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1. Introduction

Atherosclerosis is a progressive disease characterized in particular by the accumulation of lipids and fibrous elements in artery walls. Over the past decade, scientists come to appreciate a prominent role for inflammation in atherosclerosis. Atherosclerosis is characterized by dysfunction of endothelium, vasculitis and accumulation of lipid, cholesterol and cell elements inside blood vessel wall. This process develops in...
arterial walls. Atherosclerosis develops from oxidized low-density lipoprotein molecules (LDL). When oxidized LDL evolves in plaque formations within an artery wall, a series of reactions occur to repair the damage to the artery wall caused by oxidized LDL. The body’s immune system responds to the damage to the artery wall caused by oxidized LDL by sending specialized white blood cells—macrophages (Mphs) to absorb the oxidized-LDL forming specialized foam cells. Macrophages accumulate inside arterial intima. Also Smooth Muscle Cells (SMC) accumulate in the atherosclerotic arterial intima, where they proliferate and secrete extracellular matrix to form a fibrous cap (Loscalzo & Schafer, 2003). Unfortunately, macrophages are not able to process the oxidized-LDL, and ultimately grow and rupture, depositing a larger amount of oxidized cholesterol into the artery wall. The atherosclerosis process is shown in Fig. 1.

This chapter describes a completely new computer model for plaque formation and development. The first section is devoted to the LDL model of transport from the lumen to intima and detailed three-dimensional model for inflammatory and plaque progression process. The next section describes some of the benchmark examples for 2D, 2D axisymmetric and 3D model of plaque formation and development. At the end a complex specific-patient 3D model is given. Finally the main conclusions of the work are addressed.

2. Methods

In this section a continuum based approach for plaque formation and development in three-dimension is presented. All algorithms are incorporated in program PAK-Athero from University of Kragujevac (Filipovic et al., 2010).

2.1 Governing equations for modeling of LDL transport through the arterial wall

The governing equations and numerical procedures are given. The blood flow is simulated by the three-dimensional Navier-Stokes equations, together with the continuity equation

\[-\mu \nabla^2 u_l + \rho (u_l \cdot \nabla)u_l + \nabla p_l = 0\]  \hspace{1cm} (1)

\[\nabla u_l = 0\]  \hspace{1cm} (2)

where \(u_l\) is blood velocity in the lumen, \(p_l\) is the pressure, \(\mu\) is the dynamic viscosity of the blood, and \(\rho\) is the density of the blood.

Mass transfer in the blood lumen is coupled with the blood flow and modelled by the convection-diffusion equation as follows

\[\nabla \cdot (-D_l \nabla c_l + c_l \mu_l) = 0\]  \hspace{1cm} (3)

in the fluid domain, where \(c_l\) is the solute concentration in the blood lumen, and \(D_l\) is the solute diffusivity in the lumen.

Mass transfer in the arterial wall is coupled with the transmural flow and modelled by the convection-diffusion-reaction equation as follows

\[\nabla \cdot (-D_w \nabla c_w + k c_w u_w) = r_w c_w\]  \hspace{1cm} (4)

in the wall domain, where \(c_w\) is the solute concentration in the arterial wall, \(D_w\) is the solute diffusivity in the arterial wall, \(K\) is the solute lag coefficient, and \(r_w\) is the consumption rate constant.
LDL transport in lumen of the vessel is coupled with Kedem-Katchalsky equations:

\[ J_p = L_p \left( \Delta p - \sigma_d \Delta \pi \right) \]  
\[ J_s = P \Delta c + \left( 1 - \sigma_f \right) J_o \bar{c} \]  

where \( L_p \) is the hydraulic conductivity of the endothelium, \( \Delta c \) is the solute concentration difference across the endothelium, \( \Delta p \) is the pressure drop across the endothelium, \( \Delta \pi \) is the oncotic pressure difference across the endothelium, \( \sigma_d \) is the osmotic reflection coefficient, \( \sigma_f \) is the solvent reflection coefficient, \( P \) is the solute endothelial permeability, and \( \bar{c} \) is the mean endothelial concentration.

