

Biventricular Cardiac Model

Introduction

This example demonstrates how to simulate a cardiac contraction on a simplified heart geometry, where only the ventricles are considered and the stimulus starts from the atrioventricular node.

During the cardiac cycle (heartbeat) an electrical stimulus generated from the sino-atrial node propagates throughout the entire heart. When excited, cardiac cells are subjected to an electric potential jump across their membranes. The transport and accumulation of ions and additional chemical reactions cause cells to contract. At a macro-scale these chemical and electrical processes trigger the contraction of the muscle tissue, the myocardium, allowing blood to be pumped to the arteries.

The muscle tissue (myocardium) is mainly composed of myocardial cells grouped in layered sheets that are aligned in a preferential direction (fibers). The fiber orientation strongly affects the mechanical and electrical tissue properties. This arrangement suggests that the cardiac tissue can be modeled as an anisotropic hyperelastic material.

The time and spatial evolution of the electric potential is described by the *Aliev–Panfilov* equations with a nonlinear current-voltage relation. This electrochemical model describes the quick rise of the cell's transmembrane electric potential (depolarization) and the return to its resting value (repolarization).

The tissue contraction in the hyperelastic model is obtained using an additive decomposition of the stress tensor. The actual contraction at cellular level is taken into account by introducing an additional stress (active stress) in the Solid Mechanics interface. This additional stress is voltage-dependent, and its evolution is described by additional partial differential equations.

Model Definition

The simplified heart geometry shown in Figure 1 includes the two elliptical ventricular chambers. The atria are not considered.



Figure 1: Geometry of the two ventricular chambers of the heart.

The base is located at Z = 0, and the left ventricle apex is located at Z = -70 mm. The ventricle dimensions are taken from Ref. 2 and shown in Table 1.

TABLE I: LEFT AND RIGHT VENTRICLE DIMENSIONS.

| Geometry | Left ventricle | Right ventricle |
|-------------------------|----------------|-----------------|
| x-semiaxis (epicardium) | 30[mm] | 51[mm] |
| y-semiaxis (epicardium) | 30[mm] | 30[mm] |
| z-semiaxis (epicardium) | 70[mm] | 60[mm] |
| Thickness | 12[mm] | 6[mm] |

The ventricular wall (myocardium) is surrounded by the epicardium on the outside (Figure 2) and by the endocardium on the inside (Figure 3). These are not included in the model, but these terms will refer to the walls boundaries that are in contact with the myocardium.



Figure 2: Boundaries in contact with the epicardium.



Figure 3: Boundaries in contact with the endocardium.

FIBER DIRECTION

The fibers have an inclination of 60° at the endocardium (θ_{end}^{max}) and -60° at the epicardium (θ_{epi}^{max}) with respect to the horizontal basal plane (Z = 0). It is also assumed that the fibers' inclination goes to zero at the left ventricle apex (Z_{LV}^{apex}), so the expressions for the fiber angle on the myocardium boundaries read

$$\theta_{epi} = \theta_{epi}^{max} \left(1 - \frac{Z}{Z_{LV}^{apex}} \right)$$
$$\theta_{end} = \theta_{end}^{max} \left(1 - \frac{Z}{Z_{LV}^{apex}} \right)$$

Across the cardiac walls the fiber angle varies linearly as

$$\theta = \beta \theta_{end} + (1 - \beta) \theta_{epi}$$

where β is a dimensionless parameter representing the fiber distance to the epicardium boundary. It takes values between 0 (epicardium) and 1 (endocardium), and it is an expression based on the fiber distance to the epicardium, $D_{\rm epi}$, and to the endocardium, $D_{\rm end}$,

$$\beta = \frac{D_{\rm epi}}{D_{\rm epi} + D_{\rm end}} \tag{1}$$

The myocardium is modeled as an anisotropic hyperelastic material with three preferential directions: fiber direction (\mathbf{a}), sheet direction (\mathbf{s}) and normal to sheet direction (\mathbf{n}). This reference system can be approximated through a composition of coordinate systems.

The coordinate system used to define the material and electrical tissue properties is created in two steps. First, the transmural direction is obtained through the use of the **Curvilinear Coordinate** interface. It is assumed that the transmural direction coincides with the sheet direction (**s**). The resulting curvilinear system has the first basis vector oriented in the sheet direction, and the second basis vector oriented toward the apico-basal direction (Figure 4). Coordinate system volume: Base vector system



Figure 4: Intermediate coordinate system obtained with the Curvilinear Coordinate interface. The first basis vector (red) corresponds to the transmural direction. Note that the transmural direction coincides with the sheet direction in this example.

This intermediate coordinate system is rotated with an angle θ around the first basis vector (**s**) to align the third axis along the fiber direction **a**. The resulting fiber orientation is shown in Figure 5.

y ^z _ x



Figure 5: Fiber layout in the undeformed configuration. Note that the fiber angle changes along the transmural direction.

ELECTROPHYSIOLOGY MODEL

The electric potential Φ in the myocardium is described by the following equation (Ref. 1)

$$\chi_{\rm m} C_{\rm m} \frac{\partial}{\partial t} \Phi + \nabla \cdot (-D \nabla \Phi) + \chi_{\rm m} [I_{\rm ion}(\Phi, F, r_i)] = 0$$
⁽²⁾

where χ_m is the membrane surface to volume ratio, C_m is the membrane capacitance, D is the conductivity tensor, I_{ion} is the ionic current per unit area, and F the deformation gradient. The ionic current depends on the potential, the deformation gradient, and other internal variables, r_i .

In this example, the ionic current is split into an excitation-induced, purely electrical part $I_{\rm e}$, and a stretch-induced part $I_{\rm m}$, as described in Ref. 2.

$$I_{\text{ion}}(\Phi, F, r_i) = I_{\text{e}}(\Phi, r_i) + I_{\text{m}}(F)$$
(3)

The *Aliev–Panfilov* equations (Ref. 3) are used to represent the purely electrical part of the ionic current, I_e , as a function of the electric potential. These equations use one additional internal variable, and they are usually expressed in dimensionless form:

$$\tilde{I}_{\rm e}(v,r) = c\phi(\phi-\alpha)(\phi-1) + r\phi$$
$$\frac{\partial r}{\partial \tau} = \left(\gamma + \frac{\mu_1}{\mu_2 + \phi}r\right)[-r - c\phi(\phi-b-1)]$$

Here, ϕ is the dimensionless electric potential, τ is the dimensionless time, and *r* is an internal variable called *recovery variable*. Furthermore, *c*, α , γ , μ_1 , μ_2 , and *b* are material parameters whose values are taken from Ref. 2 and can be found in Table 2.

| PARAMETER | VALUE |
|-----------|-------|
| с | 8 |
| α | 0.01 |
| γ | 0.002 |
| μ_1 | 0.2 |
| μ_2 | 0.3 |
| b | 0.15 |

TABLE 2: MATERIAL PARAMETERS FOR THE ALIEV-PANFILOV EQUATION.

The a stretch-induced electric current $I_{\rm m}$ is defined as

$$\tilde{I}_{\rm m}(F) = \Theta G_s(\lambda(F) - 1)(\phi - \phi_s)$$

where λ is the stretch in the fiber direction, G_s is the maximum conductance, ϕ_s is the resting electric potential for the ion channels, and θ is an activation parameter. This expression assumes the value of 1 when the fibers are stretched, and 0 when they are compressed. The values of the additional parameters are taken from Ref. 2 and shown in Table 3.

TABLE 3: STRETCH-INDUCED CURRENT PARAMETERS.

| PARAMETER | VALUE |
|------------|-------|
| $G_{ m s}$ | 10 |
| ϕ_s | 0.6 |

A dimensional mapping of the ionic current equation (Equation 3) is used to match the experimental values of electric potential and activation time in the myocardium:

$$\Phi = \phi \beta_{\phi} + \delta_{\phi}$$
$$t = \beta_{t} \tau$$

Here, β_{ϕ} and δ_{ϕ} are selected to match experimental values: the resting potential of the heart is -80 mV and the maximum potential value is 20 mV.

