Mathematical Modelling and Optimal Control of Anthracnose

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Abstract—In this paper we propose two nonlinear models for the control of anthracnose disease. The first is an ordinary differential equation (ODE) model which represents the within-host evolution of the disease. The second includes spatial diffusion of the disease in a bounded domain. We demonstrate the well-posedness of those models by verifying the existence of solutions for given initial conditions and positive invariance of the positive cone. By considering a quadratic cost functional and applying a maximum principle, we construct a feedback optimal control for the ODE model which is evaluated through numerical simulations with the scientific software Scilab\(^\text{R}\). For the diffusion model we establish under some conditions the existence of a unique optimal control with respect to a generalized version of the cost functional mentioned above. We also provide a characterization for this optimal control.

KeyWords— Anthracnose modelling, nonlinear systems, optimal control. AMS Classification—49J20, 49J15, 92D30, 92D40.

I. INTRODUCTION

Anthracnose is a phytopathology which attacks a wide range of commercial crops, including almond, mango, banana, blueberry, cherry, citrus, coffee, hevea and strawberry. The disease has been identified in such diverse areas as Ceylon (1923), Guadeloupe (1925), Sumatra (1929), Indochina (1930), Costa Rica (1931), Malaysia (1932), Java (1933), Madagascar (1934), Cameroon (1934), Colombia (1940), Salvador (1944), Brazil (1946), Nyassaland (1949), New Caledonia (1954), and Arabia (1956)\(^[5]\). Anthracnose can affect various parts of the plant, including leaves, fruits, twigs and roots. Possible symptoms include defoliation, fruit rot, fruit fall and crown root rot, which can occur before or after harvest depending on both pathogen and host\(^[5], [28]\).

The Anthracnose pathogen belongs to the Colletotrichum species (acutatum, capsici, gloeosporioides, kahawae, lindemuthianum, musae, ...). Colletotrichum is an ascomycete fungus. It can
reproduce either asexually or sexually, but sexual reproduction is rare in nature \[28\]. Favourable growth conditions occur particularly in tropical zones. Rainfall, wetness and altitude are all conducive to sporulation and conidia spreading \[18\], \[23\]. Sources of inoculum are thought to be leaves, buds and mummified fruits.

**A. Anthracnose pathosystem**

The process of infection by *Colletotrichum* species can usually be divided into at least seven steps, depending on various factors including growth conditions, host tissues and involved species. Conidia deposited on the host attach themselves on its surface. The conidia germinate after 12–48 hours, and appressoria are produced \[5\], \[15\]. Several studies on infection chronology show that appressoria production can occur between 3–48 hours following germination under favourable conditions of wetness and temperature \[18\], \[28\]. The pathogen then penetrates the plant epidermis, invades plant tissues, produces acevuli and finally sporulates. The penetration of plant epidermis is enabled by a narrow penetration peg that emerges from the appressorium base \[7\]. In some marginal cases penetration occurs through plant tissues’ stomata or wounds. Once the cuticle is crossed, two infection strategies can be distinguished: intracellular hemibiotrophy and subcuticular intramural necrotrophy, as shown in Figure 2. Invasion of the host is led through formation of hyphae which narrow as the infection progresses. *Colletotrichum* produce enzymes that degrade carbohydrates, dissolve cell walls, and hydrolyze cuticle. Some of those enzymes are polygalacturonases, pectin lyases and proteases. Some hosts may employ various biochemical strategies to counter the pathogen. For example, the peel of unripe avocados has been found in vitro to contain a preformed antifungal diene (cis, cis-1-acetoxy-2-hydroxy-4-oxo-heneicosa-12, 15-diene) that inhibits the growth of *Colletotrichum gloeosporioides* when present above a certain concentration \[28\].

**B. Models in the literature**

Most previous mathematical studies on *Colletotrichum*-host pathosystem have focused on forecasting disease onset based on environmental factors affecting host sensitivity. DANNEBERGER et al. in \[9\] have developed a forecasting model for the annual bluegrass anthracnose severity index, using weather elements such as temperature and wetness. Their model is a quadratic regression

\[
ASI = a_0 + a_{0.1}W + a_{1.0}T + a_{1.1}T \times W + a_{0.2}T^2 + a_{2.0}W^2
\]

where $ASI$ is the anthracnose severity index, $T$ is the daily average temperature and $W$ is the average number of hours of leaves’ wetness per
day. DODD et al. in [10] have studied the relationship between temperature \((T)\), relative humidity \((H)\), incubation period \((t)\) and the percentage \((p)\) of conidia of Colletotrichum gloeosporioides producing pigmented appressoria on one month old mangoes. They used the following logistic model:

\[
\ln(p/(1-p)) = a_0 + a_{0.1}H + a_{1.0}T + a_{0.2}H^2 + a_{2.0}T^2 + b\ln(t)
\]

DUTHIE in [12] examines the parasite’s response \((R)\) to the combined effects of temperature \((T)\) and wetness duration \((W)\). That response could be the rate of germination, infection efficiency, latent period, lesion density, disease incidence or disease severity. Several models are discussed, the two principal being

\[
R(T, W) = f(T) \left[1 - \exp\left(-[b(W - c)]^d\right)\right]
\]

and

\[
R(T, W) = a \left[1 - \exp\left(-[f(T)(W - c)]^d\right)\right],
\]

where

\[
f(T) = \frac{e(1 + h)h^{\frac{b}{2}}}{(1 + \exp(g[T - f]))} \exp\left(\frac{g[T - f]}{1 + h}\right)
\]

and

\[
a > 0, \quad b > 0, \quad W \geq c \geq 0, \quad d > 0, \quad e > 0, \quad f \geq 0, \quad g > 0, \quad h > 0.
\]

MOUEN et al. attempt in [17] to develop a spatio-temporal model to analyse infection behaviour with respect to the time, and identify potential foci for disease inoculum. Logistic regression and kriging tools are used used. In addition to these references, there are several other statistical models in literature [12], [17], [18], [19], [20], [21], [28].

C. Controlling anthracnose

There are many approaches to controlling anthracnose diseases. The genetic approach involves selection or synthesis of more resistant cultivars [3], [4], [5], [14], [27]. Several studies have demonstrated the impact of cultivation practices on disease dynamics [19], [20], [21], [28]. Other tactics may be used to reduce predisposition and enhance resistance, such as pruning old infected twigs, removing mummified fruits, and shading [5]. Biological control uses microorganisms or biological substrates which interact with pathogen or induce resistance in the host [11]. Finally there is chemical control, which requires the periodic application of antifungal compounds [5], [22], [24]. This seems to be the most reliable method, though relatively expensive. The best control policy should schedule different approaches to optimize quality, quantity and cost of production. Note that inadequate application of treatments could induce resistance in the pathogen [26].

D. Organization of the paper

The remainder of this paper is organized as follows. In section [II] we propose and study a within-host model of anthracnose. We present that model and give parameters meaning in subsection [II-A]. Throughout subsection [II-B] we establish the well-posedness of the within-host model both in mathematical and epidemiological senses. The optimal control of the model is surveyed in subsection [II-C] and numerical simulations are performed in the last subsection [II-D]. We make a similar study on a spatial version of the model includdling a diffusion term in section [III]. That last model is presented in subsection [III-A]. Studies on its well-posedness and its optimal control are made respectively in subsections [III-B] and [III-C]. Finally, in section [IV] we discuss our modelling and some realistic generalizations which could be added to the model.

