



Simultaneous Identification of the Parameters in the Mathematical Model of Brain Tumor Growth Dynamics Under Treatment

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Abstract. This paper is devoted to an inverse problem for a nonlinear parabolic equation related to brain tumor dynamics. After reformulating the inverse problem as a minimization problem, we prove the existence and stability of the solution to the minimization problem. Based on the Fréchet differentiability of the objective (cost) functional, we develop an efficient iterative procedure for the numerical solution to the minimization problem. Numerical examples with noise-free and noisy data illustrate applicability and accuracy of the proposed method to some extent.

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Nomenclature

$g(c)$ Reaction term	T Final time
\mathcal{B} Spatial domain of the brain	ε Noise level
ν The outward normal vector on the boundary	c_{lim} Carrying capacity
φ Initial cell density	ρ Proliferation parameter
$D(x)$ Diffusion coefficient	d_w Diffusion coefficient for white matter
c Cell density at position x and time t	d_g Diffusion coefficient for gray matter
div Divergence operator	$k(x, t)$ Treatment term
∇ Gradient operator	α Spatial component of the treatment profile
$\partial\mathcal{B}$ Boundary of \mathcal{B}	β Temporal component of the treatment profile
h Mesh step in x direction	$\mathcal{A}_\alpha \times \mathcal{A}_\varphi$ Set of admissible solutions
τ Time step in t direction	$J(\alpha, \varphi)$ Objective functional

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ζ_{α}^k	Step size for α	S_{φ}^k	Search direction for φ
ζ_{φ}^k	Step size for φ	$\bar{\varepsilon}$	Tolerance parameter
S_{α}^k	Search direction for α	ω_i	Weight functions

1. Introduction

In this paper, we consider an inverse problem for the following problem:

$$\begin{cases} \partial_t c - \operatorname{div}(D(x)\nabla c) + k(x, t)c - g(c) = 0 & (x, t) \in \mathcal{B} \times (0, T), \\ D(x)\partial_{\nu}c = 0 & (x, t) \in \mathcal{B} \times (0, T), \\ c(x, 0) = \varphi(x) & x \in \bar{\mathcal{B}}, \end{cases} \quad (1)$$

where $\mathcal{B} \subset \mathbb{R}^d$ ($d = 1, 2, \dots$) be an open bounded domain with a sufficiently smooth boundary $\partial\mathcal{B}$, $T > 0$ is a final time, $c(x, t)$ is the tumor cell density at the time t and the spatial location x within the brain region \mathcal{B} , $k(x, t)c(x, t)$ is the treatment term describing the death of cells due to chemotherapy or radiation therapy, $D(x)$ is the diffusion coefficient of cells in the brain tissue modeling a random tumor cell movement as a diffusive flux proportional to the cell density gradient, $\operatorname{div}(D(x)\nabla c)$ captures random tumor cell movement, $g(c)$ is the reaction term that describes the growth dynamics of the tumor and ν is the outward unit normal. The boundary condition $D(x)\partial_{\nu}c = 0$ ensures tumor cells do not diffuse beyond the brain region. Additionally, the function $\varphi(x) \in L^2(\mathcal{B})$ represents the initial tumor cell density at $t = 0$. The problem (1) is a well-known model related to the growth of brain tumors under treatment

The model (1) serves as a fundamental tool for understanding brain tumor dynamics and optimizing treatment strategies through comprehensive mathematical analysis. For instance, the authors work the problem (1) with a source term such that the model has logistic growth and accomplish full 3-dimensional simulations of the tumor in time on two types of imaging data, the 3d Shepp-Logan head phantom image and an MRI T1-weighted brain scan from the Internet Brain Segmentation Repository in [15]. In [1], the authors study a treatment parameter identification problem in (1). They derive a nonlinear conjugate gradient method for the inverse problem and reconstruct the unknown parameter from additional information about the tumor taken at a fixed instance of time. In [26], the authors analyze the behavior of gliomas mechanistically, deriving their behavior from two fundamental properties, the net proliferation rate and the diffusion coefficient. They conclude that although these clearly determine the tumor's growth as an expansion of its traveling wave front, other factors (such as age and Karnofsky performance score) combine to determine the patient's prognosis, the survival time. The authors extend a mathematical model of gliomas based on proliferation and diffusion rates to incorporate the effects of augmented cell motility in white matter as compared to grey matter and they simulate model tumors on an anatomically accurate brain domain by using a detailed mapping of the white and grey matter in the brain developed for a MRI simulator,

in [24]. A simple spatio-temporal mathematical model, based on proliferation and diffusion, that incorporates the effects of radiotherapeutic and chemotherapeutic treatments has been studied in [19]. Moreover, they study the effects of different schedules of radiation therapy, including fractionated and hyperfractionated external beam radiotherapy, using a generalized linear quadratic model. In [23], the authors present a novel explicit triscale reaction-diffusion numerical model of glioblastoma multiforme tumor growth, they adopt a finite-difference time-domain method for the numerical solution and then a clinical scenario is addressed to demonstrate the workflow of a possible clinical validation procedure. Other researchers [20] present an extension of Swanson's reaction-diffusion model in order to include the effects of radiation therapy using the classic linear-quadratic radiobiological model for radiation efficacy, along with an investigation of response to various therapy schedules and dose distributions on a virtual tumor. Some of the recent developments in mathematical modeling are reviewed in [25]. A model of untreated gliomas is represented first followed by models of polyclonal gliomas that follows a chemotherapy or a surgical resection. Such reviewed models illustrate the evolution of mathematical models for glioma growth and invasion beginning in simple homogeneous tissue, with or without gross anatomical boundaries (skull and ventricles), extending to complex heterogeneous tissue, with varying proportions of grey and white matter in cerebral cortex (including the sulcal pattern), deep cerebral nuclei, brainstem and cerebellum. A patient-specific, biologically based mathematical model for glioma growth that quantifies response to XRT in individual patients in vivo is presented in [21]. This mathematical model uses net rates of proliferation and migration of malignant tumor cells to characterize the tumor's growth and invasion along with the linear-quadratic model for the response to radiation therapy.

In this paper, we study an inverse problem for the nonlinear parabolic problem (1). Inverse problems of finding coefficients, the initial distribution, sources or parameters are physically and practically very important. Recently, there has been a growing interest in inverse problems for nonlinear parabolic equations. For instance, the simultaneous reconstruction of the initial temperature and heat radiative coefficient in a heat conductive system is studied in [28]. They prove the stability of the inverse problem and then the reconstruction process is done by Tikhonov regularization. In [3], the identification of the space-dependent reaction coefficient, the initial temperature and the source term from measured temperatures at two instants t_1 , t_2 and at the final time T are investigated. In [4], an inverse problem of simultaneously identifying and reconstructing the space-dependent reaction coefficient and source term component from time-integral temperature measurements is investigated. After they prove the existence and uniqueness of the solution, the conjugate gradient method to find the numerical solution is developed, and its convergence is proved from the Lipschitz continuity of these gradients. A novel inverse problem of reconstructing the unknown time-dependent source term entering the fourth-order parabolic equation of thermal grooving by surface diffusion from a given integral observation is formulated in [5]. The author studies the determination of an unknown time-dependent source term in a Kuramoto-Sivashinsky equation from given additional integral-type measurement in [2]. In [22], some methods are developed for find-

ing the thermophysical parameters of a two-layer soil and a rational method of choosing the damping coefficient is also proposed, which provides an indicative rate of convergence of the approximate value of the functional to zero. Our paper can be considered as an addition to these studies.

This article is organized as follows: In the next section, we formulate and analyze the direct and the inverse problems. In Section 3, we present an iterative procedure based on the conjugate gradient method to solve the minimization problem. Some numerical examples with noise-free and noisy data are given to show the efficiency of the method in Section 4. The conclusions and possible directions on the problem are given in Section 5.

2. Formulations and analyses of the direct and the inverse problems

In this section, we define the direct and the inverse problems. Throughout this paper, we define $Q := \mathcal{B} \times (0, T)$ and $\mathcal{B}_T := \partial\mathcal{B} \times (0, T)$. The diffusion coefficient $D(x)$ is assumed to encompass two regions, white and grey matter, as in the papers [10, 24, 25], reflecting the spatial heterogeneity of brain tissue. Specifically, we define $D(x)$ as:

$$D(x) = \begin{cases} d_w, & \text{if } x \in \text{white matter,} \\ d_g, & \text{if } x \in \text{grey matter,} \end{cases}$$

where d_g and d_w denote the diffusion coefficients for grey and white matter, respectively, satisfying $0 < d_g \ll d_w$. For the growth function $g(c)$, we adopt a logistic growth model to ensure that the growth rate diminishes as the cell density approaches its maximum capacity. Thus, $g(c)$ is defined as:

$$g(c) = \rho c \left(1 - \frac{c}{c_{\text{lim}}} \right),$$

where ρ is the net proliferation rate and c_{lim} represents the carrying capacity. This formulation provides a more realistic representation of tumor growth dynamics compared to simple exponential models [7, 9, 24, 26, 27], which may result in unbounded growth over time.

