmer, Pushparajah, et al., 2013), non-invasive pressure measurement, tissue properties orthotropy and regionality (Lee, Wenk, et al., 2011; Xi, Lamata, et al., 2011; Marchesseau, Delingette, et al., 2013). Another important limitation is our understanding of prestrain and residual stress. Indeed, as a result of prestrain, subject-specific geometries of biological structures extracted from medical imaging do not represent the reference configuration required for mechanical analysis, even if the images correspond to an externally unloaded state. To identify the truly unstressed reference configuration, we have to solve an inverse problem that is highly non-linear.

In this paper, we propose a computational method to introduce physiological residual stress fields in subject-specific left ventricular geometries. The proposed method is embedded within the continuum theory of fictitious configurations and uses a fixed-point iteration on

the geometry itself. (Wang, Luo, et al., 2013) recently proposed an alternative approach, also using fixed-point iterations on the geometry to recover a relaxed configuration. However, they use (Shams, Destrade and Ogden, 2011)'s method to introduce residual stress in the invariant-based Holzapfel model (Holzapfel and Ogden, 2009) through additional pseudo-invariants. This implies that (i) the method is limited to invariant-based formulations, (ii) that residual stress is introduced as an *ad hoc* parameter to fit experimental data (Costa, May-Newman, et al., 1997), and (iii) that the resulting residual stress field is not strictly speaking divergence free, *i.e.*, auto-balanced. Here, the residual stress is generated mechanically by the underlying biological process, heterogeneous growth, and is naturally auto-balanced.

2. Methods

2.1. Opening angle experiment

2.1.1. Animal preparation—We performed all animal experiments in accordance with national and local ethical guidelines, including the Principles of Laboratory Animal Care, the Guide for the Care and Use of Laboratory Animals (National Research Council (US) Committee for the Update of the Guide for the Care and Use of Laboratory Animals, 2011) and the National Association for Biomedical Research (Cardon, Bailey and Bennett, 2012), and an approved Indiana University Purdue University Indianapolis IACUC protocol regarding the use of animals in research. Four Yorkshire pigs (55.1 ± 3.0 Kg body weight) of either sex were used in this study. The pigs were fasted overnight and surgical anesthesia was induced with TKX (Telazol 10 mg/kg, Ketamine 5 mg/kg, Xylazine 5 mg/kg) and maintained with 1–2% Isoflurane-balance O₂ during acute, non-aseptic surgery. An introducer sheath was placed percutaneously in the right jugular vein for administration of drugs. The chest was opened through a mid-line sternotomy and the heart was arrested with an IV bolus injection of saturated KCl solution (20 mL) under deep anesthesia and proper anticoagulation with Heparin (100 IU/kg body weight).

2.1.2. Heart preparation—The heart was excised and placed in cardioplegic solution. We cannulated and perfused the left anterior descending, the left circumflex, and the right coronary arteries with a solution (Lin and Yin, 1998) of the following composition in mmol: NaCl, 127; KH₂PO₄, 1.3; MgSO₄, 0.6; NaHCO₃, 25; KCl, 2.3; CaCl₂, 2.5; dextrose, 11.2; BDM (2,3 butanedione monoxime, Sigma, St. Louis, MO), 30. After perfusion, the aorta and pulmonary arteries, the right and left atria, and the right ventricular free wall were carefully cut, keeping an intact left ventricle in the shape of a cone.

2.1.3. Experimental procedure—After the first basal slice was discarded, three 1-cm-thick slices were cut from the left ventricle (basal, equatorial, and apical) and placed in petri dishes containing perfusion solution (see Figure 3). Two cameras were set up to photograph the cross-sectional surfaces of the slice from the top and the bottom surfaces simultaneously. A baseline picture was taken before a radial cut was made in the middle of the left ventricular free wall. Immediately after the cut was made, continuous pictures were taken every five minutes for a period of 30 minutes. The opening angle was measured with a program for image processing and analysis, ImageJ 1.44p (National Institutes of Health).

2.2. Mechanics of pretrained biological systems

2.2.1. Continuum model—We characterize prestrain using the concept of fictitious configurations and introduce a stress-free reference configuration, a residually stressed but mechanically unloaded configuration, and residually stressed mechanically loaded configuration (Rausch and Kuhl, 2013).

Figure 1 illustrates the three configurations, which can also be used to model prestrain induced by other physical mechanisms such as friction (Hild, 1998; Fagiano, Genet, et al., 2014). Under the small deformation hypothesis, this decomposition induces an additive decomposition of the strain itself: The total elastic strain is the sum of prestrain and mechanically-induced strain. Under finite deformations, this decomposition induces a multiplicative decomposition of the deformation gradient: The total elastic tensor $\underline{\underline{F}}^e$ is the product of a prestrain tensor $\underline{\underline{F}}^p$ and the mechanically-induced deformation gradient $\underline{\underline{F}}$:

$$\underline{\underline{F}}^e = \underline{\underline{F}} \cdot \underline{\underline{F}}^p \quad \text{with} \quad \underline{\underline{F}} = \underline{\underline{F}}^e \cdot \underline{\underline{F}}^{p-1} = \nabla \underline{\underline{\varphi}} \quad (1)$$

Inherent to this approach, only the mechanically-induced deformation gradient $\underline{\underline{F}} = \nabla \underline{\underline{\varphi}}$ is a gradient of a vector field, while $\underline{\underline{F}}^e$ and $\underline{\underline{F}}^p$ are generally incompatible. We can then adopt the classical finite deformation theory for nearly incompressible bodies (Holzapfel, 2000), however, now based on the total elastic tensor $\underline{\underline{F}}^e$ instead of the purely mechanically-induced deformation gradient $\underline{\underline{F}}$. This implies that we can express the strain energy potential, which is generally a function of $\underline{\underline{F}}$ and $\underline{\underline{F}}^p$, as a function of $\underline{\underline{F}}^e$ only:

$$\psi(\underline{\underline{F}}, \underline{\underline{F}}^p) = \psi^e(\underline{\underline{F}}^e) \quad (2)$$

