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A multilevel approach to cancer growth modeling

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Abstract

Cancer growth models may be divided into macroscopic models, which describe the tumor as a single entity, and microscopic ones, which consider the tumor as a complex system whose behaviour emerges from the local dynamics of its basic components, the neoplastic cells. Mesoscopic models (e.g. as based on the Local Interaction Simulation Approach (Delsanto et al., 1998)), which explicitly consider the behavior of cell clusters and their interactions, may be used instead of the microscopic ones, in order to study the properties of cancer biology that strongly depend on the interactions of small groups of cells at intermediate spatial and temporal scales. All these approaches have been developed independently, which

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limits their usefulness, since they all include relevant features and information that should be cross-correlated for a deeper understanding of the mechanisms involved.

In this contribution we consider multicellular tumor spheroids as biological reference systems and propose an intermediate model to bridge the gap between a macroscopic formulation of tumor growth and a mesoscopic one. Thus we are able to establish, as an important result of our formalism, a direct correspondence between parameters characterizing processes occurring at different scales. In particular, we analyze their dependence on an important limiting factor to tumor growth, i.e. the extra-cellular matrix pressure. Since the macro and meso-models stem from totally different roots (energy conservation and clinical observations vs cell groups dynamics), their consistency may be used to validate both approaches. It may also be interesting to note that the proposed formalism fits well into a recently proposed conjecture of growth laws universality.

Key words: Cancer growth, Multicellular Tumor Spheroids, Modeling, Scaling, Multilevel Approach, Cellular Dynamics, Universality, Energy Balance, Complexity

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1 1 INTRODUCTION

Crucial to our understanding of the development of complexity is our ability 2 to relate phenomena occurring at different scales. This is necessary not only to 3 predict the emergence of macroscopic phenomena from microscopic processes. 4 but also to relate microscopic variables to the more accessible parameters of 5 macroscopic phenomenology. In fact, the use of more realistic values for the pa-6 rameters of the microscopic (and mesoscopic) simulations may greatly enhance 7 their predictive potentiality and, therefore, their applicability for biomedical 8 clinical purposes. As a rule, numerical simulations are necessary to impleç ment microscopic or mesoscopic models, while analytical (or semi-analytical) 10 solutions are usually possible for macroscopic models. 11

Tumors are remarkable examples of complex self-organizing systems. Due 12 to their inherent complexity, it is necessary to analyze their growth at dif-13 ferent scales. In a macroscopic approach, we consider them as single entities, 14 whose behavior can be predicted in terms of their global interaction with the 15 environment and a few internal parameters. This approach has led to a host 16 of useful models of cellular population dynamics in different biological sys-17 tems (for example cell cultures (Murray, 2004), the immune system (Adam 18 and Bellomo, 1996; Perelson and Weisbuch, 1997), neoplastic masses (Adam 19 and Bellomo, 1996; Preziosi, 2003)). 20

Interest in this approach has been further rekindled by the conjecture of Guiot et al. (Guiot et al., 2003) that the ontogenetic growth law for all living organisms of West, Brown, and Enquist (WBE) (West et al., 1999, 2001) may be fruitfully extended to cancer growth. In the form proposed by WBE, this law states that two hypotheses suffice to ensure the existence of a universal growth dynamics: the conservation of energy and the presence of a fractal

distribution network for energy supply at each part of the biological system considered. These basic hypotheses lead to the well-known exponent in the relationship between metabolic rate and mass scaling, which is purportedly characteristic of all organisms.

In a *microscopic* description, one should identify individual cell properties 31 and predict tumor development from cell-cell interactions. Such an approach 32 has two drawbacks: first, cancer growth is a collective phenomenon whose 33 complexity may not emerge from the ensemble of its individual cell properties 34 alone. Second, the huge number of cells involved (typically 10^8 cells for a 1 cm^3 35 tumor) restricts considerably the feasibility of simulations taking into account 36 the behavior and dynamics of each individual cell, unless populations with a 37 small number of individuals can be considered. As a consequence, *mesoscopic* 38 models (Scalerandi et al., 1999; Scalerandi et al., 2001b; Sansone et al., 2001a; 39 Ferreira et al., 2002; Scalerandi et al., 2002; Scalerandi and Sansone, 2002; 40 Chen et al., 2003) have been proposed, in which the coarse graining of the 41 system, and the behavior of cell clusters and their interactions is considered. 42 Mesoscopic models are also better adapted to describe the influence of the 43 macroscale world on microscale phenomena and viceversa. A nice exposition 44 of the insights that mathematical modeling can yield about the mechanisms 45 underpinning the great complexity of the various phases of cancer growth is 46 presented in a recent review by Byrne et al. (Byrne et al., 2006). Among other 47 recent papers about modeling tumor growth, we can quote another review 48 paper by Alarcón et al. (Alarcón et al., 2005) or, more generally, refer to the 49 repository of mathematical models and corresponding computational codes 50 assembled within the framework of the CViT (Center for the development of 51 a Virtual Tumor) Project (http://www.cvit.org), belonging to the US NIH-52 NCI ICBP (Integrative Cancer Biology Program). 53

