## Diffusion MRI simulation of realistic neurons with SpinDoctor and the Neuron Module

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## **Table of contents**

#### Introduction to Diffusion MRI

- Three main ingredients of dMRI
- Principles of Diffusion MRI
- Two classes of imaging methods

#### 2 SpinDoctor & Neuron Module

#### 3 Numerical examples

- Validation of hypotheses
- Potential biomarkers for brain microstructure

### 4 Conclusion

## **Introduction to Diffusion MRI**

#### 1. Molecular Diffusion (Wikipedia)

Diffusion is the thermal motion of all (liquid or gas) particles at temperatures above absolute zero.



Diffusion from high concentration regions to low concentration regions



Brownian motion [Johansen 2013]

#### 2. Larmor Precession (Wikipedia)

Larmor precession is the precession of the magnetic moment of an object about an external magnetic field.



#### 3. Nuclear Magnetic Resonance (NMR) techniques

align the magnetic moment of particles



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- synchronize the Larmor precession of particles
- focus on one dMRI voxel
- measure the dMRI signal, etc.



### Diffusion MRI

Diffusion MRI is an imaging modality that can be used to probe the tissue microstructure by encoding the diffusion motion of water molecules.





Position:  $\vec{x}$ Magnetization: exp(- $i\gamma\delta \vec{g}\cdot \vec{x}$ )

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 $\vec{g}$ 



$$S(\overrightarrow{g}, \Delta) = \sum_{\overrightarrow{r}} n_r \cdot exp(i\gamma\delta \overrightarrow{g} \cdot \overrightarrow{r})$$



$$\mathcal{S}(\overrightarrow{g},\Delta) = \sum_{\overrightarrow{r}} N_{total} \mathcal{P}(\overrightarrow{r},\Delta) exp(i\gamma \delta \overrightarrow{g} \cdot \overrightarrow{r})$$



$$S(\overrightarrow{g}, \Delta) = \sum_{\overrightarrow{r}} N_{total} P(\overrightarrow{r}, \Delta) exp(i\gamma \delta \overrightarrow{g} \cdot \overrightarrow{r})$$
  
 $\overrightarrow{q} = \gamma \delta \overrightarrow{g} / 2\pi$ 



$$\boldsymbol{S}(\overrightarrow{q},\Delta) = N_{total} \sum_{\overrightarrow{r}} \boldsymbol{P}(\overrightarrow{r},\Delta) exp(2\pi i \overrightarrow{q} \cdot \overrightarrow{r})$$

Fourier relationship between the measured signal  $S(\vec{q}, \Delta)$  and the diffusion propagator  $P(\vec{r}, \Delta)$ .

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[Van 2010]



[Van 2010]

## **DSI Tractography**



[Van 2010]

## **Model-Independent Diffusion MRI**

- Diffusion Spectrum Imaging (DSI)
- Diffusion-Weighted Imaging (DWI)  $S(\overrightarrow{q}, \tau) = S_0 \cdot e^{-D_{eff} |\overrightarrow{q}|^2 \tau}$
- Diffusion Tensor Imaging (DTI)  $S(\overrightarrow{q}, \tau) = S_0 \cdot e^{-\overrightarrow{q} \cdot \tau} \mathbf{D}_{eff} \overrightarrow{q} \tau$
- Diffusion Kurtosis Imaging (DKI)

etc.



An example of DWI image [Radiopaedia.org, rID: 33859]

## Model-Independent Diffusion MRI

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etc.

#### Motivation 1

The understanding of the physical process affects the way of the image reconstruction.

We can take an alternative perspective — Bloch-Torrey equation:

$$\frac{\partial}{\partial t}M(\boldsymbol{x},t) = -i\gamma \boldsymbol{g} \cdot \boldsymbol{x} M(\boldsymbol{x},t) + \nabla \cdot (\mathcal{D}\nabla M(\boldsymbol{x},t))$$

The models of the brain microstructure



#### The models of the brain microstructure



The models of the brain microstructure



- Neurite Orientation Dispersion and Density Imaging (NODDI)
- White Matter Tract Integrity (WMTI)
- Soma And Neurite Density Imaging (SANDI)
- DIstribution of Anisotropic MicrOstructural eNvironments in Diffusion-compartment imaging (DIAMOND)
- Linearly Estimated Moments provide Orientations of Neurites And their Diffusivities Exactly (LEMONADE)
- etc.

