

Microfluidics 2019  
Sète, Octobre 2019



# Blood flow and mass transfers in brain microcirculation

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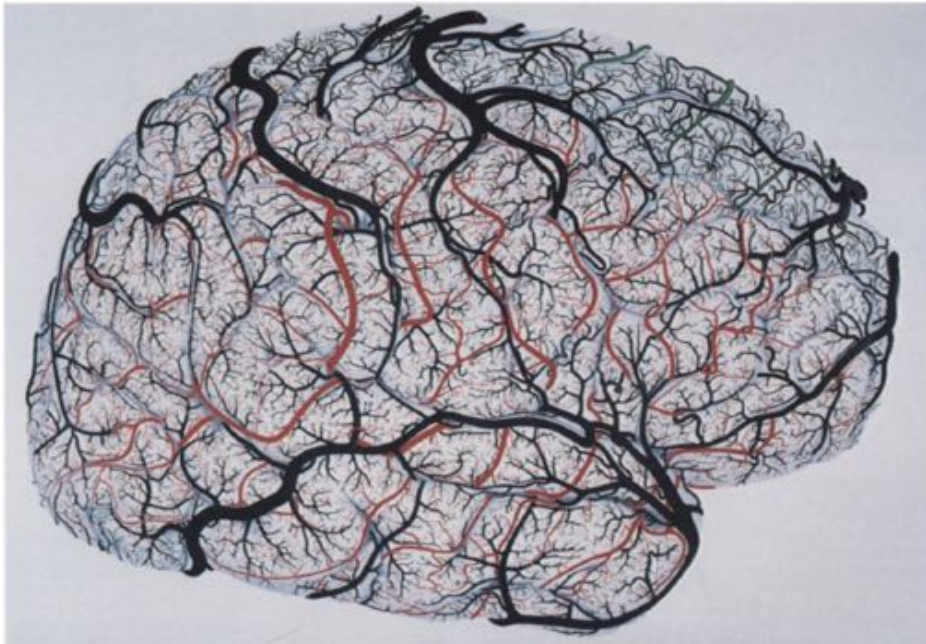


Porous and Biological Media  
Fluid Mechanics Institute of Toulouse  
UMR CNRS INPT UPS 5502

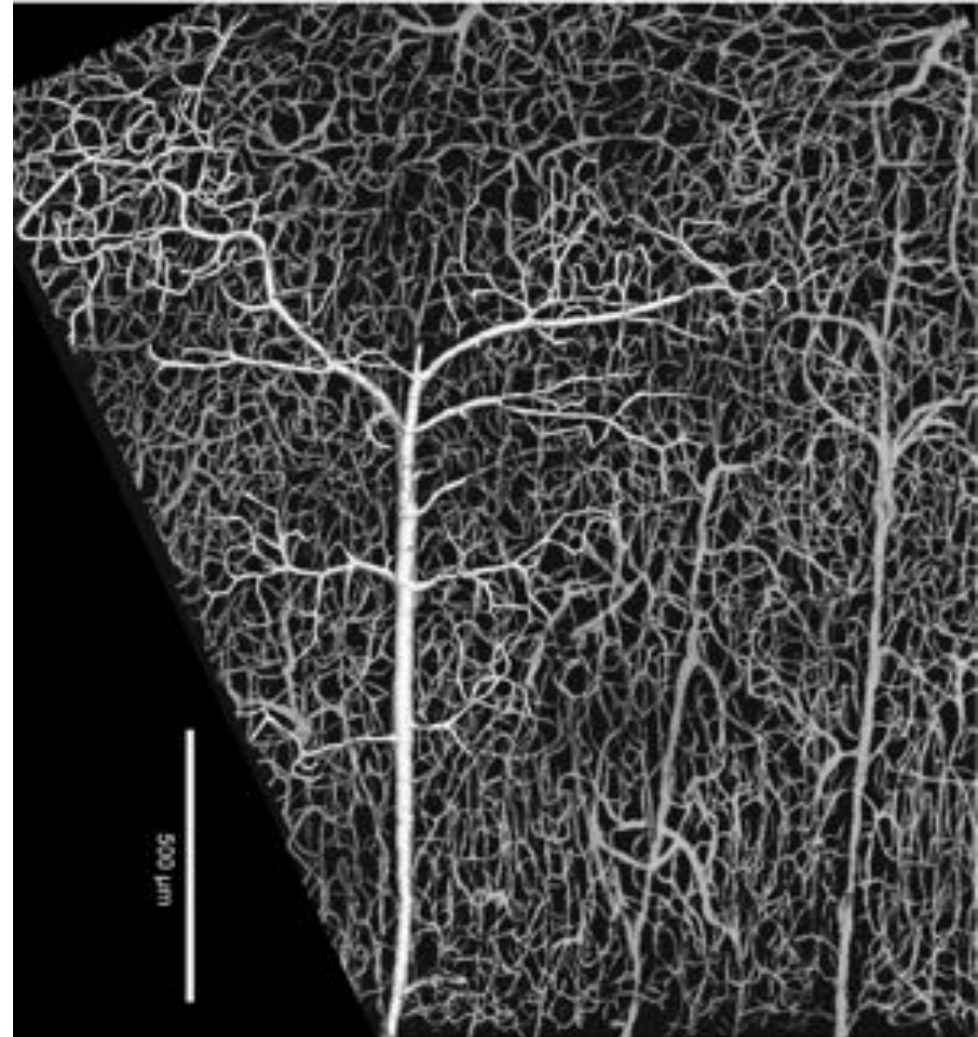
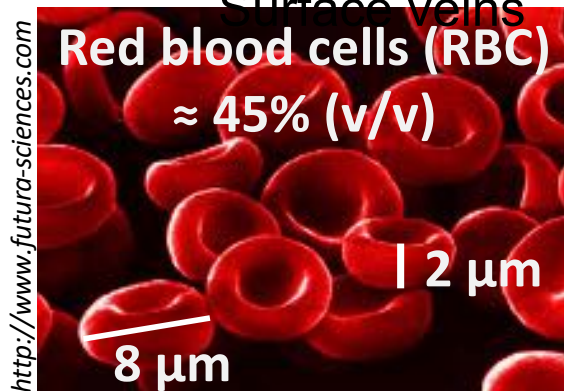
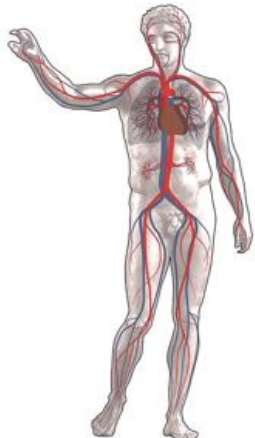


# Brain microcirculation

## Structure



Duvernoy et al. Brain Res Bull 1981 **Surface arteries**



Cassot et al. Microcirculation 2006

↓ Surface veins    ↖ Surface arteries

# Outline

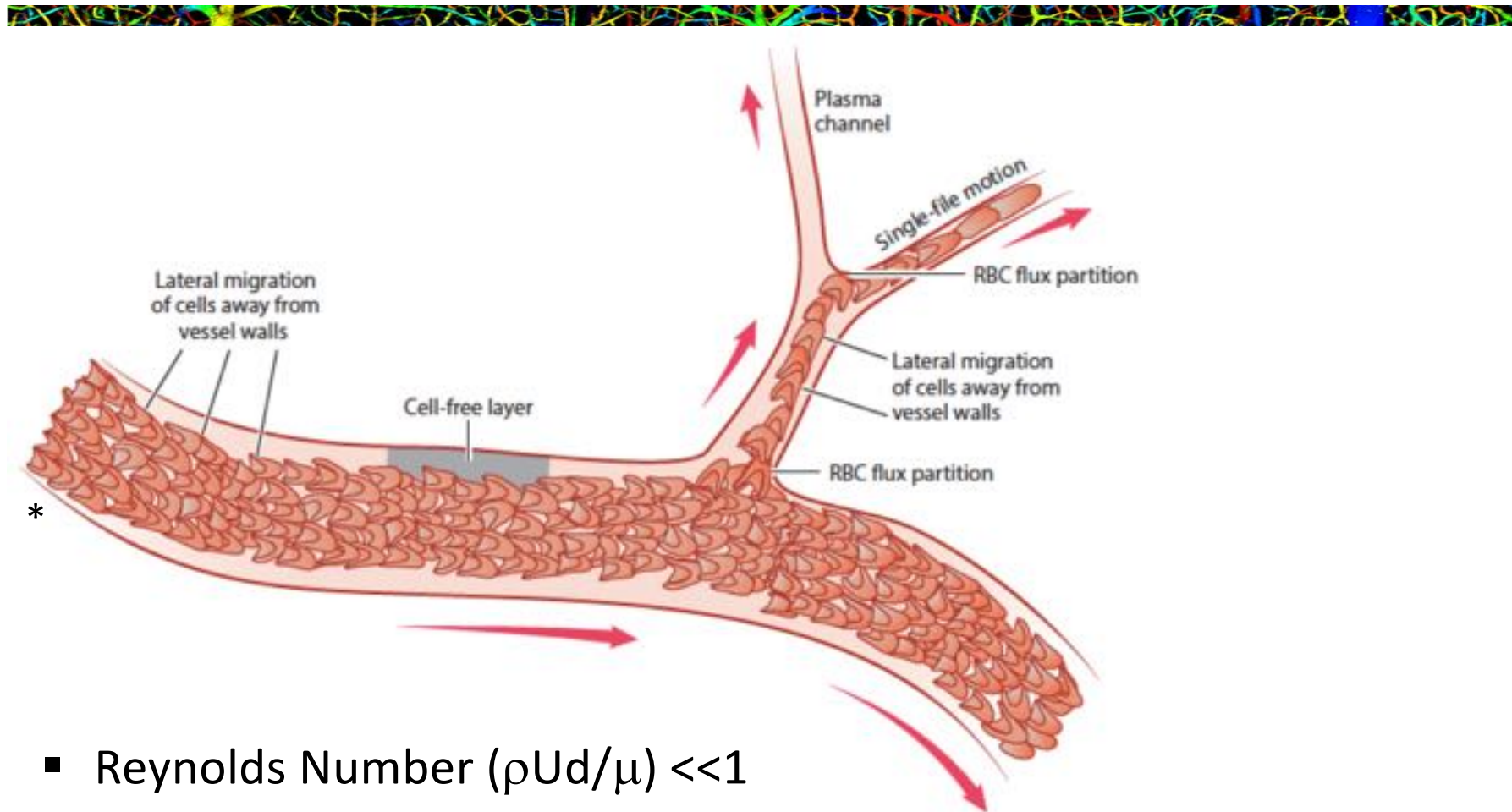


- **Brain versus other organs**
  - What is generic ?
  - What is specific ?
- **Why study brain microcirculation ?**
  - In health
  - In disease
- **Investigation tools and associated scales**
- **Blood flow in networks**
- **Mass transfers in networks**
- **Blood flow at organ scale**



# What is generic ?

## The physics of red blood cell flows

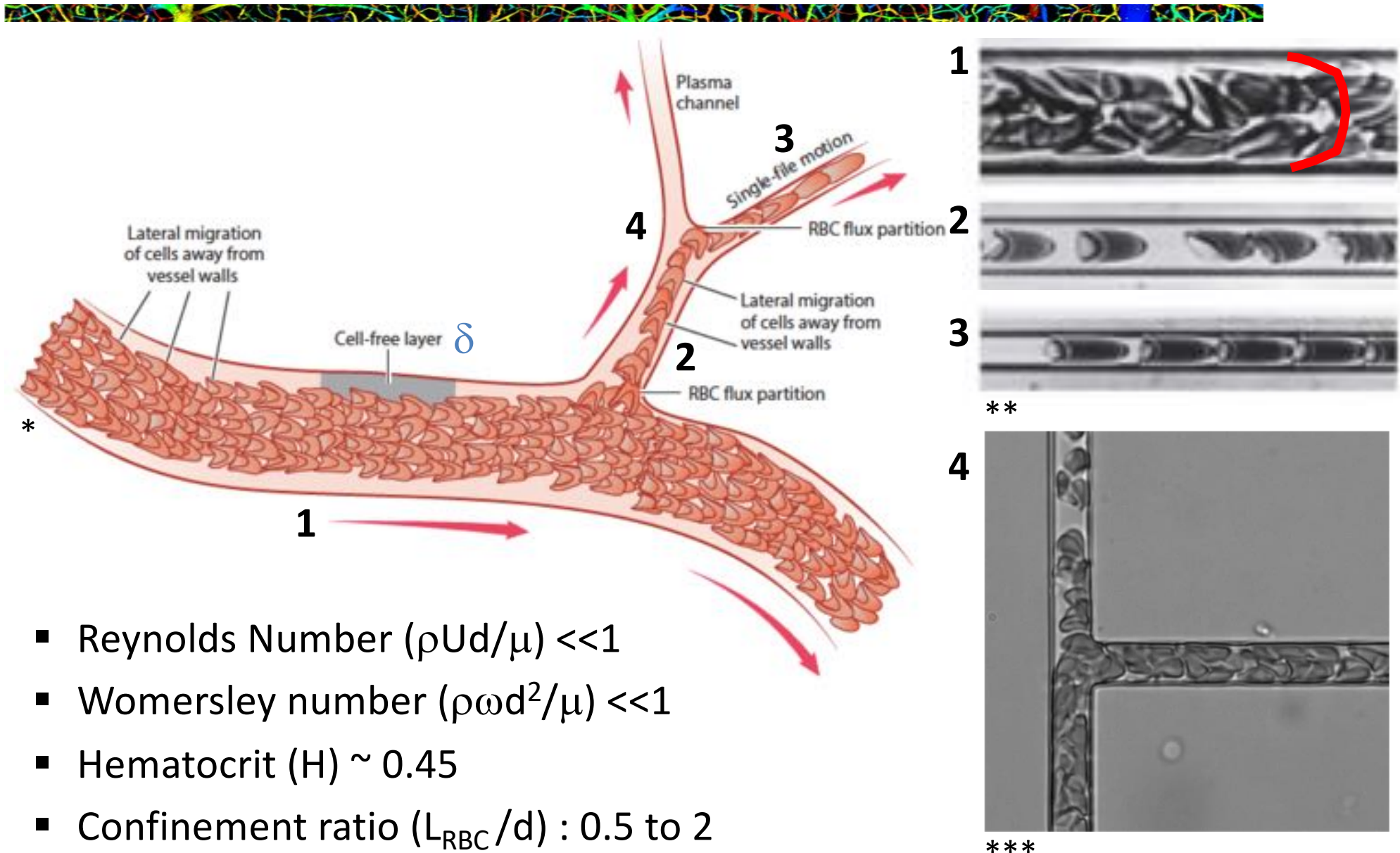


- Reynolds Number ( $\rho U d / \mu$ )  $\ll 1$
- Womersley number ( $\rho \omega d^2 / \mu$ )  $\ll 1$
- Hematocrit ( $H$ )  $\sim 0.45$
- Confinement ratio ( $L_{RBC} / d$ ) : 0.5 to 2



# What is generic ?

## The physics of red blood cell flows

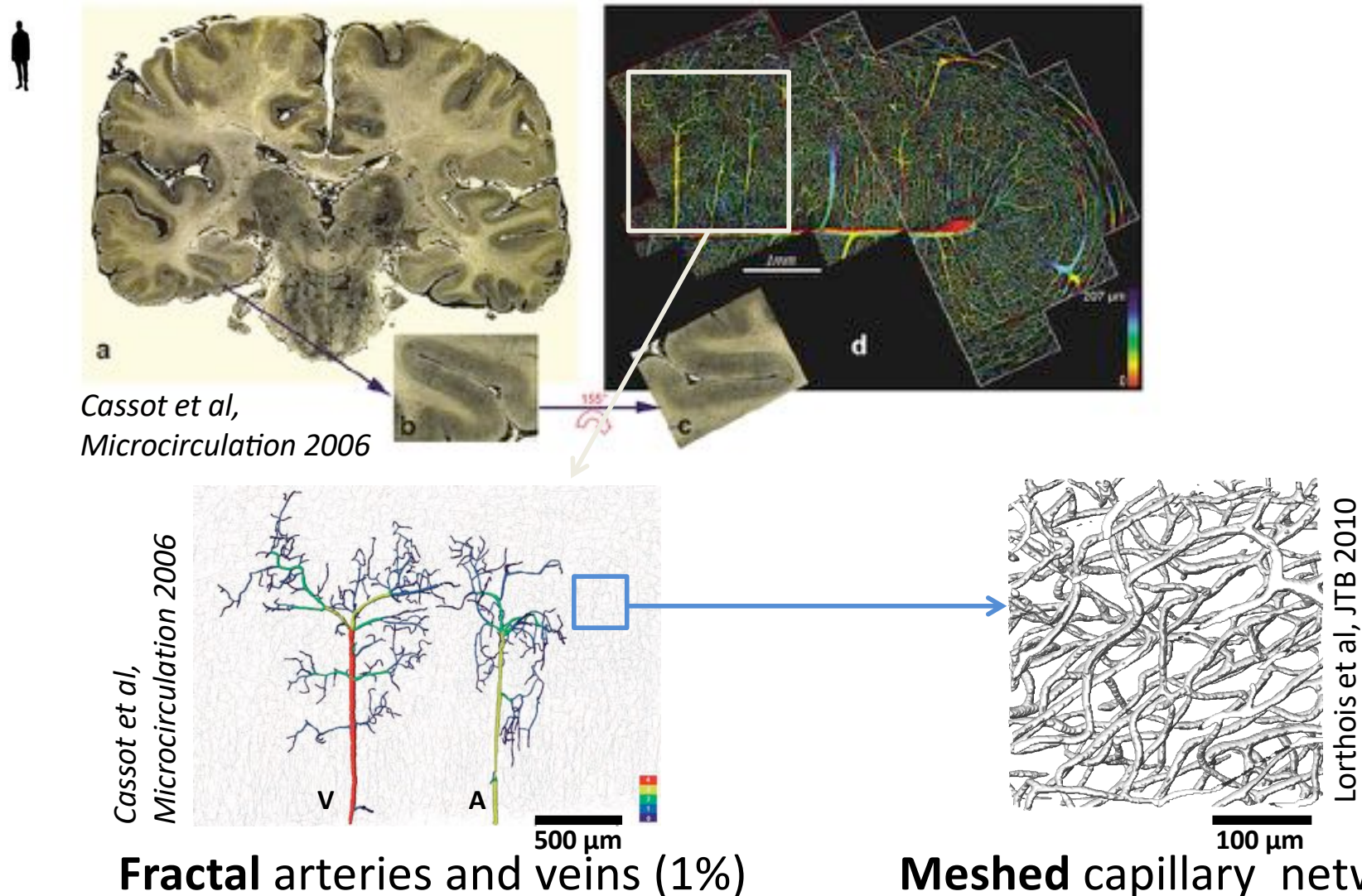


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- Hematocrit (H)  $\sim 0.45$
- Confinement ratio ( $L_{RBC} / d$ ) : 0.5 to 2

# What is generic ?

The general organization of the vascular architecture

- Brain networks are multi-scale, with two components

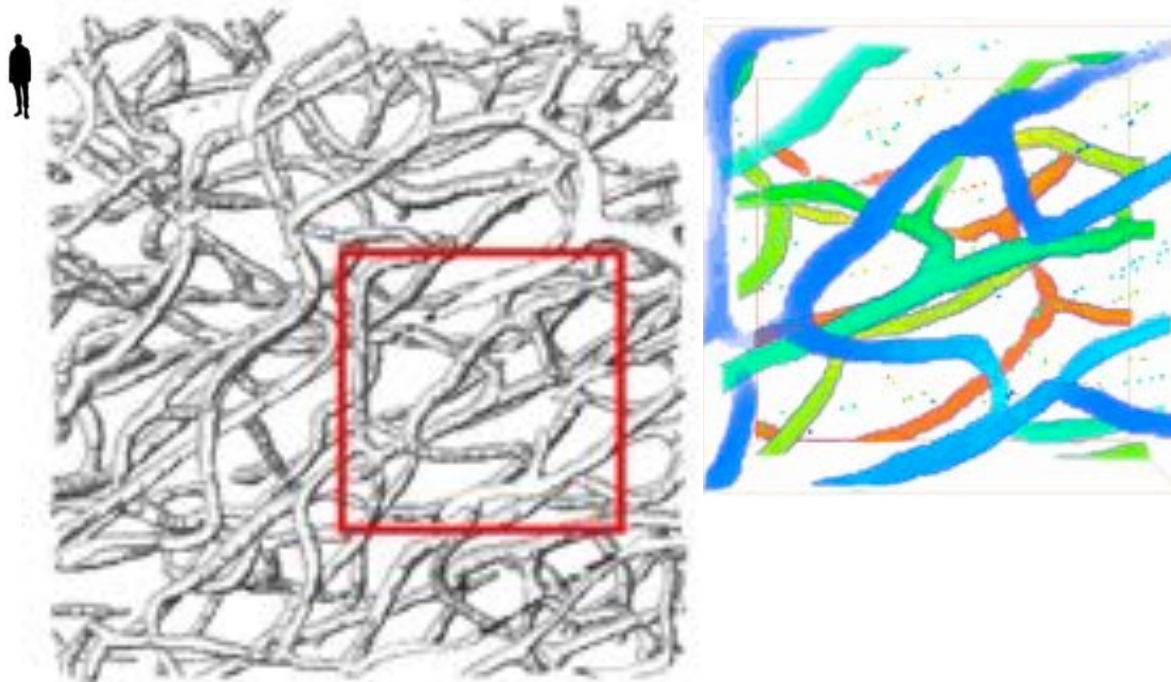




# What is generic ?

## The general organization of the vascular architecture

- Capillary networks in the brain are 3-connected, space-filling



- Looped and random at small scale
- Homogeneous & Space-filling above  $\sim 50\mu\text{m}$   
(Each point in the tissue is close to a capilla)

→ REV / Porous medium

- Volume density  
 $\sim 2\%$
- Exchange surface  
 $\sim 5\text{ mm}^2/\text{mm}^3$
- Cumulative length  
 $\sim 500\text{ mm}/\text{mm}^3$

Cassot et al. 2006

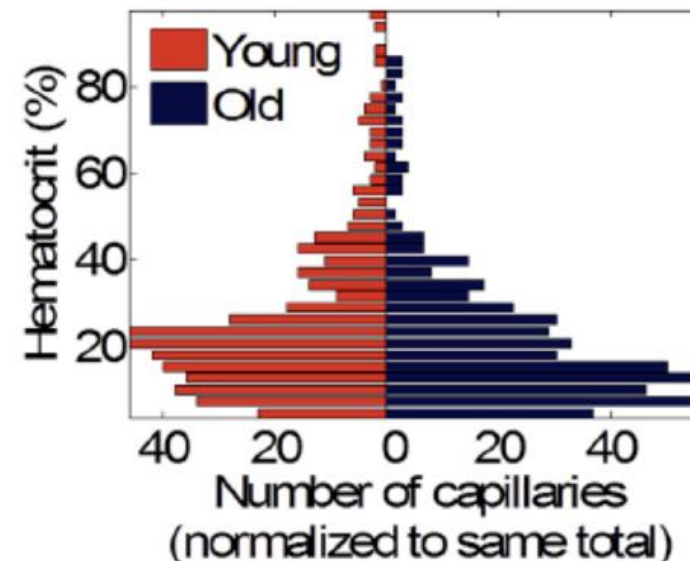
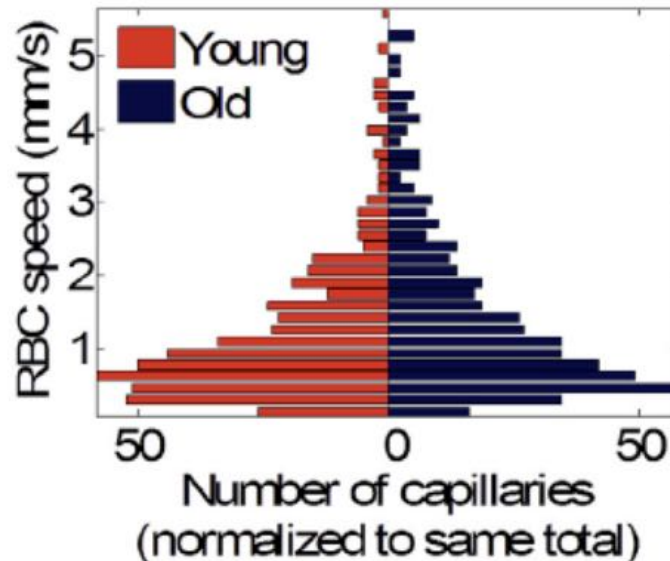
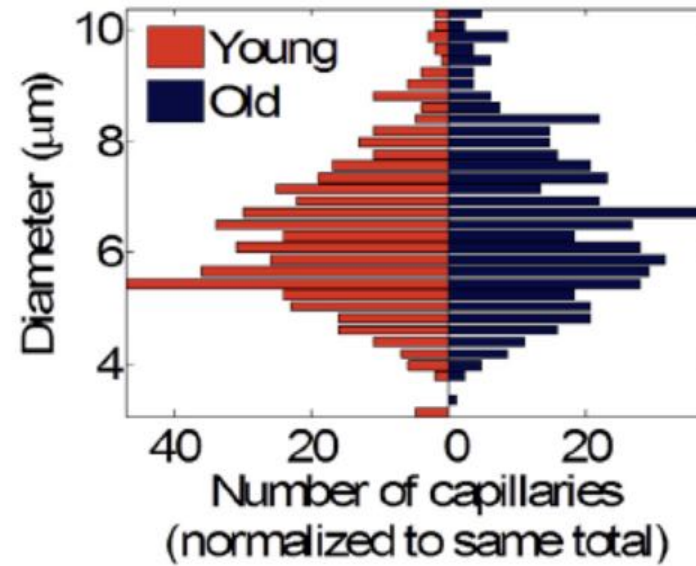
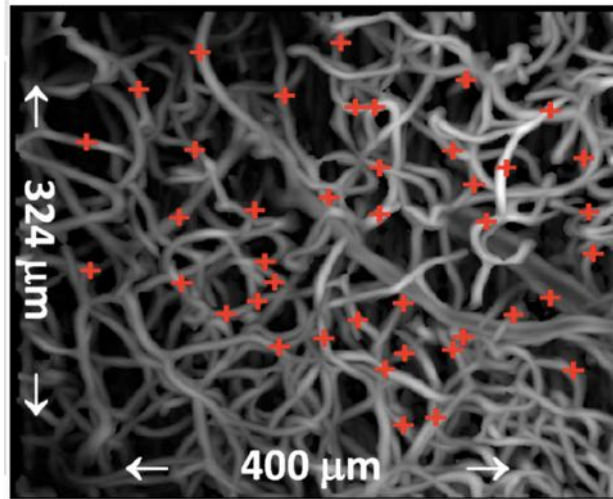
Heinzer et al. 2006

