

Sub-voxel perfusion modeling in terms of coupled $3d-1d$ problem

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Abstract. We study perfusion by a multiscale model coupling diffusion in the tissue and diffusion along the one-dimensional segments representing the vasculature. For the model equations a block-diagonal preconditioner is proposed and its robustness is demonstrated by numerical experiments. We compare our model to a macroscale model by [P. Tofts, Modelling in DCE MRI, 2012].

1 Introduction

The micro-circulation is altered in diseases such as cancer and Alzheimer's disease as demonstrated with modern perfusion MRI. In cancer, the so-called enhanced permeability and retention (EPR) effect describes the fact that the smaller vessels in a tumor are leaky, highly permeable vessels that enable the tumor cells to grow quicker than normal cells.

In Alzheimer's disease (AD), the opposite is alleged to happen. According to [7], hypoperfusion is a precursor to AD, and the cause of the pathological cell-level changes occurring in AD. This could also explain why various kinds of heart disease are risk factors for AD, as changes in the blood pressure would affect the perfusion of the brain.

The vasculature (e.g. idealized as a system of pipes) and the surrounding tissue are clearly three-dimensional. However, the fact that in many applications the radii of the vessels are negligible compared to their lengths, permits reducing their governing equations to models prescribed on (one dimensional) curves where the physical radius R enters as a parameter. In this paper we consider a coupled $3d-1d$ system

$$\begin{aligned} \frac{\partial u}{\partial t} &= D_\Omega \Delta_\Omega u + \delta_\Gamma \beta \lambda && \text{in } \Omega, \\ \frac{\partial \hat{u}}{\partial t} &= D_\Gamma \Delta_\Gamma \hat{u} - \beta \lambda && \text{in } \Gamma, \\ \lambda &= \beta(\Pi u - \hat{u}) && \text{in } \Gamma. \end{aligned} \tag{1}$$

Here u , \hat{u} are the concentrations in the tissue domain Ω and the one-dimensional vasculature representation Γ while D_Ω , D_Γ are conductivities on the respected domains. The last equation is then a generalized Starling's law; relating \hat{u} and the value Πu of u averaged over the idealized cylindrical

vessel surface centered around Γ . The equation thus represents a coupling between the domains.

Variants of the system (1) have been used to study coupling between tissue and vasculature flow in numerous applications. In [6] a steady state limit of the system is used to numerically investigate oxygen supply of the skeletal muscles. Finite differences were used for the discretization. Using the method of Green's functions [16], [17] and [2] studied oxygen transport in the brain and tumors. Transport of oxygen inside the brain was also investigated by [11] using the finite volume method (FVM) and [5] using the finite element method (FEM) for the $3d$ diffusion problem and FVM elsewhere. In these studies the $1d$ problem was transient. More recently, coupled models discretized entirely by FEM were applied to study cancer therapies, see e.g. [14] and references therein. Mathematical foundations of these works are rooted in the seminal contributions of [4], [3] where well-posedness of the following problem is analyzed

$$\begin{aligned} -\Delta_{\Omega}u + (\hat{u} - \Pi u)\delta_{\Gamma} &= f & \text{in } \Omega \\ -\Delta_{\Gamma}\hat{u} - (\hat{u} - \Pi u) &= g & \text{in } \Gamma. \end{aligned} \tag{2}$$

Here δ_{Γ} is Dirac measure on Γ .

Presence of the measure term in (2) requires use of non-standard spaces in the analysis. In [4] the variational formulation of the problem is shown well-posed using the weighted Sobolev spaces. In particular the solution u is sought in $H_{\alpha}^1(\Omega)$, $\alpha > 0$ while the test functions v for (the first equation in) (2) are taken from $H_{-\alpha}^1(\Omega)$. With this choice the right-hand side $\langle f, v \rangle$ as well as the trace operator $v|_{\Gamma}$ are well defined while the reduced regularity of u is sufficient to make sense of the average Πu . As shown in [3] FEM for the formulation in weighted spaces yields optimal rates if the computational mesh is gradually refined towards Γ (graded meshes).

