

An Introduction to Diffusion Tensor Image Analysis

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KEYWORDS

- Diffusion Tensor MRI • DTI • Brain imaging
- Tractography • Review

Diffusion tensor magnetic resonance imaging (DTI) is a relatively new technology that is popular for imaging the white matter of the brain. This article provides a basic and broad overview of DTI to enable the reader to develop an intuitive understanding of these types of data, and an awareness of their strengths and weaknesses. The authors have tried to include equations for completeness, but these are not necessary for understanding the content. Wherever possible, more in-depth technical articles or books are suggested for further reading. The authors especially recommend the new diffusion MRI textbook,¹ the introductory paper on fiber tracts and tumors,² the white matter atlas book,³ and the review of potential pitfalls in DTI analysis.⁴ The remainder of this article addresses basic questions about DTI (the what, why, and how of DTI), followed by a discussion of issues in interpretation of DTI, and finally an overview of more advanced diffusion imaging methods and future directions.

WHY DTI? A BRIEF HISTORY OF DTI AND ITS IMPACT ON CLINICAL RESEARCH

The diffusion tensor was originally proposed for use in MRI by Peter Basser in 1994.^{5,6} Before DTI, diffusion MRI^{7,8} had developed from research in diffusion nuclear magnetic resonance.⁹ Before

the diffusion tensor model was introduced, the orientation of the axons in a tissue sample had to be known to measure anisotropic diffusion, and therefore only fixed samples could be scanned, such as the axon of the giant squid.¹⁰ The introduction of the diffusion tensor model allowed, for the first time, a rotationally invariant description of the shape of water diffusion. The invariance to rotation was crucial because it enabled application of the DTI method to the complex anatomy of the fiber tracts in the human brain.¹¹ However, the diffusion tensor is not able to fully describe crossing of the fiber tracts.^{12,13}

The popularity of DTI has been enormous. It has been applied to a tremendous variety of neuroscientific studies,^{14–16} including schizophrenia,¹⁷ traumatic brain injury,¹⁸ multiple sclerosis,^{19,20} autism,²¹ and aging.²² Anatomic investigations have been undertaken regarding the structure of the language network,^{23,24} the asymmetry of the white matter in twins and siblings,²⁵ and the location, asymmetry, and variability of the fiber tracts.²⁶ Recent investigations have attempted to model the human “connectome” by analyzing structural versus functional brain connectivity as measured with DTI and functional MRI.^{27,28} DTI has also been applied for neurosurgical planning and navigation.^{29–32} A large prospective study showed that addition of preoperative DTI to

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neuronavigation increased tumor resection and survival and decreased neurologic morbidity.³³

WHAT IS DTI?

DTI is a sensitive probe of cellular structure that measures the diffusion of water molecules. The measured quantity is the diffusivity or diffusion coefficient, a proportionality constant that relates diffusive flux to a concentration gradient⁸ and has units of $\frac{mm^2}{s}$. Unlike the diffusion in a glass of pure water, which would be the same in all directions (isotropic), the diffusion measured in tissue varies with direction (is anisotropic). The measured macroscopic diffusion anisotropy is the result of microscopic tissue heterogeneity.⁶ In the white matter of the brain, diffusion anisotropy is primarily caused by cellular membranes, with some contribution from myelination and the packing of the axons.^{11,34,35} Anisotropic diffusion can indicate the underlying tissue orientation (Fig. 1).

The diffusion tensor describes the diffusion of water molecules using a Gaussian model. Technically, it is proportional to the covariance matrix of a three-dimensional Gaussian distribution that models the displacements of the molecules. The tensor is a 3×3 symmetric, positive-definite matrix, and these matrix properties mean that it has three orthogonal (mutually perpendicular) eigenvectors and three positive eigenvalues. The major eigenvector of the diffusion tensor points in the principal diffusion direction (the direction of the fastest diffusion). In anisotropic fibrous tissues, the major eigenvector also defines the fiber tract axis of the tissue,⁶ and thus the three orthogonal eigenvectors can be considered a local fiber coordinate system. (This interpretation is only strictly true in regions where fiber tracts do not cross, fan, or branch.) The three positive eigenvalues of the tensor ($\lambda_1, \lambda_2, \lambda_3$) give the diffusivity in the direction of each eigenvector. Together, the eigenvectors and eigenvalues define an ellipsoid that

