

Biological Tissue Movement

Oleksandr Hubanov
Vilnius Gediminas Technical University

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The mitral valve, also known as the bicuspid valve or left atrioventricular valve, is a valve with two flaps in the heart, that lies between the left atrium and the left ventricle. The mitral valve and the tricuspid valve are known collectively as the atrioventricular valves because they lie between the atria and the ventricles of the heart.

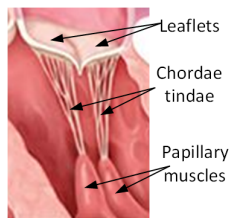


Figure 1: Mitral valve structure

Mitral valve has cyclic working conditions. The valve opens and closes because of pressure differences, opening when there is greater pressure in the left atrium than ventricle, and closing when there is greater pressure in the ventricle than atrium. In abnormal conditions, blood may flow backwards through the valve (mitral regurgitation) or the mitral valve may be narrowed (mitral stenosis). Mitral valve prolapse (MVP) is a valvular heart disease characterized by the displacement of an abnormally thickened mitral valve leaflet into the left atrium during systole. By other words, it is a condition in which the two flaps of the mitral valve doesn't close smoothly and evenly, but instead bulge (prolapse) upward into the left atrium. [Hayek2005a]

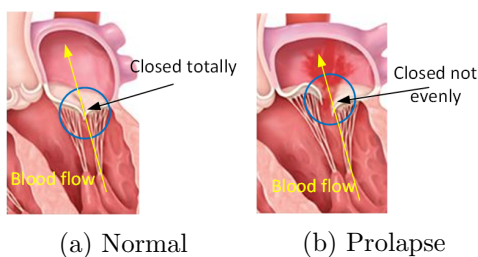


Figure 2: Prolapse in compare to normal

To solve such disease by surgical way, usually "methd" is used. link for article

Providing the surgeon with an anatomically and biomechanically accurate computational model of a particular patient's mitral heart valve could enable preoperative surgical planning and potentially improve surgical outcome.

There are a large number of different numerical modeling methods. All of them are derivatives of Finite Element Method(FEM) and Discrete Element Method(DEM). While general finite-element studies are helpful for the evaluation and development of MV repair surgery, patient-specific models are required for individual therapy planning. The patient-specific mass-spring MV model uses a segmentation of 3D TEE images for the initialization of a mass-spring model of the closed MV under systolic pressure. An iterative approach is used to adjust the spring rest-length so that the model can accurately simulate the shape of the closed MV under systolic pressure. To simulate MV annuloplasty, the model can then be deformed, according to the annuloplasty ring to be used, to create a prediction of the shape of the closed MV after surgery.

Based on the properties of the material of biological tissues and review of existing projects, the most appropriate method is Mass - Spring modeling(MSM). This method based on ideas of DEM and basic element here is very know in mechanic simple one dimensional(1D) beam.

Computational complexity of MSM is much less compare to FEM-based methods, because of less number of equations to integrate on each time step. This important advantage and physics way have method describes basic element gave to MSM very wide using in computer games for calculating reality-looks hair or cloth movement in real time. Modelling by using MSM could be parallel calculated on each time step.[1] [2]

Mathematical model

The motion of chordae as motion of physical object can be expressed by Newton's equation of motion:

$$m\ddot{x} + k\dot{x} + c\ddot{x} = \vec{F} \quad (1)$$

where F is internal force of object, m – mass of object and \ddot{x} is acceleration, k – stiffness coefficient of object, x is displacement of object and c – damping coefficient of object, \dot{x} is velocity of object.

