Comparison of mathematical tumor growth models

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Abstract—Tumor cells need a set of functional capabilities for development and growth. One of these capabilities is angiogenesis; tumor cells have to become able to force new vessel formation to ensure oxygen and nutrient supply. In the literature, there are several different growth models which aim to describe the growth characteristics. The model creation is a trade-off problem. On the one hand, the model has to be sufficiently complex to describe all the important processes which take part in the system; however, on the other hand, the model has to be sufficiently simple and manageable for real-life usability and – taking into account the applicability of biomedical control theory – for controller design. We discuss some relevant tumor growth models used in the literature showing their complexity and their modeling power.

I. INTRODUCTION

Tumor growth is a very complex process and requires a complex system which contains several key points. Hanahan and Weinberg [1] have collected the set of functional capabilities which are necessary for tumors during their development; they labeled these capabilities as the “hallmarks of cancer” (Fig. 1):

- self-sufficiency in growth signals
- insensitivity to anti-growth signals
- evading apoptosis
- limitless replicative potential
- sustained angiogenesis
- tissue invasion and metastasis

One of these key capabilities is angiogenesis, the process of new vessel formation. Three main classes of mathematical models have been created in the field of antiangiogenic therapy [2, 3]:

- temporal models [4, 5];
- spatiotemporal models [6, 7];
- multiscale models [8, 9].

The main disadvantage of these models is that they are mechanistic or semi-mechanistic [10] models built up from physical equations, and they have not been validated with in vivo data in most of the cases. Exceptions (validated models) exist, however they have other problem. The Hahnfeldt model [4] is not valid any more in the light of new medical results. A newly created and validated model posed by Gevertz [8] takes into account numerous effects and as a result it is overly difficult (it has 13 variables and 21 parameters).

The paper gives a systematic overview of the different effects taken into account in the literature and is organized as follows.

In Section II, we review the linear-exponential growth models, from the simplest exponential model to a complex one which contains exponential and linear phases as well by taking into account the phenomenon of angiogenesis.

In Section III, we discuss the sigmoid shaped models like logistic and Gompertz models, together with a model, which assumes dynamic (time-dependent) carrying capacity.

In Section IV, a tumor growth model is presented in detail, in which the apoptosis is taken into account as a volume loss mechanism of the solid tumor.

In Section V summarizes the tumor growth model inducing angiogenesis. Finally, the paper ends with the conclusions.
II. LINEAR-EXPONENTIAL MODELS

A. Exponential growth model

The simplest tumor growth model is an exponential one:

\[ N(t) = N(0) \cdot e^{\lambda t} \]

where \( N(t) \) is the current tumor size, \( N(0) \) is the initial tumor size, and \( \lambda \) is the growth parameter, which describes the speed of the growth.

B. Linear-expontential growth models

Several research articles \([11, 12, 13]\) describe the phenomenon that tumor growth dynamics contains different phases, which can be described in a linear and exponential way.

Simeoni et al. \([11]\) present a minimal pharmacokinetic-pharmacodynamic model, which describes the tumor growth in nontreated animals. They have found that the initial exponential phase is followed by a linear phase. In the study, a transit compartmental system was used to model the process of cell death at the later phase. The growth rate, which describes the above mentioned characteristics, can be formulated as a Cauchy problem \([12]\):

\[
\begin{align*}
\frac{dV}{dt} &= a_0 V, \quad t \leq \tau \\
\frac{dV}{dt} &= a_1, \quad t > \tau, \\
V(t = 0) &= V_0
\end{align*}
\]

where \( a_0 \) belongs to the exponential phase, it is the coefficient of proliferative cells times \( ln \frac{2}{T_c} \) (\( T_c \) is cell cycle length); while \( a_1 \) is the coefficient of linear phase. From Equation (2), one can determine the value of \( \tau \):

\[
\tau = \frac{1}{a_0} \log \left( \frac{a_1}{a_0 V_0} \right)
\]

