Data driven modeling of pseudopalisade pattern formation

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Abstract

Pseudopalisading is an interesting phenomenon where cancer cells arrange themselves to form a dense garland-like pattern. Unlike the palisade structure, a similar type of pattern first observed in schwannomas by pathologist J.J. Verocay (Wippold et al. in AJNR Am J Neuroradiol 27(10):2037–2041, 2006), pseudopalisades are less organized and associated with a necrotic region at their core. These structures are mainly found in glioblastoma (GBM), a grade IV brain tumor, and provide a way to assess the aggressiveness of the tumor. Identification of the exact bio-mechanism responsible for the formation of pseudopalisades is a difficult task, mainly because pseudopalisades seem to be a consequence of complex nonlinear dynamics within the tumor. In this paper we propose a data-driven methodology to gain insight into the formation of different types of pseudopalisade structures. To this end, we start from a state of the art macroscopic model for the dynamics of GBM, that is coupled with the dynamics of extracellular pH, and formulate a terminal value optimal control problem. Thus, given a specific, observed pseudopalisade pattern, we determine the evolution of parameters (bio-mechanisms) that are responsible for its emergence. Random histological images exhibiting pseudopalisade-like structures are chosen to serve as target pattern. Having identified the optimal model parameters that generate the specified target pattern, we then formulate two different types of pattern counteracting ansatzes in order to determine possible ways to impair or obstruct the process of pseudopalisade formation. This provides the basis for designing active or live control of malignant GBM. Furthermore, we also provide a simple, yet insightful, mechanism to synthesize new pseudopalisade patterns by linearly combining the optimal model parameters respon-
sible for generating different known target patterns. This particularly provides a hint that complex pseudopalisade patterns could be synthesized by a linear combination of parameters responsible for generating simple patterns. Going even further, we ask ourselves if complex therapy approaches can be conceived, such that some linear combination thereof is able to reverse or disrupt simple pseudopalisade patterns; this is investigated with the help of numerical simulations.

**Keywords**  Optimal control · Pseudopalisades · Data driven modeling · Pattern formation

**Mathematics Subject Classification**  35K59 · 35Q92 · 49J20

1 Introduction

Biological phenomena produce some of the visually most appealing patterns, but unfortunately not all of them can be associated with a beneficial outcome. For example, skin and tissue patterns of animals could actually indicate the onset/progression of a harmful process. Pseudopalisades belong to such category, wherein the microscopic cellular arrangement, although both visually and dynamically quite intriguing, indicates the most advanced stage of glioblastoma multiforme (GBM), a type of brain tumor, which in most cases is lethal. Such patterns are actually used to pathologically characterize the aggressiveness or malignancy of the tumor (Wippold et al. 2006). Unlike the highly regular palisade structure observed in schwannoma cells, pseudopalisade structures are less organized and more irregular in appearance. The initiation of such pathological structures is not very clear, but is mainly hypothesized (see Brat et al. 2004) to be a complex interaction of different biophysical processes such as: (i) rapidly proliferating neoplastic cells, (ii) cells being highly resistant to apoptosis and (iii) cells migrating away from the toxic debris formed by cellular necrosis, as a consequence of hypoxia and acidosis.

In contrast, the microenvironment surrounding pseudopalisades is fairly better understood. According to the studies (Rong et al. 2006; Wippold et al. 2006; Brat et al. 2004; Martínez-González et al. 2012) the cells made of such structures are mainly hypoxic and have less proliferating capabilities. These cells, however, show increased vascular endothelial growth factor (VEGF) expression, that results in development of microvascular structures (Zagzag et al. 2000; Plate et al. 1992). Due to the dense structure of the brain tissue, this additional vascular growth is very irregular and even results in the formation of glomeruloid bodies. The area enclosed by pseudopalisades is composed of mainly dead cells and other cellular debris forming the necrotic core. Moreover, due to the vascular aberrations there exist anisotropic oxygen gradients, with the center being hypoxic.

As hypoxia is closely associated with acidosis (Brahimi-Horn and Pouysségur 2007; Chiche et al. 2010; Jing et al. 2019), it is well possible that the migratory cells have switched to a glycolytic pathway, which in turn exacerbates the micro-acidity and promotes migratory behavior of the cells (Estrella et al. 2013; Piasentin et al. 2020). Because GBM is the most dominant type of malignant brain tumors (Dolecek et al.
and since the detection of pseudopalisades indicates a worsening condition of a glioma patient (Brat and Mapstone 2003; Kleihues et al. 1995), it is highly important to get insight into the formation and behavior of these structures. For this purpose, mathematical models have proven to be highly effective, especially for understanding and validating the dynamics of biological processes. In the context of GBM, various types of models have been proposed. Nice reviews on the chronological evolution of such models can be found in Hatzikirou et al. (2005), Harpold et al. (2007), Martirosyan et al. (2015), Alfonso et al. (2017). Broadly speaking, there are mainly two classes of models: discrete and continuous. The former mainly comprise rule-based computational models (e.g. Sander et al. 2002; Khain et al. 2011; Böttger et al. 2012; Kim and Roh 2013) that try to identify the self-organization behavior of the cells. These models take advantage of the computational power to explore different rule-configurations that could explain the phenomena. On the other hand, continuous models are based on continuous abstraction of the evolution of physical processes. The most simple, yet effective continuous models are ODE based. They not only employed to study the proliferating capabilities of glioma (Sturrock et al. 2015), but also for assessing the effects of radio- and chemotherapy (Yu et al. 2021). However, when one is interested in studying the invasive behavior of GBMs, space becomes important, thus spatial dynamics needs to be taken into account. Most of the cancer invasion models, inspired by the early works of Murray (2002), are based on reaction-diffusion based settings, see e.g., (Jbabdi et al. 2005; Swanson et al. 2011; Hatzikirou et al. 2012; Martínez-González et al. 2012; Kim and Roh 2013; Alfonso et al. 2016). They only consider movement based on random motion with very limited ability to incorporate direction/orientation information from the microenvironment, e.g. only via some anisotropic diffusion coefficient. These models were generalized in Hinow et al. (2009), Kim et al. (2009), Colombo et al. (2015), where advection/taxis terms were introduced to incorporate the relevant microenvironment information such as tissue structure, vasculature etc. Because cancer growth and spread is a complex multiscale process, mere macroscopic models fail to ilicit the outcomes of cross-scale interactions. Many cellular motion models originate at the subcellular or cellular scale by first considering the dynamics of individual cells followed by modeling the interactions with other cells and physical/chemical components of the environment. This is then upscaled to the tissue level, where experimental observations are possible. Such multiscale framework has been considered in Painter and Hillen (2013), Engwer et al. (2015, 2016a, b), Hunt and Surulescu (2016), Corbin et al. (2018), Conte et al. (2020), Conte and Surulescu (2021), Corbin et al. (2021), Dietrich et al. (2022), Conte et al. (2022) to study the invasive patterns of glioma. Thereby, mainly parabolic scaling is used to obtain a corresponding macroscopic PDE which consequently involves diffusion and reaction coefficients that are coupled with the dynamics of the lower scales. In contrast to the deterministic models, authors in Hiremath and Surulescu (2015, 2016, 2017), Hiremath et al. (2018) have used stochastic multiscale settings to ilicit transient invasive patterns of cancer. The kind of models considered so far are in some sense phenomenological descriptions that try to explain or justify in vivo or in vitro observations. This is a bottom-up approach, where theoretical reasoning is used to explain the observed data. In contrast,
one could resort to statistical techniques to infer relevant properties of the dynamics directly from the data without considering any biophysical model. Alternatively, one could simply complement the phenomenological model by statistically incorporating the observed data. Such models are called data-driven, where the data and the (bio)physics model are coupled through an optimization formulation. This type of inverse problem formulation has been used e.g., in Hogea et al. (2008), Konukoglu et al. (2010), Gholami et al. (2016) to estimate patient specific model parameters that can subsequently be used for making predictions. Similar to these approaches, in this paper we formulate an optimal control problem with the aim of gaining insight into the dynamical processes responsible for generating a specific type of pseudopalisade patterns. Unlike previous studies (Kim et al. 2009; Caiazzo and Ramis-Conde 2015; Martínez-González et al. 2012; Kumar et al. 2021, 2022) where a more or less phenomenological approach was employed, we adopt a data-driven approach where all model unknowns along with the involved model parameters are estimated from the data itself. Starting from a macroscopic model (Kumar et al. 2021), which in turn is obtained using a multiscale modeling technique, given some arbitrary initial condition and a target pseudopalisade pattern, we compute the optimal model parameters such as growth rate, diffusion coefficient, taxis direction, such that the initial tumor density optimally evolves to the final pseudopalisade pattern. The advantages of this approach are the following:

1. Given some fixed arbitrary initial condition $u_0$ and different target (final) patterns $(O_k)_{k \in \mathbb{N}}$, we can compute corresponding optimal model parameters $\theta_{O_k}$ and solutions $u_{O_k}$ (see (13)) which are able to directly explain the data. By analyzing the qualitative properties of the obtained parameters it is possible to gain insight into their interactions that eventually result in the formation of the observed structures. Furthermore, this approach also provides a way to directly compare the differences in the model parameters, and thereby also the internal microscopic dynamics, which eventually resulted in different end patterns.

2. By reversing the initial and target (final) conditions, we can identify the optimal parameters that can reverse or undo the developed pattern. A typical application of this would be in developing strategies to renormalize or neutralize the tumor microenvironment with the aim of reducing the malignancy of the developed tumor. In cases where direct intervention on the tumor is conceivable, appropriate additional parameterized equations (based on the type of intervention) can be introduced with the aim of stopping further progression of the pattern.

3. By combining the parameters $\theta_{O_k}$ which are responsible for generating simple target patterns $O_k$, in order to obtain a new parameter vector $\theta_{O'}$, we can synthesize new unseen patterns $O'$. E.g., we can define $\theta_{O'} := \sum_{k=1}^{N} c_k \theta_{O_k}$, for $N \in \mathbb{N}$, $c_k \in \mathbb{R}$, and simulate the dynamics to synthesize a new pattern $O'$. The main application of this would be that, if therapy strategies can be designed leading to simpler pseudopalisade patterns, then a similar linear combination of interventions could, putatively, also work in neutralizing complex pseudopalisade patterns. Whether this is useful for conceiving new therapy approaches remains arguable, however it can help understanding the histological patterns.
Based on the above discussion, the rest of the document is organized in the following manner. In Sect. 2 we present a multiscale mathematical model for acid modulated cancer dynamics and we formulate a terminal optimal control problem (TOCP), (13), for formation of the observed pseudopalisade pattern. In Sect. 3.2 we establish the wellposedness of the model (Theorem 5). Following that, in Sect. 3.3 we establish the existence of a solution to (13) (Theorem 8). To find a suitable minimizing sequence via a first order gradient descent method, we first establish the existence of the gradient of the objective functional w.r.t to the parameter $\theta$ (Theorem 9) and Lemma 10 and show its continuity property in Theorem 11. Next, in Theorem 12 we show that the minimizing sequence generated by Algorithm 1 is indeed the minimizing sequence for the formulated optimization problem (13). Subsequently, in Sect. 4 we implement the algorithm and investigate its performance and results. Following this, in Sect. 5 we not only discuss the application of TOCP to therapy problems, but also for the synthesis of new unseen patterns along with its plausible value for interventions. Finally, in Sect. 6 we discuss the results and draw conclusions.

2 Modeling

The goal of this section is to set up a system of equations that is able to mimick the complex interactions of cancer cells with their host tissue. Because we are interested in analyzing the influence of tissue acidity on the type of glioma patterns that emerge, we restrict the description to mainly the interactions between cancer cells and protons in the extracellular region. The latter is modeled by accounting for the dynamics of extracellular proton concentration $H$. Since protons are much smaller than cancer cells, their dynamics is much faster, thus it can be considered directly at the macroscopic level. Following this reasoning, the evolution of acid is described by the following reaction-diffusion equation:

$$\frac{\partial H}{\partial t} = D_s \Delta H - \alpha H + \beta f_2(C, H),$$

where $D_s$ is the effective proton diffusion coefficient, $\alpha$ is the effective acid removal rate by vasculature, and $\beta$ represents the effective expulsion rate of protons by cancer cells mainly as a byproduct of the glycolytic energy cycle (Gatenby and Gawlinski 2003). Here $f_2(C, H) := \frac{CH}{(1+C^2+H^2)^2}$ models the efflux of protons by cell membrane transporters such as MCT, NHE (Webb et al. 1999, 2004). Since the activity of these membrane transporters is dependent on the interaction between intra-/extracellular proton concentration, by approximating the concentration of intracellular protons to be proportional to the cell density $C$, the activity of the transporter is modelled as the interaction between the cancer cells and extracellular protons, resulting in the numerator term $CH$. However, since the activity of membrane transporters saturates with increasing ion concentrations, we introduce the denominator term to capture this behavior. Lastly, we note that the coefficients $\alpha$ and $\beta$ are considered to be functions of space and time.

On the other hand, the dynamics of cancer cells is much slower, so it can be modelled not only by considering intracellular events, but also transmembrane and extracellular
interactions. This basically results in multiscale modeling of tumor evolution, which for the case of GBM has been done previously by several authors Painter and Hillen (2013), Engwer et al. (2015), Engwer et al. (2016b), Hunt and Surulescu (2016), Swan et al. (2018). For our study, we refer to the more recent works (Kumar et al. 2021; Kumar and Surulescu 2020), particularly to the former, where the activity of proton-specific transmembrane units was considered to deduce a kinetic transport equation for the evolution of tumor density which was subsequently upscaled. The parabolic scaling procedure resulted in a myopic-diffusion-based PDE which not only translates the averaged random fluctuations at the microscopic level to the macroscopic one, but also adequately incorporates the advection term representing directed movement of cells. The resulting parabolic PDE in the non-dimensionalized form reads:

$$\partial_t C = \nabla \cdot (\nabla \cdot (\mathbb{D} C)) + \nabla \cdot (\delta(H)C \nabla H) + \mu C (1 - C)(1 - H), \quad (2)$$

where $C$ is the density of cancer cells, $\mathbb{D}$ is the anisotropic diffusion tensor, $\delta(H)$ is the pH taxis coefficient, and $\mu$ is the proliferation rate. For our study, we consider a slightly modified version of the macroscopic equation, given as:

$$\partial_t C = \nabla \cdot (\sigma(t, x) \nabla C + C \nabla \kappa(t, x)) + \nabla \cdot (\delta(t, x) C \nabla H) + \mu(t, x) f_1(C, H), \quad (3)$$

where we have reduced the diffusion tensor $\mathbb{D}$ and pH taxis coefficient $\delta(H)$ to space-time functions $\sigma$ and $\delta$, respectively. The growth term is modified to a bounded function $f_1(C, H)$ and the rate constant $\mu$ is taken be a space-time function. The intrinsic proliferation potential of the cell population is modeled by the logistic growth term. The negative effect on cell proliferation due to excess extracellular acidity (i.e. protons $H$) is modeled via the multiplicative term $(1 - H)$, assuming $H$ is already expressed as a normalized object. Altogether, the numerator $C(1 - C)(1 - H)$ models the acid modulated proliferation potential of cancer population. The denominator term serves the following purposes: (i) the $C^2$ term slows down the rate of proliferation at regions of high cell density, (ii) the $H^2$ term represents saturation of the membrane transporters/receptors with increasing $H$ concentration which consequently limits the magnitude of the rate of acid induced cell death. Additionally, we have introduced the advection term $C \nabla \kappa$ to model haptotactic movement described by a tissue-dependent time-varying function $\kappa$ (to be estimated). Let $T > 0$ and $I = (0, T] \subset \mathbb{R}_+$ be a finite time interval. Let $\mathcal{D} \subset \mathbb{R}^2$ be a bounded spatial domain with sufficiently smooth boundary. We assume there is no flux of cells or protons through the boundary. The resulting coupled PDE system is given by the following initial boundary value problem (IBVP):

