



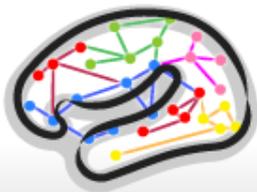
# Structural Brain Connectivity Analysis on HCP Dataset

**Zhengwu Zhang**

**Feb. 2 , 2016**

Joint work with Hongtu Zhu, Anuj Srivastava and Maxime Descoteaux

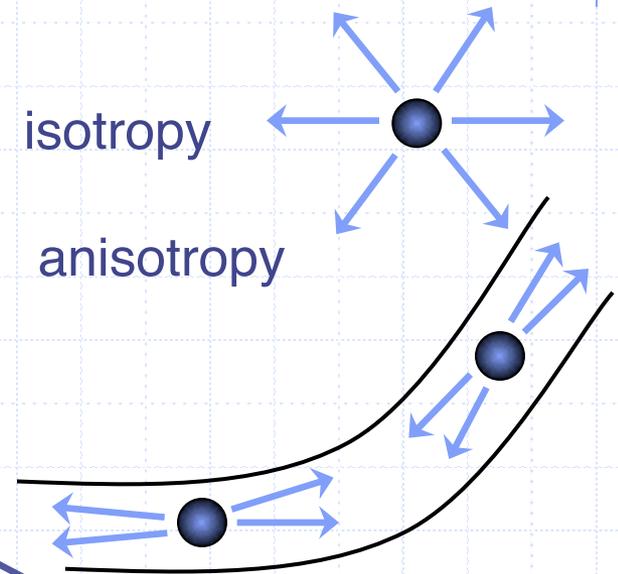
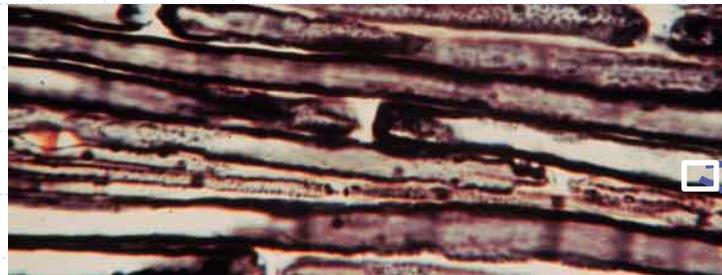




## Diffusion in Brain Tissue

➤ Water molecules in different brain tissues have different diffusion properties.

- **Gray matter:** Diffusion is unrestricted  $\longleftrightarrow$  isotropy
- **White matter:** Diffusion is restricted  $\longleftrightarrow$  anisotropy



- More diffusion along axon fibers

➤ Diffusion MRI measures the water diffusion movement inside brain



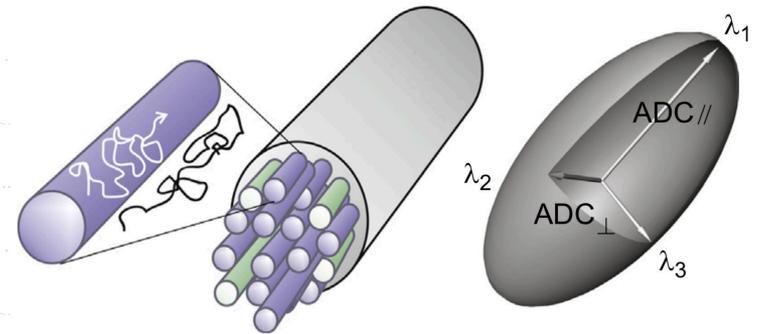
## Representation of the Diffusion Directions

- At each voxel, we want to know:
  - What is the orientation of the diffusion?
  - What is the magnitude of diffusion?

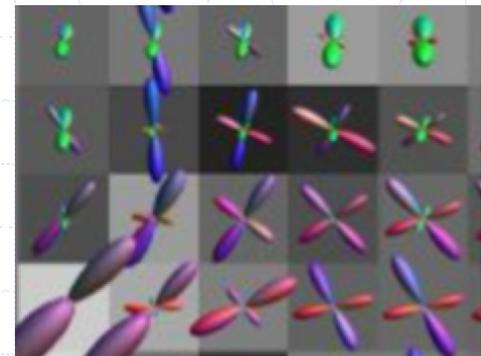
- Two popular representations:

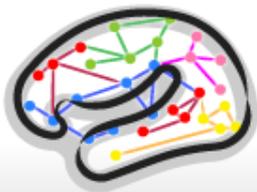
- Diffusion tensor image (DTI)

$$D = \begin{pmatrix} d_{1,1} & d_{2,1} & d_{3,1} \\ d_{2,1} & d_{2,2} & d_{3,2} \\ d_{3,1} & d_{3,2} & d_{3,3} \end{pmatrix}$$