Another approach to analysis of (2) has been recently suggested in the numerical study [1] which builds on the analysis of [8] for the elliptic problem with a 0 dimensional Dirac right-hand side. Therein, wellposedness of the problem was shown with trial spaces $W^{1,p}(\Omega)$, $p = 3 - \frac{d}{2}$ and test spaces $W^{1,q}(\Omega)$, $p^{-1} + q^{-1} = 1$ and quasi-optimal error estimates for FEM shown in the norms which excluded the R neighborhood of Γ .

In studying AD or EPR the physical parameters may vary across several orders of magnitude while small or large time steps can be desirable depending on the time scales of interest. The solution algorithm for the employed model equations is thus required to be robust with respect to these parameters as well independent of the discretization. For (the transient version of) (2) the construction of such algorithms is complicated by the non-standard spaces on the domain Ω . A potential remedy for the problem can be introduction of a Lagrange multiplier which enforces the coupling between the domains with the goal of confining the non-standard spaces to the smaller domain Γ . This idea has been used by [10] to analyze robust preconditioners

for 2d-1d coupled problems based on operators in fractional Sobolev spaces, in particular, $(-\Delta)_\Gamma^{\frac{1}{2}}$, while numerical experiments reported in [9] suggest that for suitable exponents $(-\Delta_\Gamma)^s$ defines a preconditioner for the Schur complement of a 3d-1d coupled system with a *trace* constraint¹. We note that in both cases off-the-shelf methods were used as preconditioners for the operators on Ω .

The system (1) includes an additional variable λ for the coupling constraint, cf. (2). We therefore aim to apply the techniques of [9], [10] to construct a mesh-independent preconditioner for the problem, while the ideas of operator preconditioning [13] are used to ensure robustness with respect to the physical parameters and the time-stepping.

The rest of the paper is organized as follows. Section 2 identifies the structure of the preconditioner. In §3 we discuss discretization of the proposed operator and report numerical experiments which demonstrate the robust properties. In §4 the system (1) is used to model tissue perfusion using a realistic geometry of the rat cortex. Conclusions are finally summarized in §5.

2 Preconditioner for the coupled problem

In the following we let Ω be a bounded domain in \mathbb{R}^d , $d = 2$ or 3 and Γ is a subdomain of Ω of dimension 1. By $L_2(D)$ we denote the space of square-integrable functions over D and $H^1(D)$ is the space of functions with first order derivatives in $L^2(D)$.

Discretizing (1) in time by backward-Euler discretization the problem to be solved at each temporal level is of the form

$$\mathcal{A}u = f$$

with

$$\mathcal{A} = \begin{bmatrix} I - kD_\Omega \Delta_\Omega & 0 & k\beta \Pi' \\ 0 & I - kD_\Gamma \Delta_\Gamma & k\beta I \\ k\beta \Pi & k\beta I & -k. \end{bmatrix} \quad (3)$$

and k being the time step size. Note that in order to obtain a symmetric problem the operator \mathcal{A} uses the adjoint of the *averaging* operator Π instead of the *trace*, cf. (1). The choice results in modeling error of order $\mathcal{O}(R)$.

To motivate the structure of the preconditioner let us consider a 3×3 matrix

$$A = \begin{bmatrix} 1 + \alpha_1 & 0 & \beta_1 \\ 0 & 1 + \alpha_2 & \beta_2 \\ \beta_1 & \beta_2 & -\gamma \end{bmatrix}$$

¹ Note that in (2), (1) the constraint is defined in terms of a surface averaging operator Π .

where $\alpha_1, \alpha_2, \beta_1, \beta_2$, and γ are assumed to be positive. It can then be shown that with

$$B = \begin{bmatrix} (1 + \alpha_1)^{-1} & 0 & 0 \\ 0 & (1 + \alpha_2)^{-1} & 0 \\ 0 & 0 & (\gamma + \beta_1^2 + \beta_2^2)^{-1} + (\gamma + \beta_1^2/\alpha_1 + \beta_2^2/\alpha_2)^{-1} \end{bmatrix}$$

the condition number of BA is bounded independent of the parameters. With this in mind we propose that \mathcal{A} can be preconditioned by a block-diagonal operator