Tang et al. \([13]\) modeled the three-dimensional tumor growth and angiogenesis to simulate tumor progression for chemotherapy evaluation. The whole complex model consists of three different levels: cellular, intratumoral, and tissue level. The model defines the avascular tumor pressure, and the evolution of oxygen and carbon dioxide concentration for the solid tumor growth model. The solid tumor angiogenesis model takes into account the process of hypoxia, which is a frequent phenomenon in tumor cells in the lack of oxygen. The response of tumor cells to hypoxia is the secretion of tumor angiogenesis factors (e.g. vascular endothelial growth factor, VEGF). The process when the tumor becomes able to produce angiogenesis factors is called as angiogenic switch, since thereby the tumor will possess own blood vessels. In the model, tumor vessel growth is determined by the density of the angiogenesis factors and tumor interstitial pressure. The model secludes four different phases of the growth process (Fig. 2):

- exponential growth (\( T_I \)),
- linear expansion (\( T_2 \)),
- stasis (\( T_3 \)),
- secondary growth (\( T_4 \)).

III. Sigmoid shaped models

(BASED ON THE CARRYING CAPACITY)

A. Logistic and Gompertz models

Several studies have shown that tumor growth has a sigmoid shape \([12, 14, 15]\), viz. the growth function has an inflection point. After this point, the growth slows down in time and converges to a plateau. These types of models are based on the observation that the carrying capacity of the vessels has a maximal value.

The logistic model is defined by the following equations \([12, 14]\)

\[
\begin{align*}
\frac{dV}{dt} &= aV \left( 1 - \frac{V}{K} \right) \\
V(t = 0) &= 1 \text{mm}^3
\end{align*}
\]

where \( a \) is the coefficient of cell proliferation, and \( K \) is the carrying capacity.

The Gompertz model \([12, 15]\) is described as

\[
\begin{align*}
\frac{dV}{dt} &= ae^{-\beta V} \\
V(t = 0) &= 1 \text{mm}^3
\end{align*}
\]

where \( a \) is the coefficient of the initial proliferation rate, and \( \beta \) is the coefficient of growth deceleration. The solution of the Gompertz model is

\[
V(t) = V_0 e^{\frac{a}{\beta} \left( 1 - e^{-\beta t} \right)}
\]

where \( K = V_0 e^{\frac{a}{\beta}} \) is the carrying capacity.
B. Dynamic carrying capacity based models

Assuming a dynamic (time-dependent) carrying capacity, the following equations can be written [12]

\[
\begin{align*}
\frac{dV}{dt} &= aV \log \left( \frac{K}{V} \right) \\
\frac{dK}{dt} &= bV^2
\end{align*}
\]

(7)

where \(a\) and \(b\) are coefficients, and \(K\) is the carrying capacity.

The advantage of this type of model is that it can represent the vasculature as a limit of tumor growth [4].

IV. APOPTOSIS BASED TUMOR GROWTH MODEL (VOLUME LOSS OF THE SOLID TUMOR)

A. The tumor growth model

The tumor growth model, in which the apoptosis is taken into account as a volume loss mechanism of the solid tumor was created by McElwain and Morris [16]. The model assumes that the tumor is spherical.

In the model, the diffusion of nutrient is given by

\[
\frac{k}{r^2} \frac{d}{dr} \left( r^2 \frac{d\sigma}{dr} \right)
\]

(8)

where \(k\) is the coefficient of nutrient diffusion, \(r\) is the current radius of the nodule, and \(\sigma\) is the local nutrient concentration.

The nutrient consumption \((Af(\sigma))\) and the proliferation \((sf(\sigma))\) are described as a function of the local nutrient concentration. Three different cases can be separated based on the current value of \(f(\sigma)\)

\[
f(\sigma) = \begin{cases} 
1 & \sigma^* \leq \sigma \\
\frac{\sigma}{\sigma^*} & \sigma^* < \sigma \leq \sigma^* \\
0 & \sigma \leq \sigma^*_c \end{cases}
\]

(9)

where \(\sigma^*\) is the critical level: cancer cells die when the nutrient concentration falls below this level.