**The basic relations for mass transport in the artery.** The metabolism of the artery wall is critically dependent upon its nutrient supply governed by transport processes within the blood. A two different mass transport processes in large arteries are addressed. One of them is the oxygen transport and the other is LDL transport. Blood flow through the arteries is usually described as motion of a fluid-type continuum, with the wall surfaces treated as impermeable (hard) boundaries. However, transport of gases (e.g. \( \text{O}_2, \text{CO}_2 \)) or macromolecules (albumin, globulin, LDL) represents a convection-diffusion physical process with permeable boundaries through which the diffusion occurs. In the analysis presented further, the assumption is that the concentration of the transported matter does not affect the blood flow (i.e. a diluted mixture is considered). The mass transport process is governed by convection-diffusion equation,

\[ \frac{\partial c}{\partial t} + \nabla \cdot (.cv) = D \left( \frac{\partial^2 c}{\partial x^2} + \frac{\partial^2 c}{\partial y^2} + \frac{\partial^2 c}{\partial z^2} \right) \]  

where \( c \) denotes the macromolecule or gas concentration; \( v_x, v_y, v_z \) are the blood velocity components in the coordinate system \( x, y, z \); and \( D \) is the diffusion coefficient, assumed constant, of the transported material.

**Boundary conditions for transport of the LDL.** A macromolecule directly responsible for the process of atherosclerosis is LDL which is well known as atherogenic molecule. It is also known that LDL can go through the endothelium at least by three different mechanisms, namely, receptor-mediated endocytosis, pinocytotic vesicular transport, and phagocytosis (Goldstein et al., 1979). The permeability coefficient of an intact arterial wall to LDL has been reported to be of the order of \( 10^{-8} \text{ [cm/s]} \) (Bratzler et al., 1977). The conversion of the mass among the LDL passing through a semipermeable wall, moving toward the vessel wall by a filtration flow and diffusing back to the mainstream at the vessel wall, is described by the relation

\[ c_w v_w - D \frac{\partial c}{\partial n} = K c_w \]  

where \( c_w \) is the surface concentration of LDL, \( v_w \) is the filtration velocity of LDL transport through the wall, \( n \) is coordinate normal to the wall, \( D \) is the diffusivity of LDL, and \( K \) is the overall mass transfer coefficient of LDL at the vessel wall. A uniform constant concentration \( C_0 \) of LDL is applied at the artery tree inlet as classical inlet boundary condition for eq. (7).
2.2 Finite element modeling of diffusion-transport equations

In the case of blood flow with mass transport we have domination of the convection terms due to the low diffusion coefficient (Kojic et al., 2008). Then it is necessary to employ special stabilizing techniques in order to obtain a stable numerical solution. The streamline upwind/Petrov-Galerkin stabilizing technique (SUPG) (Brooks & Hughes, 1982) within a standard numerical integration scheme is implemented. The incremental-iterative form of finite element equations of balance are obtained by including the diffusion equations and transforming them into incremental form. The final equations are

\[
\begin{bmatrix}
\frac{1}{\Delta t} \mathbf{M}_v + n^{+1} \mathbf{K}_c^{(i-1)} & n^{+1} \mathbf{F}_v^{(i-1)} + n^{+1} \mathbf{J}_v^{(i-1)} & 0 \\
0 & 0 & 0 \\
0 & 0 & \frac{1}{\Delta t} \mathbf{M}_c + n^{+1} \mathbf{K}_c^{(i-1)} + n^{+1} \mathbf{J}_c^{(i-1)}
\end{bmatrix} \times \begin{bmatrix}
\Delta \mathbf{V}^{(i)} \\
\Delta \mathbf{P}^{(i)} \\
\Delta \mathbf{C}^{(i)}
\end{bmatrix} = \begin{bmatrix}
\mathbf{N}^{+1}_K \mathbf{N}_j dV \\
\mathbf{N}^{+1}_K \mathbf{N}_j dV \\
\mathbf{N}^{+1}_K \mathbf{N}_j dV
\end{bmatrix}
\]

where the matrices are

\[
\begin{align*}
(M_v)_{ij} &= \int_V \rho N_K N_j dV, \\
(M_c)_{ij} &= \int_V N_K N_j dV \\
(n^{+1} K_c^{(i-1)})_{ij} &= \int_V D N_K N_j dV, \\
(n^{+1} K_v^{(i-1)})_{ij} &= \int_V \mu N_K N_j dV \\
(n^{+1} J_c^{(i-1)})_{ij} &= \int_V \rho N_K \nabla \cdot (\mathbf{V}^{(i-1)}) N_j dV, \\
(n^{+1} J_v^{(i-1)})_{ij} &= \int_V \rho N_K \nabla \cdot (\mathbf{V}^{(i-1)}) N_j dV \\
(n^{+1} F_c^{(i-1)})_{ij} &= \int_V \rho N_K \nabla \cdot (\mathbf{V}^{(i-1)}) N_j dV
\end{align*}
\]

and the vectors are

\[
\begin{align*}
(n^{+1} F_v^{(i-1)})_j &= n^{+1} F_j + n^{+1} F_{sc}^{(i-1)} - \frac{1}{\Delta t} M_c \left\{ n^{+1} C^{(i-1)} - n C \right\} - \\
(n^{+1} F_c^{(i-1)})_j &= \int_V N_K q^B dV \\
(n^{+1} F_{sc}^{(i-1)})_j &= \int_V D N_K V n^{+1} C^{(i-1)} \cdot \mathbf{n} dS
\end{align*}
\]

