HST.583 Functional Magnetic Resonance Imaging: Data Acquisition and Analysis Fall 2008

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Diffusion weighted imaging

Anastasia Yendiki

HMS/MGH/MIT Athinoula A. Martinos Center for Biomedical Imaging

Why diffusion imaging?

- White matter (WM) is organized in fiber bundles
- Identifying these WM pathways is important for:
 - Inferring connections
 b/w brain regions
 - Understanding effects of neurodegenerative diseases, stroke, aging, development ...



From Gray's Anatomy: IX. Neurology

Diffusion in brain tissue

- Differentiate tissues based on the diffusion (random motion) of water molecules within them
- Gray matter: Diffusion is unrestricted ⇒ isotropic



White matter: Diffusion is restricted ⇒ anisotropic



Diffusion MRI

- 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



Imaging diffusion

• Image the direction in which water molecules diffuse at each voxel in the brain

 \Rightarrow Infer WM fiber orientation at each voxel

• Clearly, direction can't be described by a usual grayscale image





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Tensors

- We express the notion of "direction" mathematically by 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 λ_1 , λ_2 , $\lambda_3 > 0$.
 - It has 3 orthogonal eigenvectors e_1 , e_2 , e_3 .

Physical interpretation

- Eigenvectors express diffusion direction
- Eigenvalues express diffusion magnitude

Isotropic diffusion:

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

Anisotropic diffusion: $\lambda_1 >> \lambda_2 \approx \lambda_3$

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

Diffusion tensor imaging (DTI)

Image:

```
A scalar intensity value f_j at each voxel j
```

Grayscale brain image removed due to copyright restrictions. **Tensor map:**

A tensor **D**_j at each voxel j

Courtesy of Gordon Kindlmann. Used with permission.

Summary measures

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

 $MD(j) = [\lambda_{I}(j) + \lambda_{2}(j) + \lambda_{3}(j)]/3$

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

 $FA(j)^{2} = \frac{3}{2} \frac{[\lambda_{l}(j)-MD(j)]^{2} + [\lambda_{2}(j)-MD(j)]^{2} + [\lambda_{3}(j)-MD(j)]^{2}}{[\lambda_{3}(j)-MD(j)]^{2}}$

 $\lambda_{l}(j)^{2} + \lambda_{2}(j)^{2} + \lambda_{3}(j)^{2}$

More summary measures

- Axial diffusivity: Greatest eigenvalue $AD(j) = \lambda_{I}(j)$
- 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

Other models of diffusion

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

- High angular resolution diffusion imaging (HARDI)
 - A mixture of the usual ("rank-2") tensors [Tuch'02]
 - A tensor of rank > 2 [Frank'02, Özarslan'03]
 - An orientation distribution function [Tuch'04]
 - A diffusion spectrum (DSI) [Wedeen'05]
- More parameters at each voxel ⇒ More data needed

Example: DTI vs. DSI

Source: Wedeen, V. J. *et al., "*Mapping complex tissue architecture with diffusion spectrum magnetic resonance imaging." *MRM* 54, no. 6 (2005): 1377-1386. Copyright (c) 2005 Wiley-Liss, Inc., a subsidiary of John Wiley & Sons, Inc. Reprinted with permission of John Wiley & Sons., Inc.

Back to the tensor

• Remember: A tensor has six unique values

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

Must estimate six times as many values at each voxel
 ⇒ Must collect (at least) six times as much data!

MR data acquisition

Measure raw MR signal (frequency-domain samples of transverse magnetization)

Reconstruct an image of transverse magnetization

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Diffusion MR data acquisition

Must acquire at least 6 times as many MR signal measurements Need to reconstruct 6 times as many values

Diffusion encoding in MRI

• Apply two gradient pulses in some direction *y*:

• Case 1: If spins aren't diffusing

No displacement in $y \Rightarrow$ No dephasing \Rightarrow No net signal change

• Case 2: If spins are diffusing

Choice 1: Gradient directions

Spin diffusion direction || Applied gradient direction
 Maximum attenuation
 Diffusion-encoding gradient g

Displacement detected

Spin diffusion direction ⊥ Applied gradient direction
 ⇒ No attenuation

Diffusion-encoding gradient *g* Displacement not detected

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

Diffusion-encoding gradient *g* Displacement partly detected

How many directions?

