

Personalized Growth Modeling of Brain Tumours

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Joint work with E. Konukoglu, E. Stretton, M. Lê,
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The Digital Patient

in vivo

Medical
Images
and
Bio-signals

Statistics

Geometry
Physics
Physiology
Cognition

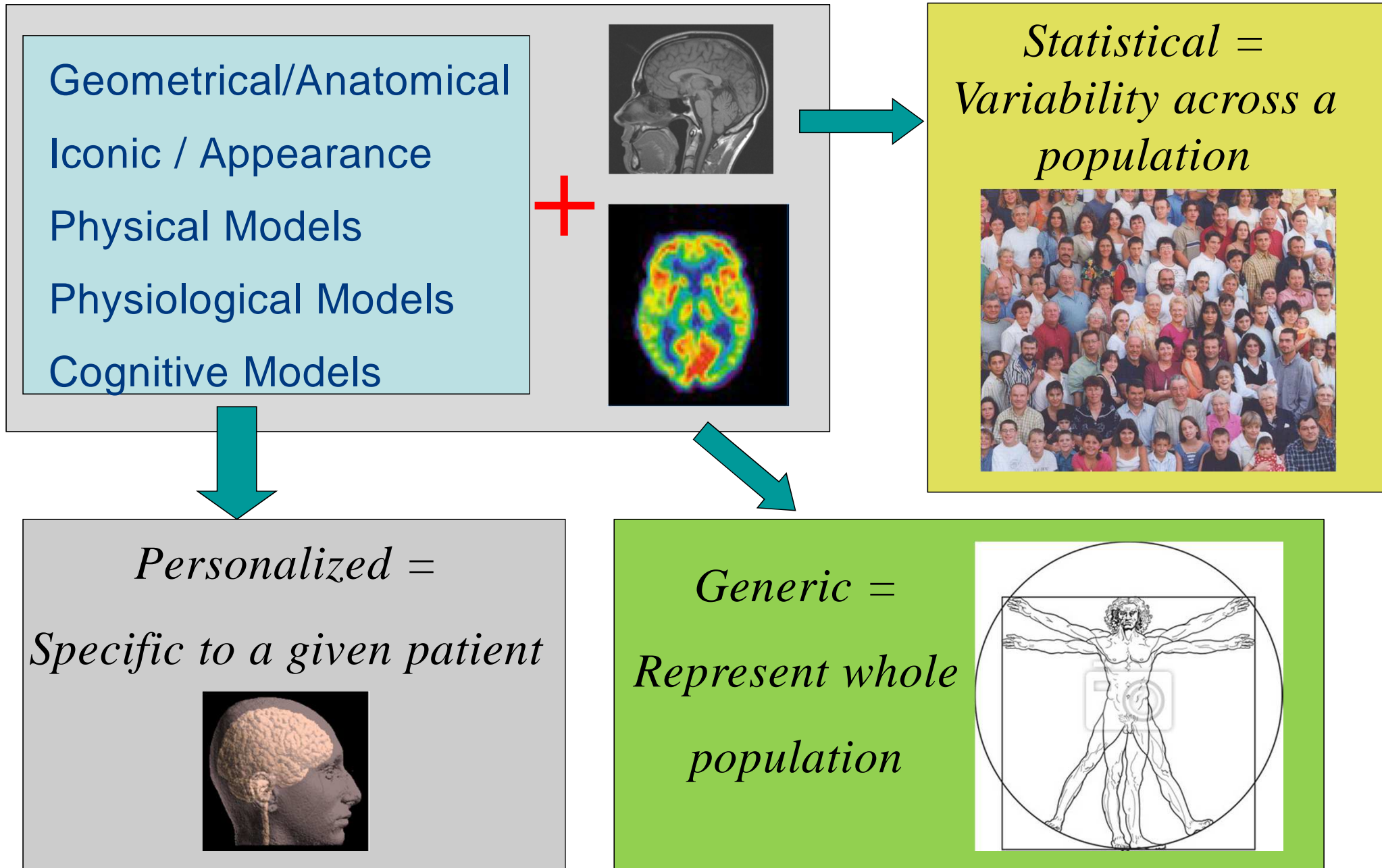
Computational
Models
&
Tools

in silico



Personnalisation

Generic, Patient Specific, Statistical Models

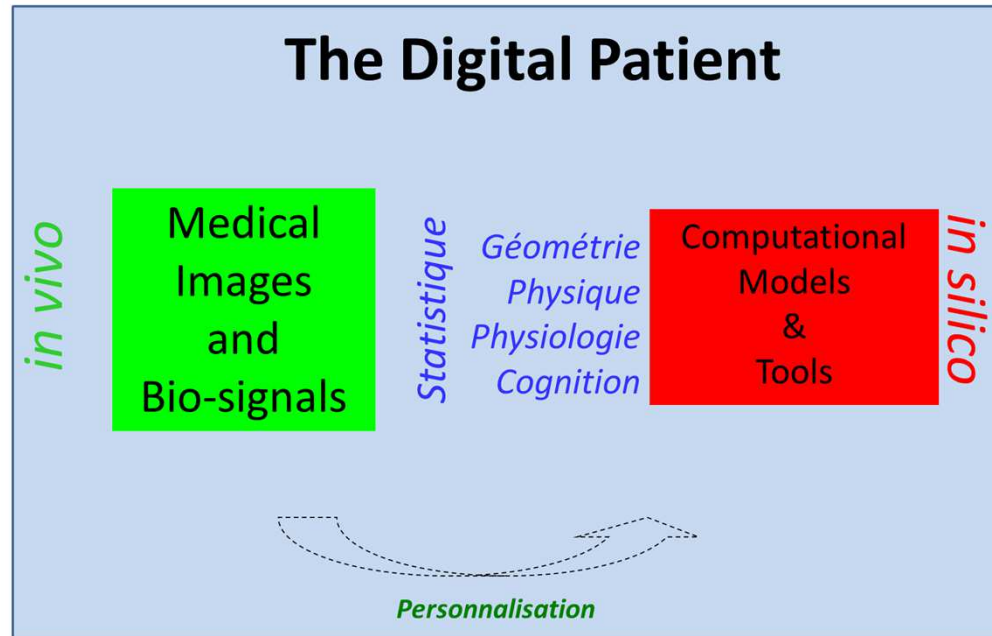


Applications of Digital Patient

*Image
Fusion*

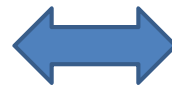
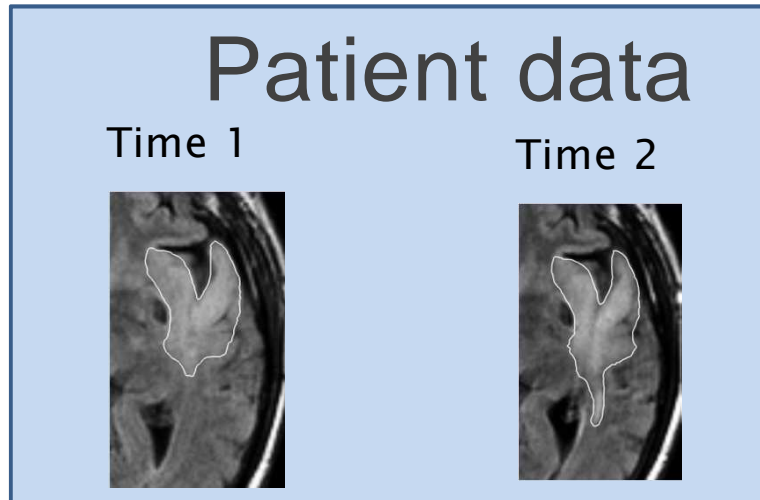
*Computer-Aided
Diagnosis*

*Therapy
Simulation*



*Therapy
Planning*

*Therapy
Guidance*



Why is the simulation different from the observation ?

Source of Errors

- Errors from the observation :
 - Noise & Artefacts
 - Errors from the computational model :
 - Computational domain (mesh)
 - Errors in the “parameters” : IC, BC
 - Errors in the implementation (bug)
 - Errors in the discretization (grid size)
 - Errors in the Model (False Hypothesis)
- Acquisition & Signal Processing Issue
- Personalization Issues
- Verification Issue
- Modeling Issue

Objectives of Model Personalization

- Model Validation (knowledge Building):
 - Model can represent observations ?
 - Yes -> can be tested for model prediction
 - No -> model should be modified
- Model Prediction
- Parameter analysis :
 - Parameter can be used for diagnosis

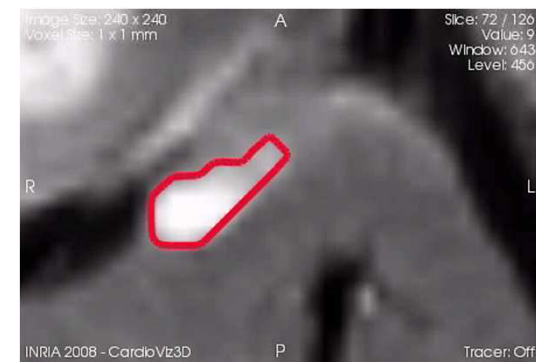
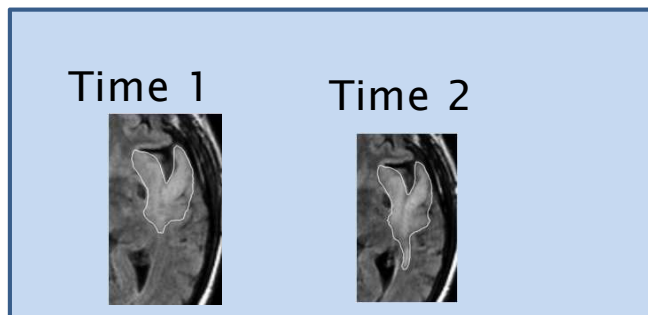
Data Regularization vs Model Personalization

- 2 sources of information :
 - Image Dataset provide observations
 - But may be noisy and partial
 - Biophysical Model based on the law of physics
 - But may not agree with the observation

**Medical
Images**

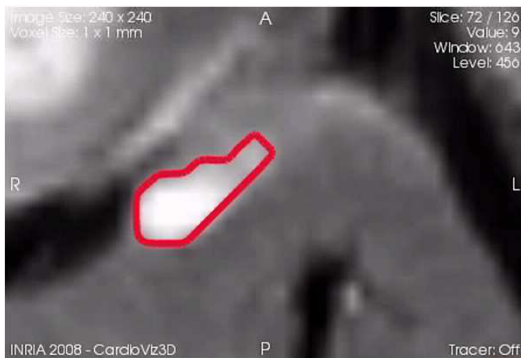
**Tumor Growth
Model**

How to combine both information ?



Data Regularization

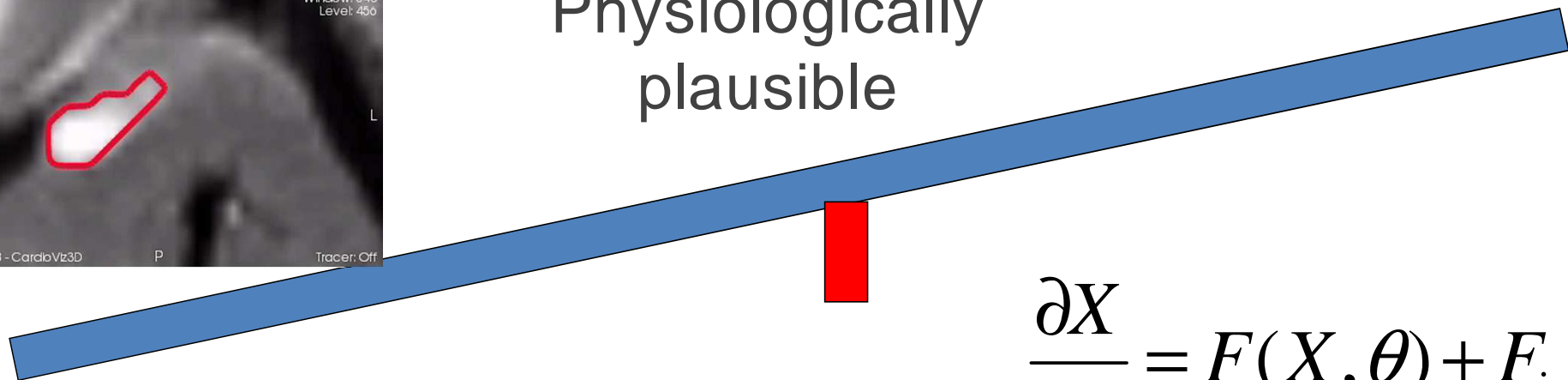
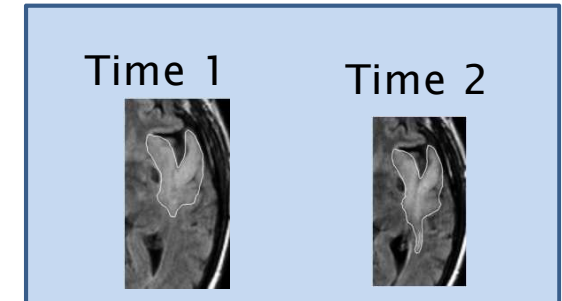
**Mostly Trust
The Data**



Data Regularization :

Filter

The data such that its
becomes
Physiologically
plausible



Data Regularization

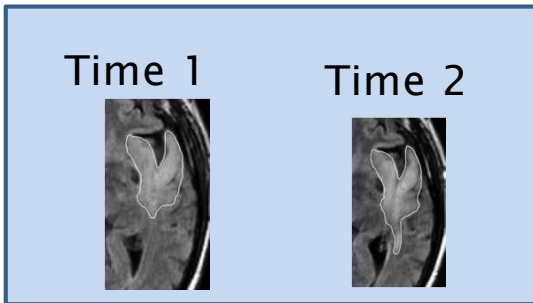
$$\underbrace{\frac{\partial X}{\partial t}}_{\text{Biophysical Model}} = \underbrace{F(X, \theta)}_{\text{Biophysical Model}} + \underbrace{F_{img}}_{\text{Image Force}}$$

Biophysical
Model

Image
Force

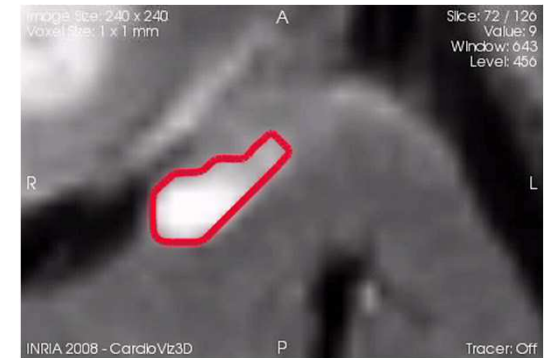
Model

Model Personalization



Model Personalization : Fit
Model Parameters, BC, IC
from data

**Mostly Trust
The Model**



Find θ^* such that

$$\frac{\partial X}{\partial t} = F(X, \theta^*) + F_{img}$$

Agrees with data

**Model
Personalization**

=

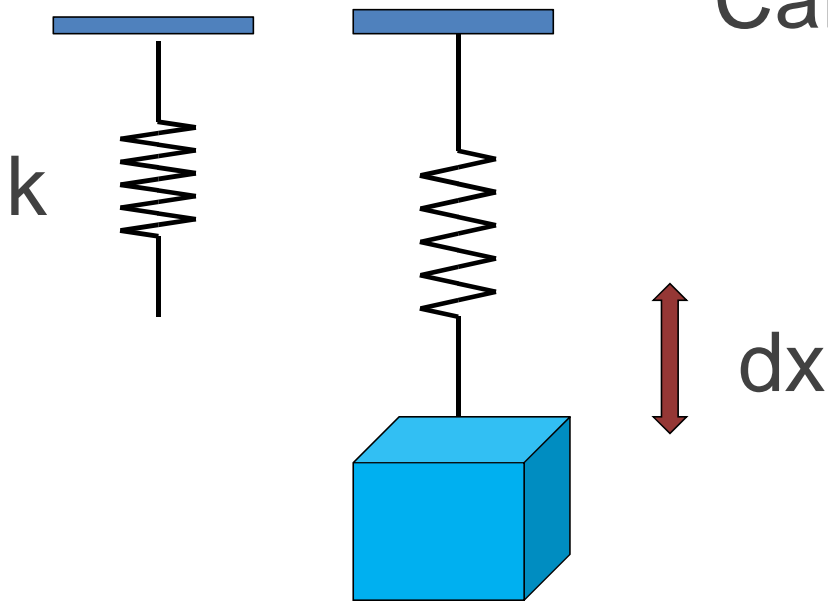
Data Assimilation

Parameter Estimation Issues

- Observability of the parameters
- Dimensionality of the parameters vs
Dimension of the observations
- Optimization Techniques

Parameter Observability

- Not all parameters can be estimated from observations

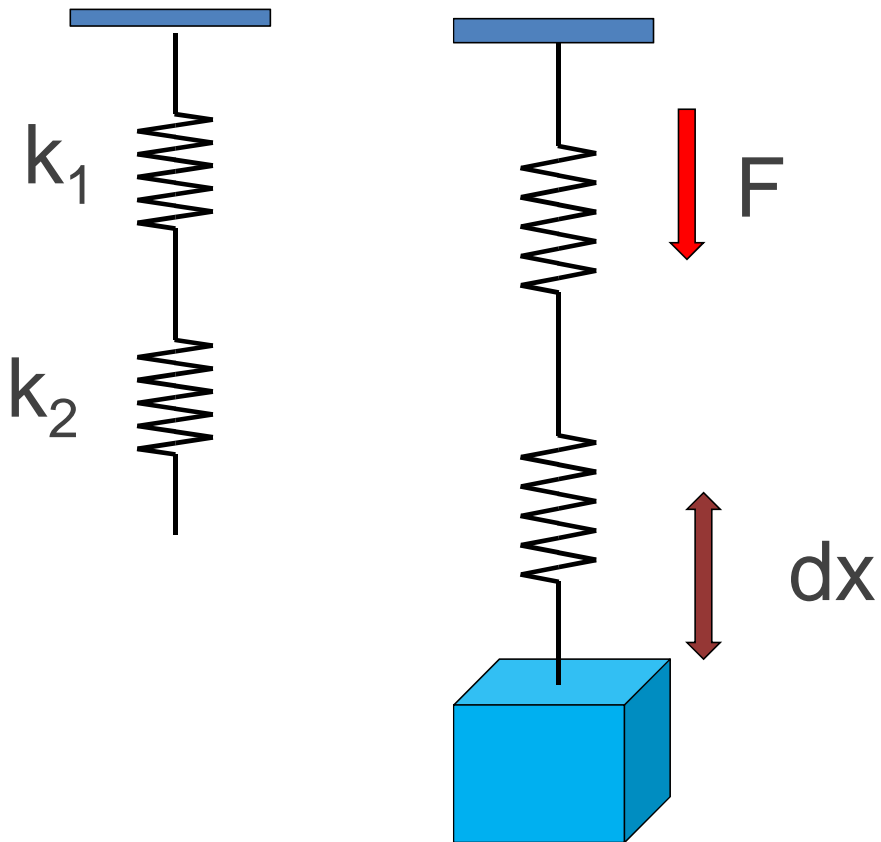


Cannot estimate spring stiffness k from dx !!

$$k = \frac{F?}{dx}$$

Parameter Observability

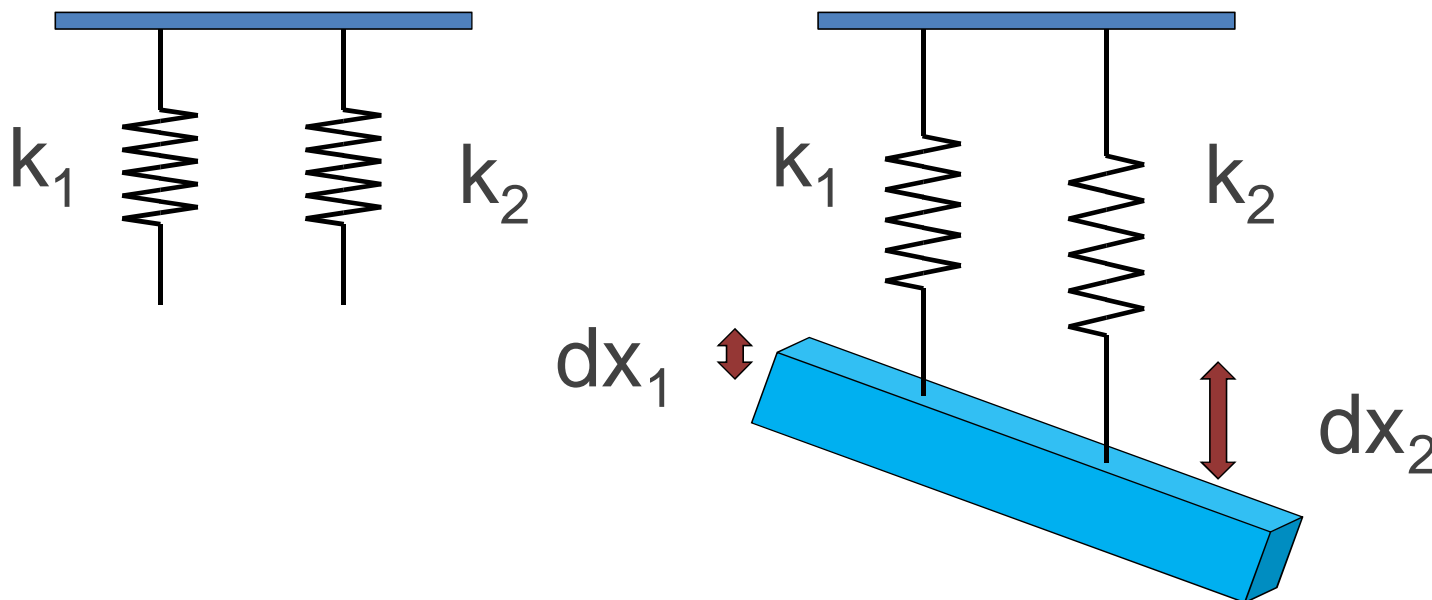
- Can estimate combination of parameters from observation



Only estimate
spring stiffness k_1+k_2
from dx and F !!

Parameter Observability

- Can estimate combination of parameters from observation



Can estimate the ratio of spring stiffness k_1/k_2 from displacements !!

Model as a tool

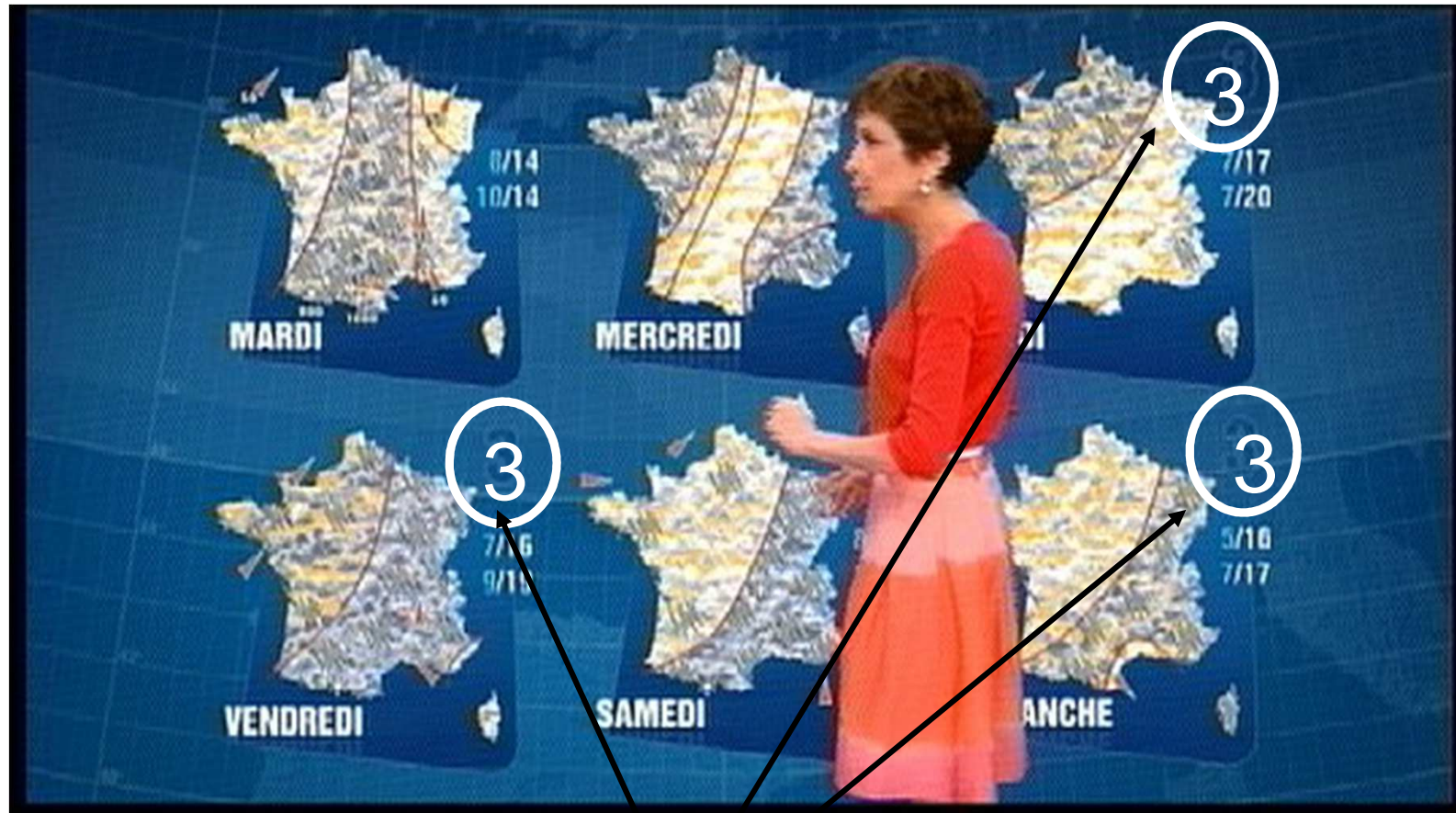
- Models should be designed to answer a given question
- Avoid overfitting of parameters :
 - Adapt model complexity to the complexity of the observations
- Follow Lex Parsimonia (Ockam's razor) : among all suitable models, select the most simple one
 - « The ideal model will be as simple as possible and as complex as necessary for the particular question raised. »

Model Selection

- A Model always fails “somewhat”:
 - Should estimate the uncertainty (covariance or posterior distribution) associated with the simulation
 - Estimate all source of errors
- How do I know a model “completely” fails ?:
 - Large discrepancy between observations & simulations after personalization
 - Large variability of parameters for subjects in similar conditions (or same subject)

Handling uncertainties

Example : Weather Forecast



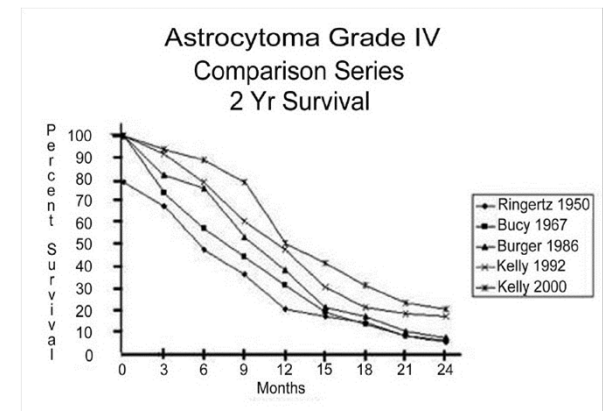
Confidence in
Weather Forecast

Glioma

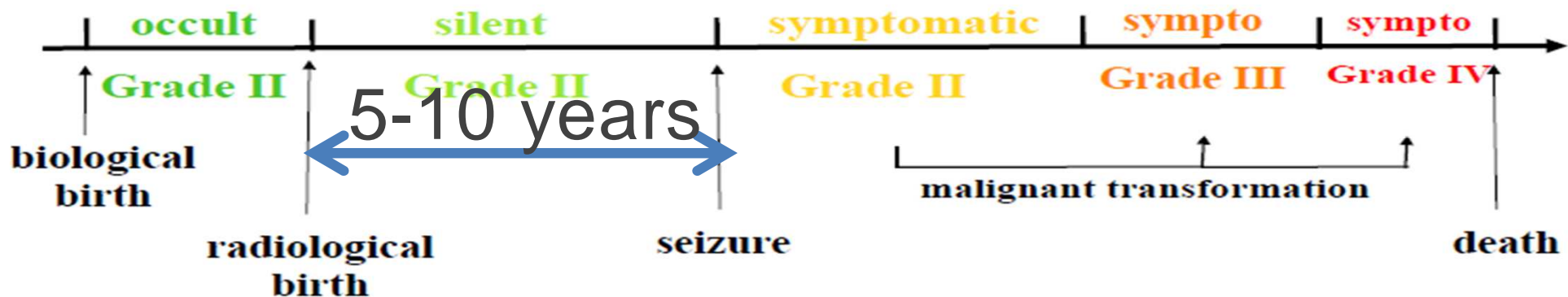
- Incidence: 5-6 cases /100.000 persons/year
 - Young adults: 3rd cause of death
 - Child: 2nd cause of cancer after leukemia
 - Peak for persons between 50-60 year old
- 80% of brain tumors
- Diagnosed with MRI & Clinics

