

Mathematical modeling in support of drug discovery

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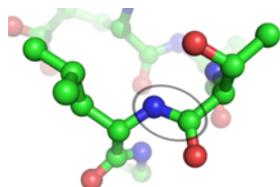
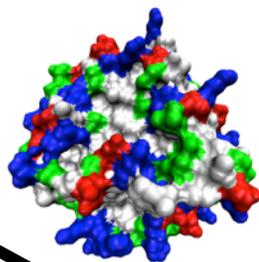
Moffitt—IMO, 15th June 2016

Drug development process

Identify disease



Identify therapeutic target (2-5 years)



Find compound effective against target (2-5 years)



Preclinical testing (1-3 years)



Human clinical testing (2-10 years)



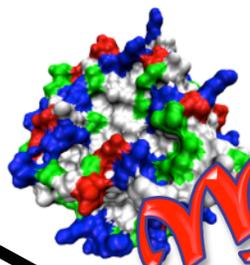
FDA approval

Drug development process

Identify disease

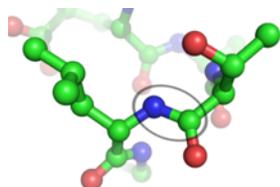


Identify therapeutic target (2-5 years)



Math

Math



Find compound effective against target (2-5 years)

Math



Preclinical testing (1-3 years)

Math



Human clinical testing (2-10 years)

Math



FDA approval

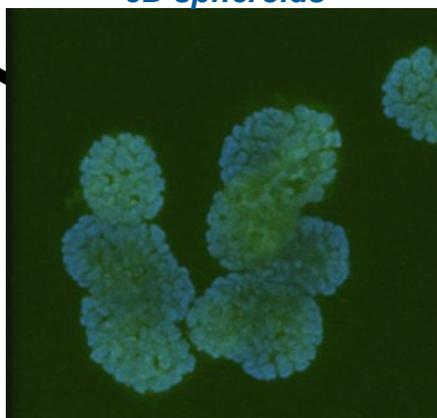
Pre-clinical tests



Petri dish



3D spheroids



mouse

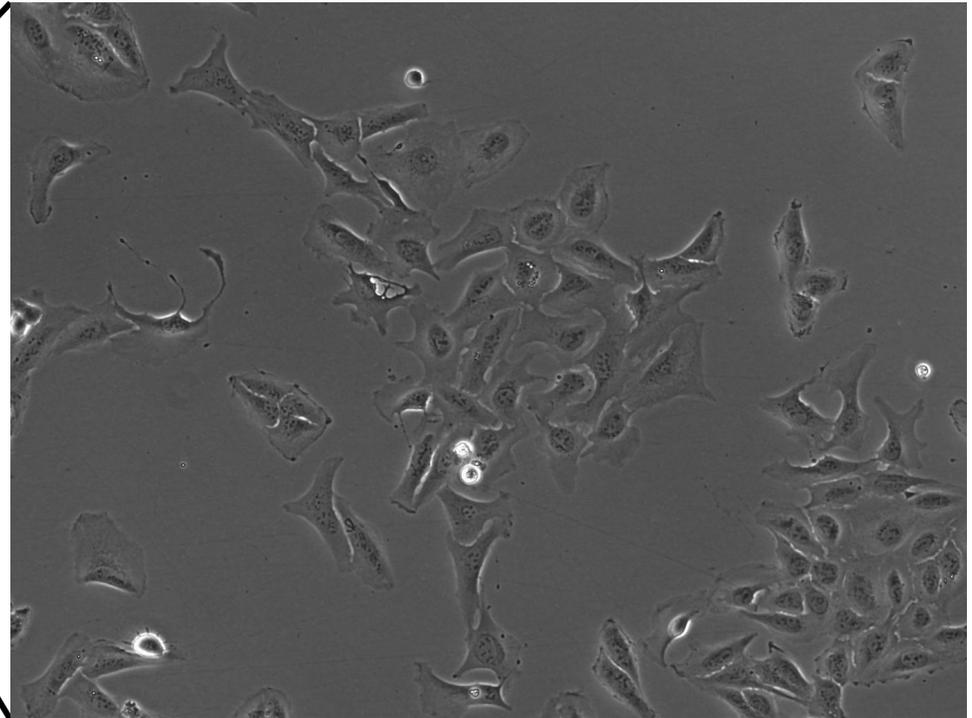


Cell colony experiments

In cancer biology new anticancer treatments are first tested in the cell culture.



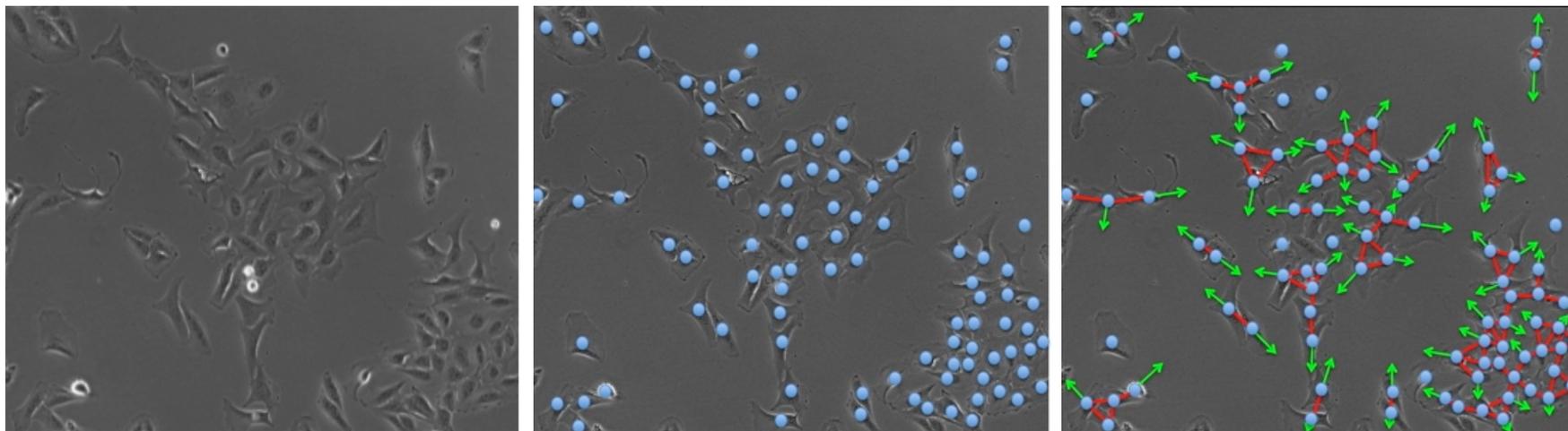
Petri dish



Part of the Petri dish seen under the microscope

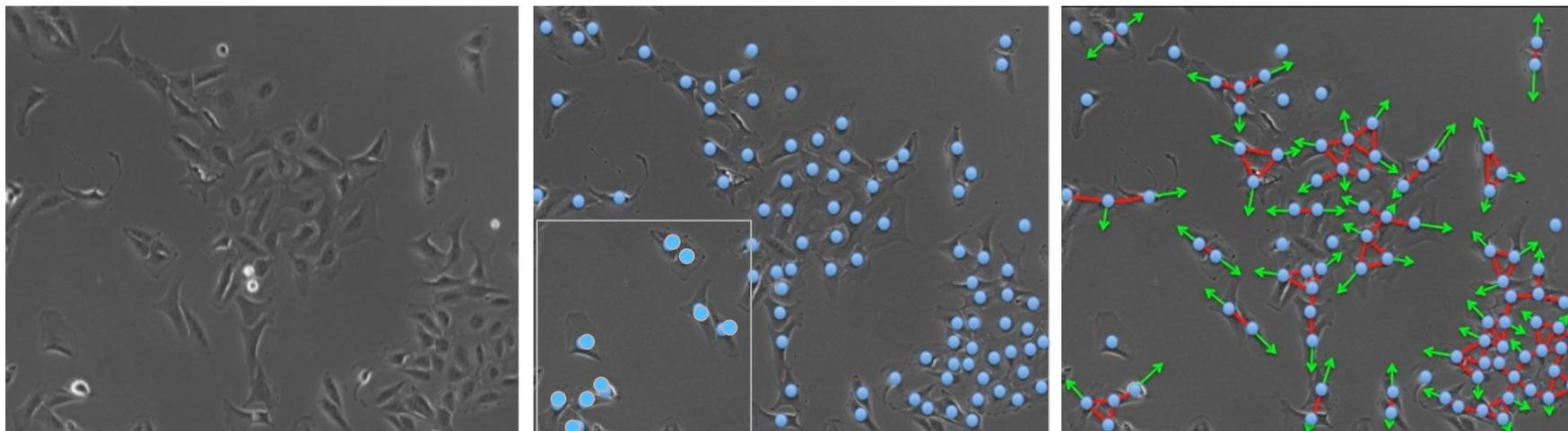
The cells grow in a two-dimensional dish in a medium that contains nutrients. Initially, the cells are seeded sparsely, but due to their divisions and movement the cell colony grows and spreads in the available space. Chemotherapeutic drugs are added to the medium in known concentrations. The total toxicity in the medium has to be controlled. The number of viable cells can be counted.

How to model these cells?

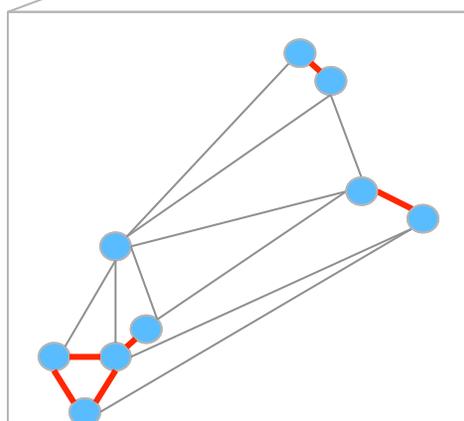


- Cell nuclei
- Spring connections
- Repulsive forces

Find cells that are near-by



- Cell nuclei
- Spring connections
- Repulsive forces



Neighborhood relation

Goal: the cells that are too close to one another should be pushed away on a distance equal to cell diameter.

Let's do some math!