The most widely used method to impose incompressibility or quasi-incompressibility consists of further decomposing the elastic tensor into volumetric and isochoric parts:

$$\underline{\underline{F}}^e = \underline{\underline{F}}^v \cdot \overline{\underline{\underline{F}}}^e \quad (3)$$

with the volumetric contribution $\underline{\underline{F}}^v = (J^e)^{1/3} \underline{\underline{1}}$ with $J^e = \det(\underline{\underline{F}}^e)$ and the isochoric contribution $\overline{\underline{\underline{F}}}^e = (J^e)^{-1/3} \underline{\underline{F}}^e$ so that $\overline{J^e} = \det(\overline{\underline{\underline{F}}}^e) = 1$. As a result of this decomposition, the independent variables are now the elastic Jacobian J^e and the isochoric elastic Cauchy-Green tensor $\overline{\underline{\underline{E}}}^e$:

$$J^e = \det(\underline{\underline{F}}^e) \quad \text{and} \quad \overline{\underline{\underline{E}}}^e = \frac{1}{2} (\overline{\underline{\underline{F}}}^{e,t} \cdot \overline{\underline{\underline{F}}}^e - \underline{\underline{1}}) \quad (4)$$

We can then split the elastic strain energy potential into volumetric and deviatoric parts:

$$\psi^e(J^e, \overline{\underline{\underline{E}}}^e) = U(J^e) + \overline{\psi}^e(\overline{\underline{\underline{E}}}^e) \quad (5)$$

For the volumetric part of the strain energy function (Auricchio, Beirão da Veiga, et al., 2013), we could suggest the following expression:

$$U(J^e) = \frac{1}{D_0} \left(\frac{1}{2} ((J^e)^2 - 1) - \ln(J^e) \right) \quad (6)$$

For the deviatoric part, we use a transversely isotropic Fung law (Fung, 1993) adapted for myocardial tissue (Guccione, McCulloch and Waldman, 1991):

$$\overline{\psi^e}(\underline{\underline{E}}^e) = \frac{1}{2} C_0 \left(\exp \left(\underline{\underline{E}}^e : \underline{\underline{B}}_0 : \underline{\underline{E}}^e \right) - 1 \right) \quad (7)$$

where $\underline{\underline{B}}_0$ is a fourth-order constitutive tensor, which takes a diagonal format in Voigt notation (Mehrabadi and Cowin, 1990; François, 1995):

$$\hat{\underline{\underline{B}}}_0 = \text{Diag} (B_{ff}, B_{ss}, B_{nn}, 2B_{fs}, 2B_{fn}, 2B_{sn}) \quad (8)$$

Its individual parameters B_{ff} , B_{ss} , B_{nn} , B_{fs} , B_{fn} , and B_{sn} weigh the influence of the normal fiber, sheet, and normal strains and of the corresponding shear components. Throughout this paper, we use the normal human myocardial properties characterized in (Genet, Lee, et al., 2014) and summarized in Table 1. Note that it is well-known that the Fung law is not polyconvex (Holzapfel, Gasser and Ogden, 2000; Holzapfel and Ogden, 2009), making the proposed model not strictly speaking thermodynamically consistent. However, it is still widely used for the myocardium; for this paper we solve our model implicitly and have not encountered any material instability. Furthermore, the framework presented here for growth-induced residual stress is independent from the choice of the strain energy potential, and a convex potential could be used instead.

These equations can, in principle, allow us to calculate the *in vivo* stress state of the heart. Unfortunately, however, neither the initial nor the grown configuration are known. Only the residually stressed mechanically unloaded configuration and the mechanically loaded configuration can be imaged *in vivo*. In what follows, we propose a fixed-point iteration to computationally retrieve the initial configuration from the *in vivo*-acquired configuration. The core assumption of this algorithm is that residual stresses are a result of biological growth.

2.2.2. Computational model—The constitutive laws have been implemented as a user subroutine into Abaqus/Standard. The subroutine is written in C++, and use the LMT++ library for linear algebra (Leclerc, 2010; Genet, 2010). The implementation is straightforward as outlined in detail in (Rausch and Kuhl, 2013).

2.3. Mechanics of growth-induced, prestrained biological systems

2.3.1. Continuum model—The origin of prestrain is still an open questions today. Here, we investigate the hypothesis that the prestrain is induced by heterogeneous growth. In order

to model the growth of biological tissues, we once again adopt the concept of fictitious configurations and decompose the total deformation multiplicatively into elastic, prestrain, and growth parts (Rodriguez, Hoger and McCulloch, 1994; Ambrosi, Ateshian, et al., 2011; Kuhl, 2014). This decomposition is a very general tool, and can also be applied for instance to model tissue activation (Rossi, Ruiz-Baier, et al., 2012; Pezzuto, Ambrosi and Quarteroni, 2014)

Figure 2 shows the different configurations and transformations. Similar to the mechanics of prestrained biological systems in Section 2.2, the elastic tensor $\underline{\underline{F}}^e$ can be multiplicatively decomposed into the prestrain $\underline{\underline{F}}^p$ and the mechanically-induced deformation $\underline{\underline{F}}'$. Yet, we can also decompose the elastic tensor $\underline{\underline{F}}^e$ into the inverse growth $\underline{\underline{F}}^g$ and the deformation gradient $\underline{\underline{F}}$:

$$\underline{\underline{F}}^e = \underline{\underline{F}}' \cdot \underline{\underline{F}}^p = \underline{\underline{F}} \cdot \underline{\underline{F}}^{g-1} \quad \text{with} \quad \underline{\underline{F}} = \underline{\underline{F}}' \cdot \underline{\underline{F}}^p \cdot \underline{\underline{F}}^g = \nabla \underline{\underline{\varphi}} \quad (9)$$

Note that in the special case where $\underline{\underline{F}} = \underline{\underline{F}}'$, we have $\underline{\underline{F}}^p = \underline{\underline{F}}^{g-1}$. However, in practice we only work with $\underline{\underline{F}}^g$, which obeys the evolution equation defined in the next paragraph, and $\underline{\underline{F}}^e$, which contains both the prestrain $\underline{\underline{F}}^p$ and load-induced deformation $\underline{\underline{F}}'$, and is used to compute stress through the elastic strain energy potential. Figure 2 also illustrates the heterogeneous growth-induced prestrain map $\underline{\underline{F}}^p$ between the incompatible grown configuration and the residually-stressed compatible configuration.