Low Grade & High Grade Glioma

- Low Grade : Astrocytoma & Oligodendroglioma
 - Median survival 10 years
- High grade Glioma : Glioblastoma
 - Median survival : 16 months
 - No major improvement in treatment



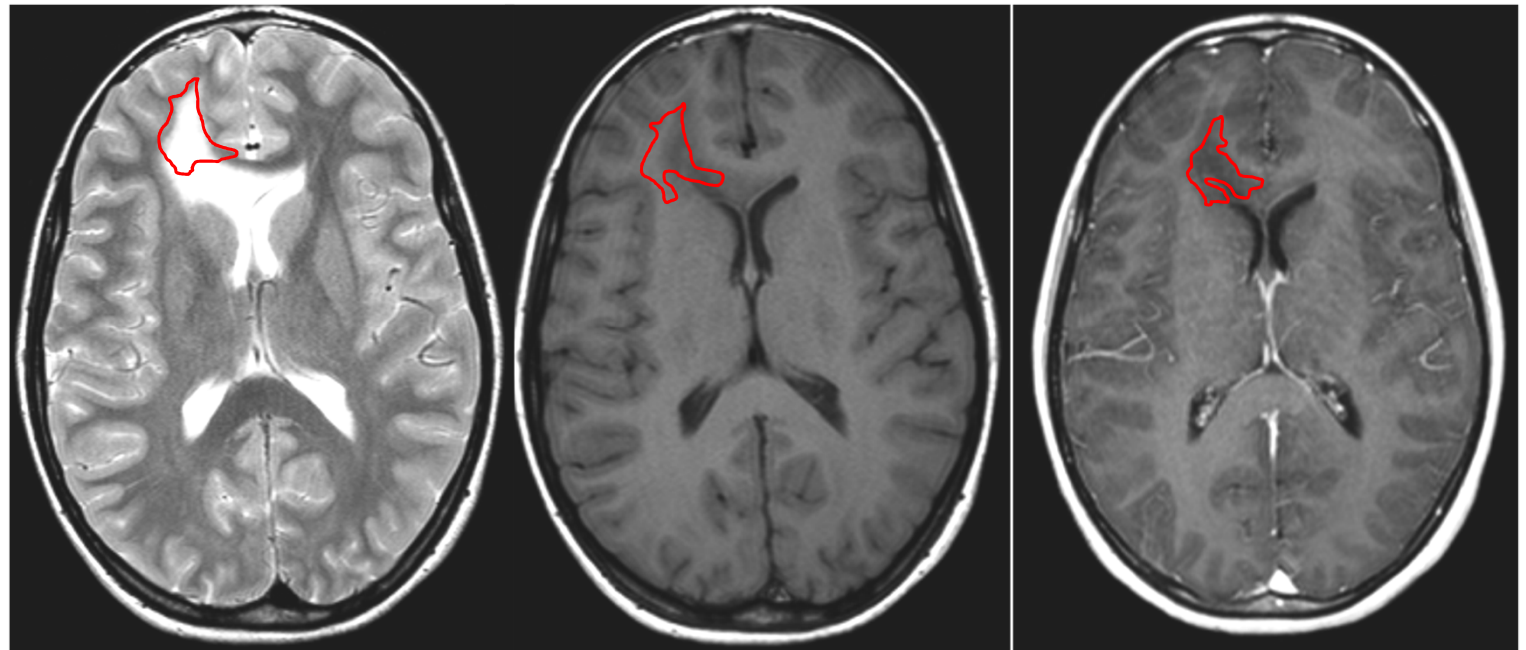
[P. Kelly Surg. Neur. Inter. 2013]



MR Sequences & Glial Tumors

- Low Grade Glioma

Legend:
+ *hyper-intense*
- *hypo-intense*
0 *no signal*



T2 / FLAIR

T1

T1+gad

Pathological
structures

edema

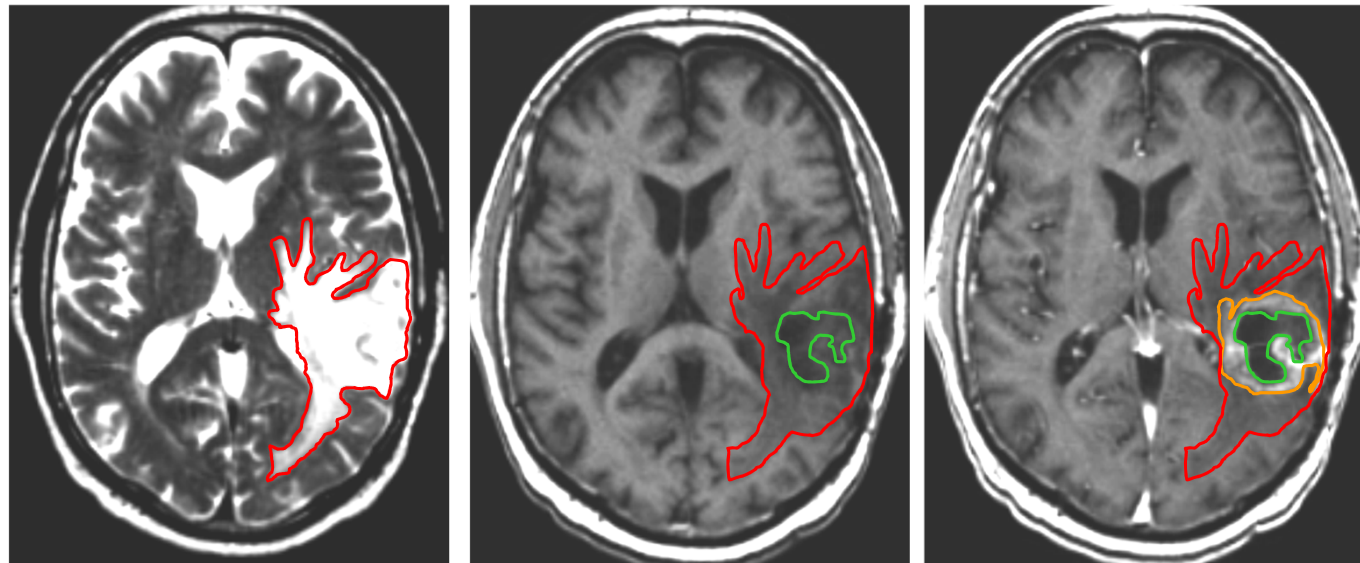
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MR Sequences & Glioma Tumors

- High Grade Glioma



Pathological structures

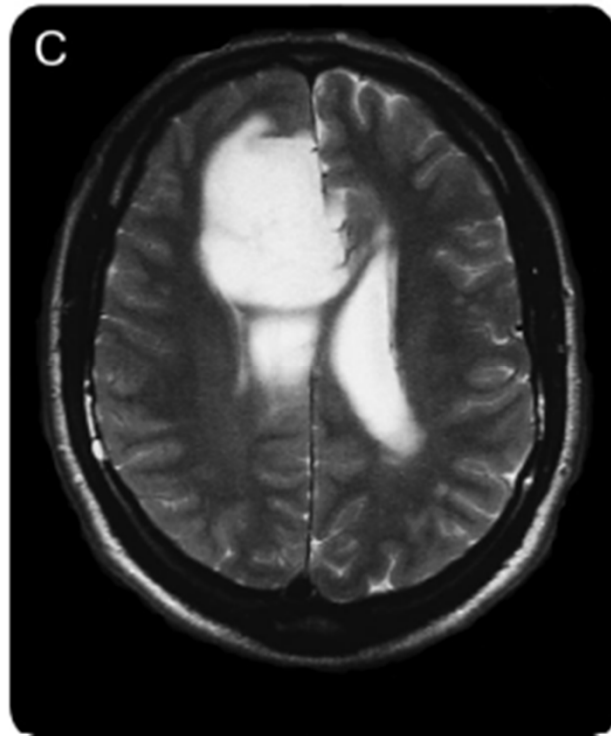
T2 / FLAIR

T1

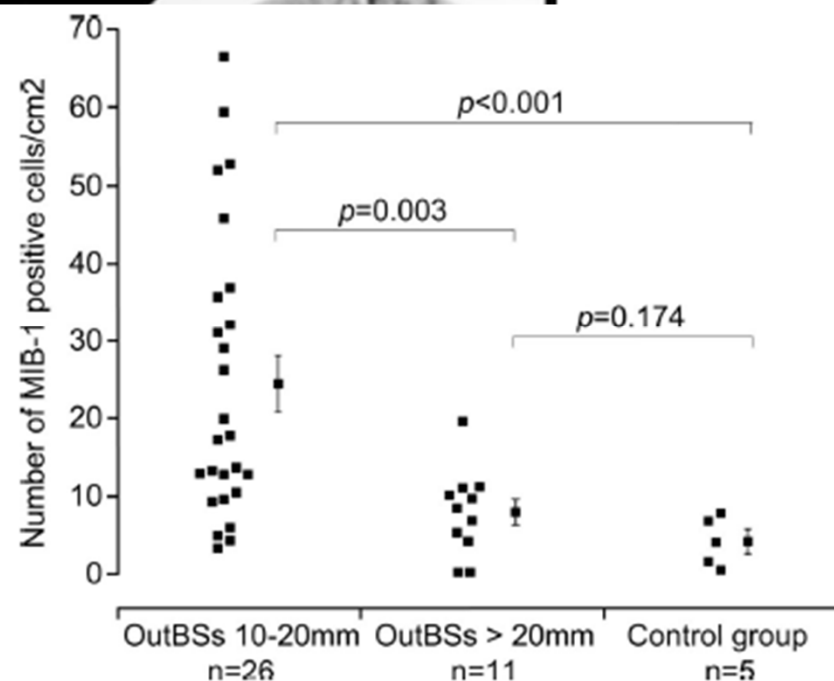
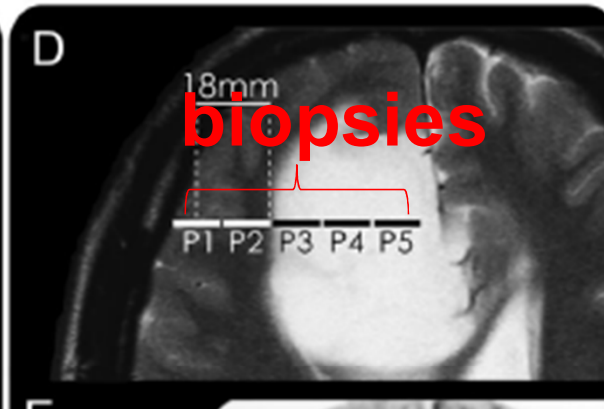
T1+gad

edema / infiltration	++	-	-
active region	++	-	++
necrotic core	++	--	--

Tumor extends beyond visible boundary



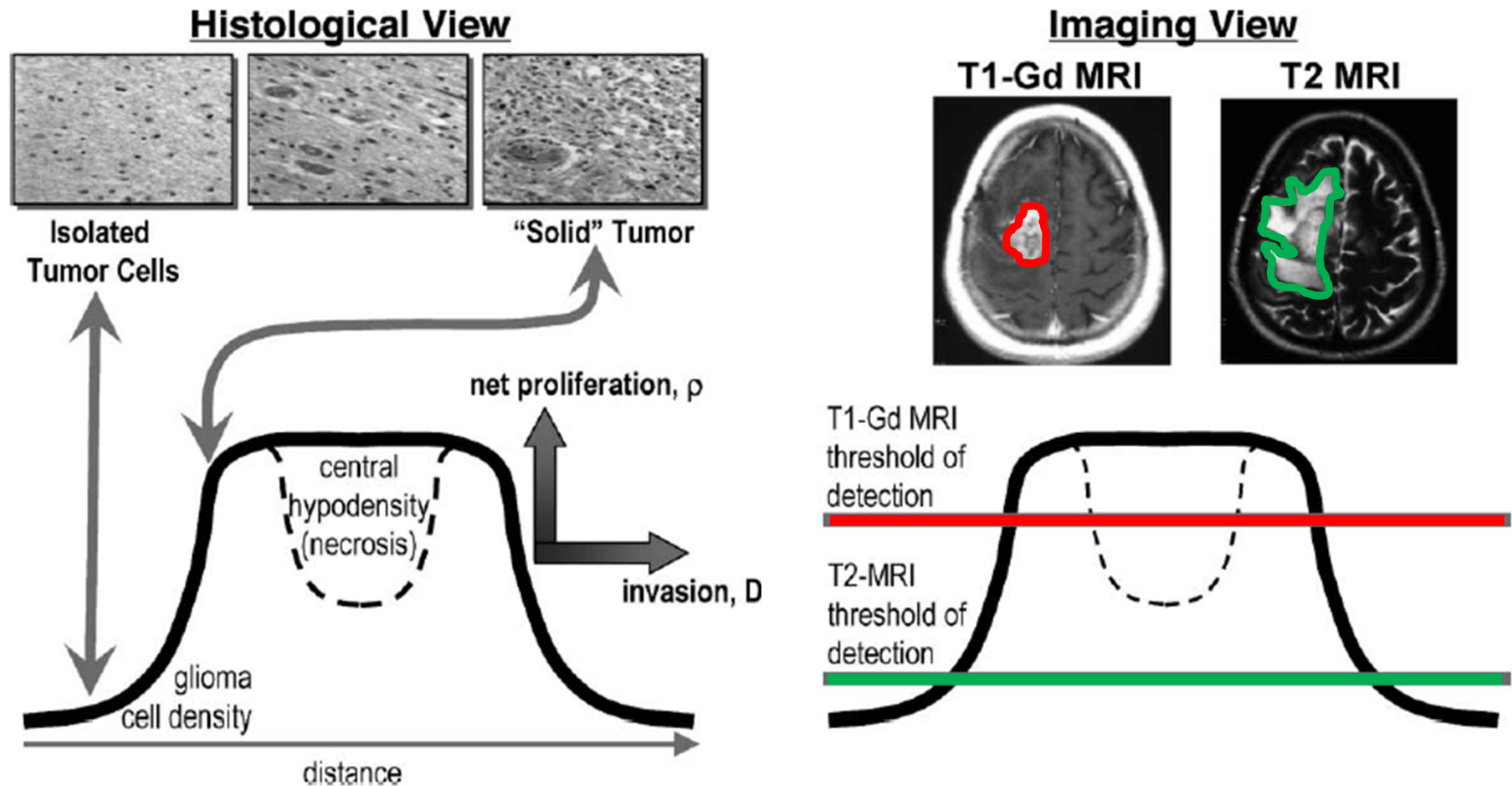
T2/ Flair



Pallud, J et al. "Diffuse Low-Grade Oligodendrogliomas Extend beyond MRI-Defined Abnormalities." *Neurology* 74.21 (2010): 1724–31.

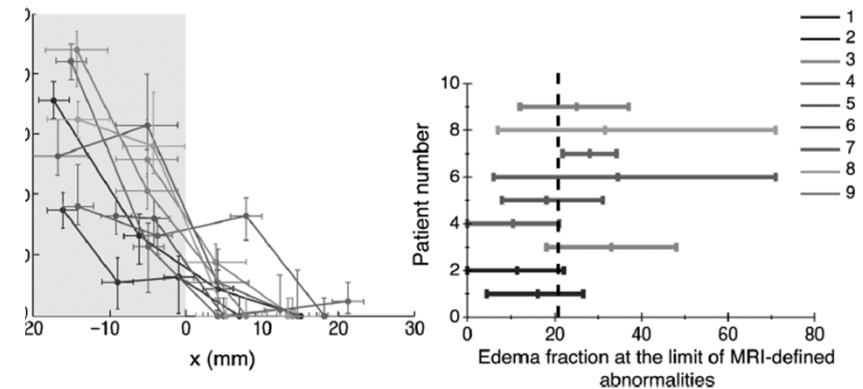
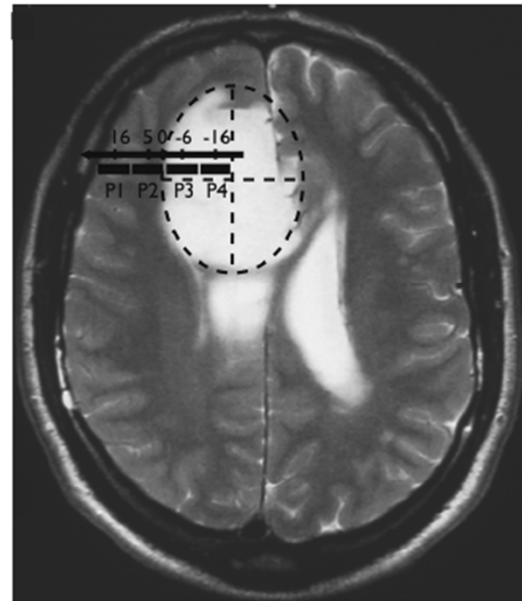
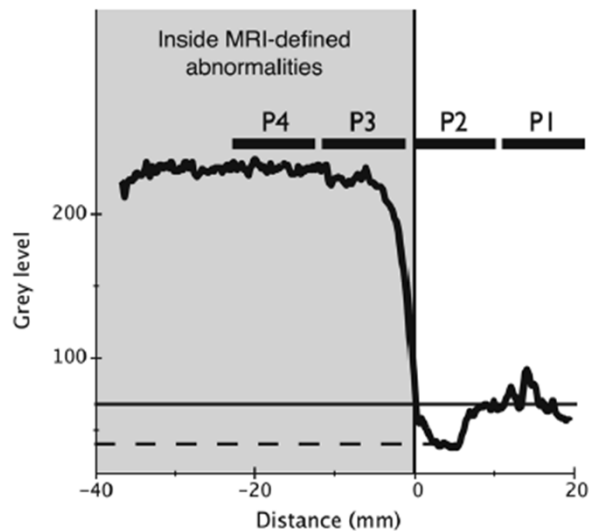
MR Sequences & Glioma Tumors

- Main Hypothesis :
visible contours = isodensity of tumor cells



MR Sequences & Glial Tumors

- This hypothesis may not be true !!



Visible Flair abnormality border correlates better with threshold on edema content

Possible Treatments

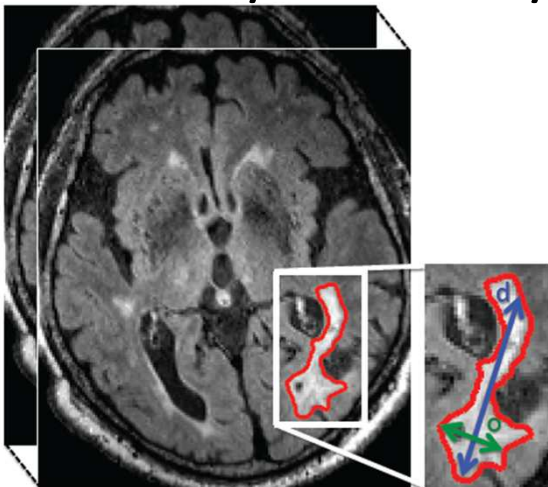
- Upon discovery :
 - Wait & watch (3-6 asymptomatic glioma / 1000 persons)
 - Surgery (awake surgery w functional mapping)
 - Radiotherapy (2cm margin)
 - Chemotherapy (anti-angiogenic drug)

3 Main Problems

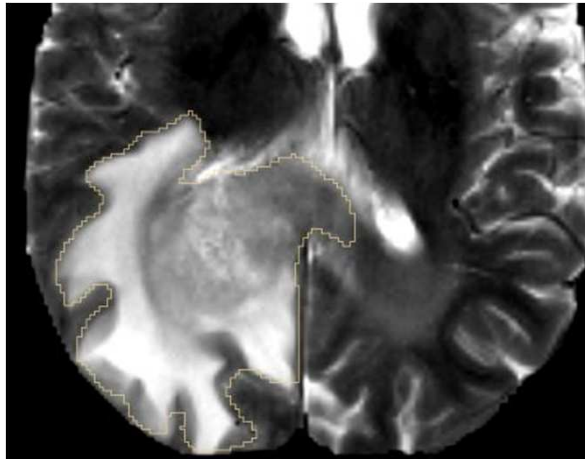
1. Quantify the extent of the tumor
2. Quantify the tumor evolution
3. Improve clinical practice

Quantification of tumor extent

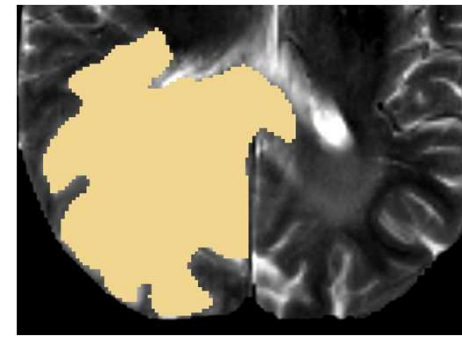
- Current clinical practice :
 - One or two largest diameter (RECIST or RANO criteria)
 - Only 1D or 2D
 - Do not differentiate between tumor compartments (core edema / white & grey matter)



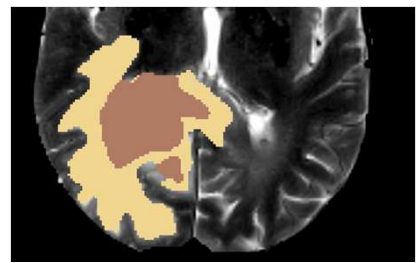
Hierarchy of compartments (Source BRATS)



High grade. T2-w.



tumor + edema
(T_2 -w, FLAIR)



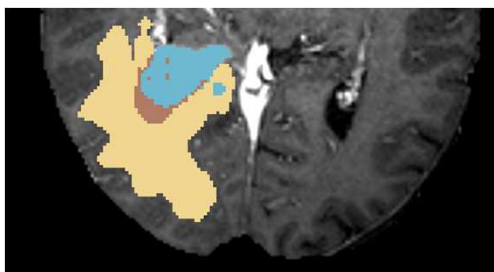
core
(abnormal in T_1 -w + Gad)

(and T_2 -w, for LG and some HG)

edema

enhancing core
(T_1 -w + Gad)

non-enhancing core



T1-w + Gad

necrotic core
(cyst)

the rest
(texture, localisation, etc.)

Automatic Segmentation ?

The Multimodal Brain Tumor Image Segmentation Benchmark (BRATS)

- Growing at

Source : B. Menze et al. "The Multimo

Jhoons H. Menze¹, Andras Jakab², Stefan Bauer³, Jayashree Kalpathy-Cramer⁴, Keyvan Farahani⁵, Justin Kirby⁶, Tulya Hazvan⁷, Nicole Poon¹, Johannes Steffensen², Erland Wass¹, Lovencio Lanius¹, Elizabeth Granger¹, Marc-Andre Weber¹, Tai Arbel, Brian H. Avants, Nicholas Ayache, Patricia Baerndt, D. Louis Collins, Nicolas Coufrier, James J. Corso, Antonio Criminisi, Tilak Dux, Hervé Delingette, Gajraj Durrani, Christopher R. Durr, Michel Dojat, Sean Doyle, Juana Feito, Florance Forbes, Huzefa Gonen, Ben Glocker, Polina Golland, Xinshao Guo, Andac Hamamci, Khan M. El-Hakim, Raj Jena, Nigel M. John, Ender Konukoglu, Daniel Lashkari, José Antonio Martí, Raphael Meier, Sergio Pereira, Debra Prasad, S. J. Price, Tammy Rakko-Raviv, Syed M. S. Rizvi, Michael Ryan, Lawrence Schwartz, Hoo-Chang Shin, Janik Shotton, Carlos A. Silva, Nuno Sousa, Nagesh K. Subhanna, Gabor Székely, Thomas J. Taylor, Owen M. Thomas, Nicholas J. Tustison, Grégoire Unal, Flor Vauzeur, Max Wistnermark, Dong Hyeon Yi, Liang Zhao, Hisheng Zhao, Darko Zikic, Marzi Prastawa¹, Mauricio Reyes⁸, Koen Van Leemput⁹

mentation

IEEE TMI 2014

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¹¹ N. Ayache, N. Coufrier, H. Delingette, and E. Granger are with Inria Sophia Antipolis, France.

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³⁶ These authors contributed equally to the work.

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- BRATS Chal

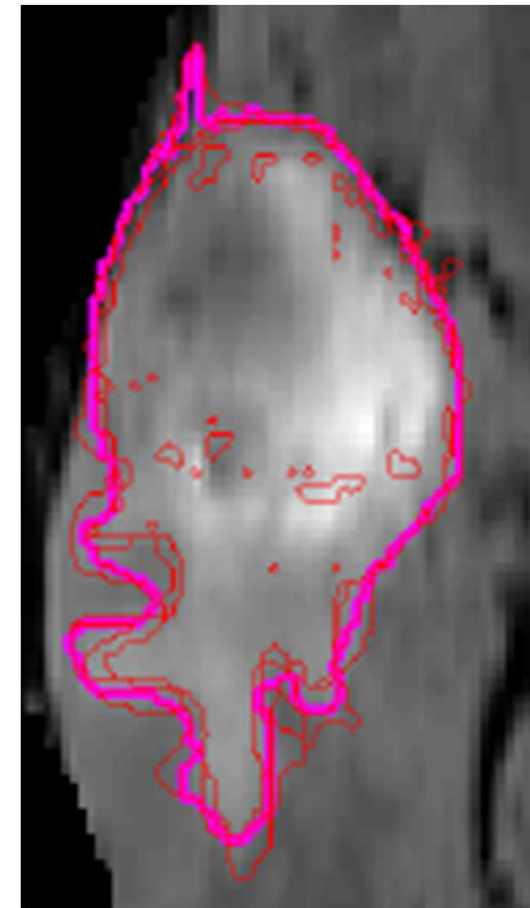
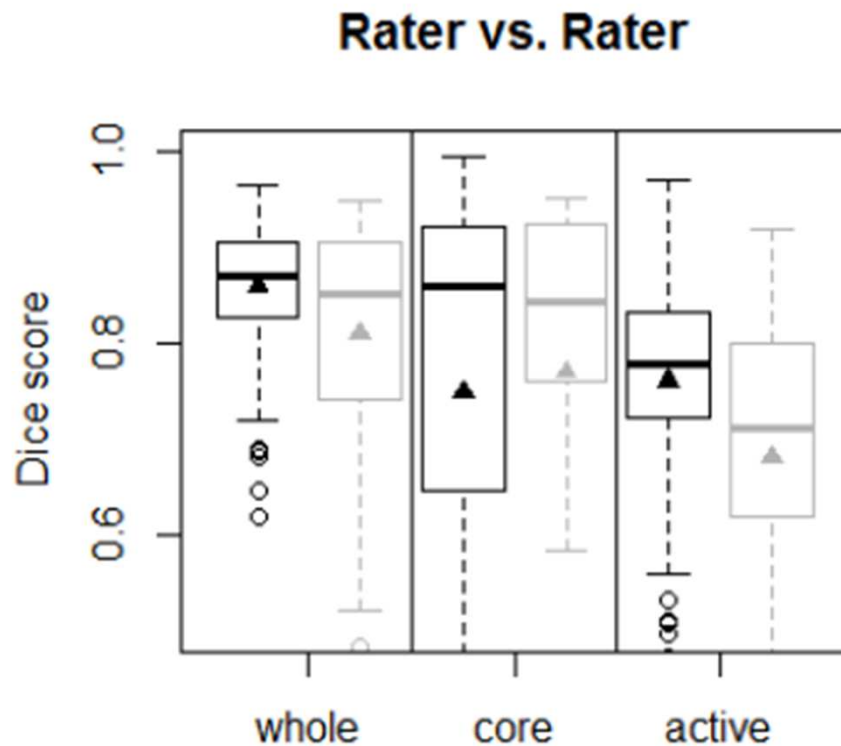
K. Farani) runi

lpathy-Cramer,

A difficult segmentation task

- Fairly large discrepancy between experts

— Single rater
— Consensus rater



Flair MR

Segmentation approaches

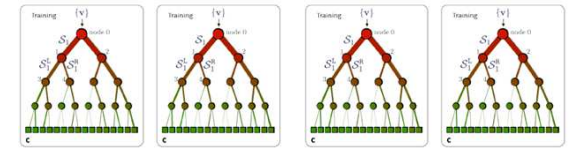
- generative or discriminative
- 2 survey papers

S. Bauer, R. Wiest, L.-P. Nolte, and M. Reyes, “A survey of MRI based medical image analysis for brain tumor studies,” *Phys Med Biol*, vol. 58, no. 13, pp. R97–R129, Jul. 2013.

