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A Mechanistic Investigation into Ischemia-Driven Distal Recurrence of Glioblastoma

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Abstract Glioblastoma (GBM) is the most aggressive primary brain tumor with a short median survival. Tumor recurrence is a clinical expectation of this disease and usually occurs along the resection cavity wall. However, previous clinical observations have suggested that in cases of ischemia following surgery, tumors are more likely to recur distally. Through the use of a previously established mechanistic model of GBM, the Proliferation Invasion Hypoxia Necrosis Angiogenesis (PIHNA) model, we explore the phenotypic drivers of this observed behavior. We have extended the PIHNA model to include a new nutrient-based vascular efficiency term that encodes the ability of local vasculature to provide nutrients to the simulated tumor. The extended model suggests sensitivity to a hypoxic microenvironment and the inherent migration and proliferation rates of the tumor cells are key factors that drive distal recurrence.

Keywords glioblastoma · hypoxia · PIHNA · ischemia · tumor growth

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1 Introduction

Glioblastoma (GBM) is the most aggressive primary brain tumor [24]. It is uniformly fatal with a median survival from diagnosis of only 15 months with standard of care treatment, consisting of a combination of resection, chemotherapy and radiotherapy [37]. Due to the sensitive location of the tumor, there is a reliance on clinical imaging to assess tumor treatment response and progression. Enhancement on T1-weighted magnetic resonance imaging (T1Gd MRI) with gadolinium contrast shows regions where gadolinium has leaked through disrupted vasculature. T2-weighted MRI (T2 MRI) shows infiltrative edema, fluid that has leaked from vasculature. Abnormalities on T1Gd MRI spatially correlate with the bulk of the tumor mass with central dark regions typically showing necrosis, whereas edema visible on T2 MRI corresponds to regions of lower tumor cell density.

An unfortunate clinical expectation following surgical resection is tumor recurrence, which usually presents on the edge of the resection cavity [9]; this is known as a local recurrence. The recurrent tumor will occasionally enhance on T1Gd MRI in a different region of the brain, away from the primary site, which is known as a distant recurrence [9]. In some cases, the T2 MRI abnormality will become much larger relative to the enhancement on T1Gd MRI, these cases are known diffuse recurrences [9]. In a retrospective study by Thiebold *et al.* it was shown that patients with GBM who had also suffered from perioperative ischemia, defined as an inadequate blood supply to a part of the brain following resection, were more likely to have a distantly and/or diffusely recurring GBM [45]. A disruption in normal vasculature can occur following GBM resection and can lead to ischemia, affecting abnormal tissue in the same way it affects the healthy tissue. By reducing available nutrients to the tumor, the tumor is forced towards a hypoxic phenotype and becomes necrotic if the reduction is sustained. Thiebold attributed the observed difference in recurrence patterns to the hypoxic conditions caused by the reduction in vasculature [45]. In retrospective analyses of patient data, Bette *et al.* found further supporting evidence that perioperative ischemia promoted aggressive GBM recurrence patterns [5,6]. Bette *et al.* showed that perioperative infarct volume was positively associated with more multifocal disease and contact to the ventricle, which have both been shown to negatively impact patient survival in a pretreatment setting [1,35].

Spatiotemporal mathematical models have been used extensively to describe the growth of GBM. These models incorporate features of tumor cells such as cell phenotype, migration, proliferation and interactions with other cells to understand how these influence observed growth behavior in GBM. Such models have the ability to provide mechanistic insight into observed tumor growth patterns and treatment effects. Mathematical models have been created to simulate GBM growth on varying spatial scales from clusters of cells [15,25,36] and murine models [16,22,32] to tissue-level scales seen throughout the presentation of the disease in patients [18,38,39,41,42,44]. Various treatments for GBM have also been modeled, such as resection [27,44], chemotherapy [2,4,7] and radiotherapy [8,22,30], which are all elements of the current standard of care. Other less widely used and experimental treatments have also been modeled such as anti-angiogenic drugs [18,33] and oncolytic virus therapy [13].

An example of a tissue-level growth model of GBM is the Proliferation Invasion Hypoxia Necrosis Angiogenesis (PIHNA) model, which has been used to

study different mechanisms of tumor development and shows similar growth and progression patterns to those seen in patient tumors [43]. Simulated hypoxic events have shown an increase in glioma growth rates in spatiotemporal models of GBM [26, 28]. We have recently found the parameters of the PIHNA model that drive faster outward growth of these simulated tumors and found that those relating to hypoxia were in some cases extremely influential [12].

To the best of our knowledge, the impact of perioperative ischemia on recurrence in GBM has not been mathematically modeled before. We aim to use the PIHNA model to gain insight into the tumor kinetics that may play a role in this behavior.

In this work, we extend a term in the PIHNA model known as the vascular efficiency term, which determines the ability of local vasculature to provide nutrients to the tumor. We carry this out through the inclusion of a nutrient-transport equation parametrized through glucose uptake rates in GBM. We apply this extended PIHNA model to a set of simulated perioperative ischemia cases to determine influential mechanisms in the model that could drive ischemia-induced distal recurrence patterns in GBM. Specifically, we vary migration and proliferation rates of normoxic and hypoxic cell phenotypes, as well as switching rates between these phenotypes. We find that simulated tumors with faster migration and slower proliferation rates are more likely to recur distantly in cases of perioperative ischemia. We see that this can be promoted by changes in switching rates between normoxic and hypoxic cell phenotypes. We have also simulated these same cases with a less intense ischemic event, and show that this in turn leads to less distantly recurring tumors. Following an initial exploration of simulated resection and ischemic injury, we present a second case example of simulated perioperative ischemia in which we observe similar recurrence patterns that depend on migration and proliferation rates of the normoxic cells. We also present two example simulations, one with and one without ischemia, to show that ischemia can offset the growth of dense tumor in simulations, which may contribute to diffuse recurrence following perioperative ischemia.

2 Methods

2.1 The PIHNA Model

To simulate glioblastoma growth and spread, we have adapted a previously established tumor growth model – the PIHNA model [12, 43]. The PIHNA model describes the growth of GBM with the interactions of vasculature in the process of angiogenesis. This model simulates five different species, that all depend on space and time, and their interactions:

- $c(\mathbf{x}, t)$ – the density of normoxic tumor cells,
- $h(\mathbf{x}, t)$ – the density of hypoxic tumor cells,
- $n(\mathbf{x}, t)$ – the density of necrotic cells,
- $v(\mathbf{x}, t)$ – the density of vascular endothelial cells,
- $a(\mathbf{x}, t)$ – the concentration of angiogenic factors.

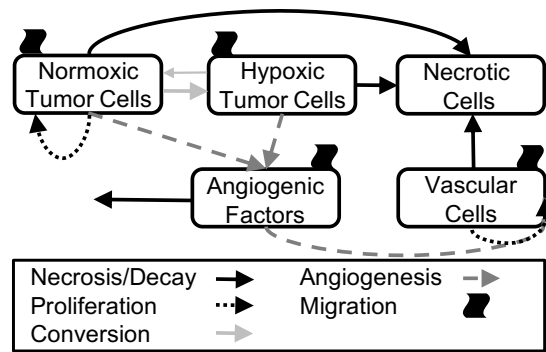


Fig. 1: A schematic for the PIHNA model. Normoxic tumor cells (c) proliferate, migrate, convert towards hypoxia and can become necrotic. Hypoxic tumor cells (h) migrate and can convert back to normoxic cells or to necrotic cells. Necrotic cells (n) accumulate as other cell types die. Angiogenic factors (a) are created in the presence of normoxic and hypoxic cells, migrate, decay and promote the local creation of vasculature. Vascular cells (v) proliferate through the facilitation of angiogenic factors and migrate.

