

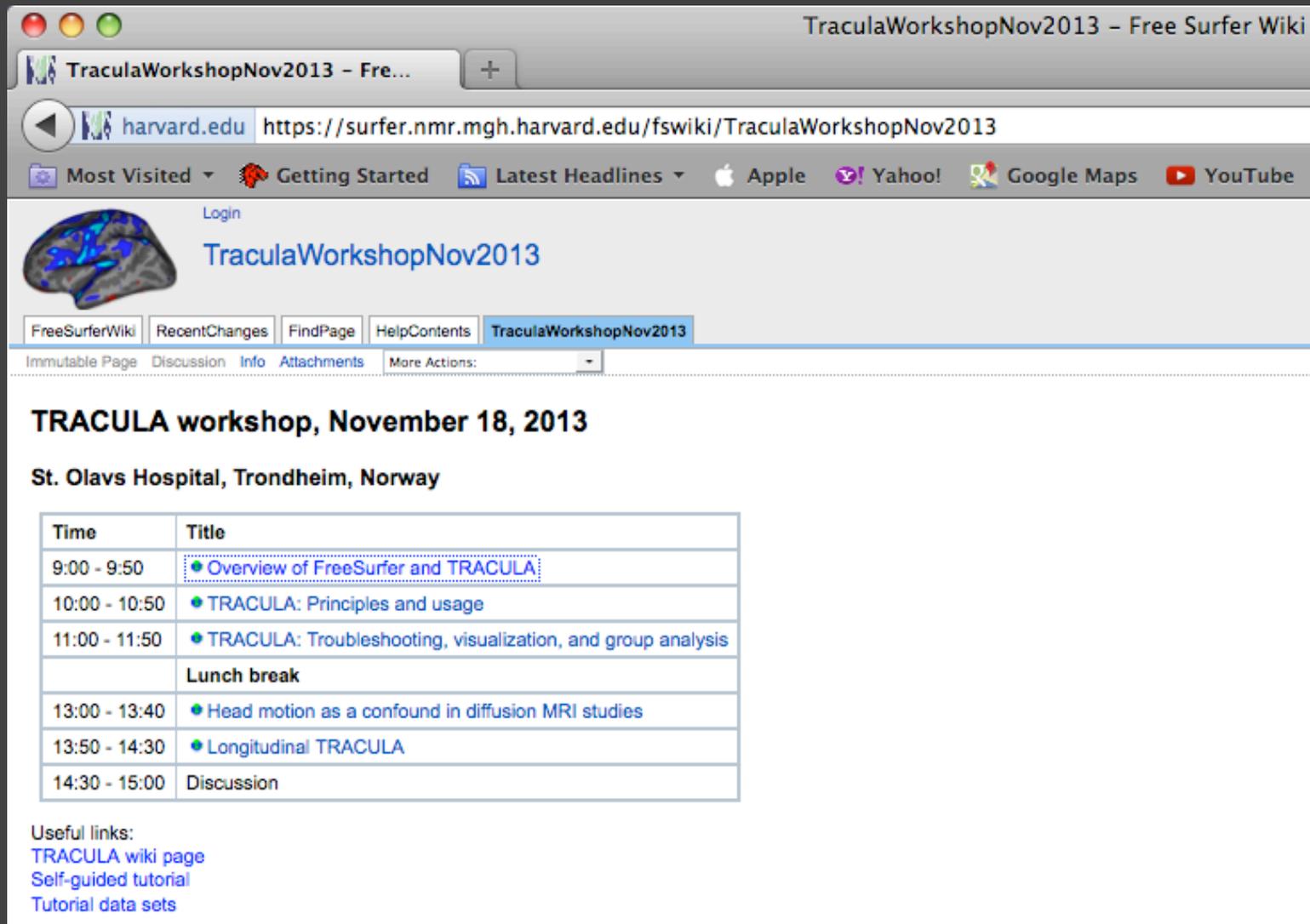
Overview of FreeSurfer and TRACULA

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Biomedical Imaging

Schedule



The screenshot shows a web browser window with the title "TraculaWorkshopNov2013 - Free Surfer Wiki". The address bar shows the URL "https://surfer.nmr.mgh.harvard.edu/fswiki/TraculaWorkshopNov2013". The page content includes a "Login" button, a brain image, and the title "TraculaWorkshopNov2013". Below the title is a navigation menu with "FreeSurferWiki", "RecentChanges", "FindPage", "HelpContents", and "TraculaWorkshopNov2013". The main content area features the heading "TRACULA workshop, November 18, 2013" and the location "St. Olavs Hospital, Trondheim, Norway". A table lists the workshop schedule, and there are "Useful links" at the bottom.

TraculaWorkshopNov2013 - Free Surfer Wiki

TraculaWorkshopNov2013 - Fre...

harvard.edu https://surfer.nmr.mgh.harvard.edu/fswiki/TraculaWorkshopNov2013

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Immutable Page Discussion Info Attachments More Actions: -

TRACULA workshop, November 18, 2013

St. Olavs Hospital, Trondheim, Norway

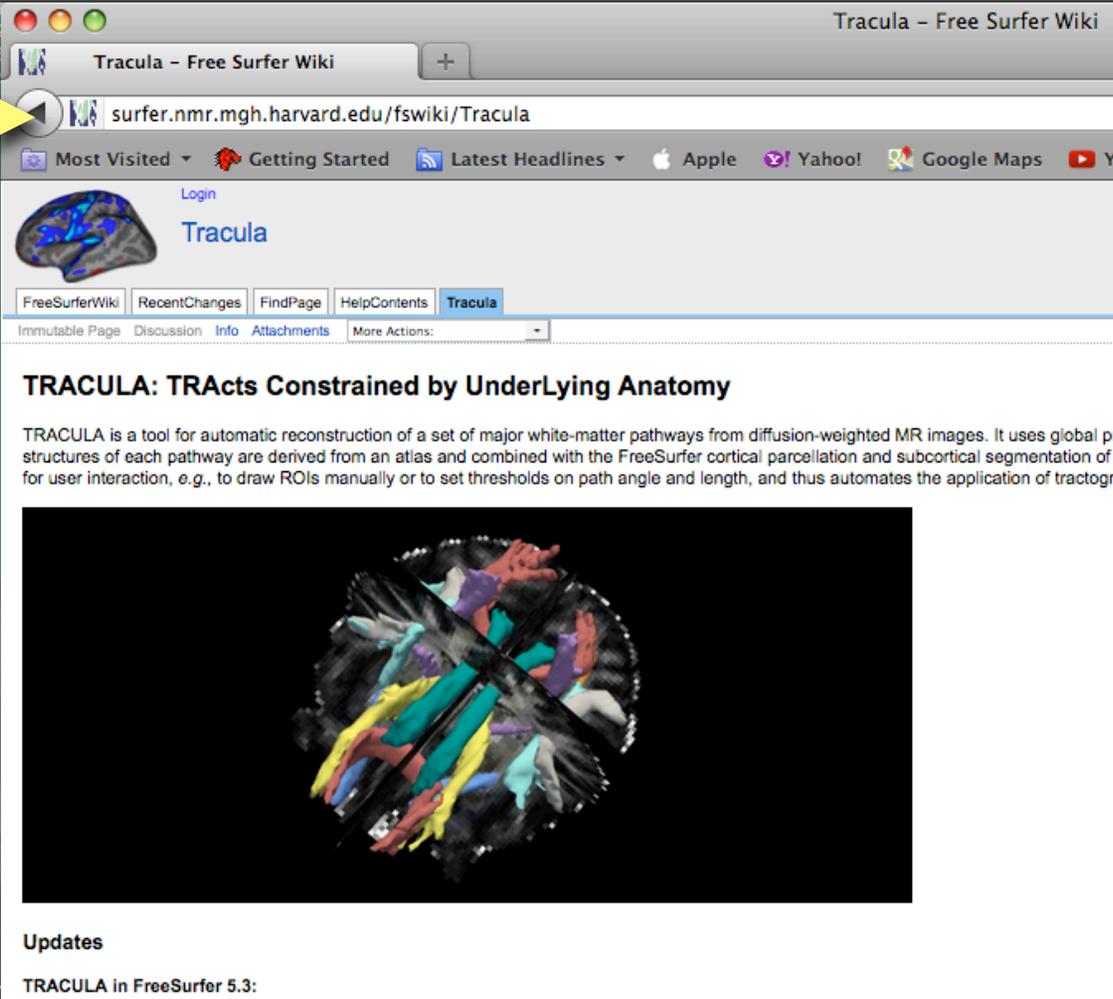
Time	Title
9:00 - 9:50	Overview of FreeSurfer and TRACULA
10:00 - 10:50	TRACULA: Principles and usage
11:00 - 11:50	TRACULA: Troubleshooting, visualization, and group analysis
	Lunch break
13:00 - 13:40	Head motion as a confound in diffusion MRI studies
13:50 - 14:30	Longitudinal TRACULA
14:30 - 15:00	Discussion

Useful links:
[TRACULA wiki page](#)
[Self-guided tutorial](#)
[Tutorial data sets](#)