Risser et al. 2007



# What is generic ?

The tremendous heterogeneity of hemodynamic parameters

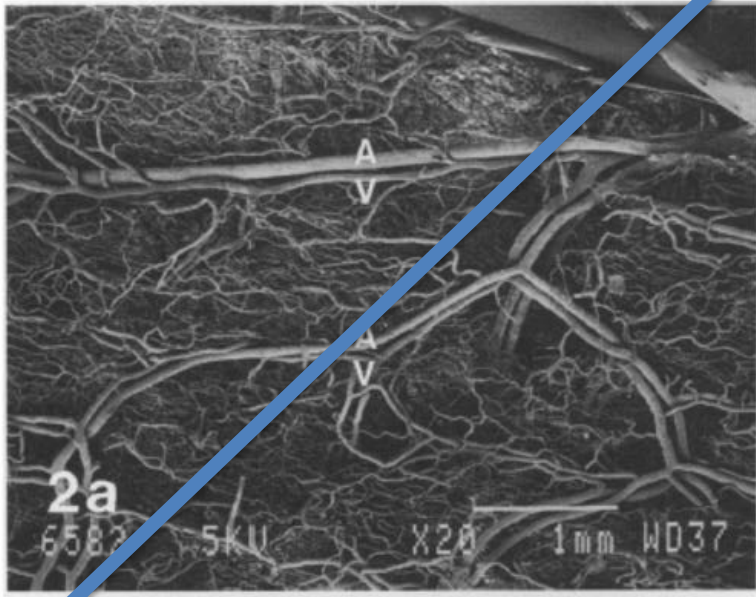


MICE

# What is specific ?

## The details of the vascular architecture

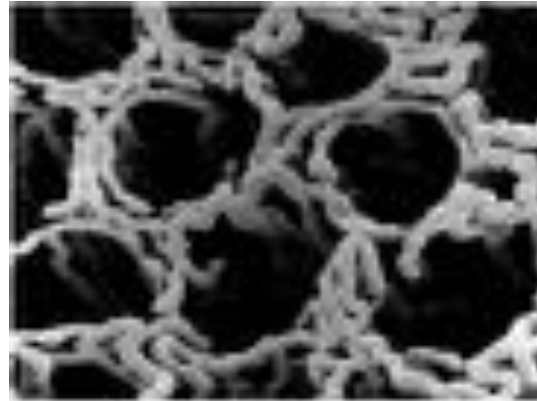
Arterioles and venules



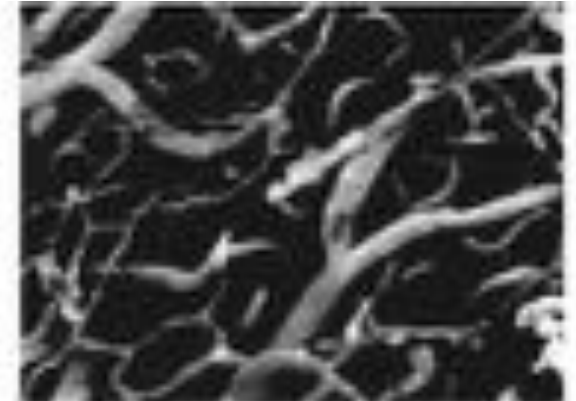
**Human temporary muscle**

Cheung et al. *J. Anatomy* 1996

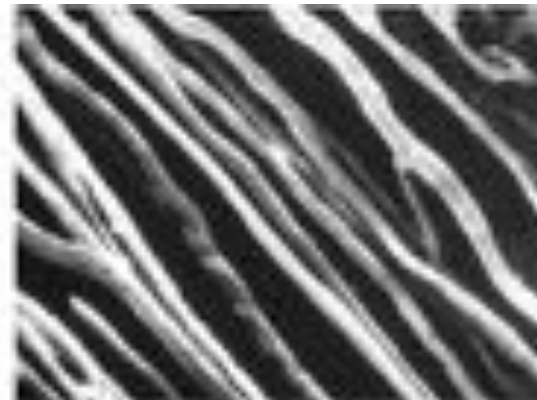
Capillary network



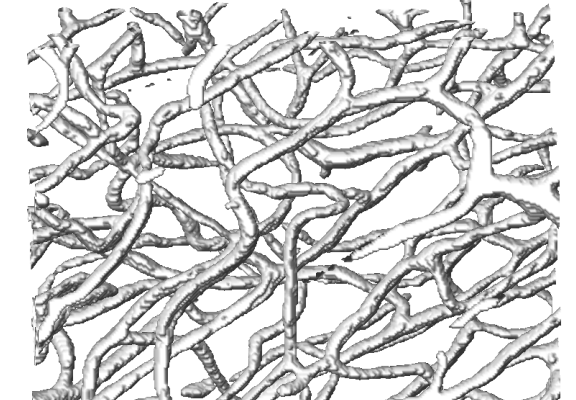
Colon



Subcutis



Skeletal muscle



Brain cortex

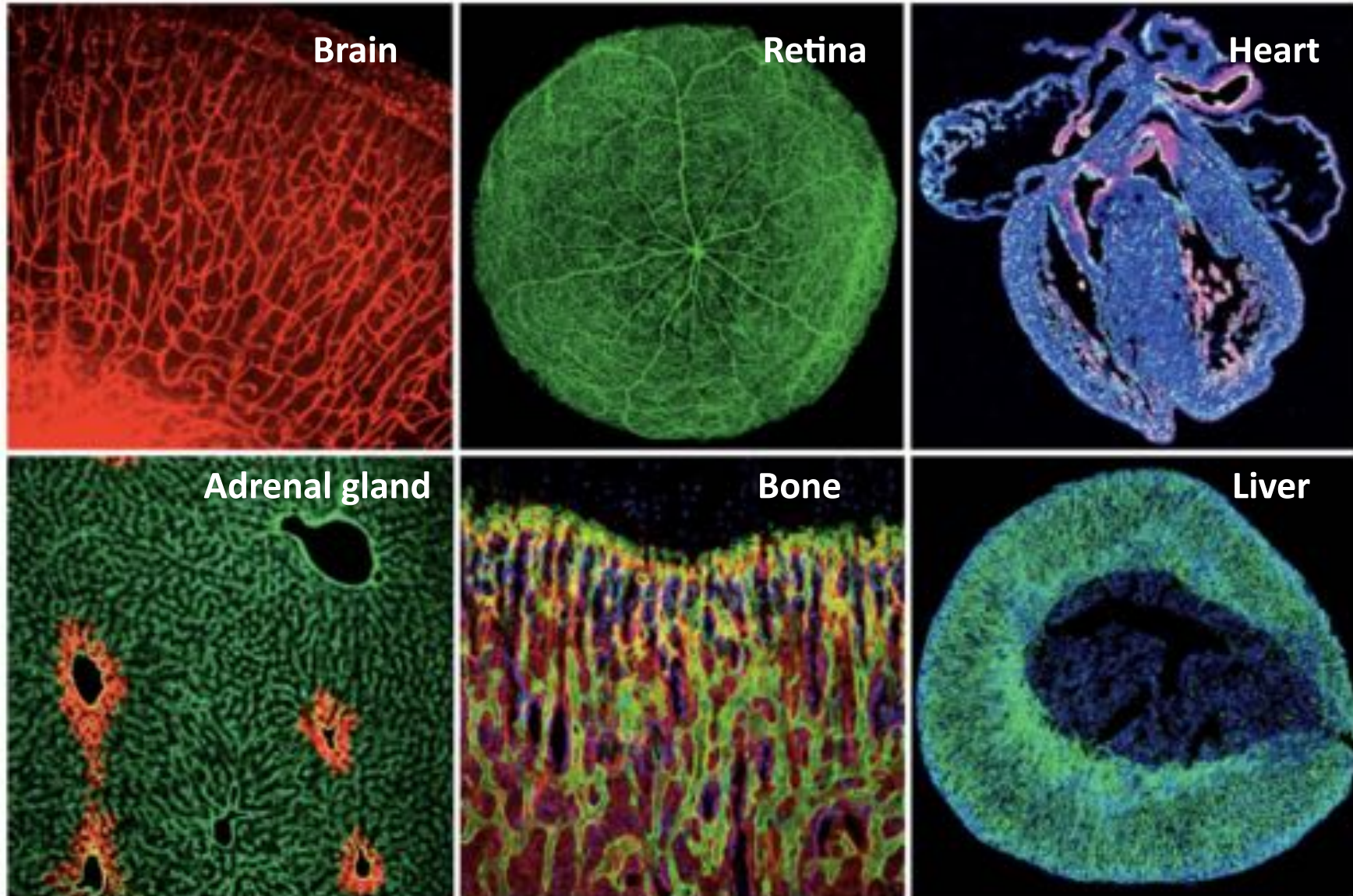
Vaupel et al. *Sem rad onc*, 2004

Cassot et al. *Microcirculation* 2006



# What is specific ?

The details of the vascular architecture





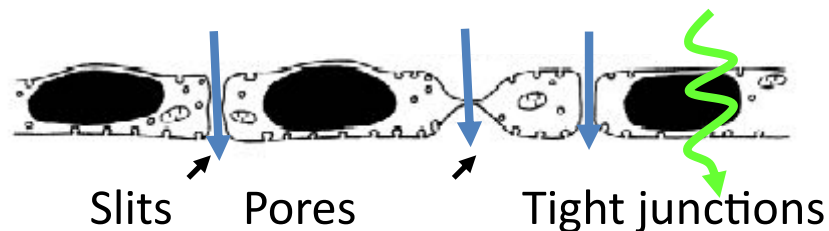
# What is specific ?

## The need for a constant microenvironment

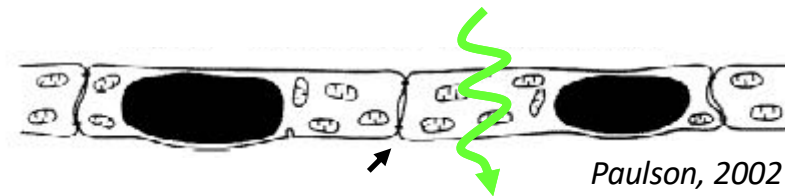
- ❑ **Ionic / Osmotic balance**
  - Neurotransmitters / Electrical impulse
  - Constant volume (Skull)
- ❑ **Protection against neurotoxicity**

### → Existence of the Blood-Brain Barrier

Ehrlich 1885, Goldman 1913



Muscle capillary endothelium



Brain capillary endothelium

# What is specific ?

## High metabolic activity / No energy storage

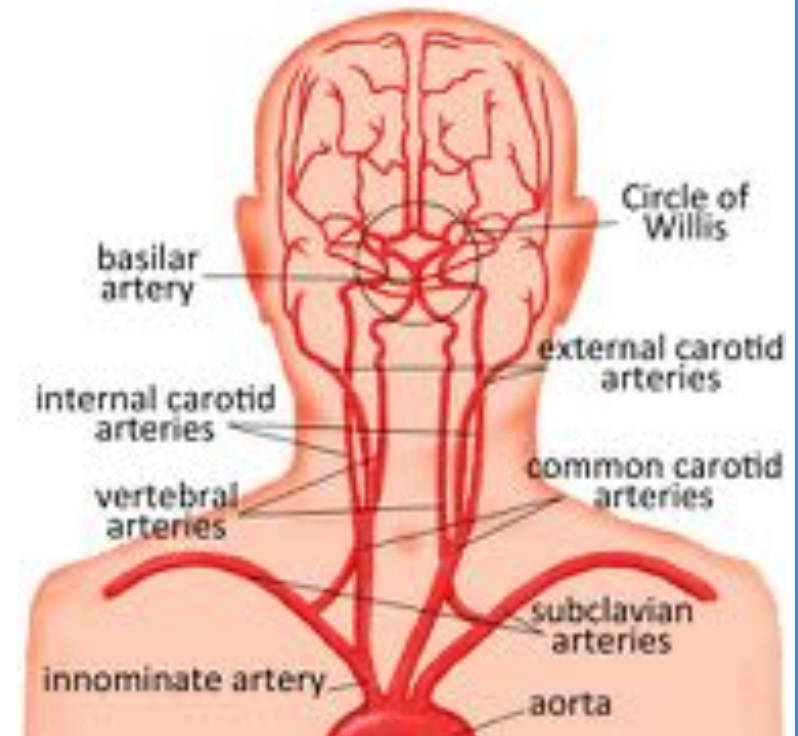
### ❑ A high metabolic activity (energy demand)

- Brain weight ~2% total weight
- Brain blood flow ~15 to 20% of total flow

### ❑ No energy storage

#### ➔ Existence of redundancies

- Circle of Willis
- Connections between cerebral arteries
- Connections between surface vessels



# What is specific ?

## High metabolic activity / No energy storage

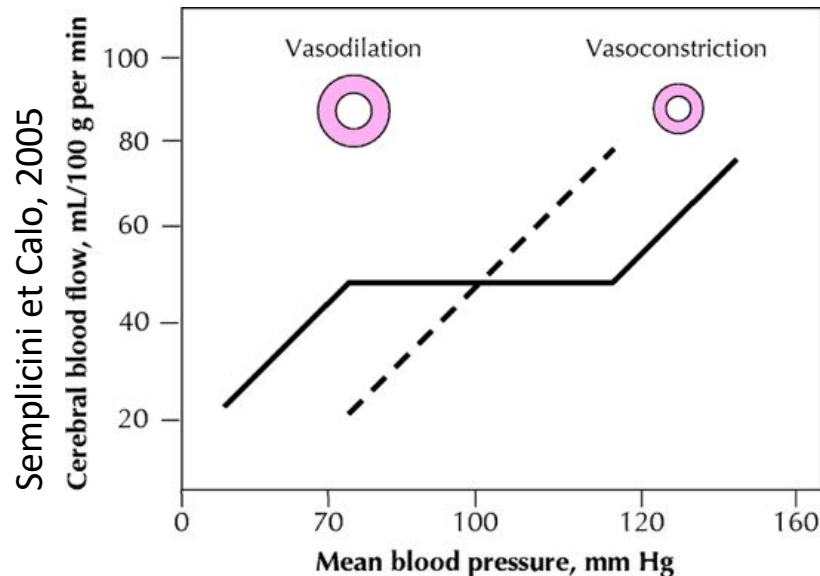
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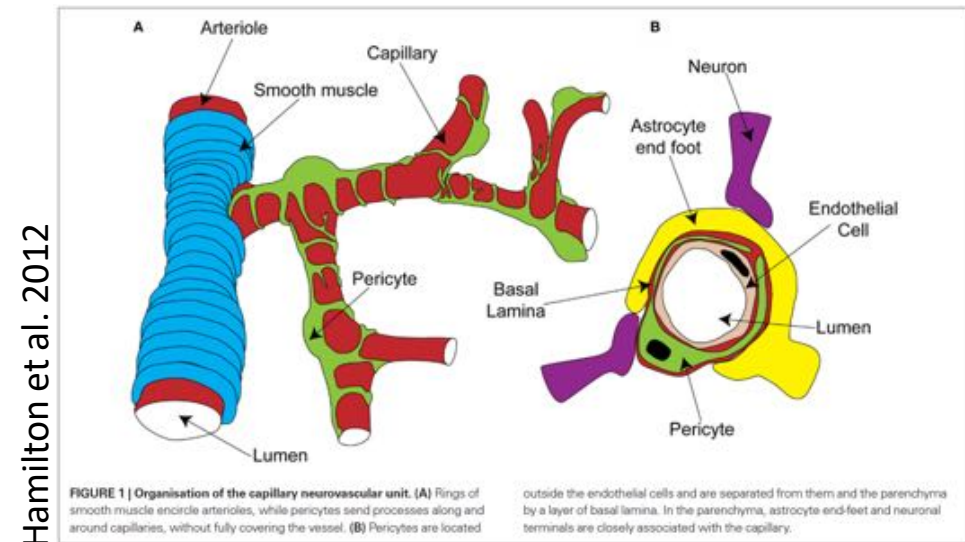
### □ No energy storage

## ➔ Two distinct regulation pathways

Brain autoregulation : global



Neurovascular coupling : local





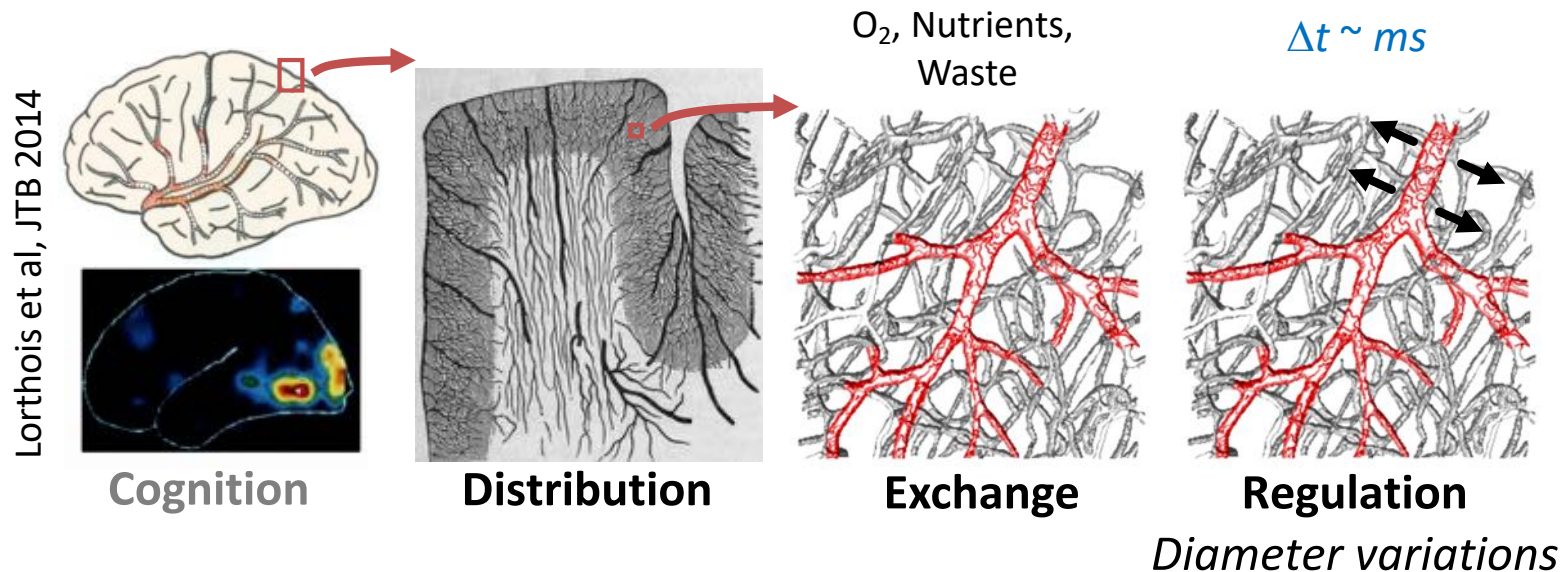
# Outline



- **Brain versus other organs**
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- **Why study brain microcirculation ?**
  - In health
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# Brain Microcirculation

- ... plays a key role in brain physiology



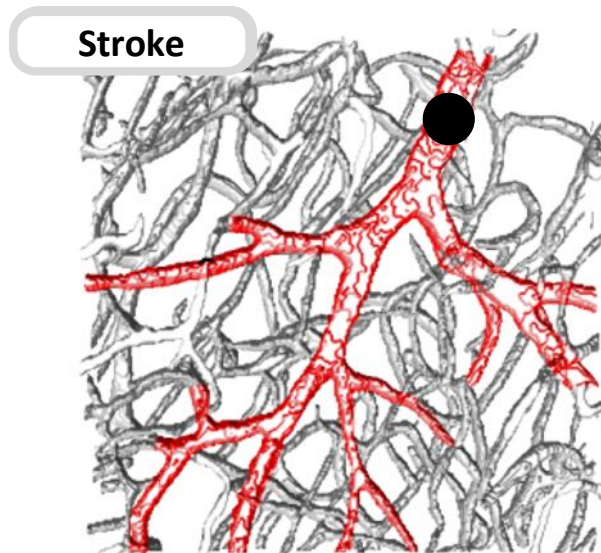
- ... and in our ability to observe the functioning brain

- Neuro-vascular coupling
- Hemodynamically-based functional imaging techniques (H<sub>2</sub>O<sup>15</sup> PET, fMRI)

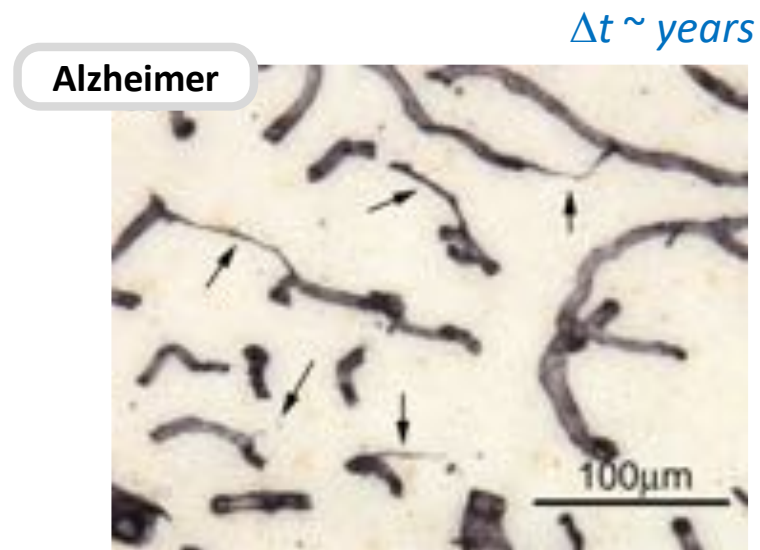
# Brain Microcirculation

## ■ ... is involved in disease

- Stroke
- Neurodegenerative disease (Alzheimer)



*Vessel occlusions*

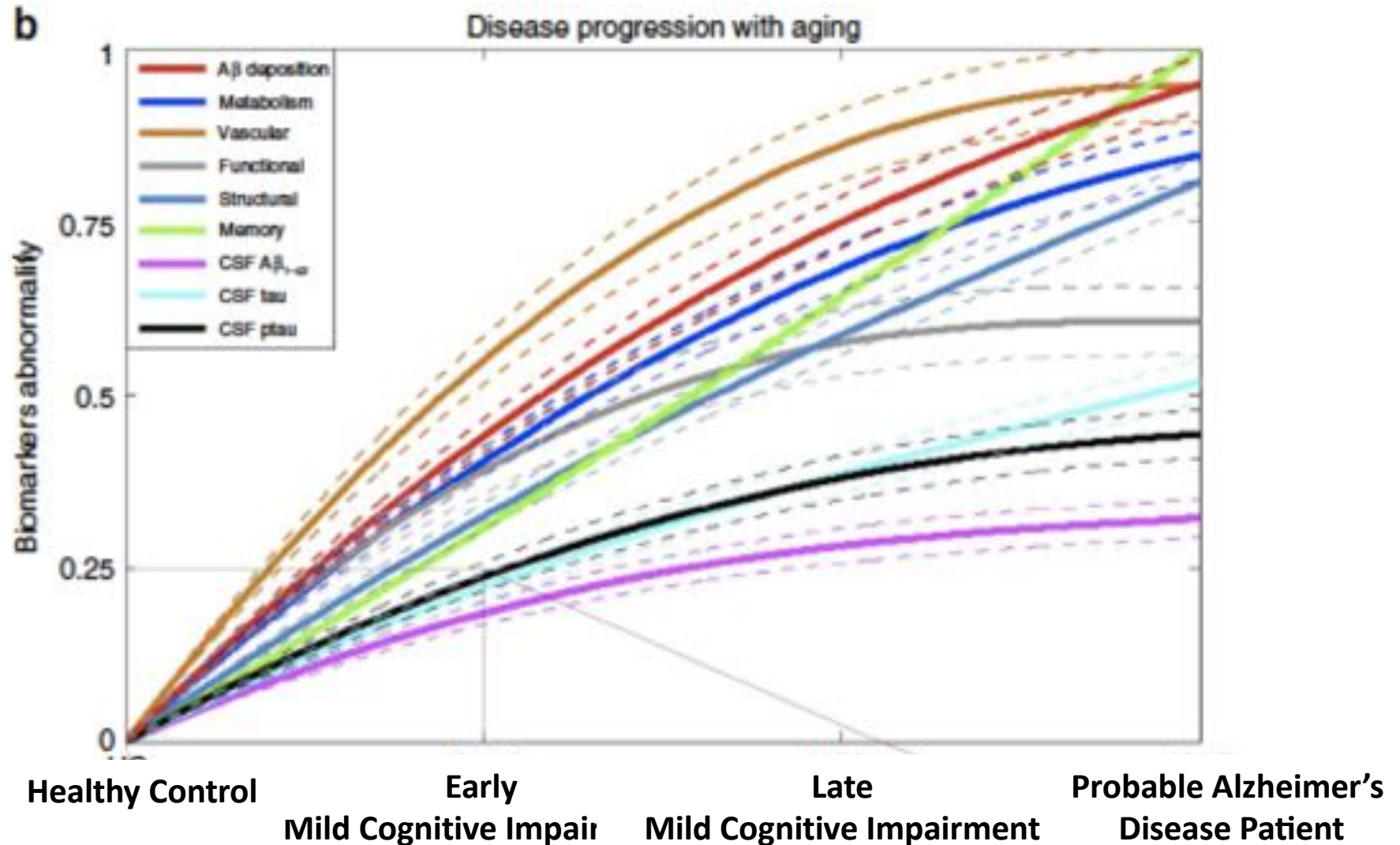


*Progressive disappearance*

Hunter et al. 2012



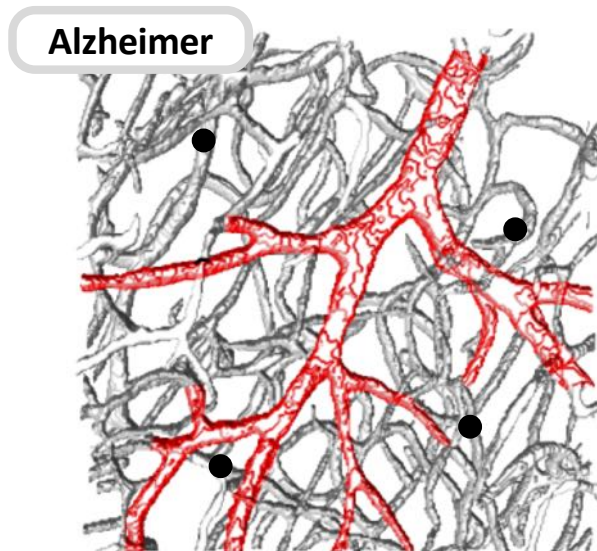
# Brain Microcirculation



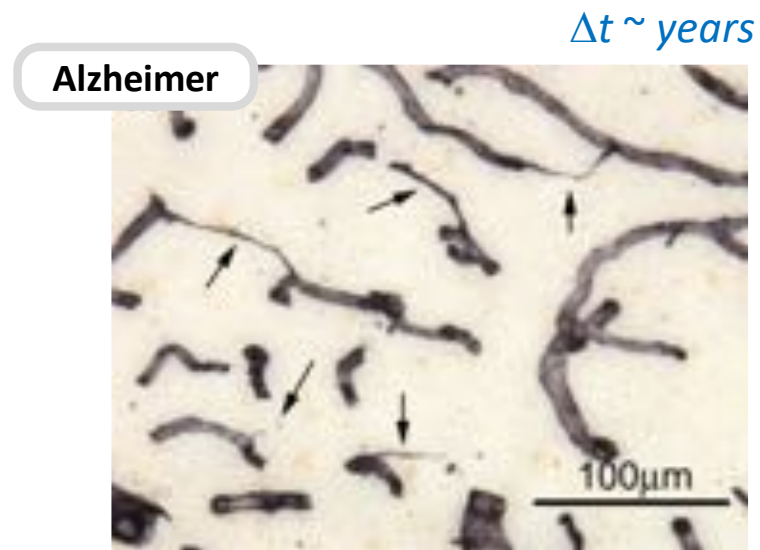
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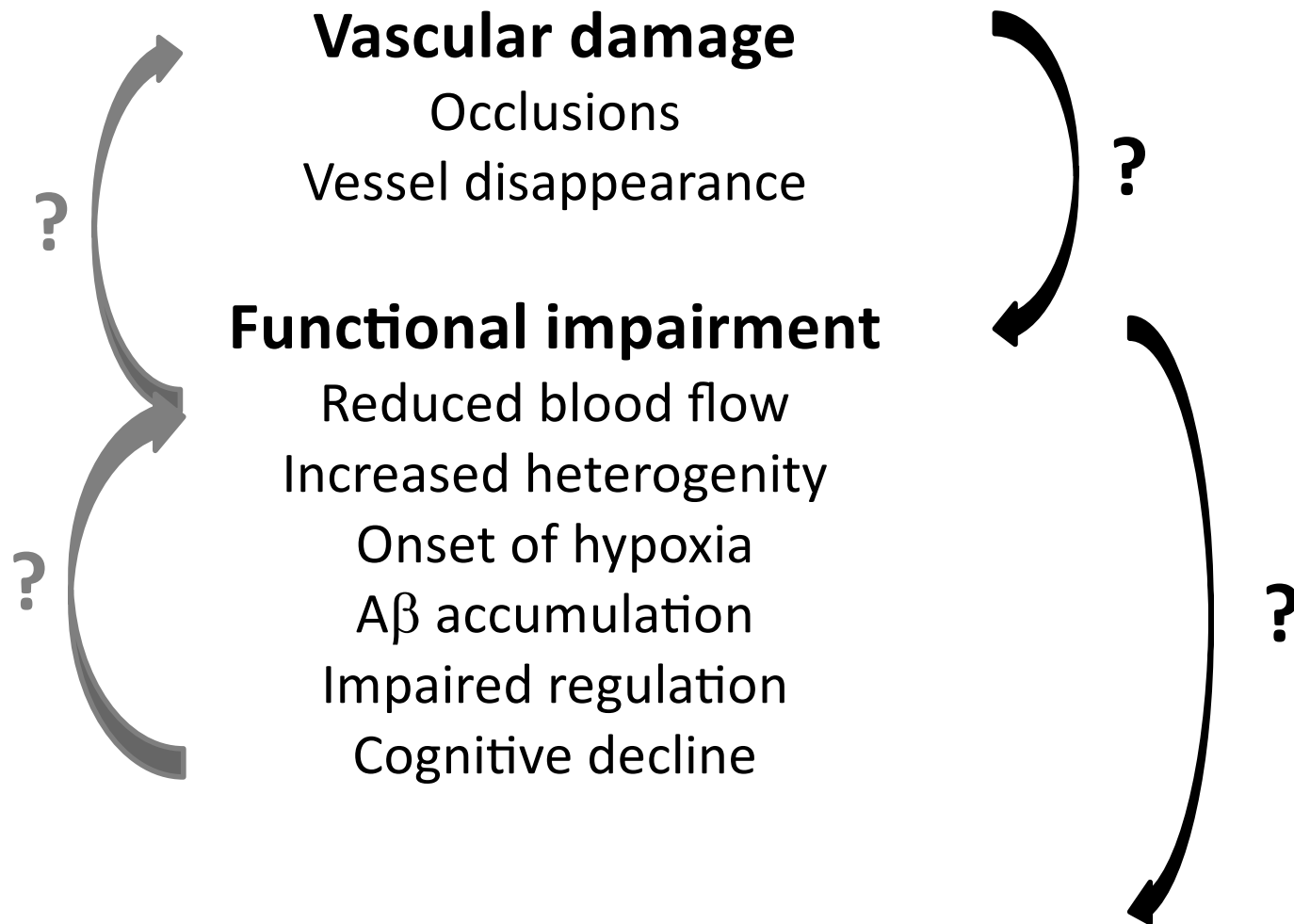