Newton law

$$F = m \cdot a =$$

$$= m \cdot \frac{v}{t} =$$

$$= \frac{m}{t} \cdot \frac{\text{displacement}}{t}$$

1) coordinates of points
 $A = (2, 1)$ $B = (6, 1)$ $C = (2, 4)$

2) distance between 2 points
 in general $\mathbf{x} = (x, y)$
 $\|A - B\| = 4$ $\|A - C\| = 3$ $\|C - B\| = 5$

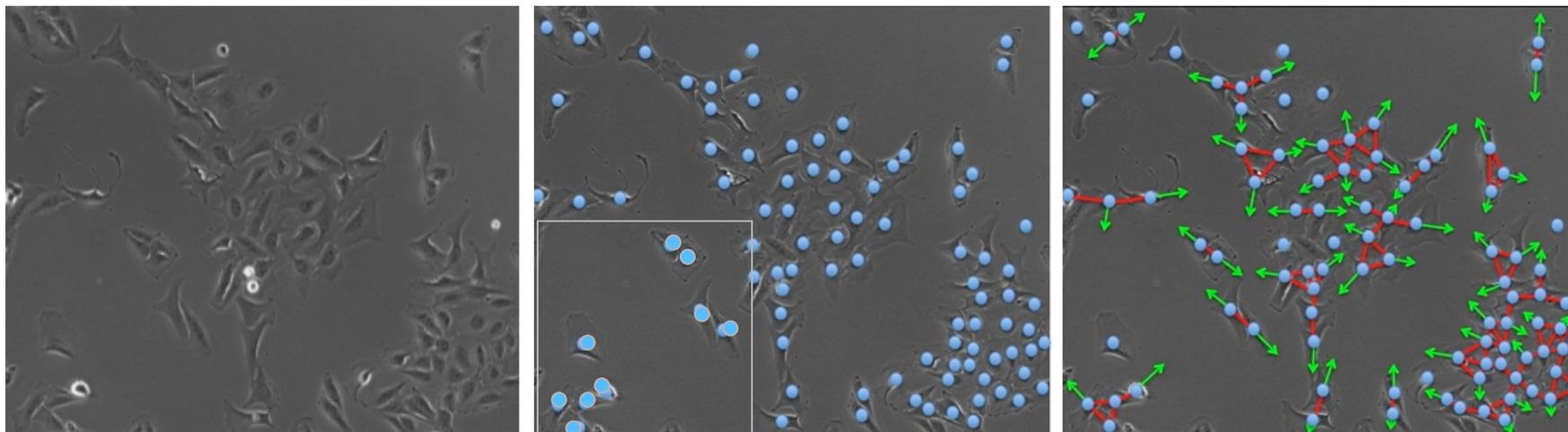
Pythagorean theorem: $a^2 + b^2 = c^2$
 in general: let $\mathbf{x}_i = (x_i, y_i)$ $\mathbf{x}_j = (x_j, y_j)$

Spring stiffness unit vector
 $\frac{\mathbf{x}_i - \mathbf{x}_j}{\|\mathbf{x}_i - \mathbf{x}_j\|}$

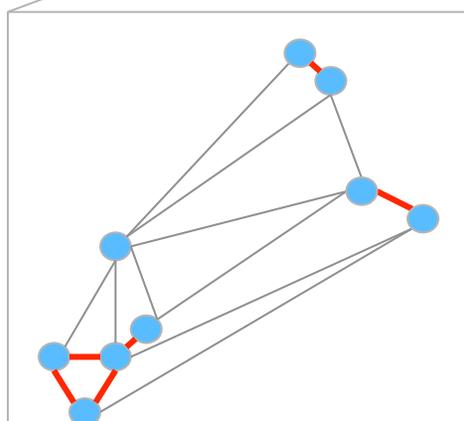
$$\|\mathbf{x}_i - \mathbf{x}_j\| = \sqrt{(x_i - x_j)^2 + (y_i - y_j)^2}$$

3) repulsive forces

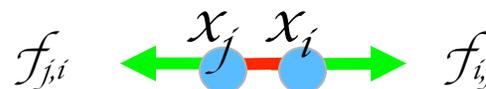
Exert force for too-close cells



- Cell nuclei
- Spring connections
- Repulsive forces



Neighborhood relation



$$f_{i,j} = \begin{cases} S (d - \|\mathbf{x}_i - \mathbf{x}_j\|) \frac{\mathbf{x}_i - \mathbf{x}_j}{\|\mathbf{x}_i - \mathbf{x}_j\|}, & \text{if } \|\mathbf{x}_i - \mathbf{x}_j\| < d, \\ 0, & \text{otherwise.} \end{cases}$$

Forces from several neighboring cells:

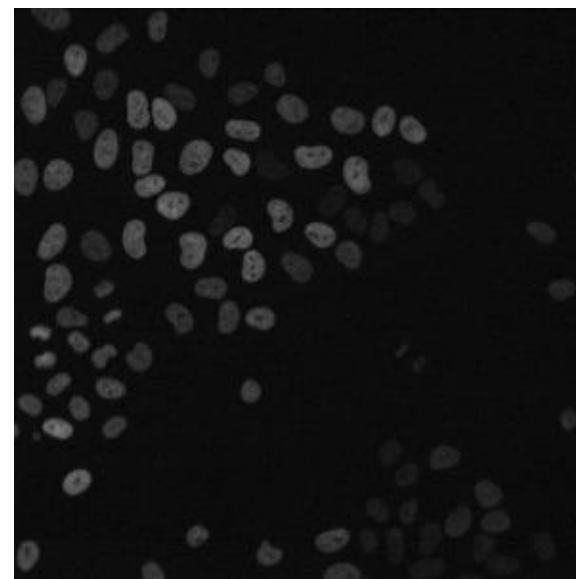
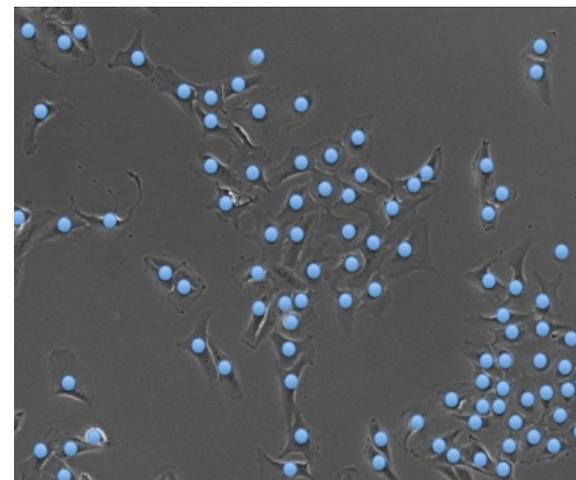
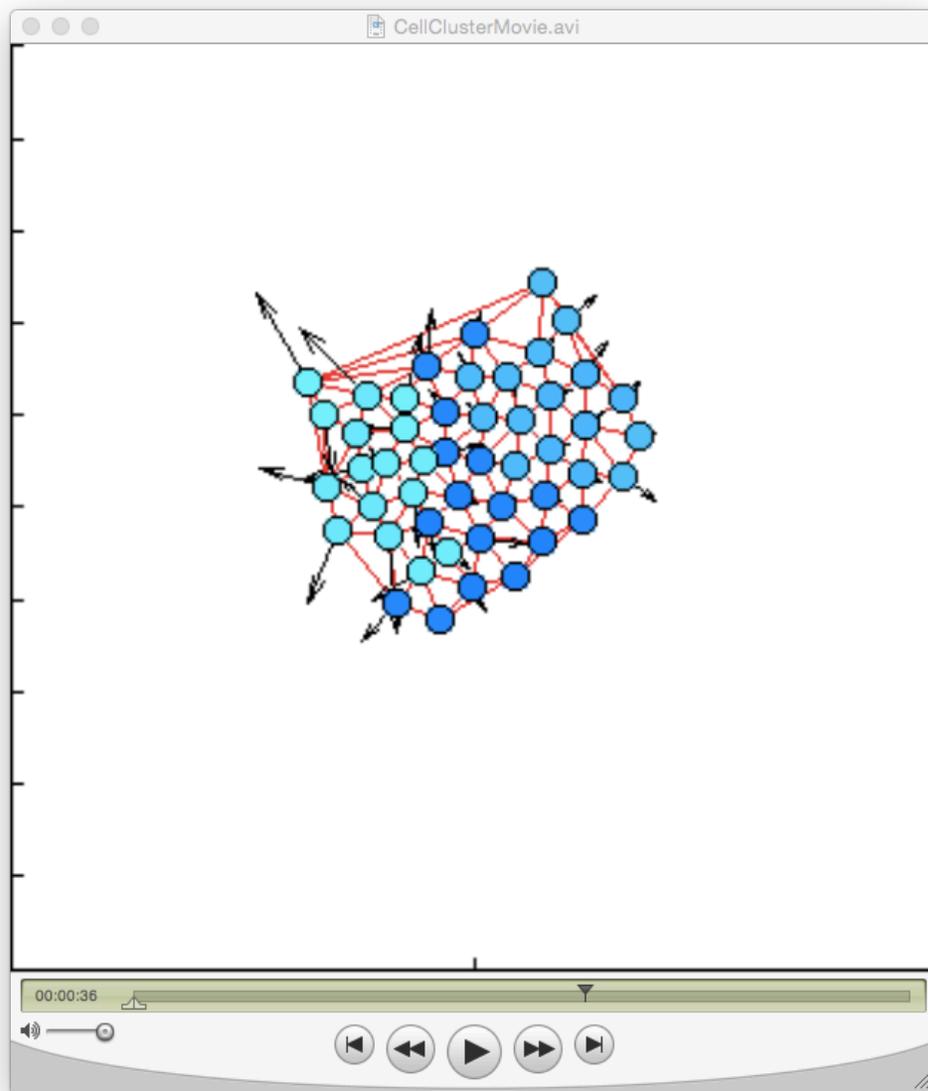
$$\begin{aligned} F_i &= f_{i,j_1} + \dots + f_{i,j_M} = \\ &= S (d - \|\mathbf{x}_i - \mathbf{x}_{j_1}\|) \frac{\mathbf{x}_i - \mathbf{x}_{j_1}}{\|\mathbf{x}_i - \mathbf{x}_{j_1}\|} + \dots + S (d - \|\mathbf{x}_i - \mathbf{x}_{j_M}\|) \frac{\mathbf{x}_i - \mathbf{x}_{j_M}}{\|\mathbf{x}_i - \mathbf{x}_{j_M}\|}. \end{aligned}$$

Change in cell position:

$$\mathbf{x}_i(t + \Delta t) = \mathbf{x}_i(t) - \frac{1}{\nu} \Delta t F_i$$

d -resting length (cell diameter),
 S -spring stiffness
 ν -media viscosity

How this model works?