Documentation and support

Wiki page: →

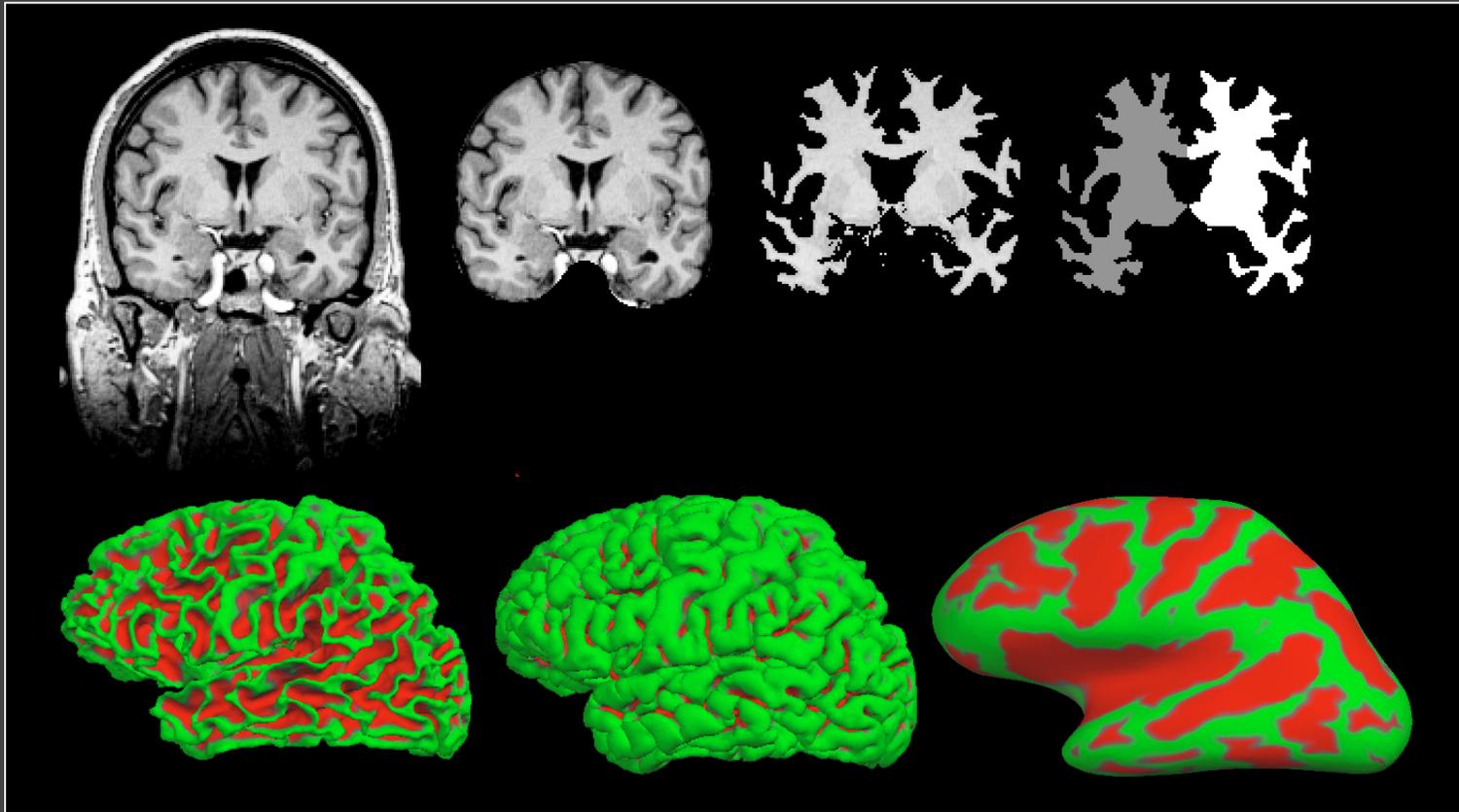
- Usage info
- Updates
- Tutorials



The screenshot shows a web browser window titled "Tracula - Free Surfer Wiki". The address bar displays "surfer.nmr.mgh.harvard.edu/fswiki/Tracula". The page content includes a "Login" button, a "Tracula" header with a brain image, and a navigation menu with options like "FreeSurferWiki", "RecentChanges", "FindPage", "HelpContents", and "Tracula". Below the navigation menu, there are links for "Immutable Page", "Discussion", "Info", "Attachments", and "More Actions". The main heading is "TRACULA: TRActs Constrained by UnderLYing Anatomy". The text below the heading describes the tool: "TRACULA is a tool for automatic reconstruction of a set of major white-matter pathways from diffusion-weighted MR images. It uses global priors and the structures of each pathway are derived from an atlas and combined with the FreeSurfer cortical parcellation and subcortical segmentation of the input image for user interaction, e.g., to draw ROIs manually or to set thresholds on path angle and length, and thus automates the application of tractography." Below the text is a 3D visualization of white-matter tracts in various colors (red, yellow, green, blue, purple) overlaid on a brain slice. At the bottom, there is an "Updates" section with the text "TRACULA in FreeSurfer 5.3:".

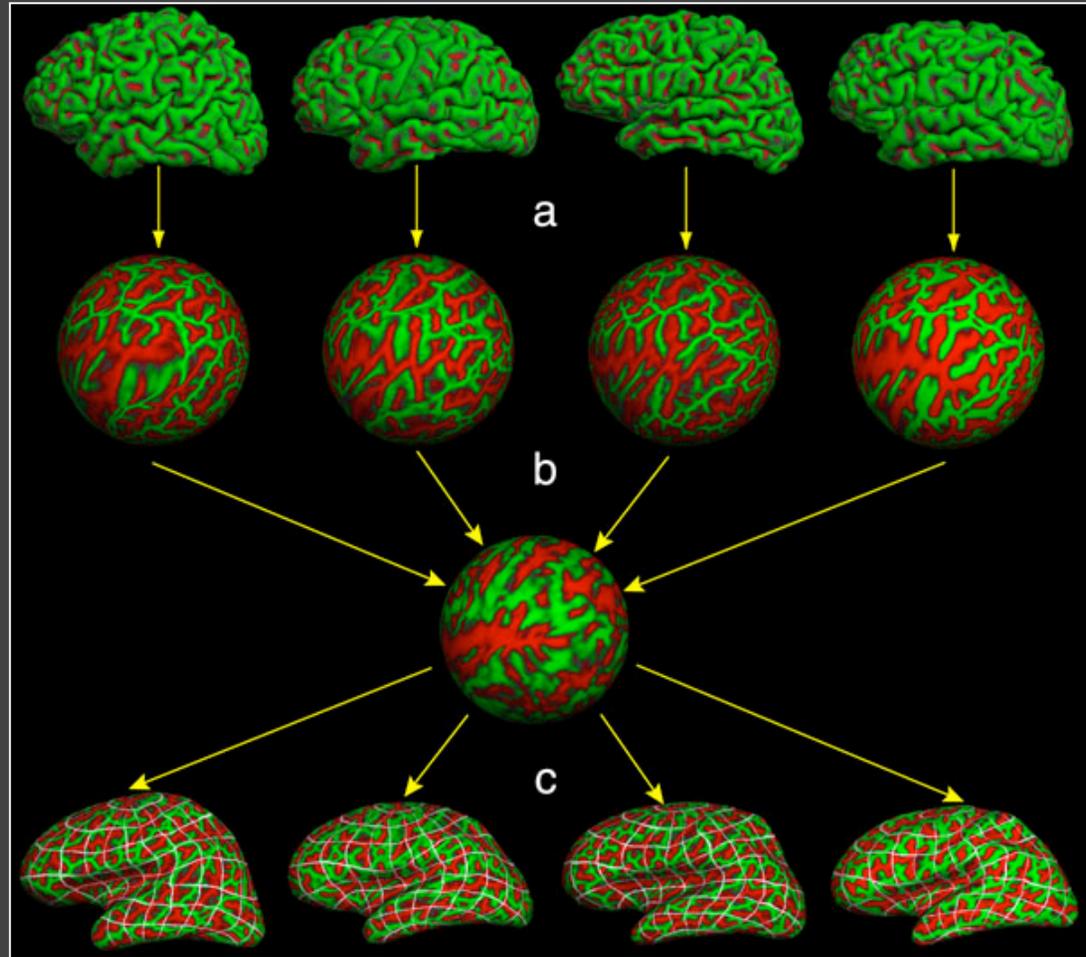
Questions: freesurfer@nmr.mgh.harvard.edu

FreeSurfer



Automated white/grey matter segmentation and surface reconstruction
[Dale 1999, Fischl 1999]

Surface-based coordinate system

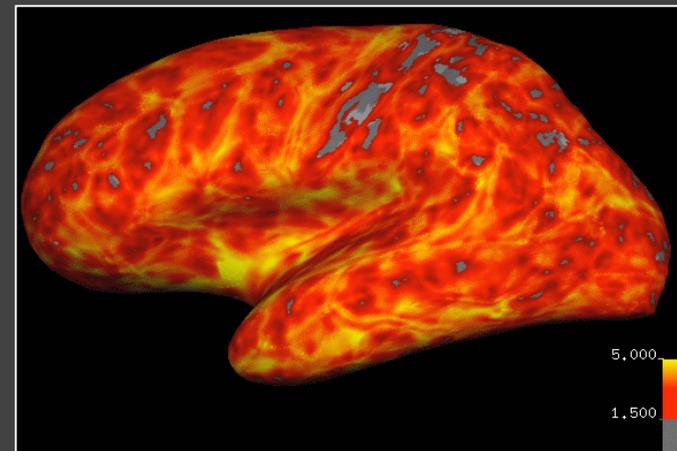


- Align individual brains based on gyral/sulcal patterns [Fischl 1999]

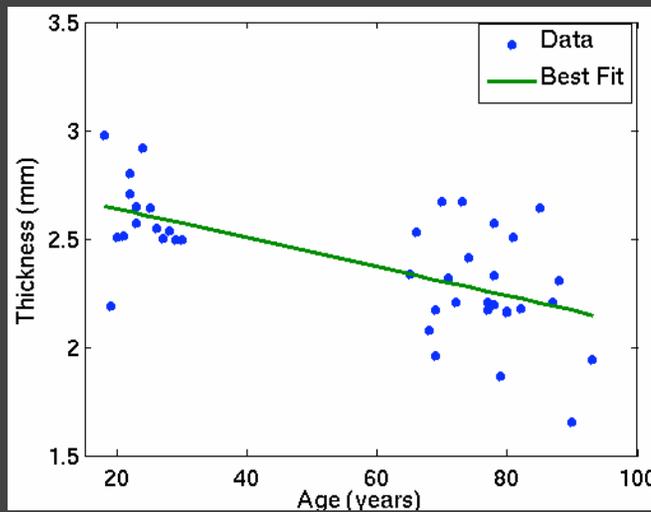
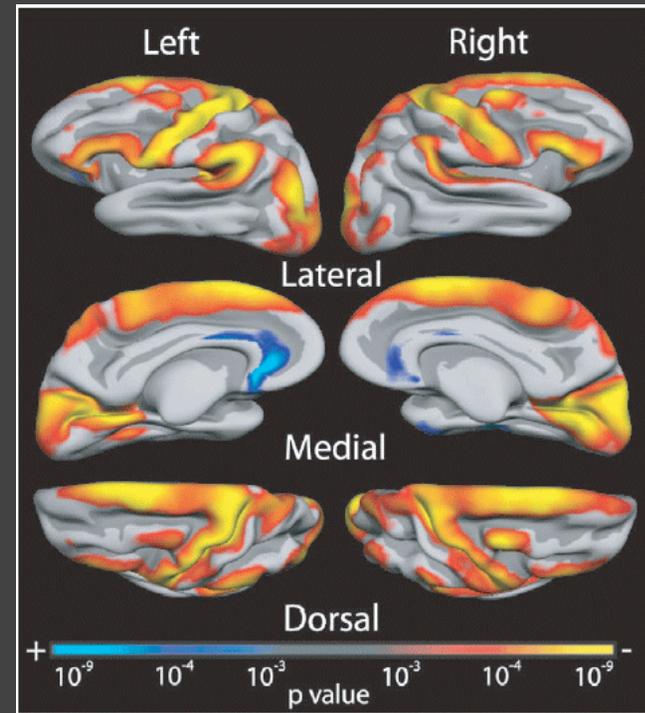
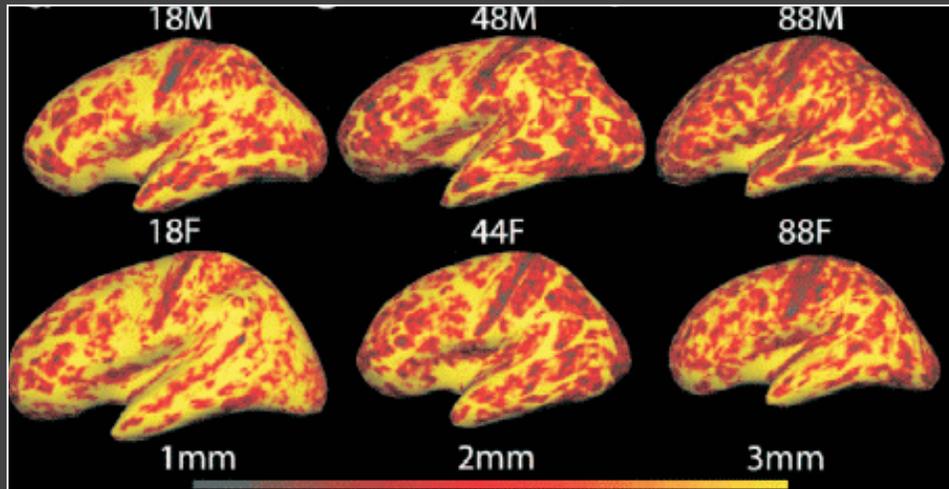
Cortical thickness



