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Upscaling mass transfer in brain capillary networks

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Hybrid continuous/network models for large scale simulation

Simulating Mass Transfer in the brain: a multi-scale problem

- Multi-scale problem (arterio-venous, capillary, parenchyma)
- Continuous representation of Parenchyma (Nicholson 2001, Kojic 2017, Holter 2017)
- Still need to explicitely resolve the vessels
- Upscale the capillary bed, including vessels and parenchyma
- Coupling with explicit arterio-venous tree is different for the capillaries and parenchyma. Two concentrations necessary.

¹ Homogenization - Equations

² Fictitious Domain Framework

Mass transfer model at capillary scale

Hypotheses

- Parenchyma and endothelium are purely diffusive domains
- \bullet Parenchyma + endothelium = tissue, $\Omega_{p} \cup \Omega_{e} = \Omega_{t}$
- \bullet D_t spacially varying (small in Ω_e)
- Red blood cells are neglected

Two-phase capillary bed model

$$
\partial_{\tau}c_{\beta} + \mathbf{u} \cdot \nabla c_{\beta} - D_{v}Δc_{\beta} = 0 \text{ in } Ω_{\beta}
$$
\n
$$
\partial_{\tau}c_{t} - D_{t}Δc_{t} = 0 \text{ in } Ω_{t}
$$
\n
$$
c_{\beta} = c_{t} \text{ on } ∂Ω_{\beta t}
$$
\n
$$
D_{v}∇c_{\beta} ⋅ \mathbf{n} = D_{t}∇c_{t} ⋅ \mathbf{n} \text{ on } ∂Ω_{\beta t}
$$

Upscaled representation of the capillary domain

Upscaled model for the two concentrations

$$
\varepsilon_{\beta} \partial_{\tau} \langle c_{\beta} \rangle^{\beta} + \left(\varepsilon_{\beta} \langle \mathbf{u} \rangle^{\beta} - \mathbf{u}_{\beta \beta} \right) \cdot \nabla \langle c_{\beta} \rangle^{\beta} - \mathbf{u}_{\beta t} \cdot \nabla \langle c_{t} \rangle^{t} + \tau_{m} (\langle c_{\beta} \rangle^{\beta} - \langle c_{t} \rangle^{t})
$$

= $\nabla \cdot (\mathbf{K}_{\beta \beta} \nabla \langle c_{\beta} \rangle^{\beta}) + \nabla \cdot (\mathbf{K}_{\beta t} \nabla \langle c_{t} \rangle^{t})$

$$
\varepsilon_t \partial_\tau \langle c_t \rangle^t - \boldsymbol{u}_{tt} \cdot \nabla \langle c_t \rangle^t - \boldsymbol{u}_{t\beta} \cdot \nabla \langle c_\beta \rangle^\beta + \tau_m (\langle c_t \rangle^t - \langle c_\beta \rangle^\beta)
$$

= $\nabla \cdot (\boldsymbol{K}_{tt} \nabla \langle c_t \rangle^t) + \nabla \cdot (\boldsymbol{K}_{t\beta} \nabla \langle c_\beta \rangle^\beta)$

- Equations solved on a domain without explicit representation of vessels
- Presence of capillary network taken into account through effective coefficients

Volume averaging

Averaging equations

$$
\langle \partial_{\tau} c_{\beta} + \mathbf{u} \cdot \nabla c_{\beta} - D_{v} \Delta c_{\beta} \rangle = 0 \text{ in } \Omega_{\beta} \langle \partial_{\tau} c_{t} - D_{t} \Delta c_{t} \rangle = 0 \text{ in } \Omega_{t}
$$

Decomposition in averaged and deviation

$$
c_{\beta} = \langle c_{\beta} \rangle^{\beta} + \tilde{c}_{\beta} \qquad \langle \tilde{c}_{\beta} \rangle = 0
$$

\n
$$
c_{t} = \langle c_{t} \rangle^{t} + \tilde{c}_{t} \qquad \langle \tilde{c}_{t} \rangle = 0
$$

- We obtain equations on $\langle c_\beta \rangle^{\beta}$ and $\langle c_t \rangle^t$
- **•** Equations depend on deviation terms \tilde{c}_β and \tilde{c}_t
- Need closure equations on \tilde{c}_β and \tilde{c}_t .
- Subtracting equations on the mean to microscopic equations leads to equations on \tilde{c}_{β} and \tilde{c}_{t} .

