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MATHEMATIK

Multiscale  
models for  
glioma  
invasion:  
proliferation  
and therapy  
aspects

Christina  
Surulescu

Motivation

A micro-meso  
setting with  
proliferation  
via go-or-grow

Proliferation  
via cell-tissue  
interactions

Therapy

# Multiscale models for glioma invasion: proliferation and therapy aspects

**Christina Surulescu**

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TU Kaiserslautern

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## Joint work with:

- Christian Engwer & Markus Knappitsch (WWU Münster)
- Alexander Hunt (TU Kaiserslautern)

## Data:

- Carsten Wolters & Felix Lucka (IBB, WWU Münster)
- Katarina Wolf (Radboud Univ. Nijmegen Medical Centre)

# Glioblastoma multiforme

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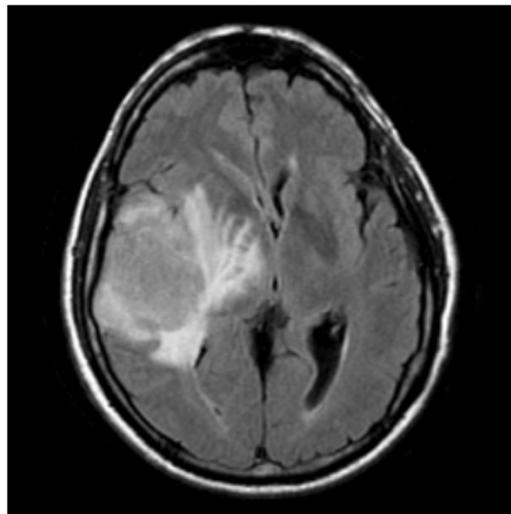
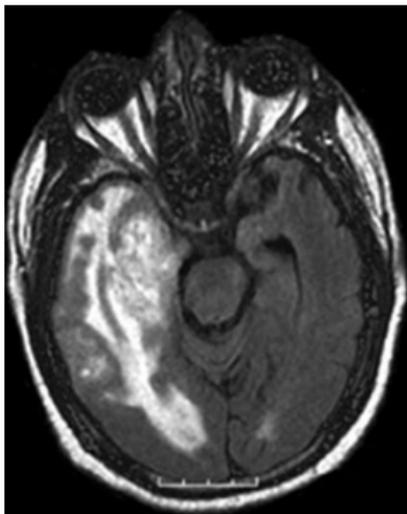
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Pictures from radiopedia.org

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Proliferation via cell-tissue interactions

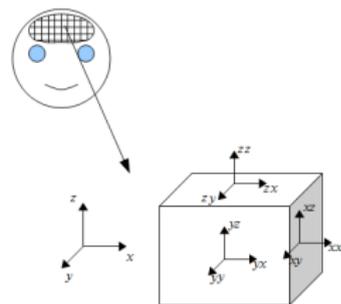
Therapy

- Variant of diffusion-weighted magnetic resonance imaging, DW-MRI.
- Measures the spatial diffusion of water molecules by MRI per volume element (voxel) .



MRI-device (Philips Chieva 3.0 T)  
<http://upload.wikimedia.org/>

Consider a single voxel:



This leads to a diffusion tensor

$$\mathbb{D}(\mathbf{x}) = \begin{pmatrix} d_{xx}(\mathbf{x}) & d_{xy}(\mathbf{x}) & d_{xz}(\mathbf{x}) \\ d_{yx}(\mathbf{x}) & d_{yy}(\mathbf{x}) & d_{yz}(\mathbf{x}) \\ d_{zx}(\mathbf{x}) & d_{zy}(\mathbf{x}) & d_{zz}(\mathbf{x}) \end{pmatrix}$$

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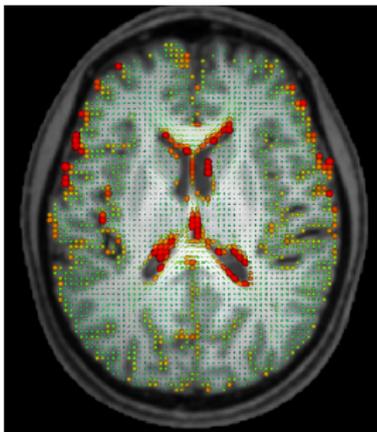
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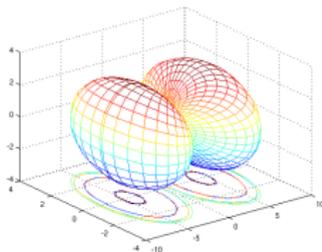
Proliferation via cell-tissue interactions

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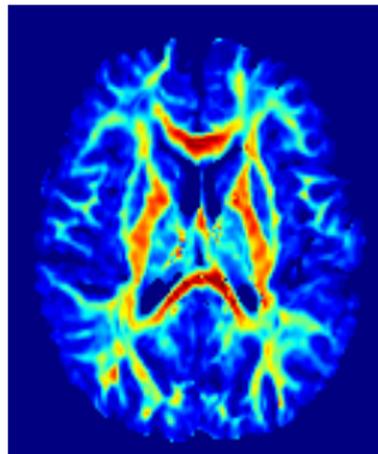
## Ellipsoids



## Peanut



## Fractional anisotropy



$$\frac{\theta_1^2}{\lambda_1} + \frac{\theta_2^2}{\lambda_2} + \frac{\theta_3^2}{\lambda_3} = 1$$

$$\mathbb{S}^2 \ni \theta \mapsto \theta^T \mathbb{D} \theta \in \mathbb{R}$$

$$\bar{\lambda} = \frac{1}{3} \sum_{i=1}^3 \lambda_i$$

$$FA(\mathbf{x}) = \sqrt{\frac{3}{2}} \frac{\sqrt{\sum_{i=1}^3 (\lambda_i - \bar{\lambda})^2}}{\sqrt{\sum_{i=1}^3 \lambda_i^2}}$$