Distance between white and pial surfaces along normal vector (1-5mm)



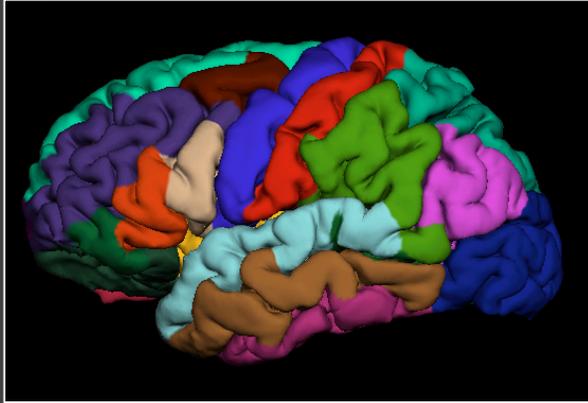
Application: Aging



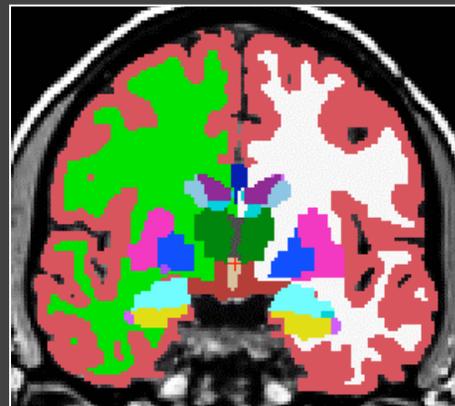
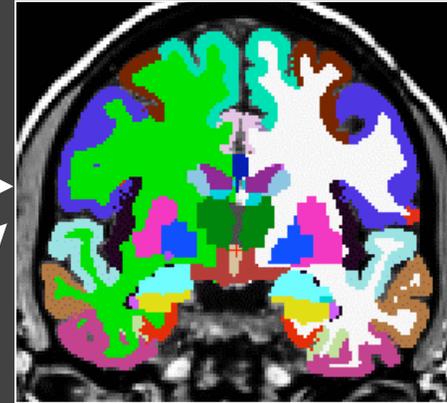
Salat *et al.*, 2004

Cortical/subcortical labels

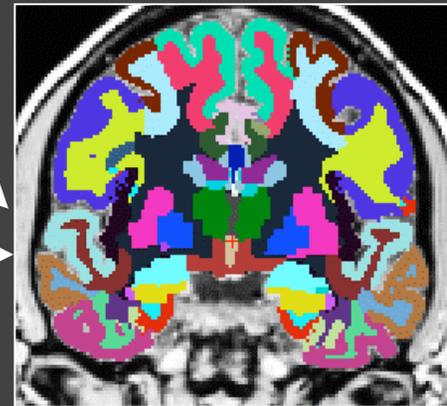
aparc



aparc+aseg



aseg



wmparc

(Nearest cortical label to any point in white matter)

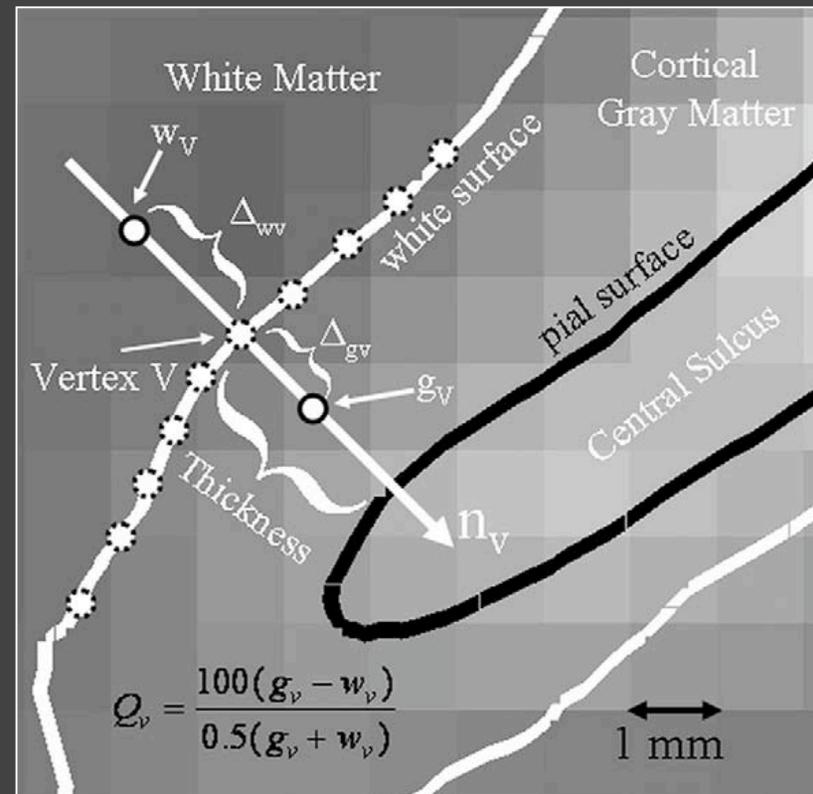
Using surfaces for intra-subject alignment

Boundary-Based Registration (BBR)

Boundary-based registration

Greve and Fischl, 2009

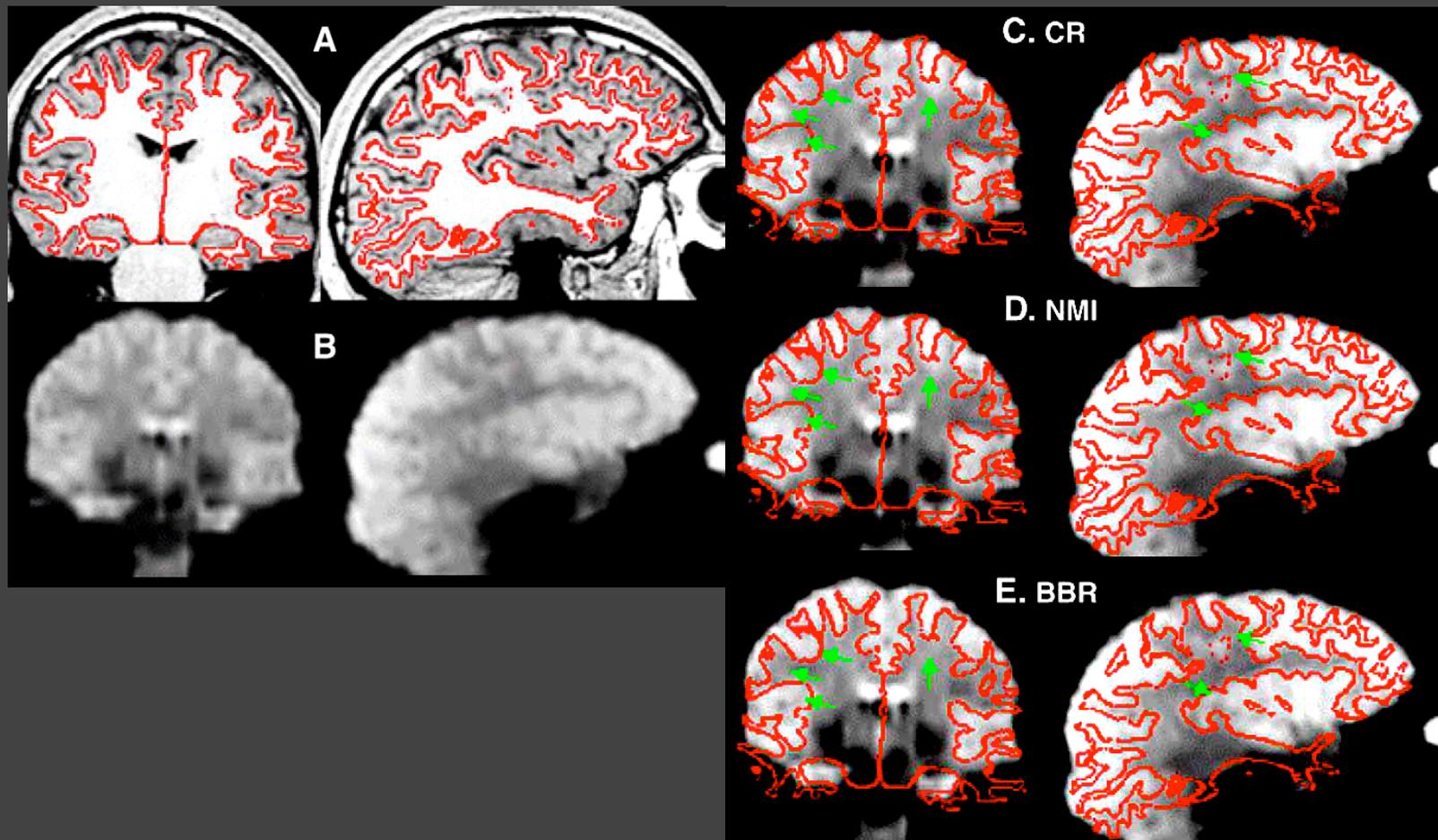
- Goal: **Register T_1** (higher resolution and SNR) **to EPI** (lower resolution and SNR) **from the same subject**
- EPI can be fMRI or a baseline (low-b) from a diffusion scan
- Most registration methods are intensity-based
- Boundary-based: Exploit the fact that T_1 gives us information on the gray/white boundary
- **Optimizes gradient of EPI gray/white contrast perpendicular to the surface**



Intensity- vs. boundary-based

Greve and Fischl, 2009

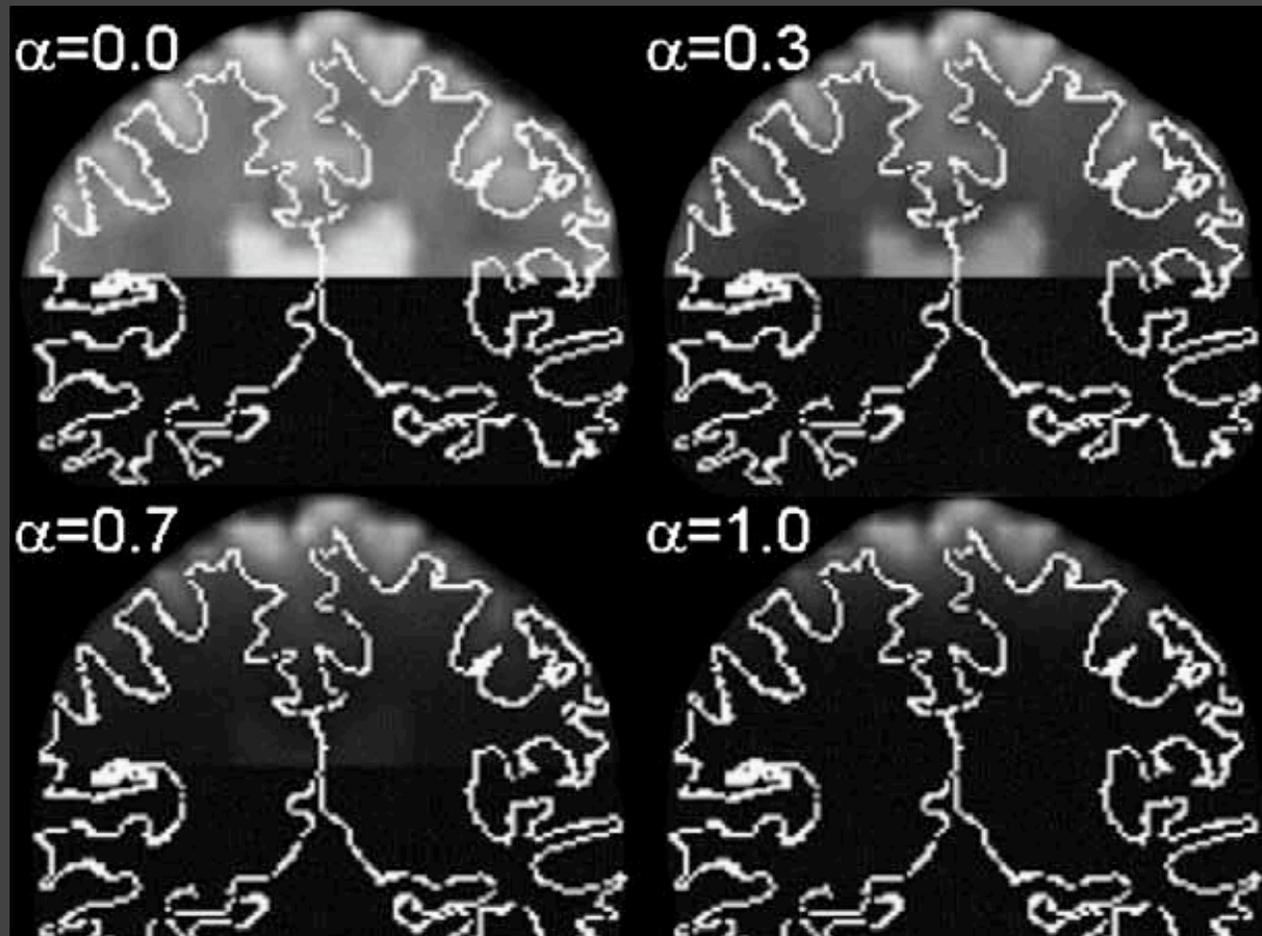
- Compare BBR to methods that use only image intensities: correlation ratio (CR), normalized mutual information (NMI)