$$\mathcal{B} = \begin{bmatrix} (I - kD_\Omega \Delta_\Omega)^{-1} & 0 & 0 \\ 0 & (I - kD_\Gamma \Delta_\Gamma)^{-1} & 0 \\ 0 & 0 & S \end{bmatrix}, \quad (4)$$

where

$$S = S_1^{-1} + S_2^{-1} \quad (5)$$

and

$$\begin{aligned} S_1 &= \gamma I + (k\beta)^2 \Pi \Pi' + (k\beta)^2 I, \\ S_2 &= \gamma I + (k\beta)^2 \Pi (-kD_\Omega \Delta_\Omega)^{-1} \Pi' + (k\beta)^2 (-kD_\Gamma \Delta_\Gamma)^{-1}. \end{aligned} \quad (6)$$

Operator \mathcal{B} could be rigorously derived within operator preconditioning [13] as a Riesz map preconditioner for \mathcal{A} viewed as an isomorphism from $V(\Omega) \times \hat{V}(\Gamma) \times Q(\Gamma)$ to its dual space. In [10] the framework was applied to a system of two elliptic problems coupled by a $2d-1d$ constraint. For (1) extension of the analysis to parabolic problems would be required. In [12] robust preconditioners for time-dependent Stokes problem were analyzed as operators between sums of (parameter) weighted Sobolev spaces. Similarly, the structure of (5) suggests that $Q = Q_1 + Q_2$ with Q_1, Q_2 being suitable interpolation spaces. However, here we shall not justify Q_1 and Q_2 (and the preconditioner \mathcal{B}) theoretically. Instead, Q_1, Q_2 are characterized and robustness of \mathcal{B} is demonstrated by numerical experiments.

3 Discrete preconditioner

Considering (4) both $(I - kD_\Omega \Delta_\Omega)^{-1}$ and $(I - kD_\Gamma \Delta_\Gamma)^{-1}$ can be realized with off-the-shelf multilevel algorithms and the crucial question is thus how to construct S efficiently. It follows from (6) that the discrete preconditioner can be realized if finite dimensional operators spectrally equivalent to $\Pi(-\Delta_\Omega)^{-1}\Pi'$ and $\Pi\Pi'$ are known.

3.1 Auxiliary operators

If $\Omega \subset \mathbb{R}^2$ and Π is understood as the trace operator the mapping properties of trace as a bounded surjective operator $H^1(\Omega) \rightarrow H^{\frac{1}{2}}(\Gamma)$ can be used to show that $\Pi(-\Delta_\Omega)^{-1}\Pi'$ is spectrally equivalent with $(\Delta_\Gamma)^{-\frac{1}{2}}$. At the

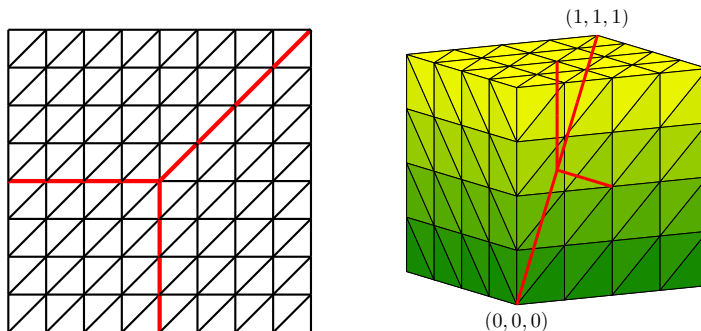


Fig. 1. Geometries used in preconditioning numerical experiments. The domain is $\Omega = [0, 1]^d$ while in order to prevent symmetries Γ (pictured in red) always features a branching point. Triangulation of Γ is made up of edges of the cells that triangulate Ω .

same time $\Pi\Pi' = \Pi(-\Delta)_\Omega^0\Pi'$ requires characterizing the space of traces of functions in $L^2(\Omega)$. We shall demonstrate by a numerical experiment that a spectrally equivalent operator is here provided by an operator $h^{-1}I_h$.