II. A WITHIN-HOST MODEL

A. Specification of the within-host model

The detrimental effects of Colletotrichum infection on fruit growth are closely related to its life cycle. It is mathematically convenient to express these effects in terms of the effective inhibition rate (denoted by \(\theta\)), which is a continuous function of time. The effective inhibition rate is defined such that the maximum attainable fruit volume is reduced by a factor \(1 - \theta\) if current infection conditions are maintained. In addition to \(\theta\), the other time-dependent variables in the model are
host fruit total volume and infected volume, denoted by \( v \) and \( v_r \) respectively. We have on the set \( S = \mathbb{R}_+ \setminus \{1\} \times \mathbb{R}_+ \times \mathbb{R}_+ \) the following equations for the time-evolution of the variables \((\theta, v, v_r)\):

\[
\begin{aligned}
    d\theta/dt &= \alpha(\theta) \left( 1 - \theta / (1 - \theta, u(t)) \right) \\
    dv/dt &= \beta(\theta) \left( 1 - v_\theta / ((1 - \theta) \eta(t) v_{\max}) \right) \\
    dv_r/dt &= \gamma(\theta) \left( 1 - v_r/v \right)
\end{aligned}
\]

(1)

The parameters in (1) have the following practical interpretations:

- \( \alpha, \beta, \gamma \) characterize the effects of environmental and climatic conditions on the rate of change of inhibition rate, fruit volume, and infected fruit volume respectively. These are all positive functions of the time \( t \) and inhibition rate \( \theta \).
- \( \gamma \) is an increasing function with respect to \( \theta \) and satisfies \( \gamma(t, 0) = 0, \forall t \geq 0 \).
- \( u \) is a measurable control parameter which takes values in the set \([0, 1]\).
- \( 1 - \theta_1 \in [0, 1] \) is the inhibition rate corresponding to epidermis penetration. Once the epidermis has been penetrated, the inhibition rate cannot fall below this value, even under maximum control effort. In the absence of control effort \((u(t) = 0)\), the inhibition rate increases towards 1.
- \( \eta \) is a function of time that characterizes the effects of environmental and climatic conditions on the maximum fruit volume. Its range is the interval \([0, \theta_2]\).
- \( v_{\max} \) represents the maximum size of the fruit.
- \( 1 - \theta_2 \in [0, 1] \) is the value of inhibition rate \( \theta \) that corresponds to a limiting fruit volume of \( \eta v_{\max} \). According to the second equation in (1), the limiting volume size is \( \eta v_{\max} (1 - \theta) / \theta_2 \leq v_{\max} \). When the volume is less than this value, it increases (but never passes the limiting value); while if the volume exceeds this value, then it decreases. This limiting value for \( v \) is less than \( \eta v_{\max} \) when \( \theta > 1 - \theta_2 \) (note \( \eta \leq \theta_2 \leq 1 \)).

Note that equations (1) are constructed so that \( v \leq v_{\max} \) and \( v_r \leq v \) as long as initial conditions satisfy these inequalities.

With the definitions

\[
A \equiv \begin{bmatrix}
    -\alpha(t, \theta) & 0 & 0 \\
    0 & -\frac{\theta_2 \beta(t, \theta)}{(1 - \theta) \eta(t) v_{\max}} & 0 \\
    0 & 0 & -\frac{\gamma(t, \theta)}{v}
\end{bmatrix},
\]

\[
B \equiv \begin{bmatrix}
    \alpha(t, \theta) & \beta(t, \theta) & \gamma(t, \theta)
\end{bmatrix}^T,
\]

and

\[
X \equiv \begin{bmatrix}
    \theta & v & v_r
\end{bmatrix}^T,
\]

then model (1) can be reformulated as

\[
dX/dt = F(t, X),
\]

(2)

where

\[
F(t, X) \equiv A(t, X, u) X + B(t, X).
\]

(3)

As indicated above, model (1) is an exclusively within-host model, and as such does not include the effects of spreading from host to host. (In Section III we propose a diffusion model for between-host spreading.) Such a model has several practical advantages. In practice, monitoring of the spreading of the fungi population is difficult. Furthermore, conidia sources and spreading mechanisms are not well-understood, although the literature generally points to mummified fruits, leaves and bark as sources of inoculum. Instead of controlling the host-to-host transmission, an alternative control method is to slow down the within-host fungi evolution process. Such an approach enables the use of statistical methods, since large samples of infected hosts may easily be obtained.

B. Well-posedness of the within-host model

In the following discussion, we demonstrate that model (1) is well-posed both mathematically and epidemiologically, under the following standard technical assumptions:

- \((H1)\) The control parameter \( u \) is measurable.
- \((H2)\) The function \( F \) is continuous with respect to the variable \( X \).
(H3) For every compact subset $K \subset S$, there is an integrable map $M_K : \mathbb{R}_+ \to \mathbb{R}_+$ such that for every $X$ in $K$ and $t$ in $\mathbb{R}_+$, $\|F(t, X)\|_S \leq M_K(t)$.

Existence of a solution is guaranteed by the following proposition, which follows from a simple application of the Carathéodory theorem.

**Proposition 1.** For every initial condition $(t_0, X_0)$ in $\mathbb{R}_+ \times S$ there is a function $X(t_0, X_0, t)$ which is absolutely continuous and satisfies (2) for almost any time $t \in \mathbb{R}_+$.

Uniqueness and smoothness of the solution may be established using the Cauchy-Lipschitz Theorem, based on properties (H2) and (H3) of the function $F$.

Next we establish positive invariance of the set $S$, and the positive invariance of a bounded subset $BS$. These results are needed to show consistency of the biological interpretation of the solution, as explained below. With the definitions

$$A_1 \equiv \begin{bmatrix} \frac{\alpha(t, \theta)}{(1-\theta)u(t)} & 0 & 0 \\ 0 & -\frac{\theta_2}{(1-\theta)\eta(t)v_{\text{max}}} & 0 \\ 0 & 0 & -\frac{1}{v} \end{bmatrix},$$

$$A_2 \equiv \begin{bmatrix} \alpha(t, \theta) & 0 & 0 \\ 0 & \beta(t, \theta) & 0 \\ 0 & 0 & \gamma(t, \theta) \end{bmatrix},$$

$$B_1 \equiv [1 \ 1 \ 1]^T,$$

and $X$ as defined above, then model (1) can be reformulated as

$$\frac{dX}{dt} = A_2 (A_1 X + B_1).$$

**Theorem 2.** The set $S$ is positively invariant for the system (4).

*Proof:* A solution to (4) satisfies for every time $t \geq 0$,

$$X(t) = \exp \left[ \int_0^t A_2(s) \cdot A_1(s) \, ds \right] X(0) + \int_0^t \exp \left[ \int_s^t A_2(\xi) A_1(\xi) \, d\xi \right] A_2(s) B_1 \, ds$$

Since $-A_2(s)A_1(s)$ is a $M$-matrix for every time $s \geq 0$, $\exp \left[ \int_t^s A_2(\xi) A_1(\xi) \, d\xi \right]$ is a positive matrix. Moreover, since $B_1$ is nonnegative, one can conclude that $X$ remain nonnegative when $X(0)$ is taken nonnegative.