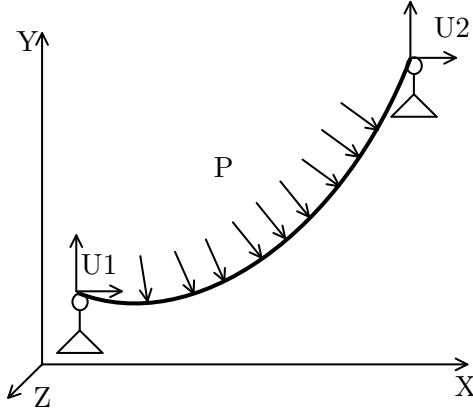


Figure 3: Chordae tindaie in global coordinate system

Initial conditions are: $x(0) = 0$ and $\dot{x}(0) = V_0$. Acceleration could be known from (1) as:

$$\ddot{x} = \vec{F}/m \quad (2)$$

Then Euler's scheme of integration can be described like:

$$\ddot{x}(t + \Delta t) = \ddot{x}(t) + \ddot{x}(t)\Delta t \quad (3)$$

$$\dot{x}(t + \Delta t) = \dot{x}(t) + \ddot{x}(t)\Delta t \quad (4)$$

Linear deformation

Internal object force becomes from physical deformation of object it self. In linear case of study, deformation of element much less compare to element dimensions. It is expressed by linear geometry equation(5), which showing relation between initial length of element and length in Δt state.

$$\varepsilon = \frac{dU}{dx} = \frac{l(\Delta t) - l(t)}{l(t)} \quad (5)$$

According to Hook law $\sigma = \varepsilon E$ and linear geometry equation (5), inner force can be described as:

$$\begin{aligned} N(x) &= \int_{Al} \varepsilon E dldA = EA \int_l \varepsilon dl \\ &= \frac{EA}{l(t)} * (l(t + \Delta t) - l(t)) \end{aligned} \quad (6)$$

where $l(\Delta t)$ is current length of chordae, $l(t)$ length of chordae at previous time moment, E – Young's modulus for chordae, A – volume of chordae. To be able to integrate equation of motion, need to express deformation in equation (6) by differences between displacements of nodes, to which chordae is connected:

$$N(x) = \frac{EA}{l(t)} * (x_i(t) - x_j(t)) \quad (7)$$

Nonlinear deformation

Nonlinearity in main mean that chordae can get huge deformation compare to its demensions. Equation of F_{elem} in this case would change to nonlinear form:

$$N(x) = \int_t \int_A \sigma dAdt \quad (8)$$

From this equation comes that cross sectional area and stiffness coefficient will get nonlinearity.

Changing of cross sectional area over time of chordae is changing its length over time. Linear geometry equation(5), showing linear relations between length, because difference in Δt state takes according initial length of chordae. In case of huge deformation need to recalculate length of chordae on each time step and take difference of displacement according to previous time step. In end of geometry equation become to nonlinear form:

$$\varepsilon = \frac{dU}{dx} = \frac{l(\Delta t) - l(t - \Delta t)}{l(\Delta t)} \quad (9)$$

Inner force(6) also become to nonlinear form:

$$\begin{aligned} N(x) &= \int_t \int_A \varepsilon E dA = EA \int_t \varepsilon \\ &= \frac{EA}{l(\Delta t)} * (l(\Delta t) - l(t - \Delta t)) \end{aligned} \quad (10)$$

where l is current length of element, l_0 length of element at $t = 0$, E – Young’s modulus for element material.

And nonlinear equation of inner force for integration:

$$N(x) = \frac{EA}{l_0} * (U_i - U_j) \quad (11)$$

Discrete model

Investigating motion of any mechanical system comes to integration motion equation of discrete parts of this system. Chordae tindaes presents itself as fibrous tissue and could be described as system of 1D rods, in case of getting mechanical equivalent of system. Example of such flat 2D system are shown on figure 5.

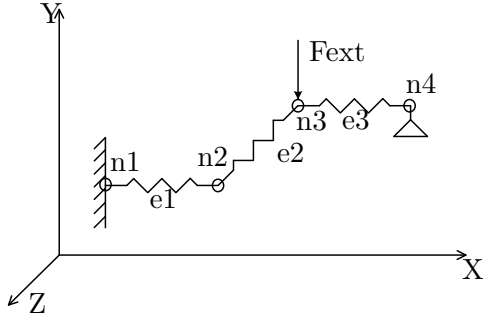


Figure 4: 1D Rod system in global coordinate system

System consists of discrete elements e_1 , e_2 and e_3 . All elements are connected to each other through nodes n_2 , n_3 and to special points through n_1 and n_4 . Each element e_n of system has own orientation in global coordinate system. From schematic representation of node comes that all vector variables of node should be calculated in global coordinate system and element variables in own local coordinate system (figure 5). For transformation between coordinate systems direction cosine matrix (DCM)(12) can be used.