Deviation terms as linear combination of averaged values and their gradients

\n
$$
\widetilde{c}_{\beta} = \mathbf{b}_{\beta\beta} \cdot \nabla \left\langle c_{\beta} \right\rangle^{\beta} + \mathbf{b}_{\beta t} \cdot \nabla \left\langle c_{t} \right\rangle^{t} - s_{\beta} \left(\left\langle c_{\beta} \right\rangle^{\beta} - \left\langle c_{t} \right\rangle^{t} \right)
$$
\n
$$
\widetilde{c}_{t} = \mathbf{b}_{t\beta} \cdot \nabla \left\langle c_{\beta} \right\rangle^{\beta} + \mathbf{b}_{tt} \cdot \nabla \left\langle c_{t} \right\rangle^{t} - s_{t} \left(\left\langle c_{\beta} \right\rangle^{\beta} - \left\langle c_{t} \right\rangle^{t} \right)
$$

Volume averaging

Decomposition of deviation terms
\n
$$
\widetilde{c}_{\beta} = \mathbf{b}_{\beta\beta} \cdot \nabla \langle c_{\beta} \rangle^{\beta} + \mathbf{b}_{\beta t} \cdot \nabla \langle c_{t} \rangle^{t} - s_{\beta} \left(\langle c_{\beta} \rangle^{\beta} - \langle c_{t} \rangle^{t} \right)
$$
\n
$$
\widetilde{c}_{t} = \mathbf{b}_{t\beta} \cdot \nabla \langle c_{\beta} \rangle^{\beta} + \mathbf{b}_{tt} \cdot \nabla \langle c_{t} \rangle^{t} - s_{t} \left(\langle c_{\beta} \rangle^{\beta} - \langle c_{t} \rangle^{t} \right)
$$

Effective coefficients associated

$$
\mathbf{u}_{\beta\beta} = \langle \tilde{\mathbf{u}}s_{\beta} \rangle - \frac{D_{\beta}}{V} \int_{\partial\Omega_{\beta t}} \mathbf{n}_{\beta t} s_{\beta} dA
$$

$$
\tau_m = \frac{1}{V} \int_{\partial\Omega_{\beta t}} \mathbf{n}_{\beta t} \cdot D_{\beta} \nabla s_{\beta} dA
$$

Closure problem on s_β and s_t coefficients:

$$
-\nabla \cdot (D_{\beta} \nabla s_{\beta}) - \varepsilon_{\beta} \tau_m + \mathbf{u} \cdot \nabla s_{\beta} = 0 \text{ on } \Omega_{\beta}
$$

\n
$$
-D_t \Delta s_t - \varepsilon_t \tau_m = 0 \text{ on } \Omega_t
$$

\n
$$
\tau_m - \frac{1}{V} \int_{\partial \Omega_{\beta t}} (\mathbf{n} \cdot D_{\beta} \nabla s_{\beta}) dV = 0
$$

\n
$$
s_{\beta} - s_t = 1 \text{ on } \partial \Omega_{\beta t}
$$

\n
$$
\mathbf{n} \cdot D_{\beta} \nabla s_{\beta} - \mathbf{n} \cdot D_t \nabla s_t = 0 \text{ on } \partial \Omega_{\beta t}
$$