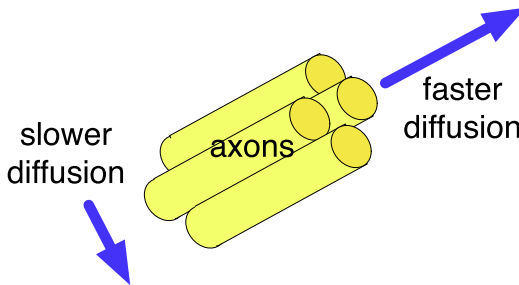


Fig. 1. Illustration of anisotropic diffusion, in the ideal case of a coherently oriented tissue. This example compares the diffusion measured parallel and perpendicular to the axons in a white matter fiber tract.

represents an isosurface of (Gaussian) diffusion probability: the axes of the ellipsoid are aligned with the eigenvectors and their lengths are $\sqrt{2\tau\lambda_i}$.⁶ Fig. 2 shows three diffusion tensors chosen from different regions of the human brain to illustrate possible shapes of the ellipsoid.

HOW IS DTI MEASURED?

To measure diffusion using MRI, magnetic field gradients are used to create an image that is sensitized to diffusion in a particular direction. Through repeating this process of diffusion weighting in multiple directions, a three-dimensional diffusion model (the tensor) can be estimated. In simplified terms, diffusion imaging introduces extra gradient pulses whose effect “cancels out” for stationary water molecules, and causes a random phase shift for molecules that diffuse. Because of their random phase, signal from diffusing molecules is lost. This loss of signal creates darker voxels (volumetric pixels), meaning that white matter fiber tracts parallel to the gradient direction will appear dark in the diffusion-weighted image for that direction (Fig. 3).

Next, the decreased signal (S_k) is compared with the original signal (S_0) to calculate the diffusion tensor (D) by solving the Stejskal-Tanner Equation (1).³⁶ This equation describes how the signal intensity at each voxel decreases in the presence of Gaussian diffusion:

$$S_k = S_0 e^{-b\hat{g}_k^T D \hat{g}_k} \quad (1)$$

In this equation, S_0 is the original image intensity at the voxel (measured with no diffusion-sensitizing gradient) and S_k is the intensity measured after the application of the k th diffusion-sensitizing gradient in the (unit) direction \hat{g}_k . The product $\hat{g}_k^T D \hat{g}_k$ represents the diffusion coefficient (diffusivity) in direction \hat{g}_k . Because the entire set of diffusion-weighted images is used (giving many

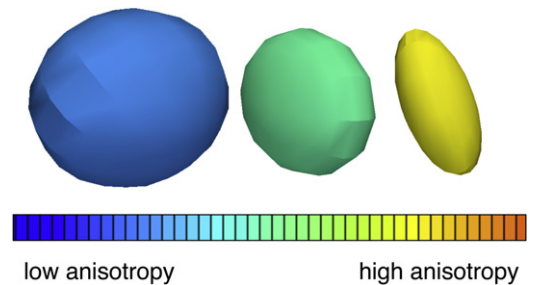


Fig. 2. Three example diffusion tensors, selected from a diffusion tensor magnetic resonance imaging scan of the human brain to illustrate differences in tensor anisotropy and orientation.

