



HUMAN
Connectome
PROJECT

“Diffuse”

*A toolbox for diffusion data processing
in BrainVISA*

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RMN – 21/12/17 – CERIMED



BrainVISA - Diffuse

- Introduction: dMRI data processing
 - Pre-processing
 - Post-processing: local models and tractography
- *Diffuse* toolbox
 - BrainVISA environment
 - Workflow
- Influence of pre-processing on post-processing analyses
 - Acquisition/Correction quality
 - Impact on co-registration
 - Impact on signal (diffusivity metrics)
- Outcome
 - Acquisition strategy
 - Centre IRM

BrainVISA - Diffuse

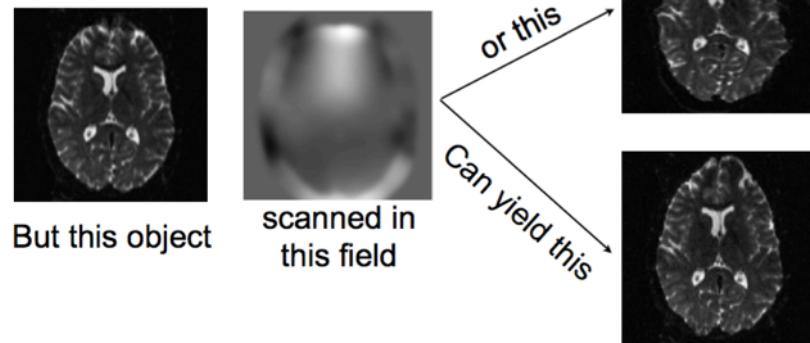
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Geometric distortions

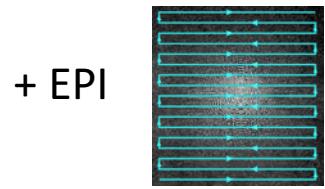
"off-resonance field"

Two origins

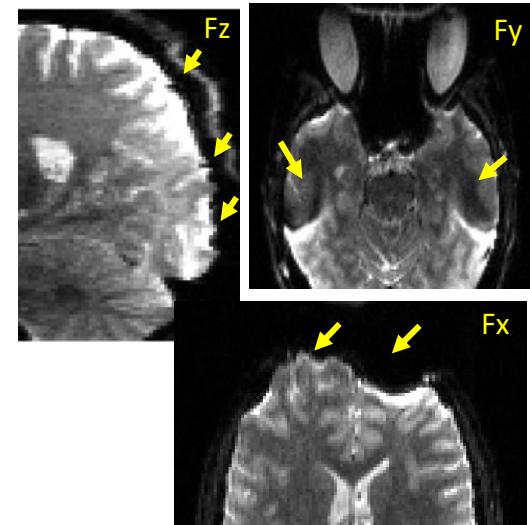
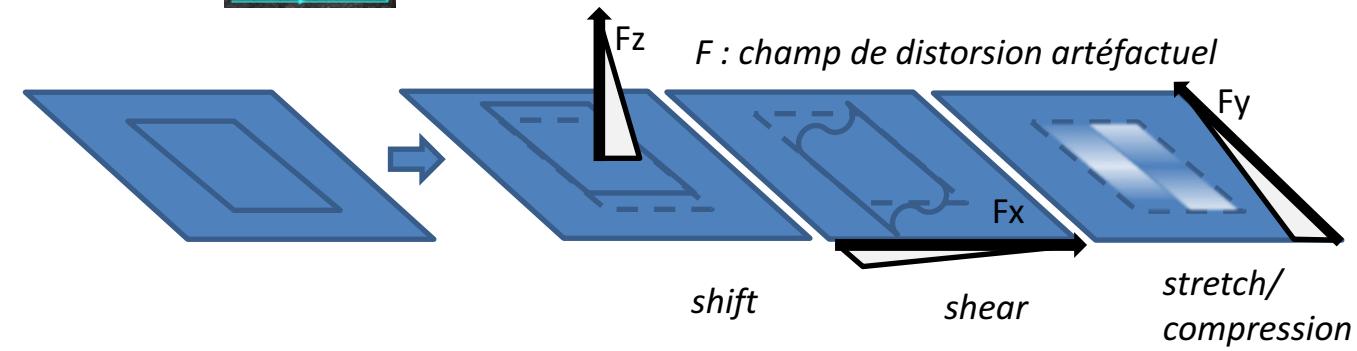
- **Susceptibility** differences between tissues
- Diffusion gradients: **Eddy currents** induced in gradient coils



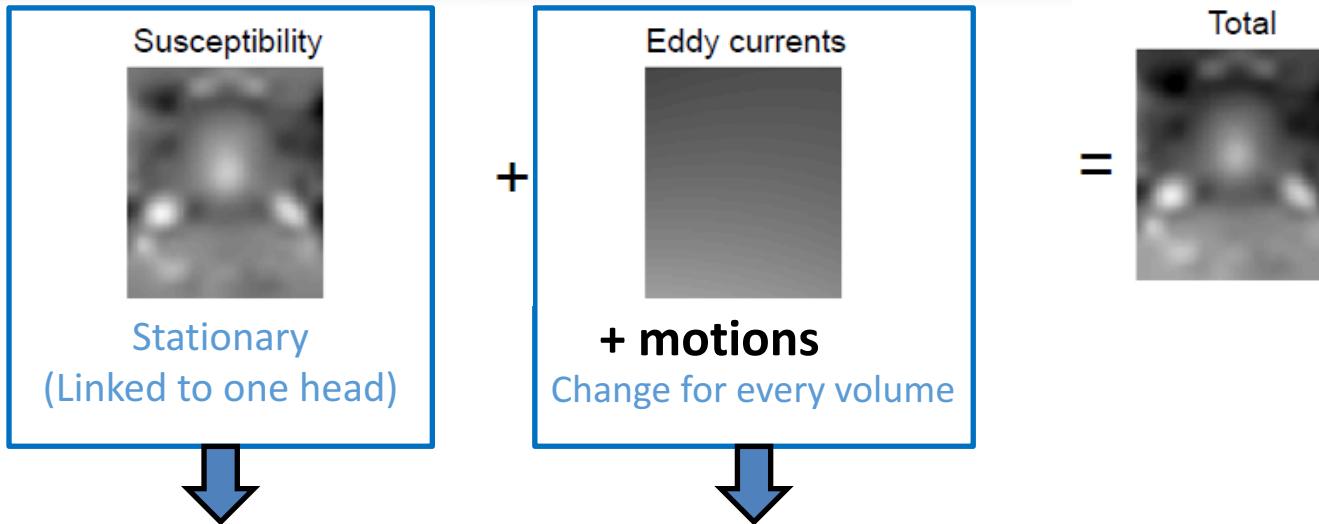
Andersson, HCP course, Boston 2016



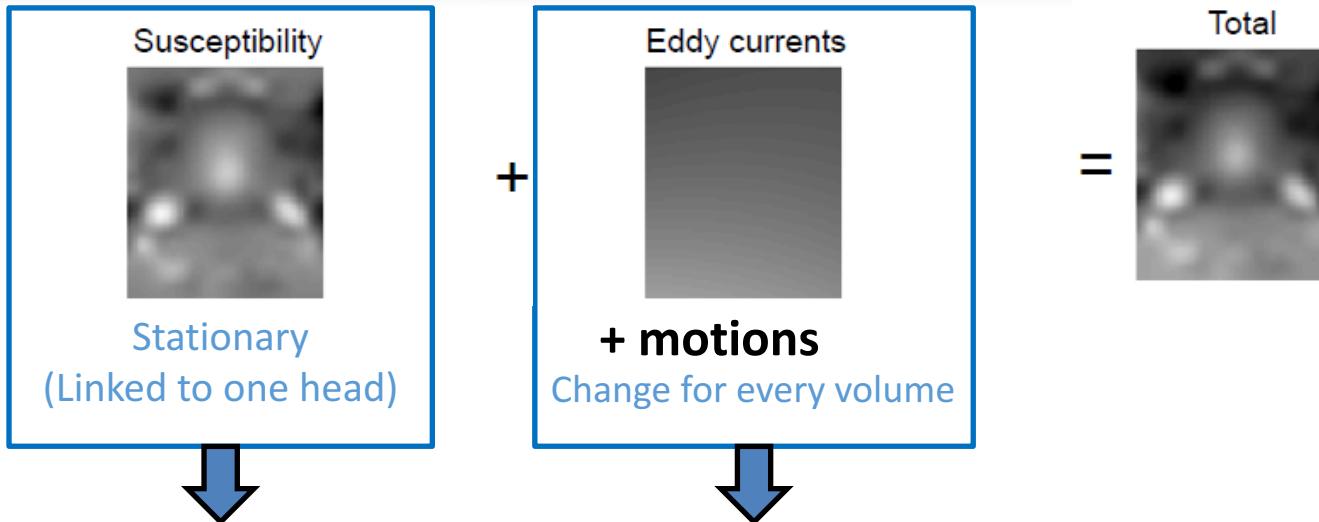
= Distortions in the phase encoding direction (low Bdw)



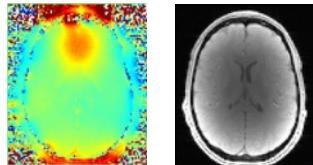
Geometric distortions



Geometric distortions

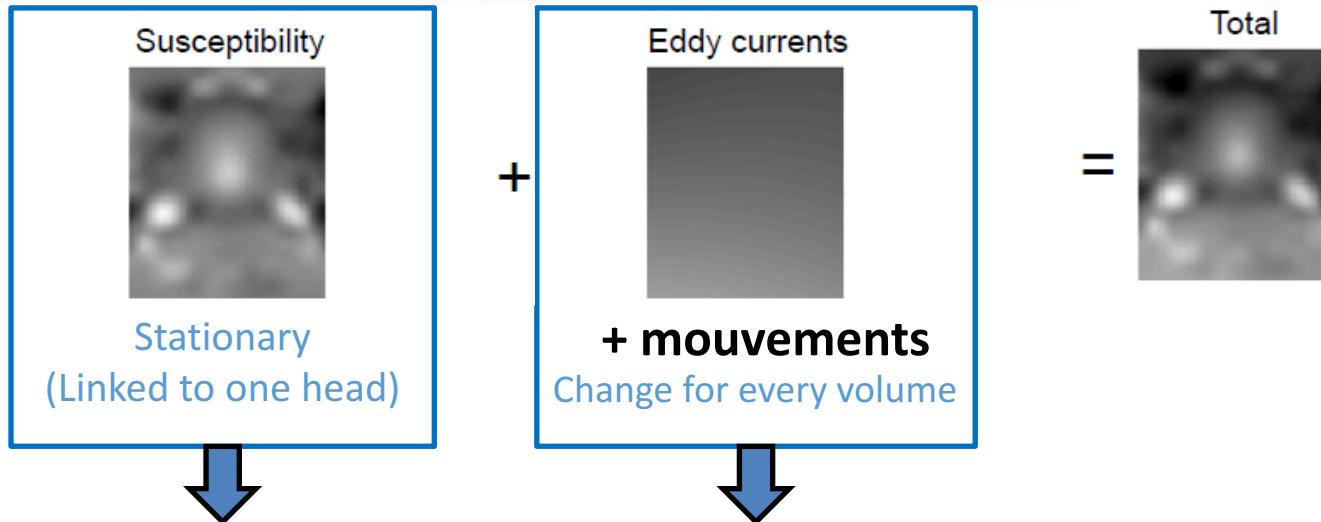


- Mesure : fieldmap

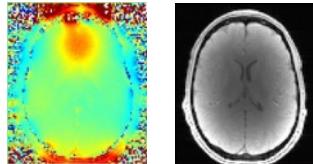


- Mesure : fieldmap
- Linear estimation: affine registration of volumes
- Estimation : *FSL-topup*
- Non-Linear estimation: *FSL-eddy*

Geometric distortions

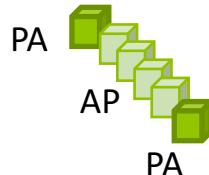


- Mesure : fieldmap
- Linear estimation: affine registration of volumes

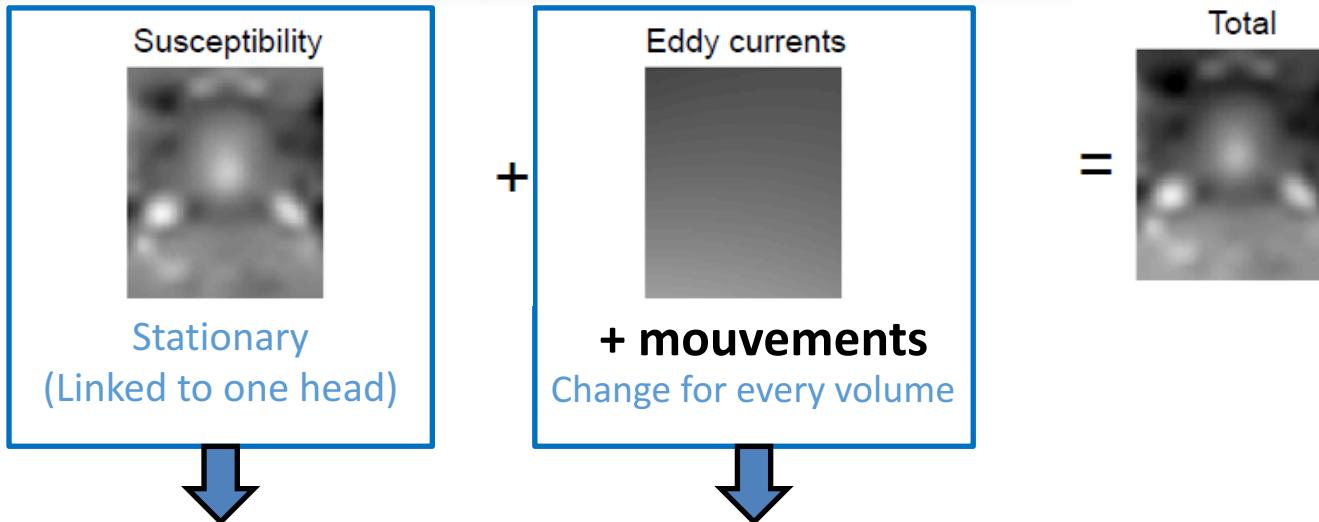


- Estimation : *FSL-topup*
- Non-Linear estimation: *FSL-eddy*

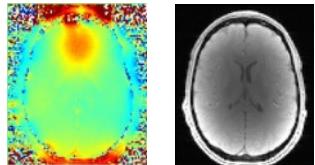
$b=0$
volumes



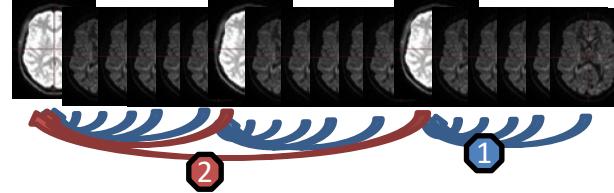
Geometric distortions



- Mesure : fieldmap

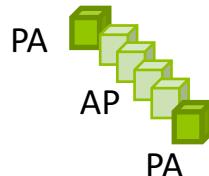


- Linear estimation: affine registration of volumes



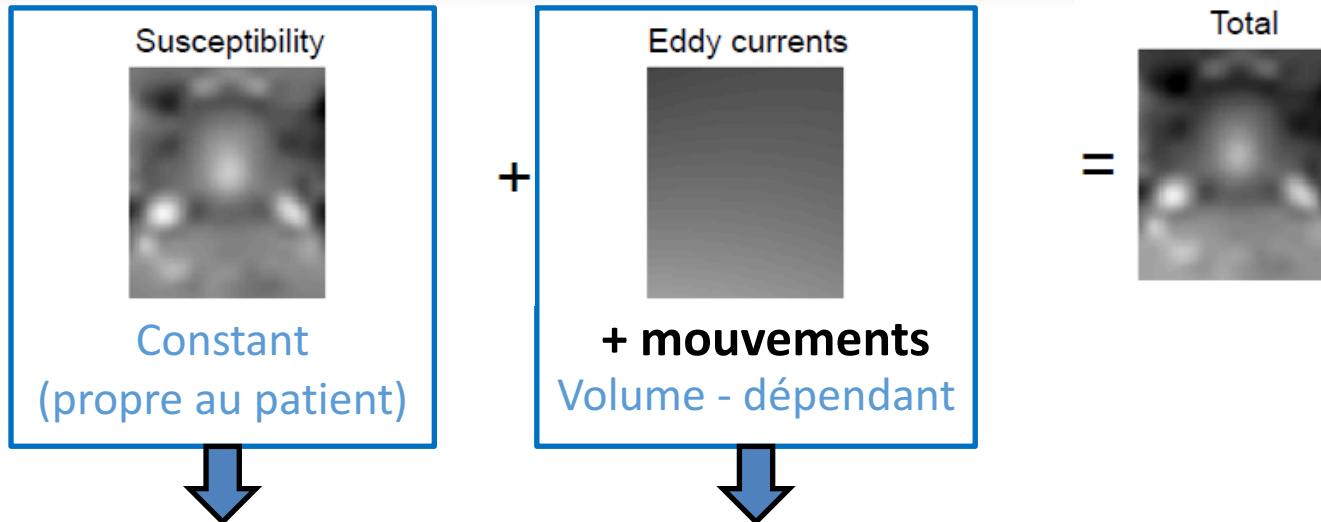
- Estimation : *FSL-topup*

$b=0$
volumes

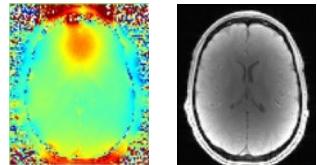


- Non-Linear estimation: *FSL-eddy*

Geometric distortions

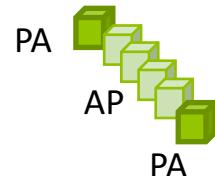


- Mesure : fieldmap

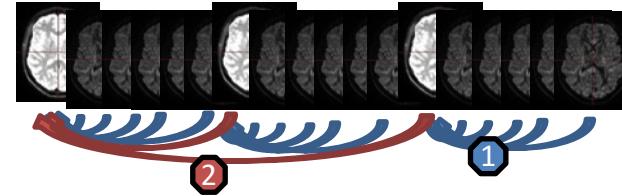


- Estimation : *FSL-topup*

Reversed
 $b=0$ volumes



- Linear estimation: affine registration of volumes



- Non-Linear estimation: *FSL-eddy*

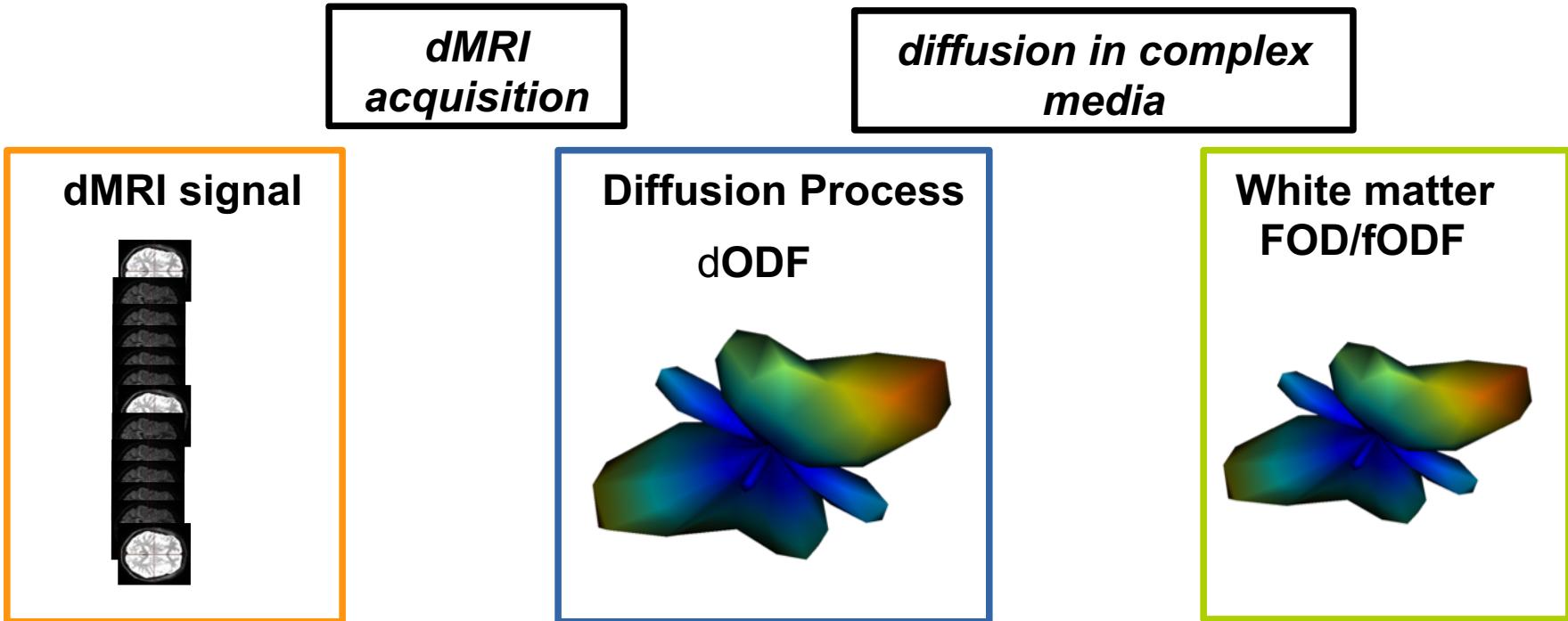
++ high gradients
 $N > 60$ & sampling
on the sphere