Partial field of view

Greve and Fischl, 2009

- Very challenging for intensity-based methods



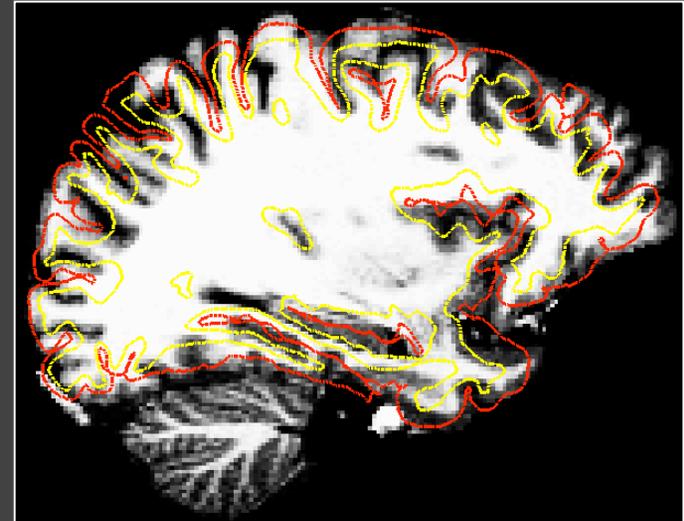
Using surfaces for inter-subject alignment

Combined Volume & Surface (CVS) registration

Inter-subject registration

Work by Lilla Zöllei and Gheorghe Postelnicu

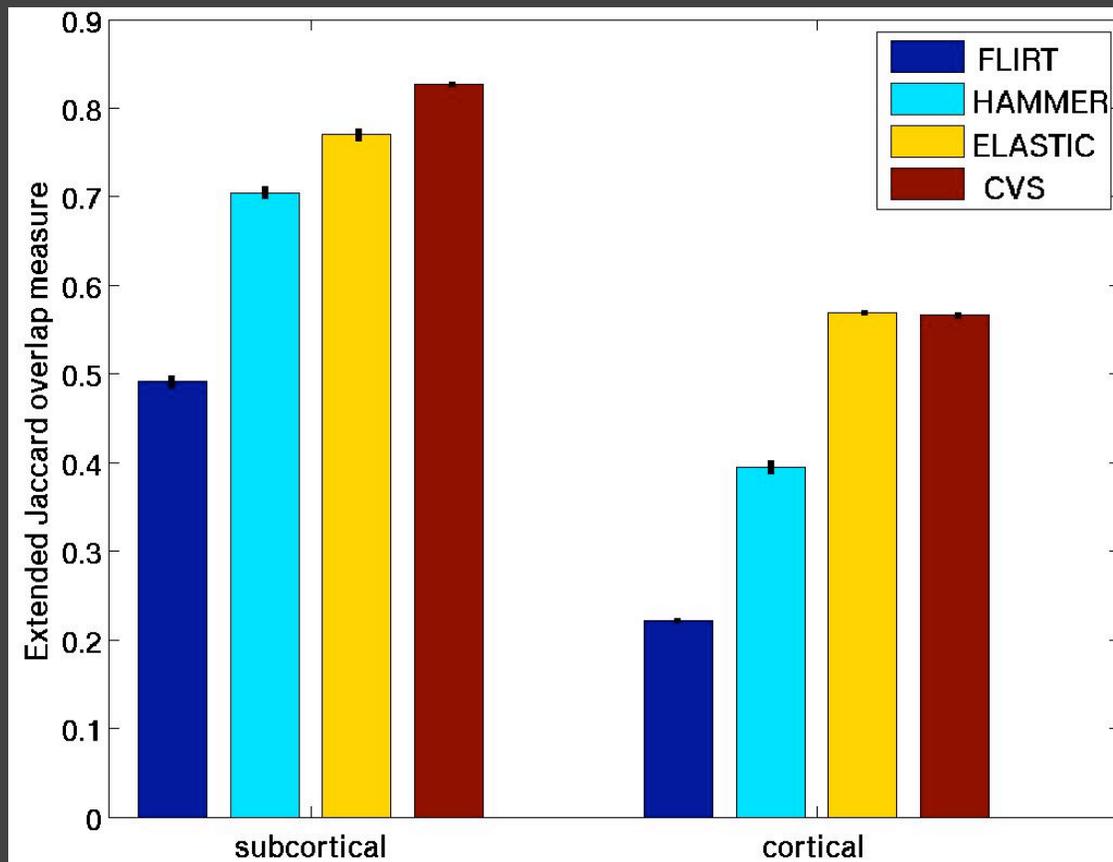
- **Volumetric (3D)**
 - Poor for cortical folds
 - + Applies to entire brain
- **Surface-based (2D)**
 - + Excellent for cortical folds
 - Doesn't apply to non-cortical structures
- **CVS: Combined Volume and Surface**
 1. Surface-based registration on the cortex
 2. Propagate into the volume via elastic transform
 3. Initialize intensity-based volumetric registration



Accuracy comparison

Slides by Lilla Zöllei

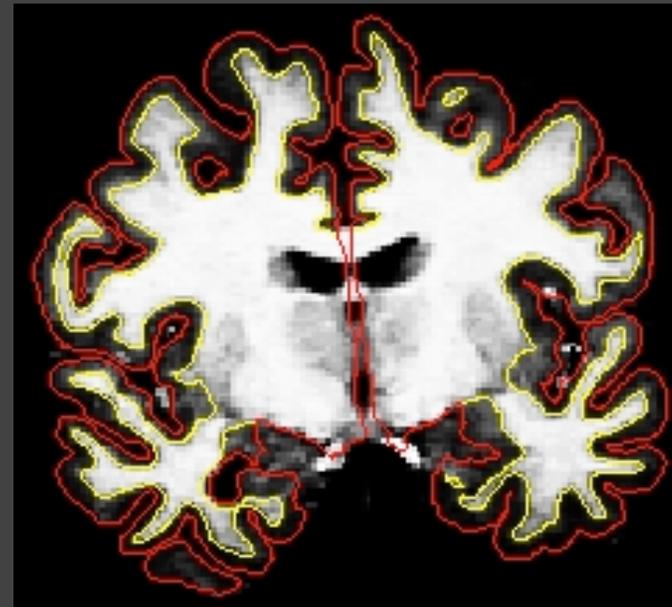
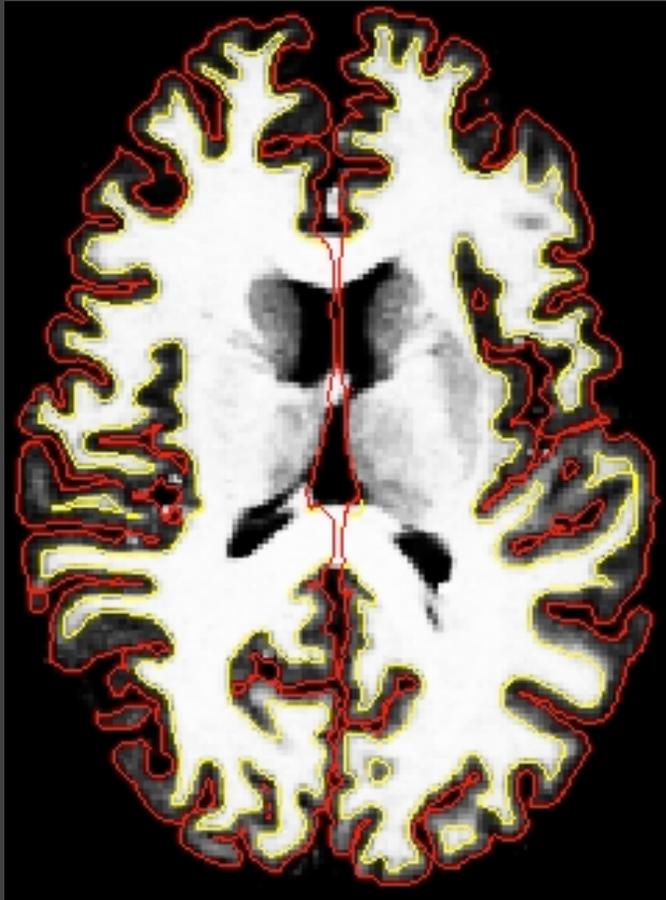
- 40 adult subjects with manually segmented cortex
- Leave-one-out, extended Jaccard overlap metric



Data courtesy of Dr.
R. Buckner, MGH

Template

Slides by Lilla Zöllei



FSL/FLIRT (affine)