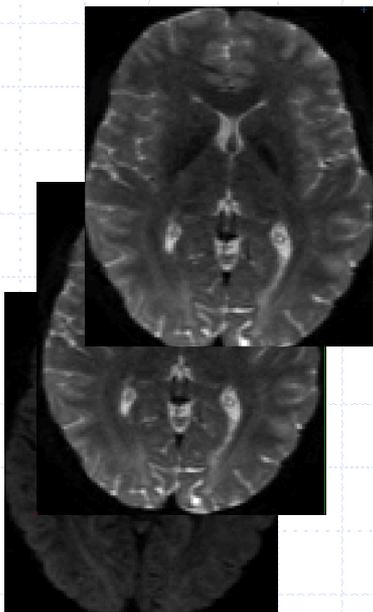
- High angular resolution diffusion imaging (HARDI)
  - Orientation distribution function [Tuch' 04]
  - Diffusion spectrum [Wedeen' 05]
  - Ball-and-stick [Behren's 03]
  - ...



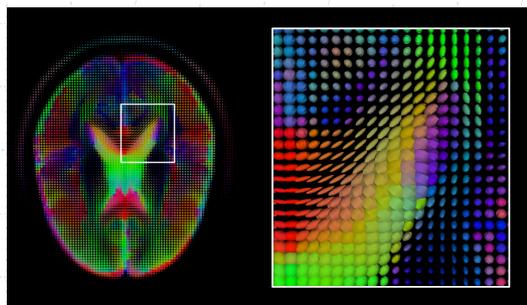


## Diffusion MRI to Connectome

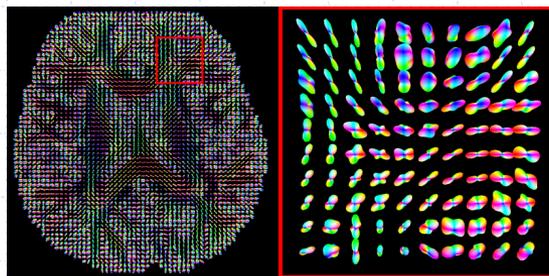
➤ From dMRI to structural connectomics



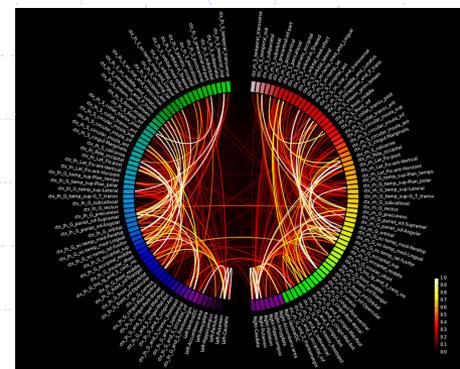
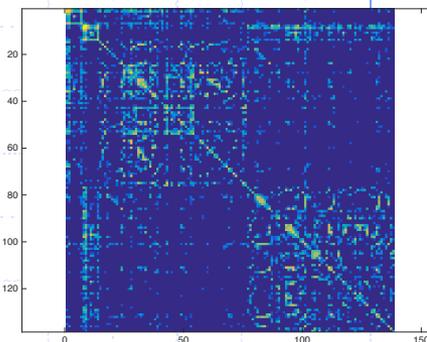
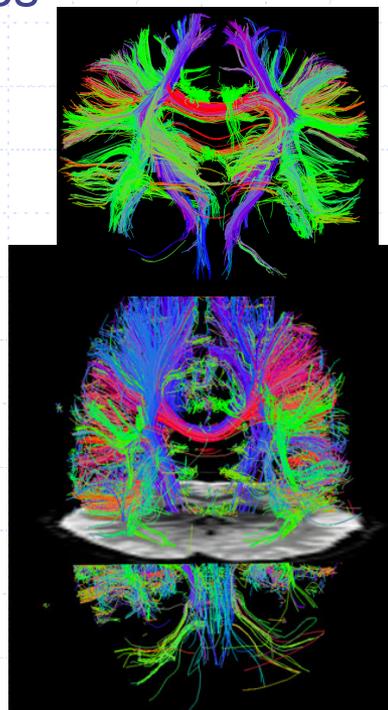
dMRI



diffusion directions

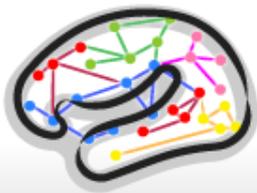


streamlines



connectivity  
matrix





## Introduction of Human Connectome Project

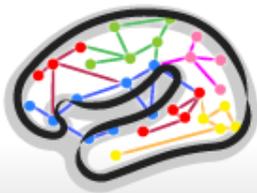
- May, 2009 : Request for Applications from NIH Blueprint
- Sept. 2010 : NIH awarded HCP grants to two consortia
  - Washington U, U of Minn, Oxford U,...
  - MGH and UCLA,...
- **Goal of HCP:** Characterize human brain connectivity and function
  - Scan ~1,200 healthy adults , ages 22-35, including
    - Twins and their non-twin siblings: MZ twins and DZ twins
- Dec. 2015: 900 subjects dataset release (**Latest release**)