Normoxic cells proliferate with rate ρ and migrate with rate D_c , whereas hypoxic cells do not proliferate and migrate with rate D_h . Cells convert from normoxic to hypoxic phenotypes (with rate β) and from hypoxic to normoxic phenotypes (with rate γ) depending on the ability of the local vascular density to provide nutrients at their location; hypoxic cells in the model become necrotic if they remain in a vasculature-poor region with rate α_h . When any other cell type meets a necrotic cell, they become necrotic with rate α_n , as necrotic cells have been shown to encourage cell death through the creation of an unfavorable microenvironment [29, 48]. Angiogenic factors migrate with rate D_a , are created by the presence of normoxic and hypoxic tumor cells (with rates δ_c and δ_h , respectively), decay naturally (λ) and are consumed through the creation and presence of vascular cells. For a more in depth justification of these model parameters, see Curtin et al. and Swanson et al. [12, 43]. The parameter values used in this work, as well as their units, can all be found in Table 1.

Following the literature [40], we have assumed that a high total relative cell density of at least 80% is visible on a T1Gd MRI through the aforementioned imaging abnormalities of enhancement and necrosis present on the image. We have also assumed a total relative density of at least 16% is visible on a T2 MRI, due to the spatial correlation between lower cell densities and edema. In the PIHNA model, this translates to the total cell density $T \geq 0.8$ being visible on a T1Gd MRI and $T \geq 0.16$ being visible on a T2 MRI. By construction, the T1Gd lesion is always less than or equal in size to the T2 lesion, which agrees with patient data [17].

We present a schematic for the PIHNA model in Figure 1 that indicates the migration and proliferation of individual species as well as the interactions between all model species. The PIHNA model itself is presented in Equations (1)-(5), which we have annotated to give a full description of each of the terms in the model.

$$\underbrace{\frac{\partial c}{\partial t}}_{\text{Change of normoxic cell density}} = \underbrace{\nabla \cdot (D_c(1-T)\nabla c)}_{\text{Net diffusion of normoxic glioma cells}} + \underbrace{\rho c(1-T)}_{\text{Net proliferation of normoxic glioma cells}} + \underbrace{\gamma h V}_{\text{Conversion of hypoxic to normoxic}} - \underbrace{\beta c(1-V)}_{\text{Conversion of normoxic to hypoxic}} - \underbrace{\alpha_n \frac{nc}{K}}_{\text{Conversion of normoxic to necrotic}} \quad (1)$$

$$\underbrace{\frac{\partial h}{\partial t}}_{\text{Change of hypoxic cell density}} = \underbrace{\nabla \cdot (D_h(1-T)\nabla h)}_{\text{Net diffusion of hypoxic glioma cells}} - \underbrace{\gamma h V}_{\text{Conversion of hypoxic to normoxic}} + \underbrace{\beta c(1-V)}_{\text{Conversion of normoxic to hypoxic}} - \underbrace{\left(\alpha_h h(1-V) + \alpha_n \frac{nh}{K}\right)}_{\text{Conversion of hypoxic to necrotic}} \quad (2)$$

$$\underbrace{\frac{\partial n}{\partial t}}_{\text{Change of necrotic cell density}} = \underbrace{\alpha_h h(1-V)}_{\text{Conversion of hypoxic to necrotic}} + \underbrace{\alpha_n \frac{n(c+h+v)}{K}}_{\text{Contact necrosis of all living cells}} \quad (3)$$

$$\underbrace{\frac{\partial v}{\partial t}}_{\text{Change of vascular cell density}} = \underbrace{\nabla \cdot (D_v(1-T)\nabla v)}_{\text{Net diffusion of vasculature}} + \underbrace{\mu \frac{a}{K_m + a} v(1-T)}_{\text{Net proliferation of vasculature}} - \underbrace{\alpha_n \frac{nv}{K}}_{\text{Conversion of vasculature to necrotic}} \quad (4)$$

$$\underbrace{\frac{\partial a}{\partial t}}_{\text{Change of angiogenic factor conc.}} = \underbrace{\nabla \cdot (D_a \nabla a)}_{\text{Net diffusion of angiogenic factor}} + \underbrace{\delta_c c + \delta_h h}_{\text{Net production of angiogenic factor}} - \underbrace{q\mu \frac{a}{K_m + a} v(1-T)}_{\text{Net consumption of angiogenic factor}} - \underbrace{\omega a v}_{\text{Decay}} - \underbrace{\lambda a}_{\text{Decay}} \quad (5)$$

where

$$V(c, h, v) = \frac{v}{v + \frac{\eta_c(D_c, \rho)c + \eta_h(D_h)h}{ps}} \quad (6)$$

and

$$T = (c + h + n + v)/K. \quad (7)$$

The term V is called the vascular efficiency and it models the relationship between the vasculature and its effect on the tumor. We let V take values in $[0, 1]$ such that it affects the switching rates between the normoxic (c), hypoxic (h) and necrotic (n) cell populations. When vasculature is abundant relative to other cells, V is close to 1 representing ample nutrient supply. Whereas when vasculature is relatively low, V is close to 0, which represents an unfavorable microenvironment of limited nutrient supply; this promotes conversion towards hypoxic and necrotic cells. In this work, we have extended the vascular efficiency term from previous iterations of the PIHNA model and present the derivation of this term in the next section.

The model equations are run on a two-dimensional slice of a realistic brain geometry from the Brainweb Database [10, 11, 20, 21], which spatially differentiates physiological structures such as white matter, grey matter, cerebrospinal fluid (CSF) and anatomical boundaries of the brain. This geometry is an average of multiple MR scans on a single patient to create a brain geometry with 1mm accuracy on and between MR slices. This gives a voxel volume of 1mm^3 , which

	Definition	Value/Range	Units	Source
D_c	Diffusion rate of normoxic cells	1 – 1000	$\frac{\text{mm}^2}{\text{year}}$	[17]
D_h	Diffusion rate of hypoxic cells	$(0.1 - 100)D_c$	$\frac{\text{mm}^2}{\text{year}}$	[17, 26]*
ρ	Proliferation rate of normoxic cells	10 – 100	1/year	[17]
β	Switching rate from normoxia to hypoxia	$0.1\rho, 0.5\rho$	1/year	[43]*
γ	Switching rate from hypoxia to normoxia	0.005, 0.05, 0.5	1/day	[43]
α_h	Switching rate from hypoxia to necrosis	0.1β	1/year	[43]*
α_n	Rate of contact necrosis	$\log(2)/50$	1/day	[31]
D_v	Diffusion rate of endothelial cells	0.18	$\frac{\text{mm}^2}{\text{year}}$	[43]
D_a	Diffusion rate of angiogenic factors	3.15	$\frac{\text{mm}^2}{\text{year}}$	[43]
δ_c	Normoxic cell production rate of angiogenic factors	2.77×10^{-13}	$\frac{\mu\text{mol}}{\text{cell} \times \text{year}}$	[43]
δ_h	Hypoxic cell production rate of angiogenic factors	5.22×10^{-10}	$\frac{\mu\text{mol}}{\text{cell} \times \text{year}}$	[43]
μ	Angiogenesis vasculature production rate	$\log(2)/15$	1/day	[43]
q	Consumption of angiogenic factors per cell	1.66	$\mu\text{mol}/\text{cell}$	[43]
λ	Natural decay rate of angiogenic factors	15.6	1/day	[43]
ω	Rate of removal of angiogenic factors by vasculature	λ/v_0	$\frac{1}{\text{cell} \times \text{day}}$	[43]
K	Maximal cell density	2.39×10^5	cells/mm ³	[43]
P_c^w	Glucose consumption ratio for normoxic cells in white matter	1.66 – 4.5	-	[14]*
P_h^w	Glucose consumption ratio for hypoxic cells in white matter	1.66 – 4.5	-	[14]*
P_c^g	Glucose consumption ratio for normoxic cells in grey matter	0.5 – 2	-	[14]*
P_h^g	Glucose consumption ratio for hypoxic cells in grey matter	0.5 – 2	-	[14]*

Table 1: Parameter definitions and values for the PIHNA model. *Parameters that we have added/alterd from the original publication of the PIHNA model [43].

we use to track tumor volume on the two-dimensional brain slice. The simulations are implemented on white and gray matter, with differences in initial vasculature density and nutrient consumption ratios between these tissues, not allowing for growth of the tumor into the CSF or past the boundaries of the brain.