*Vessel occlusions*



*Progressive disappearance*

# Brain Microcirculation in Alzheimer's disease



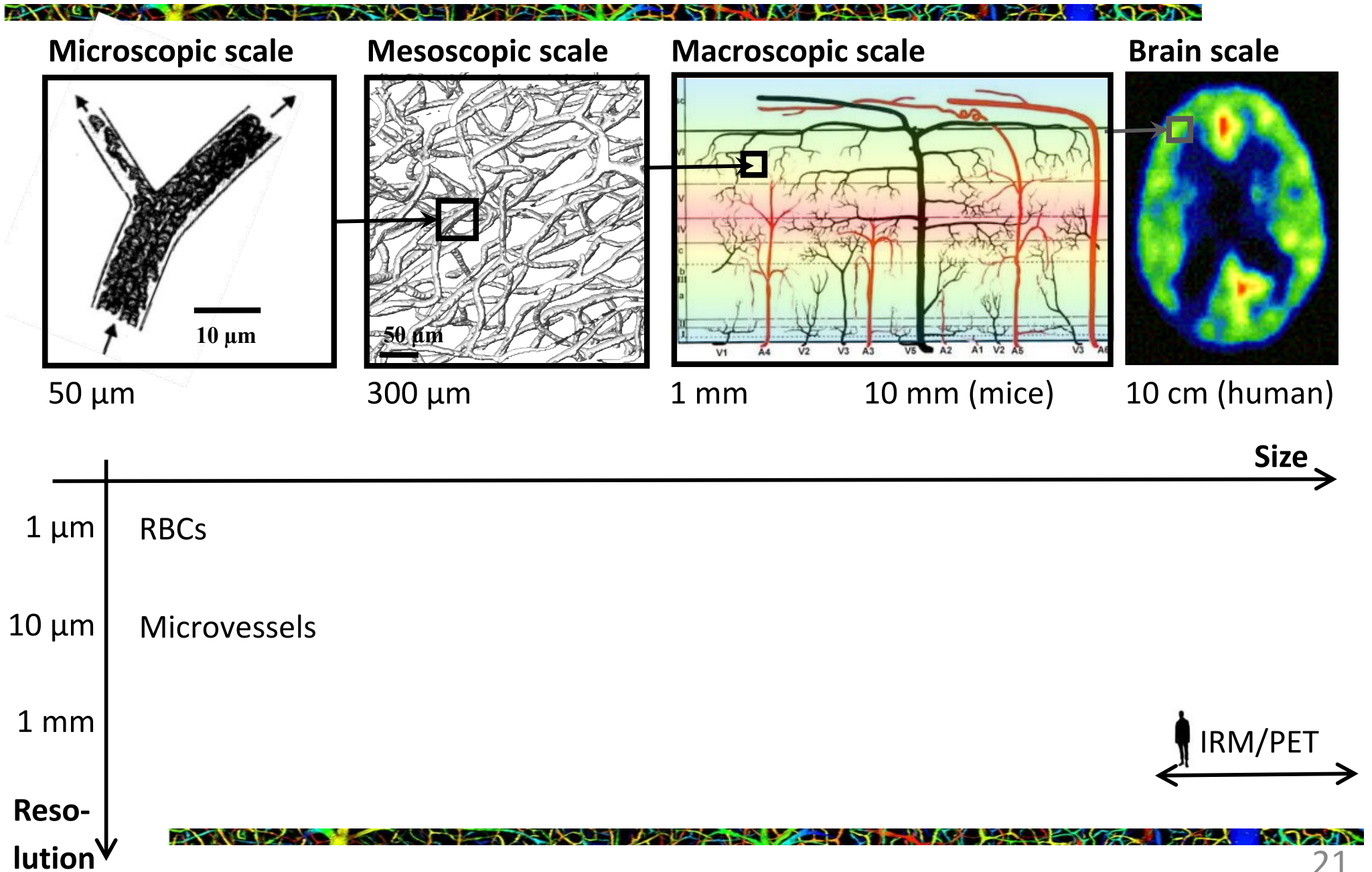


# Outline



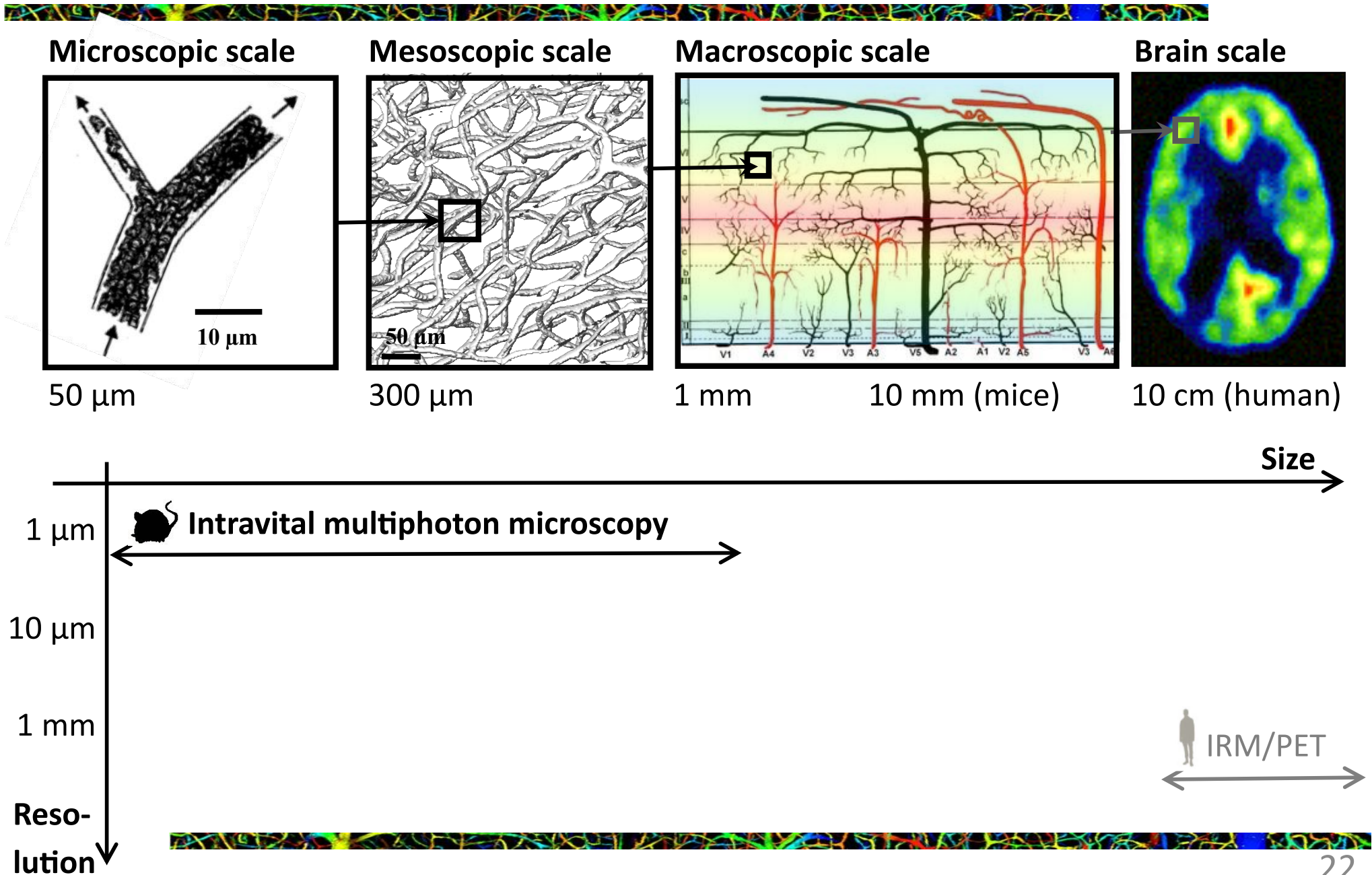
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# Investigation tools and associated scales



# Investigation tools and associated scales

## In vivo experimental methods

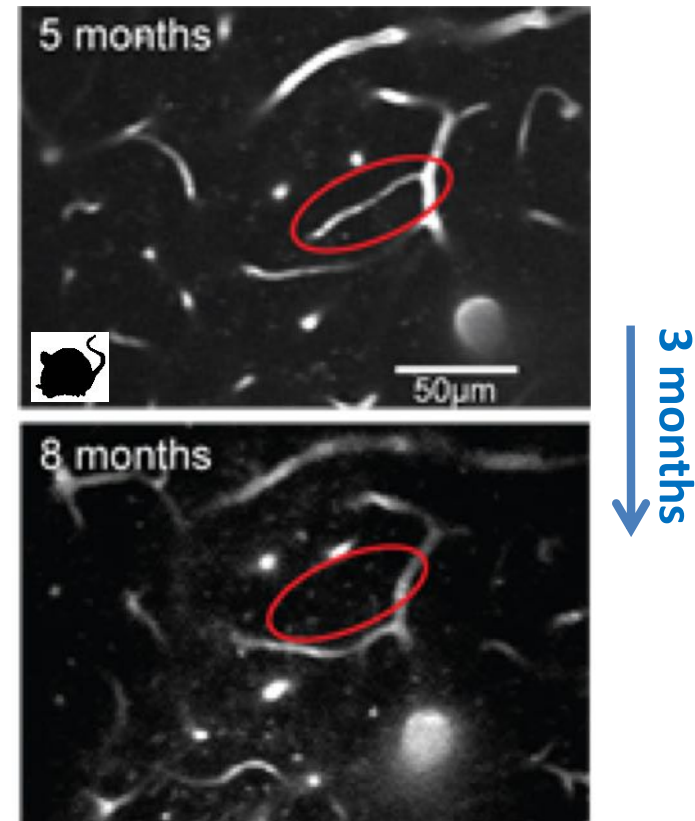
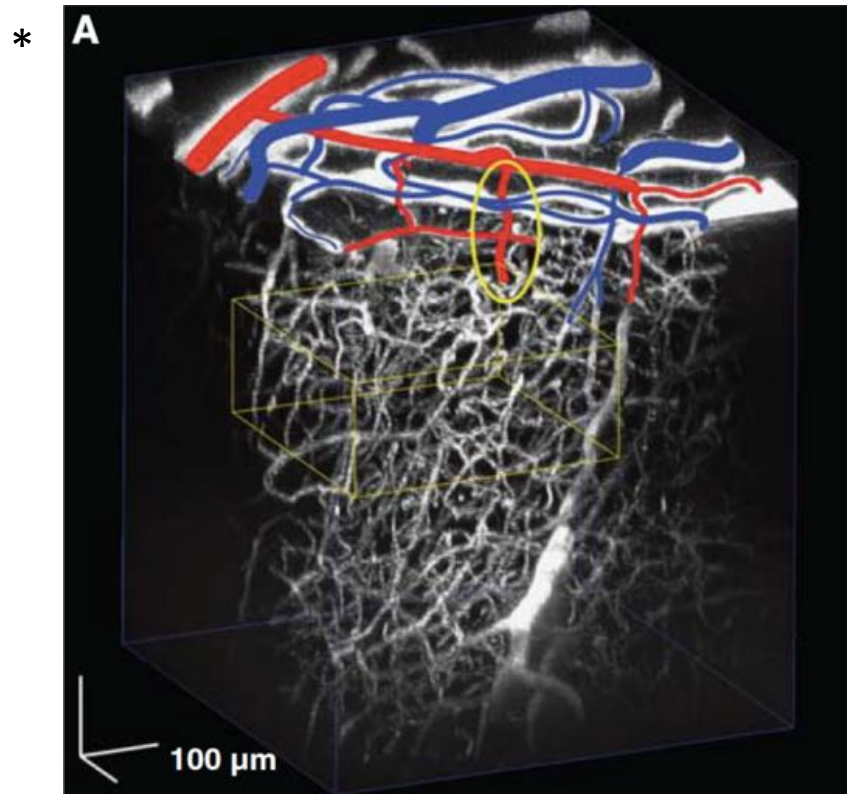




# Intravital multiphoton microscopy

## Structure

- Healthy and diseased (Alzheimer) animal models



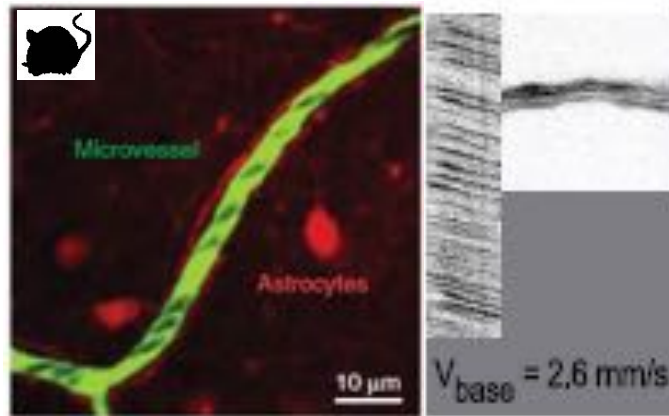
Nozomi Nishimura, Personal data

# Intravital multiphoton microscopy

## Function

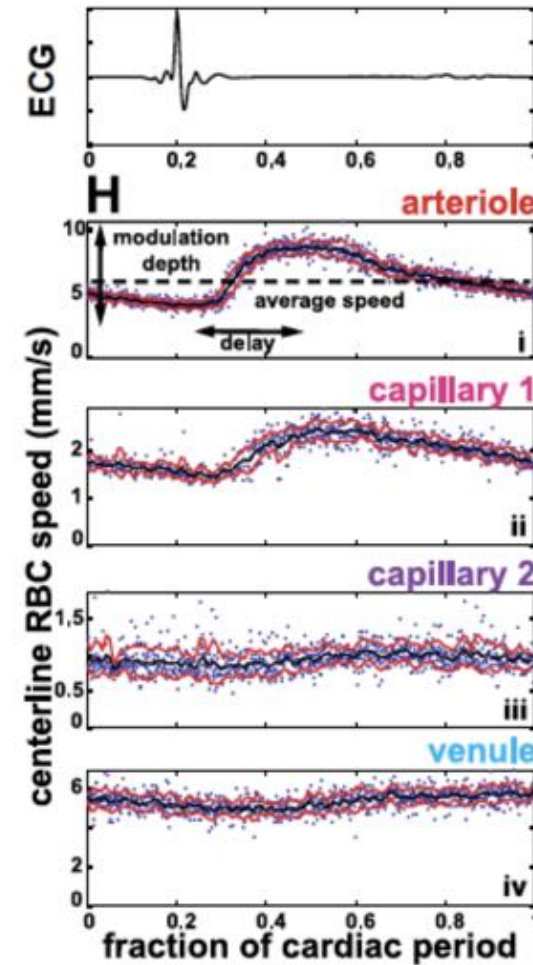
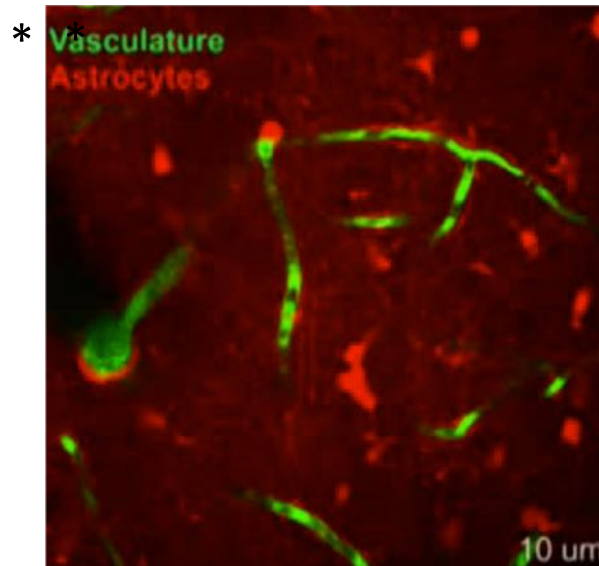
### Red Blood Cell velocity

30 s



Shih et al. 2012

Nishimura et al. 2007



Santisakultarm et al. *Am J Phys* 2011

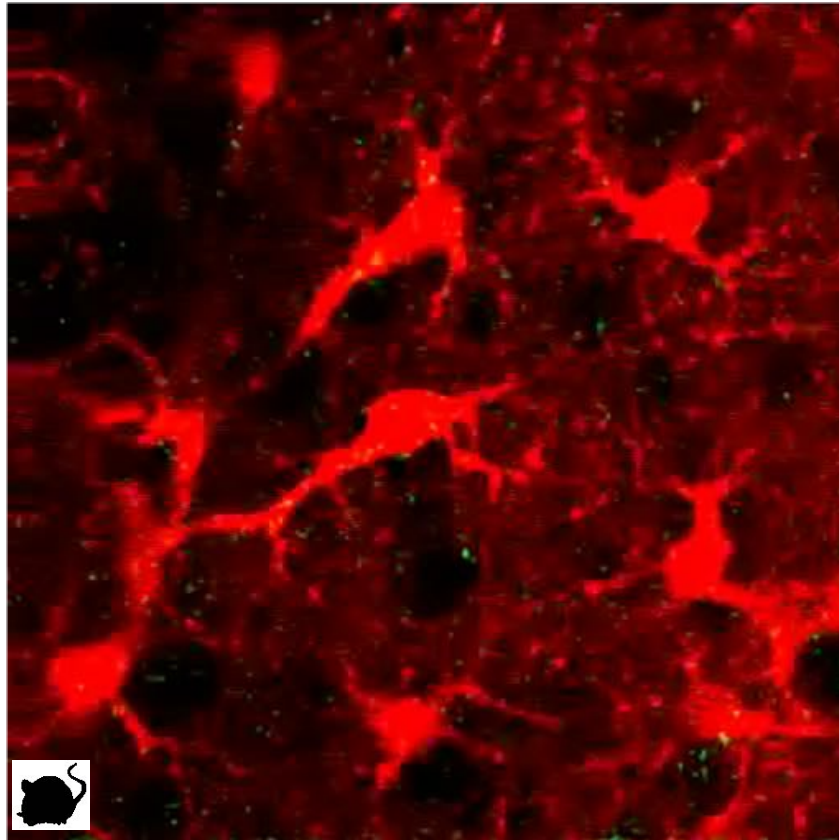
\*From Andy Shih, <https://www.youtube.com/watch?v=YV1TpKNB0S4>

# Intravital multiphoton microscopy

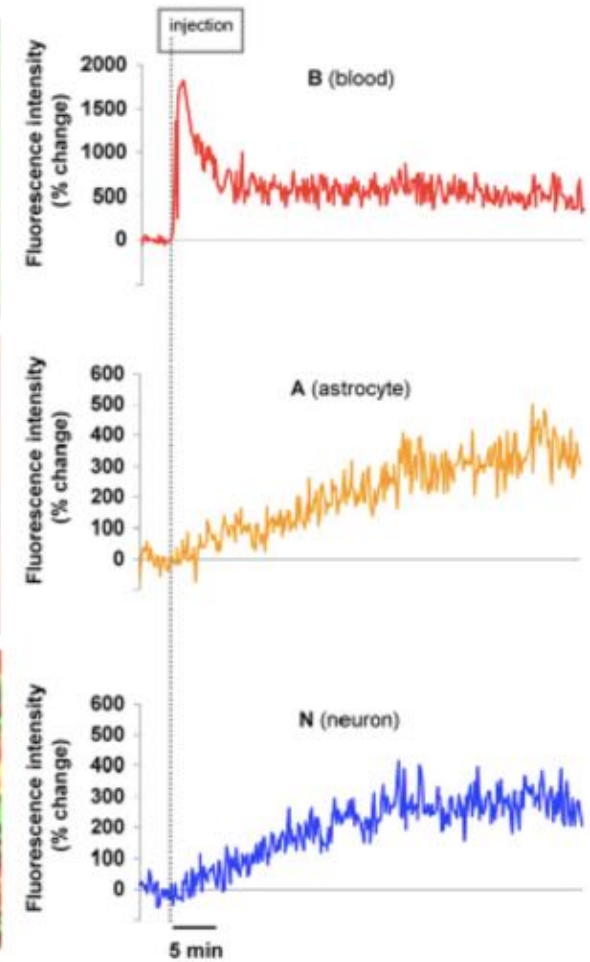
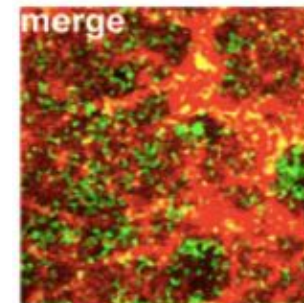
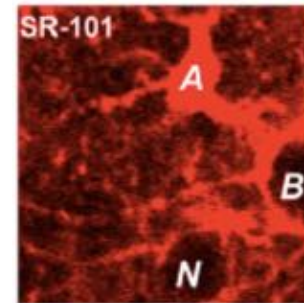
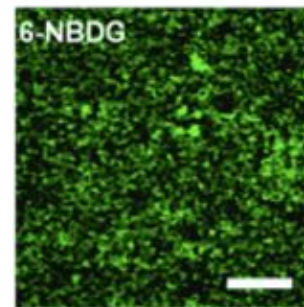
## Function



### Glucose Exchange (6NBDG)



Chuquet et al. 2010

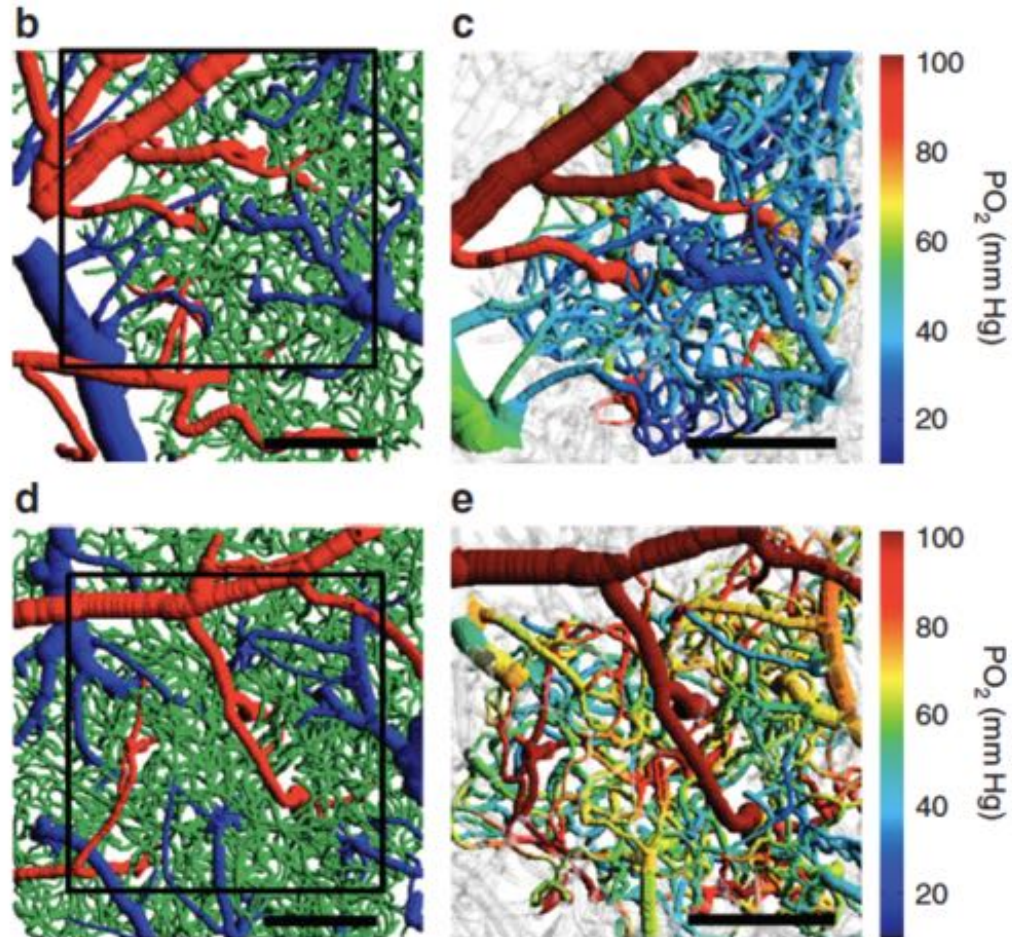




# Intravital multiphoton microscopy

## Function

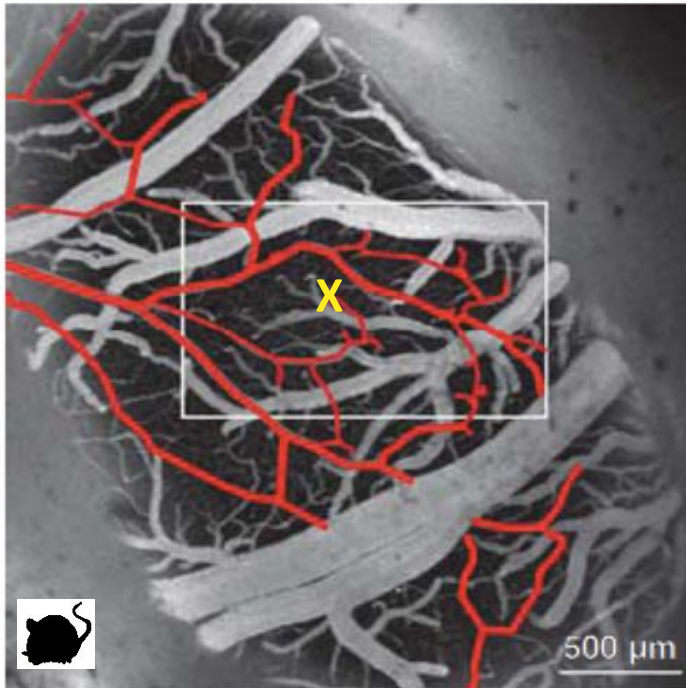
Oxygen partial pressure



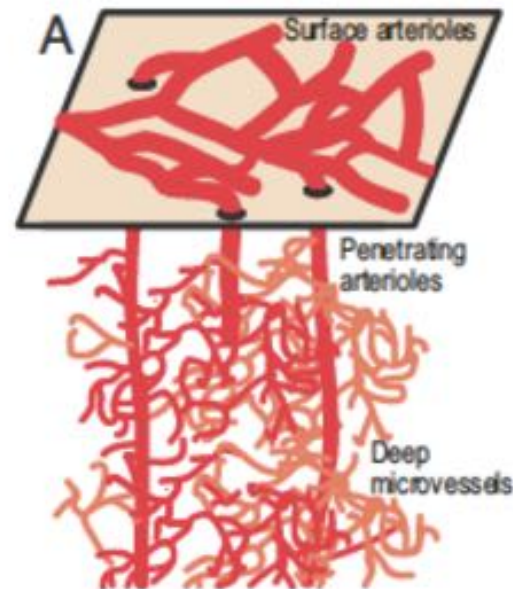
# Intravital multiphoton microscopy

## Function

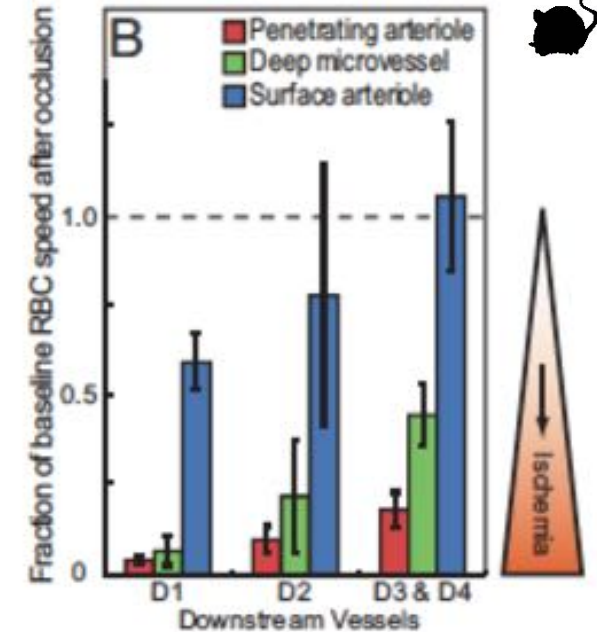
Robustness to controlled perturbations: localized obstructions