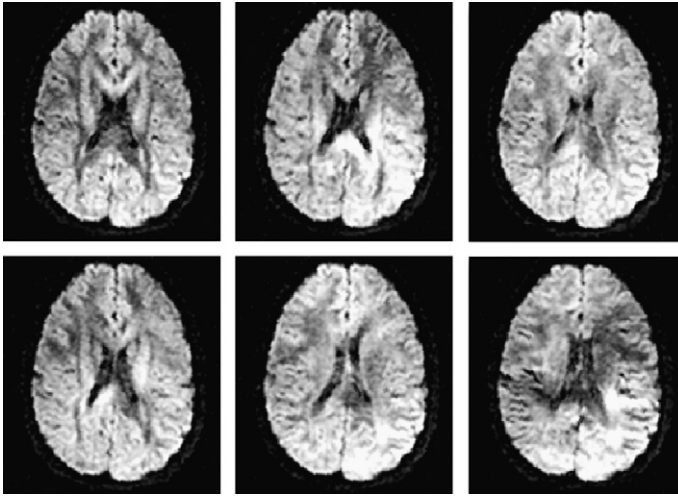


Fig. 3. Six diffusion-weighted images (the minimum number required for tensor calculation). In diffusion MRI, magnetic field gradients are used to sensitize the image to diffusion in a particular direction. The direction is different for each image, resulting in a different pattern of signal loss (*dark areas*) from anisotropic diffusion.

values for S_k and \hat{g}_k), this is actually a system of equations that is solved for D , the diffusion tensor. To calculate the six independent numbers in the 3×3 symmetric matrix D , at least seven images are needed: six diffusion-weighted images from six gradient directions (giving six values for S_k) plus one baseline image (giving S_0). However, in clinical research today, a higher number of images are almost always used. The above system of equations can be solved via the least squares method at each voxel.

Equation (1) also contains b , LeBihan's⁹ factor describing the pulse sequence, gradient strength, and physical constants. The b -factor is near $0 \frac{s}{mm^2}$ for the image S_0 , which is T2-weighted, and the b -factor is near $1000 \frac{s}{mm^2}$ for the diffusion-weighted images S_k in DTI. For rectangular gradient pulses, the b -factor is defined by $b = \gamma^2 \delta^2 (\Delta - \frac{\delta}{3}) |g|^2$, where γ is the proton gyromagnetic ratio (42 MHz/Tesla), $|g|$ is the strength of the diffusion sensitizing gradient pulses, δ is the duration of the diffusion gradient pulses, and

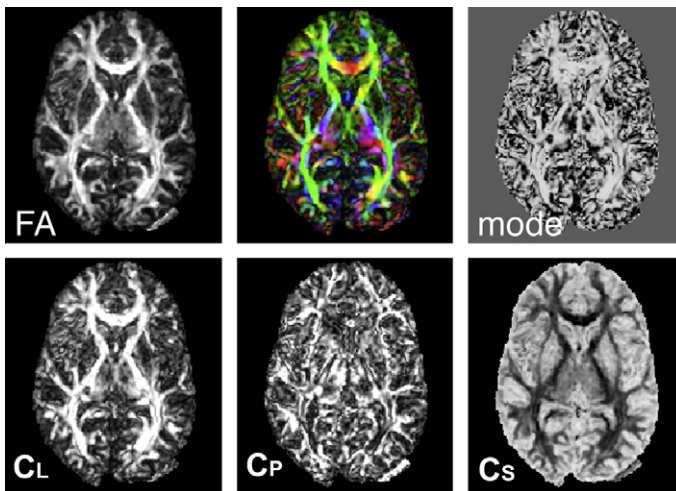


Fig. 4. Scalar measures derived from DTI include FA, mode, C_L , C_P , and C_S . Also shown (*top row, middle*) is a mapping of the major eigenvector orientation to colors. See the text for more information about the definition of these measures.

Δ is the time between diffusion gradient RF pulses.³⁷

Information on the MR physics of DTI⁸ and more information on the tensor calculation process^{5,37} are provided elsewhere. Goodlett and colleagues³⁸ provide a comparison of tensor calculation methods (including least squares and weighted least squares) in the presence of noise.

HOW IS DTI DISPLAYED?

DTI is usually displayed through condensing the information contained in the tensor (**Fig. 4**) into either one number (a scalar), or four (to give an RGB color and a brightness value). The diffusion tensor can also be viewed using glyphs, which are small three-dimensional representations of the major eigenvector or whole tensor. Finally, DTI is often viewed through estimating the course of white matter tracts through the brain using a process called *tractography*.