$$DCM = \begin{bmatrix} \cos(X, x) & \cos(X, y) & \cos(X, z) \\ \cos(Y, x) & \cos(Y, y) & \cos(Y, z) \\ \cos(Z, x) & \cos(Z, y) & \cos(Z, z) \end{bmatrix} \quad (12)$$

where $\{X, Y, Z\}$ is global coordinate system and $\{x, y, z\}$ is local coordinate system.

According to primitive scheme of node 5, mass of each node can be calculated, like sum

of half mass of each element, which acting in node. $m_n = \sum_e m_e / 2$

In case that node does not have external interrupt, like pressure or other applied force, schematic represent of node can be as on figure 5.

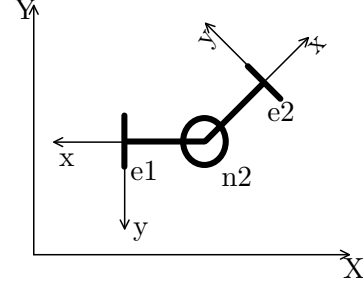


Figure 5: Extracted node from system

Mathematical model of discrete system is expressed by equations of nodes motion. All system acting in global coordinate system $\{X, Y, Z\}$ and each element acting in own local coordinate system $\{x, y, z\}$.

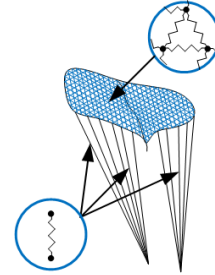


Figure 6: Displacement of papillary muscle over cardiac cycle

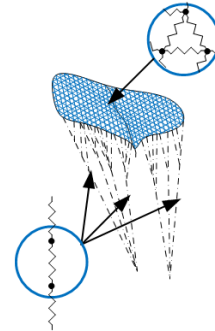


Figure 7: Displacement of papillary muscle over cardiac cycle

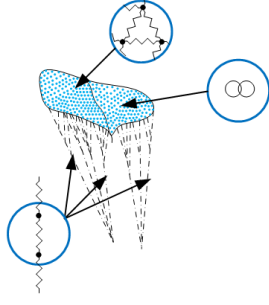


Figure 8: Displacement of papillary muscle over cardiac cycle

research aim is to compare 1 rod and rope of rods

If express kx and $c\dot{x}$ by $\sum F_{elem}$, then motion equation will become to:

$$\vec{F}_{inertia} = \vec{F}_{ext} - \sum \vec{F}_{elem} \quad (13)$$

F_{ext} is external load force, applied to node in global coordinates. Value of this force for each time step is loaded from list of loads. F_{elem} is sum of internal forces of each element, which acting in node. From each element counts only half of force to node, other half going to neighbour node. For 1D rod system F_{elem} can be expressed like:

$$F_{elem} = \sum_e N_e(x) \quad (14)$$

To words about mechanical properties of valve components, poisson ratio ν is same, 0.49 for all. Mean value of chordae length L 0,025m and diameter d 0,001m. Chordae has density $\rho = 1040 \text{ kg/m}^3$ and stiffness $E = 2000 \text{ N/m}$, when leaflets has $\rho = 1.06 \text{E}3 \text{ kg/m}^3$ and $E = 2 \text{ MPa}$ (Anterior leaflet), $E = 1 \text{ MPa}$ (Posterior leaflet)

Integration schemes

Euler first order explicit

$$\begin{aligned} v(t + \Delta t) &= v(t) + a(t) * \Delta t \\ x(t + \Delta t) &= x(t) + v(t) * \Delta t \end{aligned} \quad (15)$$

