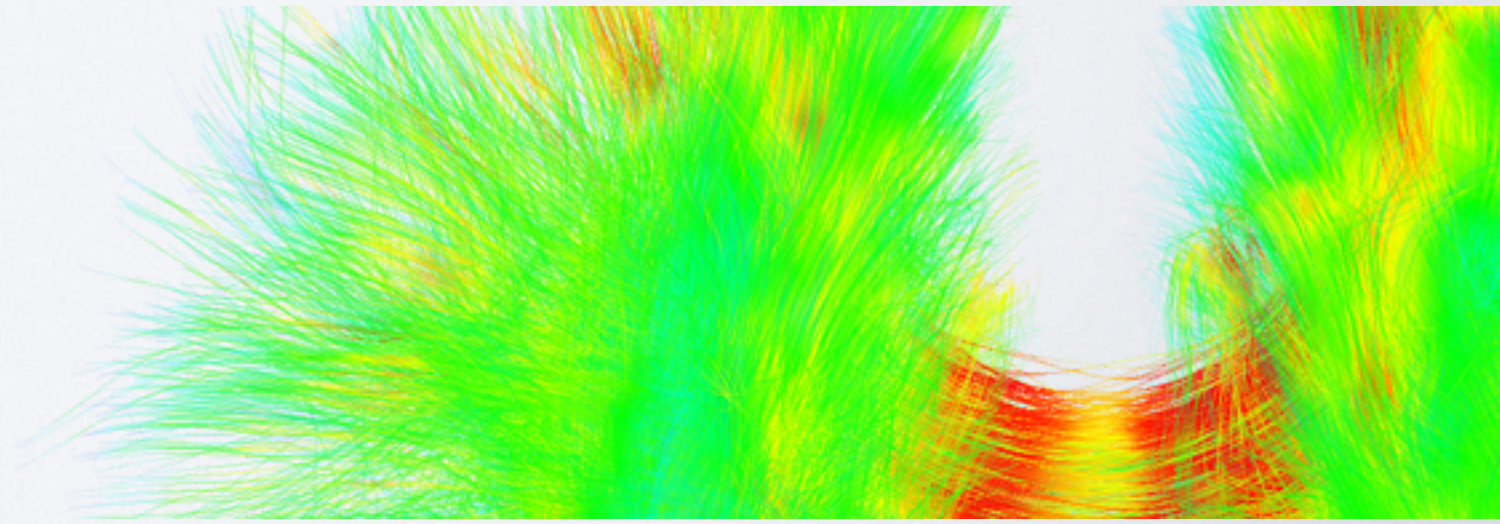
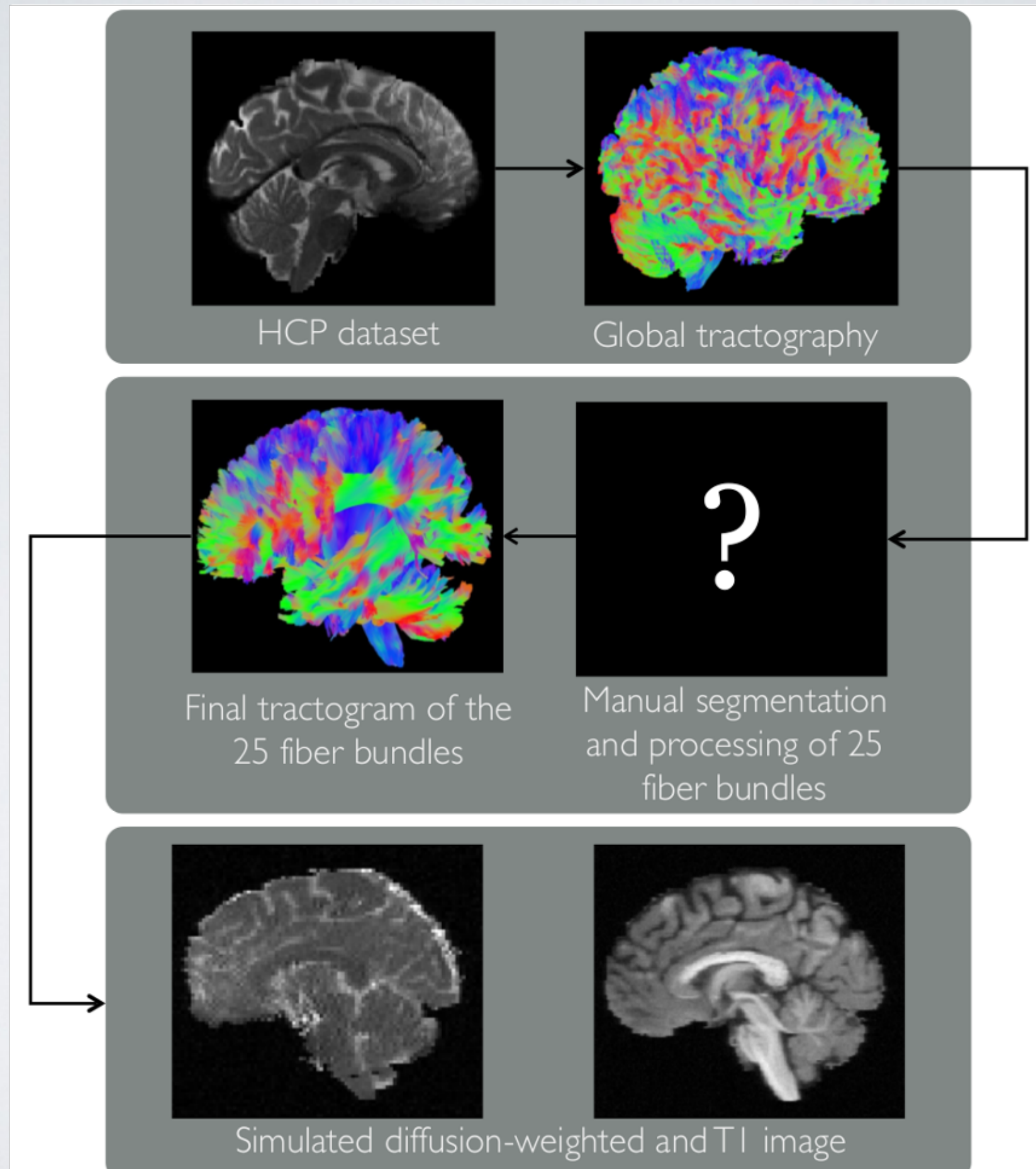


TRACTOGRAPHY CHALLENGE 2015

ISMIRM Diffusion Study Group



EXAMPLE OF THE DATA GENERATION PROCESS



IN A NUTSHELL

1. Manually segmented white matter bundles.
 - These serve as ground truth streamline fibers.
2. Clinical-like diffusion MRI dataset generated using *Fiberfox*¹:
 - 2mm isotropic, artefacts (motion, distortions, noise, etc.), 32 directions, b-value 1000 s/mm², 2 b=0 images
3. *Tractometer*² for the evaluation of fiber tracking results
 - Sensitivity & specificity with ground truth

[1] Neher, P. F., Laun, F. B., Stieltjes, B., & Maier-Hein, K. H. ***Fiberfox: Facilitating the creation of realistic white matter software phantoms.*** *MRM* 72(5), 1460-1470, 2014.

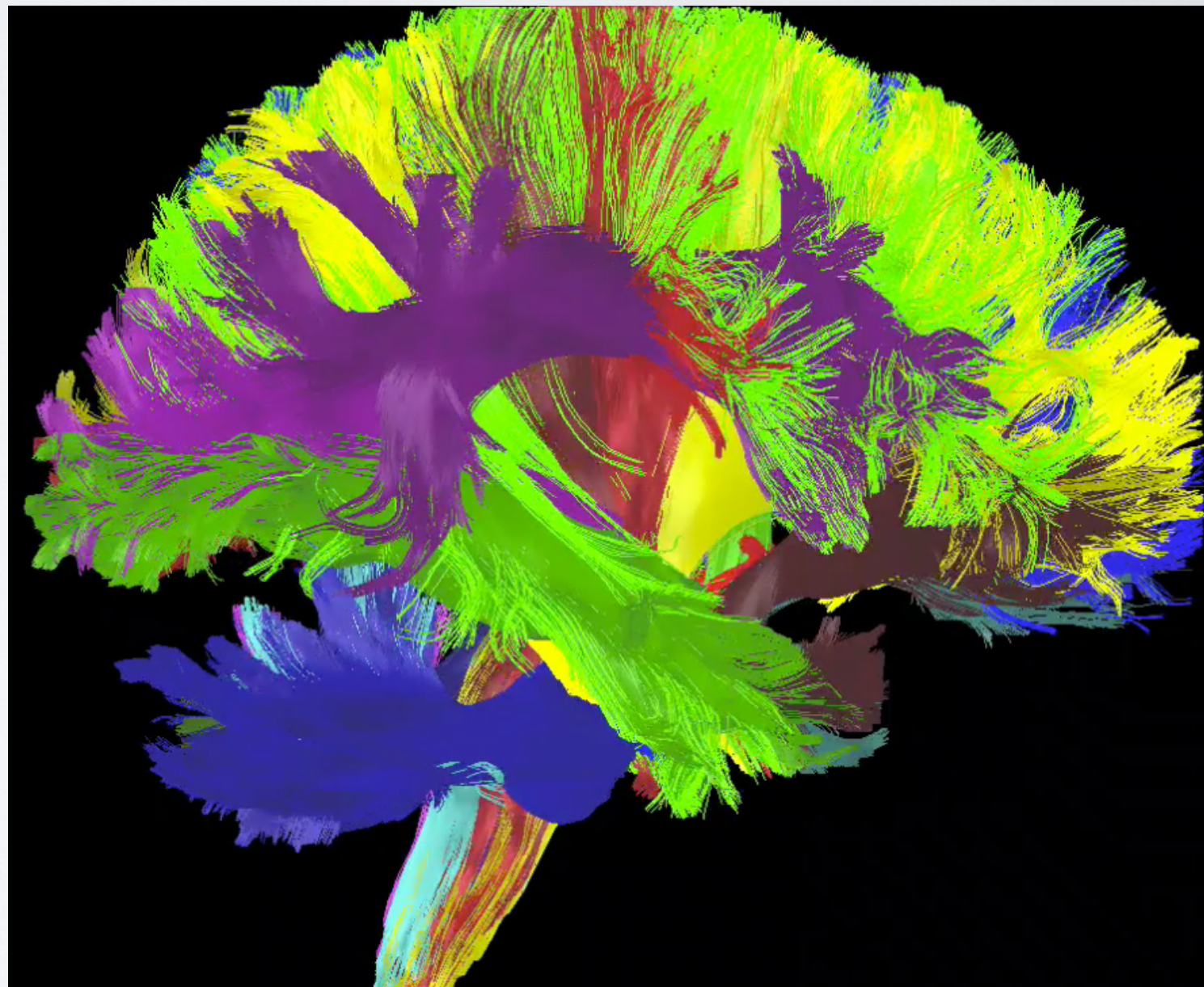
[2] Côté, M.-A., Girard, G., Boré, A., Garyfallidis, E., Houde, J.-C., & Descoteaux, M. (2013). ***Tractometer: Towards validation of tractography pipelines.*** *Medical Image Analysis*, 17(7), 857–844, 2013.