Scalars Derived from DTI

This section describes commonly used scalar quantities, which can be divided into two categories: diffusion magnitude measures and anisotropy measures. The eigenvalues of the symmetric, positive-definite diffusion tensor D are referred to using $\lambda_1 \geq \lambda_2 \geq \lambda_3 \geq 0$. Pierpaoli and colleagues¹¹ originally measured and compared several scalar measures, and the eigenvalues, in different regions of the human brain.

Measures of diffusion magnitude

The simplest and possibly most useful scalar is the average of the tensor's eigenvalues. This average may be referred to as the *mean diffusivity* (MD)³⁹; the bulk mean diffusivity ($\langle D \rangle$)⁴⁰; or the apparent diffusion coefficient (ADC) map. In clinical imaging, ADC maps may be measured using fewer diffusion gradients than needed for the tensor. A similar quantity to the MD is the sum of the eigenvalues, called the *trace* of the tensor.

The trace and MD relate to the total amount of diffusion in a voxel, which is related to the amount of water in the extracellular space. The trace is clinically useful in early stroke detection because it is sensitive to the initial cellular swelling (cytotoxic edema) that restricts diffusion.⁴¹ In the normal human brain, the trace is high in cerebrospinal fluid, around $9.6 \times 10^{-3} \frac{mm^2}{s}$, and relatively constant in normal brain parenchyma (white and gray matter), between $1.95 \times 10^{-3} \frac{mm^2}{s}$ and $2.2 \times 10^{-3} \frac{mm^2}{s}$.¹¹ For comparison, the self-diffusion coefficient of water (the diffusivity measured in pure water without any tissue) at a body temperature of 37°C is $3 \times 10^{-3} \frac{mm^2}{s}$,⁴²

which would give a trace of $9 \times 10^{-3} \frac{mm^2}{s}$. The MD and trace measured in ventricles or in edema can be higher than in water because of fluid flow or enhanced perfusion, respectively.⁷

Measures of diffusion anisotropy

Tensor anisotropy measures are ratios of the eigenvalues that are used to quantify the shape of the diffusion. These measures are useful for describing the amount of tissue organization and for locating voxels likely to contain a single white matter tract (without crossing or fanning). The following measures are normalized and all range from 0 to 1, except for the mode, which ranges from -1 to +1.

The fractional anisotropy (FA)⁴³ is the most widely used anisotropy measure. Its name comes from the fact that it measures the fraction of the diffusion that is anisotropic. This measure can be considered the difference of the tensor ellipsoid's shape from that of a perfect sphere. FA is basically a normalized variance of the eigenvalues:

$$FA = \frac{1}{\sqrt{2}} \frac{\sqrt{(\lambda_1 - \hat{\lambda})^2 + (\lambda_2 - \hat{\lambda})^2 + (\lambda_3 - \hat{\lambda})^2}}{\sqrt{\lambda_1^2 + \lambda_2^2 + \lambda_3^2}} \quad (2)$$

where $\hat{\lambda}$ is the mean diffusivity. FA is often considered a measure of white matter integrity, although changes in FA may be caused by many factors.

Three intuitive measures are C_L , C_P , and C_S , which are the linear, planar, and spherical shape measures.^{37,44} They describe whether the shape of diffusion is similar to a cigar (linear), pancake (planar), or sphere (spherical).

$$C_L = \frac{\lambda_1 - \lambda_2}{\lambda_1} \quad (3)$$

$$C_P = \frac{\lambda_2 - \lambda_3}{\lambda_1} \quad (4)$$

$$C_S = \frac{\lambda_3}{\lambda_1} \quad (5)$$

In voxels with high planar or spherical measure, the principal eigenvector will not always match an underlying fiber tract direction (where tracts cross the eigenvector may point to neither one). However, if the largest eigenvalue is much larger than the other two eigenvalues, the linear measure will be large, indicating the presence of a single fiber tract. These measures can be normalized by λ_1 , by the trace, or by $\sqrt{\lambda_1^2 + \lambda_2^2 + \lambda_3^2}$.