$$\partial_t C = \nabla \cdot (\sigma \nabla C + C \nabla \kappa + \nabla \cdot (\delta C \nabla H) + \mu f_1(C, H)) \quad \text{in } (0, T] \times \mathcal{D} \quad (4a)$$

$$\partial_t H = \Delta H - \alpha H + \beta f_2(C, H) \quad \text{in } (0, T] \times \mathcal{D} \quad (4b)$$

$$C(0) = C_0, \quad H(0) = H_0 \quad \text{in } \mathcal{D}$$

$$0 = (\sigma \nabla C + C \nabla \kappa) \cdot \hat{n} \quad \text{on } [0, T] \times \partial \mathcal{D}$$

$$0 = \nabla H \cdot \hat{n} \quad \text{on } [0, T] \times \partial \mathcal{D}.$$
The non-dimensionalized PDE system (4) serves as the abstract macroscopic model for the underlying dynamics for the evolution of pseudopalisades in GBM under the influence of acidity. Due to the nonlinear coupling of the reaction terms and interplay between different taxis terms, the resulting dynamics of GBM can be very complex and most importantly very much dependent on the qualitative and quantitative properties of the model coefficient functions (model parameters). As a result, accurately determining the model parameters for a specific observed dynamics can be very challenging. The usual way is to look for stationary solutions by means of linear stability and bifurcation analysis. This process, although very effective during the modeling phase to gain analytic insight, is, however, often unable to explain real and interesting experimental observations which are usually very complicated. In order to explain each observation accurately, it is usually required to formulate an inverse problem, which is in fact the paradigm of this paper. To this end, starting from (4) we formulate a minimization problem whose goal is to determine the optimal parameters for the model such that the final state of the tumor closely matches the real observations. This is realized by devising an optimal control problem (OCP) for which the objective function is based on the final spatial distribution of the tumor, hence it is termed as the terminal optimal control problem (TOCP). In the following section we shall first establish wellposedness of the dynamical model and then present the corresponding TOCP for which we prove the existence of a minimizer which then paves the way for performing data based analysis.

3 Analysis

3.1 Assumptions and prerequisites

Let \( I = (0, T] \subset \mathbb{R}_+ \) be a finite time interval and \( \mathcal{D} \subset \mathbb{R}^2 \) be an open bounded spatial domain with sufficiently smooth boundary. Letting \( H^2_N := \{ u \in H^2_p(\mathcal{D}) : \frac{\partial u}{\partial y} = 0 \} \), we use the following notations for the common Lebesgue and Sobolev spaces:

\[
L^p := L^p(\mathcal{D}; \| \cdot \|_{L^p}), \quad L^p_T := L^p([0, T] \times \mathcal{D}; \| \cdot \|_{L^p}), \quad L^p_T := L^p([0, T]; L^p(\mathcal{D})),
\]

\[
V := (L^2(\mathcal{D}), \langle \cdot, \cdot \rangle), \quad Z^2 := H^2_p(\mathcal{D}), \quad Z := Z^1 := H^1(\mathcal{D}), \quad W := H^1(\mathcal{D}), \quad W := W \times W,
\]

\[
Z^2(T) := L^2(I; Z^2), \quad V := V \times V, \quad Z := Z^1 \times Z^2, \quad Z(T) := Z^1(T) := L^2(I; Z),
\]

\[
W(T) := I^2(I; W), \quad V(T) := I^2(I; V), \quad Z(T) := I^2(I; Z), \quad W(T) := I^2(I; W),
\]

\[
Y := L^\infty(I \times \mathcal{D}), \quad Y(T) := L^\infty(I; Z), \quad \mathcal{C}(T) := C(I; Z),
\]

\[
X := \{ u \in L^2(I; W) : u' \in L^2(I; W') \}, \quad \mathbf{X} := \{ u \in L^2(I; W) : u' \in L^2(I; W') \}, \quad (5)
\]

with \( W' \) denoting the dual space of \( W \). Also, we denote the space of linear operators from \( U \) to \( U' \times Z \) by \( L(U; U' \times Z) \).

Finally, we define the solution space \( U \) and the parameter space \( \Theta \) as

\[
U := X \cap Y(T) \cap Z(T), \quad \text{and} \quad \Theta := \left( L^2(I; Z^2), \langle \cdot, \cdot \rangle_\Theta \right). \quad (6)
\]

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3.1.1 Formulation of the data-driven model

The data-driven model comprises two main components: the first one being the model for the dynamics of the system and the second being the objective/cost functional that couples the observation data with the system dynamics. The former is described by (4) which when coupled with a terminal type objective functional results in a terminal valued optimal control problem (TOCP). We shall now recast (4) in an abstract form so that it enables TOCP to be represented in a way that is conducive for mathematical analysis. To this end, let $\mathbf{u} = (u_1, u_2)^T$ represent the cancer density $C$ and extracellular acidity $H$, then Eq. (4) can be rewritten as

$$
\begin{align*}
\partial_t u_1 - \nabla \cdot (\sigma \nabla u_1 + u_1 \nabla k) &= \nabla \cdot (\delta u_1 \nabla u_2) + \mu f_1(u_1, u_2) \quad \text{in } (0, T) \times \mathcal{D} \quad (7a) \\
\partial_t u_2 - \Delta u_2 + \alpha u_2 &= \beta f_2(u_1, u_2) \quad \text{in } (0, T) \times \mathcal{D} \quad (7b) \\
\end{align*}
$$

where $f_1(u_1, u_2) := \frac{u_1 (1-u_1) (1-u_2)}{1+u_1^2 + u_2^2}$, $f_2(u_1, u_2) := \frac{u_1 u_2}{1+u_1^2 + u_2^2}$.

Letting $\hat{f}_1 := f_1 + \mu u_1$, the weak formulation of (7) is given as:

$$
\begin{align*}
(\partial_t u_1, \varphi) + (\sigma \nabla u_1, \nabla \varphi) + (\mu u_1, \varphi) + (\delta u_1 \nabla u_2, \nabla \varphi) &= (\mu \hat{f}_1, \varphi) - (u_1 \nabla k, \nabla \varphi), \quad (8a) \\
(\partial_t u_2, \psi) + (\nabla u_2, \nabla \psi) + (\alpha u_2, \psi) &= (\beta f_2, \psi), \\
\end{align*}
$$

$\forall \varphi, \psi \in W$, and $t \in (0, T)$. Let $\mathbf{A} : W \to \mathcal{L}(W; W')$ and $\mathbf{r} : W \to W'$ be linear and nonlinear operators, respectively, which for $\mathbf{u}, \mathbf{v}, \mathbf{w} \in W$ are defined as

$$
\begin{align*}
\mathbf{A}(\mathbf{w}) := \begin{bmatrix} A_1(w_1) & A_2(w_1) \\ 0 & A_3(w_2) \end{bmatrix}, \quad & \quad \mathbf{r}(\mathbf{u}) := \begin{bmatrix} r_1(\mathbf{u}) + r_2(\mathbf{u}) \\ r_3(\mathbf{u}) \end{bmatrix} \\
A_1(w_1) u_1, v_1 := (\sigma \nabla u_1, \nabla v_1) + (\mu u_1, v_1), \quad & \quad (r_1(\mathbf{u}), \mathbf{v}) := (\mu f_1(u_1, u_2) + \mu u_1, v_1).
A_2(w_1) u_2, v_1 := (\delta w_1 \nabla u_2, \nabla v_1), \quad & \quad (r_2(\mathbf{u}), \mathbf{v}) := (-u_1 \nabla k, \nabla v_1).
A_3(w_2) u_2, v_2 := (\nabla u_2, \nabla v_2) + (\alpha u_2, v_2), \quad & \quad (r_3(\mathbf{u}), \mathbf{v}) := (\beta f_2(u_1, u_2), v_2).
\end{align*}
$$

Then, $\forall \varphi := (\varphi_1, \varphi_2)^T \in W$, the above weak formulation can be rewritten in the more compact form

$$
\begin{align*}
(\partial_t u_1, \varphi_1) + (A_1(u_1) u_1, \varphi_1) + (A_2(u_1) u_2, \varphi_1) &= (r_1(\mathbf{u}) + r_2(\mathbf{u}), \varphi_1) \quad t \in (0, T) \quad (10a) \\
(\partial_t u_2, \varphi_2) + (A_3(u_2) u_2, \varphi_2) &= (r_3(\mathbf{u}), \varphi_2) \quad t \in (0, T) \quad (10b) \\
\mathbf{u}(0) &= \mathbf{u}_0.
\end{align*}
$$

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Based on the definition of $A$, the corresponding trilinear form $a : W \times W \times W \to \mathbb{R}$ can be defined in the following way:

$$a(w)[u, v] := (A(w)u, v)_W = (A_1(w_1)u_1, v_1) + (A_2(w_1)u_2, v_1) + (A_3(w_2)u_2, v_2)$$

$$= a_1(w_1)[u_1, v_1] + a_2(w_1)[u_2, v_1] + a_3(w_2)[u_2, v_2] \quad (11)$$

$$= a_1(w)[u, v] + a_2(w)[u, v] + a_3(w)[u, v].$$

Let the parameters appearing in (7) be represented by the vector function $\theta$ defined as $\theta := (\theta_1, \ldots, \theta_6)^\top = (\sigma, \kappa, \delta, \alpha, \beta, \mu)^\top$. Based on this, we can now formulate the TOCP. First, let the objective (or cost) functional $J$ be defined as

$$J(u, \theta) = \frac{1}{2} \|u_1(T) - O\|^2_{L^2} + \frac{\lambda}{2} \|\theta\|^2_\chi,$$

where $\lambda \in \mathbb{R}$ and $O \in H^1(\mathcal{D})$ is the image data of the observed pseudopalisade pattern. The aim of the TOCP is to find $u \in U$ and $\theta \in \Theta$ such that $u, \theta$ satisfy the state equation $G(u, \theta) = 0$ while minimizing the functional $J$. Altogether, it results in the following minimization problem: find $(u^*, \theta^*)$ with $u^* := u(\theta^*)$ and

$$u^*, \theta^* = \operatorname{argmin}_{u \in U, \theta \in \Theta} J(u, \theta) \quad \text{s.t.} \quad G(u; \theta) = 0, \quad \text{with } u \in U \text{ and } \theta \in \Xi,$$

where the equality constraint $G(u; \theta) = 0$ represents the system dynamics specified by (8). The mapping $G : U \times \Xi \to U' \times Z$ reads as

$$G(u, \theta) = \begin{pmatrix}
(\partial_t u_1, \cdot) + (\sigma \nabla u_1, \nabla \cdot) + (u_1 \nabla \kappa, \nabla \cdot) + (\delta u_1 \nabla u_2, \nabla \cdot) - (\mu f_1, \cdot) \\
(\partial_t u_2, \cdot) + (\nabla u_2, \nabla \cdot) + (\alpha u_2, \cdot) - (\beta f_2, \cdot)
\end{pmatrix}
$$

$$u(0) - u_0. \quad (14)$$

Letting $\|\cdot\|_{\Xi} := \|\cdot\|_{C(\bar{I}; Z^6)}$, the subspace $\Xi \subset \Theta$ is the set of admissible parameters defined as

$$\Xi := \left\{ \theta \in \Theta \cap C(\bar{I}; Z^6) : \theta \in C(\bar{I}; Z^6), \|\theta\|_{\Xi} \leq M_\theta \right\}.$$

**Remark 1** It is actually sufficient to define $\Xi$ as

$$\Xi := \left\{ \theta \in \Theta \cap C(\bar{I}; Z^6) : \theta_i \in C(\bar{I}; Z), \|\theta_i\|_{C(\bar{I}; Z)} \leq M_{\theta_i} \leq M_\theta, \quad i \in \{1, 2, 3\} \right\}.$$

For the sake of notational simplicity we shall avoid it here, since it does not bring any major difference in the analysis.
3.2 Model wellposedness

In this section we look at the wellposedness of the system and investigate the existence of the optimal solution. For this purpose we first introduce the following general assumptions.

3.2.1 Assumptions on $f_1, f_2$ and $\theta$

1. $f_1, f_2 \in C^\infty$, and $f_1(u, v), f_2(u, v) \in W \cap L^\infty(\Omega)$ whenever $u, v \in W, t \in I$.
2. $\partial_u f_1, \partial_v f_2$ are elements of $W \cap L^\infty(\Omega)$ for $u, v \in W, t \in I$. Thus $\|f_j\|_{C(\overline{I}, W)} \leq M_{f_j}$ for $j \in \{1, 2\}$.
3. In particular, for our application we have that $f_1(u, v) := \frac{u(1-u)(1-v)}{(1+u^2+v^2)^2}$ and $f_2(u, v) := \frac{uv}{(1+u^2+v^2)^2}$, which satisfy the above conditions.
4. The model parameter functions $\theta_i \in \{\sigma, \kappa, \delta, \alpha, \beta, \mu\}$ are such that:

   $0 < m_{\theta_i} \leq \theta_i(t, x) \leq M_{\theta_i} < \infty$, for all $t \in I, x \in \Omega$. Additionally, it is also assumed that $\|\theta_i\|_{C(\overline{I}, Z)} \leq M_{\theta_i}$.

3.2.2 Energy estimates of solutions

Lemma 1 Let $u = (u_1, u_2) \in Z$ satisfy the Eq. (7). Then its components fulfill the following energy estimates:

$$\|u_1\|^2_{W(T)} \leq k_u \|u_{1,0}\|^2_W \quad \text{and} \quad \|u_2\|^2_{W(T)} \leq k_u \|u_{2,0}\|^2_W,$$

with $k_u(T)$ and $k_v(T)$ appropriate constants.

Proof Multiplying by $u_2$ both sides of (7b) we get

$$\frac{1}{2} \frac{d}{dt} \int_{\Omega} u_2^2 + \int_{\Omega} |\nabla u_2|^2 + \alpha \int_{\Omega} u_2^2 \leq M_\beta M_{f_2} \|u_2\|^2_{W(T)} \Rightarrow \|u_2\|^2_{W(T)} \leq k_u(T) \|u_{2,0}\|^2_W \leq M_{u_2}(T),$$

with sufficiently large $M_{u_2}(T)$.