The time scaling parameter β_t is considered to be dependent on the activation time t_a , which shows better agreement with experimental observations (see Ref. 2 for details). During the cardiac cycle, the activation lapse (time between depolarization and repolarization) is not constant throughout the myocardium. Regions that are depolarized last are the first to repolarize. Therefore, the parameter β_t assumes the following expression

$$\beta_{\rm t} = (12.9 \text{ ms}) \left(1 - \tau_0 \frac{t_{\rm a} - t_0}{t_1 - t_0} \right) \tag{4}$$

where τ_0 , t_0 , and t_1 are tuning parameters (Table 4).

TABLE 4: TIME MAPPING PARAMETERS

| PARAMETER | VALUE |
|-----------|--------|
| τ_0 | 0.55 |
| t_0 | I2[ms] |
| t_1 | 75[ms] |

In this example, the activation time t_a depends on the Z coordinate and the vertical distance to the apex

$$t_{\rm a}(X, Y, Z) = (50 \,\mathrm{ms}) \left(1 - \frac{Z}{Z_{\rm LV}^{\rm apex}} \right)$$

The dimensional ionic current is then obtained from dimensionless purely electrical current $I_{e}(v, r)$, and stretch-induced current $I_{m}(F)$

$$I_{\text{ion}} = C_{\text{m}} \frac{\beta_{\phi}}{\beta_{\text{t}}} (\tilde{I}_{\text{e}} + \tilde{I}_{\text{m}})$$

The conductivity tensor in Equation 2 is decomposed into an isotropic part and an anisotropic part, which depends on the fiber direction (Ref. 2)

$$D = (d_{iso}C_m\chi_m)I + (d_{ani}C_m\chi_m)\mathbf{a}_0 \otimes \mathbf{a}_0$$

where $d_{iso} = 1 \text{ mm}^2/\text{ms}$ and $d_{ani} = 0.1 \text{ mm}^2/\text{ms}$.

ACTIVE STRESS

The active stress component, s_a, is calculated by solving the following differential equation:

$$\frac{\partial s_{a}}{\partial t} = \varepsilon(V)[k(\Phi - \Phi_{\rm r}) - s_{\rm a}]$$
(5)

where ε is a delay function introduced in Ref. 2:

$$\varepsilon(V) = \varepsilon_0 + (\varepsilon_0 - \varepsilon_1) \exp(-\exp(-\zeta(\Phi - \Phi_t)))$$

Here, k, Φ_r , ε_0 , ε_1 , ζ , and Φ_t are taken from Ref. 2 and shown in Table 5.

TABLE 5: ACTIVE STRESS PARAMETERS.

| PARAMETER | VALUE |
|----------------|---------------|
| k | 0.005[MPa/mV] |
| ε ₀ | 0.1[1/ms] |
| ε ₁ | I[I/ms] |
| ζ | I[I/mV] |
| $\Phi_{\rm r}$ | -80[mV] |
| $\Phi_{ m t}$ | 0[mV] |

The active stress obtained from Equation 5 is added to the second Piola–Kirchhoff stress in different percentages along the fiber, sheet and normal directions as described in Ref. 1:

$$S = S + s_{a}(\mathbf{a}_{0} \otimes \mathbf{a}_{0}) + 0.4s_{a}(\mathbf{s}_{0} \otimes \mathbf{s}_{0}) + 0.4s_{a}(\mathbf{n}_{0} \otimes \mathbf{n}_{0})$$
(6)

MATERIAL MODEL

The cardiac tissue is considered to be hyperelastic, and the strain energy density is split into isotropic and anisotropic contributions

$$W = W_{iso} + W_{ani}$$

A compressible Neo-Hookean strain energy density is used for the isotropic part,

$$W_{\rm iso} = \frac{1}{2}\mu(I_1 - 3) + \mu \log J_{\rm el} + \frac{1}{2}\lambda(\log J_{\rm el})^2$$

whereas the Holzapfel–Gasser–Ogden (HGO) model is used for the anisotropic contribution.

$$W_{\text{ani}} = \frac{k_1}{2k_2} (e^{k_2 (\bar{I}_a - 1)^2} - 1)$$
(7)

Here, k_1 and k_2 are material properties, and I_a is the invariant of the isochoric Green– Cauchy strain tensor along the fiber direction:

$$\overline{I}_a = \overline{I}_a(\overline{C}_{el}, \mathbf{a}_0) = \mathbf{a}_0 \cdot \overline{C}_{el} \cdot \mathbf{a}_0$$
(8)

The anisotropic hyperelastic parameters are given in Table 6.

| TABLE 6: | MATERIAL | PROPERTIES. |
|-----------|----------|-------------|
| IT OLE OF | , | |

| Material properties | Value |
|-------------------------|------------|
| λ (Ref. 2) | 0.5[MPa] |
| μ (Ref. 2) | 0.2[MPa] |
| k ₁ (Ref. 1) | 1.685[kPa] |
| k ₂ (Ref. 1) | 15.779 |

BOUNDARY AND INITIAL CONDITION

The following conditions are applied to simulate the ventricle contractions:

- The displacement of the basal surface is constrained.
- Zero flux boundary conditions are used in the additional equations for Φ , r, and s_a .
- An initial pulse of electric potential (-10 mV) is applied on a rectangular area on the basal surface between the ventricles (Figure 6).



Figure 6: Initial impulse at the atrioventricular node.

Results and Discussion

The model computes the contraction of the myocardium due to an electric stimulus. Figure 5 displays the fibers layout in the undeformed configuration.

Figure 7 shows the time variation of the electric potential distribution and the deformation of the cardiac walls. The depolarization of the cells produces an active stress that causes the myocardium to contract. There is an upward motion of the apex and a small torsion of the ventricles. The heart returns to its original state after 300 ms.



Figure 7: Depolarization and repolarization of the cardiac walls at different times during the cardiac cycle.

Figure 8 shows the activation potential at three different points. Note that the last cells to depolarize are the first one to repolarize. This effect is achieved by manipulating the dimensionless time constant as shown in Equation 4.

Figure 9 shows the internal volume variation of the left and right ventricles. Both chambers contract during the depolarization of the excited myocardial cells, thus reducing the internal volume. Then, the ventricles gradually return to their initial configuration to be again excited during the next heartbeat.



Figure 8: Plot of the activation duration at different locations.



Figure 9: Volume variation of the left and right ventricular chambers.

References

1. A.A. Bakir, A.A. Abed, M.C. Stevens, N.H. Lovell, and S. Dokos, "A Multiphysics Biventricular Cardiac Model: Simulations With a Left-Ventricular Assist Device," *Front. Physiol.*, vol. 9, p. 1259, 2018.

2. S. Göktepe and E. Kuhl, "Electromechanics of the heart: a unified approach to the strongly coupled excitation-contraction problem," *Comput. Mech.*, vol. 45, pp. 227–243, 2010.

3. M.P. Nash and A.V. Panfilov, "Electromechanical model of excitable tissue to study reentrant cardiac arrhythmias," *Prog. Biophys. Mol. Biol.*, vol. 85, pp. 501–522, 2004.

The analysis is performed in two separated studies. In the first study the fiber orientation is computed. In a second study, the cardiac contraction following an electrical excitation is analyzed. In particular, the following steps are taken:

- The fibers paths are computed using the **Wall Distance** and **Curvilinear Coordinate** interfaces.
- The contribution of the electrophysiology model (PDE interfaces) is included in the Solid Mechanics interface through the **External Stress** feature.
- The Fiber feature is set up using the curvilinear fiber orientation.
- The fiber pattern is displayed for postprocessing.

The distances D_{epi} and D_{endo} used in Equation 1 to compute the dimensionless parameter β are determined by two **Wall Distance** interfaces. In one case the wall represents the endocardium, and in the other it represents the epicardium.

Inertial terms are neglected as reported in Ref. 2. The **Structural Transient Behavior** is set to **Quasistatic** in the **Solid Mechanics** interface setting.

The active stress (Equation 6) is added to the total stress with the **External Stress** feature. The fiber coordinate system is selected in the **Coordinate System Selection** section.

In the settings for the **Fiber** feature, select the appropriate **Fiber Reference System** as a reference coordinate system. Then, in the **Orientation** section, select the orientation of the fiber (\mathbf{a}_0) to be aligned with the **third axis**.