Although FA measures how far the tensor is from a sphere, another complementary measure discriminates between linear and planar anisotropy. This information is given by the mode,

a quantity that is mathematically orthogonal to the FA measure and relates to the skewness of the eigenvalues.⁴⁵

mode

$$= \frac{(-\lambda_1 - \lambda_2 + 2\lambda_3)(2\lambda_1 - \lambda_2 - \lambda_3)(-\lambda_1 + 2\lambda_2 - \lambda_3)}{2(\lambda_1^2 + \lambda_2^2 + \lambda_3^2 - \lambda_1\lambda_2 - \lambda_1\lambda_3 - \lambda_2\lambda_3)^{3/2}} \quad (6)$$

The parallel diffusivity measure, also called the *axial diffusivity*, is equal to the largest eigenvalue. The perpendicular diffusivity measure, also called the *radial diffusivity*, is equal to the average of the two smaller eigenvalues. These measures are interpreted as diffusivity parallel and perpendicular to a white matter fiber tract, and therefore make the most sense in regions of coherently oriented axons with no fiber crossings.

Often in scientific studies, the reported measures from the diffusion tensor are not independent. However, complete sets of orthogonal (mathematically independent) scalars have been defined.^{45,46}

Colors Derived from DTI

Another type of image can represent the major eigenvector field using a mapping to colors (Fig. 5). The color scheme most commonly used to represent the orientation of the major eigenvector works as follows: blue is superior–inferior, red is left–right, and green is anterior–posterior.⁴⁷ To enhance visualization of the white matter and suppress information outside of it, the brightness of the color is usually controlled by tensor anisotropy (FA).

Glyphs Derived from DTI

Small three-dimensional objects called *glyphs* can be used to display information from each tensor

eigensystem. Example glyphs include “sticks” representing the orientation of the major eigenvector, ellipsoids related to the diffusion isoprobability surfaces,⁶ and superquadric tensor glyphs.⁴⁸

Tractography

The word *tractography* refers to any method for estimating the trajectories of the fiber tracts in the white matter. Ciccarelli and colleagues¹⁵ provide a clinical and technical overview of tractography in neurologic disorders. Tractography techniques, including explanations of common tractography artifacts and a comparison of methods, are reviewed elsewhere.^{49,50} Many methods have been proposed for tractography, and the results will vary enormously depending on the chosen method.

The most common approach is streamline tractography (Fig. 6),^{51–54} which is closely related to an earlier method for visualization of tensor fields, known as *hyperstreamlines*.⁵⁵ This method produces as output discrete curves or trajectories that are also referred to by terms such as *tracts*, *fibers*, and *traces*. The streamline tract-tracing approach works by successively stepping in the direction of the principal eigenvector (the direction of fastest diffusion). The eigenvectors are thus tangent to the trajectory that is produced. A fixed step size of 1 mm or less (smaller than a voxel) is generally used for DTI data.

Several computational methods can be used to perform basic streamline tractography, including Euler’s method (following the eigenvector or tangent for a fixed step size), second-order Runge-Kutta (also known as the *midpoint method*, in which the tangent is followed for half a step, then a new tangent is calculated at the midpoint of the interval and used to take the full step), and

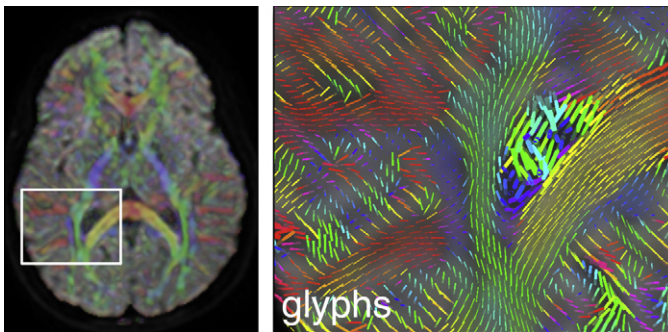


Fig. 5. An example using glyphs and colors for diffusion tensor magnetic resonance imaging visualization. On the left an axial image plane, showing the average diffusion-weighted image with semitransparent color overlay indicating the major eigenvector orientation, and a white square indicating the zoomed-in area (*right image*). In both images, the color red indicates right–left orientation, blue is superior–inferior, and green is anterior–posterior. The right image contains glyphs representing major eigenvector orientations (and scaled by the largest eigenvalue) in the region of the corpus callosum (*yellow and red*) and right lateral ventricle. The cingulum can be seen in blue, and the posterior limb of the internal capsule in green.