**Theorem 3.** Let $BS$ be the subset of $S$ defined such as

$$BS = \{(\theta, v, v_r) \in \mathbb{R}^3; 0 \leq \theta < 1, 0 < v \leq v_{\text{max}}, 0 \leq v_r \leq v\}$$

Then $BS$ is positively invariant for system (4).

*Proof:* We will show that at each point of the boundary of $BS$, the system (4) returns into $BS$. We prove this by showing that the scalar product of the system time derivative with the normal vector $n$ at each boundary point is nonpositive. It has been already shown that positive orthant is positively invariant. Let

$$F_1 \equiv \{(\theta, v, v_r) \in BS; \theta = 1\}$$

$$F_2 \equiv \{(\theta, v, v_r) \in BS; v = v_{\text{max}}\}$$

$$F_3 \equiv \{(\theta, v, v_r) \in BS; v_r = v\}$$

For all points on $F_1$, $n$ can be chosen as $(1, 0, 0)$. Since the control $u$ takes its value in $[0, 1]$ which also contains $\theta_1$, $\frac{d\theta}{dt}$ is negative and the result is obtained. For all points on $F_2$, $n$ can be chosen as $(0, 1, 0)$. Thanks to definition of $\theta_2$ and $\eta$, $\frac{dv}{dt}$ is negative and the result is obtained. For all points on $F_3$, $n$ can be chosen as $(0, -1, 1)$. $\frac{dv_r}{dt}$ is zero, and consequently $F_3$ is positively invariant.

The invariance of the set $F_3$ is biologically plausible, since once the fruit is totally rotten it remain definitely in that state, the fruit is lost. The set $BS$ is also reasonable for biological reasons: the inhibition rate is bounded, the rotten volume is no larger than the total volume, and fungus attack reduces the size of a mature fruit.

**C. Optimal control of the within-host model**

In this subsection we apply control to model (1), which we repeat here for convenience:

\begin{align}
\frac{d\theta}{dt} &= \alpha(t, \theta) (1 - \theta / (1 - \theta u(t))) \\
\frac{dv}{dt} &= \beta(t, \theta) (1 - v / (1 - \eta(t) v_{\text{max}})) \\
\frac{dv_r}{dt} &= \gamma(t, \theta) (1 - v_r / v)
\end{align}

(5)
For the control problem we focus on the first equation. This equation is controllable in $[0,1]$ since $\theta$ is continuous and $1-\theta, u(t)$ is an asymptotic threshold which can be set easily. Given a time $T>0$ (for example the annual production duration) we search for $u$ in $L^2_{loc}(\mathbb{R}^+,[0,1])$ such that the following functional (previously used in [1], [13]) is minimized:

$$J_T(u) = \int_0^T \left( ku^2(t) + \theta^2(t) \right) dt + f(\theta(T)),$$

where $k > 0$ can be interpreted as the cost ratio related to the use of control effort $u$. This functional reflects the fact that reducing inhibition rate $\theta$ will lead to increased fruit production (larger volumes with a relatively lower level of infection), while minimizing $u$ will reduce financial and environmental costs. We use the squares of $u$ and $\theta$ in the integrand because this choice facilitates the technical calculations required for minimization.

We note in passing that we could had tried to minimize the more practically relevant expression:

$$\int_0^T \left( ku(t) + \theta(t) + \left( v_{\text{max}} - v(t) \right) + v_r(t) \right) dt + \theta(T) + \left( v_{\text{max}} - v(T) \right) + v_r(T)$$

However, an exact computation of this functional would require precise expressions for $\alpha, \beta, \gamma, \eta$ in the system [1]. As far as the authors know, there is no previous study which gives those parameters. It seemed more advantageous to us to limit the random choice of parameters, so that we could perform representative simulations.

We define the set

$$U^K = \left\{ u \in C([0,T]; [0,1]) : \forall t, s \in [0,T], \quad |u(t) - u(s)| \leq K |t-s| \right\}$$

which is nonempty for every $K \geq 0$.

**Theorem 4.** Let $K \geq 0$. There is a control $u^* \in U^K$ which minimizes the cost $J_T$.

**Proof:** Since $J_T \geq 0$ it is bounded below. Let the infimum be $J^*$, and let $(u_n)_{n \in \mathbb{N}}$ be a sequence in $U^K$ such that $(J_T(u_n))_{n \in \mathbb{N}}$ converges to $J^*$. The definition of $U^K$ implies that $(u_n)_{n \in \mathbb{N}}$ is bounded and uniformly equicontinuous on $[0,T]$. By the Ascoli theorem, there is a subsequence $(u_{n_k})$ which converges to a control $u^*$. Since the cost function is continuous with respect to $u$ it follows that $J_T(u^*) = J^*$.

**Theorem 5.** Suppose that $\alpha$ depends only on time. If there is an optimal control strategy $u$ which minimizes $J_T$, then $u$ is unique and satisfies

$$u(t) = \begin{cases} 1 & \text{when } 27\alpha \theta^2 \theta p \geq 8k; \\ \frac{\theta w_3(t)-1}{\theta w_3(t)} & \text{when } 27\alpha \theta^2 \theta p < 8k \end{cases}$$

where $w_3(t)$ is the element of $[1, \min\{3/2, 1/(1-\theta)\}]$ which is the nearest to the smallest nonnegative solution of the equation

$$\alpha \theta^2 \theta p w^3 - 2kw + 2k = 0$$

**Proof:** According the maximum principle, minimizing $J_T$ is equivalent to minimizing the Hamiltonian functional

$$H(t, \theta, u) = ku^2(t) + \theta^2(t) + f(\theta(T)) + \alpha(t) p(t) (1-\theta, u(t) - \theta) / (1-\theta, u(t))$$

where the adjoint state $p$ is the solution to the following problem

$$\begin{align*}
&\frac{dp}{dt} = \alpha(t) p (1-\theta, u(t) - \theta) - 2\theta \\
&p(T) = f'(\theta(T))
\end{align*}$$

To simplify the expression, let

$$w \equiv 1 / (1-\theta, u) \in [1,1/(1-\theta, u)].$$

Then the new equivalent functional to minimize is

$$J_T^1(w) = \int_0^T \left( k \left( \frac{w(t)-1}{\theta w(t)} \right)^2 + \theta^2(t) \right) dt + f(\theta(T))$$

If and only if

$$\alpha \theta^2 \theta p w^3 - 2kw + 2k = 0.$$
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where \( w_3 (t) \) is the element of \([1, \min \{3/2, 1/(1 - \theta_1)\}]\) that is the nearest to the smallest nonnegative solution of (10). It follows from the definition of \( w \) in (9) and algebraic rearrangement that the optimal control \( u \) is given by (6), where \((p, \theta)\) is a solution to the system (7). The uniqueness of \( u \) follows from the uniqueness of the solution of the system (7).