Nishimura et al. JCBFM 2010



Nishimura et al. PNAS 2007

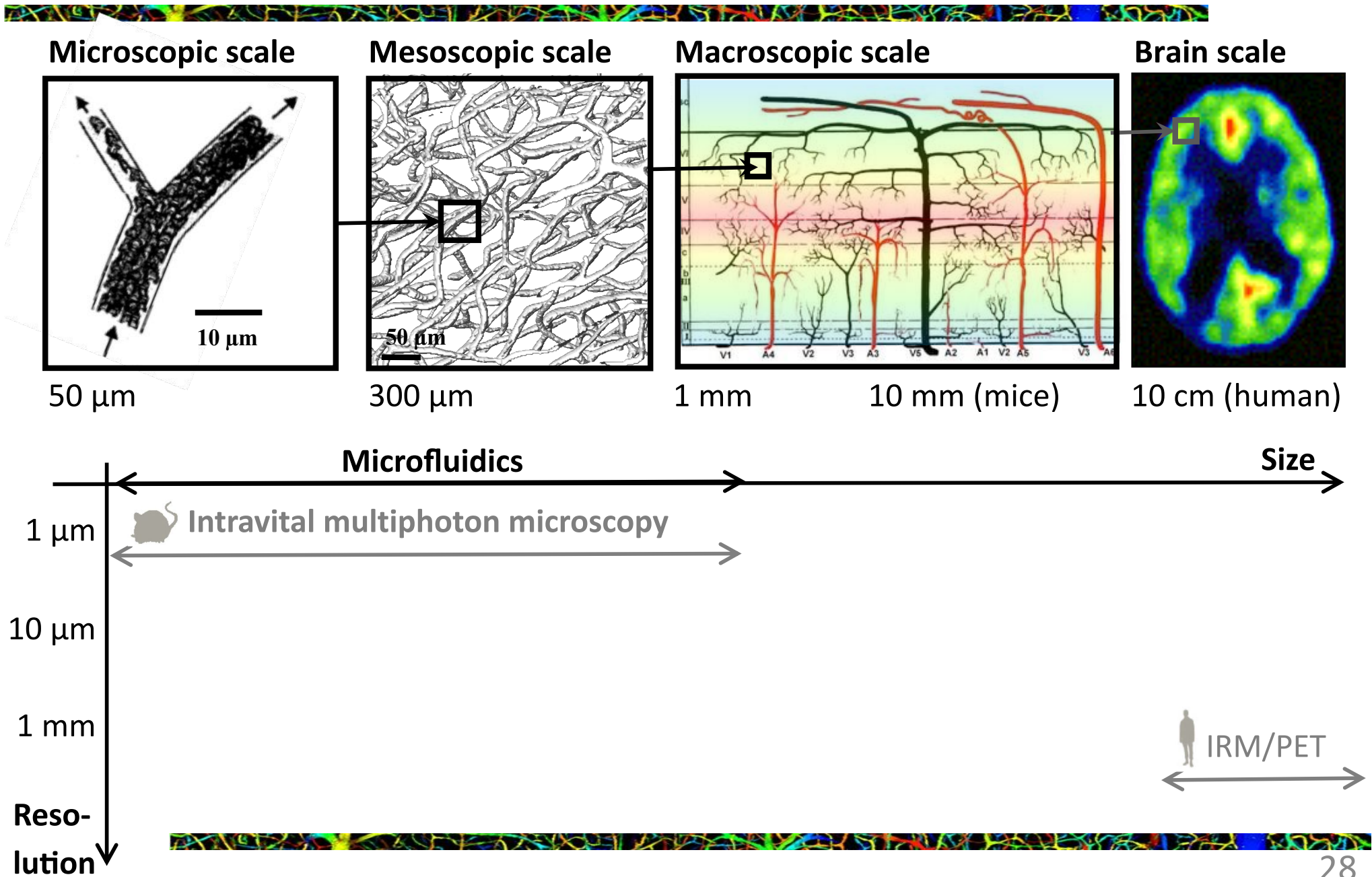


***“Penetrating arterioles are a bottleneck in the perfusion of neocortex”***



# Investigation tools and associated scales

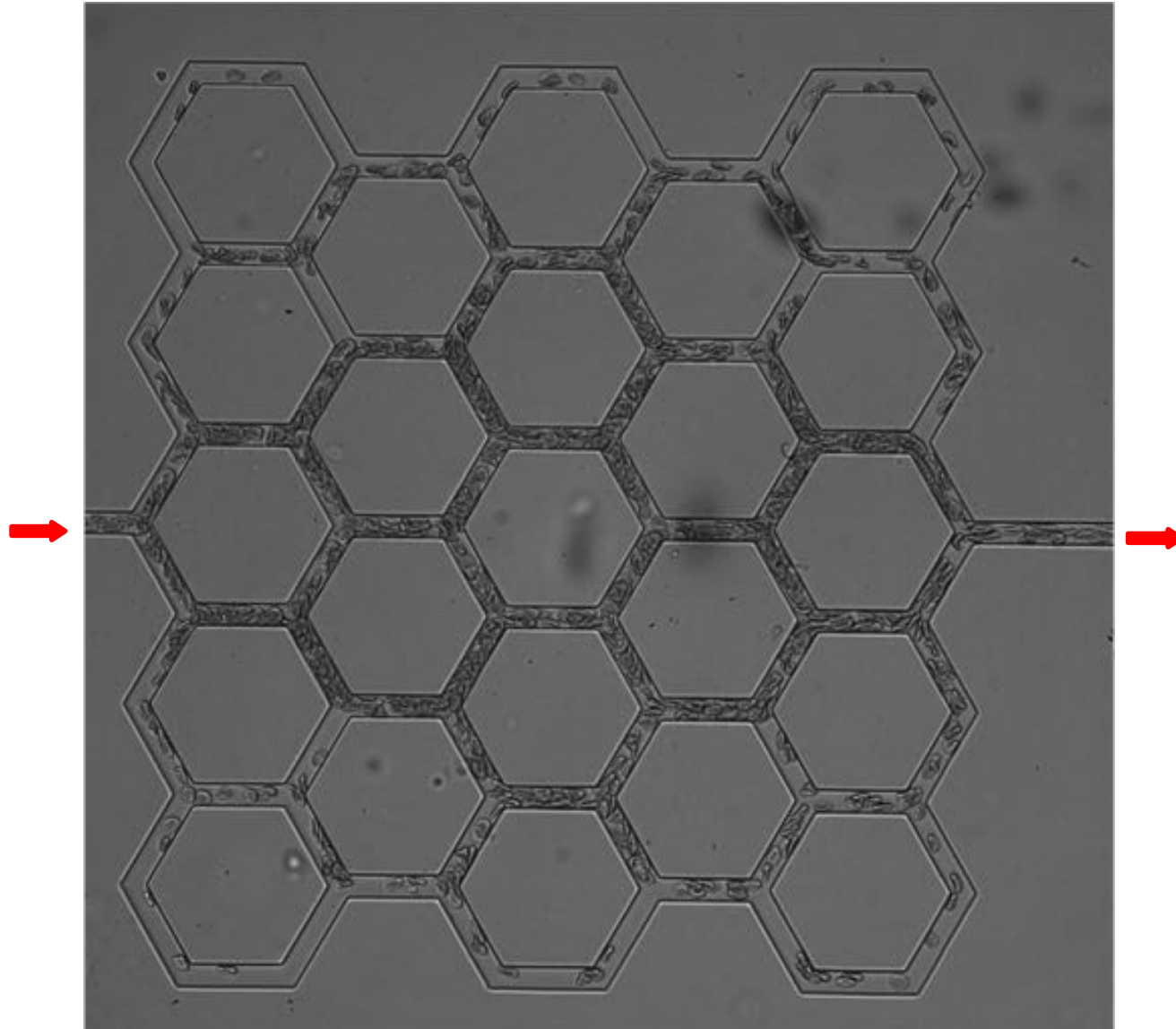
## In vitro experimental methods





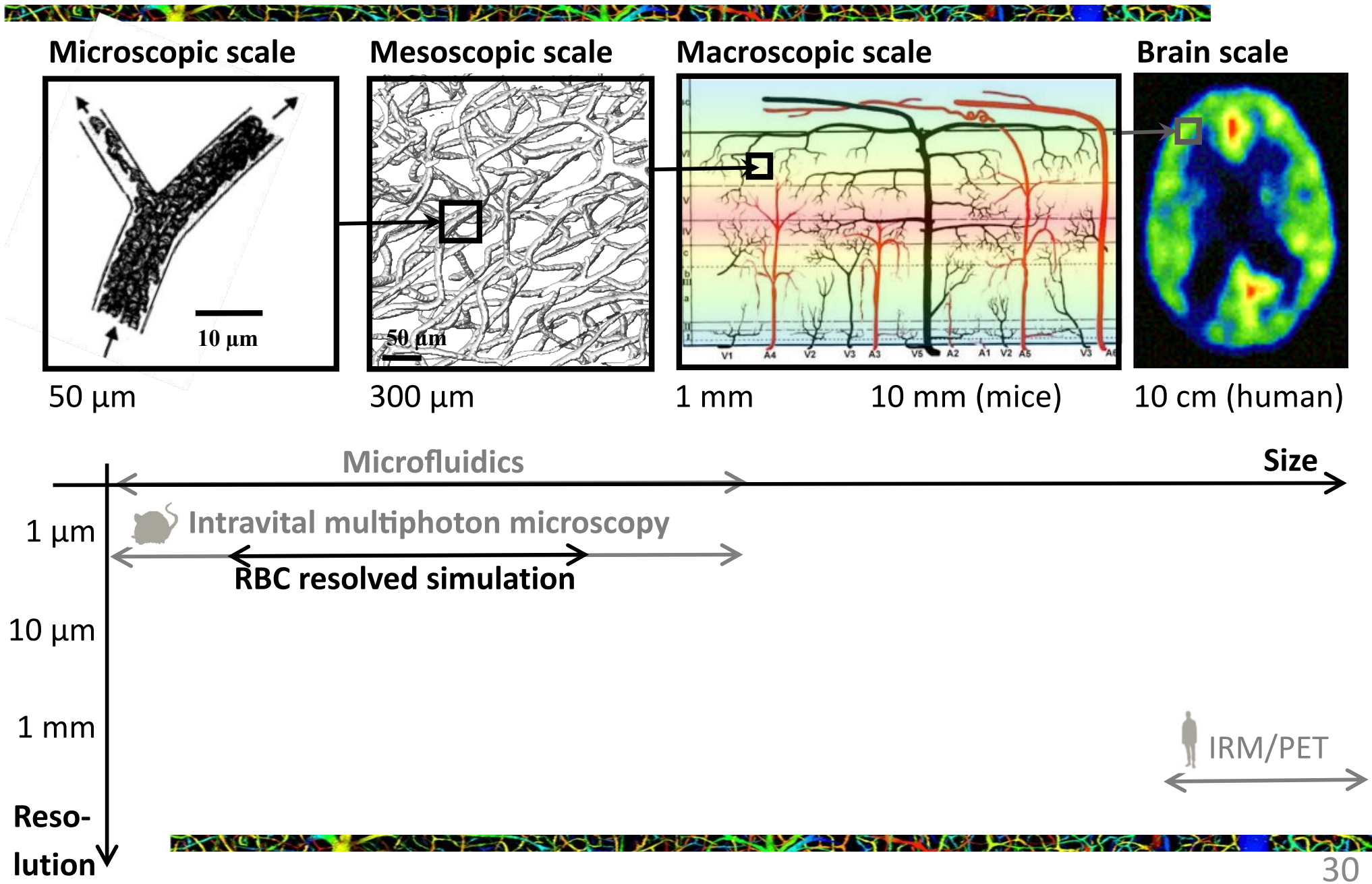
# Microfluidics

$W=10\ \mu\text{m}$   
 $L=50\ \mu\text{m}$   
 $H_t=8\%$



# Investigation tools and associated scales

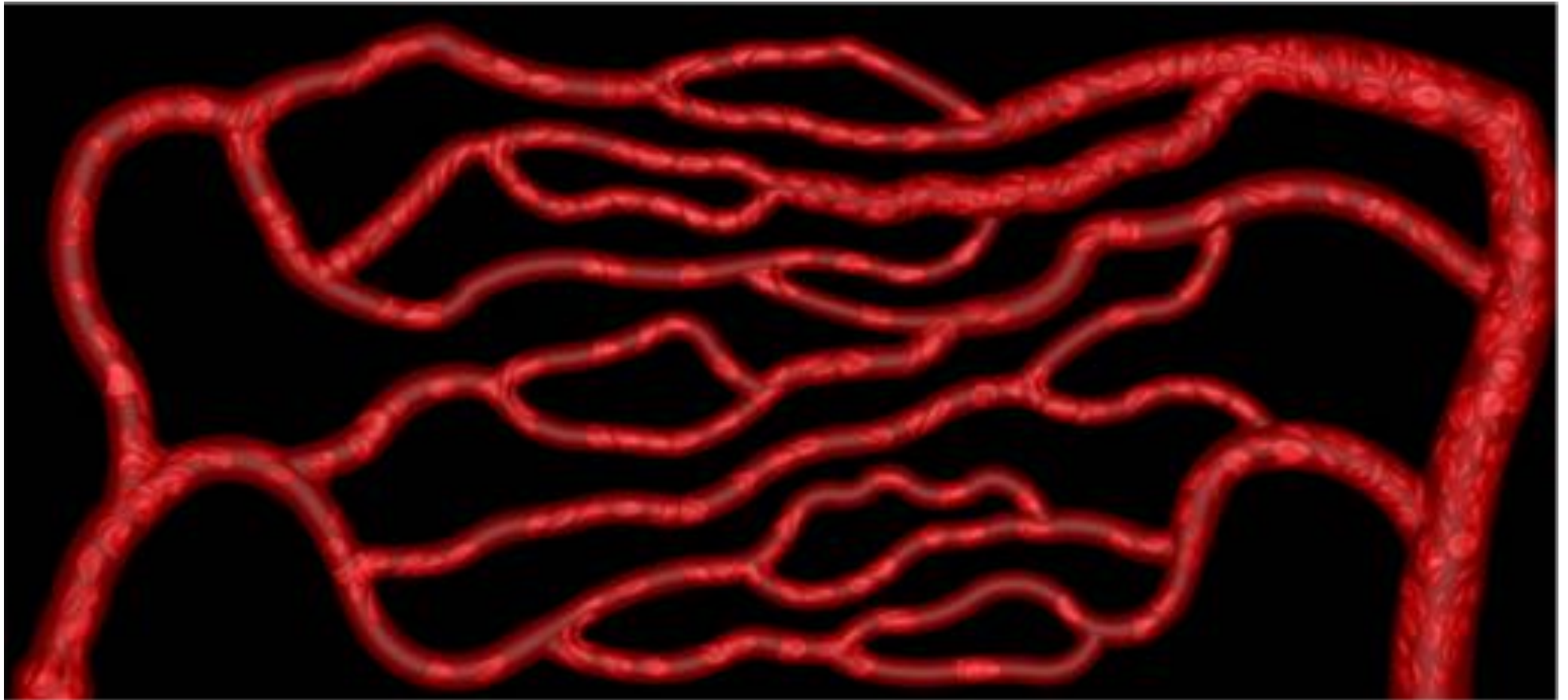
## Numerical simulation



# RBC resolved simulations

## Some examples

IBM / FV / FE / front-tracking method



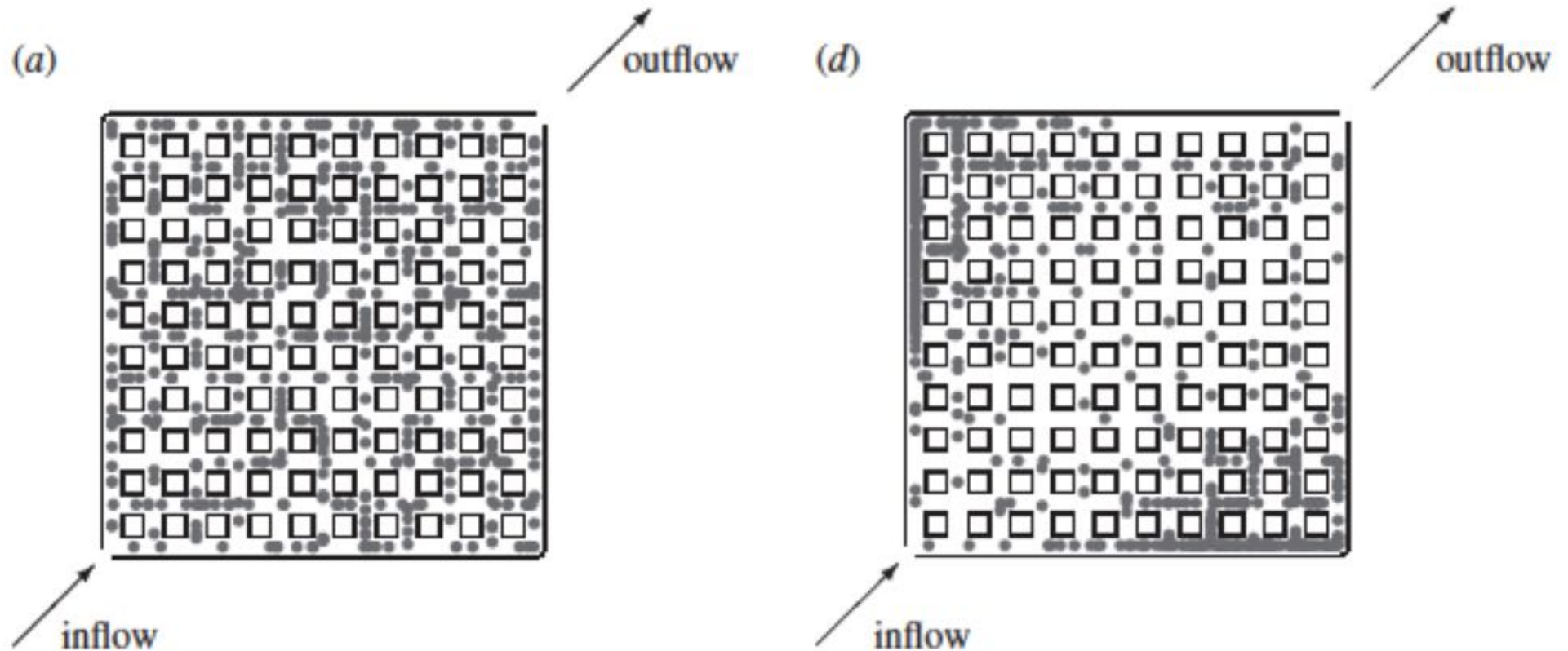
<http://bagchi.rutgers.edu>



# RBC resolved simulations

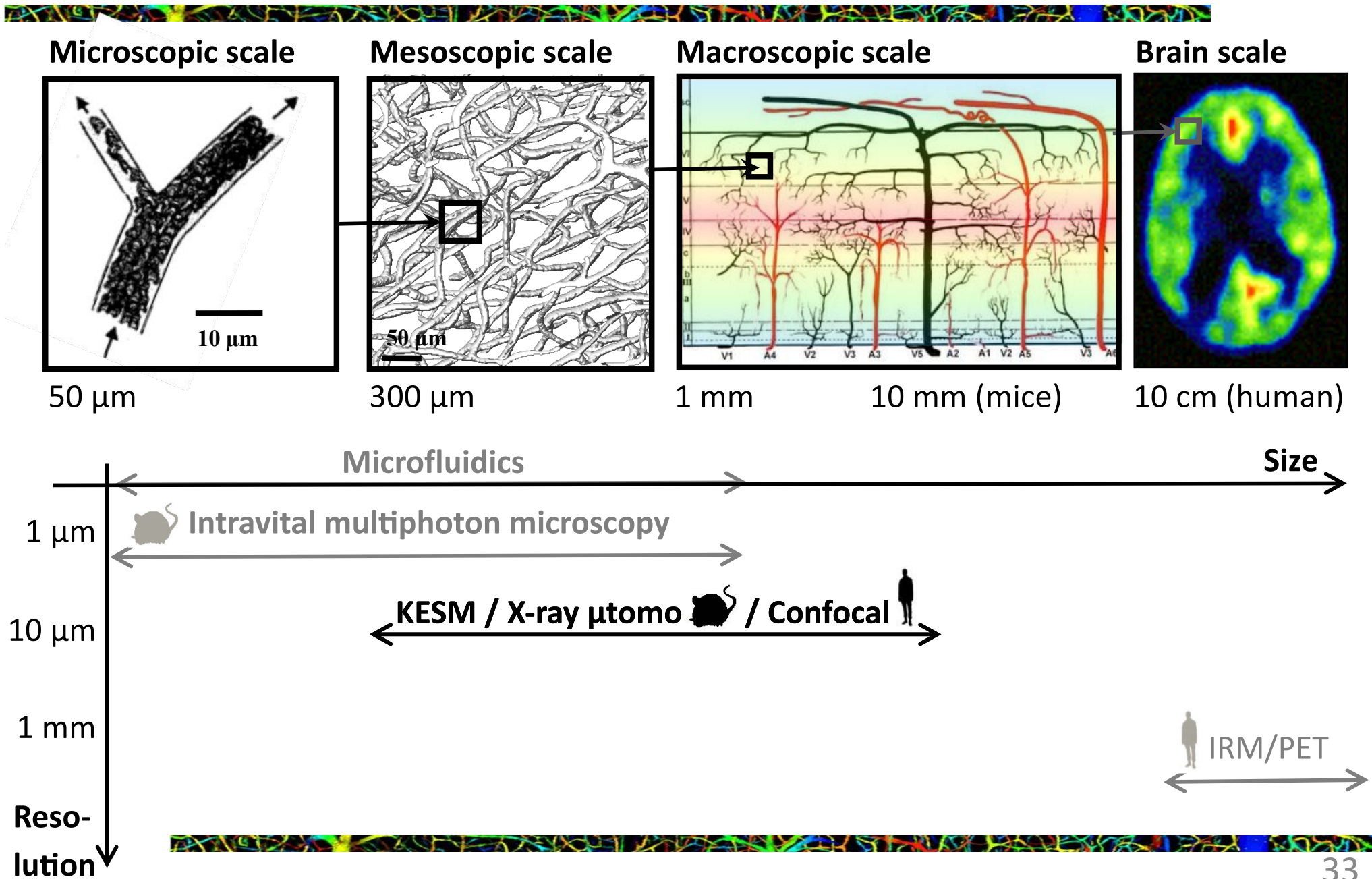
## Some examples

Lagrangian tracking of RBCs in a network

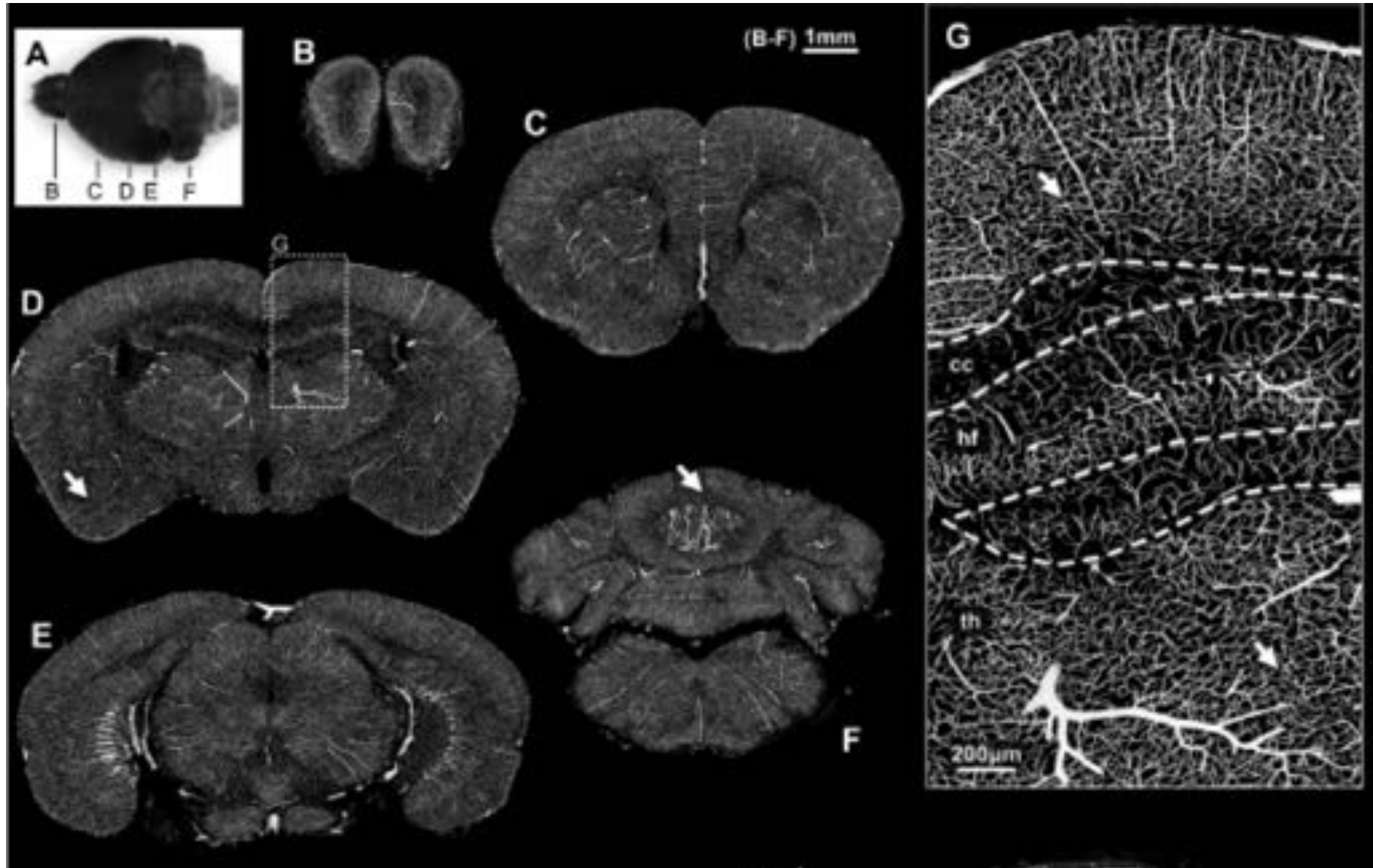


# Investigation tools and associated scales

## Post-mortem experimental methods



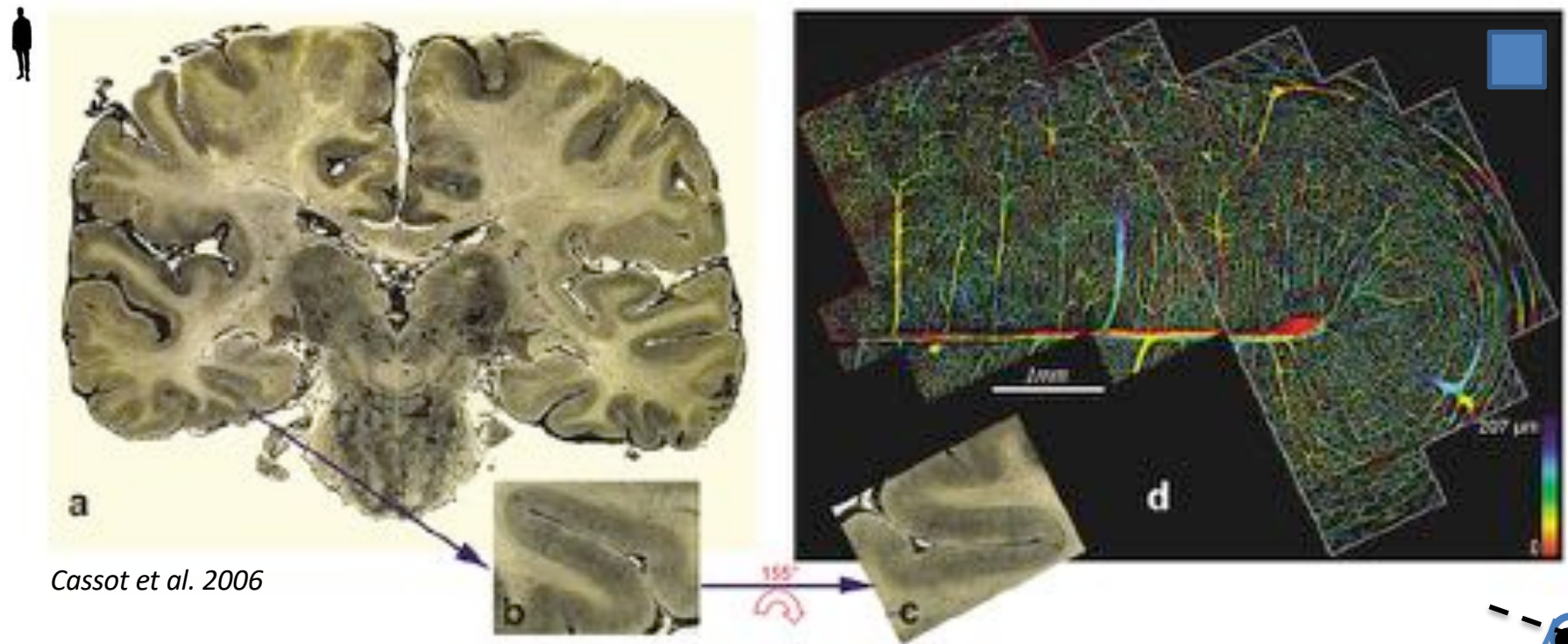
# Knife-Edge Scanning Microscopy (Micro-Optical Sectioning Tomography)



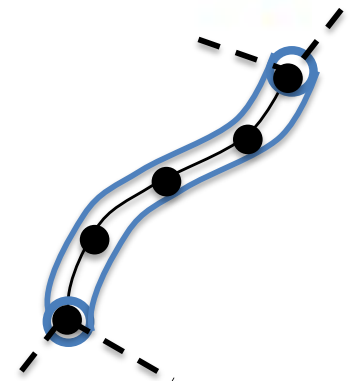


# Human anatomical database (Inserm Tonic) post mortem confocal imaging

## □ 60 year old, abdominal lymphoma

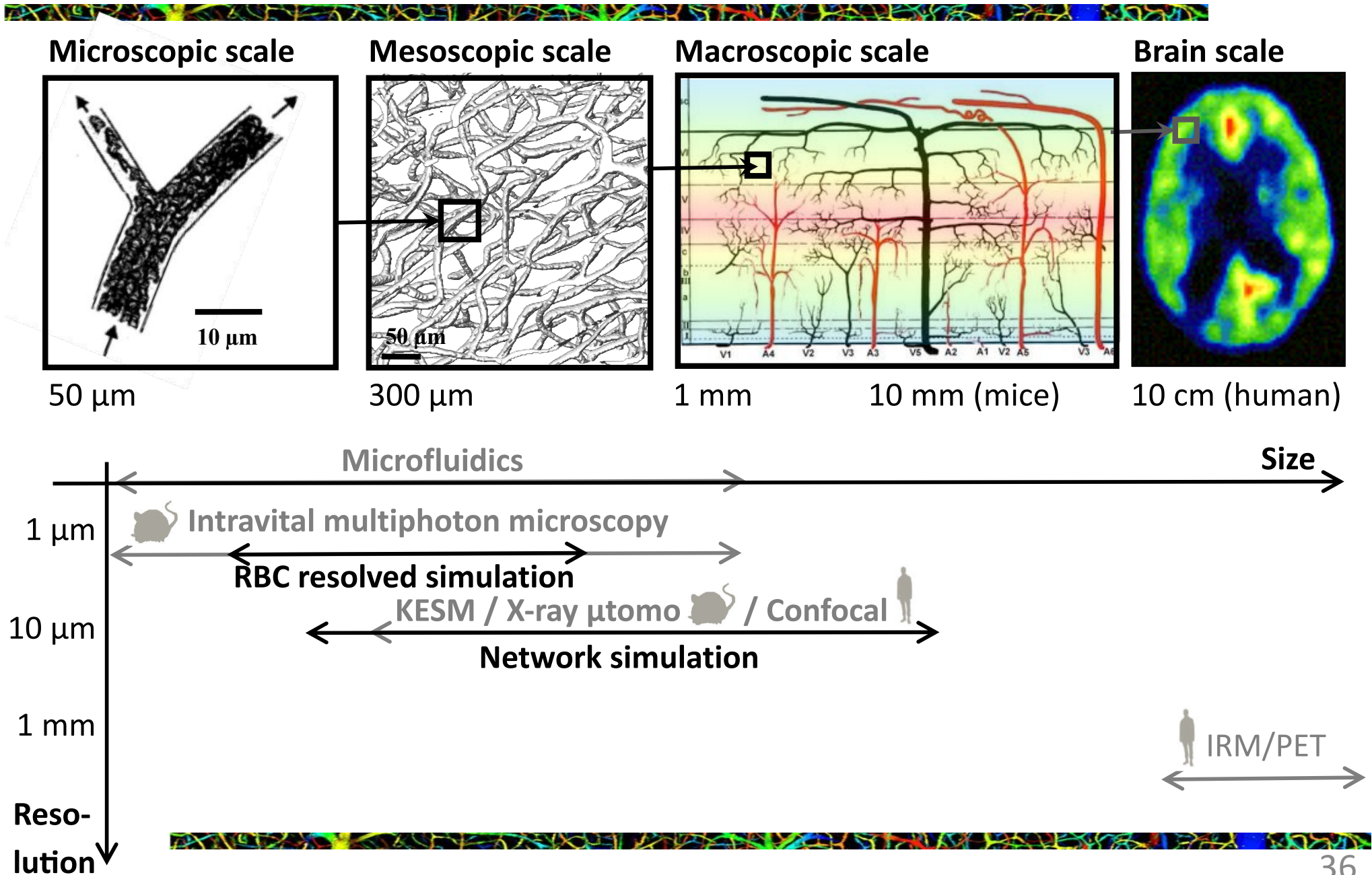


- $1.2 \times 1.2 \times 3 \mu\text{m}^3$  in areas as large as  $10 \text{ mm}^2 \times 300 \mu\text{m}$
- Vascular network graph (centerlines, radii, connectivity)
- Morphometric analysis (segments, bifurcations)\*