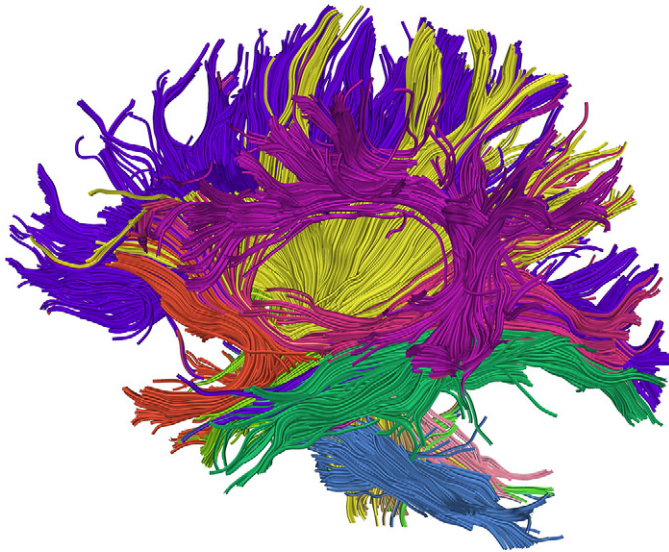


Fig. 6. Whole-brain streamline DTI tractography. Colors were assigned automatically according to an atlas-based tractography segmentation method. (Data from O'Donnell LJ, Westin CF. Automatic tractography segmentation using a high-dimensional white matter atlas. *IEEE Trans Med Imaging* 2007;26:1562–75.)

fourth-order Runge-Kutta (in which the weighted average of four estimated tangents to the curve is used when taking each step).⁵⁶ Basser and colleagues⁵¹ and Conturo and colleagues⁵² explored application of the Euler and Runge-Kutta methods to white matter tractography. Another popular method is called fiber assessment by continuous tracking.⁵³ Some related methods attempt to introduce inertia when tracking through regions of planar anisotropy (likely fiber crossings). These methods modulate the incoming tangent direction by the tensor instead of directly using the major eigenvector of the tensor.^{37,54,57,58}

Processing DTI data to display fiber tracts of interest requires expert knowledge or an automatic algorithm. After performing streamline tractography, the fiber trajectories of interest can be interactively selected through virtual dissection, in which inclusion/exclusion regions are defined and used to select trajectories.^{3,59,60} Automated methods for atlas-based tractography segmentation, which use prior knowledge to select trajectories, have also been developed.^{61–67}

In addition to streamline tractography, many other methods are available,^{49,50} including probabilistic tractography that outputs connection strengths or probabilities,^{68,69} optimization methods that use graph theory or physical models,^{70,71} region-growing and wavefront evolution methods,⁷² tractography using advanced models for fiber crossings,^{73–76} and tractography meta-analysis methods that perform clustering or fit more sophisticated tract models.^{61,77–79}

Tractography methods can produce false-positive and -negative results (discussed in next section); however, clinical validations of streamline tractography have shown accurate reconstructions (true-positive results). Tract end points, especially of the corticospinal or motor tract, have been compared with electrocortical stimulation during neurosurgery^{80,81} with good correspondence. In a study of 238 neurosurgical patients with gliomas involving the motor tract who were randomized to study or control groups, DTI was shown to increase survival and reduce postoperative motor deficits.³³

ISSUES IN INTERPRETING DTI DATA

This section presents two issues that are relevant to the clinical interpretation or meaning of the DTI data. Jones and Cercignani⁴ have provided a more thorough discussion, including additional information about scanning and processing pitfalls.