In Case1, the proliferation and nutrient consumption are “normal” (the proliferation rate and the nutrient consumption rate are not weighted).

In Case2, “nutrient deficiency phenomena” occurs (the proliferation rate and the nutrient consumption rate decreases as linear functions of the current concentration level of nutrient).

In Case3, the tumor cells die and form a “coagulative necrotic core” (the proliferation rate and the nutrient consumption rate are zero).

B. Phase I

In Phase I, there are sufficient oxygen and nutrients for the tumor cells; hence the tumor growth is not limited, it shows an exponential growth characteristics. Cell proliferation occurs with normal growth rate (Case1), and apoptosis is active as well (Fig. 3).

\[
\sigma = f(\sigma) = \begin{cases} 
\sigma(\sigma) & \rho = 0 \\
1 & \rho > 0
\end{cases}
\]

(normal growth rate)

\[
\sigma = \sigma(\sigma) = \left( \frac{\sigma_0}{\sigma^*} \right)^{3 \tau}
\]

(apoptosis)

\[
f(\sigma) = \begin{cases} 
1 & \sigma^* \leq \sigma \\
\frac{\sigma}{\sigma^*} & \sigma^* < \sigma \leq \sigma^* \\
0 & \sigma \leq \sigma^*_c
\end{cases}
\]

\[
\rho = \begin{cases} 
1 & \rho = 0 \\
\frac{A}{k\sigma^*} & \rho > 0
\end{cases}
\]

(Fig. 3. Phase I in the apoptosis based tumor growth model.)

The diffusion of nutrient in this phase can be described as

\[
\frac{k}{r^2} \frac{d}{dr} \left( r^2 \frac{d\sigma}{dr} \right) = A, \quad 0 \leq r \leq R_0,
\]

(10)

where \(A\) is the normal nutrient consumption rate, and \(R_0\) is the outer radius of the nodule. Hence, the nutrient concentration as a function of the radius can be expressed as

\[
\sigma(r) = \sigma(\sigma_0) - \frac{A}{6k} \left( r^2 - r_0^2 \right)
\]

(11)

Taking into account that the change of the tumor volume depends on the difference of the growth and death rate, one can describe that

\[
\frac{dV}{dt} = (s - \lambda)\nu,
\]

(12)

where \(s\) is the normal local proliferation rate (growth rate), and \(\lambda\) is the local rate of volume loss due to apoptosis per unit volume per unit time (death rate).

In this light, the tumor growth equation which describes the evolution of the outer radius of the nodule in time is

\[
\frac{4\pi}{3} \frac{dR_0^3}{dt} = s \frac{4\pi}{3} R_0^3 - \lambda \frac{4\pi}{3} R_0^3
\]

(13)

Considering that Equation (12) has the solution in the following form

\[
V = ce^{s - \lambda t},
\]

(14)

the solution of the tumor growth Equation (13) can be written as

\[
R_0(\tau) = \sqrt[3]{\frac{A}{k\sigma^*} R_0(0) e^{s - \lambda \tau}}
\]

(15)

where \(\tau = st\) is the scaled time, and \(\gamma = \lambda/s\) (which has to be less than unity for the growth of the nodule).

This phase stops when the nutrient concentration at the center of the nodule becomes equal to the critical concentration \((\sigma(r = 0) = \sigma^*_c)\). The value of the radius at which the tumor switches to Phase II becomes

\[
R_0 |_{\rho>0} = \sqrt[3]{\frac{A}{k\sigma^*} (\frac{\sigma(\sigma_0)}{\sigma^*})^{-1} - 1}
\]

(16)

C. Phase II

In Phase II, a new layer appears in the tumor (Fig. 4). Beside the external region which has a normal growth rate (Case1), an inner nodule is developed which slows the growth rate (Case2) due to the nutrient and oxygen deficiencies.
In this phase, the diffusion of the nutrient can be described as

$$k \frac{d}{dr} \left( r^2 \frac{d\sigma}{dr} \right) = \begin{cases} \frac{A}{\sigma^*} \sigma & 0 \leq r \leq R^* \\ \frac{A}{\sigma^*} \sigma \leq r \leq R_0 \end{cases} \quad (17)$$

where $R^* = R(\sigma = \sigma^*)$.

