

Fiber Orientation Distribution Function Estimation by Spherical Needlets

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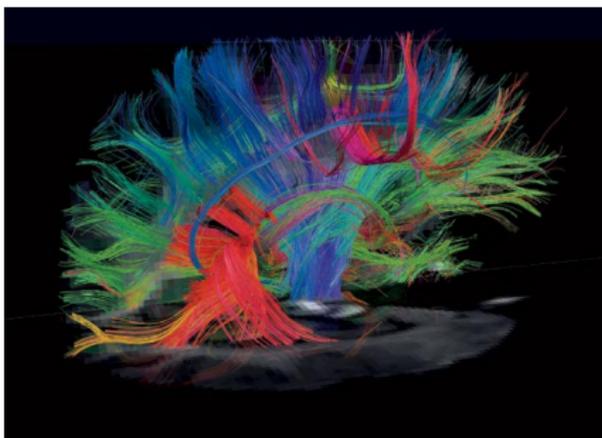
Diffusion MRI

- Diffusion MRI is a magnetic resonance imaging technology which measures *diffusion* of water molecules along a set of (predetermined) directions.
 - In vivo, non-invasive, no radiation.
- Diffusion MRI uses water diffusion as a proxy to probe the anatomy of biological tissues.
- Raw data from a D-MRI experiment:
 - Multiple grey scale images corresponding to multiple gradient directions plus a few images with no diffusion weighting.
 - Each image consists of intensities for pixels on a 3D grid (e.g., $\sim 256 \times 256 \times 59$ for a human brain).
 - Image resolution: $1 \sim 3\text{mm}$.

D-MRI Provides Information on Brain Connectivity

- Neuron axons with similar destinations form big bundles called *white matter fiber tracts*.
- When applied to human brains, diffusion MRI reveals detailed anatomy of white matter tracts such as their location, size, shape and how they are connected to each other.
 - Human connectome project
<http://www.humanconnectomeproject.org>.
- Alzheimer's Disease Neuroimaging Initiative (ADNI):
<http://adni.loni.usc.edu/>.

Tractographic Reconstruction of Neural Connections



<http://braintalks.wordpress.com/2011/12/04/10-maps-of-the-mind/>

- With diffusion information at each voxel, fiber tracts can be reconstructed using computer-aided 3D tracking techniques called tractography.

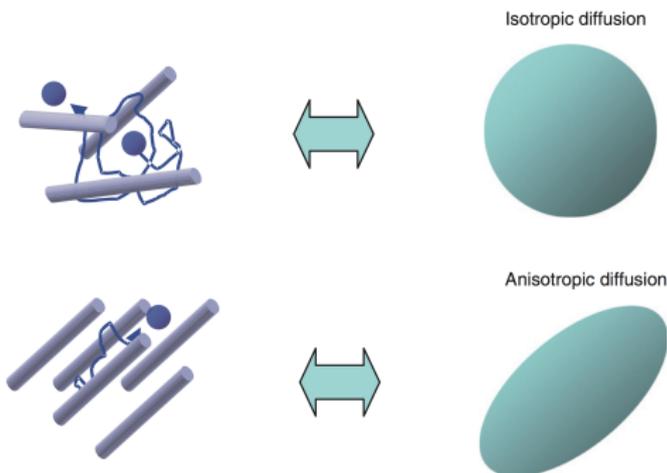
Clinical Applications of Diffusion MRI

- Detect brain abnormality in white matter regions such as specific axonal loss, deformation in brain tumors.
- Differentiate types of tumor and growth orientation.
- Measure anatomy of immature brains.
- Monitor status of specific white matter tracts.

Water Diffusion in Biological Tissues

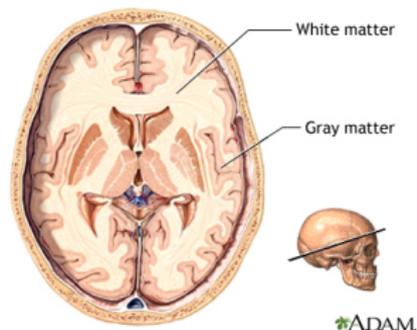
Anisotropic due to the presence of fibers with coherent orientations.

- Water tends to diffuse faster along fibers.
- Information on water diffusion may be used to probe tissue structure.



Diffusion in Brain

- White matter.
 - Astronomical number of connections: “cables” of the brain.
 - Presence of axonal bundles at image resolution \Rightarrow diffusion appears anisotropic.
- Grey matter.
 - ~ 100 billion neurons: “CPU” of the brain.
 - Lack of coherent fiber organization at image resolution ($\sim 2\text{mm}$) \Rightarrow diffusion appears isotropic.



