

# **Diffusion MRI: Principles and Applications**

# Outline

## ■ Basics of diffusion

- ▶ What is “diffusion” and why it is important
- ▶ Diffusion MRI in a nutshell

## ■ Main applications

- ▶ Characterization of *tissue properties*
- ▶ Estimation of *tissue structure*
- ▶ Reconstruction of *neuronal fiber tracts*
- ▶ Estimation of *structural brain connectivity*

## ■ Intro to local reconstruction techniques

- ▶ Diffusion Spectrum Imaging (DSI)
- ▶ Diffusion Tensor Imaging (DTI)
- ▶ Q-BALL Imaging (QBI)
- ▶ Constrained Spherical Deconvolution (CSD)

# What is diffusion?

- **Random movement of molecules** due to thermal agitation from regions of high to regions of low concentration
  - ▶ *EXAMPLE:* in a glass of water, molecules diffuse randomly and freely, only constrained by the boundaries of the container

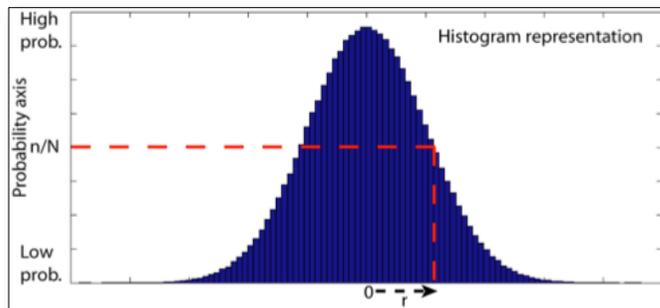


# What is diffusion?

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  - ▶ *EXAMPLE:* in a glass of water, molecules diffuse randomly and freely, only constrained by the boundaries of the container
- First noted by **Robert Brown** in 1828
  - ▶ “...random motion without any apparent cause...”
- Formally described by **Albert Einstein** in 1905

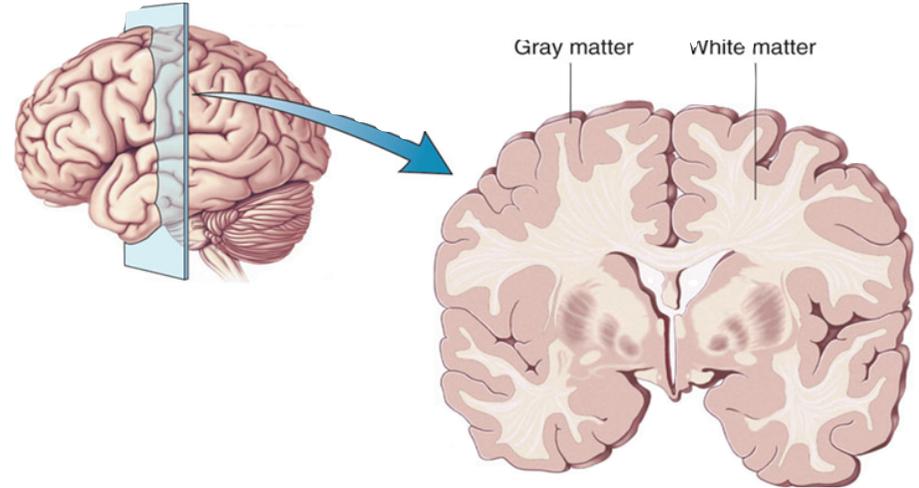
mean displacement of molecules ←  $\langle r^2 \rangle \propto t D$  → coefficient of diffusion of the medium

observation time

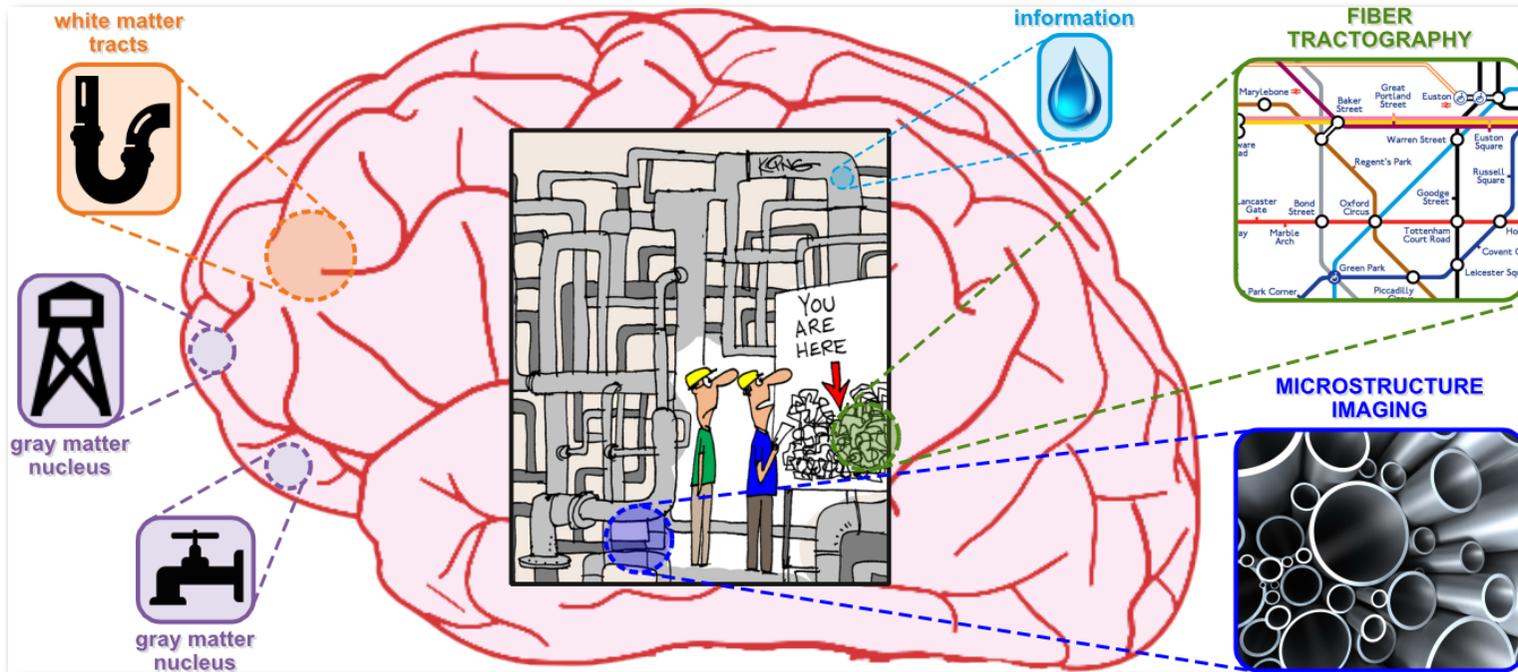


## ■ Rough brain anatomy:

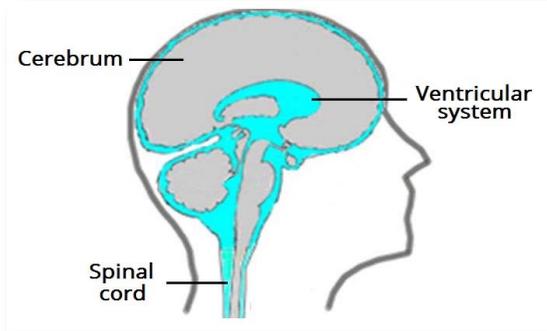
- ▶ *gray-matter* : neuronal cell bodies
- ▶ *white-matter* : mainly myelinated tracts
- ▶ *Cerebrospinal fluid (CSF)*



