

Table S1. Parameters and their values used in the model.

Parameter	Meaning	Biological Value	Normalized Value	Reference
Neoplasm Size (P)	The dimensions (in patches) of the square grid representing the neoplasm.	2mm×2mm	64×64 patches (~30 μm ×30μm/patch)	(1)
Neoplasm Cell Density (n_0)	The cell density in the neoplasm.	$3.0 \times 10^3/\text{mm}^2$	3 per patch	(2)
Initial Neoplasm Cell Number (C)	The initial number of cells at the start of the model	(derived from P and n_0)	~12,000 cells	(1, 2)
Vessel Number (V)	The total number of blood vessels.	7.5 - 150/ mm^2	30 – 600	(2)
Oxygen Input Rate (r_i)	The amount of oxygen input into the neoplasm each time step.	2.7×10^{-14} mol/ mm^2/s (estimated based on n_0, r_m)	60 units per patch per time step	Estimated
Oxygen Diffusion Coefficient (d_c)	The percent of total oxygen in a patch distributed to its eight neighbor patches in each iteration of diffusion function.	1.8×10^{-5} cm^2/s	50% (Calibrated)	(3)
Oxygen update time (t_a)	The number of time discretization per time step.	N/A, for approximating continuous behavior	10 per time step	Model specific
Hypoxia Threshold (t_h)	The amount of oxygen in a patch below which a cell signals for angiogenesis.	0.5 – 30% of the concentration in surrounding tissue	8 units	(4)
Vessel Occlusion Threshold (t_o)	The number of cells in the same patch as the blood vessel that occludes the blood vessel.	N/A, for achieving vessel turnover rate of 5 days	20 cells	Model specific
Fixed Vessel Lifespan (t_f)	The number of time steps that a blood vessel persists before it is removed from the model	48h – 250 days	4 – 500 time steps	(5)
Cell Oxygen Absorption Rate (r_a)	The maximum amount of oxygen one cell can absorb in a time step.	4.5×10^{-17} mol/cell/s (exponential state of growth)	100 units	(6)
Cell Metabolism Rate (r_m)	The amount of resources that is metabolized in one cell per time step.	8.9×10^{-18} mol/cell/s (plateau state of growth)	20 units	(6)
Cell Reproduction Threshold (n_r)	The amount of resource units that must be accumulated before a cell can divide.	N/A, for achieving tumor cell cycle of 1.5 day	240 units	(7)
Initial Migration Propensity (p)	The initial migration probability given to all cells at the start of the model.	5% of the cell population	5%	(8)
Initial Maximum Migration Distance (m)	The initial maximum distance that a cell can migrate given to all cells at the start of the model.	3 μm/h	1 patch per time step	(9)
Mutation Rate (μ)	The probability that a phenotype will change in a daughter cell when the parental cell divides	0.01 per cell division	0.01	(10)
Migration Propensity Mutation Standard Deviation (sd_p)	The standard deviation of the normal distribution from which the mutated migration propensity value is drawn.	NA	(0.01,0.025,0.1)	Parameter Sweep
Maximum Migration Distance Mutation Standard Deviation (sd_m)	The standard deviation of the normal distribution from which the mutated maximum migration distance value is drawn.	NA	(0.1,0.5,1)	Parameter Sweep

We adjusted the NetLogo diffusion coefficient and oxygen update time to mimic the oxygen gradient observed in normal tissues. Specifically, we modeled a slice of tissue of 8 by 8 patches with one microvessel in the center (~ 120 μm radius). We turned off the cell death, reproduction and movement. Then we adjusted the oxygen update time and oxygen diffusion coefficient so that the oxygen gradient is similar to the observation in (11). This also agrees well with the fact the diffusion distance is between 100 μm and 200 μm (12).

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