Similarly, multiplying by $u_2'$ both sides of (7b) we get

$$2 \|u_2'\|^2_{\overline{V}(T)} + \frac{1}{2} \frac{d}{dt} \int_{\Omega} |\nabla u_2|^2 + \alpha \frac{d}{dt} \|u_2\|^2_{\overline{V}} \leq M_\beta M_{f_2} \frac{d}{dt} \|u_2\|^2_{\overline{V}} \leq M_\beta M_{f_2} k_{u_2} \|u_{2,0}\|^2_W \Rightarrow \|u_2'\|^2_{\overline{V}(T)} + \sup_{t \in [0, T]} \|\nabla u_2\|^2_{V} \leq (1 + M_\beta M_{f_2} k_{u_2}) \|u_{2,0}\|^2_W \leq M_{u_2}(T).$$

Lastly, the second derivative can be bounded from above as follows:

$$\|\Delta u_2\|_{V} \leq (M_\alpha + M_\beta M_{f_2}) \|u_2\|_{V} + \|u_2'\|_{V}$$
\[
\begin{align*}
\leq (M_\alpha + M_\beta M_{f_2})[\|u_2\|_V + \|u_2'\|_V] \\
\leq (M_\alpha + M_\beta M_{f_2})(2k_{u_2} + 1 + M_\beta M_{f_2}k_{u_2} T)\|u_{2,0}\|_W \\
\leq M_{u_2}(T). \quad (17)
\end{align*}
\]

Altogether, we get that \(u_2 \in L^\infty([0, T]; Z)\) and \(u_2' \in L^2(I; V)\). Additionally, differentiating (7b) and multiplying with \(u_2'\) and using (18) below and boundedness of \(\partial_{u_1} f_2, \partial_{u_2} f_2\) we can get that \(u_2 \in H^1([0, T]; W)\).

Now let us consider Eq. (7a). Like above, multiplying by \(u_1\) both sides of the equation we get:

\[
\begin{align*}
\frac{1}{2} \frac{d}{dt} \int_\mathbb{D} u_1^2 + m_\sigma \int_\mathbb{D} |\nabla u_1|^2 \\
\leq (M_\kappa + M_\delta \|u_2\|_{L^\infty}) \int_\mathbb{D} u_1 \cdot \nabla u_1 + 2M_\mu M_{f_1} \int_\mathbb{D} \frac{1}{2} u_1^2 \\
\leq \frac{(M_\kappa + M_\delta \|u_2\|_{L^\infty})^2}{2m_\sigma} \|u_1\|^2 + \frac{m_\sigma}{2} \|\nabla u_1\|^2 + 2M_\mu M_{f_1} \frac{1}{2} \|u_1\|^2 \\
\Rightarrow \frac{1}{2} \frac{d}{dt} \int_\mathbb{D} u_1^2 + \frac{m_\sigma}{2} \int_\mathbb{D} |\nabla u_1|^2 \leq \left[ \frac{(M_\kappa + M_\delta \|u_2\|_{L^\infty})^2}{2m_\sigma} + 2M_\mu M_{f_1} \right] \frac{1}{2} \|u_1\|^2 \\
\Rightarrow \|u_1\|_{W(T)}^2 \leq \exp(k_{u_1} T) \|u_{1,0}\|_W^2 \text{ and } \|\nabla u_1\|_{L^2}^2 \leq \exp(k_{u_1} T) \|u_{1,0}\|_W^2 \\
\Rightarrow \|u_1\|_{W(T)}^2 \leq \exp(k_{u_1} T) \|u_{1,0}\|_W^2.
\end{align*}
\]

Now multiplying by \(u_1'\) both sides of (7a) we get

\[
\begin{align*}
\|u_1'\|^2 + \frac{m_\sigma}{2} \frac{d}{dt} \|\nabla u_1\|^2 \leq (M_\kappa + M_\delta M_{u_2})(u_1, \nabla u_1') + 2M_\mu M_{f_1} \frac{1}{2} \frac{d}{dt} \|u_1\|^2 + M_\sigma \|\nabla u_1\|^2 \\
\leq (M_\kappa + M_\delta M_{u_2})(u_1, \nabla u_1') + (2M_\mu M_{f_1} + M_\sigma) \exp(k_{u_1} T) \|u_{1,0}\|_W^2 \quad (18)
\end{align*}
\]

Using integration by parts for the time derivative i.e.

\[
\int_0^T \int_\mathbb{D} u_1 \cdot \nabla u_1' = \int_\mathbb{D} \int_0^T u_1 \cdot \nabla u_1' = \int_\mathbb{D} (u_1 \cdot \nabla u_1)|_0^T - \int_0^T \int_\mathbb{D} u_1' \cdot \nabla u_1,
\]

we get that \((M_\kappa + M_\delta M_{u_2}) \int_0^T (u_1, \nabla u_1')\) satisfies the following inequalities.

\[
\begin{align*}
\leq (M_\kappa + M_\delta M_{u_2}) \left( \epsilon \|u_1'\|^2_{V(T)} + \frac{1}{4\epsilon} \|\nabla u_1\|^2_{V(T)} + \frac{1}{2} \|u_{1,0}\|_V^2 + \|\nabla u_{1,0}\|_V^2 \right) \\
+ \|u_{1,T}\|^2_{V} + \|\nabla u_{1,T}\|^2_{V} \right) \\
\leq \frac{1}{2} \|u_1'\|^2_{V(T)} + \frac{(M_\kappa + M_\delta M_{u_2})^2}{2} \left( \|\nabla u_1\|^2_{V(T)} + \frac{1}{2} \|u_{1,0}\|_V^2 + \|\nabla u_{1,0}\|_V^2 \right) \\
+ \|\nabla u_{1,0}\|^2_{V} + \|u_{1,T}\|^2_{V} + \|\nabla u_{1,T}\|^2_{V}) \\
\leq \frac{1}{2} \|u_1'\|^2_{V(T)} + \frac{(M_\kappa + M_\delta M_{u_2})^2}{2} \left( \|u_1\|^2_{W(T)} + \frac{1}{2} \|u_{1,0}\|_W^2 + \|u_{1,T}\|_W^2 \right).
\]

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Now plugging this in (18) we get that

$$\|u_1\|_{V(T)}^2 + m_\sigma \sup_{t \in [0,T]} \|\nabla u_1\|_V^2$$

$$\leq (M_k + M_\delta M_{u_2}^2)\left(\|u_1\|_{W(T)}^2 + \frac{1}{2}(\|u_{1.0}\|_W^2 + \|u_{1.}\|_W^2)\right)$$

$$+ (4M_\mu M_{f_1} + 2M_\sigma) \exp(k_{u_1} T)\|u_{1.0}\|_W^2$$

$$\leq \left(2(M_k + M_\delta M_{u_2}^2) + 4M_\mu M_{f_1} + 2M_\sigma\right) \exp(k_{u_1} T)\|u_{1.0}\|_W^2.$$  

Additionally, if $u_1 \in L^\infty([0, T]; V)$, we also have that

$$m_\sigma \|\Delta u_1\|_V \leq M_\sigma \|\nabla u_1\|_V + M_k \|\nabla u_1\|_V + \|u_1\|_V M_k + M_\mu M_{f_1} \|u_1\|_V + \|u_1\|_V$$

$$+ M_\delta (M_{u_2} \|u_1\|_V + M_{u_2} \|\nabla u_1\|_V + \|u_1\|_{L^\infty} \|\Delta u_2\|_V)$$

$$\leq (M_\sigma + M_k + M_\mu M_{f_1} + M_\delta M_{u_2})\|u_1\|_W + M_\delta \|u_1\|_{L^\infty} \|\Delta u_2\|_V + \|u_1\|_V$$

$$\leq (M_\sigma + M_k + M_\mu M_{f_1} + M_\delta M_{u_2}) \exp(k_{u_1} T)\|u_{1.0}\|_V$$

$$+ M_\delta M_{u_2} \|u_1\|_{L^\infty} + \|u_1\|_V.$$

\[\square\]

### 3.2.3 Non-negativity of solutions

**Lemma 2** Let $u = (u_1, u_2) \in Z$ satisfy system (7). Then $u(t) \geq 0$ for all $t \in I$ if $u_0 \geq 0$.

**Proof** Let $f_1 := u_1 \tilde{f}_1$ with $\tilde{f}_1 := \frac{(1-u_1)(1-u_2)}{(1+u_1^2+u_2^2)^2}$, $q(u_1) := \begin{cases} \frac{1}{2}u_1^2 & \text{if } u_1 \in (-\infty, 0) \\ 0 & \text{else} \end{cases}$. Then the function $Q(t) = \int_{\mathcal{D}} q(u_1(t))dx$ is continuously differentiable. Its derivative (using (7a)) is given by:

$$Q'(t) = \int_{\mathcal{D}} q'(u_1) \nabla \cdot (\sigma \nabla u_1 + u_1 \nabla \kappa) + \int_{\mathcal{D}} q'(u_1) \nabla \cdot (\delta u_1 \nabla u_2) + \int_{\mathcal{D}} q'(u_1) \mu u_1 \tilde{f}_1$$

$$= -\int_{\mathcal{D}} \nabla q'(u_1) \cdot \nabla u_1 - \int_{\mathcal{D}} \nabla \kappa \cdot u_1 \nabla q'(u_1) - \int_{\mathcal{D}} \nabla q'(u_1) \cdot \delta u_1 \nabla u_2$$

$$+ \mu \int_{\mathcal{D}} q'(u_1) u_1 \tilde{f}_1$$

$$\leq -\int_{\mathcal{D}} \sigma \nabla q'(u_1) \cdot \nabla u_1 + M_k \int_{\mathcal{D}} |\nabla q'(u_1)| \cdot |u_1| + M_\delta \int_{\mathcal{D}} |\nabla q'(u_1)| \cdot u_1 \nabla u_2$$

$$+ M_\mu M_{f_1} \int_{\mathcal{D}} q'(u_1) u_1$$

$$\leq -m_\sigma \int_{\mathcal{D}} |\nabla q'(u_1)|^2 + M_k \int_{\mathcal{D}} \epsilon_1 |\nabla q'(u_1)|^2 + \frac{M_k}{4\epsilon_1} \int_{\mathcal{D}} u_1^2 + M_\delta \int_{\mathcal{D}} \epsilon_2 |\nabla q'(u_1)|^2$$

$$+ \frac{M_\delta}{4\epsilon_2} \int_{\mathcal{D}} u_1^2 |\nabla u_2|^2 + M_\mu M_{f_1} \int_{\mathcal{D}} q'(u_1) u_1 \quad \text{(using Young’s inequality)}.$$
\[ \leq \frac{M_k^2}{m_\sigma} \int_{\mathcal{D}} \frac{1}{2} u_1^2 + \frac{M_\delta^2 M_{\sigma}^2}{m_\sigma} \int_{\mathcal{D}} \frac{1}{2} u_1^2 + M_\mu M_{f_1} \int_{\mathcal{D}} \frac{1}{2} q'(u_1) u_1 \]

\[ \leq \frac{M_k^2}{m_\sigma} + \frac{M_\delta^2 M_{\sigma}^2}{m_\sigma} + M_\mu M_{f_1} \int_{\mathcal{D}} q(u_1) := k(t) Q(t) \]

\[ \Rightarrow Q(t) = 0 \quad \forall t \geq 0 \quad \text{using Gronwall's inequality.} \]

Similarly, letting \( q(u_2) \) and using the above result that \( u_1 \geq 0 \), we get that \( Q'(t) \leq M_\alpha Q(t) \). This in turn (due to Gronwall) implies that \( Q(t) = 0 \) for all \( t \geq 0 \).

**Lemma 3** Let \( u \) be a solution to Eq. (7). Then \( u \in L^\infty(I \times \mathcal{D}) \).

**Proof** In order to prove uniform boundedness of \( u \) in \( \mathcal{D}_T := [0, T] \times \mathcal{D} \), following the approach of Finotti et al. (2012), we partition \( \mathcal{D}_T \) along the time axis and show that the increment of the magnitude of \( u \) over these different partitions tends to zero. To this end, let the finite time sequence \((t_i)_i \), with \( i \in \{0, \ldots, K\} \) and \( K \in \mathbb{N} \) finite, represent the partition of \([0, T]\). Correspondingly, let \( \mathcal{D}_{t_i} := [t_{i-1}, t_i] \times \mathcal{D} \) represent the \( i^{th} \) partition of the time-partitioned space-time cylinder. Let \( u := (u, v) \) and let the norm for the time-continuous \( H^1(\mathcal{D}) \) valued functions be denoted as

\[ \|w\|_{C_{t_i}^1(\mathcal{D})} := \sup_{t \in [t_{i-1}, t_i]} \|w(t)\|_{H^1(\mathcal{D})}. \]

Based on the energy estimate (15) we have that \( \|u\|_{C_{t_i}^1(\mathcal{D})} \leq k_{u_1} \) and \( \|v\|_{C_{t_i}^1(\mathcal{D})} \leq k_{u_2} \) for every \( i \in \mathbb{N} \). Now let \( u_k = \max\{u - k, 0\} \) for \( k > k_0 := \max\{1 + \epsilon, \|u_0\|_{L^\infty}\} \). Correspondingly, the supporting sets of \( u_k \) are denoted by \( \mathcal{D}_{u_k}(t) := \{x \in \mathcal{D} : u(t, x) > k\} \) and \( \mathcal{D}_{t_i}(k) := \{(t, x) \in \mathcal{D}_{t_i} : u(t, x) > k\} \) for each \( i \in \mathbb{N} \). Now, testing (7a) with \( u_k \) and letting \( f_1 := u \tilde{f}_1 \) we get

\[ \frac{1}{2} \frac{d}{dt} \int_{\mathcal{D}} u_k^2 + m_\sigma \int_{\mathcal{D}} |\nabla u_k|^2 \leq -\int_{\mathcal{D}} u \nabla k \cdot \nabla u_k - \int_{\mathcal{D}} \nabla u_k \cdot \delta u \nabla v + M_\mu \int_{\mathcal{D}} u_k f_1 \]

\[ \leq \int_{\mathcal{D}_{u_k}} |\nabla u_k \cdot \nabla k| + \int_{\mathcal{D}_{u_k}} |\nabla u_k \cdot \delta u \nabla v| + M_\mu \int_{\mathcal{D}_{u_k}} u_k f_1 \]

\[ = \int_{\mathcal{D}_{u_k}} |\nabla u_k \cdot \nabla k| + \int_{\mathcal{D}_{u_k}} |\nabla u_k \cdot \delta u \nabla v| + M_\mu \int_{\mathcal{D}_{u_k}} u_k f_1 \]

\[ \leq \int_{\mathcal{D}_{u_k}} \epsilon_1 |\nabla u_k|^2 + \int_{\mathcal{D}_{u_k}} M_\delta^2 u_k^2 + \int_{\mathcal{D}_{u_k}} \epsilon_2 |\nabla u_k|^2 + \int_{\mathcal{D}_{u_k}} M_\mu^2 f_1^2 |\nabla v|^2 \]

\[ + M_\mu M_{f_1} \int_{\mathcal{D}_{u_k}} u_k u \]

\[ \Rightarrow \frac{1}{2} \frac{d}{dt} \int_{\mathcal{D}} u_k^2 + \frac{m_\sigma}{2} \int_{\mathcal{D}} |\nabla u_k|^2 \leq \left[ \frac{M_k^2}{m_\sigma} + \frac{M_\delta^2 u_k^2}{m_\sigma} \right] \int_{\mathcal{D}_{u_k}} u_k^2 + M_\mu M_{f_1} \int_{\mathcal{D}_{u_k}} u_k u \]
\[
\frac{1}{2} \frac{d}{dt} \int_{\mathcal{D}} u_k^2 + \frac{m_\sigma}{2} \int_{\mathcal{D}} |\nabla u_k|^2 \leq \left[ \frac{M_k^2}{m_\sigma} + \frac{M_\delta^2}{m_\sigma} M_{u_2}^2 \right] \int_{\mathcal{D} u_k} u^2 + M_\mu M_{f_1} \int_{\mathcal{D} u_k} u_k u
\]

\[
\leq \left[ \frac{M_k^2}{m_\sigma} + \frac{M_\delta^2}{m_\sigma} M_{u_2}^2 + M_\mu M_{f_1} \right] \int_{\mathcal{D} u_k} (u^2 + u u_k)
\]

\[
\leq 4 \left[ \frac{M_k^2 + M_\delta^2 M_{u_2}^2}{m_\sigma} + M_\mu M_{f_1} \right] \int_{\mathcal{D} u_k} ((u_k)^2 + k^2)
\]

\[
= k(m_\sigma, M_k, M_{u_2}, M_\delta, M_\mu, M_{f_1}) \|u_k\|_{L^2}^2 + k^2 |\mathcal{D} u_k(t)|.
\]