The **Fiber** feature automatically computes the corresponding invariants, for instance I_a as used in Equation 8, in order to compute the strain energy function W_{ani} as in Equation 7.

The option **Stiffness in tension only** is activated by default for the HGO model, and it evaluates the fibers' strain energy to zero if the fiber stretch is in compression. This means that the fibers contribute to tensile stresses only.

The model parameters taken from literature references already include the effect of the membrane surface to volume ratio, χ_m , and the membrane capacitance, C_m . For this reason, these two parameters are set equal to 1 in this example.

The internal volume is computed through the use of Gauss's theorem to convert area integrals to volume integrals.

Application Library path: Nonlinear_Structural_Materials_Module/ Hyperelasticity/biventricular_cardiac_model

Modeling Instructions

From the File menu, choose New.

NEW

In the New window, click 🔗 Model Wizard.

MODEL WIZARD

- I In the Model Wizard window, click 间 3D.
- 2 In the Select Physics tree, select Mathematics>Wall Distance (wd).
- 3 Click Add.
- 4 Click Add.
- 5 In the Select Physics tree, select Mathematics>Curvilinear Coordinates (cc).
- 6 Click Add.
- 7 Click 🔿 Study.
- 8 In the Select Study tree, select General Studies>Stationary.
- 9 Click 🗹 Done.

GEOMETRY I

- I In the Geometry toolbar, click Insert Sequence and choose Insert Sequence.
- **2** Browse to the model's Application Libraries folder and double-click the file biventricular cardiac model geom sequence.mph.
- 3 In the Geometry toolbar, click 🟢 Build All.
- **4** Click the **v Go to Default View** button in the **Graphics** toolbar.

Complete geometry instructions can be found in the Appendix - Geometry Modeling Instructions section.

GLOBAL DEFINITIONS

Heart Geometry Parameters

I In the Model Builder window, under Global Definitions click Parameters I.

2 In the Settings window for Parameters, type Heart Geometry Parameters in the Label text field.

DEFINITIONS

Create selections to identify the basal surface, the epicardium and the endocardium.

Basal Surface

- I In the **Definitions** toolbar, click **here Explicit**.
- 2 In the Settings window for Explicit, type Basal Surface in the Label text field.
- 3 Locate the Input Entities section. From the Geometric entity level list, choose Boundary.
- **4** Click the Transparency button in the Graphics toolbar.
- **5** Select Boundaries 1, 3, 9, and 17 only.

LV-Endocardium

- I In the **Definitions** toolbar, click 🗞 **Explicit**.
- 2 In the Settings window for Explicit, type LV-Endocardium in the Label text field.
- 3 Locate the Input Entities section. From the Geometric entity level list, choose Boundary.
- 4 Select Boundaries 6, 7, 12, and 16 only.

RV-Endocardium

- I In the **Definitions** toolbar, click **herefore Explicit**.
- 2 In the Settings window for Explicit, type RV-Endocardium in the Label text field.
- **3** Locate the **Input Entities** section. From the **Geometric entity level** list, choose **Boundary**.
- 4 Select Boundaries 20 and 22 only.

Epicardium

- I In the Definitions toolbar, click 🐚 Explicit.
- 2 In the Settings window for Explicit, type Epicardium in the Label text field.
- 3 Locate the Input Entities section. From the Geometric entity level list, choose Boundary.
- **4** Select Boundaries 4, 5, 10, 11, 15, 18, 19, and 21 only.

Endocardium

- I In the **Definitions** toolbar, click 💾 **Union**.
- 2 In the Settings window for Union, type Endocardium in the Label text field.
- 3 Locate the Geometric Entity Level section. From the Level list, choose Boundary.
- 4 Locate the Input Entities section. Under Selections to add, click + Add.

- 5 In the Add dialog box, in the Selections to add list, choose LV-Endocardium and RV-Endocardium.
- 6 Click OK.

GLOBAL DEFINITIONS

Structural Mechanics Parameters

- I In the Home toolbar, click Pi Parameters and choose Add>Parameters.
- 2 In the Settings window for Parameters, type Structural Mechanics Parameters in the Label text field.
- 3 Locate the Parameters section. Click 📂 Load from File.
- **4** Browse to the model's Application Libraries folder and double-click the file biventricular_cardiac_model_mechanical_passive_param.txt.

Electrical Parameters

- I In the Home toolbar, click Pi Parameters and choose Add>Parameters.
- 2 In the Settings window for Parameters, type Electrical Parameters in the Label text field.
- 3 Locate the Parameters section. Click 📂 Load from File.
- 4 Browse to the model's Application Libraries folder and double-click the file biventricular_cardiac_model_electrical_param.txt.

Active Stress Parameters

- I In the Home toolbar, click **P**i Parameters and choose Add>Parameters.
- 2 In the Settings window for Parameters, type Active Stress Parameters in the Label text field.
- 3 Locate the Parameters section. Click 📂 Load from File.
- 4 Browse to the model's Application Libraries folder and double-click the file biventricular_cardiac_model_active_stress_param.txt.

Conversion Factors Parameters

- I In the Home toolbar, click Pi Parameters and choose Add>Parameters.
- 2 In the Settings window for Parameters, type Conversion Factors Parameters in the Label text field.
- 3 Locate the Parameters section. Click 📂 Load from File.
- **4** Browse to the model's Application Libraries folder and double-click the file biventricular_cardiac_model_conversion_param.txt.

To compute the orientation of the fibers in any point in the myocardium it is necessary to know its distance from both the epicardium and the endocardium.

WALL DISTANCE: EPICARDIUM

- I In the Model Builder window, under Component I (compl) click Wall Distance (wd).
- 2 In the Settings window for Wall Distance, type Wall Distance: Epicardium in the Label text field.

Wall I

- I In the Physics toolbar, click 🔚 Boundaries and choose Wall.
- 2 In the Settings window for Wall, locate the Boundary Selection section.
- **3** From the Selection list, choose Epicardium.

WALL DISTANCE: ENDOCARDIUM

- I In the Model Builder window, under Component I (compl) click Wall Distance 2 (wd2).
- 2 In the Settings window for Wall Distance, type Wall Distance: Endocardium in the Label text field.
- 3 In the Physics toolbar, click 📄 Boundaries and choose Wall.

Wall I

- I In the Settings window for Wall, locate the Boundary Selection section.
- 2 From the Selection list, choose Endocardium.

Assuming that the sheets are oriented perpendicularly to the wall, their direction can be found by computing the transmural direction.

SHEET DIRECTION

- I In the Model Builder window, under Component I (comp1) click Curvilinear Coordinates (cc).
- 2 In the Settings window for Curvilinear Coordinates, type Sheet Direction in the Label text field.
- 3 Locate the Settings section. Select the Create base vector system check box.

Coordinate System Settings I

- I In the Model Builder window, under Component I (comp1)>Sheet Direction (cc) click Coordinate System Settings I.
- 2 In the Settings window for Coordinate System Settings, locate the Settings section.
- 3 From the Second basis vector list, choose z-axis.

Diffusion Method I

In the Physics toolbar, click 🔚 Domains and choose Diffusion Method.

Inlet I

- I In the Physics toolbar, click 层 Attributes and choose Inlet.
- 2 In the Settings window for Inlet, locate the Boundary Selection section.
- 3 From the Selection list, choose Endocardium.

Diffusion Method I

In the Model Builder window, click Diffusion Method I.

Outlet I

- I In the Physics toolbar, click 🔙 Attributes and choose Outlet.
- 2 In the Settings window for Outlet, locate the Boundary Selection section.
- 3 From the Selection list, choose Epicardium.

MESH I

- I In the Model Builder window, under Component I (compl) click Mesh I.
- 2 In the Settings window for Mesh, locate the Physics-Controlled Mesh section.
- **3** From the **Element size** list, choose **Finer**.
- 4 Click 📗 Build All.

The first study is used to obtain the sheets' direction and the distance from the walls. These quantities can be used to find the fibers' direction.

STUDY: FIBER DIRECTION

- I In the Model Builder window, click Study I.
- 2 In the Settings window for Study, type Study: Fiber Direction in the Label text field.
- **3** In the **Home** toolbar, click **= Compute**.