Slides by Lilla Zöllei



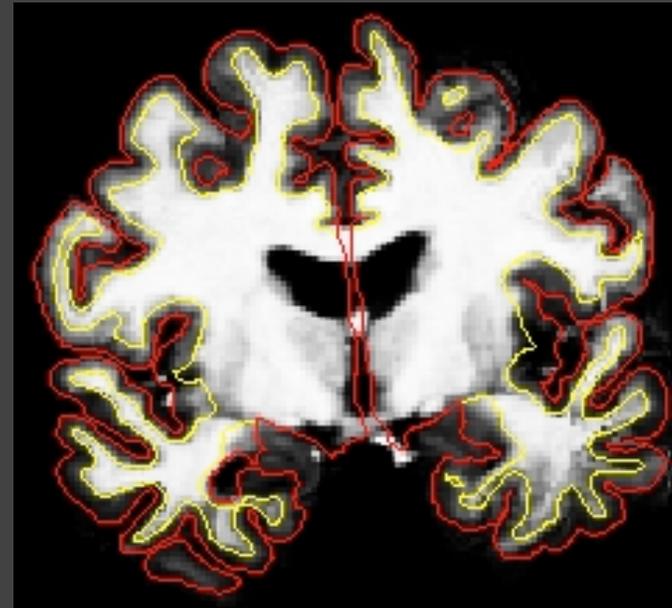
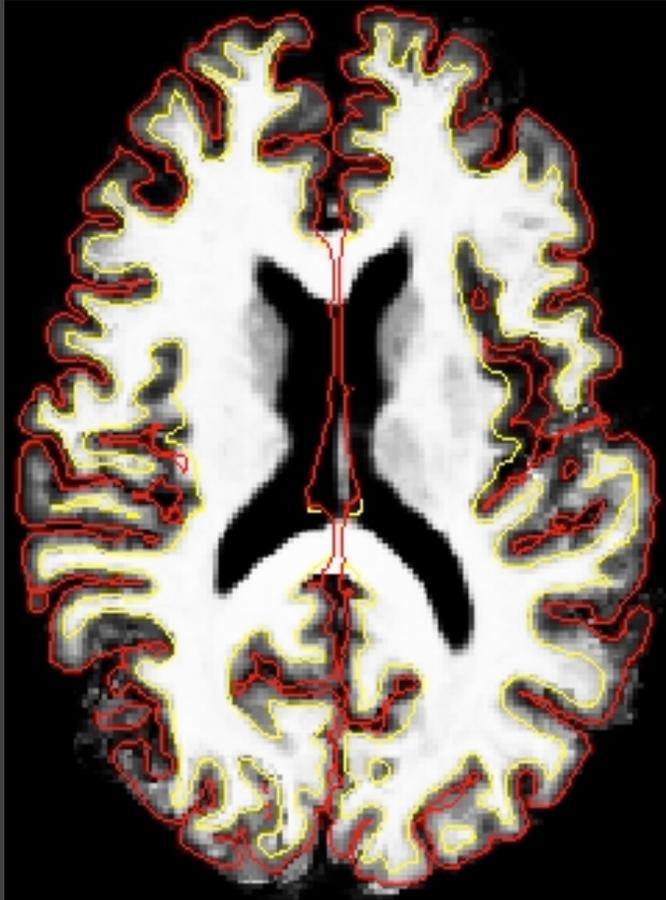
HAMMER (volumetric + gray/white)

Slides by Lilla Zöllei



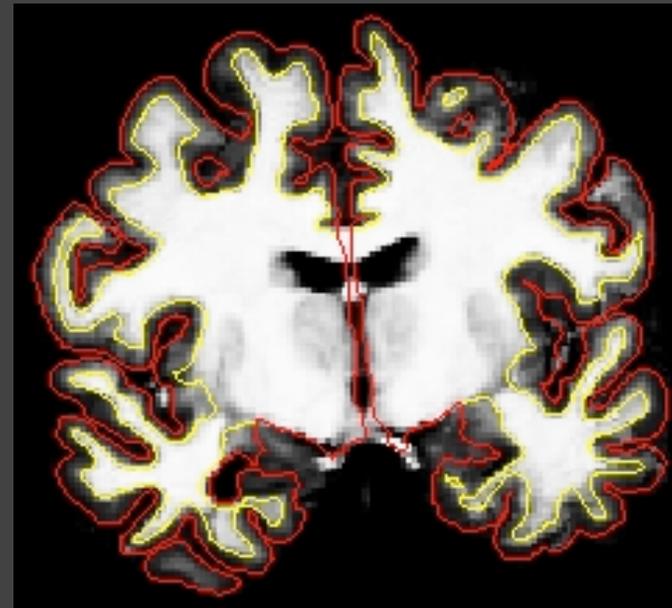
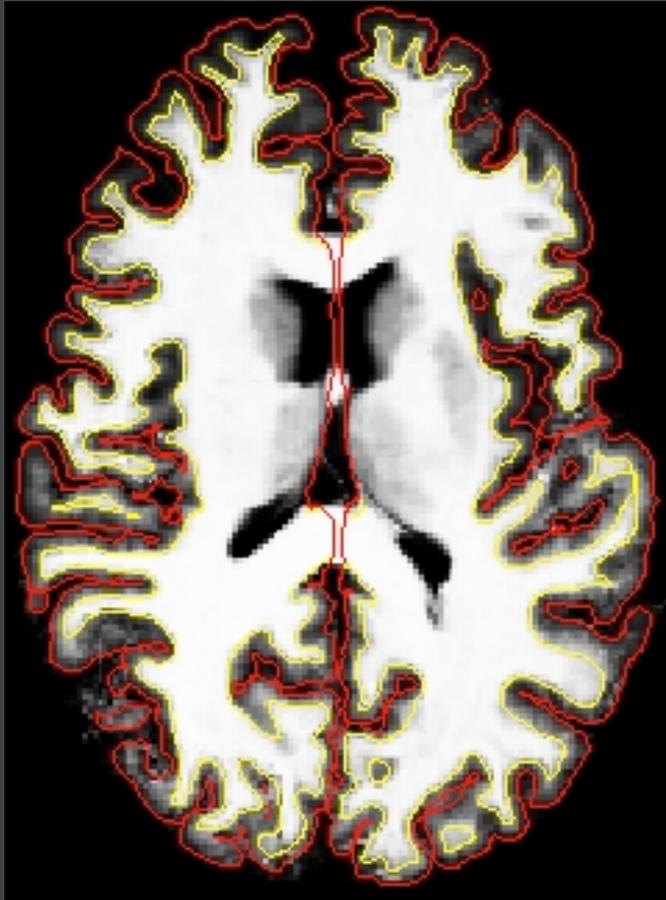
Elastic only

Slides by Lilla Zöllei



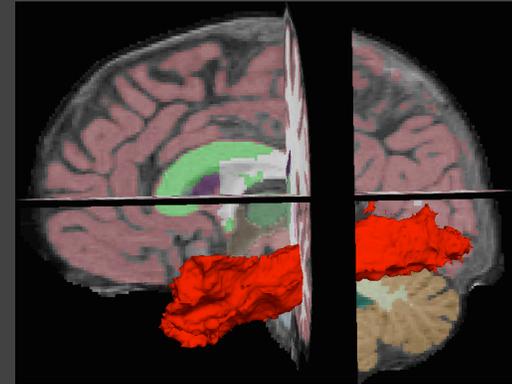
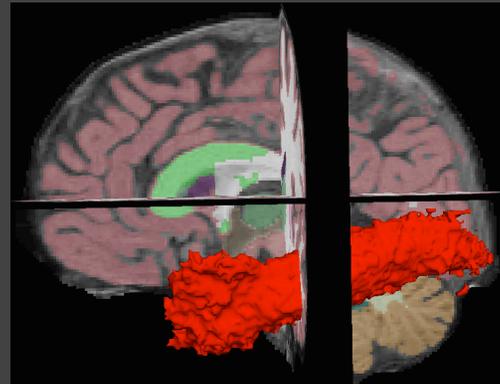
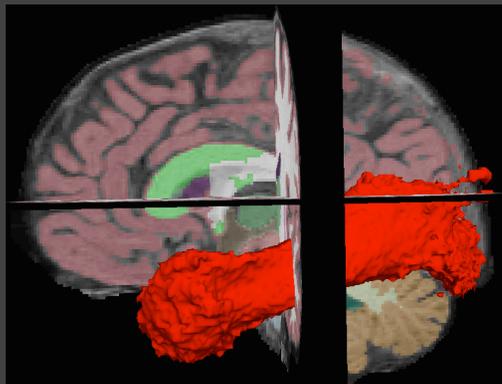
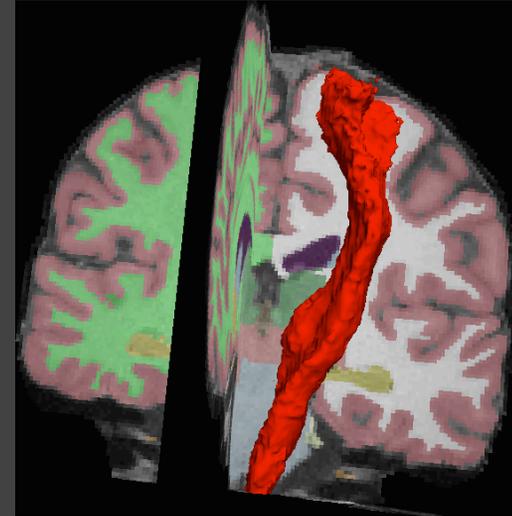
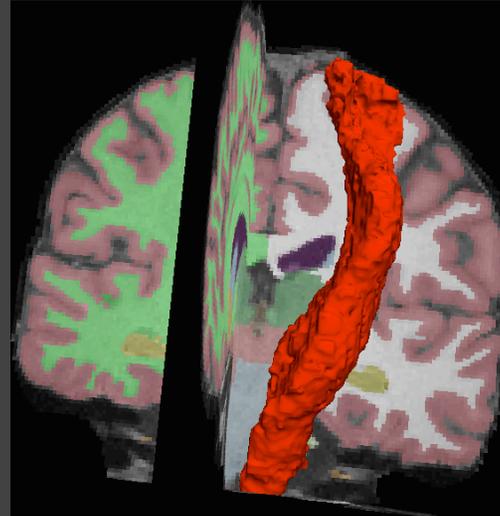
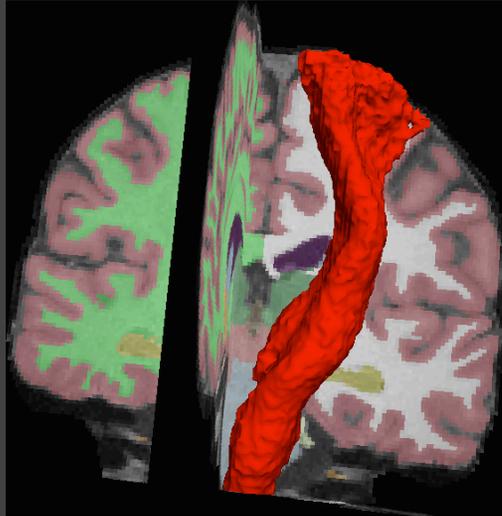
CVS (elastic + intensity-based)

Slides by Lilla Zöllei



Alignment of training paths

Zöllei et al, 2010



FLIRT (affine)

TBSS/FNIRT (nonlinear)

CVS (nonlinear)