# Investigation tools and associated scales

## Numerical simulation



# Outline



- **Brain versus other organs**
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# Blood flow in networks

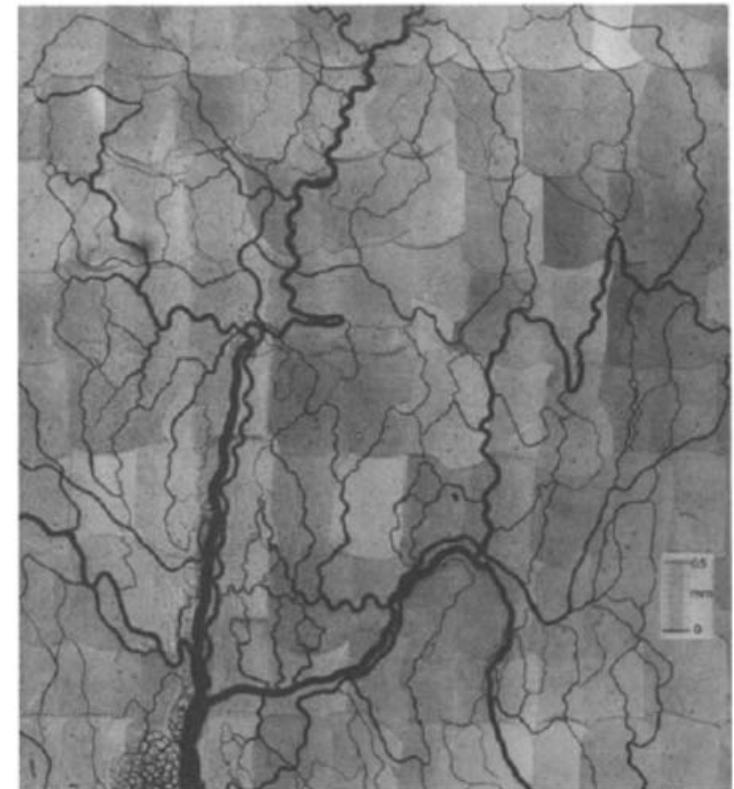
## Modeling framework

### Blood Flow in Microvascular Networks

#### Experiments and Simulation

A.R. Pries, T.W. Secomb, P. Gaehtgens, and J.F. Gross

- In vivo experiments
  - Rat mesentery
  - Structure (R, L, Connectivity)
  - Hemodynamic data ( $Q$ ,  $H_{\text{tube}}$ )



# Blood flow in networks

## Modeling framework



### Blood Flow in Microvascular Networks

#### Experiments and Simulation

A.R. Pries, T.W. Secomb, P. Gaehtgens, and J.F. Gross

#### □ Simulations

- The vascular network is described as a graph
- The blood is viewed as an equivalent fluid (i.e. at vessel scale)
- Its non-linear rheological properties are described by empirical relationships
- A first guess on their coefficient is based on experiments (in vivo or in glass tubes)

#### □ Simulation/Experiments

- The values of these coefficients are adjusted to best match the *in vivo* measurements
- This gives insights on biophysical aspects of blood flow in the microvasculature

# The blood as an equivalent fluid

## Apparent viscosity and relative viscosity

If the fluid is Newtonian and the flow is laminar, we have the Hagen-Poiseuille formula [Eq. (8) of Sec. 3.3]:

$$\frac{\Delta p}{\Delta L} = \frac{8\mu}{\pi a^4} \dot{Q}, \quad (1)$$

where  $\Delta p$  is the pressure drop in length  $\Delta L$ ,  $\mu$  is the coefficient of viscosity of the fluid,  $a$  is the radius of the tube, and  $\dot{Q}$  is the volume rate of flow. If the fluid is blood, this equation does not apply; but we can still measure  $\Delta p/\Delta L$  and  $\dot{Q}$ , and use Eq. (1) to calculate a coefficient  $\mu_{app}$ :

$$\mu_{app} = \frac{\pi a^4}{8} \frac{1}{\dot{Q}} \frac{\Delta p}{\Delta L}. \quad (2)$$

The  $\mu$  so computed is defined as the *apparent coefficient of viscosity for the circular cylindrical tube*,

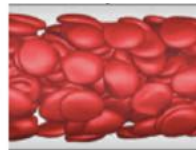
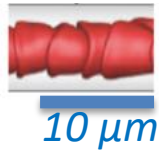
Eq. (2) is formally identical to Poiseuille equation but this does not imply a parabolic velocity profile (nor any other assumption on the shape of the velocity profile)



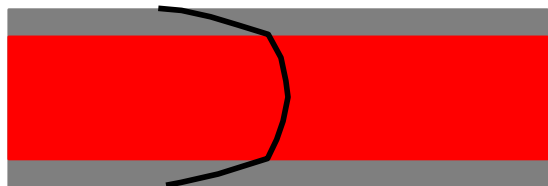
# The blood as an equivalent fluid

## Apparent viscosity and relative viscosity

Apparent and relative viscosities are not intrinsic properties of the blood; they are properties of the blood and blood vessel interaction, and depend on the data reduction procedure. There are as many definitions for apparent viscosities as there are good formulas for well-defined problems. Examples are: Stokes flow around a falling sphere, channel flow, flow through an orifice, and flow in a cylindrical tube.



The simplest well-defined model for blood : a two-fluid model with fixed  $\delta$



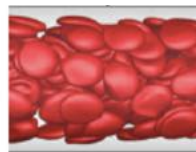
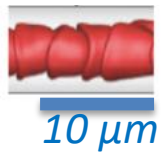
$$\mu_{app} = \mu_{out} / [(1 - \delta/R)^4 (\mu_{out}/\mu_{in} - 1) + 1]$$

$$\mu_{in} du/dy]_- = \mu_{out} du/dy]_+$$

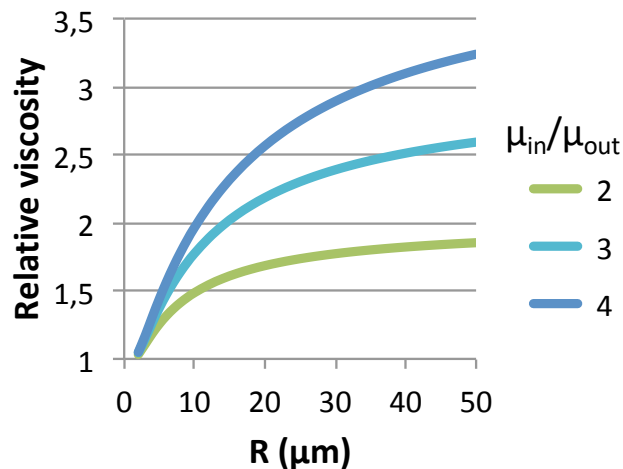
# The blood as an equivalent fluid

## Apparent viscosity and relative viscosity

Apparent and relative viscosities are not intrinsic properties of the blood; they are properties of the blood and blood vessel interaction, and depend on the data reduction procedure. There are as many definitions for apparent viscosities as there are good formulas for well-defined problems. Examples are: Stokes flow around a falling sphere, channel flow, flow through an orifice, and flow in a cylindrical tube.



The simplest well-defined model for blood : a two-fluid model with fixed  $\delta$

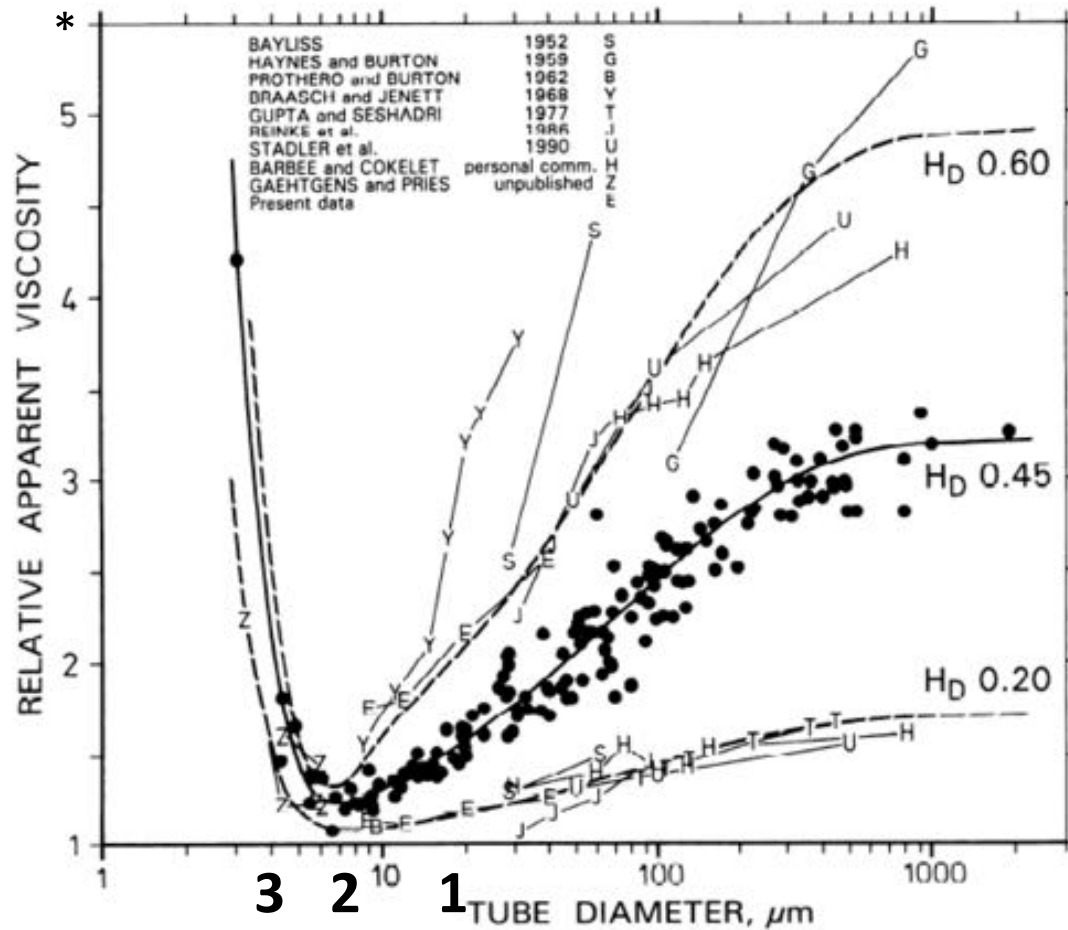


$$\mu_{app} = \mu_{out} / [(1 - \delta/R)^4 (\mu_{out}/\mu_{in} - 1) + 1]$$

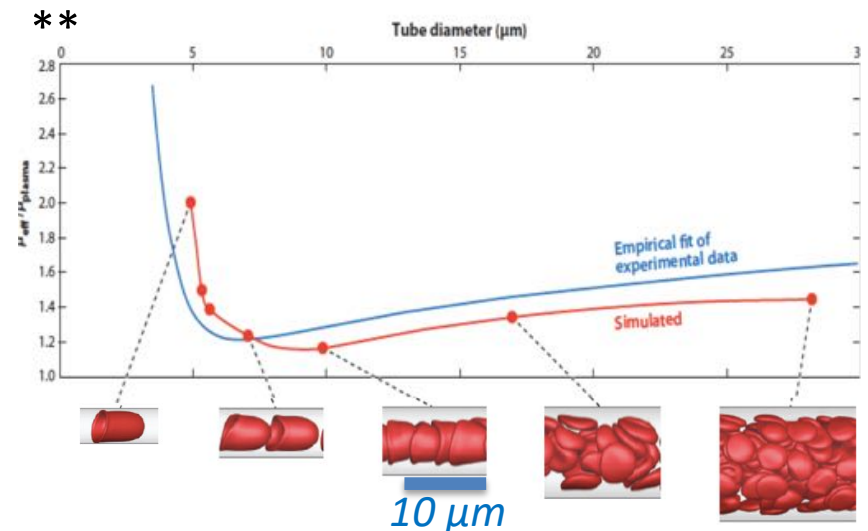
$$\mu_{rel} = \mu_{app} / \mu_{plasma} = [Q_{plasma} / Q_{blood}]_{same \Delta P}$$

# The blood as an equivalent fluid

## Apparent viscosity and relative viscosity



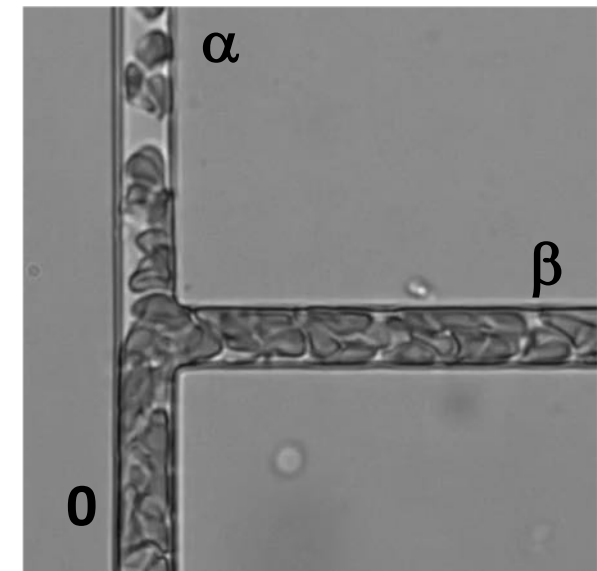
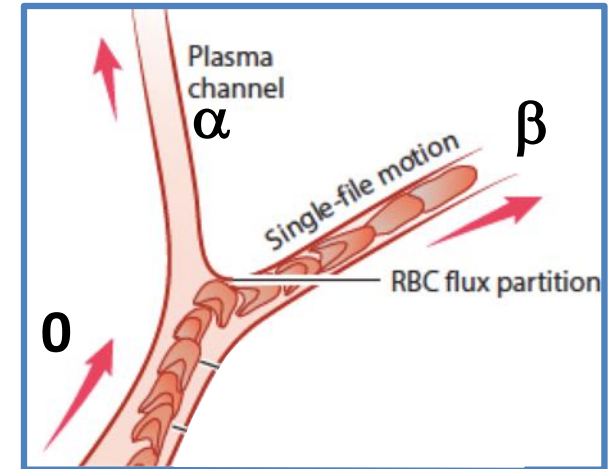
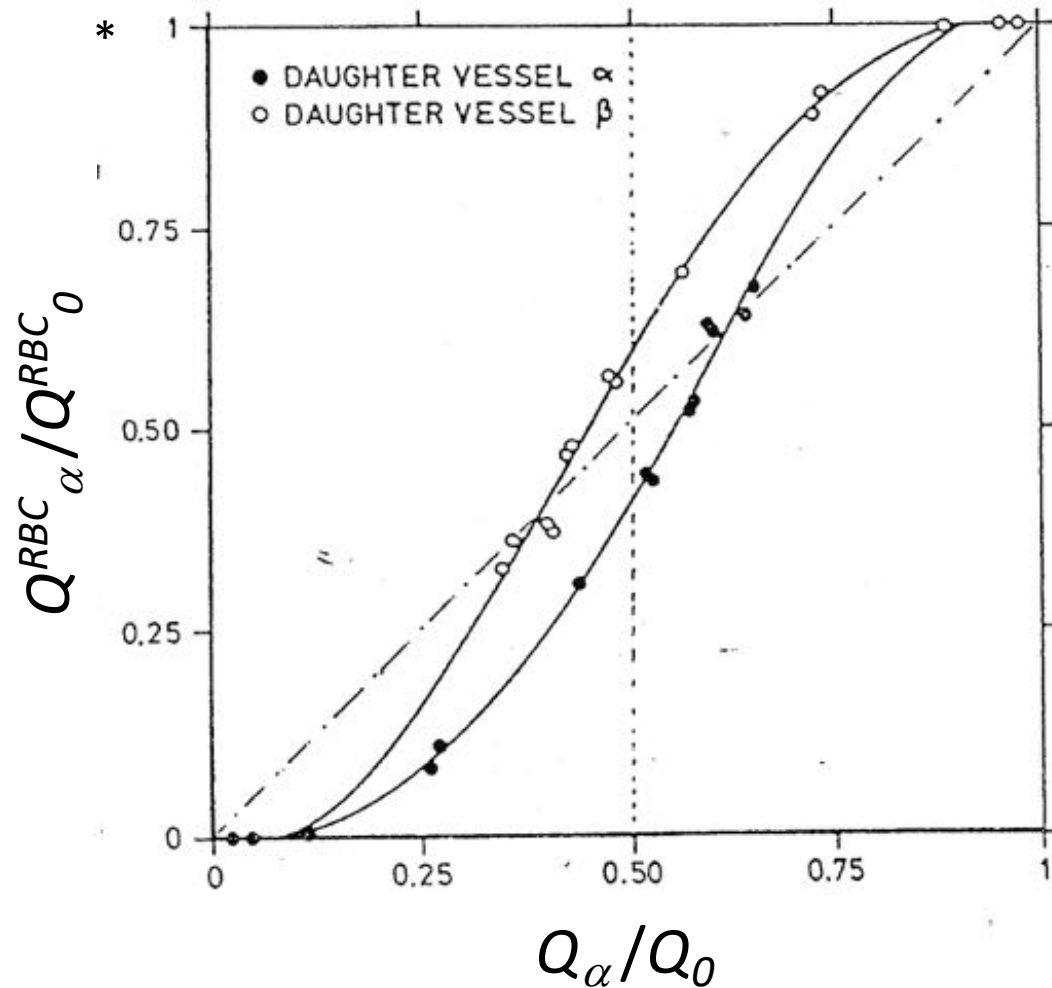
$$\mu_{rel} = f(d, H)$$





# The blood as an equivalent fluid

## Bifurcation law



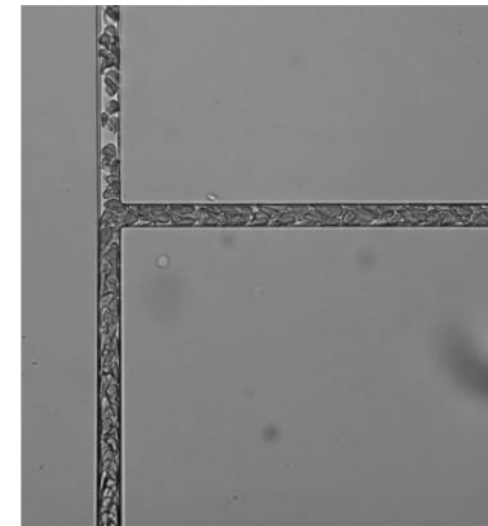
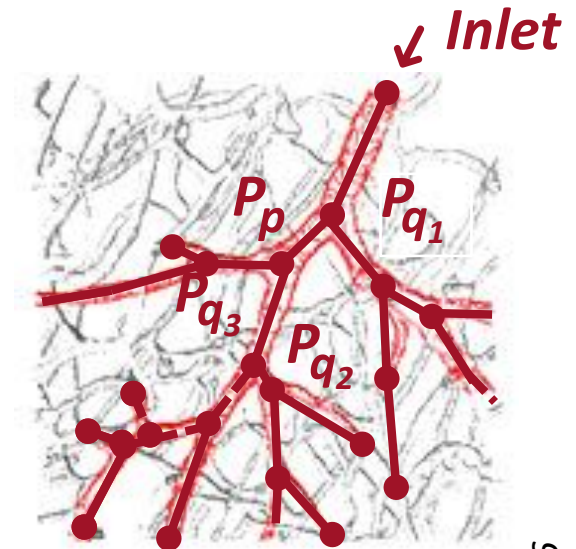
\*\*

$$Q_{\alpha}^{RBC} / Q_0^{RBC} = g(Q_{\alpha} / Q_0, H_0, d_0, d_{\alpha} / d_0, d_{\beta} / d_0)$$

# Blood flow in networks

## Modeling framework

- ❑ The vascular network is represented as a graph, with prescribed BCs
- ❑ A uniform hematocrit distribution ( $H=0.45$ ) is assumed
- ❑ The viscosity  $\mu_{app}$  in each vessel depends on its diameter  $d$  and hematocrit  $H$  (Pries et al. 1996)
- ❑ Their resistance  $R$  is deduced
$$R = (128 \mu_{app} L) / (\pi d^4)$$
- ❑ Mass conservation ( $\sum_i [R_{pq_i}^{-1} (P_{q_i} - P_p)] = 0$ ) is solved yielding the **pressures** at each node
- ❑ **Flow rate** in each vessel is deduced using the linear relationship  $Q_{pq_i} = R_{pq_i}^{-1} (P_{q_i} - P_p)$
- ❑ **Hematocrit** in each vessel is deduced using a bifurcation law for phase separation (Pries et al. 1996)



Roman et al. Biomicrofluidics 2016

# Blood flow in networks

## Modeling framework



### Blood Flow in Microvascular Networks

#### Experiments and Simulation

A.R. Pries, T.W. Secomb, P. Gaehtgens, and J.F. Gross

#### □ Simulations

- The vascular network is described as a graph
- The blood is viewed as an equivalent fluid (i.e. at vessel scale)
- Its non-linear rheological properties are described by empirical relationships
- A first guess on their coefficient is based on experiments (in vivo or in glass tubes)

#### □ Simulation/Experiments

- The values of these coefficients are adjusted to best match the *in vivo* measurements
- This gives insights on biophysical aspects of blood flow in the microvasculature



# Blood flow in networks

## Modeling framework

- Flow direction

913 seg.

546 seg.

436 seg.

TABLE 2. Evaluation of Model Simulation in the Rat Mesentery Under Six Different Sets of Input Conditions

Condition	$H_D$	Viscosity law	Network A		Network B		Network C	
			$N_{INV}$	$r^2_{H_D}$	$N_{INV}$	$r^2_{H_D}$	$N_{INV}$	$r^2_{H_D}$
1	Predicted	In vitro	44 (4.8%)	0.060	30 (5.5%)	0.133	13 (3.0%)	0.177
2	Predicted	Uniform	25 (2.7%)	0.080	24 (4.4%)	0.163	7 (1.6%)	0.226
3	Predicted	Modified	15 (1.6%)	0.148	25 (4.6%)	0.183	2 (0.5%)	0.333
4	Measured	In vitro	69 (7.6%)	...	47 (8.6%)	...	22 (5.0%)	...
5	Measured	Uniform	25 (2.7%)	...	24 (4.4%)	...	7 (1.6%)	...
6	Measured	Modified	20 (2.2%)	...	29 (5.3%)	...	3 (0.7%)	...

$H_D$ , discharge hematocrit;  $N_{INV}$ , segments in which the predicted flow direction was inverted relative to observation;  $r^2_{H_D}$ , the squared coefficient of correlation between predicted and measured discharge hematocrits in all vessel segments.  $H_D$  was either predicted from the phase-separation effect or measured in vivo.

(Pries *et al.* 1990)

- Minimisation of the number of vessels with wrong flow direction

➔ *In vivo* viscosity law

# Blood flow in networks

## Modeling framework

### ■ In vivo viscosity law

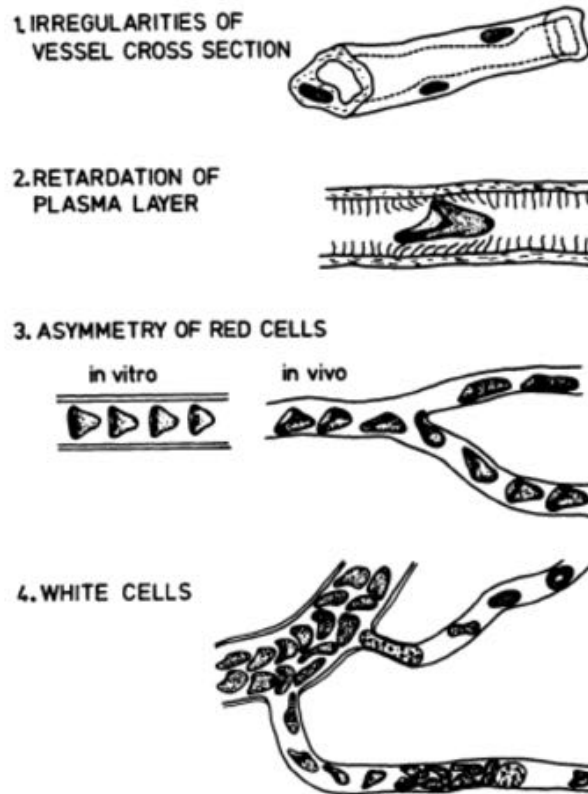
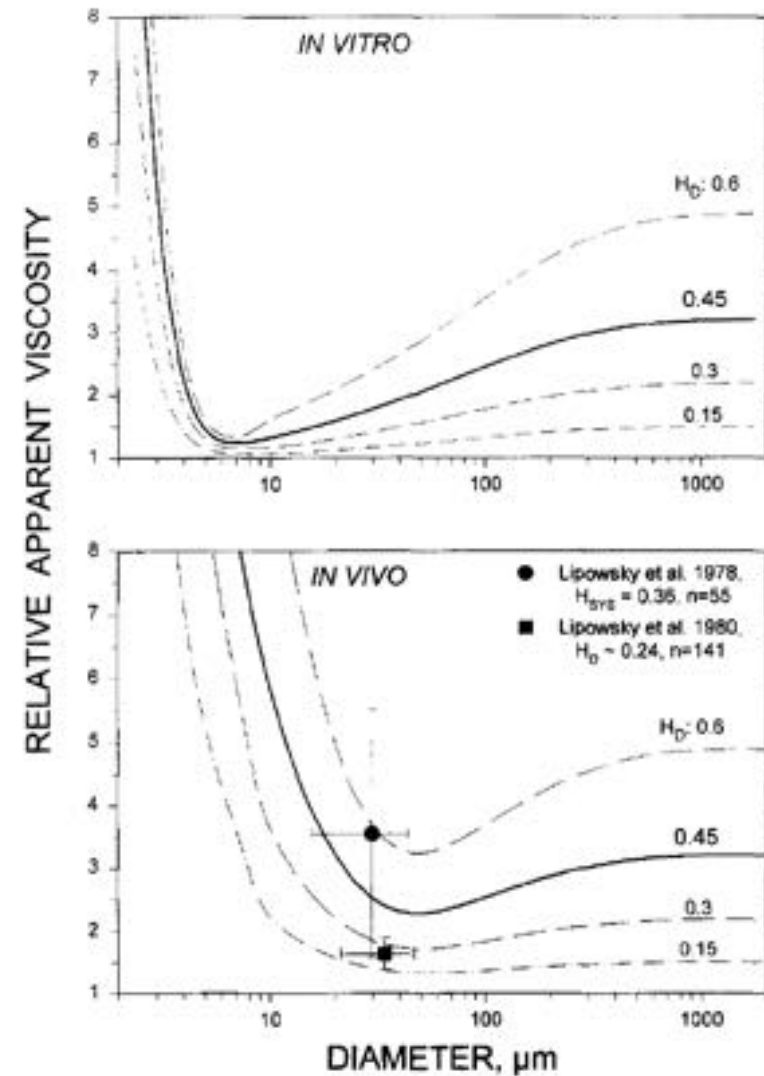


FIGURE 8. Schematic drawings showing possible reasons for high resistance in small microvessels.