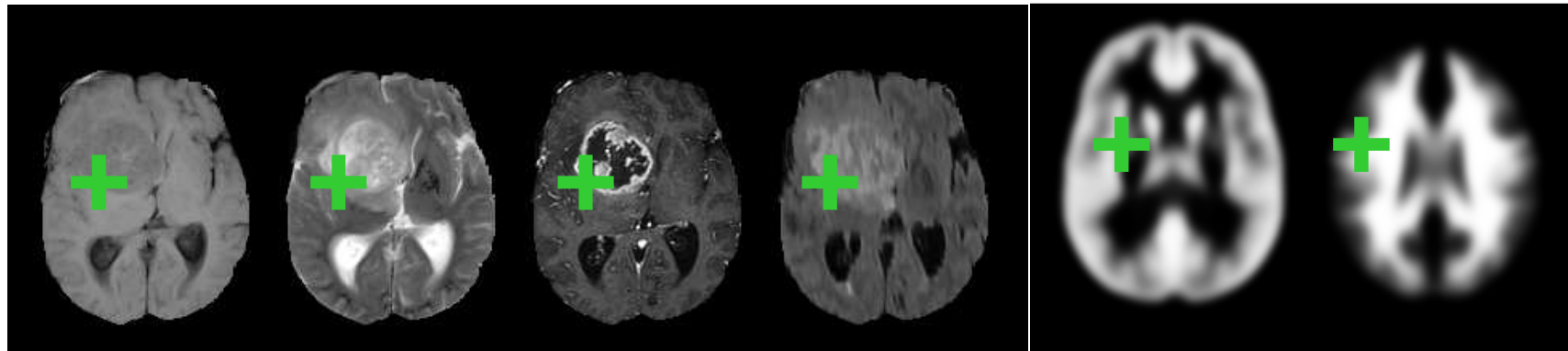
Angelini E., Clatz O., Mandonnet E., Konukoglu E., Capelle L. and Duffau H. , *Glioma Dynamics and Computational Models: A review of Segmentation, Registration and In Silico Growth Algorithms and their Clinical Applications*, Current Medical Imaging Reviews, 2007

Examples of segmentation method

- Random Forest: Features



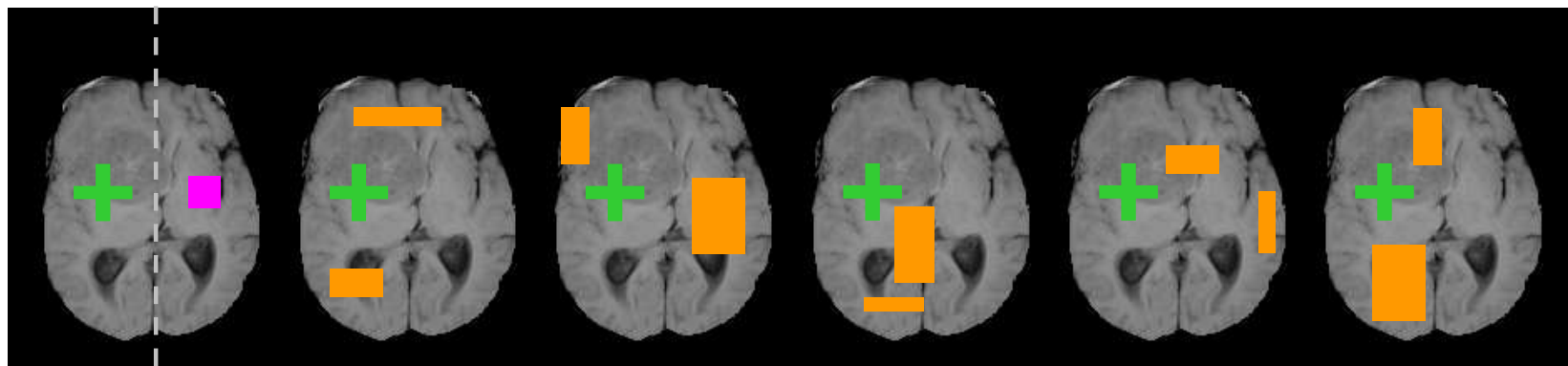
Local



Multi-channel MRIs
T1, T2, T1+Gd, Flair

Spatial priors
GM, WM

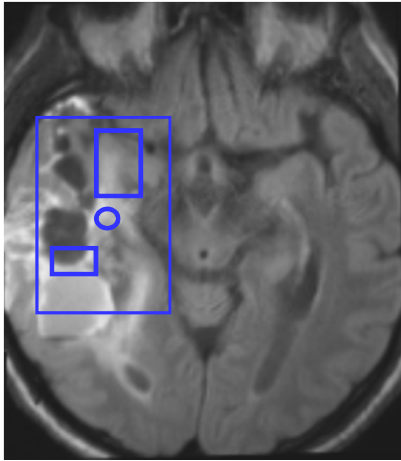
Contextual



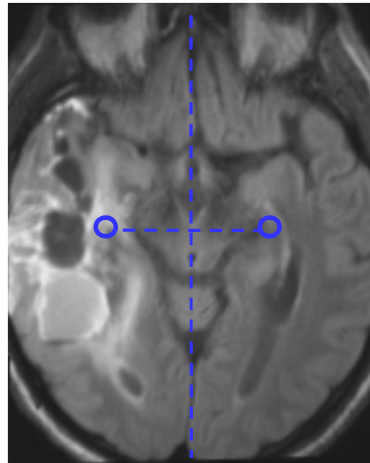
asymmetry

Randomized context-rich features

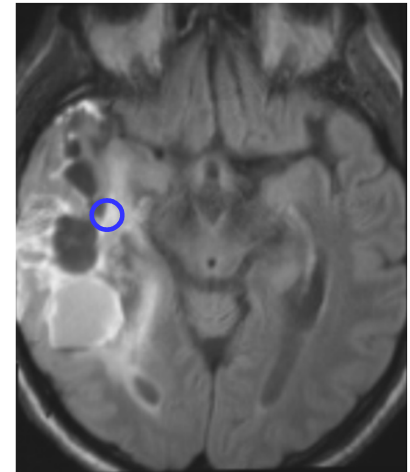
Selected Features



41 %
Contextual
features

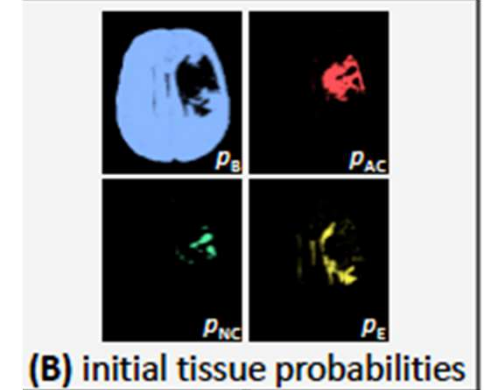
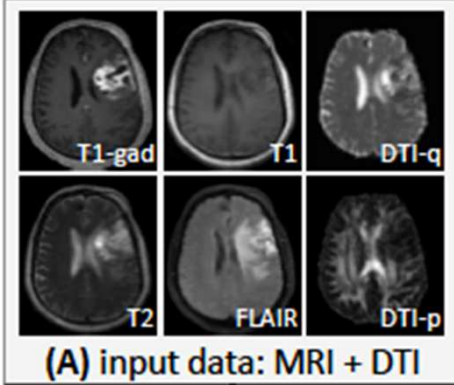


36 %
Asymmetry



23 %
Local Features

Typical Results



Input Image							
Ground Truth							
Random Forest							

Necrotic Core

Proliferating Cells

Edema

Parenchyma

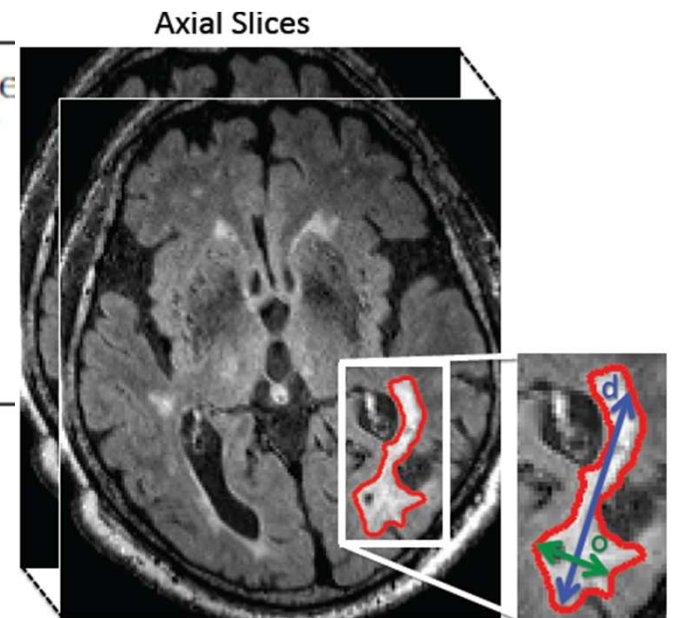
E. Geremia, D. Zikic, O. Clatz, B.H. Menze, B. Glocker, E. Konukoglu, J. Shotton, O.M. Thomas, S.J. Price, T. Das, R. Jena, N. Ayache, and A. Criminisi, *Classification Forests for Semantic Segmentation of Brain Lesions in Multi-Channel MRI*, in Decision Forests for Computer Vision and Medical Image Analysis, Springer, 2013

3 Main Questions

1. Quantify the extent of the tumor
2. Quantify the tumor evolution
3. Improve clinical practice

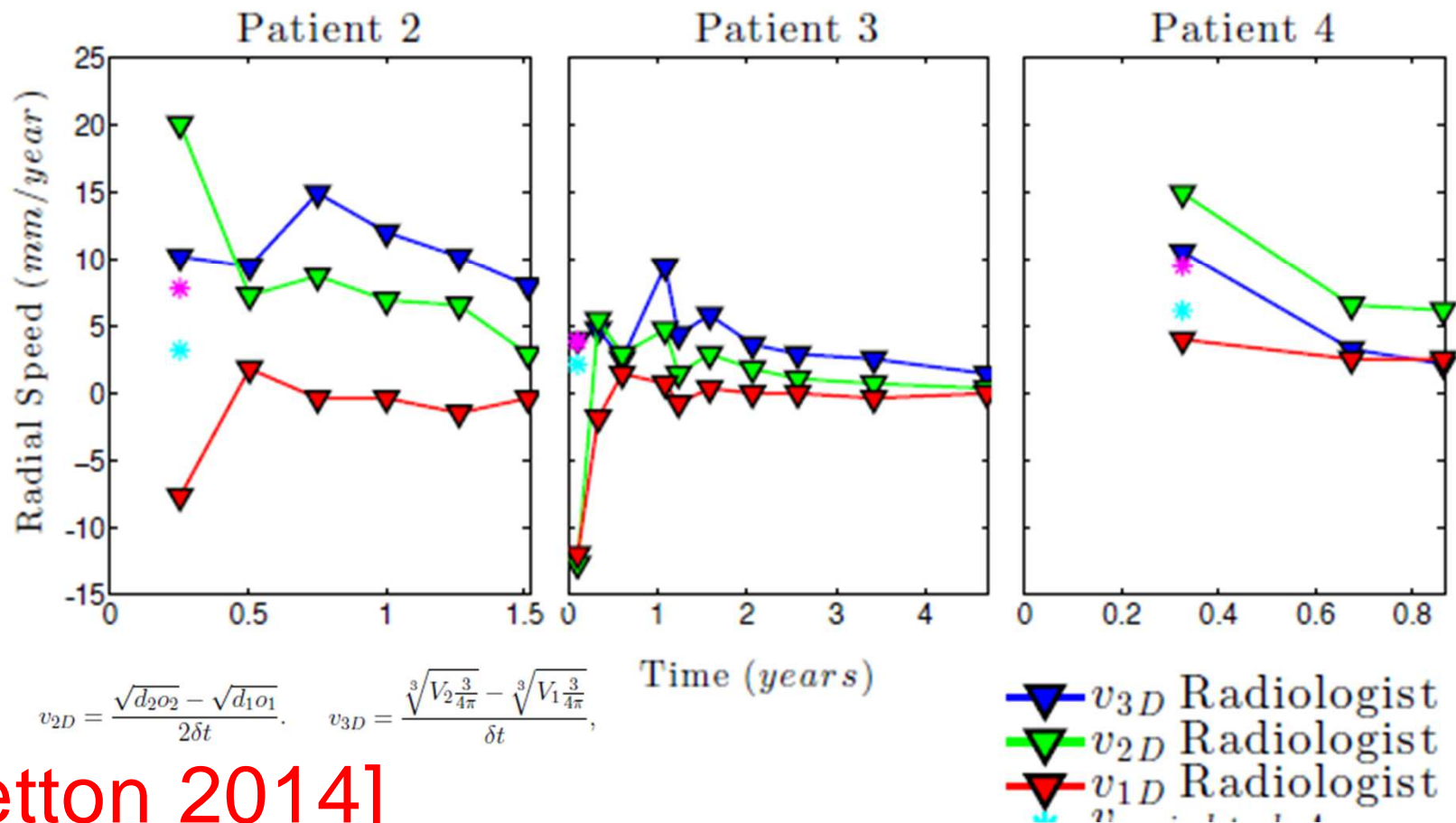
Quantification of tumor growth

<i>Imaging Modality</i>	<i>RECIST (1D)</i>	<i>Macdonald (2D)</i>	<i>RANO (2D)</i>
<i>T1-gad</i>	$\geq 20\%$ increase in sum of maximal diameters; confirm at 4 weeks	$\geq 25\%$ increase in product of orthogonal diameters; confirm at 4 weeks	$\geq 25\%$ increase in product of orthogonal diameters; confirm at 4 weeks
<i>T2/FLAIR</i>	$\geq 20\%$ increase in sum of maximal diameters; confirm at 4 weeks	N/A	Significant increase



Discrepancy between growth speed

- Growth rate based on 1D, 2D or 3D measurements from the same images



$$v_{1D} = \frac{d_2 - d_1}{2\delta t}, \quad v_{2D} = \frac{\sqrt{d_{2o2}} - \sqrt{d_{1o1}}}{2\delta t}, \quad v_{3D} = \frac{\sqrt[3]{V_{2\frac{3}{4\pi}}} - \sqrt[3]{V_{1\frac{3}{4\pi}}}}{\delta t}$$

[Stretton 2014]

Can we do better ?

Maybe using a personalized tumor growth model

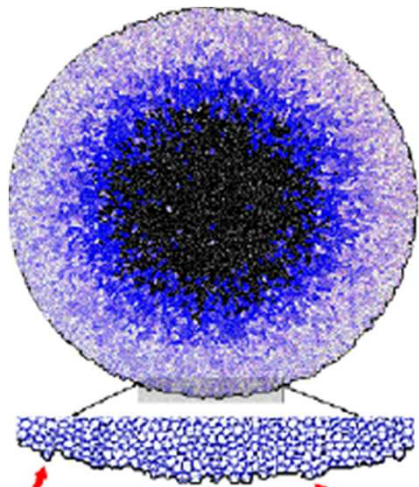
- Estimate glioma growth speed in grey and white matter
- Take into account anatomical barriers
- Can extrapolate future evolution

Tumor Growth Modeling

Extremely complex phenomena

genes ↔ proteins ↔ enzymes ↔ cells ↔ tissues ↔ organs

AVascular Growth



Drasdo et al., Phys. Biol. 2005

Angiogenesis

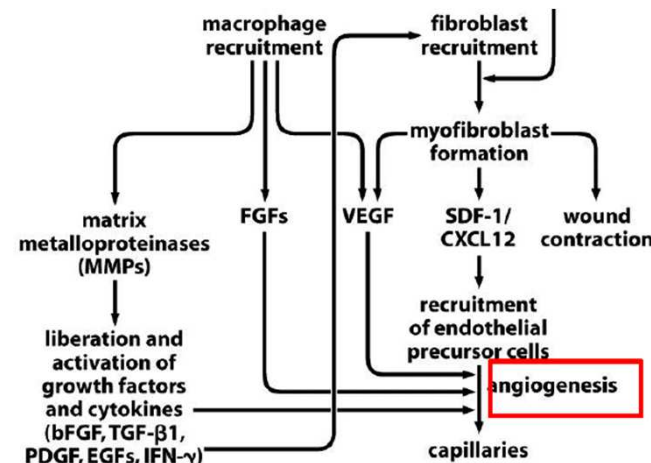
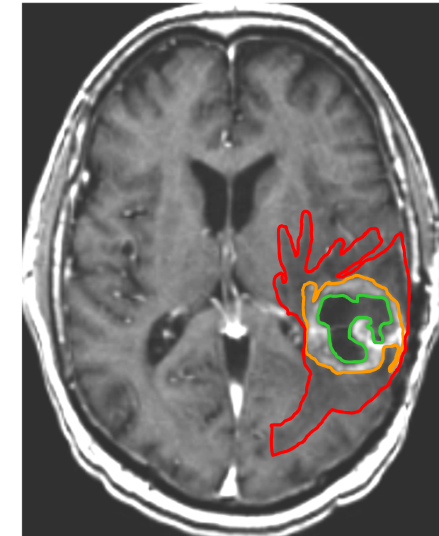


Figure 13-10 The Biology of Cancer (© Garland Science 2007)

M. Brady, IHP, 2012

Vascular Growth



T1+ gad

T S Deisboeck and G Stamatakos, editors, *Multiscale Cancer Modeling*, CRC Press, 2010.

Mathematical Tumor Growth Models

Microscopic Models

Macroscopic Models

Alarcon [Prog. Biophys. Mol. Biol. 2004]
Araujo [Bull. Math. Biol. 2004]
Athale-Deisboeck [JTB 2006]
Byrne [Math Med Bio 2003, MMMAS 2006]
Beward [Bull. Math. Biol. 2004]
Chaplain [NeuroOncology 2000]
Drasdo [Phys. Biol. 2005]
Frieboes-Cristini [NeuroImage 2007]
Maini [Tissue Eng. 2004]
Mantzaris [J Math. Biol. 2004]
Lloyd-Szekely [MICCAI 2007]
Plank [Bull. Math. Biol. 2004]
Zhang [J Th. Biol. 2007]

- *in-vivo* and *in-vitro* experiments
- Cellular level dynamics: interactions between different cells, different chemicals secreted, nutrition/oxygen sources,...
- Large variety of mathematical methods: PDEs, cellular automata, statistical methods
- Stochastic nature of the tumor growth
- Complex models, large number of parameters

Mathematical Tumor Growth Models

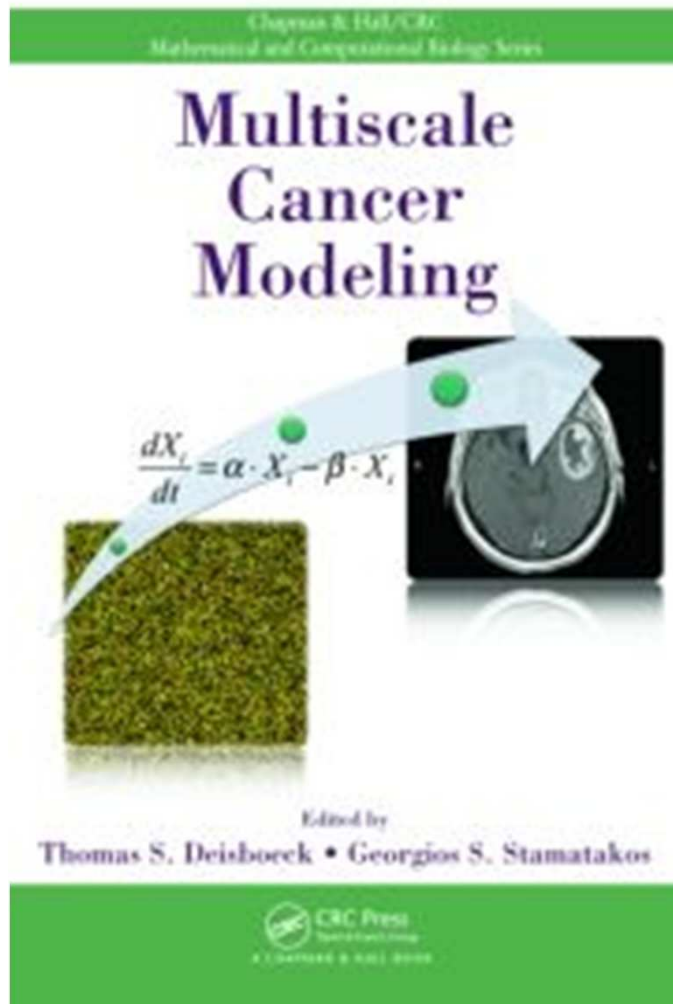
Microscopic Models

Macroscopic Models

- Observations at the macroscopic scale e.g. medical images
- Average behavior of tumor cells and their interactions with the tissues (white matter, gray matter, ...)
- simpler formulations, smaller number of variables
- Identification through manual fitting with observations

Ashraf-Davatzikos [Media 2006]
Clatz [IEEE TMI 2005]
Hogea [MICCAI 2006, 2007]
Jbadbi-Benali [MRM 2005]
Garg-Miga [SPIE 2008]
Mohamed [MedIA 2006]
Murray [Mathematical Biology 2002]
Prastawa-Gerig [MICCAI 2005, MedIA 2008]
Sierra-Szekely [MedIA 2005]
Stamatakis [Brit. J. Rad. 2006]
Stein [J Biophys. 2007]
Swanson [Br. J. Cancer 2002, 2008]
Tracqui [Cell Proliferation 1995]

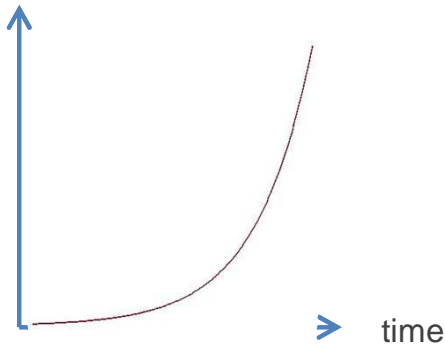
Multiscale Cancer Modeling



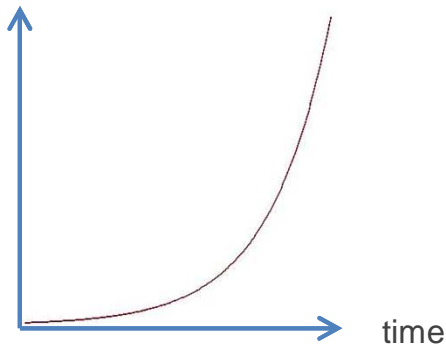
T S Deisboeck and G Stamatakos,
editors, *Multiscale Cancer Modeling*,
CRC Press, 2010.

Simple Global Growth models

Tumor cell density



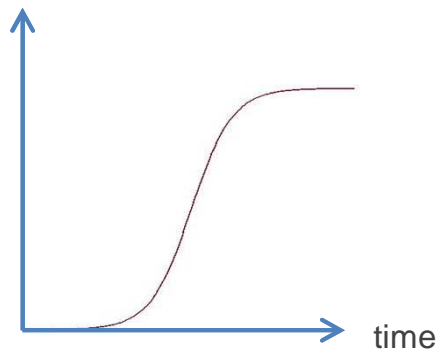
Tumor cell speed growth



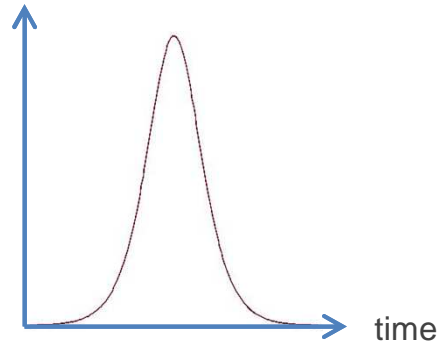
Exponential

$$\frac{dc}{dt} = \rho c$$

Tumor cell density



Tumor cell speed growth

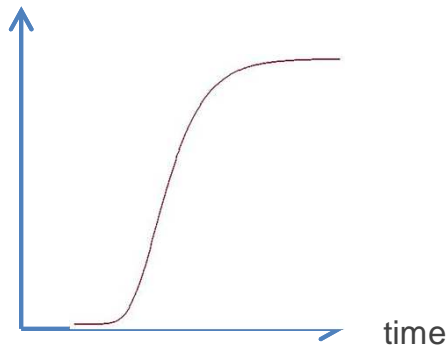


Logistic

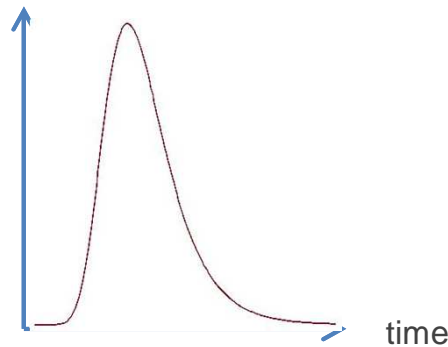
$$\frac{dc}{dt} = \rho c(1 - c)$$

Simple Global Growth models

Tumor cell density



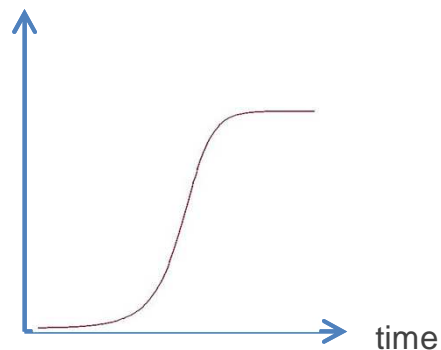
Tumor cell speed growth



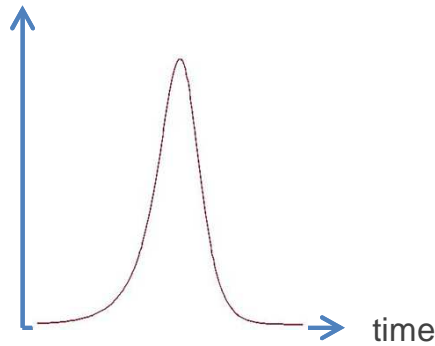
Gompertz

$$\frac{dc}{dt} = \rho c \log(K/c)$$

Tumor cell density



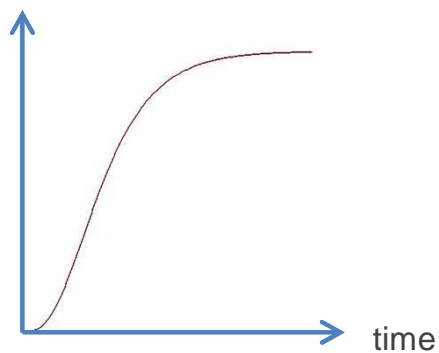
Tumor cell speed growth



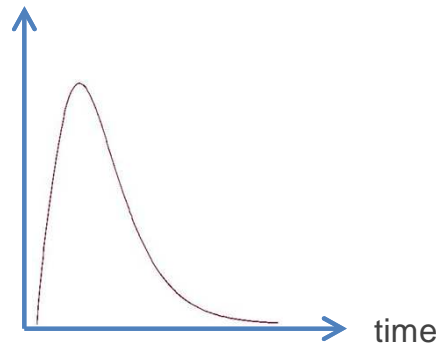
Generalized Logistic

$$\frac{dc}{dt} = \frac{\rho}{\nu} c (1 - c^\nu)$$

Tumor cell density



Tumor cell speed growth



Universal Growth

[West et al. 2001]

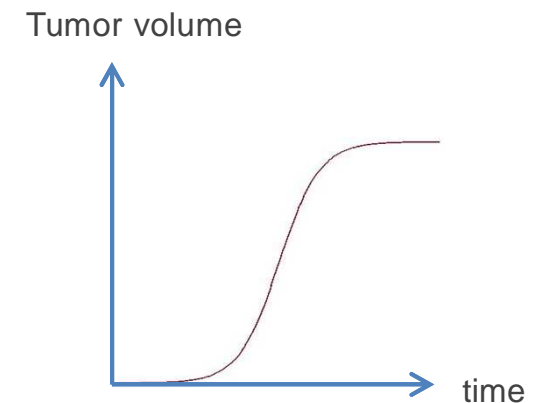
$$\frac{dc}{dt} = \frac{\rho}{\nu} c^{\frac{3}{4}} (1 - c^{1/4})$$

Limitation of Global Models

- Model evolution of global volume

$$\frac{dc}{dt} = \rho c(1 - c)$$

- Limitations :
 - heterogeneity of image
 - Anisotropy of phenomena
 - Natural barriers
- Adding spatial information

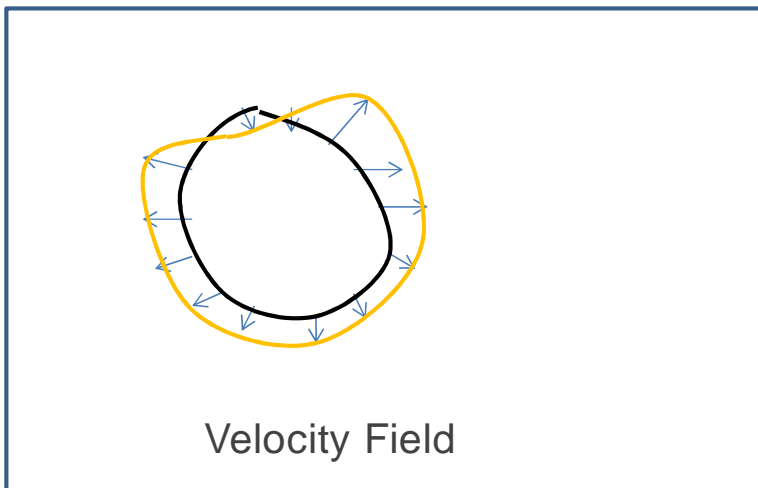


From Global to Local Models

$$\frac{dc}{dt} = \underbrace{v \cdot \nabla c}_{\text{Advection / Transport}} + \underbrace{\rho c(1 - c)}_{\text{reaction}}$$

$$\frac{dc}{dt} = \underbrace{d\nabla \cdot \nabla c}_{\text{Diffusion}} + \underbrace{\rho c(1 - c)}_{\text{reaction}}$$

$$\frac{\partial}{\partial t} \int_{\Omega} c dV = \int_{\Omega} \rho c(1 - c) dV$$



[Hogea et al. 2006]
[T. Colin, O. Saut et al.]