Multicellular Tumor Spheroids (MTSs) represent convenient experimental 54 systems for analyzing and comparing tumor growth models at different lev-55 els. They are spherical aggregations of tumor cells that may be grown in 56 vitro under strictly controlled conditions, mimicking some of the important 57 features of solid tumors developing in vivo (Hamilton, 1998; Thomson and 58 Byrne, 1999; Mueller-Klieser, 2000; Chignola et al., 2000; Kelm et al., 2003). 59 Due to their simple geometry, the possibility of culturing them in large quan-60 tities and of controlling relevant parameters, such as the porosity and stiffness 61 of the surrounding environment, they are excellent systems upon which to 62 test the applicability of various models (Marusic et al., 1994: Delsanto et al., 63 2004, 2005a). Experimental spheroid setups are designed to provide a suitable 64 amount of oxygen and other nutrients, which diffuse to the outer edges of 65 the MTS, and then to the interior. Due to consumption by the outer region, 66 nutrient concentration decreases towards the center. Consequently, as it has 67 been observed, proliferating cells are usually present in the outermost shell, 68 quiescent (non reproductive) cells dominate in the interior, and, eventually at 69 a later stage of growth, a necrotic core is formed by dead cells. An MTS is 70 thus an heterogeneous cellular system of considerable complexity, but whose 71 properties and growth can be carefully monitored and modeled. 72

Using MTS's as working models (but with the expectation that our con-73 clusions may be applied also to the study of some aspects of *in vivo* tumor 74 growth), we review in Sections 2 and 3 two recently proposed growth mod-75 els at the mesoscopic (Delsanto et al., 2005b) and macroscopic (Guiot et al., 76 2003; Delsanto et al., 2004) levels, respectively. The latter is extended in order 77 to take into account the pressure exerted by the growing tumor against its 78 environment. This term plays a crucial role in the mechanisms involved in 79 tumoral invasion (Guiot et al., 2006b), which ultimately represents the proce-80

dure which allows further growth of tumors *in vivo*. Then, in Sec. 4, we show that it is possible to formulate a coherent picture embodying both descriptions by means of an Intermediate Model (IM) (Delsanto et al., 2005a). A detailed description of the IM and of the relationship (mediated by the IM) between the mesoscopic and macroscopic parameters represents the main goal of the present contribution.

87 2 A LINK TO CELL DYNAMICS: MESOSCOPIC MODEL

In this Section we review the mesoscopic MTS model of Ref. (Delsanto 88 et al., 2005b). In this model space is divided into concentric isovolumetric 89 shells n = 0, ..., N (n = 0 labels the central sphere of radius r_0). Each shell 90 has a volume $V_0 = (4/3)\pi r_0^3$, and a correspondence is established between 91 the shell system and a one-dimensional grid. The center of the shell system 92 coincides with the location of the implanted spheroid seed. The MTS growth is 93 controlled by local nutrient availability and proceeds according to the following 94 rules (see Fig. 1): 95

⁹⁶ a) Feeding: nutrient is absorbed by each shell at a rate γc_n , where c_n is the ⁹⁷ number of live cancer cells in the n^{th} shell.

b) Reproduction: cancer cells reproduce at rates $\rho_n c_n$, only in the shells where the number ν_n of locally available nutrient units exceeds a given threshold Q_R .

c) Migration: cells migrate to adjacent shells with a flux $\mu_n c_n$ per unit of grid cell surface if the number of nutrient units falls below a threshold Q_M . Since nutrients diffuse inwards, migration will usually proceed towards outer shells. We may plausible assume that outwards migration is also favored by mechanical stress gradients (Gordon et al., 2003). Due to the bias introduced

by the radial gradients, we distinguish between outwards (μ^+) and inwards (μ^-) migration fluxes.

d) Death: cell death occurs (at rates $\delta_n c_n$) where the number of nutrient units falls below the threshold Q_D .

The thresholds satisfy the conditions $Q_D < Q_M < Q_R$. Figure 1 summarizes in a simple scheme the set of thresholds associated to the different biological/biophysical phenomena and the corresponding value ranges for the rates. As reported in (Griffa et al., 2004) the actual implementation of the model makes use of sigmoidal functions for representing the rates dependence on nutrient concentration instead of Heaviside functions (i.e. thresholds).