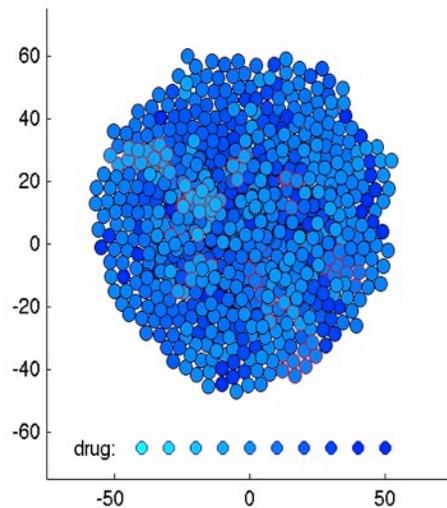


Let's do some experiments

Question:

What is the best **drug schedule** that will allow us to **suppress growth** of the whole tumor colony, but will use the **smallest drug concentration** (minimal toxicity)?

dose: 450, # cells:763, total drug:12

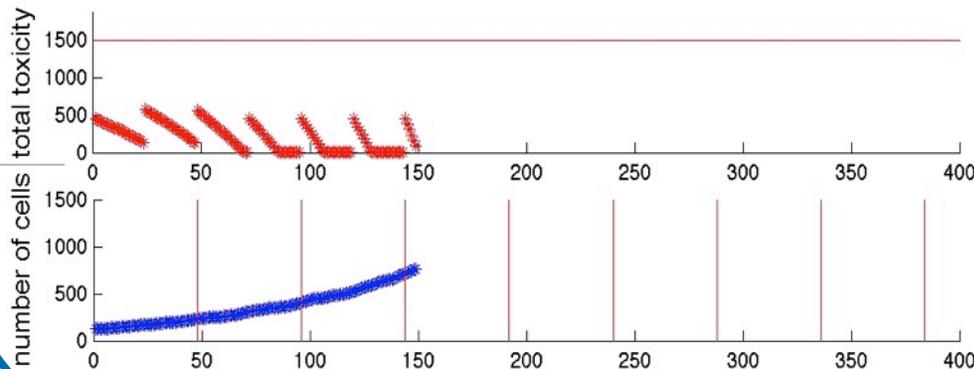


Model of a discrete colony of cells

Level of absorbed drug – color coded
Mutated cells have red perimeter

Drug schedule can be chosen:

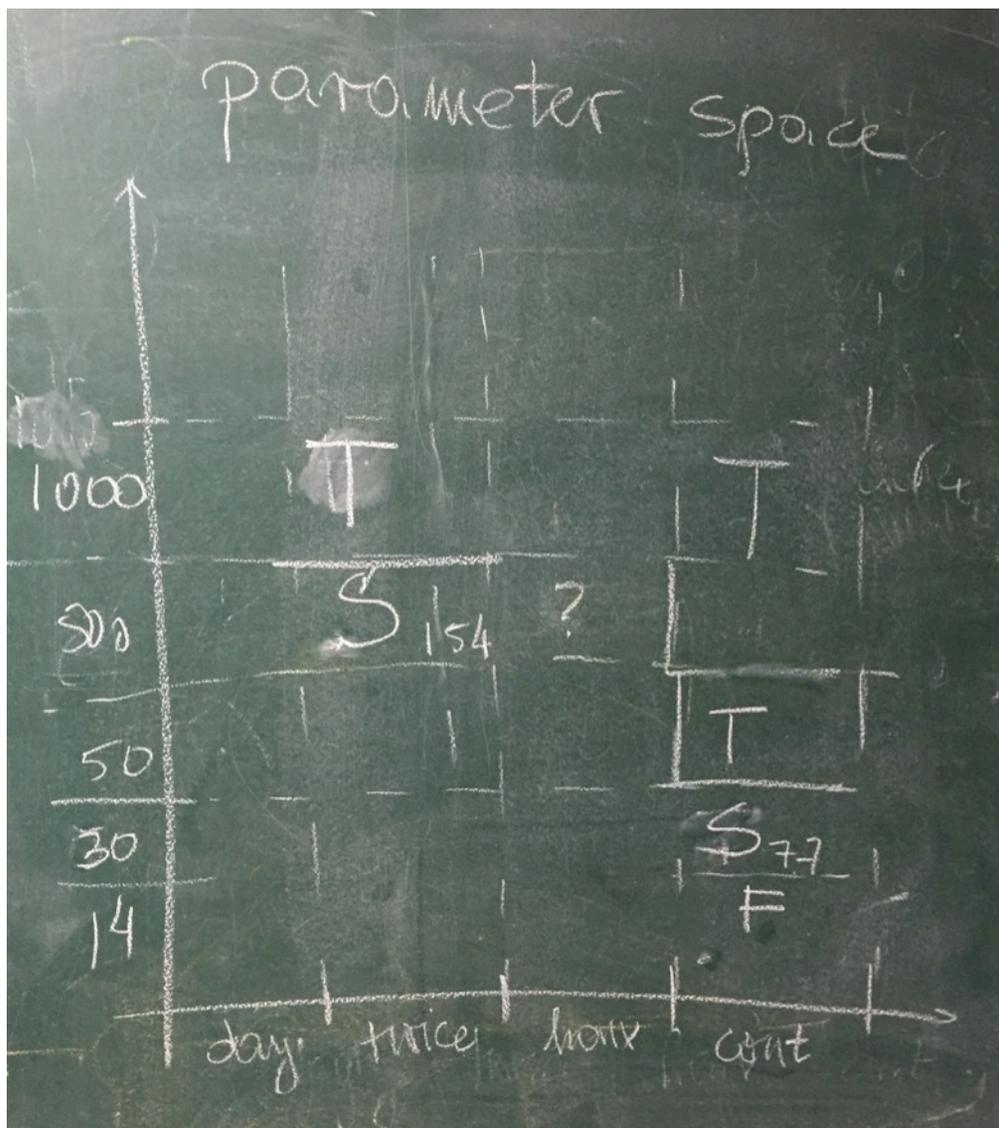
```
*****
possible treatment schedule:
1. daily
2. twice daily
3. hourly
4. continuous
```



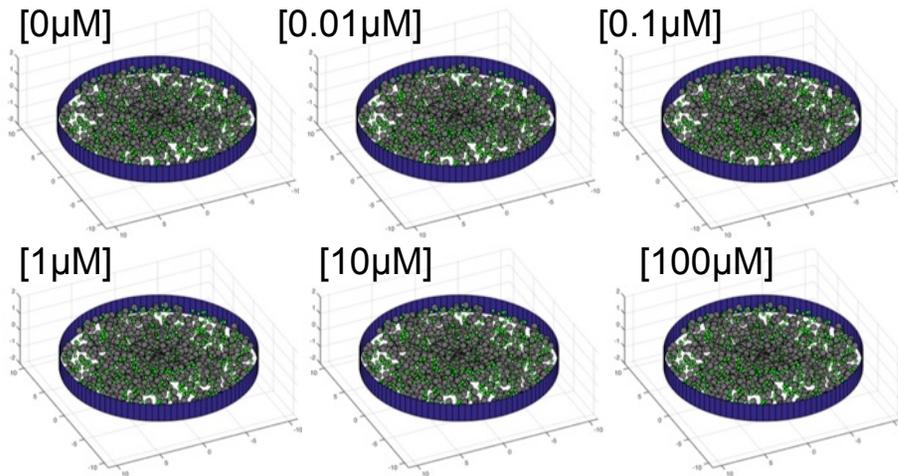
Control total toxicity

Control the number of cells

Let's do some experiments!



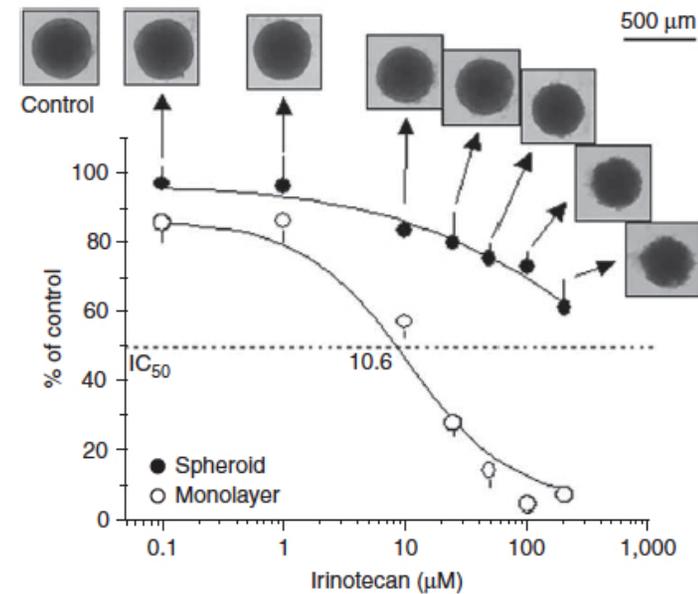
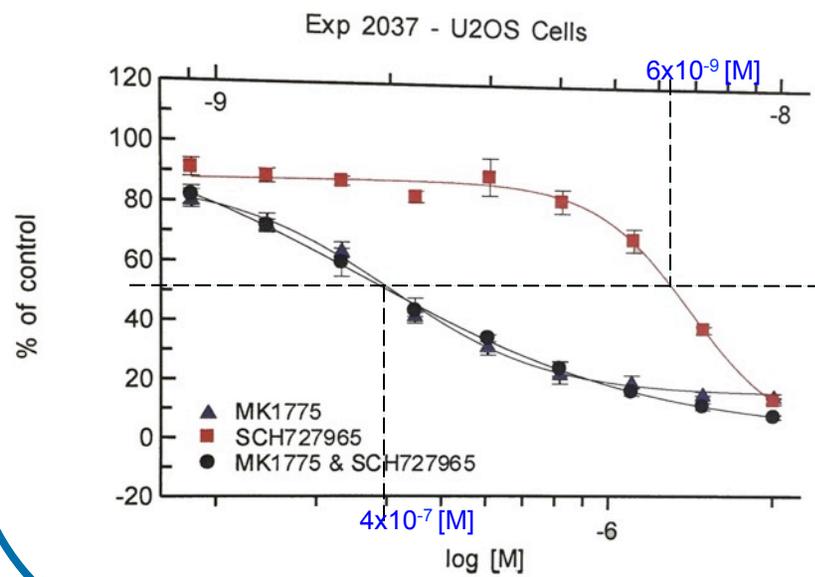
Drug response curves (IC_{50})



IC_{50} curve

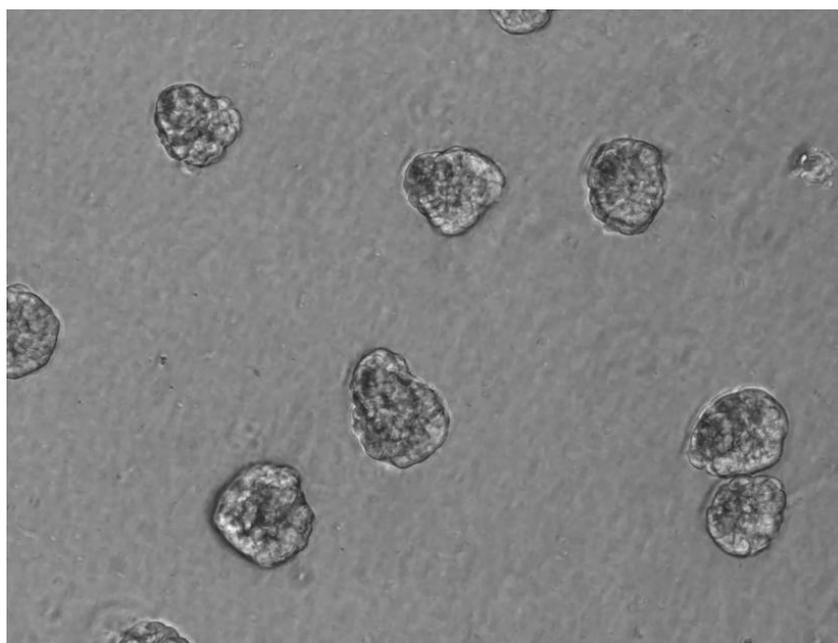
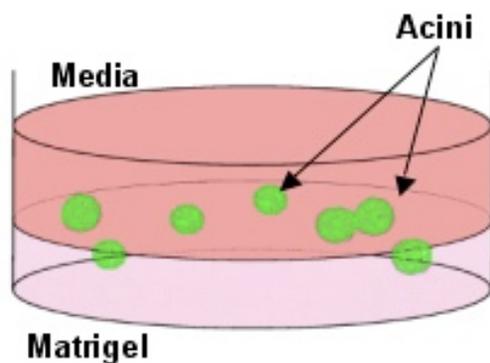
This quantitative measure indicates how much of a particular drug is needed to inhibit a given biological process by half.