\n
$$
\langle s_{\beta} \rangle^{\beta} = 0
$$

\n
$$
\langle s_t \rangle^t = 0
$$

Take-home idea

Obtaining the effective coefficients necessitates to solve PDEs on a Representative Elementary Volume of the micro-structure

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Computational Framework

Solution proposed:

- \bullet C++/Parallel Finite element library ¹FEEL++
- Use a distance function field to capture the domain geometries
	- \blacktriangleright Mesh adaptation from distance function
	- \triangleright Sub-domain localization to solve Stokes problem
	- \blacktriangleright Fictitious domain to solve advection/diffusion problem

 1 Prud'homme et. al. (2016), feelpp.org

¹ Homogenization - Equations

² Fictitious Domain Framework

Network geometries

 $(300 \text{ }\mu\text{m})^3$ network. Ross (2005), Tsai (2009), Kleinfeld (2011), Kaufhold 2012, Blinder (2013)

- Segment networks extracted from biological images (center-line detection)
- Need to mesh inner and outer domains (vessels and tissue)
- Difficult to mesh automatically with CAD software (design of bifurcations)
- Geometry representation needs to be robust and automated
- Solution: use a fictitious domain method

Levelset field

Defining ϕ_i , the distance field to centerline S_i of a vessel of radius r_i The distance field to the closest vessel boundary of the network is:

$$
\phi(\boldsymbol{x}) = \min_{i=0}^{\text{Nb Vessels}} \text{dist}(\boldsymbol{x}, S_i) - r_i
$$

Properties of ϕ :

- $\phi(\mathbf{x}) < 0$ if $\mathbf{x} \in \Omega_{\beta}$
- $\phi(\mathbf{x}) > 0$ if $\mathbf{x} \in \Omega_t$
- $\phi(\mathbf{x}) = 0$ if $\mathbf{x} \in \partial \Omega_{\beta t}$

Mesh adaptation from distance function

Target mesh size field calculated from levelset field

$$
h(\boldsymbol{x}) = \begin{cases} h_{\min}, & |\phi(\boldsymbol{x})| \leq R_1, \\ \alpha |\phi(\boldsymbol{x})| + \beta, & R_1 \leq |\phi(\boldsymbol{x})| \leq R_2, \\ h_{\max}, & |\phi(\boldsymbol{x})| \geq R_2 \end{cases}
$$

Transport equations

Two domain problem

$$
\partial_{\tau}c_{\beta} + \mathbf{u} \cdot \nabla c_{\beta} - D_{v}\Delta c_{\beta} = 0 \text{ in } \Omega_{v}
$$

\n
$$
\partial_{\tau}c_{t} - D_{t}\Delta c_{t} = 0 \text{ in } \Omega_{t}
$$

\n
$$
c_{\beta} = c_{t} \text{ on } \partial\Omega_{\beta t}
$$

\n
$$
D_{\beta}\nabla c_{\beta} \cdot \mathbf{n} = D_{t}\nabla c_{t} \cdot \mathbf{n} \text{ on } \partial\Omega_{\beta t}
$$

\nPrescribed B.Cs for c_{β} on $\partial\Omega_{\beta} \setminus \partial\Omega_{\beta t}$
\nPrescribed B.Cs for c_{t} on $\partial\Omega_{t} \setminus \partial\Omega_{\beta t}$

One domain problem with spacialized coefficients
 $\partial_\tau c + \bm{u} \cdot \bm{\nabla} c - D(\phi) \Delta c = 0 \quad \text{in Ω}$

$$
\partial_{\tau}c + \mathbf{u} \ \nabla c - D(\phi)\Delta c = 0 \quad \text{in } \Omega
$$

Prescribed B.Cs for c on $\partial\Omega$

- \bullet Velocity \boldsymbol{u} is obtained by solving Stokes equation on the vessel domain only.
- Diffusion coefficient spatialized to take into account the BBB.