Definition 1. *The weak solution $c \in L^2(0, T; H^1(\mathcal{B}))$ of the problem (1) is defined as the solution of the following problem:*

$$\int_0^T \int_{\mathcal{B}} \left(\partial_t c(x, t) \psi + D(x) \nabla c(x, t) \cdot \nabla \psi + k(x, t) c(x, t) \psi \right) dx dt = \int_0^T \int_{\mathcal{B}} g(c) \psi dx dt, \quad (2)$$

for all $\psi \in L^2(0, T; H^1(\mathcal{B}))$.

For given inputs $D(x)$, $k(x, t)$, $\varphi(x)$ and $T > 0$, the problem (1) is called the direct problem. Like most direct problems of the mathematical physics, the problem (1) is well posed, see [1, 16] for details.

Here the inverse problem consists of simultaneously determining the treatment profile $k(x, t)$ and the initial tumor cell density φ , by means of the time-integral measurements of tumor cell density (measured data) defined as follows:

$$\phi_1 := \int_0^T \omega_1(t) c(x, t) dt \quad \text{and} \quad \phi_2 := \int_0^T \omega_2(t) c(x, t) dt, \quad x \in \mathcal{B}, \quad (3)$$

where $\omega_1(t)$ and $\omega_2(t)$ denote two distinct weight functions. We note that since time-integral measurements of tumor cell density do not typically reveal time-dependent profiles, we consider the function $k(x, t) \in L^\infty(Q)$ in the following separable form:

$$k(x, t) = \alpha(x) \beta(t), \quad (4)$$

where $\alpha(x) \in L^\infty(\mathcal{B})$ represents the spatial component of the treatment profile $k(x, t)$ and describes how the treatment efficacy varies across different spatial locations within the brain region. On the other hand, $\beta(t) \in L^\infty(0, T)$ is the temporal component of the treatment profile and captures how the treatment effectiveness evolves over time. In the inverse problem considered here, our focus shifts to scenarios where the spatial component $\alpha(x)$ is unknown, while the temporal component $\beta(t)$ is known. By taking (4) into account in the problem (1), we have

$$\begin{cases} \partial_t u - \operatorname{div}(D(x)\nabla u) + \alpha^*(x) \beta(t) u - g(u) = 0 & \text{in } Q, \\ D(x) \partial_\nu u = 0 & \text{on } \mathcal{B}_T, \\ u(x, 0) = \varphi^*(x) & \text{in } \bar{\mathcal{B}}. \end{cases} \quad (5)$$

The focus of our investigation in this article revolves around the simultaneous determination of the response treatment parameter $\alpha^*(x)$ and the initial cell density $\varphi^*(x)$ from two time-integral measurements of u defined by (3).

We first define the sets of admissible solutions to analyze the direct problem (5). A set $\mathcal{A}_\alpha \times \mathcal{A}_\varphi$ defined by

$$\begin{aligned} \mathcal{A}_\alpha &:= \{\alpha(x) : \alpha \in L^\infty(\Omega), 0 < \underline{\alpha} \leq \alpha(x) \leq \bar{\alpha}, \text{ a.e } x \in \Omega\}, \\ \mathcal{A}_\varphi &:= \{\varphi(x) : \varphi \in L^\infty(\Omega), |\varphi(x)| \leq \bar{\varphi}, \text{ a.e } x \in \Omega\}, \end{aligned}$$

is called the set of admissible solutions where $\underline{\alpha}$, $\bar{\alpha}$, and $\bar{\varphi}$ are predetermined positive constants. For each pair (α, φ) belonging to $\mathcal{A}_\alpha \times \mathcal{A}_\varphi$, we denote the solution to problem (5) by $c(\alpha, \varphi)$. In real-world scenarios, observed data ϕ_1 and ϕ_2 are given with some error including measurement errors, biological variability, spatial heterogeneity and temporal dynamics inherent in tumor growth and treatment response. To overcome these effects, we introduce perturbed measurements ϕ_1^ε and ϕ_2^ε as follows:

$$\|\phi_i^\varepsilon - \phi_i\|_{L^\infty(\Omega)} \leq \varepsilon, \quad \text{for } i = 1, 2, \quad (6)$$

where $\varepsilon > 0$ represents the magnitude of noise in the data. Consequently, the quasi-solution, see [14], of the inverse problem under consideration is obtained by minimizing the following least-squares objective functional:

$$J(\boldsymbol{\alpha}, \boldsymbol{\varphi}) := \frac{1}{2} \left\| \int_0^T \omega_1(t) c(\boldsymbol{\alpha}, \boldsymbol{\varphi}) dt - \phi_1^\varepsilon \right\|_{L^2(\Omega)}^2 + \frac{1}{2} \left\| \int_0^T \omega_2(t) c(\boldsymbol{\alpha}, \boldsymbol{\varphi}) dt - \phi_2^\varepsilon \right\|_{L^2(\Omega)}^2. \quad (7)$$

Therefore, the inverse problem addressed in this article can be reformulated and modeled by the following minimization problem:

$$\begin{cases} \text{Find } (\boldsymbol{\alpha}^*, \boldsymbol{\varphi}^*) \in \mathcal{A}_\alpha \times \mathcal{A}_\varphi \text{ such that} \\ J(\boldsymbol{\alpha}^*, \boldsymbol{\varphi}^*) \leq J(\boldsymbol{\alpha}, \boldsymbol{\varphi}), \quad \forall (\boldsymbol{\alpha}, \boldsymbol{\varphi}) \in \mathcal{A}_\alpha \times \mathcal{A}_\varphi. \end{cases} \quad (8)$$

It is worth noting that in this paper, we adopt the regularization before discretization strategy, as opposed to the alternative approach of discretization before regularization discussed in [3, 11].

Theorem 1. *There exists at least one minimizer for the optimization problem (8).*

Proof. Since $J(\boldsymbol{\alpha}, \boldsymbol{\varphi})$ is nonnegative, we know that $\inf J(\boldsymbol{\alpha}, \boldsymbol{\varphi})$ is finite over $\mathcal{A}_\alpha \times \mathcal{A}_\varphi$. Thus, there exists a minimizing sequence $\{\boldsymbol{\alpha}_n, \boldsymbol{\varphi}_n\}_{n \geq 0}$ from $\mathcal{A}_\alpha \times \mathcal{A}_\varphi$ such that

$$\lim_{n \rightarrow \infty} J(\boldsymbol{\alpha}_n, \boldsymbol{\varphi}_n) = \inf_{\substack{\boldsymbol{\alpha} \in \mathcal{A}_\alpha \\ \boldsymbol{\varphi} \in \mathcal{A}_\varphi}} J(\boldsymbol{\alpha}, \boldsymbol{\varphi}).$$

This implies the boundedness of $\{\boldsymbol{\alpha}_n, \boldsymbol{\varphi}_n\}_{n \geq 0}$ in $L^\infty(\mathcal{B}) \times L^2(\mathcal{B})$ and therefore there is a subsequence (still denoted as $\{\boldsymbol{\alpha}_n, \boldsymbol{\varphi}_n\}_{n \geq 0}$) such that both $\{\boldsymbol{\alpha}_n\}_{n \geq 0}$ and $\{\boldsymbol{\varphi}_n\}_{n \geq 0}$ converge weakly to $\boldsymbol{\alpha}^* \in L^\infty(\mathcal{B})$ and $\boldsymbol{\varphi}^* \in L^2(\mathcal{B})$, respectively, since the sets \mathcal{A}_α and \mathcal{A}_φ are closed and convex. We shall prove first that $(\boldsymbol{\alpha}^*, \boldsymbol{\varphi}^*)$ is indeed a minimizer of (8). Since each pair $(\boldsymbol{\alpha}^n, \boldsymbol{\varphi}^n)$ corresponds to a solution $c^n := c(\boldsymbol{\alpha}^n, \boldsymbol{\varphi}^n)$ to (5) with $\boldsymbol{\alpha} = \boldsymbol{\alpha}^n$ and $\boldsymbol{\varphi} = \boldsymbol{\varphi}^n$, by letting $\psi = c^n$ in (2) and using $\frac{1}{2} \partial_t c^2 = c_t c$, it follows immediately that the sequence $\{c^n := c(\boldsymbol{\alpha}^n, \boldsymbol{\varphi}^n)\}$ is also bounded in $L^2(0, T; H^1(\mathcal{B}))$. This indicates the existence of some $c^* \in L^2(0, T; H^1(\mathcal{B}))$ and a subsequence of $\{c^n\}$, still denoted by $\{c^n\}$, such that

$$c^n := c(\boldsymbol{\alpha}^n, \boldsymbol{\varphi}^n) \rightharpoonup c^* \quad \text{in } L^2(0, T; H^1(\mathcal{B})). \quad (9)$$

We claim $c^* = c(\boldsymbol{\alpha}^*, \boldsymbol{\varphi}^*)$. The continuity of $g(c)$ with respect to c together with (9) imply

$$\partial_t c(\boldsymbol{\alpha}^n, \boldsymbol{\varphi}^n) \rightharpoonup \partial_t c^*, \quad \nabla c(\boldsymbol{\alpha}^n, \boldsymbol{\varphi}^n) \rightharpoonup \nabla c^* \quad \text{in } L^2(Q).$$

Letting $n \rightarrow \infty$ in (2) with $\boldsymbol{\alpha} = \boldsymbol{\alpha}^n$ and $\boldsymbol{\varphi} = \boldsymbol{\varphi}^n$ yields

$$\int_0^T \int_{\mathcal{B}} (\partial_t c^* \psi + D(x) \nabla c^* \cdot \nabla \psi + \boldsymbol{\alpha}^*(x) \beta(t) c^* \psi) dx dt = \int_0^T \int_{\mathcal{B}} g(c^*) \psi dx dt, \quad (10)$$

for all $\psi \in L^2(0, T; H^1(\mathcal{B}))$.