D. Computer simulations of the controlled within-host model

We performed simulations in order to demonstrate the practical controllability of the system. For these simulations we used an inhibition pressure of the form

\[
\alpha(t) = a(t - b)^2(1 - \cos(2\pi t/c)),
\]

with \( b \) and \( c \) in \([0, 1]\). This function reflects the seasonality of empirically-based severity index models found in the literature [9], [10], [12]. The particular values used in the simulation were \( a = 4, b = 0.75, c = 0.2 \) and \( k = 1 \). We also took \( f(\theta(T)) = \theta(T) \), so that \( p(T) = f'(\theta(T)) = 1 \). In this case and the shooting method can be used to numerically estimate the value of \( p_0 \) which produces a solution to (8).

Fig1: Optimum control effort over a one-year period \( \theta_0 = 0.2, \theta_1 = 1 - 0.4 \).

Fig2: Evolution of inhibition rate over a one-year period with \( \theta_0 = 0.2, \theta_1 = 1 - 0.4 \).

Fig3: Optimum control effort over a one-year period \( \theta_0 = 0.5, \theta_1 = 1 - 0.4 \).

Fig4: Evolution of inhibition rate over a one-year period \( \theta_0 = 0.5, \theta_1 = 1 - 0.4 \).

The above figures show how the control strategy adapts itself in response to inhibition pressure represented by \( \alpha \). Figures 1-2 correspond to the case
of low initial effective inhibition rate \((\theta_0 = 0.2,\) corresponding to a low initial level of infection),
while Figures 3-4 correspond to the case of high initial effective inhibition rate \((\theta_0 = 0.5).\) The
simulations also show in different cases the effectiveness of the optimal strategy as compared to
taking no control action. Regardless of whether the initial inhibition rate is above or below the
threshold \(1 - \theta_1,\) the dynamics are sensitive to the control effort.

III. A DIFFUSION MODEL

In this section, we will model the geographical spread of the disease via diffusive factors such as
the movement of inoculum through ground water and wind.

A. Specification of the diffusion model

We include the effect of diffusive factors on the spread of infection by adding a diffusion term to
the within-host equation for \(\theta\) from system (1). Together with boundary conditions, the model is
\[
\frac{\partial \theta}{\partial t} = \alpha(x) \left( (1 - \theta) - (1 - \theta) u(x) \right) + \text{div} \left( A(x) \nabla \theta \right)
\]
\[\text{on } \Omega, \quad \theta(0,x) = \theta_0(x) \geq 0 \quad \text{for each } x \in \Omega,
\]
\[\alpha(x) \left( \langle A(x) \nabla \theta, n \rangle = 0 \right) \quad \text{on } \partial \Omega,
\]
\[
\langle A(x) \nabla \theta, n \rangle = 0 \quad \text{on } \partial \Omega,
\]
\[\theta(x) = \theta_0(x) \geq 0 \quad x \in \Omega.
\]

where \(\Omega\) is an open bounded subset of \(\mathbb{R}^3\) with a continuously differentiable boundary \(\partial \Omega,\) and \(\theta_0 \in [0,1].\) For a given element \((t,x,\theta),\) \(A\) is a \(3 \times 3\)-matrix \((a_{ij}(t,x,\theta)).\) The functions \(\alpha\) and \(a_{ij}\) are assumed to be nonnegative. Since \(\theta\) depends
on \((t,x),\) the functions \(\alpha\) and \(a_{ij}\) can be identified with elements of the set
\(C([0,T]; H^1(\Omega))\). The function \(u \in C([0,T]; H^1(\Omega))\) designates the control, which takes its values in the set \([0,1].\)

We may thus study the system on each interval separately, and assume that all parameters are constant. We also assume that functions \(\alpha\) and \(A\) do not depend on \(\theta.\) This leads to the following simplified model,
\[
\frac{\partial \theta}{\partial t} = \alpha(x) \left( (1 - \theta) - (1 - \theta) u(x) \right) + \text{div} \left( A(x) \nabla \theta \right)
\]
\[\text{on } [0,T] \times \Omega,
\]
\[\langle A(x) \nabla \theta, n \rangle = 0 \quad \text{on } [0,T] \times \partial \Omega,
\]
\[\theta(0,x) = \theta_0(x) \geq 0 \quad x \in \Omega.
\]

In order to formalize the model, we define the Hilbert space
\[E = \{ \theta \in H^2(\Omega) ; \theta \text{ satisfies } \theta(0,x) = \theta_0(x) \geq 0 \quad x \in \Omega \}\]
provided with the inner product
\[\langle f, g \rangle_E = \int_U (fg + \langle \nabla f, \nabla g \rangle + \Delta f \cdot \Delta g) \, dx\]

Define also the linear unbounded operator \(L : D(L) = E \subset L^2(\Omega) \rightarrow L^2(\Omega)\) as
\[\langle L\theta, f \rangle = \alpha(x) \frac{\theta - u \cdot \theta}{1 - \theta} - \text{div} (A(x) \nabla \theta)
\]

Then equation (14) takes the following form
\[
\frac{\partial \theta}{\partial t} + L \theta = \alpha.
\]

We also introduce the following condition, which we will use to ensure that the system has realistic solutions:

\((H4)\) There exists a constant \(C > 0\) such that for almost every \(x \in \Omega, \) \(A(x)\) is symmetric, positive definite and
\[\langle v, A(x) v \rangle \geq C \langle v, v \rangle, \quad \forall v.
\]

B. Well-posedness of the diffusion model

We are now ready to prove that our model has been a mathematically and epidemiologically well-
posed. In other words, we show that exists a unique solution \(0 \leq \theta(t,x) \leq 1\) to the system (14) — (16).
This shall follow from the Hille-Yosida theorem but first we need the following proposition.

**Proposition 6.** If \( A(x) \) is a positive semidefinite bilinear form for almost every \( x \in \Omega \), then the linear operator \( L \) is monotone on \( E \). Moreover, if \( A(x) \) satisfies (H4), then \( L \) is maximal.

**Proof:**

(ii) To show \( L \) is monotone, we let \( \theta \in E \) and compute:

\[
\int_{\Omega} (\theta + L \theta) \times \varphi \, dx \\
= \int_{\Omega} \varphi \theta (1 + \alpha - \theta, u) / (1 - \theta, u) \, dx \\
- \int_{\Omega} \text{div} \left( A(x) \nabla \theta \right) \varphi \, dx \\
+ \int_{\Omega} \langle A(x) \nabla \theta, \nabla \varphi \rangle \, dx \\
- \int_{\partial \Omega} \langle A(x) \nabla \theta, n(x) \rangle \varphi \, ds \\
= \int_{\Omega} \varphi \theta (1 + \alpha - \theta, u) / (1 - \theta, u) \\
+ \int_{\Omega} \langle A(x) \nabla \theta, \nabla \varphi \rangle \, dx \\
\equiv p(\theta, \varphi),
\]

where \( p \) is a symmetric continuous and coercive bilinear form on \( H^1(\Omega) \). The Lax-Milgram theorem implies that there is a unique \( \theta \in H^1(\Omega) \) such that \( \theta + L \theta = f \).