Diffusion Weighted Signals

At voxel \mathbf{s} , along direction \mathbf{q} , diffusion weighted signal:

$$S_0(\mathbf{s}) \int_{R^3} p_{\mathbf{s}, \Delta t}(\mathbf{r}) \exp(i\gamma\delta\mathbf{q} \cdot \mathbf{r}) d\mathbf{r}.$$

- $S_0(\mathbf{s})$: signal intensity without diffusion weighting at voxel \mathbf{s} .
- $p_{\mathbf{s}, \Delta t}(\cdot)$: p.d.f. of water displacement in time duration Δt at voxel \mathbf{s} .
- Δt : time between “dephasing” and “rephasing”.
- δ : duration of dephasing/rephasing.
- γ : gyromagnetic ratio.

Diffusion weighted signal is the inverse Fourier transform of the diffusion probability density function.

[▶ details](#)

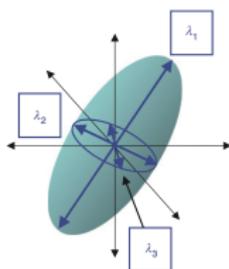
Gaussian Diffusion and Single Tensor Model

$$p_{\mathbf{s}, \Delta t}(\mathbf{r}) = \frac{1}{(2\pi)^{3/2}} |\mathbf{D}(\mathbf{s}) \Delta t|^{-\frac{1}{2}} \exp\left(-\frac{\mathbf{r}^T \mathbf{D}(\mathbf{s})^{-1} \mathbf{r}}{2\Delta t}\right), \quad \mathbf{r} \in \mathbb{R}^3.$$

- $\mathbf{D}(\mathbf{s})$: **diffusion tensor**, a 3×3 p.d. matrix. Its principal eigenvector captures the fiber orientation within the voxel.
- DWI signal along gradient direction \mathbf{u} :

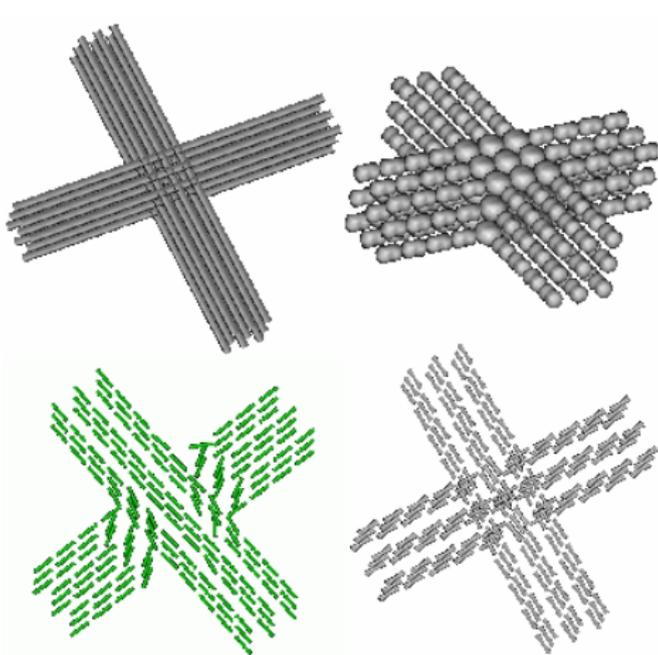
$$S(\mathbf{u}) = S_0 \exp(-b \mathbf{u}^T \mathbf{D} \mathbf{u}), \quad b = \frac{\gamma^2 \delta^2 \|\mathbf{q}\|^2 \Delta}{2}.$$

- $\mathbf{D}(\mathbf{s})$ can be recovered with as few as 6 distinct gradient directions.



Crossing Fibers

More than 30% voxels have multiple fiber bundles with distinct orientations (under D-MRI image resolution).



Limitations of Single Tensor Model

- Single tensor model can not resolve multiple fiber orientations within a voxel since a tensor only has one principal direction. It may incorrectly lead to:
 - Oblate ($\lambda_1 \approx \lambda_2 \gg \lambda_3$) tensor estimation.
 - Low anisotropy and random diffusion directions.
 - Consequently, early termination of fiber tracking or bias/switching of fiber tracking.
- Tensor Mixture Model.
- **Nonparametric methods using HARDI data.**

HARDI Techniques

High angular resolution diffusion imaging (HARDI) techniques enable the detection of multi-modal diffusion signals.

- Q-ball imaging: Gradients are sampled from a single spherical shell of a particular radius (a single b value).
 - Diffusion orientation distribution function (ODF) (Tuch, 2004, Descoteaux et al., 2007).
 - Fiber orientation density (FOD) function (Tournier et al., 2004, 2007).