The central difference scheme, also known as velocity Verlet is a very widely used integration method of second order. This was initially proposed by Cundall and Strack (1979)

and adopted by several other authors. Velocities in the upcoming time step $t + \Delta t/2$ and positions at $t + \Delta t$ are calculated as

$$\begin{aligned} v(t + \Delta t/2) &= v(t + \Delta t/2) + a(t - \Delta t/2) * \Delta t \\ x(t + \Delta t) &= x(t) + v(t + \Delta t/2) * \Delta t \end{aligned} \quad (16)$$

predictor-corrector schemes very commonly used for molecular dynamics and discrete element applications are Gear's schemes. They are based on three stages, whereas in addition to the predictor and corrector step known from the Adams-method an evaluation step is added. Schemes considered here are the third order Gear's method (GPC3) and the fourth order Gear's method (GPC4). In the prediction step positions and their higher derivatives are calculated based on Taylor series expansions as

$$\begin{aligned} c(t + \Delta t, p) &= c(t) \\ b(t + \Delta t, p) &= b(t) + c(t) * \Delta t \\ a(t + \Delta t, p) &= a(t) + b(t) * \Delta t^2 + \frac{1}{6} * c(t) * \Delta t^2 \\ v(t + \Delta t, p) &= v(t) + a(t) * \Delta t + \frac{1}{2} * b(t) * \Delta t^2 + \\ &\quad \frac{1}{6} * c(t) * \Delta t^3 \\ x(t + \Delta t, p) &= x(t) + v(t) * \Delta t + \frac{1}{2} * a(t) * \Delta t^2 + \\ &\quad \frac{1}{6} * b(t) * \Delta t^3 + \frac{1}{24} * c(t) * \Delta t^4 \end{aligned} \quad (17)$$

with the first and second derivative of the accelerations calculated as

for GPC3:

$$b(t) = \frac{\Delta a(t)}{\Delta t} \quad (18)$$

$$c(t) = 0$$

for GPC4:

$$\begin{aligned} b(t) &= \frac{\Delta a(t)}{\Delta t} \\ c(t) &= \frac{\Delta b(t)}{\Delta t} \end{aligned} \quad (19)$$

In the evaluation step the difference in the accelerations calculated based on the acceleration $a(t + \Delta t, p)$ and the acceleration $a(t + \Delta t)$

calculated from positions $x(t + \Delta t, p)$ and velocities $v(t + \Delta t, p)$ is obtained by

$$\Delta a = a(t + \Delta t) - a(t + \Delta t, p) \quad (20)$$

In the following, correction step positions and their higher derivatives are calculated based on their values from the previous time step and the obtained difference in acceleration as

$$\begin{aligned} x(t + \Delta t) &= x(t + \Delta t, p) + k_1 * \Delta a * \Delta t^2 \\ v(t + \Delta t) &= v(t + \Delta t, p) + k_2 * \Delta a * \Delta t \\ a(t + \Delta t) &= a(t + \Delta t, p) + k_3 * \Delta a \\ b(t + \Delta t) &= b(t + \Delta t, p) + k_4 * \frac{\Delta a}{\Delta t} \\ c(t + \Delta t) &= c(t + \Delta t, p) + k_5 * \frac{\Delta a}{\Delta t^2} \end{aligned} \quad (21)$$

Gear's scheme parameters k1-k5

for GPC3: $k_1 = 1/12, k_2 = 5/12, k_3 = 1, k_4 = 1, k_5 = 0$

for GPC4: $k_1 = 19/240, k_2 = 3/8, k_3 = 1, k_4 = 3/2, k_5 = 1$

Results

List of literature

- [1] Allan Rasmusson, Jesper Mosegaard, and Thomas Sangild Sørensen. “Exploring parallel algorithms for volumetric Mass-Spring-Damper models in CUDA”. In: *Lecture Notes in Computer Science (including subseries Lecture Notes in Artificial Intelligence and Lecture Notes in Bioinformatics)*. 2008. ISBN: 3540705201. DOI: 10.1007/978-3-540-70521-5_6.
- [2] Ronan Mendonça Amorim et al. “An electro-mechanical cardiac simulator based on cellular automata and mass-spring models”. In: *Lecture Notes in Computer Science (including subseries Lecture Notes in Artificial Intelligence and Lecture Notes in Bioinformatics)*. 2012. ISBN: 9783642333491. DOI: 10.1007/978-3-642-33350-7-45.