¹¹⁶ Nutrients diffuse from the n^{th} shell to the adjacent ones at a rate $\alpha \nu_n$ per ¹¹⁷ unit area. Since cellular displacement and molecular diffusion between shells ¹¹⁸ are proportional to the areas of the separating surfaces, the diffusion terms ¹¹⁹ across the interface between the (n-1)-th and *n*-th shells will be proportional ¹²⁰ to $n^{2/3}$. The model equations are then written directly in their time-discretized ¹²¹ forms:

$$c_n^* = c_n (1 - \tau \delta_n + \tau \rho_n) + \tau r_0^2 [n^{2/3} (\mu_{n-1}^+ c_{n-1} - \mu_n^- c_n) + (n+1)^{2/3} (\mu_{n+1}^- c_{n+1} - \mu_n^+ c_n)],$$
(1)

$$l_n^* = d_n + \tau \delta_n c_n - \tau \lambda_n d_n, \tag{2}$$

$$\nu_n^* = \nu_n - \tau \gamma c_n + \tau \alpha r_0^2 \left[n^{2/3} (\nu_{n-1} - \nu_n) + (n+1)^{2/3} (\nu_{n+1} - \nu_n) \right], \qquad (3)$$

where τ is the time step and the asterisk means that the corresponding quantity must be evaluated at the time $t + \tau$, instead of at the time t. In Eq.(2) d_n is the number of dead cells in the n^{th} shell; the last term, with a coefficient λ_n , has been added to account for possible disintegration of dead cells releasing intracellular fluid (Frieboes et al., 2006).

¹²⁷ Numerical simulations based on the above model show that, in agreement
¹²⁸ with experimental observations, in a first stage the spheroid is fully populated

by proliferating cells. Then, in the inside, cells become quiescent, i.e. alive but 129 not proliferating, with a "wavefront" of proliferating cells which have greater 130 probability of undergoing an outward migration due to the second type of 131 movement mechanism implemented in the model, the one based on a smoothed 132 cell density-per each shell threshold, as explained in Ref. (Griffa et al., 2004). 133 This in silico mechanism corresponds to the biomechanical stress-mediated 134 transport of cells towards regions with lower levels of cell density, packing and 135 deformation, due to both passive (purely elastic) and cell-mediated responses 136 (Vernon et al., 1992; Dembo and Wang, 1999; Gordon et al., 2003; Deisboeck 137 et al., 2005) to mechanical deformations. Finally, at later times, a predomi-138 nantly necrotic core develops. According to the model implementation, when 139 the total number of living-plus-dead cells in the n-th shell is larger than a 140 given threshold, the probability per unit of time of migration towards the 141 nearest neighbor shells arises, so that some cells leave that shell. The thresh-142 old value is a function of the cell mean radius and of r_0 . Thus, the mechanical 143 stress-driven migration is essentially triggered by the unbalance between the 144 volume of each shell and the one occupied by the living and dead cells. If 145 the latter is too large, a higher level of packing and cell deformation occurs 146 with mass transport as consequence. Different weights are assigned in com-147 puting the volume occupied by the living vs. dead cells, in order to account 148 for their different deformation. Also, a rigid Heaviside function is replaced by 149 a sigmoid. See Ref. (Griffa et al., 2004) for the mathematical details. Figure 2 150 shows the radial distribution of viable cells within the simulated spheroid at 151 three different time steps belonging respectively to the three cited stages of 152 growth. The simulation aims at reproducing the typical layout of an MTS in 153 terms of cell state spatial distribution. 154

155 3 A LINK TO PHENOMENOLOGY: MACROSCOPIC MODEL

Both the discussion in the Introduction and Fig. 2 suggest the use of a three-156 layer macroscopic model for the tumor. Using the WBE model, as extended 157 to tumors in Ref. (Guiot et al., 2003), we assume that a central core of dead 158 cells (region Z_0) is surrounded by a first layer Z_1 of quiescent cells, and by an 159 outer layer Z_2 of active cells and neglect, as temporary, any mixing, i.e. the 160 presence of cells in the "wrong" regions. We label the three cell species 0, 1, 161 and 2 respectively, and call their corresponding masses m_i (i = 0, 1, 2). It is 162 important to remark that the central core Z_0 and the two layers Z_1 and Z_2 163 need not to be spherical, i.e. the macroscopic model may be used to describe 164 not only MTS's, but also almost any kind of pre-vascular in vivo solid cancers. 165 Since energy is transported to the tumor cells by the diffusing nutrients, we 166 assume that it is proportional to the amount of the latter. Therefore, applying 167 the law of energy conservation as in Ref. (Delsanto et al., 2004), we may write 168 the energy balance for region Z_2 as, 169