RESULTS

Wall Distance: Epicardium

In the **Settings** window for **3D Plot Group**, type Wall Distance: Epicardium in the **Label** text field.

Wall Distance: Endocardium

I In the Model Builder window, under Results click 3D Plot Group 2.

2 In the Settings window for 3D Plot Group, type Wall Distance: Endocardium in the Label text field.

Coordinate system (cc), Vector Field (cc), Wall Distance: Endocardium, Wall Distance: Epicardium

- In the Model Builder window, under Results, Ctrl-click to select
 Wall Distance: Epicardium, Wall Distance: Endocardium, Vector Field (cc), and
 Coordinate system (cc).
- 2 Right-click and choose Group.

Study: Fiber Direction

In the Settings window for Group, type Study: Fiber Direction in the Label text field.

Add variables related to the fibers orientation.

DEFINITIONS

Fiber Orientation

- I In the Model Builder window, expand the Component I (compl)>Definitions node.
- 2 Right-click **Definitions** and choose **Variables**.
- 3 In the Settings window for Variables, type Fiber Orientation in the Label text field.
- 4 Locate the Variables section. Click 📂 Load from File.
- 5 Browse to the model's Application Libraries folder and double-click the file biventricular_cardiac_model_fibers.txt.

The fibers reference system is obtained by rotating the curvilinear coordinate system found in the previous study.

Rotated System 2 (sys2)

- I In the Definitions toolbar, click \sum_{x}^{y} Coordinate Systems and choose Rotated System.
- 2 In the Settings window for Rotated System, locate the Rotation section.
- **3** Find the **Euler angles (Z-X-Z)** subsection. In the β text field, type theta.

Fiber Reference System

- I In the Definitions toolbar, click \sum_{x}^{y} Coordinate Systems and choose Composite System.
- 2 In the Settings window for Composite System, type Fiber Reference System in the Label text field.
- 3 Locate the Input Systems section. From the Base system list, choose Curvilinear System (cc) (cc_cs).

4 From the Relative system list, choose Rotated System 2 (sys2).

Create a cylindrical system for postprocessing purposes.

Cylindrical System 4 (sys4)

- I In the Definitions toolbar, click \sum_{x}^{y} Coordinate Systems and choose Cylindrical System.
- 2 In the Settings window for Cylindrical System, locate the Coordinate Names section.
- 3 From the Frame list, choose Material (X, Y, Z).

Boundary System 1 (sys1), Curvilinear System (cc) (cc_cs), Cylindrical System 4 (sys4), Fiber Reference System (sys3), Rotated System 2 (sys2)

- I In the Model Builder window, under Component I (comp1)>Definitions, Ctrl-click to select Boundary System I (sys1), Curvilinear System (cc) (cc_cs), Rotated System 2 (sys2), Fiber Reference System (sys3), and Cylindrical System 4 (sys4).
- 2 Right-click and choose Group.

Coordinate Systems

- I In the Settings window for Group, type Coordinate Systems in the Label text field.
- 2 In the Model Builder window, collapse the Coordinate Systems node.

Now we have all information needed to simulate the contraction. Add a Solid Mechanics interface and three PDE interfaces. The three PDEs will be used to solve the monodomain equation and to compute the active stress.

ADD PHYSICS

- I In the Home toolbar, click 🙀 Add Physics to open the Add Physics window.
- 2 Go to the Add Physics window.
- 3 In the tree, select Structural Mechanics>Solid Mechanics (solid).
- 4 Find the Physics interfaces in study subsection. In the table, clear the Solve check box for Study: Fiber Direction.
- 5 Click Add to Component I in the window toolbar.
- 6 In the tree, select Mathematics>PDE Interfaces>Coefficient Form PDE (c).
- 7 In the table, clear the Solve check box for Study: Fiber Direction.
- 8 Click Add to Component I in the window toolbar.
- 9 In the tree, select Mathematics>PDE Interfaces>Coefficient Form PDE (c).
- IO In the table, clear the Solve check box for Study: Fiber Direction.
- II Click Add to Component I in the window toolbar.

12 In the tree, select Mathematics>PDE Interfaces>Coefficient Form PDE (c).

I3 In the table, clear the Solve check box for Study: Fiber Direction.

14 Click Add to Component 1 in the window toolbar.

- IS In the tree, select Mathematics>PDE Interfaces>Coefficient Form PDE (c).
- I6 In the table, clear the Solve check box for Study: Fiber Direction.
- 17 In the Home toolbar, click 🙀 Add Physics to close the Add Physics window.

SOLID MECHANICS (SOLID)

- I In the Settings window for Solid Mechanics, locate the Structural Transient Behavior section.
- 2 From the list, choose Quasistatic.
- **3** Click to expand the **Discretization** section. From the **Displacement field** list, choose **Linear**.

Hyperelastic Material I

- I Right-click Component I (comp1)>Solid Mechanics (solid) and choose Material Models> Hyperelastic Material.
- 2 In the Settings window for Hyperelastic Material, locate the Domain Selection section.
- **3** From the Selection list, choose All domains.

Fiber 1

In the Physics toolbar, click 层 Attributes and choose Fiber.

MATERIALS

Material I (mat1)

- I In the Model Builder window, under Component I (comp1) right-click Materials and choose Blank Material.
- 2 In the Settings window for Material, locate the Material Contents section.
- **3** In the table, enter the following settings:

| Property | Variable | Value | Unit | Property group |
|--------------------------|----------|-------------|-------|----------------------------|
| Lamé parameter λ | lambLame | lambda_lame | N/m² | Lamé parameters |
| Lamé parameter μ | muLame | mu_lame | N/m² | Lamé parameters |
| Density | rho | rhos | kg/m³ | Basic |
| Fiber stiffness | kIHGO | af | Pa | Holzapfel-Gasser- Ogden |

| Property | Variable | Value | Unit | Property group |
|------------------|----------|-------|------|----------------------------|
| Model parameter | k2HGO | bf | I | Holzapfel-Gasser- Ogden |
| Fiber dispersion | k3HGO | 0 | I | Holzapfel-Gasser- Ogden |

ELECTROPHYSIOLOGY: TRANSMEMBRANE POTENTIAL (PHI)

- I In the Model Builder window, under Component I (compl) click Coefficient Form PDE (c).
- 2 In the Settings window for Coefficient Form PDE, type Electrophysiology: Transmembrane Potential (Phi) in the Label text field.
- **3** Locate the Units section. Click i Define Dependent Variable Unit.
- 4 In the Dependent variable quantity table, enter the following settings:

| Dependent variable quantity | Unit |
|-----------------------------|------|
| Custom unit | V |

5 In the Source term quantity table, enter the following settings:

| Source term quantity | Unit |
|----------------------|-------|
| Custom unit | A/m^3 |

- 6 Click to expand the Discretization section. From the Element order list, choose Linear.
- 7 From the Frame list, choose Material.
- 8 Click to expand the Dependent Variables section. In the Field name text field, type Phi.
- 9 In the **Dependent variables** table, enter the following settings:

Phi

ELECTROPHYSIOLOGY: CONDUCTANCE OF SLOW PROCESSES (R)

- I In the Model Builder window, under Component I (comp1) click Coefficient Form PDE 2 (c2).
- 2 In the Settings window for Coefficient Form PDE, type Electrophysiology: Conductance of Slow Processes (r) in the Label text field.
- **3** Click to expand the **Discretization** section. Locate the **Units** section. In the **Source term quantity** table, enter the following settings:

| Source term quantity | Unit |
|----------------------|------|
| Custom unit | 1/s |

- 4 Locate the Discretization section. From the Element order list, choose Linear.
- 5 From the Frame list, choose Material.
- 6 Locate the Dependent Variables section. In the Field name text field, type r.
- 7 In the **Dependent variables** table, enter the following settings:

r

ACTIVE STRESS (SA)

- I In the Model Builder window, under Component I (comp1) click Coefficient Form PDE 3 (c3).
- 2 In the Settings window for Coefficient Form PDE, type Active Stress (Sa) in the Label text field.
- 3 Locate the Units section. Click 🖬 Define Dependent Variable Unit.
- 4 In the **Dependent variable quantity** table, enter the following settings:

| Dependent variable quantity | Unit | |
|-----------------------------|-------|--|
| Custom unit | N/m^2 | |

5 In the Source term quantity table, enter the following settings:

| Source term quantity | Unit |
|----------------------|---------|
| Custom unit | N/m^2/s |

- 6 Locate the Discretization section. From the Element order list, choose Linear.
- 7 From the Frame list, choose Material.
- 8 Locate the Dependent Variables section. In the Field name text field, type Sa.
- 9 In the Dependent variables table, enter the following settings:

Sa

DEFINITIONS

Electrophysiology Variables

- I In the Model Builder window, under Component I (compl) right-click Definitions and choose Variables.
- 2 In the Settings window for Variables, type Electrophysiology Variables in the Label text field.
- **3** Locate the Variables section. Click *b* Load from File.

