## BSD

# Diffusion Tractography

- Goal of tractography
- Estimating Fibre Orientations BEDPOSTX
- Probabilistic Tractography PROBTRACKX
- ProbtrackX outputs
- Tractography limitations





### DTI tractography

**v**<sub>1</sub> map Principal Diffusion Direction



Principal Diffusion Direction





#### Assumption:

Direction of maximum diffusivity (in anisotropic voxels) is an <u>estimate</u> of the major fibre orientation.



### But is WM always coherently organised within a voxel?



Unfortunately not, complex fibre patterns (e.g. crossings) are very common at the voxel scale.

Williams, Gluhbegovic, and Jew, "The Human Brain: Dissections of the Real Brain", Virtual Hospital, University of Iowa, 1997



#### Predictions from the tensor model no crossing fibres









### How good is the DTI Model in regions with crossing fibres?

- In voxels containing two crossing bundles, the tensor ellipsoid is pancake-shaped (oblate, planar tensor).
- In voxels containing three crossing bundles, the tensor ellipsoid is spherical.
- In these areas, DTI  $\mathbf{v}_1$  is meaningless.





#### **Uncertainty on DTI Fibre Orientation Estimates**

Repeat an acquisition many times and obtain the variability in  $v_1$  from the different datasets.



Cones of uncertainty on DTI  $v_1$ 

Jones, 2002



Do we have to use the DTI model to estimate orientations? Not really, many models exist



#### Ball & Sticks Model Unlike the DT model, it can represent many orientations



- Anisotropic tensors (sticks) with isotropic background (ball)
- Fibre Orientations modelled explicitly and separated from isotropic partial volumes





### Predictions from the ball and sticks model crossing fibres



#### Markov Chain - Monte Carlo (MCMC) Sampling





#### Output in Each voxel = Distributions of Parameters







#### Ball & Sticks Model Selection

- Model selection problem: One, two or more fibres within a voxel?
- Automatic Relevance Determination: Only estimate complexity that is supported by the data





#### Modelling Complex Fibre Architectures Automatic Relevance Determination (A.R.D.)



ARD1

**Measured Signal** 



Modelling Complex Fibre Architectures Automatic Relevance Determination (A.R.D.)

- After running BedpostX all voxels will have estimated parameters for the maximum number of sticks requested.
- But due to ARD, the sticks that are not supported in a voxel will have an almost zero volume fraction.
- We use a threshold (e.g. >5%) to **exclude sticks with tiny volume fraction**.





#### **Ball & Sticks Orientations**

All sticks, with secondary ones thresholded (*f<sub>n</sub>*>5%)





#### **DTI vs Ball & Sticks Orientations**

DTI









A large portion of the WM supports crossing fibres

Coherence in orientations shows that we are not over-fitting (the ARD works)



#### Multi-Shell Diffusion Acquisitions Why bother?



Higher b value gives us more angular contrast!!!



#### Multi-Shell Diffusion Acquisitions Why bother?







#### Generalised Ball & sticks Model Gets best of both worlds

- Multi-shell model (or model=2) in Bedpostx options.
- Allows representation of multiple diffusivities within a voxel (rather than just one).
- More accurate model for multi-shell data & partial volume effects.



#### Human Connectome Project Data

\*Jbabdi, Sotiropoulos et al, MRM 2012 \* Sotiropoulos, Jbabdi et al, NeuroImage 2013

### Faster bedpostx on GPUs



Hernandez et al, Plos One 2013