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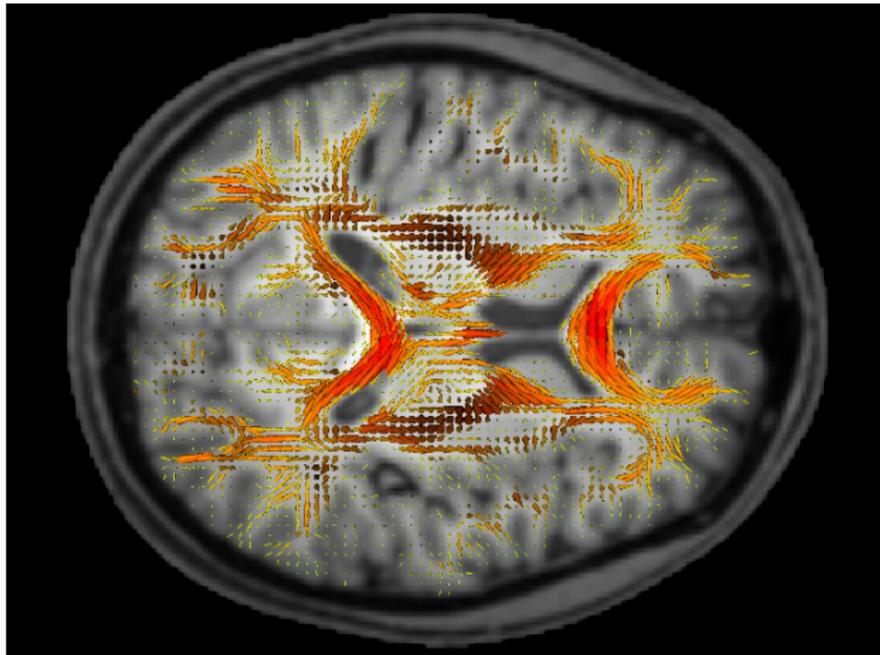
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Visualization of the main direction of diffusion by a vector pointing in the direction of the tensor's leading eigenvalue



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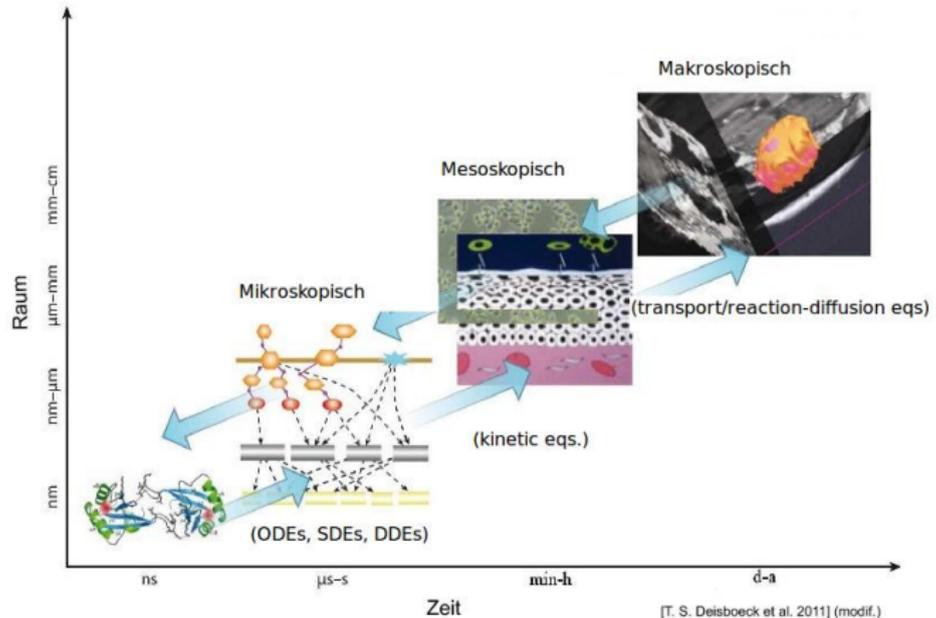
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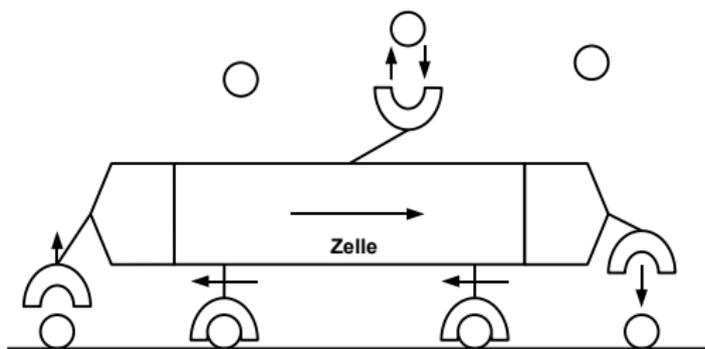
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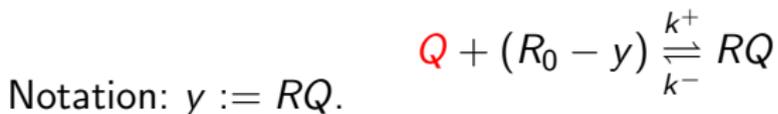


**Goal: multiscale descriptions**

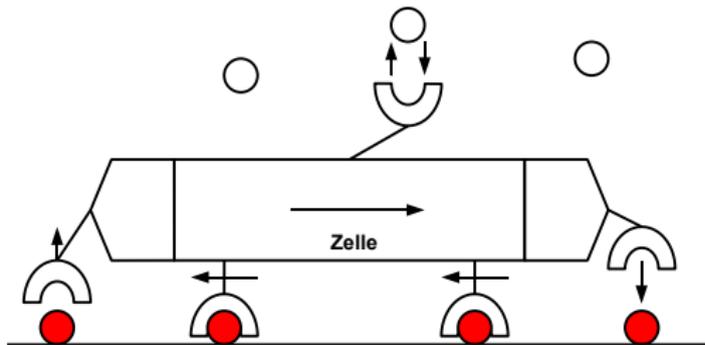
- Cells interact with the neighbouring tissue in order to move forward (contact guidance)



- Receptor binding to unsoluble components  $Q$



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- Receptor binding to unsoluble components  $Q$



# Individual variables ( $N$ cells)

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|                |   |
|----------------|---|
| Position       | $\mathbf{x}^{(j)} \in \mathbb{R}^n$               |
| Velocity       | $\mathbf{v}^{(j)} \in V = s\mathbb{S}^{n-1}$      |
| Receptor state | $\mathbf{y}^{(j)} \in Y \quad (j = 1, \dots, N).$ |

- Newton's law (in the absence of reorientations)

$$\frac{d\mathbf{x}^{(j)}}{dt} = \mathbf{v}^{(j)}, \quad \frac{d\mathbf{v}^{(j)}}{dt} = 0$$

- ODE for receptor dynamics

$$\frac{d\mathbf{y}^{(j)}}{dt} = G(\mathbf{y}^{(j)}, Q(t, \mathbf{x}^{(j)}))$$

- **Goal:** derive a kinetic equation for the cell density  $\rho(t, \mathbf{x}, \mathbf{v}, \mathbf{y})$ .
- In the absence of reorientations:

$$\frac{\partial \rho}{\partial t} + \underbrace{\mathbf{v} \cdot \nabla_{\mathbf{x}} \rho}_{\text{Transport with velocity } \mathbf{v}} + \underbrace{\nabla_{\mathbf{y}} \cdot (\mathbf{G}(\mathbf{y}, Q)\rho)}_{\text{From receptor dynamics}} = 0$$

- Changes in orientation (and speed) have to be incorporated in the right-hand side.