4 Browse to the model's Application Libraries folder and double-click the file biventricular_cardiac_model_electrophysiology.txt.

To compute the internal volume of the chamber during the contraction using Gauss' theorem we need to integrate over the internal boundaries.

Integration 1 (intop1)

- I In the Definitions toolbar, click 🖉 Nonlocal Couplings and choose Integration.
- 2 In the Settings window for Integration, type intLV in the Operator name text field.
- **3** Locate the **Source Selection** section. From the **Geometric entity level** list, choose **Boundary**.
- 4 From the Selection list, choose LV-Endocardium.

Integration 2 (intLV2)

- I Right-click Integration I (intLV) and choose Duplicate.
- 2 In the Settings window for Integration, type intRV in the Operator name text field.
- 3 Locate the Source Selection section. Click 🚺 Clear Selection.
- 4 Select Boundaries 19–22 only.

Ventricular Internal Volume

- I In the Model Builder window, right-click Definitions and choose Variables.
- 2 In the Settings window for Variables, type Ventricular Internal Volume in the Label text field.
- 3 Locate the Variables section. In the table, enter the following settings:

| Name | Expression | Unit | Description |
|----------|-------------------------------|------|------------------------------------|
| volumeLV | <pre>intLV(-x*solid.nx)</pre> | m³ | Left ventricle internal volume |
| volumeRV | <pre>intRV(-x*solid.nx)</pre> | m³ | Right ventricle internal volume |

Define the partial differential equations to solve in order to obtain the active stress.

ELECTROPHYSIOLOGY: TRANSMEMBRANE POTENTIAL (PHI) (C)

Coefficient Form PDE 1

- I In the Model Builder window, expand the Component I (comp1)> Electrophysiology: Transmembrane Potential (Phi) (c) node, then click Coefficient Form PDE I.
- 2 In the Settings window for Coefficient Form PDE, locate the Diffusion Coefficient section.

- 3 From the list, choose Symmetric.
- **4** In the *c* table, enter the following settings:

| D_iso+DfibXX | DfibXY | DfibXZ |
|--------------|--------------|--------------|
| DfibXY | D_iso+DfibYY | DfibYZ |
| DfibXZ | DfibYZ | D_iso+DfibZZ |

- **5** Locate the **Source Term** section. In the *f* text field, type Chi_m*(Ie+Im).
- 6 Locate the Damping or Mass Coefficient section. In the d_a text field, type Chi_m*Cm.

Initial Values 1

- I In the Model Builder window, click Initial Values I.
- 2 In the Settings window for Initial Values, locate the Initial Values section.
- 3 In the Phi text field, type Phir+70[mV]*(Z>-1e-4[mm])*(X<31[mm])*(X>0[mm])*
 (Y>-10[mm])*(Y<10[mm]).</pre>

ELECTROPHYSIOLOGY: CONDUCTANCE OF SLOW PROCESSES (R) (C2)

Coefficient Form PDE 1

- I In the Model Builder window, expand the Component I (comp1)> Electrophysiology: Conductance of Slow Processes (r) (c2) node, then click Coefficient Form PDE 1.
- 2 In the Settings window for Coefficient Form PDE, locate the Diffusion Coefficient section.
- **3** In the *c* text field, type **0**.
- 4 Locate the Absorption Coefficient section. In the a text field, type (1/betat)*(gamma+(mu1/(phi+mu2))*c*phi*(phi-b-1)).
- 5 Locate the Source Term section. In the f text field, type (1/betat)*(-gamma*c*phi* (phi-b-1)-mu1/(phi+mu2)*r^2).

ACTIVE STRESS (SA) (C3)

Coefficient Form PDE 1

- I In the Model Builder window, expand the Component I (compl)>Active Stress (Sa) (c3) node, then click Coefficient Form PDE I.
- 2 In the Settings window for Coefficient Form PDE, locate the Diffusion Coefficient section.
- **3** In the *c* text field, type **0**.
- **4** Locate the **Absorption Coefficient** section. In the *a* text field, type eps_delay.
- **5** Locate the **Source Term** section. In the *f* text field, type **eps_delay***kT*(**Phi-Phir**).

SOLID MECHANICS (SOLID)

In the Model Builder window, expand the Component I (compl)>Solid Mechanics (solid) node.

Fiber 1

- I In the Model Builder window, expand the Component I (comp1)>Solid Mechanics (solid)> Hyperelastic Material I node, then click Fiber I.
- 2 In the Settings window for Fiber, locate the Coordinate System Selection section.
- 3 From the Coordinate system list, choose Fiber Reference System (sys3).
- 4 Locate the **Orientation** section. From the **a** list, choose **Third axis**.

The coupling between the potential and the deformation is obtained through the active stress. The active stress is added as an external stress. Use the fiber coordinate system to easily input the components.

Hyperelastic Material I

In the Model Builder window, click Hyperelastic Material I.

External Stress 1

- I In the Physics toolbar, click 📃 Attributes and choose External Stress.
- 2 In the Settings window for External Stress, locate the Coordinate System Selection section.
- 3 From the Coordinate system list, choose Fiber Reference System (sys3).
- 4 Locate the External Stress section. From the list, choose Diagonal.
- **5** In the S_{ext} table, enter the following settings:

| vs*Sa | 0 | 0 |
|-------|-------|-------|
| 0 | vn*Sa | 0 |
| 0 | 0 | vf*Sa |

Prescribed Displacement I

- I In the Physics toolbar, click 🔚 Boundaries and choose Prescribed Displacement.
- **2** In the **Settings** window for **Prescribed Displacement**, locate the **Boundary Selection** section.
- 3 From the Selection list, choose Basal Surface.
- **4** Locate the **Prescribed Displacement** section. Select the **Prescribed in x direction** check box.
- **5** Select the **Prescribed in y direction** check box.
- 6 Select the Prescribed in z direction check box.

ADD STUDY

- I In the Home toolbar, click 2 Add Study to open the Add Study window.
- 2 Go to the Add Study window.
- 3 Find the Physics interfaces in study subsection. In the table, clear the Solve check boxes for Wall Distance: Epicardium (wd), Wall Distance: Endocardium (wd2), and Sheet Direction (cc).
- 4 Find the Studies subsection. In the Select Study tree, select General Studies> Time Dependent.
- 5 Click Add Study in the window toolbar.
- 6 In the Home toolbar, click Add Study to close the Add Study window.

STUDY: EXCITATION-CONTRACTION

- I In the Model Builder window, click Study 2.
- 2 In the Settings window for Study, type Study: Excitation-Contraction in the Label text field.

Set up the solver and initialize the solution to generate the Fiber default plot.

Step 1: Time Dependent

- I In the Model Builder window, under Study: Excitation-Contraction click Step I: Time Dependent.
- 2 In the Settings window for Time Dependent, locate the Study Settings section.
- 3 From the Time unit list, choose ms.
- 4 In the **Output times** text field, type range(0,5,320).
- 5 Click to expand the Values of Dependent Variables section. Find the Values of variables not solved for subsection. From the Settings list, choose User controlled.
- 6 From the Method list, choose Solution.
- 7 From the Study list, choose Study: Fiber Direction, Stationary.

RESULTS

Study: Excitation-Contraction/Solution 2 (sol2)

- I In the Model Builder window, expand the Results>Datasets node, then click Study: Excitation-Contraction/Solution 2 (sol2).
- 2 In the Settings window for Solution, locate the Solution section.