Here, in order to focus on growth-induced residual stress and not on the growth itself, we will use a simple isotropic and strain-driven growth law. Indeed, even though it is now established that maladaptive hypertrophic processes are largely anisotropic (Taber, 1995; Frey and Olson, 2003), physiological growth must involve both longitudinal and transverse growth, and isotropic growth is the simplest growth model that can generate residual stresses. Moreover, even though the hormonal mechanisms underlying tissue growth are complex and beyond the scope of this study, it is now accepted that hormonal signals are strongly regulated by mechanical signals, and that mechanotransduction, especially stretch-sensitivity, is one of the fundamental processes underlying tissue growth (Maillet, van Berlo and Molkenkin, 2013). Note that elastic strain and stress are coupled through the constitutive relationship and not mutually independent, so a stress-driven law could be used as well. To characterize the kinematics of growth we can thus parameterize the growth tensor $\underline{\underline{F}}^g$ in terms of a single scalar-valued growth multiplier ϑ :

$$\underline{\underline{F}}^g = \vartheta^g \underline{\underline{1}} \quad (10)$$

To characterize the kinetics of growth, we assume that growth evolves in time according to the following evolution law:

$$\dot{\vartheta}^g = \frac{1}{\tau} \text{tr}(\underline{\underline{E}}^e) \quad (11)$$

where τ is a time constant for growth. Since it is the only time constant of the problem, we simply set it to 1, and consider all other time-dependent quantities in terms of it. This implies that the driving force for growth is the trace of the elastic deformation $\underline{\underline{E}}^e$.

2.3.2. Computational model—Similar to the elasticity part in Section 2.2, we implemented the growth model as a user subroutine into the finite element package Abaqus/Standard. Since growth is driven by the elastic strains $\underline{\underline{E}}^e$, which are in turn affected by growth as $\underline{\underline{F}}^e = \underline{\underline{F}} \cdot \underline{\underline{F}}^{g-1}$, we iteratively determine the growth multiplier \mathcal{G} at the integration point level at every global iteration step. Rather than using a consistent Newton-Raphson procedure, which requires a consistent algorithmic linearization (Göktepe, Abilez and Kuhl, 2010; Göktepe, Abilez, et al., 2010), we adopt a local fixed-point iteration, supplemented by Aitken's relaxation (Genet, Marcin and Ladeveze, 2013). For the time integration, we use an implicit mid-point rule. This results in the following algorithm to determine the local growth multiplier \mathcal{G} :

```

given:  $\vartheta_t^g ; \underline{\underline{F}}_{t+1} ; \Delta t$ 
initialize growth rate and iteration counter:  $\dot{\vartheta}_{t+1/2}^g = 0 ; n_{iter} = 0$ 
loop
  update iteration counter:  $n_{iter} = n_{iter} + 1$ 
  update growth:  $\vartheta_{t+1}^g = \vartheta_t^g + \dot{\vartheta}_{t+1/2}^g \Delta t$ 
  calculate elastic deformation:  $\underline{\underline{E}}_{t+1}^e = \frac{1}{2} \left( \frac{1}{(\vartheta_{t+1}^g)^2} \underline{\underline{F}}_{t+1}^t \cdot \underline{\underline{F}}_{t+1} - \underline{\underline{1}} \right)$ 
  evaluate residual:  $\begin{cases} \text{if } (n_{iter} > 1) \text{ then } r^{old} = r \\ r = \dot{\vartheta}_{t+1/2}^g - \frac{1}{\tau} \text{tr} \left( \frac{\underline{\underline{E}}_{t+1}^e + \underline{\underline{E}}_{t+1}^e}{2} \right) \end{cases}$ 
  check convergence:  $\text{if } (r < \text{tolerance}) \text{ then exit loop}$ 
  calculate relaxation:  $\begin{cases} \text{if } (n_{iter} = 0) \text{ or } (r = r^{old}) \text{ then } \alpha = 1 \\ \text{else } \alpha = -\alpha r^{old} / (r - r^{old}) \end{cases}$ 
  update growth rate:  $\dot{\vartheta}_{t+1/2}^g = \dot{\vartheta}_{t+1/2}^g - \alpha r$ 
end loop

```

For a tolerance of $\text{tol} = 10^{-6}$, the algorithm converges rapidly in only a few iterations.

2.3.3. Example of growth-induced, prestrained left ventricle—Throughout this paper, we will illustrate our proposed method on a patient-specific left ventricular model of a healthy human (Genet, Lee, et al., 2014). This model characterizes healthy human ventricular mechanics. Briefly, the geometry was extracted from magnetic resonance images by manual segmentation, and triangulated using linear hexahedrons. A generic myofiber orientation field, with linearly varying helix angle from $+60^\circ$ at the endocardium to -60° at the epicardium (Streeter and Bassett, 1966; Streeter, Spotnitz, et al., 1969), was assigned to the mesh. We use a standard displacement finite element method, with fully implicit time stepping. In terms of boundary conditions, we constrain normal displacement of the basal nodes, and apply pressure loading at the endocardial faces.

To introduce residual stresses in a subject-specific geometry, we solve the following problem: We load the left ventricle to a given ventricular pressure P , then keep the pressure constant for a time t to allow the ventricle to grow, and then gradually remove the pressure. Growth duration t is usually expressed as a function of the growth time constant τ , which is the only time constant of the problem. As a result of heterogeneous growth, the final configuration is unloaded, but not stress free, *i.e.*, residually stressed. Our goal is now to iteratively determine the reference configuration for which the *in vivo* grown geometry for given a given P and time t matches our subject-specific imaging data. The following section describes how we identify the reference configuration using fixed-point iterations.

2.4. Computation of stress-free reference configuration of biological systems

2.4.1. Fixed-point iteration—As discussed in Section 2.3 and illustrated in Figure 2, the mechanically unloaded configuration from medical imaging is not a true reference configuration in the continuum mechanics sense since it is residually stressed. The stress free reference configuration is unknown and has to be determined iteratively. A common approach to this problem is to solve the so-called inverse elastic problem (Govindjee and Mihalic, 1996, 1998). Yet, this approach requires non-standard formulations and solvers on the continuum and discrete levels. Here we suggest an alternative fixed-point-based approach, which requires only modular modifications to the original forward problem. Fixed-point iterations have originally been proposed by (Sellier, 2011), and have recently been adopted for in the context of cardiac mechanics by (Wang, Gao, et al., 2013; Wang, Luo, et al., 2013; Krishnamurthy, Villongco, et al., 2013). However, previous approaches have only used the fixed-point iteration to identify the reference configuration associated with a loaded *in vivo* configuration. Here, we use it to find the reference configuration associated with an unloaded *in vivo* configuration. We assume that this unloaded structure contains heterogeneous growth-induced residual stresses.

