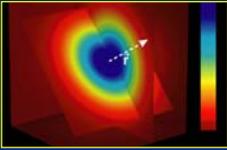
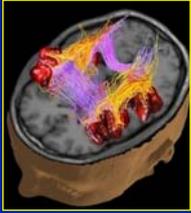


From Diffusion MR Image Analysis to Whole Brain Connectivity Simulation



Jean-Philippe Thiran

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EPFL - Lausanne

**– HPC in life sciences
at EPFL**

- *The Blue Brain project:*
 - create a biologically accurate, functional model of the brain using IBM's Blue Gene supercomputer
 - 8192 processors
- *Several other projects*
- *Including MR image analysis*





**Blue
Brain
Project**

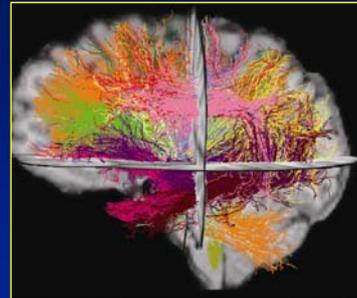



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Motivation

- **MR imaging** is now established as the most important modality in **neuro-imaging**
- Recent advances allow to obtain *in vivo* information on the **architecture** of the (brain) tissues
 - New MR sequences
 - New algorithmic developments
- This opens **new perspectives in fundamental neurosciences** as well as in clinical practice



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Outline

- **Basics of diffusion**
- **Imaging the diffusion by MRI**
- **From Diffusion MRI to Fiber Tracking**
- **Applications to Brain Connectivity Analysis**



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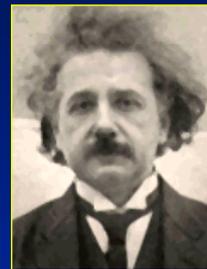


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Basics of diffusion

- Molecular diffusion – Brownian motion
 - First formally described in 1905 by A. Einstein
 - We will consider **water molecule** diffusion
 - *Example: in a glass of water, molecules diffuse randomly and freely, only constrained by the boundaries of the container*

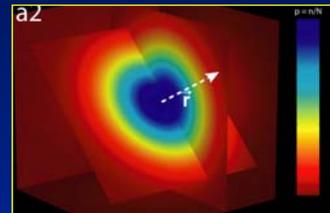


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Basics of diffusion

- For a **3D volume**,
 - 3D diffusion *pdf*
 - we use here the color-coding of the *pdf*
- For **homogeneous media**, this diffusion is an **isotropic Gaussian**
 - With $\sigma^2=2D\Delta$
 - **D** is the diffusion coefficient
 - Function of medium viscosity, t° , ...

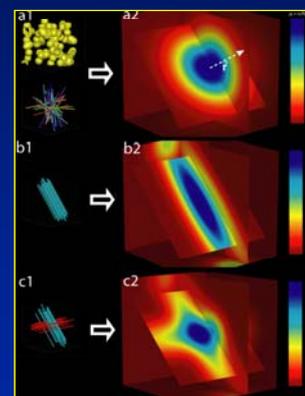
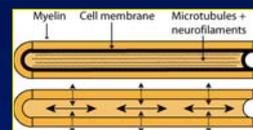


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Diffusion in complex media

- **Neuronal tissues: fibrillar structure**
 - Tightly packed, coherently aligned axons
 - Diffusion more restricted in direction perpendicular to the axonal orientation than along its parallel direction
 - This is called anisotropic diffusion, as opposed to isotropic diffusion
- fiber crossings:
 - Certainly not a Gaussian pdf then.