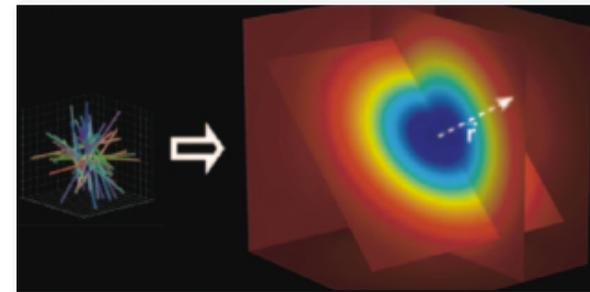
## ■ Metaphor: “connectivity” as “water supply network”



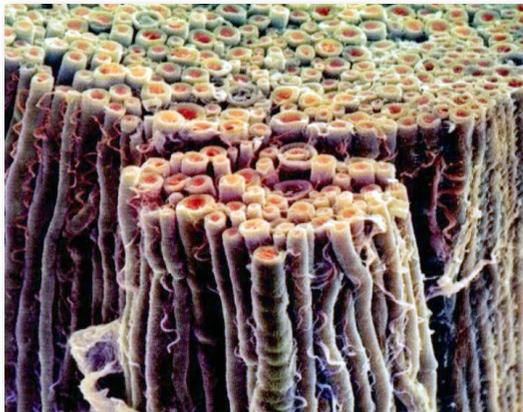
## ■ Cerebrospinal Fluid (CSF)



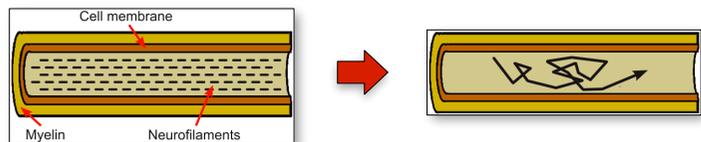
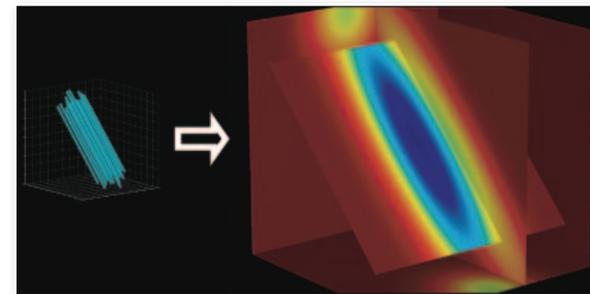
- ▶ Displacements are *isotropic*
- ▶ *Variance* depends on the fluid's properties



## ■ Neuronal tracts (white-matter)



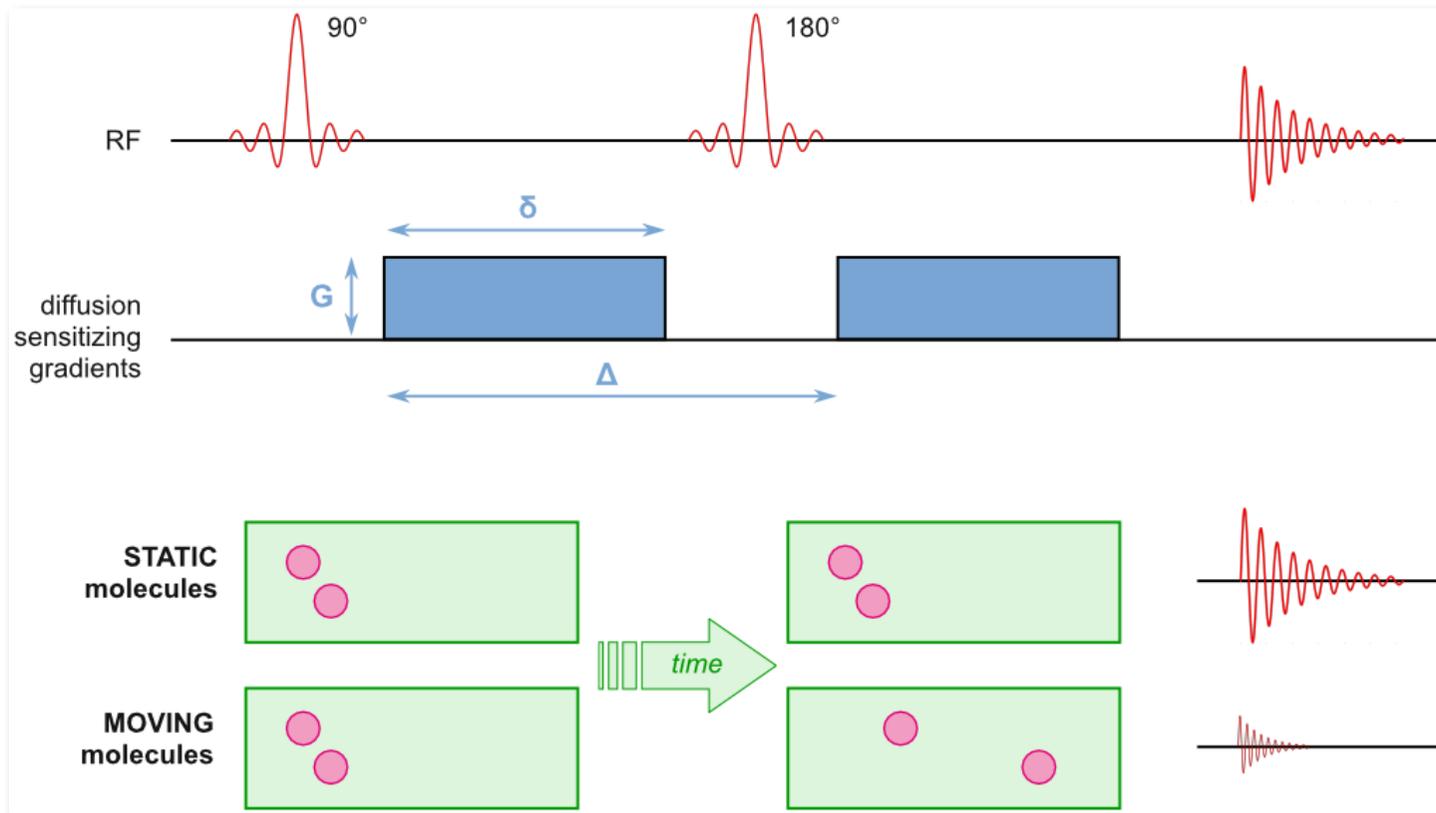
- ▶ Diffusion more *restricted* perpendicular to the tracts
- ▶ *Degree of restriction* depends on tissue properties