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$$\partial_t p + \nabla_{\mathbf{x}} \cdot (\mathbf{v} p) + \partial_y (G(y)p) = -\lambda(y)p + \lambda(y) \int_V K(\mathbf{x}, \mathbf{v}, \mathbf{v}') p(\mathbf{v}') d\mathbf{v}'$$

with  $p(t, \mathbf{x}, \mathbf{v}, y)$  cell density at time  $t$ , position  $\mathbf{x} \in \mathbb{R}^n$ , with velocity  $\mathbf{v} \in V \subset \mathbb{R}^n$ , and internal state  $y \in Y \subseteq [0, R_0]$ .

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**Subcellular dynamics:**  $\frac{d}{dt} y(t) = G(y(t), Q(t, \mathbf{x}))$ .

$Q(t, \mathbf{x})$ : volume fraction of tissue fibres.

$$\partial_t p + \nabla_{\mathbf{x}} \cdot (\mathbf{v} p) + \partial_y (G(y) p) = -\lambda(y) p + \lambda(y) \int_V K(\mathbf{x}, \mathbf{v}, \mathbf{v}') p(\mathbf{v}') d\mathbf{v}'$$

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$$\dot{y} = k^+(R_0 - y)Q - k^- y.$$

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- Moving cancer cells:

$$\partial_t p + \nabla_{\mathbf{x}} \cdot (\mathbf{v}p) + \partial_y (G(y, Q)p) = \mathcal{L}[\lambda]p - a(\mathbf{x})p + \frac{bq}{\omega}r - \ell(N)p$$

# Proliferation: Go-or-grow

- Moving cancer cells:

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- Non-moving (proliferating) cancer cells:

$$\partial_t r = a(\mathbf{x}) \int_{\mathbf{V}} p d\mathbf{v} - br + g(N)r - \ell(N)r.$$

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- Total cell density (macroscopic):

$$N(t, \mathbf{x}) = \int_V \int_Y p(t, \mathbf{x}, \mathbf{v}, y) dy d\mathbf{v} + \int_Y r(t, \mathbf{x}, y) dy$$

# Receptor dynamics in a static field



Steady state:  $y^* = \frac{k^+ QR_0}{k^+ Q + k^-}$ .

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# Receptor dynamics in a static field

Steady state:  $y^* = \frac{k^+ QR_0}{k^+ Q + k^-}$ .

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**Turning rate:**  $\lambda(z) = \lambda_0 - \lambda_1 z \geq 0$ , with adequate  $\lambda_0, \lambda_1 > 0$ .

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Write the transport equations w.r.t.  $z$  and consider the moments w.r.t.  $z$  and  $\mathbf{v}$ .

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Write the transport equations w.r.t.  $z$  and consider the moments w.r.t.  $z$  and  $\mathbf{v}$ . Wanted:

$$M(t, \mathbf{x}) := \iint_{\mathbf{v} \times Z} p(t, \mathbf{x}, \mathbf{v}, z) dz d\mathbf{v}$$

$$w(t, \mathbf{x}, \mathbf{v}) := \int_Z r(t, \mathbf{x}, \mathbf{v}, z) dz$$

to recover the macroscopic cell density  $N(t, \mathbf{x})$ .

Assume the internal dynamics equilibrates rapidly, s.t. the system is close to steady-state.

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# Parabolic scaling

Assume the internal dynamics equilibrates rapidly, s.t. the system is close to steady-state. **Scaling:**  $\hat{t} = \varepsilon^2 t$ ,  $\hat{x} = \varepsilon x$ .

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**Assumption:** the time scale on which birth and death events occur is much slower than the (biased) random walk process.

$$g(N) \rightarrow \varepsilon^2 \hat{g}(\hat{N})$$

$$\ell(N) \rightarrow \varepsilon^2 \hat{\ell}(\hat{N}).$$

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Then

$$\begin{aligned} \partial_t N_0 - \nabla \cdot \left( \frac{1}{\lambda_0 + a(\mathbf{x})} \nabla \cdot \left( \frac{b}{a(\mathbf{x}) + b} \mathbb{D}_{\mathcal{T}(\mathbf{x})} N_0 \right) \right) \\ + \nabla \cdot \left( \frac{\lambda_1}{\lambda_0 + a(\mathbf{x})} \gamma(\mathbf{x}) f'(Q) \frac{b}{a(\mathbf{x}) + b} \mathbb{D}_{\mathcal{T}(\mathbf{x})} \cdot \nabla Q N_0 \right) \\ = \frac{a(\mathbf{x})}{a(\mathbf{x}) + b} g(N_0) N_0 - N_0 \ell(N_0), \end{aligned}$$

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with the **tumor diffusion tensor**  $\mathbb{D}_T(\mathbf{x}) = \frac{1}{\omega} \int_V \mathbf{v} \mathbf{v}^t q(\hat{\mathbf{v}}) d\mathbf{v}$ .

# Effective equations on the macroscale

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With the logistic growth choice  $g(N_0) = c_g$ ,  $\ell(N_0) = c_\ell N_0$ , where  $N_0 = \frac{a+b}{b} M_0$ , we get

$$\begin{aligned} \partial_t N_0 - c_D(\mathbf{x}) \nabla \nabla (\mathbb{D}_T(\mathbf{x}) N_0) - \lambda_1 c_D(\mathbf{x}) \nabla (\mathbf{u}(\mathbf{x}) N_0) \\ = \frac{a}{a+b} c_g N_0 - c_\ell N_0^2, \end{aligned}$$

with  $c_D(\mathbf{x}) = \frac{b}{(\lambda_0 + a(\mathbf{x}))(a(\mathbf{x}) + b)}$  and the drift velocity

$$\mathbf{u}(\mathbf{x}) = \gamma(\mathbf{x}) f'(Q(\mathbf{x})) \mathbb{D}_T(\mathbf{x}) \nabla Q,$$

where  $\gamma(\mathbf{x}) = (k^+ Q + k^- + \lambda_0 + a)^{-1}$  and  $f(Q(\mathbf{x})) = \frac{k^+ Q(\mathbf{x}) R_0}{k^+ Q(\mathbf{x}) + k^-}$ .