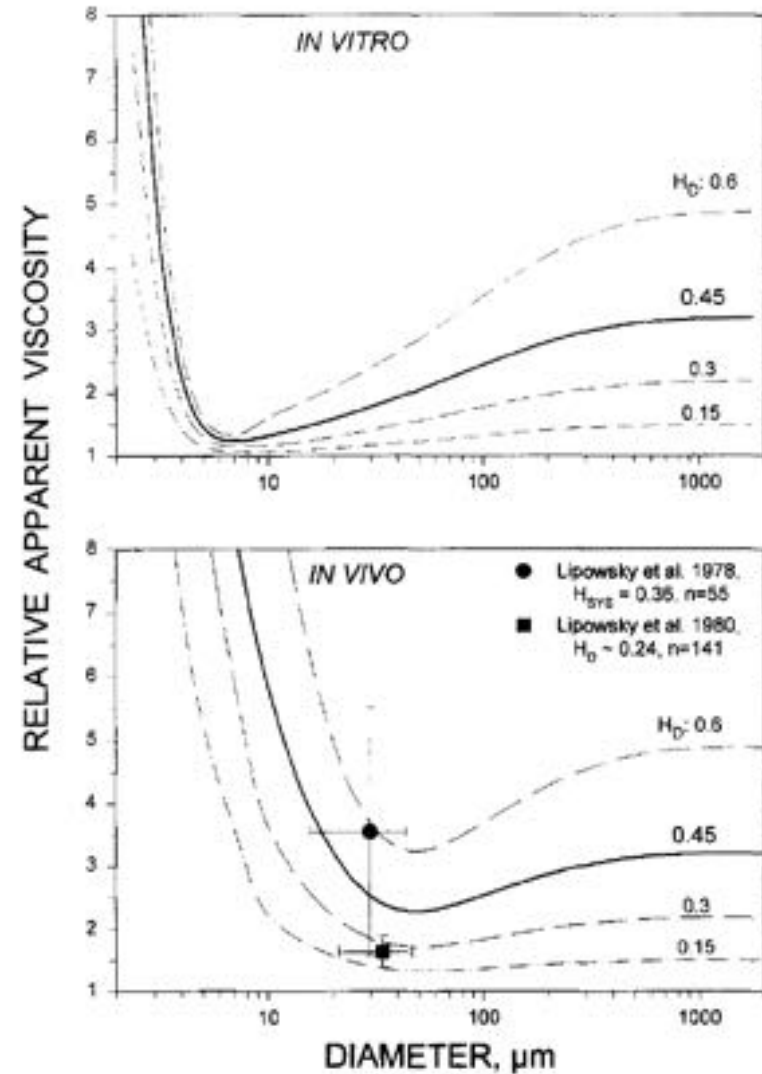


# Blood flow in networks

## Modeling framework

- In vivo viscosity law

comparison with the original viscosity law (Figure 1) shows a marked viscosity increase in the low diameter range. It should be cautioned, however, that this viscosity relation cannot be considered as a precise, quantitative representation of effective blood viscosity in vivo. Model results with comparable values of  $N_{INV}$  and  $r^2_{H_0}$  could be achieved with a number of viscosity relations that exhibit significant quantitative differences to those described by Equation 7. All these relations, however, are qualitatively similar in predicting a viscosity increase with decreasing vessel diameter below about 15  $\mu\text{m}$ . The results presented

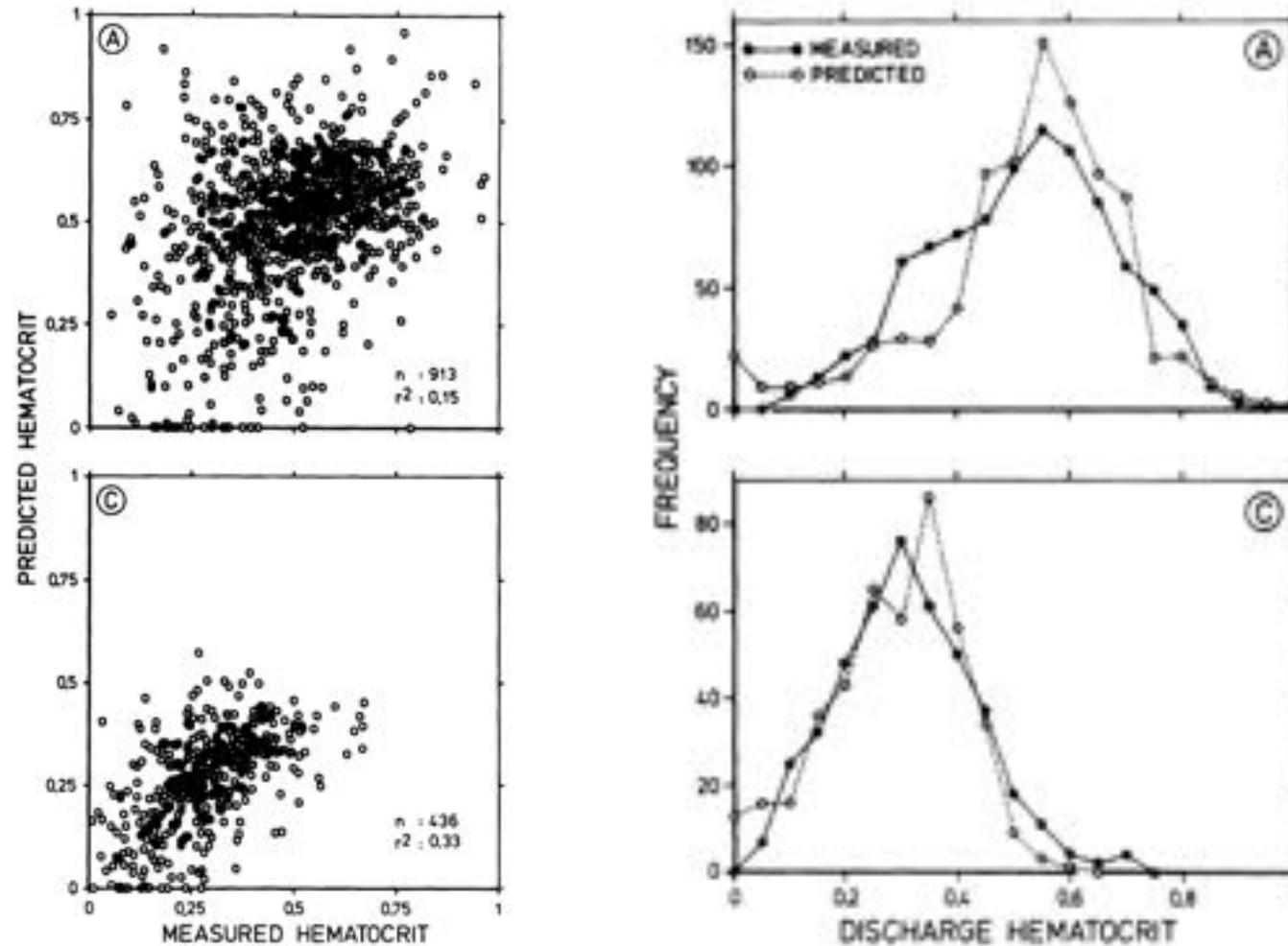




# Blood flow in networks

## Validation against in vivo data

### □ Hematocrit

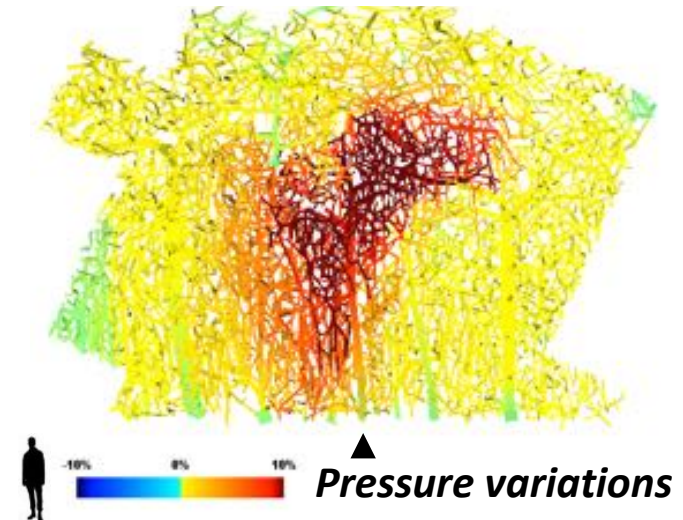
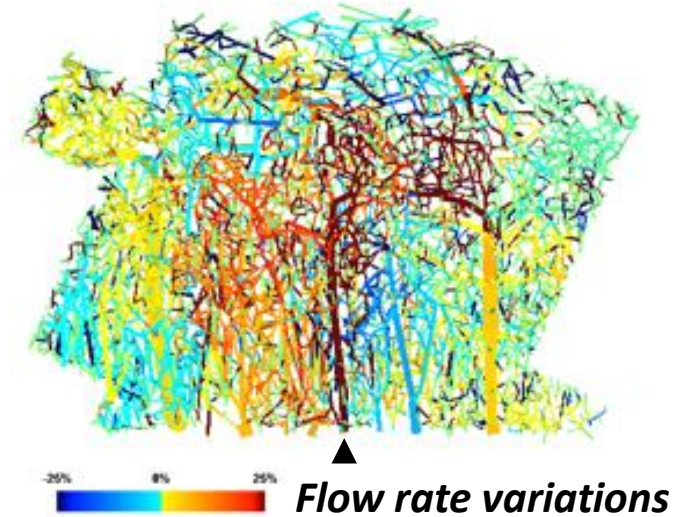


# Blood flow in networks

## Back to the brain...

### □ Strengths\*

- Changes in vascular structure (vessel occlusions, vessel dilations) can be easily imposed
- Post-processing is versatile
  - Spatial maps : baseline parameters, parameter variations induced by changes in structure
  - Spatial averages
  - Passive tracer injection : vascular territories, transit time distributions, lagrangian analysis
  - Correlations between variables: e.g. flow variation =  $f(\text{topological distance})$

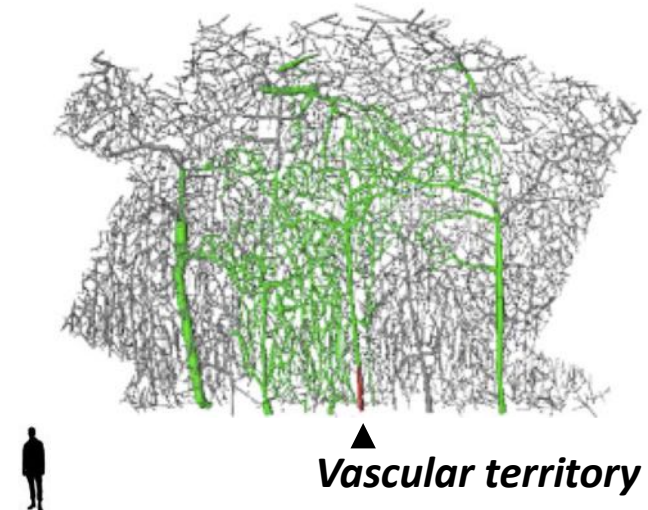
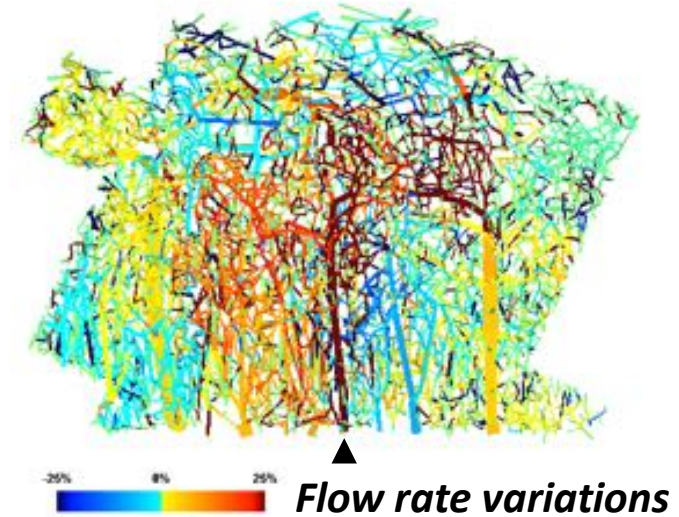


# Blood flow in networks

## Back to the brain...

### □ Strengths\*

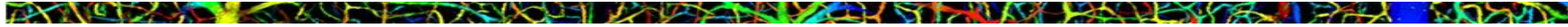
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  - Correlations between variables: e.g. flow variation =  $f(\text{topological distance})$





# Blood flow in networks

Back to the brain...



## □ Weaknesses

- Need of comprehensive high resolution anatomical data
- Sensitivity to vessel diameters, boundary conditions, domain size ?
  - Critical for the exploitation of human data (post-mortem, slices)



# Blood flow in networks

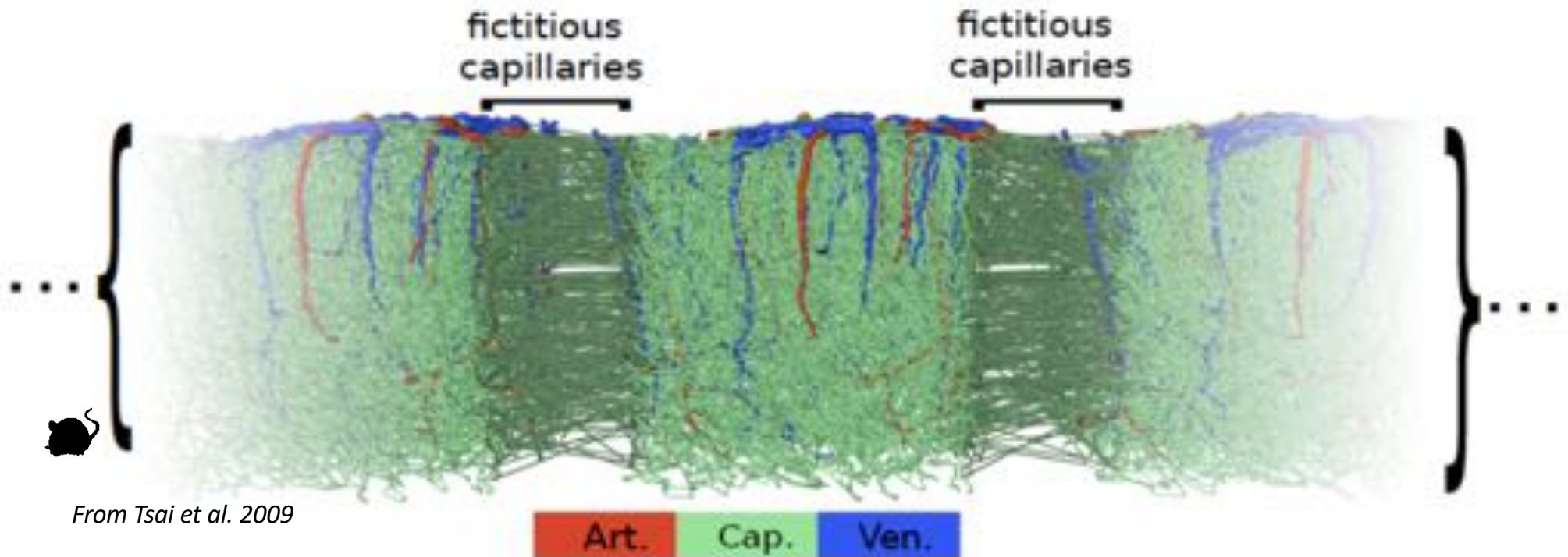
Back to the brain...



## □ Weaknesses

- Need of comprehensive high resolution anatomical data
- Sensitivity to vessel diameters, boundary conditions, domain size ?
  - ➔ Critical for the exploitation of human data (post-mortem, slices)

## □ Validation vs. mice experimental data in large domains

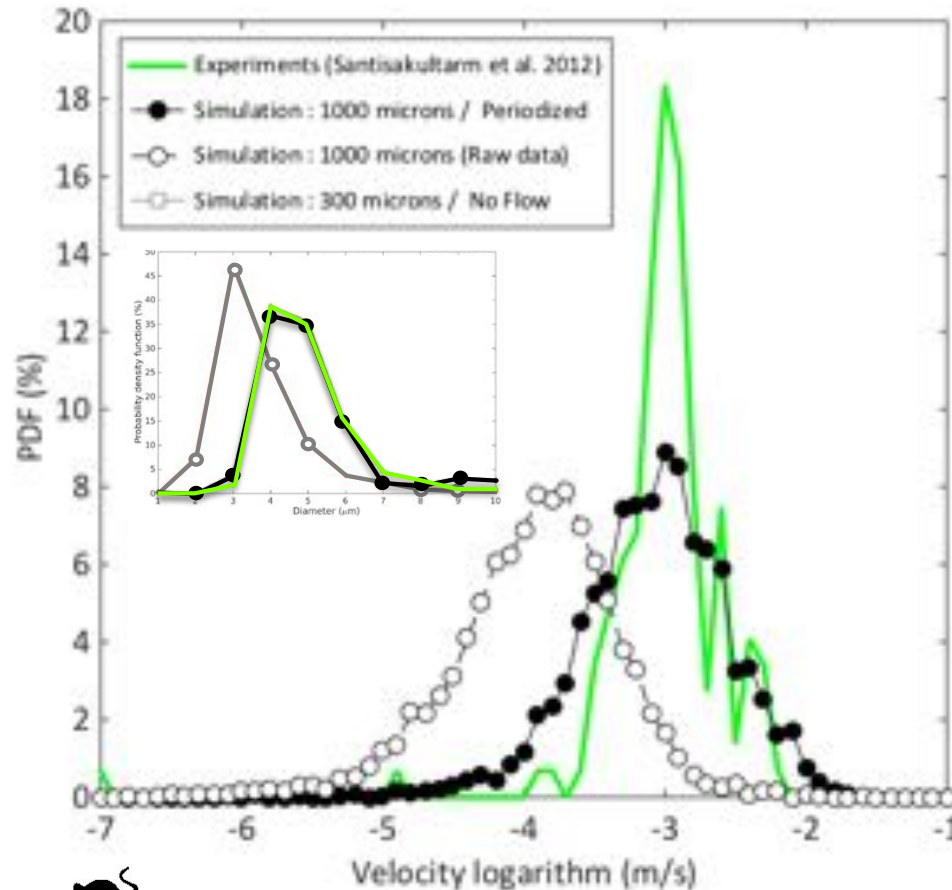


# Blood flow in networks

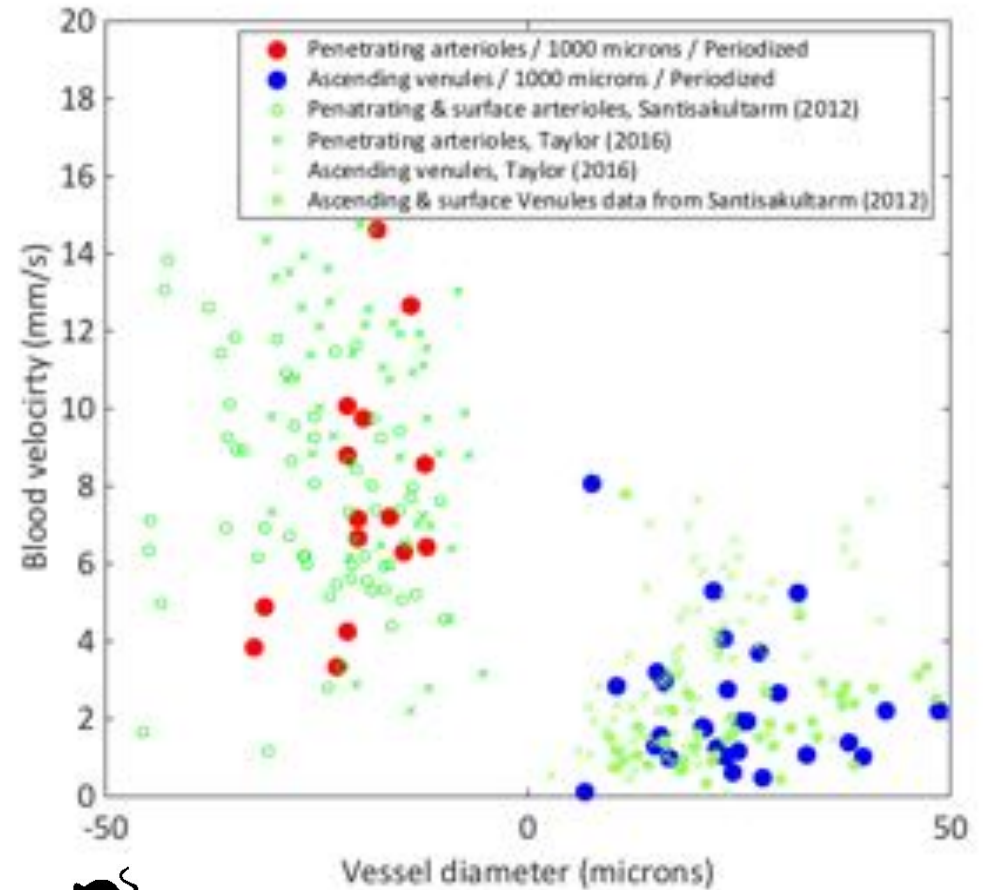
Back to the brain...



## Comparison of velocity distributions with TPSLM experiments



**In capillaries  
(less than 300  $\mu\text{m}$  under brain surface)**



**In penetrating arterioles  
and ascending venules**

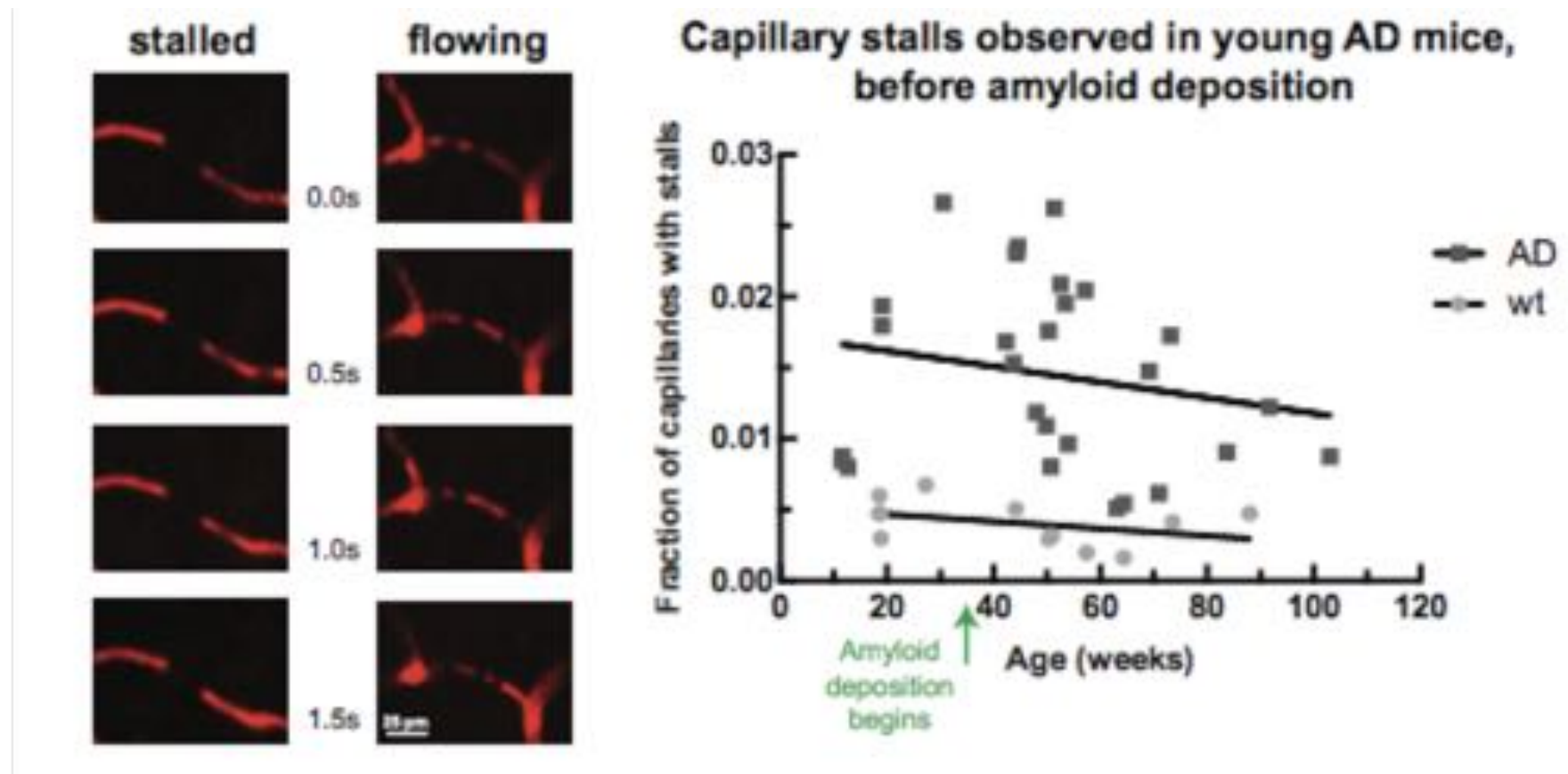




# Blood flow in networks

## Vascular component of AD: impact of small initial perturbation

- In APP/PS1 mice, ~2% capillaries are stalled by leucocytes\*



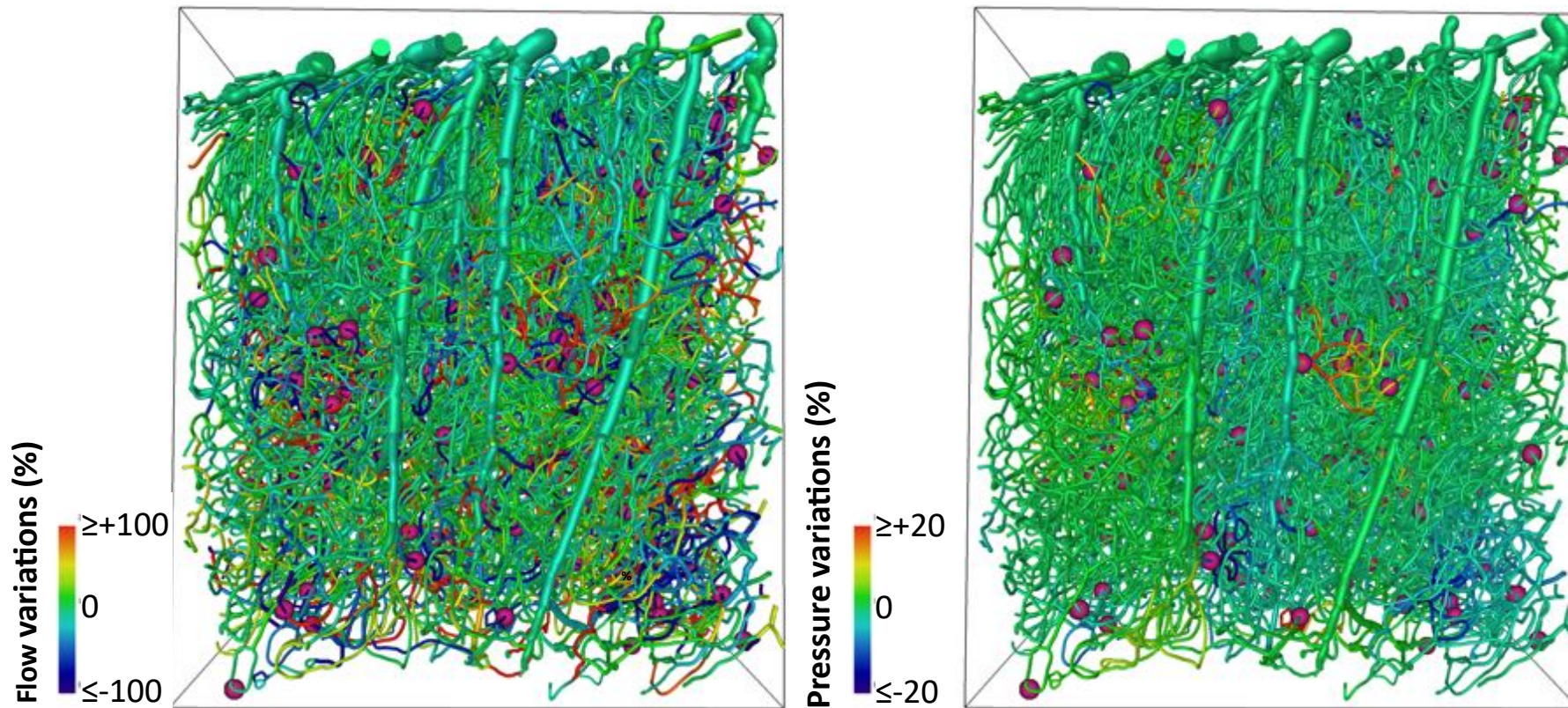
- Pharmacological removal of stalls increases blood flow and improves memory performance

# Blood flow in networks

## Vascular component of AD: impact of small initial perturbation

### □ Effect of random capillary occlusions on blood flow patterns

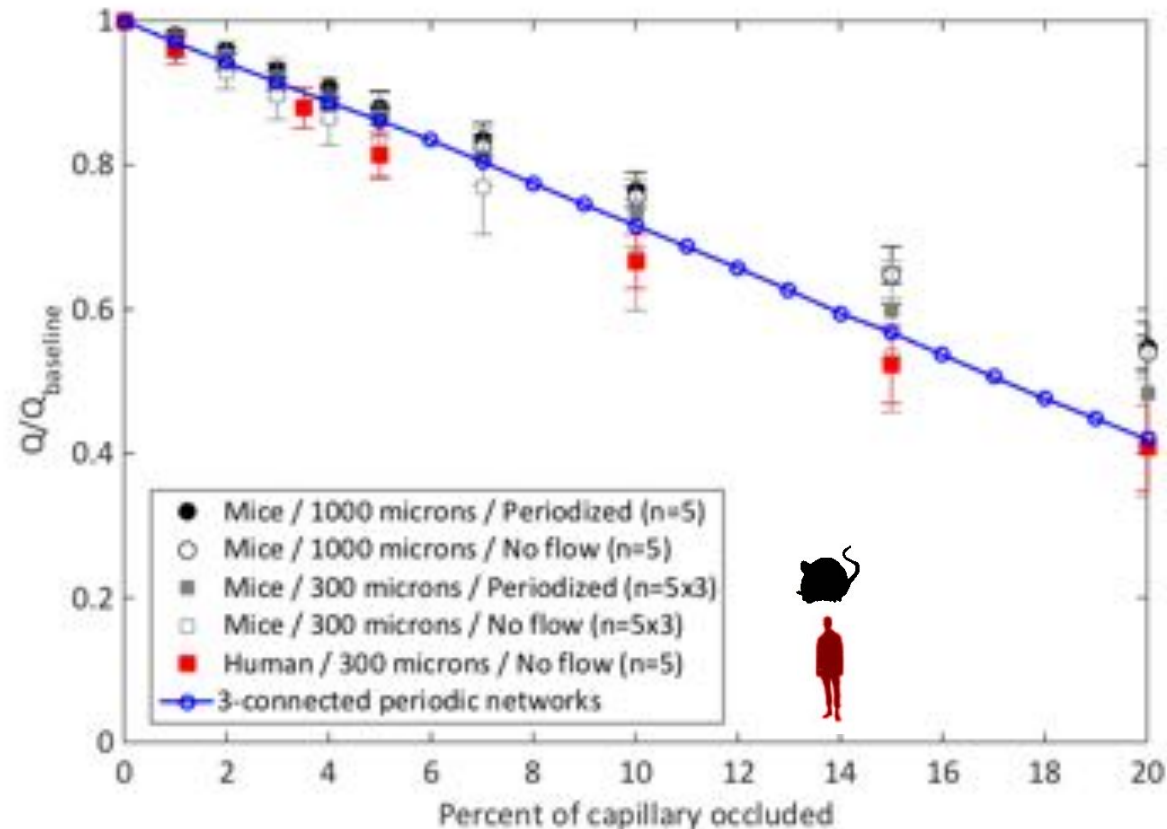
- 2% capillary occlusions
- Assuming no changes in perfusion pressure, no diameter variations