Image-based Glioma Growth Models

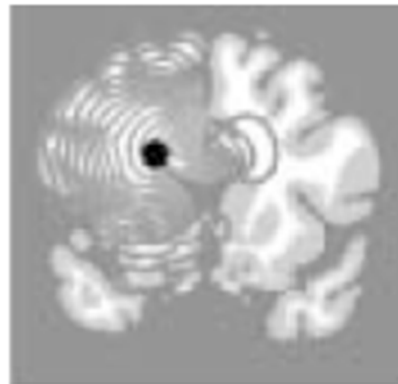
2D, Isotropic,
homogeneous



1995

P. Tracqui et al.

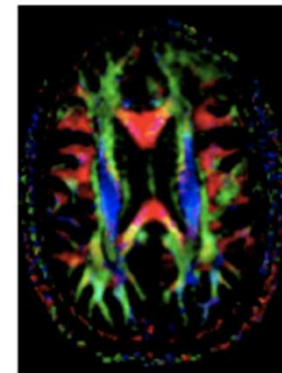
3D, Isotropic,
Non-homogeneous



2002

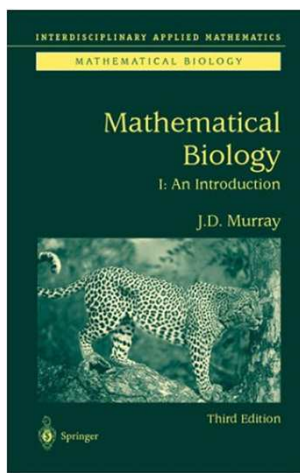
K. Swanson et al.

3D, anisotropic,
Non-homogeneous



2005

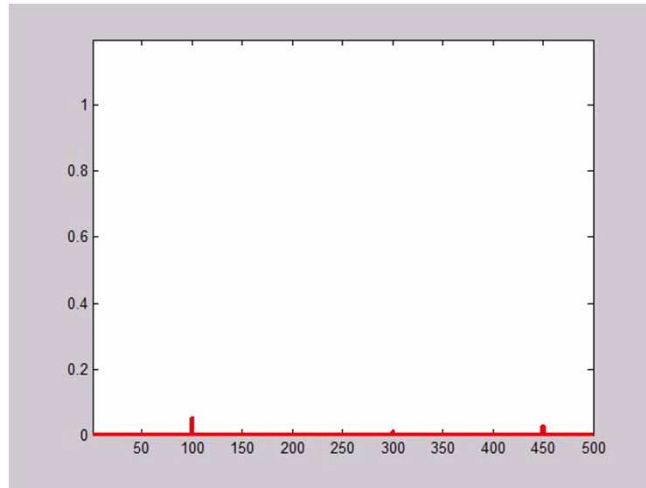
O. Clatz et al.
Jbabdi et al.



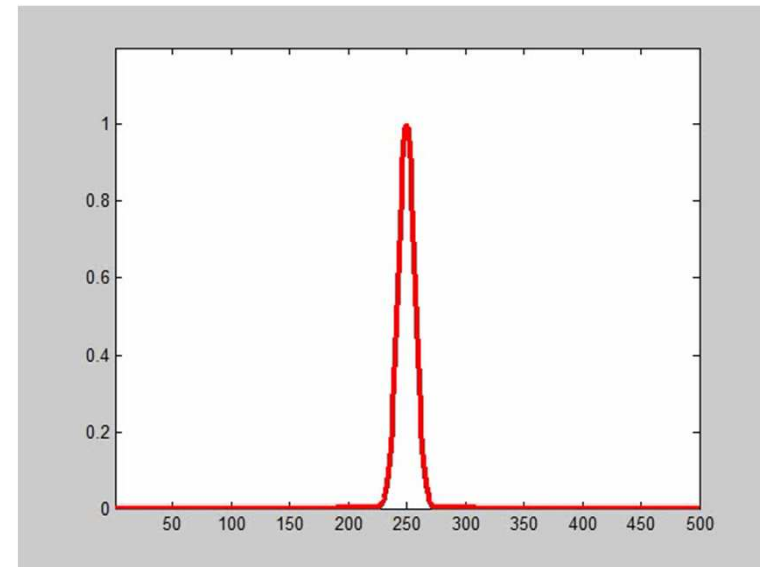
James Murray
Univ. Of Washington

Giese, Alf M. et al., "Migration of Human Glioma Cells on Myelin", *Neurosurgery*, April 1996 - Volume 38 - Issue 4 - pp 755-764

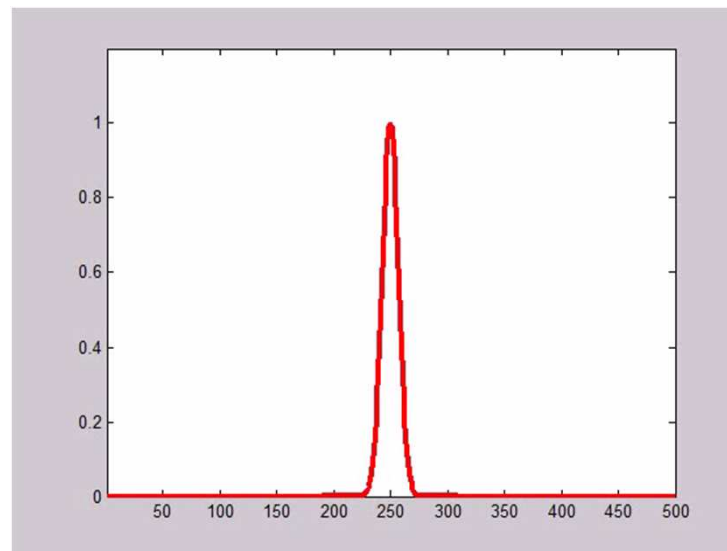
Understanding Reaction diffusion



Reaction Only



Diffusion Only



Reaction-Diffusion

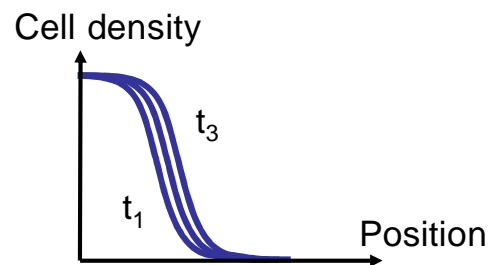
Fisher Equation

$$\frac{\partial c}{\partial t} = \nabla \cdot (D \nabla c) + \rho c(1 - c)$$

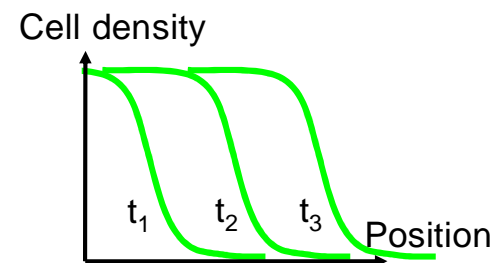
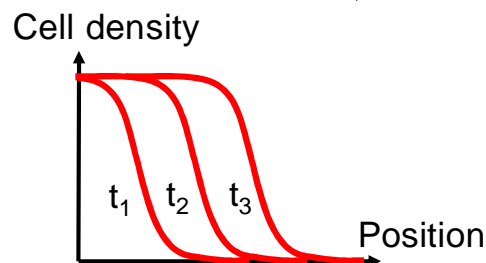
Product controls growth speed

$$D\rho$$

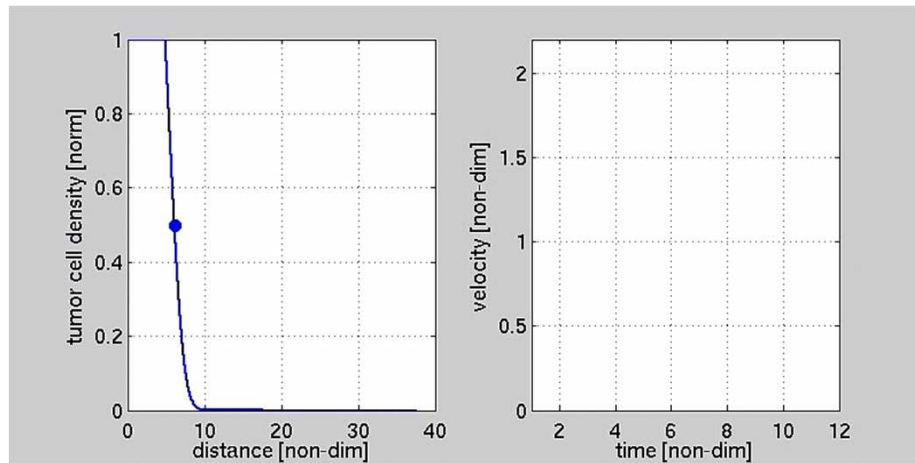
$$\text{Velocity index} = 2\sqrt{D\rho}$$



$$D_1\rho_1 \gg D_2\rho_2 \gg D_3\rho_3$$

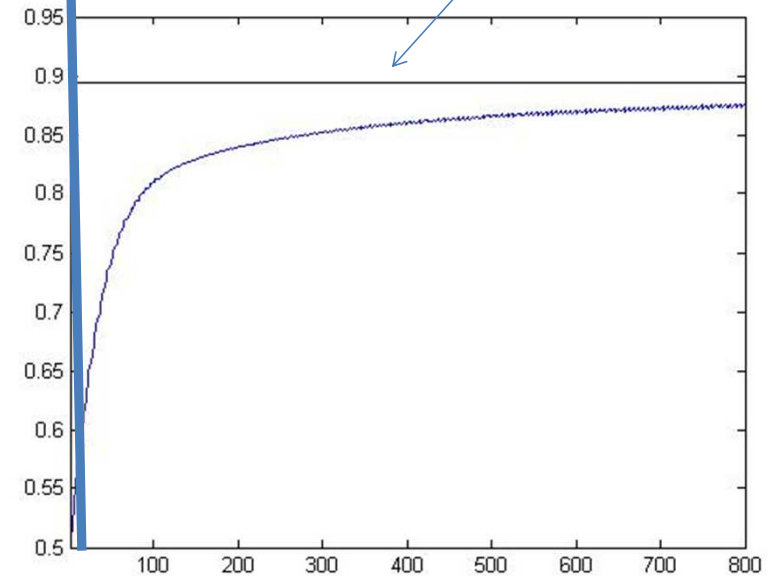


Speed of moving front



velocity

Asymptotic speed



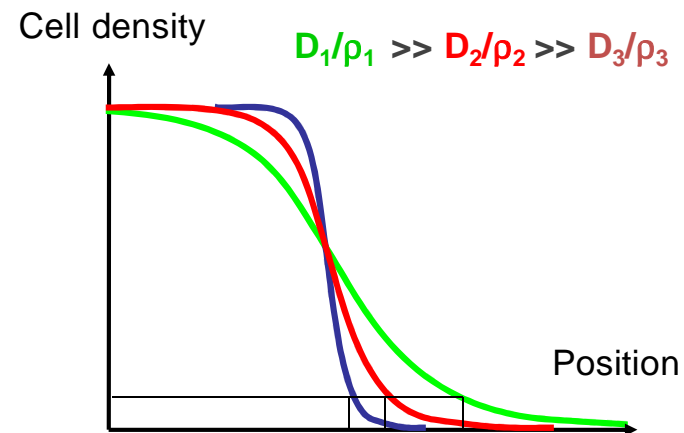
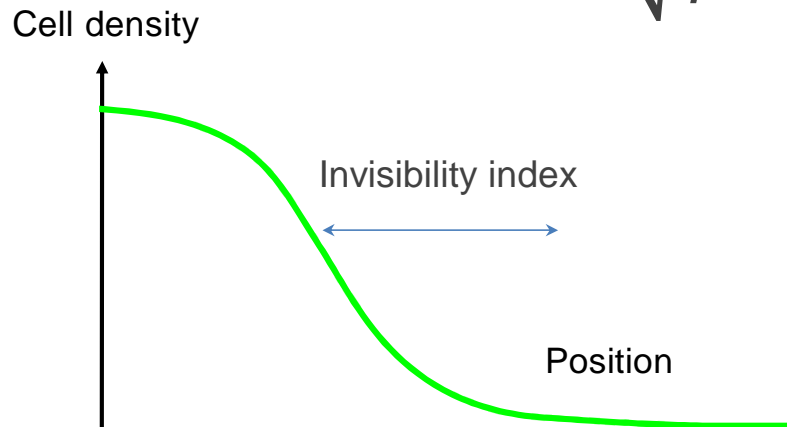
Fisher Equation

$$\frac{\partial c}{\partial t} = \nabla \cdot (D \nabla c) + \rho c(1 - c)$$

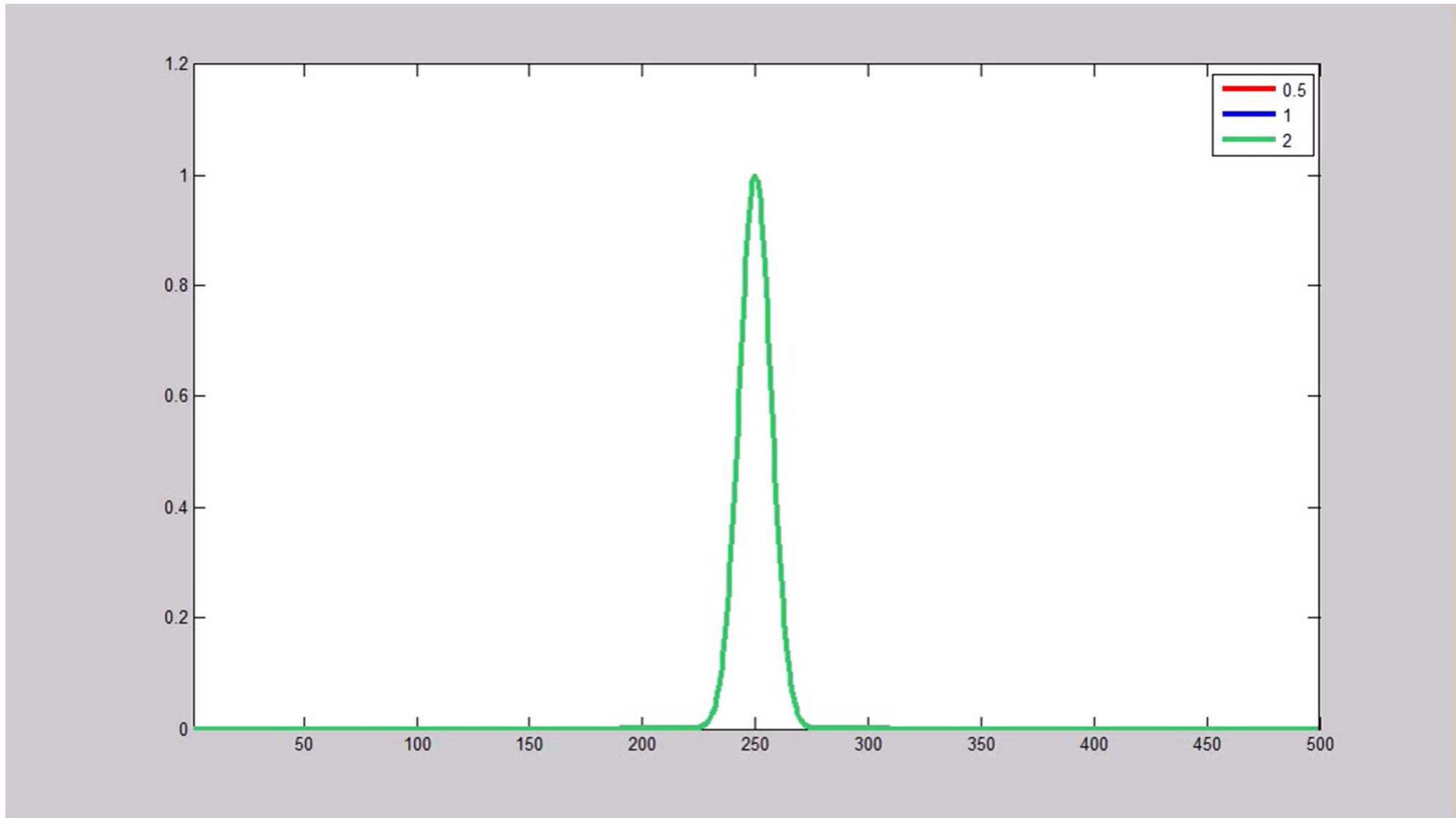
Ratio controls the extent of infiltration

$$\frac{D}{\rho}$$

$$\text{Invisibility index} = \sqrt{\frac{D}{\rho}}$$

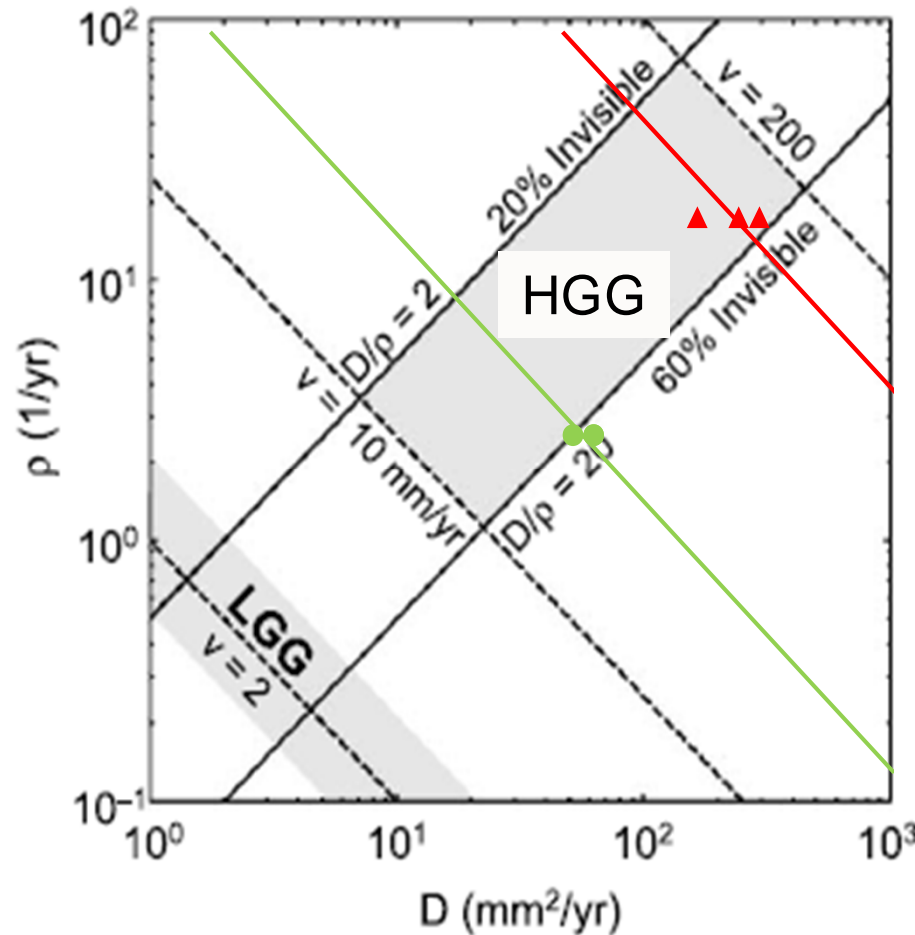


Understanding the Fisher Equation



Infiltration vs. propagation

$$v = 2(D\rho)^{1/2}$$



High grade,
 $v_w = 110\text{mm/yr}$

Low grade,
 $v_w = 2\text{-}4\text{mm/yr}$

$$i = D/\rho$$

From Swanson et al.

State space Fisher equation

A Multilevel Model of Tumor Growth

- Proposed by Clatz et al. (2005)
- Includes
 - Geometry
 - Statistics (Atlas)
 - Biomechanics
 - Physiopathology



March



September

Coll. CAL-CHU (Nice) & SPL BWH (Harvard)

O. Clatz, M. Sermesant, P.-Y. Bondiau, H. Delingette, S. Warfield, G. Malandain, N. Ayache. Realistic Simulation of the 3D Growth of Brain Tumors in MR Images Including Diffusion and Mass Effect. *IEEE Transactions on Medical Imaging*. 24(10):1334-1346, Oct. 2005.

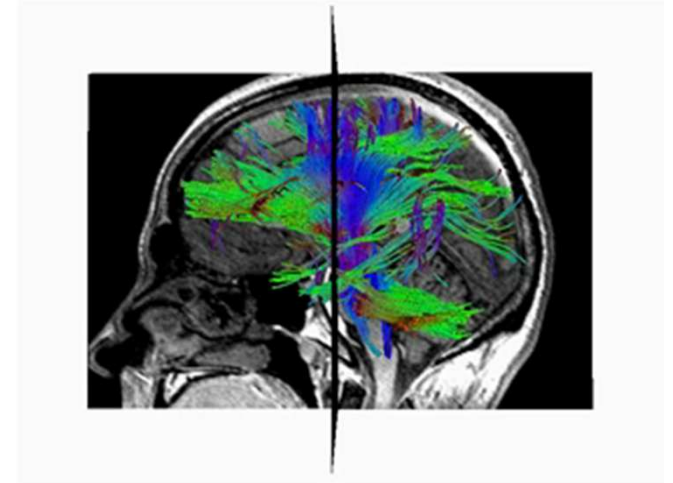
Anisotropic Diffusion

Fisher Kolmogorov

$$\frac{\partial c}{\partial t} = \nabla \cdot (D \nabla c) + \rho c(1 - c)$$

Different Cell Motility depending on tissue

- White matter -> High diffusion
- Grey Matter -> Low diffusion



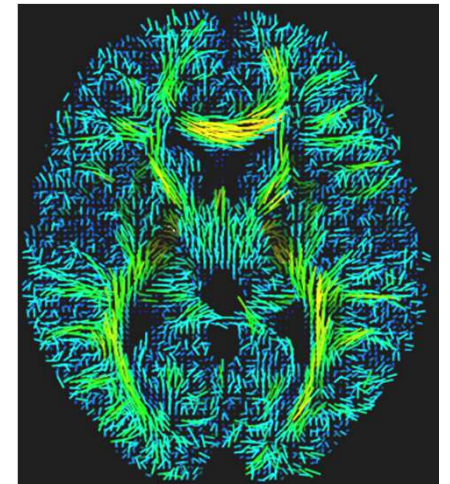
D= Tumor Diffusion Tensor

In grey matter $D = d_g Id$

In white matter $D = d_w \frac{D_{water}}{d_0}$

DTI Image

Normalization factor

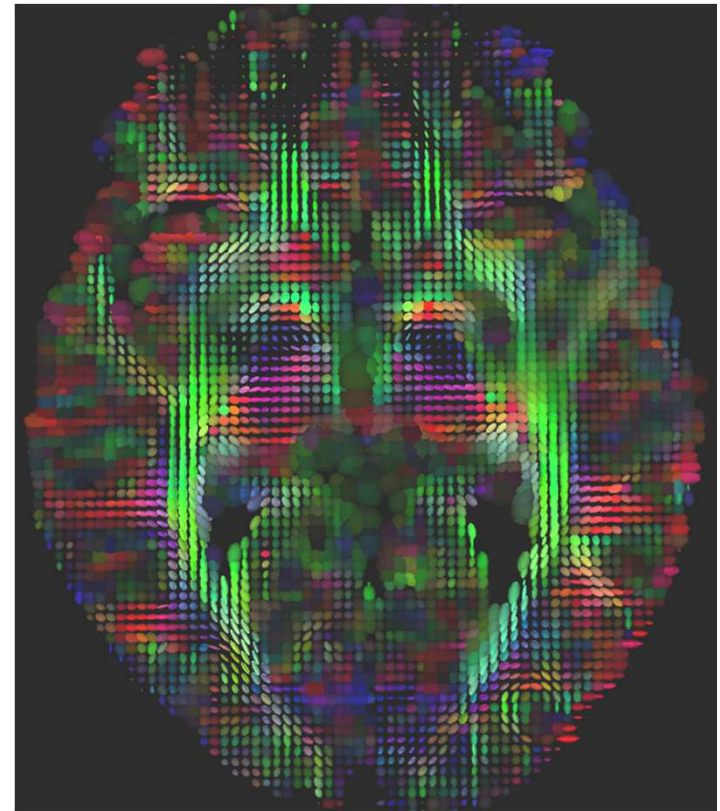


Other methods for TDT : [Jbabdi 2005], [Swanson et al.]

1. Geometrical Model



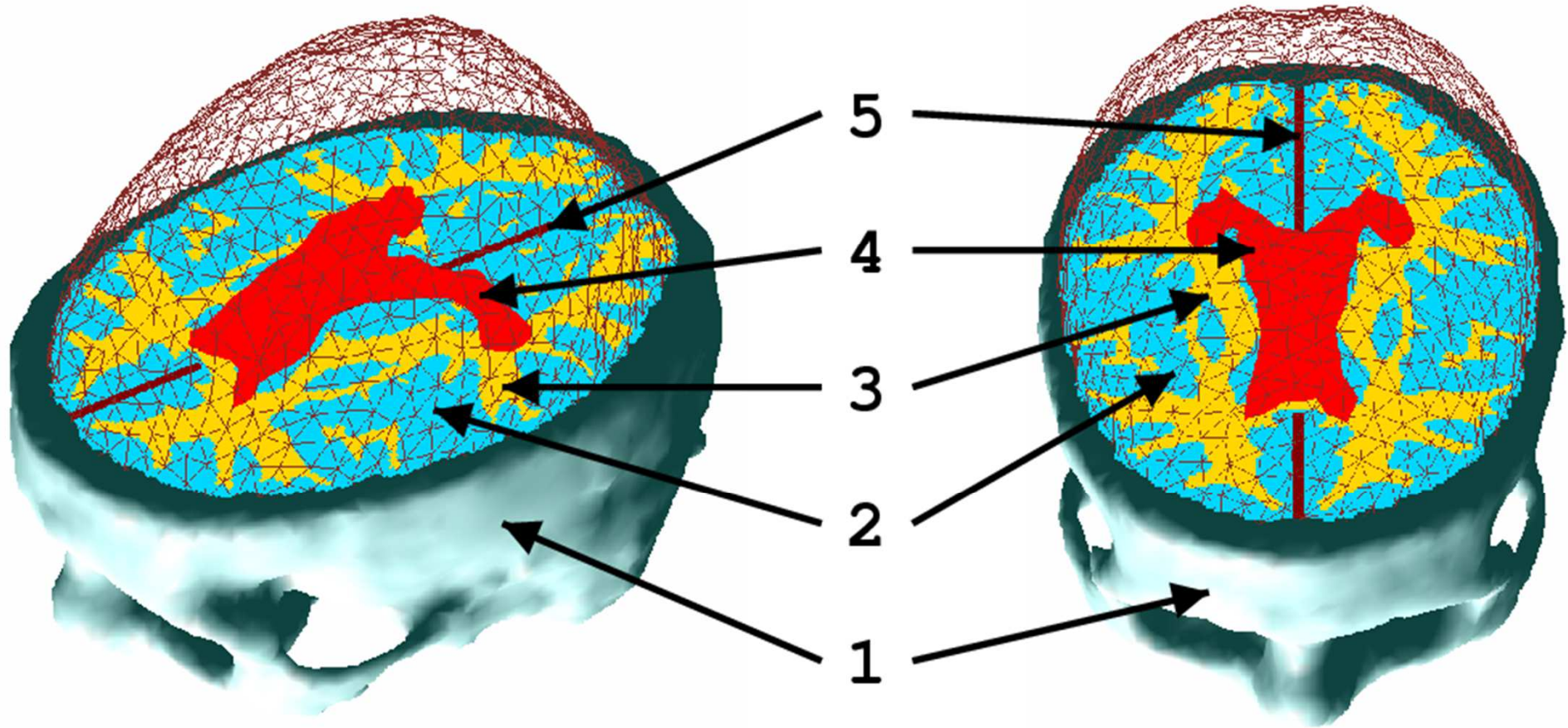
Segmented T1 MRI
(Brainweb)
GM, WM, CSF, Skull, etc.