We initiate the simulation with a small normoxic cell population that decreases spatially from a point with coordinates (x_0, y_0)

$$c(\mathbf{x}, 0) = 1000e^{-100R^2}, \quad (8)$$

where $R^2 = (x - x_0)^2 + (y - y_0)^2$. The initial seeding locations are described in Section 2.3 and can be seen for the first location in Figure 2.

The initial vascular cell densities are heterogeneous, set to 3% and 5% of the carrying capacity, K , in white and grey matter respectively; these values fall within the values for cerebral blood volume found from the literature [47]. All other spatio-

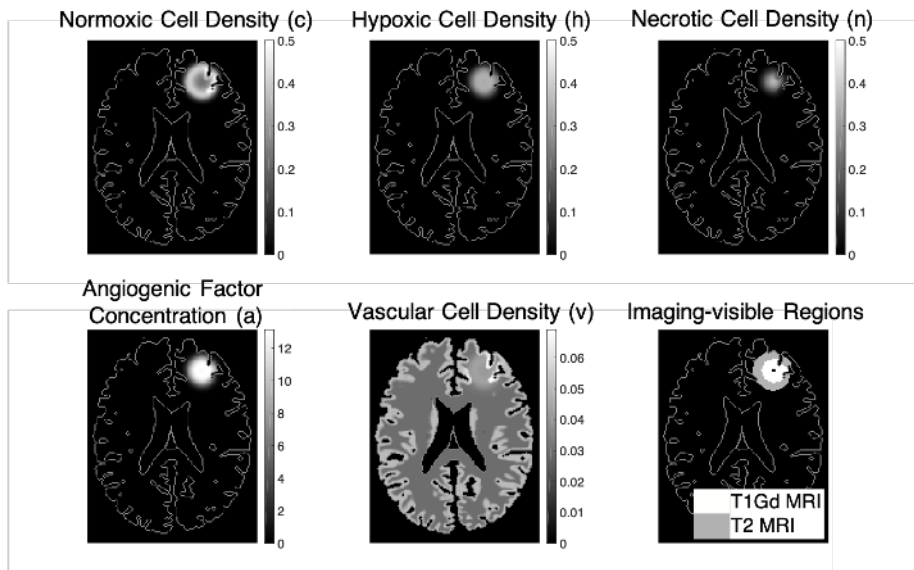


Fig. 2: An example simulation shown at a size equivalent to a circle of 1cm radius on simulated T1Gd MRI. We show all cell densities divided by K and the angiogenic factor concentration divided by K_M . We see how normoxic cells lead the outward growth of the simulated GBM, followed by hypoxic cells and necrotic cells. Angiogenic factors are mostly found in the hypoxic cell region. We also show the regions that are assumed visible on T1Gd MRI ($T \geq 0.8$) and T2 MRI ($T \geq 0.16$) as well as the point where the tumor is initiated (black pixel). In this simulation, $D_h/D_c = 10$, $D_c = 10^{1.5} \text{mm}^2/\text{year}$, $\rho = 100/\text{year}$, $\beta = 0.5\rho$ and $\gamma = 0.05/\text{day}$.

temporal variables are initially set to zero. There are no-flux boundary conditions on the outer boundary of the brain for all variables as well as on the CSF that do not allow growth outside of the brain or into CSF regions.

2.2 Nutrient-Based Vascular Efficiency

The extended vascular efficiency term uses a reaction-transport equation to model the nutrient consumption by the tumor cells. Using this reaction-transport equation for the movement and consumption of nutrient, f^1 , the derivation of the vascular efficiency term goes as follows:

$$\frac{\partial f}{\partial t} = \nabla \cdot (D_f \nabla f) + psv(f_{blood} - f) - \eta_c(D_c, \rho)cf - \eta_h(D_h)hf, \quad (9)$$

where p is the permeability of the blood brain barrier to nutrient, s is the vascular surface area per unit volume, f_{blood} is the concentration of nutrient in the blood

¹ We denote this as f to represent fuel for the cells, to avoid reusing n which is already assigned to necrotic cells.

which is assumed fixed², η_c is the rate of nutrient consumption by normoxic cells and η_h is the nutrient consumption rate by hypoxic cells. We have let η_c depend on the diffusion (D_c) and proliferation (ρ) rates of normoxic cells, as these processes require energy. A larger D_c and ρ will require more energy as the tumor cells migrate and proliferate relatively quickly. Similarly, we have set η_h to depend on the value of D_h , as faster migrating tumor cells require more energy and in turn more nutrient.

Now if we assume that in the timescale of interest, the nutrient concentration rapidly reaches steady state, and that the nutrient is consumed much faster than it diffuses, we can eliminate those terms to be left with

$$0 = 0 + psv(f_{blood} - f) - \eta_c(D_c, \rho)cf - \eta_h(D_h)hf \quad (10)$$

and rearrange to get

$$\frac{f}{f_{blood}} = \frac{v}{v + \frac{\eta_c(D_c, \rho)c + \eta_h(D_h)h}{ps}}. \quad (11)$$

We assign this expression as the vascular efficiency term, V , as it corresponds to the ability of the vasculature to provide nutrients to the tumor. This term is similar to that seen in the original formulation of the PIHNA model but now includes the nutrient consumption and extravasation of nutrients from the blood [43].

To estimate the parameters η_c , η_h and ps , we used Fludeoxyglucose (FDG) Positron Emission Tomography (PET) data from a paper by Delbeke *et al.* [14]. FDG is analogous to glucose and can be detected on PET scans. We have chosen glucose as an estimate for our generic nutrient due to the availability of imaging data that we could use to parametrise our vascular efficiency term. Due to the increase in anaerobic respiration of cancer cells compared with normal tissue, known as the Warburg effect [23,46], we might expect oxygen uptake ratios to be lower than glucose.

We note that to parametrize our nutrient-based vascular efficiency term, we only need to consider the ratio between $\eta_c : ps$ and $\eta_h : ps$. As both of these expressions are in the same units of $\text{mm}^3/\text{cell}/\text{year}$, their ratio is dimensionless. Delbeke presents the uptake ratios between tumor and healthy tissue within both white and grey matter. To make use of these values, we assume that in a homeostatic healthy brain, the rate of glucose being used by healthy tissue that is not vasculature is equal to the rate of glucose entering from the vasculature. We do not, however, model healthy tissue in the current formulation of the PIHNA model. For the benefit of this section, let us introduce unaffected healthy tissue u_0 , with glucose uptake rate η_u , we assume

$$psv_0 = \eta_u(u_0 - v_0), \quad (12)$$

where v_0 is the initial background vascular cell density in the PIHNA model, and u_0 is the healthy tissue density. We then have $ps = \eta_u(u_0/v_0 - 1)$, which will always be positive in PIHNA simulations as vasculature takes up a small percentage of

² It is well known that nutrient concentrations in blood (such as glucose concentration) fluctuates throughout a single day, however we are interested in modeling tumor growth over many days and months, so only consider the average nutrient concentration across these daily fluctuations.