$$B_2 dt = N_2 \xi_2 dt + \epsilon dN_2 + \chi dN_2 \tag{4}$$

171 where

$$\chi = \left(\frac{\kappa PM}{\zeta}\right) \tag{5}$$

172

and N_2 is the total number of cancer cells in Z_2 at time t, P is the hydrostatic pressure on the spheroid wall, B_2 is the net nutrient-associated energy inflow into Z_2 during the interval dt, ζ is the mass density, assumed to be uniform, ξ_2 is the metabolic rate for a single cancer cell, and ϵ is the energy required to create a new cell in a "soft" environment. If M is the mass of a single cell, $m_2 = MN_2$.

In Eq. (4) the last term on the right-hand-side is new with respect to Ref. 179 (Delsanto et al., 2004), and represents the amount of energy required by the 180 volumetric expansion. It includes not only the mechanical work done by the 181 biologically growing system against its environment at pressure P, i.e. (from 182 thermodynamics) PdV_2 , which is comparatively small (and would correspond 183 to a value of $\kappa = 1$), but also the excess energy required to create new biological 184 material in a stressed environment. Such excess energy is the result of complex 185 biological processes and can be assumed (at least as a first approximation) to 186 be proportional to the volume increase, $dV_2 = M dN_2$. 187

Two plausible and complementary assumptions can be formulated regarding 188 the pressure dependence, as a function of the constitutive laws of the matrix 189 and tumor materials. In a recent paper (Guiot et al., 2006b), the interface pres-190 sure increases due to the tumor elastic growth, until a characteristic strength 191 of the matrix is reached, the stress released and a new annular region of matrix 192 colonized by the tumor. In this model a perfectly plastic constitutive law for 193 the matrix is also assumed, corresponding to a strain flow at a given value 194 P. For such a case the pressure P is expected to be constant, representing 195 the yielding strength of the matrix material. Obviously these two assumptions 196 represent limit conditions of a more complex reality, but the introduction of 197 P represents an important novelty with respect to previous models (for which 198 P = 0, due to the role of mechanical stress in triggering and controlling var-199 ious biomechanical and biophysical processes (Chaplain, 2006) both at the 200 single cell level (remodulation of intracellular structures by the cell deforma-201 tion and consequent change in gene expression and protein synthesis (Ingber 202 et al., 1995) and at the multicellular one (change of intercellular communica-203 tions via release of molecules, apoptosis and cell proliferation control via the 204 change of cell adhesion (Shraiman, 2005; Hufnagel et al., 2007; Helmlinger 205

²⁰⁶ et al., 1997)).

The left-hand-side of Eq. (4), B_2dt , describes the energy necessary for the 207 overall thermodynamic processes occurring within the cell population occupy-208 ing the region Z_2 . This energy, as previously stated, results from the balance 209 of two different flows of nutrients molecules (the carriers of energy for every 210 metabolic process): the total diffusive inflow towards the spheroid, which is 211 proportional to a power p_2 of its whole mass, according to the general law 212 relating the mass of a biological system and the rate of energy inflow needed 213 for maintenance and proliferation (West et al., 1999, 2001; Guiot et al., 2003), 214 and the nutrient molecules flow towards the inner region Z_1 , which produces 215 an energy leakage rate for the Z_2 population, but an energy acquisition rate 216 for the Z_1 one proportional to the p_1 power of its mass, according to the same 217 law. Thus, 218

²¹⁹
$$B_2 = B_{02}(m_0 + m_1 + m_2)^{p_2} - B_{01}(m_0 + m_1)^{p_1},$$
 (6)

where B_{01} and B_{02} are constants. The assumption that cellular feeding is 220 controlled only by diffusive processes, i.e. that the tumor is in a pre-vascular 221 stage, would imply that $p_1 = p_2 = 2/3$, because the inflow through the sur-222 face is proportional to the 2/3 power of the volume. By contrast WBE's law 223 assumes that resources are transported to the cells through a fractal hierar-224 chical branching network (West et al., 1999), which implies p = 3/4. In fact, 225 the power law dependence of B on m has been recognized for a long time 226 (Kleiber, 1932), but the value of the exponent and its ultimate meaning are 227 still sources of controversy (Dodds et al., 2001; Makarieva et al., 2003). It has 228 been recently pointed out that our model (and in particular a careful moni-220 toring of the p exponent) may enable us to predict the progression of a tumor 230 (Guiot et al., 2006a; Carpinteri and Pugno, 2005; Guiot et al., 2006b). 231