Scale of Diffusion Measurements Versus Axons

The measured diffusion effects are averaged over a voxel (three-dimensional pixel), complicating the biophysical interpretation of the diffusion tensor.^{40,82} For example, in scientific studies FA is often interpreted as “white matter integrity”; however, many factors (eg, cell death, change in myelination, increase in extracellular or

Table 1
The scale of diffusion tensor magnetic resonance imaging and the brain: neuron sizes and quantities, and water diffusion times and distances

Quantity	Measurement	References
Axon packing density (pyramidal tract)	60,000–70,000/mm ²	11
Axon packing density (corpus callosum)	338,000/mm ²	11
Axon diameter (pyramidal tract)	26 μm	11
Axon diameters in central nervous system	0.2–20 μm	83
Neuron cell body diameter	≥50 μm	83
Typical voxel size in diffusion MRI	2.5 × 2.5 × 2.5 mm	
Typical diffusion time in diffusion tensor magnetic resonance imaging	30–100 ms	34,42
Mean water diffusion distance	1–15 μm (in 50–100 ms)	42
Number of neurons in human brain	100 billion (10 ¹¹)	83
Synaptic connections per axon	Up to 1,000	83

intracellular water) may cause changes in FA. Overall, this difficulty in interpreting DTI is because the scale at which diffusion is measured with DTI is very different from the size scale of individual axons. To illustrate the complexity of the human brain and the size/time scales of the diffusion imaging experiment, **Table 1** lists relevant quantities, such as the number of neurons in the brain (10¹¹) and the distance over which water diffuses during an imaging experiment (1–15 μm, a distance similar to the diameter of an axon).

False-Positive or -Negative Tractography

Because the tensor model is only able to represent one major fiber direction in a voxel, DTI tractography can be confounded by regions of crossing fibers, as shown schematically in **Fig. 7**. A significant fraction of white matter voxels in the brain contain multiple fiber bundles oriented in different directions, where the diffusion tensor model is not reliable.⁷³ Other factors can confound tractography. Partial volume effects, in which two types of

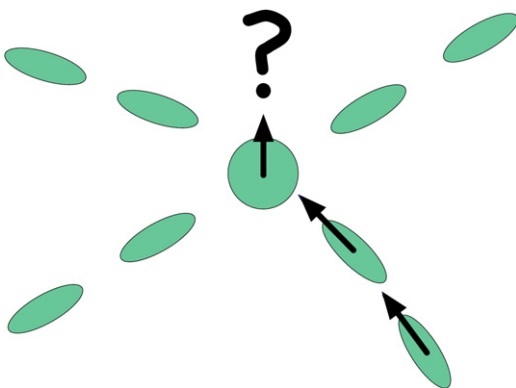


Fig. 7. The major eigenvector may not be aligned with a fiber tract in the case of crossing fibers.

tissue are present in a voxel, can produce a tensor that represents neither tissue well.⁸² Also, crossing, “kissing,” and “fanning” fiber tracts¹² are not represented well at the voxel level by the diffusion tensor. Finally, in standard streamline tractography, all decisions are made locally, and therefore errors can accumulate.

These issues may cause false-positive and -negative connections (**Figs. 8** and **9**). Common false-positive connections include trajectories from the corona radiata that cross the corpus callosum, and trajectories from the corona radiata that cross at the pons and ascend in the corona radiata of the other hemisphere. Common false-negatives with DTI tractography are the lateral lip/hand connections of the corticospinal tract^{49,73} and lateral connections of the corpus callosum.⁷⁴

The particular errors depend strongly on the tractography algorithm used, and on the type of diffusion data used (DTI vs higher-order models). However, no perfect method exists, and perfect tractography is unlikely to be possible because even perfect diffusion MRI data would not solve the peak selection problem. At each step, a direction must be chosen to follow for the next step, and with a more detailed model than the tensor, this requires some logical heuristic, such as choosing the closest direction to the current direction.

ADVANCES BEYOND THE DIFFUSION TENSOR

New diffusion models, scanning paradigms, and analysis methods are continually being developed for diffusion MRI.