Let $\underline{X}_{\text{ref}}$ denote the nodal positions of the residually stressed, mechanically unloaded configuration obtained, e.g., from *in vivo* imaging. The objective of the fixed-point iteration is to find the nodal positions \underline{X} of the stress-free, incompatible configuration that will deform into $\underline{X}_{\text{ref}}$ through the loading-growth-unloading scenario described in Section 2.3. Similar to the fixed-point iteration solver used for the growth law in Section 2.3.2, we adopt Aitken's relaxation (Aitken, 1950) to improve convergence. For time integration, we use here an implicit backward Euler scheme. The algorithm to iteratively determine the nodal positions \underline{X} in the stress-free configuration looks as follows:

```

given:  $\underline{X}_{ref}$  ;  $\underline{X}_{init}$ 
initialize configuration and iteration counter:  $\underline{X} = \underline{X}_{init}$  ;  $n_{iter} = 0$ 
loop
  update iteration counter:  $n_{iter} = n_{iter} + 1$ 
  for given  $\underline{X}$ ,  $\bar{P}$  and  $\bar{t}$ ,
  solve growth problem :  $\underline{X}_{def}$ 
  evaluate residual:  $\begin{cases} \text{if } (n_{iter} > 1) \text{ then } \underline{R}^{old} = \underline{R} \\ \underline{R} = \underline{X}_{def} - \underline{X}_{ref} \end{cases}$ 
  check convergence:  $\text{if } (\max(\underline{R}) < \text{tolerance}) \text{ then exit loop}$ 
  calculate relaxation:  $\begin{cases} \text{if } (n_{iter} = 0) \text{ or } (\underline{R} = \underline{R}^{old}) \text{ then } \alpha = 1 \\ \text{else } \alpha = -\alpha \frac{\underline{R}^{old} \cdot (\underline{R} - \underline{R}^{old})}{(\underline{R} - \underline{R}^{old}) \cdot (\underline{R} - \underline{R}^{old})} \end{cases}$ 
  update configuration:  $\underline{X} = \underline{X} - \alpha \underline{R}$ 
end loop

```

Here \underline{X}_{init} are the initial nodal position of the stress-free reference configuration, and \underline{X}_{def} are the deformed nodal position after the loading-growth-unloading procedure. This implies that \underline{X}_{init} could, but does not have to, be equal to \underline{X}_{ref} . Similar to the algorithm in Section 2.3.2, the fixed-point iteration converges rapidly in a few iterations for a tolerance of 10^{-1} mm.

2.4.2. Computational model—We implemented the fixed-point iteration as a python script, which solves the growth problem inside the inner loop using Abaqus/Standard and then post-processes the nodal positions.

3. Results and Discussion

3.1. Opening angle experiment

Figure 3 illustrates our opening angle experiment. It shows an isolated heart slice before (left) and after (right) cutting the slice radially in the middle of the left ventricular free wall.

Table 2 summarizes the opening angles of the basal, equatorial, and apical slices of all four hearts. Average opening angles increase from base to apex (from 8.7° to 16.6°) with an increasing standard deviation (from $\pm 1.8^\circ$ to $\pm 13.7^\circ$). The variability in opening angle may be attributed to a combination of geometric and structural differences between the two control slices, and the location of the radial cut. Nevertheless, opening angles are significantly lower than values found for rat hearts (45°) (Omens and Fung, 1990), hence potentially revealing a size-effect in residual stress development.

3.2. Generating growth-induced prestrain

Figure 4 illustrates our protocol to generate prestrain through heterogeneous growth in a patient specific left ventricular geometry (Genet, Lee, et al., 2014). For the sake of illustration, we used a nominal ventricular pressure P of 1 mmHg, and a nominal growth duration t of one growth time constant τ . During the strain-driven growth step, the myofiber stress field becomes more and more homogeneous. While homogeneous isotropic growth does not generate residual stress, homogeneous anisotropic growth (Rausch, Dam, et al., 2011; Kerckhoffs, 2012) and heterogeneous isotropic growth (Göktepe, Abilez and Kuhl,

2010; Göktepe, Abilez, et al., 2010) do. Thus, the final configuration is residually stressed: the sub-endocardial region is in compression, the sub-epicardial region in in tension.

This is in contrast to some models in the literature (Kroon, Delhaas, et al., 2007, 2009), which hypothesize that the growth-induced residual stresses are relieved over time, such that the grown configuration is always assumed to be stress free. Our opening angle experiment indicates though that the inner layer of the ventricle, similar to arteries, is in compression and the outer layer is in tension. This result is compatible with strain-driven growth-induced prestrain, the underlying hypothesis in this paper.

We have used the trace of the elastic Green-Lagrange strain tensor as the thermodynamical force to drive growth. Other authors have used the fiber stretch (Lee, Genet, et al., 2014a); however, the fiber stretch is largest in the mid-wall and not in the sub-endocardial region (Choi, D'hooge, et al., 2010; Lee, Genet, et al., 2014a), and the associated residual stress fields are not physiological. Others have use the Kirchhoff (Klepach, Lee, et al., 2012) or Mandel (Göktepe, Abilez and Kuhl, 2010; Göktepe, Abilez, et al., 2010) stresses, which are more “isotropic” quantities. Actually, in the case of isotropic growth and compressible elasticity, the Mandel stress is the energy conjugate of the growth state variable; however, in the case of quasi-incompressible elasticity considered here, the energy conjugate of the isotropic growth state variable is basically the isostatic pressure, which does not seem to be a reasonable driving force. In other words, the distribution of residual stresses constrains the functional form of the growth law. This is why, after systematically testing a collection of other driving forces for growth, we decided to select the trace of the elastic deformation.

As for growth kinetic equation itself, for the lack of appropriate experiments, we use a simple linear evolution law. More complex evolution laws have been proposed, including thresholds and limiting values (Kuhl, 2014), and even multiple domains (Lee, Genet, et al., 2014a) to predict both growth and reverse growth. The growth law we have selected here is the simplest law to predict heterogeneous growth.

3.3. Growth-induced prestrain in patient-specific left ventricle

Figure 5 illustrates the fixed-point iteration method to solve for the reference configuration associated with a given amount of prestrain, once again generated with a nominal ventricular pressure of 1 mmHg and a nominal growth duration of one growth time constant. For each iteration (*i.e.*, each column), we show the reference stress-free configuration (top row) and the final (after loading, growth, and unloading) configuration (bottom row). During the fixed-point iterations, we compute the error between the reference configuration (obtained from medical imaging) and the deformed configuration (after loading-growth-unloading). The very last geometry (last column, bottom row) is identical to the very first (first column, top row); this geometry is the *in vivo* unloaded geometry extracted from medical imaging), but it contains heterogeneous growth-induced residual stress.