3 From the Frame list, choose Material (X, Y, Z).

Active Stress (Sa), Electrophysiology: Conductance of Slow Processes (r), Electrophysiology: Transmembrane Potential (Phi), Stress (solid)

I In the Model Builder window, under Results, Ctrl-click to select Stress (solid), Electrophysiology: Transmembrane Potential (Phi), Electrophysiology: Conductance of Slow Processes (r), and Active Stress (Sa).

Remove defaults plots that are not used.

2 Right-click and choose **Delete**.

Here the steps to obtain the model thumbnail are shown.

3 In the Home toolbar, click **Markov Add Predefined Plot**.

ADD PREDEFINED PLOT

- I Go to the Add Predefined Plot window.
- 2 In the tree, select Study: Excitation-Contraction/Solution 2 (sol2)>Solid Mechanics> Fibers (solid)>Fiber, Hyperelastic Material I (solid).
- **3** Click **Add Plot** in the window toolbar.
- **4** In the **Home** toolbar, click **I** Add **Predefined Plot**.

RESULTS

Fiber

- I In the Model Builder window, under Results click Fiber, Hyperelastic Material I (solid).
- 2 In the Settings window for 3D Plot Group, type Fiber in the Label text field.
- 3 Click to expand the Title section. In the Title text area, type Fiber orientation.
- **4** In the **Fiber** toolbar, click **I** Plot.

Endocardium

- I In the Model Builder window, expand the Fiber node, then click Fiber I.
- 2 In the Settings window for Streamline, type Endocardium in the Label text field.
- **3** Locate the **Streamline Positioning** section. From the **Positioning** list, choose **On selected boundaries**.
- 4 From the Point distribution list, choose Mesh based.
- 5 In the **Element refinement** text field, type 2.
- 6 Locate the Selection section. Click to select the 🔲 Activate Selection toggle button.
- 7 Select Boundaries 1–3, 8, 9, 13, 14, and 17 only.

8 Locate the Coloring and Style section. Find the Line style subsection. From the Type list, choose Line.

Filter I

Right-click Endocardium and choose Filter.

Color Expression

- I In the Model Builder window, expand the Results>Fiber>Endocardium node, then click Color Expression.
- 2 In the Settings window for Color Expression, locate the Expression section.
- 3 In the Expression text field, type theta.
- **4** In the **Unit** field, type deg.
- 5 Locate the Coloring and Style section. Click Change Color Table.
- 6 In the Color Table dialog box, select Rainbow>Rainbow in the tree.
- 7 Click OK.
- 8 In the Settings window for Color Expression, click to expand the Range section.
- 9 Select the Manual color range check box.
- **IO** In the **Minimum** text field, type -60.
- II In the Maximum text field, type 60.

Filter I

- I In the Model Builder window, click Filter I.
- 2 In the Settings window for Filter, locate the Element Selection section.
- **3** In the Logical expression for inclusion text field, type Beta>0.95.

Myocardium

- I In the Model Builder window, right-click Endocardium and choose Duplicate.
- 2 In the Settings window for Streamline, type Myocardium in the Label text field.
- **3** Locate the Selection section. Click Clear Selection.
- 4 Select Boundaries 2, 8, 13, and 14 only.
- 5 Locate the Streamline Positioning section. In the Element refinement text field, type 1.
- 6 Click to expand the Inherit Style section. From the Plot list, choose Endocardium.

Filter 1

- I In the Model Builder window, expand the Myocardium node, then click Filter I.
- 2 In the Settings window for Filter, locate the Element Selection section.

3 In the Logical expression for inclusion text field, type (Beta>0.45)*(Beta<0.55)*(Z<-cL/9).

Epicardium

- I In the Model Builder window, right-click Myocardium and choose Duplicate.
- 2 In the Settings window for Streamline, type Epicardium in the Label text field.
- 3 Locate the Selection section. Click 📉 Clear Selection.
- 4 Select Boundaries 2, 3, 5, 8–11, 13–15, 17, and 18 only.
- **5** Locate the **Streamline Positioning** section. In the **Element refinement** text field, type **3**.

Filter 1

- I In the Model Builder window, expand the Epicardium node, then click Filter I.
- 2 In the Settings window for Filter, locate the Element Selection section.
- 3 In the Logical expression for inclusion text field, type Beta<0.05*((sys4.phi>100[deg])*(sys4.phi<190[deg])*(Z>-cL/3) || (Z<-cL/3)).</pre>
- **4** In the **Fiber** toolbar, click **I** Plot.
- **5** Click the **Show Grid** button in the **Graphics** toolbar.
- 6 Click the **Zoom Extents** button in the **Graphics** toolbar.

Create a 3D plot to display the active potential distribution while the simulation runs.

Potential

- I In the Home toolbar, click 🚛 Add Plot Group and choose 3D Plot Group.
- 2 In the Settings window for 3D Plot Group, type Potential in the Label text field.
- 3 Locate the Data section. From the Dataset list, choose Study: Excitation-Contraction/ Solution 2 (sol2).

Volume 1

- I Right-click Potential and choose Volume.
- 2 In the Settings window for Volume, locate the Expression section.
- 3 In the Expression text field, type Phi.
- 4 From the **Unit** list, choose **mV**.
- 5 Click to expand the Range section. Select the Manual color range check box.
- 6 In the Minimum text field, type -80.
- 7 In the Maximum text field, type 20.

Deformation I

- I Right-click Volume I and choose Deformation.
- 2 In the Settings window for Deformation, locate the Scale section.
- 3 Select the Scale factor check box. In the associated text field, type 1.

STUDY: EXCITATION-CONTRACTION

- I In the Model Builder window, click Study: Excitation-Contraction.
- 2 In the Settings window for Study, locate the Study Settings section.
- 3 Clear the Generate default plots check box.

Step 1: Time Dependent

- I In the Model Builder window, expand the Study: Excitation-Contraction node, then click Step I: Time Dependent.
- **2** In the **Settings** window for **Time Dependent**, click to expand the **Results While Solving** section.
- **3** Select the **Plot** check box.
- 4 From the Plot group list, choose Potential.
- **5** In the **Home** toolbar, click **= Compute**.

RESULTS

Contraction, Snapshots

- I In the Home toolbar, click 🚛 Add Plot Group and choose 3D Plot Group.
- 2 In the Settings window for 3D Plot Group, type Contraction, Snapshots in the Label text field.
- 3 Locate the Data section. From the Dataset list, choose Study: Excitation-Contraction/ Solution 2 (sol2).
- 4 Click to expand the Title section. From the Title type list, choose Manual.
- 5 In the Title text area, type Potential and deformation.
- 6 Clear the Parameter indicator text field.
- 7 Click to expand the Plot Array section. Select the Enable check box.
- 8 From the Array shape list, choose Square.
- 9 From the Array plane list, choose xz.
- **IO** Click the $\begin{bmatrix} xz \end{bmatrix}$ **Go to XZ View** button in the **Graphics** toolbar.

Slice 1

- I Right-click Contraction, Snapshots and choose Slice.
- 2 In the Settings window for Slice, locate the Data section.
- 3 From the Dataset list, choose Study: Excitation-Contraction/Solution 2 (sol2).
- 4 From the Time (ms) list, choose 5.
- 5 Locate the Expression section. In the Expression text field, type Phi.
- 6 From the **Unit** list, choose **mV**.
- 7 Locate the Plane Data section. From the Plane list, choose ZX-planes.
- 8 In the Planes text field, type 1.
- 9 Click to expand the Plot Array section. Select the Manual indexing check box.

Deformation 1

- I Right-click Slice I and choose Deformation.
- 2 In the Settings window for Deformation, locate the Scale section.
- **3** Select the **Scale factor** check box. In the associated text field, type 1.
- **4** In the **Contraction, Snapshots** toolbar, click **I Plot**.

Slice 2

- I In the Model Builder window, under Results>Contraction, Snapshots right-click Slice I and choose Duplicate.
- 2 In the Settings window for Slice, click to expand the Title section.
- **3** From the **Title type** list, choose **None**.
- 4 Locate the Plane Data section. From the Plane list, choose XY-planes.
- **5** In the **Planes** text field, type 1.
- 6 Locate the Coloring and Style section. Clear the Color legend check box.
- 7 Click to expand the Inherit Style section. From the Plot list, choose Slice 1.