Integrating w.r.t. \( t \in [0, t_1] \), with \( t_1 > 0 \) small enough such that

\[
t_1 \sup_{t \in [0, t_1]} k(m_\sigma, M_k, M_{u_2}, M_\delta, M_\mu, M_{f_1})(t) < 1/2,
\]

we get that

\[
\|u_k\|_{C^1_{\mathcal{D} t_1}}^2 \leq 2k(m_\sigma, M_k, M_{u_2}, M_\delta, M_\mu, M_{f_1})k^2 \eta_k, \quad \text{with } \eta_k = |\mathcal{D} t_1| = \int_0^{t_1} |\mathcal{D} k(t)| dt
\]

\[
\Rightarrow \|u_k\|_{C^1_{\mathcal{D} t_1}} \leq \theta_1 k^{\frac{1}{2}}\eta_k. \tag{19}
\]

Let \( N_0 := n_0 \hat{k} \) for some \( n_0 > 1 \), let \( k_i = N_0(2 - 2^{-i}) \) for \( i \in \mathbb{N}_0 \), then \( \eta_k \) fulfills the following inequality:

\[
\begin{align*}
\frac{1}{2} \left( \int_{\mathcal{D}_{k_i+1}} \frac{1}{2} \int_{\mathcal{D}_{k_i+1}} \frac{1}{2} \right)^2 \left( \int_{\mathcal{D}_{k_i}} k_{i+1} \right) &= \left( \int_{\mathcal{D}_{k_i}} k_{i+1} \right) \left( \int_{\mathcal{D}_{k_i}} \frac{1}{2} \int_{\mathcal{D}_{k_i}} \frac{1}{2} \right)^2 \\
&= \left( \int_{\mathcal{D}_{k_i}} \frac{1}{2} \int_{\mathcal{D}_{k_i}} \frac{1}{2} \right)^2 \left( \int_{\mathcal{D}_{k_i}} k_{i+1} \right) \\
&= \left( \int_{\mathcal{D}_{k_i}} \frac{1}{2} \int_{\mathcal{D}_{k_i}} \frac{1}{2} \right)^2 \left( \int_{\mathcal{D}_{k_i}} k_{i+1} \right)
\end{align*}
\]

\[
\Rightarrow (k_{i+1} - k_i) \eta_k \leq \left( \int_{\mathcal{D}_{k_i}} \frac{1}{2} \int_{\mathcal{D}_{k_i}} \frac{1}{2} \right)^2 \left( \int_{\mathcal{D}_{k_i}} k_{i+1} \right) \quad \text{since } k_i > \hat{k} > 1 \text{ are constants}
\]

\[
\leq \left( \int_{\mathcal{D}_{k_i}} \frac{1}{2} \int_{\mathcal{D}_{k_i}} \frac{1}{2} \right)^2 \left( \int_{\mathcal{D}_{k_i}} k_{i+1} \right) \\
\leq \left( \int_{\mathcal{D}_{k_i}} \frac{1}{2} \int_{\mathcal{D}_{k_i}} \frac{1}{2} \right)^2 \left( \int_{\mathcal{D}_{k_i}} k_{i+1} \right)
\]

\[
\Rightarrow (k_{i+1} - k_i) \eta_k \leq \left( \int_{\mathcal{D}_{k_i}} \frac{1}{2} \int_{\mathcal{D}_{k_i}} \frac{1}{2} \right)^2 \left( \int_{\mathcal{D}_{k_i}} k_{i+1} \right) \leq \|u_k\|_{L^r(\mathcal{D}_{t_1})}. \tag{21}
\]
Using the Sobolev embedding inequality for $2(v + 1) =: r \in [2, \frac{2d}{d-2}]$ with $v > 0$ we have that

$$\|u_k\|_{L_r(\mathcal{D}_1)} \leq \gamma_0 \|u_k\|_{H^1(\mathcal{D})} \leq \gamma_0 \vartheta_1 k_i \eta^\gamma_{k_i}, \quad \forall i \in \mathbb{N}_0, \gamma_0 > 1$$

$$\Rightarrow \eta^\gamma_{k_{i+1}} \leq \frac{\gamma_0 \vartheta_1 k_i}{(k_{i+1} - k_i)} \eta^\gamma_{k_{i+1}} \leq 4 \gamma_0 \vartheta_1 2^i \eta^\gamma_{k_i} \leq 4 \gamma_0 \vartheta_1 2^i (\eta^\gamma_{k_i})^{1+v},$$

$$\forall i \in \mathbb{N}_0, \text{ (since } \frac{v + 1}{r} = \frac{1}{2}).$$

In particular, by taking $k_i := \hat{k}$ and $k_{i+1} := N_0$ and defining $\eta^0 := \eta^\gamma_{k_0}$ we have that $\eta^\gamma_{k_i} \leq \frac{\gamma_0 \vartheta_1}{n_0 - 1} (T|\mathcal{D}|)^{\frac{1}{2}}$. Thus for $n_0 > 1 + \gamma_0 \vartheta_1 (T|\mathcal{D}|)^{\frac{1}{2}} (4 \gamma_0 \vartheta_1)^{\frac{1}{2}} 2^\frac{1}{v}$ we get that $\eta^\gamma_{k_i} \leq (4 \gamma_0 \vartheta_1)^{\frac{1}{2}} 2^\frac{1}{v}$. Thus invoking Theorem 2.4.1 of Zacher (2010) we get that $(\eta^\gamma_{k_i})^{\frac{1}{2}} \to 0$ as $i \to \infty$. In particular, we get that $\eta^\gamma_{k_\infty} := \eta^\gamma_{k_\infty} = 0$. Consequently, we get that $u \leq c_1 := k_\infty = 2 \eta_0^{\hat{k}}$ on $\mathcal{D}_1$. Now taking $N \in \mathbb{N}$ such that $\cup_{i=1}^N [t_{i-1}, t_i] = [0, T]$ we get that $u \leq \sum_{i=1}^N c_i < \infty$ on $\mathcal{D}_T$. For the $v$ component of $u$, by repeating the above steps, we also get that $v \leq \sum_{i=1}^N c_{v, i}$ in $\mathcal{D}_T$. Consequently, we get that $u$ is bounded in $\mathcal{D}_T$. 

\[ \square \]

### 3.2.4 Properties of the operators $a$ and $r$

**Lemma 4** Let $w, u, v \in W$ be non-negative, $\theta \in \Xi \subset \Theta$. Then there exist $M_a, M_r, m_a \in (0, \infty)$ such that operators $a$ and $r$ appearing in (9) satisfy the following inequalities:

$$|a[w](u, v)| \leq M_a \|w\| W \|u\| W \|v\| W$$

$$a(w)[u, u] \geq m_a \|u\|^2 W$$

$$|a(w^1)[u, v] - a(w^2)[u, v]| \leq M_a \|w^1 - w^2\| W \|u_2\| W \|v_2\| W$$

$$\|r(u)\| W \leq M_r \|u\| W$$

$$\|r(u^1) - r(u^2)\| W \leq M_r \|u^1 - u^2\| W.$$

**Proof** First let us recall the definition of $a$ and $r$ (see (9), (11)). For any $u, v, w \in W$ they are defined as

$$a(w)[u, v] := a_1(w)[u, v] + a_2(w)[u, v] + a_3(w)[u, v], \quad r(u) := \left[ \begin{array}{c} r_1 + r_2 \\ r_3 \end{array} \right],$$

$$a_1(w)[u, v] := (\sigma \nabla u_1, \nabla v_1) + (\mu u_1, v_1), \quad r_1(u)[v] := (\mu_1 f_1(u_1, u_2) + \mu u_1, v_1),$$

$$a_2(w)[u, v] := (\delta \nabla u_2, \nabla v_1), \quad r_2(u)[v] := (-u_1 \nabla \kappa, \nabla v_1),$$

$$a_3(w)[u, v] := (\nabla u_2, \nabla v_2) + (\alpha u_2, v_2), \quad r_3(u)[v] := (\beta f_2(u_1, u_2), v_2).$$

Based on this, now let us verify the properties of the operator $a : W \times W \times W \to \mathbb{R}$. For the boundedness property we have that

$$a_1(w)[u, v] = (\sigma \nabla u_1, \nabla v_1) + (\mu u_1, v_1)$$

$$\leq M_\sigma \|\nabla u_1\| \|\nabla v_1\| + M_\mu \|u_1\| \|v_1\|.$$
with $0 < M_a < \infty$ being a large enough generic constant independent of any $u, v, w \in W$. Next, for the coercivity property let $w = (w_1, w_2)^\top$ be such that $w_1 \geq 0$, then for any $u \in W$ we have that:

$$a(w)[u, u] = a_2(w)[u, u] + a_2(w)[u, u] + a_3(w)[u, u]$$

$$= \langle \sigma \nabla u_1, \nabla u_1 \rangle + \langle \mu u_1, u_1 \rangle + \langle \nabla u_2, \nabla u_2 \rangle + \langle \alpha u_1, u_1 \rangle + \langle \delta w_1 \nabla u_2, \nabla u_2 \rangle$$

$$= \langle \sigma, \|\nabla u_1\|^2 \rangle + \|\nabla u_2\|^2 + \|\mu u_1\|^2 + \|\alpha u_2\|^2 + \|\delta w_1, \|\nabla u_2\|^2\rangle$$

$$\geq m_1 \langle 1, \|\nabla u_1\|^2 \rangle + \|\nabla u_2\|^2 + m_2 \|u_1\|^2 + m_3 \|u_2\|^2 + m_4 (w_1, \|\nabla u_2\|^2)$$

$$\geq k(\sigma, \alpha, \mu, \delta) \langle \|\nabla u_1\|^2 + \|\nabla u_2\|^2 + \|u_1\|^2 + \|u_2\|^2 + (w_1, \|\nabla u_2\|^2) \rangle$$

$$\Rightarrow a(w)[u, u] \geq m_a \|u\|^2_W + \|u_2\|^2_W$$

with $0 < m_a < \infty$ is a small enough constant independent of any $u, w \in W$. Next, for the Lipschitz continuity property let $w_1, w_2 \in W$, then we have that:

$$|a(w_1)[u, v] - a(w_2)[u, v]| = |a_2(w_1)[u, v] - a_2(w_2)[u, v]| = |(w_1 - w_2) \nabla u_2, \nabla u_2|$$

$$\leq M_5 \|w_1 - w_2\|_{L^2} \|\nabla u_2\|_{L^4} \|\nabla u_2\|_{L^6}$$

$$\leq M_5 \|w_1 - w_2\| \|\nabla u_2\| \|u_2\| w_2 \|w\|$$

$$\leq M_a \|w_1 - w_2\| \|\nabla u_2\| \|w_2\| w_2 \|w\|$$

$$\Rightarrow |a(w_1)[u, v] - a(w_2)[u, v]| \leq M_a \|w_1 - w_2\| \|w\|$$

with $0 < M_a < \infty$ beign a large enough generic constant independent of any $u, v, w \in W$.

Now, for the nonlinear operator $r : W \to W'$ let $u, v \in W$. Then we have that

$$|r(u)[v]| \leq |r_1(u)[v]| + |r_2(u)[v]| + |r_2(u)[v]| \leq |(\mu f_1(u_1, u_2) + \mu u_1, v_1)|$$

$$+ |\beta(f_2(u_1, u_2), v_2)| + |(-u_1 \nabla \kappa, \nabla v_1)|$$

$$\leq M_\mu (1 + M_{f_1}) \|u_1\| \|w\| \|v_1\| w + M_\beta M_{f_2} \|u_2\| \|w\| \|v_2\| w + M_\kappa \|u_1\| \|w\| \|v_1\| w$$
\[ \leq k(\kappa, \mu, \alpha, \beta, f_1, f_2) \|u\|_W \|v\|_W \]
\[ \Rightarrow \|r(u)\|_W \leq M_r \|u\|_W \]

with \( 0 < M_r < \infty \) begin a big enough constant independent of any \( u, v \in W \).

Next for the Lipschitz continuity, let \( u^1, u^2 \in W \) then we have that
\[
\|(r(u^1) - r(u^2))(v)\| \leq |(r_1(u^1) - r_1(u^2))(v_1)| + |(r_3(u^1) - r_3(u^2))(v_2)| \\
\leq (M_\mu \|\nabla f_1\|_{L^\infty} + M_\beta \|\nabla f_2\|_{L^\infty}) \|u^1 - u^2\|_W \|v\|_W + M_\mu \|u^1 - u^2\|_W \|v_1\|_W \\
\leq k(\kappa, \mu, \alpha, \beta, f_1, f_2) \|u^1 - u^2\|_W \|v\|_W \\
\Rightarrow \|r(u^1) - r(u^2)\|_W \leq M_r \|u^1 - u^2\|_W
\]

with \( 0 < M_r < \infty \) being a large enough generic constant independent of any \( u^1, u^2 \in W \).

\[ \Box \]

### 3.2.5 Existence of a unique solution

**Theorem 5** For every \( 0 \leq u_0 \in Z \) and \( \theta \in \Theta \), the pseudopalisade system (7) possesses a unique non-negative weak-solution

\[ u \in U \cap Y \cap C(T), \quad u \geq 0, \]

where \( T > 0 \) is dependent on \( \|u_0\|_Z \leq M_{u_0} \) and \( \|\theta\|_Z \leq M_\theta \). Moreover, \( u \) satisfies the following inequality:

\[ \|u\|_Y + \|u\|_{C(T)} + \|u\|_X \leq k_u(M_{u_0}, M_\theta). \]

**Proof** This is a direct consequence of Lemmas 1, 2, 3, 4 and Theorem 5.10 in Yagi (2009). \( \Box \)

### 3.3 Existence of an optimal parameter function

Let us recall that the state equation \( G(u, \theta) = 0 \) (the weak form of Eq. (7)) is given as:

\[ G(u, \theta) = \begin{pmatrix}
  (\partial_t u_1, \cdot) + (\sigma \nabla u_1, \nabla \cdot) + (u_1 \nabla \kappa, \nabla \cdot) + (\delta u_1 \nabla u_2, \nabla \cdot) - (\mu f_1, \cdot) \\
  (\partial_t u_2, \cdot) + (\nabla u_2, \nabla \cdot) + (a u_2, \cdot) - (\beta f_2, \cdot)
\end{pmatrix}, \quad \theta \in \Xi. \]

**Lemma 6** For a fixed \( \theta \in \Xi \), the operator \( G : U \times \Xi \to U' \times Z \) is infinitely Fréchet differentiable. The partial derivative \( G_u := \partial_u G \) of \( G \) with respect to \( u \) at point \( (u, \theta) \)

\[
\begin{align*}
\n\end{align*}
\]