Fiber Orientation Density Function

FOD is a symmetric p.d.f. on \mathbb{S}^2 which describes the distribution of fiber orientations (corresponding to coherently oriented fiber bundles) at a voxel.

- Example. K distinct fiber bundles:

$$F(\theta, \phi) = \sum_{k=1}^K w_k \delta_{\theta_k, \phi_k}(\theta, \phi), \quad \theta \in [0, \pi], \quad \phi \in [0, 2\pi),$$

where $w_k > 0$, $\sum_{k=1}^K w_k = 1$ are the volume fractions and θ_k (polar angle) and ϕ_k (azimuthal angle) are the spherical coordinates of the k -th fiber direction.

Assumptions

- DWI signals are the summation of signals originated from distinct fiber bundles.
 - No water exchange between distinct fiber bundles.
 - No water exchange between orientationally distinct segments of the same fiber bundle.
- Diffusion characteristics along all fiber bundles are (i) identical no matter the direction or abundance of the fiber bundle, and (ii) *axially symmetric* around the fiber direction.
 - DWI signal from a single coherently oriented fiber bundle can be represented by an axially symmetric *response function*.
 - The response function is identical across fiber bundles.

Response Function

- An axially symmetric kernel

$$R : [-1, 1] \rightarrow \mathbb{R}$$

which describes DWI signal resulting from water diffusion along a single fiber bundle aligned with the z-axis.

- Estimation of the response function. Assume response function is identical across voxels.
 - Fit the single tensor model to every voxel.
 - Identify voxels with high FA values and find their principal eigenvectors.
 - For each such voxel, rotate the DWI signals such that the principal eigenvectors are aligned with the z-axis.
 - Average the rotated DWI profiles across these voxels.

- Example. Gaussian diffusion with $\lambda_1 = \lambda_2 < \lambda_3$:

$$R(\cos(\theta)) = S_0 \exp^{-b\mathbf{q}(\theta,\phi)^T \Lambda \mathbf{q}(\theta,\phi)} = S_0 \exp^{-b(\lambda_1 \sin^2 \theta + \lambda_3 \cos^2 \theta)},$$

where $\theta \in [0, \pi]$, $\mathbf{q}(\theta, \phi) = (\sin \theta \cos \phi, \sin \theta \sin \phi, \cos \theta)^T$ and $\Lambda = \text{Diag}(\lambda_1, \lambda_2, \lambda_3)$.

- Since water diffuses fastest along the dominant fiber direction, the response function is attenuated the most along the z-axis.

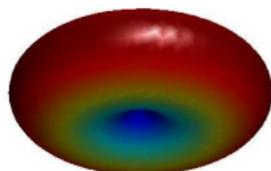


Figure: Gaussian diffusion response function with $\lambda_1 = \lambda_2 = 20, \lambda_3 = 1000$.

Spherical Convolution Model of Diffusion Signals

- DWI signal $S(\cdot)$ is the *spherical convolution* of the response function $R(\cdot)$ with the FOD $F(\cdot)$:

$$S(\mathbf{x}) = R \star F(\mathbf{x}) = \int_{\mathbb{S}^2} R(\mathbf{x}^T \mathbf{y}) F(\mathbf{y}) d\mathbf{y}, \quad \mathbf{x} \in \mathbb{S}^2.$$

- The FOD $F(\cdot)$ can be obtained by performing the *spherical deconvolution* of the response function $R(\cdot)$ from the DWI signal function $S(\cdot)$.
- Spherical deconvolution can be achieved through *spherical harmonic representation*.

Connection between the FOD model and the multi-tensor model.

- In the multi-tensor model, if the tensors D_k 's have the same set of eigenvalues satisfying $\lambda_1 = \lambda_2 < \lambda_3$, then it corresponds to the FOD model with:
 - Response function

$$R(\cos(\theta)) = S_0 \exp^{-b(\lambda_1 \sin^2 \theta + \lambda_3 \cos^2 \theta)}, \quad \theta \in [0, \pi],$$

- FOD

$$F(\theta, \phi) = \sum_{k=1}^K w_k \delta_{\theta_k, \phi_k}(\theta, \phi), \quad \theta \in [0, \pi], \quad \phi \in [0, 2\pi),$$

where (θ_k, ϕ_k) denotes the principal eigenvector of the tensor D_k ($k = 1, \dots, K$).