# Determine explicit forms of the coefficients



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Choices of  $Q(\mathbf{x})$ :

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Choices of  $Q(\mathbf{x})$ :

- Fractional anisotropy (FA), from data.

# Determine explicit forms of the coefficients

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## Choices of $Q(\mathbf{x})$ :

- Fractional anisotropy (FA), from data.

**Problem:** assumes  $Q$  to be high where tissue is strongly aligned  $\rightsquigarrow$  also true in regions of isotropic (non-aligned) and densely packed tissue??

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- Estimated  $Q$  via free path length from diffusivity measured by DTI:

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- **Estimated  $Q$  via free path length from diffusivity measured by DTI:**

Characteristic (diffusion) length:

$$l_c = \sqrt{Dt_c},$$

with  $D$  a diffusion-related coefficient and  $t_c$  the characteristic (diffusion) time.

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- **Fractional anisotropy (FA)**, from data.  
**Problem:** assumes  $Q$  to be high where tissue is strongly aligned  $\rightsquigarrow$  also true in regions of isotropic (non-aligned) and densely packed tissue??

- **Estimated  $Q$  via free path length from diffusivity measured by DTI:**

Characteristic (diffusion) length:

$$l_c = \sqrt{Dt_c},$$

with  $D$  a diffusion-related coefficient and  $t_c$  the characteristic (diffusion) time.

**Choice of  $D$ :**  $tr(\mathbb{D}_W)$ .

# Determine explicit forms of the coefficients

Multiscale  
models for  
glioma  
invasion:  
proliferation  
and therapy  
aspects

Christina  
Surulescu

Motivation

A micro-meso  
setting with  
proliferation  
via go-or-grow

Proliferation  
via cell-tissue  
interactions

Therapy

Choice of characteristic time  $t_c$ : expected exit time of  
Brownian motion from a ball with minimal radius surrounding  
the voxel of length  $h$ , hence  $t_c = \frac{h^2}{4}$ .

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This estimate is valid for  $\mathcal{N}(0, t - s)$ -distributed increments, and ours are  $\mathcal{N}(0, \sigma \cdot (t - s))$ -distributed, where  $\sigma$  is some estimation of the diffusion speed. We choose  $\sigma = l_1$ , where  $l_1$  is the largest eigenvalue of  $\mathbb{D}_W$ .

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Volume fraction of tissue fibers:  $l_c = \sqrt{\frac{h^2 \text{tr}(\mathbb{D}_W)}{4l_1}}$ .

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Volume fraction of tissue fibers:  $l_c = \sqrt{\frac{h^2 \text{tr}(\mathbb{D}_W)}{4l_1}}$ .

The free volume fraction of one voxel is  $l_c^3/h^3$ . So the occupied volume is  $Q = 1 - \frac{l_c^3}{h^3}$ .

# Volume fraction of tissue fibers

Multiscale  
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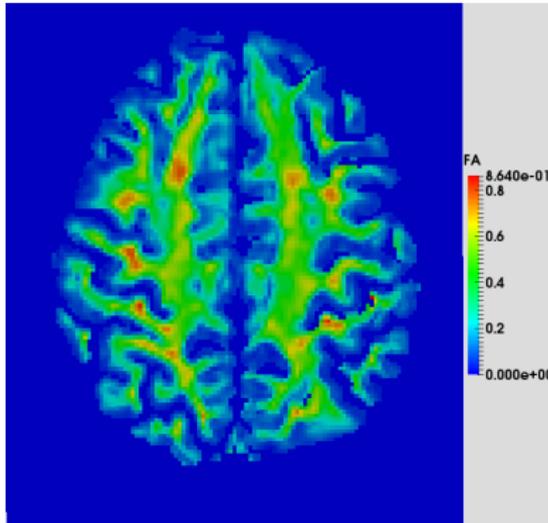
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Surulescu

Motivation

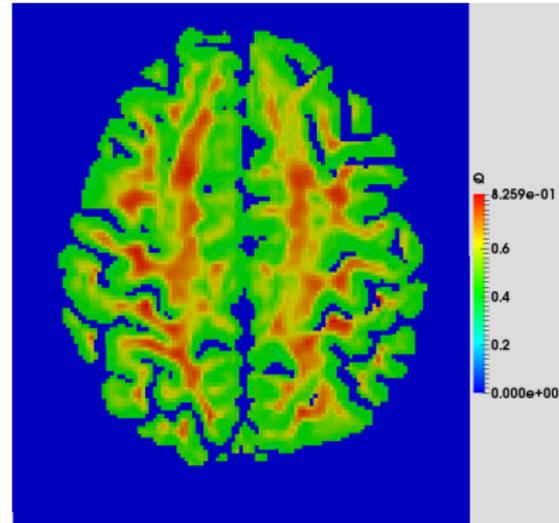
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FA



Estimated Q

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Choices of  $q(\mathbf{x}, \theta)$ :

- Peanut:  $q(\mathbf{x}, \theta) = \frac{n}{|\mathbb{S}^{n-1}| \text{tr} \mathbb{D}_W(\mathbf{x})} \theta^t \mathbb{D}_W(\mathbf{x}) \theta.$

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- orientation distribution function (ODF):

$$q(\mathbf{x}, \theta) = ODF(\theta) := \int_0^\infty \Pi(r\theta) r^2 dr, \text{ with } \Pi(r\theta):$$

displacement probability of a spatial point in spherical coordinates. It can be shown (Aganj et al 2010) that

$$q(\mathbf{x}, \theta) = \frac{1}{4\pi |\mathbb{D}_W(\mathbf{x})| (\theta^t \mathbb{D}_W(\mathbf{x})^{-1} \theta)^{\frac{3}{2}}}$$

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The tumor diffusion tensor can be explicitly computed for each of these choices.