\( \Box \ Springer \)
is represented as a mapping \((\mathbf{u}, \theta) \mapsto \mathbf{G}_\mathbf{u}[\mathbf{u}, \theta]. \mathbf{G}_\mathbf{u} : U \times \Xi \to L(U; U' \times Z).\) Its exact form is given by

\[
\mathbf{G}_\mathbf{u}[\mathbf{u}, \theta](v_1, v_2) = \left( \begin{array}{c}
(\partial_t v_1, \cdot) + (\sigma \nabla v_1, \nabla \cdot) + (v_1 \nabla k, \nabla \cdot) + (\partial_t v_1 \nabla u_2, \nabla \cdot) - (\mu \partial_{u_1} v_1, \cdot) + (\partial u_1 \nabla v_2, \nabla \cdot) - (\mu \partial_{u_2} f_1 v_2, \cdot) \\
(\partial_t v_2, \cdot) + (\nabla v_2, \nabla \cdot) + (\alpha v_2, \cdot) - (\partial_{u_1} f_2 v_1, \cdot) - (\beta \partial_{u_2} f_2 v_2, \cdot) \\
\end{array} \right) \bigg|_{v_1(0)} \bigg|_{v_2(0)}.
\]

(22)

Similarly, the partial derivative \(\mathbf{G}_\theta := \partial_\theta \mathbf{G}\) of \(\mathbf{G}\) with respect to \(\theta\) at the point \((\mathbf{u}, \theta)\) is represented as a mapping \((\mathbf{u}, \theta) \mapsto \mathbf{G}_\theta[\mathbf{u}, \theta]. \mathbf{G}_\theta : U \times \Xi \to L(\Xi; U' \times Z).\) Its exact form is given as

\[
\mathbf{G}_\theta[\mathbf{u}, \theta](\phi_1, \ldots, \phi_6) = \left( \begin{array}{c}
(\phi_1 \nabla u_1, \nabla \cdot) + (u_1 \nabla \phi_2, \nabla \cdot) + (\phi_3 u_1 \nabla u_2, \nabla \cdot) + (\phi_4 f_1(\mathbf{u}), \cdot) \\
(\phi_5 u_2, \cdot) + (-\phi_6 f_2(\mathbf{u}), \cdot) \\
0 \\
0
\end{array} \right).
\]

(23)

**Proof** By following the technique of Lemma 1.17 of Hinze et al. (2008) and by using equation (1.95) of Yagi (2009) (due to Lemma 3) we can obtain the first partial Fréchet derivatives \(\mathbf{G}_\mathbf{u}\) and \(\mathbf{G}_\theta\) as given in (22) and (23), respectively. For the latter, it is clear that the higher-order derivatives \(\partial_\theta^{(k)} \mathbf{G}, k > 1\) are equal to 0. However, the second derivative \(\partial_{\mathbf{u}}^{(2)} \mathbf{G}\) reads

\[
\mathbf{G}_{\mathbf{u}}[\mathbf{u}, \theta](v_1, v_2)(v_3, v_4) = \left( \begin{array}{c}
(\delta u_1 \nabla v_4, \nabla \cdot) + (\delta v_4 \nabla v_2, \nabla \cdot) - (\delta u_1 v_1 v_3, \cdot) - (\mu \delta_{u_1} v_1 f_1 v_2 v_3, \cdot) \\
- (\mu \delta_{u_2} v_1 f_1 v_2 v_3, \cdot) - (\mu \delta_{u_2} f_1 v_2 v_3, \cdot) - (\beta \delta_{u_1} f_1 v_2 v_3, \cdot) \\
0 \\
0
\end{array} \right).
\]

Now, further derivatives of \(\mathbf{G}\) with respect to \(\mathbf{u}\) are just multiples of the partial derivatives \(\partial_{\mathbf{u}}^{(k)} f_1\) and \(\partial_{\mathbf{u}}^{(k)} f_2\). Since \(f_1\) and \(f_2\) are \(C^\infty\) functions, we get that \(\mathbf{G}\) is infinitely Fréchet differentiable. \(\square\)

**Lemma 7** The operator \(\mathbf{G}_\mathbf{u} \in L(U \times \Theta; U' \times Z)\) has a bounded inverse.

**Proof** Let \(U' \ni \mathbf{g} = (g_1, g_2)\) and \(Z \ni \mathbf{v}_0\), let \(\mathbf{G}_\mathbf{u}\) be the operator \(\mathbf{G}_\mathbf{u}\) linearized at \(\mathbf{u} \in U, \theta \in \Theta\). Then \(\mathbf{G}_\mathbf{u}\) is said to have an inverse if there exists a unique \(\mathbf{v} = (v_1, v_2)\) which satisfies the equation \(\mathbf{G}_\mathbf{u}(\mathbf{v}) = \mathbf{g}\). This is to say that \(\mathbf{v}\) satisfies the following PDE in weak form:

\[
(\partial_t v_1, \cdot) + (\sigma \nabla v_1 + \nabla k v_1, \nabla \cdot) + (\delta v_1 \nabla u_2, \nabla \cdot) + (\delta u_1 \nabla v_2, \nabla \cdot) = (g_1 + \mu \partial_{u_1} f_1 v_1 + \mu \partial_{u_2} f_1 v_2, \cdot)
\]

\[
(\partial_t v_2, \cdot) + (\nabla v_2, \nabla \cdot) + (\alpha v_2, \cdot) = (g_2 + \beta \partial_{u_2} f_2 v_2 + \partial_{u_1} f_2 v_1, \cdot)
\]

\(\mathbf{v}(0) = \mathbf{v}_0,\)

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which can be rewritten as

\[(\partial_t v_1, \cdot) + (\sigma \nabla v_1 + v_1[\nabla \kappa + \delta \nabla u_2], \nabla \cdot) + (\delta u_1 \nabla v_2, \nabla \cdot) = (g_1 + \mu \partial_{u_1} f_1 v_1 + \mu \partial_{u_2} f_1 v_2, \cdot) \]

\[(\partial_t v_2, \cdot) + (\nabla v_2, \nabla \cdot) + (\alpha v_2, \cdot) = (g_2 + \beta \partial_{u_2} f_2 v_2 + \partial_{u_1} f_2 v_1, \cdot) \]

\[v(0) = v_0.\]

This can be further simplified to

\[(\partial_t v_1, \cdot) + (\sigma \nabla v_1 + v_1 \tilde{k}, \nabla \cdot) + (\delta u_1 \nabla v_2, \nabla \cdot) = (\tilde{g}_1(u, v, \theta), \cdot) \]

\[(\partial_t v_2, \cdot) + (\nabla v_2, \nabla \cdot) + (\alpha v_2, \cdot) = (\tilde{g}_2(u, v, \theta), \cdot) \]

\[v(0) = v_0.\]

where \(\tilde{k} := \nabla \kappa + \delta \nabla u_2\), \(\tilde{g}_1(u, v, \theta) := g_1 + \mu \partial_{u_1} f_1 v_1 + \mu \partial_{u_2} f_1 v_2\) and \(\tilde{g}_2(u, v, \theta) := g_2 + \beta \partial_{u_2} f_2 v_2 + \beta \partial_{u_1} f_2 v_1\). Since \(u \in U\) (cf. Theorem 5) and since \(f, \nabla f\) are elements of \(Z \cap Y\) for all \(t \in I\), Lemmas 4, 3 can be used to invoke Theorem 4.7 of Yagi (2009) for obtaining the existence of a unique \(v\) that satisfies the equation \(G_u[u, \theta](v) = g\). Specifically, for \(g \in U' \cap V(T)\), by performing computations similar to Lemma 1, it follows that the solution \(v \in X \cap Y\) satisfies the following inequality:

\[\|v\|_{W(T)} \leq k_{G_u}(u, \theta)(\|v_0\|_V + \|g\|_{V(T)}) \leq k_{G_u}(\|v_0\|_Z + \|g\|_{V(T)})\]

\[\Rightarrow \|G_u^{-1}\|_{L(U', U)} \leq k_{G_u}(u, \theta). \tag{24}\]

As a consequence of this, we can apply the implicit function theorem to interpret \(u\) as a function of \(\theta\) via the mapping \(\theta \mapsto u(\theta)\). Moreover, it also follows that \(u(\theta)\) is infinitely differentiable, and thus also Lipschitz, with respect to \(\theta\). Now we are ready to establish the existence of an optimal parameter \(\theta\).

**Theorem 8** Let \(T \in (0, \infty), u_0 \in Z\) and \(u_0\) be non-negative. Then there exists an optimal parameter vector \(\theta \in \Theta\) minimizing the functional \(J(\theta)\) from (12).

**Proof** Clearly, by definition of \(J(\theta)\), we have that \(J \geq 0\) thus \(\inf_{\theta} J(\theta)\) exists. Due to continuity of the mapping \(\theta \mapsto J(\theta)\) and closedness of \(\Theta\), there exists a sequence \((\theta_n)_{n \in \mathbb{N}} \subset \Theta\) such that \(J(\theta_n) \to J(\theta^*) = \inf_{\theta} J(\theta)\). Since \(\Theta \subset \Theta\) is bounded, the sequence \((\theta_n)_{n \in \mathbb{N}}\) is also bounded in \(\Theta\). Hence, we can extract a weakly convergent sub-sequence \((\theta_{n_j})_{j \in \mathbb{N}}\) such that \(\theta_{n_j} \to \theta^*\) in \(\Theta\). Now, since the mapping \(\Theta \ni \theta \mapsto u(\theta) \in X\) is a bounded operator (due to Lemma 3 and Theorem 5) we can let \((u_{n_j})_{j \in \mathbb{N}}\) be a sequence of solutions corresponding to the parameter sequence \((\theta_{n_j})_{j \in \mathbb{N}}\), where, due to Lemma 3, the sequence \((u_{n_j})_{j \in \mathbb{N}}\) is bounded in \(X\), there exists a sub-sequence \((u_{n_{j_k}})_{k \in \mathbb{N}}\) such that \(u_{n_{j_k}} \to u^*\) in \(W(T)\). Moreover, we also have that \((u'_{n_{j_k}})_{k \in \mathbb{N}} \in W(T)'\) is a bounded sequence. Now, applying the Lions-Aubin compactness theorem (see e.g., Necas et al. 1996) we get that:

\[u_{n_{j_k}} \to u^* \text{ in } V(T), \quad \nabla u_{n_{j_k}} \rightharpoonup \nabla u^* \text{ in } V(T), \quad \text{and } u'_{n_{j_k}} \rightharpoonup u''^* \text{ in } W(T)'\].
Due to uniqueness, we get that $u^*$ is the solution to (7) corresponding to $\theta^*$, i.e. $\theta^* \mapsto u^*(\theta^*)$. Finally, due to the weak lower semicontinuity of the norm and the weak convergence of $u(\theta_{j_k})$ to $u^*(\theta^*)$ in $V(T)$, we have that

$$J(u^*(\theta^*), \theta^*) = \liminf_{k \to \infty} J(u_{j_k}, \theta_{j_k}) = \inf_{\theta} J(u(\theta), \theta) \leq J(u^*(\theta^*), \theta^*).$$

Thus $J(u(\theta^*), \theta^*)$ is indeed equal to $\inf_{\theta} J(u(\theta), \theta)$.

Next we shall construct the minimizing sequence by deducing the adjoint equation and the necessary optimality condition.

**Theorem 9** Let $(\tilde{u}, \tilde{\theta})$ be an optimal solution to problem (13). Then there exists an adjoint state $\tilde{p} \in U \subset U''$ s.t. the following optimality condition holds

$$G(\tilde{u}, \tilde{\theta}) = 0 \quad (25a)$$

$$G_u^*(\tilde{u}, \tilde{\theta})\tilde{p} = -J_u(\tilde{u}, \tilde{\theta}) \quad (25b)$$

$$(J_\theta(\tilde{u}, \tilde{\theta}) + G_{\theta}^*(\tilde{u}, \tilde{\theta})\tilde{p}, \theta - \bar{\theta})_\Theta \geq 0 \quad (25c)$$

where $G_{\theta}^*$ and $G_u^*$ denote the adjoint operator of $G_{\theta}$ and $G_u$, respectively.

**Proof** Due to Theorem 5 and Lemma 7 we can invoke Theorem 1.48 of Hinze et al. (2008) for the reduced cost function $\hat{J}(\theta) := J(u(\theta), \theta)$, which ensures that the local solution $\tilde{\theta} \in \Xi$ satisfies the following variational inequality:

$$\langle \hat{J}'(\tilde{\theta}), \theta - \tilde{\theta} \rangle_{\Theta', \Theta} \geq 0 \quad \forall \theta \in \Xi.$$

For $\bar{\theta} := \theta - \tilde{\theta}, \theta \in \Xi$ we have that

$$\langle \hat{J}'(\bar{\theta}), \bar{\theta} \rangle_{\Theta', \Theta} = \langle J_u(\tilde{u}, \tilde{\theta}), \bar{u}' \bar{\theta} \rangle_{U', U} + \langle J_\theta(\tilde{u}, \tilde{\theta}), \bar{\theta} \rangle_{\Theta', \Theta}$$

$$= \langle (\tilde{u})^* J_u(\tilde{u}, \tilde{\theta}), \bar{\theta} \rangle_{\Theta', \Theta} + \langle J_\theta(\tilde{u}, \tilde{\theta}), \bar{\theta} \rangle_{\Theta', \Theta}$$

Based on the state equation $\hat{G}(\theta) := G(u(\theta), \theta) = 0$, we have that $G_u u'(\theta) + G_{\theta} = 0$. This implies $u'(\theta) = -G_{u}^{-1}G_{\theta}$. Consequently, we get that $u'(\theta)^* J_u = -G_{\theta}^*(G_u^*)^{-1} J_u$. Defining $p := -(G_u^*)^{-1} J_u$ we get the following adjoint equation:

$$G_u^* p = -J_u \text{ in } U'.$$

Explicitly, the equation can be written as

$$\begin{pmatrix}
-\partial_t - \nabla \cdot (\alpha \nabla) + (\nabla \kappa + \delta \nabla u_2) \cdot \nabla - \nabla \cdot (\delta u_1 \nabla) & 0 \\
0 & -\partial_t - \Delta + \alpha
\end{pmatrix}
\begin{pmatrix}
p_1 \\
p_2
\end{pmatrix}
= 
\begin{pmatrix}
\mu \delta u_1 f_1(u) & \mu \delta u_2 f_1(u) \\
\beta \delta u_1 f_2(u) & \beta \delta u_2 f_2(u)
\end{pmatrix}
\begin{pmatrix}
p_1 \\
p_2
\end{pmatrix}
= 
\begin{pmatrix}
\nabla_n p_1 = 0, \quad \nabla_n p_2 = 0, \quad p_1(T) = u_1(T) - O, \quad p_2(T) = 0.
\end{pmatrix}$$

\[\square\]
Proof

\[
\begin{align*}
&- (\delta_t p_1, \varphi) + (\sigma \nabla p_1, \nabla \varphi) + ((\nabla \kappa + \delta \nabla u_1) \cdot \nabla p_1, \varphi) + (\delta u_1 \nabla p_2, \varphi) \\
&= - (\mu \delta u_1 f_1 p_1, \varphi) - (\nu \delta u_2 f_1 p_2, \varphi), \\
&- (\delta_t p_2, \psi) + (\nabla p_2, \nabla \psi) + (\alpha p_2, \psi) = - (\beta \delta u_2 f_2 p_2, \psi) - (\delta u_1 f_2 p_1, \psi), \\
p_1(T) = u_1(T) - O, \quad p_2(0) = 0.
\end{align*}
\]

(28)

In light of (26) and using the Riesz isomorphism associated to the corresponding duality pairing, the optimality condition reads as:

\[
(G_\theta^* \mathbf{p} + J_\theta, \theta - \tilde{\theta})_\Theta \geq 0, \quad \forall \theta \in \Xi.
\]

(29)

We now establish the stability result for the adjoint equation

**Lemma 10** Let \( p^1, \ p^2 \) be two solutions to Eq. (27) generated by two \( u^1, \ \theta^1 \) and \( u^2, \ \theta^2 \) respectively. Then inequality (31) holds.