# Diffusion MRI in a nutshell

■ MRI sequences are **sensitive to diffusion** by inserting **two additional *magnetic field gradient pulses***

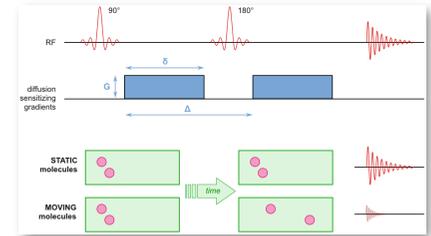
- ▶ The goal is to **change phase** of moving molecules
- ▶ Movement  $\Rightarrow$  phase differences  $\Rightarrow$  **signal cancels out/drops**



# Diffusion MRI in a nutshell

## ■ MRI sequences are **sensitive to diffusion** by inserting two additional *magnetic field gradient pulses*

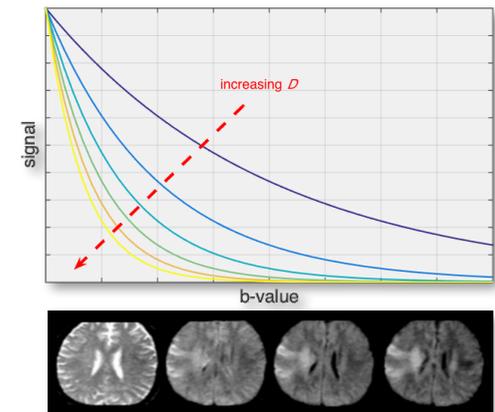
- ▶ The goal is to **change phase** of moving molecules
- ▶ Movement  $\Rightarrow$  phase differences  $\Rightarrow$  **signal cancels out/drops**



## ■ Signal decays as:

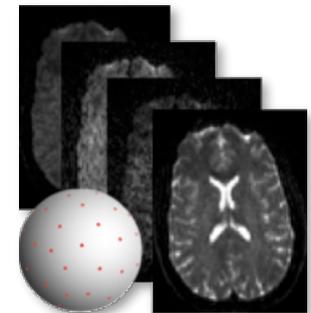
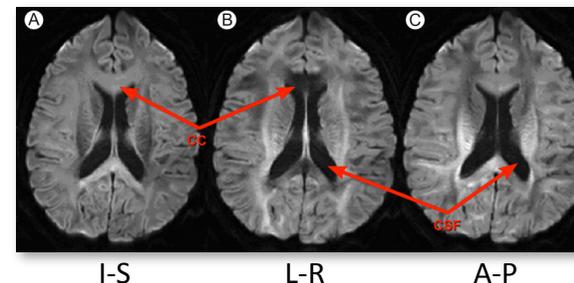
$$\text{signal} \propto e^{-bD}$$

- ▶  $D$  : *diffusion coefficient* of the tissue
- ▶  $b$  : *diffusion weighting/contrast* of the images ( $\delta$ ,  $\Delta$ ,  $|\mathbf{G}|$ )



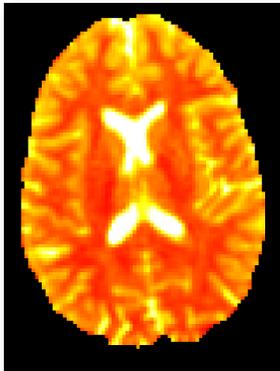
## ■ **NOTE:** signal strongly depends on

- ▶ the *b-value* ( $b$ )
- ▶ the *diffusion coefficient* ( $D$ )
- ▶ the *gradient direction* ( $\mathbf{G}$ )

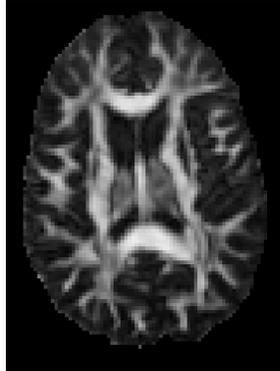


## ■ Scalar maps

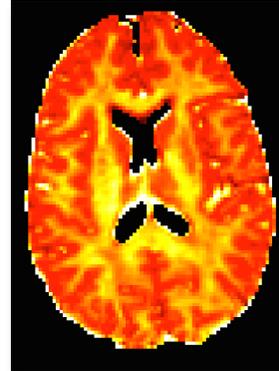
- ▶ Estimate *local features* of the tissue



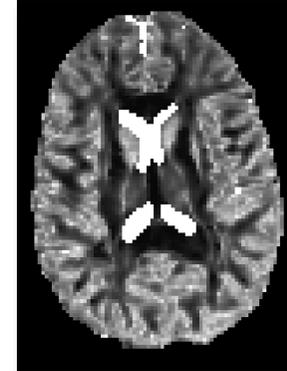
diffusion coefficient



diffusion anisotropy



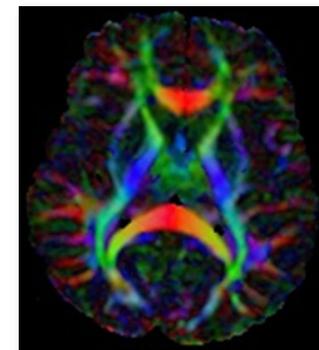
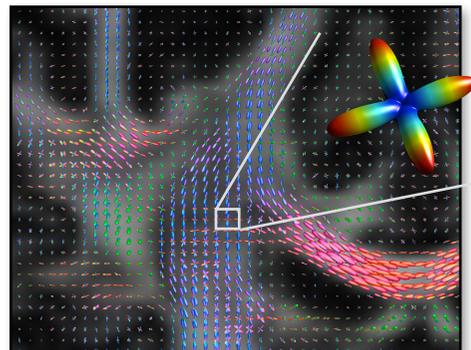
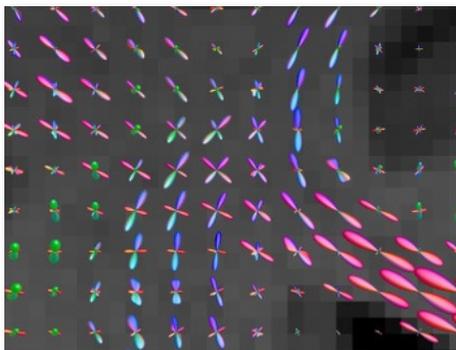
axonal density



