

## Fractional bioheat equation

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### Abstract

In this work we develop a new mathematical model for the Pennes' bioheat equation assuming a fractional time derivative of single order. A numerical method for the solution of such equations is proposed, and, the suitability of the new model for modelling real physical problems is studied and discussed.

*Key words: Time-fractional diffusion equation, Caputo derivative, bioheat equation, stability, convergence*

## 1 Introduction

Pennes' [1] bioheat transfer equation, which describes the thermal distribution in human tissue, taking into account the influence of blood flow, (see Fig. 1) is given by,

$$\rho_t c_t \frac{\partial T(x, t)}{\partial t} = k \frac{\partial^2 T(x, t)}{\partial x^2} + W_b c_b (T_a - T) + q_m, \quad t > 0, \quad 0 < x < L, \quad (1)$$

where  $\rho_t$ ,  $c_t$  are constants representing the density [ $kg/m^3$ ] and the specific heat [ $J/(kg \text{ } ^\circ C)$ ], respectively, and  $k$  is the tissue thermal conductivity [ $J/(s.m \text{ } ^\circ C)$ ];  $W_b$  is the mass flow rate

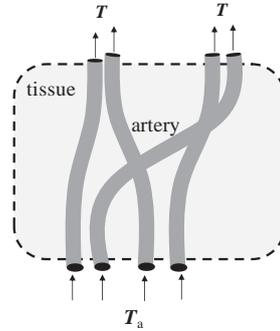


Figure 1: Heat transfer between blood vessels and tissue.

of blood per unit volume of tissue  $[kg/(s.m^3)]$ ;  $c_b$  is the blood specific heat;  $q_m$  is the metabolic heat generation per unit volume  $[J/(s.m^3)]$ ;  $T_a$  represents the temperature of arterial blood  $[^\circ C]$ ;  $T$  is the temperature and the term  $W_b c_b (T_a - T)$  represents the blood perfusion.

Although this model presents some limitations: the blood velocity field is not taken into account; assumes that thermal equilibration occurs in the capillaries; the blood leaving the tissue does not influence the temperature of the medium; it has been used by several researchers working in different research fields. The reason for that is that the model is simple, attractive and the new models proposed are complex presenting a large number of parameters that are difficult to obtain experimentally.

In order to overcome some of the model limitations, in this work we propose a new bio-heat model, where the classical time derivation is substituted by a time-fractional derivative.

The bioheat equation presented before (Eq.1) is now written, using the time-fractional derivative instead of the the first-order time derivative,  $\frac{\partial T(x,t)}{\partial t}$ , generalizing in this way the original equation derived by Harry Pennes:

$$\frac{\partial^\alpha T(x,t)}{\partial t^\alpha} = A \frac{\partial}{\partial x} \left( k(x) \frac{\partial T(x,t)}{\partial x} \right) - BT(x,t) + C \quad 0 < t < T^*, \quad 0 < x < L, \quad (2)$$

where  $\frac{\partial^\alpha}{\partial t^\alpha}$  is the fractional Caputo derivative given by [2],

$$\frac{\partial^\alpha T(x,t)}{\partial t^\alpha} = \frac{1}{\Gamma(1-\alpha)} \int_0^t (t-s)^{-\alpha} \frac{\partial T(x,s)}{\partial s} ds \quad (3)$$

with  $0 < \alpha < 1$ , and  $A = \frac{1}{\rho_t c_t \tau^{\alpha-1}}$ ,  $B = \frac{W_b c_b}{\rho_t c_t \tau^{\alpha-1}}$ , and  $C = \frac{W_b c_b T_a + q_m}{\rho_t c_t \tau^{\alpha-1}}$ . Note that  $k(x)$  is a function of  $x$ , meaning that we can deal with possible anisotropy. Also, it is worth-mentioning the fact that we have added a new parameter  $\tau$  [s] to the equation, so that it becomes dimensionally consistent.

## 2 Numerical solution

For the numerical solution of Eq. 2, we consider a uniform space mesh on the interval  $[0, L]$ , defined by the gridpoints  $x_i = i\Delta x$ ,  $i = 0, \dots, N$ , where  $\Delta x = \frac{L}{N}$ , and we approximate the space derivative by a second order finite difference.