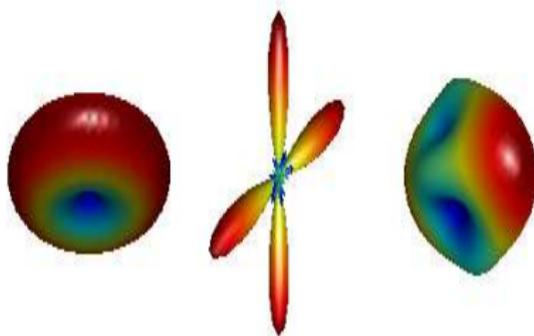


Figure: Spherical Convolution. Left to right: response function $R(\cdot)$, FOD $F(\cdot)$ and the DWI signal $S(\cdot)$.

Spherical Harmonics

- Spherical harmonics, denoted by $\tilde{\Phi}_{l,m}$:

$$\tilde{\Phi}_{l,m}(\theta, \phi) = \sqrt{\frac{2l+1}{4\pi} \frac{(l-m)!}{(l+m)!}} P_l^m(\cos(\theta)) \exp(im\phi), \quad \theta \in [0, \pi], \phi \in [0, 2\pi).$$

- $l (\geq 0)$ denotes the harmonic order and $m (-l \leq m \leq l)$ denotes the phase factor.
- Angular frequency increases with order l . Harmonics with even l are symmetric and those with odd l are anti-symmetric.
- P_l^m is an associated Legendre polynomial of order (l, m) .

Real Symmetric Harmonic Basis

- For $l = 0, 2, \dots, l_{\max}$ and $m = -l, \dots, 0, \dots, l$

$$\Phi_{l,m} = \begin{cases} \frac{\sqrt{2}}{2} (\tilde{\Phi}_{l,m} + (-1)^m \tilde{\Phi}_{l,-m}) & \text{if } 0 < m \leq l \\ \Phi_l^0 & \text{if } m = 0 \\ \frac{\sqrt{2}}{2i} ((-1)^{m+1} \tilde{\Phi}_{l,m} + \tilde{\Phi}_{l,-m}) & \text{if } -l \leq m < 0 \end{cases}$$

- $\Phi_{l,m}$ form an orthonormal basis for real symmetric square-integrable functions (including R and F) defined on \mathbb{S}^2 .

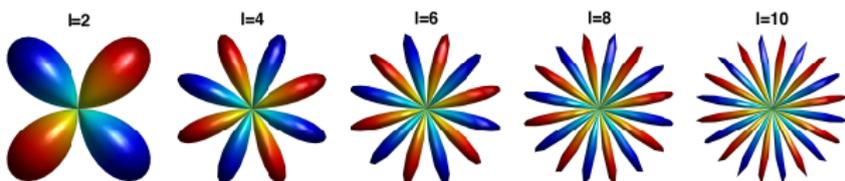


Figure: Real symmetric spherical harmonics.

- Since $S = R \star F$, so

$$s_{lm} = \sqrt{\frac{4\pi}{2l+1}} r_l f_{lm}, \quad l = 0, 2, 4, \dots, m = -l, \dots, 0, \dots, l,$$

where $s_{lm} = \langle S, \Phi_{l,m} \rangle$, $r_l = \langle R, \Phi_{l,0} \rangle$ and $f_{lm} = \langle F, \Phi_{l,m} \rangle$ are the spherical harmonics (rotational harmonics) coefficients of S , R and F , respectively.

- Assume that $S(\cdot)$, $R(\cdot)$, $F(\cdot)$ can be represented by finite-order spherical harmonics functions $\{\Phi_{l,m} : -l \leq m \leq l\}_{l=0,2,\dots,l_{\max}}$.
- The number of SH functions: $L = (l_{\max} + 1)(l_{\max} + 2)/2$.

Regression Model For DWI Measurements

The observed D-MRI signals:

$$\mathbf{y} = \mathbf{\Phi} \mathbf{R} \mathbf{f} + \boldsymbol{\epsilon}.$$

- $\mathbf{y} = (y(\theta_1, \phi_1), \dots, y(\theta_n, \phi_n))^T$ is the $n \times 1$ vector of observed DWI measurements.
- $\mathbf{\Phi}$ is the $n \times L$ matrix of the SH functions evaluations at the n gradient directions $\{(\theta_i, \phi_i)\}_{i=1}^n$.
- \mathbf{R} is an $L \times L$ diagonal matrix with diagonal elements $\sqrt{4\pi/(2l+1)}r_l$ (SH coefficients of the response function) in blocks of size $2l+1$ for $l = 0, 2, \dots, l_{\max}$.
- \mathbf{f} is the $L \times 1$ vector of SH coefficients of the FOD F .