Next, we shall prove $c^*(\cdot, 0) = \varphi^*$, which with (10) implies $c^* = c(\alpha^*, \varphi^*)$. In doing so, we take ϑ in $C^1[0, T]$, with $\vartheta(0) \neq 0$ and $\vartheta(T) = 0$ and $\zeta \in L^2(\mathcal{B})$. We employ the classical integration by parts formula to get

$$\int_0^T \int_{\mathcal{B}} \partial_t c(\alpha^n, \varphi^n) \psi \zeta \, dx \, dt = - \int_{\mathcal{B}} \varphi^n \psi(0) \zeta \, dx - \int_0^T \int_{\mathcal{B}} c(\alpha^n, \varphi^n) \partial_t \psi \zeta \, dx \, dt. \quad (11)$$

Letting $n \rightarrow \infty$ in (11), we conclude that

$$\int_0^T \int_{\mathcal{B}} \partial_t c^* \psi \zeta \, dx \, dt = - \int_{\mathcal{B}} \varphi^* \psi(0) \zeta \, dx - \int_0^T \int_{\mathcal{B}} c^* \partial_t \psi \zeta \, dx \, dt. \quad (12)$$

On the other hand, we have:

$$\int_0^T \int_{\mathcal{B}} \partial_t c^* \psi \zeta \, dx \, dt = - \int_{\mathcal{B}} c^*(\cdot, 0) \psi(0) \zeta \, dx - \int_0^T \int_{\mathcal{B}} c^* \partial_t \psi \zeta \, dx \, dt, \quad (13)$$

which, together with (12), implies that $c^*(\cdot, 0) = \varphi^*$. Therefore, $c^* := c(\alpha^*, \varphi^*)$.

By $\alpha^n \rightharpoonup \alpha^*$ and $\varphi^n \rightharpoonup \varphi^*$ in $L^\infty(\mathcal{B}) \subset L^2(\mathcal{B})$ and $L^2(\mathcal{B})$, respectively, we use the lower semi-continuity of the L^2 -norm to conclude that

$$\begin{aligned} J(\alpha^*, \varphi^*) &= \frac{1}{2} \left\| \int_0^T \omega_1(t) c(\alpha^*, \varphi^*) \, dt - \phi_1^\varepsilon \right\|_{L^2(\mathcal{B})}^2 + \frac{1}{2} \left\| \int_0^T \omega_2(t) c(\alpha^*, \varphi^*) \, dt - \phi_2^\varepsilon \right\|_{L^2(\mathcal{B})}^2 \\ &\leq \liminf_{n \rightarrow \infty} \frac{1}{2} \left\| \int_0^T \omega_1(t) c(\alpha^n, \varphi^n) \, dt - \phi_1^\varepsilon \right\|_{L^2(\mathcal{B})}^2 + \frac{1}{2} \left\| \int_0^T \omega_2(t) c(\alpha^n, \varphi^n) \, dt - \phi_2^\varepsilon \right\|_{L^2(\mathcal{B})}^2 \\ &\leq \liminf_{n \rightarrow \infty} J(\alpha^n, \varphi^n) = \inf_{\substack{\alpha \in \mathcal{A}_\alpha \\ \varphi \in \mathcal{A}_\varphi}} J(\alpha, \varphi). \end{aligned}$$

Hence, the proof is complete.

After that, we prove the stability of (8) in the sense that the minimization problem (8) effectively stabilizes the considered inverse problem against perturbations in the measured data. For this purpose, let $\{\phi_{1,\ell}^\varepsilon, \phi_{2,\ell}^\varepsilon\}_{\ell=0}^\infty$ denote a sequence of measurements of $\{\phi_1^\varepsilon, \phi_2^\varepsilon\}$ in $L^2(\mathcal{B}) \times L^2(\mathcal{B})$. For each $\ell \in \mathbb{N}$, we denote the solution of the minimization problem by $(\alpha_\ell, \varphi_\ell)$. Then, we have the following minimization problem:

$$\min_{\substack{\alpha \in \mathcal{A}_\alpha \\ \varphi \in \mathcal{A}_\varphi}} J_\ell(\alpha, \varphi), \quad J_\ell(\alpha, \varphi) := \frac{1}{2} \sum_{k=1}^2 \left\| \int_0^T \omega_k(t) c(\alpha, \varphi) \, dt - \phi_{k,\ell}^\varepsilon \right\|_{L^2(\mathcal{B})}^2, \quad \ell = 0, 1, \dots \quad (14)$$

The following theorem states the convergence of the sequence $\{\alpha_\ell, \varphi_\ell\}_{\ell \geq 0}$ when

$$\phi_{k,\ell}^\varepsilon \longrightarrow \phi_k^\varepsilon \quad \text{in } L^2(\mathcal{B}) \quad \text{as } \ell \longrightarrow \infty, \quad \text{for } k = 1, 2. \quad (15)$$

Theorem 2. *If (15) holds, then the sequence $\{\alpha_\ell, \varphi_\ell\}_{\ell \geq 0}$ of minimizers of the problems (14) converges weakly in $L^\infty(\mathcal{B}) \times L^2(\mathcal{B})$ to a minimizer, associated with the measured data $\{\phi_1^\varepsilon, \phi_2^\varepsilon\}$, of the optimization problem (8).*

Proof. The existence of each pair $(\alpha_\ell, \varphi_\ell)$ is guaranteed by Theorem 1. By definition, we have:

$$J_\ell(\alpha_\ell, \varphi_\ell) \leq J_\ell(\alpha, \varphi), \quad \forall (\alpha, \varphi) \in \mathcal{A}_\alpha \times \mathcal{A}_\varphi,$$

that implies the uniform boundedness of $(\alpha_\ell, \varphi_\ell)$ in $L^\infty(\mathcal{B}) \times L^2(\mathcal{B})$. Consequently, there exist $\alpha^* \in L^\infty(\mathcal{B})$, $\varphi^* \in L^2(\mathcal{B})$, and a subsequence (still denoted as $\{\alpha_\ell, \varphi_\ell\}_{\ell \geq 0}$), such that:

$$(\alpha_\ell, \varphi_\ell) \rightharpoonup (\alpha^*, \varphi^*) \quad \text{in } L^\infty(\mathcal{B}) \times L^2(\mathcal{B}) \quad \text{as } \ell \rightarrow \infty.$$

It suffices to show that (α^*, φ^*) is indeed a minimizer of (8). Repeating the same argument as that in the proof of Theorem 1, we derive:

$$c(\alpha_\ell, \varphi_\ell) \rightharpoonup c(\alpha^*, \varphi^*) \quad \text{in } L^2(0, T; H^1(\mathcal{B})) \quad \text{as } \ell \rightarrow \infty,$$

up to taking a further subsequence. Combining this convergence with (15), we obtain:

$$\int_0^T \omega_k(t) c(\alpha_\ell, \varphi_\ell) - \phi_{k,\ell}^\varepsilon dt \rightharpoonup \int_0^T \omega_k(t) c(\alpha^*, \varphi^*) - \phi_k^\varepsilon dt \quad \text{in } L^2(\mathcal{B}) \quad \text{as } \ell \rightarrow \infty, \quad \text{for } k = 1, 2.$$

Therefore, for $k = 1, 2$, we have:

$$\left\| \int_0^T \omega_k(t) c(\alpha^*, \varphi^*) dt - \phi_k^\varepsilon \right\|_{L^2(\mathcal{B})}^2 \leq \liminf_{\ell \rightarrow \infty} \left\| \int_0^T \omega_k(t) c(\alpha_\ell, \varphi_\ell) dt - \phi_{k,\ell}^\varepsilon \right\|_{L^2(\mathcal{B})}^2.$$

By using the lower semi-continuity of the L^2 -norm, we deduce that:

$$\begin{aligned} J(\alpha^*, \varphi^*) &= \frac{1}{2} \sum_{k=1}^2 \left\| \int_0^T \omega_k(t) c(\alpha^*, \varphi^*) dt - \phi_k^\varepsilon \right\|_{L^2(\mathcal{B})}^2 \\ &\leq \frac{1}{2} \liminf_{\ell \rightarrow \infty} \sum_{k=1}^2 \left\| \int_0^T \omega_k(t) c(\alpha_\ell, \varphi_\ell) dt - \phi_{k,\ell}^\varepsilon \right\|_{L^2(\mathcal{B})}^2 \\ &\leq \frac{1}{2} \lim_{\ell \rightarrow \infty} \sum_{k=1}^2 \left\| \int_0^T \omega_k(t) c(\alpha, \varphi) dt - \phi_{k,\ell}^\varepsilon \right\|_{L^2(\mathcal{B})}^2 \\ &= \frac{1}{2} \sum_{k=1}^2 \left\| \int_0^T \omega_k(t) c(\alpha, \varphi) dt - \phi_k^\varepsilon \right\|_{L^2(\mathcal{B})}^2 = J(\alpha, \varphi), \quad \forall (\alpha, \varphi) \in \mathcal{A}_\alpha \times \mathcal{A}_\varphi. \end{aligned}$$

Thus, the proof is complete.