brain volume compared to other tissue. We assume that in healthy white matter tissue there is 3% vasculature and in grey there is 5%, so we let $v_0/u_0 = 0.03$ in white matter and $v_0/u_0 = 0.05$ in grey matter; these values fall within realistic values for cerebral blood volume [47]. Now the ratios of glucose uptake rates by tumor to the glucose uptake rates by healthy tissue given by Delbeke can be considered as various values of $P_c = \eta_c/\eta_u$ and $P_h = \eta_h/\eta_u$ in the PIHNA model. So Equation 11 is now expressed as

$$V = \frac{v}{v + \frac{P_c \eta_u c + P_h \eta_u h}{\eta_u (u_0/v_0 - 1)}}, \quad (13)$$

and the η_u terms cancel to give

$$V = \frac{v}{v + \frac{P_c c + P_h h}{u_0/v_0 - 1}}. \quad (14)$$

We noted that in the work by Delbeke et al.[14] there was a spread of relative tumor uptake values for high grade gliomas within cortical and white matter tissue across 20 patients. As an approximation, we attributed these differences to the nutritional demands of the individual high grade gliomas. We assign normoxic cells with high (low) D_c and high (low) ρ in the PIHNA model with the higher (lower) glucose uptake rates from the literature, which also vary between white and grey matter. We assign hypoxic cells with high (low) D_h high (low) glucose uptake rates in the same manner as the normoxic cells. The values in between the extremes are assigned using a log linear scale, due to the large range of D_c , D_h and ρ values used in PIHNA simulations. The ratios P_c^w , P_c^g , P_h^w and P_h^g are then given by

$$P_c^{w,g}(D_c, \rho) = G_{max}^{w,g} - (G_{max}^{w,g} - G_{min}^{w,g}) \frac{\log_{10}\left(\frac{D_{c_{max}}}{D_c}\right) + \log_{10}\left(\frac{\rho_{max}}{\rho}\right)}{\log_{10}\left(\frac{D_{c_{max}}}{D_{c_{min}}}\right) + \log_{10}\left(\frac{\rho_{max}}{\rho_{min}}\right)} \quad (15)$$

and

$$P_h^{w,g}(D_h) = G_{max}^{w,g} - (G_{max}^{w,g} - G_{min}^{w,g}) \frac{\log_{10}(D_{h_{max}}/D_h)}{\log_{10}(D_{h_{max}}/D_{h_{min}})}, \quad (16)$$

where we use the extremes of tumor to normal tissue uptake ratios in white matter (taken as $G_{min}^w = 1.66$ and $G_{max}^w = 4.5$) and the extremes of observed uptake ratio in grey matter (taken as $G_{min}^g = 0.5$ and $G_{max}^g = 2$) as the minimum to maximum glucose uptake ratios G_{min} and G_{max} . The maximum and minimum D_c , D_h and ρ values are equal to the maximal and minimal rates that we run in our simulations, see Table 1. This along with the values of v_0/u_0 give the parametrisation of the nutrient-based vascular efficiency term.

2.3 Modeling Resection and Ischemia

Using the PIHNA model, we have simulated a resection that occurs once the tumor has grown to a shape with a volume equivalent to a disc of 1cm radius on simulated T1Gd imaging. The tumor was initiated at $x_0 = 100$, $y_0 = 150$ on the 85th axial slice of the brain geometry. Post resection, zero-flux boundary conditions are added around the simulated frontal lobectomy (green outline in Figure 3) so that

regrowth into the resection cavity is not possible. This boundary condition was chosen as recurrence into the resection cavity itself is unlikely in GBM. GBM cells have a migratory nature and need structure and nutrients to migrate and grow [3]. As there is a lack of tissue and vasculature within the resection cavity, it is not an environment conducive to GBM growth. Every resection is the same, in that the same region of brain geometry is removed, which removes all of the enhancing T1Gd region. To incorporate the potential reality that surgery could induce a nearby ischemic event (red outline in Figure 3), we add subsequent ischemia through a transient reduction in the vasculature term, v , to a region adjacent to the resection cavity wall. We have modeled ischemia as a reduction only at the time point of resection, the vessels then continue to follow the model equations. We reduced the vasculature to 1% of its value at the time of resection thus simulating a near complete ischemic event in the region noted in red in Figure 3, which is also the value used in all recurrence pattern figures presented in the main text. In further simulations, we have also reduced the vasculature to 10% to see the impact of this less intense simulated ischemia on recurrence locations; results of these simulations are presented in Appendix Figures 12 - 14.

We also present a second location with a recurrence that occurs when the tumor has reached a volume of a disc of 0.25cm radius on T1Gd imaging. In this setting, the tumor location was initiated at $x_0 = 44$, $y_0 = 132$ on the 89th coronal slice of the brain geometry. This resection cavity was simulated as a 10 pixel radius around the initial tumor location, with ischemia as a further 5 pixel radius. Zero flux boundary conditions are added to this resection cavity and perioperative ischemia is simulated in the surrounding tissue.

2.4 Virtual Experiments

We run simulations for different values of normoxic cell migration (D_c with range $1 - 1000\text{mm}^2/\text{year}$) and proliferation (ρ with range $10 - 100/\text{year}$), as well as test two values of β (0.1ρ and 0.5ρ), which is the switching rate from the normoxic cell density towards hypoxic cell density, three values of γ (0.005 , 0.05 , $0.5/\text{day}$), which is the switching rate back from hypoxic cell density to a normoxic cell density, and the rate of hypoxic to normoxic cell migration, D_h/D_c (1 , 10 or 100). We vary the ratio of hypoxic to normoxic cell migration due to evidence that GBM cells migrate faster in hypoxic conditions [19,49]. These parameters were chosen as they represent the tumor's response to hypoxic stress. In previous work, we have observed that all of these parameters (except for β) influence the outward growth rate of PIHNA simulations, which is another consideration of the effects of hypoxia on GBM [12]. We note that the varying tumor kinetics (migration rates D_c , D_h and proliferation rate ρ) affect the nodularity of the simulated tumors. Simulations with higher ratios of migration to proliferation will be more infiltrative tumors, whereas those with higher ratios of proliferation to migration will be denser tumor masses with less infiltration and more well-defined tumor cell density boundaries. Examples of this effect can be seen in Figure 3. We ran further simulations of a second tumor location with $\beta = 0.5\rho$, $\gamma = 0.5/\text{day}$ and $D_c = 10D_h$, as a validation that the observed dependency of recurrence patterns on migration and proliferation rates were not simply a function of the resection and ischemic regions that we chose in the first location.

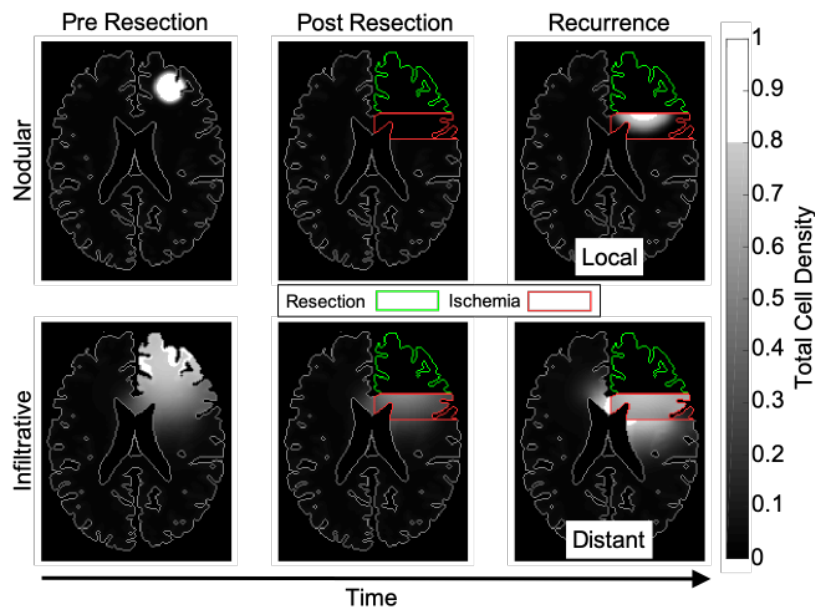


Fig. 3: The tumor undergoes resection that removes the T1Gd imageable tumor cell density at 1cm radius (assumed at 80% of the maximum cell density and shown in white) as well as the surrounding tissue. In these two examples, the nodular tumor (top row) recurs locally, whereas the infiltrative tumor (bottom row) recurs distantly. In these simulations, $\beta = 0.5\rho$, $\gamma = 0.05/\text{day}$ and $D_h = 10D_c$. For the nodular tumor, $D_c = 10^{0.5}\text{mm}^2/\text{year}$ and $\rho = 10^{1.5}/\text{year}$. For the infiltrative tumor, $D_c = 100\text{mm}^2/\text{year}$ and $\rho = 10/\text{year}$.