## Data Acquisition

### (1) Imaging Data:

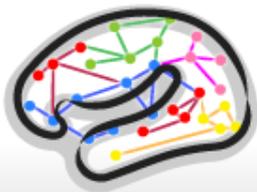
- 4 modalities, 1200 subjects, 1 customized 3T scanner (WashU)
  - Structural MRI (T1-weighted, T2-weighted)
  - Resting-state fMRI (rfMRI)
  - Task fMRI (tfMRI)
  - Diffusion MRI (dMRI)
- Improved scanners, pulse sequences: producing high quality data



## Data Acquisition

(1) Imaging Data:

(2) MEG/EEG Data:



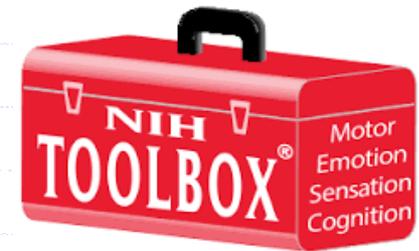
## Data Acquisition

(1) Imaging Data:

(2) MEG/EEG Data:

(3) Behavioral Data:

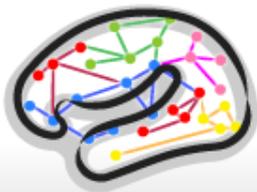
- Measures that have the potential to covary with brain connectivity and function:
  - NIH Toolbox ; Penn Neuropsychological Battery
- Diverse phenotypes  
Cognition; Emotional health; Motor skills; Sensory;  
Personality; Fluid intelligence; Family environmental factors
- Demographic, physical data
- Psychiatric status, substance use





## Data Acquisition

- (1) Imaging Data:
- (2) MEG/EEG Data:
- (3) Behavioral Data:
- (4) **Genetic Data:**

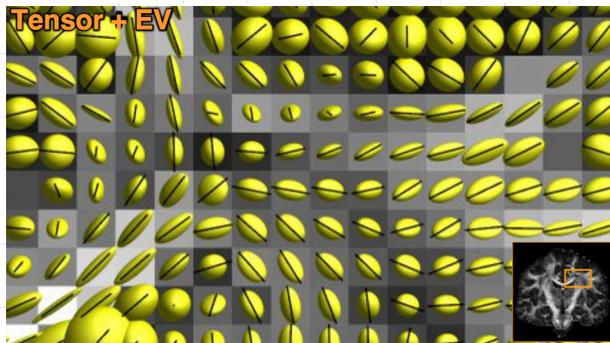


## Connectome Extraction

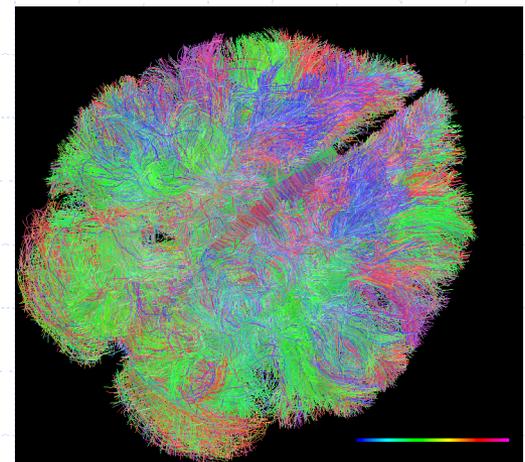
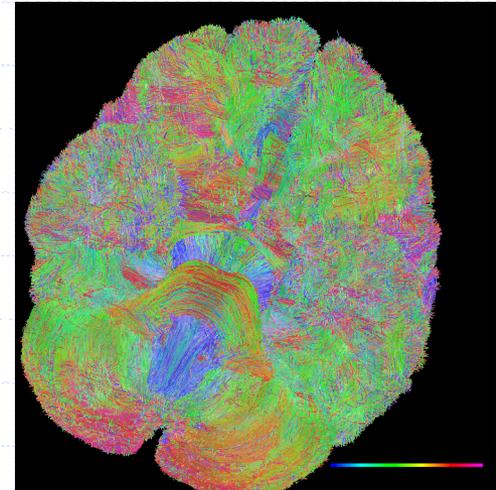
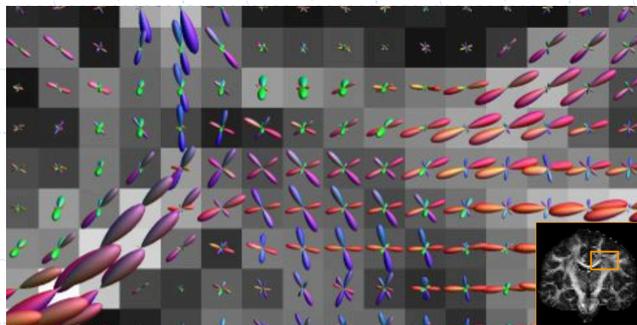
➤ We are interested in extracting the connectome from dMRI in HCP dataset.