The half maximal inhibitory concentration (IC_{50}) is a measure of the effectiveness of a substance in inhibiting a specific biological or biochemical function.

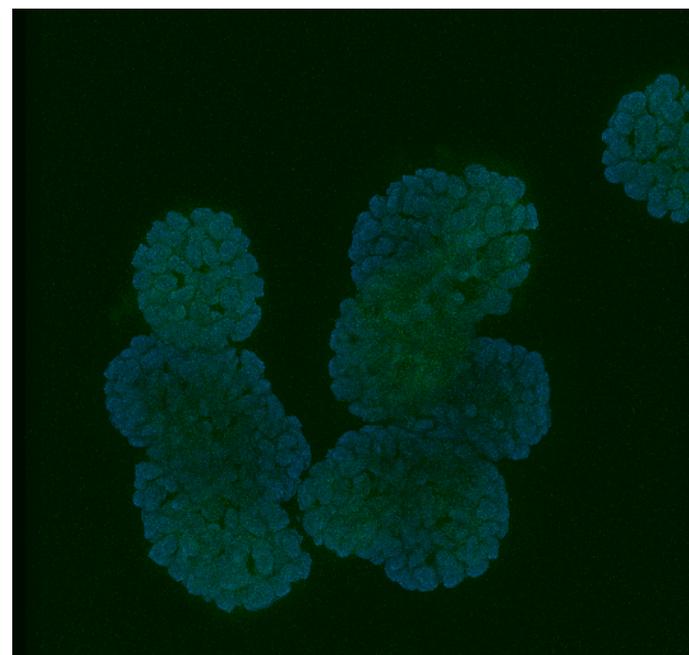


Friedrich et al. Nature Protocols, 2009

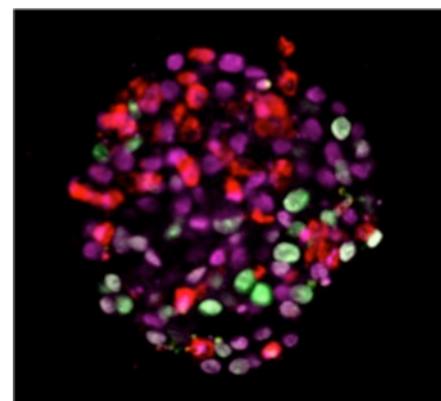
3D tumor spheroids



Tumor cells in 3D culture (top view)



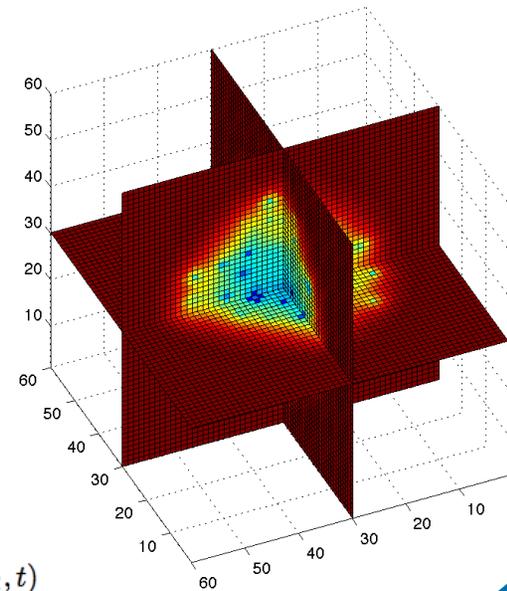
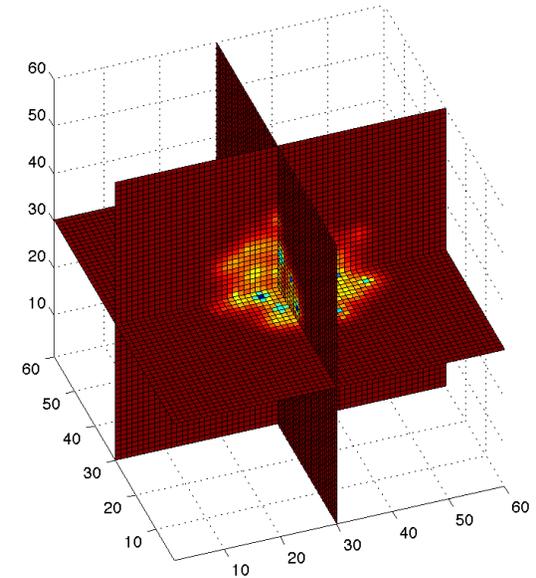
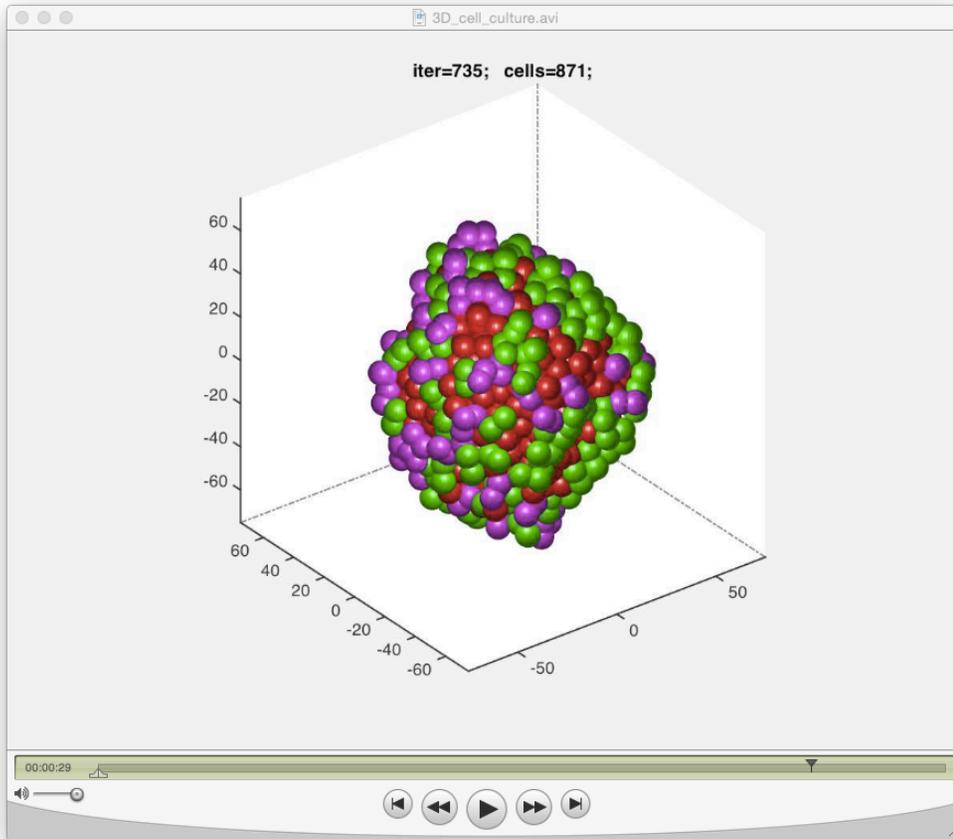
Tumor cells in 3D culture (side view)



Central cross section

3D model of tumor spheroids

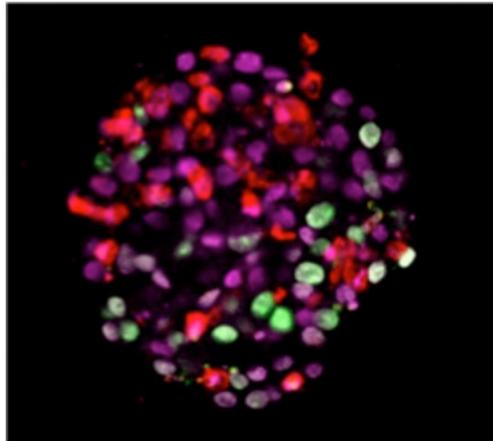
Branton Huffstutler (USF)



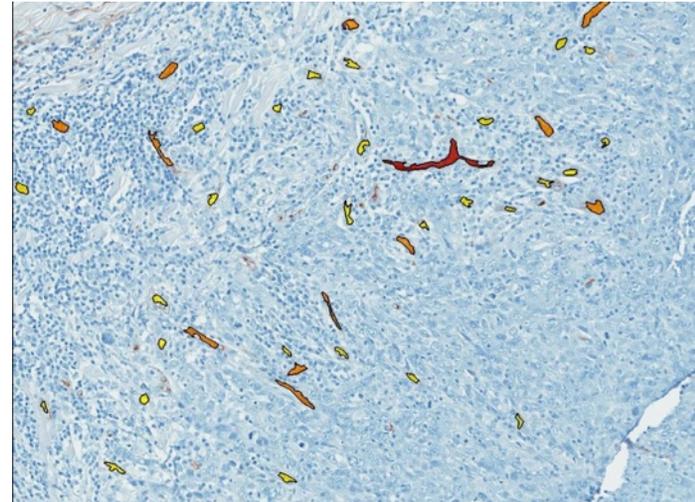
Diffusion equation with cellular uptake

$$\frac{\partial \gamma(x, y, z, t)}{\partial t} = \frac{\partial^2 \gamma(x, y, z, t)}{\partial x^2} + \frac{\partial^2 \gamma(x, y, z, t)}{\partial y^2} + \frac{\partial^2 \gamma(x, y, z, t)}{\partial z^2} - \alpha \gamma(X_{cell}, Y_{cell}, Z_{cell}, t)$$