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Modeling dMRI signal at voxel level



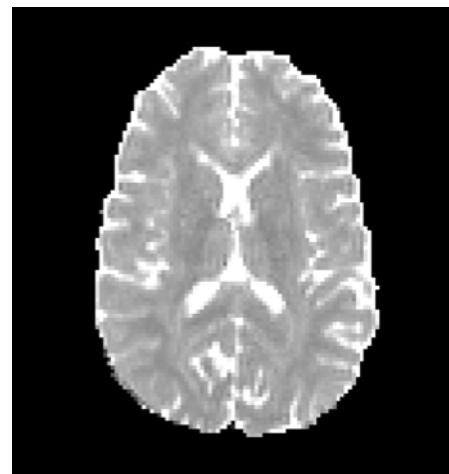
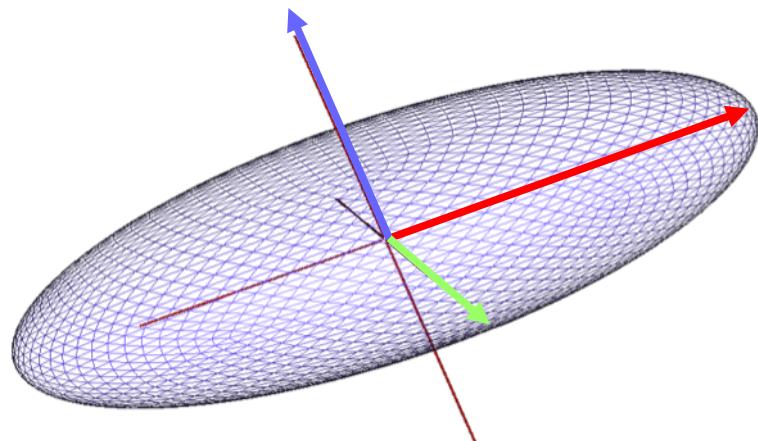
ODF: Orientation Distribution Function. Angular distribution of quantity of interest (on a sphere for diffusion).

dODF: diffusion ODF

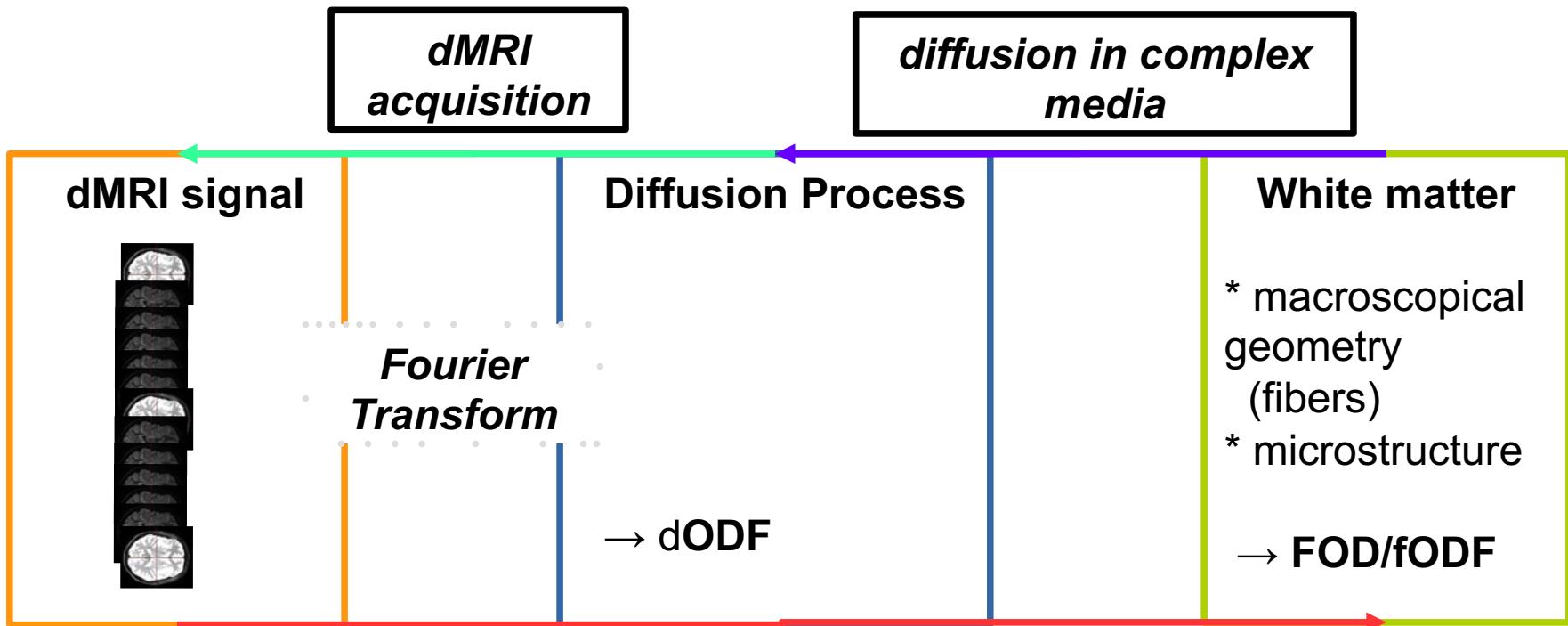
FOD: Fiber Orientation Distribution = fiber ODF

Diffusion Tensor model

- Signal in the voxel is assumed to derive from 3D non degenerated gaussian

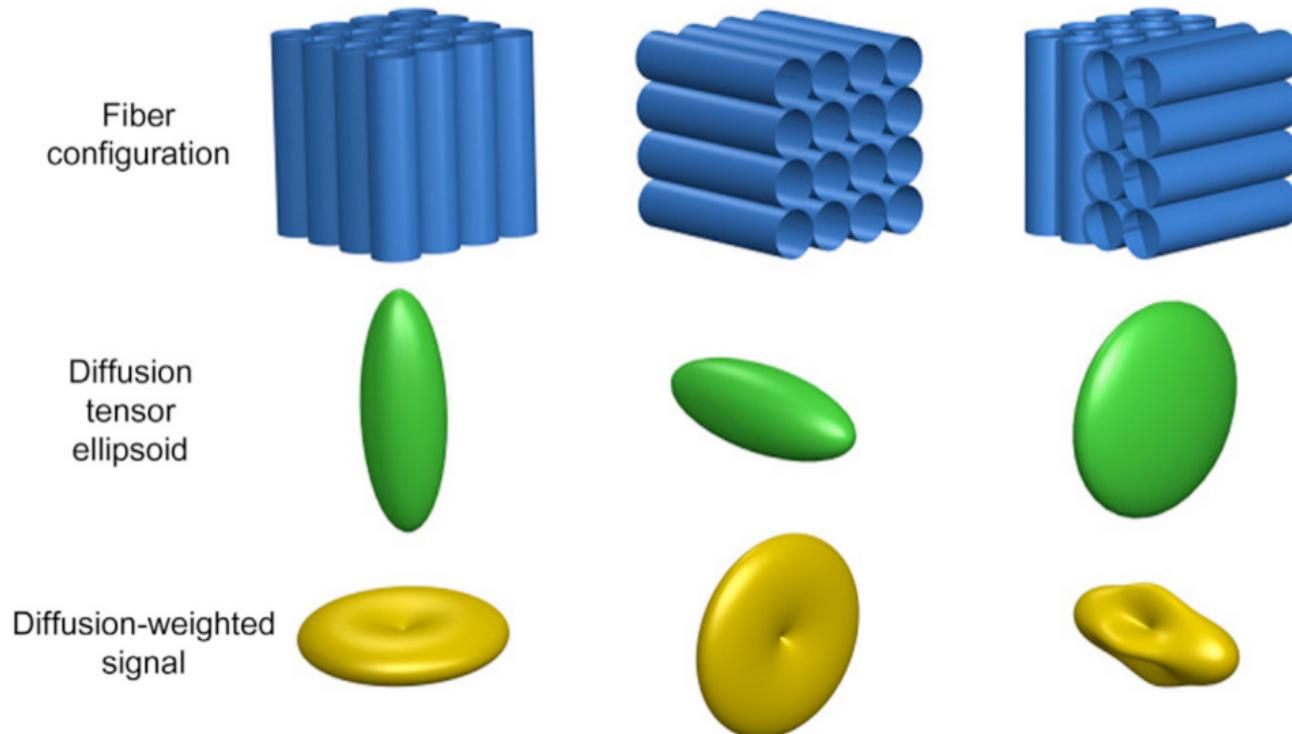


Modeling dMRI signal at voxel level



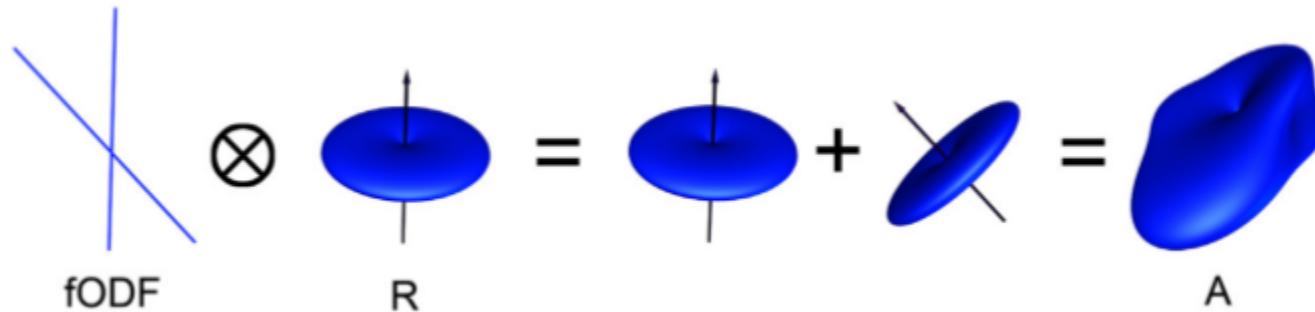
Modeling dMRI signal at voxel level

- Signal in the voxel is a mixture of fiber with identical signal (impulsionnal response)



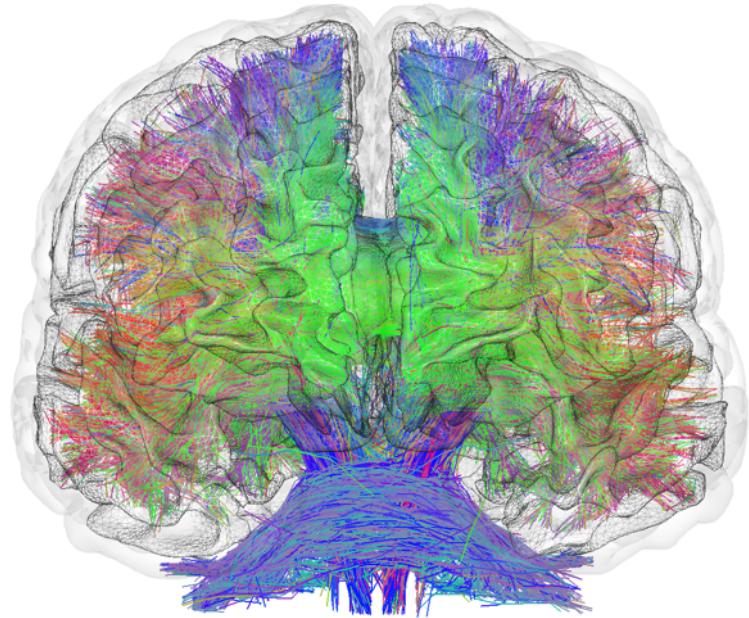
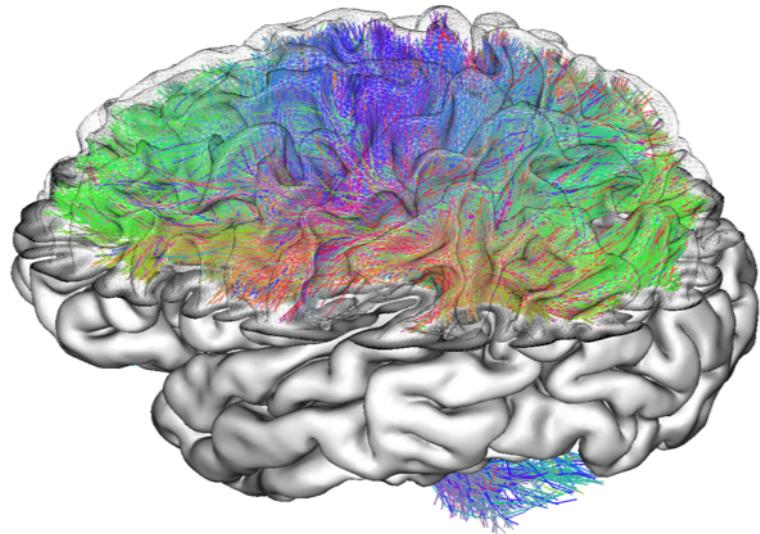
Constrained Spherical Deconvolution

- Signal in the voxel is a mixture of fiber with identical signal (impulsionnal response)

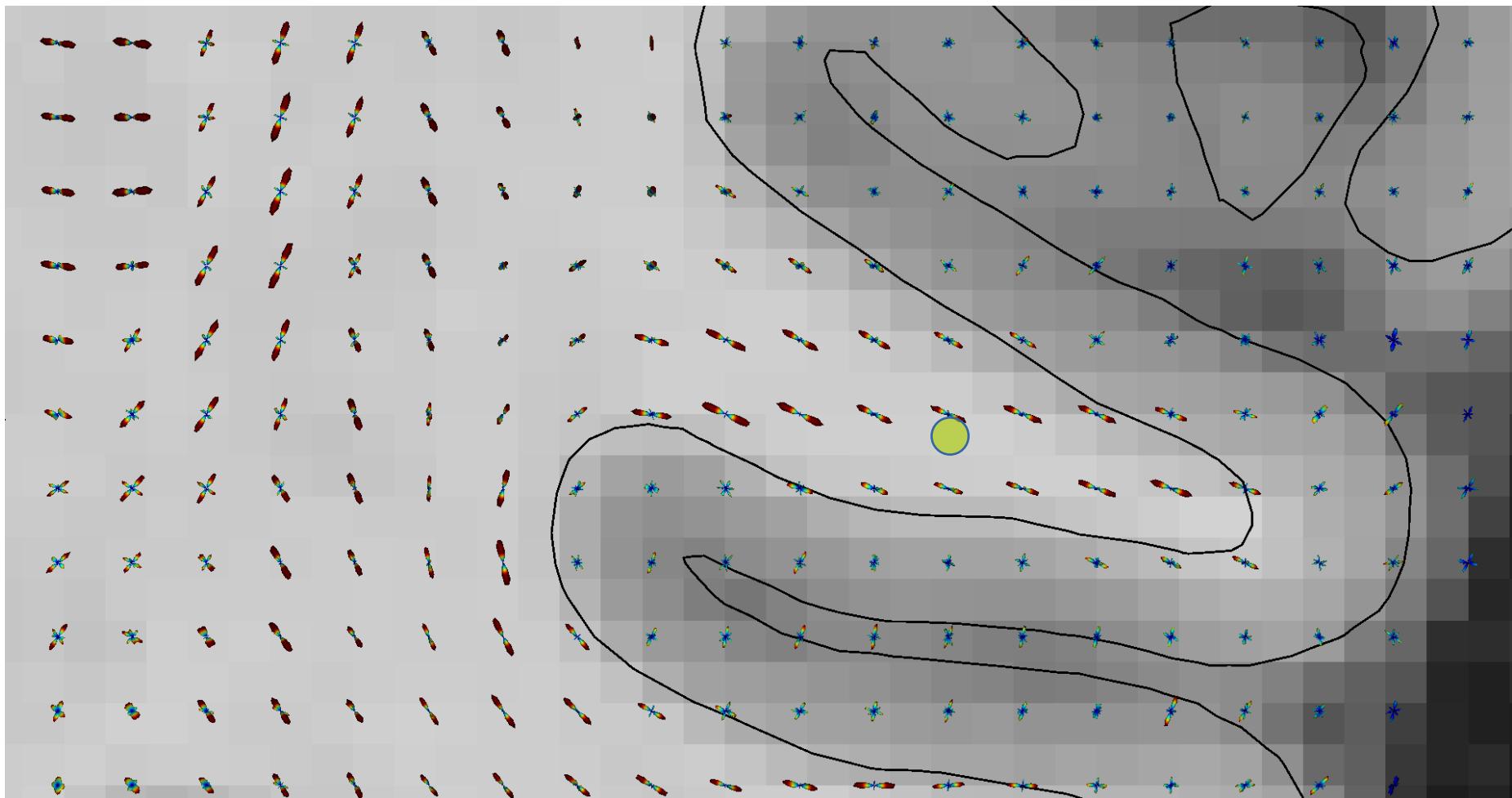


- Impulsionnal response is estimated from « single fiber voxels » [Tax, Tournier]
- Spherical deconvolution in the Fourier domain is an ill posed problem and must be regularized [Tournier, Cheng] to avoid negative signal values, e.g. regularization sphere