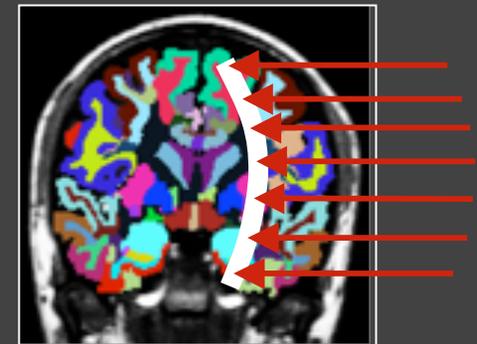
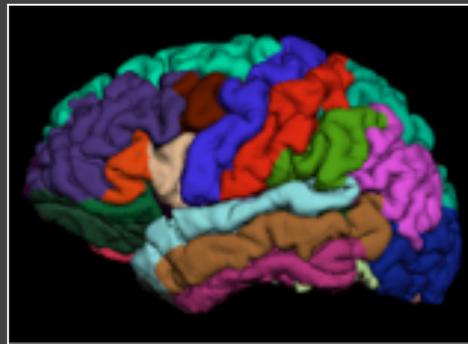
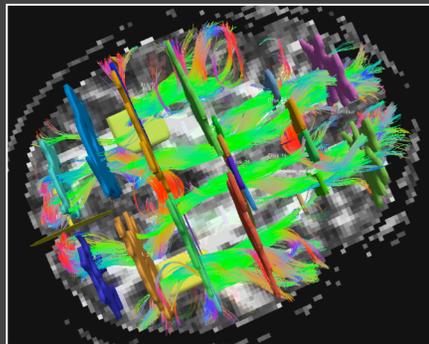
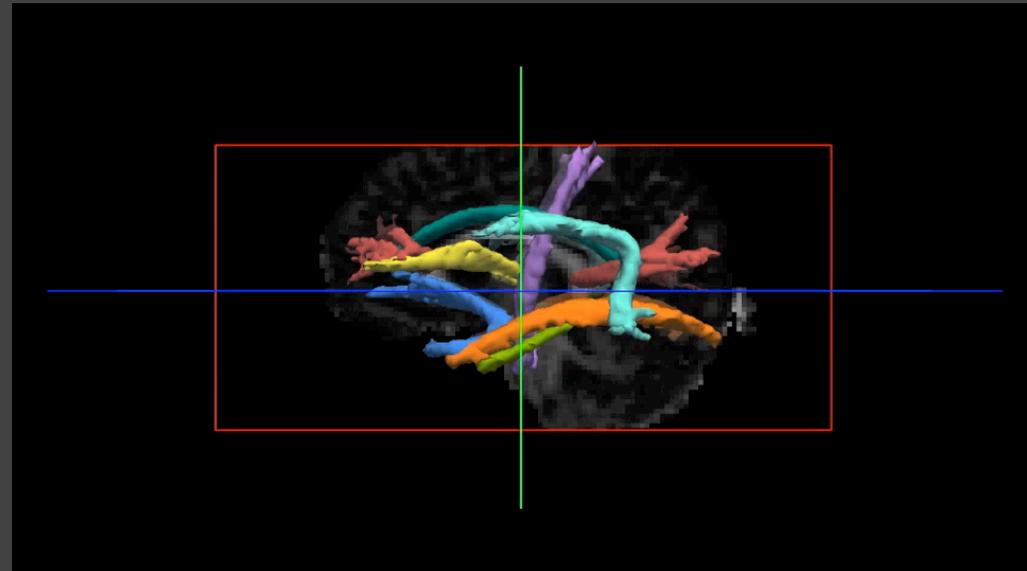
Using anatomical segmentation for automated tractography

TRActs Constrained by UnderLying Anatomy
(TRACULA)

Automated pathway reconstruction

[Yendiki et al 2011]

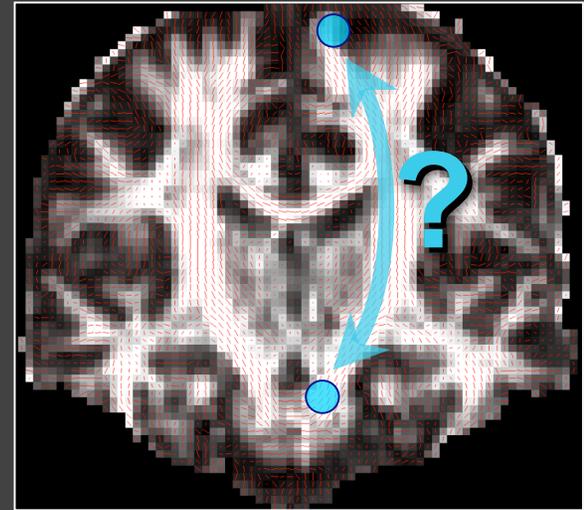
- **TRACULA** (TRActs
Constrained by
UnderLying Anatomy)
- Algorithm trained on the
likelihood of each
pathway passing by each
anatomical structure



Tractography studies

- **Exploratory tractography:**

- Example: *Show me all regions that the motor cortex is connected to*
- Seed region can be anatomically defined (motor cortex) or functionally defined (region activated in an fMRI finger-tapping task)



- **Tractography of known pathways:**

- Example: *Show me the corticospinal tract*
- Use prior anatomical knowledge of the pathway's terminations and trajectory (connects motor cortex and brainstem through capsule)

- TRACULA is for the latter type of study

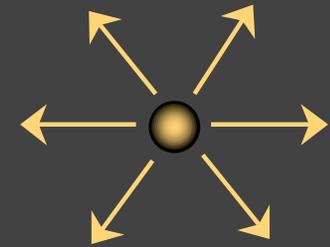
Tractography methods

- Every tractography method can be characterized by:
 - Which model of diffusion does it use?
 - Representation of local orientation of diffusion at every voxel that is fit to image data (tensor, ball-and-stick, etc.)
 - Is it deterministic or probabilistic?
 - Deterministic estimates only the most likely orientation
 - Probabilistic also estimates the uncertainty around that
 - Is it local or global?
 - Local fits the pathway to the data one step at a time
 - Global fits the entire pathway at once
- TRACULA does global probabilistic tractography with the ball-and-stick model

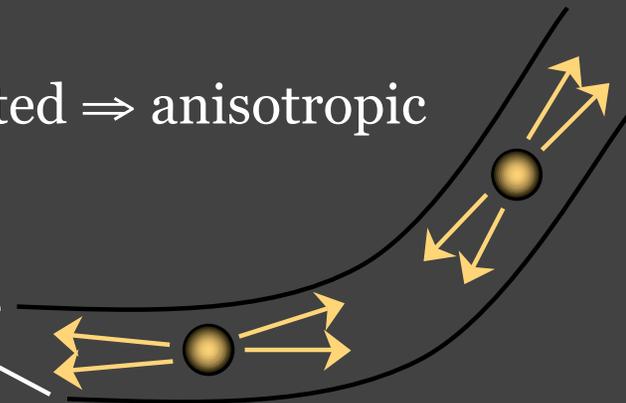
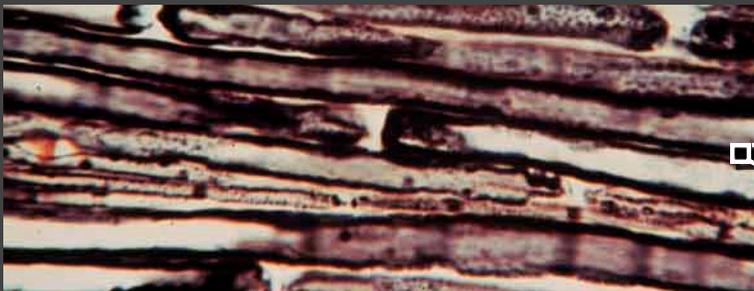
Diffusion in brain tissue

- Differentiate between tissues based on the diffusion (random motion) of water molecules within them

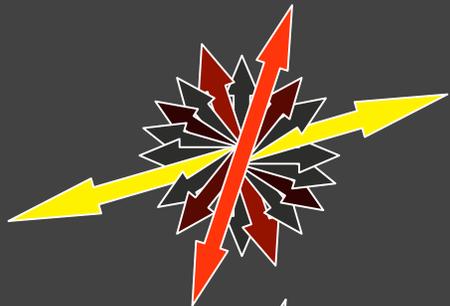
- Gray matter: Diffusion is unrestricted \Rightarrow isotropic



- White matter: Diffusion is restricted \Rightarrow anisotropic

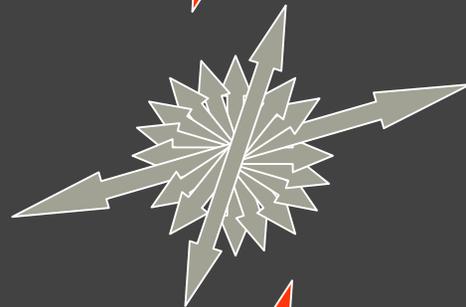


Models of diffusion



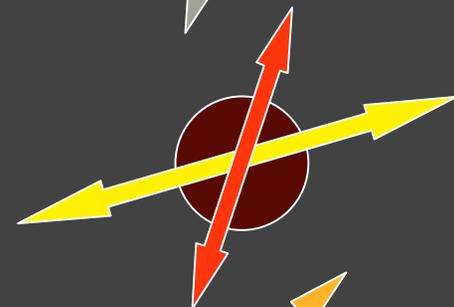
Diffusion spectrum:
Full distribution of orientation and magnitude

DSI



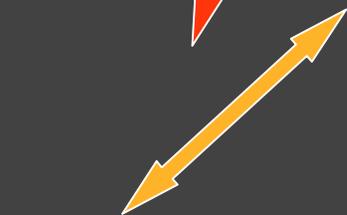
Orientation distribution function (ODF):
No magnitude info

Q-ball



Ball-and-stick:
Orientation and magnitude for up to N anisotropic compartments (default N=2)

FSL
(bedpostX)

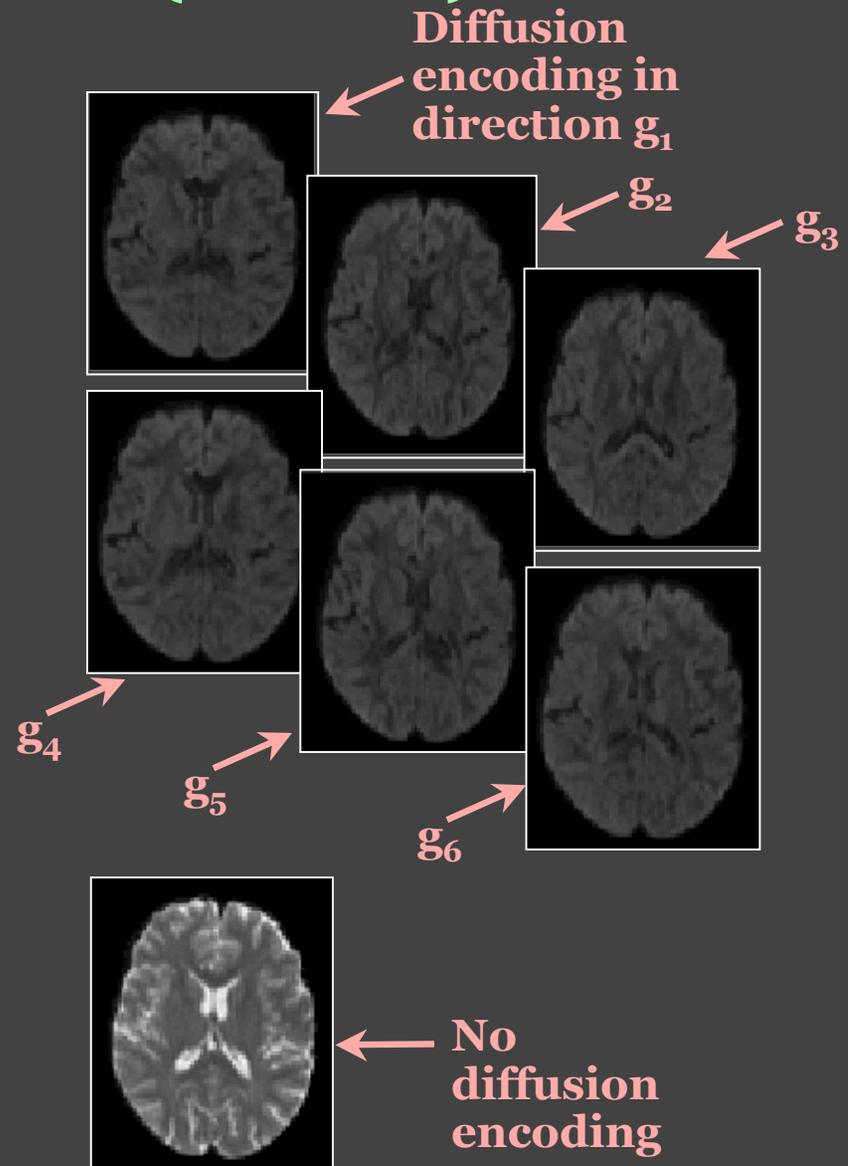


Tensor:
Single orientation and magnitude

DTI

Diffusion MRI (dMRI)