axonal dispersion

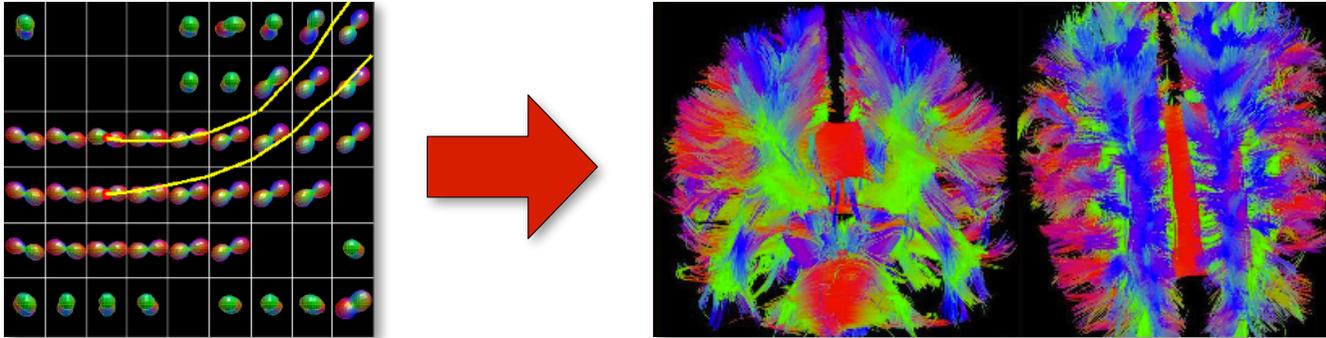
## ■ Intra-voxel fiber structure

- ▶ Estimate the *number and orientation* of fiber populations in each voxel



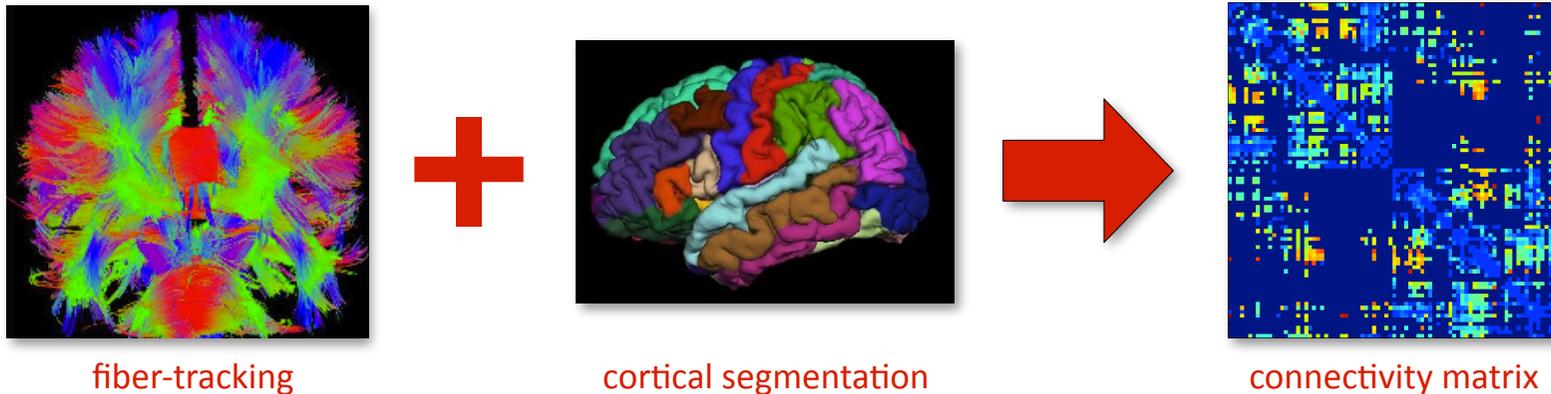
## ■ Fiber-tracking

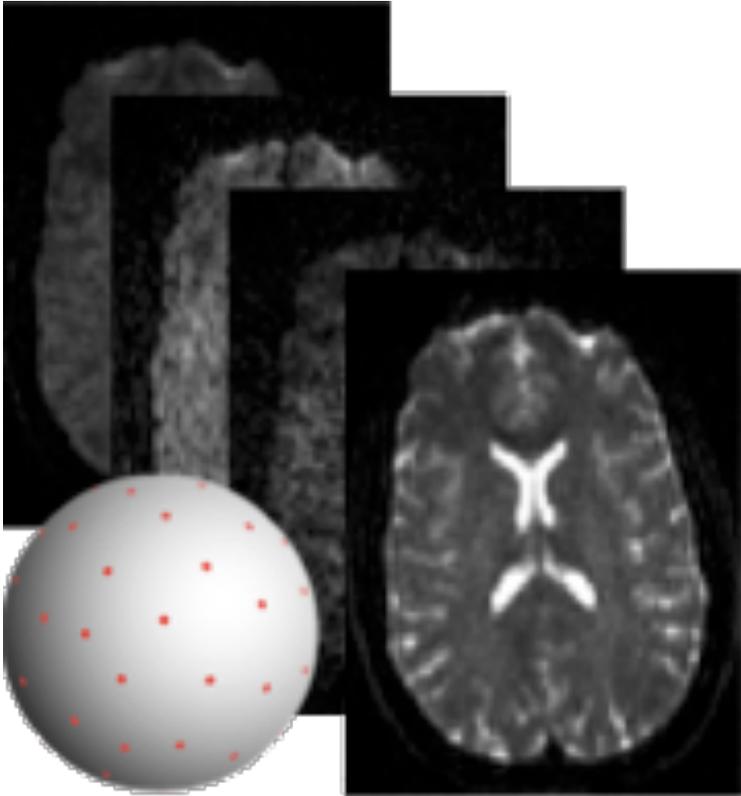
- ▶ Infer *axonal trajectories* by exploiting the diffusion information in each voxel