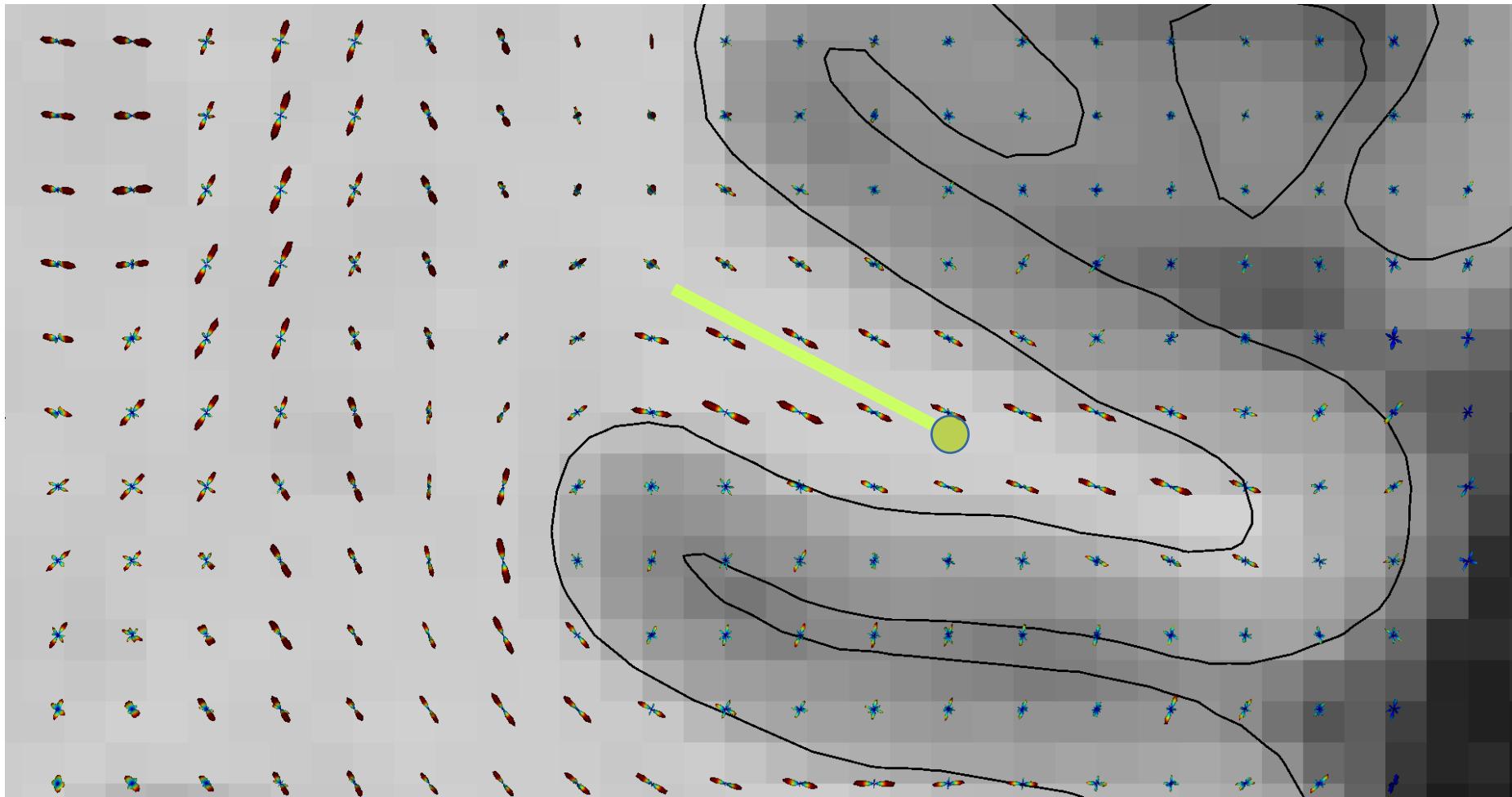
Tractography : from local to global



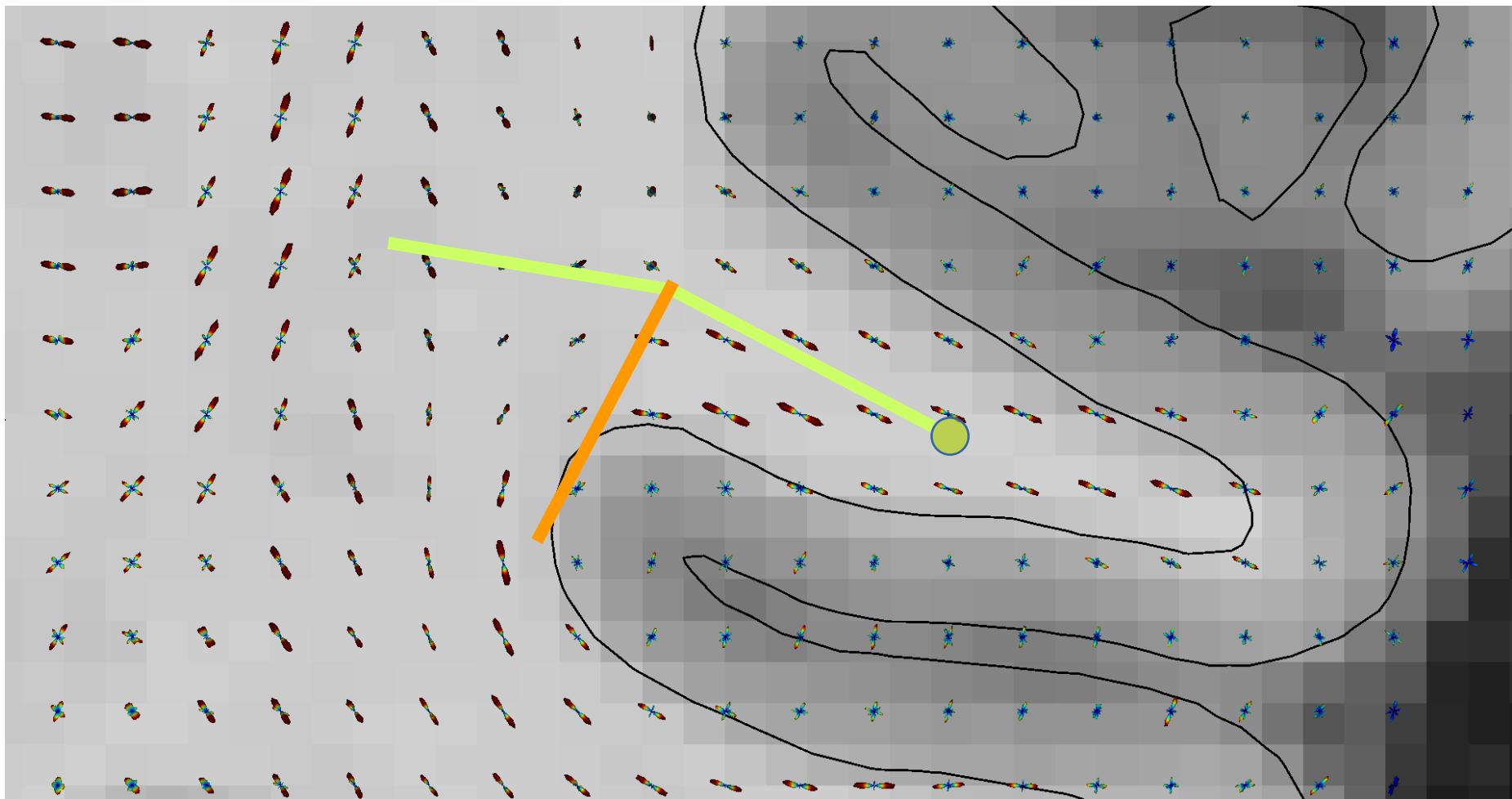
Tractography : principle



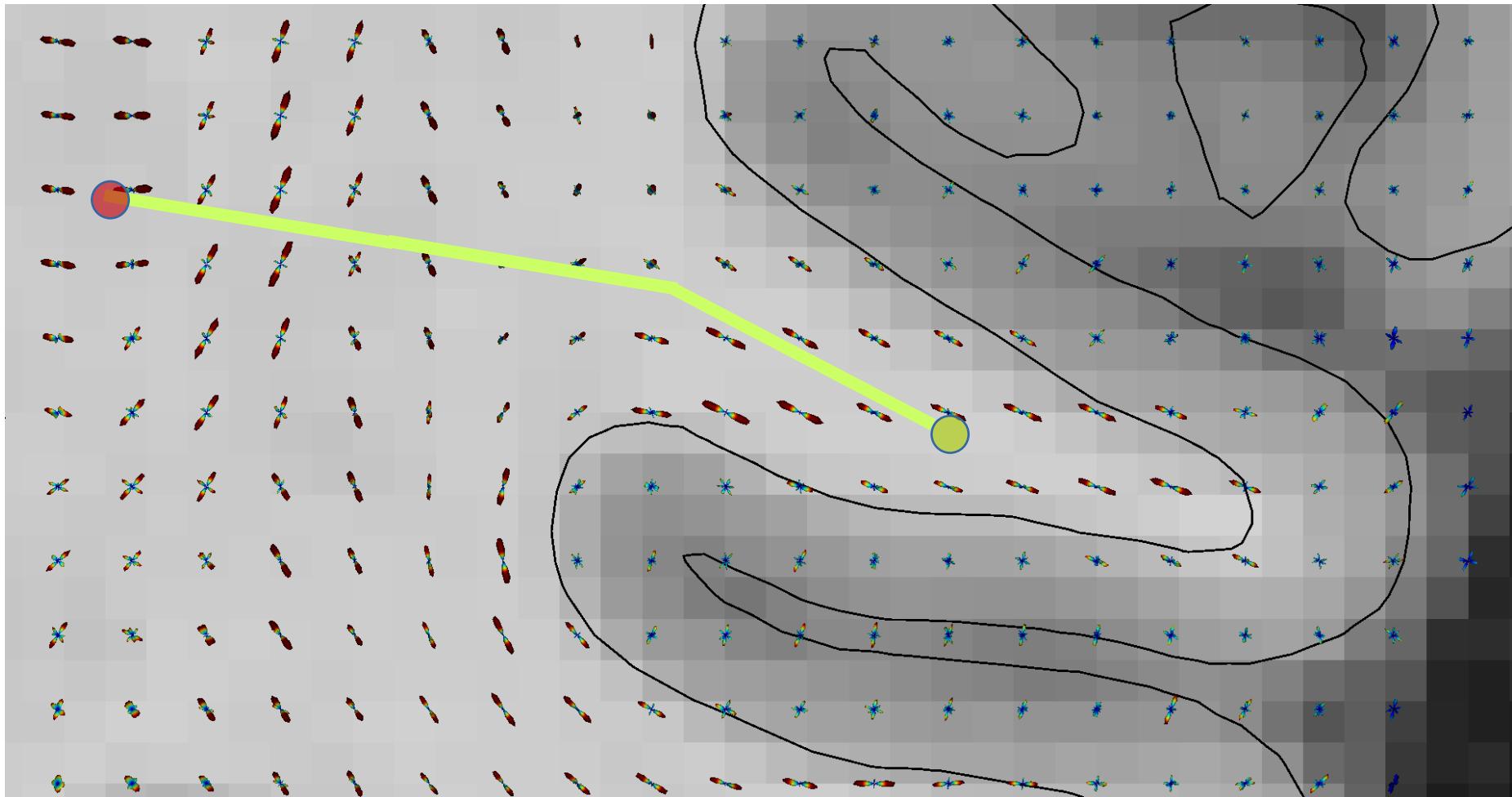
Tractography : principle



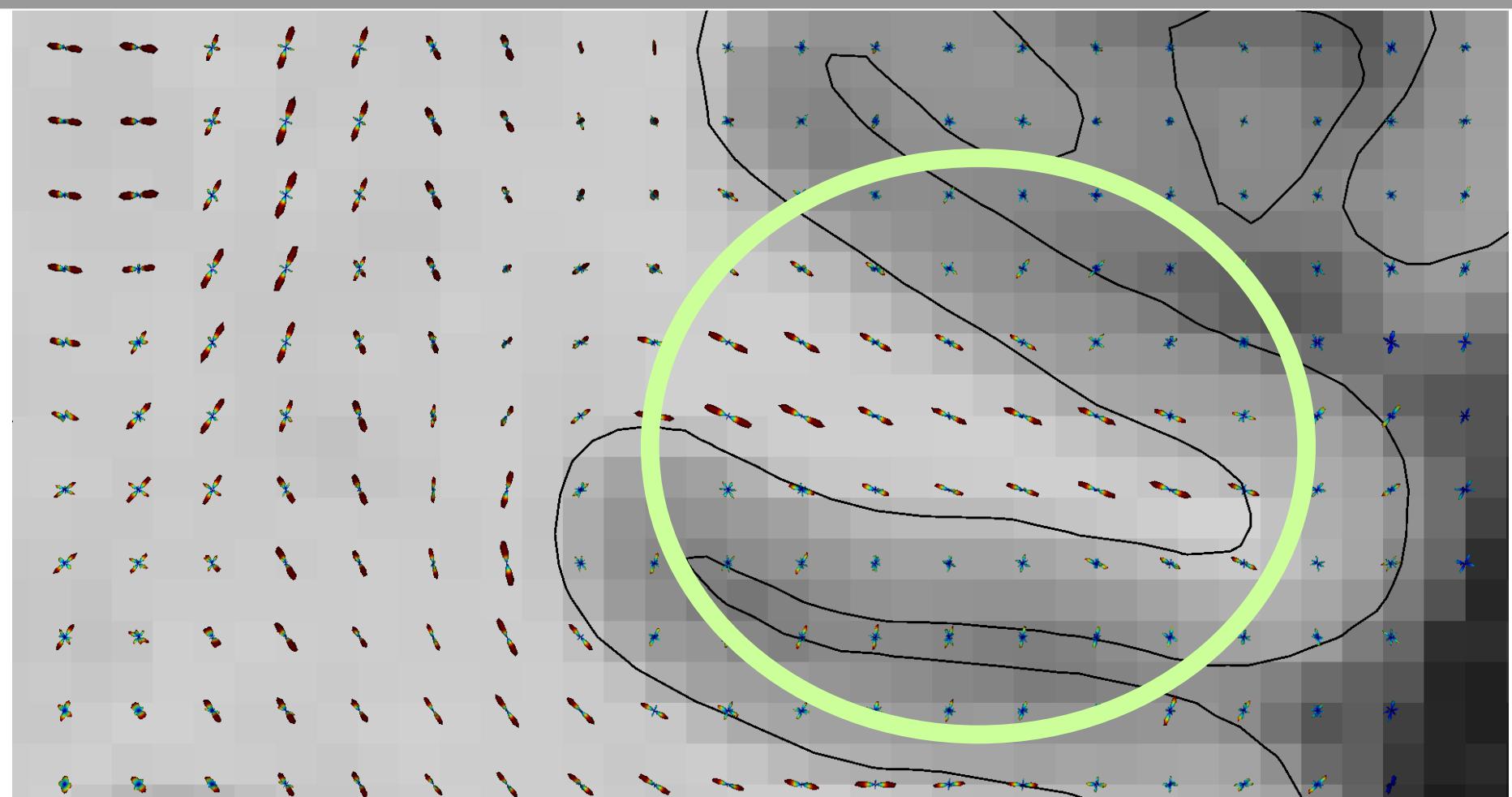
Tractography : principle



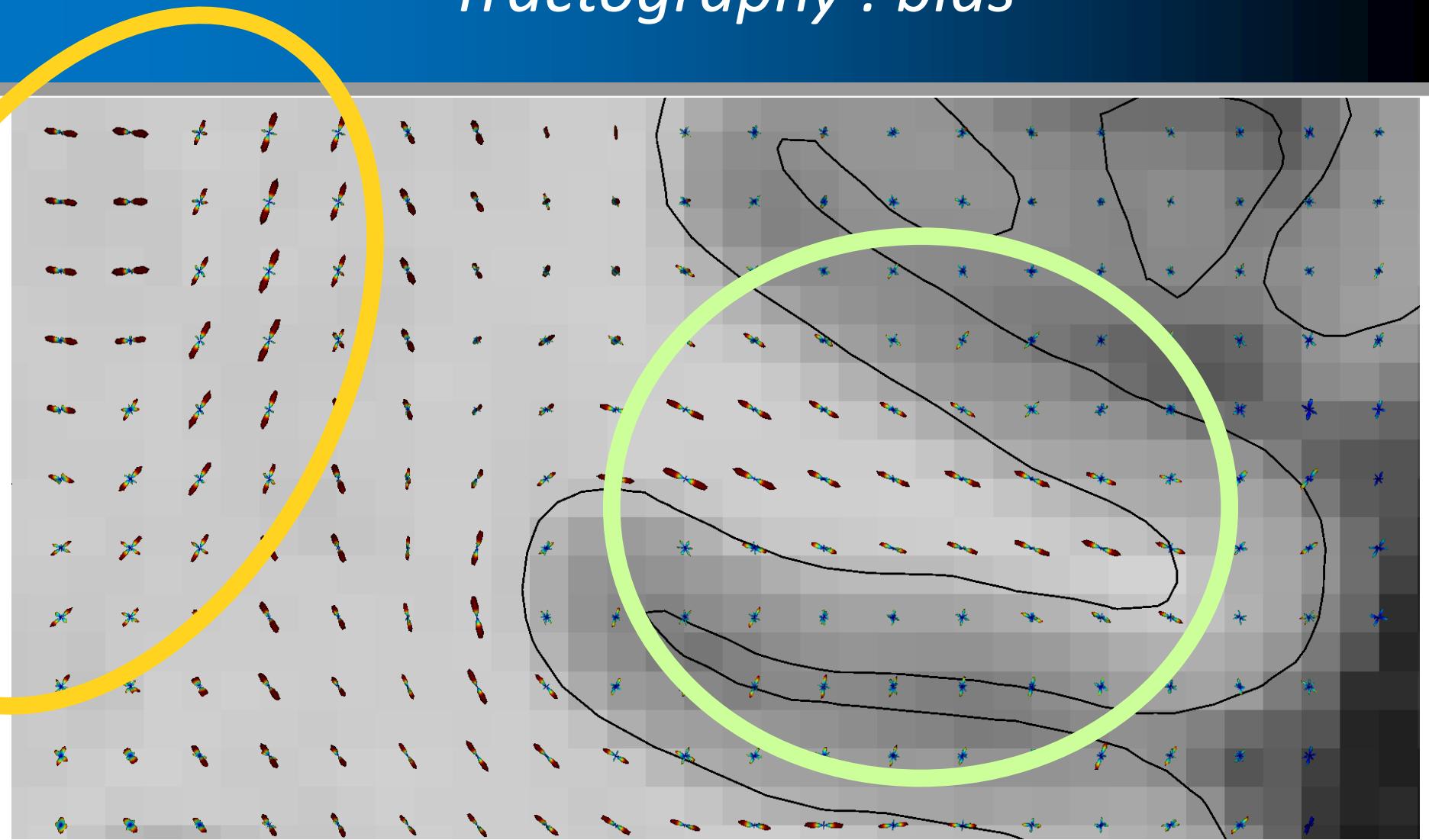
Tractography : principle



Tractography : principle



Tractography : bias



Tractography : from local to global

- Aim : Estimate macroscopic fiber bundle geometry by integrating local diffusion information
- What is needed ?
 - start point i.e. **seed** (surfacic / volumic strategy ; random /deterministic position)
 - end point (ending criteria)
 - integration method (probabilistic, deterministic, first order, second order and beyond)
 - integration step
 - interpolation method (for ODF) (trilinear in the spherical harmonics basis)
 - a priori:
 - geometrical constraints e.g on curvature [ref Descoteaux , Jones]
 - anatomical constraints e.g no streamline ending into cerebro-spinal fluid [ref ACT]
 - microstructural constraints ? e.g. axonal radius. [ref MIT]

Tractography : from local to global

- Known bias: a short list. See [Ref Jones, Descoteaux]
- Gyral bias: Due to sharp angle and partial volume effect, streamlines end easily near gyral crests.
- Seeding bias : 1 seed = 1 streamline thus longer streamlines are over-represented.
- Growing litterature on the topic, solutions are set up by methodologists :
- Gyral bias → priors on local fiber orientation [Achille Teillac, SET]
- Seeding bias → Seeding from grey/white interface, post-tractography filtering (SIFT, SIFT2, COMMIT)
- Complex configuration → global tractography with dynamic priors [Mangin]

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Diffuse Toolbox

- Toolbox for processing Diffusion MRI

- Guided and adapted for each acquisition type
- Automated
- Allowing interface with anatomical data



<http://brainvisa.fr/web/index.html>

BrainVISA 4.5.1 (on frioul03)

BrainVISA Support View

Toolboxes

- BabySeg
- Diffusion and Tracking
- Cortical Surface
- Data Management
- DataMind
- Diffuse**
- Freesurfer
- Morphologist
- Structural analysis
- TMS
- Tools
- Viewers

Diffuse

- Constrained Spherical Deconvolution Reconstruction
- Diffusion Preprocessing Pipeline
- Diffusion Tensor Imaging (DTI) Reconstruction
- Local Tractography Pipeline
- + distortion correction
- + import
- + local modeling
- + registration
- + tools
- + tractography
- + viewers

Documentation

Search site:

Diffuse toolbox

Presentation

Diffuse regroups processes to handle properly and in a user-friendly way the processing of diffusion weighted MRI (DWI) scans. Diffuse is developed by the [Methods and Computational Anatomy \(MeCA\)](#) research group of Institut de Neurosciences de la Timone (INT).

This toolbox mostly relies on the [FSL](#) algorithms and on [Dipy](#) library.