- Magnetic resonance imaging can provide “diffusion encoding”
- Magnetic field strength is varied by gradients in different directions
- Image intensity is attenuated depending on water diffusion in each direction
- Compare with baseline images to infer on diffusion process



Tensors

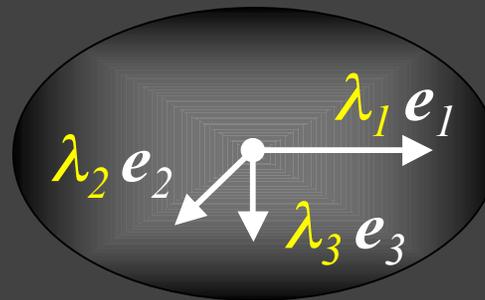
- One way to express the notion of direction is a **tensor** D
- A tensor is a 3x3 symmetric, positive-definite matrix:

$$D = \begin{bmatrix} d_{11} & d_{12} & d_{13} \\ d_{12} & d_{22} & d_{23} \\ d_{13} & d_{23} & d_{33} \end{bmatrix}$$

- D is symmetric 3x3 \Rightarrow It has 6 unique elements
- Suffices to estimate the upper (lower) triangular part

Eigenvalues & eigenvectors

- The matrix D is positive-definite \Rightarrow
 - It has 3 real, positive eigenvalues $\lambda_1, \lambda_2, \lambda_3 > 0$.
 - It has 3 orthogonal eigenvectors e_1, e_2, e_3 .



$$D = \lambda_1 e_1 \cdot e_1' + \lambda_2 e_2 \cdot e_2' + \lambda_3 e_3 \cdot e_3'$$

eigenvalue

eigenvector

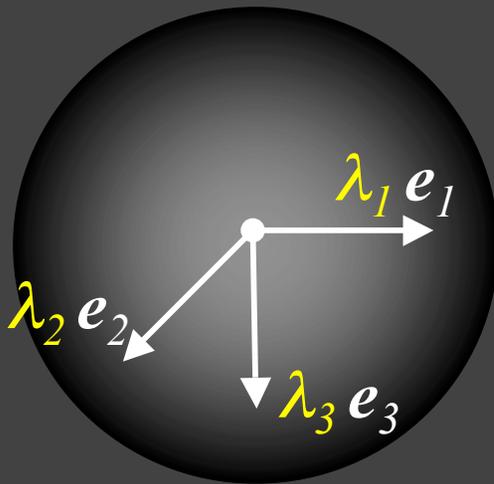
$$e_i = \begin{bmatrix} e_{ix} \\ e_{iy} \\ e_{iz} \end{bmatrix}$$

Physical interpretation

- Eigenvectors express diffusion direction
- Eigenvalues express diffusion magnitude

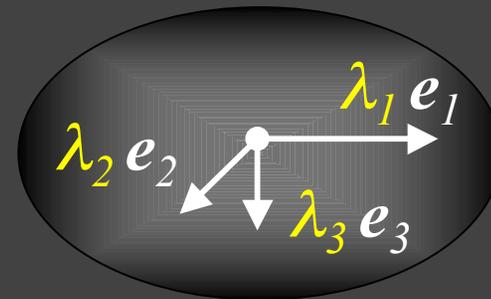
Isotropic diffusion:

$$\lambda_1 \approx \lambda_2 \approx \lambda_3$$



Anisotropic diffusion:

$$\lambda_1 \gg \lambda_2 \approx \lambda_3$$

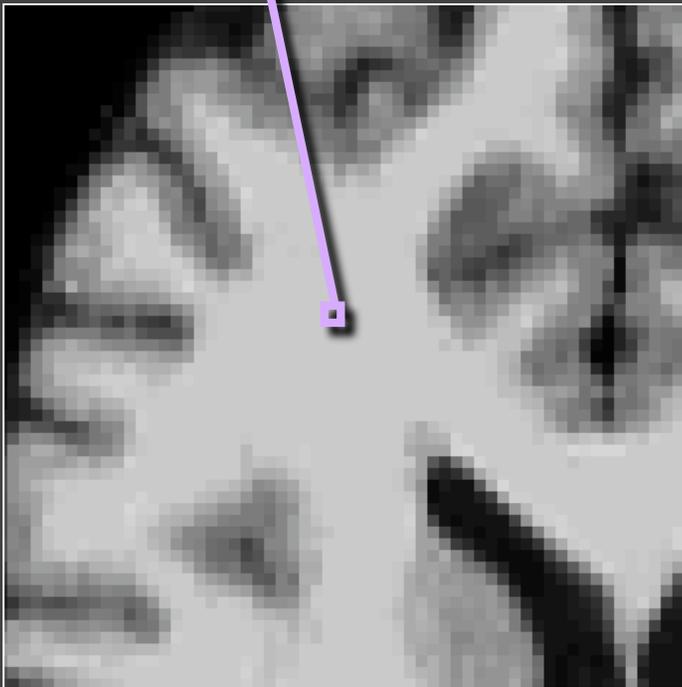


- One such ellipsoid at each voxel: Likelihood of water molecule displacements at that voxel

Diffusion tensor imaging (DTI)

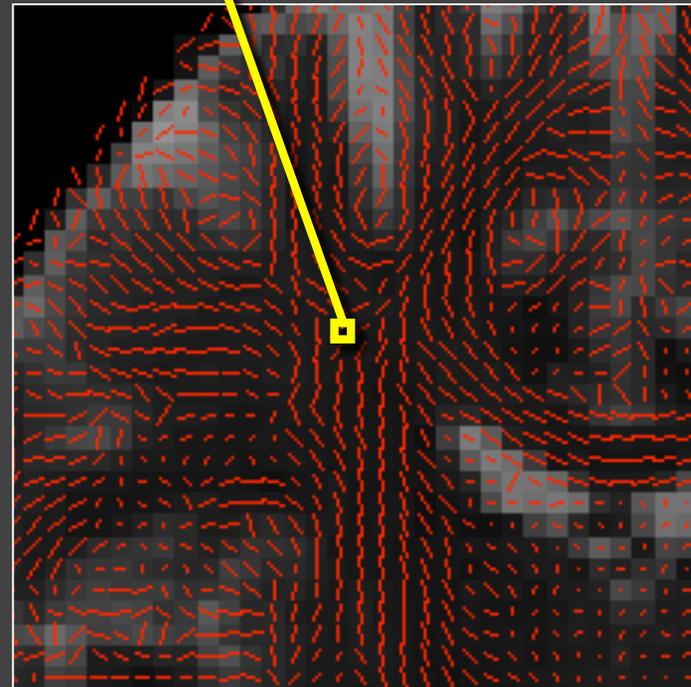
Image:

An intensity value at each voxel



Tensor map:

A tensor at each voxel

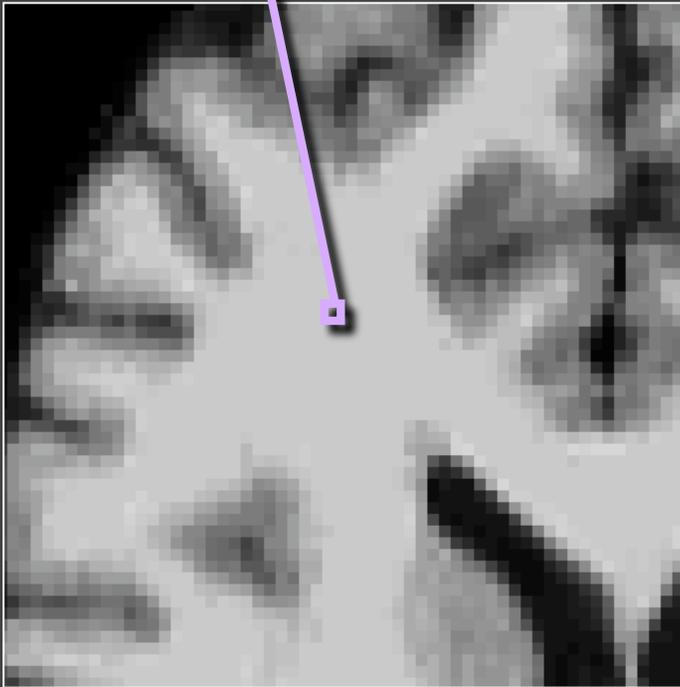


Direction of eigenvector corresponding to greatest eigenvalue

Diffusion tensor imaging (DTI)

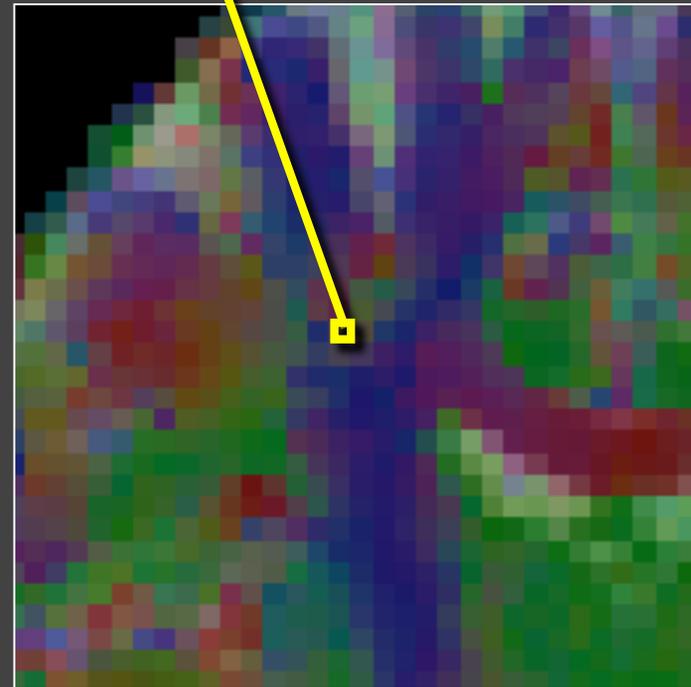
Image:

An intensity value at each voxel



Tensor map:

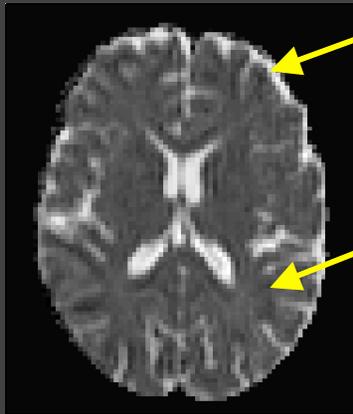
A **tensor** at each voxel



Direction of eigenvector corresponding to greatest eigenvalue

Red: L-R, Green: A-P, Blue: I-S

Tensor-based measures of diffusion

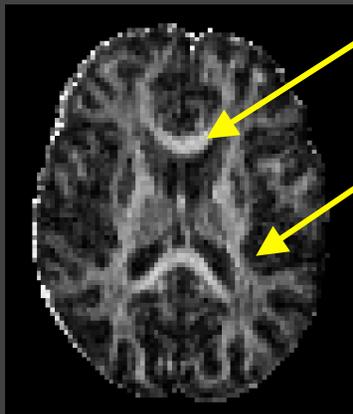


Faster diffusion

Slower diffusion

- Mean diffusivity (MD):
Mean of the 3 eigenvalues

$$MD(j) = [\lambda_1(j) + \lambda_2(j) + \lambda_3(j)]/3$$



Anisotropic diffusion

Isotropic diffusion

- Fractional anisotropy (FA):
Variance of the 3 eigenvalues,
normalized so that $0 \leq (FA) \leq 1$

$$FA(j)^2 = \frac{3}{2} \frac{[\lambda_1(j) - MD(j)]^2 + [\lambda_2(j) - MD(j)]^2 + [\lambda_3(j) - MD(j)]^2}{\lambda_1(j)^2 + \lambda_2(j)^2 + \lambda_3(j)^2}$$