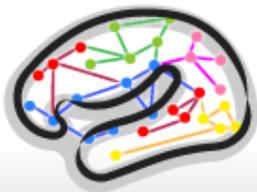
- **Step1. Construct HARDI:**  
better than DTI, can handle fiber crossing

DTI:



HARDI:

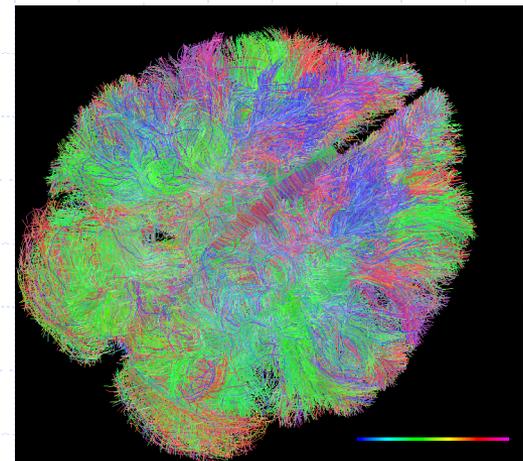
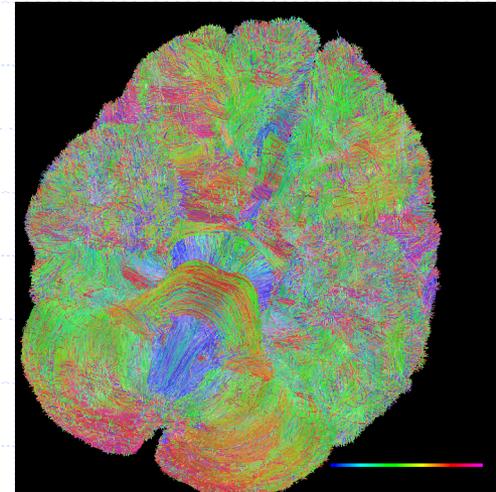


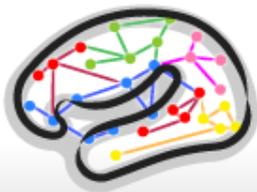


## Connectome Extraction

➤ We are interested in extracting the connectome from dMRI in HCP dataset.

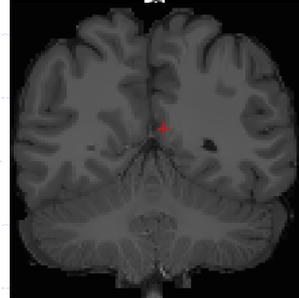
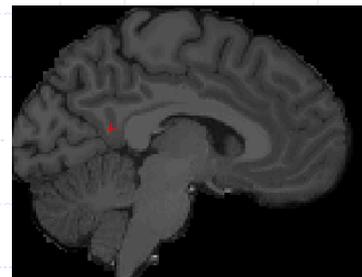
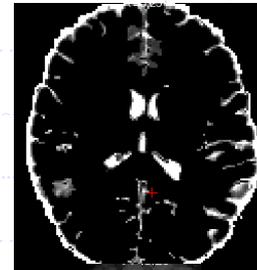
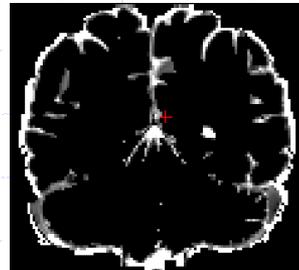
- **Step 1. Construct HARDI:**  
better than DTI, can handle fiber crossing
- **Step 2. Fiber tracking:**
  - Masking
  - Seeding
  - Streamline growing





## Masking and Seeding

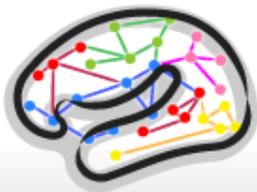
- Masking is used to
  - decide the propagation of a streamline
  - include the stopped streamlines



Included mask:

Excluded mask:

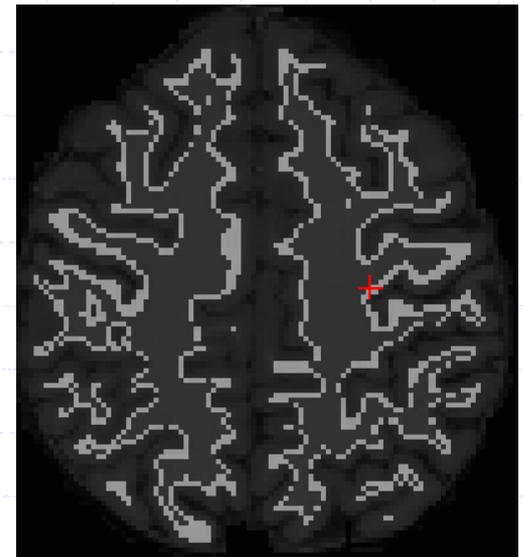
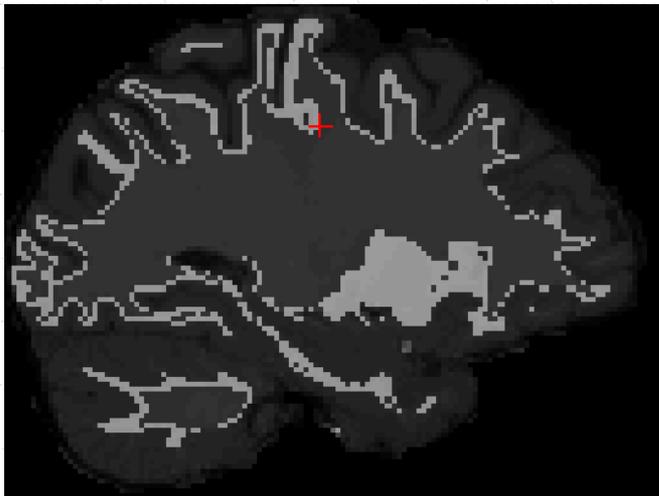
T1 brain:



## Masking and Seeding

- Seeding is used to
  - reduce the bias of streamlines

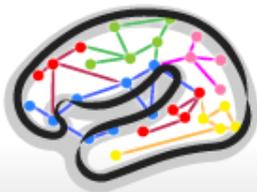
Seeding mask (the interface between GM and WM):





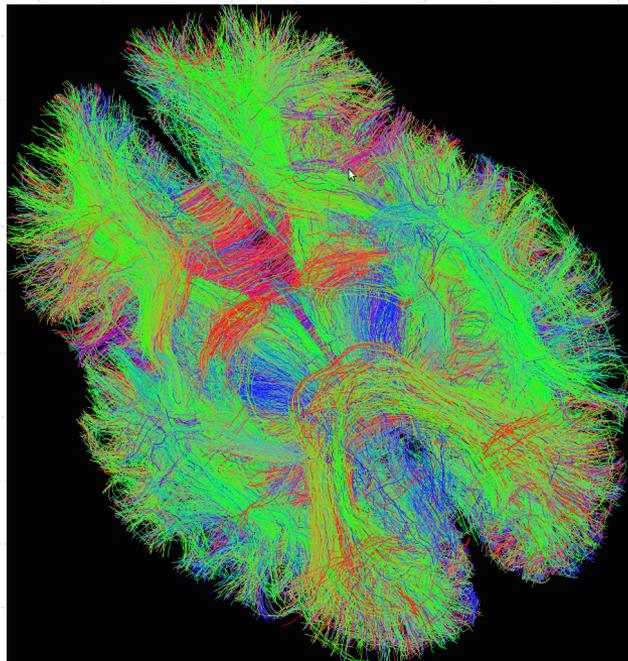
## Streamline Growing

- Streamlines growing
  - (1) probabilistic
  - (2) deterministic
  - (3) .....

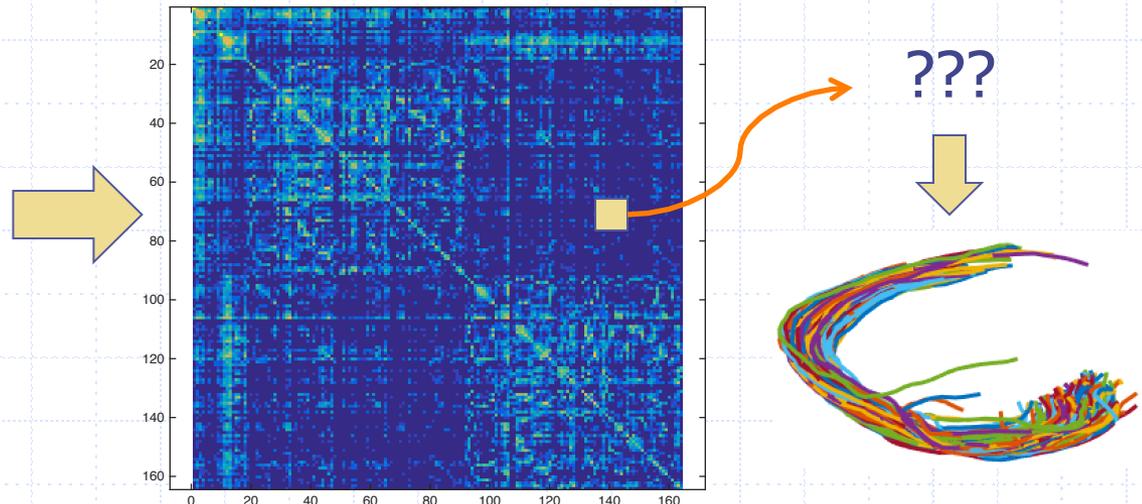


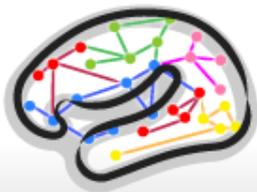
## Low Dimensional Representations

- Whole brain tractography is complicated
- A low dimensional representation is necessary for statistical inference