The Diffuse toolbox organization

Diffuse Toolbox

- Toolbox for processing Diffusion MRI
 - Guided and adapted for each acquisition type
 - Automated
 - Allowing interface with anatomical data
- Pre-processing : assembly of FSL tools in pipelines
- Post-processing : integration of Dipy functions



<http://brainvisa.fr/web/index.html>



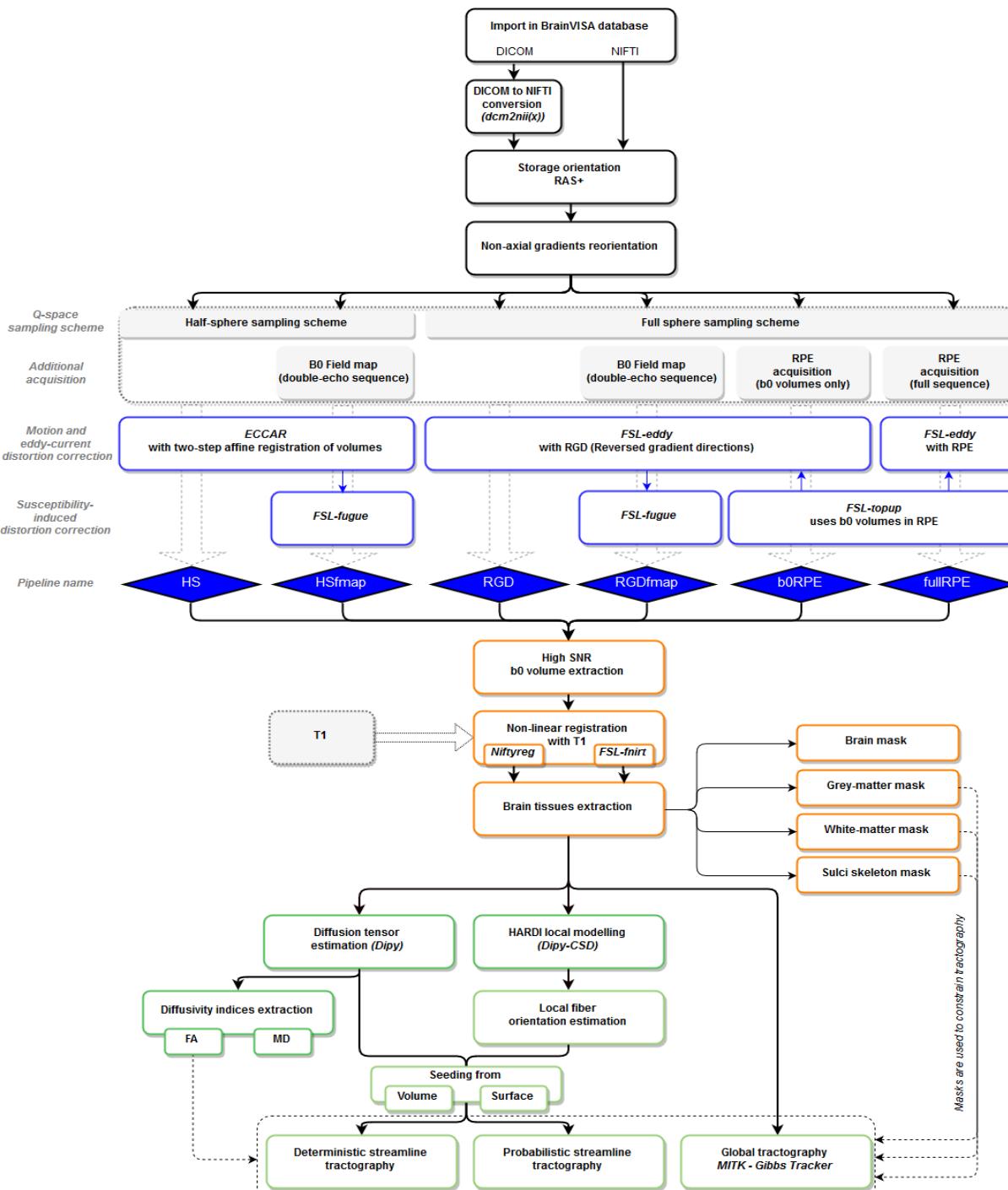
<http://fsl.fmrib.ox.ac.uk/fsl/fslwiki/>

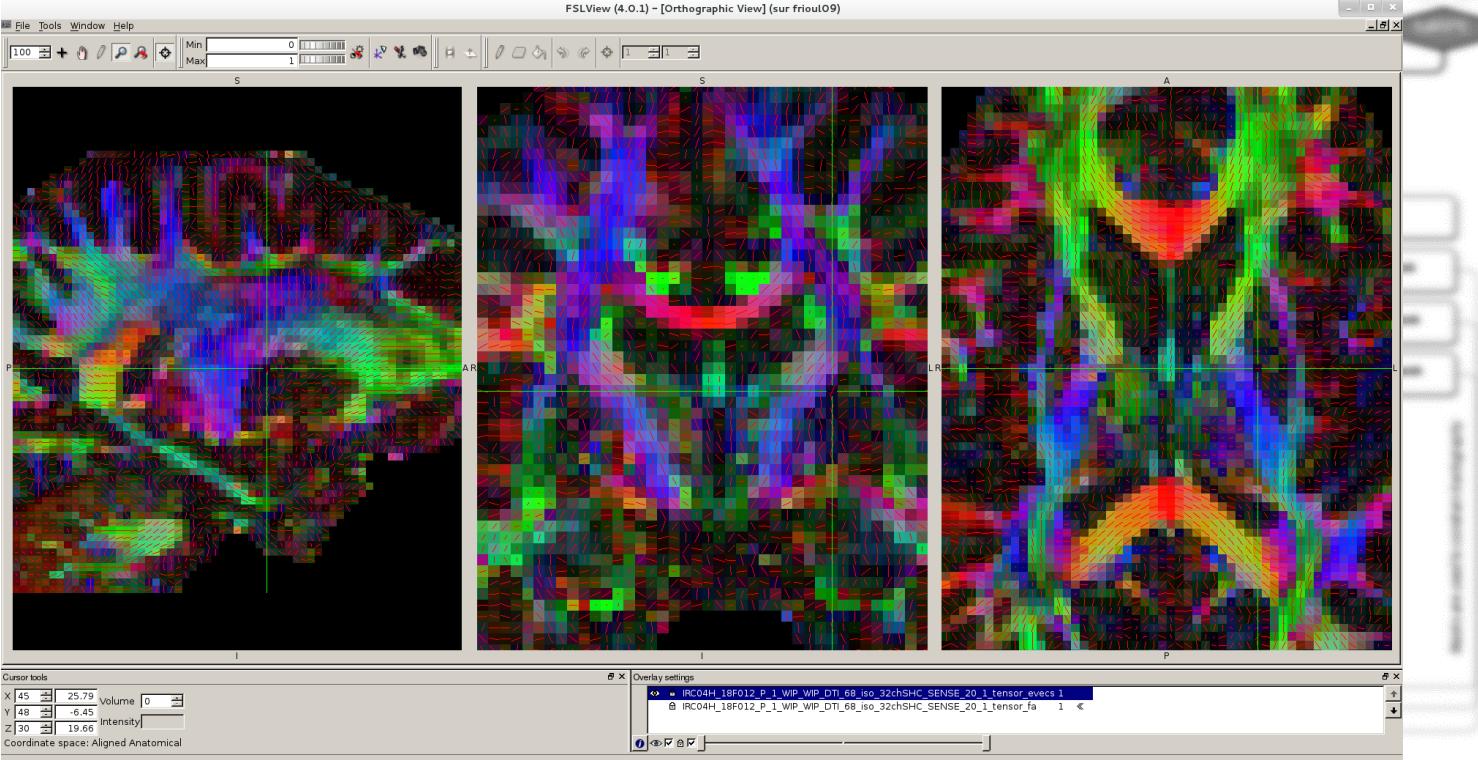
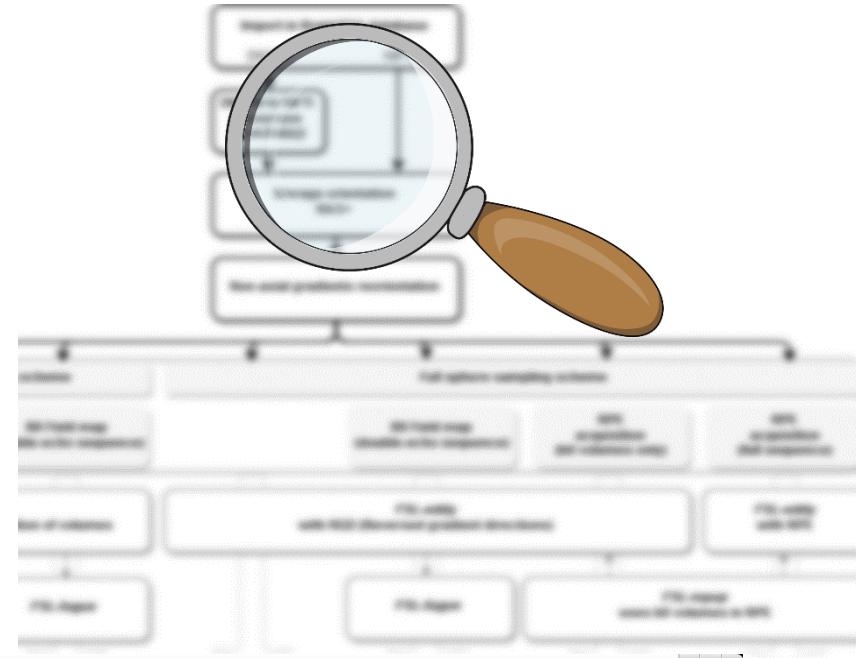
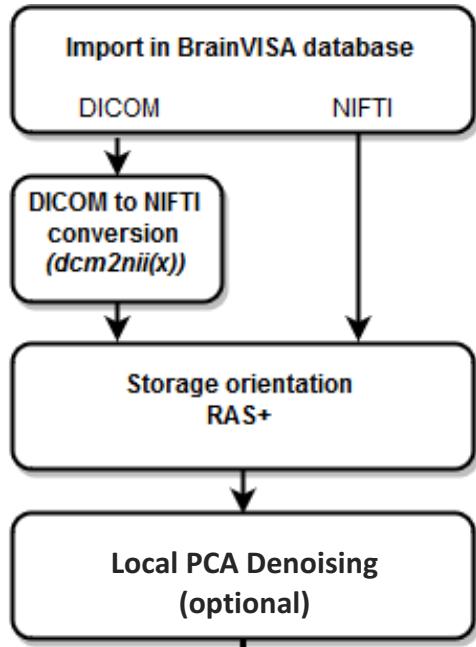


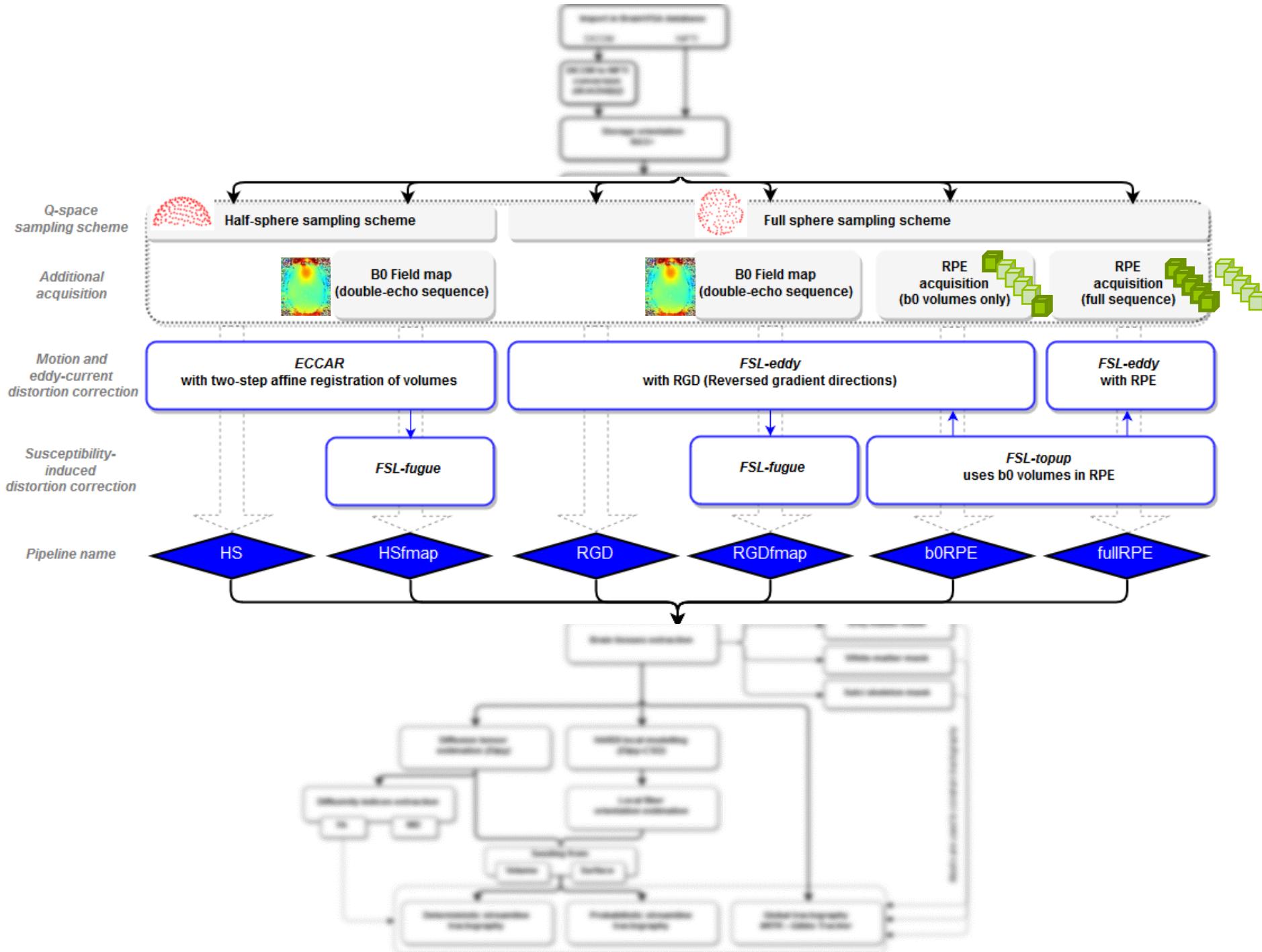
<http://nipy.org/dipy/>

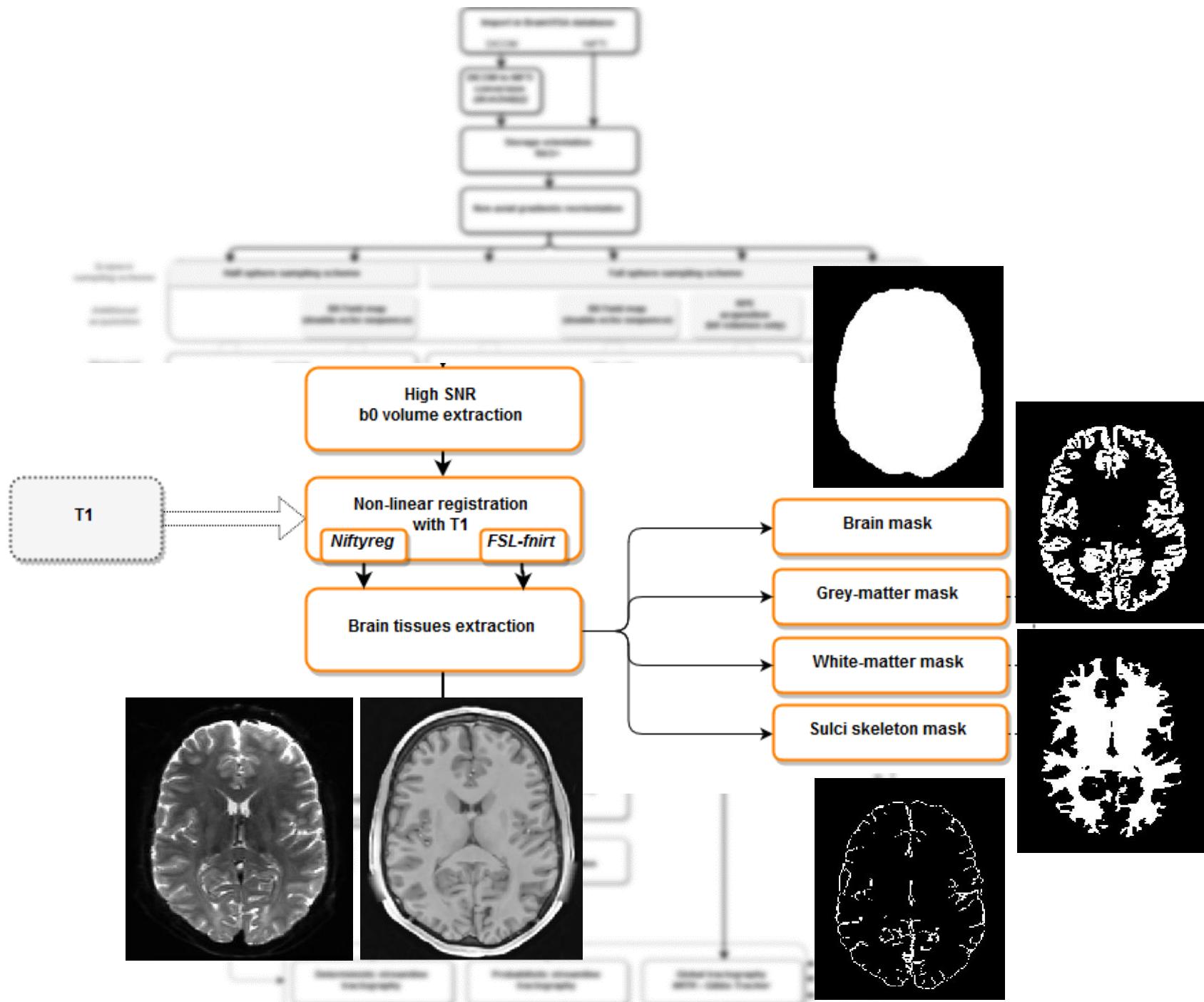
BrainVISA - Diffuse

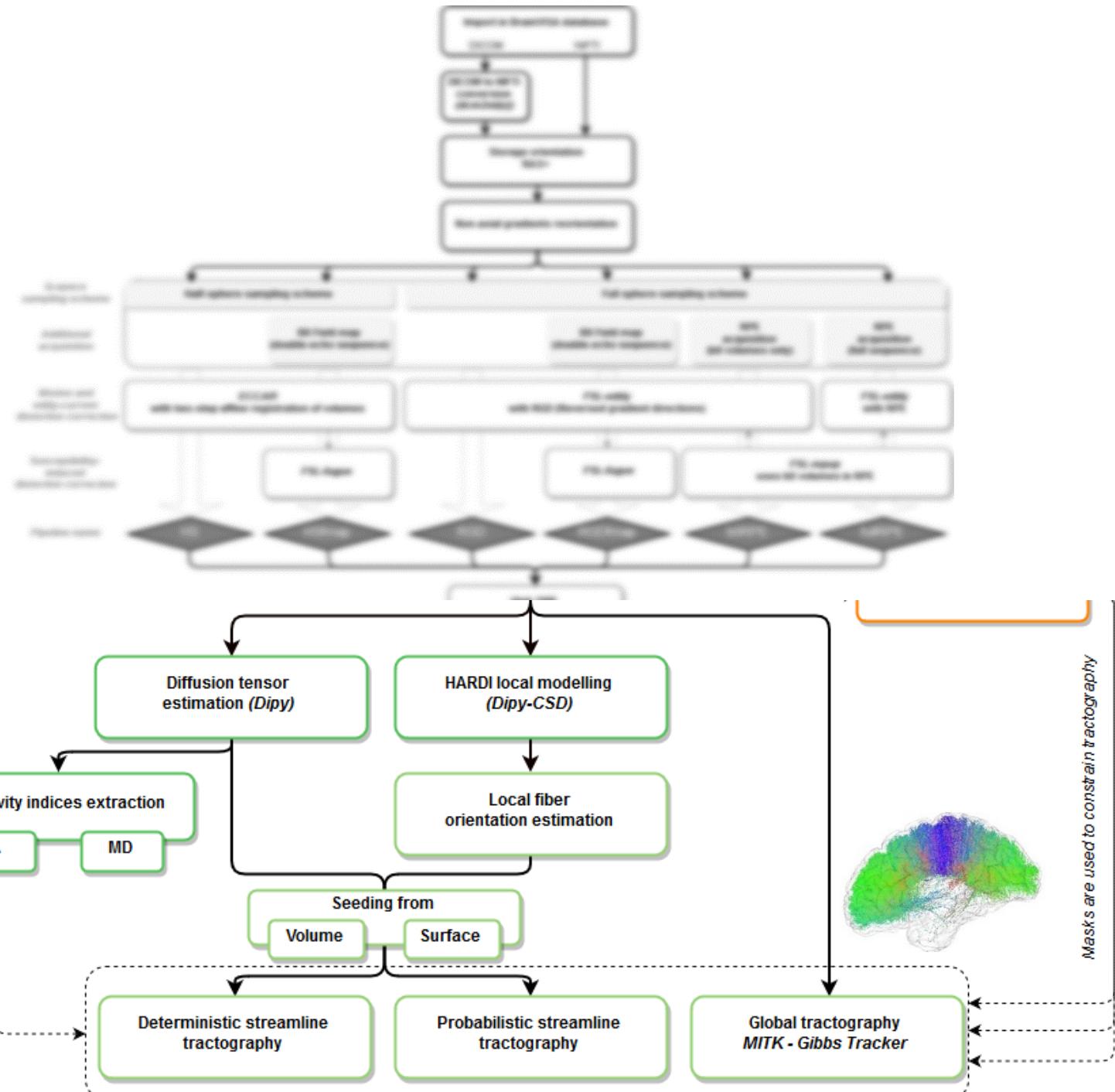
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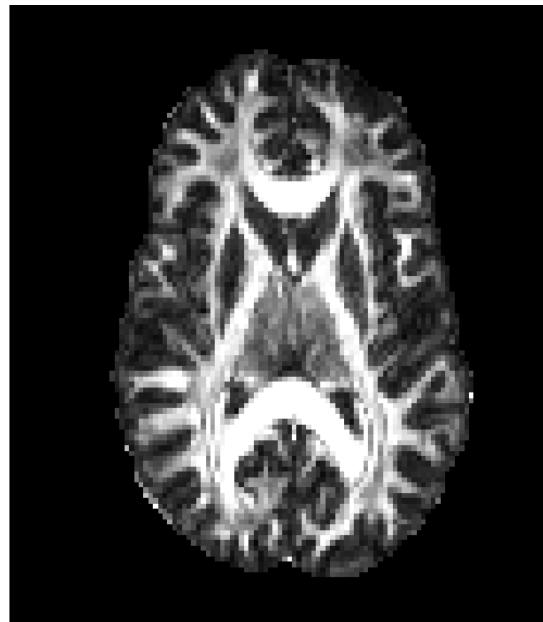