Annotation I

- I In the Model Builder window, right-click Contraction, Snapshots and choose Annotation.
- 2 In the Settings window for Annotation, locate the Annotation section.
- 3 In the **Text** text field, type t=5 ms.
- **4** Locate the **Position** section. In the **Z** text field, type **20**.
- 5 Click to expand the Plot Array section. Select the Manual indexing check box.

Annotation 1, Slice 1, Slice 2

- I In the Model Builder window, under Results>Contraction, Snapshots, Ctrl-click to select Slice I, Slice 2, and Annotation I.
- 2 Right-click and choose **Duplicate**.

Slice 3

- I In the Settings window for Slice, click to expand the Title section.
- 2 From the Title type list, choose None.
- **3** Locate the **Coloring and Style** section. Clear the **Color legend** check box.
- 4 Locate the Inherit Style section. From the Plot list, choose Slice I.
- 5 Locate the Data section. From the Time (ms) list, choose 35.
- 6 Locate the Plot Array section. In the Column index text field, type 1.

Slice 4

- I In the Model Builder window, click Slice 4.
- 2 In the Settings window for Slice, locate the Data section.
- 3 From the Time (ms) list, choose 35.
- 4 Locate the Plot Array section. In the Column index text field, type 1.

Annotation 2

- I In the Model Builder window, click Annotation 2.
- 2 In the Settings window for Annotation, locate the Annotation section.
- 3 In the Text text field, type t=35 ms.
- **4** Locate the **Plot Array** section. In the **Column index** text field, type 1.
- **5** In the **Contraction, Snapshots** toolbar, click **I Plot**.
- 6 Click the 4 Zoom Extents button in the Graphics toolbar.

Annotation 2, Slice 3, Slice 4

- I In the Model Builder window, under Results>Contraction, Snapshots, Ctrl-click to select Slice 3, Slice 4, and Annotation 2.
- 2 Right-click and choose **Duplicate**.

Slice 5

- I In the Settings window for Slice, locate the Data section.
- 2 From the Time (ms) list, choose 55.
- 3 Locate the Plot Array section. In the Column index text field, type 2.

Slice 6

- I In the Model Builder window, click Slice 6.
- 2 In the Settings window for Slice, locate the Data section.
- 3 From the Time (ms) list, choose 55.
- 4 Locate the Plot Array section. In the Column index text field, type 2.

Annotation 3

- I In the Model Builder window, click Annotation 3.
- 2 In the Settings window for Annotation, locate the Annotation section.
- 3 In the Text text field, type t=50 ms.
- 4 Locate the Plot Array section. In the Column index text field, type 2.
- 5 In the Contraction, Snapshots toolbar, click 💽 Plot.
- **6** Click the |+ **Zoom Extents** button in the **Graphics** toolbar.

Annotation 3, Slice 5, Slice 6

- I In the Model Builder window, under Results>Contraction, Snapshots, Ctrl-click to select Slice 5, Slice 6, and Annotation 3.
- 2 Right-click and choose Duplicate.

Slice 7

- I In the Settings window for Slice, locate the Data section.
- 2 From the Time (ms) list, choose 100.
- 3 Locate the Plot Array section. In the Column index text field, type 3.

Slice 8

- I In the Model Builder window, click Slice 8.
- 2 In the Settings window for Slice, locate the Data section.
- 3 From the Time (ms) list, choose 100.
- 4 Locate the Plot Array section. In the Column index text field, type 3.

Annotation 4

- I In the Model Builder window, click Annotation 4.
- 2 In the Settings window for Annotation, locate the Annotation section.
- 3 In the **Text** text field, type t=100 ms.
- 4 Locate the Plot Array section. In the Column index text field, type 3.
- 5 In the Contraction, Snapshots toolbar, click **I** Plot.

6 Click the + Zoom Extents button in the Graphics toolbar.

Annotation 4, Slice 7, Slice 8

- I In the Model Builder window, under Results>Contraction, Snapshots, Ctrl-click to select Slice 7, Slice 8, and Annotation 4.
- 2 Right-click and choose Duplicate.

Slice 9

- I In the Settings window for Slice, locate the Data section.
- 2 From the Time (ms) list, choose 145.
- 3 Click to expand the Plot Array section. In the Row index text field, type -1.
- **4** In the **Column index** text field, type **0**.

Slice 10

- I In the Model Builder window, click Slice 10.
- 2 In the Settings window for Slice, locate the Data section.
- 3 From the Time (ms) list, choose 145.
- 4 Locate the Plot Array section. In the Row index text field, type -1.
- **5** In the **Column index** text field, type **0**.

Annotation 5

- I In the Model Builder window, click Annotation 5.
- 2 In the Settings window for Annotation, locate the Annotation section.
- 3 In the Text text field, type t=145 ms.
- 4 Locate the **Position** section. In the **Z** text field, type -80.
- 5 Click to expand the Plot Array section. In the Row index text field, type -1.
- 6 In the Column index text field, type 0.
- 7 In the Contraction, Snapshots toolbar, click **I** Plot.
- 8 Click the 4 Zoom Extents button in the Graphics toolbar.

Annotation 5, Slice 10, Slice 9

- I In the Model Builder window, under Results>Contraction, Snapshots, Ctrl-click to select Slice 9, Slice 10, and Annotation 5.
- 2 Right-click and choose **Duplicate**.

Slice 11

I In the Settings window for Slice, locate the Data section.

- 2 From the Time (ms) list, choose 200.
- 3 Locate the Plot Array section. In the Column index text field, type 1.

Slice 12

- I In the Model Builder window, click Slice 12.
- 2 In the Settings window for Slice, locate the Data section.
- 3 From the Time (ms) list, choose 200.
- 4 Locate the Plot Array section. In the Column index text field, type 1.

Annotation 6

- I In the Model Builder window, click Annotation 6.
- 2 In the Settings window for Annotation, locate the Annotation section.
- 3 In the **Text** text field, type t=200 ms.
- 4 Locate the Plot Array section. In the Column index text field, type 1.
- 5 In the Contraction, Snapshots toolbar, click 💿 Plot.
- 6 Click the **Zoom Extents** button in the **Graphics** toolbar.

Annotation 6, Slice 11, Slice 12

- I In the Model Builder window, under Results>Contraction, Snapshots, Ctrl-click to select Slice 11, Slice 12, and Annotation 6.
- 2 Right-click and choose Duplicate.

Annotation 7, Slice 13, Slice 14

- I In the Model Builder window, under Results>Contraction, Snapshots, Ctrl-click to select Slice 13, Slice 14, and Annotation 7.
- 2 In the Settings window for Slice, locate the Data section.
- 3 From the Time (ms) list, choose 250.
- 4 Locate the Plot Array section. In the Column index text field, type 2.

Slice 14

- I In the Model Builder window, click Slice 14.
- 2 In the Settings window for Slice, locate the Data section.
- 3 From the Time (ms) list, choose 250.
- 4 Locate the Plot Array section. In the Column index text field, type 2.

Annotation 7

I In the Model Builder window, click Annotation 7.

- 2 In the Settings window for Annotation, locate the Plot Array section.
- **3** In the **Column index** text field, type **2**.
- 4 Locate the Annotation section. In the Text text field, type t=250 ms.
- 5 In the Contraction, Snapshots toolbar, click 🗿 Plot.
- **6** Click the $4 \rightarrow$ **Zoom Extents** button in the **Graphics** toolbar.

Annotation 7, Slice 13, Slice 14

- I In the Model Builder window, under Results>Contraction, Snapshots, Ctrl-click to select Slice 13, Slice 14, and Annotation 7.
- 2 Right-click and choose **Duplicate**.

Slice 15

- I In the Settings window for Slice, locate the Data section.
- 2 From the Time (ms) list, choose 300.
- 3 Locate the Plot Array section. In the Column index text field, type 3.

Slice 16

- I In the Model Builder window, click Slice 16.
- 2 In the Settings window for Slice, locate the Data section.
- 3 From the Time (ms) list, choose 300.
- 4 Locate the Plot Array section. In the Column index text field, type 3.