DTI
Main fiber bundles of WM

2. Biomechanical Model

Inhomogeneous Anisotropic Linear Elastic



1 Skull. 2 Grey matter 3 White matter.
4 Ventricles. 5 Falx cerebri

3. Physiopathological Model

Evolution of tumor density c

Fisher Kolmogorov

$$\frac{\partial c}{\partial t} = \nabla \cdot (D \nabla c) + \rho c (1 - c)$$

+CL

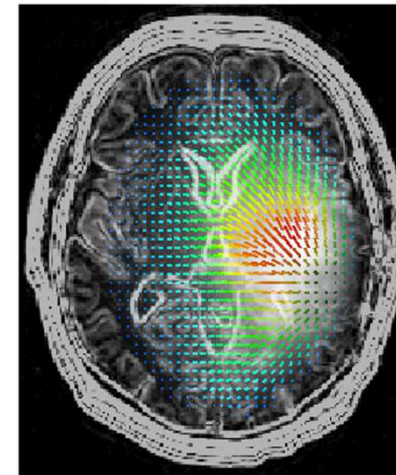
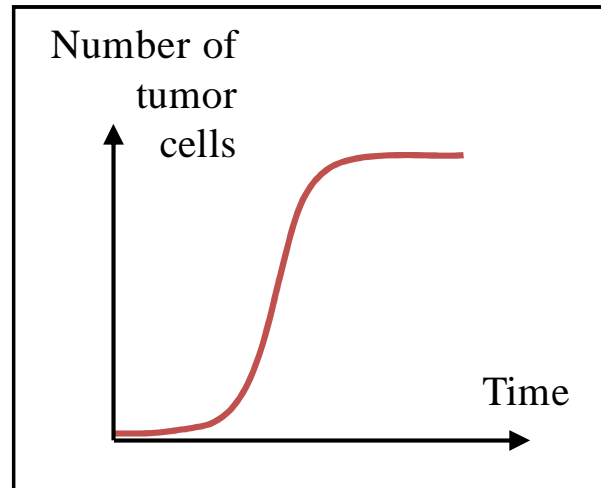
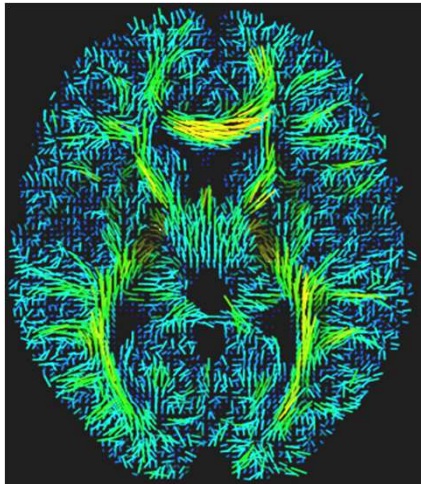
Anisotropic migration

Logistic proliferation

Biomechanical coupling

$$\text{div}(\sigma - \alpha c I_3) + Fe = 0$$

Mass effect

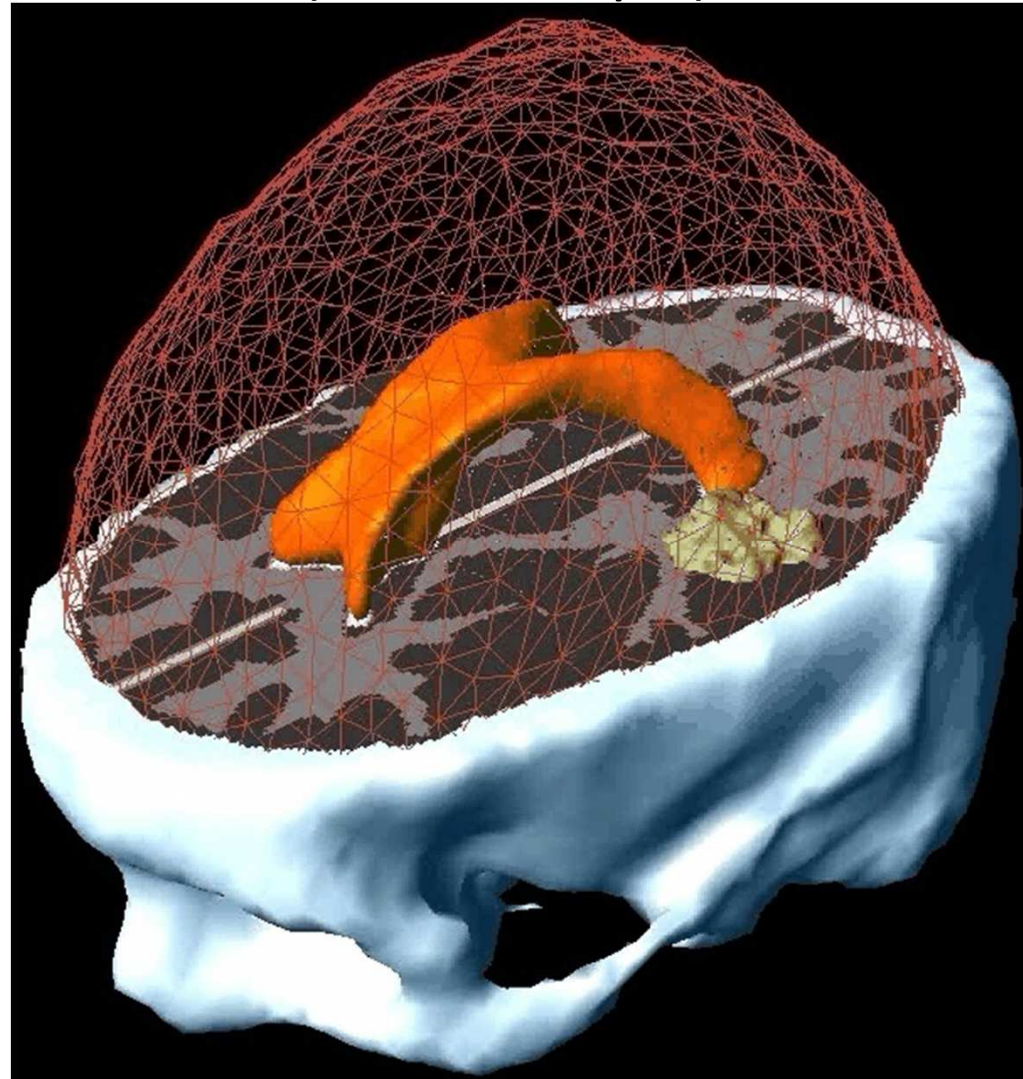


Collaboration Centre Antoine Lacassagne & Harvard



Simulated Growth

(Mars – Sept.)



O Clatz, PY Bondiau, H Delingette, M Sermesant, SK Warfield, G. Malandain,
N.A.. *Brain Tumor Growth Simulation*. [IEEE-TMI 2005](#)

Solving the Fisher equation :

- Several schemes :

- Explicit schemes

- Lattice-Boltzman Method

- Semi-implicit scheme : “smooth and grow”

} Easily parallelizable

- Parameters :

- Diffusivity in grey matter d_g + white matter d_w

- Proliferation factor : ρ

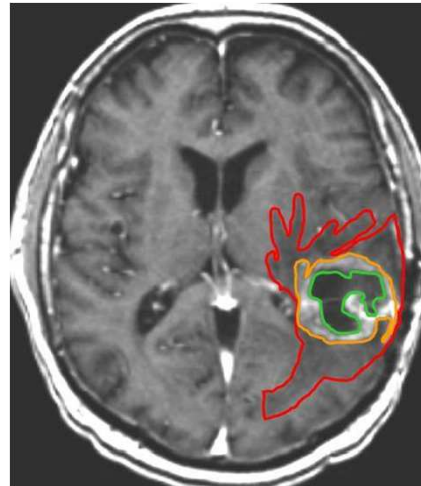
Necrosis and vascularisation

- **3 types** of tumor cells

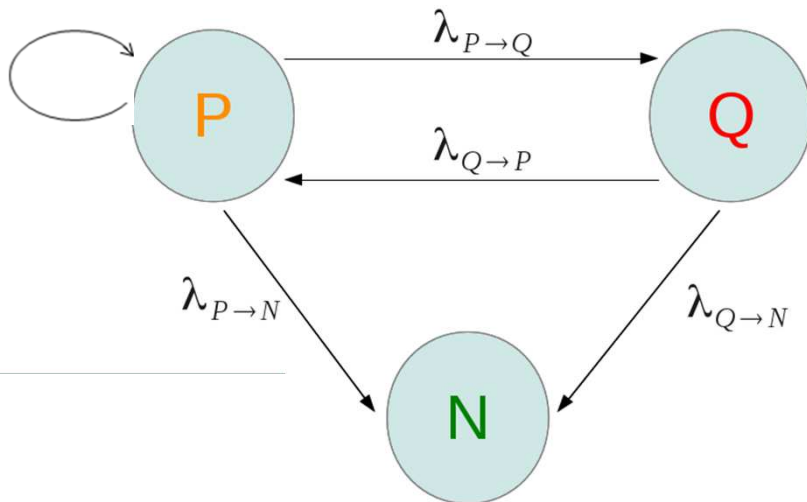
Proliferating : multiplication and diffusion

Quiescent : strong diffusion

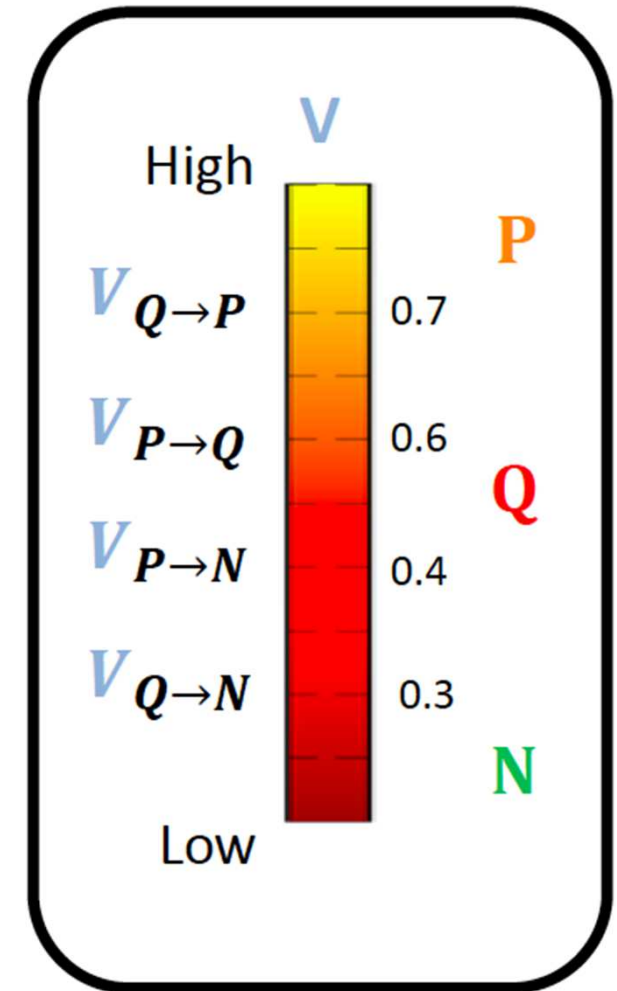
Necrosed : static



- **Transitions** : depend on vascularisation



- **Vascularisation** : function of P & N



T. Colin, O. Saut et al., 2012, M. Le 2012

Necrosis and vascularisation

DIFFUSION

PROLIFERATION

TRANSITIONS

$$\frac{\partial P}{\partial t} = \nabla \cdot (D_P(1 - T)\nabla P)$$

$$+\rho P(1 - T)$$

$$-\lambda_{P \rightarrow Q}P - \lambda_{P \rightarrow N}P + \lambda_{Q \rightarrow P}Q$$

$$\frac{\partial Q}{\partial t} = \nabla \cdot (D_Q(1 - T)\nabla Q)$$

$$-\lambda_{Q \rightarrow P}Q - \lambda_{Q \rightarrow N}Q + \lambda_{P \rightarrow Q}P$$

$$\frac{\partial N}{\partial t} =$$

$$+\lambda_{P \rightarrow N}P + \lambda_{Q \rightarrow N}Q$$

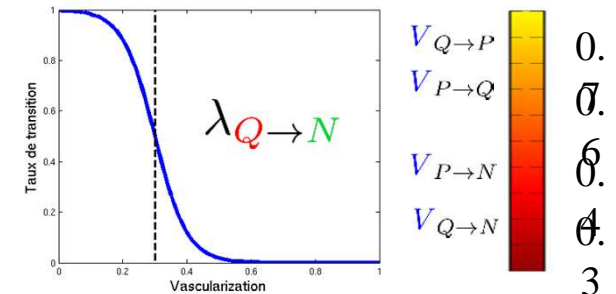
$$T = P + Q + N$$

ANGIOGENESIS

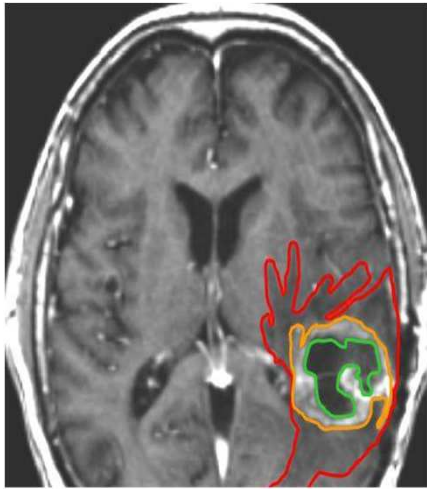
DEGRADATION

$$\frac{\partial V}{\partial t} = \alpha P(1 - V)$$

$$-\beta NV$$



Necrosis and vascularisation



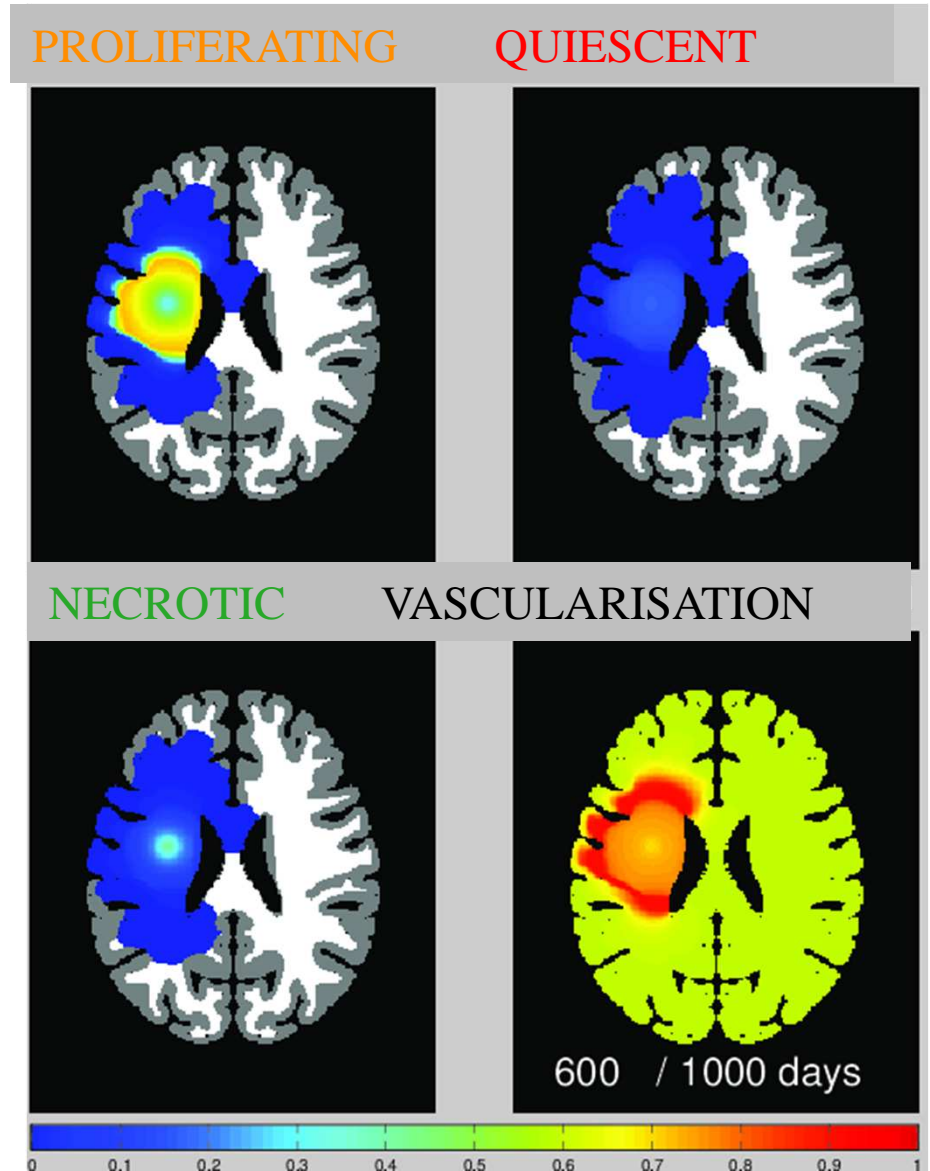
3 types of cells

Proliferating

Quiescent

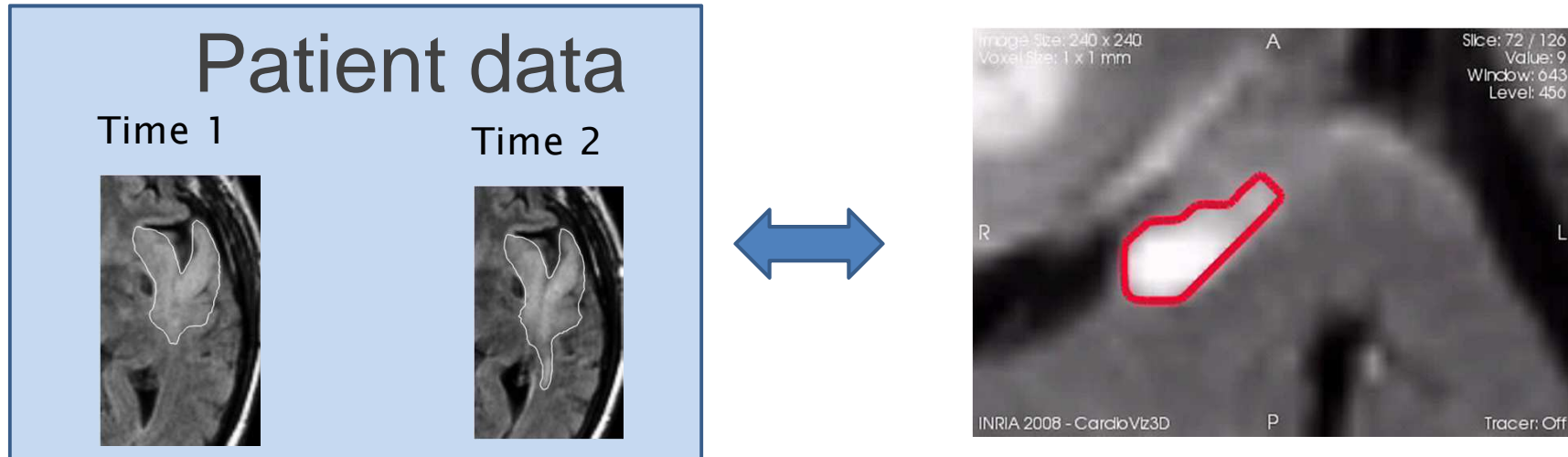
Nécrotic

- **Vascularisation** : *function of P and N*



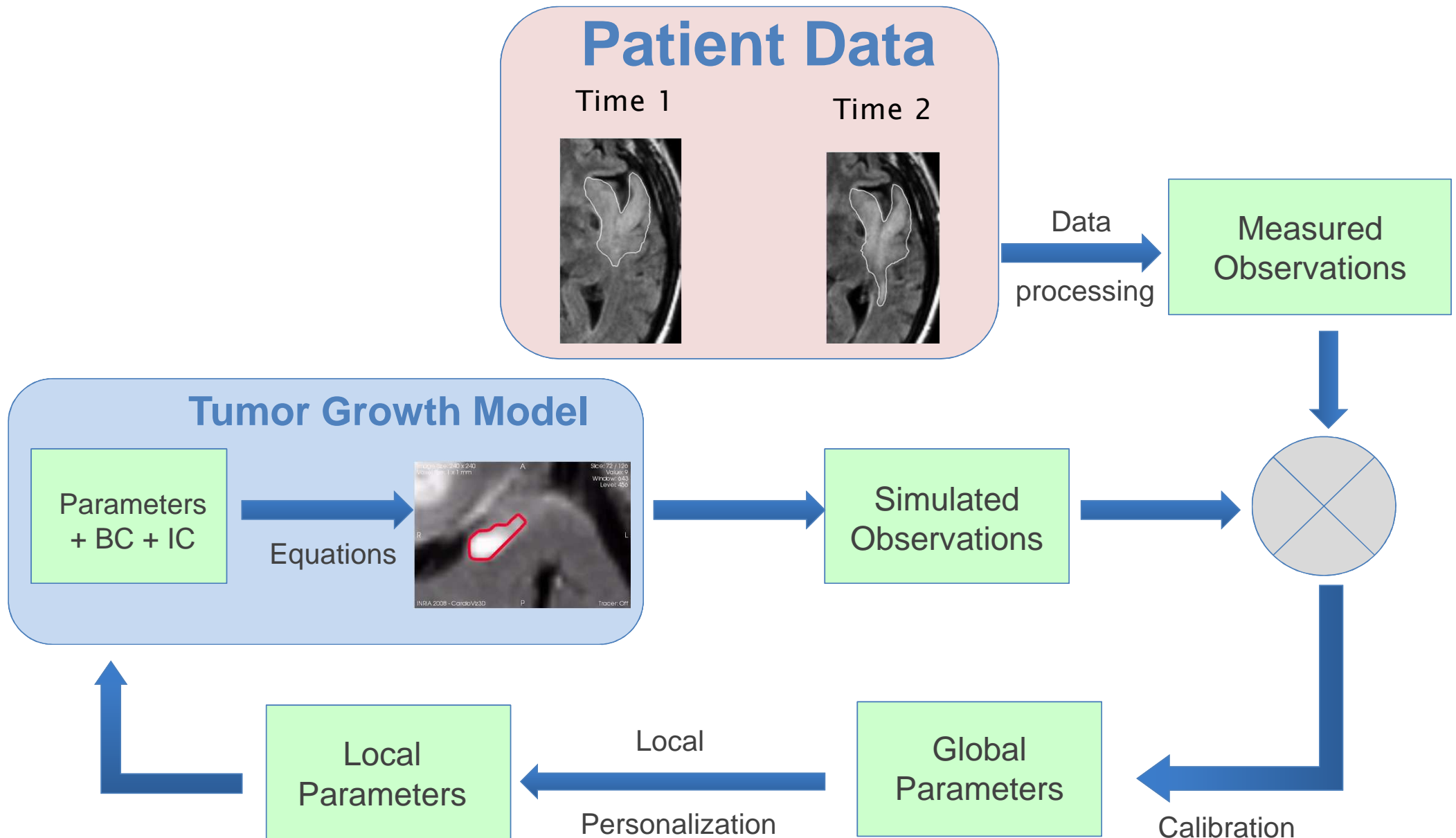
T. Colin, O. Saut et al., M. Le 2012

Personalization of Glioma Growth Models



- Why Personalization of Models ?
 - Understanding the limits of models
 - Quantification of tumor growth
 - Therapy planning

Model Personalization



Personnalization

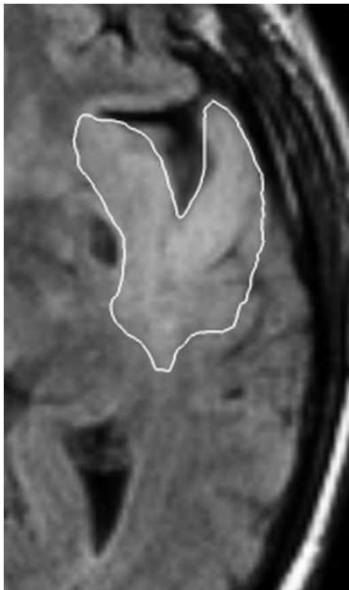
- 1. Growth speed
- 2. Invisibility Index

A simpler problem

- Only observe the evolution of 2 contours

➔ Estimate only the speed of front

Time 1



Time 2



Time N



A simpler Model

- Approximate the RD equation for an isodensity surface \longrightarrow Asymptotic analysis

Search solution as $c(x - vt)$

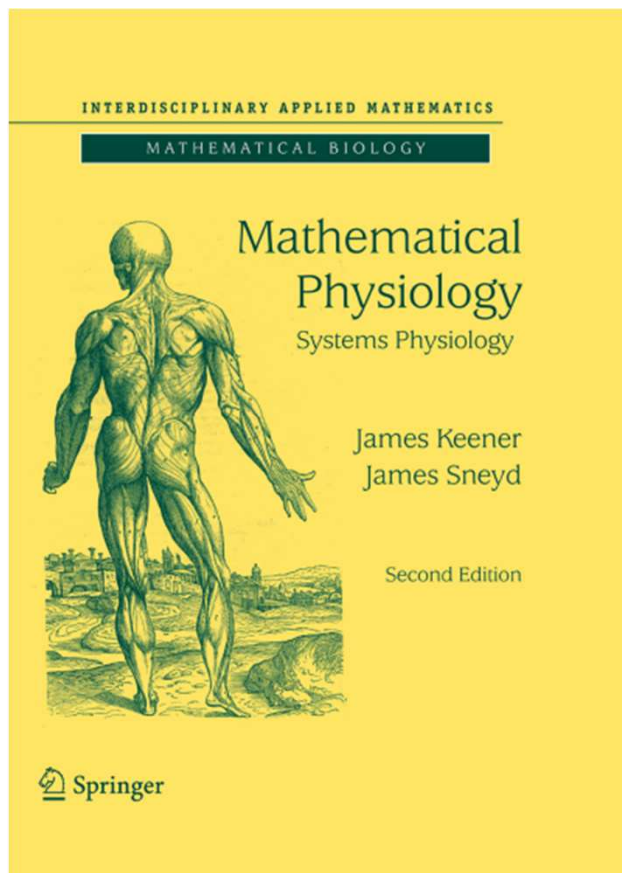
Leads to Eikonal equation

Isotropic propagation $F\nabla T = 1$

Anisotropic propagation

$$2\sqrt{\rho} \sqrt{\nabla T^T D \nabla T} = 1$$

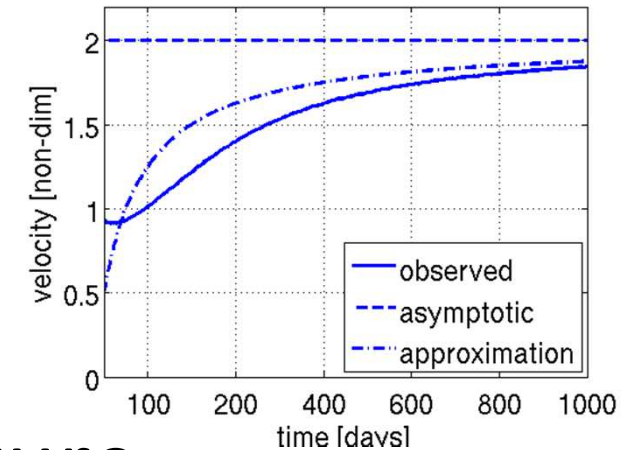
Solved with (Anisotropic) Fast Marching



Advanced Asymptotic Model of Tumor Growth

- Include dependence on time

$$\left(\frac{4\rho T - 3}{2T\sqrt{\rho}} \right) \sqrt{\nabla T' \mathbf{D} \nabla T} = 1$$