### Pros

- Neurite Density
- Dendrites orientation distribution
- Soma size and density

#### Cons

- Sensitive to the noise
- Failed in the grey matter
- Lack of validation

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- etc.

#### Motivation 2

The model of the brain microstructure also matters. Realistic neuron models should be built and applied for the future research.

## SpinDoctor & Neuron Module

SpinDoctor is a Matlab-based dMRI framework which employs the finite element method (FEM) to solve the Bloch-Torrey equation. [Li 2019] Available on https://github.com/jingrebeccali/SpinDoctor

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- Generate high quality FE meshes
- Analyse the compartments individually



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## SpinDoctor

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Features:

- Generate high quality FE meshes
- Analyse the compartments individually
- Support permeable interface
- Support many types of diffusion-encoding pulse sequences
- 100x faster than the classical Monte-Carlo simulator Camino



Camino logo

Neuron Module is a new module of SpinDoctor which contains 65 embedded realistic neuron meshes.

The embedded neurons are built from the morphological descriptions published in NeuroMorpho.

NeuroMorpho's morphological descriptions obtained from microscopic measurement.



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Features:

 30x faster than the GPU-based Monte-Carlo simulation

Computational time (s)		neuron	
$\delta = 10ms, \Delta = 43ms$	b	$E_{max}$	t(s)
SpinDoctor	1000	0.16	17.1
	4000	0.34	26.0
GPU Monte-Carlo	1000	0.14	537.8
	4000	0.60	1895.9
GPU MC/	1000		31
SpinDoctor ratio	4000		72

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- Separated neuron compartments



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# What difference can we make?

## Validation of hypotheses



#### The power law hypothesis

For tubular structures such as cylinders, the diffusion direction averaged signal,  $S_{ave}(b)$ , is linear in  $\frac{1}{\sqrt{b}}$ :

$$\mathcal{S}_{ave}(b) \equiv \int_{\|oldsymbol{u}_{oldsymbol{g}}\|=1} \mathcal{S}_{oldsymbol{u}_{oldsymbol{g}}}(b) doldsymbol{u}_{oldsymbol{g}} \sim c_0 + c_1 rac{1}{\sqrt{b}}$$

with  $b = \gamma^2 \|\boldsymbol{g}\|^2 \delta^2 (\Delta - \delta/3)$  for PGSE sequences.

## Validation of the power law

#### The indispensable Neuron Module features

- Realistic neuron meshes
- Separated neuron compartments
- High angular resolution
- High efficiency



## Validation of the power law



- The power law is valid for dendrite branches.
- Due to the presence of the soma, the power law is no longer valid for the whole neuron.

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## Potential biomarkers

By leveraging the collection of the realistic neuron meshes, we statistically show the deviation from the power law has the potential to serve as biomarkers for revealing the soma size.



## Potential biomarkers

We performed a statistical study of the above 6 candidate biomarkers on the collection of the 65 neurons.



•  $x_0$ ,  $c_0$ ,  $c_1$ ,  $\mathcal{E}$  and w exhibit exponential relationship with the soma volume  $v_{soma}$ .

## **Potential biomarkers**

We performed a statistical study of the above 6 candidate biomarkers on the collection of the 65 neurons.



- $x_0, c_0, c_1, \mathcal{E}$  and w exhibit exponential relationship with the soma volume  $v_{soma}$ .
- $y_0$ ,  $c_0$  and  $\mathcal{E}$  seem capable of indicating the lower bound for the soma volume fraction  $f_{soma}$ .

## Conclusion

## Conclusion

- SpinDoctor provides an accurate and fast dMRI simulator for solving the Bloch-Torrey equation.
- Neuron Module which is built on SpinDoctor's FEM solver, allows us to solve the Bloch-Torrey equation on 65 embedded realistic neurons and their compartments.
- The combination of the advanced dMRI simulator and the realistic neuron models enables fast prototype validation and the design for new imaging methods.

Github repo: https://github.com/jingrebeccali/SpinDoctor Paper: https://doi.org/10.1016/j.neuroimage.2020.117198

## Thanks for your attention!

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