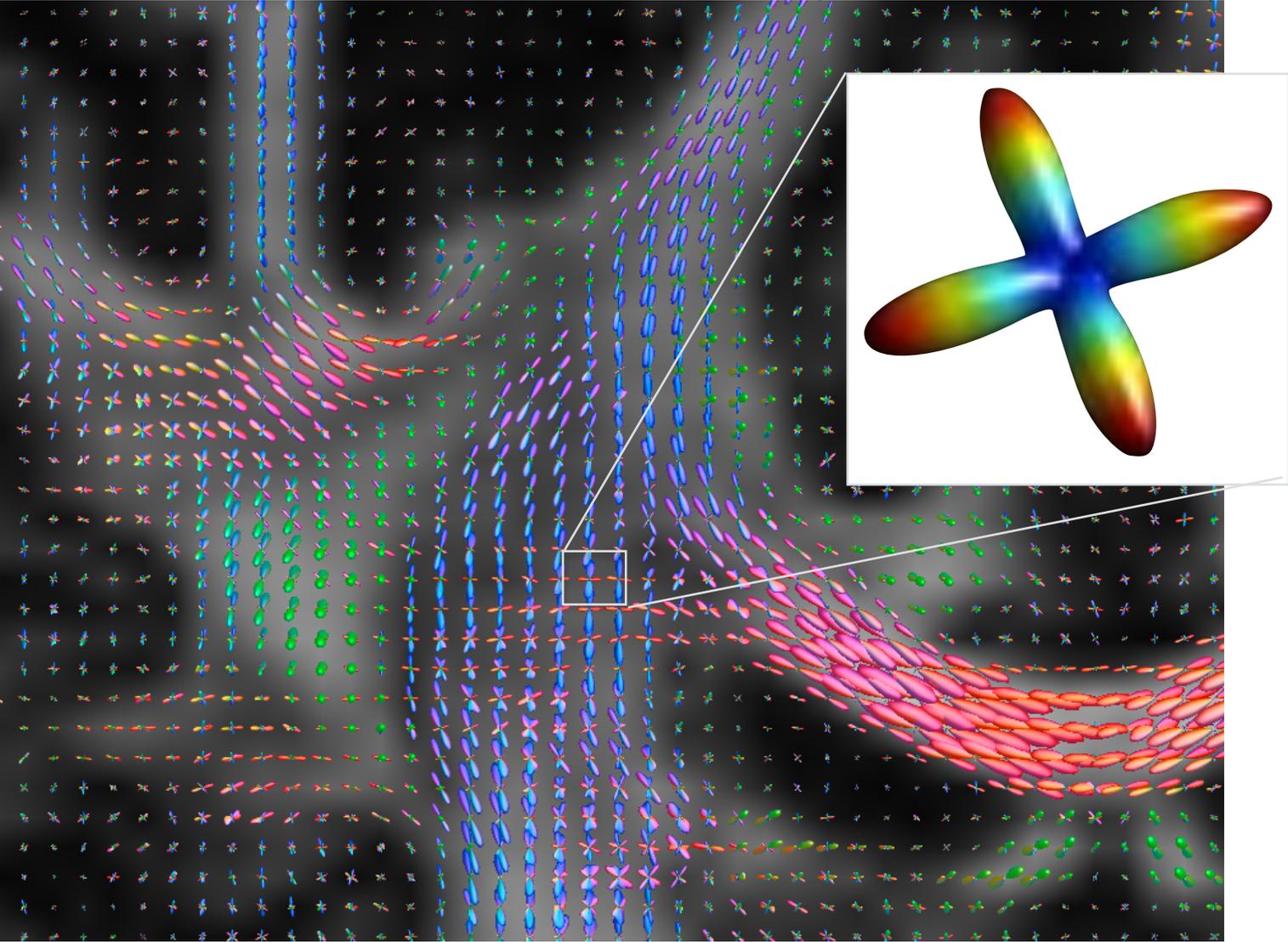


## ■ Connectivity analysis

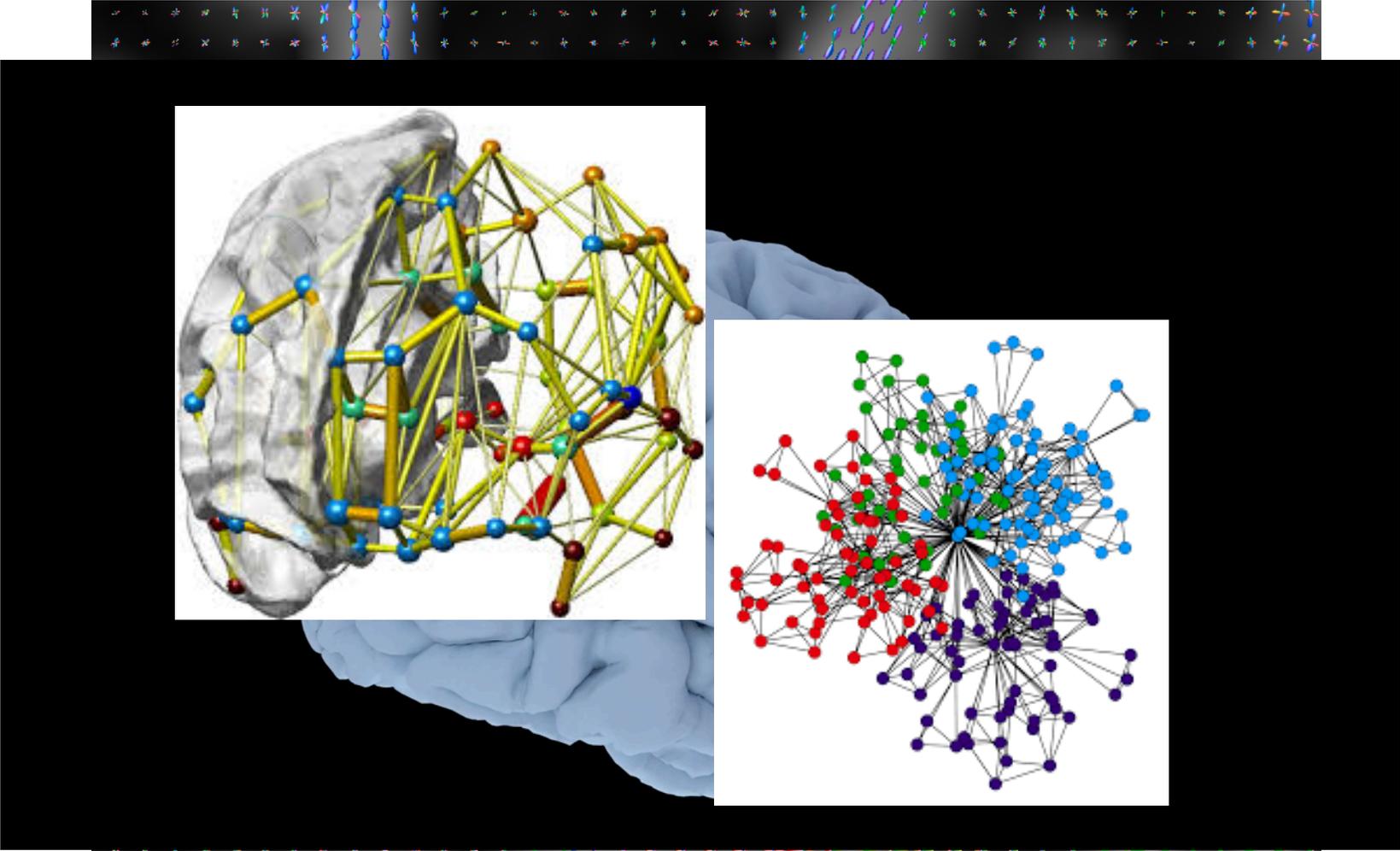
- ▶ *In-vivo* and *non-invasive* assessment of structural wiring of the brain