Note that \( \hat{N}_j \) are the interpolation functions for pressure (which are taken to be for one order of magnitude lower then interpolation functions \( N_j \) for velocities). The matrices \( M_c \) and \( K_c \) are the ‘mass’ and convection matrices; \( K_v \) and \( J_c \) correspond to the convective terms of equation (7); and \( \mathbf{F}_c \) is the force vector which follows from the convection-diffusion equation in (7) and linearization of the governing equations.
**2.3 Modeling of plaque formation and development**

The inflammatory process was solved using three additional reaction-diffusion partial differential equations (Calvez et al., 2008; Boynard et al., 2009):

\[
\begin{align*}
\frac{\partial_t}{t} Ox &= d_2 \Delta Ox - k_4 Ox \cdot M \\
\frac{\partial_t}{t} M + \text{div}(v_w M) &= d_1 \Delta M - k_1 Ox \cdot M + S / (1 - S) \\
\frac{\partial_t}{t} S &= d_3 \Delta S - \lambda S + k_2 Ox \cdot M + \gamma (Ox - Ox^{\text{dir}})
\end{align*}
\]  

(12)

where \( Ox \) is the oxidized LDL or \( c_w \) - the solute concentration in the wall from eq. (7); \( M \) and \( S \) are concentrations in the intima of macrophages and cytokines, respectively; \( d_1,d_2,d_3 \) are the corresponding diffusion coefficients; \( \lambda \) and \( \gamma \) are degradation and LDL oxidized detection coefficients; and \( v_w \) is the inflammatory velocity of plaque growth, which satisfies Darcy’s law and continuity equation (Kojic et al., 2008; Filipovic et al., 2004, 2006a, 2006b):

\[
\begin{align*}
v_w - \nabla \cdot (p_w) &= 0 \\
\nabla v_w &= 0
\end{align*}
\]  

(13, 14)

in the wall domain. Here, \( p_w \) is the pressure in the arterial wall.

**3. Results**

**3.1 2D model of plaque formation and development**

For the first example of a two-dimensional model of the mild stenosis, a fully developed parabolic steady velocity profile was assumed at the lumen inlet boundary

\[
u(r) = 2U_0 \left( 1 - \left( \frac{2r}{D} \right)^2 \right) \]

(15)

where \( u(r) \) is the velocity in the axial direction at radial position \( r \); \( D \) is the inlet diameter; and \( U_0 \) is the mean inlet velocity. At the lumen side of the endothelial boundary, a lumen-to-wall transmural velocity in the normal direction was specified:

\[
t^l \cdot u_l = 0, \quad u_l n_l = f_r
\]

(16)

where \( t^l \) and \( n_l \) are the tangential and normal unit vectors of fluid subdomain, respectively. Oxidized LDL distribution for a mild stenosis is shown in Fig. 2.

**3.2 2D axi-symmetric model of plaque formation and development**

The plaque formation and development is modeled through an initial straight artery in 2D axi-symmetric model with mild constriction of 30%. The inlet artery diameter \( d_0=0.4 \ [cm] \). Blood was modeled as a Newtonian fluid with density \( \rho = 1.0 \ [g/cm^3] \) and viscosity \( \mu = 0.0334 \ [P] \). The steady state conditions for fluid flow and mass transport are assumed. The entering blood velocity is defined by the Reynolds number \( Re \) (calculated using the mean blood velocity and the artery diameter).
Fig. 2. Oxidized LDL distribution for a mild stenosis (30% constriction by area)

Fig. 3. a) Velocity distribution for an initial mild stenosis 30% constriction by area
b) Velocity distribution at the end of stenosis process after $10^7$ sec [units m/s]
Fig. 4. a) Pressure distribution for an initial mild stenosis 30% constriction by area
b) Pressure distribution at the end of stenosis process after $10^7$ sec [units Pa]