On heuristically justified grounds, we can also apply the same formulation to describe cell evolution in Z_0 and Z_1 , using the same approach and the same type of rules which connect the energy rate consumed by a cell population to its total mass through a power law. By extending the definition of B_{02} and ξ_2 to the other two regions and using Eqs. 4 and 6, the subsequent equation for the temporal evolution of the masses of each cell population is obtained:

$$\frac{dm_i}{dt} = a_i \left(\sum_{j=0}^i m_j\right)^{p_i} - a_{i-1} \left(\sum_{j=0}^{i-1} m_j\right)^{p_{i-1}} - b_i m_i,\tag{7}$$

239

where m_i is the total mass of the *i*-th population, for i = 0, 1, 2, and

$$a_i = \frac{MB_{0i}}{\epsilon + \kappa PM/\zeta},\tag{8}$$

(9)

 $b_i = \frac{\xi_i}{\epsilon + \kappa P M / \zeta},$

with $a_{-1} = 0$. The growth of the tumor mass in each region is proportional 242 to the difference between the net energy input (first term minus second term 243 in Eq. 7 and the amount of energy used for cell maintenance (third term). 244 Equations (7) to (9) define a consistent phenomenological model whose pa-245 rameters may be evaluated from the results of macroscopic experiments. They 246 belong to a class, called U2, of a recently proposed classification scheme for 247 phenomenological universalities in growth problems (Castorina et al., 2006). 248 U2 includes, as special cases, the WBE, the logistic and all the other previ-249 ously proposed growth models. The introduction of the expansion term in Eq. 250 (4) leads to a reduction in the size of most of the coefficients, a fact whose 251 consequences will be explored later. The outer shell coefficients a_2 and b_2 can 252 be obtained from experimental observations of the whole spheroid (Condat 253 and Menchón, 2006). As indicated in the Appendix, the coefficients a_0 and b_0 254 can be obtained form observations of the necrotic core. 255

If we assume that necrotic cells do not consume energy $\xi_0 = 0$. Then $b_0 = 0$ and Eq. (7) for m_0 can be straightforwardly solved, yielding,

$$m_0(t) = \left[(\tilde{m})^{1-p_0} + a_0(1-p_0)(t-t_0) \right]^{1/(1-p_0)}$$
(10)

where $p_0 \neq 1$ and \tilde{m} is the mass at time t_0 . If we choose t_0 as the time of onset of the necrotic core, $\tilde{m} = 0$. At long times the necrotic mass increases as a power law; in particular, if $p_0 = 2/3$, $m_0(t) \sim t^3$. For $p_0 = 1$, there is exponential growth at all times. Since a_0 is expected to decrease with increasing pressure, the necrotic core (and the rest of the tumor) will grow more slowly under conditions of higher pressure. A tumor steady state can be obtained from Eqs. (7) only if $b_0 \neq 0$.

266 4 THE BRIDGE: INTERMEDIATE MODEL

To formulate an auxiliary model that allows us to bridge the gap between the mesoscopic and macroscopic descriptions, we start by rewriting the equation describing the evolution of the number of cancer cells in the n^{th} mesoscopic shell as a first order differential equation that explicitly exhibits the cell fluxes. Assuming that the time step τ is very small, we write,

The mesoscopic Eq.(1) then becomes

 $c_n^* - c_n \approx \tau \frac{dc_n}{dt}$

$$\frac{dc_n}{dt} = \beta_n c_n + \Phi_{n-1} - \Psi_{n-1} + \Psi_n - \Phi_n, \qquad (12)$$

where Φ_n and Ψ_n stand for the outward and inward cell flows, respectively

$$\Phi_n = (n+1)^{2/3} r_0^2 \mu_n^+ c_n, \tag{13}$$

(11)

$$\Psi_n = (n+1)^{2/3} r_0^2 \mu_{n+1}^- c_{n+1}, \tag{14}$$

where $\beta_n = \rho_n - \delta_n$ is the difference between the reproduction and death rates, except in Z_0 where $\beta_n = -\lambda_n$.