SH-ridge Estimator of FOD

- Penalized regression:

$$\hat{\mathbf{f}} = \arg \min_{\mathbf{f}} \|\mathbf{y} - \Phi \mathbf{R} \mathbf{f}\|_2^2 + \lambda \mathbb{E}(F), \quad F = \sum_{l,m} f_{l,m} \Phi_{l,m}$$

- Laplace-Beltrami penalty, $\mathbb{E}(F)$, a measure of roughness.

$$\mathbb{E}(F) := \int_{\Omega} (\Delta_b F)^2 d\Omega = \mathbf{f}^T \mathbf{P} \mathbf{f},$$

where \mathbf{P} is a diagonal matrix with entries $l^2(l+1)^2$ in blocks of size $2l+1$.

- SH-ridge estimator:

$$\hat{\mathbf{f}}^{LB} = (\mathbf{R}^T \Phi^T \Phi \mathbf{R} + \lambda \mathbf{P})^{-1} \mathbf{R}^T \Phi^T \mathbf{y}, \quad \hat{F}^{LB} = \sum_{l,m} \hat{f}_{l,m}^{LB} \Phi_{l,m}.$$

sCSD Sharpening

Suppress small values of the estimated FOD and sharpens the peak(s) of the FOD estimator.

1. Initial step: Get an initial estimator $\hat{\mathbf{f}}_0$ by SH-ridge.
2. At the $k + 1$ updating step

$$\hat{\mathbf{f}}_{k+1} = \arg \min_{\mathbf{f}} \|\mathbf{y} - \Phi \mathbf{R} \mathbf{f}\|^2 + \lambda \|\mathbf{P}_k \mathbf{f}\|^2 \quad (1)$$

where \mathbf{P}_k is an $n \times L$ matrix,

$$\mathbf{P}_{k,i,(l,m)} := \begin{cases} \Phi_{i,(l,m)} & \text{if } \hat{F}_{k,i} < \tau \\ 0 & \text{if } \hat{F}_{k,i} > \tau \end{cases} \quad (2)$$

where $\hat{F}_{k,i}$ is the i the element of the vector $\hat{\mathbf{F}}_k = \Phi \hat{\mathbf{f}}_k$

Tournier et al.(2007).

Spherical Needlets

- Spherical needlets $\psi_{j,k}$ s are constructed from spherical harmonics functions (Nrcowich et al., 2006).
- Needlets are spatially localized with exponential concentration with respect to the frequency index j .
- Needlets provide sparse representations for spherical functions with small spatial scale features.

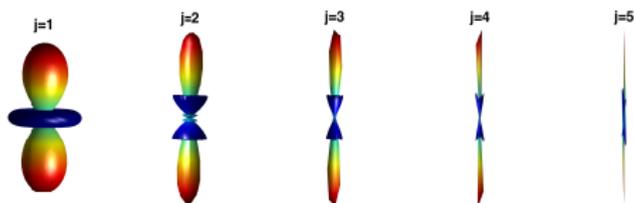


Figure: Real symmetric spherical needlets.

Needlets-lasso Estimator of FOD

- Assume that $S(\cdot)$, $R(\cdot)$, $F(\cdot)$ can be represented by finite-order spherical harmonics functions.
- Then they can be represented by finite needlets functions $\{\psi_{k,j} : k \in \chi_j\}_{j=0, \dots, j_{\max}}$.
- The number of symmetric needlets: $N = 2^{2j_{\max}+1} - 1$.
- There is a transition matrix \mathbb{C} such that the spherical harmonics coefficients

$$\mathbf{f} = \mathbb{C}\boldsymbol{\beta},$$

where $\boldsymbol{\beta}$ is the corresponding spherical needlets coefficients which are expected to be sparse.

- The observed D-MRI signals:

$$\mathbf{y} = \Phi \mathbf{R} \mathbb{C} \boldsymbol{\beta} + \boldsymbol{\epsilon}.$$

ℓ_1 penalized regression with nonnegativity constraints:

$$\hat{\beta} = \arg \min_{\beta: \tilde{\Phi} \mathbb{C} \beta \geq 0} \| \mathbf{y} - \Phi \mathbf{R} \mathbb{C} \beta \|_2^2 + \lambda \| \beta \|_1 .$$

- λ is a tuning parameter which controls the degree of sparsity.
- $\hat{F}^{NL} = \tilde{\Phi} \mathbb{C} \beta \geq 0$ ensures that the estimated FOD is nonnegative on the evaluation grid.
- This is a constrained convex minimization problem and can be solved by the ADMM algorithm.