- Acquiring with more gradient directions leads to:
 - + More reliable estimation of diffusion measures
 - Increased imaging time ⇒ 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]
- HARDI/DSI:
 - Usually a few 100's of directions

Choice 2: The b-value

- The b-value depends on acquisition parameters: $b = \gamma^2 G^2 \delta^2 (\Delta - \delta/3)$
 - $-\gamma$ the gyromagnetic ratio
 - *G* the strength of the diffusion-encoding gradient
 - $-\delta$ the duration of each diffusion-encoding pulse
 - ⊿ 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 ⇒ Less reliable estimation of diffusion measures in practice
- DTI: $b \sim 1000 \text{ sec/mm}^2$
- HARDI/DSI: b ~ 10,000 sec/mm²
- Data can be acquired at multiple b-values for trade-off
- Repeat acquisition and average to increase signal-tonoise ratio

Estimating the tensor

• $f_i^{b,g} = f_i^0 e^{-bg' \cdot D_j \cdot g}$

where the D_i the diffusion tensor at voxel j

- Design acquisition:
 - *b* the diffusion-weighting factor
 - g the diffusion-encoding gradient direction
- Acquire images:
 - $f_j^{b,g}$ image acquired with diffusion-weighting factor b and diffusion-encoding gradient direction g
 - $-f_j^0$ "baseline" image acquired without diffusionweighting (b=0)
- Estimate unknown diffusion tensor *D_i*

Noise in diffusion-weighted images

- Due to signal attenuation by diffusion encoding, signal-to-noise ratio in DW images can be an order of magnitude lower than "baseline" image
- Eigenvalue decomposition is sensitive to noise, may result in negative eigenvalues

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

Distortions: Eddy currents

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

Source: Le Bihan *D., et al.* "Artifacts and pitfalls in diffusion MRI." *JMRI* 24, no. 3 (2006): 478-488. Copyright © 2006 Wiley-Liss, Inc., A Wiley Company. Reprinted with permission of John Wiley & Sons., Inc.

Distortion correction

Post-process images to reduce distortions due to field inhomogeneities and eddy-currents:

Either register distorted DW images to an undistorted (non-DW) image

[Haselgrove'96, Bastin'99, Horsfield'99, Andersson'02, Rohde'04, Ardekani'05, Mistry'06]

Or use information on distortions from separate scans (field map, residual gradients)
 [Jezzard'98, Bastin'00, Chen'06; Bodammer'04, Shen'04]

Tractography

- What does one do with diffusion data?
 - Statistical analysis on MD, FA, tensors...
 - Tractography: Given the diffusion data, determine "best" pathway between two brain regions
- Challenges in tractography:
 - Noisy, distorted images
 - Pathway crossings
 - High-dimensional space
- Many methods to overcome them...

Deterministic vs. probabilistic

- Deterministic methods: Model geometry of diffusion data, e.g., tensor/eigenvectors [Conturo '99, Jones '99, Mori '99, Basser '00, Catani '02, Parker '02, O'Donnell '02, Lazar '03, Jackowski '04, Pichon '05, Fletcher '07, Melonakos '07, ...]
- Probabilistic methods: Also model statistics of diffusion data [Behrens '03, Hagmann '03, Pajevic '03, Jones '05, Lazar '05, Parker '05, Friman '06, Jbabdi '07, ...]

Local vs. global

- Local: Uses local information to determine next step, errors propagate from areas of high uncertainty
- **Global:** Integrates information along the entire path

Local tractography

- Define a "seed" voxel or ROI to start the tract from
- Trace the tract by small steps, determine "best" direction at each step
- Deterministic: Only one possible direction at each step

• **Probabilistic:** Many possible directions at each step (because of noise), some more likely than others

Some issues

- Not constrained to a connection of the seed to a target region
- How do we isolate a specific connection? We can set a threshold, but how?
- What if we want a nondominant connection? We can define waypoints, but there's no guarantee that we'll reach them.
- Not symmetric between tract "start" and "end" point

Global tractography

- Define a "seed" voxel or ROI
- Define a "target" voxel or ROI
- **Deterministic:** Only one possible path
- **Probabilistic:** Many possible paths, find their probability distribution
- Constrained to a specific connection
- Symmetric between seed and target regions

Application: Huntington's disease

Data courtesy of Dr. D. Rosas, MGH

Healthy

Huntington's (premanifest)

Huntington's (advanced)

CST

SLF2

SLF1

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Vlean FA

Vean FA

0.6

0.4

0.2

0.4

0.2

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