Let Ω_h, Γ_h be triangulations of Ω and Γ such that Γ_h consists of a subset of edges of the elements Ω_h , cf. Figure 1. Further let V_h, Q_h be finite element spaces of continuous linear Lagrange elements on Ω_h and Γ_h respectively. Finally we consider the eigenvalue problem: Find $\lambda \in \mathbb{R}$, $u \in V_h$, $p \in Q_h$ such that

$$\begin{aligned} \int_{\Omega} uv + \int_{\Gamma} p\Pi v &= \lambda \int_{\Omega} uv & \forall v \in V_h, \\ \int_{\Gamma} q\Pi u &= \lambda \int_{\Gamma} h^{-1}pq & \forall q \in Q_h. \end{aligned} \quad (7)$$

Table 1 shows the spectral condition number $\kappa = \max |\lambda| / \min |\lambda|$ of linear systems (7). For all the considered resolutions h the value of κ is bounded.

As the mapping properties of $\Pi\Pi'$ and $\Pi(-\Delta_\Omega)^{-1}\Pi'$ in case $\Omega \subset \mathbb{R}^3$ are not trivially obtained from the continuous analysis we again resort to finding the suitable approximations by numerical experiments. Similar to the two dimensional case the first operator with Π having the constant radius $R = 0.02$ is found to be spectrally equivalent with $h^{-1}I_h$. Following [9] an approximation to $\Pi(-\Delta_\Omega)^{-1}\Pi'$ is searched for as a suitable power $s < 0$ of $(-\Delta_\Gamma + I_\Gamma)$. More precisely, we look for the exponent yielding the most h -stable condition number of the eigenvalue problem: Find $\lambda \in \mathbb{R}$, $u \in V_h$,

	$\Omega \subset \mathbb{R}^2$						$\Omega \subset \mathbb{R}^3$					
	1/h						1/h					
	32	64	128	256	512	1024	4	8	16	32	64	128
eq. (7)	4.36	4.33	4.35	4.36	4.36	4.35	4.26	5.22	5.58	4.44	4.85	4.85
eq. (8)	8.80	8.84	8.84	8.86	8.88	8.87	11.04	8.25	7.44	9.02	10.30	11.25

Table 1. Spectral condition numbers of the eigenvalue problems related to approximations of $\Pi\Pi'$ (eq (7)) and $\Pi(-\Delta_\Omega)\Pi'$ (eq (8)). In the two-dimensional case (8) uses $s = -\frac{1}{2}$ in agreement with the mapping properties of the continuous trace operator. Results for $s = -0.55$ are reported in the three dimensional case. On the finest triangulation $\dim V_h \sim 10^6$ and $\dim Q_h \sim 10^3$ when $d = 2$ and $\dim Q_h \sim 10^2$ for $d = 3$.

$p \in Q_h$ such that

$$\begin{aligned} \int_{\Omega} \nabla u \cdot \nabla v + \int_{\Gamma} p \Pi v &= \lambda \int_{\Omega} \nabla u \nabla v + uv & \forall v \in V_h, \\ \int_{\Gamma} q \Pi u &= \lambda \int_{\Gamma} p (-\Delta + I)^s q & \forall q \in Q_h. \end{aligned} \quad (8)$$

In (8) the powers are computed using the spectral decomposition of the operator $-\Delta + I$.

We shall not present here the results for the entire optimization problem and only report on the optimum which is found to be $s = -0.55$. For this value Table 1 shows the history of the condition numbers of system (8) using again $R = 0.02$. There is a slight growth by a constant increment on the finer meshes, however, the final condition number is comparable with that obtained on the coarsest mesh. We note that we have not investigated the behaviour of the exponent on different curves or with variable radius.

3.2 Discrete preconditioner for the coupled problem

Applying the proposed preconditioner (4) of the coupled $3d-1d$ problem (3) requires evaluating the inverse of operators S_1 and S_2 in (6). The former is readily computed since, due to the suggested equivalence of $\Pi\Pi'$ and $h^{-1}I_h$, the matrix representation of S_1 is essentially a rescaled mass matrix. For S_2 we show that if $(-\Delta + I)^s$ is computed from the spectral decomposition then the inverse S_2^{-1} can be computed in a closed form.