The nutrient concentration when $0 \leq r \leq R^*$ is

$$\sigma(r) = \sigma^* \frac{R^*}{r} \sinh \left( \frac{A}{k\sigma^*} r \right) \quad (18)$$

while the nutrient concentration when $R^* \leq r \leq R_0$ can be expressed as

$$\sigma(r) = \sigma(R_0) + \frac{A}{6k} \left( r^2 - R_0^2 \right) + \left( \sigma^* - \sigma(R_0) \right) \frac{A}{6k} \left( R^* \frac{r - R_0}{R^* - R_0} \right) \quad (19)$$

The tumor growth equation can be described by

$$\frac{4\pi}{3} \frac{dR_0}{dt} = s \frac{4\pi}{3} \left( R_0^3 - R_{**}^3 \right) - \lambda \frac{4\pi}{3} R_0^3 + \frac{4\pi}{3} \int_0^r \frac{\sigma(r)}{\sigma^*} r^2 dr \quad (20)$$

and the solution of this growth function is

$$\frac{A}{k\sigma^*} R_0^2 \frac{d}{dt} \left( \frac{A}{k\sigma^*} R_0 \right) = \frac{1}{3} \left( 1 - \gamma \right) \left( \frac{A}{k\sigma^*} R_0 \right)^3 + \left( \frac{\sigma(R_0)}{\sigma^*} - 1 \right) \frac{A}{6k} \left( \frac{A}{k\sigma^*} R_0^3 - \frac{A}{k\sigma^*} R_{**}^3 \right) \frac{A}{k\sigma^*} \left( R_0 - R_{**} \right) \quad (21)$$

The process of slowed growth can result in two different outcomes (Fig. 5). If the volume growth due to cell division is balanced by the volume loss due to apoptosis, a steady state (dormant state) can be attained. Otherwise, if at the center of the nodule the nutrient concentration becomes equal to $\sigma_i$, viz. $\sigma(r = 0) = \sigma_i$, a necrotic core formation will start and the tumor switches to Phase III.

D. Phase III

In Phase III, the tumor has a three layer structure (Fig. 6). In the outermost shell, there is a normal growth with exponential growth characteristics (Case1). In the middle shell, the growth is slowed due to nutrient deficiencies (Case2), while in the inner region tumor cells die and form a coagulative necrotic core (Case3).
The diffusion of the nutrient in the three different regions is

\[
k \frac{d}{dr} \left[ r^2 \frac{d\sigma}{dr} \right] = \frac{A}{\sigma} \frac{\sigma}{\sigma^*} R_i \leq r \leq R^*, \quad (22)
\]

where \( R_i \) is the radius of the inner necrotic region.

The solution of the nutrient concentration when \( 0 \leq r \leq R_i \) is

\[
\sigma(r) = \frac{\sigma_i}{\sigma}, \quad (23)
\]

when \( R_i \leq r \leq R^* \), the nutrient concentration becomes

\[
\sigma(r) = \frac{\sigma_i}{\sqrt{A/k\sigma^*}} \sin \left( \frac{A}{k\sigma^*} (r - R_i) \right) + \frac{A}{k\sigma^*} R_i \cos \left( \frac{A}{k\sigma^*} (r - R_i) \right), \quad (24)
\]

and finally, the nutrient concentration when \( R^* \leq r \leq R_0 \) results as

\[
\sigma(r) = \sigma(R_0) + \frac{A}{6k} \left( r^2 - R_0^2 \right) + \frac{A}{k\sigma^*} \left( 1 - \frac{1}{R_0} \right), \quad (25)
\]

where \( F \) causes \( \sigma \) to satisfy \( \sigma(R^*) = \sigma^* \).