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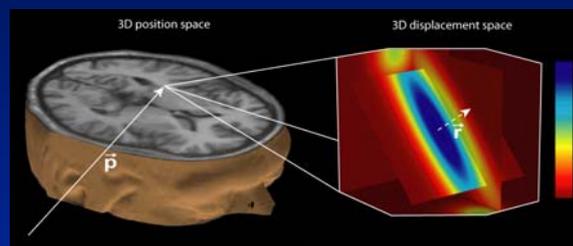


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Diffusion imaging

- Imaging the diffusion of the 3D volume would ideally give a 6D data set

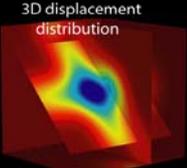
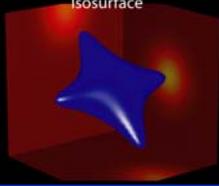


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Diffusion imaging

- But this information is too complex to be analyzed and visualized
- Simplified representations
 - Isosurfaces
 - Orientation Distribution Functions (ODF)
 - + color code for the orientation


Data reduction




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Diffusion imaging

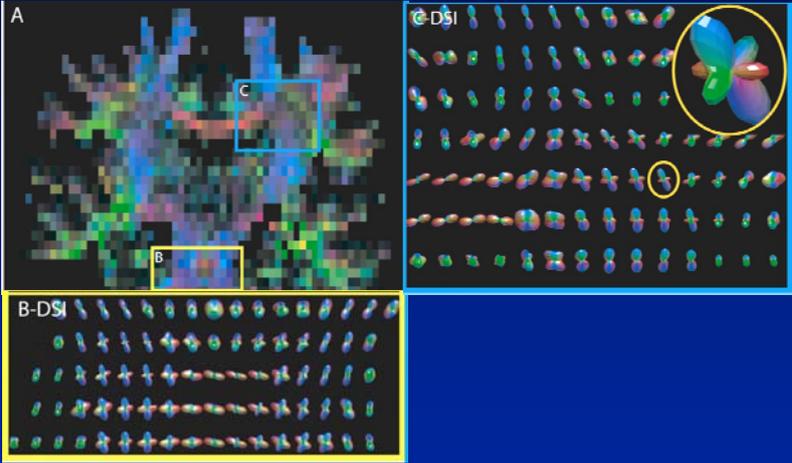





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Diffusion imaging



The figure displays diffusion imaging results. Panel A shows a 3D fiber tracking visualization of white matter tracts in a brain slice, with regions B and C highlighted by yellow and blue boxes respectively. Panel B-DSI shows a grid of diffusion tensor maps corresponding to region B. Panel C-DSI shows a grid of diffusion tensor maps corresponding to region C, with a magnified view of a single tensor map showing its principal components as colored lobes.

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Outline

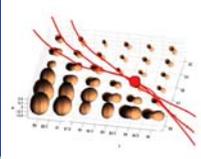
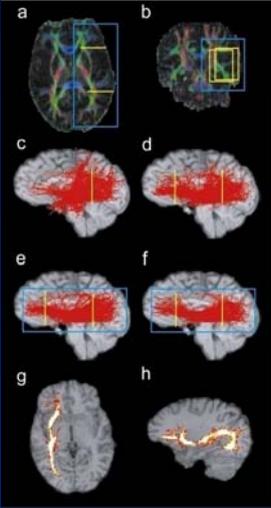
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 - Diffusion-Tensor MRI
- **From Diffusion MRI to Fiber Tracking**
- Applications to Brain Connectivity Analysis

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Tractography

- **Fibre tracking**
 - Infer axonal trajectories i.e. brain connectivity
 - Computation of trajectories following principal directions of diffusion
- **Whole brain simulation**
 - Trajectories are initiated all over the brain's WM
 - Result is an estimate of the whole brain connectivity (~100'000 lines)
- **Tract selection, virtual dissection**
 - Fibre selection using ROIs