# Simulations (with FA and peanut)

Multiscale models for glioma invasion: proliferation and therapy aspects

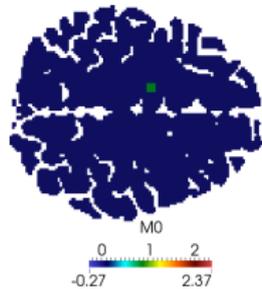
Christina Surulescu

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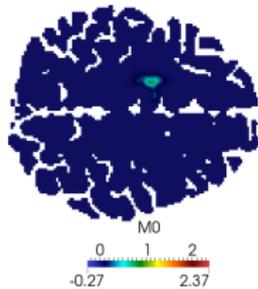
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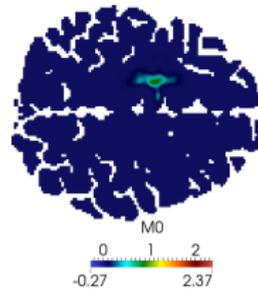
Therapy



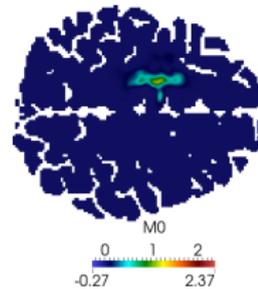
$t = 0$



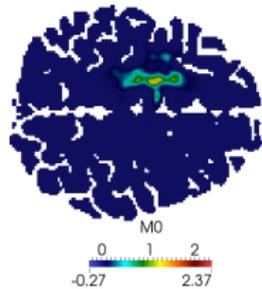
$t = 100 \cdot 10^4 s$



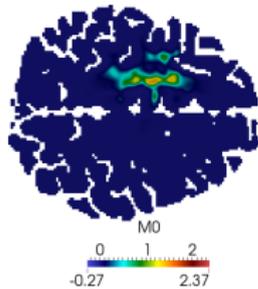
$t = 200 \cdot 10^4 s$



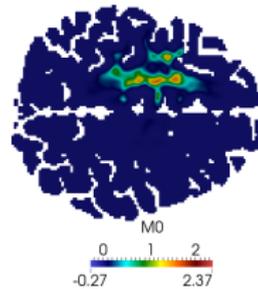
$t = 300 \cdot 10^4 s$



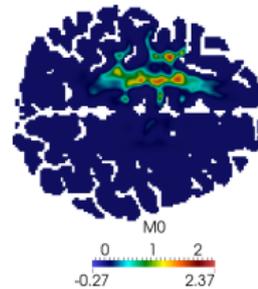
$t = 400 \cdot 10^4 s$



$t = 500 \cdot 10^4 s$



$t = 600 \cdot 10^4 s$



$t = 700 \cdot 10^4 s$

# Alternative modeling of proliferation: interactions with tissue

FELIX KLEIN  
ZENTRUM FÜR  
MATHEMATIK

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$$\mathcal{P}(p) = \underbrace{\mu(\mathbf{x}, M, \mathbf{v})}_{\text{growth rate}} \int_Z \chi(\mathbf{x}, z, z') p(t, \mathbf{x}, \mathbf{v}, z') Q(\mathbf{x}) dz',$$

$$\text{with } M(t, \mathbf{x}) = \int_V \int_Z p(t, \mathbf{x}, \mathbf{v}, z) dz d\mathbf{v}.$$

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The kernel  $\chi$  characterizes the transition from the state  $z'$  to the state  $z$  during a proliferative action.

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Then our kinetic transport equation becomes

$$\begin{aligned} \partial_t p + \nabla_{\mathbf{x}} \cdot (\mathbf{v}p) - \partial_z(((k^+ Q + k^-)z - f'(Q)\mathbf{v} \cdot \nabla Q)p) \\ = \mathcal{L}[\lambda_0]p - \mathcal{L}[\lambda_1]zp + \mathcal{P}(p). \end{aligned}$$

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**Remark:** KTAP by Bellomo assumes cell-cell interactions

$$P_i[p](t, z) = \sum_{h=1}^n \sum_{k=1}^n \mu_{hk} \iint_{Z \times Z} \chi_{hk}^i(z', z''; z) p_h(t, z') p_k(t, z'') dz' dz''$$

Assume  $\mu$  does not depend on  $\mathbf{v}$ . Then by doing again a **parabolic scaling** we get

$$\partial_t M_0 - \nabla \nabla : (\mathbb{D}_T M_0) + \nabla \cdot (g(Q(\mathbf{x})) \mathbb{D}_T \nabla Q M_0) = Q(\mathbf{x}) \mu(\mathbf{x}, M_0) M_0$$

with the drift velocity

$$\mathbf{u}(\mathbf{x}) = \frac{1}{\lambda_0 \omega} \int_V \mathbf{v} \otimes \mathbf{v} \nabla q \, d\mathbf{v}$$

and the tumor diffusion tensor  $\mathbb{D}_T(\mathbf{x}) = \frac{1}{\lambda_0 \omega} \int_V q \mathbf{v} \otimes \mathbf{v} \, d\mathbf{v}$ .

# Simulation results (with peanut)

Multiscale models for glioma invasion: proliferation and therapy aspects

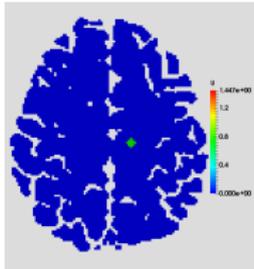
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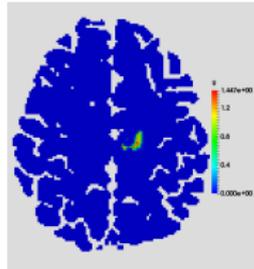
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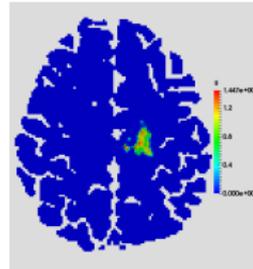
Therapy