3. Conjugate gradient method

In this section, we develop an efficient iterative procedure based on the conjugate gradient method to solve the minimization problem (8). Like all effective iterative methods for optimization problems, our approach requires information about the derivatives of the objective function. We start by deriving the Fréchet derivatives $J'_\alpha(\alpha, \varphi)$ and $J'_\varphi(\alpha, \varphi)$ of the objective functional $J(\alpha, \varphi)$ with respect to the unknown terms α and φ , respectively. Subsequently, we establish the conjugate gradient method, which utilizes these gradients to reconstruct the unknown coefficients simultaneously. We refer the readers to [13] for convergence of this algorithm.

3.1. Fréchet Derivatives

Following the methodology in [3, 4, 17], we can reduce the computational costs for the Fréchet derivatives by introducing the adjoint system of (5), which represents a backward time diffusion equation:

$$\begin{cases} -\partial_t v - \operatorname{div}(D(x)\nabla v) + (\alpha(x)\beta(t) - g'_{c(\alpha, \varphi)})v = \sum_{k=1}^2 \omega_k(t) \left(\int_0^T \omega_k(\tau) c(\alpha, \varphi) d\tau - \phi_k^\varepsilon \right) & \text{in } Q, \\ D(x)\partial_\nu v = 0 & \text{on } \mathcal{B}_T, \\ v(x, T) = 0 & \text{in } \bar{\mathcal{B}}, \end{cases} \quad (16)$$

where $g'_{c(\alpha, \varphi)}$ represents the Fréchet derivative of g at $c(\alpha, \varphi)$. The system (16) is linear and it is a well-posed problem, see [6]. For simplicity, we denote $c(\alpha, \varphi)$ by c .

Theorem 3. *The objective functional $J(\alpha, \varphi)$ is Fréchet differentiable and the derivatives $J'_\alpha(\alpha, \varphi)$ and $J'_\varphi(\alpha, \varphi)$ at $(\alpha, \varphi) \in \mathcal{A}_\alpha \times \mathcal{A}_\varphi$ are given by:*

$$\begin{cases} J'_\alpha(\alpha, \varphi) = - \int_0^T \beta(t) c(x, t) v(x, t) dt, \\ J'_\varphi(\alpha, \varphi) = v(x, 0), \end{cases} \quad (17)$$

where v is the solution to the adjoint problem (16).

We need the following two sensitivity problems to prove theorem 3.

Sensitivity problem associated with the treatment parameter. Assume $c(x, t)$ is perturbed by $\epsilon \eta_\alpha(x, t)$ and the therapy parameter $\alpha(x)$ is perturbed by $\epsilon \alpha_0(x)$, where $\epsilon > 0$ is a small number and $\alpha_0 \in \mathcal{A}_\alpha$. Substituting the perturbed cell density $c(x, t) + \epsilon \eta_\alpha(x, t)$ and the perturbed therapy parameter $\alpha(x) + \epsilon \alpha_0(x)$ into the original problem (5) yields a perturbed problem. The resulting sensitivity problem is given by

$$\begin{cases} \partial_t \eta_\alpha - \operatorname{div}(D(x)\nabla \eta_\alpha) + \alpha(x)\beta(t)\eta_\alpha - g'_c \eta_\alpha = -\beta(t)\alpha_0(x)c & \text{in } Q, \\ D\partial_\nu \eta_\alpha = 0 & \text{on } \mathcal{B}_T, \\ \eta_\alpha(x, 0) = 0 & \text{in } \bar{\Omega}. \end{cases} \quad (18)$$

Sensitivity problem associated with initial activation of the tissue. Similarly, assume $\mathbf{c}(x, t)$ is perturbed by $\epsilon \eta_\varphi(x, t)$ and the initial activation of the tissue, denoted as $\varphi(x)$, is perturbed by $\epsilon \varphi_0(x)$, where $\epsilon > 0$ is a small number and $\varphi_0 \in \mathcal{A}_\varphi$. Substituting the perturbed cell density $\mathbf{c}(x, t) + \epsilon \eta_\varphi(x, t)$ and perturbed initial activation of the tissue $\varphi(x) + \epsilon \varphi_0(x)$ into the original problem (5) yields a perturbed problem. The resulting sensitivity problem is given by

$$\begin{cases} \partial_t \eta_\varphi - \operatorname{div}(D(x) \nabla \eta_\varphi) + \alpha(x) \beta(t) \eta_\varphi - g'_c \eta_\varphi = 0 & \text{in } Q, \\ D \partial_\nu \eta_\varphi = 0 & \text{on } \mathcal{B}_T, \\ \eta_\varphi(x, 0) = \varphi_0(x) & \text{in } \bar{\Omega}. \end{cases} \quad (19)$$

Note that the problems (18) and (19) are linear, and their well-posedness is classical, see [6]. These sensitivity problems are important to determine the step sizes in the descent direction in the conjugate gradient method.

Proof of theorem 3. Let $\epsilon > 0$ and $\alpha_0 \in \mathcal{A}_\alpha$ be given, then direct calculations yield

$$\begin{aligned} J(\alpha + \epsilon \alpha_0, \varphi) - J(\alpha, \varphi) &= \epsilon \left\{ \sum_{k=1}^2 \int_Q \eta_\alpha(x, t) \omega_k(t) \left(\int_0^T \omega_k(\tau) \mathbf{c}(x, \tau) d\tau - \phi_k^\epsilon \right) dx dt \right\} \\ &\quad + \frac{\epsilon^2}{2} \left\{ \sum_{k=1}^2 \left(\left\| \int_0^T \omega_k(t) \eta_\alpha dt \right\|_2^2 \right) \right\}. \end{aligned} \quad (20)$$

Therefore, the directional derivative of $J(\alpha, \varphi)$ with respect to α in the direction $\alpha_0(x)$ is given by

$$\begin{aligned} J'_\alpha(\alpha, \varphi) \cdot \alpha_0 &= \lim_{\epsilon \rightarrow 0} \frac{J(\alpha + \epsilon \alpha_0, \varphi) - J(\alpha, \varphi)}{\epsilon} \\ &= \sum_{k=1}^2 \int_Q \eta_\alpha(x, t) \omega_k(t) \left(\int_0^T \omega_k(\tau) \mathbf{c}(x, \tau) d\tau - \phi_k^\epsilon \right) dx dt. \end{aligned}$$

Multiplying both sides of the first equation in (16) by η_α , integrating over Q , using the Green formula and the fact that $D(x) \partial_\nu v = 0$ on \mathcal{B}_T , we obtain

$$\begin{aligned} & - \int_Q \partial_t v \eta_\alpha dx dt + \int_Q D(x) \nabla v \nabla \eta_\alpha dx dt + \int_Q (\alpha(x) \beta(t) - g'_c) v \eta_\alpha dx dt \\ &= \sum_{k=1}^2 \int_Q \omega_k(t) \eta_\alpha \left(\int_0^T \omega_k(\tau) \mathbf{c}(x, \tau) d\tau - \phi_k^\epsilon \right) dx dt. \end{aligned}$$

Applying the Green's and the integration by parts formulas in the above identity and taking into account that $\eta_\alpha(x, 0) = 0$ and $v(x, T) = 0$ for all $x \in \bar{\Omega}$, we get

$$\begin{aligned} & \int_Q (\partial_t \eta_\alpha - \operatorname{div}(D(x)\nabla \eta_\alpha) + \alpha(x) \beta(t) \eta_\alpha - g'_c \eta_\alpha) v \, dx \, dt \\ &= \sum_{k=1}^3 \int_Q \omega_k(t) \eta_\alpha \left(\int_0^T \omega_k(\tau) \mathbf{c}(x, \tau) \, d\tau - \phi_k^\varepsilon \right) \, dx \, dt. \end{aligned}$$

The first equation in (18) leads to

$$\sum_{k=1}^2 \int_Q \omega_k(t) \eta_\alpha \left(\int_0^T \omega_k(\tau) \mathbf{c}(x, \tau) \, d\tau - \phi_k^\varepsilon \right) \, dx \, dt = - \int_Q \beta(t) \alpha_0(x) \mathbf{c}(x, t) v(x, t) \, dx \, dt.$$

Therefore, the directional derivative of $J(\alpha, \varphi)$ with respect to α in the direction $\alpha_0(x)$ is given by

$$J'_\alpha(\alpha, \varphi) \cdot \alpha_0 = - \int_Q \beta(t) \alpha_0(x) \mathbf{c}(x, t) v(x, t) \, dx \, dt. \quad (21)$$

Similarly, by using the weak formulations of the adjoint problem (16) and the sensitivity problem (19), we can derive

$$J'_\varphi(\alpha, \varphi) \cdot \varphi_0 = \int_\Omega \varphi_0(x) v(x, 0) \, dx.$$

The proof is complete.