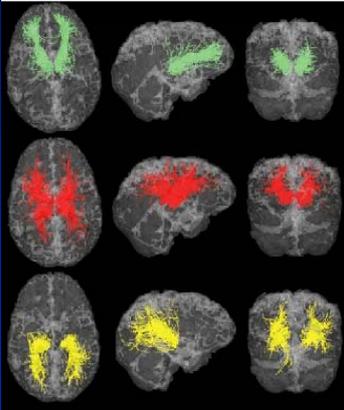
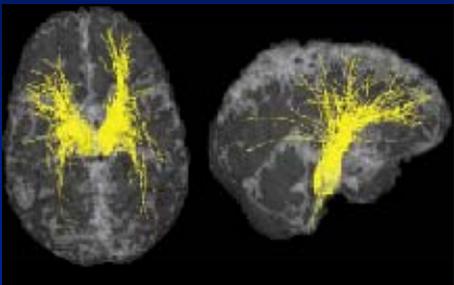



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Fibre-tracking, results

- Thalamic projections
- Cortico-spinal and cortico-bulbar tracts

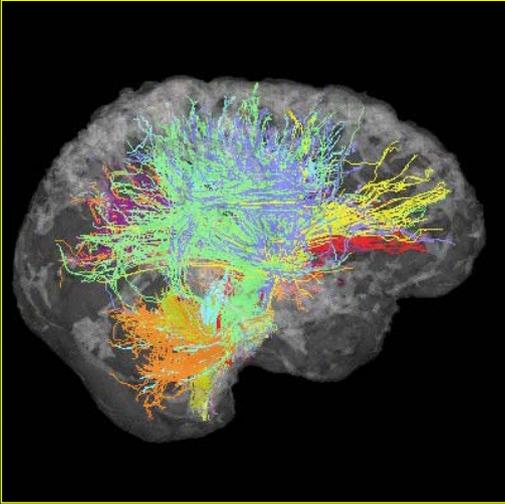






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Fibre-tracking, results

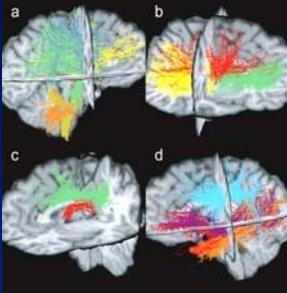




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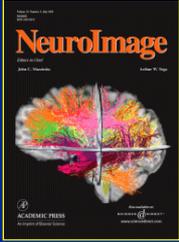


References





Hagmann et al. 2003





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Outline

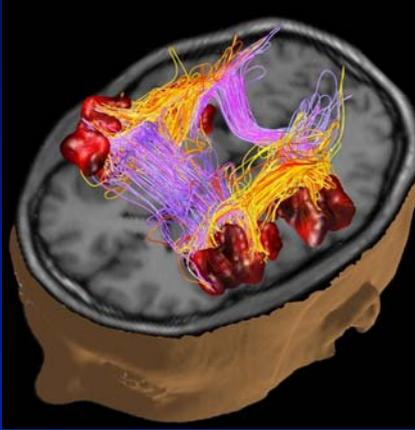
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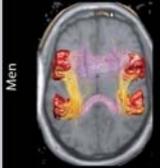

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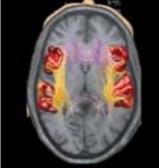
Application: study of language networks