Using regularization methods similar those used in Theorem 9.26 of [6], it follows that \( \theta \in E \).

Given that the linear operator \( L \) is maximal monotone and \( \theta_0 \) is in \( E \), then by the Hille-Yosida theorem there is a unique function \( \theta \in C^1([0, T] ; L^2(\Omega)) \cap C([0, T] ; E) \) which satisfies (14)–(16), and \( \forall (t, x) \in [0, T] \times \Omega \) we have

\[
\theta(t, x) = (S_L(t) \theta_0)(x) + \int_0^t (S_L(t - s) \alpha)(x) \, ds,
\]

where \( S_L(t) \) is the contraction semigroup generated by \( -L \).

Now that we have established existence and uniqueness of the solution \( \theta \), we now prove that \( 0 \leq \theta(t, x) \leq 1 \) for all \((t, x)\) in the domain. Assuming that \( A(x) \) satisfies the condition (H4). We define

\[
m \equiv \inf_{\partial \Omega} \theta_0, \quad M \equiv \max \left\{ \sup_{\partial \Omega} \theta_0, \, \sup_{\Omega} (1 - \theta, u) \right\}, \quad v \equiv 1 / (1 - \theta, u).
\]

Note that \( M \leq 1 \) as long as \( \theta_0 \leq 1 \) and \( 0 \leq \theta_1u \leq 1 \).

Let \( E_+ \) designate the set of elements in \( E \) which are nonnegative almost everywhere on \( \Omega \). The following theorem gives sufficient conditions under which the solution \( \theta \) of (14)–(16) is bounded by \( M \leq 1 \) and the positive cone \( E_+ \) of \( E \) is positively invariant.

**Theorem 7.** If \( A(x) \nabla \alpha = 0 \) for every \((t, x) \in [0, T] \times \Omega \), then for almost every \( x \) in \( \Omega \),

\[
m \leq e^{\alpha(t, x)} \theta(t, x), \quad t \in [0, T].
\]

Moreover if \( A(x) \nabla v = 0 \) for every \((t, x) \in [0, T] \times \Omega \), then

\[
\theta(t, x) \leq M
\]

In particular, (18) and (19) hold when \( \alpha \) and \( u \) do not depend on the space variable \( x \).

**Proof:** Let \( G \in C^1(\mathbb{R}) \) such that

\[
(i) \ G(s) = 0, \ \forall s \leq 0, \text{ and}
\]

\[
(ii) \ G(s) \geq 0, \ \forall s > 0.
\]
We observe that
\[ \phi \]
We may also compute
\[ H(s) \equiv \int_0^s G(\sigma) \, d\sigma, \forall s \in \mathbb{R}, \]
\[ \varphi_1(t) \equiv \int_\Omega H(m - e^{t \varphi}(t, x)) \, dx, \]
\[ \varphi_2(t) \equiv \int_\Omega H(e^{t \varphi}(\theta(t, x) - 1/v)) \, dx. \]

We observe that
\[ \varphi_1, \varphi_2 \in C([0, T]; \mathbb{R}) \cap C^1([0, T]; \mathbb{R}) , \]
\[ \varphi_1, \varphi_2 \geq 0 \text{ on } [0, T], \]
\[ \varphi_1(0) = \varphi_2(0) = 0. \]

We may also compute
\[ \varphi_1'(t) = -\int_\Omega e^{t \varphi}(m - e^{t \varphi}(\theta) - (v \alpha \theta + \partial \theta/\partial t) \, dx \]
\[ = -\int_\Omega e^{t \varphi}(m - e^{t \varphi}(\theta)) - (\alpha - \mathcal{L} \theta + v \alpha \theta) \, dx \]
\[ = -\int_\Omega \alpha \varphi(m - e^{t \varphi}(\alpha \theta)) \, dx \]
\[ + \int_\Omega \langle A(x) \nabla \theta, \nabla e^{t \varphi}(m - e^{t \varphi}(\theta)) \rangle \, dx \]
\[ = -\int_\Omega \alpha \varphi(m - e^{t \varphi}(\alpha \theta)) \, dx \]
\[ - \int_\Omega e^{2t \varphi} G'(m - e^{t \varphi}(\theta) \langle A(x) \nabla \theta, \nabla \theta \rangle \, dx \]
\[ + \int_\Omega (G(m - e^{t \varphi}(\alpha \theta)) - e^{t \varphi} G'(m - e^{t \varphi}(\theta))) \]
\[ \times \langle A(x) \nabla \theta, \nabla e^{t \varphi} \rangle \, dx \]
\[ \leq 0. \]

Since \( \varphi_1' \leq 0 \) on \( \mathbb{R}_+^* \), \( \varphi_1 \) is identically zero on \( \mathbb{R}_+ \) and therefore almost everywhere in \( \Omega \).

If \( A(x) \nabla v = 0 \) for every \((t, x) \in [0, T] \times \Omega \), then \( \varphi_1'(t) \)
\[ = \int_\Omega G(e^{t \varphi} (\theta - 1/v)) (\alpha - v \alpha \theta) \, dx \]
\[ = \int_\Omega e^{t \varphi} G(e^{t \varphi}(\theta - 1/v)) (\alpha - v \alpha \theta) \, dx \]
\[ = -\int_\Omega \langle A(x) \nabla \theta, \nabla e^{t \varphi}(\theta - 1/v) \rangle \, dx \]
\[ - \int \langle A(x) \nabla \theta, \nabla e^{t \varphi}(\theta - 1/v) \rangle \, dx \]
\[ = -\int_\Omega G(e^{t \varphi}(\theta - 1/v)) \langle A(x) \nabla \theta, \nabla \theta \rangle \, dx \]
\[ - \int_\Omega e^{t \varphi}(\theta - 1/v) G'(e^{t \varphi}(\theta - 1/v)) \, dx \]
\[ \times \langle A(x) \nabla \theta, \nabla e^{t \varphi} \rangle \, dx \]
\[ \leq 0. \]

Since \( \varphi_1' \leq 0 \) on \( \mathbb{R}_+^* \), \( \varphi_1 \) is identically zero on \( \mathbb{R}_+ \) and therefore almost everywhere in \( \Omega \).

The following theorem proves boundedness of \( \theta \) under more general conditions.