Figure 6 shows, for multiple growth durations ranging from 1 to 6 growth time constants, the reference configuration computed using the fixed-point iteration method (top row) as well as the prestrained configuration. The reference configuration is always stress-free, but the geometry varies. On the contrary, the prestrained configuration has always the same

geometry (the one extracted from medical imaging data), but different levels of residual stress.

By assuming that the early-diastolic ventricular geometry extracted from medical imaging is unloaded, we implicitly neglected the ventricular pressure at the beginning of diastole.

However, this hypothesis is not necessary for using the fixed-point iterations. In principle, it is possible to prescribe any loading at the end of the forward step of the inverse problem, thus taking into account the mechanical load applied to the tissues.

3.4. Virtual opening angle experiment

After generating residual stresses in the subject-specific left ventricular geometry, we slice the virtual heart longitudinally and then cut it open. The result is illustrated in Figure 7, for a nominal ventricular pressure of 1 mmHg, and a nominal growth duration of six growth time constants. To reproduce the experimental conditions, the nodes belonging to the upper and lower faces can only move in the sliding plane. To remove rigid body motions, the nodes of one face of the cut can only move in the cutting plane.

Figure 8 shows the relationship between the maximum principal residual strain in the slice and the opening angle. Different levels of residual strain are generated using different growth durations, ranging from 1 to 6 growth time constants, while a 1 mmHg ventricular pressure is always used. The black line and the gray box mark the average and standard deviation from the physical opening angle experiment on the mid-level slice in Section 3.1.

It is not necessary to shrink the ventricle to an infinitesimally small size to then induce enough residual stress through growth to generate a realistic opening angle. Our proposed method thus allows us to distinguish between the residual stress that is released, and the residual stress that remains within the tissues. Moreover, the fact that some of the residual stress is released justifies the original assumption that there exists a stress-free compatible configuration, which is described in Figure 2, and computed using the fixed-point algorithm described Section 2.4.

3.5. Influence of the prestrain on the ventricular mechanical response

An important consequence of prestrain is the influence of the residual stress on the mechanical response of the heart. Figure 9 shows the passive pressure-volume relationship for different levels of prestrain associated with different opening angles. Once again, the different levels of prestrain are generated using different growth durations, here ranging from 1 to 6 growth time constants, and the ventricular pressure during growth is kept constant a 1 mmHg. Because of the nature of residual stresses in growing circular cross sections, with compressive stresses at the sub-endocardial region and tensile stresses at the sub-epicardial region, prestrain tends to make the ventricle more compliant. For instance, for a maximal principal prestrain of 15.34%, a fiber residual stress varying from -0.12kPa to $+0.055\text{kPa}$, and an opening angle of 17.1° , *i.e.*, close to the average experimental value, the ventricular pressure after a passive filling corresponding to an unloaded ventricular volume (*i.e.*, a 50% ejection fraction) is reduced by 25.8% compared to the prestrain-free ventricle. As a consequence of this increased compliance, the ventricle requires less energy to fill. In

other words, besides the well-known effect of homogenizing the ventricular wall stress (Omens and Fung, 1990), residual stresses could be a mechanism to directly improve ventricular diastolic function.

4. Summary and Perspectives

In this paper, we investigated the hypothesis that residual stress in the beating heart is induced by growth of the organ. We used the finite growth framework, together with a fixed-point iteration on the geometry itself, to introduce residual stress in a subject-specific left ventricular geometry. We showed that the residual stress fields induced by heterogeneous growth are compatible with experimental results, both qualitatively and quantitatively. It can successfully predict the physiological range of opening angles. We showed that the prestrain can have a significant impact on the passive mechanical response of the ventricle, making it much more compliant.

Some more general perspectives follow from this work. First, in addition to studying the influence of the prestrain on the passive ventricular response alone, we will characterize the sensitivity of the active response, and, eventually, of the entire pressure-volume relationship, for different prestrain levels. Another important follow up study will be to incorporate the algorithm introduced here within a personalized left ventricular model (Genet, Lee, et al., 2014), and compute personalized passive and active cardiac mechanical properties by taking into account physiological prestrain levels. Finally, we will study the interactions between physiological residual stress and residual stress induced by the injection of biomaterials in diseased myocardium as a potential treatments (Lee, Wall, et al., 2013, 2014), and their respective influence on ventricular mechanics. We believe that prestrain could be the key to explain various discrepancies in the literature between *in vivo* and *ex vivo* measurements and computational simulations. Including the effects of prestrain have the potential to make subject-specific modeling more reliable and more applicable to today's cardiac health problems.

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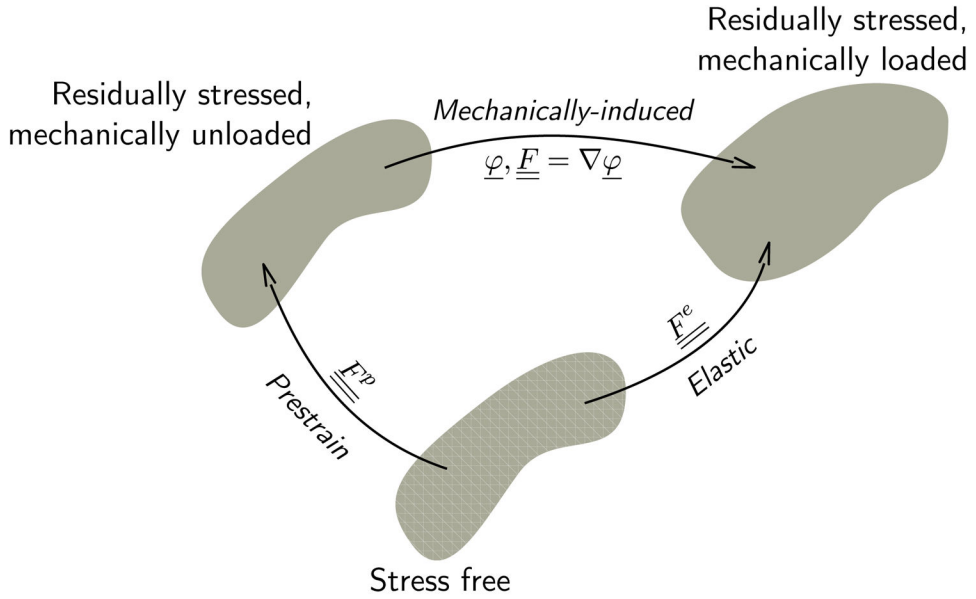


Figure 1.