MANUAL TRACT SEGMENTATION

- Fiber tracking algorithm ran on high-resolution 1.25mm isotropic HCP data
- Bundle segmentation manually done by expert radiologist using anatomically placed regions of interest
- 26 bundles extracted covering association, projection and commissural fibers across the whole brain

26 MANUALLY SEGMENTED WHITE MATTER BUNDLES

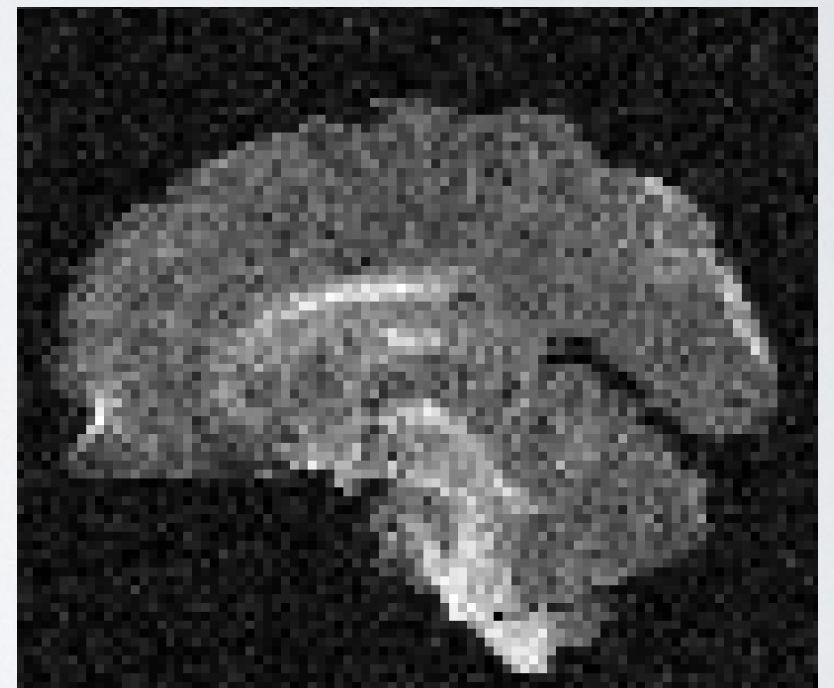
- These 26 bundles serve as ground truth models. They are used as **artificial fibers** to generate the raw diffusion MRI dataset
- This is just another way to generate a phantom dataset based on realistic looking streamline fibers



DIFFUSION MRI SIMULATION

- MOTIVATION: **Clinical-like dataset**

- 2 mm isotropic, 2 $b=0$ images
- 32 directions, b -value 1000 s/mm^2
- Noise added
- Many types of artefacts



Example of simulated $b=1000 \text{ s/mm}^2$
of typical clinical imaging protocol

DATA FOR PARTICIPANTS

1. Diffusion MRI dataset:

Diffusion.nii.gz, Diffusion.bvals, Diffusion.bvecs
(1st image $b=0$ and then 32 dwi)