# Blood flow in networks

## Vascular component of AD: impact of small initial perturbation

### □ Effect of random capillary occlusions on global blood flow



➔ No threshold (versus Hudetz, Microvasc. Res. 1993)

➔ Similar blood flow reduction mice, humans and 3-connected mesh

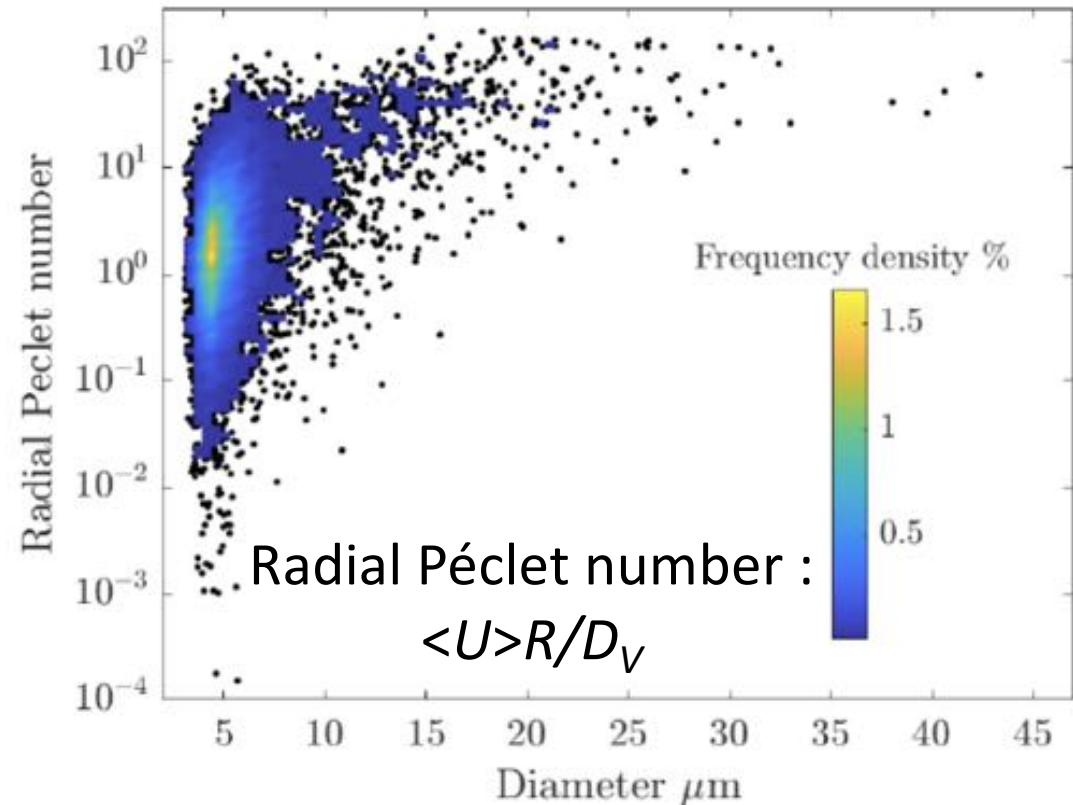
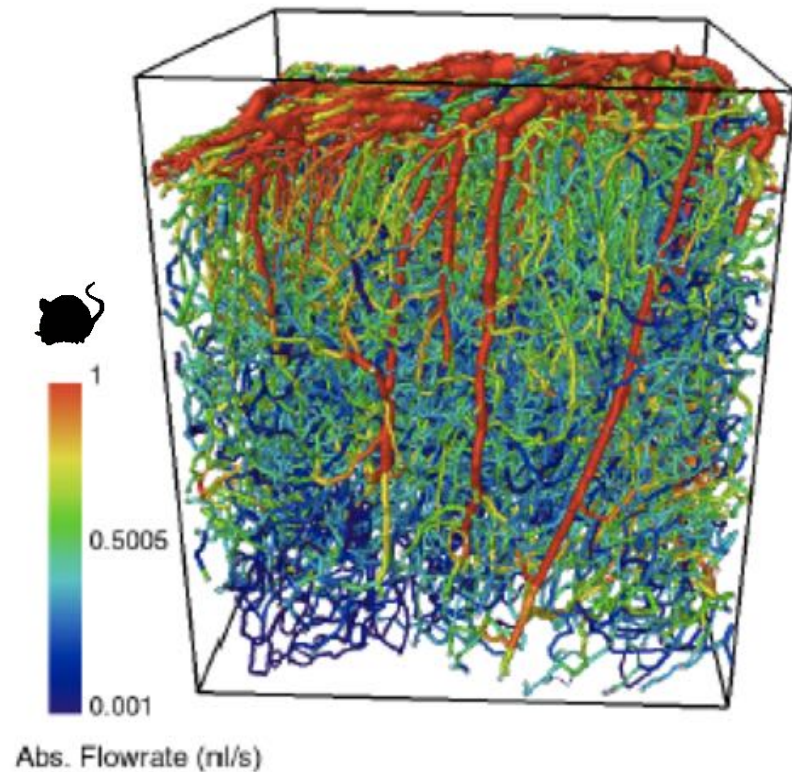


# Outline



- **Brain versus other organs**
  - What is generic ?
  - What is specific ?
- **Why study brain microcirculation ?**
  - In health
  - In disease
- **Investigation tools and associated scales**
- **Blood flow in networks**
- **Mass transfers in networks**
- **Blood flow at organ scale**

# Distribution of Péclet numbers



## Extension to mass transfers :

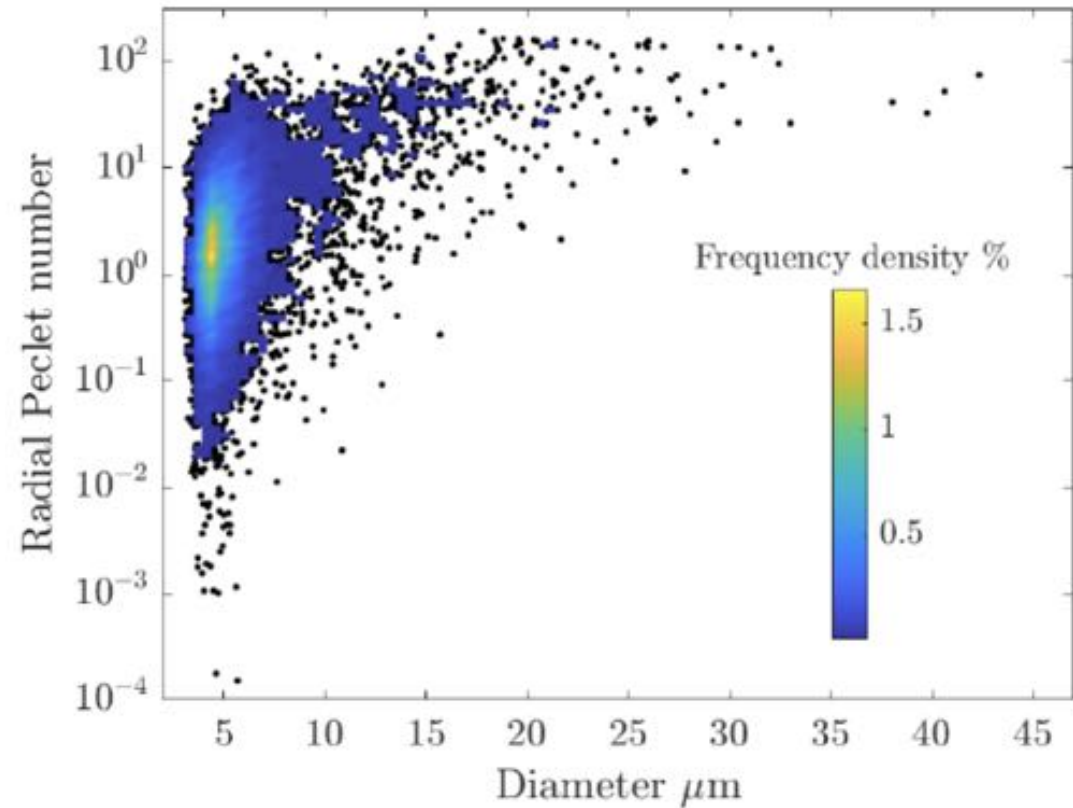
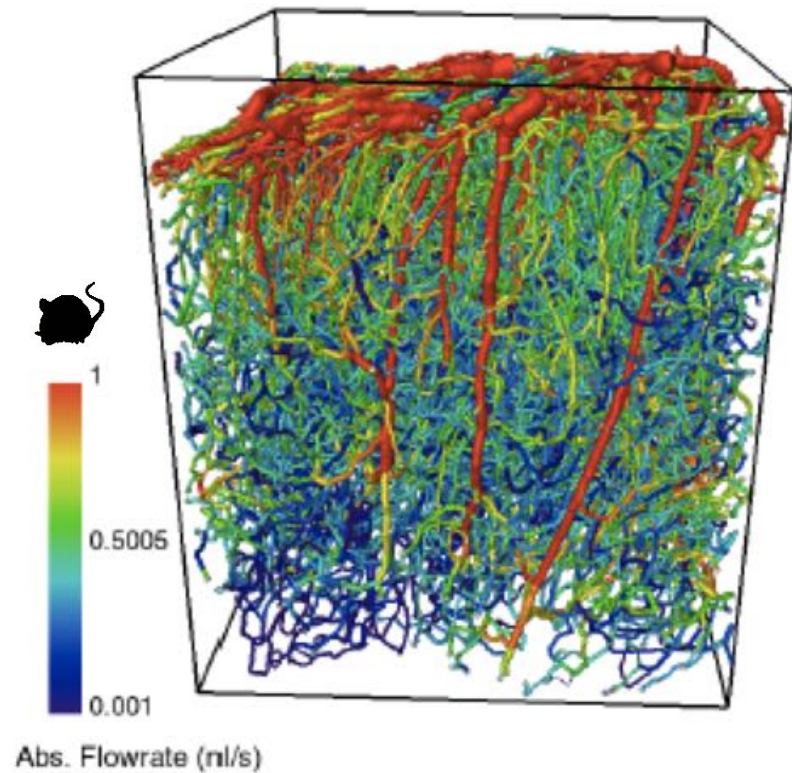
What are the errors of a network approach where radial concentration gradients are neglected in vessels

(well-mixed model\*) ?

PhD Maxime Berg

# Extension to mass transfers :

## Distribution of Péclet numbers



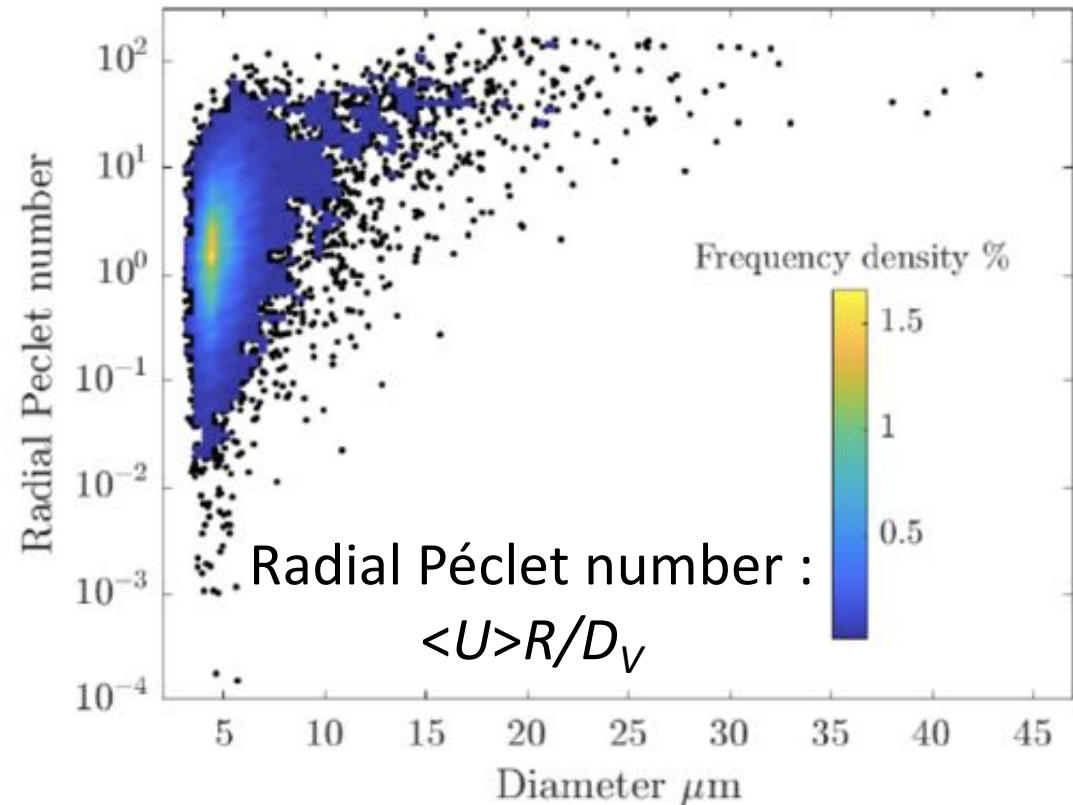
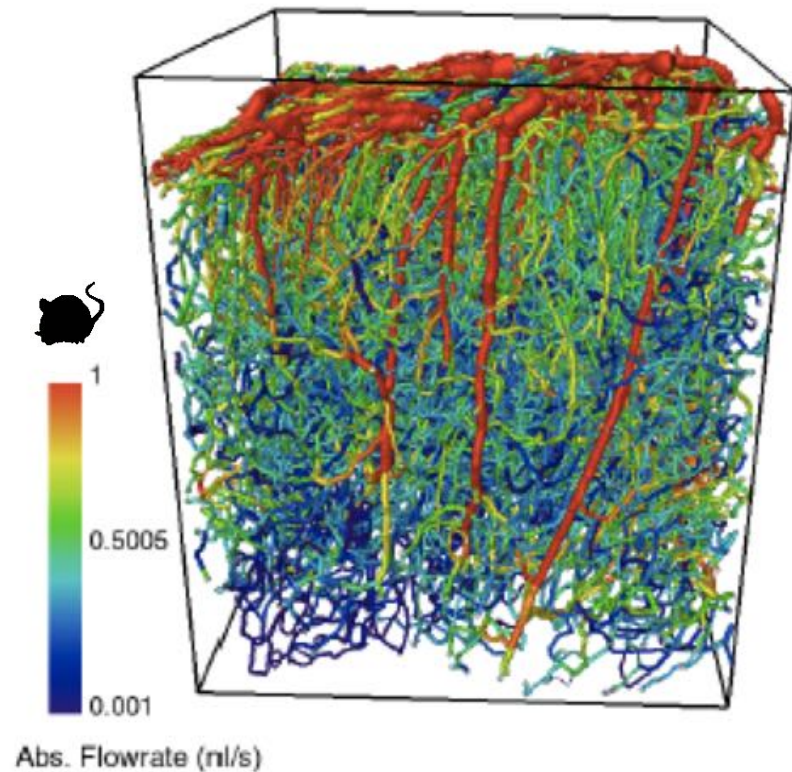
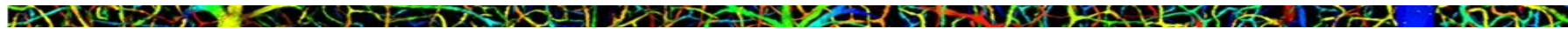
Radial Péclet number :  $Pe_R = \langle U \rangle R / D_V$   
Length for complete mixing :  $L \sim Pe_R R$





# Extension to mass transfers :

## Distribution of Péclet numbers



What are the errors of a network approach where radial concentration gradients are neglected in vessels (well-mixed model\*) ?

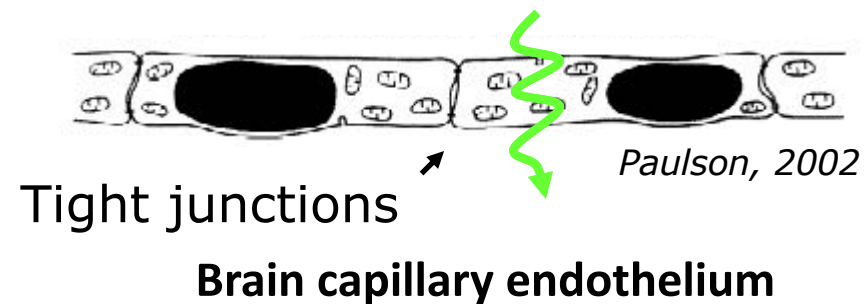
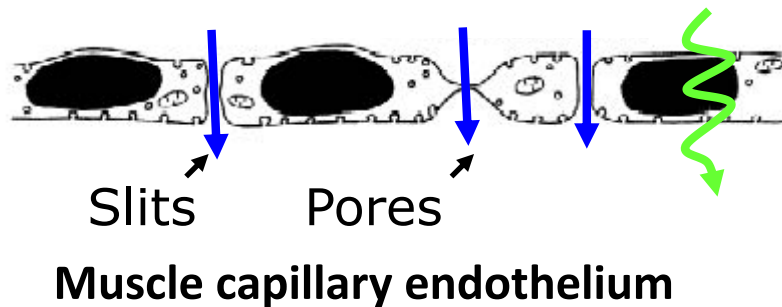
PhD Maxime Berg

# Mass transfers at vessel scale

## Phenomenology and simplifications

### □ Blood brain barrier

- Ionic/Osmotic balance (neuronal communication / constrained volume)
- Protection against neuro-toxics



Damköhler number :

### □ Boundary conditions at vessel surface

- Flux continuity:  $D_V \nabla C_V \cdot \mathbf{n} = D_{\text{tissue}}^* \nabla C_{\text{tissue}} \cdot \mathbf{n}$
- Membrane condition :  $D_V \nabla C_V \cdot \mathbf{n} = -K_m (C_V - \lambda C_{\text{tissue}})$

# Mass transfers at vessel scale

## A 1D effective equation within vessels ( $\epsilon \ll 1$ )

### □ Asymptotic limit of small tissue concentration

- Membrane condition :  $\mathcal{D}_V \nabla C_V \cdot \mathbf{n} = -K_m (C_V - \lambda C_{\text{tissue}})$

→ Robin condition :  $\mathcal{D}_V \nabla C_V \cdot \mathbf{n} = -K_m C_V$

### □ Volume averaging with closure (coll. Y. Davit, M. Quintard)

$$\frac{\partial \langle C_V \rangle}{\partial t} + U_{eff} \cdot \nabla \langle C_V \rangle - D_{eff} \nabla^2 \langle C_V \rangle + K_{eff} \langle C_V \rangle = 0$$

where  $U_{eff} = Pe (1 + U_+)$

$$\epsilon = \frac{R}{L}$$

$$D_{eff} = 1 + \frac{(\epsilon Pe)^2}{Pe_c^2}$$

$$Pe = \frac{\langle U \rangle L}{D_V}$$

$$K_{eff} = \frac{8\epsilon^{-1} Da_m}{\epsilon Da_m + 4}$$

$$Da_m = \frac{K_m L}{D_V}$$

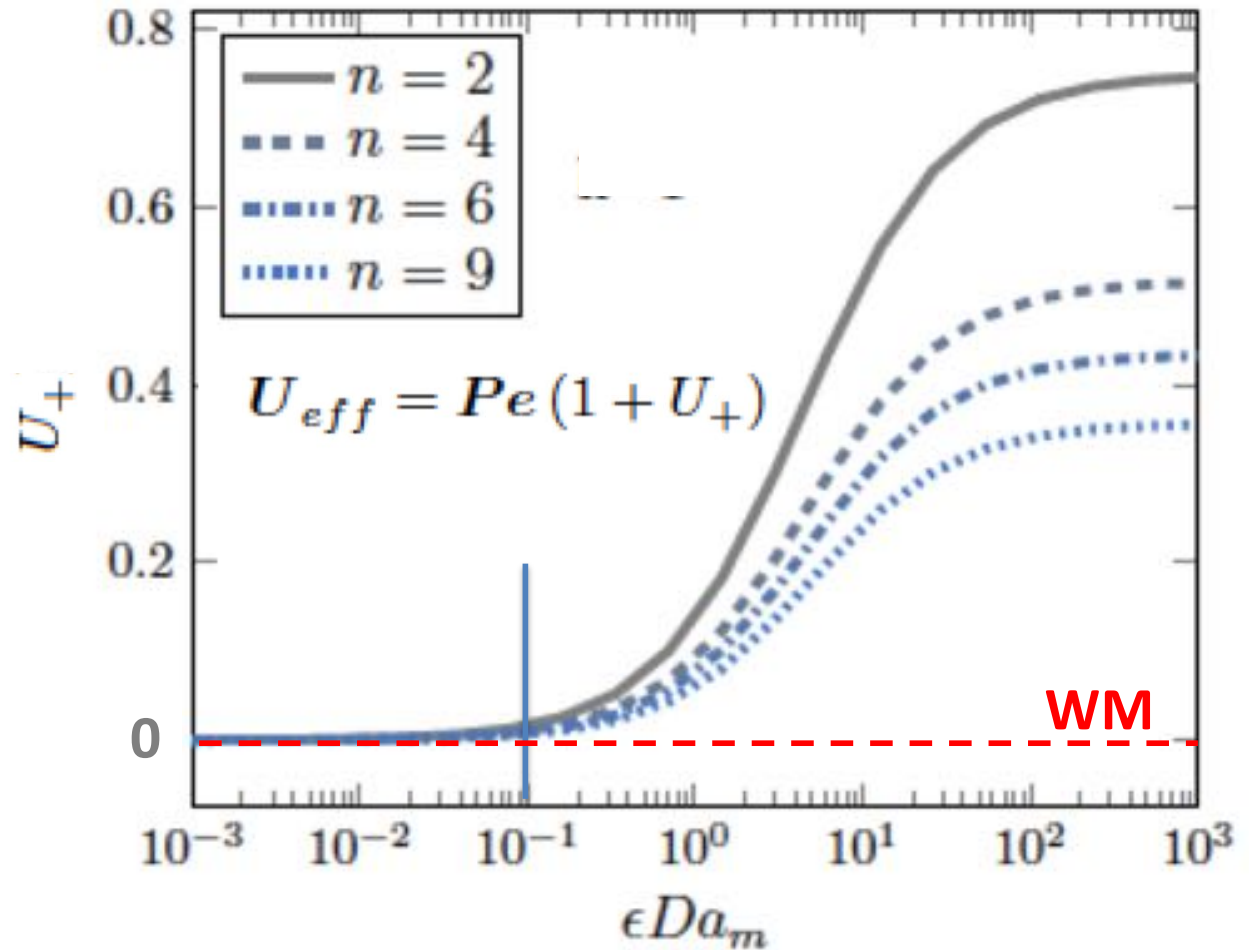
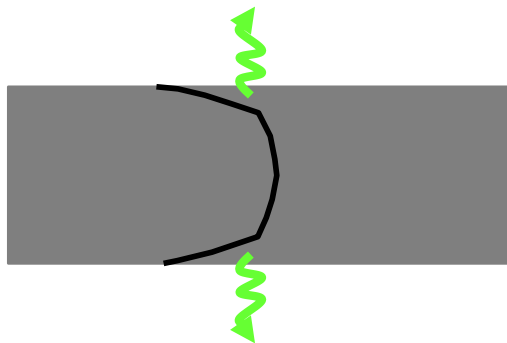
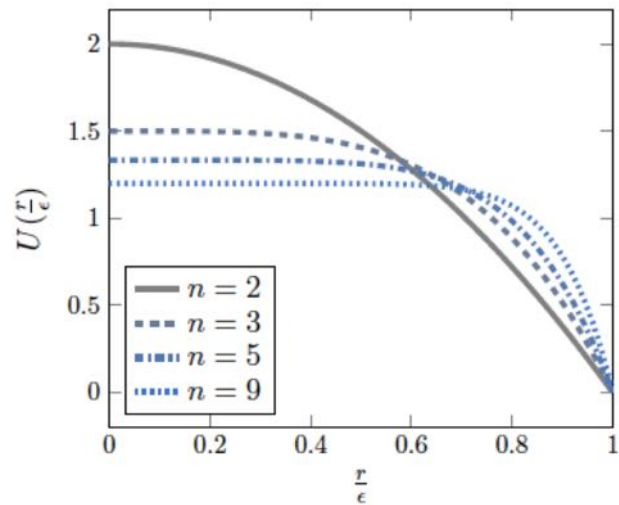
and  $U_+$  and  $Pe_c$  depend on  $\epsilon$ ,  $Da_m$  and the shape of the velocity profile.



# Mass transfers at vessel scale

A 1D effective equation within vessels ( $\varepsilon \ll 1$ )

## □ Effective velocity

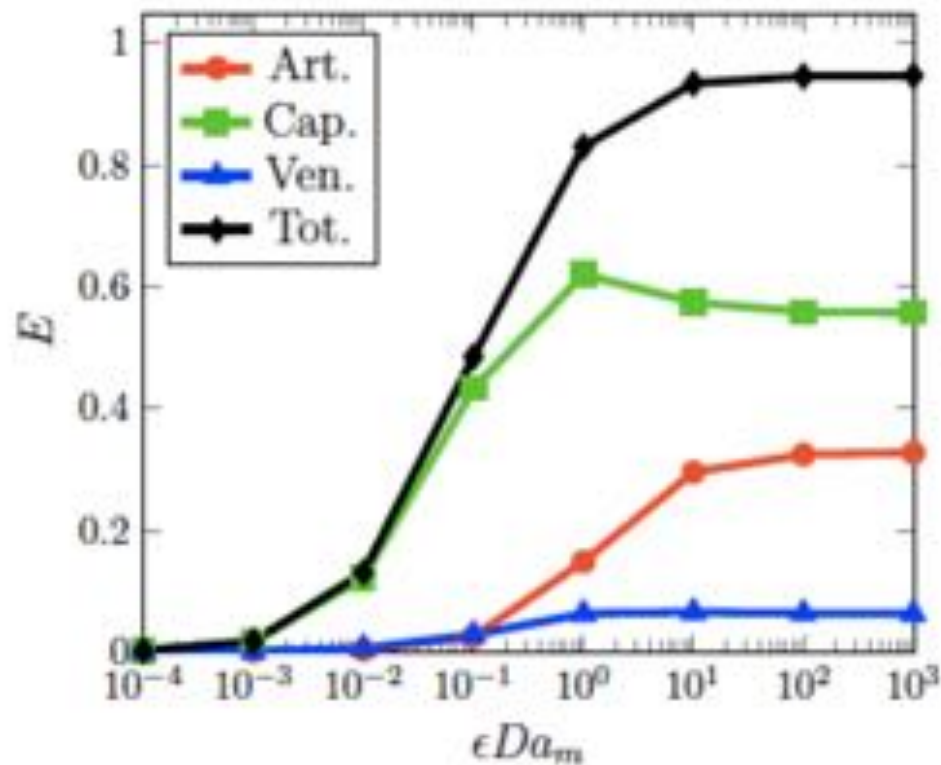


# Impact at network scale

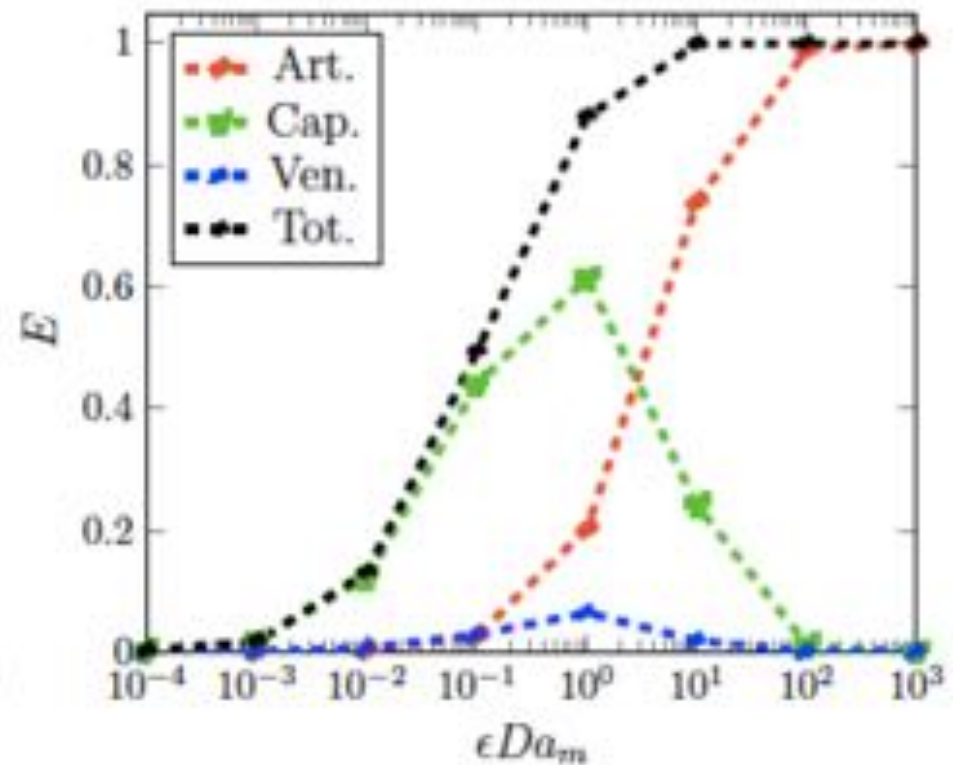
## Poiseuille velocity profiles

- Stationnary extraction (flux through vessel walls / entry flux)