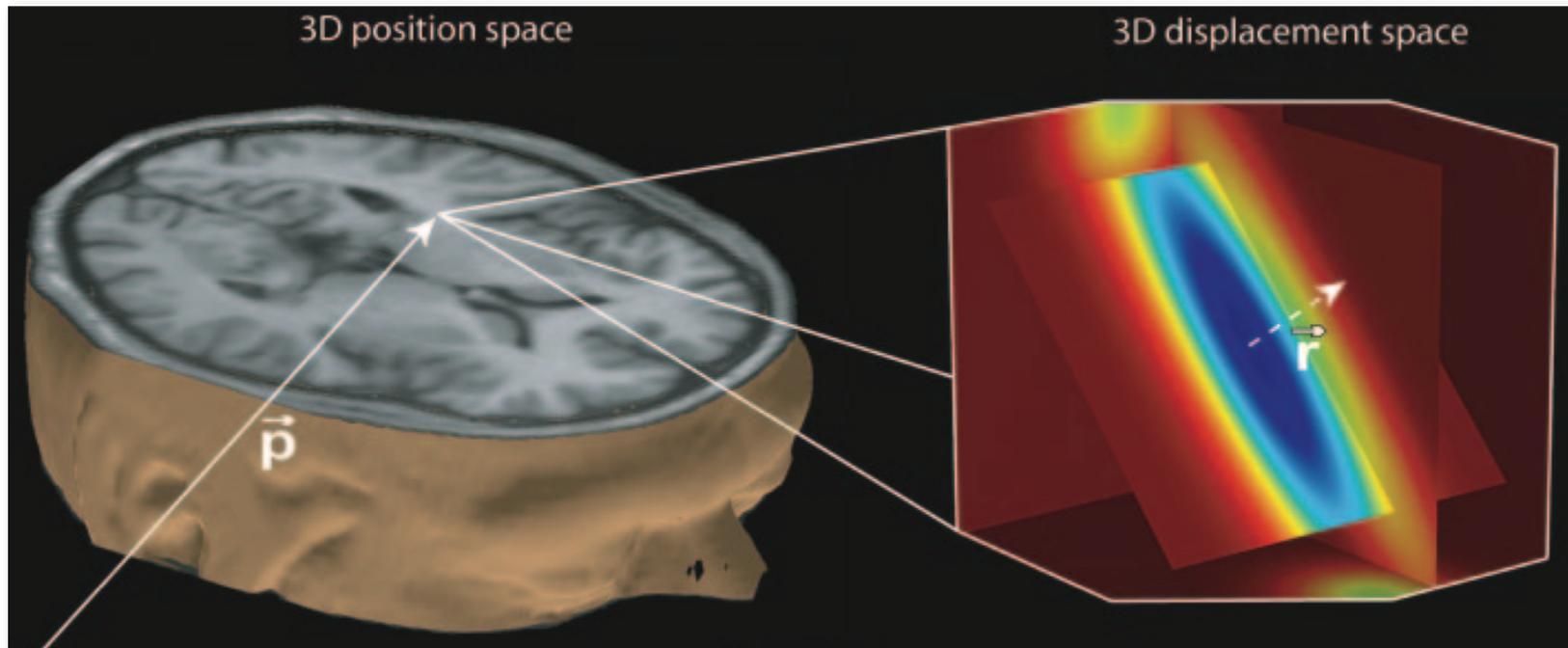
# EAP, ODF and fODF

## ■ EAP (*Ensemble Average Propagator*)

- ▶ In each voxel, 3D PDF giving the *probability of water displacements* → diffusion MRI is a **6D modality**
- ▶ Related to the *signal attenuation* by a **3D FFT**:

$$P(\vec{r}) = \int_{\mathbb{R}^3} E(\vec{q}) e^{-2\pi i \vec{q} \cdot \vec{r}} d\vec{q}$$

← **q-space**



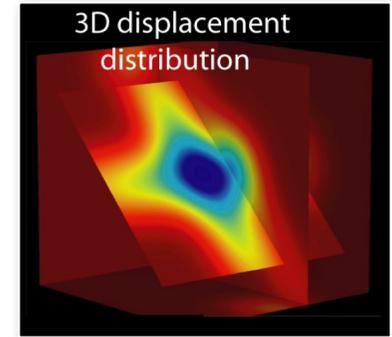
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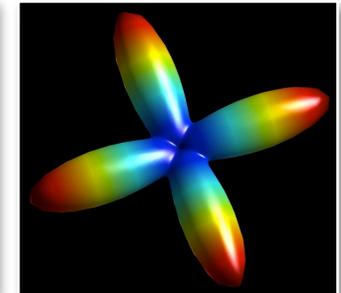
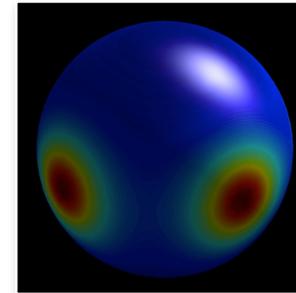


## ■ ODF or dODF (*Orientation Distribution Function*)

- ▶ Probability of diffusion *along a given direction*:

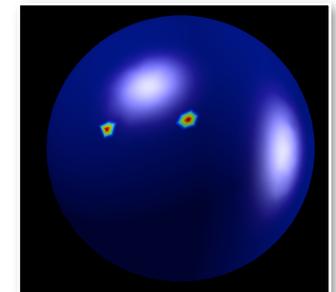
$$\text{ODF}(\hat{r}) = \int_{\mathbb{R}_+} P(r, \hat{r}) r^2 dr$$

- ▶ Function on the sphere



## ■ FOD or fODF (*fiber ODF*)

- ▶ Probability of having a fiber population *along a given direction*
- ▶ Function on the sphere



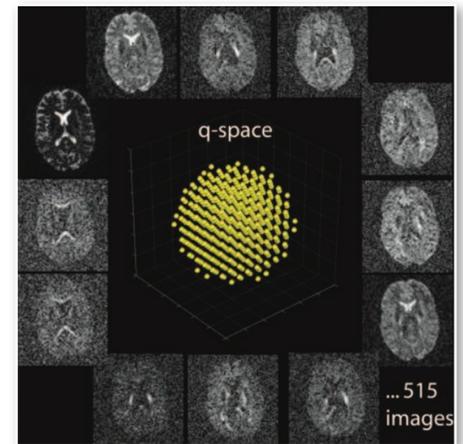
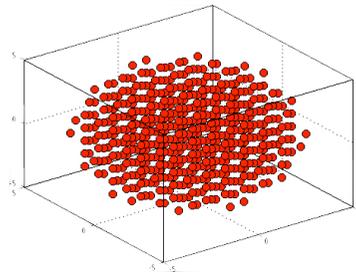
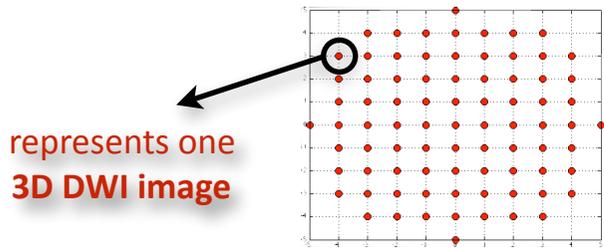
- Exploits the **3D Fourier Transform relationship** between the *MR signal* and the *displacement distribution (EAP)*

$$P(\vec{r}) = \int_{\mathbb{R}^3} E(\vec{q}) e^{-2\pi i \vec{q} \cdot \vec{r}} d\vec{q}$$