2. Diffusion with reverse-phase
encoded $b=0$ image

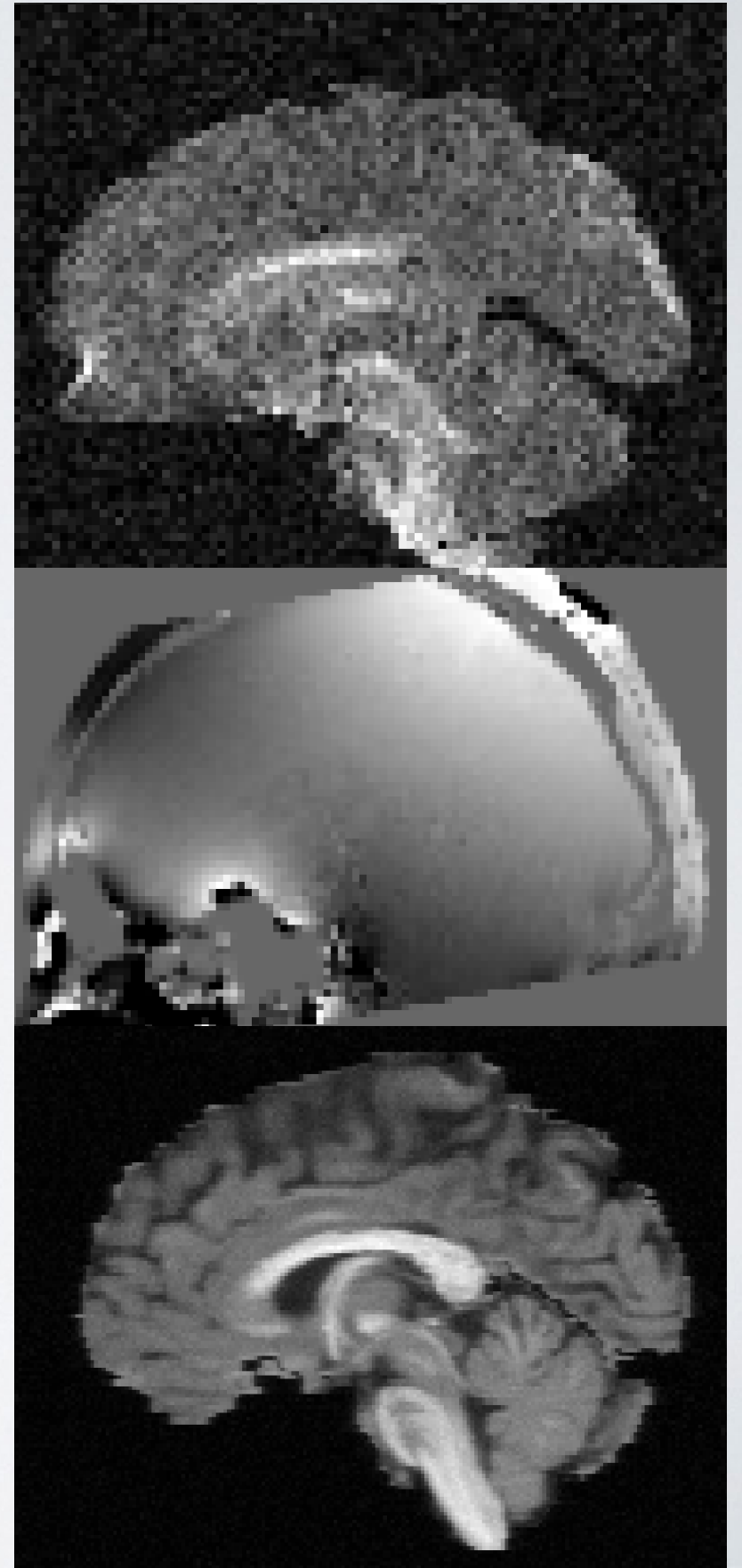
Diffusion_WITH_REVERSEPHASE.nii.gz

3. B0 FieldMap for this dataset:

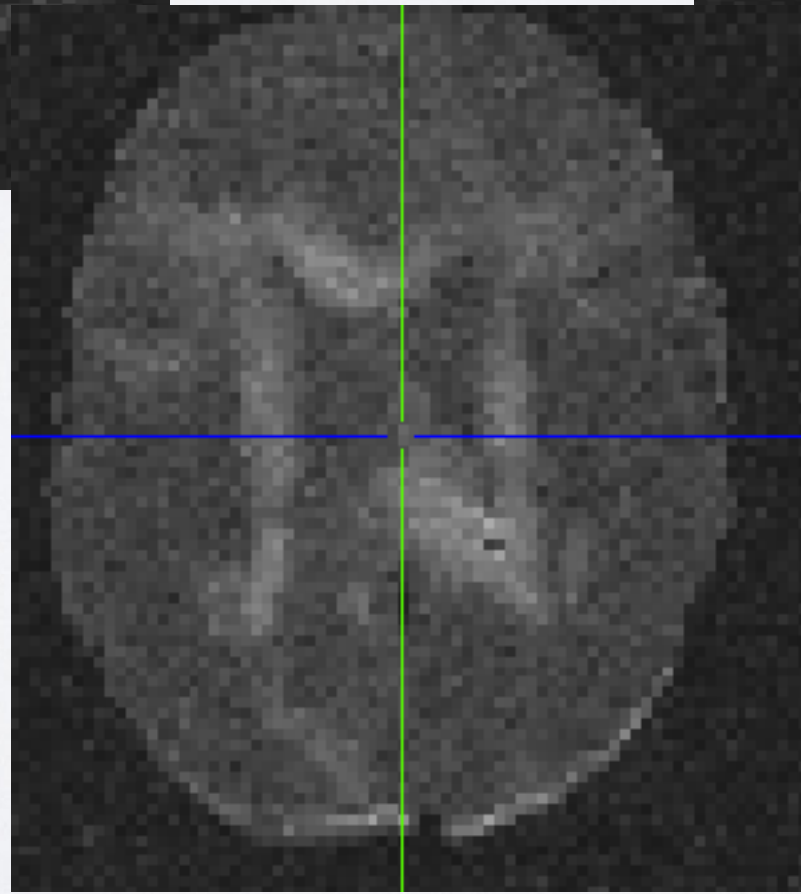
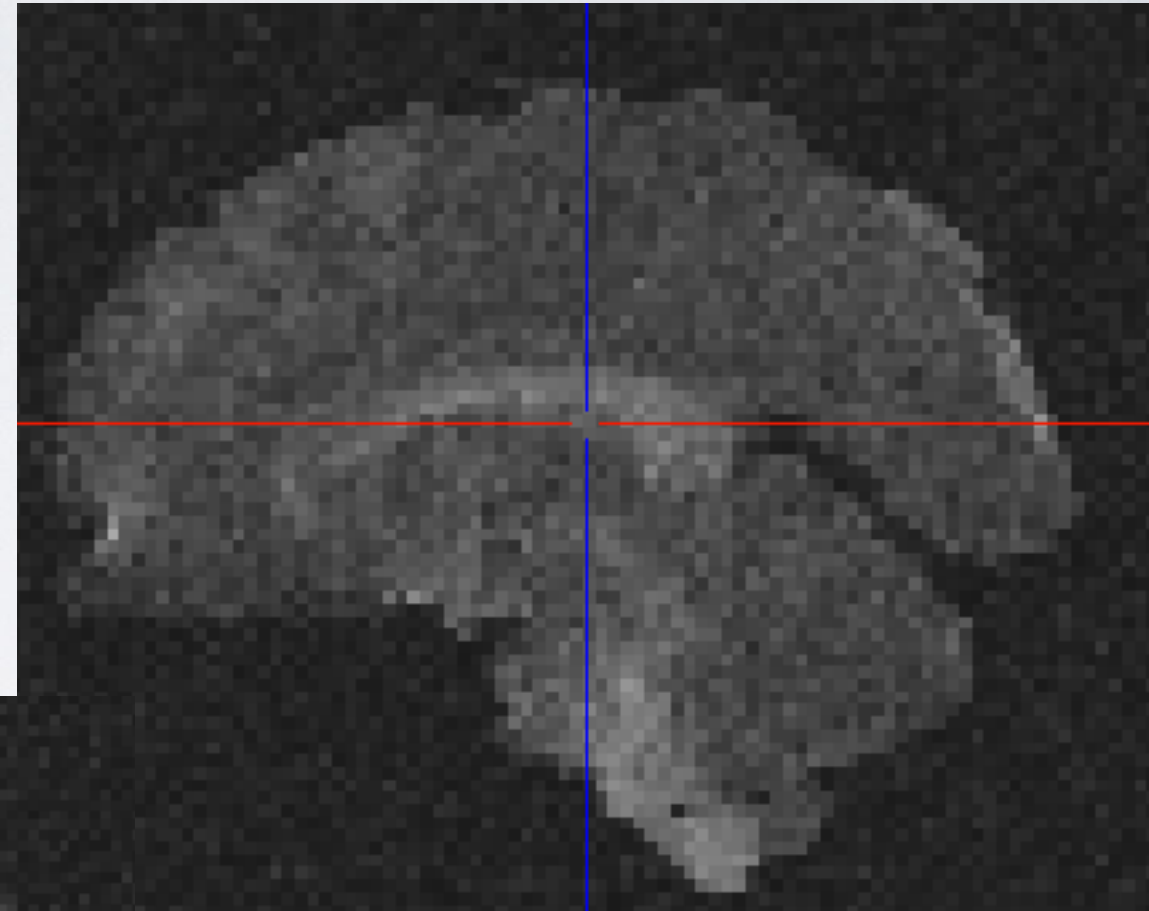
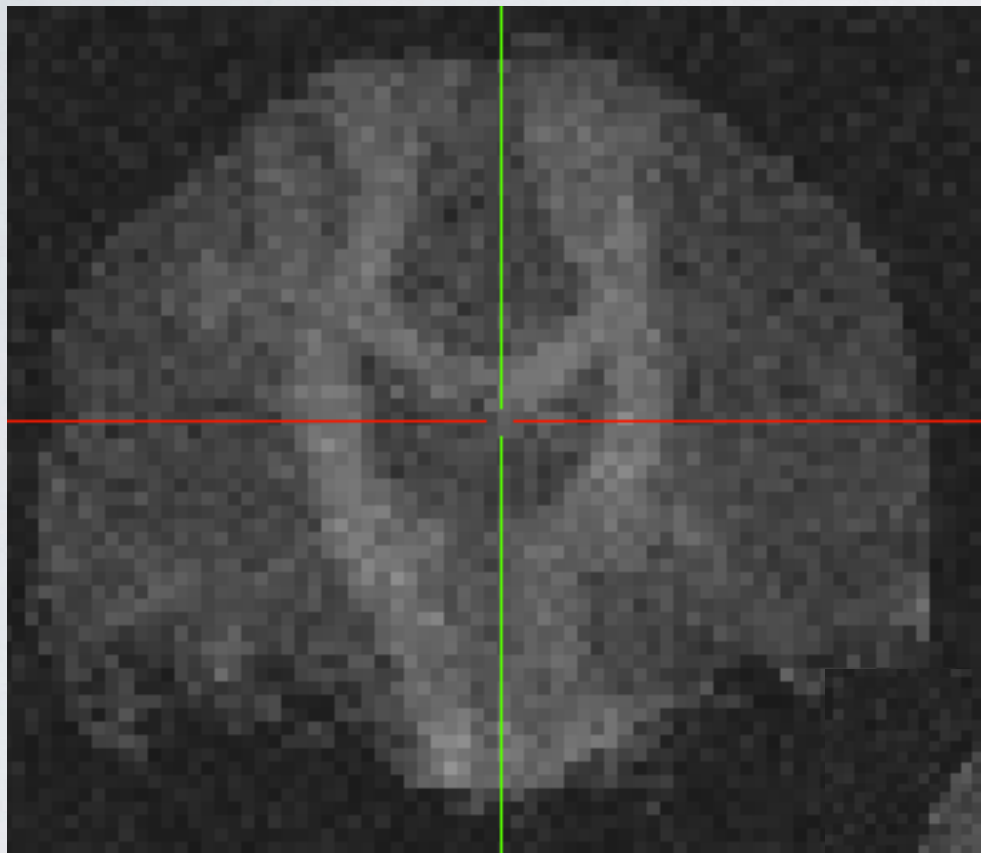
fmap_RadPerSec.nii.gz,
fmap_Hz.nii.gz

4. T1-like image:

T1.nii.gz

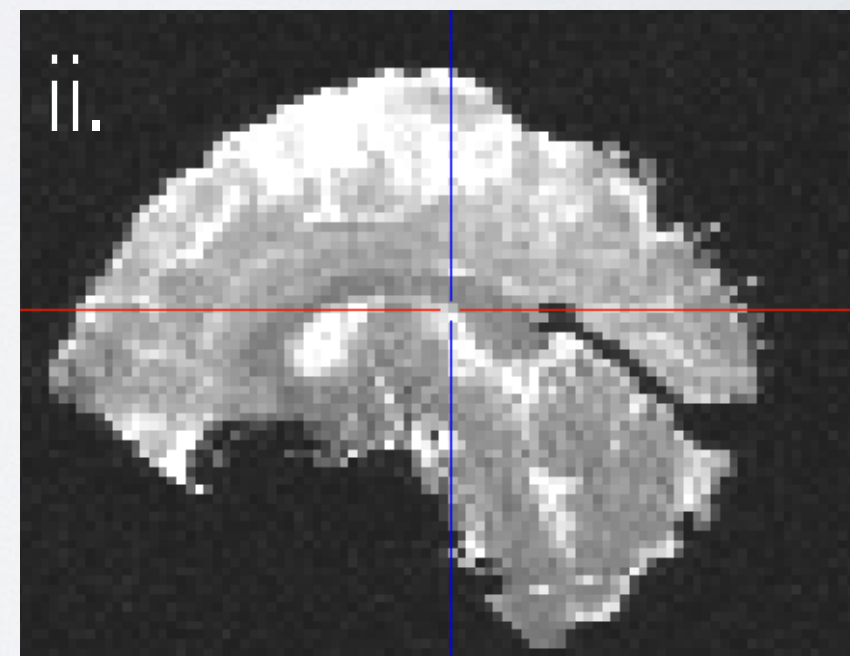
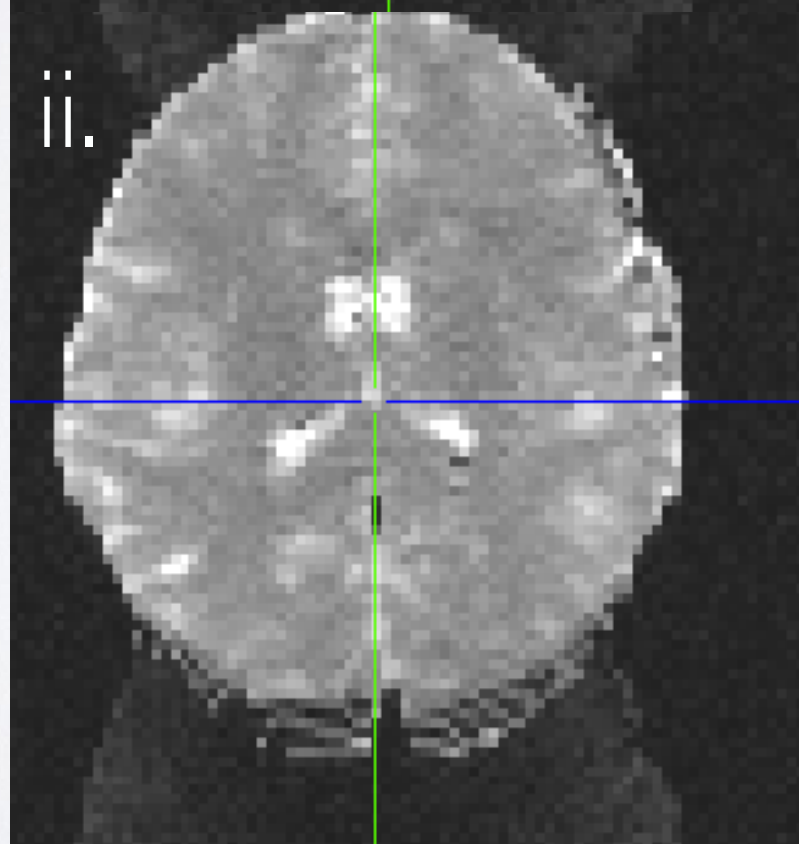
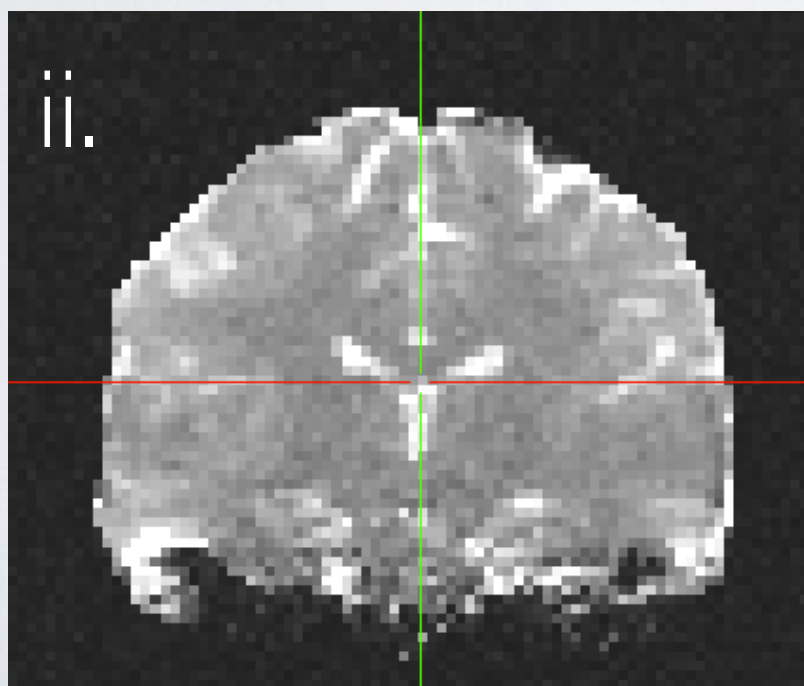
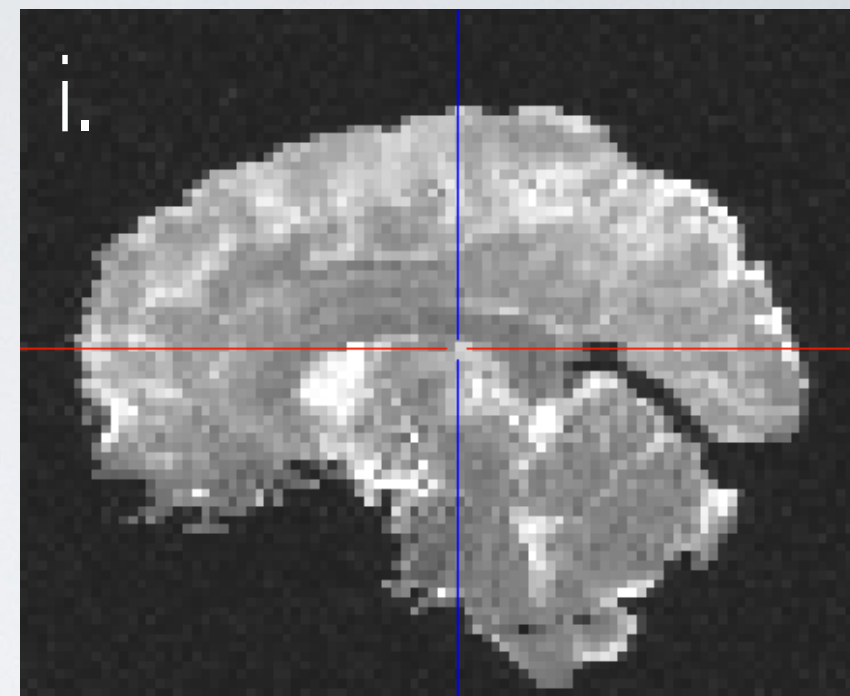
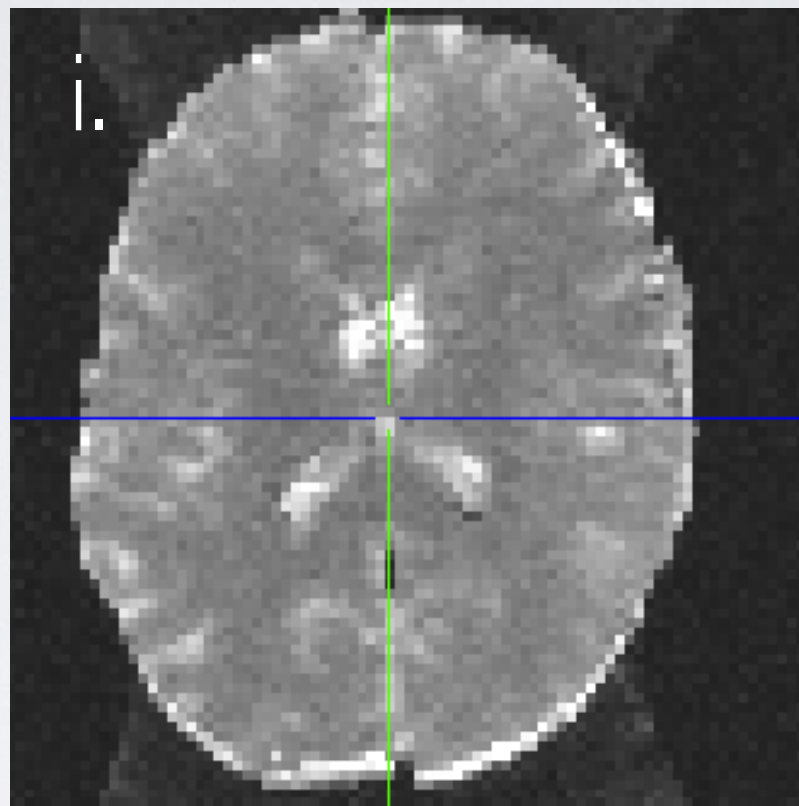
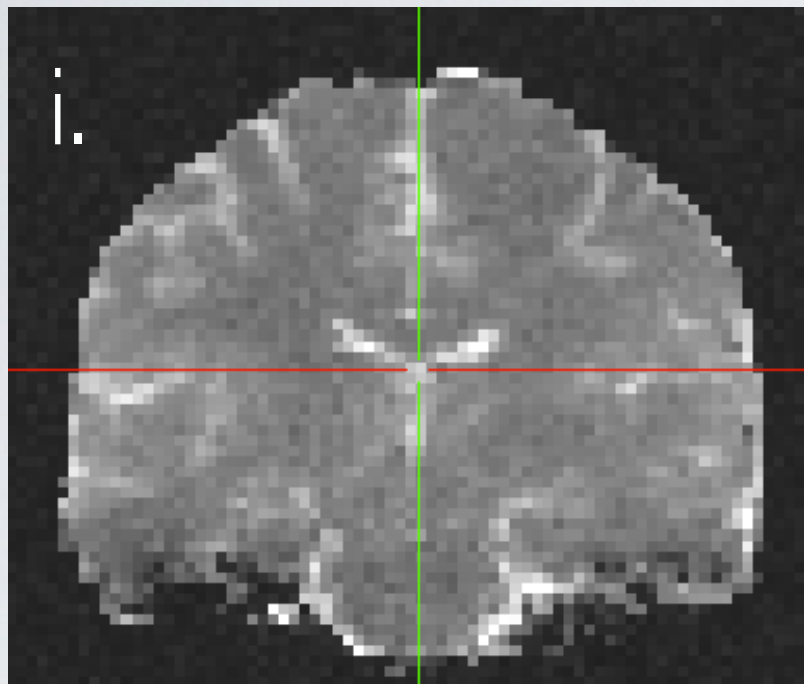