Diffuse Toolbox : DTI model

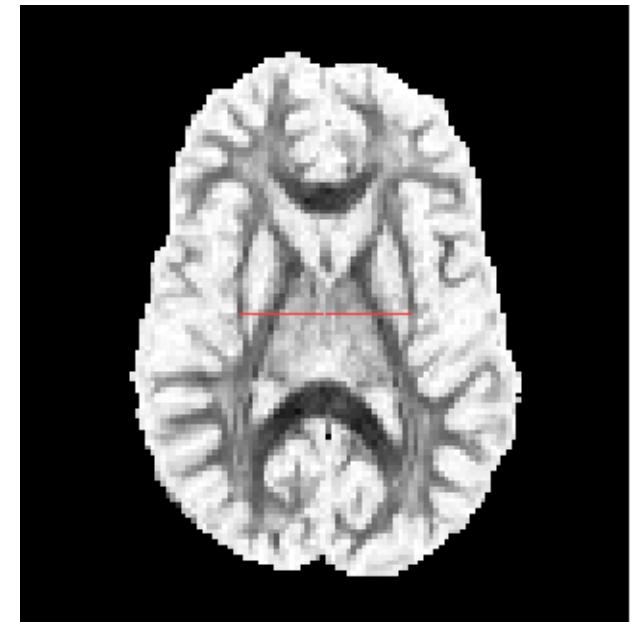
- Several estimation methods are available (Least Square, Weighted Least Square, Non Linear Least Square)



Mean diffusivity



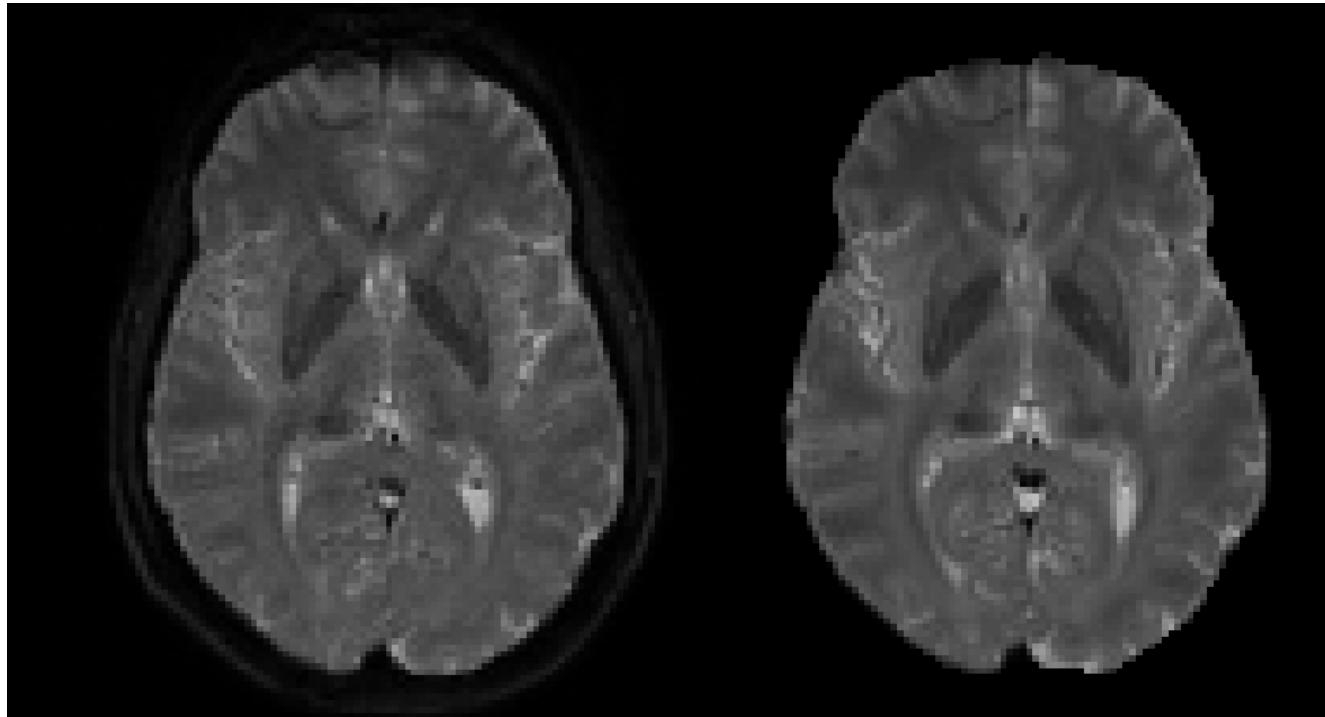
**Fractionnal
Anisotropy**



Sphericity

Diffuse Toolbox : DTI model

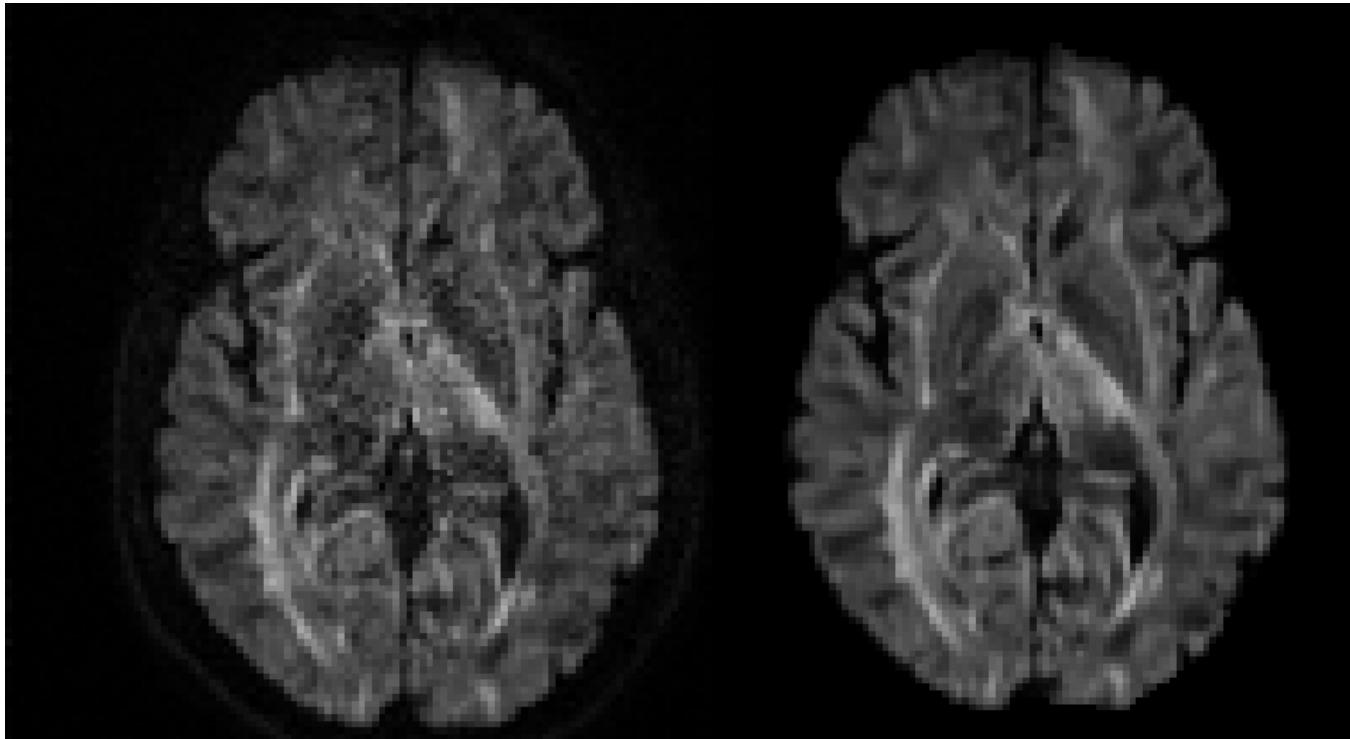
From tensor coefficients back to signal : dMRI signal prediction



b=0 : left (original), right (predicted)

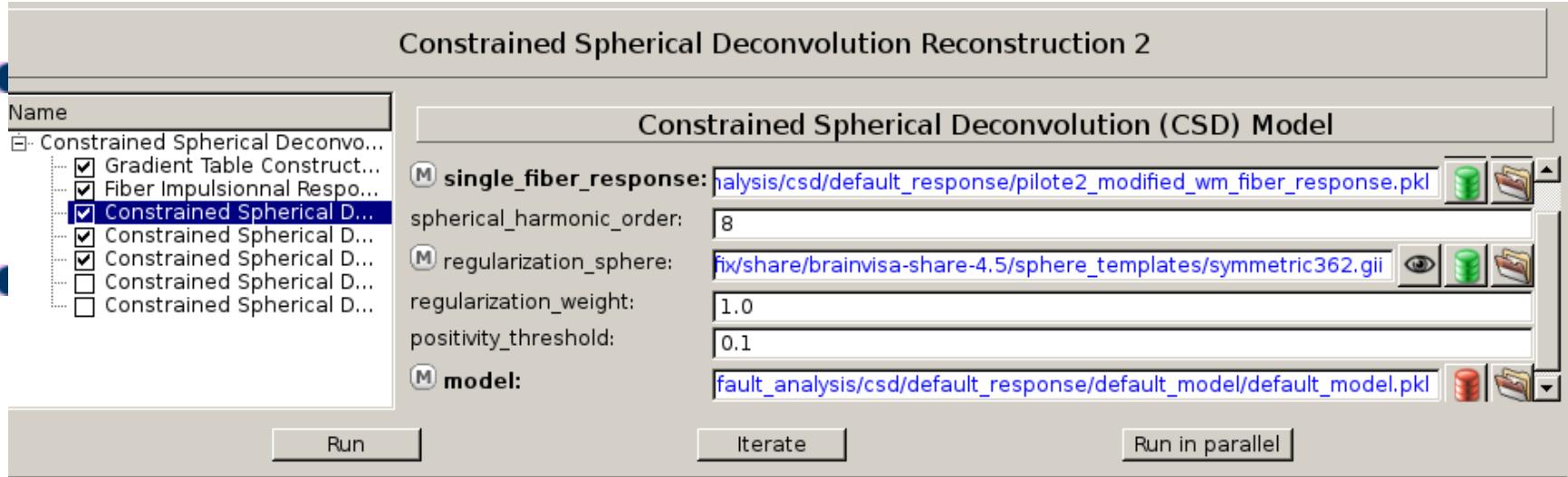
Diffuse Toolbox : DTI model

From tensor coefficients back to signal : dMRI signal prediction



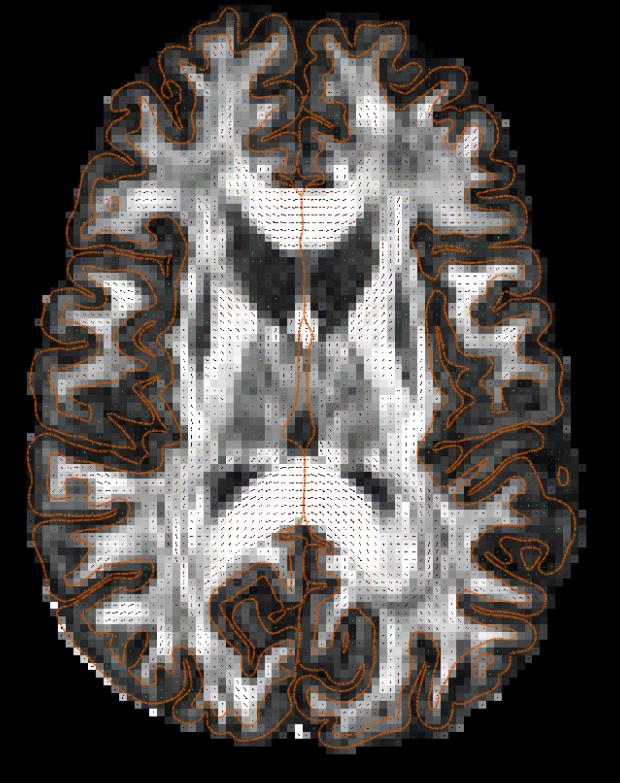
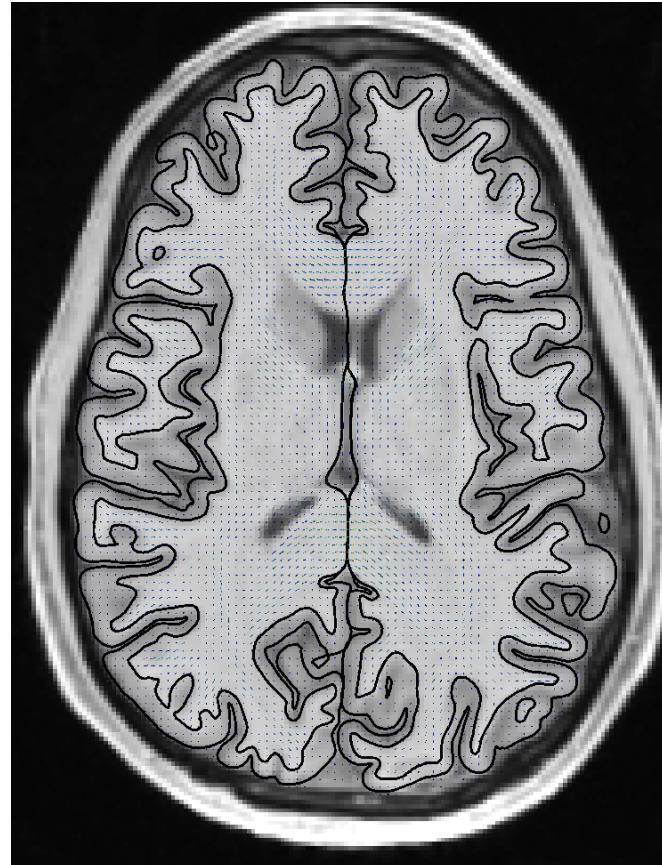
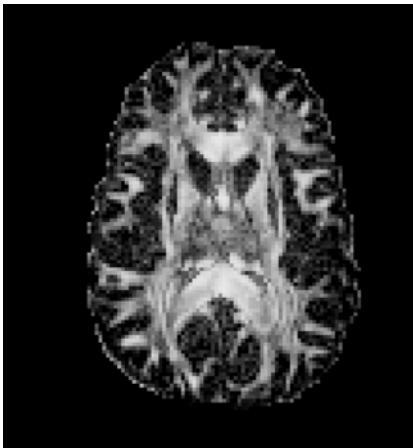
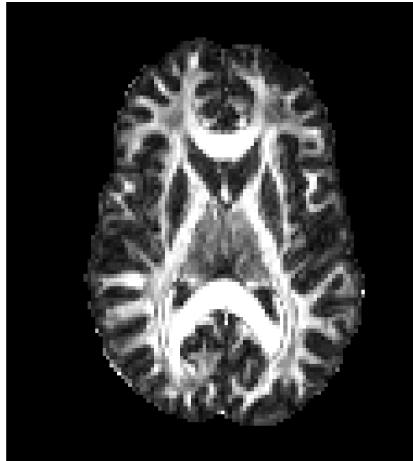
b=2000 : left (original), right (predicted)

Diffuse Toolbox : CSD model

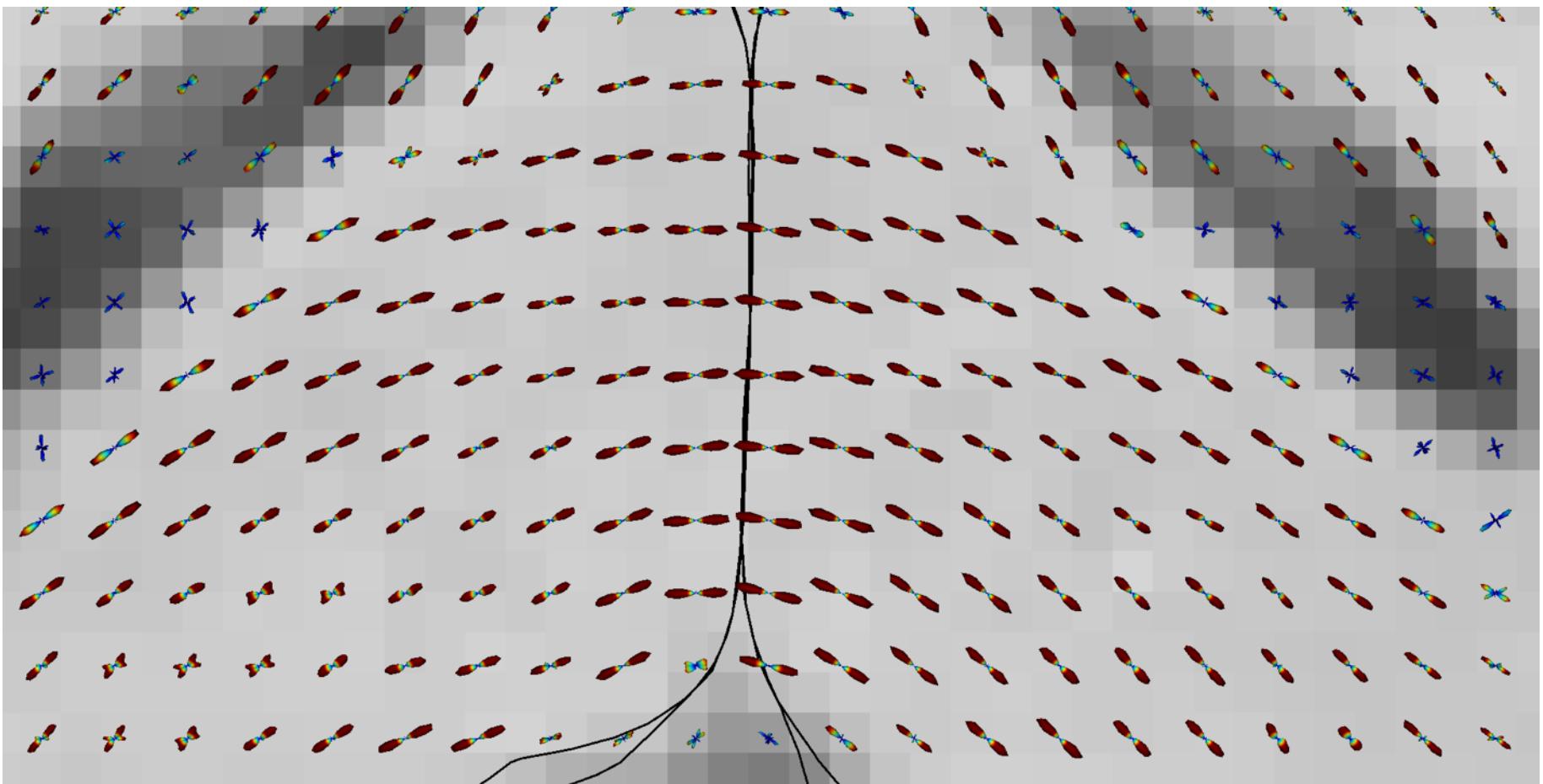


Estimation parameters such as spherical harmonic order and regularization sphere are setup automatically according to diffusion data sequence but can also be tuned using graphical user interface.