3.2. Iterative algorithm

In this subsection, we employ a conjugate gradient method to approximate the minimizer of the functional $J(\alpha, \varphi)$ defined by (7). Let (α^k, φ^k) be the k th approximation of the solution (α, φ) . The iterations are given by the following equations:

$$\alpha^{k+1} = \alpha^k + \zeta_\alpha^k S_\alpha^k \quad \text{and} \quad \varphi^{k+1} = \varphi^k + \zeta_\varphi^k S_\varphi^k, \quad k = 0, 1, 2, \dots \quad (22)$$

In (22), the index k denotes the number of iterations, $\alpha^0(x)$ and $\varphi^0(x)$ are the initial guesses for $\alpha(x)$ and $\varphi(x)$, respectively. The terms ζ_α^k and ζ_φ^k are the step sizes, and S_α^k and S_φ^k represent the search directions which are determined by the following equations:

$$S_\alpha^k = \begin{cases} -J_\alpha^0, & k = 0, \\ -J_\alpha^k + \vartheta_\alpha^k S_\alpha^{k-1}, & k \geq 1, \end{cases} \quad \text{and} \quad S_\varphi^k = \begin{cases} -J_\varphi^0, & k = 0, \\ -J_\varphi^k + \vartheta_\varphi^k S_\varphi^{k-1}, & k \geq 1, \end{cases} \quad (23)$$

where $J_\alpha^k = J'_\alpha(\alpha^k, \varphi^k)$ and $J_\varphi^k = J'_\varphi(\alpha^k, \varphi^k)$ denote the derivatives of $J(\alpha, \varphi)$ with respect to α and φ , respectively, at point (α^k, φ^k) . The terms ϑ_α^k and ϑ_φ^k are the conjugate coefficients obtained using the Fletcher-Reeves formula [12]

$$\vartheta_\alpha^k = \frac{\|J_\alpha^k\|_2}{\|J_\alpha^{k-1}\|_2} \quad \text{and} \quad \vartheta_\varphi^k = \frac{\|J_\varphi^k\|_2}{\|J_\varphi^{k-1}\|_2}, \quad k = 1, 2, \dots \quad (24)$$

Building upon the preceding discussions, it is noteworthy that all parameters are explicitly expressed, except the search step sizes ζ_α^k and ζ_φ^k . The determination of these parameters is accomplished through the utilization of the line search method. Specifically, the values for ζ_α^k and ζ_φ^k are obtained by minimizing the following functional:

$$J(\alpha^{k+1}, \varphi^{k+1}) := \frac{1}{2} \sum_{i=1}^2 \left(\left\| \int_0^T \omega_i(t) u(\alpha^{k+1}, \varphi^{k+1}) dt - \phi_i^\varepsilon \right\|_2^2 \right).$$

Since the search step sizes ζ_α^k and ζ_φ^k are implicit in the expression of $J(\alpha^{k+1}, \varphi^{k+1})$, we adopt a similar approach to [3, 4] and linearize it such that these parameters become explicit in the new formulation. Hence $c(\alpha^{k+1}, \varphi^{k+1})$ becomes as follows:

$$c(\alpha^{k+1}, \varphi^{k+1}) \approx c(\alpha^k, \varphi^k) + \zeta_\alpha^k \eta_\alpha^k + \zeta_\varphi^k \eta_\varphi^k, \tag{25}$$

where η_α^k and η_φ^k are obtained by solving the sensitivity problems (18) and (19) with $(\alpha, \varphi) = (\alpha^k, \varphi^k)$ and $(\alpha_0, \varphi_0) = (S_\alpha^k, S_\varphi^k)$. We now define

$$c_i^k = \int_0^T \omega_i c^k dt, \quad \eta_{\alpha,i}^k = \int_0^T \omega_i \eta_\alpha^k dt \quad \text{and} \quad \eta_{\varphi,i}^k = \int_0^T \omega_i \eta_\varphi^k dt, \quad \text{for } i = 1, 2,$$

with $c^k = c(\alpha^k, \varphi^k)$. Then, from (7) and (25), we have

$$J(\alpha^{k+1}, \varphi^{k+1}) = \frac{1}{2} \sum_{i=1}^2 \int_\Omega (c_i^k + \zeta_\alpha^k \eta_{\alpha,i}^k + \zeta_\varphi^k \eta_{\varphi,i}^k - \phi_i^\varepsilon)^2 dx. \tag{26}$$

It follows

$$\frac{\partial J(\alpha^{k+1}, \varphi^{k+1})}{\partial \zeta_\alpha^k} := \mathcal{R}_1 \zeta_\alpha^k + \mathcal{R}_2 \zeta_\varphi^k - \mathcal{Y}_1 \quad \text{and} \quad \frac{\partial J(\alpha^{k+1}, \varphi^{k+1})}{\partial \zeta_\varphi^k} := \mathcal{R}_2 \zeta_\alpha^k + \mathcal{R}_3 \zeta_\varphi^k - \mathcal{Y}_2$$

, where

$$\begin{aligned} \mathcal{R}_1 &= \sum_{i=1}^2 \|\eta_{\alpha,i}^k\|_2^2, & \mathcal{R}_2 &= \sum_{i=1}^2 \int_{\mathcal{B}} \eta_{\alpha,i}^k \eta_{\varphi,i}^k dx, & \mathcal{R}_3 &= \sum_{i=1}^2 \|\eta_{\varphi,i}^k\|_2^2, \\ \mathcal{Y}_1 &= \sum_{i=1}^2 \int_{\mathcal{B}} \eta_{\alpha,i}^k (\phi_i^\varepsilon - c_i^k) dx, & \text{and} & & \mathcal{Y}_2 &= \sum_{i=1}^2 \int_{\mathcal{B}} \eta_{\varphi,i}^k (\phi_i^\varepsilon - c_i^k) dx. \end{aligned}$$

By setting

$$\frac{\partial J(\alpha^{k+1}, \varphi^{k+1})}{\partial \zeta_\alpha^k} = \frac{\partial J(\alpha^{k+1}, \varphi^{k+1})}{\partial \zeta_\varphi^k} = 0,$$

one can deduce that the search step sizes ζ_α^k and ζ_φ^k satisfy the following linear system

$$\begin{cases} \mathcal{R}_1 \zeta_\alpha^k + \mathcal{R}_2 \zeta_\varphi^k = \mathcal{Y}_1, \\ \mathcal{R}_2 \zeta_\alpha^k + \mathcal{R}_3 \zeta_\varphi^k = \mathcal{Y}_2. \end{cases}$$

Therefore, direct calculations yield

$$\zeta_{\alpha}^k = \frac{\mathcal{R}_3 \mathcal{Y}_1 - \mathcal{R}_2 \mathcal{Y}_2}{\mathcal{R}_3 \mathcal{R}_1 - \mathcal{R}_2^2} \quad \text{and} \quad \zeta_{\varphi}^k = \frac{\mathcal{R}_1 \mathcal{Y}_2 - \mathcal{R}_2 \mathcal{Y}_1}{\mathcal{R}_3 \mathcal{R}_1 - \mathcal{R}_2^2}. \quad (27)$$

In the context of iterative algorithms, it is widely recognized that determining a suitable stopping criterion is crucial. Addressing this concern, the present study employs the discrepancy principle to conclude the iteration process. According to this principle, the iteration procedure if the following condition is fulfilled:

$$J(\alpha^k, \varphi^k) \leq \bar{\varepsilon}, \quad (28)$$

where $\bar{\varepsilon}$ is a small positive value, such as $\bar{\varepsilon} = 10^{-6}$ for exact measurements. In cases where the measurements exhibit noise,

$$\bar{\varepsilon} = \frac{1}{2} \sum_{i=1}^2 \|\phi_i^{\varepsilon} - \phi_i\|_2^2.$$

Building upon (6), it is established that $\bar{\varepsilon} \leq \varepsilon^2$. Consequently, the primary steps of our reconstruction approach are summarized in the following algorithm:

Algorithm 1 (Step 1). Set $k = 0$ and choose initial guesses α^0 and φ^0 for the unknown coefficients α and φ , respectively.

Step 2. Solve the direct problem (5) with $(\alpha, \varphi) = (\alpha^k, \varphi^k)$ to get $c^k = c(\alpha^k, \varphi^k)$.