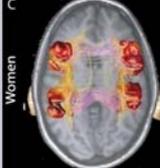
A Right Hander



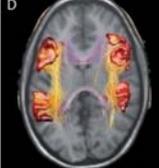
B Left Hander



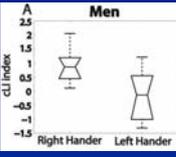
C Right Hander



D Left Hander

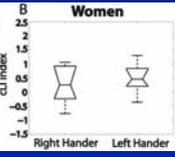


cLI index



Right Hander Left Hander

cLI index



Right Hander Left Hander

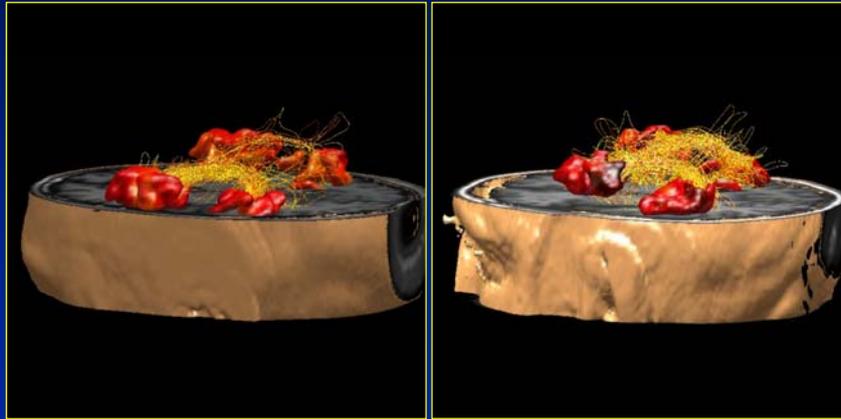



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Application: study of language networks

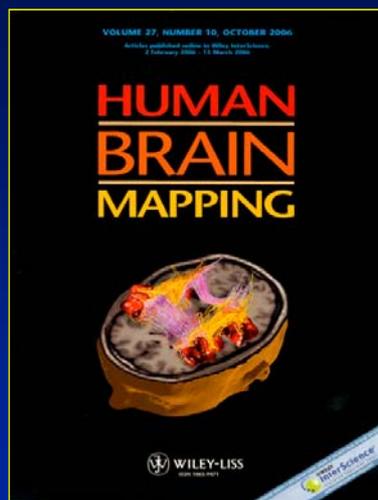
- Language networks in right and left handers



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Application: study of language networks



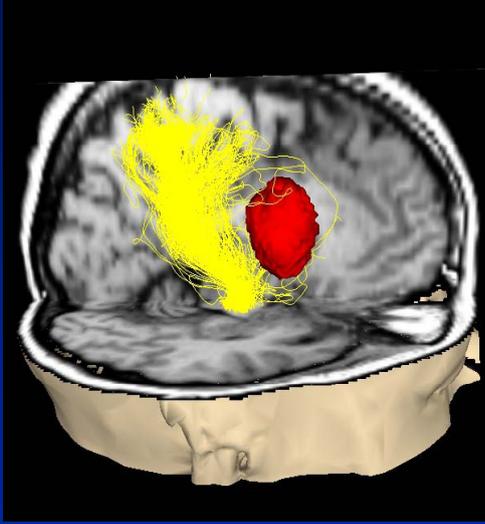
Hagmann et al. Human Brain Mapping, 2006



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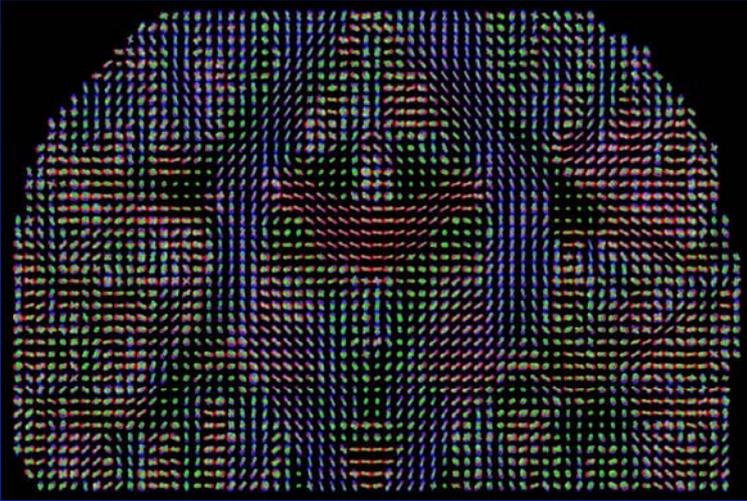
Application: diagnosis and therapy