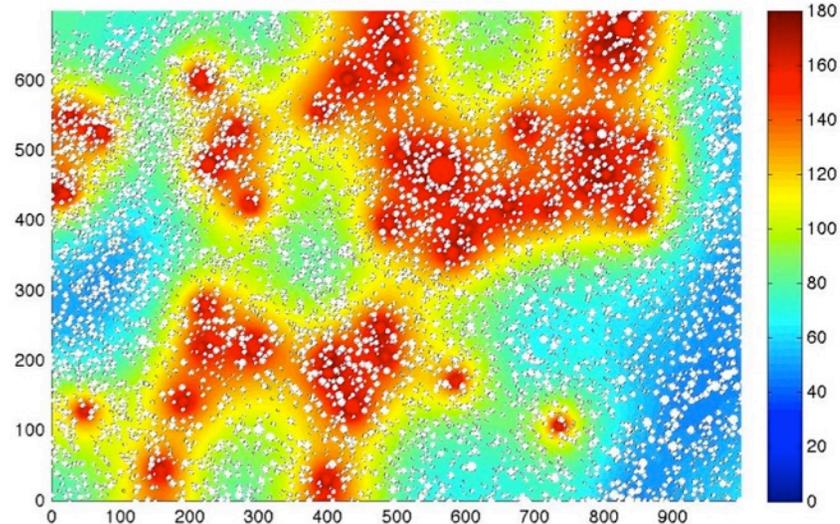
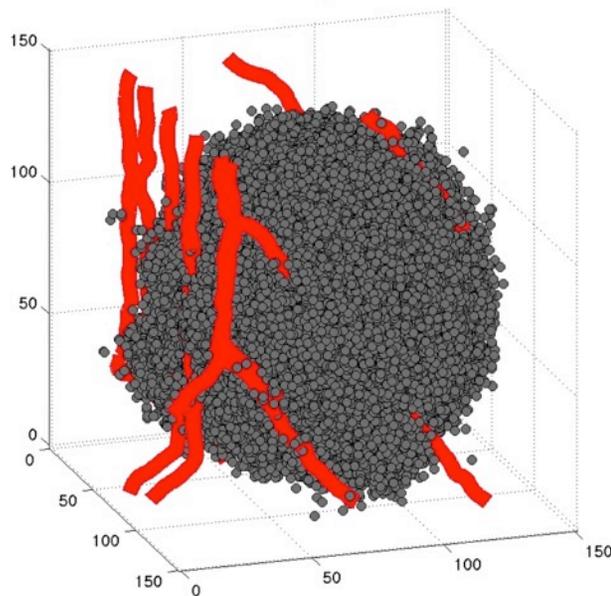
Models of tumor tissue



**Cross section
through 3D spheroid**



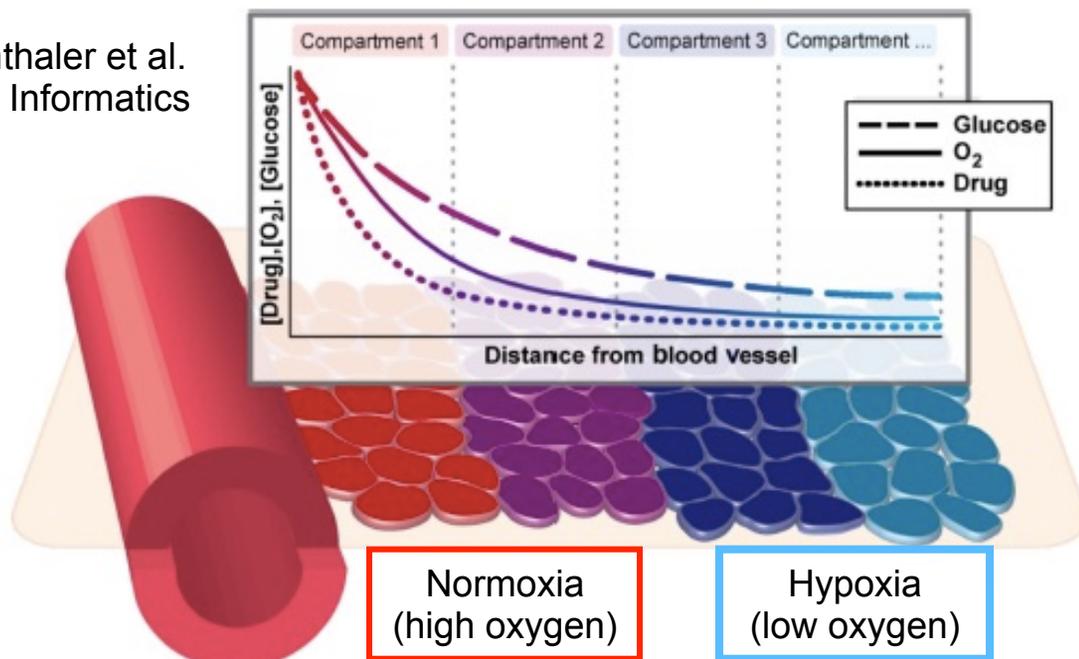
**Cross section
through the tumor tissue**



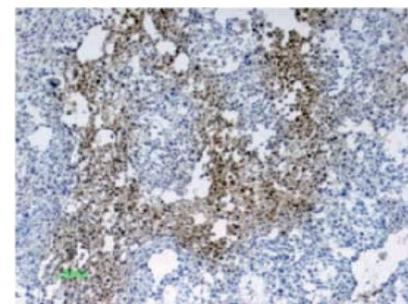
Mathematical models

Hypoxia-activated drugs

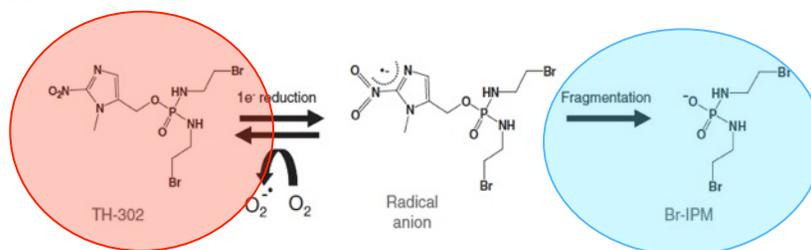
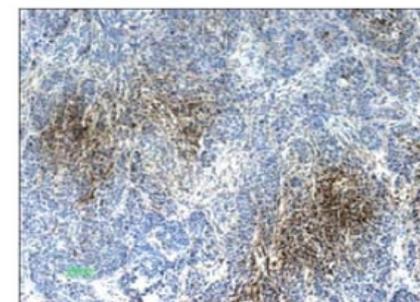
Mumenthaler et al.
Cancer Informatics
2015



Hs766t



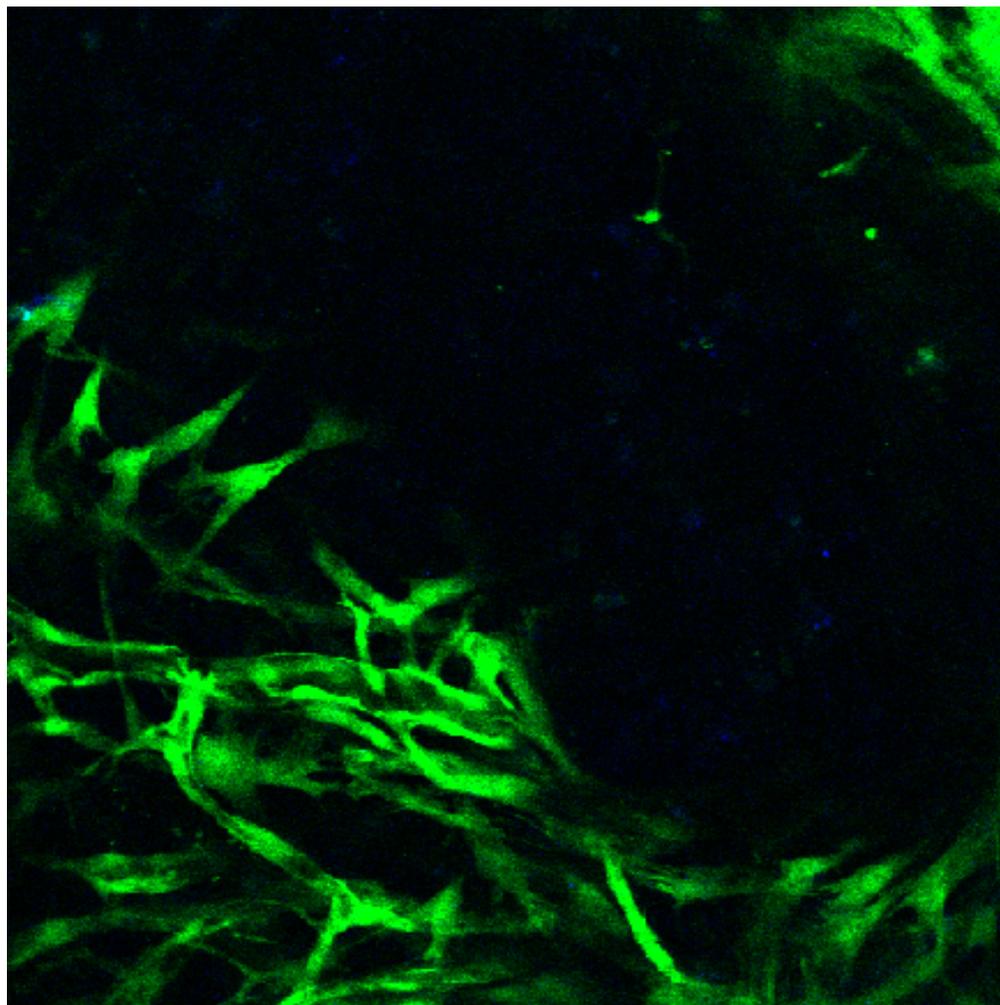
BxPC-3



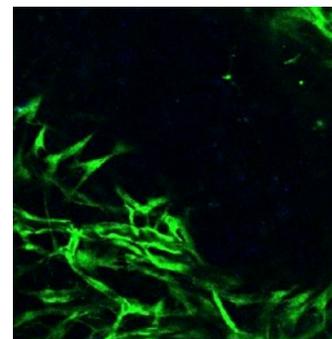
Meng et al. *Mol Cancer Ther*, 2012

Pimonidazole (PIMO) staining
for hypoxia
Sun et al. 2012

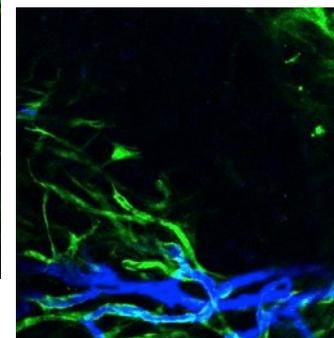
Intravital microscopy model (mice)



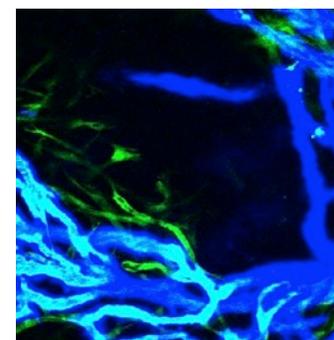
Green – blood vessels
Blue - dextran



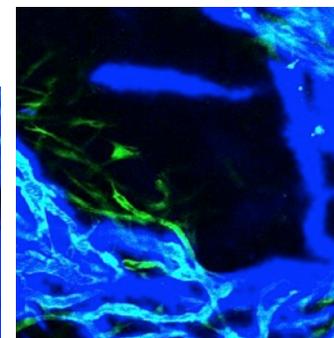
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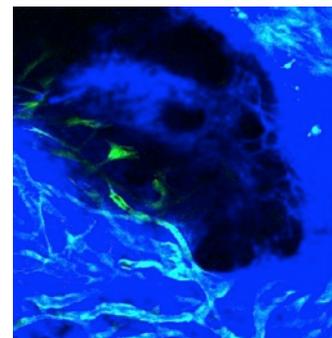
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2