Step 3. Solve the adjoint problem (16) and evaluate the gradients $J_{\alpha}^{\prime,k} = J'_{\alpha}(\alpha^k, \varphi^k)$ and $J_{\varphi}^{\prime,k} = J'_{\varphi}(\alpha^k, \varphi^k)$ given in theorem 3.

Step 4. Calculate the conjugate coefficients ϑ_{α}^k and ϑ_{φ}^k by (24) and the directions S_{α}^k and S_{φ}^k by (23).

Step 5. Calculate the step sizes ζ_{α}^k and ζ_{φ}^k by (27).

Step 6. Update α^{k+1} and φ^{k+1} by (22).

Step 7 If the condition (28) is satisfied, then go to **Step 8**. Otherwise set $k = k + 1$ and go to **Step 2**.

Step 8 End.

4. Numerical experiments

We set $T = 1$ and $\mathcal{B} = (0, 1)$ in all numerical examples. We rely on the finite difference method as our primary tool to deal with not only the direct problem but also its associated sensitive and adjoint variants. Setting a structured approach, we designate $h = 0.02$ as the step size and $\tau = 0.005$ as the time increment, ensuring a balanced exploration of the problem domain both spatially and temporally. Moreover, to handle integrals within our computations, we employ the trapezoidal rule, a widely recognized numerical technique that provides reliable approximations. All numerical experiments are carried out in MATLAB (version R2017a).

In the context of noisy observations, we enhance the fidelity of our simulations by incorporating Gaussian additive noise to ϕ_1^ε and ϕ_2^ε , as detailed in (6). This noise, with amplitude σ , is dynamically adjusted based on a specified percentage p representing the noise level in the data. Utilizing the term "random(1)", we introduce variability representative of real-world conditions by generating random values from a Gaussian distribution with a mean of zero and a standard deviation of unity. Specifically, the noisy observations ϕ_i^ε are generated according to:

$$\phi_i^\varepsilon = \phi_i + \sigma \times \text{random}(1), \quad i = 1, 2,$$

where σ is calculated as $\sigma = \frac{p}{100} \times \max_{x \in \bar{\Omega}} \{|\phi_1(x)|, |\phi_2(x)|\}$. This approach ensures that our computational model captures the inherent uncertainties present in real-world measurements, enhancing the robustness of our simulations. The accuracy errors $E_\alpha(k)$ and $E_\varphi(k)$, as functions of the iteration number k , for the treatment parameter $\alpha(x)$ and the initial cell density $\varphi(x)$, respectively, are defined as

$$E_\alpha(k) := \|\alpha^k - \alpha\|_2, \quad (29)$$

$$E_\varphi(k) := \|\varphi^k - \varphi\|_2, \quad (30)$$

where α^k and φ^k are the CGM iterates, and α and φ are the analytical expressions of the treatment parameter and the initial cell density.

Throughout this section, we maintain a consistent temporal component $\beta(t)$ within the treatment profile, specified as $\exp(-t)$. Furthermore, our initial guesses for the treatment parameter α^0 and initial cell density φ^0 are both set to 1, establishing a standardized starting point for our analysis. Expanding upon our problem configuration, the parameters governing the logistic reaction function $g(c)$ and the diffusion term are listed in Table 1.

Parameter	Value	References
Growth rate ρ	0.0012	Cock et al. [8]
Diffusion coefficient d_g	0.0013	Tracqui et al. [27]
Diffusion coefficient d_ω	0.0065	Swanson et al. [24]
Carrying capacity c_{lim}	0.625	Noviantri et al. [18]

Table 1: Numerical values of physical coefficients.

Example 1. In this example, we aim to evaluate the performance of our algorithm in identifying the treatment parameter α and initial cell density φ within a controlled setting. These two functions are assumed to exhibit smooth behavior, reflecting typical characteristics observed in many practical applications. By selecting smooth functions for α and φ , we aim to establish a baseline assessment of the algorithm's capability to accurately reconstruct these essential parameters under ideal conditions. This test serves as a crucial step in validating the effectiveness and reliability of our computational approach before extending it to more complex scenarios. In doing so, we choose the weight functions as $\omega_1(t) = 1$ and $\omega_2(t) = e^{-t}$, tailoring our algorithm to specific characteristics of the problem domain. With these choices, we proceed to test our algorithm's ability to reconstruct a treatment parameter α and initial cell density φ defined as

$$\alpha(x) = 2x + \sin(\pi x) + 1 \quad \text{and} \quad \varphi(x) = \frac{\cos(4\pi x) + 1}{2}.$$

In Figure 1, we show the behavior of the objective functional $J(\alpha^k, \varphi^k)$, as defined in (7), with respect to the iteration number k . This functional characterizes the simultaneous recovery of the two unknown parameters $\alpha(x)$ and $\varphi(x)$ in the noise free case ($p = 0$), as well as with noise levels $p = 1$ and $p = 2$. From this figure, it can be seen that the objective functional (7) is monotonic decreasing and convergent. Additionally, we observe rapid convergence of the functional to a small positive value, underscoring the algorithm's efficiency in minimizing the objective and thereby refining the parameter estimates.

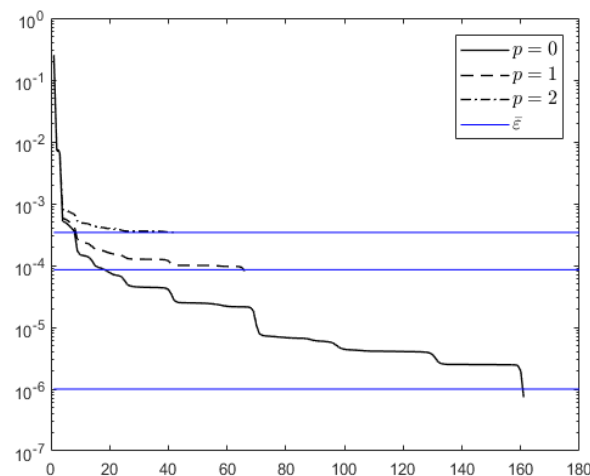


Figure 1: The objective functional $J(\alpha^k, \varphi^k)$, for different noise levels $p \in \{0, 1, 2\}$, in **Example 1**.

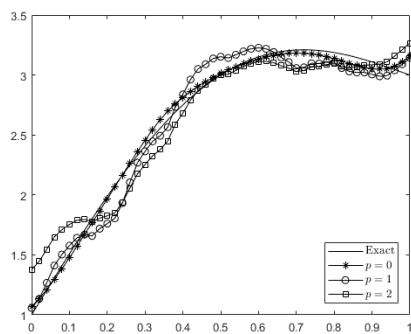
Based on the discrepancy principle (28), Table 2 presents a comprehensive overview of key metrics for each chosen noise level $p \in \{0, 1, 2\}$. Specifically, the table lists the stopping iteration numbers k^* , the threshold $\bar{\varepsilon}$, the errors (29) associated with the unknowns $\alpha(x)$

and $\varphi(x)$, as well as the norms of the Fréchet gradients $\|J'_{\alpha^{k^*}}\|_2$ and $\|J'_{\varphi^{k^*}}\|_2$.

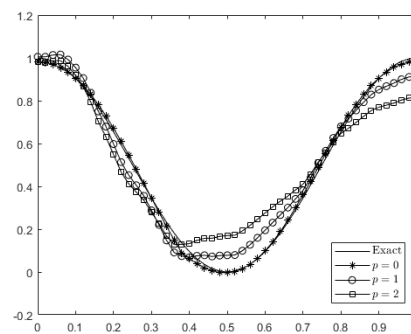
	$\bar{\varepsilon}$	k^*	$E_{\alpha}(k^*)$	$E_{\varphi}(k^*)$	$\ J'_{\alpha^{k^*}}\ _2$	$\ J'_{\varphi^{k^*}}\ _2$
$p = 0$	$1.00e - 06$	161	$4.41e - 02$	$3.11e - 02$	$2.09e - 04$	$3.34e - 04$
$p = 1$	$8.54e - 05$	66	$1.21e - 01$	$1.17e - 01$	$3.04e - 04$	$3.63e - 04$
$p = 2$	$3.41e - 04$	43	$2.21e - 01$	$1.56e - 01$	$1.24e - 03$	$1.45e - 03$

Table 2: Choices of noise levels p and their respective threshold values, the corresponding stopping iteration numbers k^* , L^2 -errors, and norms of the Fréchet gradients in **Example 1**.

From Table 2, it is evident that the numerical solutions are reasonably accurate for both $\alpha(x)$ and $\varphi(x)$. Furthermore, the norms of the Fréchet gradients indicate that the Conjugate Gradient Method (CGM), with iterations stopped by the discrepancy principle (28), serves as a semi-convergent regularization method. In Figure 2, we illustrate the numerical solutions for the treatment parameter $\alpha(x)$ and the initial cell density $\varphi(x)$ at the stopping iteration numbers k^* determined from Table 2. These solutions are shown for noise levels p in $\{0, 1, 2\}$. From Figure 2, it is evident that for noise-free data ($p = 0$), the analytical and numerical solutions overlap, making them indistinguishable graphically. However, as the level of noise increases, the accuracy of the solutions decreases. Overall, it can be observed that stable and accurate solutions are obtained for both coefficients $\alpha(x)$ and $\varphi(x)$.