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Fast-Marching Tractography (Fast-TraC)



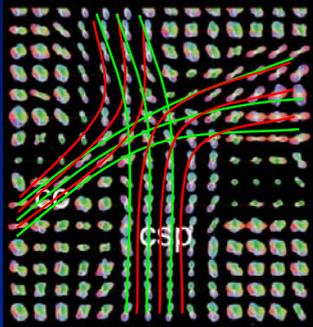
Orientation Distribution Function map (coronal section)

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Fast-Marching Tractography (Fast-TraC)

- **Motivation**
 - Classical tractography algorithms: not accurate in complex fiber crossings regions, because fibers follow locally the most linear trajectories.
- **Goal**
 - Design a tractography algorithm able to map complex fiber crossings in the brain WM using a Fast-Marching approach.



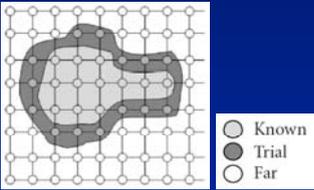


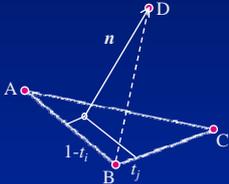

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Fast-Marching Tractography (Fast-TraC)

- **Front propagation**
 - 3D Anisotropic Fast Marching (shifted grid, 26 neighbors, Qingfen Lin 2003)







$$u_D = (1-t_i)u_A + (t_i-t_j)u_B + t_j u_C + \min\{|n \times s_1|, \dots, |n \times s_N|\}$$

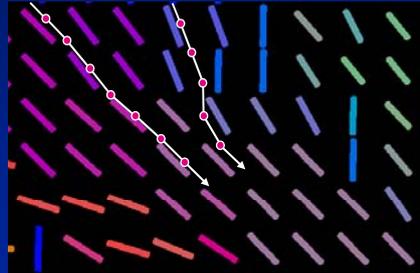



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Fast-Marching Tractography (Fast-TraC)

- Back propagation of the fibers
 - Gradient descent through cost map $u(\mathbf{r})$
 - Initialization points: one in each GM voxel

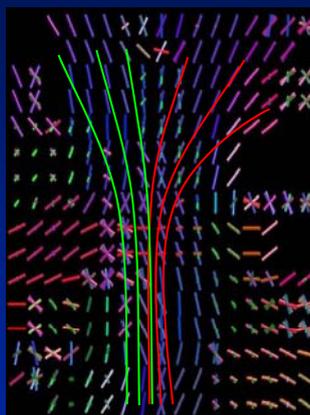


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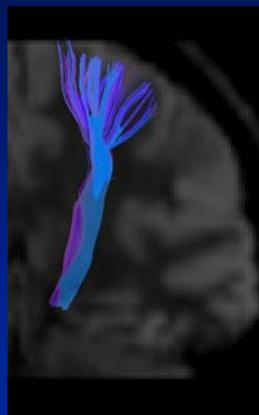


Fast-Marching Tractography (Fast-TraC)

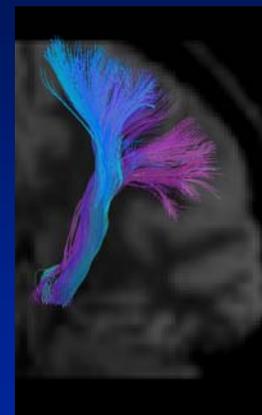
- Results: cortico-spinal tract



Local maxima
of the ODF map



Classical streamline
tractography (posterior view)