**Theorem 8.** Suppose that \( v \) and \( \alpha \theta \) are increasing functions \( h \) and \( g \) of the state \( \theta \), and there is a constant \( K > 0 \) such that
\[ a \theta' \leq K (1 + a \theta') \exp(a \theta), \forall a \geq 0. \]

Then for every time \( t \in [0, T] \) and almost every \( x \) in \( \Omega \),
\[ m \leq e^{t \varphi}(t, x) \]
and
\[ \theta(t, x) \leq M. \]

**Proof:** Let \( G \in C^1(\mathbb{R}) \) such that
(i) \( G(s) = 0, \forall s \leq 0, \) and
(ii) \( KG(s) \leq G'(s) \leq C, \forall s > 0. \)

Using [20] and the fact that operator \( A \) is monotone, we have
\[ \langle A(x) \nabla \theta, \nabla v \rangle = \langle A(x) \nabla \theta, \nabla h(\theta) \rangle \]
\[ = h'(\theta) \langle A(x) \nabla \theta, \nabla \theta \rangle \]
\[ \geq 0, \]
\[ \langle A(x) \nabla \theta, \nabla e^{t \nu} \rangle = \langle A(x) \nabla \theta, \nabla \exp(tg(\theta)) \rangle \\
= t g'(\theta) \exp(tg(\theta)) \langle A(x) \nabla \theta, \nabla \theta \rangle \geq 0, \]

and
\[ \langle A(x) \nabla \theta, \nabla e^{t \nu}G(m - e^{t \nu} \theta) \rangle \\
= \langle A(x) \nabla \theta, \nabla \exp(tg(\theta))G(m - \theta \exp(tg(\theta))) \rangle \\
= t g'(\theta) \exp(tg(\theta))G(m - \theta \exp(tg(\theta))) \langle A(x) \nabla \theta, \nabla \theta \rangle \\
- \exp(2tg(\theta))(1 + t g'(\theta))G'(m - \theta \exp(tg(\theta))) \langle A(x) \nabla \theta, \nabla \theta \rangle \leq (t g'(\theta) - K(1 + t g'(\theta)) \exp(tg(\theta))) \exp(tg(\theta)) \times G(m - \theta \exp(tg(\theta))) \langle A(x) \nabla \theta, \nabla \theta \rangle \leq 0. \]

Define
\[ H(s) \equiv \int_{0}^{s} G(\sigma) d\sigma, \forall s \in \mathbb{R}, \]
\[ \varphi_1(t) \equiv \int_{\Omega} H(m - e^{t \nu} \theta(t,x)) dx, \]
\[ \varphi_2(t) \equiv \int_{\Omega} H(e^{t \nu} \theta(t,x) - 1/w) dx. \]

Note that as in Theorem 7 we have
\[ \varphi_1, \varphi_2 \in C([0,T]; \mathbb{R}) \bigcap C^1([0,T]; \mathbb{R}), \]
\[ \varphi_1, \varphi_2 \geq 0 \text{ on } [0,T], \]
\[ \varphi_1(0) = \varphi_2(0) = 0. \]

As in Theorem 7 we may compute
\[ \varphi_1'(t) \]
\[ = - \int_{\Omega} e^{t \nu} G(m - e^{t \nu} \theta)(w \alpha \theta + \partial \theta/\partial t) dx \]
\[ = - \int_{\Omega} e^{t \nu} G(m - e^{t \nu} \theta)(\alpha - \mathcal{L} \theta + v \alpha \theta) dx \]
\[ = - \int_{\Omega} \alpha G(m - e^{t \nu} \theta) dx \]
\[ + \int_{\Omega} \langle A(x) \nabla \theta, \nabla e^{t \nu} G(m - e^{t \nu} \theta) \rangle dx \]
\[ \leq \int_{\Omega} \langle A(x) \nabla \theta, \nabla e^{t \nu} G(m - e^{t \nu} \theta) \rangle dx \leq 0. \]

Since \[ \varphi_1' \leq 0 \text{ on } \mathbb{R}_+^*, \] \varphi_1 is identically null on \[ [0,T] \] and therefore almost everywhere in \[ \Omega \]
\[ m \leq e^{t \nu} \theta(t,x). \]

We also have
\[ \varphi_2'(t) = \int_{\Omega} G(e^{t \nu} \theta - 1/w)(-\alpha + v \alpha \theta + \partial \theta/\partial t) dx \]
\[ = \int_{\Omega} e^{t \nu} G(e^{t \nu} \theta - 1/w)(-\mathcal{L} \theta + v \alpha \theta) dx \]
\[ = - \int_{\Omega} \langle A(x) \nabla \theta, \nabla e^{t \nu} G(e^{t \nu} \theta - 1/w) \rangle dx \]
\[ = - \int_{\Omega} e^{2t \nu} G'(e^{t \nu} \theta - 1/w) \langle A(x) \nabla \theta, \nabla \theta \rangle dx \]
\[ - \int_{\Omega} e^{2t \nu} G'(e^{t \nu} \theta - 1/w) \langle A(x) \nabla \theta, \nabla e^{t \nu} \theta \rangle dx \]
\[ - \int_{\Omega} e^{2t \nu} G'(e^{t \nu} \theta - 1/w) \langle A(x) \nabla \theta, e^{t \nu} \theta \rangle dx \]
\[ \leq 0. \]

Since \[ \varphi_2' \leq 0 \text{ on } [0,T] \setminus \{0\}, \] \varphi_2 is identically null on \[ [0,T] \] and therefore almost everywhere in \[ \Omega \]
\[ \theta(t,x) \leq M. \]

Condition 20 of Theorem 8 is satisfied in particular when \[ g \geq 0. \] Using the same arguments as in the proof of Proposition 6, there is a unique equilibrium \[ \theta^*, \] for the system (14) - (16). \[ \theta^* \] is asymptotically stable if and only if all the eigenvalues of the linear operator \[ \mathcal{L} \] have non-negative real parts. Stability of the equilibrium \[ \theta^* \] has the advantage that the disease inhibition is maintained in its neighborhood, which enables easier control strategies. In particular, the norm of \[ \theta^* \] is a decreasing function of the control \[ u. \]

**Proposition 9.** The real number \[ \lambda \] is not an eigenvalue of \[ \mathcal{L} \] if at least one of the following conditions is satisfied:

(i) \[ \alpha \geq \lambda(1 - \theta \cdot u) \text{ almost everywhere in } \Omega \]
and that inequality is strict on a nonnegligible subset of \[ \Omega. \]

(ii) There exists a real \[ k \geq 0 \text{ such that for every } \theta \in E \]
\[ \int_{\Omega} \langle A(x) \nabla \theta, \nabla \theta \rangle dx \geq k \| \theta \|_{H^2(\Omega)}, \]
and
\[ (\alpha - \lambda(1 - \theta \cdot u)) / (1 - \theta \cdot u) > -k \]
almost everywhere in \[ \Omega. \]
**Proof:** Let \( \theta, \varphi \in E \). Then we may compute
\[
\int_{\Omega} (\mathcal{L} \theta - \lambda \theta) \times \varphi \, dx = \int_{\Omega} \varphi \theta (\alpha - \lambda (1 - \theta, u)) / (1 - \theta, u) \, dx - \int_{\Omega} \operatorname{div} (A(x) \nabla \theta) \varphi \, dx = \int_{\Omega} \varphi \theta (\alpha - \lambda (1 - \theta, u)) / (1 - \theta, u) + \langle A(x) \nabla \theta, \nabla \varphi \rangle \, dx \equiv p_1 (\theta, \varphi).
\]

If either of the two conditions of the proposition is satisfied, we may use the Lax-Milgram theorem to obtain the desired result. \( \square \)

**Corollary 10.** The principal spectrum of \( -\mathcal{L} \) is contained in \( D_0^* \equiv \{ \lambda \in \mathbb{C}^* ; \operatorname{Re} (\lambda) \leq 0 \} \).