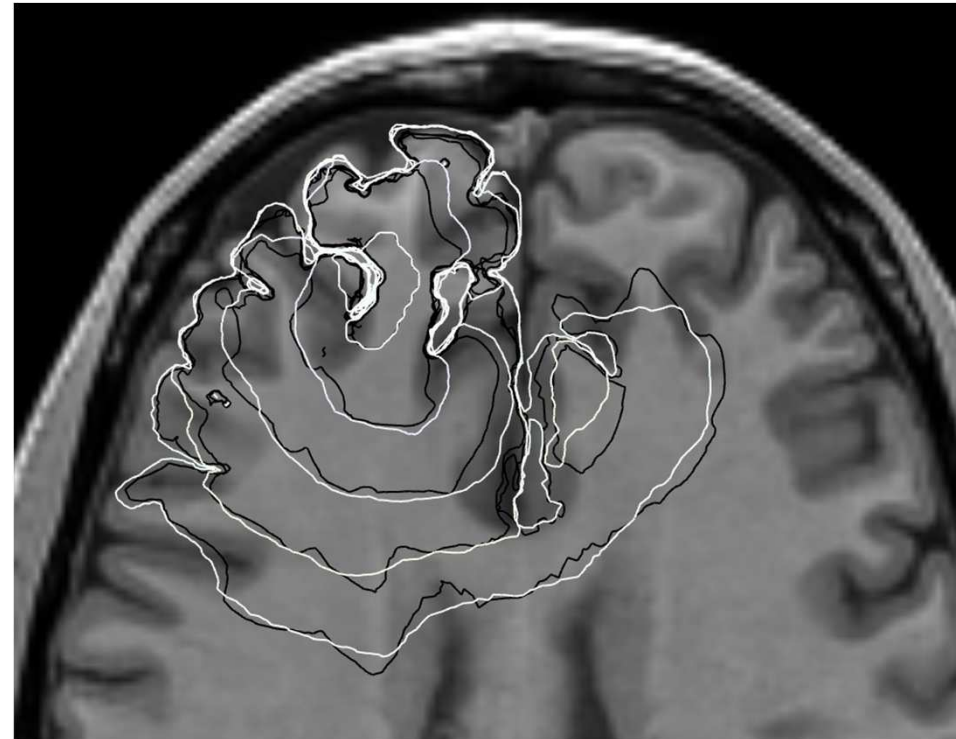
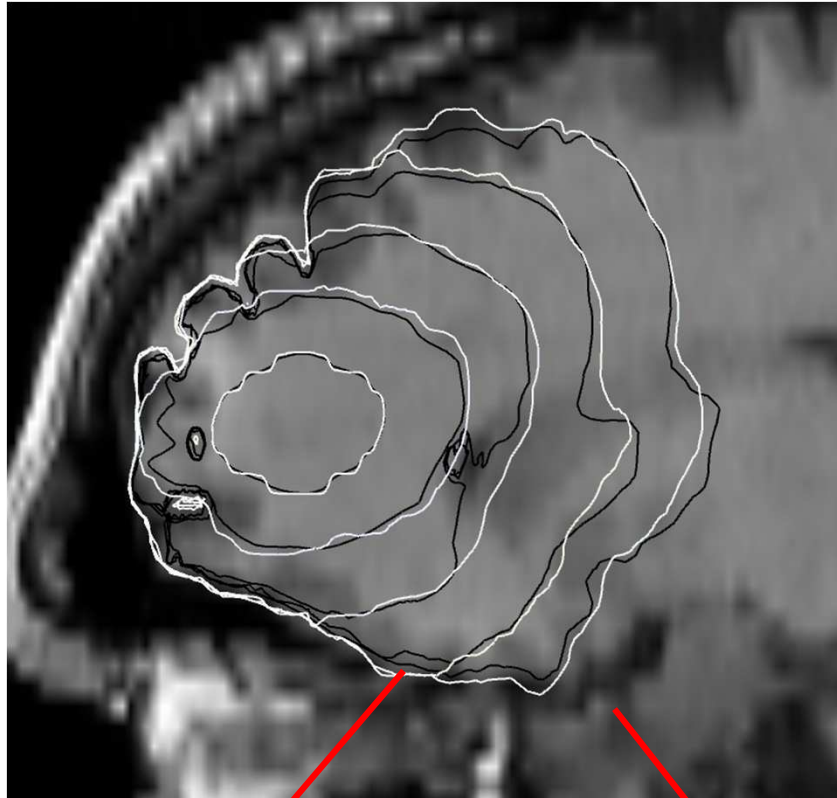
- Include dependence on curvature

$$\left\{ \frac{4\rho T - 3}{2T\sqrt{\rho}} - 0.3\sqrt{\rho} \left(1 - e^{-|\kappa_{eff}|/(0.3\sqrt{\rho})} \right) \right\} \sqrt{\nabla T' \mathbf{D} \nabla T} = 1, \quad \kappa_{eff} = \nabla \cdot \frac{\mathbf{D} \nabla T}{\sqrt{\nabla T' \mathbf{D} \nabla T}}$$

Eikonal-Curvature equation

Solved by iterative anisotropic fast-marching

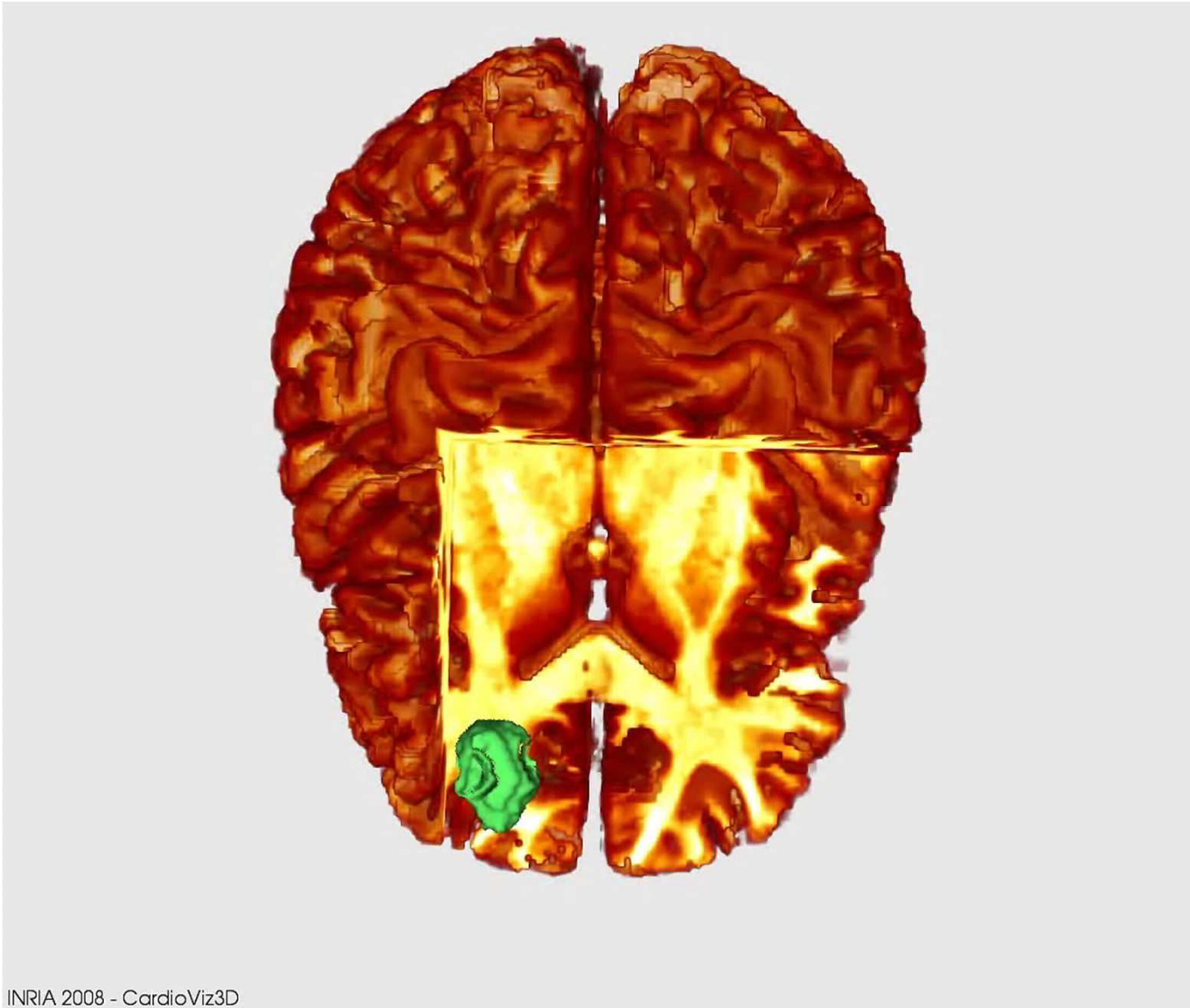
Comparing Reaction-Diffusion with Traveling Time Formulation



$u=0.4$ iso-density contour at days 400,600,800,1000 and 1200

Black: Traveling Time

White Fisher PDE



INRIA 2008 - CardioViz3D

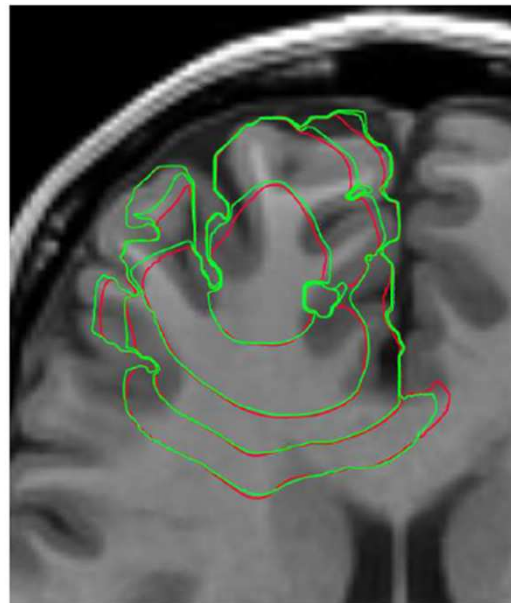
Parameters of tumor growth model

- Diffusion of tumor cells in
 - White matter : d_w
 - Grey matter : d_g
 - ~~• Proliferation rate : ρ~~
- Time of initial appearance of the tumor : T_0

Observability issue
Cannot estimate both ρ
and D

Set value of ρ

	Red	Green
d_w	0.273	0.153
d_g	0.024	0.014
ρ	0.012	0.0185



But speed
 $v \approx 2\sqrt{\rho D}$ depends
On both ρ and D

Personalization

- **Input**

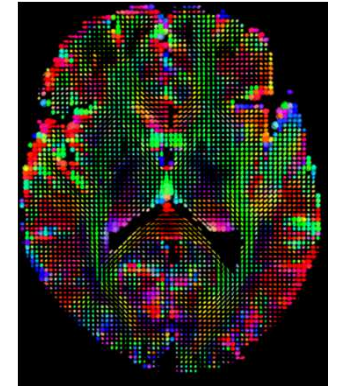


Tumor segmentation

Time between exams

$$\Delta t_1, \Delta t_2, \dots, \Delta t_{N-1}$$

DTI



$\mathbf{D}_{\text{water}}$

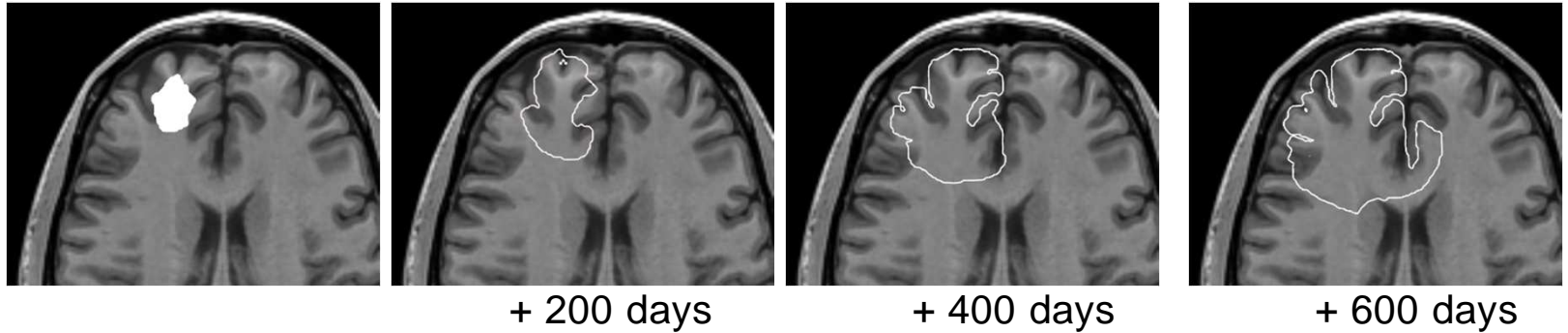
- **Output :** $(\rho d_w, \rho d_g, T_0)$

- Speed GM and WM,
- Time T0

$$\mathbf{D} = \begin{cases} \alpha d_w \mathbf{D}_{\text{water}}, & \text{white matter} \\ d_g \mathbf{I}_3, & \text{gray matter} \end{cases}$$

Forward & Backward Simulation

For a set of parameters $(\rho d_w, \rho d_g, T_0)$



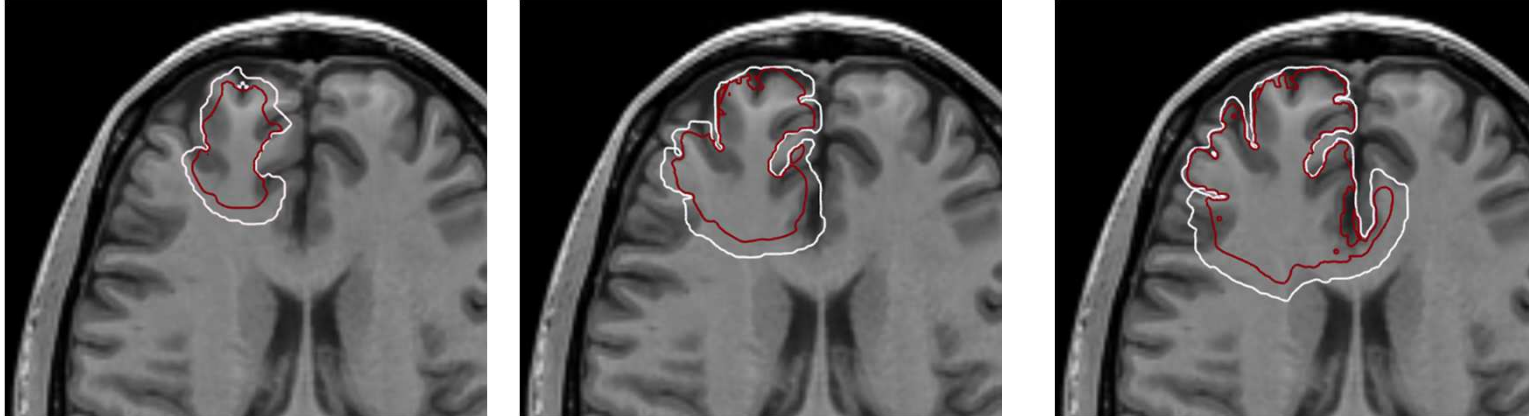
Forward Simulation
Estimation of $(\rho d_w, \rho d_g)$



Backward Simulation
Estimation of T_0

Error Criterion

For a set of parameters $(\rho d_w, \rho d_g, T_0)$



Difference between real and simulated contours

$$C_1 = \sum_{i=1, \dots, N-1} \text{dist}(\Gamma_i, \hat{\Gamma}_i)^2$$

Difference between estimated and simulated time to shrink to a single voxel

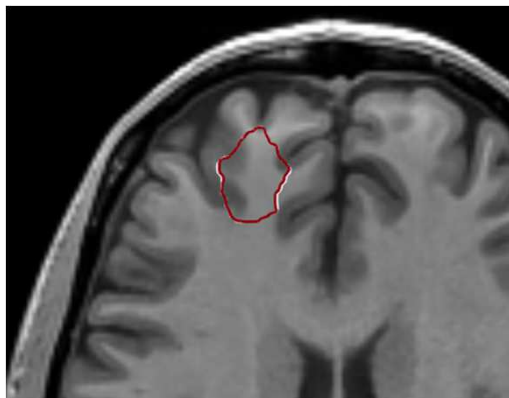
$$C_2 = \left(v_{\min} |T_{\min} - T_0| \right)^2$$

Optimisation

$$C = C_1 + C_2$$

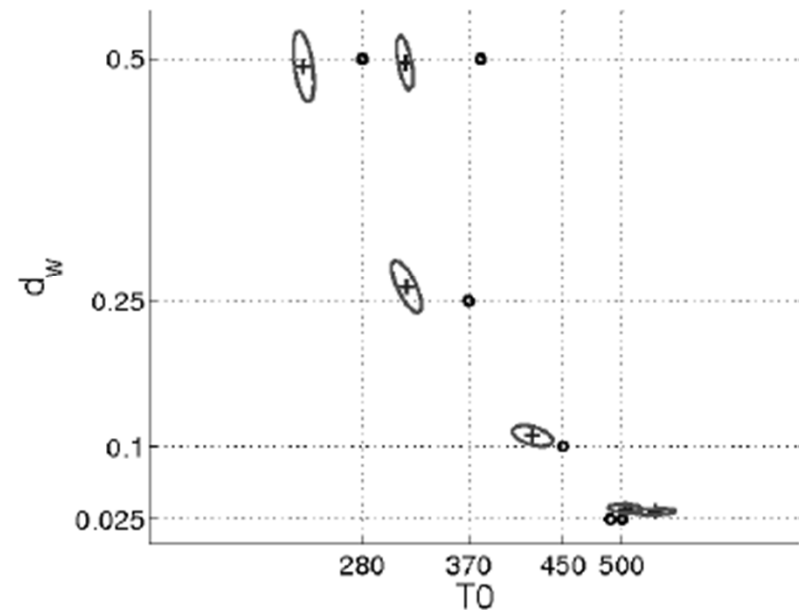
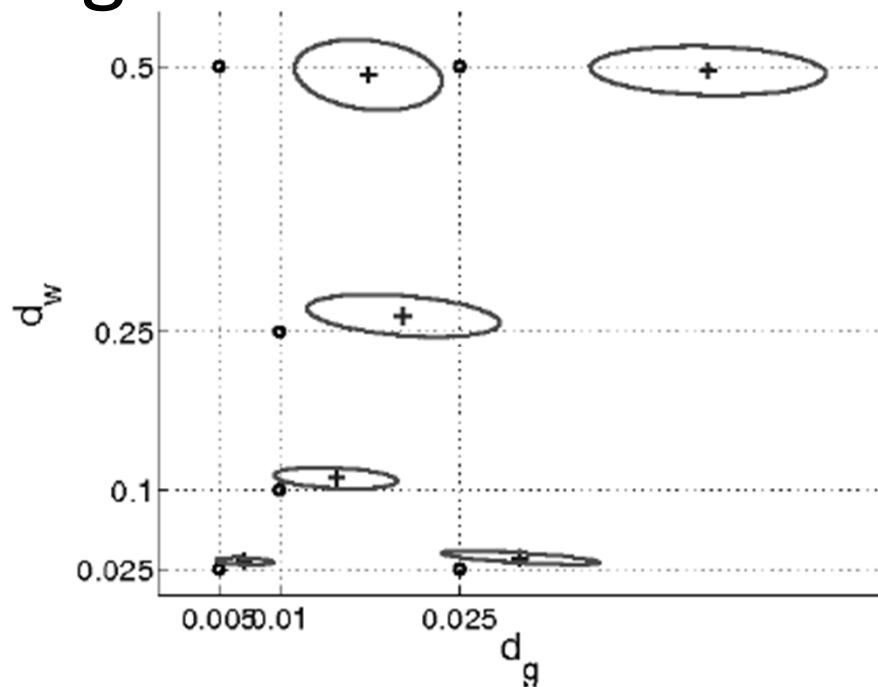
- Minimisation of criterion
 - Gradient-Free Algorithm **Powell UoByQa**

$$(\rho d_w^*, \rho d_g^*, T_0^*)$$



Test on synthetic cases

- Recover well d_w (constant relative error)
- Bias + large uncertainty for d_g
- Estimate parameter uncertainty due to segmentation uncertainty



Two real cases

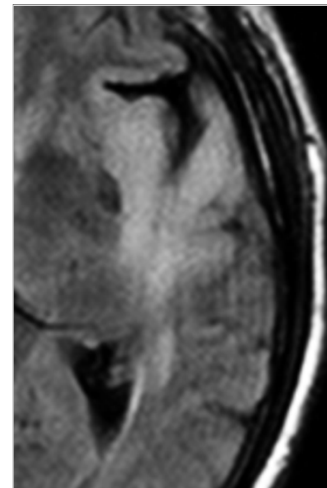
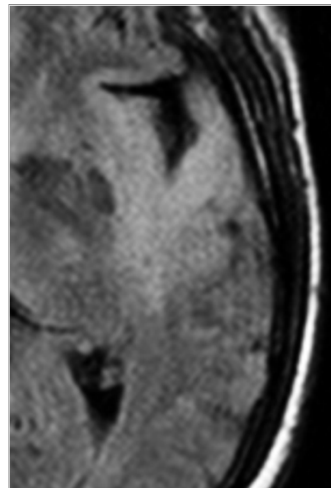
Time 1



Time 2



High grade glioma evolution



Low grade glioma evolution

High Grade Glioma



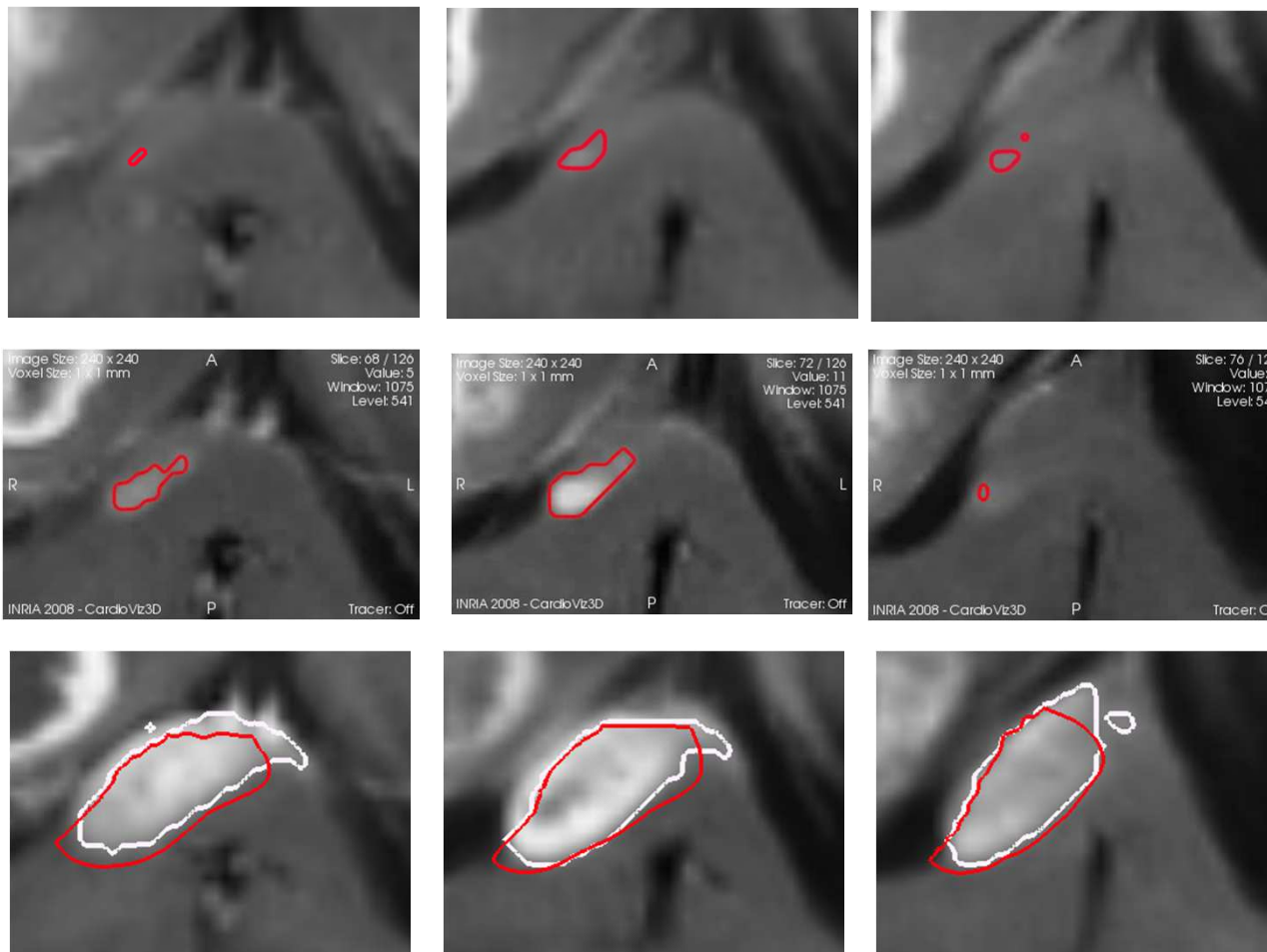
Day 1

Day 21

Day 67

time

Space



learn

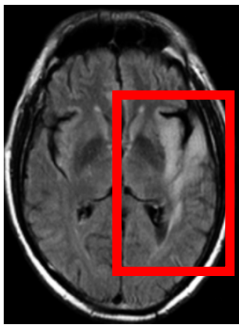
predict

MRI T1 Gd, 0.5*0.5*6.5mm
3 time points

MR DTI : 2.5mm (time 2)