Let A, M be the $n \times n$ matrix representations of Galerkin approximations of $-\Delta + I$ and I in the space $Q_h \subset H^1(\Gamma)$. Following [10] the matrix representation of $(-\Delta + I)_h^s$ is $H_s = MU\Lambda^s(MU)'$ where matrices U, Λ solve the generalized eigenvalue problem $AU = MU\Lambda$ such that $U'MU = I$. Using H_s it is easily established that the matrix representation of S_2 is

$$\gamma H_0 + (k\beta)^2 H_{-\frac{1}{2}} + (k\beta)^2 k^{-1} H_{-1}.$$

As $H_s^{-1} = UA^{-s}U'$ the inverse of the S_2 matrix is given by

$$U \left(\gamma A^0 + (k\beta)^2 A^{-\frac{1}{2}} + (k\beta)^2 k^{-1} A^{-1} \right) U'. \quad (9)$$

Using the spectral decomposition the cost of setting up the preconditioner S is determined by the cost of solving the generalized eigenvalue problem for U and A . This practically limits the construction to systems where $\dim Q_h \sim 10^3$. However, for the problems considered further this limitation does not present an issue. In particular, the preconditioner can be setup on the discretization of vasculature of the cortex tissue used in §4 which contains approximately twenty thousand vertices.

With a discrete approximation of S we can finally address the question of \mathcal{B} being a robust preconditioner for (3). Motivated by (1) we do not vary all the parameters, and instead, set $\gamma = 1$ in \mathcal{A} . Moreover, the conductivity on Ω is taken as unity and only variable $D_\Gamma > 1$ is considered mimicking the expected faster propagation along the one-dimensional domain. Finally, the time step k and the coupling constant β are taken between 10^{-4} – 10^{-8} .

For a fixed choice of parameters D_Γ , β , k the preconditioned problem $\mathcal{B}\mathcal{A}x = \mathcal{B}f$ is considered on geometries from Figure 1 and discretized with continuous linear Lagrange elements. The resulting linear system is then solved with the MinRes method where the iterations are stopped once the preconditioned residual norms is less than 10^{-10} .

The observed iteration counts are reported in Table 2. For both 2d-1d and 3d-1d coupled problems the iterations can be seen to be bounded in the discretization parameter. Moreover the preconditioner performs almost uniformly in the considered ranges of D_Γ and β while there is a clear boundness in k as well. We note that these conclusions are not significantly altered if the ranges for k and β are extended to 1.

Having demonstrated the numerical stability of our model, we next test it on the same problem as [18], namely a bloodborne tracer perfusing and later being cleared from tissue. While [18] considers this problem on a macroscopic scale, we model it on the micro-scale, where individual blood vessels can be resolved as part of our 1d domain.

4 Perfusion experiment

In [15], a $0.7\text{mm} \times 0.7\text{mm} \times 0.7\text{mm}$ piece of mouse brain microvasculature was imaged using two-photon microscopy. To obtain a realistic geometry for our model, we used this data to generate a 3d mesh of the extravascular space in which vessel segments corresponded to 1d mesh edges.

In our model, this corresponds to initial conditions of $u, \hat{u} = 0$, followed by a boundary condition of $\hat{u} = 1$ on the part of the boundary corresponding to inlet vessels. To model clearance, the inlet boundary condition was swapped from $\hat{u} = 1$ to $\hat{u} = 0$ after a third of the simulation time had passed.