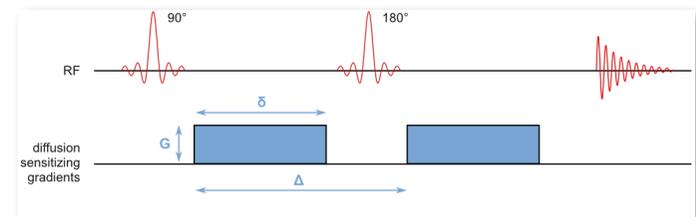
- The  $q$ -space must be **properly sampled**

- Data must be sampled in a *dense 3D cartesian grid*
- Usual protocol: 515 samples with  $b_{\max} \approx 8000 \text{ s/mm}^2$



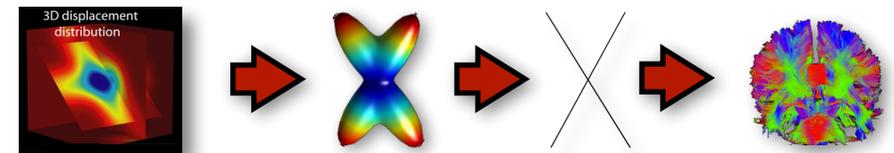
- Measures *directly* water displacements making **“almost” no assumptions**

- “almost”  $\Rightarrow$  *short pulse condition* is required



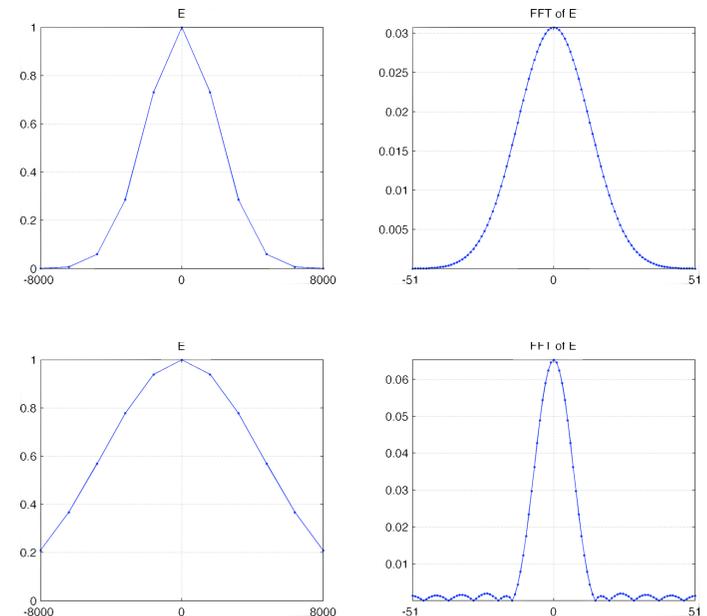
## Advantages:

- ▶ Model free
- ▶ Complex fiber configurations recovered
- ▶ Recovers the EAP  
(even though radial information is usually ignored!!!)



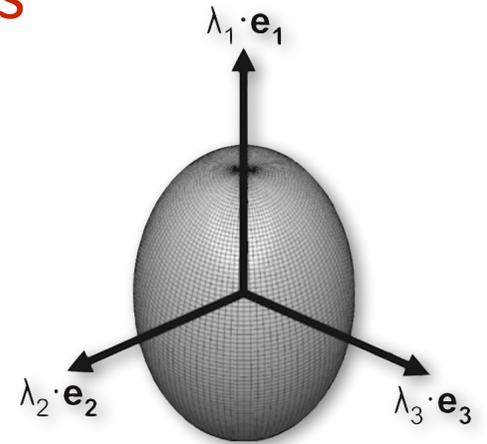
## Limitations:

- ▶ Long acquisitions ( $\approx 30-40$  min)
- ▶ Motion sensitive
- ▶ High b-values + long TE  $\Rightarrow$  low SNR
- ▶ Short pulses never met  $\Rightarrow$  smooth EAP
- ▶ No useful maps from the EAP
- ▶ Severe truncation artifacts
  - Inherent to FFT and “relatively low” b-values
  - Hanning filter mitigates but introduces blurring

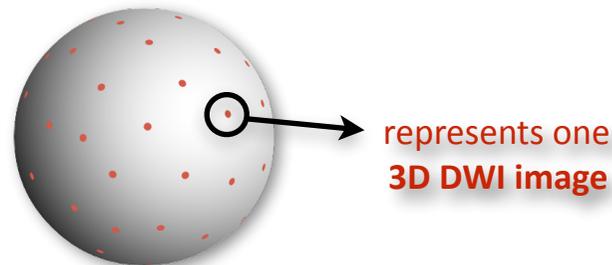


## ■ Assumption: displacements of water molecules follow a *multivariate gaussian distribution*

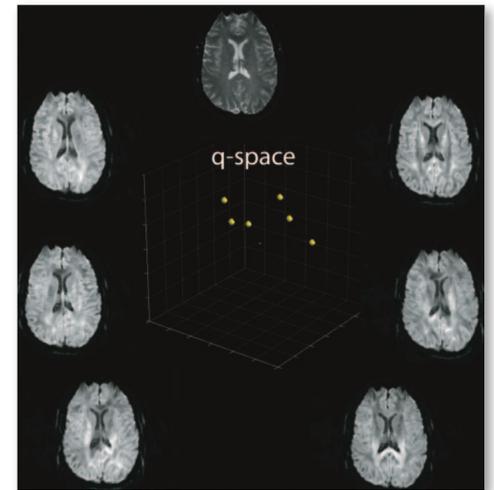
- ▶ Process fully characterized by its **covariance matrix** (3x3 symmetric positive semi-definite matrix)
- ▶ Usually represented as an ellipsoid
- ▶ 6 degrees of freedom (3 rotations + 3 variances)



## ■ This matrix, i.e. **diffusion tensor**, can be estimated acquiring *6+1 DWI images* (at least)



- ▶ Usual protocol: 6-32 directions with  $b \approx 1000 \text{ s/mm}^2$



## Advantages:

- ▶ *Fast* acquisitions ( $\approx$  4-5 min)  $\Rightarrow$  clinically feasible
- ▶ Does not require special hardware
- ▶ Useful *scalar maps*, e.g.