### Connectivity Matrix



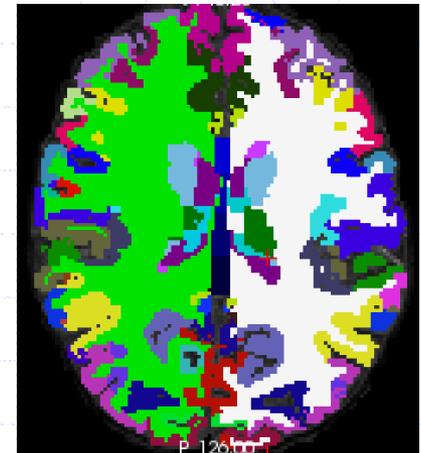
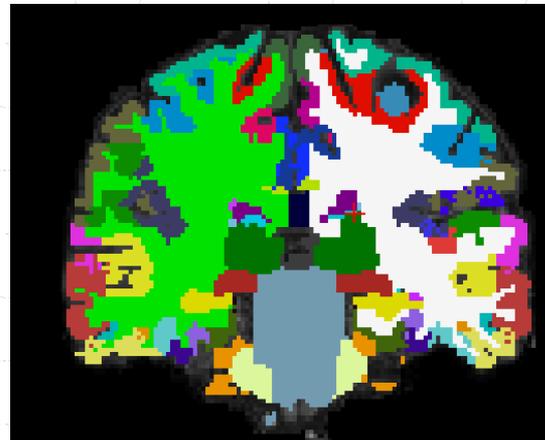


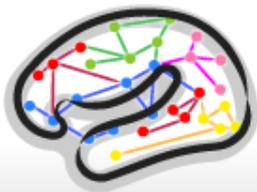
## Connectivity Matrix

- Connectivity matrix is a summarization of the brain connections
- Given streamlines:

Step 1. Brain parcellation – *freesurfer / other software*

- Use Destrieux atlas in FreeSurfer: ~ 170 ROIs





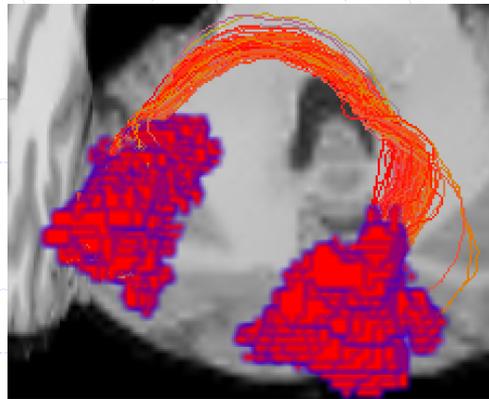
## Connectivity Matrix

- Connectivity matrix is a summarization of the brain connections
- Given streamlines:

Step 1. Brain parcellation – *freesurfer / other software*

Step 2. Find the fibers connecting each pair of regions

$CM(41,160) =$



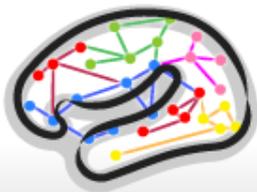
In order to include more streamlines:

- ROI dilation
- Streamline ending points expansion



## Problems with the Connectivity Matrix

- The connectivity matrix contains millions of fibers
  - saving, loading, and analyzing are difficult
    - Compression is needed
- Summarize the connectivity matrix
  - Extract robust measure(s) of connectivity strength



## Compression of Connectivity Matrix

- Examples of fibers in CM(1,160) for different subjects

*1 - left-lateral-ventricle*

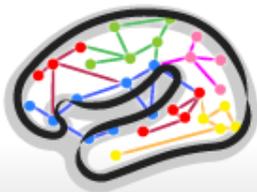
*160 - ctx-rh-S-parieto-occipital*



- Observations:

- They have similar **shapes** after removing the translation, rotation, scaling and re-paramterization.