**Proof:** From assumption (H4), \( \mathcal{L} \) is maximal monotone and \( S_\mathcal{L} \) is a contraction semigroup. Since \( S_\mathcal{L} \) is a contraction semigroup, the resolvent set \( \rho (-\mathcal{L}) \) of \( -\mathcal{L} \) contains \( \mathbb{R}_+ \ni \{ 0 \} \) and \( \| S_\mathcal{L} (t) \| \leq 1, \forall t \in [0, T] \). Therefore, the spectral radius of \( S_\mathcal{L} (t) \) is less than one. On the other hand \( 0 \notin \exp (t \sigma_p (-\mathcal{L})) \subseteq \sigma_p (S_\mathcal{L} (t)) \subseteq \{ 0 \} \cup \bigcup \exp (t \sigma_p (-\mathcal{L})) , \forall t \in [0, T] \). Clearly, if \( \lambda = \operatorname{Re} (\lambda) + i \operatorname{Im} (\lambda) \) is an element of the principal spectrum of \( -\mathcal{L} \) then \( \exp (\lambda t) \) is an element of the principal spectrum of \( S_\mathcal{L} (t) \) and \( | \exp (\lambda t) | = \exp (\operatorname{Re} (\lambda) t) \| S_\mathcal{L} (t) \| \leq 1 \). It follows that \( \operatorname{Re} (\lambda) \leq 0 \). \( \square \)

**Corollary 11.** The equilibrium \( \theta^* \) is stable. Moreover if all the complex eigenvalues \( \lambda \) of the operator \( \theta \mapsto \operatorname{div} (A(x) \nabla \theta) \) satisfy \( \alpha \geq (1 - \theta, u) \operatorname{Re} (\lambda) \) almost everywhere in \( \Omega \), then \( \theta^* \) asymptotically stable.

**C. Optimal control of the diffusion model**

In the previous section we have seen that the equilibrium of system (14) - (16) was conditionally asymptotically stable. Whether or not the equilibrium \( \theta^* \) is asymptotically stable, the disease progression shall be contained with respect to some criteria given in terms of costs. The aim of this section is to control the system such that the following cost functional is minimized:
\[
J_1^3 (u) = \int_0^T \int_\Omega (\theta^2 + k_1 (x) u^2) \, dx dt + \int_\Omega k_2 (x) \theta^2 (T, x) \, dx,
\]
where \( k_1 > 0, k_2 \geq 0 \) are bounded penalization terms. The function \( k_1 (x) \) can be interpreted as the cost ratio related to the use of control effort \( u \); while \( k_2 \) is the cost ratio related to the magnitude of the final inhibition rate \( \theta (T, \cdot) \). In practice, \( k_1 \) reflects the spatial dependence of environmental sensitivity to control means, while \( k_2 \) reflects geographical variations in the cost of the inhibition rate of *Colletotrichum* at the end of the control period.

In order to establish the optimal control, we will first need to define \( U^{K,C} \) as the set of controls \( u \in C ([0, T] ; H^1 (\Omega) ; [0, 1]) \) such that for every \( t, s \in [0, T] \), \( \| u (t, \cdot) - u (s, \cdot) \|_{H^1 (\Omega)} \leq K | t - s | \) and \( \| \nabla u (t, \cdot) \|^2 \leq C \). For every \( K, C \geq 0 \), \( U^{K,C} \) is nonempty.

**Theorem 12.** Let \( K, C \geq 0 \). Then there is a control \( v \in U^{K,C} \) which minimizes the cost \( J_1^3 \).

**Proof:** Since \( J_1^3 \) is greater than zero it is bounded below. Let that infimum be \( J^* \). There is a sequence \( (u_n)_{n \in \mathbb{N}} \) such that the sequence \( (J_1^3 (u_n))_{n \in \mathbb{N}} \) converges to \( J^* \). Using definition of \( U^{K,C} \) the \( (u_n)_{n \in \mathbb{N}} \) is bounded and uniformly equicontinuous on \( [0, T] \). By the Ascoli theorem, there is a subsequence \( (u_{n_k}) \) which converges to a control \( v \). Since the cost function is continuous with respect to \( u \) it follows that \( J_1^3 (v) = J^* \).

We first look the linearized system in the neighborhood of \( (\theta, u) = (\varepsilon, 0) \), where \( \varepsilon \) depends on \( x \). Indeed, this case is of practical significance since the monitoring is assumed to be continuous year-round, and the endemic period corresponds to particular conditions. Thus the outbreak of the disease is "observable" at the moment of onset. The linearized version of (14) is
\[
\partial \theta / \partial t = \alpha - \alpha \theta - \alpha \varepsilon u \theta + \operatorname{div} (A(x) \nabla \theta), \text{ on } [0, T] \times \Omega
\]
(23)

Note that if \( \varepsilon = 0 \) the linearized system is not controllable.
Let $L_1 \theta = -\alpha \theta + \text{div} (A(x) \nabla \theta)$. Equation (14) becomes
\[ \frac{\partial \theta}{\partial t} = L_1 \theta - \alpha \varepsilon \theta + \alpha, \quad \text{on } ]0, T[ \times \Omega. \]

**Theorem 13.** The linearized version of (14) -- (16) has an optimal control in $C([0, T]; L^2(\Omega))$ given by
\[ u(t, \cdot) = (1/k_1) BP (T - t, \cdot) \theta(t, \cdot) + 1/\varepsilon \theta(t, \cdot), \quad t \in [0, T] \]
where the linear operator $P$ is solution to the following Riccati equation:
\[ P = L_1 P + P L_1 - (1/k_1) PB^2P + I, \quad P(0) = k_2 I. \]

In that equation $I$ is the identity linear operator and $B$ is the linear operator $\alpha \varepsilon \theta I$.

**Proof:** (Sketch)
If we set $v = u - 1/\varepsilon \theta_1$ then equation (23) becomes
\[ \frac{\partial \theta}{\partial t} = L_1 \theta - \alpha \varepsilon v \theta_1, \quad \text{on } ]0, T[ \times \Omega. \]
The rest of the proof is similar to the proof in [29] concerning linear regulators.

If $S_{L_1}$ is the contraction semigroup generated by $L_1$, then we have $\forall t \in [0, T]
\[ P(t) = S_{L_1}(t) P(0) S_{L_1} f + \int_0^t S_{L_1}(t-s) (I - (1/k_1) PB^2 P ) S_{L_1}(t-s)f ds \]
Let now consider the nonlinear equation (14). Let $L_u$ be the operator $L$ corresponding to control strategy $u$ and let $S_{L_u}$ be the contraction semigroup generated by $-L_u$.

Let $\Omega = \{ u \in C([0, T]; H^1(\Omega; [0, 1])); \forall t \in [0, T], \ S_{L_u}(t) \text{ is invertible} \}.

Some necessary and sufficient conditions for a semigroup of operators to be embedded in a group of operators are given in [25].