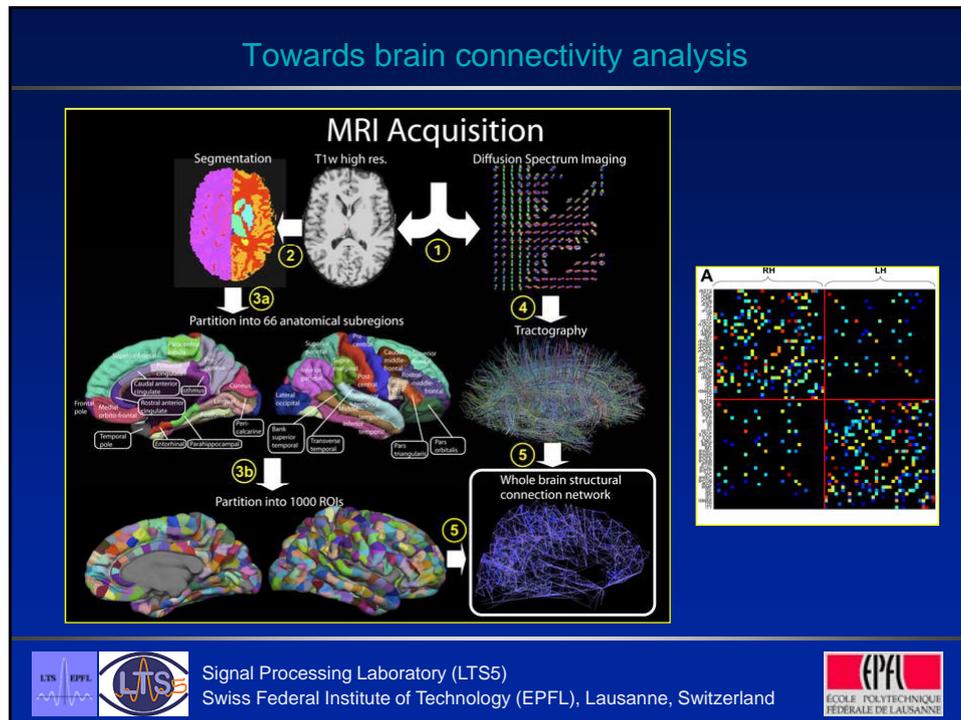


Fast-TraC
(posterior view)



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Computational complexity

- **Front propagation**
 - Volume of interest (WM): $\sim 450\text{cm}^3$
 - Resolution: $1 \times 1 \times 1 \text{ mm}^3$
 - => Number of voxels : $\sim 450'000$
 - In each of these voxels, we have to minimize u_D
 - Regions of Interest: ~ 1000
 - One front propagation per ROI
 - => 998 front propagations
- **Back propagation**
 - Computed trajectories: 1 – 20 millions

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Computational complexity

- **Computation time**
 - Approx. 20 hours (P4 3.8GHz, C++ code)
 - Each front propagation is independent, the algorithm is highly parallelizable
- **What would we gain with HPC ?**
 - Ideally, we should perform **one front propagation for every voxel** of the WM-GM interface
 - WM-GM interface: ~200'000 voxels
 - Estimated computation time without parallelization: 4000 hours!



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Fast-Marching Tractography (Fast-TraC)

- **Discussion**
 - New methodology for whole brain tractography, able to solve fiber crossings and branchings
 - Increases the global connectivity in the brain WM
 - Preliminary results obtained on the premotor cortex show a strong improvement of tractography results compared to classical streamline
 - Need for HPC to improve the resolution of the tractography



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Conclusions

- Diffusion MRI & adapted tractography are useful tools to study brain connectivity analysis *in vivo*
- **The *connectome*** gives both a visual and quantitative representation of the brain connectivity
 - Normal subjects
 - Pathologies
 - Schizophrenia
 - Epilepsy
 - ...
 - Brain development



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 - Dr Xavier Bresson
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 - Gilles Puy
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 - Prof. Reto Meuli
 - Dr Claudio Pollo
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(Martinos Center for Biomedical Imaging, Massachusetts General Hospital and Harvard Medical School)
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 - Siemens
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 - Lausanne-Geneva Center for Biomedical Imaging (CIBM)



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