Kinematics of prestrained biological systems. Prestrain maps the stress-free reference configuration onto the residually stressed but mechanically unloaded configuration through the prestrain tensor F^p . Mechanical loading maps the residually stressed but mechanically unloaded configuration into the mechanically loaded configuration through the deformation gradient F . The total elastic deformation F^e is a result of prestrain F^p and mechanical deformation F . Only the mechanical deformation $F = \nabla \varphi$ is the gradient of a vector field, while the prestrain tensor F^p and the elastic tensor F^e are incompatible. Only the elastic tensor F^e generates stress.

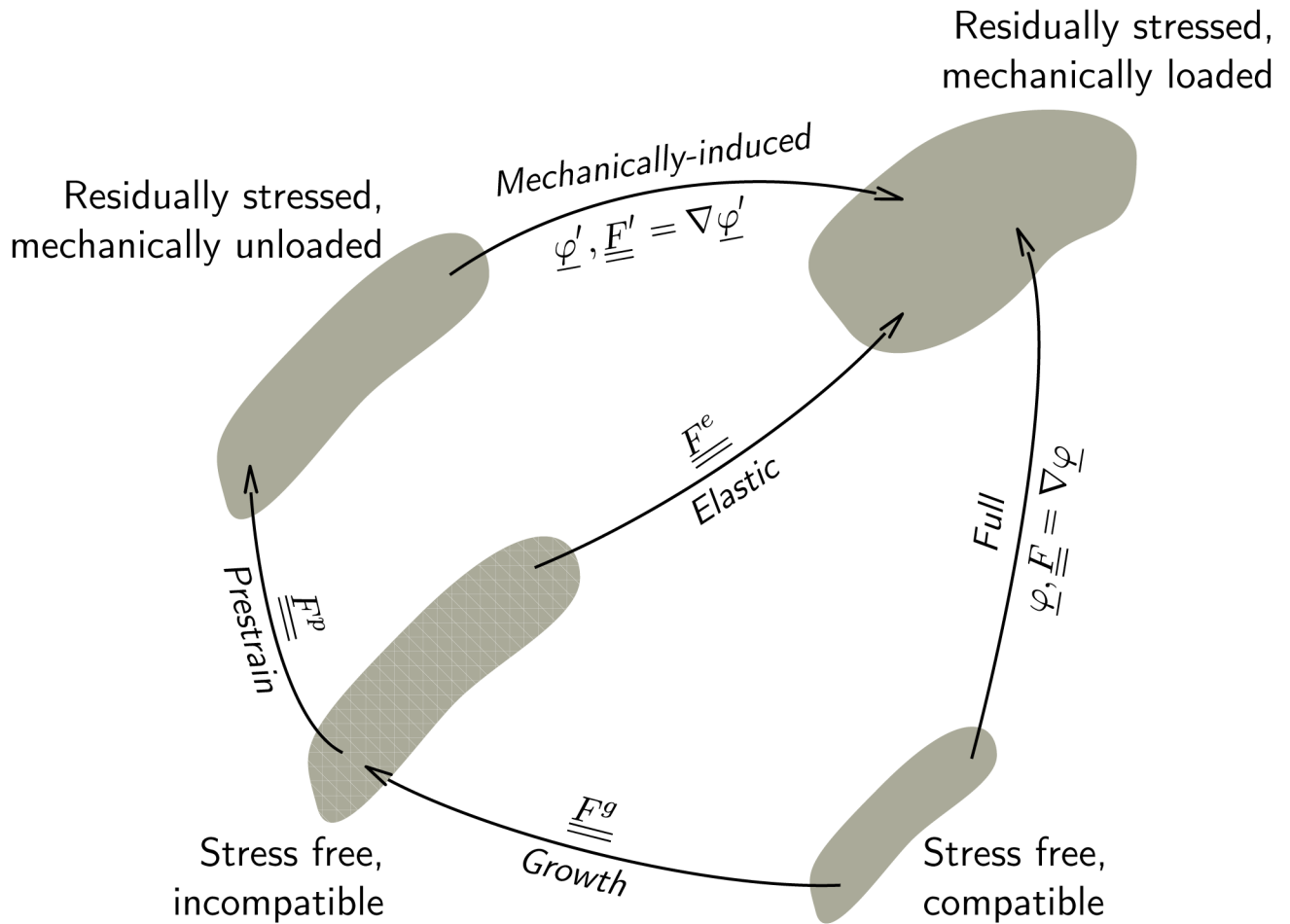


Figure 2.

Kinematics of growth-induced, prestrained biological systems. Growth turns the initial, stress-free and compatible configuration into the grown, stress-free but incompatible configuration. Prestrain maps the grown, stress-free configuration onto the residually stressed but mechanically unloaded configuration through the prestrain tensor \underline{F}^p . Mechanical loading maps the residually stressed but mechanically unloaded configuration into the mechanically loaded configuration through the deformation gradient \underline{F}' . The total elastic deformation \underline{F}^e is a result of prestrain \underline{F}^p and mechanical deformation \underline{F}' . The full deformation \underline{F} is a result of growth \underline{F}^g , prestrain \underline{F}^p , and mechanical deformation \underline{F}' . Only the full deformation $\underline{F} = \nabla \underline{\varphi}$ and mechanically-induced deformation $\underline{F}' = \nabla \underline{\varphi}'$ are gradients of a vector field, while the growth tensor \underline{F}^g , the prestrain tensor \underline{F}^p , and the elastic tensor \underline{F}^e are incompatible. Note that Figure 2 becomes identical to Figure 1 if $\underline{F}^g = \underline{1}$.

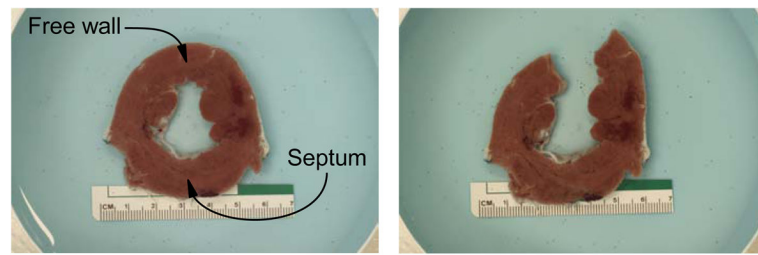


Figure 3. Opening angle experiment. Isolated heart slice before (left) and after (right) radial cutting in the middle of the left ventricular free wall.