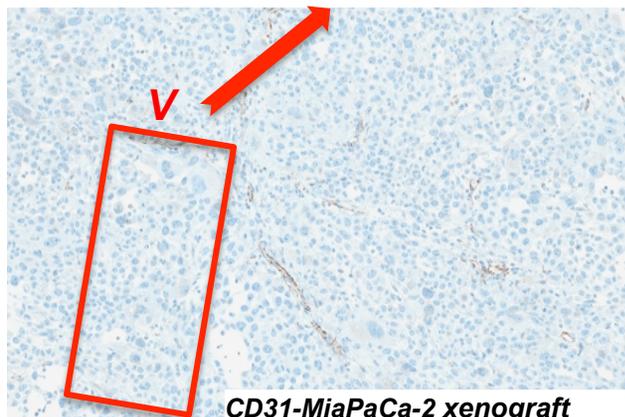
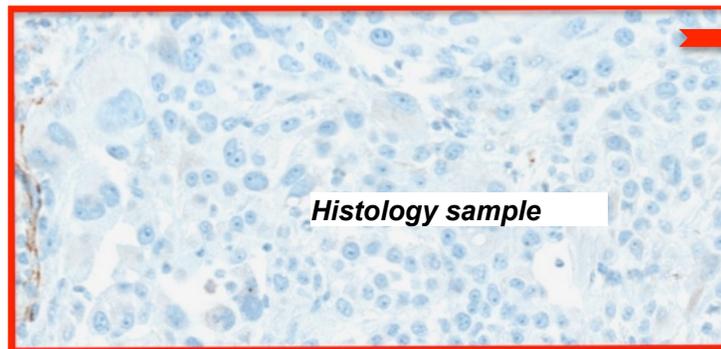
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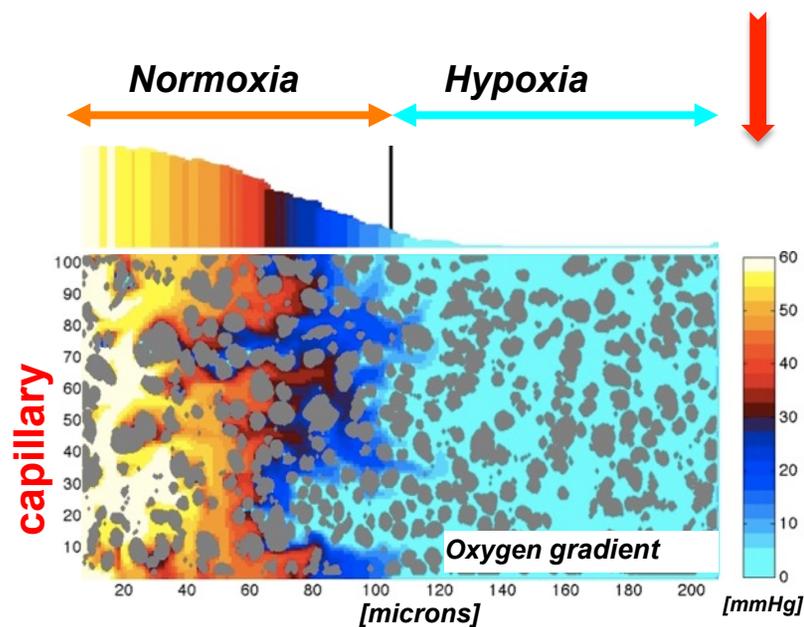
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In silico tissue morphology & tissue oxygenation

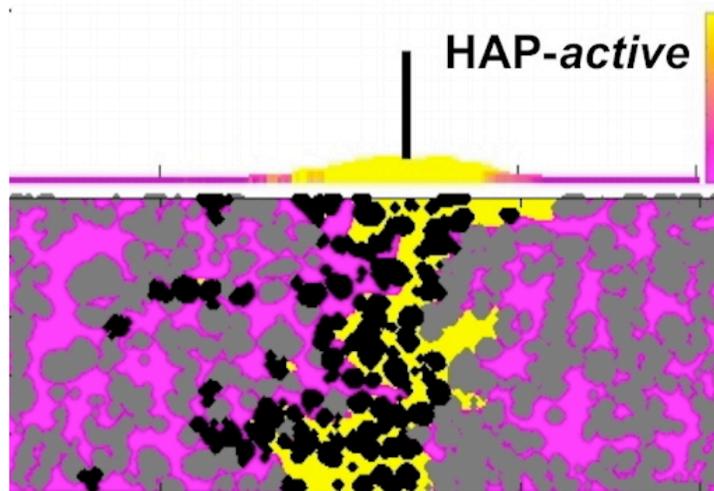
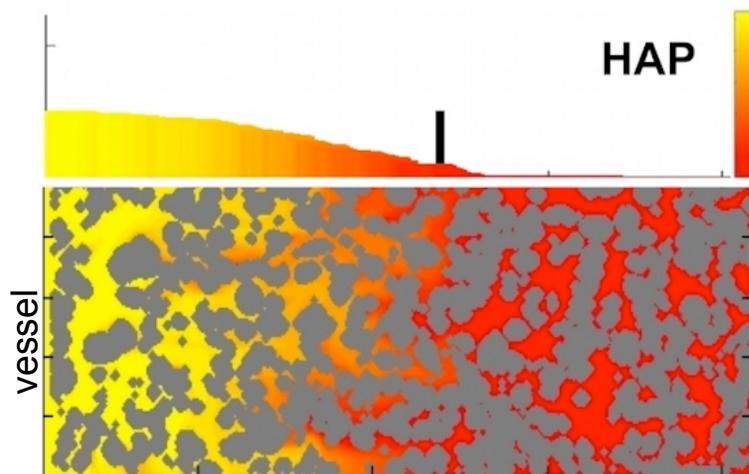
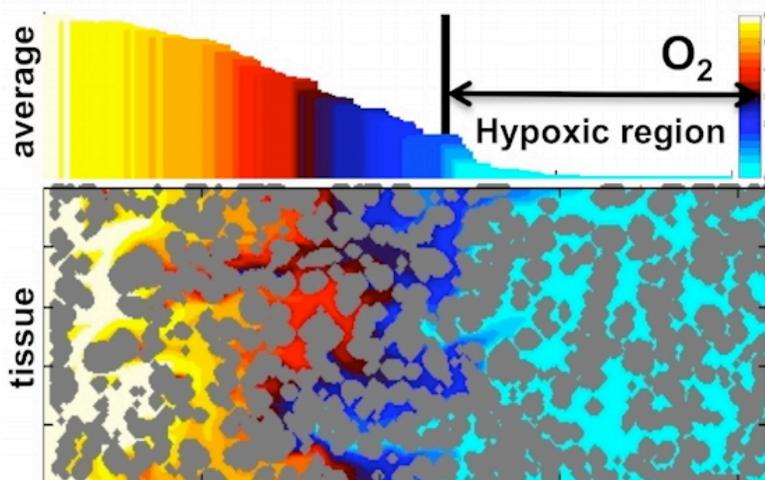
MiaPaCa-2
Human pancreatic
carcinoma cell line



J. Wojtkowiak,
R. Gillies Lab



Normoxia/Hypoxia border



Region of HAP activity— dead cells

$$\mu \Delta \mathbf{u}(\mathbf{x}) = \nabla p(\mathbf{x}) - \mathbf{f}(\mathbf{x}) \quad \text{and} \quad \nabla \cdot \mathbf{u}(\mathbf{x}) = 0 \quad (1)$$

$$\text{where } \mathbf{f}(\mathbf{x}) = f_0 \phi_\varepsilon(\mathbf{x} - \mathbf{x}_0), \quad \text{and} \quad \phi_\varepsilon(\mathbf{x}) = \frac{2\varepsilon^4}{\pi(\|\mathbf{x}\|^2 + \varepsilon^2)} \quad (2)$$

$$\frac{\partial c(\mathbf{x}, t)}{\partial t} = \underbrace{\mathcal{D}_c \Delta c(\mathbf{x}, t)}_{\text{diffusion}} - \underbrace{\mathbf{u}(\mathbf{x}, t) \cdot \nabla c(\mathbf{x}, t)}_{\text{advection}} - \underbrace{\frac{\kappa_m c(\mathbf{x}, t)}{\kappa_n + c(\mathbf{x}, t)}}_{\text{cellular uptake}} \chi(\Omega_\Gamma) \quad (3)$$

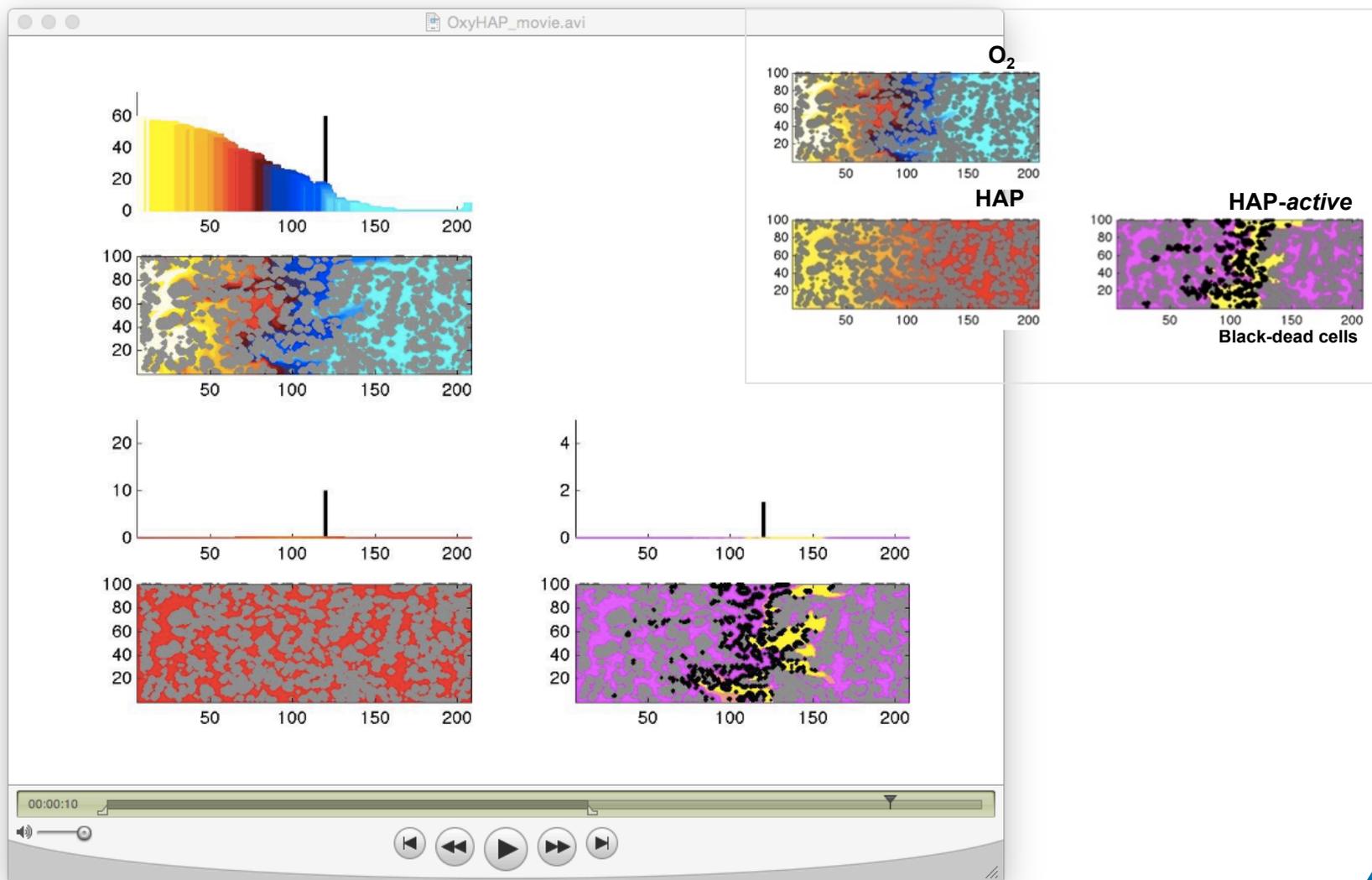
$$\frac{\partial \eta_i(\mathbf{x}, t)}{\partial t} = \underbrace{\mathcal{D}_{\eta_i} \Delta \eta_i(\mathbf{x}, t)}_{\text{diffusion}} - \underbrace{\mathbf{u}(\mathbf{x}, t) \cdot \nabla \eta_i(\mathbf{x}, t)}_{\text{advection}} - \underbrace{\xi(c(\mathbf{x}, t)) \eta_i(\mathbf{x}, t)}_{\text{activation}} - \underbrace{\omega_i \eta_i(\mathbf{x}, t)}_{\text{decay}} \quad (4)$$