**Proof** Let \( q := p^1 - p^2, \ v := u^1 - u^2 \) and \( \theta := \theta^1 - \theta^2 \)

\[
\begin{align*}
&\left( -\delta_t - \nabla \cdot (\sigma \nabla v) + (\nabla \kappa + \delta \nabla u_1^2) \cdot \nabla + \mu \delta u_1 g_11 - \nabla \cdot (\delta u_1 \nabla v) + \mu \delta u_1 g_12 \right) (\nu q_1) = (R_1, \ R_2) \\
&\nabla_n q_1 = 0, \quad \nabla_n q_2 = 0, \quad q_1(T) = v_1(T), \quad q_2(T) = 0.
\end{align*}
\]

(30)

where \( R_1 := \nabla \cdot (\delta_1 \nabla p_1^2) - (\nabla \theta_2 + \delta \nabla v_2 + \vartheta \nabla u_2^2) \cdot \nabla p_1^2 + \nabla \cdot (\delta \nabla v_1^2 \nabla p_2^2 + \delta_3 \nabla u_1^2 \nabla p_2^2) + g^3 \)

\[
\begin{align*}
R_2 &:= \delta_t p_2^2 + g^4, \\
g^3(v, \theta) &:= g_1^3(v) + g_2^3(\theta), \quad g^4(v, \theta) := g_1^3(v) + g_2^4(\theta), \\
\theta^1 &:= \theta^1 \mu_f f_1(u^1), \quad g^1 \mu_f f_1(u^1) = g_1^1(v) + g_2^1(\theta), \\
g^2 v_1 &:= \delta \nabla u_2 f_2(u^1), \quad g^2 v_2 := \delta \nabla u_2 f_2(u^2),
\end{align*}
\]

Since \( p^1, p^2 \in U \subset U'' \), \( u^1, u^2 \in U \) and \( \theta^1, \theta^2 \in \Theta \), the RHS terms \( R_1 \) and \( R_2 \) are elements of \( Z(T) \). Thus, letting \( R := (R_1, R_2) \) and \( G_u^* \) denote the adjoint operator of \( G_u \) the above Eq. (30) can be abstractly written as:

\[
G_u^* q = R \ in \ U'
\]

Due to the invertibility of \( G_u \) we get the existence of a unique solution to (30). Consequently, we have that

\[
\begin{align*}
\|q\|_U &\leq \|(G_u^*)^{-1} R\|_U \leq \|(G_u^*)^{-1} \|_{L(U', U)} \|R\|_U' \\
&\leq \|G_u^{-1}\|_{L(U', U)} \|R\|_U' \\
&\leq kG_u(u, \theta) \|R\|_Z.
\end{align*}
\]

(31)
Now we shall provide the smoothness (in terms of $\theta$) result for the cost functional $J$.

**Theorem 11** Let $u \in Y$, $p \in Y$ and $\theta \in \Theta$, then the cost functional $\theta \mapsto \hat{J}(\theta) := J(u(\theta), \theta)$ is infinitely Frechet differentiable. Moreover, the mapping $\theta \mapsto \nabla \hat{J}(\theta)$, $\nabla \hat{J} : Y \times Y \times \Xi \rightarrow \Xi$ is Lipschitz continuous.

**Proof** The infinite differentiability of $\hat{J}$ follows from the facts that $\theta \mapsto u(\theta)$ is a smooth mapping (due to Lemma 7) and $J$ is a quadratic functional of $u$ and $\theta$. Moreover, as already mentioned above, $\nabla \hat{J}$ takes the following form:

$$\nabla \hat{J} = \lambda \theta + F(\theta), \quad F(\theta) := (G^*_\theta p)(\theta) = -(G^*_\theta(G^*_\theta)^{-1} J_u)(\theta).$$

Moreover, based on (32), $\nabla \hat{J}$ can be explicitly written as

$$\nabla \hat{J} := \begin{pmatrix}
\nabla u_1 \cdot \nabla p_1 + \lambda \theta_1 \\
\nabla \cdot (u_1 \nabla p_1) + \lambda \theta_2 \\
u_1 \nabla u_2 \cdot \nabla p_1 + \lambda \theta_3 \\
\partial_x f_1(u)p_1 + \lambda \theta_4 \\
u_2 p_2 + \lambda \theta_5 \\
\partial_x f_2(u)p_2 + \lambda \theta_6
\end{pmatrix}.$$  

(32)

Since $u$, $p \in Y$, we get that $F(\theta) \in \Theta$. Due to the stability result of the adjoint $p$ (see Lemma 10) and the linear structure of $J_u$, for the Lipschitz continuity of $\nabla J$ it suffices to only consider the operator $G^*_\theta p$ and establish its stability with respect to $u$, $p$ and $\theta$. To this end we shall consider each component of the Jacobian vector function (32)

**Component 1:**

$$\|\nabla u_1 \nabla p_1 - \nabla u_2 \nabla p_2\|_V = \|\nabla u_1 (\nabla p_1 - \nabla p_2) + \nabla p_2 (\nabla u_1 - \nabla u_2)\|_V \\
\leq \|\nabla u_1\|_{L^4} \|\nabla p_1 - \nabla p_2\|_{L^4} + \|\nabla p_2\|_{L^4} \|\nabla u_1 - \nabla u_2\|_{L^4} \\
\leq \|u_1\|_{Z} \|p_1 - p_2\|_{Z} + \|p_2\|_{Z} \|u_1 - u_2\|_{Z}$$

$\Rightarrow \|\nabla u_1 \nabla p_1 - \nabla u_2 \nabla p_2\|_{V(T)} \leq \|u_1\|_{Y(T)} \|p_1 - p_2\|_{Z(T)} \|p_2\|_{Y(T)} \|u_1 - u_2\|_{Z(T)}$

**Component 2:**

$$\|\nabla \cdot (u_1 \nabla p_1 - u_2 \nabla p_2)\|_V = \|\nabla \cdot \left(u_1 (\nabla p_1 - \nabla p_2) + p_2 (\nabla u_1 - \nabla u_2)\right)\|_V \\
\leq \|\nabla \cdot \left(u_1 (\nabla p_1 - \nabla p_2) + p_2 (\nabla u_1 - \nabla u_2)\right)\|_V \\
\leq \|\nabla \cdot \left(u_1 (\nabla p_1 - \nabla p_2)\right)\| + \|\nabla \cdot \left(p_2 (\nabla u_1 - \nabla u_2)\right)\|_V \\
\leq \|\nabla u_1 \cdot (\nabla p_1 - \nabla p_2)\| + \|u_1\|_{L^\infty} \|\Delta p_1 - \Delta p_2\|_V \\
+ \|\nabla u_1 \cdot (\nabla u_1 - \nabla u_2)\| + \|u_1\|_{L^\infty} \|\Delta u_1 - \Delta u_2\|_V \\
\leq \|u_1\|_{Z} \|p_1 - p_2\|_{Z} + \|u_1\|_{L^\infty} \|p_1 - p_2\|_{Z} + \|u_1\|_{L^\infty} \|u_1 - u_2\|_{Z} + \|u_1\|_{L^\infty} \|u_1 - u_2\|_{Z}.$$
+ \|p_2\|_{L^\infty} \|u_1 - u_2\|_Z \\
\leq (\|u_1\|_Z + \|u_1\|_{L^\infty}) \|p_1 - p_2\|_Z + (\|p_2\|_Z + \|p_2\|_{L^\infty}) \|u_1 - u_2\|_Z \\
\leq \|u_1\|_Z \|p_1 - p_2\|_Z + \|u_1\|_{L^\infty} \|p_1 - p_2\|_Z + \|p_2\|_Z \|u_1 - u_2\|_Z \\
+ \|p_2\|_Z \|u_1 - u_2\|_Z \\
\leq 2\|u_1\|_Z \|p_1 - p_2\|_Z + 2\|p_2\|_Z \|u_1 - u_2\|_Z \\
\Rightarrow \|\nabla \cdot (u_1 \nabla p_1 - u_2 \nabla p_2)\|_{V(T)} \leq 2\|u_1\|_{Y(T)} \|p_1 - p_2\|_{Z(T)} + 2\|p_2\|_{Y(T)} \|u_1 - u_2\|_{Z(T)}. \\

Component 3:

\|u_1 \nabla u_1 \nabla p_1 - u_2 \nabla u_2 \nabla p_2\|_V = \|u_1 \nabla u_1 (\nabla p_1 - \nabla p_2) + u_1 \nabla p_2 (\nabla u_1 - \nabla u_2) \\
+ \nabla u_2 \nabla p_2 (u_1 - u_2)\|_V \\
\leq \|u_1 \nabla u_1 \|_{L^4} (\|\nabla p_1 - \nabla p_2\|_{L^4} + \|u_1 \nabla p_2\|_{L^4} (\|\nabla u_1 - \nabla u_2\|_{L^4} \\
+ \|\nabla u_2 \nabla p_2\|_V \|u_1 - u_2\|_{L^\infty} \\
\leq \|u_1 \|_{L^2} \|\nabla u_1 \|_{L^4} \|p_1 - p_2\|_W + \|u_1 \nabla p_2\|_{L^4} \|\nabla u_1 - \nabla u_2\|_W \\
+ \|\nabla u_2 \nabla p_2\|_V \|u_1 - u_2\|_{L^\infty} \\
\leq \|u_1 \|_{L^2} \|\nabla u_1 \|_{L^4} \|p_1 - p_2\|_W + \|u_1 \|_{L^\infty} \|\nabla p_2\|_{L^4} \|\nabla u_1 - \nabla u_2\|_W \\
+ \|\nabla u_2 \|_{L^2} \|\nabla p_2\|_{L^4} \|u_1 - u_2\|_{L^\infty} \\
\leq \|u_1 \|_{L^2} \|\nabla u_1 \|_{L^4} \|p_1 - p_2\|_W + \|u_1 \|_{L^\infty} \|\nabla p_2\|_{L^4} \|\nabla u_1 - \nabla u_2\|_W \\
+ \|\nabla u_2 \|_{L^2} \|\nabla p_2\|_{L^4} \|u_1 - u_2\|_{L^\infty} \\
\leq \|u_1 \|_{L^2} \|\nabla u_1 \|_{L^4} \|p_1 - p_2\|_W + \|u_1 \|_{L^\infty} \|\nabla p_2\|_{L^4} \|\nabla u_1 - \nabla u_2\|_W \\
+ \|\nabla u_2 \|_{L^2} \|\nabla p_2\|_{L^4} \|u_1 - u_2\|_{L^\infty} \\
\Rightarrow \|u_1 \nabla u_1 \nabla p_1 - u_2 \nabla u_2 \nabla p_2\|_{V(T)} \leq \|u_1\|_{Y(T)} \|p_1 - p_2\|_{Z(T)} + \|p_2\|_{Y(T)} \|u_1 - u_2\|_{Z(T)} \\
+ \|u_2\|_{Y(T)} \|u_1 - u_2\|_{Z(T)}. \\

Component 4:

\|\partial_u f^1_1 p_1 - \partial_u f^2_1 p_2\|_V \leq \|\partial_u f^1_1 p_1 - \partial_u f^1_1 p_2 + \partial_u f^1_1 p_2 - \partial_u f^2_1 p_2\| \\
\leq \|\partial_u f^1_1 p_1 - \partial_u f^1_1 p_2\|_V + \|\partial_u f^1_1 p_2 - \partial_u f^2_1 p_2\| \\
\leq \|\partial_u f^1_1 \|_W \|p_1 - p_2\|_W + \|p_2\|_W \|\partial_u f^1_1\|_W - \|\partial_u f^1_1\|_W \\
\leq \|\partial_u f^1_1 \|_W \|p_1 - p_2\|_W + \|p_2\|_W \|\partial_u f^1_1\|_W \|u_1 - u_2\|_Z \\
\Rightarrow \|\partial_u f^1_1 p_1 - \partial_u f^2_1 p_2\|_{V(T)} \leq M_{f_1} \|p_1 - p_2\|_{Z(T)} + \|p_2\|_{Y(T)} M_{f_1} \|u_1 - u_2\|_{Z(T)} \\

Component 5:

\|u_1 p_1 - u_2 p_2\|_V = \|u_1 (p_1 - p_2) + p_2 (u_1 - u_2)\|_V \\
\leq \|u_1\|_{L^4} \|(p_1 - p_2)\|_{L^4} + \|p_2\|_{L^4} \|u_1 - u_2\|_{L^4} \\
\leq \|u_1\|_W \|p_1 - p_2\|_W + \|p_2\|_W \|u_1 - u_2\|_W \\
\Rightarrow \|u_1 p_1 - u_2 p_2\|_V \leq \|u_1\|_{Y(T)} \|p_1 - p_2\|_{Z(T)} + \|p_2\|_{Y(T)} \|u_1 - u_2\|_{Z(T)}.
Component 6:

\[
\| \partial u_2 f_1^2 p_1 - \partial u_2 f_2^2 p_2 \| \leq \| \partial u_2 f_1^1 p_1 - \partial u_2 f_2^1 p_2 + \partial u_2 f_2^1 p_2 - \partial u_2 f_2^2 p_2 \|
\]

\[
\quad \leq \| \partial u_2 f_1^1 p_1 - \partial u_2 f_2^1 p_2 \| v + \| \partial u_2 f_2^1 p_2 - \partial u_2 f_2^2 p_2 \| v
\]

\[
\quad \leq \| \partial u_2 f_1^1 \| w \| p_1 - p_2 \| w + \| p_2 \| w \| \partial u_2 f_2^1 - \partial u_2 f_2^2 \| w
\]

\[
\quad \leq \| \partial u_2 f_2^1 \| w \| p_1 - p_2 \| w + \| p_2 \| w \| \partial u_2 f_2^1 \| L_{\infty} \| u^1 - u^2 \| Z
\]

\[
\Rightarrow \| \partial u_2 f_2^1 p_1 - \partial u_2 f_2^2 p_2 \| \leq M_{f_2} \| p_1 - p_2 \| w + \| p_2 \| \gamma(T) M_{f_2} \| u^1 - u^2 \| Z(T)
\]

Finally, we need the following result for the numerical solution of the minimization problem.

**Theorem 12** Let \( \theta^* \in \Xi \) be the solution of system (25). Then Algorithm 1., i.e. the projected gradient descent method, generates a minimizing sequence \((\theta_n)_{n \in \mathbb{N}} \in \Xi\) that converges to \( \theta^* \) in \( \Theta \).