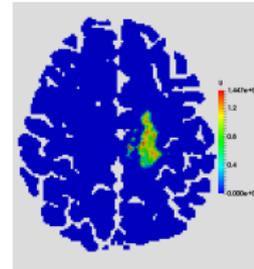
day 0, FA



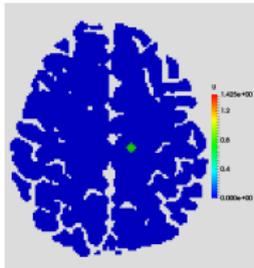
day 200, FA



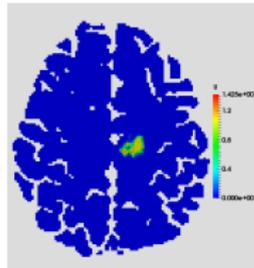
day 400, FA



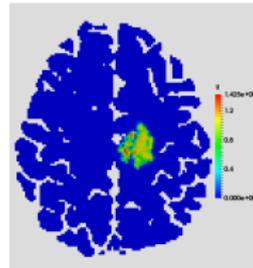
day 600, FA



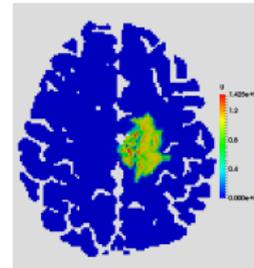
day 0, Q



day 200, Q



day 400, Q



day 600, Q

# Simulation results in 3D (with estimated $Q$ and peanut)

Multiscale models for glioma invasion: proliferation and therapy aspects

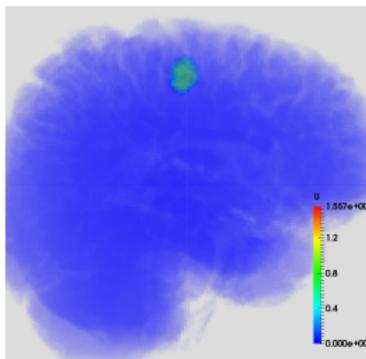
Christina Surulescu

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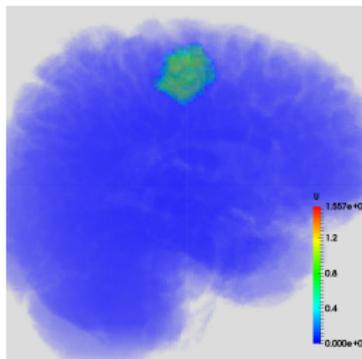
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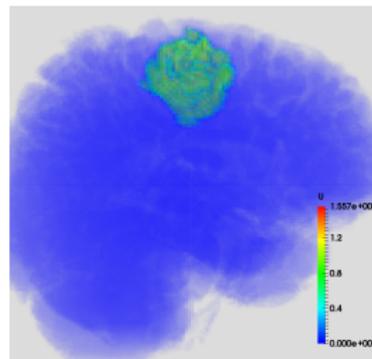
Therapy



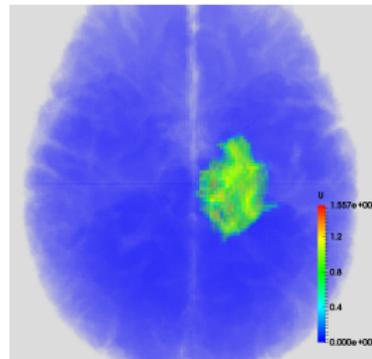
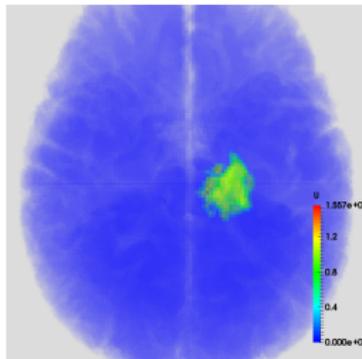
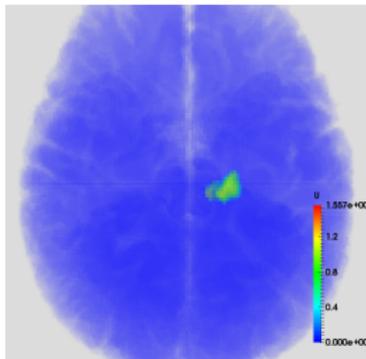
day 200



day 400



day 600



# Comparison with a pure macroscopic model

Multiscale models for glioma invasion: proliferation and therapy aspects

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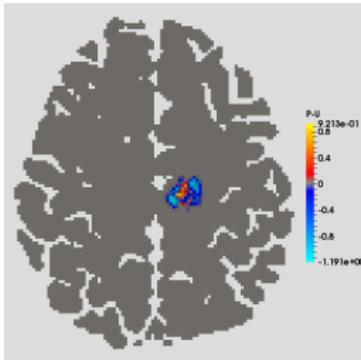
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A micro-meso setting with proliferation via go-or-grow

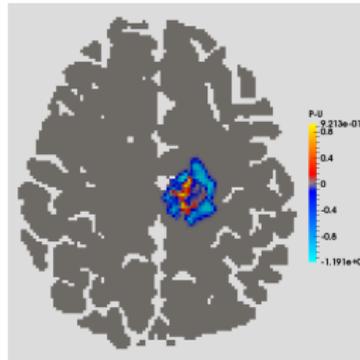
Proliferation via cell-tissue interactions

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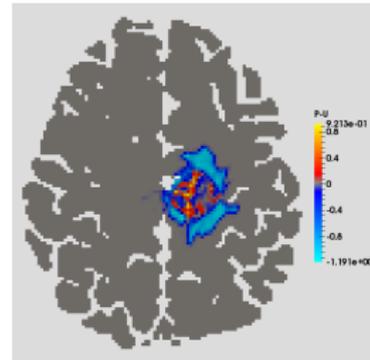
$$\partial_t M_0 - \nabla \cdot (\mathbb{D}_T(\mathbf{x}) \nabla M_0) = Q(\mathbf{x}) \mu(M_0) M_0.$$



day 200, Q



day 400, Q



day 600, Q

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Therapy

$$\begin{aligned} & \partial_t p + \nabla_{\mathbf{x}} \cdot (\mathbf{v}p) + \partial_y (G(y, Q, d_c, d_r)p) \\ & = \mathcal{L}[\lambda(y)]p - a(\mathbf{x}, d_c)p + b(\mathbf{x}, d_c) \frac{q(\hat{\mathbf{v}})}{\omega} r - L_1(N, \alpha_1, d_r)p \end{aligned}$$

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$$\partial_t r = a(\mathbf{x}, d_c) \int_V p(\mathbf{v}) d\mathbf{v} - b(\mathbf{x}, d_c)r + g(N, d_c)r - L_2(N, \alpha_2, d_r)r$$

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$$\partial_t r = a(\mathbf{x}, d_c) \int_{\mathcal{V}} p(\mathbf{v}) d\mathbf{v} - b(\mathbf{x}, d_c)r + g(N, d_c)r - L_2(N, \alpha_2, d_r)r$$

$$\text{with } L_l(N, \alpha_l, d_r) := \ell_l(N) + R_l(\alpha_l, d_r) \quad (l = 1, 2).$$

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$$\dot{y} = G(y, Q, d_c, d_r) = k^+(d_c)(R_0 - y)Q S(\alpha_3, d_r) - k^-(d_c)y.$$

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$$\dot{y} = G(y, Q, d_c, d_r) = k^+(d_c)(R_0 - y)Q S(\alpha_3, d_r) - k^-(d_c)y.$$

$$R_j(\alpha_j, d_r) = \sum_{i=1}^{\nu} (1 - S(\alpha_j, d_r)) \eta_{\delta}(t - t_i), \quad t_i \in \text{radiotherapy},$$

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$$\partial_t r = a(\mathbf{x}, d_c) \int_V p(\mathbf{v}) d\mathbf{v} - b(\mathbf{x}, d_c)r + g(N, d_c)r - L_2(N, \alpha_2, d_r)r$$

with  $L_l(N, \alpha_l, d_r) := \ell_l(N) + R_l(\alpha_l, d_r)$  ( $l = 1, 2$ ).