2.5 Defining Recurrence Location

Recurrence location of a tumor is classified as the reappearance of the tumor on T1Gd MR scans as is done clinically [50]. If a tumor initially reappears outside of the simulated ischemic region above a certain thresholded size (a disc of radius 2mm on simulated T1Gd MRI) before appearing anywhere else, it is classified as appearing anywhere else, it is classified as distant. Whereas if it appears within the ischemic region along the cavity wall above the same threshold before anywhere else, it is classed as a local recurrence. Examples of these cases can be seen in Figure 3. We do not consider the nature of T2 signal in our definition of distant recurrence. We define a mixed recurrence when the tumor appears on simulated T1Gd MRI both inside and outside the ischemic region before the size threshold within either region is reached. This method of defining distant, local and mixed recurrence patterns is applied to both tumor locations presented in this work.

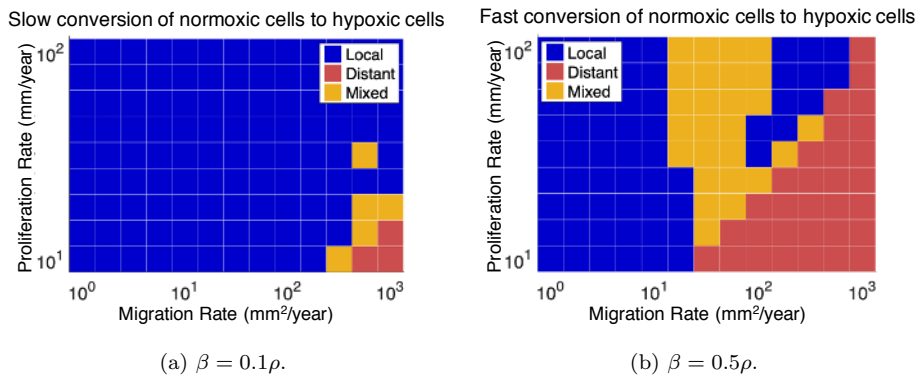


Fig. 4: Recurrence location classified for various D_c , ρ and β for $D_h = 10D_c$ and $\gamma = 0.05/\text{day}$. We see that higher values of β (the conversion rate from normoxic to hypoxic cells) lead to a larger proportion of distant recurrences in D_c and ρ parameter space. Higher migration rates, D_c , and lower proliferation rates, ρ , lead to more distantly recurring simulated tumors.

3 Results

3.1 Individual tumor kinetics affect recurrence location following perioperative ischemia

Extending on the paper by Thiebold *et al.*, that suggests distant recurrence can occur through ischemia and subsequent hypoxia [45], the PIHNA model suggests that tumor kinetics also play a role. Figure 3 shows two simulated tumors, one nodular and the other infiltrative, that go through resection and subsequently recur. The recurrence pattern for the nodular tumor is local, whereas the infiltrative tumor recurs distantly. Such distantly recurring tumors remain in lower cell densities within the ischemic region and appear outside on simulated T1Gd imaging as they continue to increase their cell density outside of the ischemic region. The only differences between the two simulations presented in Figure 3 are the migration and proliferation rates of the tumor cells.

3.2 Tumor response to hypoxic conditions affects recurrence location

By varying individual tumor kinetics (D_c and ρ with $D_h = 10D_c$) and the maximal rate at which tumor cells become hypoxic and in turn necrotic (β), with a fixed vascular ischemia post resection, we are able to show differing tumor recurrence locations, see Figure 4. We also varied the maximal rate at which hypoxic tumor cells returned to a normoxic state, γ . Changing this parameter also had an effect on the recurrence patterns, see Figure 5. A low level of γ promotes more distant recurrence, while a high level of γ promotes local recurrence. Recurrence location is classified as the first reappearance of a tumor on T1Gd MR imaging as described in the previous section.

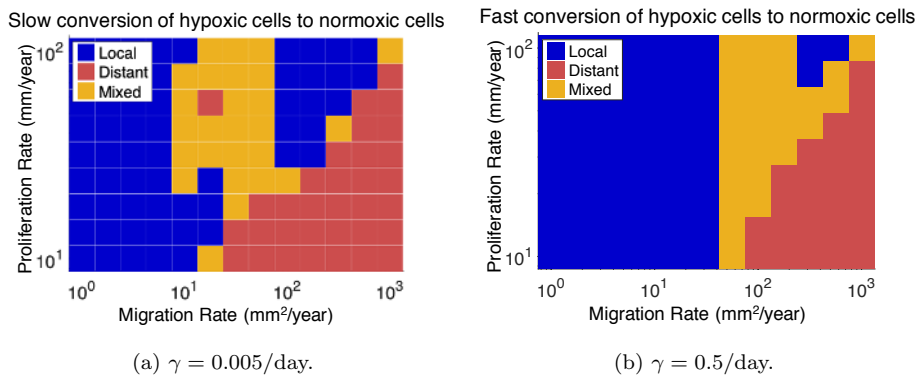


Fig. 5: Recurrence location classified for various D_c , ρ and γ for $D_h = 10D_c$ and $\beta = 0.5\rho$. We see the higher (lower) values of γ lead to a lower (higher) proportion of distant recurrences in D_c and ρ parameter space. Higher migration rates, D_c , and lower proliferation rates, ρ , lead to more distantly recurring simulated tumors.

An increase in β leads to more sensitivity in the tumors to ischemia, which causes them to become more hypoxic and therefore less proliferative within the ischemic region. They are more likely to become more dense, and therefore imageable on simulated T1Gd MRI, outside of the ischemic region and be seen as a distant recurrence. Conversely, an increase in γ , the conversion rate from a hypoxic cell phenotype back to normoxic, hinders this effect as it limits the impact of hypoxia on the growth of the simulated tumor. We present other simulation results of varying β and γ in Appendix Figure 10.

3.3 Faster Hypoxic Cell Migration Rates Promote Distant Recurrence

Following this initial analysis, we also varied the hypoxic diffusion rate relative to the normoxic counterpart, D_h/D_c . Along with the simulations where $D_h = 10D_c$ described in the previous section, we have set $D_h/D_c = 1$ and $D_h/D_c = 100$ (see Figure 6). We see that the higher the D_h/D_c value, the more distantly recurring tumors occur for fixed values of β and γ . The effect of an increase in D_h/D_c is more pronounced for tumors that are more sensitive to the hypoxic environment caused by the ischemia, see Appendix Figures 9 - 11. In previous work, we have shown that an increase in D_h/D_c increases the outward growth rate of PIHNA-simulated GBM [12]. With faster hypoxic migration rates, the simulated tumor cell densities are able to travel through the hypoxic region faster. These tumors can then reach the region of the brain slice unaffected by ischemia and develop into a dense tumor mass before the tumor develops within the ischemic region.