DIFFUSION - DWI #3



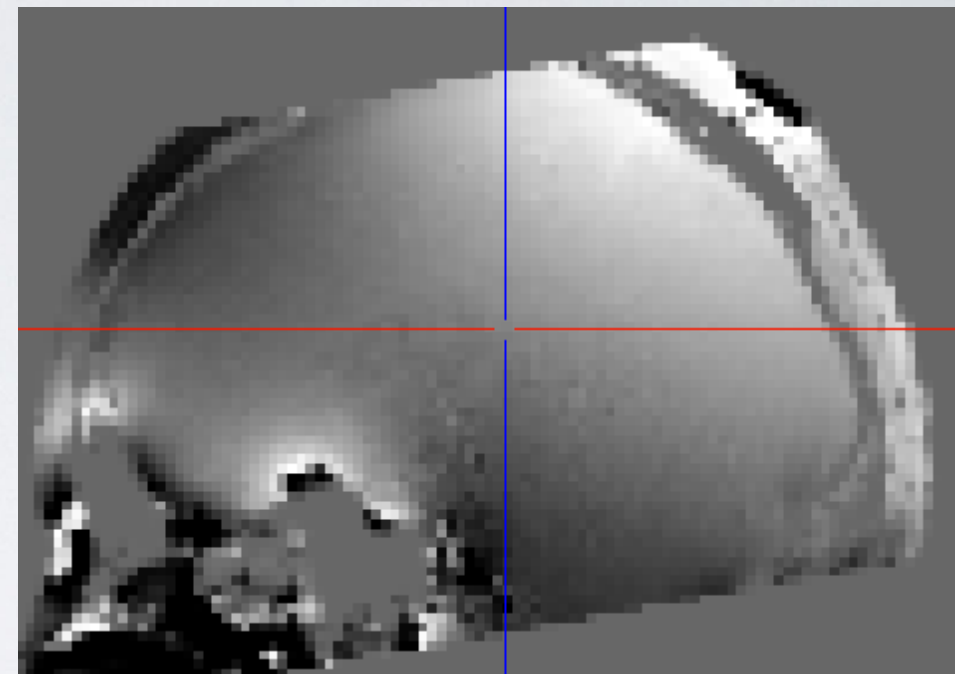
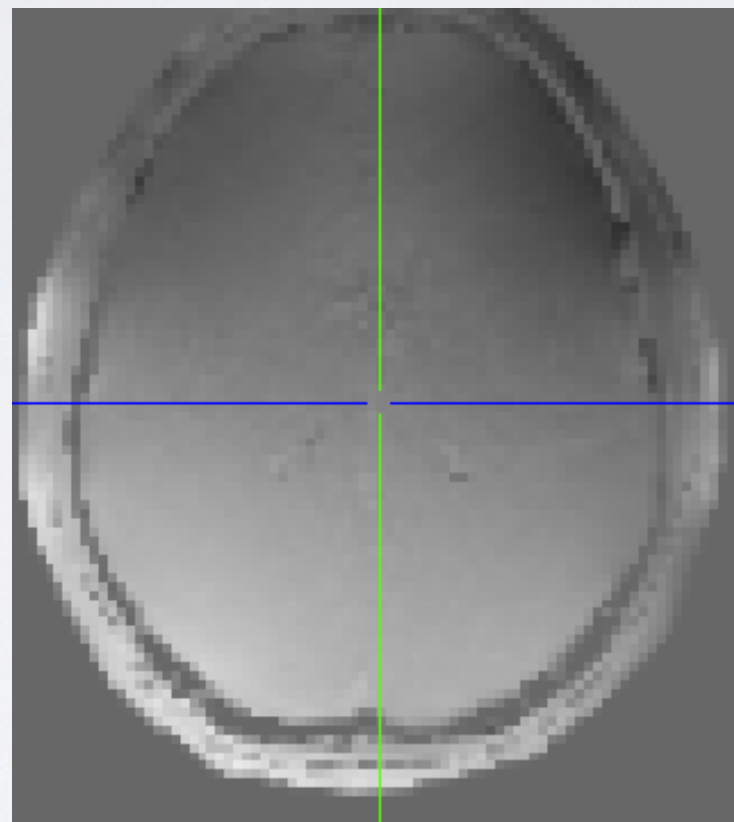
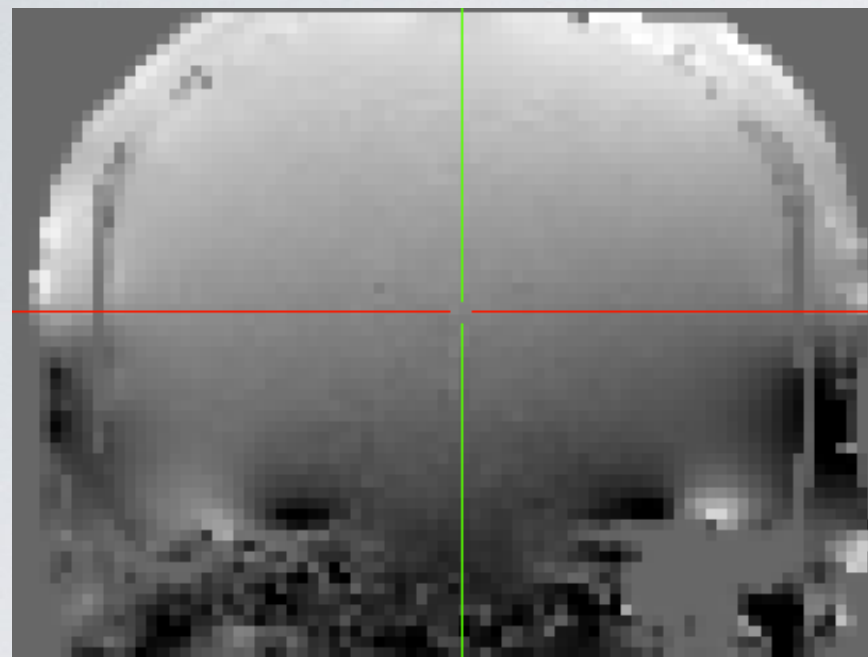
REVERSE PHASE