Simulation Setting

- FODs with two distinct fiber bundles:

$$F(\theta, \phi) = w_1 \delta(\theta_1, \phi_1) + w_2 \delta(\theta_2, \phi_2).$$

- $w_1 = w_2 = 0.5$.
- Separation angle between the two fiber bundles:
 $\theta_{sep} = 45, 60, 75, 90$ degrees.
- Gaussian diffusion response function with $\frac{\lambda_3}{\lambda_1} = 50$.
- $n = 81$ gradient directions, sampled from an equal angle grid.
- $bvalue = 1000s/mm^2, 3000s/mm^2$.
- $SNR = \frac{S_0}{\sigma} = 20$, where S_0 is the b_0 image intensity and σ is the Rician noise standard deviation.

The number of gradient directions, SNR and bvalue are typical/close to those in real D-MRI studies.

Simulation Results

- FODs are estimated by SH-ridge with $l_{\max} = 8$ ($L = 45$ SH functions), by sCSD sharpening, and by needlets-lasso with $j_{\max} = 4$ ($N = 511$ needlets functions).
- Tuning parameters in SH-ridge and needlets-lasso are chosen by BIC. Those of sCSD are set as the recommended values by the paper.
- The needlets-lasso estimator has much sharper peaks.
- sCSD is sensitive to the penalty parameter τ as well as l_{\max} and thus is hard to automate in the real data setting where there are hundreds of thousands voxels with different fiber population characteristics.

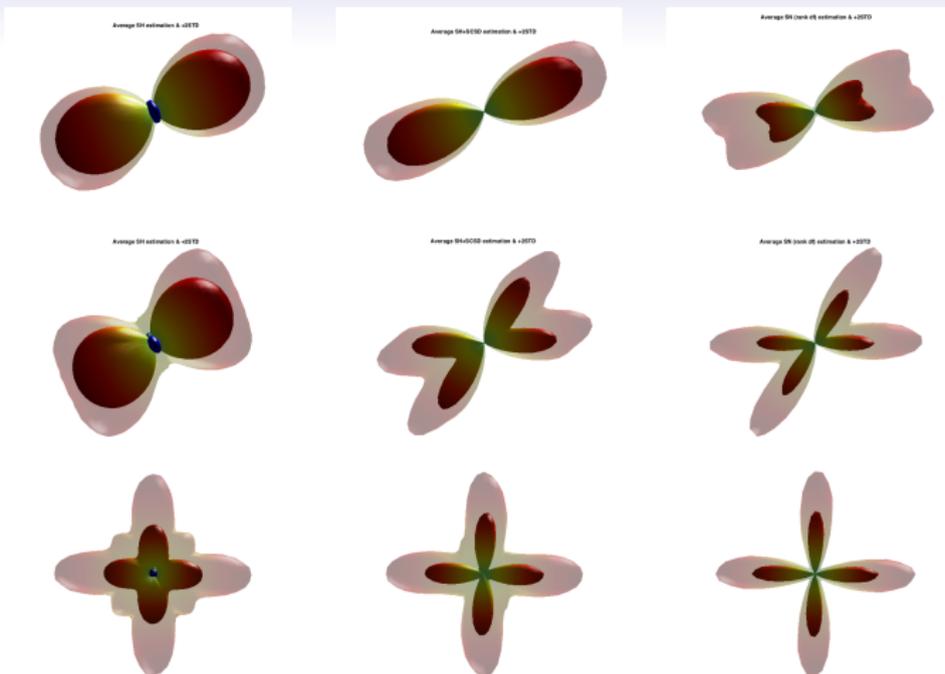


Figure: Mean plus 2-SD plots across 100 replicates. $bvalue=1000$. Top: $\theta_{sep} = 45$; Middle: $\theta_{sep} = 60$; Bottom: $\theta_{sep} = 90$. Left: SH-ridge; Middle: sCSD; Right: needlets-lasso.

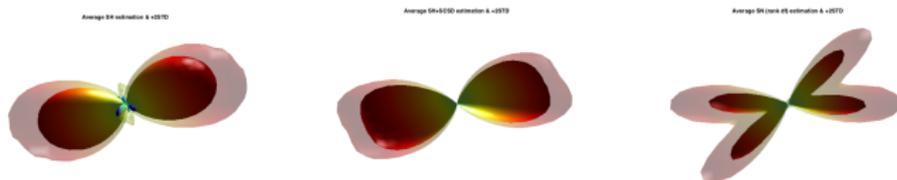


Figure: Mean plus 2-SD plots across 100 replicates. $bvalue=3000$. $\theta_{sep} = 45$. Left: SH-ridge; Middle: sCSD; Right: needlets-lasso.

Three fiber bundles simulation.