**Proof** Since \( \Theta \) is a Hilbert space and \( \Xi \subset \Theta \) is a closed convex set, the optimality condition can be written as

\[
\theta^* = \mathcal{P}_\Xi(\theta^* - \gamma \nabla \hat{J}(\theta^*))
\]

(33)

where, \( \gamma > 0 \) is some arbitrary fixed constant, \( \mathcal{P}_\Xi(\theta) = \text{argmin}_{\theta \in \Xi} \| \hat{\theta} - \theta \|_\Theta \) being the projection operator onto the convex subset \( \Xi \). First we notice that any arbitrary \( \theta \) obtained via the Eq. (33) is an element of \( \Xi \). Without loss of generality, letting \( \gamma := \lambda^{-1} \) we have that

\[
\theta^* = \mathcal{P}_\Xi(\theta^* - \lambda^{-1} \nabla \hat{J}(\theta^*)) = \mathcal{P}_\Xi(-\lambda^{-1} F(\theta^*))
\]

\[
\Rightarrow \| \theta^* \| \leq \| \mathcal{P}_\Xi(0) + \theta^* - \mathcal{P}_\Xi(0) \|
\]

\[
\leq \| \mathcal{P}_\Xi(0) \| + \| \mathcal{P}_\Xi(-\lambda^{-1} F(\theta^*)) - \mathcal{P}_\Xi(0) \|
\]

\[
\leq \| \mathcal{P}_\Xi(0) \| + \lambda^{-1} \| F(\theta^*) \|
\]

The last inequality follows from the nonexpansiveness property of the projection operator. Since \( F(\cdot) \in \Theta \) we get that \( \theta^* \in \Theta \) and the projection operator \( \mathcal{P}_\Xi \) ensures \( \theta^* \in \Xi \). Finally, due to Lipschitz continuity of \( F \), we can invoke Theorem 2.4 from Hinze et al. (2008) to conclude that the projected gradient descent generates a minimizing sequence \( (\theta)_k \subset \Xi \).

□

This completes the analysis section. Next we perform numerical simulations and discuss the obtained results.

### 4 Numerical simulations

In this section we present a numerical method to compute the optimal parameter vector \( \theta \in \Theta \) that generates a specified target state which is provided as an input data.
There are two main paradigms for numerically solving the OCP (13): *first optimize then discretize* and *first discretize then optimize*. In the case of a pure Galerkin approximation, both techniques produce the same outcome. The former, however, not only results in a strongly consistent scheme (in general), but also offers superior asymptotic approximation, both techniques produce the same outcome. The former, however, not only

\[ \begin{align*}
(\partial_t u_1, \varphi) + (\sigma \nabla u_1 + u_1 \nabla \kappa, \nabla \varphi) + (\delta u_1 \nabla u_2, \nabla \varphi) &= (\mu f_1, \varphi), \quad \forall \varphi \in W \\
(\partial_t u_2, \psi) + (\nabla u_2, \nabla \psi) + (\alpha u_2, \psi) &= (\beta f_2, \psi), \quad \forall \psi \in W
\end{align*} \]

\[ u_1(0) = u_{1,0}, \quad u_2(0) = u_{2,0}. \]

Similarly, letting \( O \) represent the final (target) value of \( u_1 \), i.e. \( u_1(T) = O \), with \( O \) being the specified data, the adjoint Eq. (25b) reads

\[ \begin{align*}
-(\partial_t p_1, \varphi) + (\sigma \nabla p_1, \nabla \varphi) + ((\nabla \kappa + \delta \nabla u_1) \cdot \nabla p_1, \varphi) &= -(\delta u_1 \nabla p_2, \varphi) - (\mu \partial u_1 f_1 p_1, \varphi) - (\nu \partial u_2 f_2 p_2, \varphi), \\
-(\partial_t p_2, \psi) + (\nabla p_2, \nabla \psi) + (\alpha p_2, \psi) &= -(\beta \partial u_2 f_2 p_2, \psi) - (\partial u_1 f_2 p_1, \psi),
\end{align*} \]

\[ p_1(T) = u_1(T) - O, \quad p_2(0) = 0 \]

Together, the above two equations can be compactly written as:

\[ \begin{align*}
(\partial_t u, \varphi) + (A(u; \theta) u, \varphi) &= (f(u; \theta), \varphi), \quad \forall \varphi \in W \\
-(\partial_t p, \psi) + (A^*(p; \theta, u)p, \psi) &= (g(u; \theta)p, \psi), \quad \forall \psi \in W
\end{align*} \]

\[ u(0) = u_0, \quad p(T) = p_T \]

System (34) is numerically solved by discretizing it both spatially and temporally. For the spatial discretization we use the finite element method. Consequently, we replace the space \( W \) by a finite-dimensional subspace \( W_h \subset W \) which consists of continuous piecewise polynomial functions of degree 1, spanned by a nodal basis \( \{ \varphi_j \} \) with \( \dim(W_h) = N_h \). The time interval \( I := [0, T], T \in \mathbb{R}^+ \) is divided into \( N_T \) subintervals, each having width \( \tau := \frac{T}{N_T} \). Based on this, the temporal grid points are denoted by \( I_T := \{ t_n \} \) with \( t_n := n \tau \) for \( n = 1, \ldots, N_T \). Finally, let \( u_n^h := u(t_n)^h, \theta_n^h := \theta(t_n)^h, p_n^h := p(t_n)^h \) denote the finite dimensional approximations of \( u, p, \theta \) at time point \( t_n \), respectively. Then for all \( \varphi^n, \psi^n \in W_h \), the discrete version of (34) is given as:

\[ \begin{align*}
(u_{n+1}^h, \varphi^h) + \tau (A(u_n^h; \theta_n^h) u_n^h, \varphi^h) &= (u_n^h, \varphi^h) + (f(u_n^h; \theta_n^h), \tau \varphi^h), \\
(p_{n+1}^h, \psi^h) + \tau (A^*(p_n^h; \theta_n^h, u_n^h) p_n^h, \psi^h) &= (p_n^h, \psi^h) + (g(u_n^h; \theta_n^h)p_n^h, \tau \psi^h).
\end{align*} \]
Fig. 1 Target pseudopalisade patterns

Given $\theta_{h,\tau} := (\theta^h_n)_{n \in \mathbb{N}, n < N_\tau}$, the finite-element scheme (35a) can be used to obtain an approximate solution $u_{h,\tau} := (u^h_n)_{n \in \mathbb{N}, n < N_\tau}$ of the state equation. Analogously, given $\theta_{h,\tau}$ and $u_{h,\tau}$, the finite element scheme (35b) can be used to generate an approximate solution $p_{h,\tau} := (p^h_n)_{n \in \mathbb{N}, n < N_\tau}$ of the adjoint equation. Subsequently, the approximates $u_{h,\tau}$ and $p_{h,\tau}$ can be used to compute a new $\theta_{h,\tau}$ based on the optimality relation (29). This basically leads to the following iterative method, commonly known as the projected gradient descent method, for computing the optimal parameter function $\theta^\ast$. The sequential steps of the procedure are described in Algorithm 1. The algorithm can be viewed as a mapping $(O, u_0) \mapsto E(O, u_0) = (\hat{u}, \theta^\ast)$, which takes an initial value $u_0$ and a final (target) value $O$ and computes the optimal solution $\hat{u} \in U$ and optimal parameter $\hat{\theta} \in \Theta_{ad}$. Since Algorithm 1, (i.e. the mapping $E$), is a numerical method, it is clear that $\hat{u}$ and $\hat{\theta}$ are the discrete representatives of the corresponding true optimal functions $u^\ast$ and $\theta^\ast$ (Tables 1, 2).

Algorithm 1: PGD

Data: $O \in H^2(\mathcal{D}), u_0 \in H^2(\mathcal{D}), \epsilon > 0, \tau, h > 0, N_\tau \in \mathbb{N}, T > 0$

$\theta^0_{h,\tau} := \theta^0$

for $k = 1, \ldots$ do

$u_{h,\tau}^{k+1} = S^h_{\tau}(\theta^k_{h,\tau})$ using (35a)

$p_{h,\tau}^{k+1} = (S^h_{\tau})^* (u_{h,\tau}^{k+1} , \theta^k_{h,\tau})$ using (35b)

$-\nabla J(u_{h,\tau}^{k+1}, p_{h,\tau}^{k+1}, \theta^k_{h,\tau}) = -G^\ast(u_{h,\tau}^{k+1}, \theta^k_{h,\tau})(p_{h,\tau}^{k+1} - \lambda \theta^k_{h,\tau})$

$\theta^k_{h,\tau} + \gamma_k p_{h,\tau}^{k+1} \in \mathbb{P}_{\Omega}(\theta^k_{h,\tau} - \gamma_k \nabla J(u_{h,\tau}^{k+1}, p_{h,\tau}^{k+1}, \theta^k_{h,\tau}))$, $\gamma_k \in \{1, \frac{1}{2}, \frac{1}{4}, \ldots\}$

end

if $J(\theta^{k+1}_{h,\tau}) < \epsilon$

exit

A histological image $I$ of taken at a specific time $T$ serves as the target state $O \approx \hat{u}(T)$ for the terminal optimal control problem (13). It represents the observation data, based on which the above algorithm computes the optimal parameters for the model. The raw images $I$ cannot be used directly in the optimization algorithm, but instead each need to be transformed into an image that represents the non-dimensionalized tumor...
### Table 1 Simulation parameters

#### Numerical parameters

| Parameter                  | Value     |
|----------------------------|-----------|
| T (Total time)             | 10        |
| τ (Temporal step size)     | 0.1       |
| $h_{x_1}$ (Spatial step size along $x_1$) | 0.1       |
| $h_{x_2}$ (Spatial step size along $x_2$) | 0.1       |
| $N_{x_1}$ (Grid resolution along $x_1$) | N (image col size) |
| $N_{x_2}$ (Grid resolution along $x_2$) | M (image row size) |
| $u_{1,0}$ (initial value for $u_1$) | 0.2       |
| $u_{2,0}$ (initial value for $u_2$) | 0.5       |
| $\theta^0_i$ (initial value for $\theta_i$ for $i \in \{1, \ldots, 6\}$) | [0.001, 0, 0, 0.02, 0.5, 0.01]$^T$ |
| $\lambda_i$ (regularization parameter for $\theta_i$ for $i \in \{1, \ldots, 6\}$) | $10^{-4}$ |

### Table 2 Model parameters

#### Scalar factors for migration coefficients

**Phenomenological relevance**

| Factor | Value     |
|--------|-----------|
| $\gamma_\kappa$ | Speed of pH-taxis for cancer cells | .01 |
| $\gamma_\delta$ | Speed of advection for cancer cells | .001 |
| $\gamma_{pH}$ | Constant diffusion coefficient for protons | .01 |

**Box constraints**

| Parameter | Value     |
|-----------|-----------|
| $\sigma$ | ∈ [0.001, 0.01] |
| $\kappa$ | ∈ [−0.01, 0.01] |
| $\delta$ | ∈ [−0.01, 0.01] |
| $\mu$ | ∈ [0.001, 10] |
| $\alpha$ | ∈ [0.001, 10] |
| $\beta$ | ∈ [0.001, 10] |

...This pre-processing step is performed using Algorithm 2. Once again, the algorithm can be viewed as a mapping $I \mapsto \Psi(I) =: O$, $\Psi : L^2(\Omega) \to H^2(\Omega)$, that takes a raw data $I \in L^2(\Omega)$ as input and transforms it to an observation variable $O \in H^2(\Omega)$. The final processed image data $O$ represents normalized volumetric concentration of the cancer cells. Thus it serves as a valid measurement for the non-dimensionalized model (7).

#### 4.1 Evaluation of the optimization algorithm

In this section we numerically investigate the minimizing properties of the Algorithm 1. To this end we consider different noisy perturbations of a fixed target image and evaluate the obtained outputs of the Algorithm 1. For the target image we consider the processed image $O = \Psi(I)$ (Fig. 2a) obtained after applying the Algorithm 2 to the raw image $I$ (Fig. 1a). Let $G_{k,s}$ be the discrete Gaussian filter with kernel size $k$ and sigma (standard deviation) $s$ and $D_n$ to be the $n$-fold down sampling filter. Then a different perturbed version $O_k$ of $O$ is obtained applying $G_{k,s}$ and $D_n$ for
Algorithm 2: Preprocessing steps

**Data:** \( T \): an RGB image of the tissue, with size \( M \times N \)

1. \( I_g = \text{gray}(T) \). Convert RGB to grayscale image
2. \( Ig_s = G * Ig \). Smoothen the image using a Gaussian filter \( G \)
3. \( Ig_s = M * Ig_s \). Remove ‘salt and pepper’ noise by applying a median filter
4. Generate an image mask \( m \) by applying binary thresholding and performing morphological operation:
   - 4.1 apply binary thresholding to extract dominant features
   - 4.2 perform morphological open operation to remove isolated features. This results in the required mask \( m \)
5. \( Igsm = Ig_s(m) \land Ig_s(m) \). Perform bitwise ‘and’ operation of the smoothened gray image with itself using the mask \( m \).
6. \( Igsmi = 1 - Igsm / 255 \). Normalize the image.
7. \( O = Igsmi[::hx, ::hy] \). Downsample the \( M \times N \) image to an \( m \times n \) image.

different values of \( k, s \) and \( n \). Based on this, Fig. 2b is obtained as \( O_1 = G_{1.5}^1(D_4(O)) \). Similarly, Fig. 2c, d are obtained as \( O_3 = G_{3.3}^3(D_4(O)) \) and \( O_5 = G_{5.1}^5(D_4(O)) \) respectively. Due to the smoothing property of the Gaussian filter, increasing the kernel size and sigma results in smoother images, i.e. dampens spatial noise. As a result, we obtain that \( O_5 \) is smoother than \( O_3 \), which is in turn smoother than \( O_1 \).

Now applying the minimization algorithm \( E \) (Algorithm 1) to these perturbed inputs we can gauge its performance. To this end, by letting \( (\hat{u}_k, \hat{\theta}_k) = E(O_k) \), we define the following error metrics:

\[
\begin{align*}
    e_{2}^k &= \| \hat{u}_k(T) - O_k \|_{L^2}, &
    e_{\infty}^k &= \| \hat{u}_k(T) - O_k \|_{L^\infty} \\
    e_{\text{rel}}^k &= \frac{\| \hat{u}_k(T) - O_k \|_{L^2}}{\| O_k \|_{L^2}}, &
    e_{\Delta}^k(\epsilon) &= \frac{1}{|\Delta|} \int_{\Delta} \mathbb{1}_{\{e_2^k > \epsilon\}} \, dx.
\end{align*}
\]

Figure 3 depicts the error reduction profiles corresponding to the noisy target images \( O_1, O_3, O_5 \) (Fig. 2b–d). Based on this we can infer the following:

(1) as can be seen from Fig. 3b, the absolute error \( e_2^k \) tends to a stable low value for each \( O_k \). It holds that \( e_2^{k_1} < e_2^{k_2} \) when \( O_{k_1} \) is smoother than \( O_{k_2} \).

(2) according to Fig. 3a, for smoother target images the error reduction is relatively faster, especially for \( e_\infty \) and \( e_{\Delta} \).

The deterioration of error reduction for increased noise levels is expected and justified since (based on Theorem 5) we expect \( \hat{u} \in U \), i.e. \( \hat{u}(t) \in H^2(\Delta) \) for all \( t \in [0, T] \). Thus for a noisy target pattern, the optimization can only be suboptimal due to the violation of spatial smoothness.