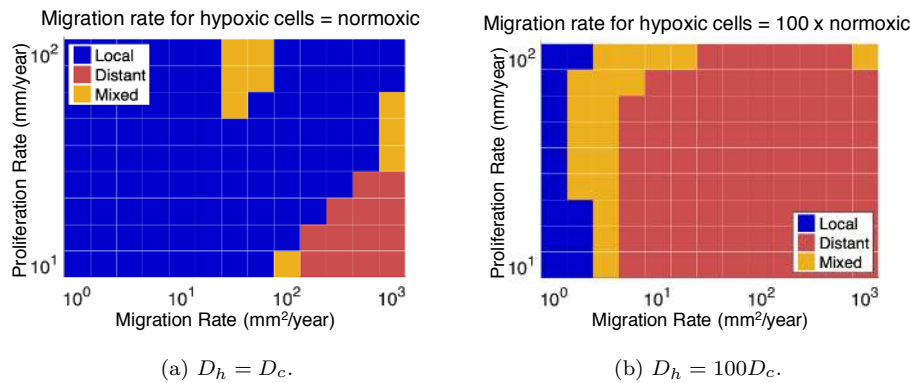


Fig. 6: Recurrence location classified for various D_c , ρ and D_h/D_c for $\beta = 0.5\rho$. We see that the higher (lower) level D_h/D_c lead to a larger (smaller) proportion of distant recurrences in D_c and ρ parameter space. Higher migration rates, D_c , and lower proliferation rates, ρ , lead to more distantly recurring simulated tumors.

3.4 Diffuse Recurrence Present Through an Ischemia-Induced Reduction in Dense Tumor

The paper by Thiebold considered distal recurrence as either diffuse or distant [45]. Diffuse recurrence presents as a marked increase in T2 signal without an accompanying increase in T1Gd enhancement. To explore how diffuse recurrence may occur in the PIHNA model, we present a plot of T1Gd and T2 volume over time for a simulated case of perioperative ischemia, compared with the same simulation without such ischemia. This case recurred distantly and is an example of parameters used in Subfigure 5a, with $D_c = 10^{2.5} \text{mm}^2/\text{year}$, $\rho = 10^{1.25}/\text{year}$, $D_h = 10D_c$, $\beta = 0.5\rho$ and $\gamma = 0.005/\text{day}$. As can be seen in Figure 7, we see a reduction in T1Gd signal as a result of the perioperative ischemia for a period of growth, while T2 signal is less affected. If the tumor with perioperative ischemia was observed in this window, the relative T2 signal compared with T1Gd enhancement would appear larger, resulting in a higher probability of the tumor being considered diffuse at recurrence. After some further growth, the T1Gd radii of both simulations meet, as the tumor within the ischemic region eventually grows to a high-enough density to be visible on T1Gd MRI.

3.5 Extent of Vasculature Reduction Impacts Recurrence Patterns

PIHNA model simulations were run with a lower level of reduction in functional vasculature to 10% within the same geometry of perioperative ischemia shown in Figure 3. This was implemented for all changes in β , γ and D_h presented in the main text. In this setting, all recurrence location results in Figures 4 - 6 showed a shift towards more local recurrence within the values of D_c and ρ that were tested. We present these corresponding recurrence location figures in the Appendix (Figures 12 - 14).

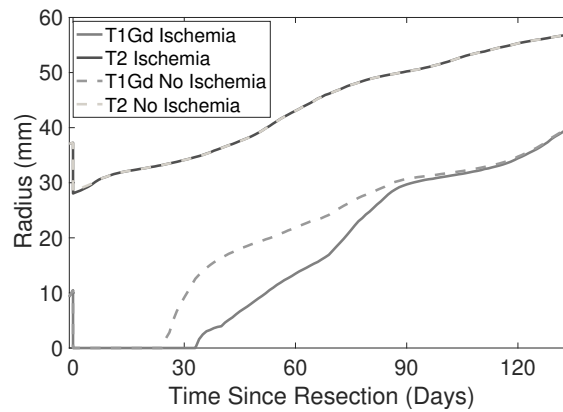


Fig. 7: We present the imageable tumor radii as a function of time for two simulations. The only difference between these simulations is that one has perioperative ischemia and the other does not. Note that the growth of the T1Gd imageable radius is delayed by the ischemia, whereas the T2 imageable radius is minimally affected. This offset of dense tumor growth may be a contributing factor to diffuse recurrence following perioperative ischemia. In these simulations, $D_c = 10^{2.5} \text{mm}^2/\text{year}$, $\rho = 10^{1.25}/\text{year}$, $D_h = 10D_c$, $\beta = 0.5\rho$ and $\gamma = 0.005/\text{day}$.

3.6 Validation in a Second Tumor Location

To explore the simulated recurrence location further, we ran PIHNA simulations for a second tumor location in a coronal view. This tumor was initiated closer to the surface of the brain and resected at a smaller T1Gd imageable tumor volume (equivalent to a disc of 0.25cm radius). We present recurrence locations and example simulations for $\beta = 0.5\rho$, $\gamma = 0.5/\text{day}$ and $D_h = 10D_c$ with varying migration and proliferation rates in Figure 8. Other than the different location and size at resection, these simulations have the same parameters as Subfigure 5b. As we observed in the first location, tumors with more diffuse characteristics recurred distantly, whereas those with more nodular characteristics and lower migration rates tended to recur locally or mixed. In this second location, we observe some simulated tumors with faster proliferation rates but mid-range migration rates that also recur distantly, which we also saw some evidence of in the first location, present in Subfigures 4b, 5a and 6a.

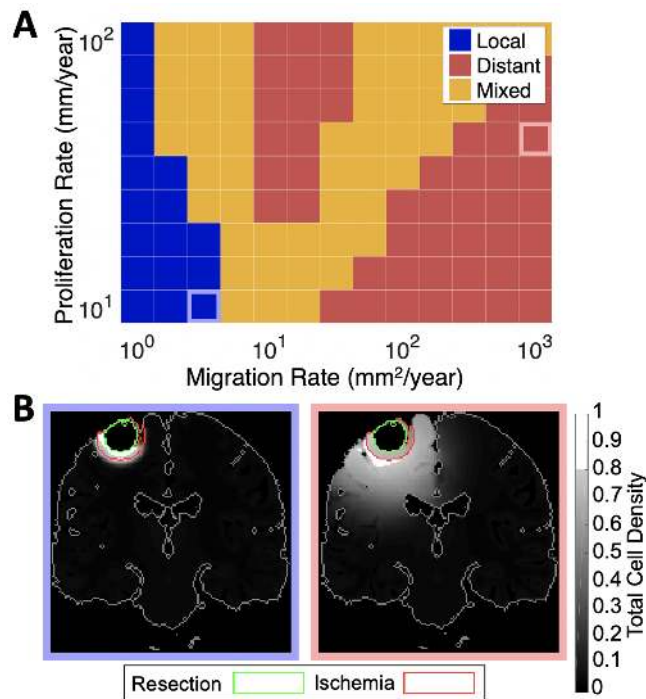


Fig. 8: A) Recurrence location classified for the second tumor location in coronal view, for various D_c and ρ values. As in the first location, tumors with more diffuse characteristics recur distantly, while those with lower migration rates and faster proliferation rates tended to be more mixed. In these simulations, $\beta = 0.5\rho$, $\gamma = 0.5/\text{day}$ and $D_h = 10D_c$, parameter values are equivalent to Subfigure 5b. B) Two example simulations, with local recurrence (left) and distant recurrence (right), showing the resection (inner green outline) and ischemic (outer red outline) regions. Migration and proliferation rates of these example simulations are indicated on Subfigure A.