i. $b=0$ blipup



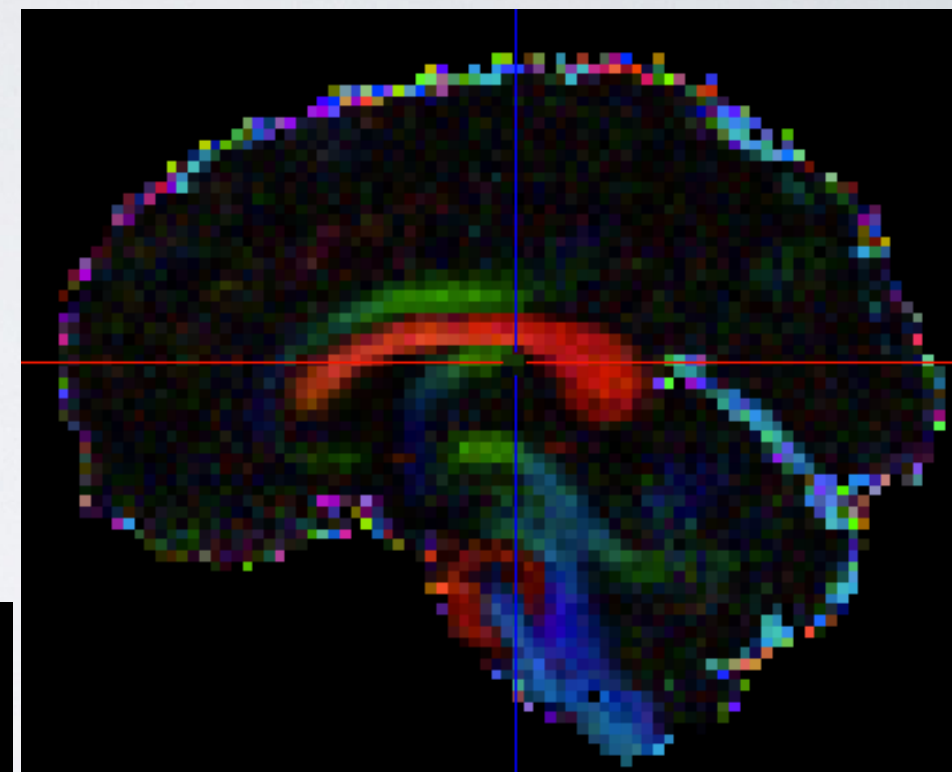
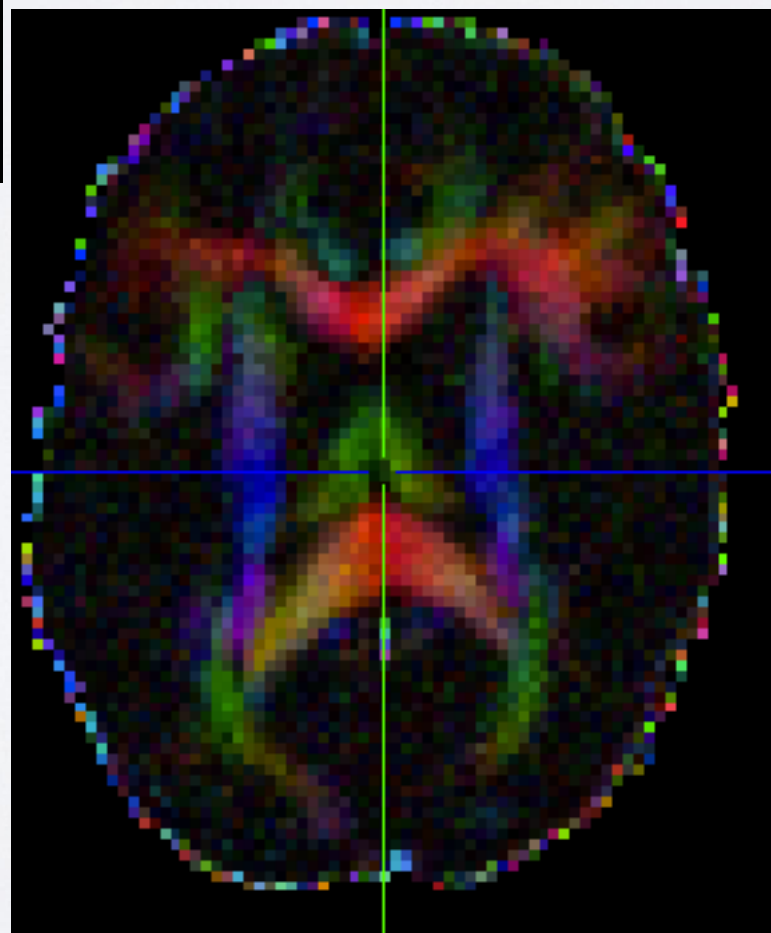
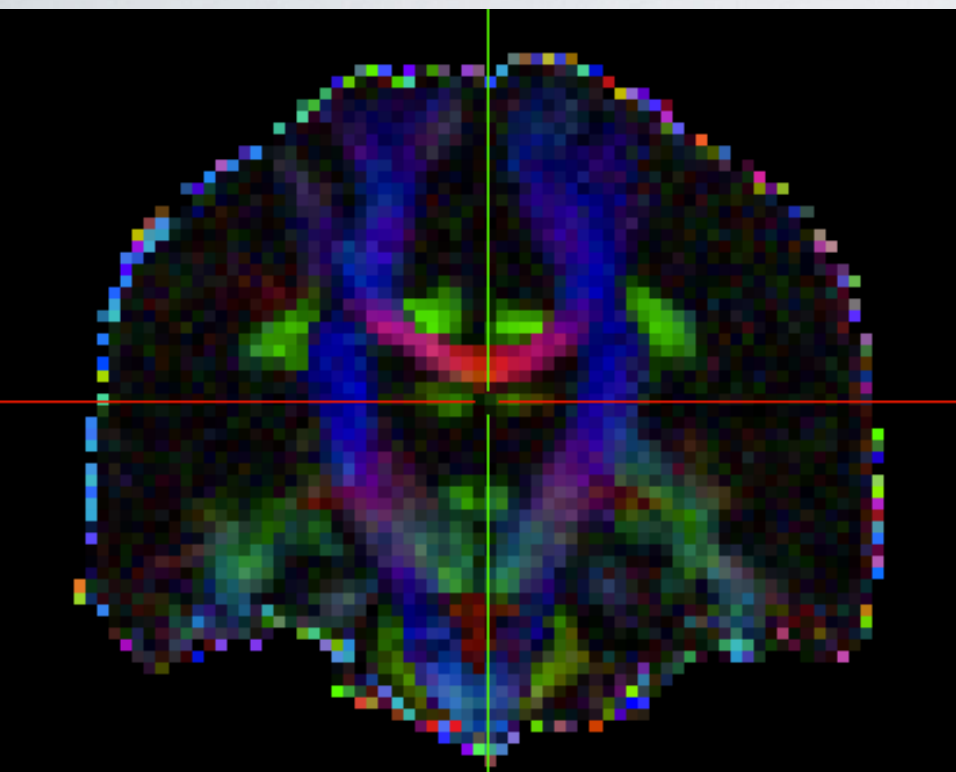
ii. $b=0$ blipdown