$$\dot{y} = G(y, Q, d_c, d_r) = k^+(d_c)(R_0 - y)Q S(\alpha_3, d_r) - k^-(d_c)y.$$

$$R_j(\alpha_j, d_r) = \sum_{i=1}^{\nu} (1 - S(\alpha_j, d_r)) \eta_{\delta}(t - t_i), \quad t_i \in \text{radiotherapy},$$

$$\text{supp } \eta_{\delta} \subset (-\delta, \delta), \quad \delta \ll 1, \quad j = 1, 2, 3.$$

$$\begin{aligned} \partial_t p + \nabla_{\mathbf{x}} \cdot (\mathbf{v}p) + \partial_y (G(y, Q, d_c, d_r)p) \\ = \mathcal{L}[\lambda(y)]p - a(\mathbf{x}, d_c)p + b(\mathbf{x}, d_c) \frac{q(\hat{\mathbf{v}})}{\omega} r - L_1(N, \alpha_1, d_r)p \end{aligned}$$

$$\partial_t r = a(\mathbf{x}, d_c) \int_V p(\mathbf{v}) d\mathbf{v} - b(\mathbf{x}, d_c)r + g(N, d_c)r - L_2(N, \alpha_2, d_r)r$$

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Survival fractions (LQ model):

$$S(\alpha_j, d_r) = \exp(-\nu(\alpha_j \hat{d}_r + \beta_j \hat{d}_r^2)) = \exp(-\alpha_j d_r (1 + \hat{d}_r / (\alpha_j / \beta_j)))$$

Multiscale  
models for  
glioma  
invasion:  
proliferation  
and therapy  
aspects

Christina  
Surulescu

Motivation

A micro-meso  
setting with  
proliferation  
via go-or-grow

Proliferation  
via cell-tissue  
interactions

Therapy

$$\text{Remember } N(t, \mathbf{x}) = \int_V \int_Y p(t, \mathbf{x}, \mathbf{v}, y) dy d\mathbf{v} + \int_Y r(t, \mathbf{x}, y) dy.$$

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$$\begin{aligned} \partial_t N_0 - \nabla \cdot \left( \frac{1}{\lambda_0 + a} \nabla \cdot \left( \frac{b}{a + b} \mathbb{D}_{\mathcal{T}(\mathbf{x})} N_0 \right) \right) \\ + \nabla \cdot \left( \frac{\lambda_1 f'(Q)}{\gamma(\mathbf{x})} \frac{b}{(\lambda_0 + a)(b + a)} \mathbb{D}_{\mathcal{T}(\mathbf{x})} \nabla Q N_0 \right) \\ = \left( (g(N_0) - L_2(N_0)) \frac{a}{a + b} - L_1(N_0) \frac{b}{a + b} \right) N_0, \end{aligned}$$

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where  $\gamma(\mathbf{x}) := k^+ Q S + k^- + \lambda_0 + a$ .

- **Strategy 1:** resection (2 weeks after start), no further therapy.
- **Strategy 2:** resection (2 weeks after start), followed after 3 weeks by radiotherapy (weekends excluded) for 6 weeks.
- **Strategy 3:** resection (2 weeks after start), followed after 3 weeks by concurrent chemotherapy and radiotherapy (weekends excluded) for 6 weeks.

Multiscale models for glioma invasion: proliferation and therapy aspects

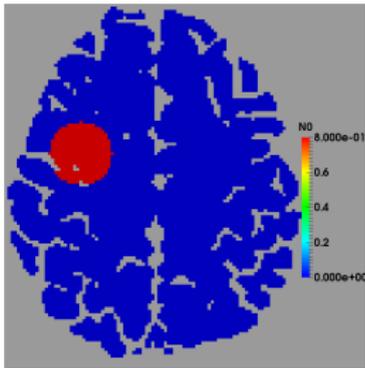
Christina Surulescu

Motivation

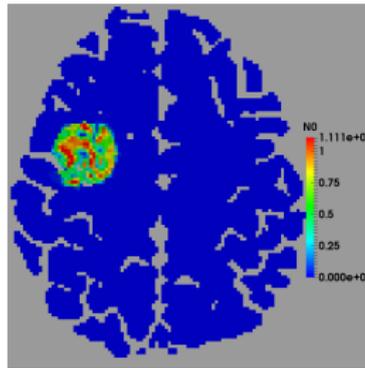
A micro-meso setting with proliferation via go-or-grow

Proliferation via cell-tissue interactions

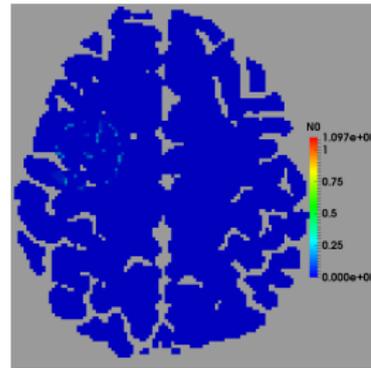
Therapy



start



before resection



after resection

# Results

Multiscale models for glioma invasion: proliferation and therapy aspects

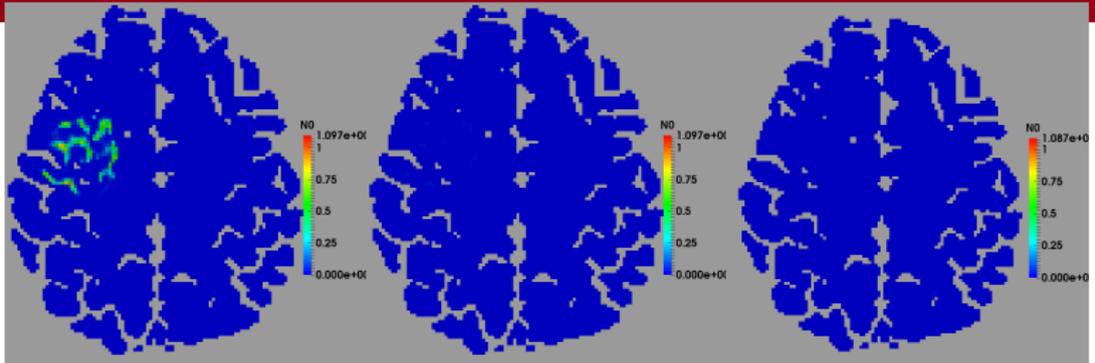
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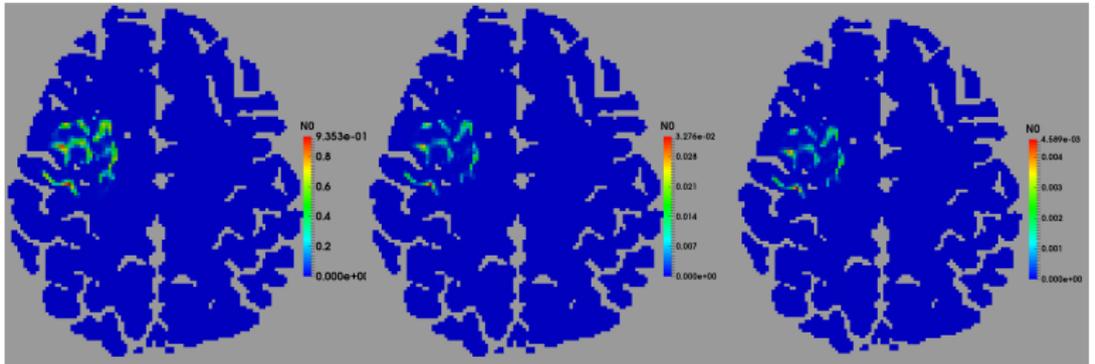
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Therapy