$\rho(\text{set})$	d_w	d_g
0.05 1/day	0.66 mm ² /day	0.0013 mm ² /day

Simulation/Reality

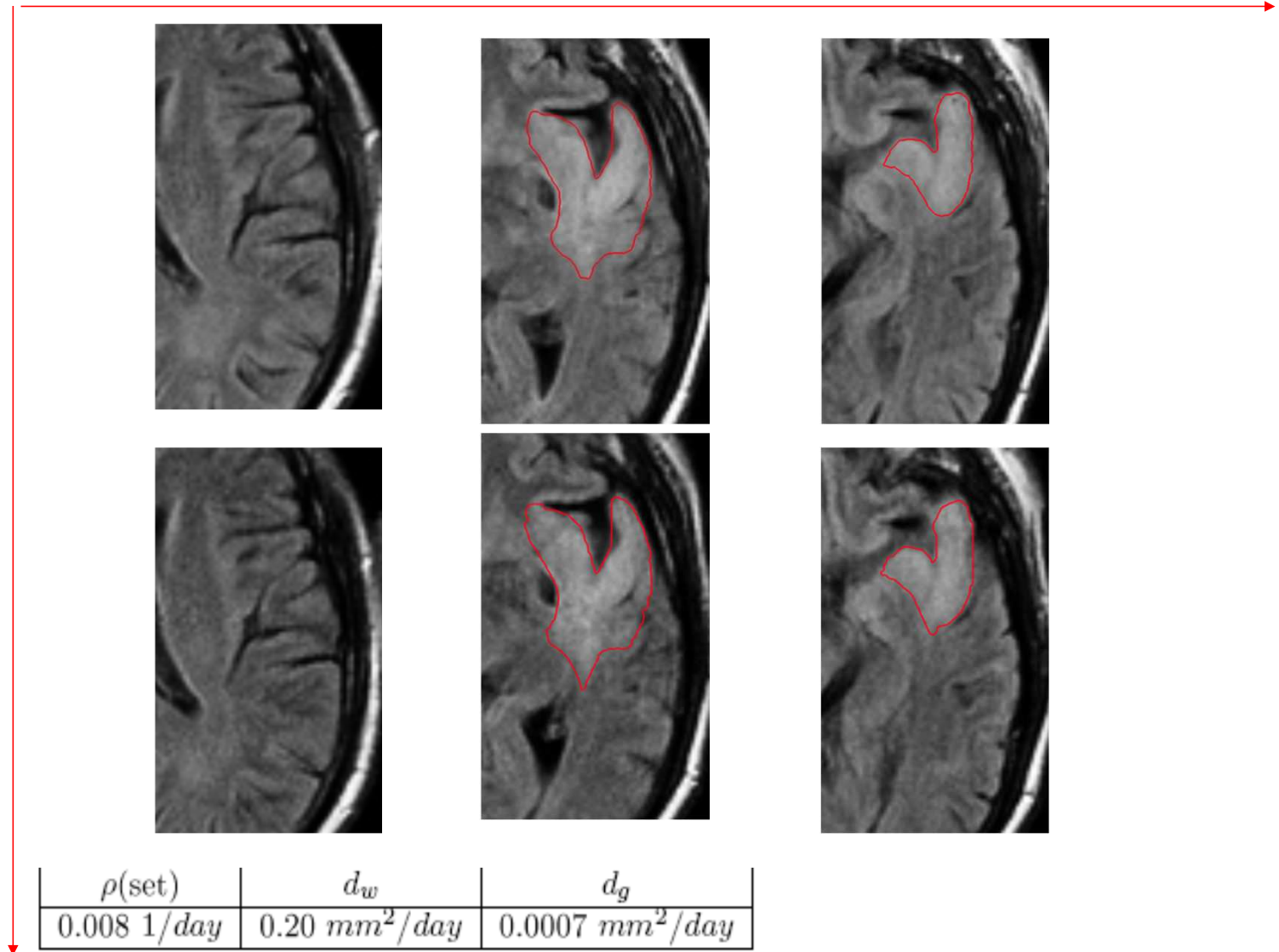


Low Grade Glioma

Space

Learn
on
4 months

time

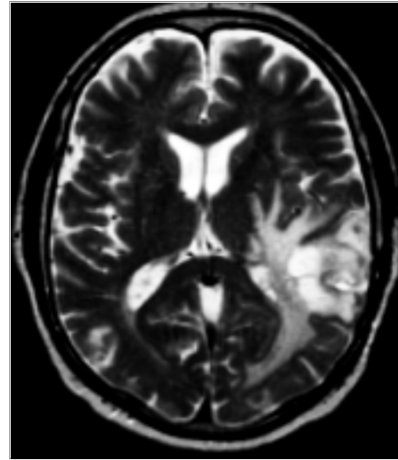


IRM T2 Flair
0.5*0.5*6.5mm
5 time points
MR DTI : 2.5mm (T0)

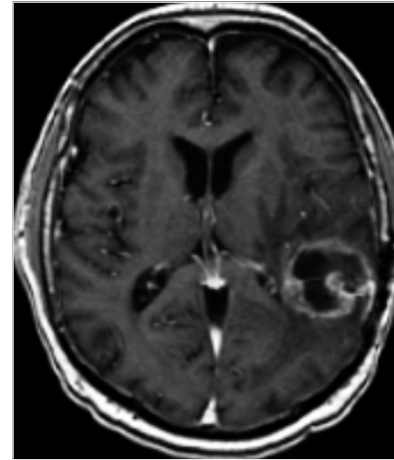
Personnalisation

- 1. Growth speed
- 2. Invisibility Index

T2 and T1 Gd Images

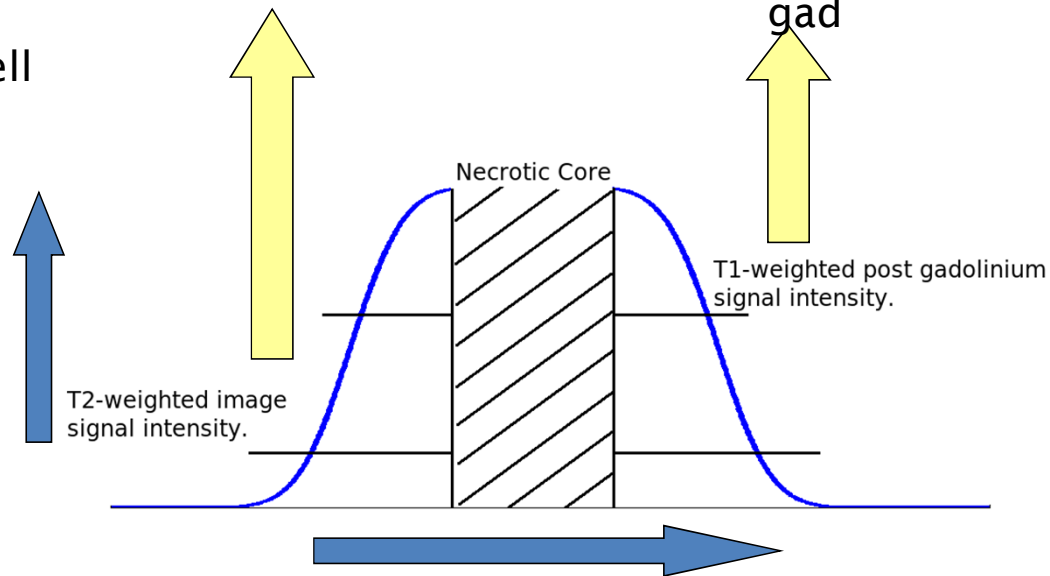


T2w



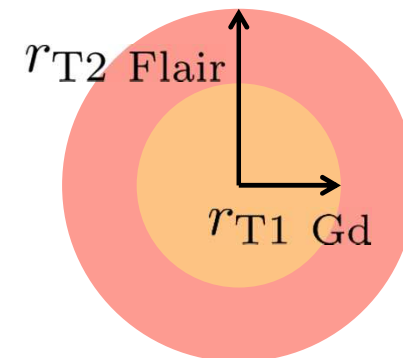
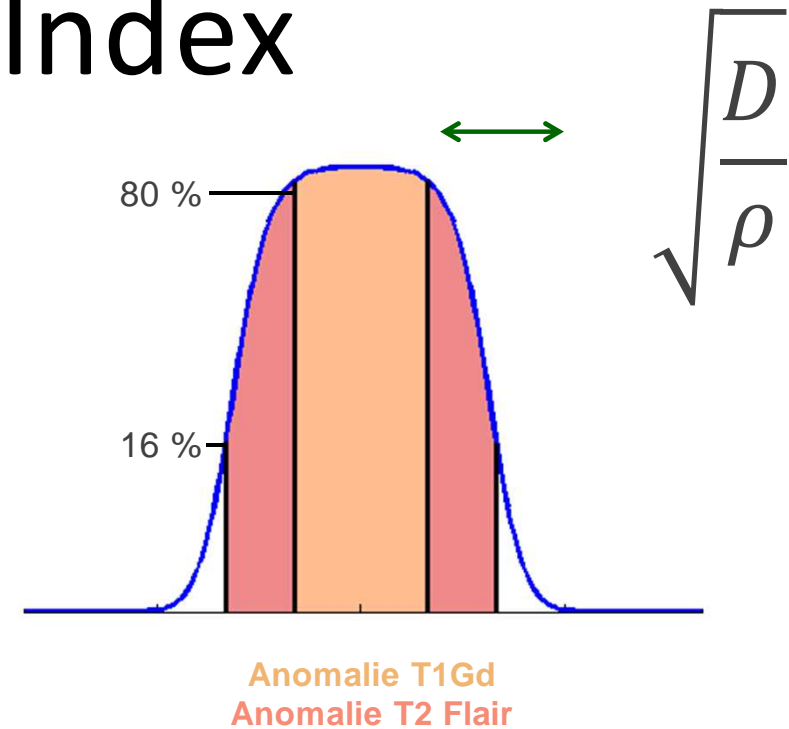
T1w post-gad

Tumor cell density



Invisibility Index

- Empirical formula to estimate Invisibility index from 2 radii of T1Gd & Flair images



Swanson et al.

Predict Infiltration

- Given invisibility index
- Solve an Eikonal equation

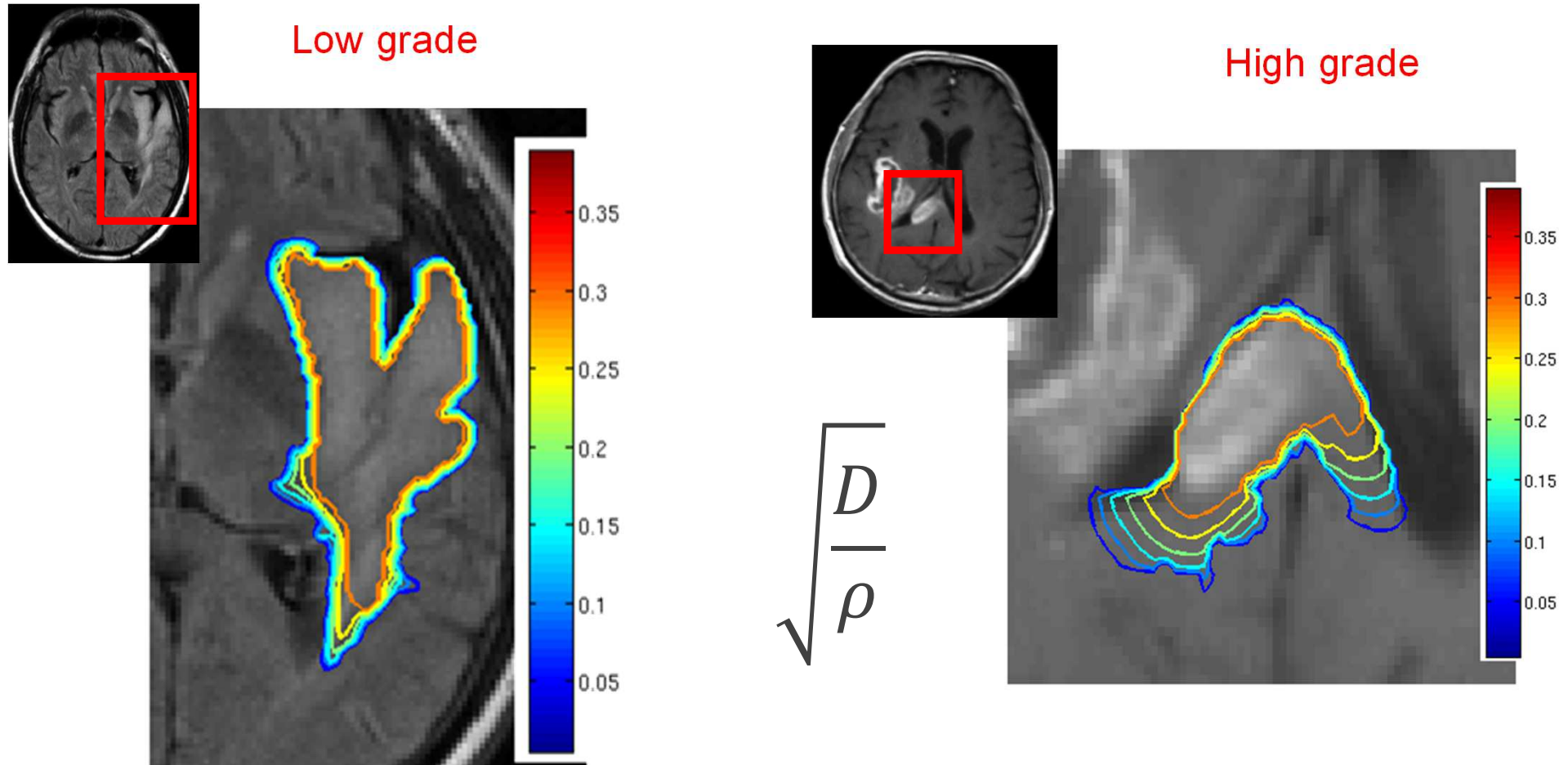
$$\frac{\sqrt{\nabla \tilde{u} \cdot (\mathbf{D} \nabla \tilde{u})}}{\sqrt{\rho \tilde{u} (1 - \sqrt{\tilde{u}})}} = 1, \tilde{u}(\Gamma) = u_0$$

- Anisotropic Fast Marching Algorithm

E. Konukoglu, O. Clatz, P.Y. Bondiau, H. Delingette, N. Ayache. *Extrapolating Glioma Invasion Margin in Brain MRI: Suggesting New Irradiation Margins*. Medical Image Analysis 2010.

Predict Invisible infiltration

Predicted Isodensities between 40% and 1%



E. Konukoglu, O. Clatz, P.Y. Bondiau, H. Delingette, N. Ayache. *Extrapolating Glioma Invasion Margin in Brain MRI: Suggesting New Irradiation Margins.* Medical Image Analysis 2010.

Limitations

- Difficult to have patient DT-MRI
- Assume DT-MRI undisturbed during tumor growth
- Uncertainty in the tumor delineation
- Only model growth no shrinkage
- No mass effect
- Prediction : assumes no effect of therapy

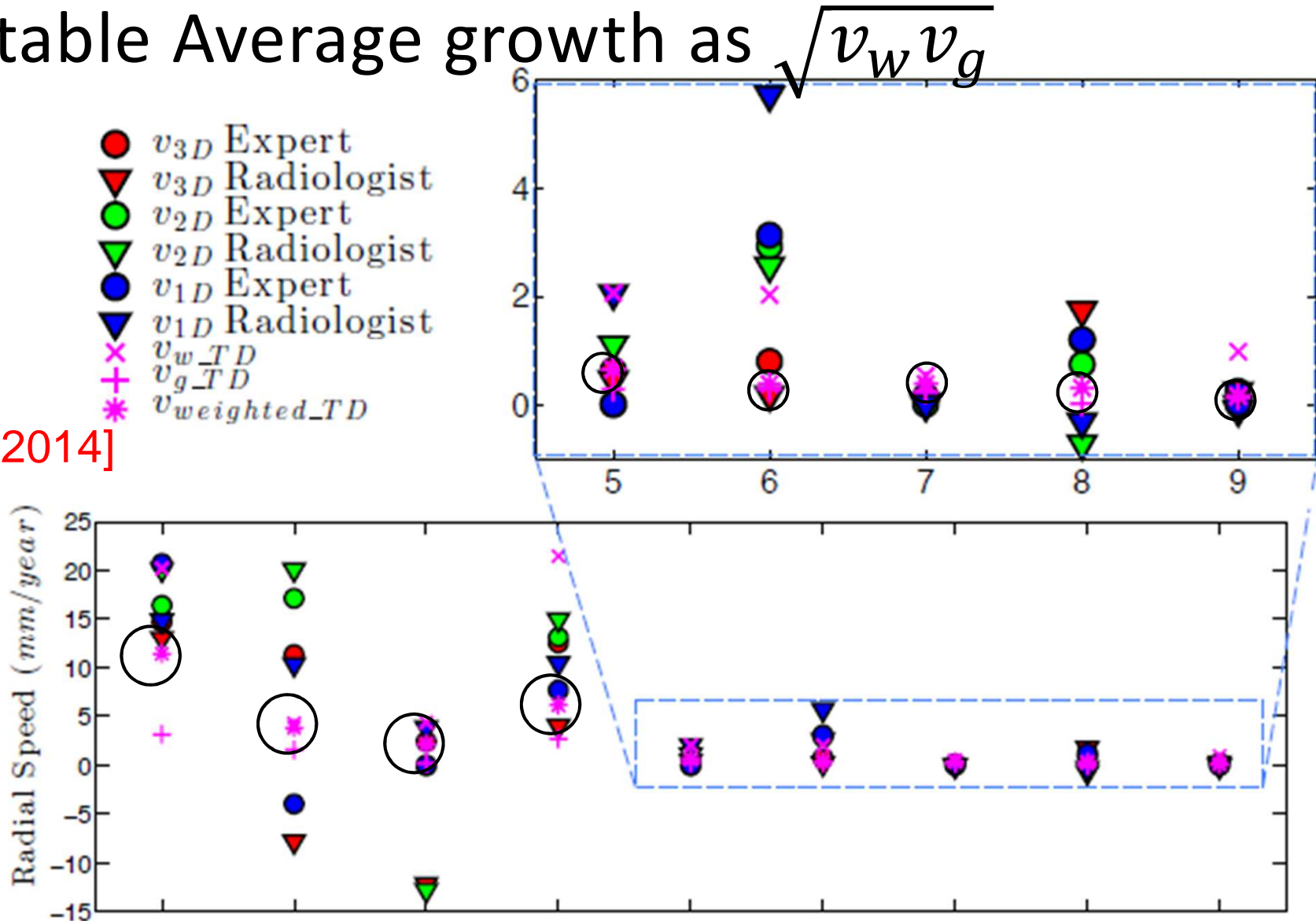
3 Main Problems

1. Quantify the extent of the tumor
2. Quantify the tumor evolution
3. Improve clinical practice
 - Diagnosis
 - Surgery
 - Radiotherapy

Quantification of tumor growth

- Stable Average growth as $\sqrt{v_w v_g}$

[Stretton 2014]

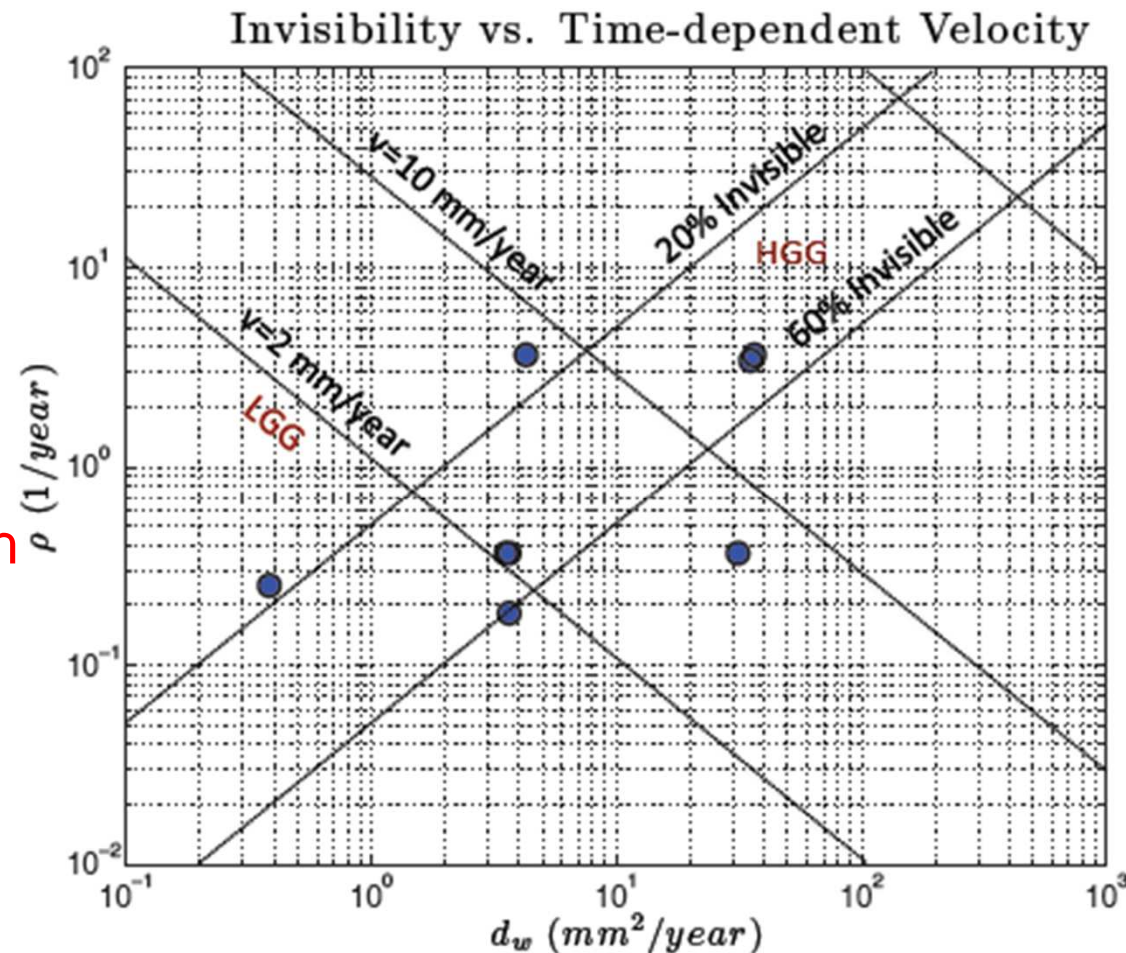


Quantification of tumor growth

- Grading Tumor growth

[Stretton 2014]

Collaboration with
DKFZ



Quantification of tumor growth

- Better prediction of Time-To-Progression than current practice (Recist)

<i>Manual and Model TTP Predictions in 1D</i>			
	Retrospectively Calculated	Radiologist Predicted	Model Predicted
<i>Patient No.</i>	<i>TTP Interval in 1D (days)</i>	<i>TTP_{v1D} (days)</i>	<i>TTP_{v_weighted_A} (days)</i>
1	[1 92]	58	47
2	[92 184]	60	149
3	[221 396]	328	260
4	[315)	174	188
5	[1154)	961	704
6	[189)	78	637
7	[724)	undefined	1171
8	[112)	-1046-	159
9	[915)	2428	1248

[Stretton 2014]

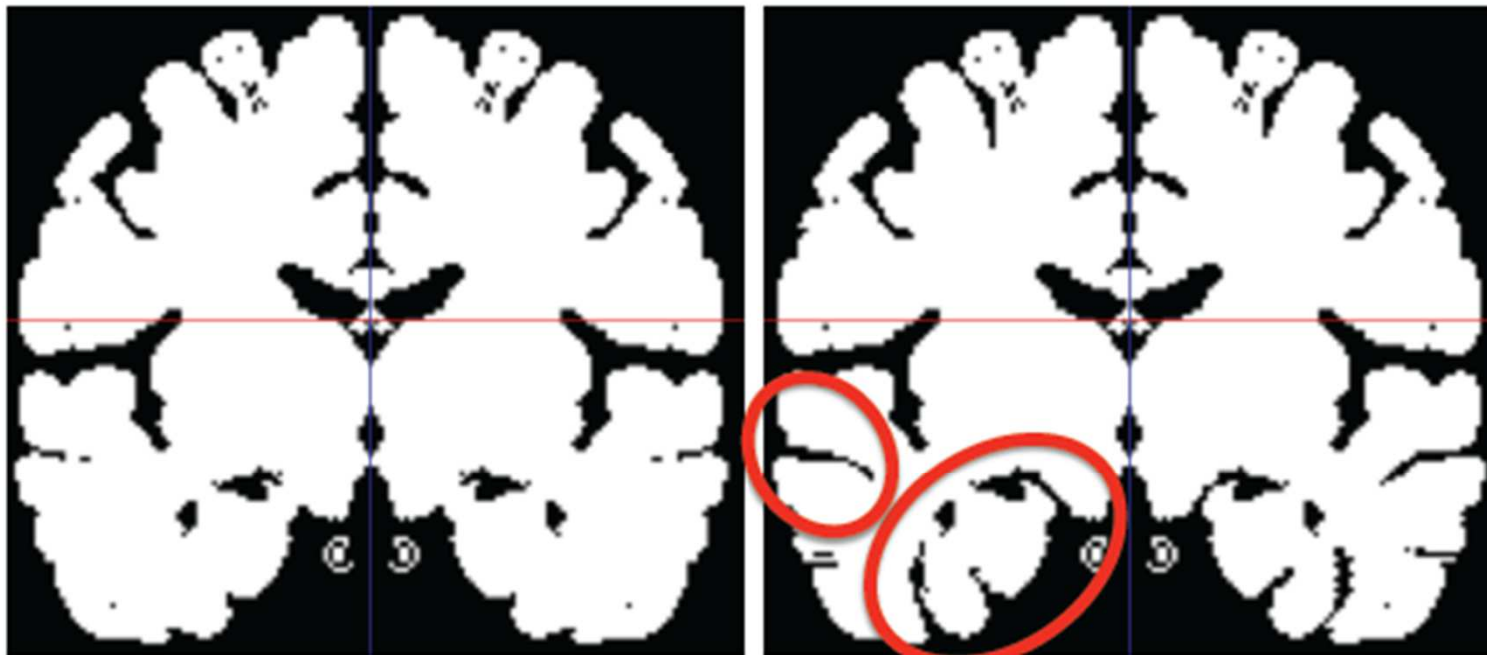
Collaboration with
DKFZ

Definition of anatomical barriers

- Atlas with well defined anatomical barrier

Compare BMs

[Amelot et al.-Mandonnet 2014]