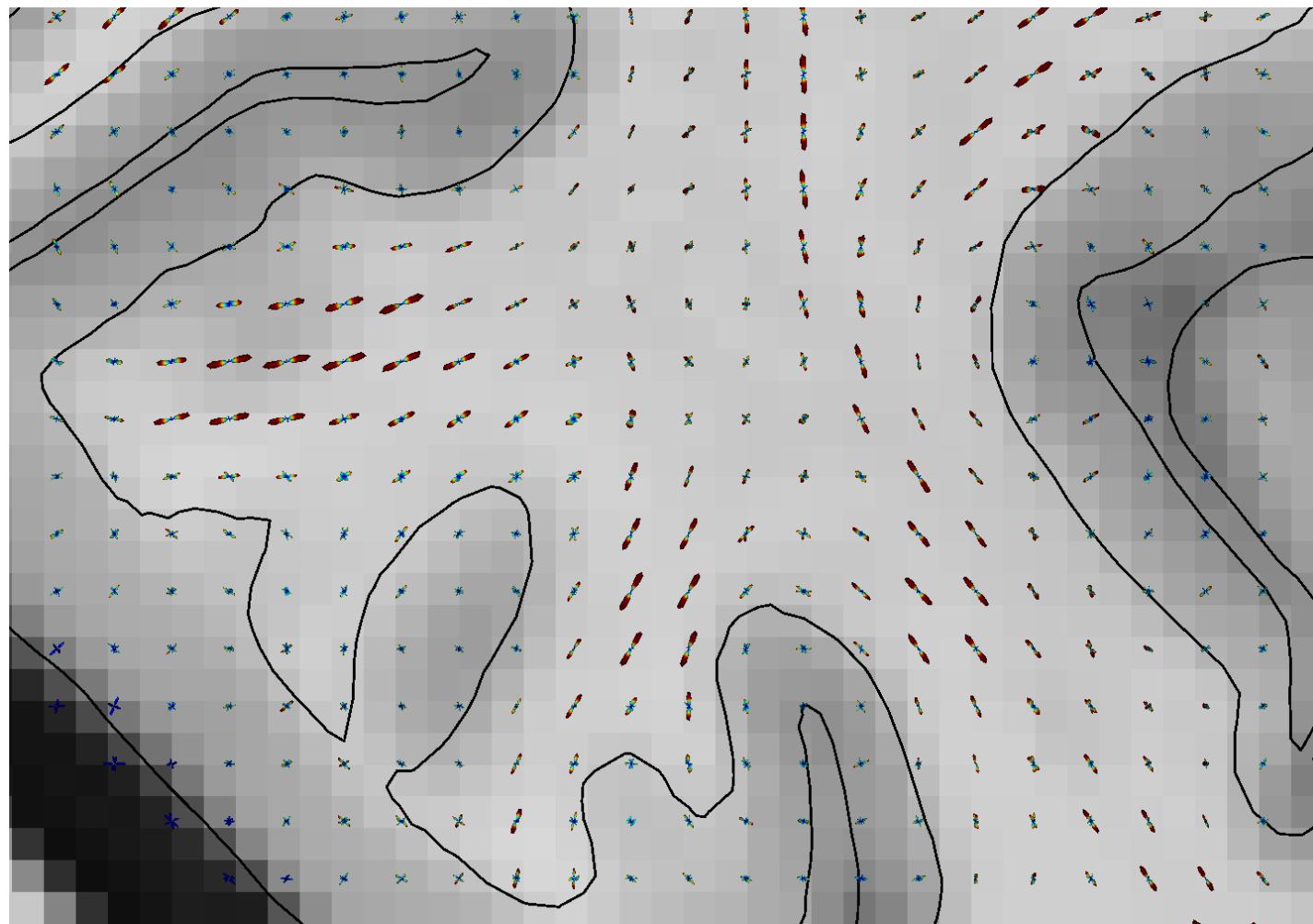
Diffuse Toolbox : CSD model



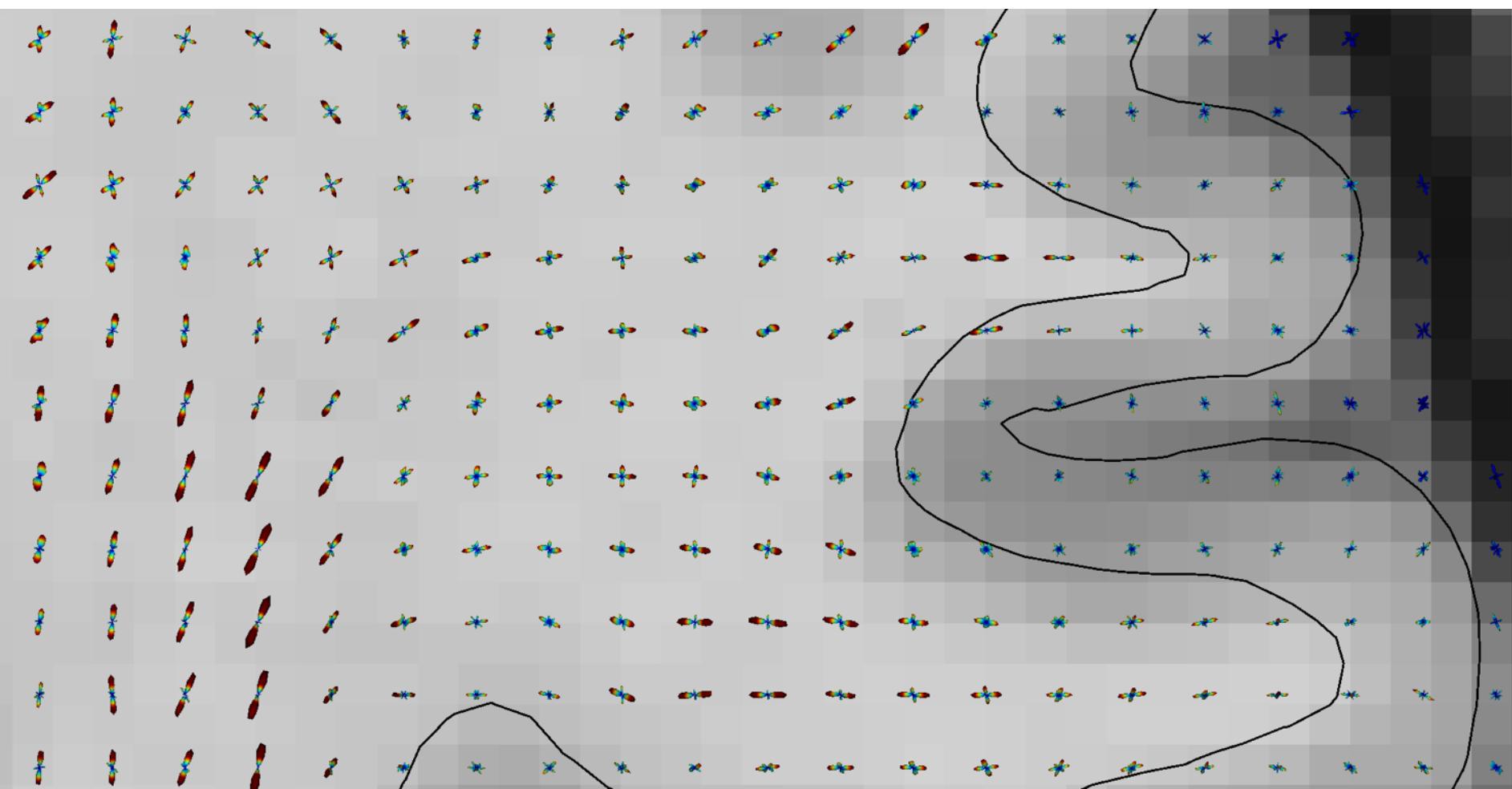
Diffuse Toolbox : CSD model



Diffuse Toolbox : CSD model

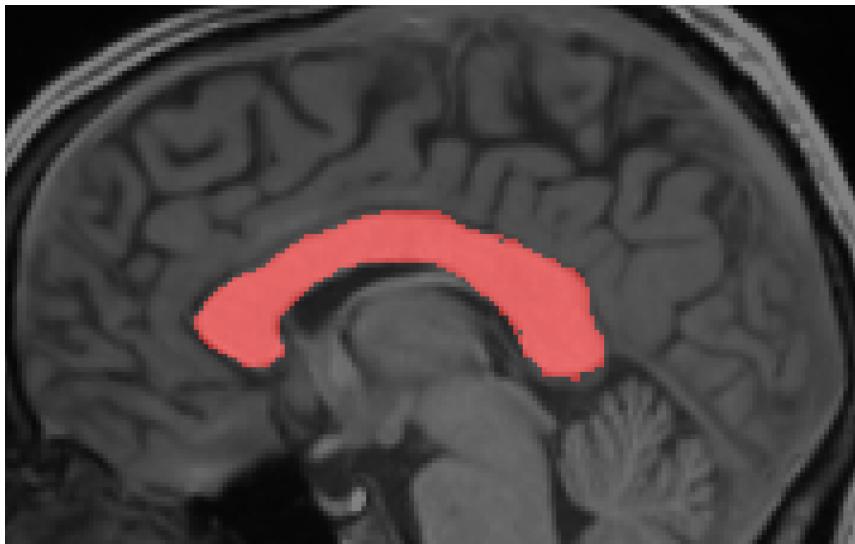


Diffuse Toolbox : CSD model



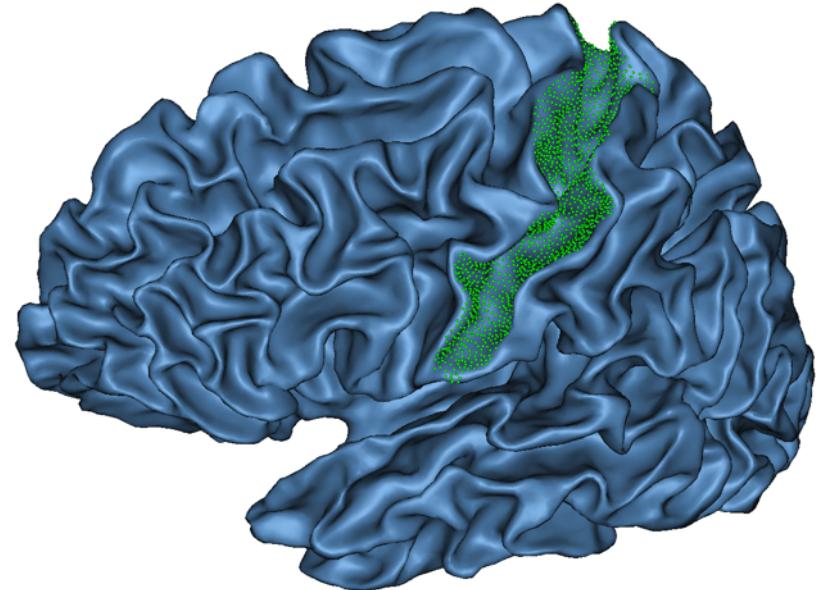
Diffuse Toolbox : Seeding strategy

- **Volumic Seeding**



Manually extracted corpus callosum
ROI from T1, diffusion space, subject
pilote2 from IRM center.

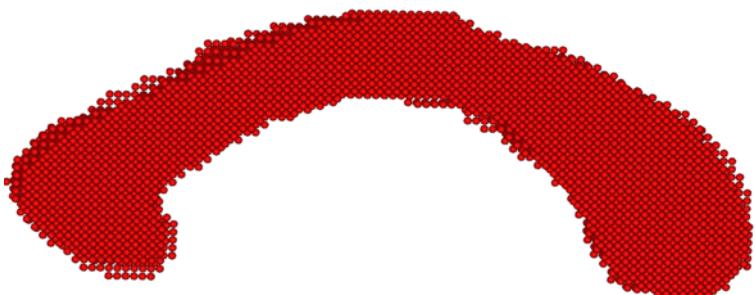
- **Surfacic seeding**



Central sulcus area (green) manually drawn
on left white mesh, diffusion space.

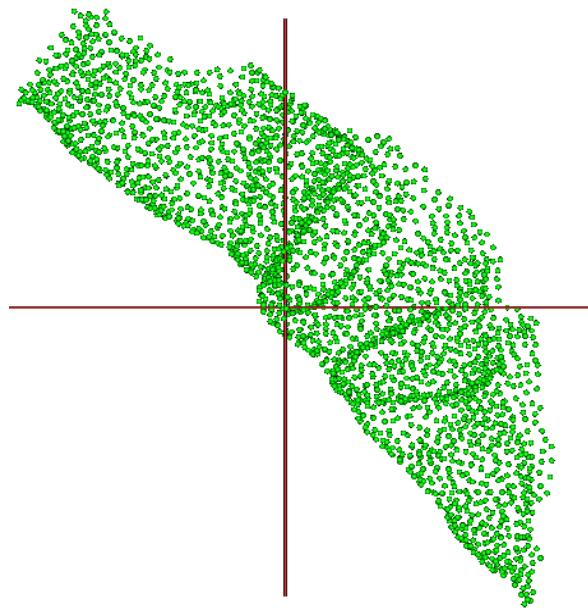
Diffuse Toolbox : Seeding strategies

- **Volumic Seeding**



Seeds from ROI mask, 1 seed/voxel
Deterministic placement.

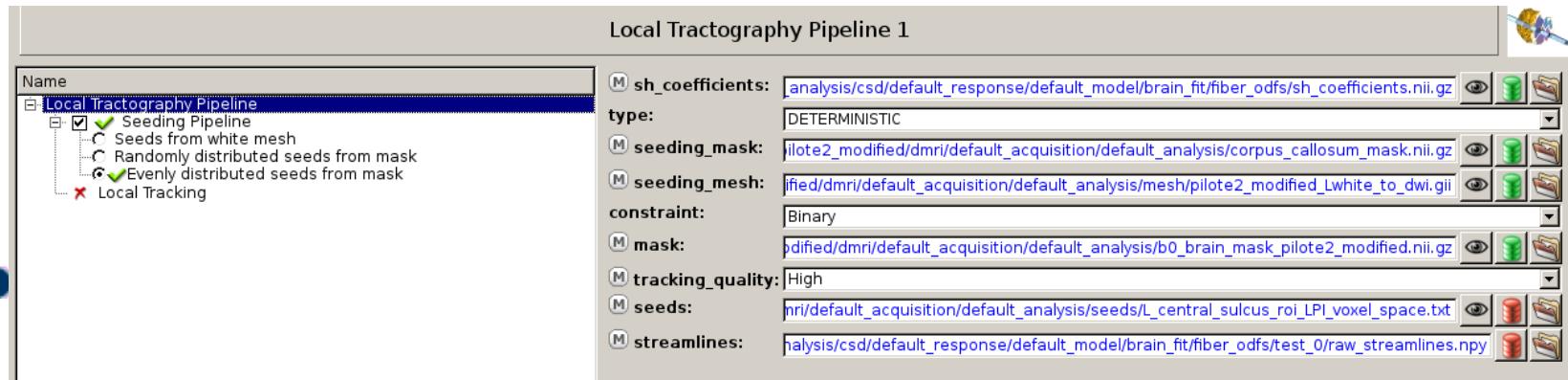
- **Surfacic seeding**



Seeds from surfacic ROI,
1seed/vertex. Refinement is possible

**These two mechanism can easily be combined to generate hybrid seeds
(e.g. volumic from subcortical grey volume and surfacic from white mesh)**

Diffuse Toolbox : Tractography parameters



Diffuse Toolbox : Tractography parameters

Local Tractography Pipeline 1

Name	Local Tractography Pipeline
Seeding Pipeline	<input checked="" type="checkbox"/> Seeds from white mesh <input type="checkbox"/> Randomly distributed seeds from mask <input checked="" type="checkbox"/> Evenly distributed seeds from mask
Local Tracking	

sh_coefficients: analysis/csd/default_response/default_model/brain_fit/fiber_odfs/sh_coefficients.nii.gz
type: DETERMINISTIC
seeding_mask: /pilot2_modified/dmri/default_acquisition/default_analysis/corpus_callosum_mask.nii.gz
seeding_mesh: /pilot2_modified/dmri/default_acquisition/default_analysis/mesh/pilot2_modified_Lwhite_to_dwi.gii
constraint: Binary
mask: /pilot2_modified/dmri/default_acquisition/default_analysis/b0_brain_mask_pilot2_modified.nii.gz
tracking_quality: High
seeds: /pilot2_modified/dmri/default_acquisition/default_analysis/seeds/L_central_sulcus_roi_LPI_voxel_space.txt
streamlines: analysis/csd/default_response/default_model/brain_fit/fiber_odfs/test_0/raw_streamlines.npy

Local Tractography Pipeline 1

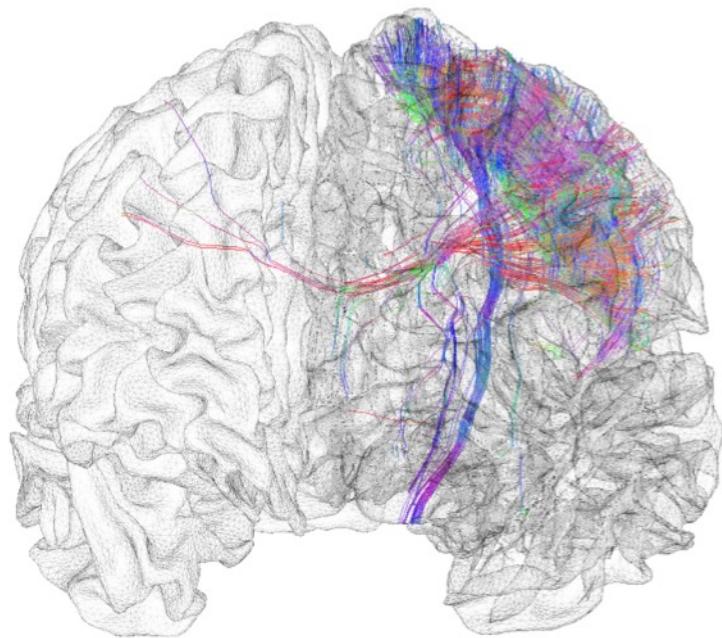
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Local Tracking

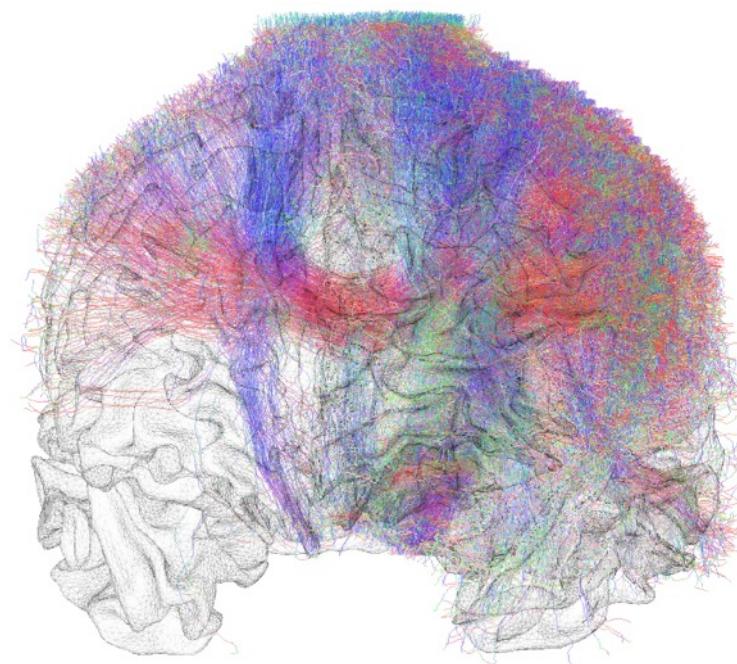
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type: DETERMINISTIC
constraint: Binary
mask: /pilot2_modified/dmri/default_acquisition/default_analysis/b0_brain_mask_pilot2_modified.nii.gz
sphere: /nvisia_compil_clust/build/bug_fix/share/brainvisa-share-4.5/sphere_templates/symmetric362.gii
seeds: /pilot2_modified/dmri/default_acquisition/default_analysis/seeds/L_central_sulcus_roi_LPI_voxel_space.txt
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relative_peak_threshold: 0.1
min_separation_angle: 5.0
crossing_max:
step_size: 0.1
fixed_step:
nb_iter_max: 500
return_all:
streamlines: default_analysis/csd/default_response/default_model/brain_fit/fiber_odfs/test_0/raw_streamlines.npy

Diffuse Toolbox : Tractography examples

- **Deterministic tracking**



- **Probabilistic tracking**



Surfacic seeds, left central sulcus,
angle=45 degrees, step size=0.1

BrainVISA - Diffuse

- Introduction: dMRI data processing
 - Pre-processing
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 - BrainVISA environment
 - Workflow
 - Demo
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 - Acquisition/Correction quality
 - Impact on co-registration
 - Impact on signal (diffusivity metrics)
- Outcome
 - Acquisition strategy
 - CentreIRM

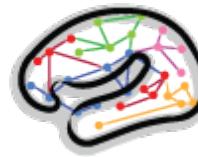
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Données

HCP Data - 900 Subjects

- 20 sujets sains, 25-30 ans
- 1,25 x 1,25 x 1,25 mm
- $b = 1000, 2000, 3000 \text{ s/mm}^2$
- 3 x 95 directions (sphère entière, régulière)
- Double encodage de phase LR/RL
- Fieldmap



HUMAN
Connectome
PROJECT

3T Prisma/Skyra

TR = 5520 ms
TE = 89.5 ms
MB = 3
FOV = 210 x 188

Centre IRM - INT

- 1 sujet sain, 25 ans
- 1,5 x 1,5 x 1,5 mm
- $b = 2000 \text{ s/mm}^2$
- 60 directions (sphère entière, régulière)
- Double encodage de phase AP/PA
- Fieldmap