4 Discussion

Through mathematical modeling, we have found a possible mechanism for distal GBM recurrence in response to ischemia. If the tumor has an invasive phenotype, it can remain unimageable on simulated T1Gd MRI as it travels through the ischemic region (using our assumed threshold of 80% total cell density). Once it reaches healthy intact vasculature, it will return to a normoxic phenotype and proliferate to an imageable density outside of the ischemic region before it does so next to the cavity wall. We see that the switching rate from normoxic cells to hypoxic cells plays a role in this behavior, increasing this rate leads to more distantly recurring tumors within the parameter range of D_c and ρ that we have used (Figure 4). Conversely, increasing the recovery rate from a hypoxic cell phenotype to a normoxic cell phenotype leads to less distantly recurring tumors (Figure 5).

We also note that an increase in the rate of hypoxic cell migration relative to normoxic cell migration promotes distantly recurring tumors (Figure 6).

The migratory nature of GBM cells is a key limitation of conventional treatment efficacy and contributes to tumor recurrence [34]. The dependency of distant recurrence on cell migration shown in these simulations suggests that the use of anti-migratory drugs may reduce the cases of distal recurrence, especially in instances of perioperative ischemia. However, this result is purely theoretical at this stage. These results may also be suggestive of tumor response to hypoxic conditions more generally. Future work may explore patient data to compare pre-operative infiltration patterns with distance to recurrence.

We have shown the intensity of the ischemia plays a role in the observed simulated recurrence patterns (see Appendix Figures 12 - 14). A reduction in functional vasculature to 10% of its pre-resection value does not promote distant recurrence for as many values of D_c and ρ as the lower value of 1%. As the value of 10% does not promote quite as much hypoxia in the ischemic region, more simulations are able to reach a T1Gd imageable density inside this region and recur locally.

Furthermore, we have shown that similar simulated recurrence patterns occur in a second tumor location, that was located closer to the surface of the brain than the first and in a more functionally-important region. We also simulated the resection of this second location at a smaller size, yet saw similar patterns of recurrence as the first location. This suggests that the observed model behavior is not simply a function of tumor location or the geometries that we chose for the first resection and ischemic regions. We also ran this simulation in a coronal view of the brain to highlight that this recurrence behavior can occur in any plane. In the future, we can bypass this by moving the model to a more realistic 3D space.

Simulated T1Gd MRI volumes are inhibited within the ischemic region, which may explain why diffuse as well as distant recurrences are observed in patients with perioperative ischemia. If the hypoxic cell phenotype were maintained following exposure to ischemia, the tumor as a whole could remain more diffuse in a clinical sense of a large T2 volume relative to T1Gd. Utilizing the PIHNA model may be a useful tool in our effort to understand patterns of recurrence in GBM and understanding the role of ischemia in recurrence and growth patterns more broadly.

Acknowledgements

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A Recurrence Results of Other PIHNA Simulations

We present the results of PIHNA simulations that were not shown in the main text. The trends in distant recurrence patterns that we observe in the main text all hold in these simulations, supporting our observations regarding D_h/D_c , β , γ , D_c and ρ .

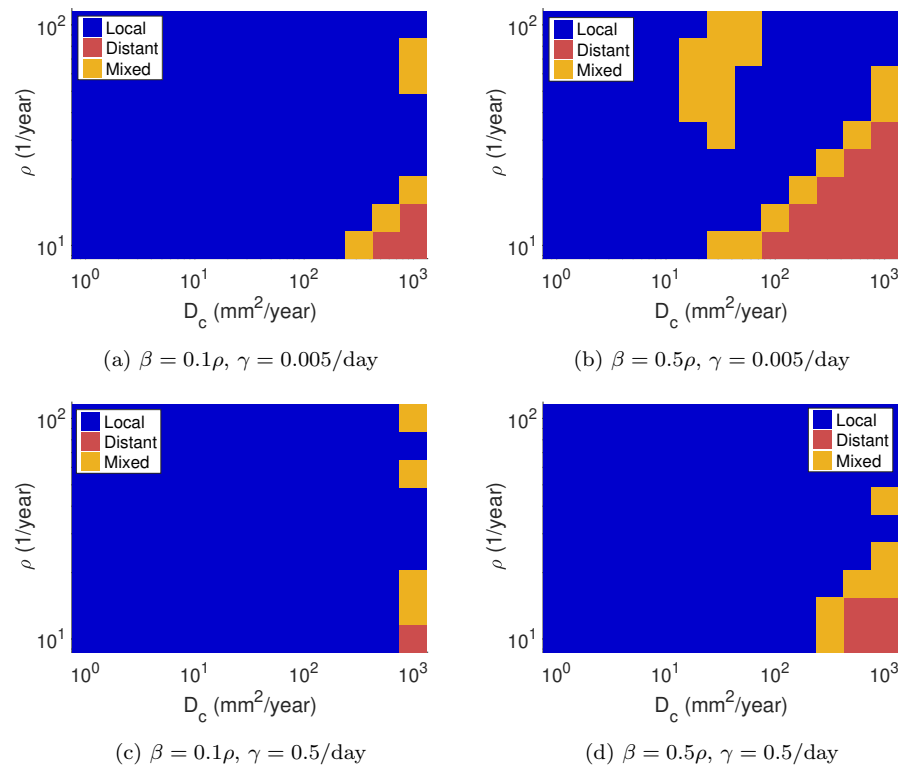


Fig. 9: Recurrence location classified for various D_c , ρ , β and levels of ischemia for $D_h = D_c$ for $\gamma = 0.005/\text{day}$ and $\gamma = 0.5/\text{day}$. We see that higher values of β and lower levels of γ lead to a larger proportion of distant recurrences in D_c and ρ parameter space. Higher migration rates, D_c , and lower proliferation rates, ρ , lead to more distantly recurring simulated tumors.

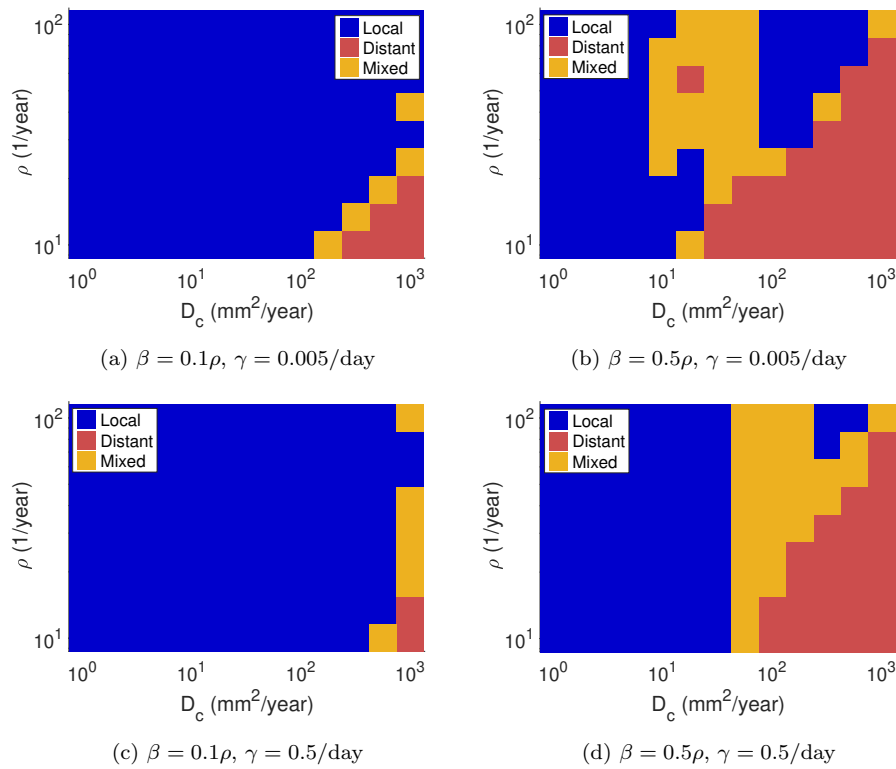


Fig. 10: Recurrence location classified for various D_c , ρ , β and levels of ischemia for $D_h = 10D_c$ for $\gamma = 0.005/\text{day}$ and $\gamma = 0.5/\text{day}$. We see that higher values of β and lower levels of γ lead to a larger proportion of distant recurrences in D_c and ρ parameter space. Higher migration rates, D_c , and lower proliferation rates, ρ , lead to more distantly recurring simulated tumors.