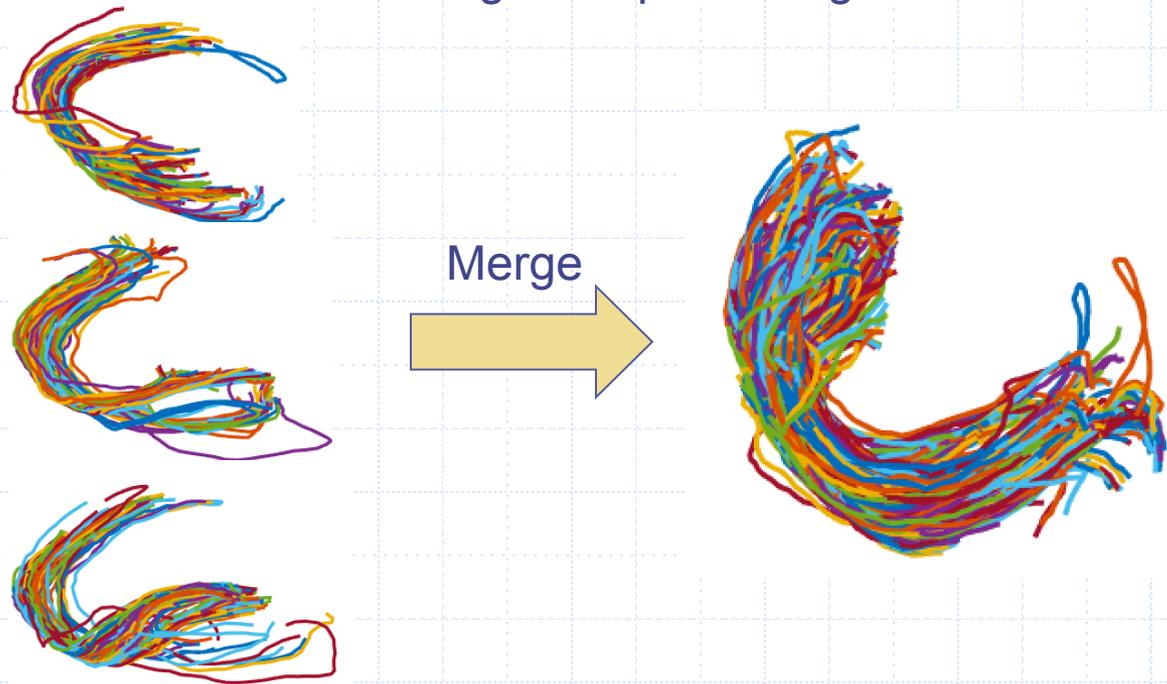


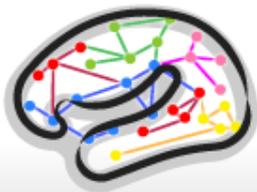
## Compression of Connectivity Matrix

- Compression happens at efficiently representing the **shapes of fibers**
- Learn basis functions to represent the fibers connecting a pair of regions

Step 1. Generate **atlas** for fibers connecting each pair of regions

Randomly select  
healthy subjects:





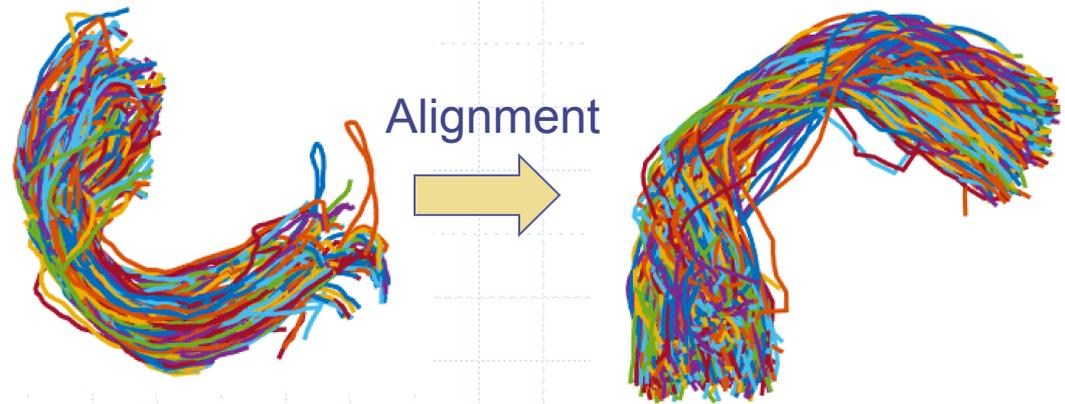
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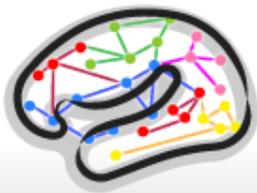
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Step 1. Generate **atlas** for fibers connecting each pair of regions

Step 2. Alignment using the **Elastic Shapes Analysis** framework (Srivastava et al. 2012)

- **rotation**
- **translation**
- **scaling**
- **re-parameterization**





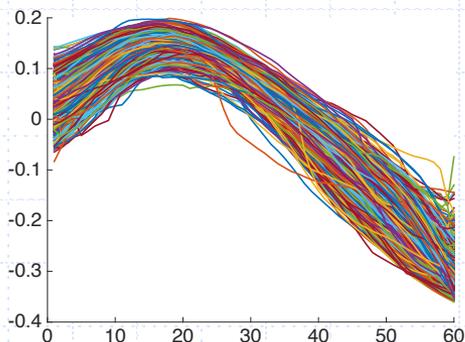
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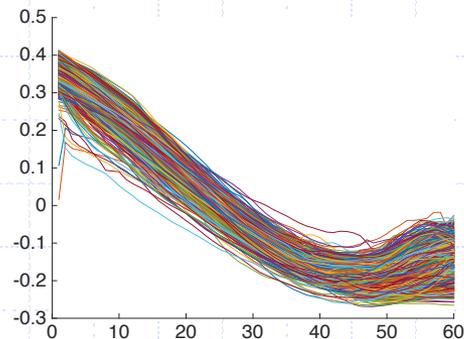
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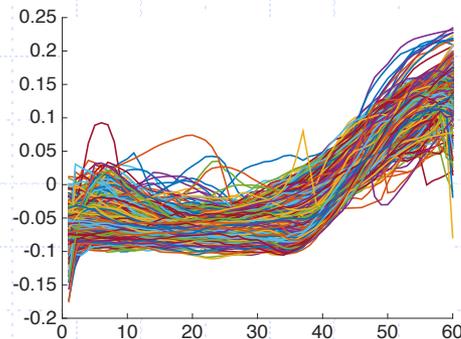
Step 3. Use **fPCA** to learn basis functions for each component