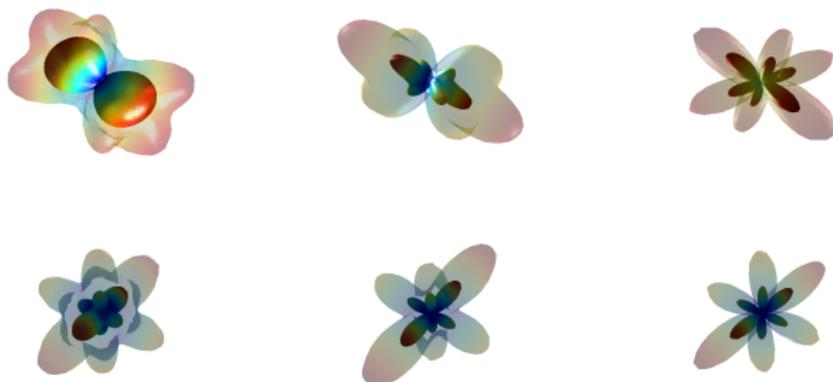


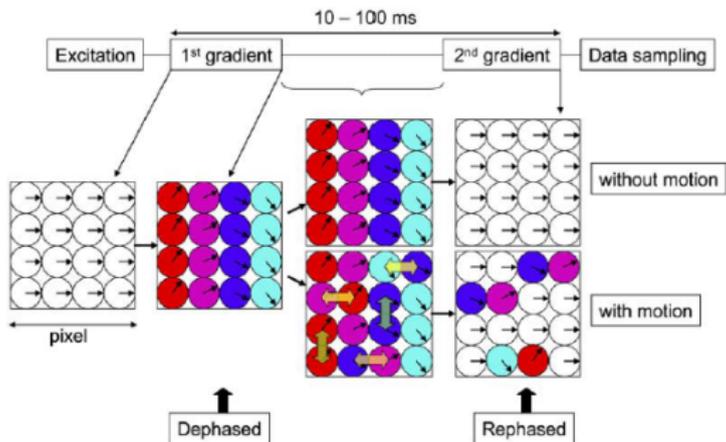
Figure: Mean plus 2-SD plots across 100 replicates. $bvalue=1000$. Top: $\theta_{sep} = 75$; Bottom: $\theta_{sep} = 90$. Left: SH-ridge; Middle: sCSD; Right: needlets-lasso.

Discussion

- Borrow information from neighboring voxels.
- Inference based on bootstrapping.
- Feature extraction and multiscale analysis.

Sensitize MRI Signal by Water Diffusion

- *Excitation*: Apply a strong homogeneous field \Rightarrow water molecules resonate at the same frequency and phase.
- *Dephasing* : Apply a linearly inhomogeneous gradient field \Rightarrow water molecules resonate at different frequencies depending on their locations.
- Apply the homogeneous field \Rightarrow water molecules resonate at the same frequency again, but signal is still out of phase.
- *Rephasing*: Apply an opposite gradient field.
 - If water molecules had moved along the gradient direction, then there would be a disruption of phase \Rightarrow signal loss and signals are *diffusion weighted*.



Mori and Zhang, 2006, Neuron 51

MRI can not measure the phase of individual water molecules, but it can detect imperfect rephasing through signal loss.

- During MR measurements, the amount of water molecular displacement is about $1 \sim 20 \mu m$, depending on sample, temperature, duration of experiment, etc.
- DT-MRI experiments are designed such that this amount of diffusion leads to $10 \sim 90\%$ signal loss.

Diffusion Weighted Signals

- “Dephasing” encodes the location information of water molecules through their signal phase.
- MR can not measure the phase of individual water molecules, but it can detect imperfect rephasing through signal loss.
 - Perfect rephasing only happens when water molecules remain stationary between the two applications of gradients.
 - If water moved between “dephasing” and “rephasing”, there will be a disruption of phase across the sample.
 - Then after rephasing, some of the molecules that moved will have different phases from the stationary molecules. This leads to an overall signal attenuation – signals are *diffusion weighted*.
- Water motion along directions perpendicular to the gradient direction will not cause signal loss and thus can not be detected.
 - Multiple gradient directions need to be applied if water diffuse anisotropically.

Model Diffusion Signals

- Signal loss equals to the summation (across locations within a voxel) of the sinusoid waves with shifted signal phases weighted by the proton density at their corresponding location.
- Applying a gradient field \mathbf{q} with duration δ introduces a *phase shift* in space:

$$\gamma\delta\mathbf{q} \cdot \mathbf{r}, \quad \mathbf{r} \in R^3$$

which is proportional to the projected distance on \mathbf{q} .

- $\mathbf{q} \in R^3$ – gradient field: $\|\mathbf{q}\|$ – field strength, $\mathbf{u} = \mathbf{q}/\|\mathbf{q}\|$ – gradient direction.
- $\mathbf{r} \in R^3$: displacement vector.
- δ : duration of dephasing/rephasing (assumed to be short, so ignore water movement during δ).
- γ : gyromagnetic ratio.