PDEs solved by our framework

The frameworks of fictitious network allows to solve:

The microscopic problem (DNS)

Closure equations to get the effective coefficients

Upscaled model

- Monolitic solver used for the upscaled model
- Mesh is a simple box

¹ Homogenization - Equations

² Fictitious Domain Framework

Test Case, order 3 network.

Representative Elementary Volume, size $(2 \times 2 \times 2)$

Comparison DNS - Homogenized model

Direct Numerical Simulation (DNS)

Darcy Scale

DNS

- Number of elements (tetrahedrons): 8.5×10^{6}
- Computational Cost: 1h on 100 cores

Darcy Scale

- Number of elements (tetrahedrons): 3×10^4
- Computational Cost: 10 min on 20 cores

 $\frac{2}{9}$

Quantitative comparison

Adimentional coefficients

Péclet number $P_e^{\text{REV}} = \frac{\langle u \rangle_z^{\beta} L^{\text{REV}}}$ D_v

• Membrane Permeability
$$
K_m^* = \frac{D_e}{D_v} \frac{r}{e}
$$

Quantitative comparison

Péclet number $P_e^{\text{REV}} = \frac{\langle u \rangle_z^{\beta} L^{\text{REV}}}{D}$ D_v

• Membrane Permeability
$$
K_m^* = \frac{D_e}{D_v} \frac{r}{e}
$$

Quantitative comparison

Conclusion and perspective

Conclusion

- Two-phase representation of the capillaries and tissue
- Sucessful upscaling of the capillary bed
- Framework using fictitious domain for DNS and Closure equations

In the future...

- Coupling with arterio-venous tree (1D-3D) or (3D-3D)
- Include the RBCs
- Include non linear consumption in tissue (Michaelis-Menten like)

Thank you!

Use of the distance function

Distance function used to:

define a sub-domain:

$$
\Omega_{v_h} \subset \Omega_h, \ \Omega_{v_h} = \bigcup_i \left(\Omega_{e_i} \mid \pi_{\Omega_{e_i}}^{P_{dh}^0}(\phi) < 0 \right)
$$

An example of projection function on a elements is given by:

$$
\pi_{\Omega_{e_i}}^{P_{dh}^0} = \left\{ \begin{array}{ll} -1, & \text{if } \phi < 0 \text{ on at least one dof of } \Omega_{e_i} \\ 1 & \text{else} \end{array} \right.
$$

define continuously quantities having a different value in the domains thanks to a smoothed Heaviside function (level set method):

$$
H_{\varepsilon}(\phi) = \begin{cases} 0, & \phi \leq -\varepsilon, \\ \text{smooth variations}, & -\varepsilon \leq \phi \leq \varepsilon, \\ 1, & \phi \geq \varepsilon \end{cases}
$$

Adimentional coefficients

Péclet number $P_e^{\text{REV}} = \frac{\langle u \rangle_z^{\beta} L^{\text{REV}}}$ D_v

• Membrane Permeability
$$
K_m^* = \frac{D_e}{D_v} \frac{r}{e}
$$

Adimentional coefficients

Péclet number $P_e^{\text{REV}} = \frac{\langle u \rangle_z^{\beta} L^{\text{REV}}}$ D_v

• Membrane Permeability
$$
K_m^* = \frac{D_e}{D_v} \frac{r}{e}
$$

 $P_e^{\text{REV}} = 1, K_m = 3 \times 10^{-3}$

Adimentional coefficients

Péclet number $P_e^{\text{REV}} = \frac{\langle u \rangle_z^{\beta} L^{\text{REV}}}$ D_v

• Membrane Permeability
$$
K_m^* = \frac{D_e}{D_v} \frac{r}{e}
$$

Adimentional coefficients

Péclet number $P_e^{\text{REV}} = \frac{\langle u \rangle_z^{\beta} L^{\text{REV}}}{D}$ D_v

• Membrane Permeability
$$
K_m^* = \frac{D_e}{D_v} \frac{r}{e}
$$