B0 FIELD MAP



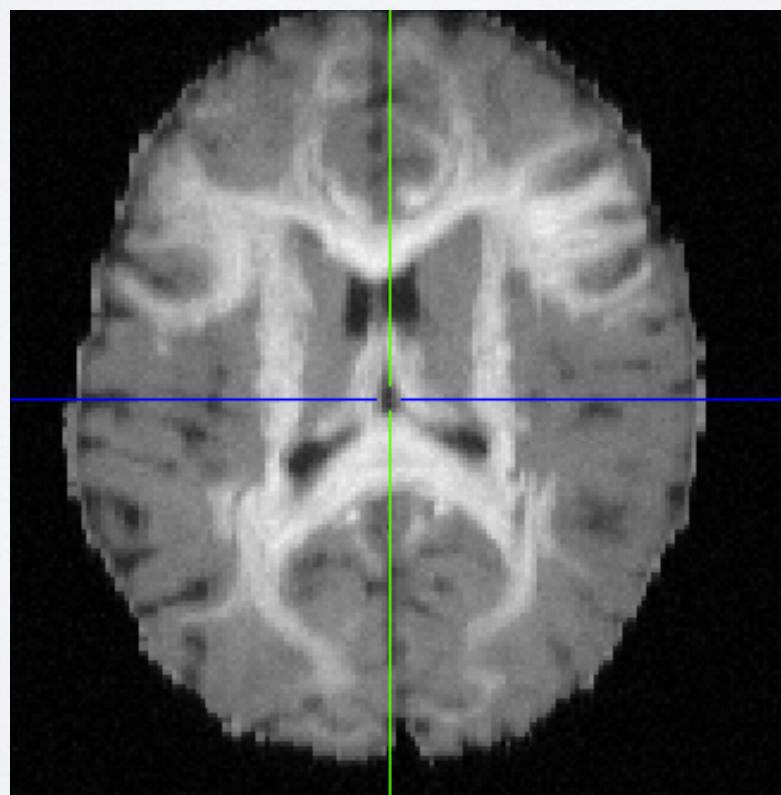
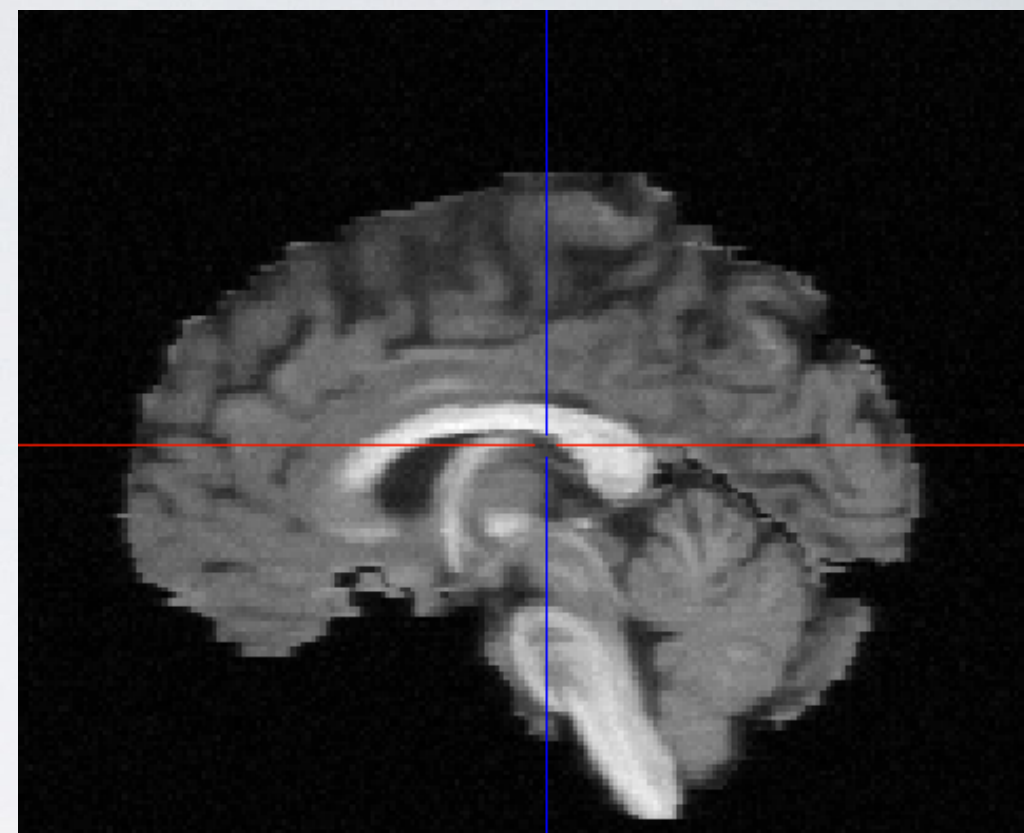
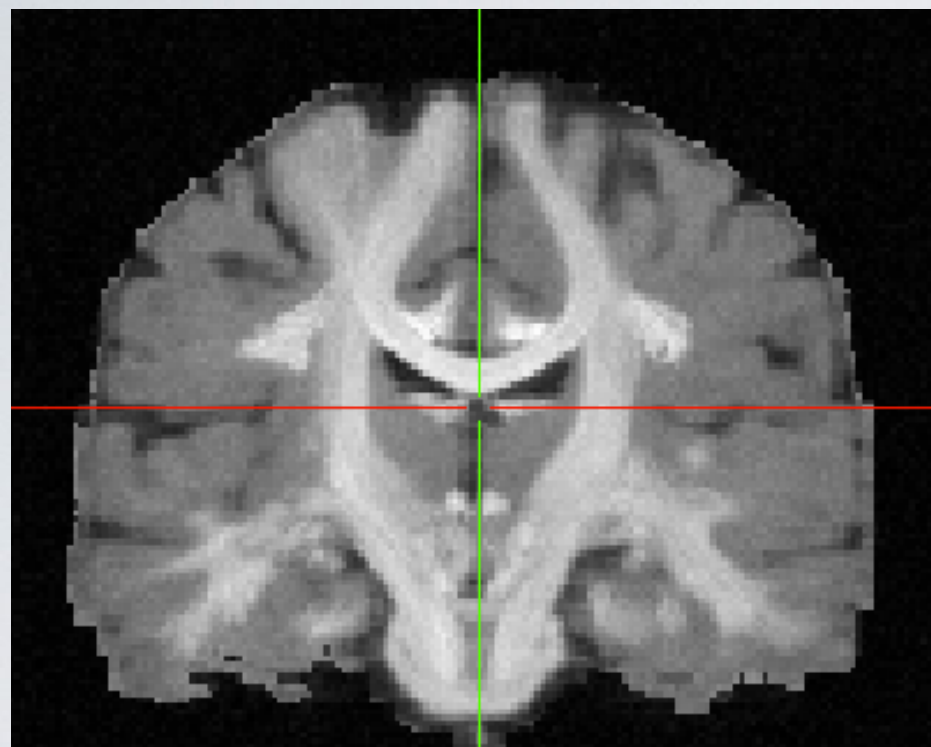
- The same field map is applied to all image volumes, regardless of head motion
- Field map available in rad/sec or Hz

DIFFUSION MRI



Basic
DTI reconstruction
(no correction)

T1



EXPECTED SUBMISSION

- A single whole brain tractogram (max 2Gb). Accepted format:
 - .trk (TractVis)
 - .tck (MRtrix)
 - .vtk
- Challenge website will gives further indications for fileFormat conversion

EVALUATION WITH THE *TRACTOMETER*

- Valid connections
- Invalid connections
- No connections
(stopping in white matter or ventricles)
- Overlap with ground truth streamline fibers
- ***Tractometer***. [Coté et al Descoteaux. MEDIA 2013]