$$\frac{\partial \eta_a(\mathbf{x}, t)}{\partial t} = \underbrace{\mathcal{D}_{\eta_a} \Delta \eta_a(\mathbf{x}, t)}_{\text{diffusion}} - \underbrace{\mathbf{u}(\mathbf{x}, t) \cdot \nabla \eta_a(\mathbf{x}, t)}_{\text{advection}} + \underbrace{\xi(c(\mathbf{x}, t)) \eta_i(\mathbf{x}, t)}_{\text{activation}} - \underbrace{\alpha \eta_a(\mathbf{x}, t)}_{\text{cellular uptake}} \chi(\Omega_\Gamma) \quad (5)$$

initial conditions: $\mathbf{u}(\mathbf{x}, t_0) = \mathbf{u}_0(\mathbf{x}), \quad c(\mathbf{x}, t_0) = c_0(\mathbf{x}),$

vessel boundary conditions: $\mathbf{u}(\mathbf{x}_0, t) = \mathbf{u}^0, \quad c(\mathbf{x}_0, t) = c^0, \quad \eta_i(\mathbf{x}_0, t) = \eta_i^0$

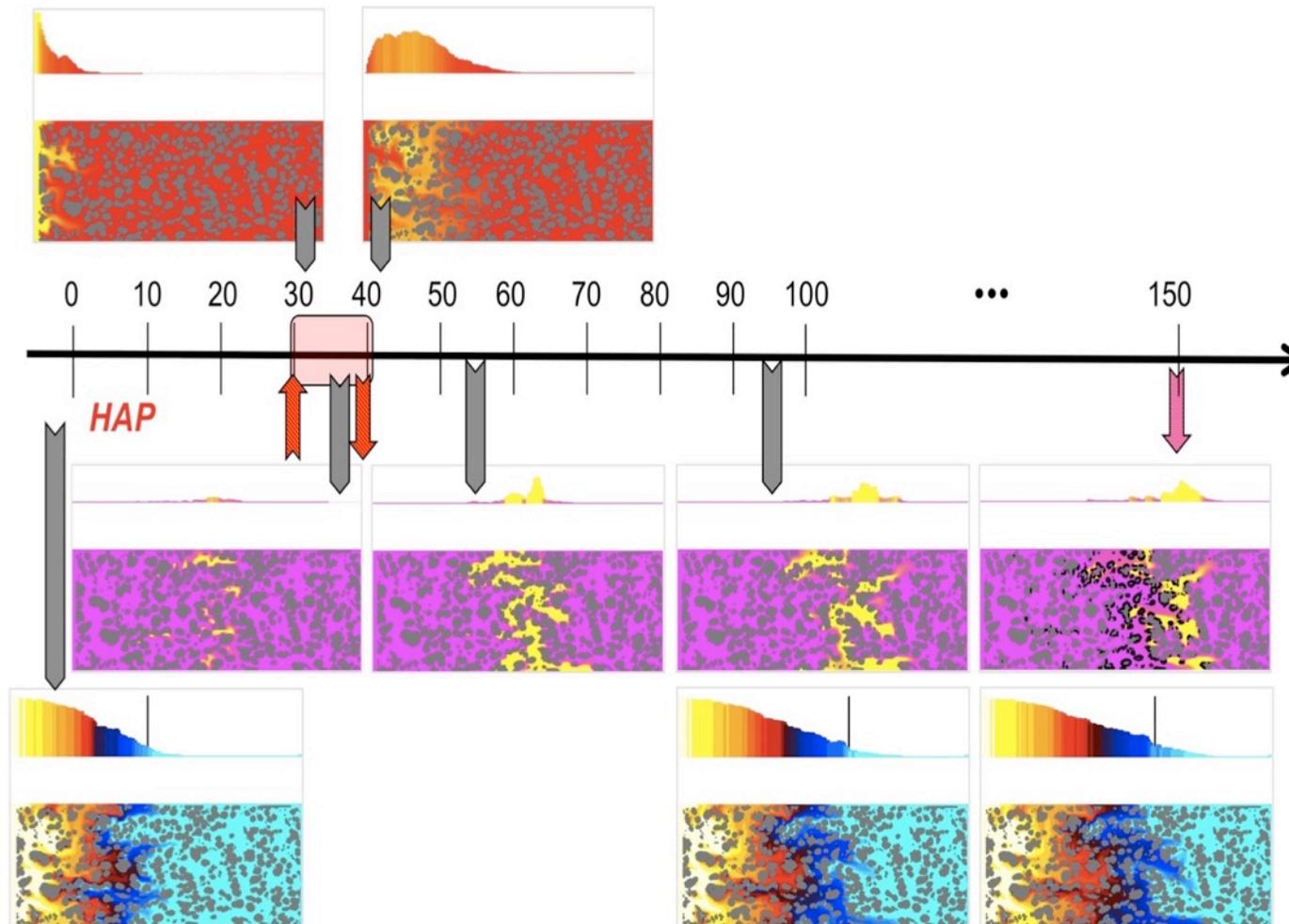
Tissue metabolic landscape under HAP



Inactive drug

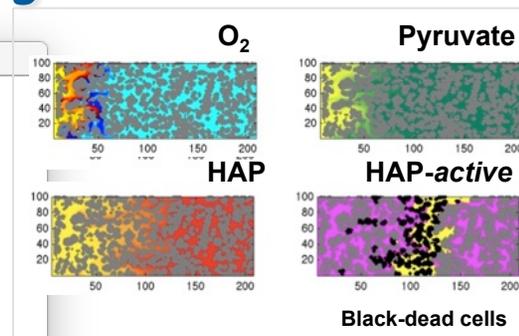
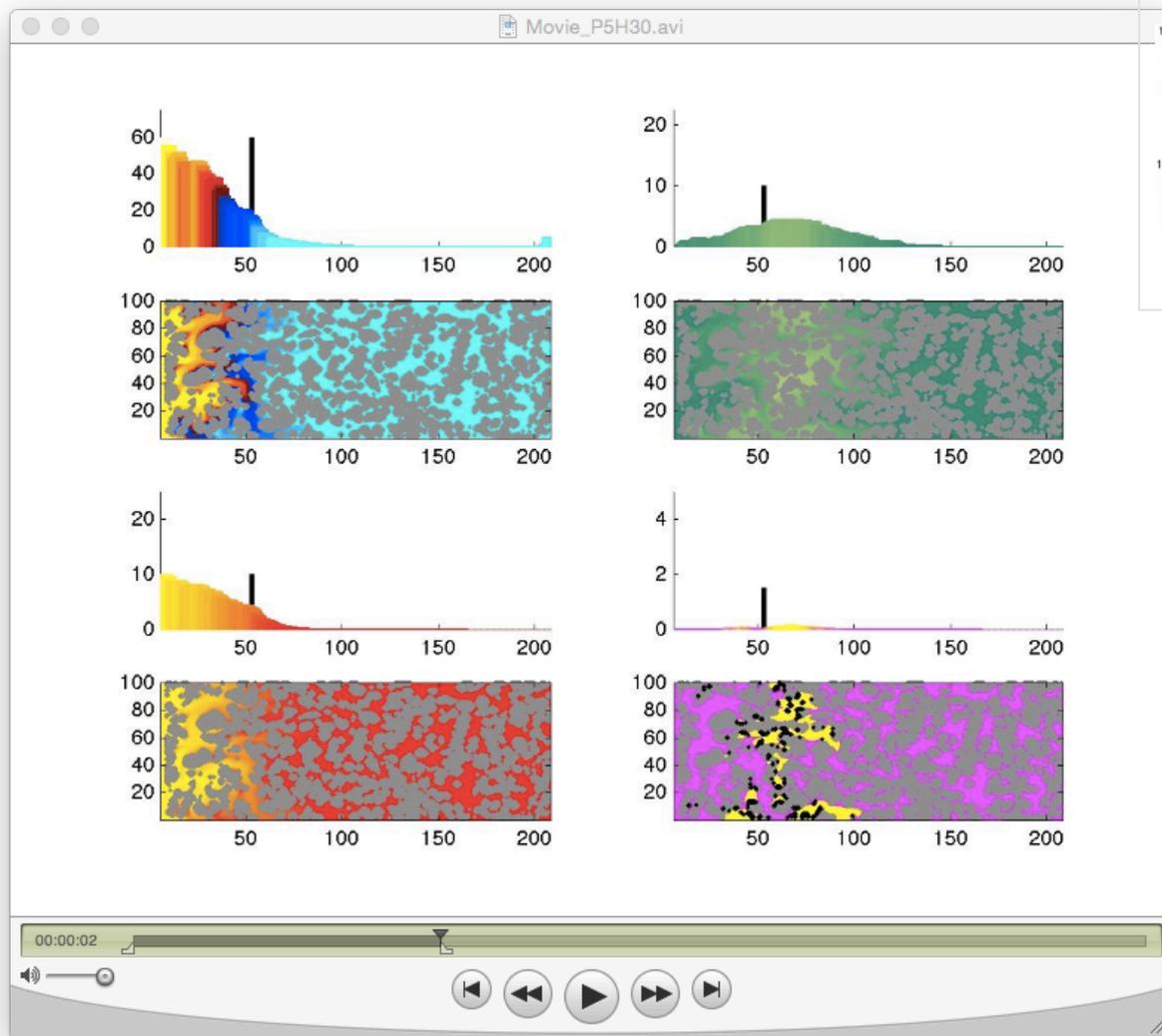
activated drug

HAPs-O₂ interplay



HAP injection can perturb the metabolic landscape of a tissue and increase the O₂ levels due to diminished oxygen uptake as a result of cell death. The steady gradient of O₂ (control case) can be shifted after HAP application.