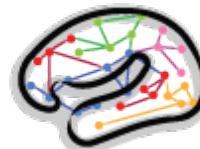
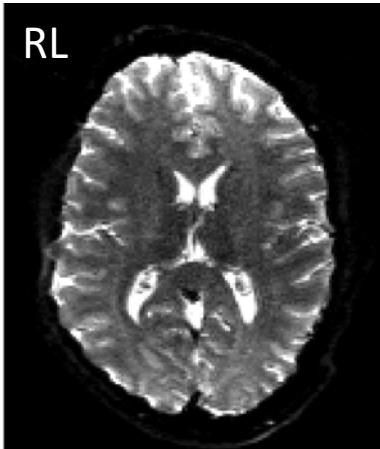
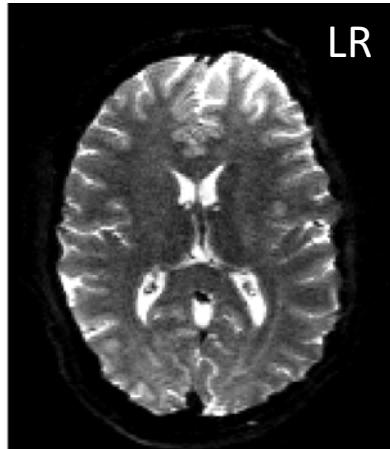


3T Siemens Prisma

TR/TE = 3270/87ms
MB = 4
FOV = 210²

Données

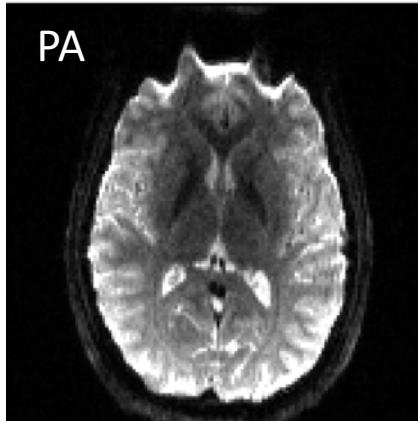
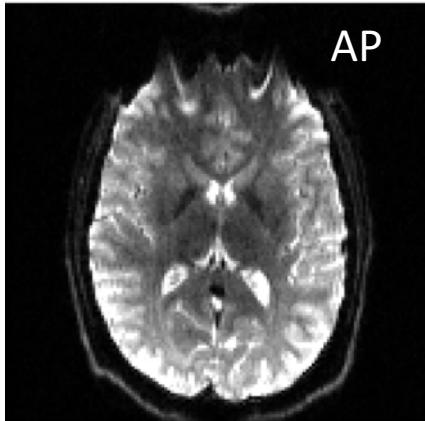
HCP Data - 900 Subjects



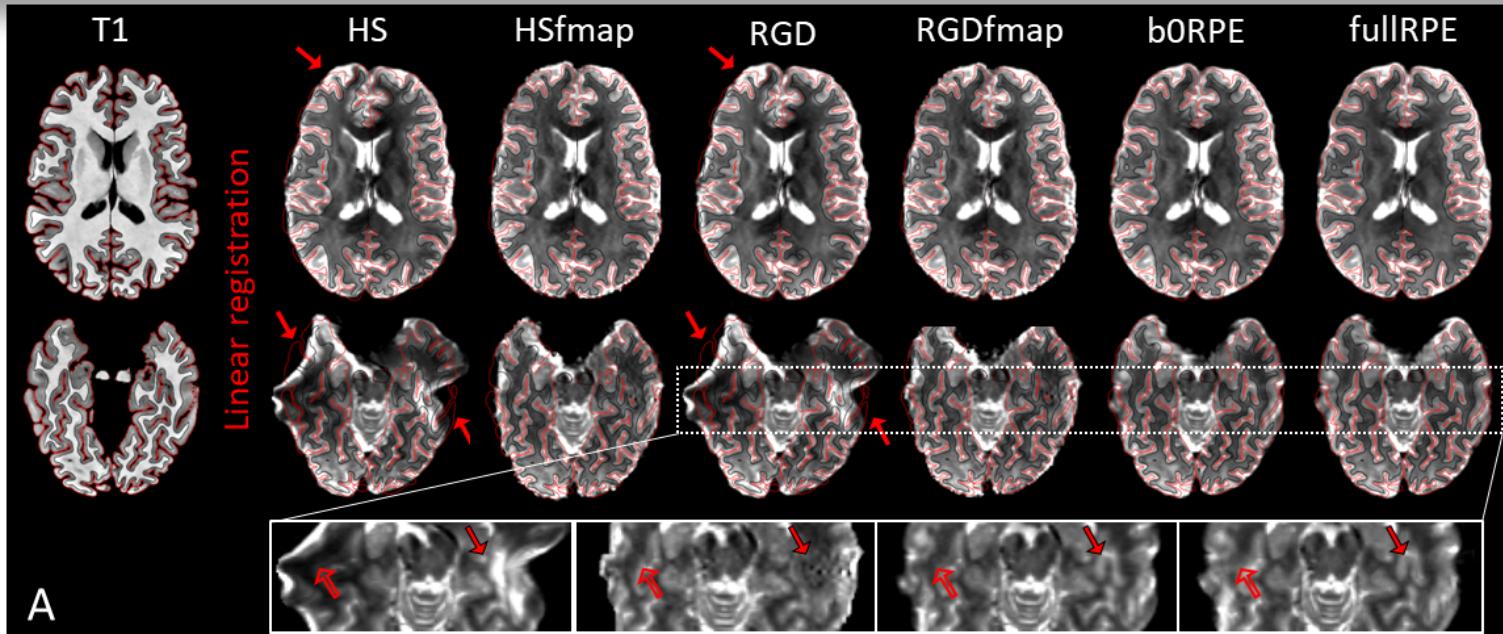
HUMAN
Connectome
PROJECT

3T Siemens Prisma

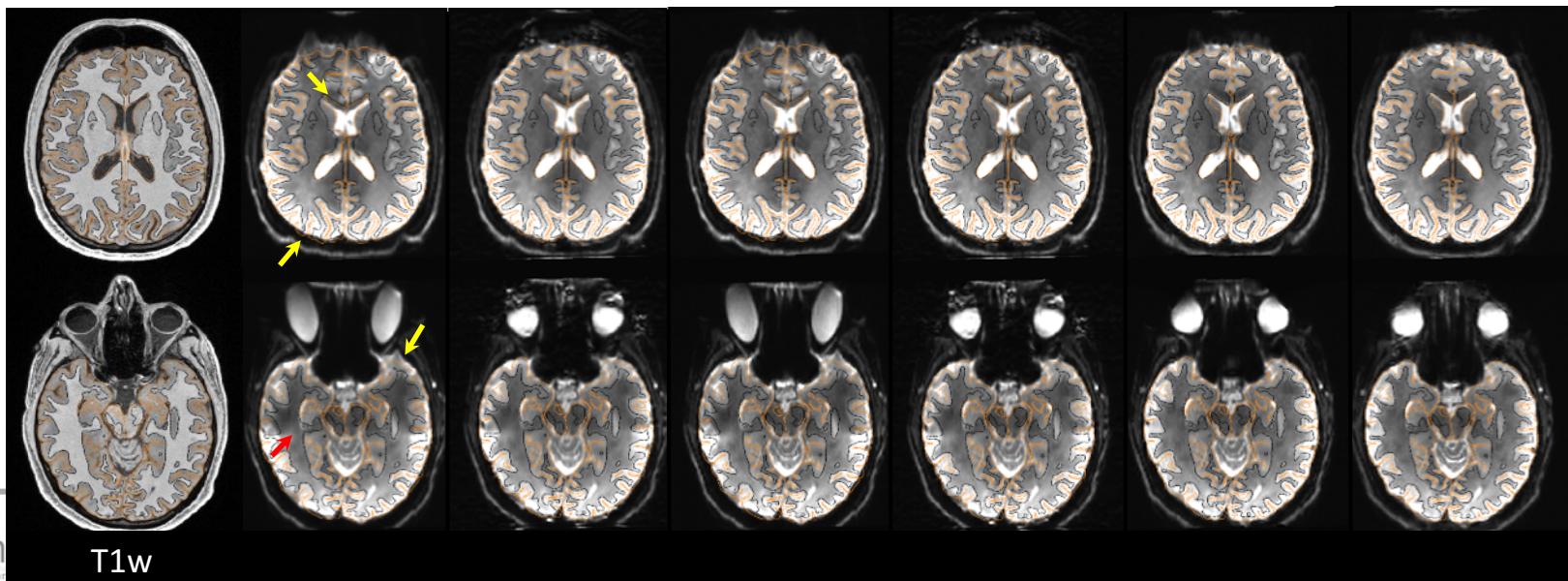
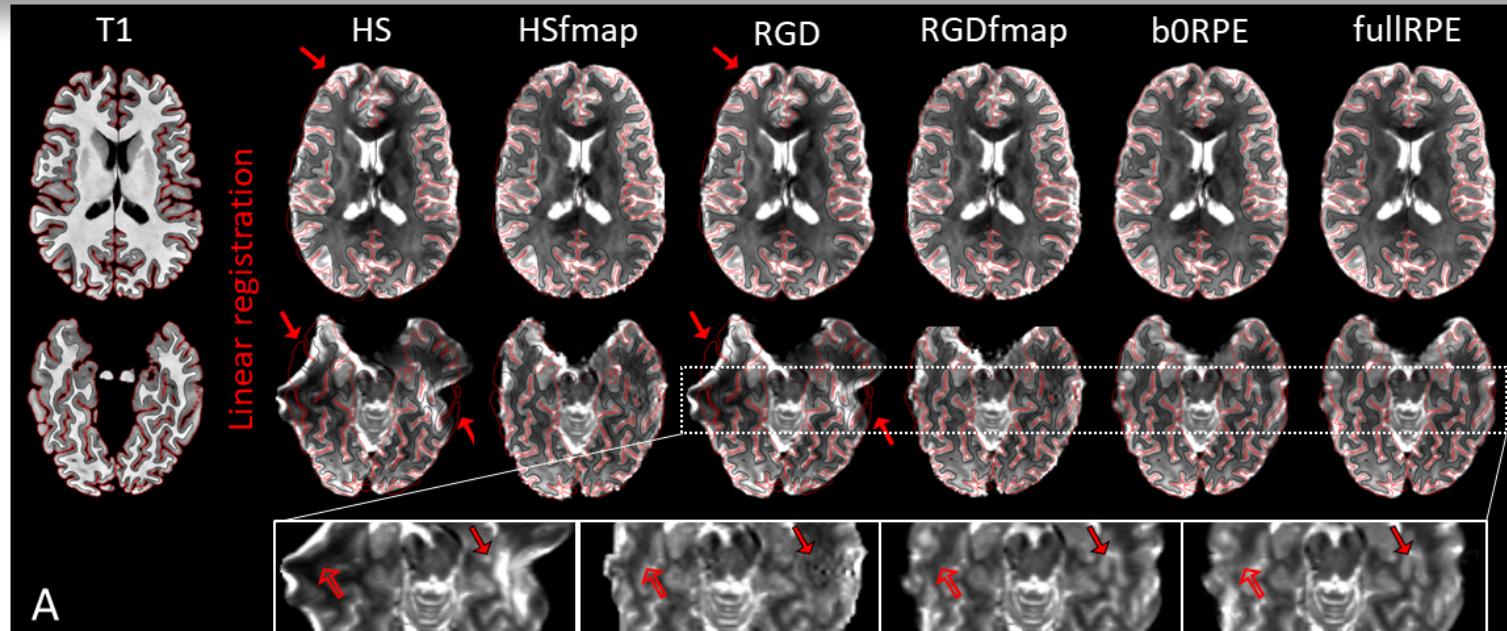
Centre IRM - INT



Comparison: distortion correction quality



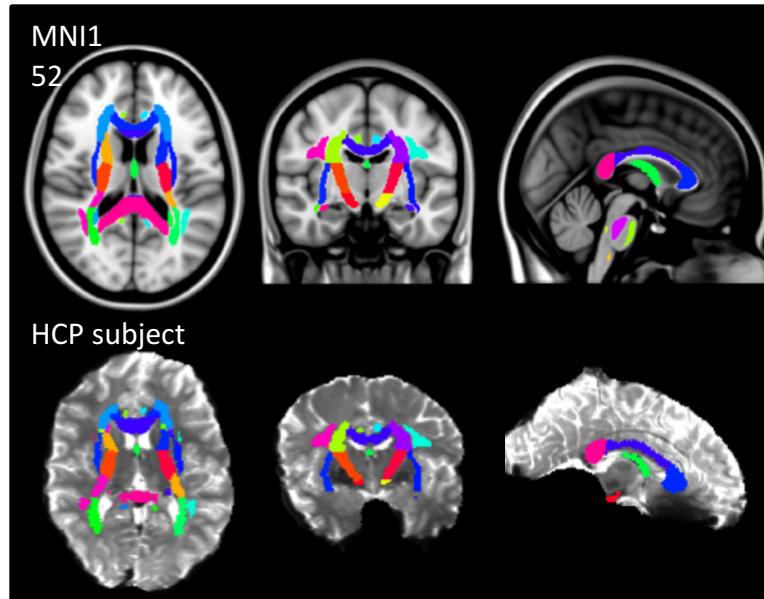
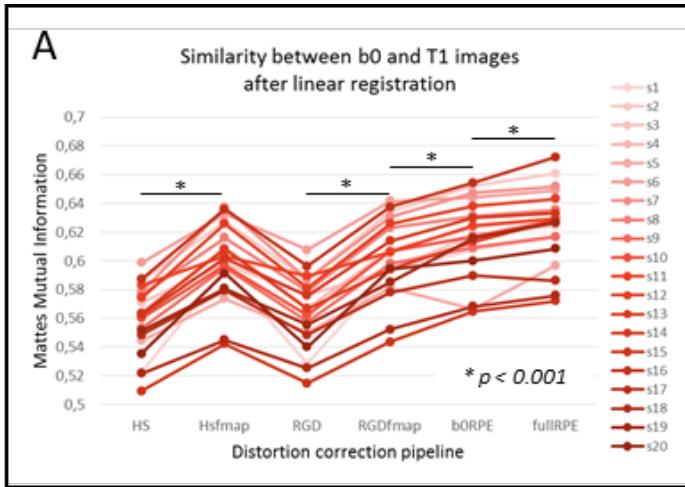
Comparison: distortion correction quality



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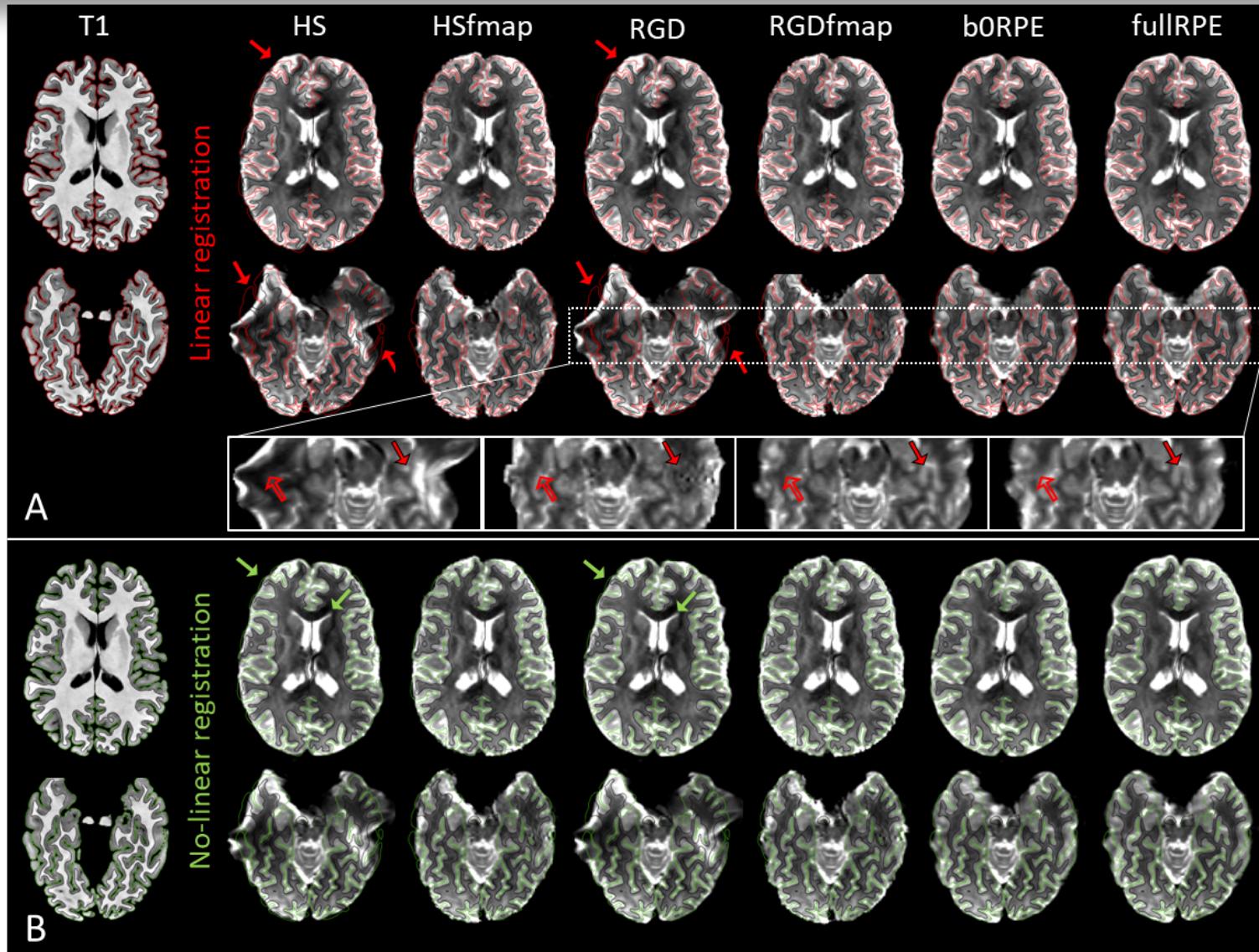
Comparison: co-registration quality



- Quality of distortion correction: is the true brain geometry recovered ?
- Quality of co-registration with T1
 - Impact on brain masking/segmentation
 - Impact on **seeding strategy !**

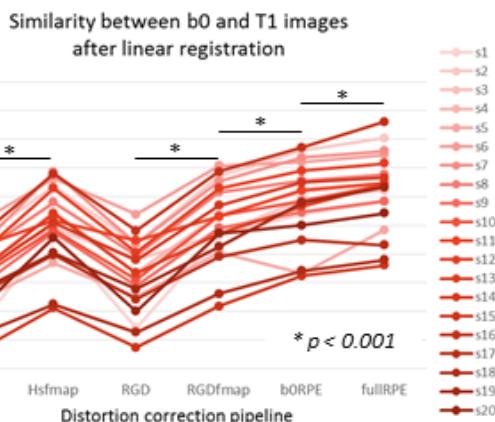
⇒ **Use non-linear co-registration**

Comparison: non-linear registration

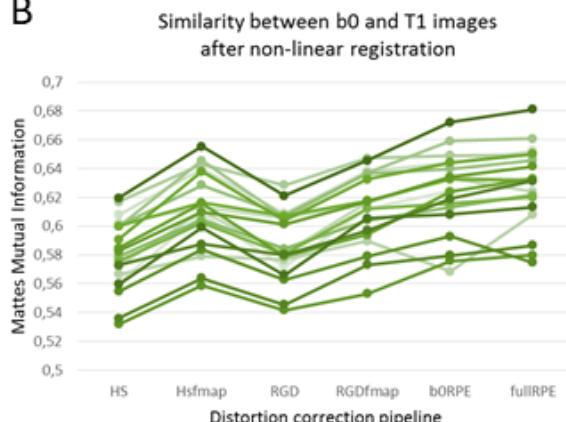


Comparison: non-linear registration

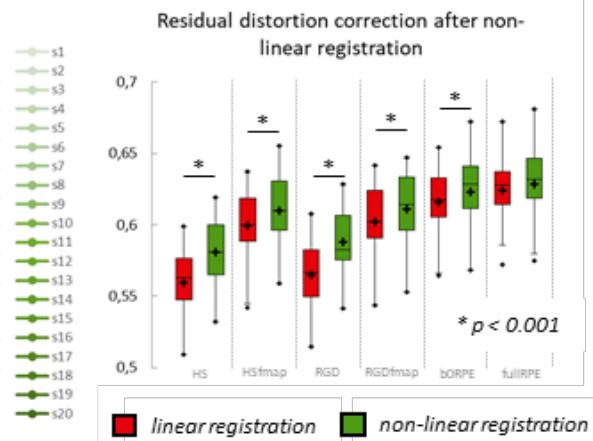
A



B



Residual distortion correction after non-linear registration

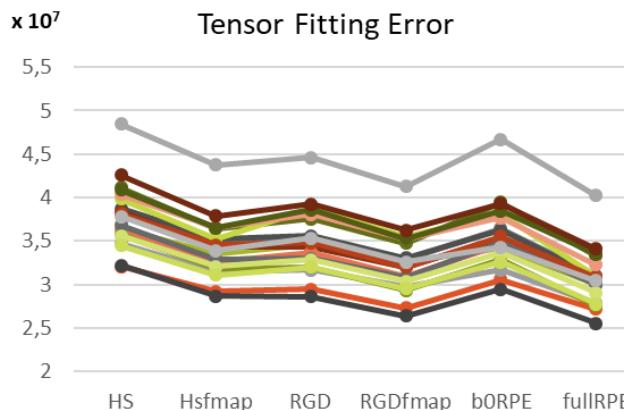


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Comparison: impact on diffusivity metrics