D_r	β	k	$2d-1d$						$3d-1d$					
			$1/h$						$1/h$					
			32	64	128	256	512	1024	4	8	16	32	64	128
10^0	10^{-8}	10^{-8}	11	10	10	8	7	7	8	13	12	11	11	10
		10^{-6}	15	13	12	10	8	8	10	14	16	16	15	13
		10^{-4}	16	11	12	13	13	14	12	16	18	16	13	14
	10^{-6}	10^{-8}	11	10	10	8	7	7	8	13	12	11	11	10
		10^{-6}	15	13	12	10	8	8	10	14	16	16	15	13
		10^{-4}	15	11	12	13	13	14	12	16	18	16	13	14
	10^{-4}	10^{-8}	11	10	10	8	7	7	8	13	12	11	11	10
		10^{-6}	15	13	12	10	8	8	10	14	16	16	15	13
		10^{-4}	16	11	12	13	13	14	12	16	18	16	13	14
10^2	10^{-8}	10^{-8}	11	10	10	9	7	7	8	13	12	11	11	10
		10^{-6}	15	13	12	10	8	8	10	14	16	16	15	13
		10^{-4}	15	11	12	13	13	13	11	16	18	13	13	14
	10^{-6}	10^{-8}	11	10	10	9	7	7	8	13	12	11	11	10
		10^{-6}	15	13	12	10	8	8	10	14	16	16	15	13
		10^{-4}	15	11	12	13	13	13	11	16	18	13	13	14
	10^{-4}	10^{-8}	11	10	10	9	7	7	8	13	12	11	11	10
		10^{-6}	15	13	12	10	8	8	10	14	16	16	15	13
		10^{-4}	15	11	12	13	13	13	12	16	18	13	13	14
10^4	10^{-8}	10^{-8}	11	10	10	8	7	7	8	13	12	11	11	10
		10^{-6}	15	13	12	10	8	8	10	14	16	16	15	13
		10^{-4}	15	11	12	13	12	10	11	16	18	13	13	14
	10^{-6}	10^{-8}	11	10	10	8	7	7	8	13	12	11	11	10
		10^{-6}	15	13	12	10	8	8	10	14	16	16	15	13
		10^{-4}	15	11	12	13	12	10	11	16	18	13	13	14
	10^{-4}	10^{-8}	11	10	10	8	7	7	8	13	12	11	11	10
		10^{-6}	15	13	12	10	8	8	10	14	16	16	15	13
		10^{-4}	15	11	12	13	12	10	11	16	18	13	13	14
10^6	10^{-8}	10^{-8}	11	10	10	8	7	7	8	13	12	11	11	10
		10^{-6}	15	13	12	10	8	8	10	14	16	16	15	13
		10^{-4}	14	10	10	9	8	5	11	15	18	13	11	11
	10^{-6}	10^{-8}	11	10	10	8	7	7	8	12	12	11	11	10
		10^{-6}	15	13	12	10	8	8	10	14	16	16	15	13
		10^{-4}	14	10	10	9	8	5	11	15	18	13	10	11
	10^{-4}	10^{-8}	11	10	10	8	7	7	8	12	12	11	11	10
		10^{-6}	15	13	12	10	8	8	10	14	16	16	15	13
		10^{-4}	14	10	10	9	8	5	11	16	18	13	10	11

Table 2. Number of iterations of MinRes method on (3) using as preconditioner (4) with S approximated using (9). (Left) $2d-1d$ coupled problem and (right) $3d-1d$ coupled problem from Figure 1 are considered. Convergence criterion with residual of 10^{-10} is used.



Fig. 2. The $1d$ domain used in perfusion experiment. The $3d$ domain is a bounding box.

As parameters, we use $D_\Gamma = 6.926 \times 10^7 \mu\text{m}^2/\text{s}$, $D_\Omega = 1.87 \times 10^2 \mu\text{m}^2/\text{s}$ and $\beta = 50 \mu\text{m}/\text{s}$. We used $k = 1\text{s}$ after verifying that reducing it did not significantly affect our results. In this experiment, it was unnecessary to use the preconditioner described in §3 because the problem size was small enough that we could use a direct solver.

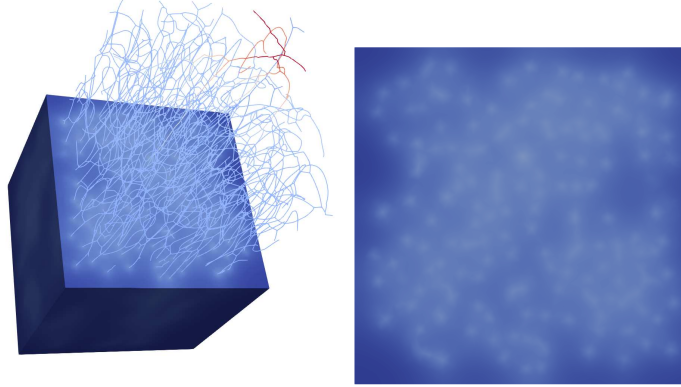


Fig. 3. Example results shown on a (left) clip of the 3d domain and (right) on a slice. Notice the 'halos' of increased concentration immediately around the vessels.