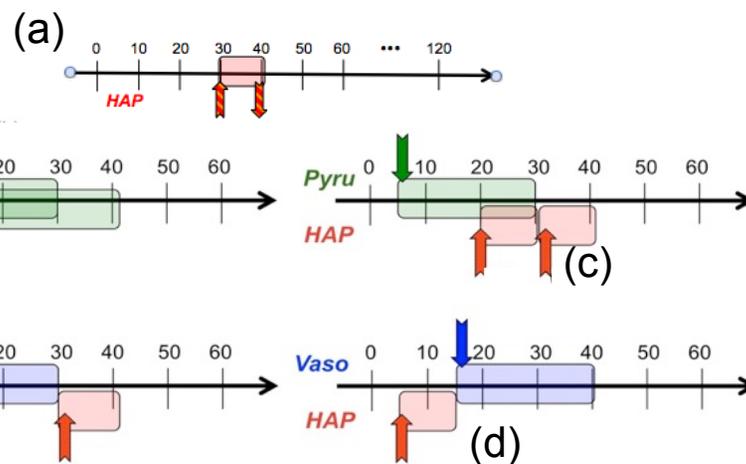
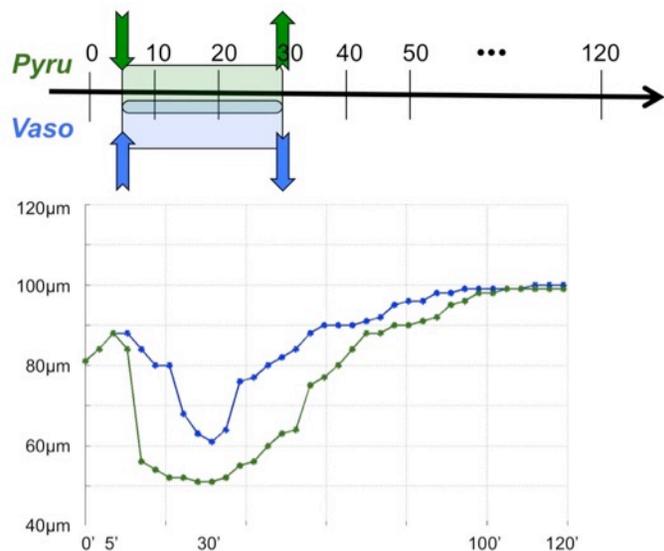
HAP+Pyruvate simulation Combination therapy



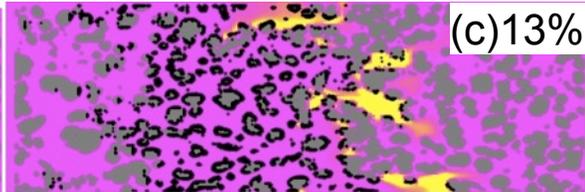
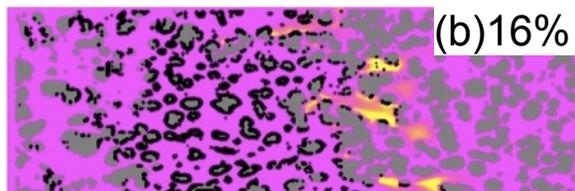
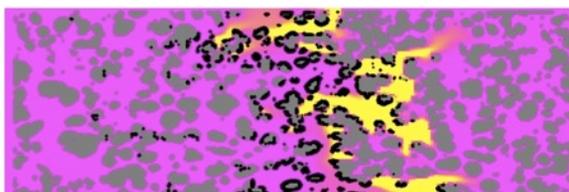
Inactive drug

activated drug

Enhancing HAPs efficacy by combination therapy scheduling



(a) 7%



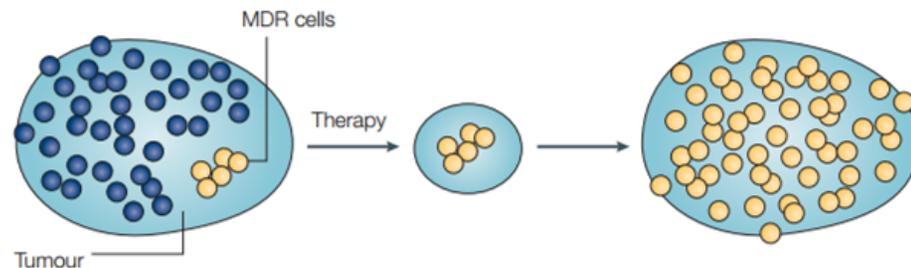
Drug resistance



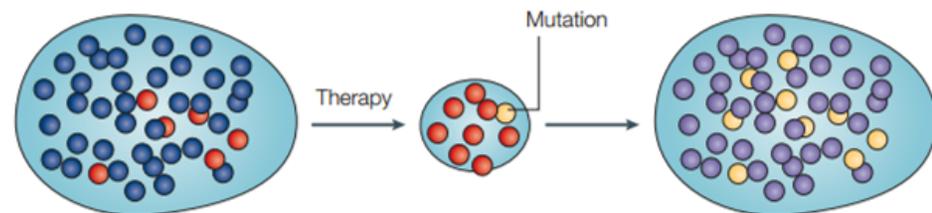
Clinically, drug resistance is defined as a reduced effectiveness of treatment during or after the course of therapy.

As a result, a cancer can progress even months or years after treatment, leading to tumor recurrence.

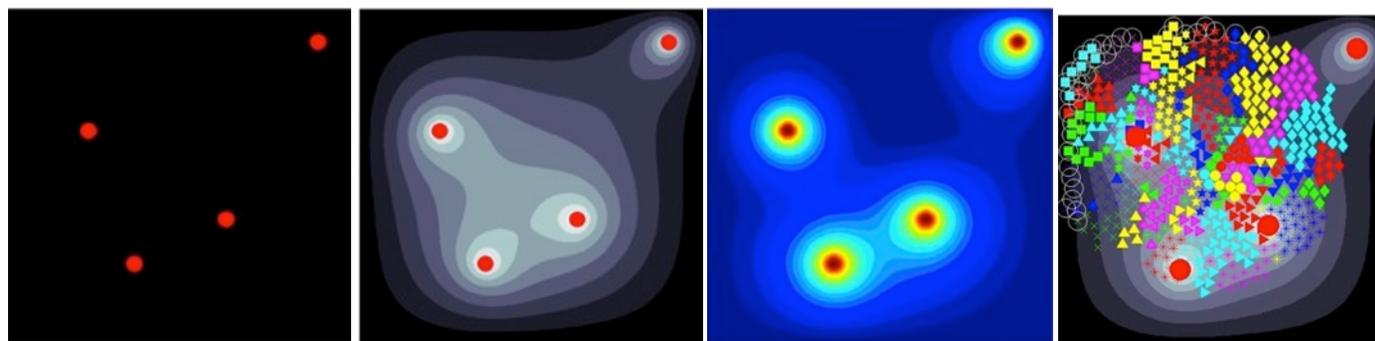
- ***Pre-existing resistance***



- ***Acquired resistance***



Model of drug resistance

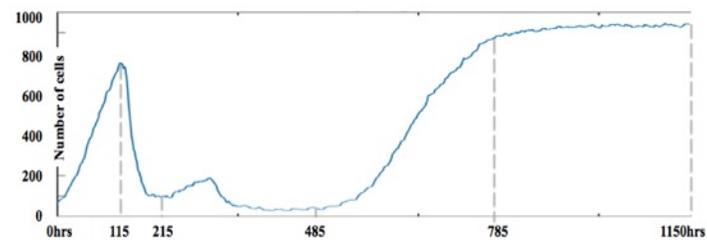
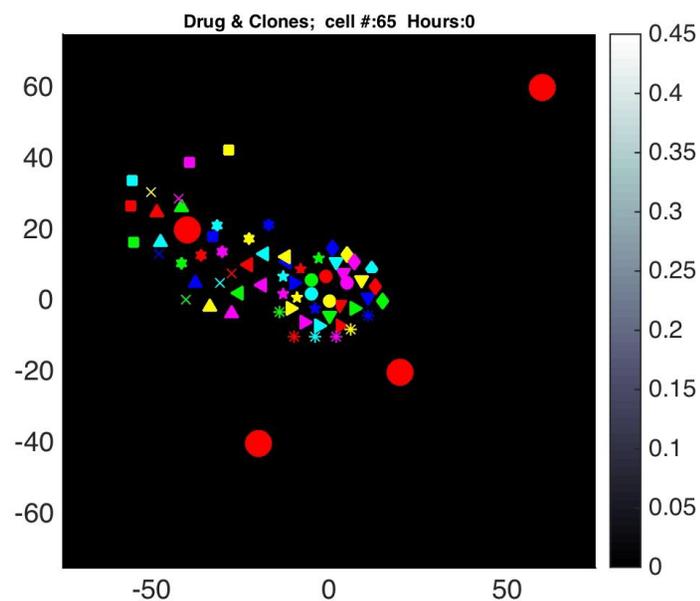


vessels

oxygen gradient

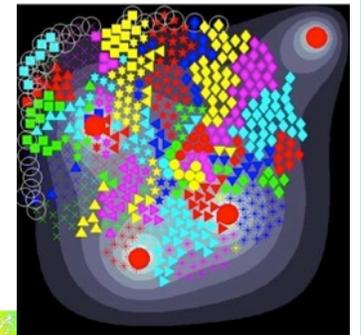
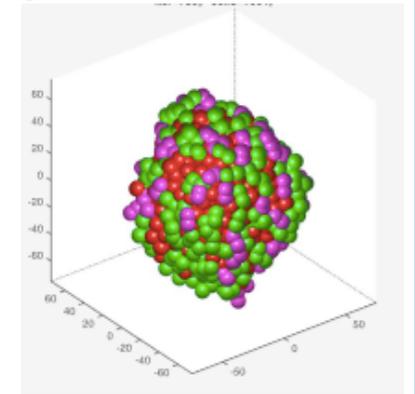
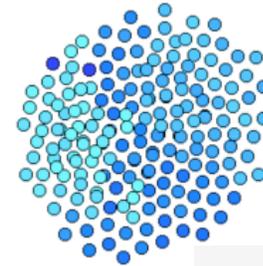
drug gradient

cell clones

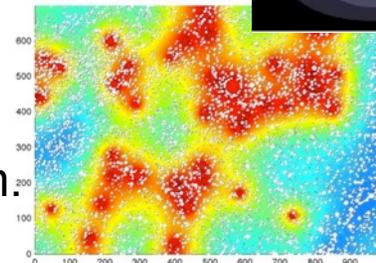


Drug delivery modeling

1. Initial experiments are done in 2D cell cultures to determine whether the drug can kill the cells or suppress their growth.
2. All cell in the 2D culture are exposed to the drug, but in 3D environments (in the tissue or in spheroids) only cells close to the source of drug (vasculature or medium) are exposed to the drug.
3. Combinations of drugs (or other therapeutic treatments) often work better than mono-therapy, however, it is difficult to know what the right order, timing and dosing of the drugs is.
4. Cells exposed to drugs may stop responding to the drug, and become resistant; It is not known how avoid drug resistance.



Math models are helping to address these problems via simulations and analysis, and can help to solve them.



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