Effective model



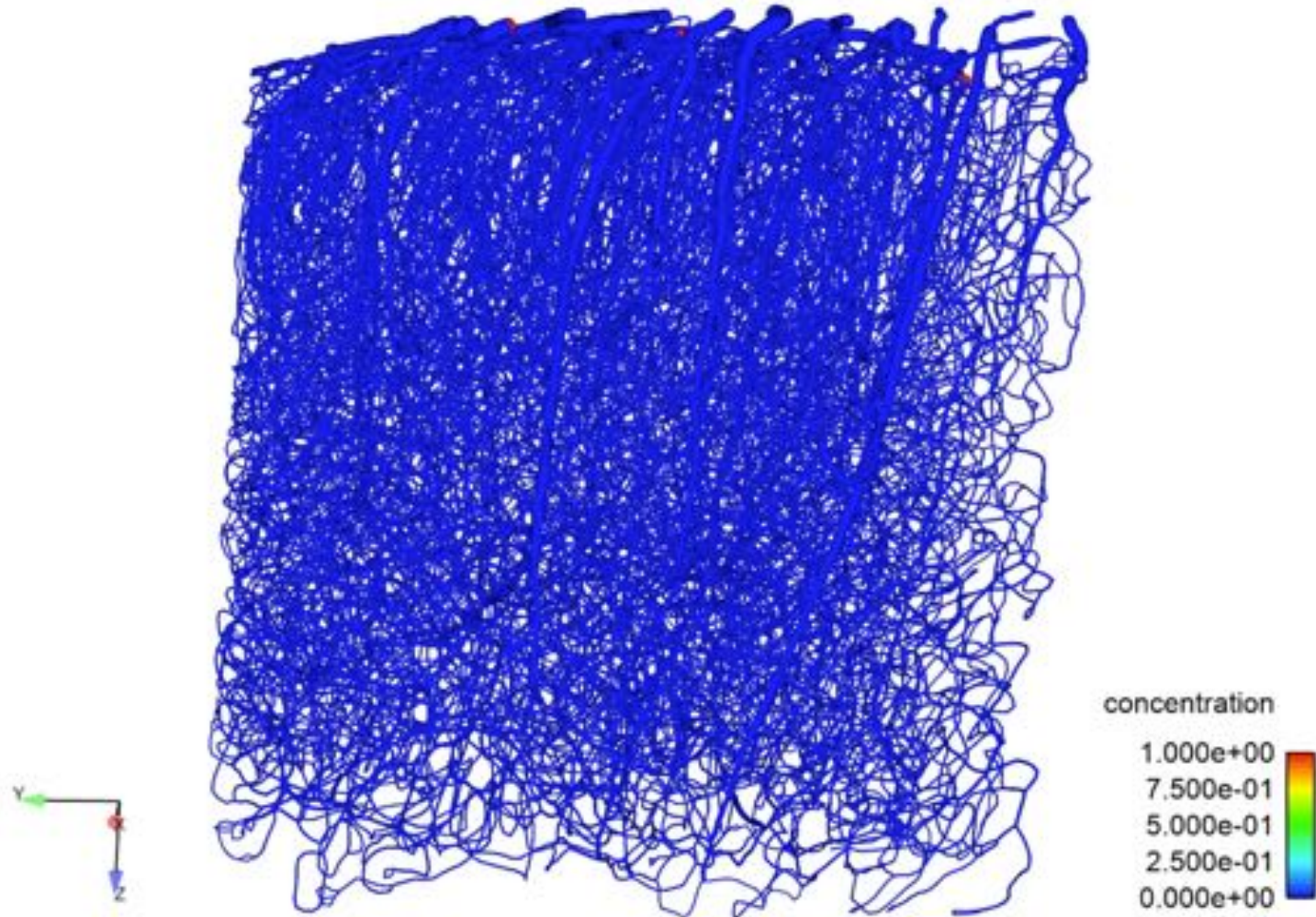
WM model



# Mass transfers at network scale

## Network model

### □ Intravascular tracer



### □ Coupling with tissue



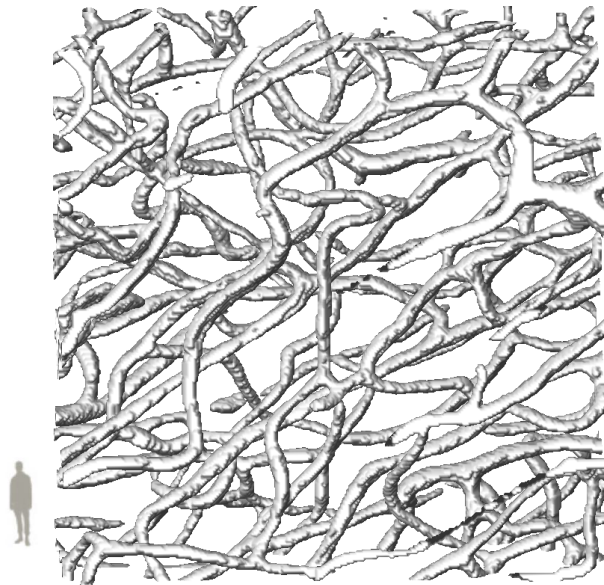
# Outline



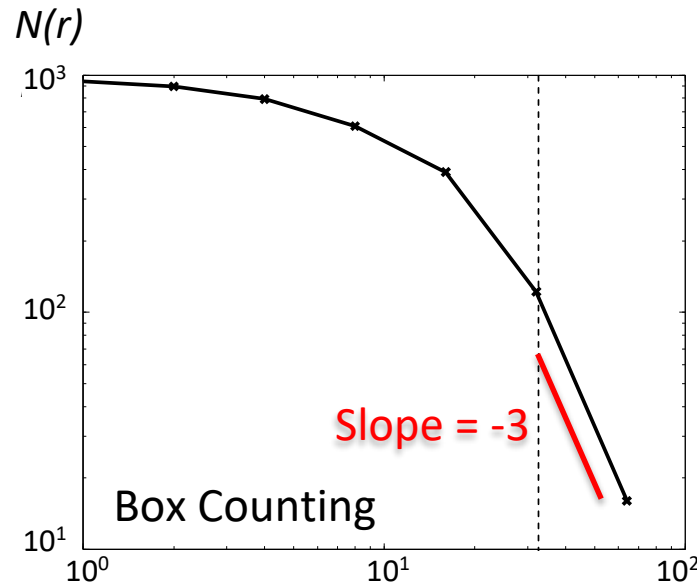
- **Why study brain microcirculation ?**
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  - In disease
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# Extension to larger scales

## Model reduction at mesoscale



**Anatomical data\***



**Multi-scale analysis\***



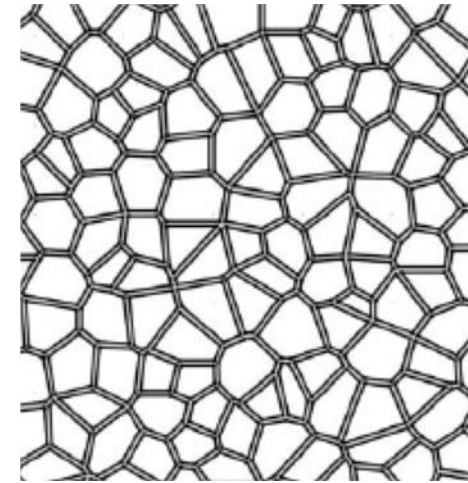
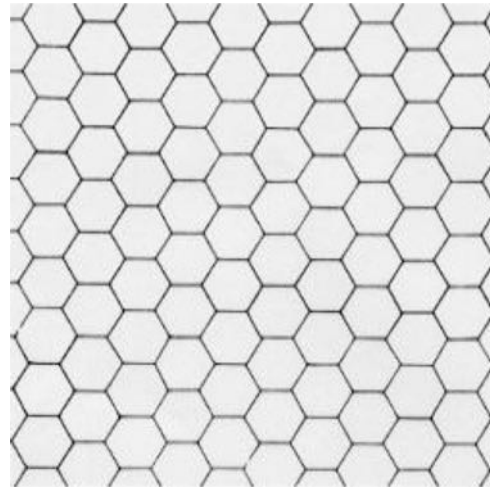
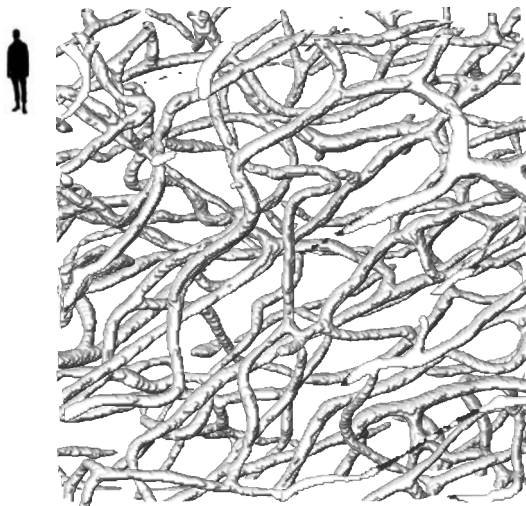
**Equivalent continuum**



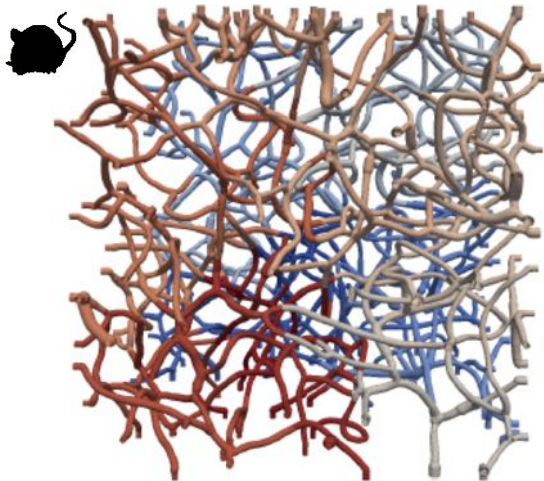
\*Lorthois & Cassot, Journal of Theoretical Biology 2011, \*\* Peyrounette et al. PLOS ONE 2018

# Note : Model capillary networks

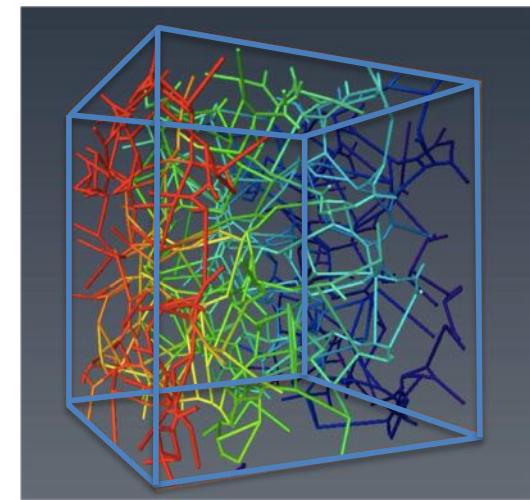
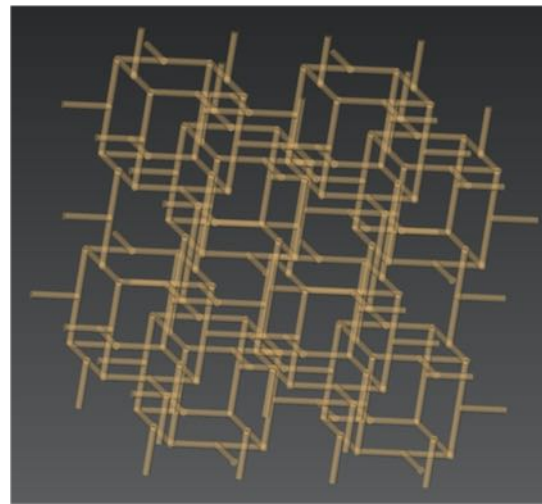
## □ Space-filling, 3-connected, looped networks



Lorthois et al, JTB 2010



Courtesy Tsai, Blinder, Kleinfeld

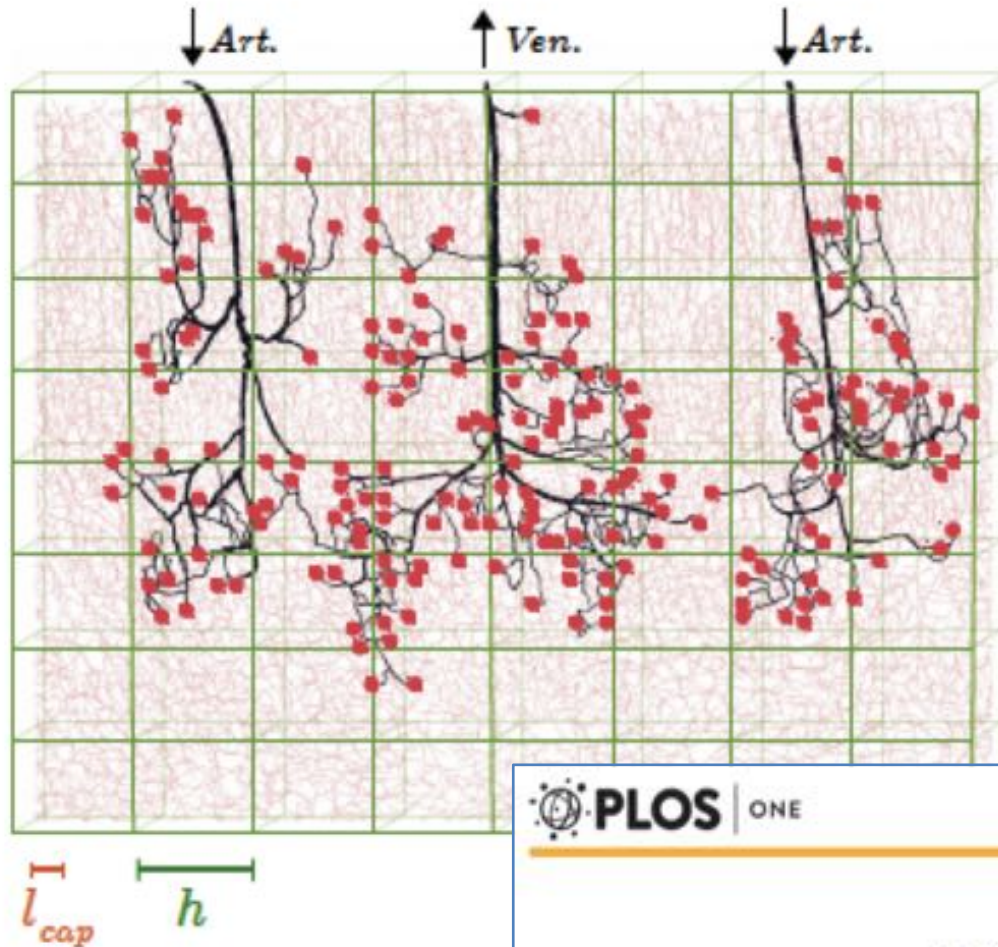


Smith et al. tbs



# Extension to larger scales

## Hybrid network/continuous approach for blood flow



- ❑ Network approach in the **arteriolar and venular trees**
- ❑ Replacing the **capillary bed** by a **continuum** discretized using FV method
- ❑ Developing a multiscale coupling model at the coupling points ●

 PLOS ONE

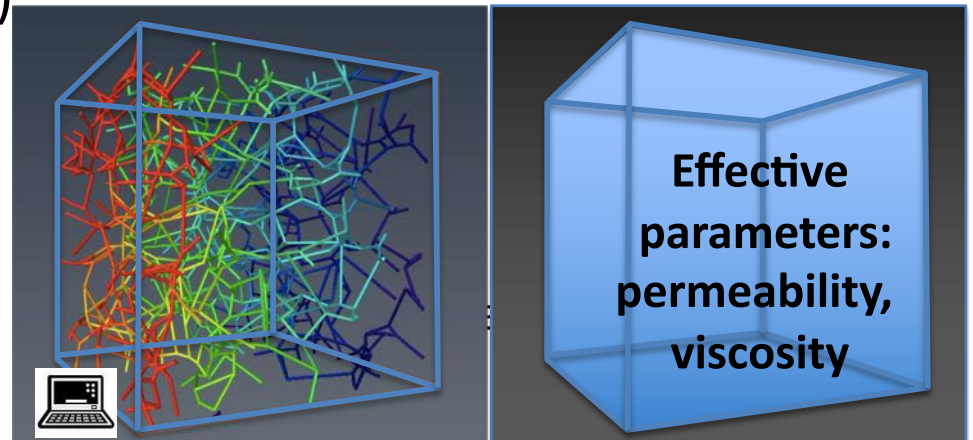
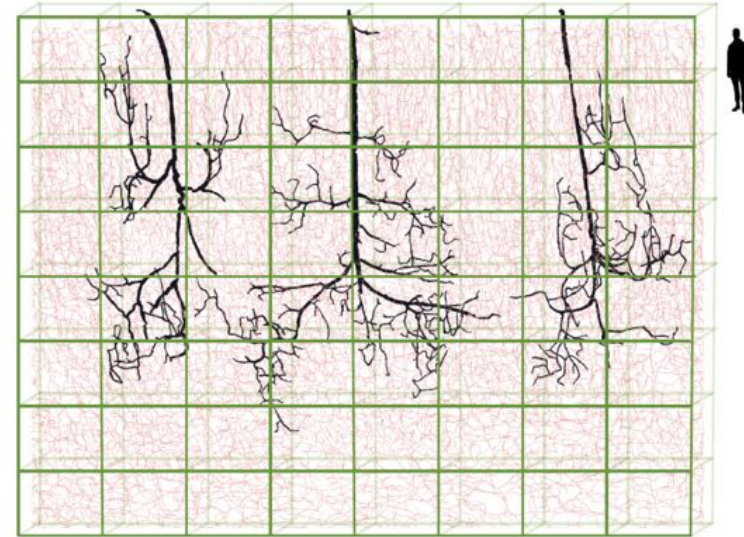
RESEARCH ARTICLE

Multiscale modelling of blood flow in cerebral microcirculation: Details at capillary scale control accuracy at the level of the cortex

# Blood flow at network scale

## Hybrid network/continuous approach

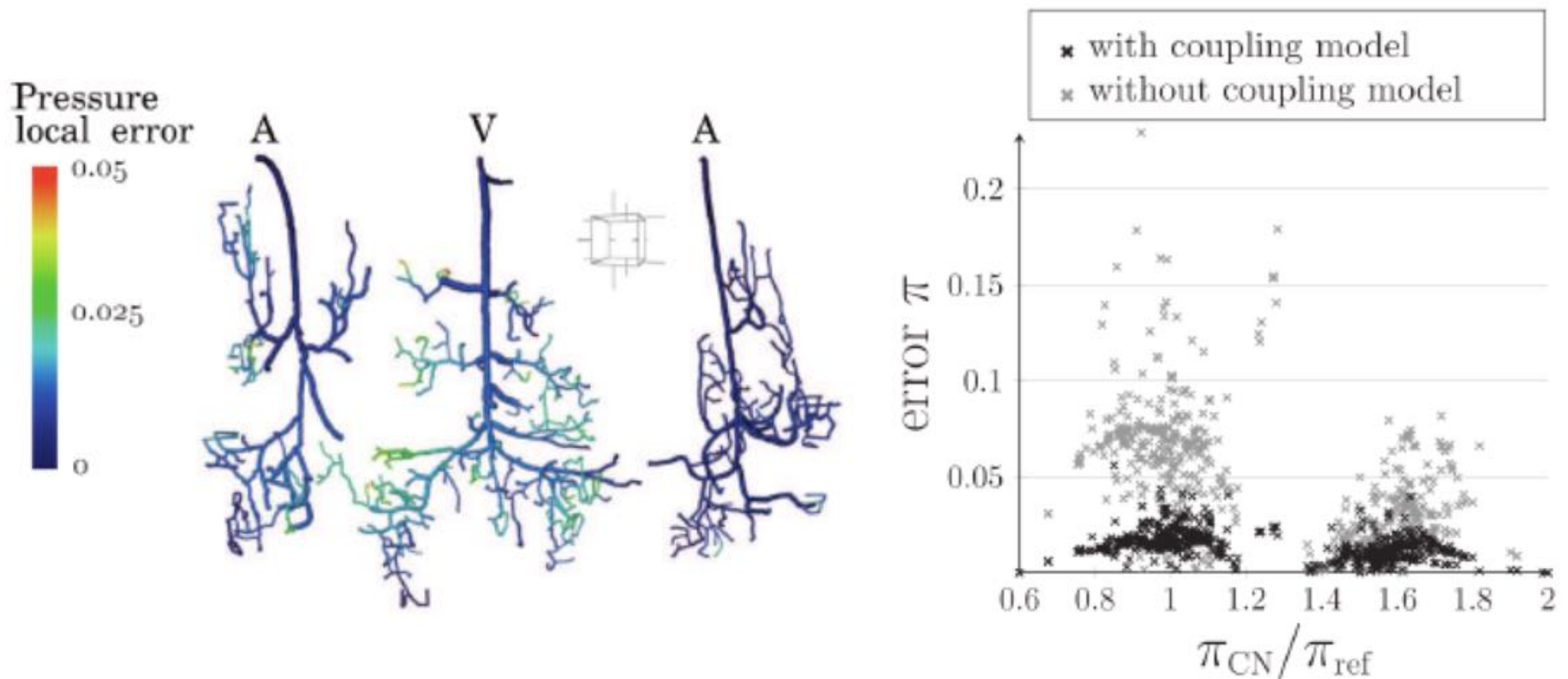
- Uniform hematocrit
- Art / Ven : Human data  
(Cassot et al. 2006)
- Capillaries :  
Synthetic networks
  - ➔ Equivalent continuum  
(Permeability, effective viscosity)
- Multi-scale coupling condition
- 750 000 vessel-equivalent
  - 1200 coarse meshes:  
Darcy equation (Finite Volume)
  - 600 Art / Ven:  
Linear network approach
  - 275 coupling points



# Blood flow at network scale

## Hybrid network/continuous approach

- Validation by comparison with network approach



- ➔ 1D / coarse 3D multi-scale coupling condition is are needed\*



# Blood flow at network scale

## Hybrid network/continuous approach

### □ Why a non-trivial coupling condition is needed ?

- Strong pressure gradients around coupling points
- Poor scale separation



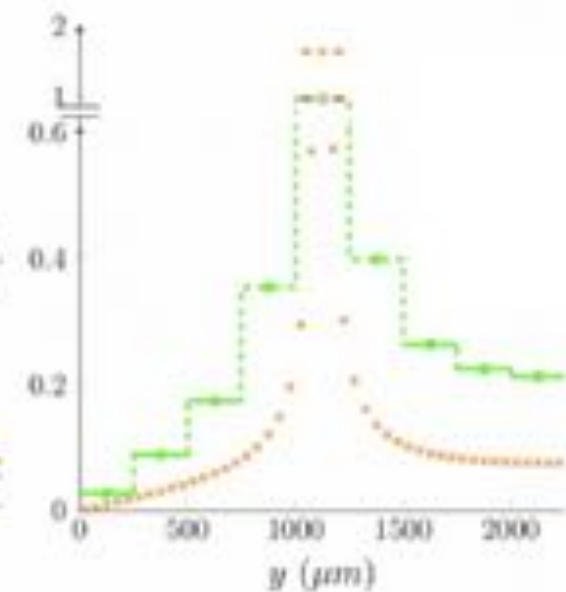
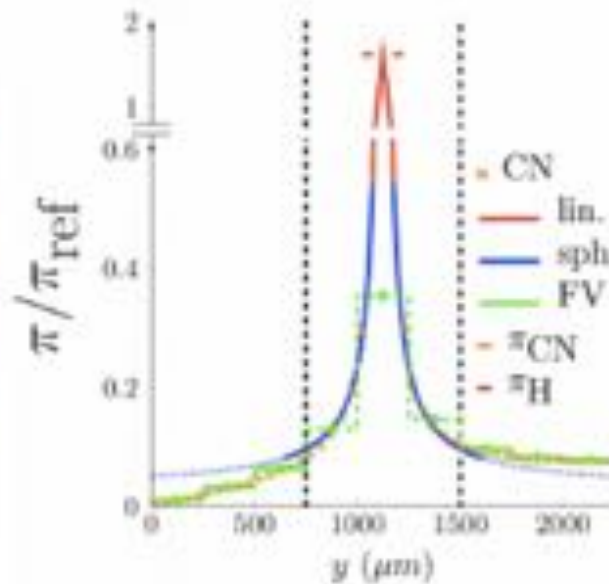
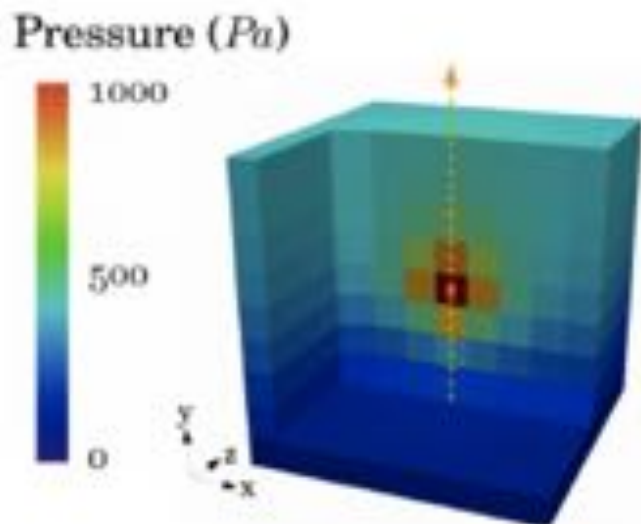
$$\pi_s = f(P_s^i, q_s)$$

vs.

$$\pi_s = P_s$$

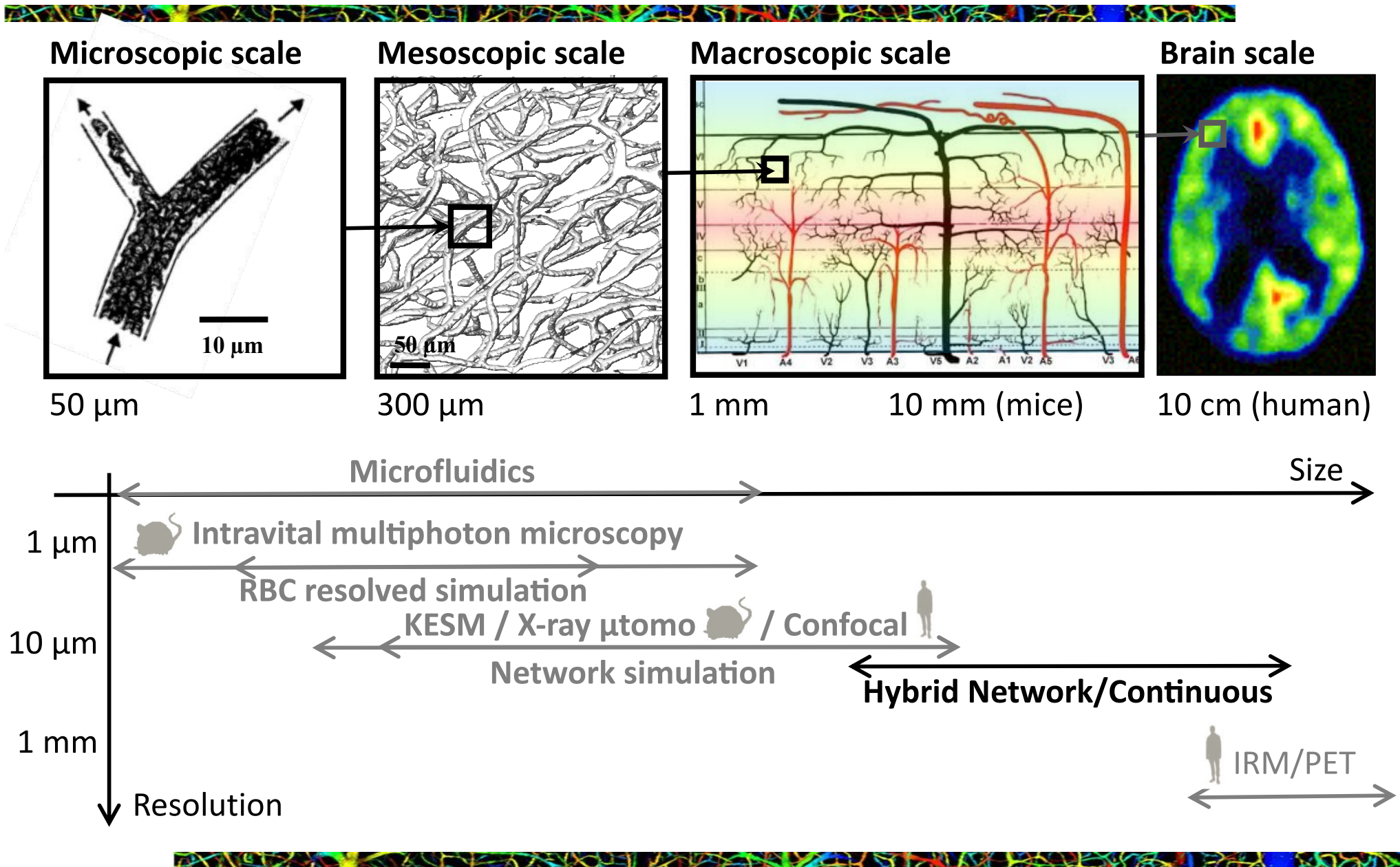
Multiscale Hybrid

Simple Hybrid



# Investigation tools and associated scales


## Numerical simulation



# Conclusions

## Focused on methods and didn't answer much questions



- **Tremendous progress of the experimental investigation tools...**
    - Structure, RBC velocity, pulsatility, glucose, oxygen, ...
  - **... has been driven by fundamental questions in neuroscience**
    - Neuro-vascular coupling
    - Interpretation of functional imaging
    - Role of microcirculation in brain disease
  - **This generated a huge amount of high resolution data...**
    - In various species, age and disease conditions at various scales
  - **... which need modeling for coherent understanding**
  - **There is a huge potential**
    - For understanding the brain, with perspectives in the clinics
    - For understanding fundamental questions in hemophysics
- 



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