Next we regroup the shells with $n \leq n_0$, $n_0 < n \leq n_1$ and $n_1 < n \leq n_2 = N$ 278 in the three regions, Z_0 , Z_1 and Z_2 , respectively. Although the equation de-279 scribing the nutrient evolution in the mesoscopic model (Eq. 3) is not explic-280 itly considered, the role of the nutrient distribution is clear: since the nutrient 281 comes from outside and is progressively consumed (or stored for later con-282 sumption) by the viable cells inside, the above stratification is a reasonable 283 approximation (Mueller-Klieser, 2000). Of course, due to the randomness in-284 herent to the involved processes, we would expect a smoother distribution in 285 the experimental data. 286

In order to set the model in a form adequate for the comparison, we proceedas follows:

1. At each discretization step we will conceptually separate the description 289 of growth into two stages. In the first stage cells multiply, migrate and die, but 290 the volumes of the various zones are kept constant. In the second stage, the 291 zone radii are allowed to vary in order to restore cell concentration uniformity. 292 Therefore the rescaled zone volumes "recapture" the cells that left each zone 293 during the first step; this corresponds to a loss of mass $-dm_i$ for the zone Z_i . 294 2. Since we are solely interested in the number of cells inside each region, 295 we can sum over the contributions of all its shells. Internal migrations in each 296 region do not affect the number of cells therein and are therefore irrelevant. 297 The only migration terms that matter are those across the region boundaries, 298 that is to the shells $n = n_0$, n_1 and $n_2 = N$. 299

300 3. By definition, all cells are included in the MTS; thus, no cells can enter 301 from the outside: $\Psi_{n_2} = 0$. We will disregard the centripetal migrations Ψ_{n_0} 302 and Ψ_{n_1} since they would amount to including active cells in Z_1 or quiescent

ones in Z_0 (which is forbidden by our definition of the regions). Operationally, this implies setting the transport coefficients $\mu_{n_i}^- = 0$ at the region edges. Alternatively, we may redefine $\bar{\Phi}_n$ as the net balance $\Phi_n - \Psi_n$.

4. As a coarse grain approximation, the rates δ_n , ρ_n and λ_n may be rea-306 sonably equated in each region to their asymptotic values or to 0 as a conse-307 quence of the sigmoidal functions implementing their dependence from nutri-308 ent concentrations. This corresponds to selecting very steep sigmoidal func-309 tions (Griffa et al., 2004). It follows that $\delta_n = \rho_n = 0$ in the first (necrotic) 310 region, since dead cells cannot die or reproduce; we also fix $\lambda_n = \lambda$ in the 311 same region. In Z_1 we set $\delta_n = \delta$ and $\rho_n = 0$, while in Z_2 we write $\delta_n = 0$ and 312 $\rho_n = \rho$, with both ρ and δ being the asymptotic values of the corresponding 313 variables. 314

With the above assumptions, we can use the set of Eqs. (12) to obtain equations for the total numbers of cells $\bar{c}_i = \sum c_j$ in their respective regions Z_i (i = 0, 1, 2). Summing Eqs. (12) over n for all the shells included in each of the three regions Z_i , we obtain

$$\frac{d\bar{c}_2}{dt} = \rho\bar{c}_2 + \bar{\Phi}_{n_1} - \bar{\Phi}_{n_2},\tag{15}$$

$$\frac{d\bar{c}_1}{dt} = -\delta\bar{c}_1 + \bar{\Phi}_{n_0} - \bar{\Phi}_{n_1},$$
(16)

(17)

 $\frac{d\bar{c}_0}{dt} = -\bar{\Phi}_{n_0} - \lambda\bar{c}_0,$

The outward cell flux from the outermost shell of each region Z_i (i = 0, 1, 2)across the surface separating it from the next region (the matrix in the case i = 2) can be renamed as $\bar{\Phi}_i$: $\bar{\Phi}_0 = \bar{\Phi}_{n_0}$, $\bar{\Phi}_1 = \bar{\Phi}_{n_1}$ and $\bar{\Phi}_2 = \bar{\Phi}_{n_2}$, where $n_2 = N$.

In Eq. (16) $\delta \bar{c}_1$ is the total dying cell rate. Since cells in the necrotic core may only be reabsorbed, the only contributions to cell variation in Z_0 are given by the flux $-\bar{\Phi}_0$ and the absorption $-\lambda \bar{c}_0$.

Next we evaluate the fluxes $\bar{\Phi}_i$. The factor $r_0^2(n+1)^{2/3}$ in Eqs. (13) and (14), 328 for $n = n_i$ (i = 0, 1, 2), corresponds to the square of the external radius, R_i^2 , of 329 the corresponding zone, Z_i . Because migrations to other regions start only from 330 the outermost shell in each region, the fluxes will be proportional to $\eta \mu_i$ (i = 331 (0, 1, 2), where $\mu_i \equiv \mu_{n_i}^+$ is the local mobility and $\eta = c_{n_i}$ is the mean number 332 of cells in the n_i -th isovolumetric shell. We keep η approximately constant 333 because of the migration mechanism based on the mechanical stress-driven 334 mass transport. In fact the outer shell of each zone is subject to an outwards 335 chemotaxis-based cell migration due to the gradient in the concentration of 336 nutrients (Dorie et al., 1982, 1986; McElwain and Pettet, 1993) (which tends 337 to increase local cell density), but also to the mechanical opposition of the 338 external cells (or the extracellular matrix in the case of the proliferant rim Z_2). 339 The intermediate increase of cell density is then compensated by the stress-340 driven migration (invasion in the case of the proliferative rim), in order not to 341 increase the cell density above its maximum allowed value (Deisboeck et al., 342 2005). The constant cell density assumption in the formulation of the IM is 343 in agreement with many experimental observations ((Freyer and Sutherland, 344 1986b, 1985, 1980), although other recent investigations (Nirmala et al., 2001) 345 show that it may slightly vary in time and space due to the local heterogeneities 346 at smaller scales within the MTS. The fluxes can be written as, 347