VALID CONNECTION

Connects 2 connected regions

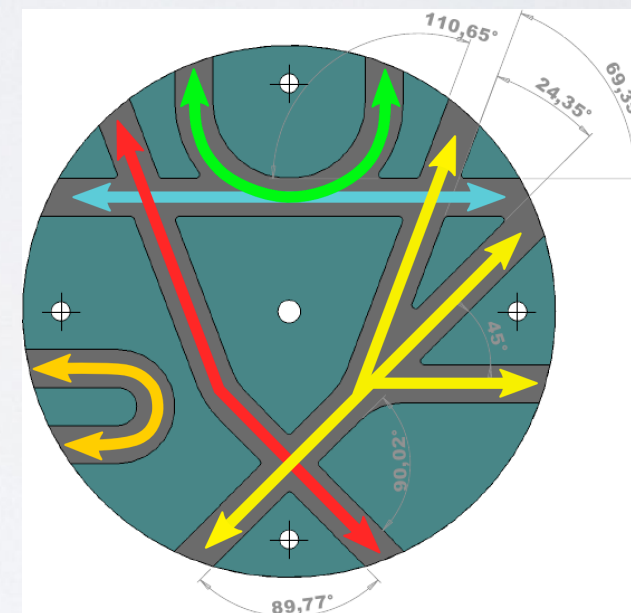
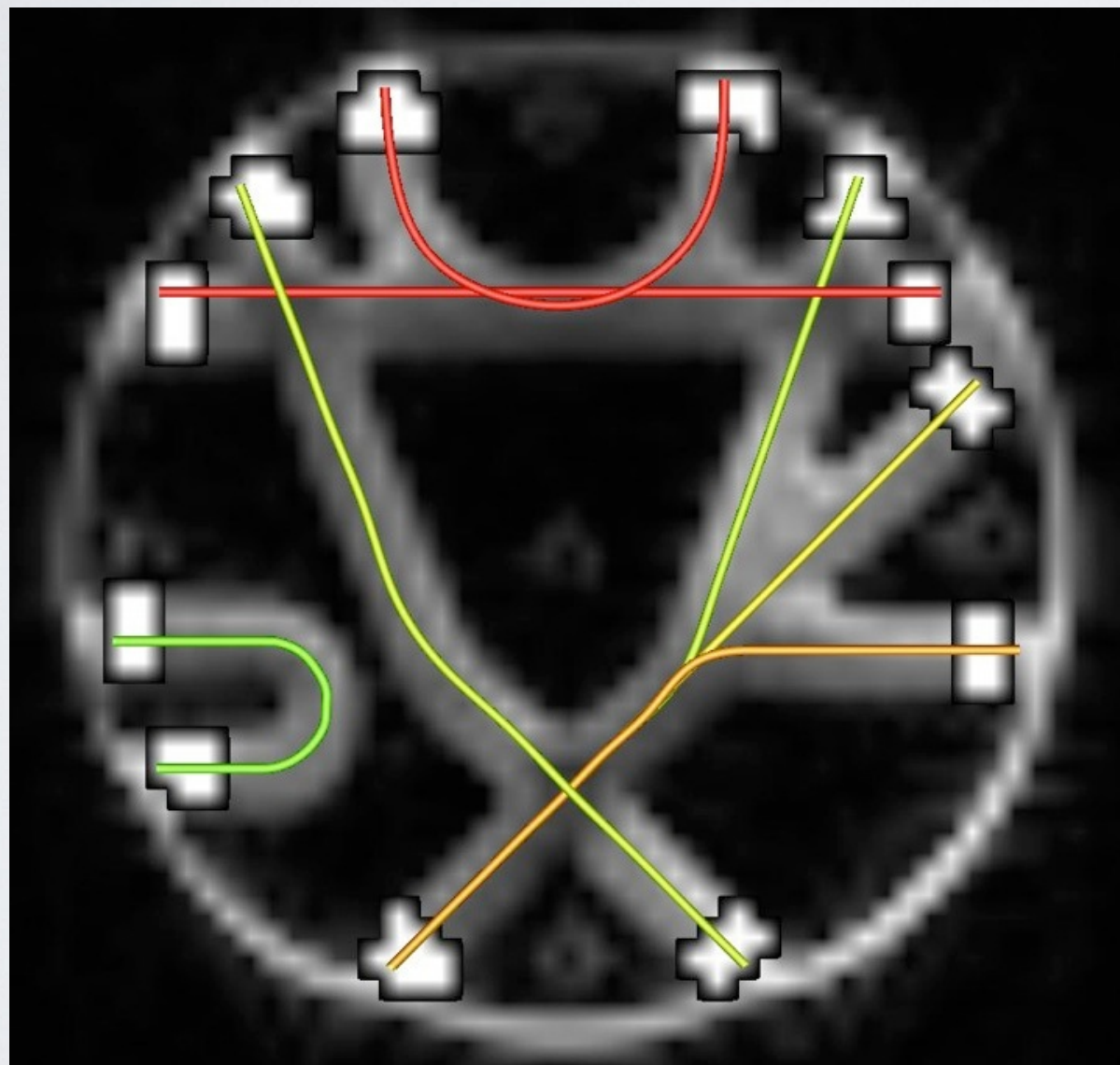
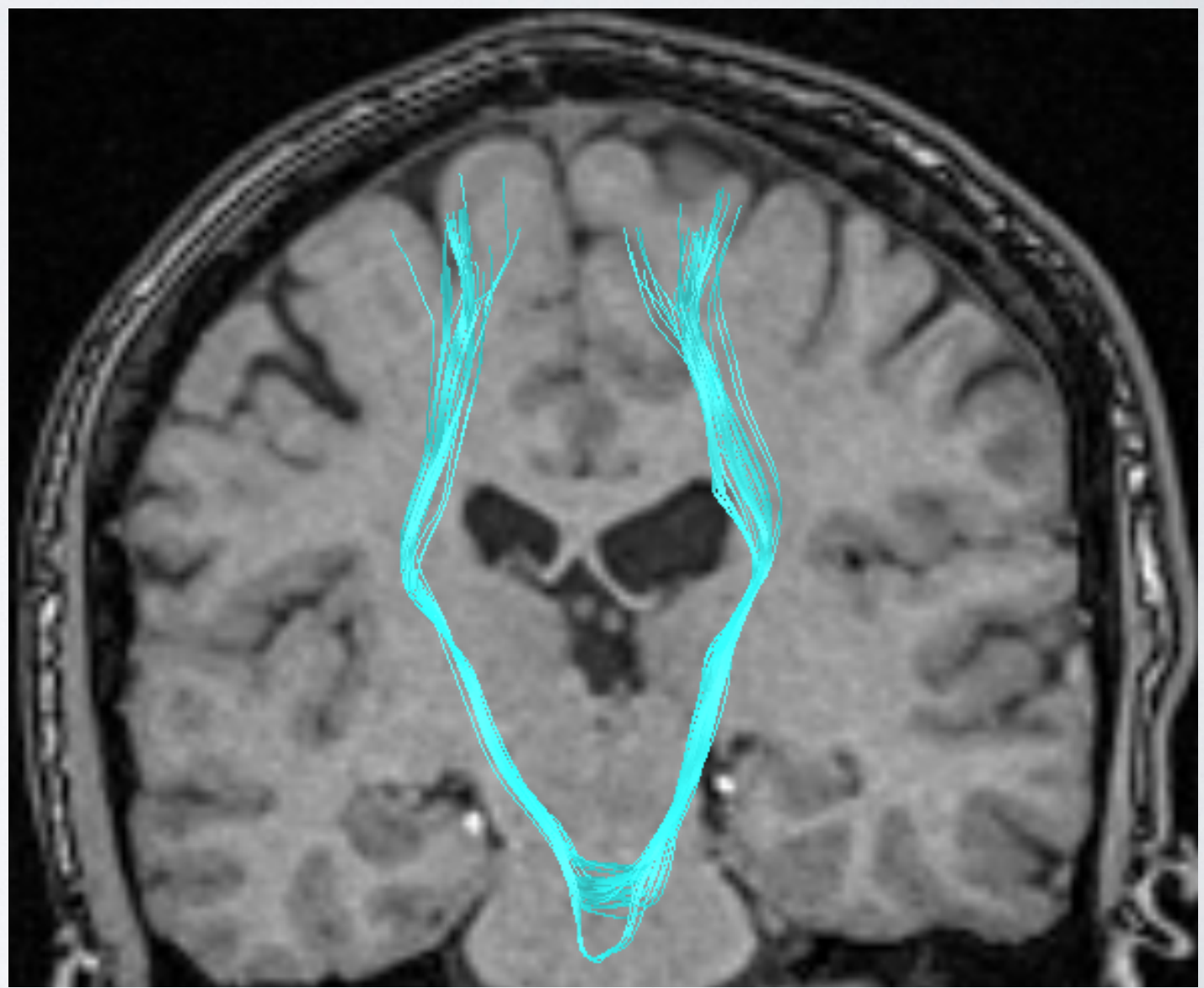
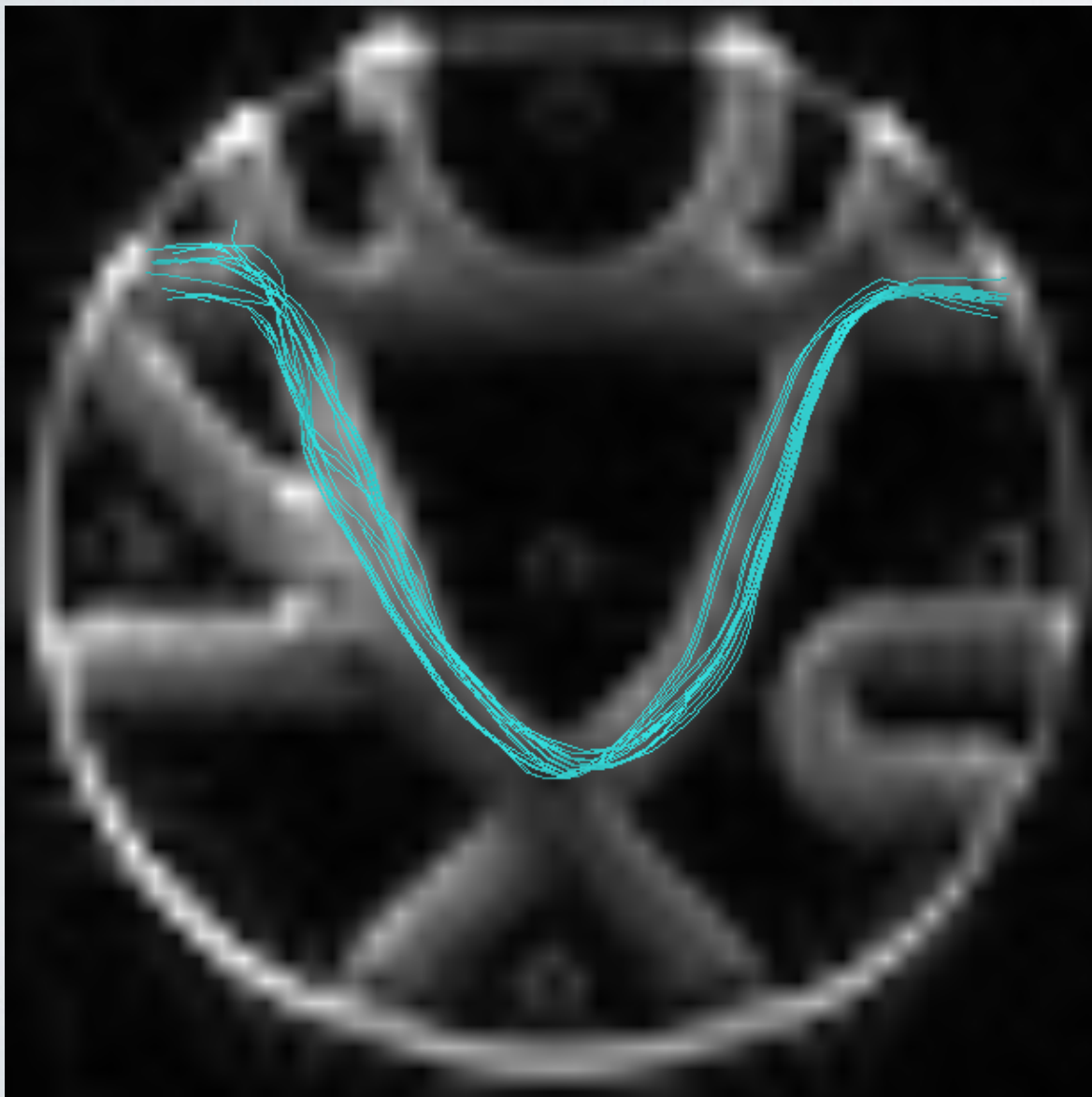


Illustration with the *FiberCup* dataset
MICCAI Challenge 2009. [Fillard, Descoteaux et al NeuroImage 2010]