The tumor growth equation in Phase III is

\[
\frac{4\pi dR_0}{3} = \frac{4\pi}{3} \left( R_0^3 - R^* \right) - \frac{4\pi}{3} \left( R_i^3 - R_i^* \right) + 4\pi \int \frac{\sigma(r)}{k} \sigma^* r^2 dr, \quad (26)
\]

and the solution of this growth function becomes

\[
\left( \frac{A}{k\sigma^*} R_0 \right)^3 \frac{d}{d\tau} \left( \frac{A}{k\sigma^*} R_0 \right) = \frac{1}{3} \left( 1 - \gamma \right) \left( \frac{A}{k\sigma^*} R_0 \right)^3 + \frac{1}{3} \int \frac{\sigma(r)}{k} \sigma^* r^2 dr + \frac{A}{k\sigma^*} \left( 1 - \frac{A}{k\sigma^*} R_0^3 - A R^* \right) \left( \frac{A}{k\sigma^*} R_0 R^* \right), \quad (27)
\]

V. ANGIogenesIs BASEd Tumor GROWTh MODEL

Yang [17] performed a mathematical model which describes the tumor growth inducing angiogenesis. The modeling concept was to describe an organ without tumor first, when normal cells and blood vessels are in a steady state; followed by the modeling of the organ with cancer cells.

The model discusses a pre-angiogenesis stage, where the cells are formed by the “mass action law” [18]. This stage is continued by the VEGF expression and finally the endothelial sprouting. The model assumes logistic growth for normal, cancer and epithelial cells as well.

The tumor growth model which takes into account angiogenesis, is described by the following system of equations

\[
\frac{dC}{dt} = \alpha C \left[ 1 - \frac{C}{k_1(E)} \right] - \mu C - C \delta_0(t) \quad (28)
\]

\[
\frac{dE}{dt} = \alpha E \left[ 1 - \frac{E}{k_1(C)} \right] - \eta E - \mu E \quad (29)
\]

\[
\frac{dT}{dt} = C \delta_0(t) + \alpha A T \left[ 1 - \frac{T}{k_3} \right] + \alpha_3 T \left[ 1 - \frac{T}{k_4} \right] - \mu T \quad (30)
\]

\[
\frac{dP}{dt} = \gamma E - \partial P - \mu P \quad (31)
\]

\[
\frac{dA}{dt} = \partial A + \eta A \left( 1 - \frac{A}{k_5} \right) - \mu A \quad (32)
\]

The equations contain the following parameters.

- \( C, E, T, P \) and \( A \) describe the concentration of normal, epithelial, cancer, pre-angiogenesis and angiogenesis cells, respectively.
- \( \alpha, \alpha_2, \alpha_3, \alpha_4 \) mean the intrinsic growth rate of normal, epithelial, cancer and angiogenesis cells, respectively.
- \( \mu, \mu_2, \mu_3, \mu_4 \) and \( \mu_5 \) denote the mortality rate of normal, epithelial, cancer, pre-angiogenesis and angiogenesis cells, respectively.
- \( k_1, k_2, k_3 \) and \( k_4 \) are the carrying capacity of normal, epithelial, cancer and angiogenesis cells, respectively.
- \( \delta \) is the transfer rate from pre-angiogenesis to angiogenesis cells, and \( \gamma \) is the epithelial sprouting rate.
- \( \beta_1 \) describes the rate of inhibition of normal cells by cancer cells, and \( \beta_2 \) describes the rate of inhibition of cancer cells by normal cells.

VI. CONCLUSION

There is a strong need to create a mathematical model which describes the tumor growth dynamics under angiogenic inhibition. This model has to take into account the previously mentioned models and their results, but it has to be sufficiently simple to be manageable for both real-life applicability and controller design. Based on the supporting data of our research results we believe that such a model can be created [19-22].
REFERENCES