$$\bar{\Phi}_i = \mu_i c_{n_i} R_i^2 = \mu_i \eta \left(\frac{3}{4\pi\zeta} \sum_{j=0}^i m_j \right)^{2/3}.$$
(18)

Next, we correlate Eqs. (7) for the mass variations with Eqs.(15-17) for the variations in cell numbers. Using the two-stages model at each discretization step, as discussed before, from $m_i = M\bar{c}_i$, it follows

$$\frac{d\bar{c}_i}{dt} = -\frac{1}{M}\frac{dm_i}{dt} \tag{19}$$

Now we can combine Eqs. (15) to (19) to write the equations for the mass 353 variations in each of the three zones: 354

$$\frac{dm_2}{dt} = -\rho m_2 - \sigma \mu_1 (m_0 + m_1)^{2/3} + \sigma \mu_2 (m_0 + m_1 + m_2)^{2/3}, \tag{20}$$

$$\frac{dm_1}{dt} = -\delta m_1 - \sigma \mu_0 m_0^{2/3} + \sigma \mu_1 (m_0 + m_1)^{2/3}, \qquad (21)$$

(22)

and 355

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$$\frac{dm_0}{dt} = -\lambda m_0 + \sigma \mu_0 m_0^{2/3},$$
(22)
where
$$\sigma = M\eta \left(\frac{4\pi}{3}\zeta\right)^{-2/3}.$$
(23)

By comparing Eqs. (7) with Eqs. (20)-(22), we find that both models coin-359 cide if the following parameter identification is performed: 360

$$a_0 = \sigma \mu_0, \qquad a_1 = \sigma \mu_1, \qquad a_2 = \sigma \mu_2,$$
 (24)
 $b_0 = -\lambda, \qquad b_1 = -\delta, \qquad b_2 = \rho$ (25)

and 361

362

$$p_0 = p_1 = p_2 = 2/3 \tag{26}$$

We have thus related the parameters corresponding to the mesoscopic and 363 macroscopic formulations. That $p_i = 2/3$ was to be expected by the con-364 struction of the mesoscopic model, but the other relations provide us with 365 information about the dependence of the mesoscopic parameters on pressure. 366 For instance, the interzone migration fluxes must decrease with increasing 367 pressure, 368

$$\mu_i = \frac{MB_{0i}}{\sigma \left(\epsilon + \frac{\kappa PM}{\zeta}\right)}.$$
(27)

This is reasonable, because migration is hindered by the increased cell concentration. The reproduction rate is decreased by the same factor:

$$\rho = \frac{\xi_2}{\epsilon + \frac{\kappa PM}{\zeta}},\tag{28}$$

which is also consistent with predictions put forward on the basis of exper-373 imental data (Helmlinger et al., 1997). Since it is not a priori clear what the 374 dependence of b_0 and b_1 on P should be, we cannot draw general conclusions 375 on the mesoscopic coefficients λ and δ . However, if we assume that the depen-376 dence of b_1 on P is the same as that of b_2 , then we can conclude that the death 377 rate is decreased by increasing pressure, a somewhat surprising result, which 378 however, agrees with the results of Helmlinger and co-workers, who observed 379 that solid mechanical stress decreases the apoptotic rate (Helmlinger et al., 380 1997). This decrease in the apoptotic rate is likely to be due to increased 381 packing and concomitantly enhanced cell-cell interactions, which trigger the 382 suppression of apoptotic cell death. Moreover, $\delta > 0$ implies $b_1 < 0$, that is, 383 the proliferating - to - quiescent bulk transformation is faster than the quies-384 cent - to - dead bulk transformation. Similarly, $\lambda > 0$ implies that the bulk 385 dying rate must be larger than the reabsorption rate. 386

387 5 CONCLUSIONS

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Through the introduction of an intermediate model, we have proved the consistency of two very different models for heterogeneous MTS growth: a macroscopic model, based on the ontogenetic growth model of West, Brown and Enquist, and a mesoscopic one, based on the coarse-graining of the cell system. Besides its intrinsic importance as a bridging tool, this unification helps us to establish a correspondence between hard-to-measure microscopic

³⁹⁴ parameters and their more accessible macroscopic counterparts (see e.g. Fig.³⁹⁵ 3).