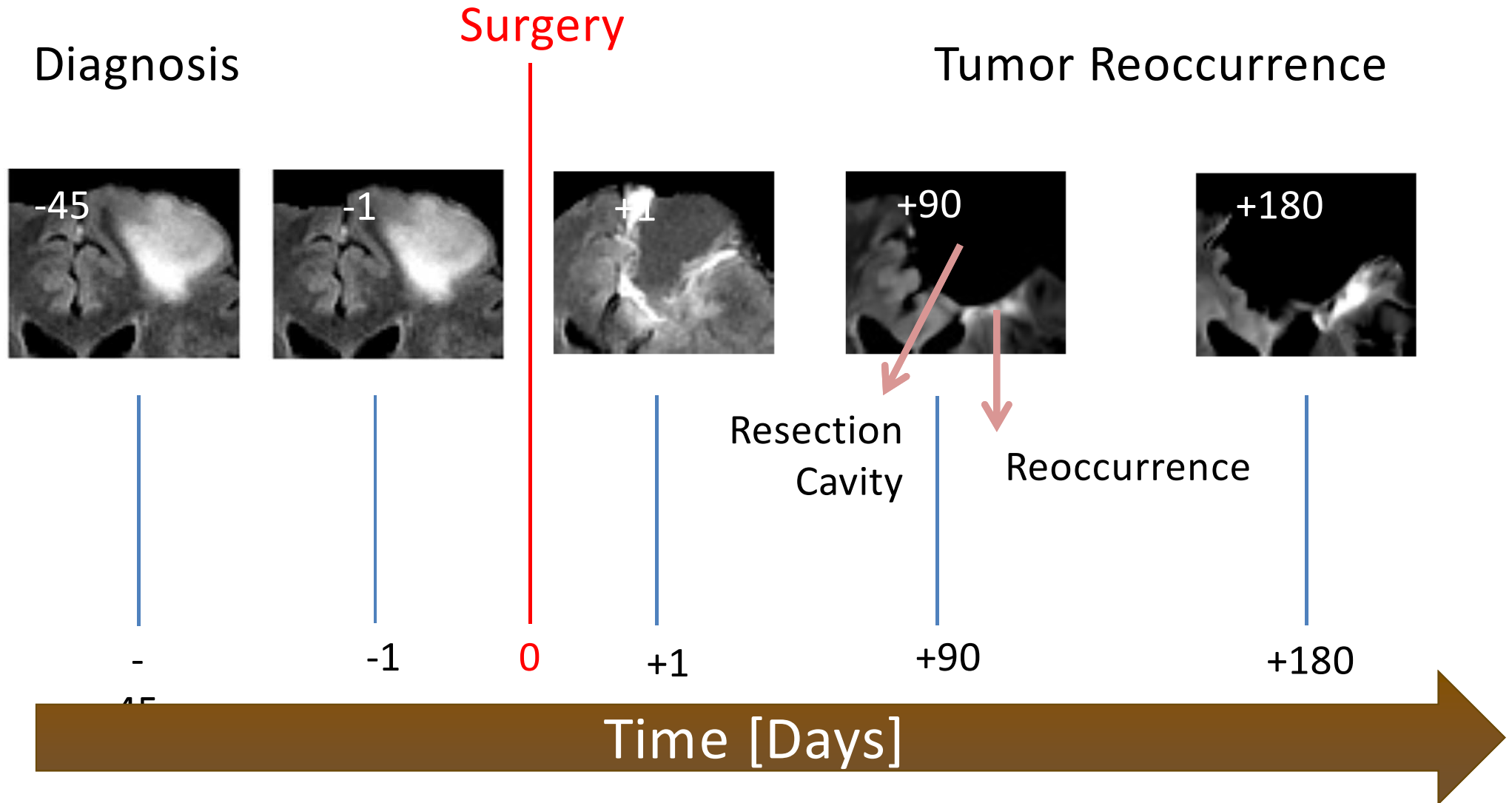


MNI Original BM

Corrected BM

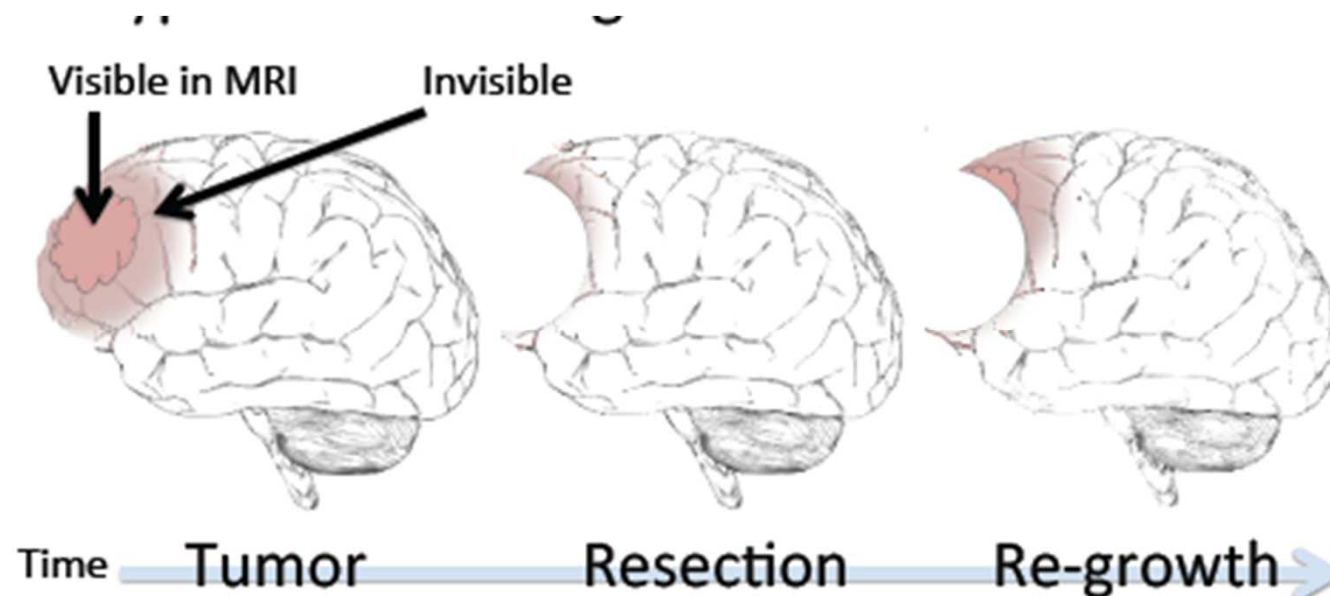
Based on MNI 152

Recurrence after surgery



Recurrence after surgery

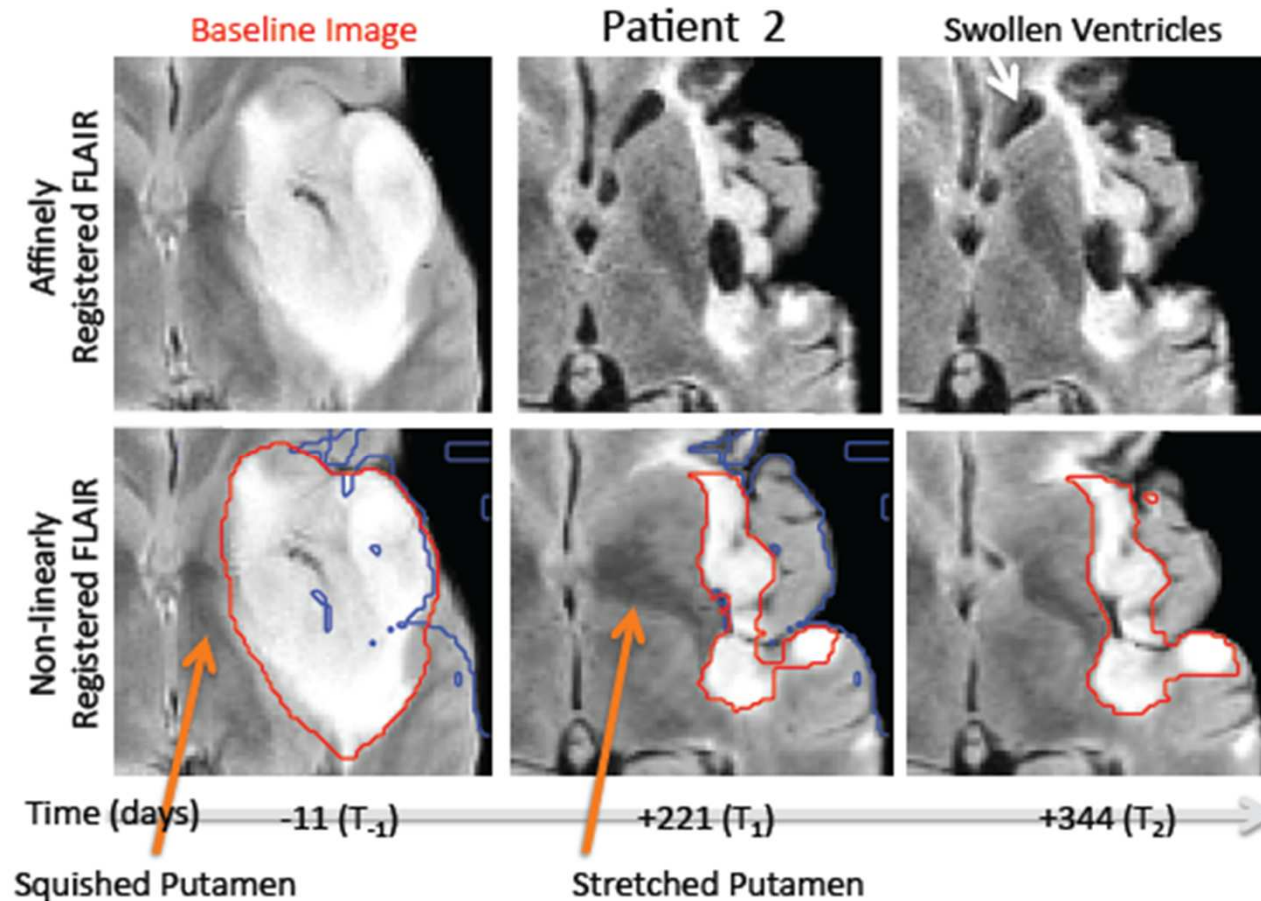
- Is it possible to :
 - Predict the location of recurrence ?
 - Predict the time of recurrence ?
 - Predict the extent of infiltration ?



Recurrence after surgery

- Difficulty due to large deformation after surgery
- Effect of therapy

[Stretton 2014]



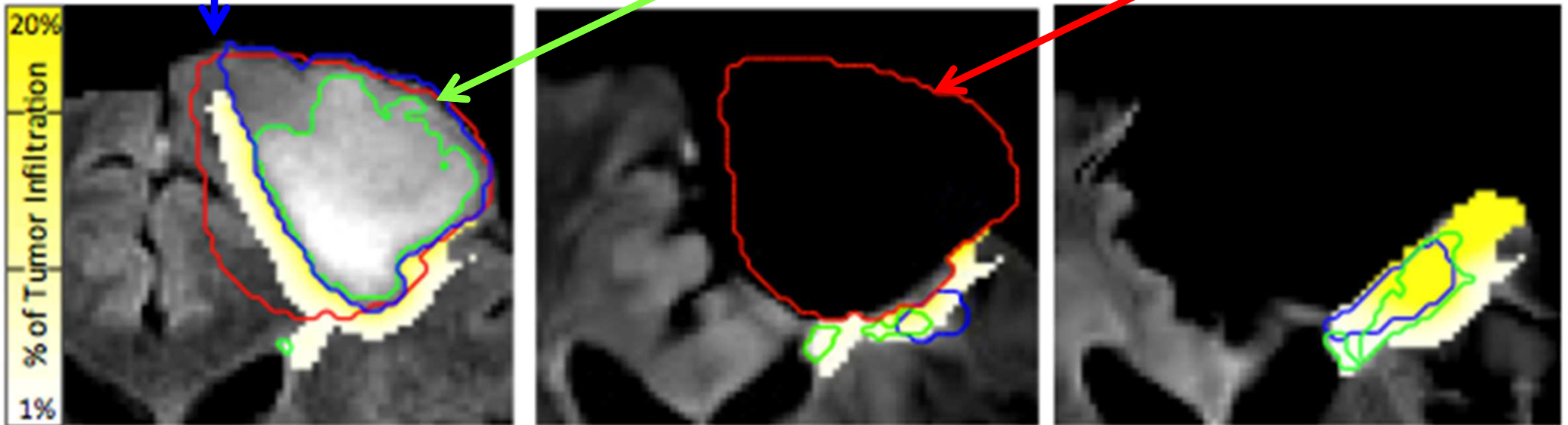
Qualitative Results

[Stretton 2014]

Neurosurgeon's
Tumor Segmentation

Hyper-Intense
Thresholded Voxels

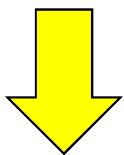
Neurosurgeon's
Cavity Segmentation



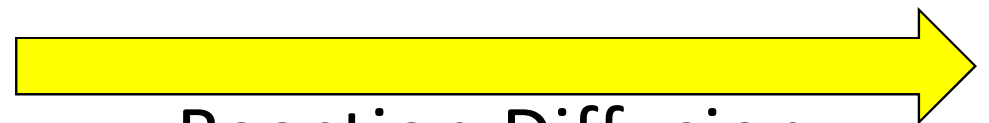
(a) Day -1

(b) Day +74

(c) Day +172



Cell Density Initialization



Reaction Diffusion
Algorithm

Image guided therapy

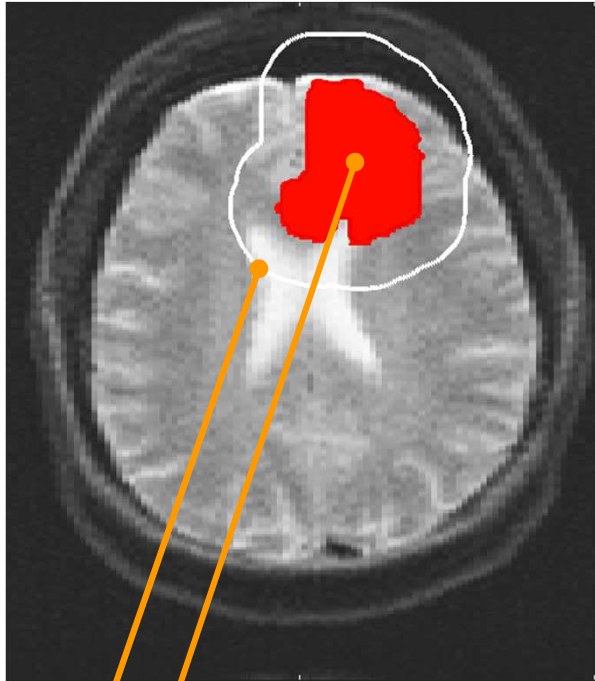


Cyberknife au CAL, Nice



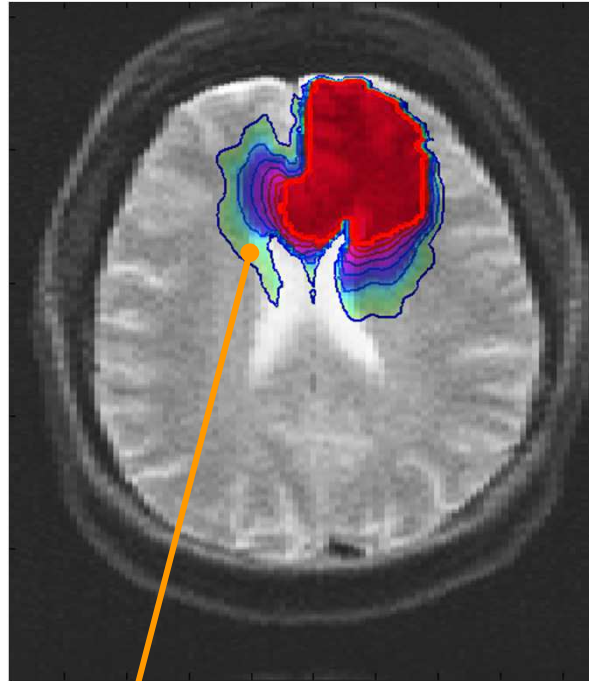
**MGH
Boston**

Dosimetry planning

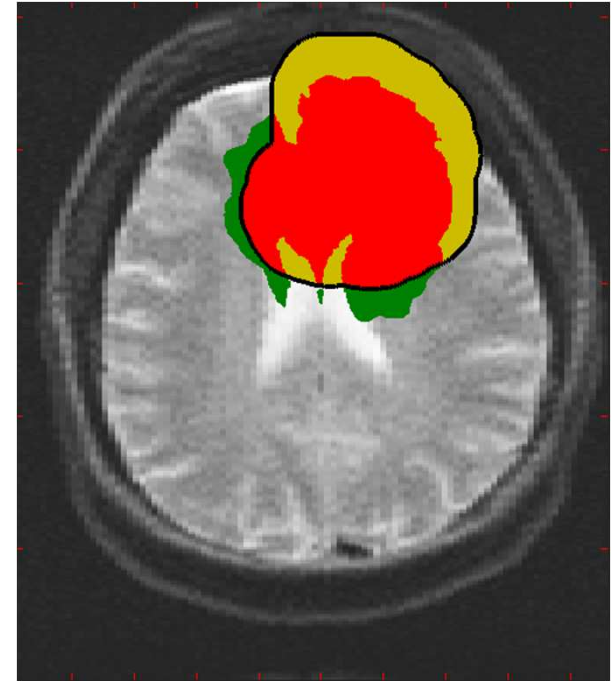


Visible Tumor

Constant Margin (2cm)



Invisible Infiltration



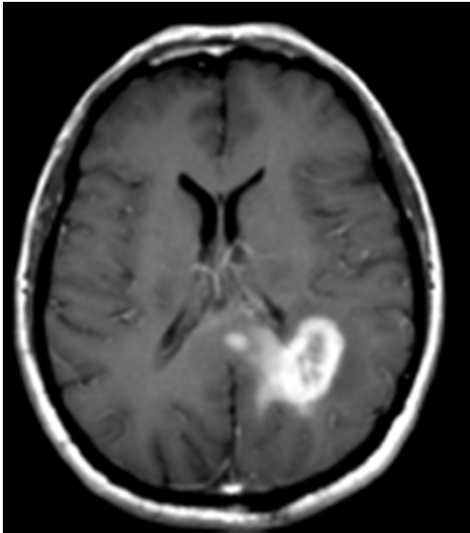
Targeted Tumors

Targeted Healthy Tissue

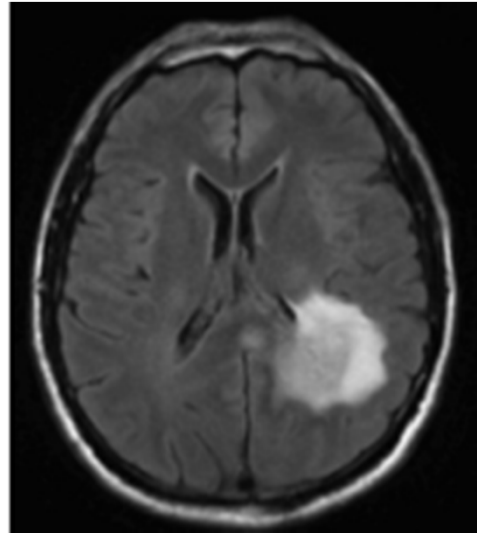
Non targeted tumor

Predicting infiltration

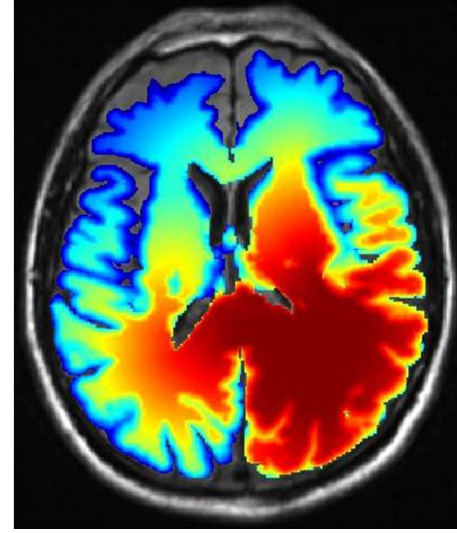
$$\frac{\partial u}{\partial t} = \nabla \cdot (D \nabla u) + \rho u(1 - u)$$



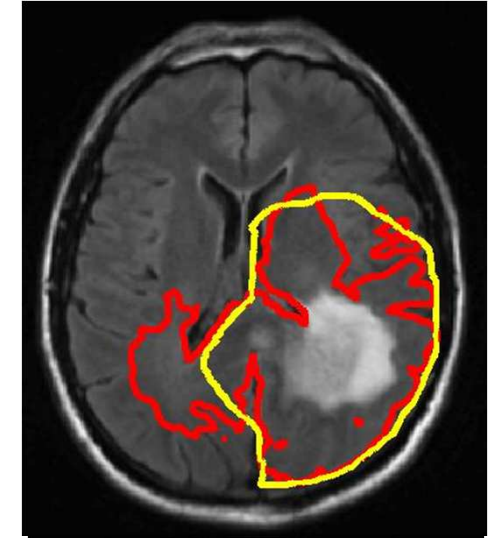
T1 Gd



T2 Flair



Log-density
Of predicted
tumor cells



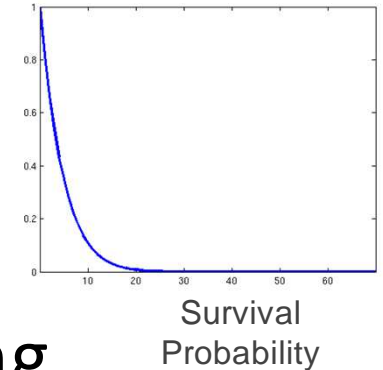
Simulated
Target Volume

Standard
Target Volume

J Unkelbach B Menze, E Konukoglu, F Dittmann, N Ayache, H Shi, *Radiotherapy planning for glioblastoma based on a tumor growth model: implications for spatial dose redistribution*. *Physics in Medicine and Biology*, December 2013.

Same
Volume

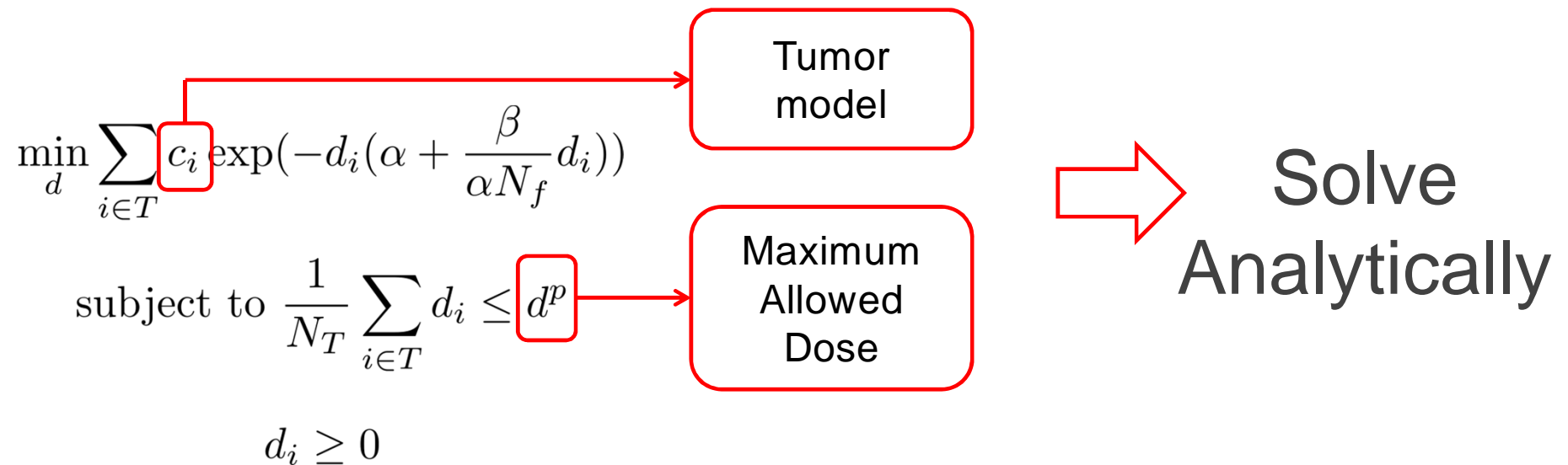
Radiotherapy Dose Planning



- Linear Quadratic Model

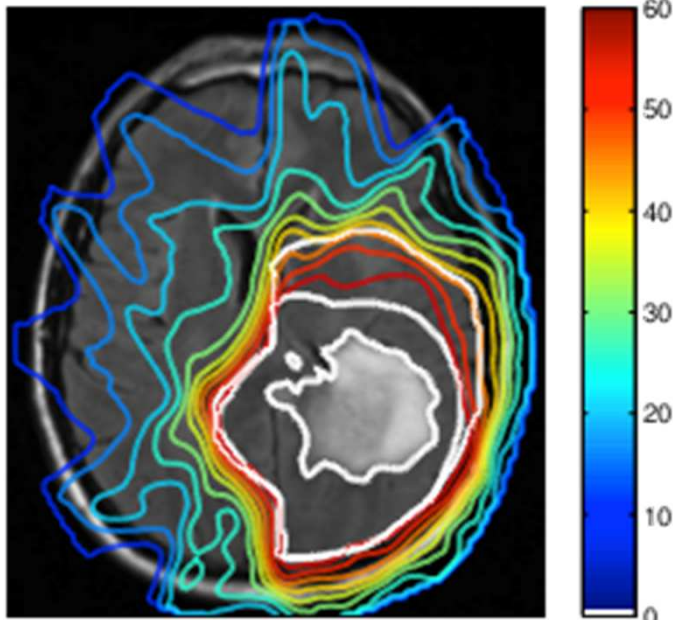
$$P(\text{cell survival} | \text{Dose } d, \text{ Fractions } N_f) \approx \exp\left(-d \left(\alpha + \frac{\beta}{N_f} d\right)\right)$$

Optimize d_i at each voxel to minimise surviving tumoral cells



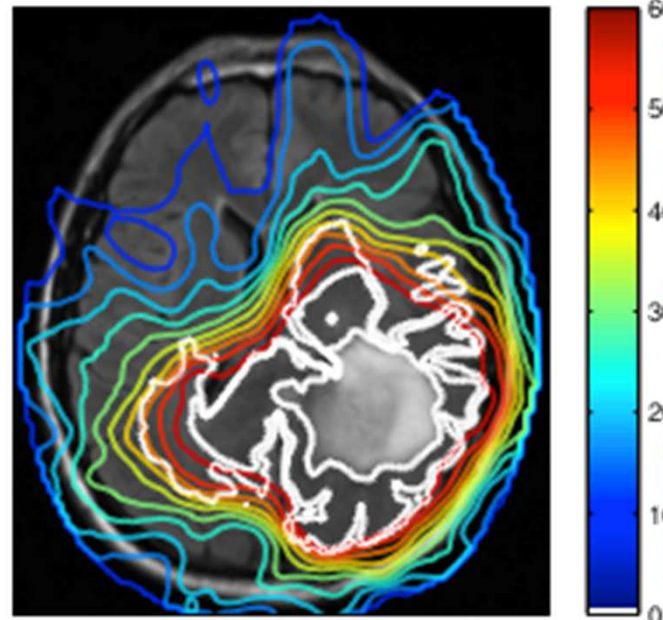
Radiotherapy

Standard Dosimetry
Tumor Contours
+ margin



GTV : 60 Gy - T1Gd + 2cm
CTV 46 Gy - T2Flair +
1.5cm

Optimized Dosimetry
For the personalized
growth model



Same total dose

MGH
Boston

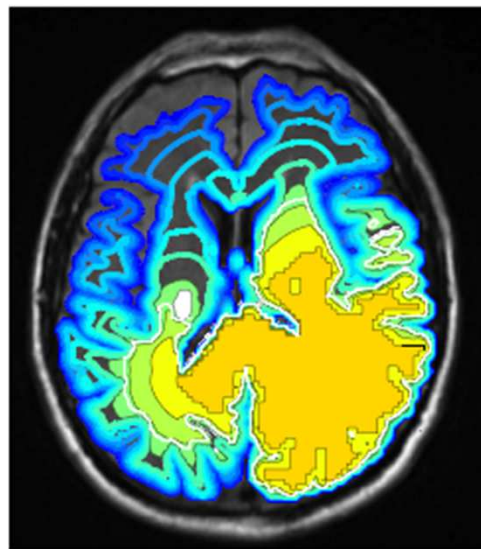


J Unkelbach, B Menze, E Konukoglu, F Dittmann, M Le, N Ayache, H Shi. *Radiotherapy planning for glioblastoma based on a tumor growth model: improving target volume delineation.*

Physics in Medicine and Biology, December 2013.

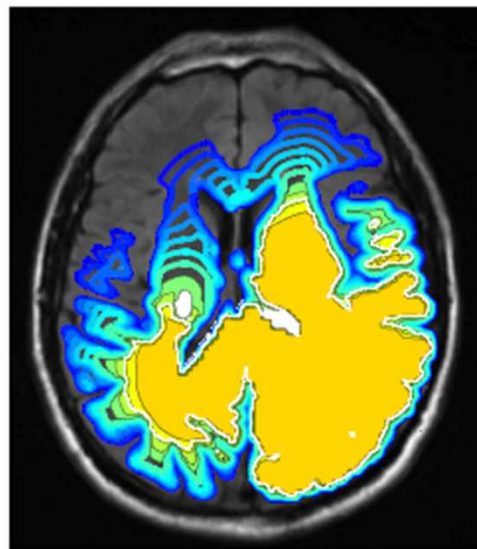
Impact of invisibility index

$$\sqrt{\frac{D}{\rho}} = 4,2 \text{ mm}$$

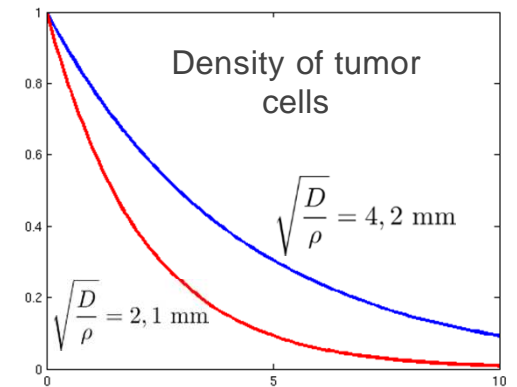


Computed Dose (Gray)

$$\sqrt{\frac{D}{\rho}} = 2,1 \text{ mm}$$



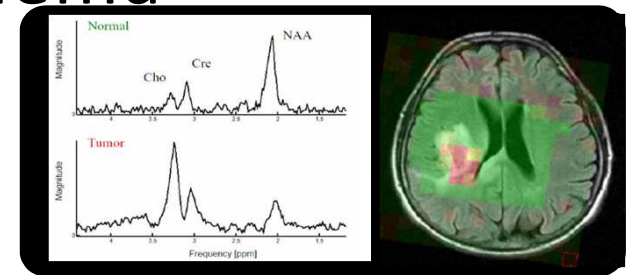
Compute dose(Gray)



$$\frac{\partial u}{\partial t} = \nabla \cdot (D \nabla u) + \rho u (1 - u)$$

Challenges

- Understanding Imaging of brain tumor :
 - vasogenic edema vs infiltrated edema
 - Imaging tumor cell density



MR Spectroscopy

- Multiple compartment tumor growth models & more imaging (PET, perfusion)
- Better account of anisotropic diffusion
- Apply on Large cohort of patients

Take Home Messages

- Objectives of Model Personalization
- Data Assimilation vs Data Regularization
- Notion of parameter Observability
- Model as a tool
- Glioma growth models
- Personalization of speed and invisibility index
- Clinical applications for radiotherapy

Acknowledgments

- Colleagues : B. Ribba, J. Unkelbach, O. Clatz, B. Menze, N. Ayache
- Phd Students : E. Konukoglu, M. Lê, N. Cordier, E. Stretton
- Medical Collaborators : E. Mandonnet,
- Funding : Inria, ERC

Take Home Messages

- Geometric & Biophysical Personalization
- Notion of parameter Observability
- Sensitivity Analysis
- Calibration & Parameter Optimization techniques
- Model as a tool
- Uncertainty estimation