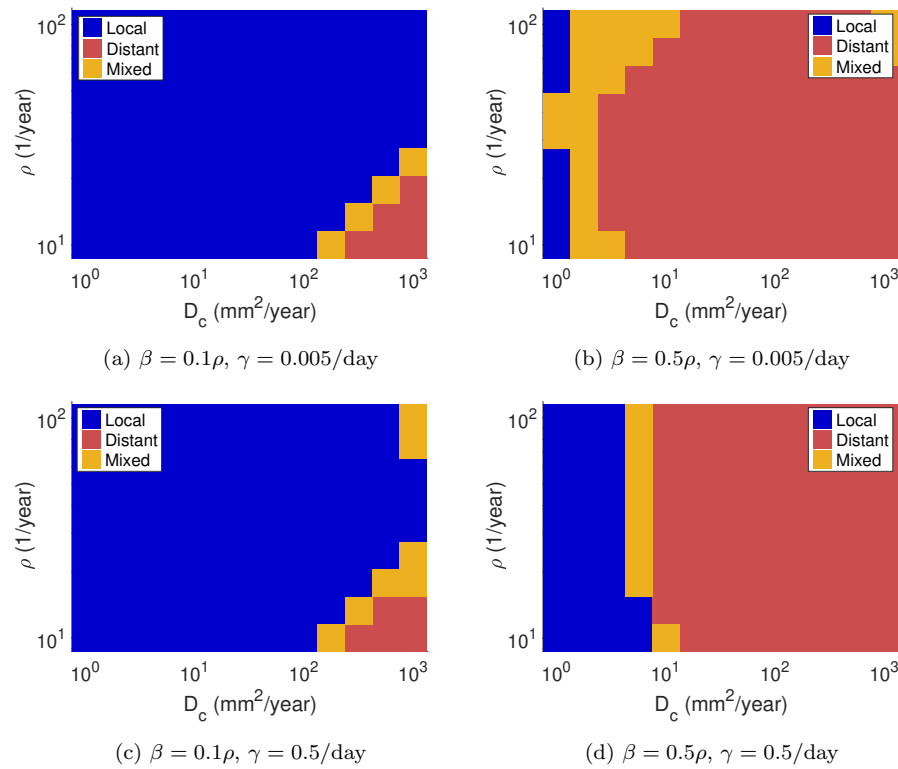


Fig. 11: Recurrence location classified for various D_c , ρ , β and levels of ischemia for $D_h = 100D_c$ for $\gamma = 0.005/\text{day}$ and $\gamma = 0.5/\text{day}$. We see that higher values of β and lower levels of γ lead to a larger proportion of distant recurrences in D_c and ρ parameter space. Higher migration rates, D_c , and lower proliferation rates, ρ , lead to more distantly recurring simulated tumors.

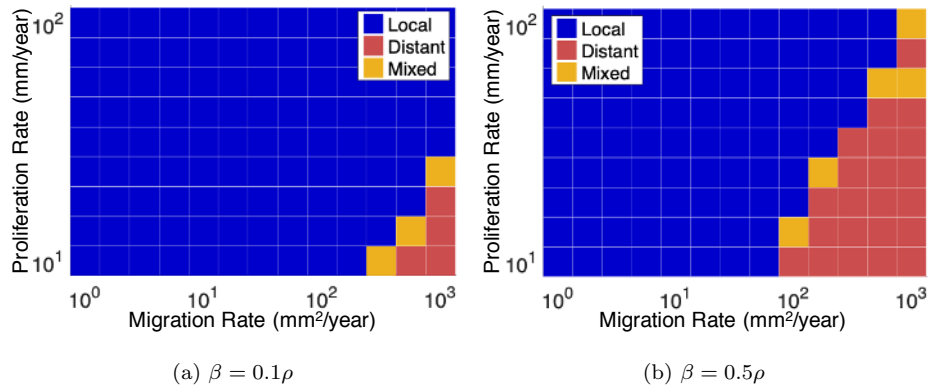


Fig. 12: Recurrence location classified for various D_c , ρ , β and levels of ischemia for $D_h = 10D_c$ for $\gamma = 0.05$ /day. In these simulations, perioperative ischemia was set at 10% of the pre-resection value. We see a larger proportion of local recurrence in these figures compared with those in the main text (see Figure 4).

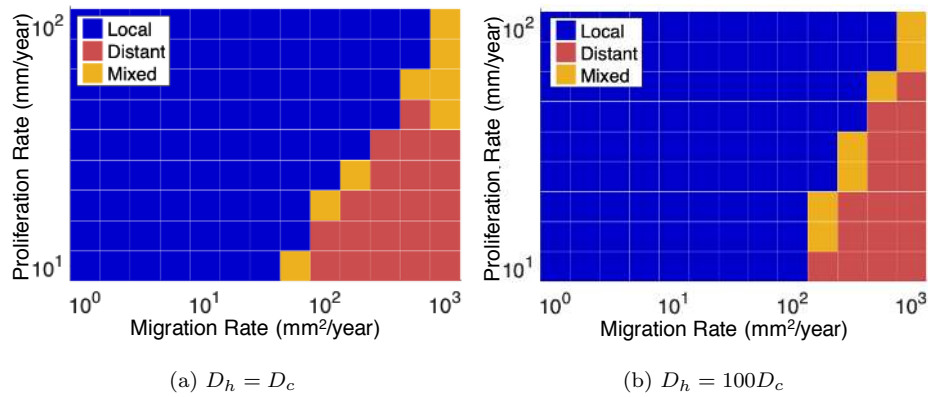


Fig. 13: Recurrence location classified for various D_c , ρ , β and levels of ischemia for $D_h = 10D_c$ for $\gamma = 0.005$ /day. In these simulations, perioperative ischemia was set at 10% of the pre-resection value. We see a larger proportion of local recurrence in these figures compared with those in the main text (see Figure 5).

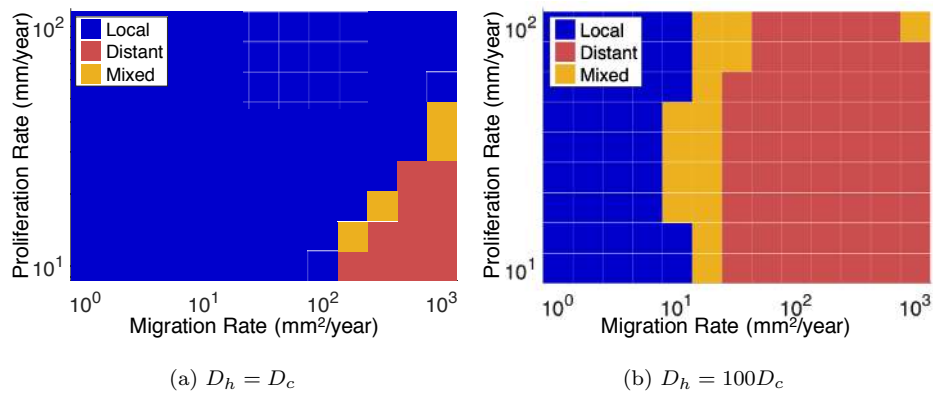


Fig. 14: Recurrence location classified for various D_c , ρ and D_h levels for $\beta = 0.5\rho$ and $\gamma = 0.05/\text{day}$. In these simulations, perioperative ischemia was set at 10% of the pre-resection value. We see a larger proportion of local recurrence in these figures compared with those in the main text (see Figure 6).