Tofts [18] assumes a relation

$$\frac{\partial C_t}{\partial t} = \frac{K}{\nu} (C_v - C_t) \quad (10)$$

between the pixel tissue concentration C_t and the pixel vessel concentration C_v for some constant K . Here, ν is the vascular volume fraction. Our geometry is of a size comparable to a single pixel in [18], so C_t and C_v correspond to the normalized averages

$$C_t = \frac{\int_\Omega u}{\int_\Omega 1} \quad \text{and} \quad C_v = \frac{\int_\Gamma \pi R^2 \hat{u}}{\int_\Gamma \pi R^2}.$$

We computed K by solving for u, \hat{u} using our model, and then computing C_t, C_v as given above, and defining K such that equation (10) holds at each time point. This makes K a function of time, with units seconds⁻¹.

4.1 Discussion of perfusion experiment

Our value for K is not entirely constant. There is a small variation in time, as perfusion seems to be somewhat faster when the extravascular space is

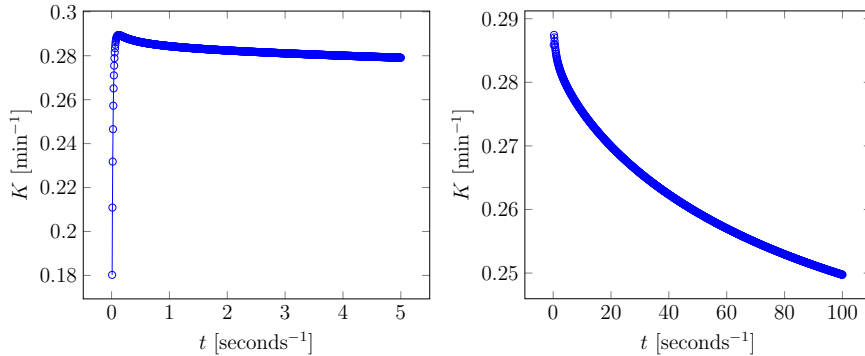


Fig. 4. Behavior of K on short and long time scales.

completely empty of the tracer. As it starts getting saturated, perfusion slows down somewhat. This translates into K decreasing by about 20% over a period of about 5 minutes, from 0.28 min^{-1} to 0.23 min^{-1} .

This is a bit lower than [18], whose numbers are between 0.46 and 1.3. There are several possible explanations for this. One might be that in our model, vascular transport is modeled as exceptionally fast diffusion for convenience, whereas in [18] (and in reality) it occurs by convection. However, in both cases $1d$ transport is very fast compared to the $3d$ transport and the $1d$ - $3d$ exchange. Further, K is defined in terms of the $1d$ - $3d$ exchange alone, so non-extreme variations in the $1d$ transport seem unlikely to be relevant.

Another possibility might be that the underlying biology are taken from human kidney tissue, whereas our vasculature is from mouse brain. Given the different physiological functions of the organs, it seems reasonable that brain vessels and kidney vessels would have different rates of blood to tissue exchange. A third reason might be our diffusion constants not exactly matching the Gadolinium tracer used in [18].

In further work, it would be interesting to incorporate convective transport into the model. One way of doing so is to introduce an unknown for the blood pressure \hat{p} on the $1d$ domain, and assume that the blood flow \hat{q} in a segment is laminar and follows Poiseuille's law

$$\hat{q} = R^4 C \nabla \hat{p}$$

Further, assuming that blood is incompressible and does not escape from the vessels yields that $\nabla \cdot \hat{q} = 0$, which using the relation above translates to a Laplace-type equation for \hat{p} alone. This then determines \hat{q} , which can be put into a convection-diffusion equation for the $1d$ tracer transport with the same $1d$ - $3d$ exchange term as before.

5 Conclusions

A coupled $3d$ - $1d$ system with an additional unknown enforcing the coupling between the domains was used as a model of tissue perfusion. For the system we proposed a robust preconditioner and demonstrated its properties through numerical experiments. Further, we have shown that the model can be applied to a physiological problem with reasonable results.

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