**Theorem 14.** Assume that there is a bounded admissible control $u^* \in U$ which minimizes the cost function $J^3_T$. Let $\tilde{\theta}$ be the absolutely continuous solution of (14) -- (16) associated with $u^*$. Then
\[ \int_\Omega \left( \frac{\partial \tilde{\theta}(t_0, x)}{\partial t}^2 + k_1(x) (u^*(t_0, x))^2 - p(t) L_{u^*} \tilde{\theta}(t, x) \right) dx, \]
where $p$ is the absolutely continuous solution on $[0, T]$ of the adjoint state problem
\[ \left\{ \begin{array}{l}
\frac{\partial p}{\partial t} = L_u p - 2 \tilde{\theta}, \quad (t, x) \in R^+_T \times \Omega \\
A(x) \nabla p, n = 0, \quad \text{on } \partial \Omega \\
p(T) = 2k_2 \tilde{\theta}(T, \cdot) \end{array} \right. \quad (24) \]

**Proof:** We give a proof following the maximum principle proof in [29].
For an arbitrary control $w$ and sufficiently small $h \geq 0$, define the needle variation of $u^*$ as
\[ u^h(t) = \begin{cases} u^*(t), & t \in [0, T] \\
u^w(t), & t \in [0, T] \end{cases} \]
Let $\theta^h$ be the output corresponding to $u^h$. Since $u^*$ minimizes $J^3_T$; $J^3_T(\theta^h) > J^3_T(\theta^0)$ and $\partial^+ J^3_T(\theta^0)/\partial h > 0$. Then
\[ \partial^+ \theta^0(t_0, \cdot)/\partial h \]
\[ \lim_{h \to 0} \frac{1}{h} \left[ \theta^h(t_0, \cdot) - \theta^0(t_0, \cdot) \right] \]
\[ \lim_{h \to 0} \frac{1}{h} \int_{t_0-h}^{t_0} \left( L_{u^0} \tilde{\theta}(s, \cdot) - L_{u^h} \theta^h(s, \cdot) \right) ds \]
\[ (L_{u^0} - L_w) \theta^0(t_0, \cdot) \]
Since $v^h(t) = u^*(t)$ on $[t_0, T]$, for almost $t$ in $[t_0, T]$, $\partial \theta^h/\partial t = \alpha - L_{u^0} \theta^h$. Then
\[ \partial \left( \frac{\partial^+ \theta^0(t, \cdot)}{\partial h} \right)/\partial t \]
\[ \partial^+ \theta^0(t, \cdot)/\partial h \]
\[ \partial^+ \theta^0(t, \cdot)/\partial h \]
\[ \partial^+ \theta^0(t, \cdot)/\partial h \]
\[ \partial^+ \theta^0(t, \cdot)/\partial h \]
Therefore,
\[ \partial^+ \theta^0(t, \cdot)/\partial h = S_{L_{u^0}} (t) (S_{L_{u^0}} (t_0))^{-1} (L_{u^0} - L_w) \theta^0(t_0, \cdot) \]
Consequently,
\[
\partial^+ \left( \int_{\Omega} k_2(x) \left( \theta^0(T, x) \right)^2 \, dx \right) / \partial h \\
= 2 \int_{\Omega} k_2(x) \theta^0(T, x) \partial^+ \theta^0(T, x) / \partial h \, dx \\
= 2 \int_{\Omega} k_2(x) \theta^0(T, x) S_{L_{\theta}}(T) (S_{L_{\theta}}(t_0))^{-1} \\
\times (L_{w^0} - L_w) \theta^0(t_0, x) \, dx \\
= 2 \int_{\Omega} (S_{L_{\theta}}(t_0))^{-1} S_{L_{\theta}}(t) k_2(x) \theta^0(T, x) \\
\times (L_{w^0} - L_w) \theta^0(t_0, x) \, dx,
\]
and in the same manner
\[
\partial^+ J^2_T (\theta^0) / \partial h \geq 0,
\]
we have
\[
2 \int_0^T \int_{\Omega} \left( \partial^+(\theta^0(t, x))^2 \right) \, dx \, dt \\
+ 2 \int_0^T \int_{\Omega} \left( \partial^+(\theta^0(t, x))^2 \right) / \partial h \, dx \, dt \\
= 2 \int_0^T \int_{\Omega} (S_{L_{\theta}}(t_0))^{-1} S_{L_{\theta}}(t) \theta^0(t, x) \\
\times (L_{w^0} - L_w) \theta^0(t_0, x) \, dx \, dt.
\]
Since \( \partial^+ J^2_T (\theta^0) / \partial h \geq 0 \), we have
\[
2 \int_0^T \int_{\Omega} \left( \partial^+(\theta^0(t, x))^2 \right) \, dx \, dt \\
+ 2 \int_0^T \int_{\Omega} \left( \partial^+(\theta^0(t, x))^2 \right) / \partial h \, dx \, dt \\
\geq 2 \int_0^T \int_{\Omega} (S_{L_{\theta}}(t_0))^{-1} S_{L_{\theta}}(t) \theta^0(t, x) \\
\times (L_{w^0} - L_w) \theta^0(t_0, x) \, dx \, dt
\]
which is analogous to (10) for the within-host model. Then we can adopt the following corresponding strategy
\[
u^*(t) = \begin{cases} 
1 & \text{when } 27\alpha\theta_2\theta_3 > 8k_1, \\
w_3(t) & \text{when } 27\alpha\theta_2\theta_3 < 8k_1,
\end{cases}
\]
where \( w_3(t) \) is the element of \( \left[0, \min \left\{ \frac{1}{8k_1}, 1 \right\} \right] \) which is the nearest to the smallest nonnegative solution of the equation (26).

**IV. CONCLUSION**

In this paper two models of anthracnose control have been surveyed. These models both have the general form
\[
\partial \theta / \partial t = f(t, \theta, u) + g(t),
\]
where \( f \) is linear in the state \( \theta \) but not necessarily in the control \( u \). As far as the authors know, this type of control system has not been extensively studied. This may be due to the fact that physical control problems usually do not take this form. The majority of such problems tend to use "additive" controls (see [8], [16] for literature on models). But in models of population dynamics, "multiplicative" control are often more realistic.

Our first model characterizes the within-host behaviour of the disease. We were able to explicitly calculate an optimal control strategy that effectively reduces the inhibition rate compared to the case where no control is used. In our second model we take into account the spatial spread of the disease by adding a diffusion term. That makes the model more interesting but considerably more difficult to analyze. Moreover, visual evaluation appears more difficult because in this case the state of the system is a function of three spatial variables.
plus time. Although we have provided equations satisfied by the optimal control (for the linearized system), in this paper we do not give a practical method for computing the optimal control. It is possible that adapted gradient methods may be used [2]: this is a subject of ongoing research.

Our models seems quite theoretical, but could be used for practical applications if the needed parameters were provided. Indeed, in the literature [9], [10], [12], [17] there are several attempts to estimate these parameters. The principal advantage of our abstract approach is that it can be used to set automatic means to control the disease which are able to adapt themselves with respect to the host plant and to the parameters values.

Obviously our models can be improved. In particular, several results are based on some conditions of smoothness of parameters, and the control strategy is also very regular. In practice parameters are at most piecewise continuous, and some control strategies are discontinuous. For instance, cultural interventions in the farm are like pulses with respect to a certain calendar. The application of antifungal chemical treatments are also pulses, and the effects of these treatments though continuous are of limited duration. We are currently investigating a more general model that takes into account those irregularities.

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