- After the “dephasing” stage, water molecules start to move during the time period Δt ($\gg \delta$). Their final locations are distributed according to a *diffusion probability density function*:

$$p_{\Delta t}(\mathbf{r}), \quad \mathbf{r} \in R^3$$

– density of protons having a displacement \mathbf{r} in time duration Δt .

- Δt : time between “dephasing” and “rephasing”. Typically, the molecular displacement is $1 \sim 20 \mu m$.
- The probability of water molecules having displacements \mathbf{r} and $-\mathbf{r}$ is the same:

$$p_{\Delta}(\mathbf{r}) = p_{\Delta}(-\mathbf{r}).$$

▶ back

Microscopic vs. Macroscopic

- The average distance that water molecules move during the MR measurement is $1 \sim 20\mu m$.
- Thus only barriers that have a smaller dimension may cause anisotropy in water diffusion, this includes microscopic cellular architecture ($< 10\mu m$) such as protein filaments, cell membranes, myelin sheaths.
- However, the image resolution is much coarser ($\sim 2mm$). Therefore information is averaged within each voxel.
- Thus **diffusion MRI provides information on “macroscopic coherent arrangement of anisotropic microscopic anatomy”** (Mori, 2007). Only when both factors exist in a voxel, one can observe diffusion anisotropy.

Spherical Needlets Construction

- Two main ideas: (i) discretization of the sphere by an exact quadrature formula; (ii) Littlewood-Paley decomposition.
- The quadrature formula discretizes the sphere into cubature points and cubature weights:

Theorem 1. Denote \mathcal{H}_l as the space spanned by $\{Y_{lm} : m = -l, \dots, l\}$, and let $\mathcal{K}_l = \bigoplus_{k=0}^l \mathcal{H}_k$. For any $l \in \mathbb{N}$, there exist a finite subset $\mathcal{X}_l = \{\xi_{lk} : k = 1, \dots, n_l\}$ of \mathbb{S}^2 and positive real numbers $\{\lambda_{lk} : k = 1, \dots, n_l\}$ such that

$$\int_{\mathbb{S}^2} f(x) dx = \sum_{k=1}^{n_l} \lambda_{lk} f(\xi_{lk}), \quad (7)$$

for any $f \in \mathcal{K}_l$. Here ξ_{lk} and λ_{lk} are called cubature points and cubature weights, respectively.

Spherical Needlets Construction (Cont'd)

- Given a frequency $j \in \mathbb{N}_0$ and cubature points ξ_{jk} and weights λ_{jk} , the spherical needlets with frequency j ($B > 1$):

$$\psi_{jk}(x) = \sqrt{\lambda_{jk}} \sum_{l=\lfloor Bj^{-1} \rfloor}^{\lceil Bj^{+1} \rceil} b\left(\frac{l}{B^j}\right) \sum_{m=-l}^l \tilde{\Phi}_{l,m}(\xi_{jk}) \bar{\tilde{\Phi}}_{l,m}(x), \quad x \in \mathbb{S}^2.$$

- $b(\cdot)$ is a window function satisfying: (i) $\text{supp}(b) = [1/B, B]$; (ii) $\sum_{j=0}^{\infty} b^2(t/B^j) = 1$, for any $t > 0$; (iii) $b \in C^{(M)}$ for some $M \geq 1$.
- $b(\cdot)$ decomposes the frequency domain into several overlapping intervals (B^{j-1}, B^{j+1}) .
- ξ_{jk} determines the location of the needlets ψ_{jk} and λ_{jk} determines to what extent ψ_{jk} is localized.
- Varying ξ_{jk} and j is analogous to translation and dilation in multiscale analysis.

Spherical Needlets Properties

- Needlets are real-valued spherical functions.
- They are localized in the frequency domain since the window function has compact support.
- Needlets are spatially localized with exponential concentration with respect to the frequency index j .
- Spherical needlets provide sparse representation of spherical functions with sharp local peaks.
- Needlets (together with the first spherical harmonics $\tilde{\Phi}_{00}$) form a tight frame on $L^2(\mathbb{S}^2)$:

$$\|f\|_{L^2}^2 = \sum_{j,k} |\langle f, \psi_{jk} \rangle|^2 + a_{00}^2.$$

- They are almost orthogonal: for $|j - j'| \geq 2$, $\langle \psi_{jk}, \psi_{j'k'} \rangle = 0$.
- The spherical needlets ψ_{jk} and $\psi_{j'k'}$ are asymptotically uncorrelated as the frequency j increases and the distance between them remains fixed.