4.2 Pseudopalisade specific results

In this section we evaluate the ability of the model to replicate different realistic pseudopalisade patterns \( O_n \) as shown in Fig. 1. We also consider other target patterns which are displayed in pre-processed form. The generated optimal final distribution
Fig. 2 Pre-processed version of the raw image \( I \) (cf. Figure 1a) and its noisy perturbations (see text)

Fig. 3 Error reduction for noisy target images obtained using different standard deviation parameter \( sd = \sqrt{s} \in \{.2, .6, 1, 1.4\} \) in the smoothing kernel \( G \) and downscalings \( D_n \) with \( n \in \{1, 3, 5, 7\} \)

of tumor cells corresponding to \( O_n \) is as shown in Figs. 4, 5, 6, and 7. Based on these outputs we can observe the following:

1. The optimization algorithm is able to accurately generate/recreate the target pseudopalisade pattern. This is evident by looking at the fourth column of the Figs. 4, 5, 6, and 7, where we depicted the \( L^2 \)-norm of the error i.e. the difference between the estimated final tumor density and the required target density. The difference is mainly in the 2nd decimal and only for very small volume fractions. Also, it is important to notice that the estimated final tumor cell density is much smoother when compared to the required target density. This is a consequence of the wellposedness Theorem 5 which stipulates that the solution \( \hat{u} \) lies in \( U \) with \( \hat{u}(t) \in H^2(\mathcal{D}) \) for every \( t \in [0, T] \).

2. Along with the cancer cell density, the algorithm also estimates the acid distribution. This is depicted in the third column of Figs. 4, 5, 6, and 7. Based on this we see that, at the regions of higher cancer cell density, in particular at areas of pseudopalisade formation, the proton concentration is relatively low compared to that of the surroundings. This supports the common hypothesis that the center of a pseudopalisade is a necrotic region, with poor acid removal mechanisms, which results in relatively low pH.

3. Another interesting observation is that certain localized areas in the interior region encompassed by a pseudopalisade structure, show relatively high proton concentration. This suggests that these localized regions were the sites of high tumor activity which was likely to be a consequence of increased glycolysis activity followed by growth and expansion of the tumor front. As a consequence of the excess acid produced and the expanding tumor periphery, acid gets accumulated, primarily in areas of poor vasculature such as the core of the pseudopalisade.
In order to get a deeper understanding about the formation process of pseudopalisade structures, we look at the estimated model parameter function $\theta$. We do so for a fixed target pattern, namely for Pattern-B (see Figs. 1a and 5). The obtained model parameter functions are depicted in Figs. 8, 9. Based on the dynamics of the parameters itself we can infer the following:

1. The tumor growth rate $\mu$ and acid expulsion rate $\alpha$ resemble structurally very much the target pattern. Initially the growth and expulsion rates are relatively high and later near the end time, when the tumor distribution is approaching the required pseudopalisade pattern, these rates stabilize. Moreover, it can be observed that $\mu$ and $\alpha$ are positively correlated i.e. higher $\mu$ implies higher $\alpha$, at least at the beginning of tumor evolution. This positive feedback of growth rate and acid buffering
indicates the presence of (reminiscent) vasculature and the supporting microenvironment to facilitate removal excess acid. Disruption of this vital supporting element during tumor progression results in the formation of necrotic regions like those appearing toward the end time.

2. The acid production rate $\beta$ is higher mainly in the regions where there is less tumor cell density. These regions of higher $\beta$ mainly happen to be the area of tissue necrosis, leading to the eventual accumulation of acid. These above average acid production rates could be attributed to the neoplastic transformation in those regions where excessive glycolysis takes place, in order to fulfill the energy requirements for proliferation.

3. Looking at the migration parameters we see that the diffusion coefficient $\sigma$ is lower in the sparsely populated tumor regions which are the main candidate areas for the necrotic core formation. As the tumor progresses, the diffusion coefficient mainly homogenizes and can be approximated to be spatially constant.

4. The advection coefficient $\kappa$ is initially more pronounced at the outer margin of the necrotic core which later progresses to the inner region of the core. This indicates that an unfavorable region is likely to generate an aggressive stimulus making the tumor cells more mobile.

5. The pH-taxis coefficient $\delta$ is mainly at the outer edges of areas with acid accumulation, which corresponds to necrosis. Thus, pH-taxis seems to act mainly at the interfacing/intersecting layer of high-density and low-density regions of tumor. This suggests that acidity facilitates travelling-wave like behavior of tumor-host interface.

6. Finally, the clear distinction between regions where the taxis and growth parameters are dominant supports the hypothesis of grow-or-go dichotomy in glioma tumor progression (Höring et al. 2012).

5 Pattern synthesis and disruption

In this section we consider an application of the OCP (7), aiming to shed light on pattern dynamics under various influences. One of the main advantages of data-based parameter estimation is its potential use in patient-specific therapy design. The diagnostic histological samples obtained from a patient can be used to estimate the model parameters which approximately characterize the (microscopic) dynamics of GBM progression for that specific patient. Based on this, one can then design or hypothesize different intervention mechanisms that can mitigate the development of GBM. One such plausible way would be to understand how pseudopalisade pattern formation can be disrupted or reversed. To incorporate such effects, we adapt the above model as follows:

$$\begin{align*}
d_t u_1 - \nabla \cdot (\sigma \nabla u_1 + u_1 \nabla \kappa) &= \nabla \cdot (\delta u_1 \nabla u_2) + \mu f_1(u_1, u_2) - \xi_1 u_1 \\
d_t u_2 - \Delta u_2 + \alpha u_2 &= \beta f_2(u_1, u_2) + \xi_2 u_2 \\
(\sigma \nabla u_1 + u_1 \nabla \kappa) \cdot \hat{n} &= 0, \quad \nabla \cdot u_2 = 0, \quad u_1(0) = u_{10}, \quad u_2(0) = u_{20}
\end{align*}$$
Here, $\xi = (\xi_1, \xi_2) \in L^2(I; V^{\times 2})$ with $\xi_1 \geq 0$ is a disturbance term that models the disruption of pattern formation mechanisms. Consequently, $\xi$ is responsible for neutralization or renormalization of the tumor microenvironment, which altogether impedes tumor development. Motivated by this we refer to $\xi$ as a pattern neutralizing function. Having estimated the model parameters $\theta$ for different target patterns (see pictures in Fig. 1) we can now ask what kind of external signal is needed to revert or neutralize the cancerous microenvironment. This entails solving the following modified optimization problem:

$$u^*, \xi^* = \arg\min_{u, \xi} J(u, \theta, \xi) \quad \text{s.t.} \quad G(u; \theta, \xi) = 0,$$

where $J$ is the quadratic cost function analogous to (12) and $G(u; \theta, \xi) = 0$ represents the state Eq. (36) in abstract form. Now, given a non-cancerous or a neutral tissue pattern as the target state, we can solve for the optimal pattern neutralizing function $\xi^*$. To be more precise, let $\theta_O$ be the optimal model parameter vector associated to the target pattern Pattern-$O$ where $O \in \{A, B, C, E\}$ i.e. one of the target patterns depicted in Fig. 1. Then given some starting value $u_0$ (representing a non-cancerous initial state) and fixing the model parameters $\theta_O$, we apply Algorithm 1 to determine $\hat{\xi}$ that can counteract the effects of $\theta_O$ and result in a target state which corresponds to a neutral non-cancerous tissue. Figures 10 and 11 depict the result of the application of the neutralizing function $\xi^*$ that is able to counteract the effects of the parameters.
Fig. 9 Evolution of tumor motility parameters. The columns are sorted according to time (in days). The leftmost column is the tumor state at $t = 20$, the middle left column is for $t = 50$, the middle right one for $t = 70$, and the rightmost column is for $t = 90$. The rows represent different parameter functions of the model (7). Arranged from top to bottom these are: the diffusion coefficient $\sigma$ (1st row), advection coefficient $\kappa$ (2nd row) and pH-taxis coefficient $\delta$ (bottom row).

Fig. 10 Final pseudopalisade patterns resulting after applying the pattern neutralizing function $\hat{\xi}$ with the corresponding pattern specific estimated model parameters $\hat{\theta}$.

Fig. 11 Final proton distribution corresponding to the pseudopalisade patterns in Fig. 10.

Alternatively, instead of applying the pattern neutralizing function that directly acts on the cancer cells, one could also look for an indirect method which aims at manipulating the micro-environment as a means to hinder tumor progression. This
means that instead of having two control variables $\xi_1$ and $\xi_2$ in (36) we have only one control variable $\hat{\xi}_2$ that aims to disrupt the pseudopalisade pattern by appropriately regulating the tissue acidity. Consequently, $\hat{\xi}_2$ is referred to as pH neutralizing function. Following the above steps we can find the optimal $\hat{\xi}_2$ that alone can mitigate the effects of pseudopalisade forming parameters $\hat{\theta}_0$. Based on the results depicted in Figs. 12 and 13, we can see that neutralizing the tissue acidity can also serve as an effective pseudopalisade disruption mechanism. This seems to be in line with therapeutic approaches aiming at tumor alkalinization, e.g. see (Amiri et al. 2016; Yang et al. 2020).

5.1 Synthesis of new patterns

In this section we illustrate how new patterns can be synthesized by combining, linearly or nonlinearly, the optimal parameters $\hat{\theta}_O$ corresponding to different target pseudopalisade patterns $O$. We let $\hat{\theta}_{KL} := (\hat{\theta}_K + \hat{\theta}_L)/2$, where $\hat{\theta}_K$ and $\hat{\theta}_L$ are the estimated optimal model parameters for target patterns Pattern-K and Pattern-L, respectively. The patterns generated by different such combinations are depicted in Figs. 15, and 16. They indicate that different complex patterns can arise by linear combination of processes responsible for generating simpler patterns. This is again of particular importance for therapy applications, where one can ask the question whether a similar linear combination of pattern neutralizing functions can still be effective for disrupting the new pattern. That is, if $\hat{\xi}_K$ and $\hat{\xi}_L$ are the pattern neutralizing functions that can counteract the effects of $\hat{\theta}_K$ and $\hat{\theta}_L$ respectively, would their combination $\hat{\xi}_{KL} := (\hat{\xi}_K + \hat{\xi}_L)/2$ be able to counteract the effects of $\hat{\theta}_{KL}$? Based on the obtained numerical results, as shown in Figs. 17, 18, we can conclude that the same linear combination of pattern neutralizing functions does indeed prove to be effective in counteracting the combined
effects of the processes responsible for generating simpler pseudopalisade patterns (Fig. 13).

6 Summary and conclusion

In this paper we have formulated a terminal valued optimal control problem to understand the process involved in the formation of pseudopalisade structures during the progression of GBMs based on observed data. Starting from the state of the art multiscale model (Kumar et al. 2021) we proposed a modified model for the dynamics of pseudopalisade structures under the influence of tissue acidity which served as a dynamical state equation for the optimization problem. We then performed a well-
posedness study based on which we are able to establish the existence of an optimal solution to the TOCP. This paved the way to performing numerical simulations for which we used the well known projected gradient descent method. For solving the TOCP problem the required data was obtained by taking experimentally observed images from available publications. Here we also proposed an algorithm to convert raw experimental images to model-specific non-dimensionlized volumetric/concentration data. By using these processed images, we were able to successfully recreate the target patterns by estimating the optimal model parameter functions. Because most of the developed mathematical models only rely on simulation results to reproduce experimentally observed qualitative behavior, the proposed procedure provides an effective alternative approach to validate the model using real data. Based on the target-specific optimal parameters we were able to shed light not only on the dynamical interplay between reaction and migration terms, but also on the relationship between tumor progression and acidity. This type of data-specific analysis of dynamics could be of particular interest to medical professionals to perform patient-specific diagnosis and in turn design patient-specific treatment. From this perspective, we also showed the feasibility of different methods to normalize the tissue structure and obstruct tumor progression. The computed pattern neutralizing function achieves this not only by modifying the tissue acidity, but also by directly acting on the cancer cell population. Additionally, we highlighted the strength of this data-based approach by its ability to synthesize different unobserved pseudopalisade patterns, by simply combining already known optimal parameter functions computed from specific observed data. This can be further used to determine probable pattern-neutralization functions for a new unobserved pseudopalisade pattern by combining, in a similar way, the pattern-neutralizing functions of the simpler ones. Further direction of research would be to devise an active control strategy and a corresponding control problem for obstructing GBM progression.

**Supplementary Information**  The online version contains supplementary material available at https://doi.org/10.1007/s00285-023-01933-5.

**Declarations**

**Conflict of interest**  The authors declare that they have no competing interests.
Appendix

In the following we display the evolution of the optimal parameters $\theta^*$ for Patterns-A, C and E. Furthermore, the figures and results presented in this work are submitted as supplementary files (See Figs. 19, 20, 21, 22, 23, 24).

**Fig. 19** Evolution of growth and decay parameter functions for target Pattern-A (see Fig. 4). The columns are sorted according to time (in days). The leftmost column is the tumor state at $t = 20$, the middle left column is for $t = 50$ days, the middle right one for $t = 70$ days, and the rightmost column for $t = 90$. The rows represent different parameter functions of the model (7). Arranged from top to bottom these are: the growth coefficient $\mu$ (1st row), acid removal rate $\alpha$ (2nd row), acid production rate $\beta$ (3rd row).
**Fig. 20** Evolution of tumor motility parameters for target Pattern-A (see Fig. 4). The columns are sorted according to time (in days). The leftmost column is the tumor state at $t = 20$, the middle left column is for $t = 50$, the middle right one for $t = 70$, and the rightmost column is for $t = 90$. The rows represent different parameter functions of the model (7). Arranged from top to bottom these are: the diffusion coefficient $\sigma$ (1st row), advection coefficient $\kappa$ (2nd row) and pH-taxis coefficient $\delta$ (bottom row).

**Fig. 21** Evolution of growth and decay parameter functions for target Pattern-C (Fig. 1b). The columns are sorted according to time (in days). The leftmost column is the tumor state at $t = 20$, the middle left column is for $t = 50$ days, the middle right one for $t = 70$ days, and the rightmost column for $t = 90$. The rows represent different parameter functions of the model (7). Arranged from top to bottom these are: the growth coefficient $\mu$ (1st row), acid removal rate $\alpha$ (2nd row), acid production rate $\beta$ (3rd row).
Fig. 22 Evolution of tumor motility parameters for target Pattern-C (Fig. 1b). The columns are sorted according to time (in days). The leftmost column is the tumor state at $t = 20$, the middle left column is for $t = 50$, the middle right one for $t = 70$, and the rightmost column is for $t = 90$. The rows represent different parameter functions of the model (7). Arranged from top to bottom these are: the diffusion coefficient $\sigma$ (1st row), advection coefficient $\kappa$ (2nd row) and pH-taxis coefficient $\delta$ (bottom row).

Fig. 23 Evolution of growth and decay parameter functions for target Pattern-E (Fig. 1c). The columns are sorted according to time (in days). The leftmost column is the tumor state at $t = 20$, the middle left column is for $t = 50$ days, the middle right one for $t = 70$ days, and the rightmost column for $t = 90$. The rows represent different parameter functions of the model (7). Arranged from top to bottom these are: the growth coefficient $\mu$ (1st row), acid removal rate $\alpha$ (2nd row), acid production rate $\beta$ (3rd row).
Fig. 24 Evolution of tumor motility parameters for target Pattern-E (Fig. 1c). The columns are sorted according to time (in days). The leftmost column is the tumor state at $t = 20$, the middle left column is for $t = 50$, the middle right one for $t = 70$, and the rightmost column is for $t = 90$. The rows represent different parameter functions of the model (7). Arranged from top to bottom these are: the diffusion coefficient $\sigma$ (1st row), advection coefficient $\kappa$ (2nd row) and pH-taxis coefficient $\delta$ (bottom row).

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