(a) The treatment parameter $\alpha(x)$



(b) The initial cell density $\varphi(x)$

Figure 2: The numerical results for **Example 1** for $p \in \{0, 1, 2\}$. Left: the treatment parameter $\alpha(x)$. Right: the initial cell density $\varphi(x)$.

Example 2. In this example, we test our algorithm to recover a smooth treatment parameter and a non-smooth initial cell density, driven by considerations in brain tumor growth dynamics. Typically, treatment parameters in such scenarios exhibit smooth variations, while initial cell densities may feature sharp changes or irregularities. By addressing these characteristics, we aim to enhance the applicability of our algorithm in real-world

medical contexts, where accurate characterization of tumor growth dynamics is crucial for effective treatment planning and monitoring. Moreover, investigating the algorithm’s performance in recovering both smooth and non-smooth parameters provides valuable insights into its robustness and versatility across diverse tumor growth patterns.

To demonstrate the effectiveness of our proposed iterative method in this scenario, we define the exact treatment parameter $\alpha(x)$ and initial cell density $\varphi(x)$ to be reconstructed as follows:

$$\alpha(x) = \frac{1 + \sin(4\pi x)}{2} \text{ and } \varphi(x) = \begin{cases} 0, & \text{if } x \in [0, 0.2] \cup [0.8, 0.1], \\ 2 \sin(\pi x) - 1, & \text{if } x \in [0.2, 0.5], \\ -\frac{10x}{3} + \frac{8}{3}, & \text{Otherwise.} \end{cases}$$

The measured data ϕ_i^ε , for $i = 1, 2$, are generated using the weight functions $\omega_1(t) = 1 + t^2$ and $\omega_2(t) = \exp(-2t)$. Figure 3 illustrates the objective functional $J(\alpha^k, \varphi^k)$ given by (7) for the simultaneous reconstruction of the unknown coefficients with noise levels $p \in \{0, 1, 2\}$. From this figure, it is evident that the objective function rapidly decreases monotonically and thus it is convergent.

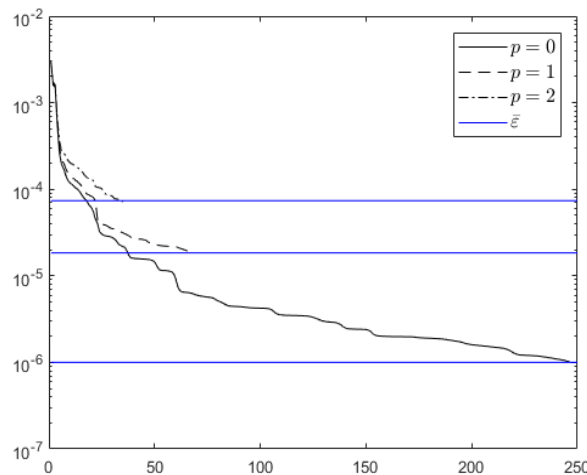


Figure 3: The objective functional $J(\alpha^k, \varphi^k)$, for different noise levels $p \in \{0, 1, 2\}$, in **Example 2**.

The choices of p in the test and the corresponding numerical performances are listed in table 3.

	$\bar{\varepsilon}$	k^*	$E_{\alpha}(k^*)$	$E_{\varphi}(k^*)$	$\ J'_{\alpha, k^*}\ _2$	$\ J'_{\varphi, k^*}\ _2$
$p = 0$	$1.00e - 06$	167	$1.14e - 02$	$1.33e - 01$	$3.43e - 04$	$2.58e - 05$
$p = 1$	$3.18e - 02$	17	$9.05e - 02$	$2.16e - 01$	$1.17e - 01$	$1.01e - 01$
$p = 2$	$1.27e - 01$	9	$1.06e - 01$	$3.07e - 01$	$3.30e - 01$	$1.22e - 01$

Table 3: Choices of noise levels p alongside their respective threshold values, the corresponding stopping iteration numbers k^* , L^2 -errors, and norms of the Fréchet gradients in **Example 2**.

In comparison with **Example 1**, Table 3 highlights the impact of the non-smoothness of the initial cell density on the accuracy of our method. The corresponding numerical solutions for the treatment parameter $\alpha(x)$ and the initial cell density $\varphi(x)$ at the stopping iteration numbers obtained based on the discrepancy principle (28) are illustrated in Figure 4.

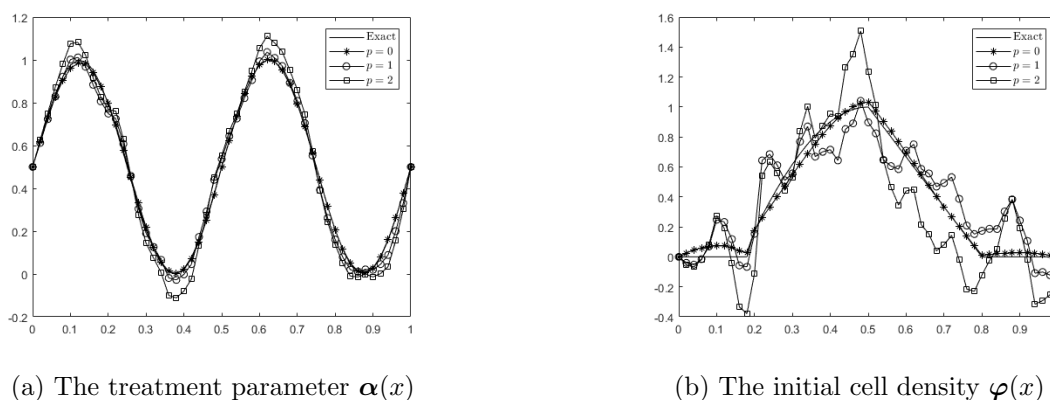


Figure 4: The numerical results for **Example 2** for different noise levels $p \in \{0, 1, 2\}$. Left: the treatment parameter $\alpha(x)$. Right: the initial cell density $\varphi(x)$.

Similar to the previous test, Figure 4 demonstrates that the efficiency of our algorithm diminishes with increasing noise level p . Specifically, the numerical results exhibit high accuracy when no noise is present (0% noise added in the measured data ϕ_i , for $i = 1, 2$). It is noteworthy that the reconstruction of smooth functions displays higher accuracy compared to non-smooth cases. However, stable and accurate solutions are attained for both coefficients overall.

Example 3. In this example, we test our algorithm to identify a non-smooth treatment parameter and a smooth initial cell density. This experimental setup stems from insights gained in brain tumor dynamics, where treatments such as radiation therapy or chemotherapy often induce localized variations in treatment effectiveness, resulting in non-smooth patterns in the treatment parameter. Conversely, the initial distribution of healthy and tumor cells within the brain tissue tends to exhibit a more uniform or grad-

ual pattern, reflecting the inherent composition and structure of the brain. By exploring this contrast in parameter characteristics, we aim to further enhance the adaptability and effectiveness of our algorithm in diverse tumor growth scenarios. For this purpose, we set $\omega_1(t) = \log(1 + t^2)$ and $\omega_2(t) = 2$, and evaluate the algorithm's performance in recovering:

$$\alpha(x) = \max(1, 1 - (1 - x^2) \cos(5\pi x)) \text{ and } \varphi(x) = \frac{\cos(\pi x) - x^2 + 2x}{2}.$$

The objective functional $J(\alpha^k, \varphi^k)$ defined by (7) is shown in Figure 5. It is clear that the objective function decreases steadily and monotonically with the iteration numbers k and thus it is convergent.

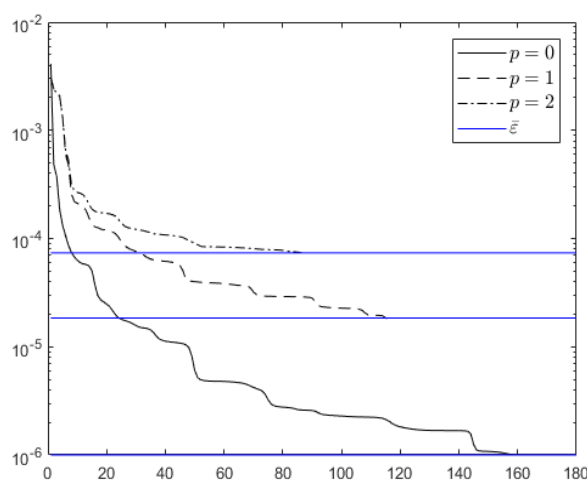


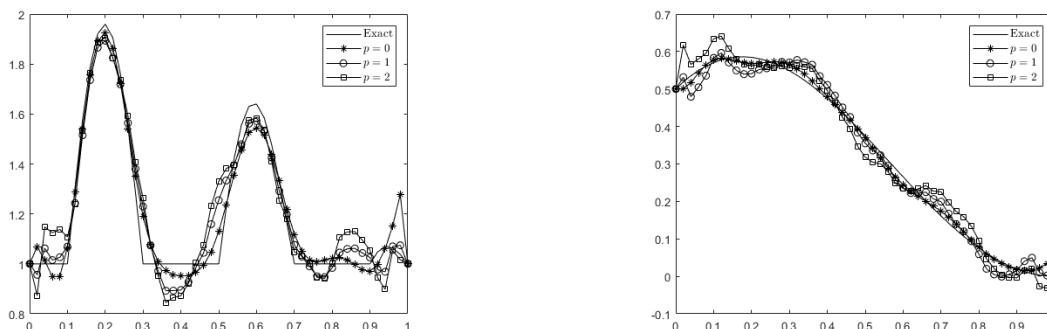
Figure 5: The objective functional $J(\alpha^k, \varphi^k)$, for different noise levels $p \in \{0, 1, 2\}$, in **Example 3**.