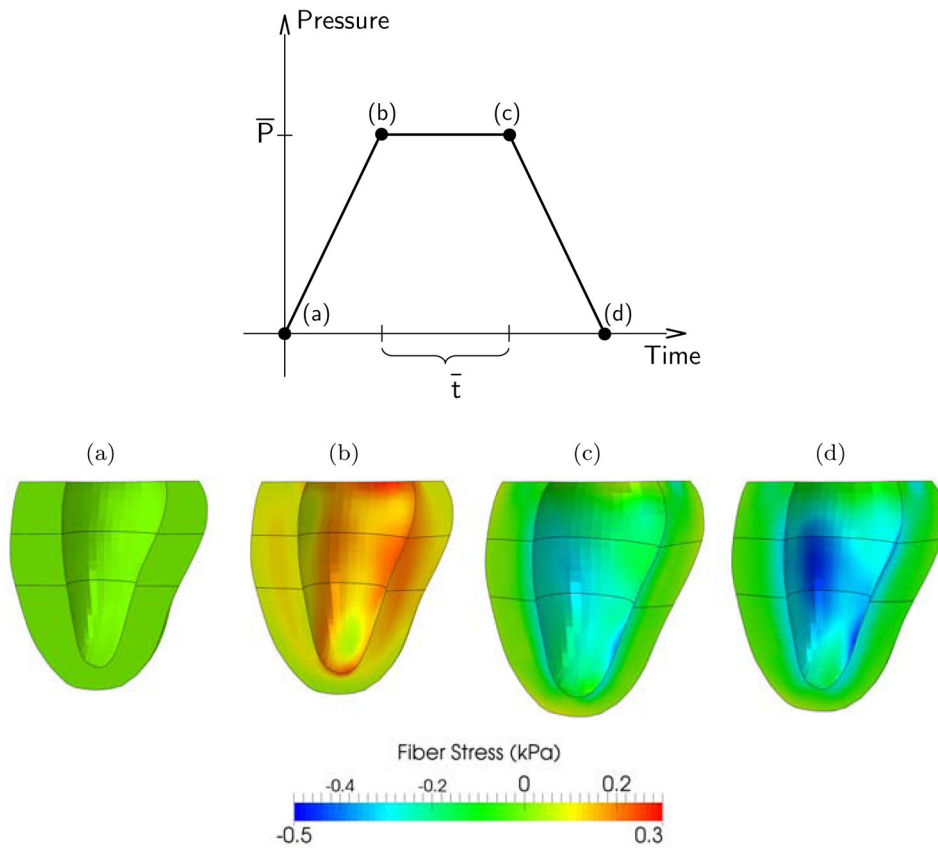


Figure 4.

Protocol to generate heterogeneous growth-induced residual stress in a patient specific left ventricular geometry. The ventricle is loaded to a ventricular pressure of \bar{P} , (a)–(b), then allowed to grow for a duration \bar{t} , (b)–(c), and then unloaded, (c)–(d). The strain-driven growth drives the growing configuration into a state with a more homogeneous myofiber stress and reduces regional stress variations (c). The final configuration is unloaded, but not stress free, with the sub-endocardial region in compression, and the sub-epicardial region in tension (d).

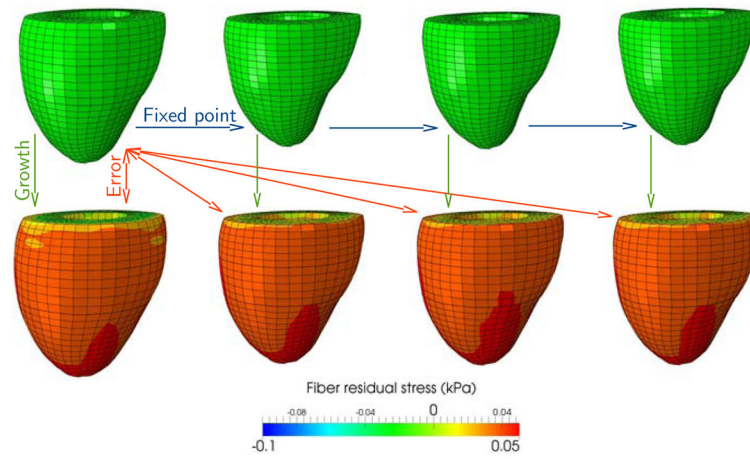


Figure 5. Fixed-point iteration method to compute the prestrained, mechanically unloaded reference configuration (last column, top row) so that the prestrained, mechanically loaded configuration (last column, bottom row) matches the *in vivo* geometry extracted from magnetic resonance images (first column, top row) but contains auto-balanced residual stresses.

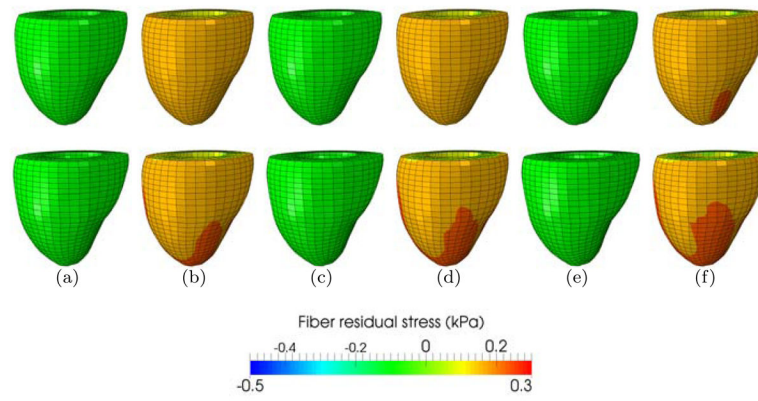


Figure 6. Prestrained configurations for varying prestrain levels. Prestrain is generated by heterogeneous growth. The reference configurations are computed using a fixed-point iteration, so that the prestrained configurations (bottom row) match the *in vivo* geometry extracted from magnetic resonance images (top row).