The correspondence between the two models is remarkable, since they stem from completely different approaches, as stated in the Introduction. It is important to note that:

1. The IM model in its present form is not self-standing, i.e. it cannot by
itself be used for actual simulations, since it implicitly depends on the evolving
nutrient distribution (provided by the Eq. (3) in the mesoscopic model).

2. The consumption rate of nutrients due to the metabolism, which is explicit in the mesoscopic model is only implicit in the IM, via the "ignored" Eq. 3. Thus the metabolism parameters b_i correspond to the "cell activity" parameters in β_i in the IM.

3. According to Eqs. (20) to (22), the parameters p_i are all equal to 2/3, which indicates that feeding must be diffusion-controlled to ensure consistency between the models. However, the values of the parameters p_i depend on the nature of nutrient transport. For instance, MTS internal vascularization is likely to change their values.

The present treatment lends itself to suggest, through simulations and math-411 ematical analysis, the relevance of the effects of different tumor microenviron-412 ments, some of which are easier to study in vitro or by implanting MTS in 413 model laboratory animals (Oudar, 2000). For example, we could extend it to 414 the case of underfeeding (Delsanto et al., 2004; Griffa and Scalerandi, 2005) 415 and to the development of angiogenesis. In the latter case, it would be useful 416 to find a connection between the various extant mesoscopic model (Dodds 417 et al., 2001; Byrne and Chaplain, 1995; Byrne et al., 2006), and the simple 418 predictions about the dependence of the growth rate on the instantaneous 419 mass furnished by the macroscopic approach (Menchón and Condat, 2007). 420

One could also explicitly consider the presence of growth inhibitors or consider 421 the role of the necrotic mass in regulating the number of viable quiescent cells 422 at the interface between the Z_0 and Z_1 regions. It is well known that necrotic 423 death is closely related to the release of cytotoxic intra-cellular substances in 424 the extra-cellular microenvironment (Greenspan, 1974; Freyer, 1988; Groebe 425 and Mueller-Klieser, 1996). This process deserves to be considered with care. 426 To conclude, we have used the simplest possible form for the dependence of 427 the extra term in the WBE equation on the pressure. Our results could be 428 easily generalized to more complicated cases by replacing κP in Eq. (4) with 429 a suitable positive, monotonically increasing function F(P). This would not 430 introduce any qualitative changes in the results obtained here. Finally, we 431 must remark that in our model the growth process is controlled by nutrient 432 consumption and the possible influence of growth promoters/inhibitors is ne-433 glected. We also have neglected part of the cell-cell interactions and cell-cycle 434 details (Jiang et al., 2005). 435

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444 FIGURE CAPTIONS

Fig.1: Schematic representation of the basic cellular processes in the n-th MTS shell as a function of the number of available nutrient units ν_n . The three lines below the ν_n axis imply that cellular death occurs only for $\nu_n < Q_D$, migration for $\nu_n < Q_M$ and cell reproduction for $\nu_n > Q_R$, respectively. Correspondingly, the three parameters δ_n , μ_n and ρ_n are different from zero (and equal to δ,μ and ρ) only for $\nu_n < Q_M$, $\nu_n < Q_M$ and $\nu_n > Q_R$, respectively.

Fig. 2: The three stages of development of a MTS, as obtained from a mesoscopic simulation at three successive times. In the first stage all cells are proliferating. In the second one a region of quiescent cells emerges and soon occupies most of the interior. In the third one a necrotic core develops, which, however, includes some viable cells in the process of dying (especially at its rim). The mesoscopic parameters for the simulations are $\mu = 0.003$, $\rho = 0.025$, $\delta = 0.01$, $\sigma = 1.7$ in the nonzero zone defined in Fig.1.

Fig. 3: Comparison between the results of a macroscopic (dashed line) and a mesoscopic (full dots) simulation with the experimental data (empty squares, ref. (Freyer and Sutherland, 1986a)) referring to a MTS made of EMT6/Ro mouse mammary carcinoma cells grown in a culture medium. The parameters for the macroscopic simulations are: $a_1 = 0.15$, $a_2 = 0.48$, $a_3 = 0.59$, $b_0 =$ -0.012, $b_1 = 0.0016$, $b_2 = 0.037$; the mesoscopic parameters are the same as in Fig.2.

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Fig. 2.



Fig. 3.