High angular resolution diffusion imaging (HARDI) includes methods that acquire diffusion data using many more than 6 diffusion directions (eg, ≥ 32).⁸⁴ These methods generally use a higher

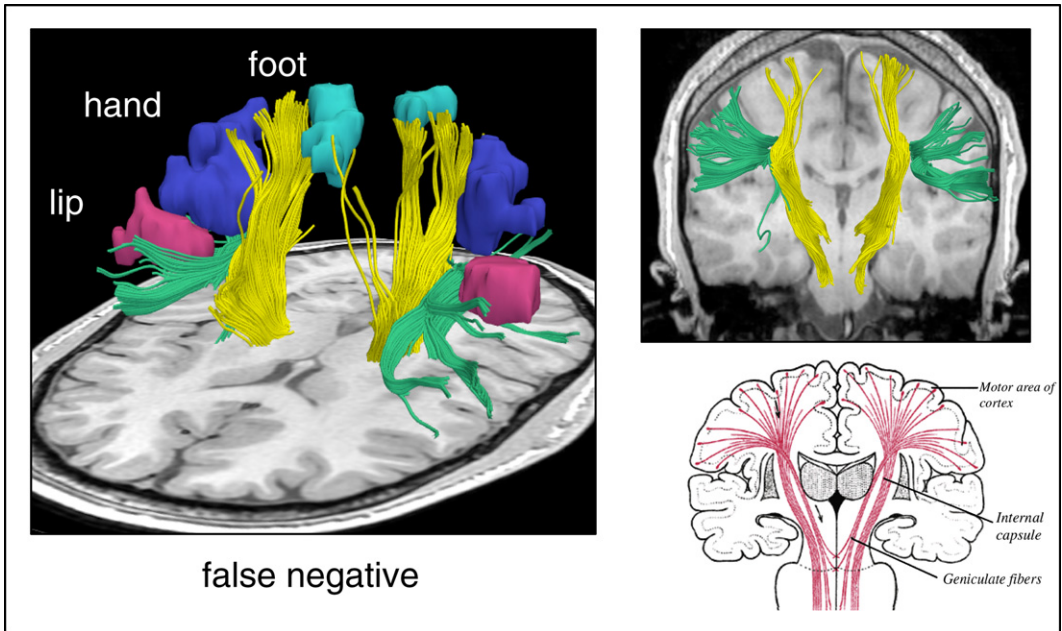


Fig. 8. Example false-negative streamline tractography error. The motor fibers (yellow) do not reach all functional magnetic resonance (fMRI) motor activations (cyan, blue, and pink) due to the presence of the superior longitudinal fasciculus (green) that runs perpendicular to the motor tract. In the right column are coronal views of the typical streamline tractography result (top) and expected anatomy (bottom).

b-value than the standard 1000 for DTI, or multiple b-values (multiple “shells” of data). Multi-shell acquisitions enable description of the full diffusion function using measures such as displacement, zero-probability, and kurtosis that are highly sensitive to myelin.^{85,86} Another type of multiple b-value acquisition is diffusion spectrum imaging (DSI).⁸⁷ Diffusion models that go beyond DTI have been proposed to extract important biomarkers, such as compartmentalization^{13,82,88,89} and axon

diameter.⁹⁰ Higher-rank tensor models have been proposed to extend DTI.⁹¹ Use of multiple pairs of diffusion gradients (double pulsed field gradient diffusion MRI) has been shown to increase sensitivity to small size scales.^{92,93} Diffusion MRI data analysis has benefited from the introduction of novel tractography methods, many types of white matter atlas,^{61–67} advanced tract-based quantification methods,^{77–79,94} new visualization methods,⁹⁵ and new scalar measures.^{45,96}



Fig. 9. False-positive streamline tractography error. In the left image, fibers (yellow with black dotted line) have traced parts of two anatomic structures by incorrectly crossing from one to the other (at arrow). In the right image, both structures (arcuate fasciculus in magenta and corona radiata in yellow) can be seen.

SUMMARY

DTI is an increasingly prevalent and popular imaging modality that has been applied to numerous scientific studies and clinical problems since its invention a little more than 15 years ago. The field is expected to benefit from many future advances in diffusion imaging and analysis.

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