Based on the discrepancy principle (28) for each chosen noise level $p \in \{0, 1, 2\}$, we present the choices of p in this test along with the corresponding numerical performances in Table 4.

	$\bar{\varepsilon}$	k^*	$E_\alpha(k^*)$	$E_\varphi(k^*)$	$\ J'_\alpha{}^{k^*}\ _2$	$\ J'_\varphi{}^{k^*}\ _2$
$p = 0$	$1.00e - 06$	162	$6.75e - 02$	$2.46e - 02$	$6.36e - 05$	$1.01e - 04$
$p = 1$	$1.83e - 05$	116	$7.92e - 02$	$5.71e - 02$	$6.62e - 04$	$6.94e - 04$
$p = 2$	$7.35e - 05$	88	$1.08e - 01$	$7.27e - 02$	$1.06e - 03$	$5.77e - 03$

Table 4: Choices of noise levels p and their respective threshold values, the corresponding stopping iteration numbers k^* , L^2 -errors, and norms of the Fréchet gradients in **Example 3**.

Similar to the previous scenarios, we conclude this test by illustrating the comparisons of the recovered solutions with the exact ones in Figure 6.



(a) The treatment parameter $\alpha(x)$

(b) The initial cell density $\varphi(x)$

Figure 6: The numerical results for **Example 3** for different noise levels $p \in \{0, 1, 2\}$. Left: the treatment parameter $\alpha(x)$. Right: the initial cell density $\varphi(x)$.

As it is expected from the previous test, Figure 6 confirms that the efficiency of our algorithm is sensitive to both the noise level and the smoothness of the coefficient being reconstructed.

Example 4. In brain tumor growth, various factors such as genetic mutations, cellular interactions, and microenvironmental conditions can contribute to the formation of complex spatial patterns in both the treatment response and the distribution of tumor cells. These patterns may include regions of high and low treatment effectiveness, as well as areas of dense and sparse tumor cell populations, resulting in non-smooth variations in these coefficients. According to this observation, in this test, we focus on the identification of the treatment parameter and the initial cell density in more complex scenarios, where both coefficients to be reconstructed are represented by non - smooth functions. In doing so, we choose the weight functions as $\omega_1(t) = 1 + t + t^2$ and $\omega_2(t) = t^3$, and we test our iterative procedure to identify the following two coefficients:

$$\alpha(x) = 2 - |1 - 2x| \text{ and } \varphi(x) = \begin{cases} 0, & \text{if } x \in [0, 0.2] \cup [0.8, 0.1], \\ \frac{10x}{3} - \frac{2}{3}, & \text{if } x \in [0.2, 0.5], \\ -\frac{10x}{3} + \frac{8}{3}, & \text{if } x \in [0.5, 0.8]. \end{cases}$$

Following the same steps as in the previous tests, we start this analysis by illustrating the evolution of the objective functional $J(\alpha^k, \varphi^k)$ in Figure 7. This depicts the simultaneous recovery of the treatment parameter $\alpha(x)$ and the initial cell density $\varphi(x)$, considering noise levels $p = \{0, 1, 2\}$. Notably, the objective function exhibits a monotonically decreasing trend with k , converging rapidly to a small positive value, as expected.

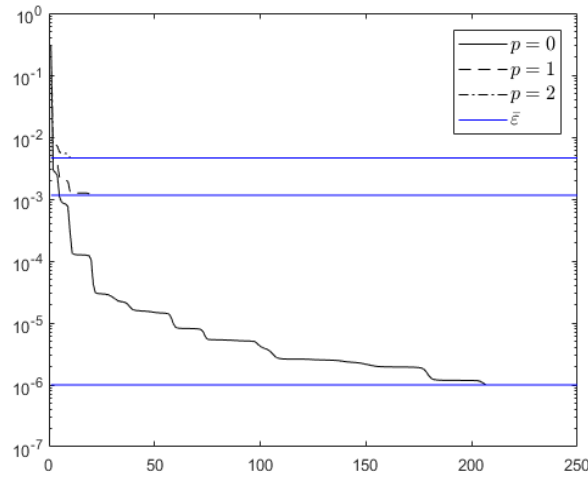


Figure 7: The objective functional $J(\alpha^k, \varphi^k)$, for different noise levels $p \in \{0, 1, 2\}$, in **Example 4**.

The choices of noise levels p along with their corresponding numerical performances are detailed in Table 5.

	$\bar{\varepsilon}$	k^*	$E_\alpha(k^*)$	$E_\varphi(k^*)$	$\ J'_\alpha{}^{k^*}\ _2$	$\ J'_\varphi{}^{k^*}\ _2$
$p = 0$	$1.00e - 06$	207	$6.23e - 02$	$1.52e - 02$	$6.21e - 04$	$1.61e - 04$
$p = 1$	$1.15e - 03$	20	$1.11e - 01$	$4.90e - 02$	$1.01e - 03$	$7.27e - 03$
$p = 2$	$4.60e - 03$	11	$1.58e - 01$	$7.38e - 02$	$1.09e - 03$	$6.52e - 03$

Table 5: Choices of noise levels p alongside their respective threshold values, the corresponding stopping iteration numbers k^* , L^2 -errors, and norms of the Fréchet gradients in **Example 4**.

Utilizing the stopping iteration numbers k^* , obtained according to the discrepancy principle (28), we present the comparisons between the exact solutions and the recovered ones for each chosen noise level p in Figure 8.

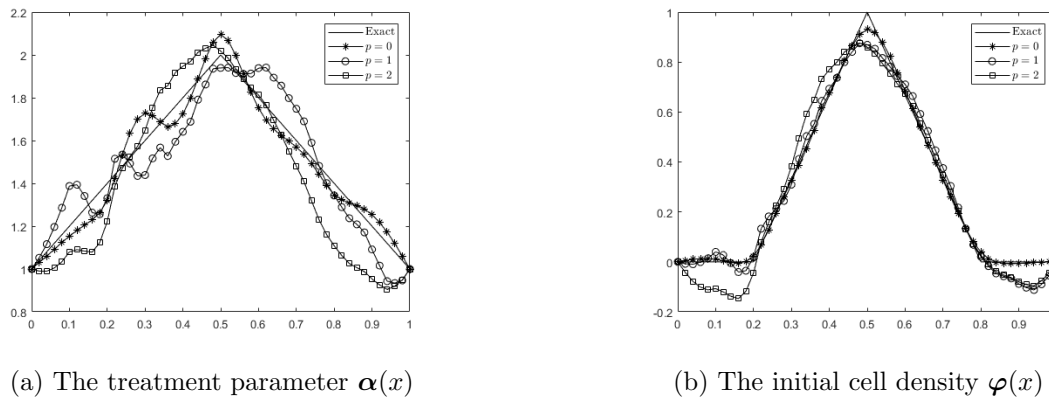


Figure 8: The numerical results for **Example 4** for different noise levels $p \in \{0, 1, 2\}$. Left: the treatment parameter $\alpha(x)$. Right: the initial cell density $\varphi(x)$.

In comparison with all previous tests, it is evident from Figure 8 that the accuracy and efficiency are reduced in this scenario due to the non-smoothness of the functions to be reconstructed. However, overall, the obtained results are acceptable.

5. Conclusions

In this paper, we addressed an inverse problem involving the simultaneous determination of the treatment response parameter and the initial tumor cell density in a nonlinear parabolic problem related to brain tumor dynamics. By reformulating the inverse problem as a minimization problem, we established the existence and initial stability of the solution. We proved the Fréchet differentiability of the objective (cost) functional and presented explicit formulas for the derivatives by using the solution to related adjoint problem. Based on the Fréchet differentiability of the objective (cost) functional, we developed an efficient iterative procedure using the conjugate gradient method combined with Morozov's discrepancy principle to solve the variational problem. Numerical examples, with both noise-free and noisy data, demonstrated the applicability and accuracy of the proposed method. On the other hand, several mathematical issues of high interest have not been discussed in this paper. The stability problem is one of them. The full stability issue is, however, to the best of our knowledge, still an open problem which deserves further investigation. Additionally, the uniqueness analysis of the considered inverse problem remains an area for future research. These aspects will be the focus of forthcoming studies to enhance the robustness and reliability of the proposed methods in practical applications.

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