Annotation 8

- I In the Model Builder window, click Annotation 8.
- 2 In the Settings window for Annotation, locate the Annotation section.
- 3 In the Text text field, type t=300 ms.
- 4 Locate the Plot Array section. In the Column index text field, type 3.

Slice 2

Click the \longleftrightarrow Zoom Extents button in the Graphics toolbar.

Contraction, Snapshots

- I Click the Transparency button in the Graphics toolbar.
- 2 Click the **1** Orthographic Projection button in the Graphics toolbar.
- 3 In the Model Builder window, click Contraction, Snapshots.
- 4 In the Contraction, Snapshots toolbar, click on Plot.

Volume Change

- I In the Home toolbar, click 🚛 Add Plot Group and choose ID Plot Group.
- 2 In the Settings window for ID Plot Group, type Volume Change in the Label text field.
- 3 Locate the Data section. From the Dataset list, choose Study: Excitation-Contraction/ Solution 2 (sol2).
- 4 Locate the Plot Settings section. Select the y-axis label check box.
- 5 Click to expand the Title section. From the Title type list, choose None.
- 6 Locate the Plot Settings section. In the y-axis label text field, type Current volume (ml).
- 7 Locate the Legend section. From the Position list, choose Upper middle.

Global I

- I Right-click Volume Change and choose Global.
- 2 In the Settings window for Global, locate the y-Axis Data section.
- **3** In the table, enter the following settings:

| Expression | Unit | Description |
|------------|------|---------------------------------|
| volumeRV | ml | Right ventricle internal volume |
| volumeLV | ml | Left ventricle internal volume |

- 4 Click to expand the Legends section. From the Legends list, choose Manual.
- **5** In the table, enter the following settings:

Legends

Right ventricular chamber

Left ventricular chamber

6 In the Volume Change toolbar, click **I** Plot.

Point at Z=-1mm

- I In the **Results** toolbar, click **Cut Point 3D**.
- 2 In the Settings window for Cut Point 3D, type Point at Z=-1mm in the Label text field.
- 3 Locate the Data section. From the Dataset list, choose Study: Excitation-Contraction/ Solution 2 (sol2).
- 4 Locate the **Point Data** section. In the **X** text field, type 25.
- **5** In the **Y** text field, type **0**.
- 6 In the Z text field, type -1.

Point at Z=-40mm

- I Right-click Point at Z=-Imm and choose Duplicate.
- 2 In the Settings window for Cut Point 3D, type Point at Z=-40mm in the Label text field.
- 3 Locate the **Point Data** section. In the X text field, type 20.
- **4** In the **Y** text field, type 0.
- **5** In the **Z** text field, type -40.

Point at Z=-60mm

- I Right-click Point at Z=-40mm and choose Duplicate.
- 2 In the Settings window for Cut Point 3D, type Point at Z=-60mm in the Label text field.
- 3 Locate the Point Data section. In the X text field, type 0.
- **4** In the **Y** text field, type 0.
- **5** In the **Z** text field, type 60.

Activation Time

- I In the **Results** toolbar, click \sim **ID Plot Group**.
- 2 In the Settings window for ID Plot Group, type Activation Time in the Label text field.
- 3 Click to expand the Title section. From the Title type list, choose None.

Point Graph 1

- I Right-click Activation Time and choose Point Graph.
- 2 In the Settings window for Point Graph, locate the Data section.
- 3 From the Dataset list, choose Point at Z=-Imm.
- 4 Locate the y-Axis Data section. In the Expression text field, type Phi.
- 5 From the **Unit** list, choose **mV**.
- 6 Click to expand the Legends section. Select the Show legends check box.
- 7 From the Legends list, choose Manual.
- 8 In the table, enter the following settings:

Legends

Point at Z=-1 mm

Point Graph 2

- I Right-click Point Graph I and choose Duplicate.
- 2 In the Settings window for Point Graph, locate the Data section.
- 3 From the Dataset list, choose Point at Z=-40mm.

- 4 Click to expand the Title section. From the Title type list, choose None.
- 5 Locate the Legends section. In the table, enter the following settings:

Legends

Point at Z=-40 mm

Point Graph 3

- I Right-click Point Graph 2 and choose Duplicate.
- 2 In the Settings window for Point Graph, locate the Data section.
- **3** From the **Dataset** list, choose **Point at Z=-60mm**.
- 4 Locate the Legends section. In the table, enter the following settings:

Legends

Point at Z=-60 mm

5 In the Activation Time toolbar, click 💿 Plot.

Heart Deformation

I In the **Results** toolbar, click **Animation** and choose **Player**.

Create an animation to see the excitation path.

- 2 In the Settings window for Animation, type Heart Deformation in the Label text field.
- 3 Locate the Scene section. From the Subject list, choose Potential.
- 4 Locate the Frames section. In the Number of frames text field, type 50.
- **5** Click the \longleftrightarrow **Zoom Extents** button in the **Graphics** toolbar.

Appendix - Geometry Modeling Instructions

From the File menu, choose New.

NEW

In the New window, click 🔗 Model Wizard.

MODEL WIZARD

I In the Model Wizard window, click 间 3D.

2 Click **M** Done.

GLOBAL DEFINITIONS

Heart Geometry Parameters

- I In the Model Builder window, under Global Definitions click Parameters I.
- 2 In the Settings window for Parameters, type Heart Geometry Parameters in the Label text field.
- 3 Locate the Parameters section. Click 📂 Load from File.
- 4 Browse to the model's Application Libraries folder and double-click the file biventricular_cardiac_model_geom_param.txt.

GEOMETRY I

- I In the Model Builder window, under Component I (compl) click Geometry I.
- 2 In the Settings window for Geometry, locate the Units section.
- **3** From the **Length unit** list, choose **mm**.

Ellipsoid: Left Ventricle

- I In the Geometry toolbar, click \bigoplus More Primitives and choose Ellipsoid.
- 2 In the Settings window for Ellipsoid, type Ellipsoid: Left Ventricle in the Label text field.
- 3 Locate the Size and Shape section. In the a-semiaxis text field, type aL.
- 4 In the **b-semiaxis** text field, type bL.
- 5 In the c-semiaxis text field, type cL.
- 6 Click to expand the Layers section. In the table, enter the following settings:

| Layer name | Thickness (mm) |
|------------|----------------|
| Layer 1 | tL |

Ellipsoid: Right Ventricle

- I Right-click Ellipsoid: Left Ventricle and choose Duplicate.
- 2 In the Settings window for Ellipsoid, type Ellipsoid: Right Ventricle in the Label text field.
- 3 Locate the Size and Shape section. In the a-semiaxis text field, type aR.
- 4 In the **b-semiaxis** text field, type bR.
- 5 In the c-semiaxis text field, type cR.

6 Locate the Layers section. In the table, enter the following settings:

| Layer name | Thickness (mm) |
|------------|----------------|
| Layer 1 | tR |

Partition Domains I (pard I)

- I In the Geometry toolbar, click 📃 Booleans and Partitions and choose Partition Domains.
- **2** Click the Transparency button in the Graphics toolbar.
- 3 On the object elp2, select Domains 1, 3, 5, 6, and 8 only.
- 4 In the Settings window for Partition Domains, locate the Partition Domains section.
- 5 From the Partition with list, choose Faces.
- 6 On the object elp1, select Boundaries 16 and 22 only.

Delete Entities 1 (del1)

- I In the **Geometry** toolbar, click **III Delete**.
- 2 In the Settings window for Delete Entities, locate the Entities or Objects to Delete section.
- 3 From the Geometric entity level list, choose Domain.
- 4 On the object elp1, select Domains 2, 4, 5, 7, and 9 only.
- 5 On the object pard I, select Domains 1–7, 9, and 10 only.
- 6 Click 🟢 Build All Objects.

7 Click the **Zoom Extents** button in the **Graphics** toolbar.



Use virtual operations to ignore faces and edges to simplify the geometry and avoid unnecessary mesh refinement zones.

Ignore Faces 1 (igf1)

- I In the Geometry toolbar, click 🏠 Virtual Operations and choose Ignore Faces.
- 2 On the object fin, select Boundaries 11 and 20 only.

3 In the **Geometry** toolbar, click 📳 **Build All**.