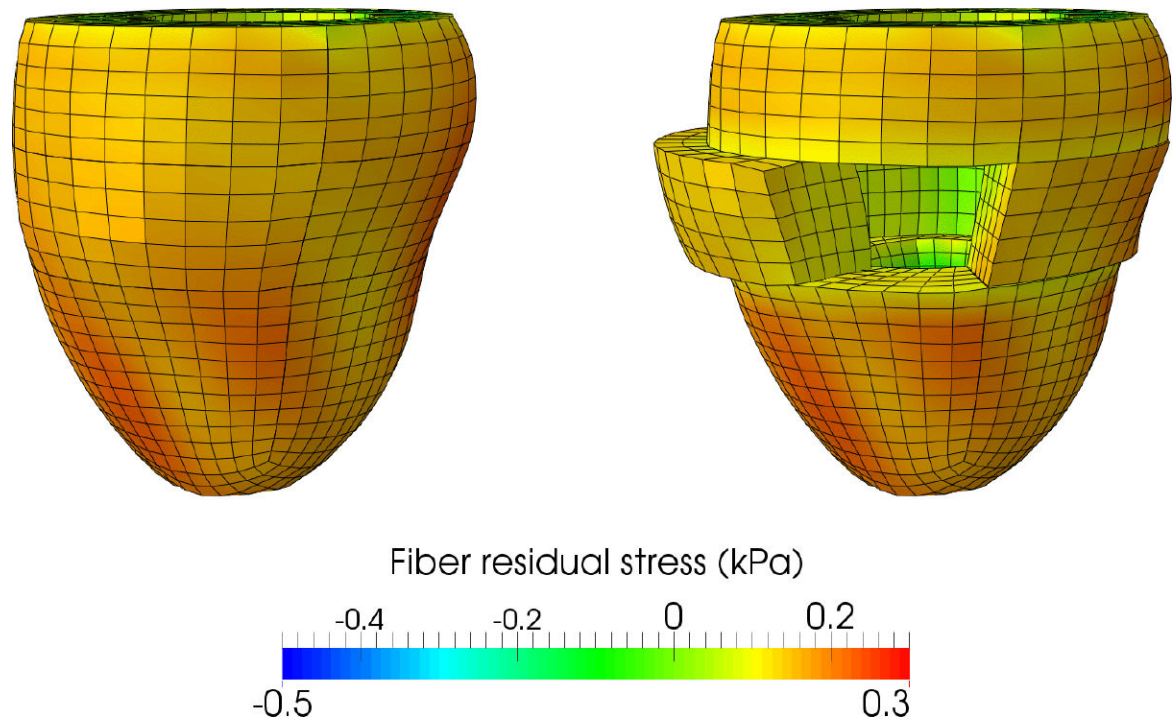


Figure 7. Virtual opening angle experiment. The prestrained configuration is sliced and then cut. The slice naturally springs open, similar to the physical experiment illustrated in Figure 3. Cutting the slice relieves some, but not all residual stress. Colors represent residual fiber stress, in kPa.

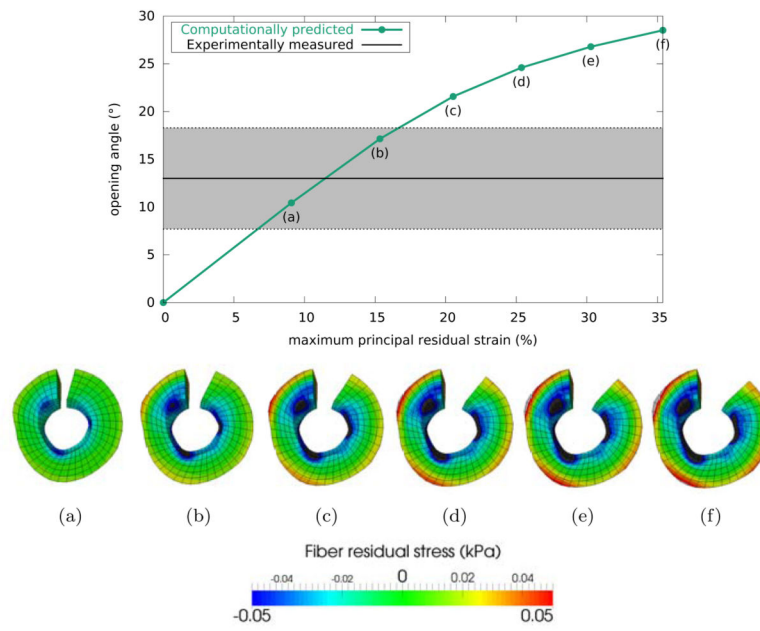


Figure 8.

Opening angles for varying prestrain levels. The green curve summarizes the computationally simulated opening angles for a pressure of $P = 1$ mmHg. The black line and gray box mark the experimentally measured opening angle of $13 \pm 5.3^\circ$ for the equatorial slice. Computationally simulated opening angles lie within the range of experimentally measured opening angles.

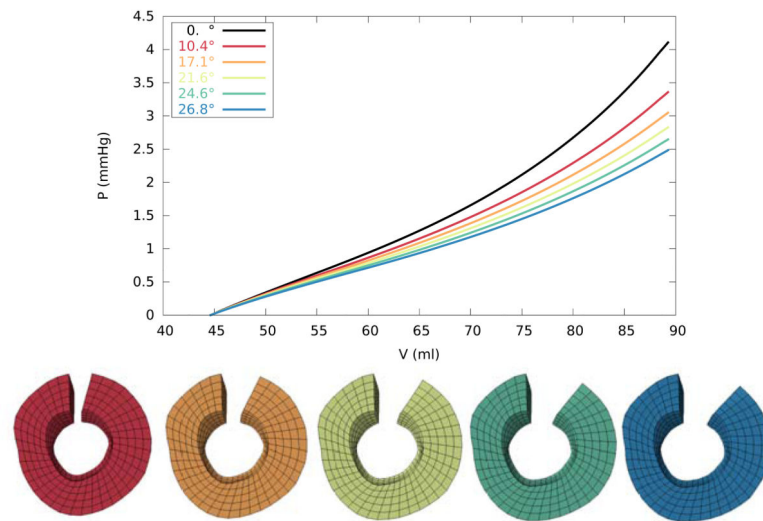


Figure 9. Passive pressure-volume response of the left ventricle for varying prestrain levels. Increasing the prestrain level increases the ventricular compliance: for an opening angle of 17.1° , *i.e.*, close to the average experimental value, the ventricular pressure after passive filling corresponding to a 50% ejection fraction is reduced by 25.8% compared to the prestrain-free case.

Normal human myocardial properties (Genet, Lee, et al., 2014) used in this paper for the ventricular simulations.

Table 1

D_0 (kPa ⁻¹)	C_0 (kPa)
10^{-3}	0.115

B_{ff} ()	B_{ss} ()	B_{mt} ()	B_{fs} ()	B_{jr} ()	B_{sn} ()
14.4	5.76	5.76	5.04	5.04	2.88

Table 2

Opening angle experimental data. All values are reported in degrees.

	Basal slice	Equatorial slice	Apical slice
Heart 1	9.3	8.7	9.8
Heart 2	9.8	19.9	16.5
Heart 3	9.7	9.0	4.2
Heart 4	6.1	14.5	35.7
Mean±SD	8.7±1.8	13.0±5.3	16.6±13.7

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