Fig. 5. a) Shear stress distribution for an initial mild stenosis 30% constriction by area
b) Shear stress distribution at the end of stenosis process after $10^7$ sec [units dyn/cm$^2$]
Fig. 6. a) Lumen LDL distribution for an initial mild stenosis 30% constriction by area
b) Lumen LDL distribution at the end of stenosis process after $10^7$ sec [units mg/mL]
Fig. 7. a) Oxidized LDL distribution in the intima for an initial mild stenosis 30% constriction by area b) Oxidized LDL distribution in the intima at the end of stenosis process after $10^7$ sec [units mg/mL]
Fig. 8. a) Intima wall pressure distribution for an initial mild stenosis 30% constriction by area b) Intima wall pressure distribution at the end of stenosis process after 10^7 sec[units Pa]
Fig. 9. a) Macrophages distribution in the intima for an initial mild stenosis 30% constriction by area b) Macrophages distribution in the intima at the end of stenosis process after $10^7$ sec [units mg/mL]
3.3 3D model of plaque formation and development
In order to make benchmark example for three-dimensional simulation we tested simple middle stenosis with initial 30% constriction for time period of $t=10^7$ sec (approximately 7 years) and compare results with 2D axi-symmetric model. The results for velocity distribution for initial and end stage of simulations are presented in Fig. 11a and Fig. 11b.
The pressure and shear stress distributions for start and end time are given in Fig. 12 and Fig. 13. Concentration distribution of LDL inside the lumen domain and oxidized LDL inside the intima are presented in Fig. 14 and Fig. 15. The transmural wall pressure is presented in Fig. 16. Macrophages and cytokines distributions are shown in Fig. 17 and Fig. 18. The diagram of three-dimensional plaque volume growing during time is given in Fig. 19. It can be seen that time period for developing of stenosis corresponds to data available in the literature (Goh et al., 2010).

Fig. 11. a) Velocity distribution for an initial mild stenosis 30% constriction by area  
b) Velocity distribution at the end of stenosis process after $10^7$ sec [units m/s]
Fig. 12. a) Pressure distribution for an initial mild stenosis 30% constriction by area
b) Pressure distribution at the end of stenosis process after $10^7$ sec [units Pa]
Fig. 13. a) Shear stress distribution for an initial mild stenosis 30% constriction by area
b) Shear stress distribution at the end of stenosis process after $10^7$ sec[units/dyn/cm$^2$]
Fig. 14. a) Lumen LDL distribution for an initial mild stenosis 30% constriction by area
b) Lumen LDL distribution at the end of stenosis process after $10^7$ sec[units mg/mL]
Fig. 15. a) Oxidized LDL distribution in the intima for an initial mild stenosis 30% constriction by area b) Oxidized LDL distribution in the intima at the end of stenosis process after $10^7$ sec [units mg/mL]
Fig. 16. a) Intima wall pressure distribution for an initial mild stenosis 30% constriction by area b) Intima wall pressure distribution at the end of stenosis process after $10^7$ sec [units Pa]
Fig. 17. a) Macrophages distribution in the intima for an initial mild stenosis 30\% constriction by area b) Macrophages distribution in the intima at the end of stenosis process after $10^7$ sec[units mg/mL]
Fig. 18. a) Cytokines distribution in the intima for an initial mild stenosis 30% constriction by area b) Cytokines distribution in the intima at the end of stenosis process after $10^7$ sec [units mg/mL]
Fig. 19. Plaque progression during time (computer simulation)

From above figures it can be observed that during time plaque is progressing and all the variables as velocity distribution, shear stress, macrophages, cytokines are increasing. Also from Fig. 19 it can be seen that plaque progression in volume during time corresponds to clinical findings (Verstraete et al., 1998).

The last example is a model of the patient specific Left Anterior Descending (LAD) coronary artery for steady flow conditions. Computed concentration of LDL indicates that there is a newly formed matter in the intima, especially in the flow separation region in the LAD artery (Fig. 20).
Fig. 20. LDL concentration distribution in the left anterior descending coronary artery

4. Discussion and conclusions

Full three-dimensional model was created for plaque formation and development, coupled with blood flow and LDL concentration in blood. The models for plaque initiation and plaque progression are developed. These two models are based on partial differential equations with space and times variables and they describe the biomolecular process that takes place in the intima during the initiation and the progression of the plaque. The model for plaque formation and plaque progression despite some difficulties concerning the different time scales that are involved and the different blood velocities in the lumen and in the intima, its numerical treatment is developed by using decomposition techniques together with finite elements methods and by splitting the numerical scheme into three independent parts: blood flow and LDL transfer, inflammatory process and atheromatous plaque evolution.
Determination of plaque location and progression in time for a specific patient shows a potential benefit for future prediction of this vascular disease using computer simulation. The understanding and the prediction of the evolution of atherosclerotic plaques either into vulnerable plaques or into stable plaques are major tasks for the medical community.

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