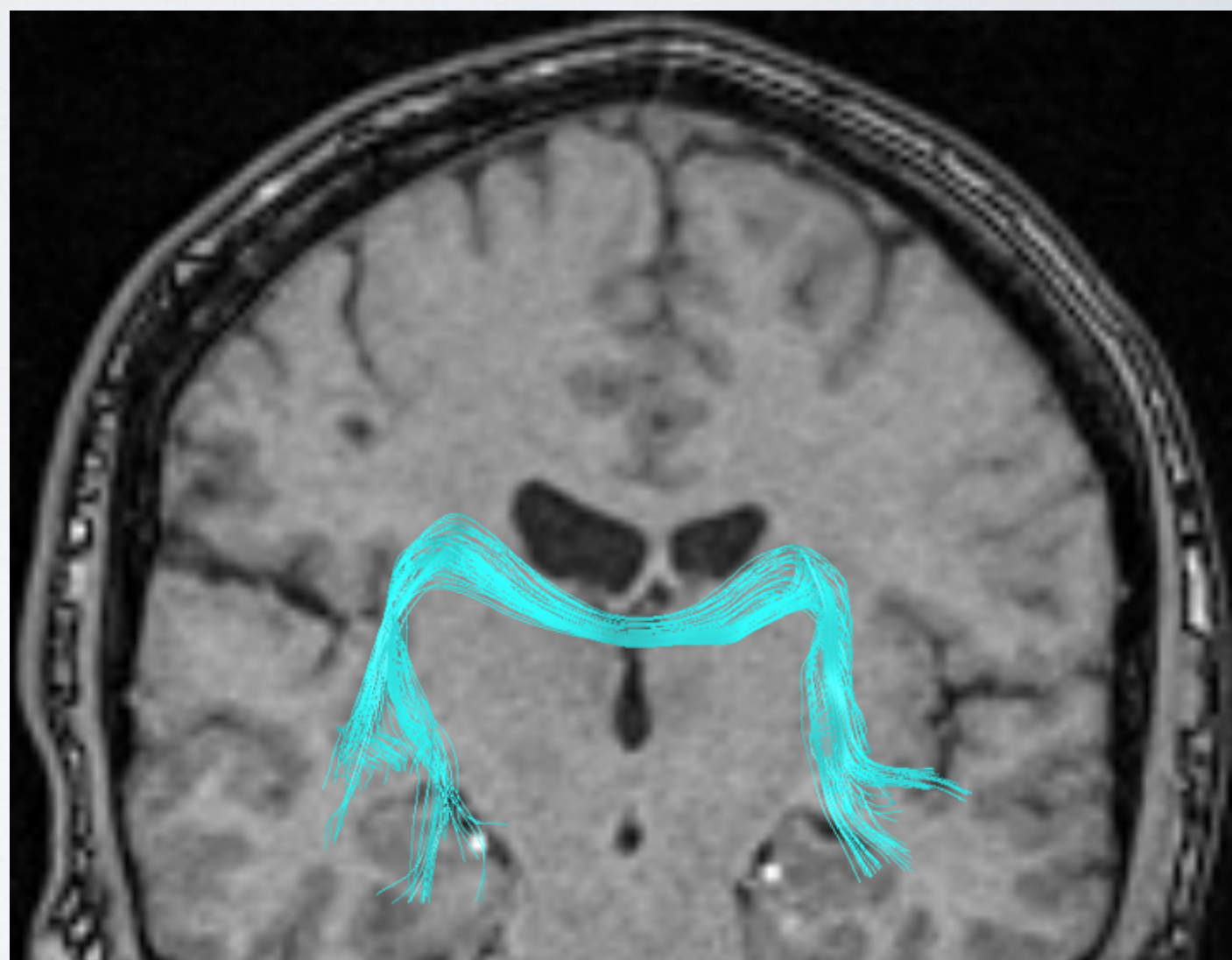
INVALID CONNECTION

Connects 2 valid regions through a wrong path



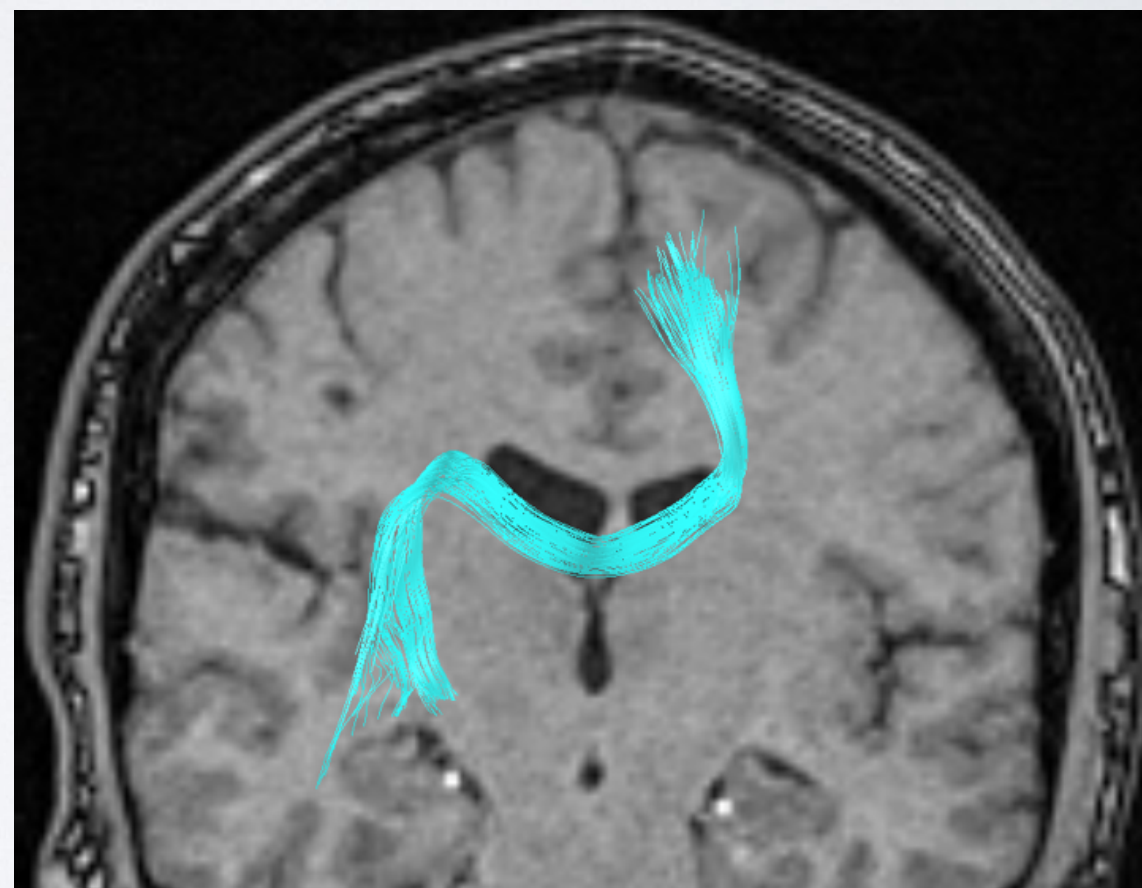
INVALID CONNECTION

Connects 2 regions that should not be connected



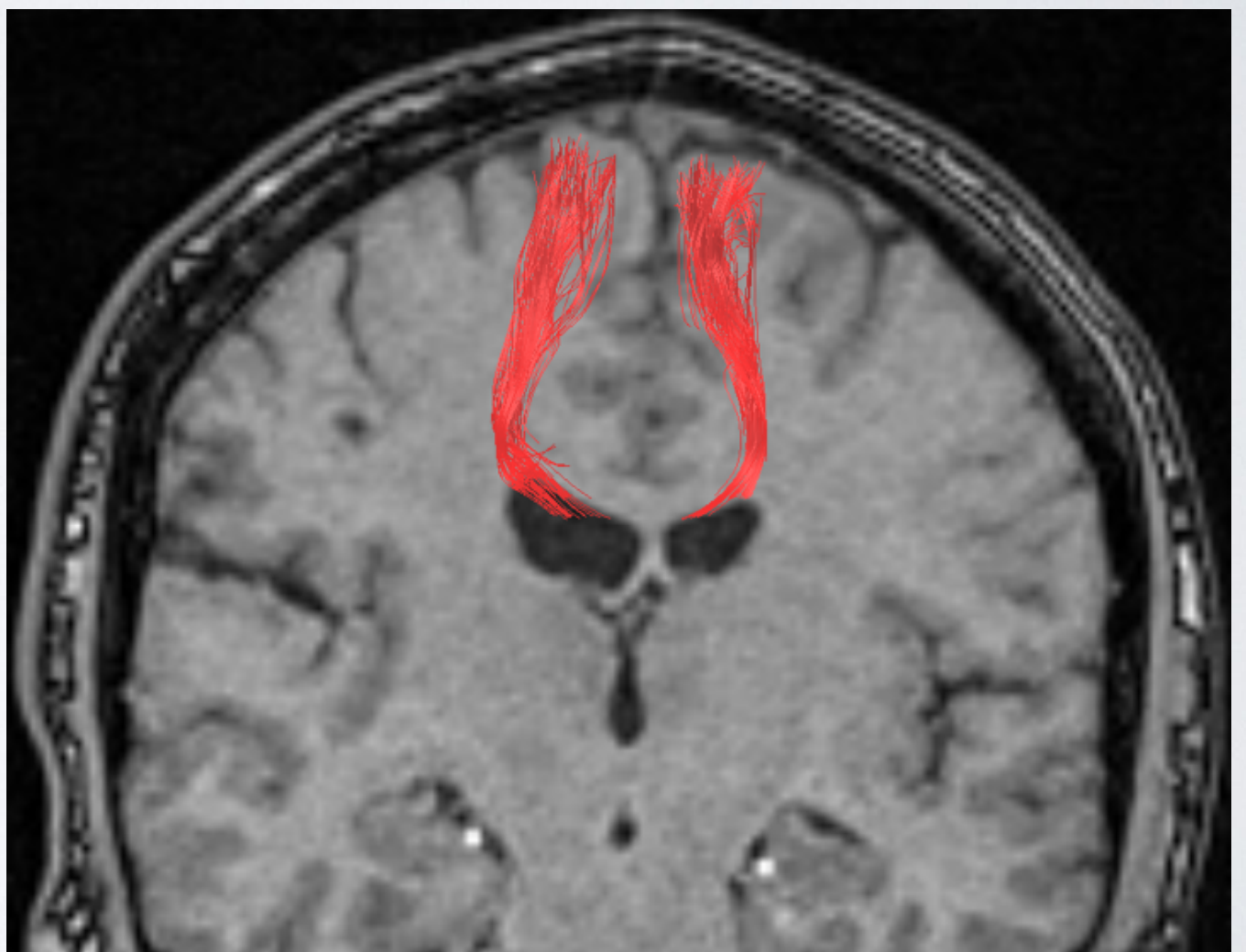
INVALID CONNECTIONS

Connects 2 regions that should not be connected



NO CONNECTION

Does not connect 2 regions
(stops prematurely in white matter or ventricles)



OVERLAP



Ground truth streamlines

Estimated streamlines



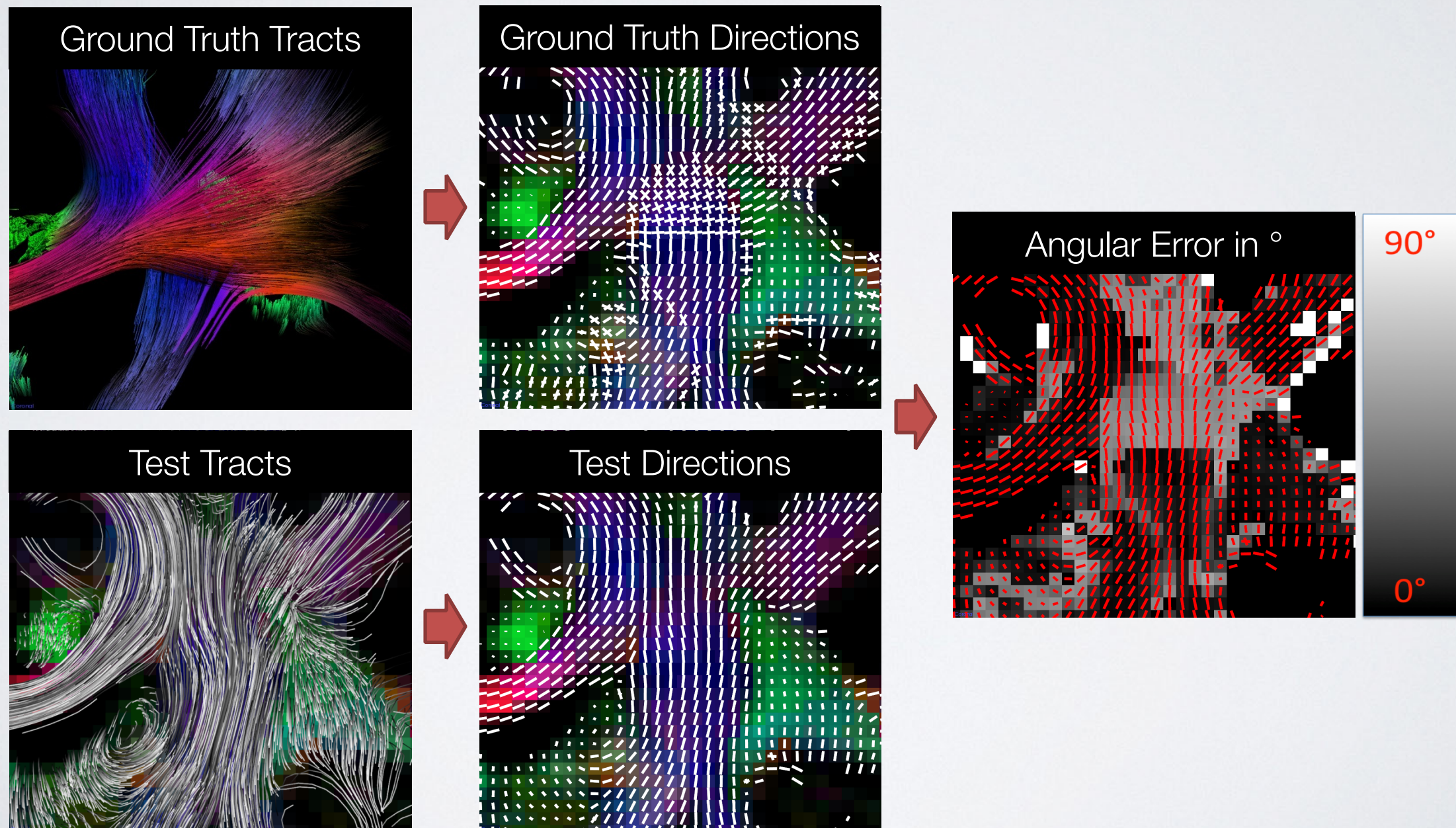
Overlapping voxels

Non-overlapping voxels

(Example on the CST)

LOCAL ANGULAR ERROR

Voxel-wise tractogram directions compared to ground truth directions.



QUESTIONS?