More tensor-based measures

- Axial diffusivity: Greatest of the 3 eigenvalues

$$AD(j) = \lambda_1(j)$$

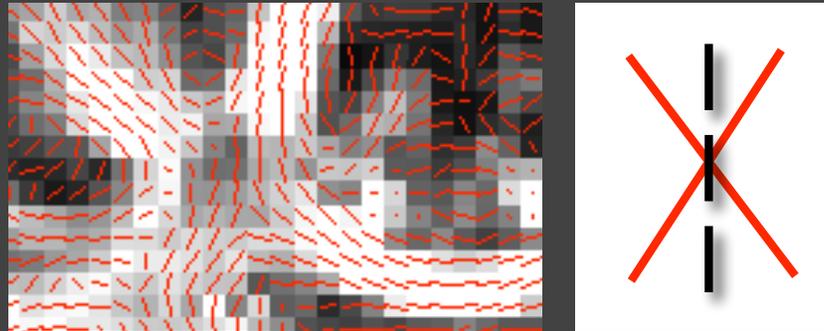
- Radial diffusivity: Average of 2 lesser eigenvalues

$$RD(j) = [\lambda_2(j) + \lambda_3(j)]/2$$

- Inter-voxel coherence: Average angle b/w the major eigenvector at some voxel and the major eigenvector at the voxels around it

Beyond the tensor

- The tensor is an imperfect model: What if more than one major diffusion direction in the same voxel?



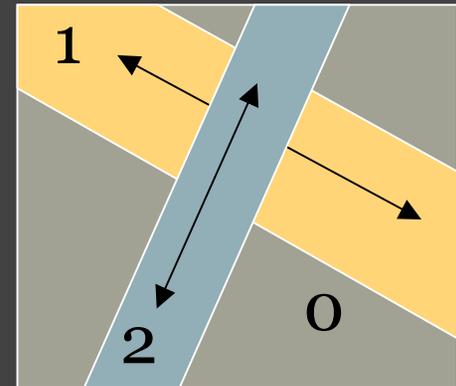
- High angular resolution diffusion imaging (HARDI): More complex models to capture more complex microarchitecture
 - Mixture of tensors [Tuch'02]
 - Higher-rank tensor [Frank'02, Özarslan'03]
 - Ball-and-stick [Behrens'07]
 - Orientation distribution function [Tuch'04]
 - Diffusion spectrum [Wedeen'05]

Non-tensor measures of diffusion

- From the orientation distribution function (ODF) [Tuch'04]:
 - Peak directions
 - Generalized fractional anisotropy (GFA)

$$\text{GFA}^2 = \frac{n \sum_i [\text{ODF}(\theta_i) - \sum_i \text{ODF}(\theta_i)/n]^2}{(n-1) \sum_i \text{ODF}(\theta_i)^2}$$

- From the ball-and-stick model [Behrens'07]:
 - Orientation angles of each anisotropic compartment (1, 2, ...)
 - Volume of each anisotropic compartment
 - Overall diffusivity in the voxel



Choice 1: Gradient directions

- True diffusion direction \parallel Applied gradient direction
 \Rightarrow Maximum attenuation



- True diffusion direction \perp Applied gradient direction
 \Rightarrow No attenuation



- To capture all diffusion directions well, gradient directions should cover 3D space uniformly



How many directions?

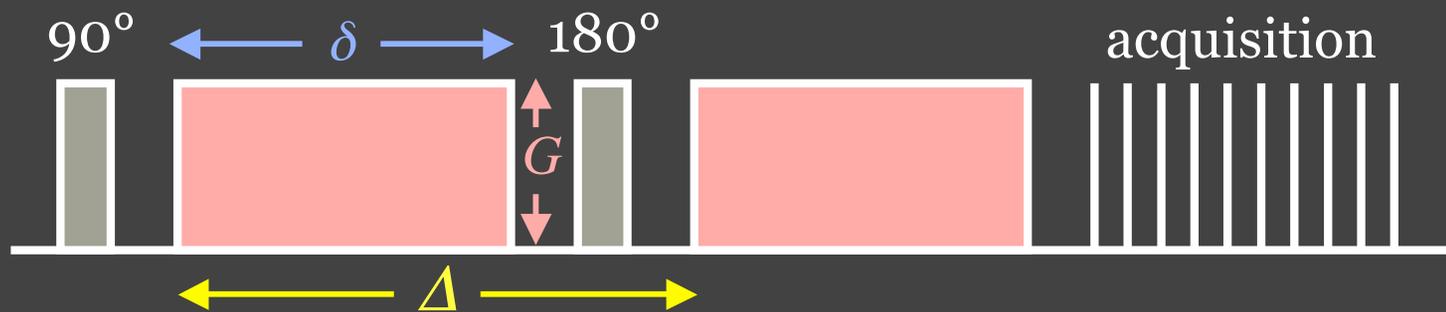
- Acquiring data with more gradient directions leads to:
 - + More reliable estimation of diffusion measures
 - Increased imaging time \Rightarrow Subject discomfort, more susceptible to artifacts due to motion, respiration, etc.
- DTI:
 - Six directions is the minimum
 - Usually a few 10's of directions
 - Diminishing returns after a certain number [Jones, 2004]
- DSI:
 - Usually a few 100's of directions

Choice 2: The b-value

- The b-value depends on acquisition parameters:

$$b = \gamma^2 G^2 \delta^2 (\Delta - \delta/3)$$

- γ the gyromagnetic ratio
- G the strength of the diffusion-encoding gradient
- δ the duration of each diffusion-encoding pulse
- Δ the interval b/w diffusion-encoding pulses



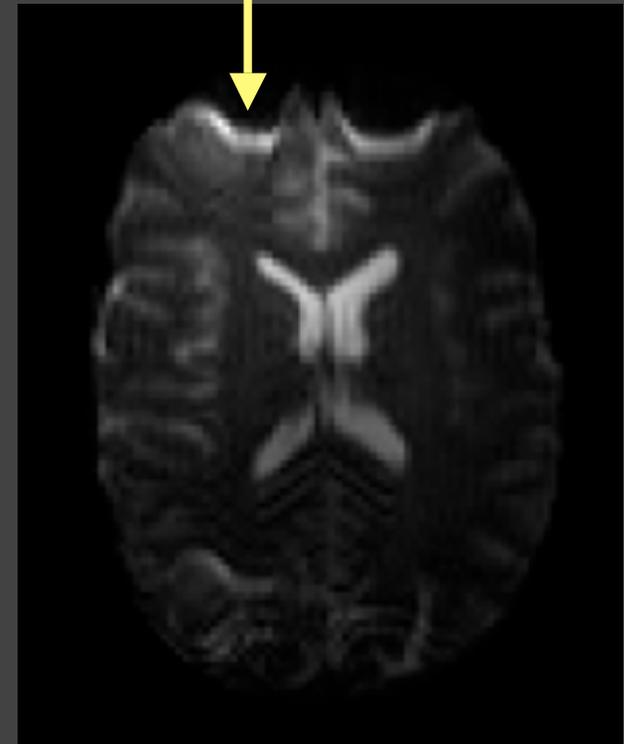
How high b-value?

- Increasing the b-value leads to:
 - + Increased contrast b/w areas of higher and lower diffusivity in principle
 - Decreased signal-to-noise ratio \Rightarrow Less reliable estimation of diffusion measures in practice
- DTI: $b \sim 1000 \text{ sec/mm}^2$
- DSI: $b \sim 10,000 \text{ sec/mm}^2$
- Data can be acquired at multiple b-values for trade-off
- Repeat acquisition and average to increase signal-to-noise ratio

Distortions: Field inhomogeneities

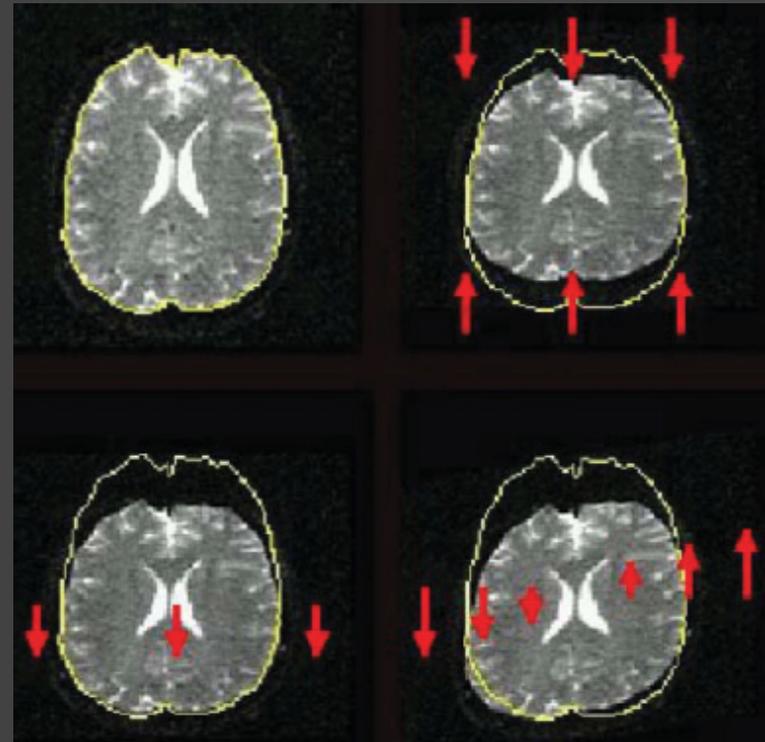
- Causes:
 - **Scanner-dependent** (imperfections of main magnetic field)
 - **Subject-dependent** (changes in magnetic susceptibility in tissue/air interfaces)
- Results:
 - Signal loss in interface areas
 - Geometric distortions (warping) of the entire image

Signal loss



Distortions: Eddy currents

- Cause: Fast switching of diffusion-encoding gradients induces eddy currents in conducting components
- Eddy currents lead to residual gradients that shift the diffusion gradients
- The shifts are **direction-dependent**, *i.e.*, different for each DW image
- Result: Geometric distortions



From Le Bihan *et al.*, Artifacts and pitfalls in diffusion MRI, JMIR 2006

Data analysis steps

- Pre-process images to reduce distortions
 - Either register distorted DW images to an undistorted (non-DW) image
 - Or use information on distortions from separate scans (field map, residual gradients)
- Fit a diffusion model at every voxel
 - DTI, DSI, Q-ball, ...
- Compute measures of anisotropy/diffusivity and compare them between populations
 - Voxel-based (*e.g.*, TBSS), ROI-based, tract-based
- For tract-based: Reconstruct pathways