For the discretization of the fractional derivative we also assume uniform meshes, with a time step  $\Delta t = Time/R$  ( $R$  is the number of divisions of the grid) and time gridpoints  $t^l$ ,  $l = 0, 1, \dots, R$ , and, we use the backward finite difference formula provided by Diethelm [2] ( $\mathcal{O}((\Delta t)^{2-\alpha_j})$ ). Denoting  $T(x_i, t_l)$  by  $T_i^l$ , and  $k(x_i \pm \frac{\Delta x}{2})$  by  $k_{i \pm \frac{1}{2}}$  and neglecting the  $\mathcal{O}((h)^2)$  and  $\mathcal{O}((\Delta t)^{2-\alpha_j})$  terms, the finite difference scheme is then given by,

$$\frac{(\Delta t)^{-\alpha}}{\Gamma(2-\alpha)} \sum_{m=0}^l a_{m,l}^{(\alpha)} (T_i^{l-m} - T_i^0) = A \frac{k_{i+\frac{1}{2}} T_{i+1}^l - (k_{i+\frac{1}{2}} + k_{i-\frac{1}{2}}) T_i^l + k_{i-\frac{1}{2}} T_{i-1}^l}{(\Delta x)^2} + f(x_i, t_l, T_i^l) \quad i = 2, \dots, N-2, \quad l = 1, \dots, R \tag{4}$$

with  $f(x_i, t_l, T_i^l) = -BT_i^l + C$ .

For consistency with the order of the spatial discretization at grid points  $i = 2, \dots, N-2$ , we also assume a second a order approximation for the Neumann boundary conditions. For that, a second order forward and backward finite difference scheme was used.

### Stability and Convergence

In this section we provide two useful theorems for the stability and convergence of the numerical mehtod proposed.

**Theorem (stability) 1.** *Let  $0 < \varepsilon \leq \Delta t$ , the scheme given by Eq. 4 is unconditionally stable with respect to the initial conditions.*

**Theorem (convergence) 1.** *Let  $0 < \varepsilon \leq \Delta t$ , if the solution of (2) is of class  $C^2$  with respect to  $t$  and of class  $C^4$  with respect to  $x$ , then there exists a constant  $C_0$  independent of  $\Delta x$  and  $\Delta t$  such that,*

$$\| \mathbf{e}^l \|_\varepsilon \leq C_0 \left( (\Delta t)^{2-\alpha} + (\Delta x)^2 \right), \quad l = 0, 1, \dots \tag{5}$$

where  $\mathbf{e}^l = [e_1^l, e_2^l, \dots, e_{N-1}^l]$ ,  $l = 1, 2, \dots$ , is vector of the errors at time step  $l$ , with  $e_i^l = T(x_i, t_l) - T_i^l$   $l = 1, 2, \dots, i = 1, \dots, N-1$  the error at each point of the mesh.

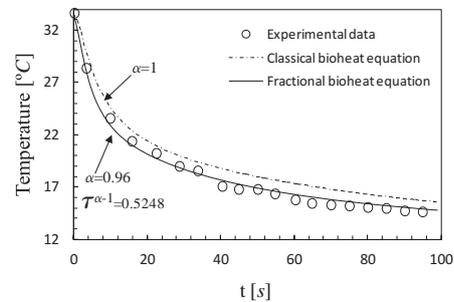


Figure 2: Fitting experimental data.

### 3 Modelling

In order to test the fractional bioheat model, we used the experimental data provided by Barcroft and Edholme [3] for the temperature variation inside a human arm. One of their experiments consisted of measuring the temperature decrease of the subcutaneous tissue (1 cm below the skin surface) when the forearm is submersed in a 12°C water bath.

In Fig. 2, we show that the proposed fractional bioheat equation can be used to improve the accuracy of the classical numerical predictions, when compared with classical models.

### 4 Conclusions

In this work we have derived a numerical method for the solution of the fractional bioheat equation of single order with a variable diffusion coefficient. We observed that the new bioheat model can be used to better predict subdiffusion processes.

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