Quantitative metrics

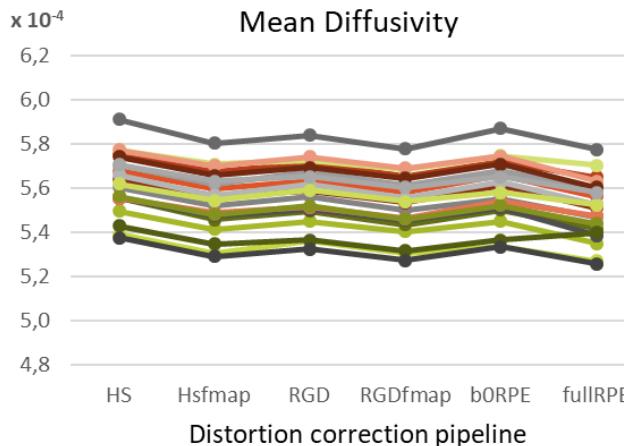


$$TFE = \sum_{i=1}^N (S_{mi} - S_{fi})^2$$

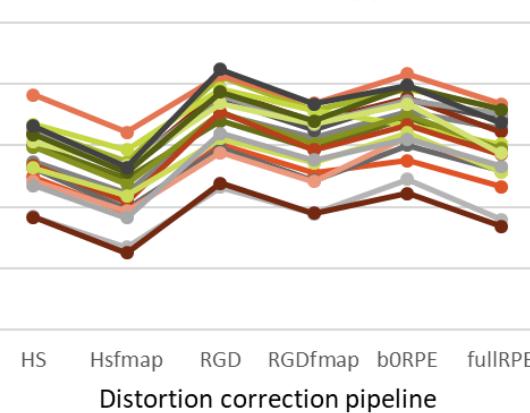
$$MDI = \frac{1}{n(S)} \sum_{x \in S} \sqrt{\frac{\lambda_2 + \lambda_3}{2\lambda_1}}$$

$$FA = \sqrt{\frac{3}{2}} \frac{\sqrt{(\lambda_1 - \hat{\lambda})^2 + (\lambda_2 - \hat{\lambda})^2 + (\lambda_3 - \hat{\lambda})^2}}{\sqrt{\lambda_1^2 + \lambda_2^2 + \lambda_3^2}}$$

Qualitative metrics



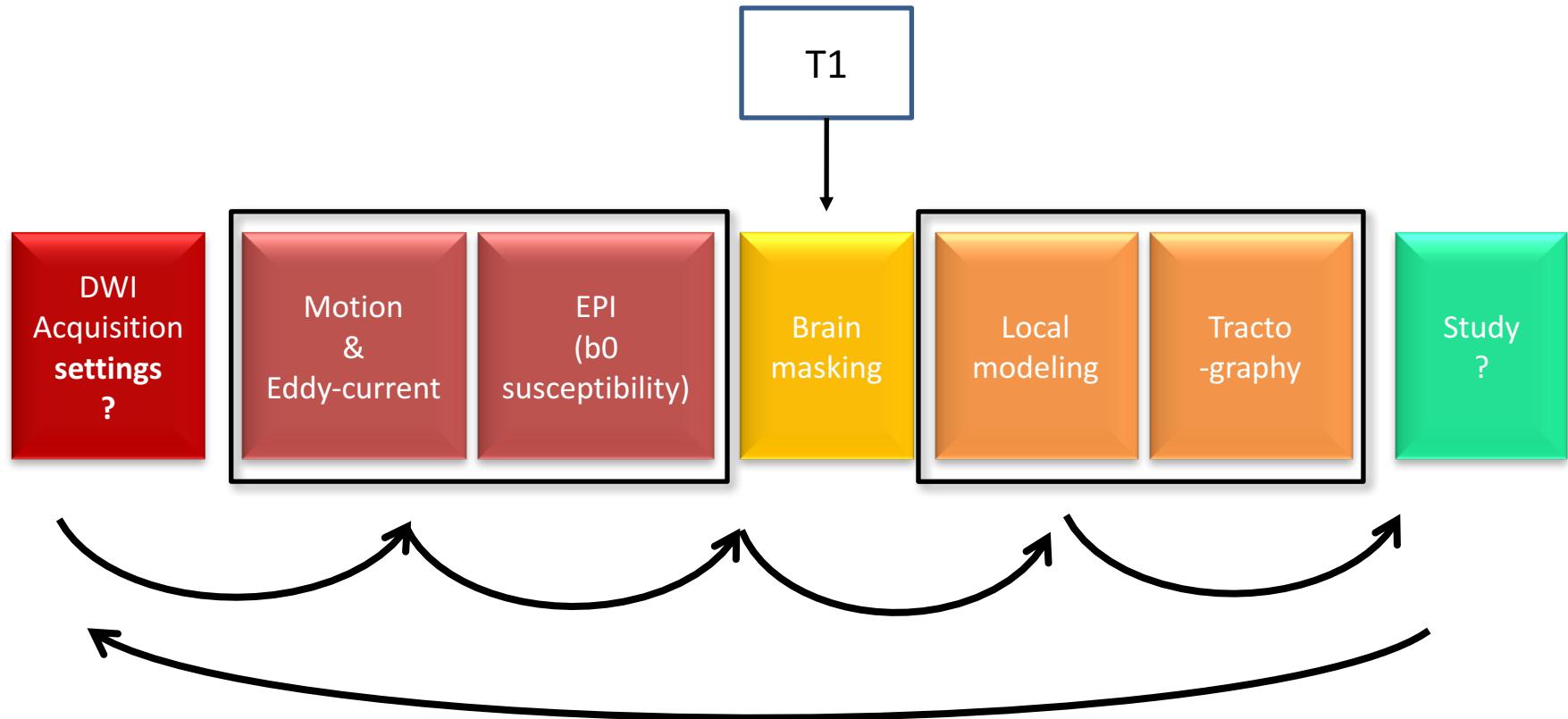
Fractional Anisotropy



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Prétraitements et corrections des distorsions : quelle stratégie d'acquisition ?



Prétraitements et corrections des distorsions : quelle stratégie d'acquisition ?

On the menu of the MRI Center

Fast and less Furious

3min23s for AP acquisition : 76 volumes (32b100 + 32 b2000 + 6 b300 + 6 b0)
Resolution 1.8mm iso, MB4

Acquisition of only 6b0 in PA possible to gain time (acquisition time: 30s)

Less Fast and More Furious

5min39s for AP acquisition: 110 volumes (64b2000 + 32 b1000 + 6 b300 + 8b0)
Resolution 1.8mm iso, MB4
Acquisition of only 8b0 in PA possible to gain time (acquisition time: 40s)

A la carte

Possibility to chose any b-values , multishell , NODDI, DSI, HARDI...





INT - MeCA

Olivier COULON
Lucile Brun
Alexandre PRON

INT - SCALP

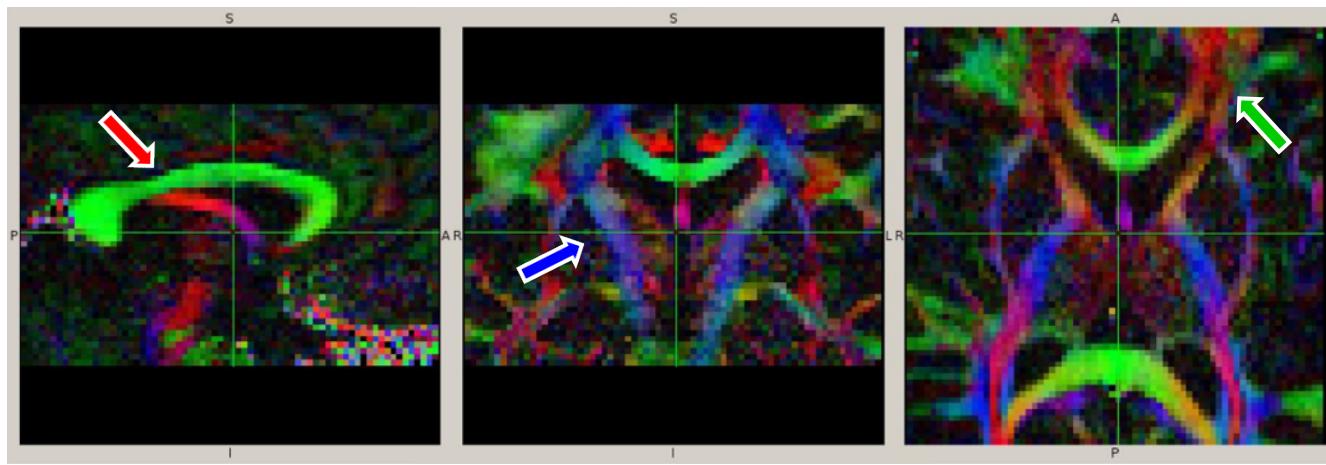
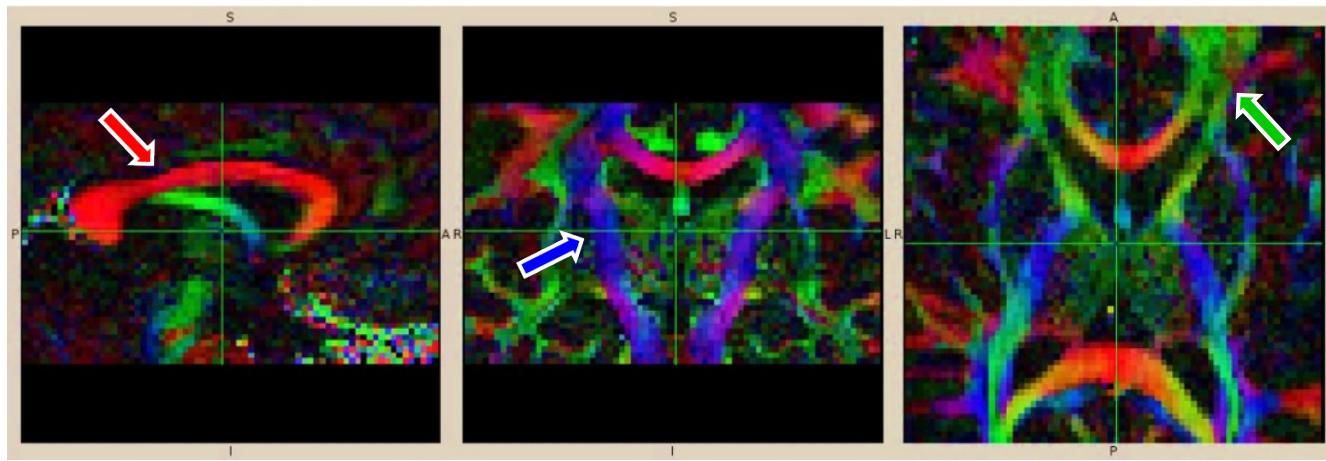
Christine DERUELLE

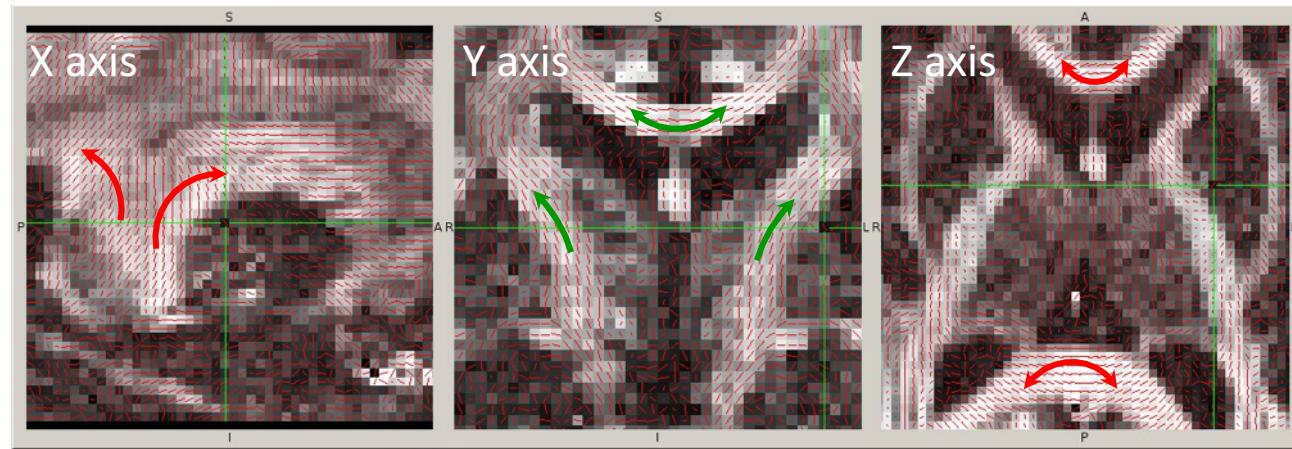
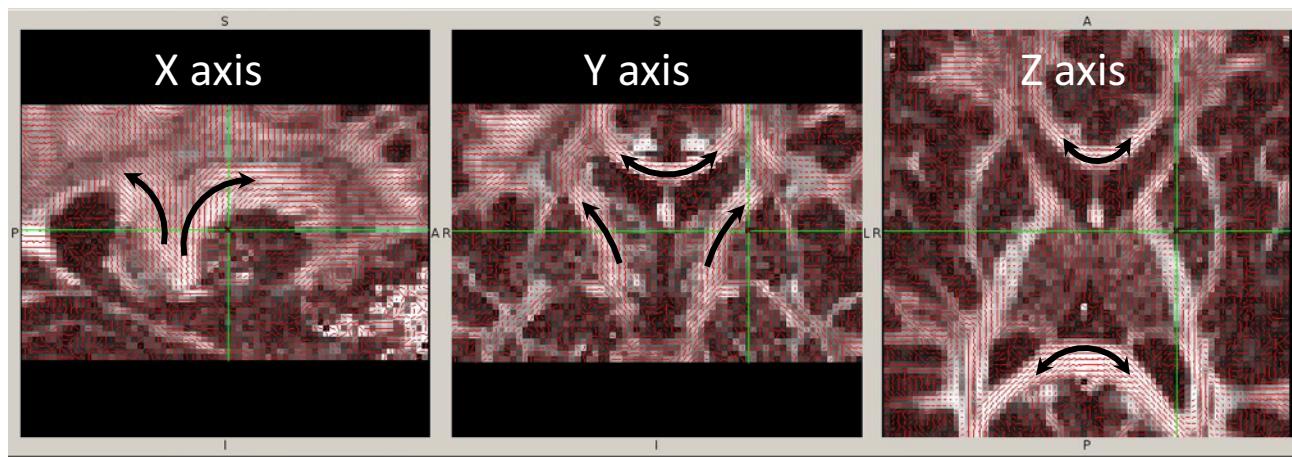


Centre IRM – INT @ CERIMED

Jean-Luc ANTON
Pascal BELIN
Bruno NAZARIAN
Julien SEIN



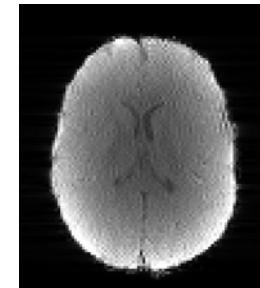
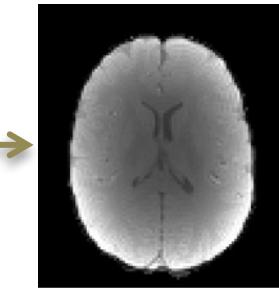
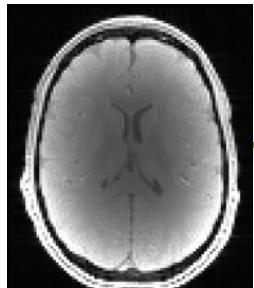




Magnetic susceptibility

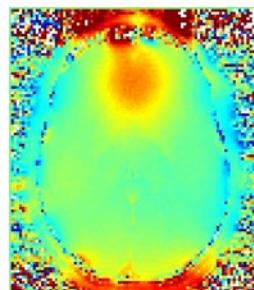
Cusack et al, NeuroImage 2003

Magnitude



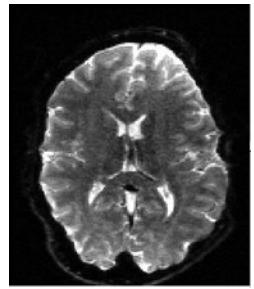
warped
magnitude

Fieldmap



Forward
warping
FSL-fugue

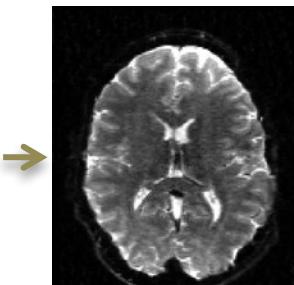
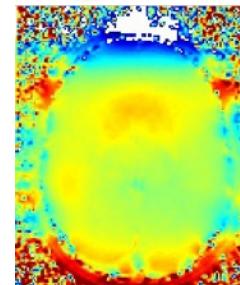
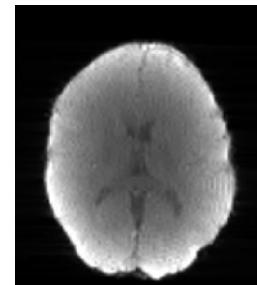
DWI



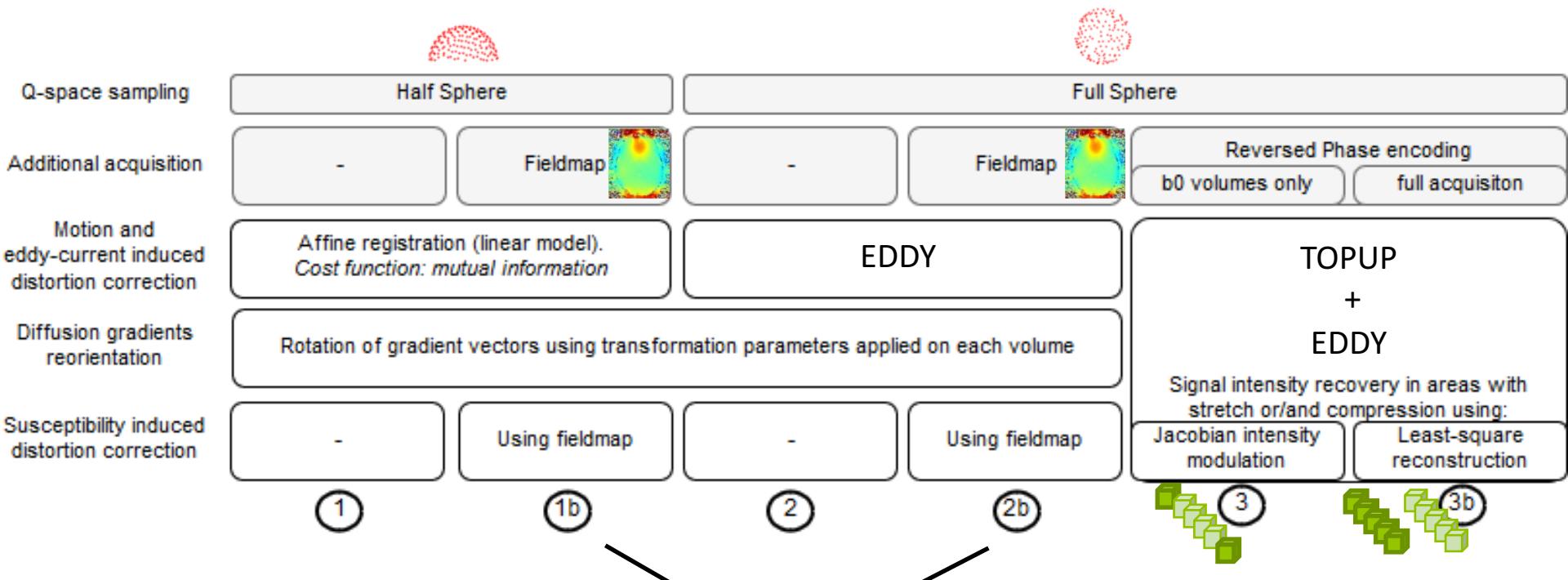
FSL-flirt
12dof

Récupération du
signal compressé
impossible !

Unwarping
FSL-fugue unwarped
data



Acquisition strategy



- Low computing time
- Two resampling steps

- Meilleure estimation du champ de distorsion des EC
- Un seul ré-échantillonnage de l'image finale