end of therapy



end of therapy, scaled

# Results

Multiscale models for glioma invasion: proliferation and therapy aspects

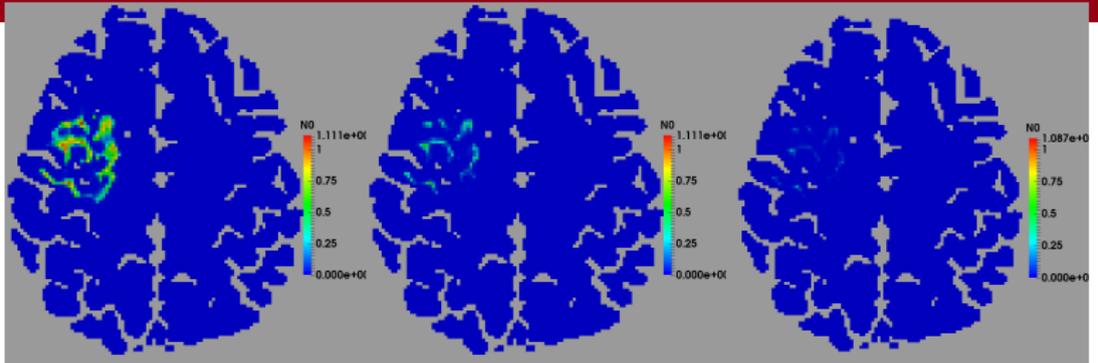
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Motivation

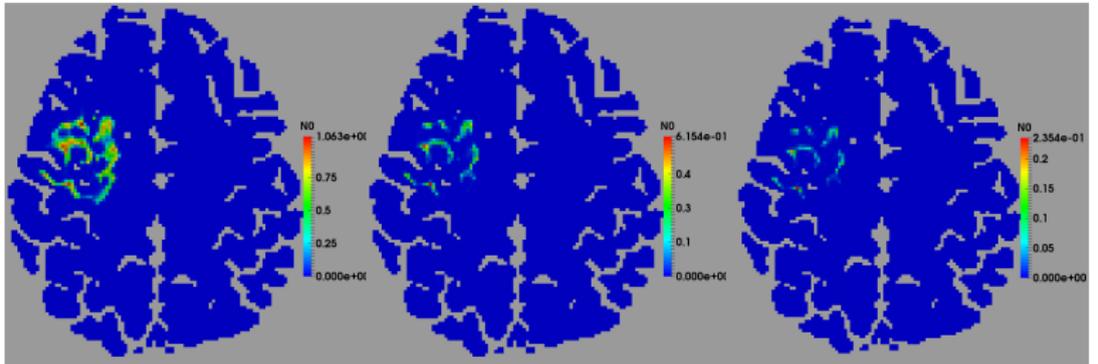
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follow-up after two months



follow-up, scaled

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## Multiscale models:

- allow testing the influence of many factors;
- are more difficult to handle numerically and analytically:
  - high dimensionality;
  - different scales both w.r.t. space and time;
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**Outlook:** tissue degradation, **therapy** Meral, Stinner & S. (JCS Multiscale Dyn. 2015) .

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## INI Isaac Newton Institute for Mathematical Sciences

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### New Mathematical and Computational Problems involved in Cell Motility, Morphogenesis and Pattern Formation

Isaac Newton Institute for Mathematical Sciences

Science

Scientific Programmes

Coupling Geometric PDEs with Physics for Cell Morphology, Motility and Pattern Formation

Workshops and Other Events

**New Mathematical and Computational Problems involved in Cell Motility, Morphogenesis and Pattern Formation**

- > Overview
- > [Invited Speakers](#)
- > [Participants](#)

7th December 2015 to 11th December 2015

**Organisers:** Alan Champneys (*Bristol*), [John King](#) (*Nottingham*), John Mackenzie (*Strathclyde*) and Christina Surulescu (*TU Kaiserslautern*)

#### Workshop Theme

Cell motility, morphogenesis, and pattern formation are essential features of cell dynamics. The involved biochemical processes and biomechanical properties range from the intracellular level over cell surface dynamics, cell-cell and cell-tissue interactions up to the scale of cell population behaviour influencing organ formation and functioning.

Mathematical models handling biological events taking place on one or several such scales can provide a powerful framework to understand these phenomena, test experimentally suggested conjectures, and make predictions about the behaviour of the studied system. Current modelling approaches are often continuous, involving systems of partial differential equations of various kinds (e.g., reaction-diffusion-transport, taxis, kinetic transport, population balance), possibly coupled to ordinary, random, or stochastic differential equations. Furthermore, the so-called agent-based approaches (e.g., cellular automata, Potts models, etc.) characterize the behaviour of individual cells or intracellular particles in a discrete way, permitting rather detailed descriptions of motions, interactions etc. Yet other model types are hybrids between discrete and continuous descriptions. Applications include, but are not restricted to embryogenesis, tumour growth and invasion, wound healing, tissue bioengineering, biofilms, etc. The models lead to highly complex analytical and numerical problems, which often call for the development of new mathematical tools or for the enhancement of existing ones. At the same time recent mathematical developments for example in nonlinear waves and coherent structures, in solid mechanics and in dynamical systems theory can help shed light on generic mechanisms; as well as the biology providing challenges to the mathematical state of the art.

Therefore, the aim of this workshop is to bring together scientists working on these timely and challenging topics of mathematical biology, analysis and numerics. It will provide both an international framework and motivation to further develop the modelling of the mentioned biological phenomena and to strengthen the synergies between the involved branches of applied mathematics, but also between mathematics and life sciences.

**Deadline for applications: 27 September 2015**