- Mean Diffusivity (**MD**) :

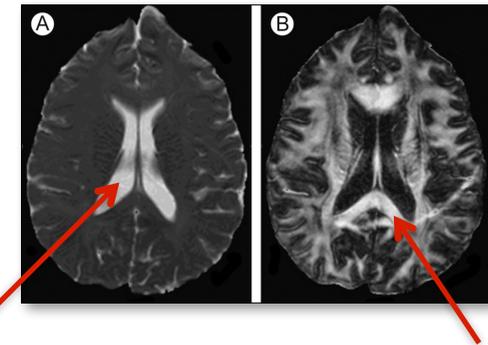
$$\bar{\lambda} = \frac{\lambda_1 + \lambda_2 + \lambda_3}{3}$$

- Fractional Anisotropy (**FA**) :

$$FA = \sqrt{\frac{3}{2}} \sqrt{\frac{(\lambda_1 - \bar{\lambda})^2 + (\lambda_2 - \bar{\lambda})^2 + (\lambda_3 - \bar{\lambda})^2}{\lambda_1^2 + \lambda_2^2 + \lambda_3^2}}$$

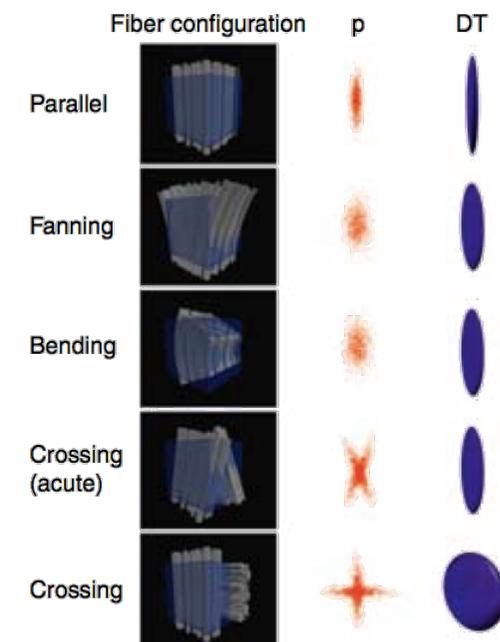
## Limitation(s):

- ▶ Complex fiber configurations *cannot be modeled*
  - majority of voxels in the brain
  - Enough for characterization of major bundles, but inadequate for whole brain connectivity analyses
  - NB: acquiring more data does overcome this limitation!



**Mean Diffusivity:**  
high values = fast diffusion

**Fractional Anisotropy:**  
high values = fiber bundles

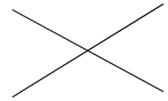


## High angular resolution diffusion imaging (HARDI):

- ▶ Focus on *angular information*:
  - The radial component is discarded/averaged
  - No access to tissue micro-structural features, e.g. axonal diameter and density
- ▶ Usually based on *spherical sampling* (at least 60 samples in q-space)



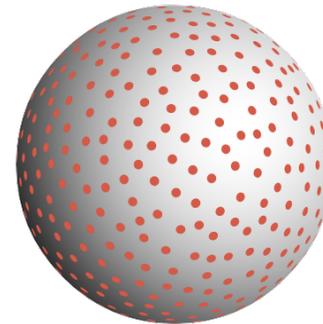
dODF



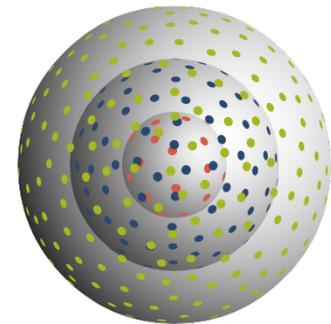
fODF

## Vast literature of methods:

- ▶ Multi-Tensor fitting (Tuch et al, 2002)
- ▶ Q-BALL (QBI) (Tuch, 2004)
- ▶ Q-BALL in Constant Solid Angle (QBI<sub>CSA</sub>) (Aganj et al, 2010)
- ▶ Constrained Spherical Deconvolution (CSD) (Tournier et al, 2004)
- ▶ Diffusion Orientation Transform (DOT) (Ozarslan et al, 2006)
- ▶ ...

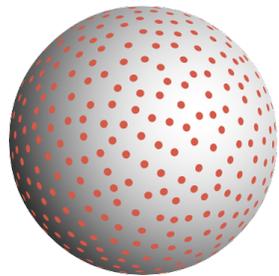


1-shell



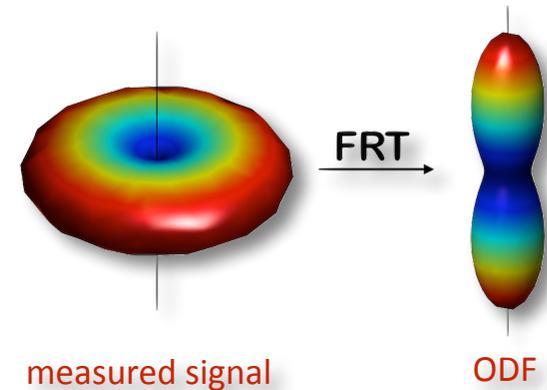
multi-shell

- **IDEA:** data is acquired on a *single shell* and the ODF is approximated by means of *Funk-Radon Transform (FRT)*



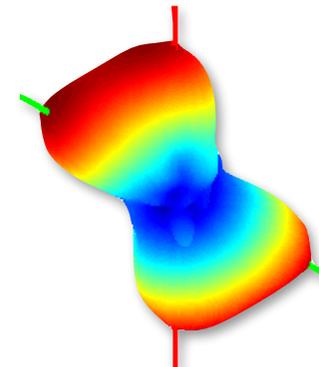
Samples distributed on a single-shell

Same idea behind CT, but on the sphere

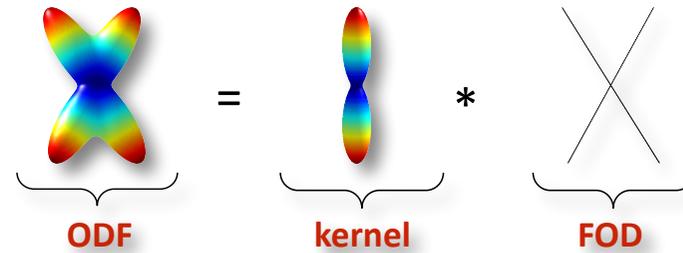


## Notes:

- ▶ Model free
  - ▶ FRT induces blurring
  - ▶ ODF is only approximated (missing  $r^2$  term in the integral)
  - ▶ Usual protocol:  $\geq 60$  directions with  $b \approx 3000 \text{ s/mm}^2$
- } ODF are smooth



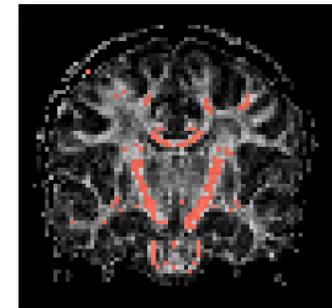
- Assumption: ODF can be seen as a convolution on the sphere



ODF = kernel \* FOD

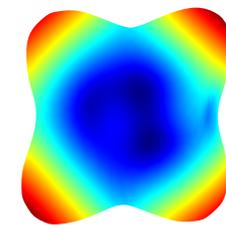
- The **kernel** characterizes the diffusion response-function of a single fiber; can be estimated from the data:

- ▶ Identify known areas with only one fiber population
- ▶ Fit a tensor in each voxel and average



## Notes:

- ▶ Model based
- ▶ High angular accuracy (i.e. sharper profiles), but sensitive to noise
- ▶ Assumes the *same diffusion properties* across the whole brain
- ▶ Usual protocol:  $\leq 60$  directions with  $b \approx 3000 \text{ s/mm}^2$



Q-BALL



CSD