$j = 1$



$j = 2$



$j = 3$



$$\mu_j$$
$$\{\phi_{i,j}\}$$



## Compression of Connectivity Matrix

- Compression happens at efficiently representing the **shapes of fibers**
- Learn basis functions to represent the fibers connecting a pair of regions
- Encoding: **given a new fiber**  $f$

Step 1. Align  $f$  to the mean fiber in the atlas

$$\operatorname{argmin}_{O \in SO(3), C \in \mathbb{R}^3} \|O * (f - C) - \mu\|$$

$$g = \boxed{O} * (f - \boxed{C})$$

Step 2. Represent the aligned fiber using basis functions

$$\hat{g}_j = \mu_j + \sum_{i=1}^M \boxed{c_{j,i}} \phi_{j,i}$$

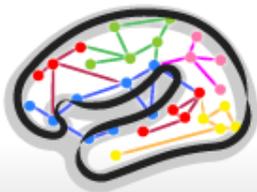
Parameters to save



## Compression of Connectivity Matrix

- Compression happens at efficiently representing the **shapes of fibers**
- Learn basis functions to represent the fibers connecting a pair of regions
- Encoding
- Decoding: reconstruct  $f$

$$\hat{f} = O' * \hat{g} + C$$



## Compression of Connectivity Matrix

- Compression happens at efficiently representing the **shapes of fibers**
- Learn basis functions to represent the fibers connecting a pair of regions
- Encoding
- Decoding
- Compression Ratio: **~ 90-98%**

$$\rho = 100 * \left(1 - \frac{N_c}{N_r}\right)$$

$N_c$  # parameters after compression

$N_r$  # parameters before compression



## Robust Coupling Strength Measures

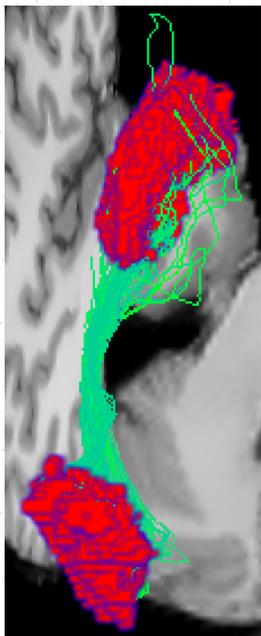
1. What is a good measure of the connectivity strength between two ROIs?

Right-Putamen (16)

ctx\_rh\_G\_occipital\_sup(115)

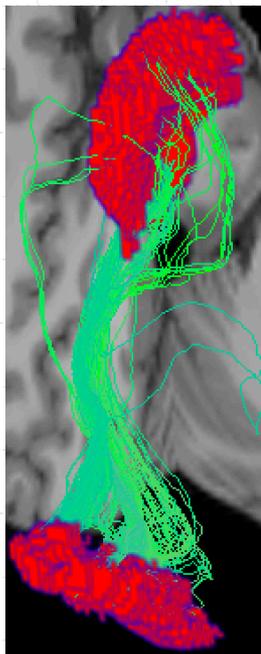
ID: 104820

$|CM(16,115)| = 483$



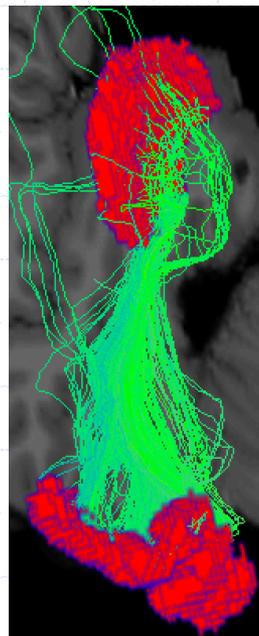
ID: 101006

$|CM(16,115)| = 758$



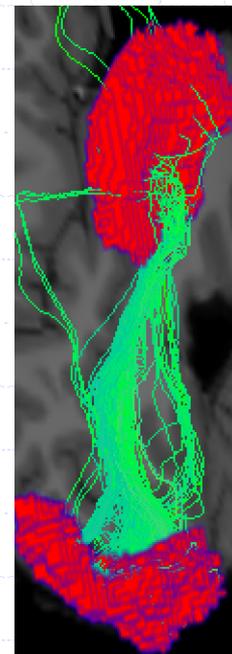
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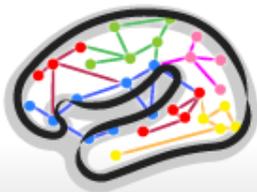
$|CM(16,115)| = 614$



ID: 145836

$|CM(16,115)| = 429$





## Robust Coupling Strength Measures

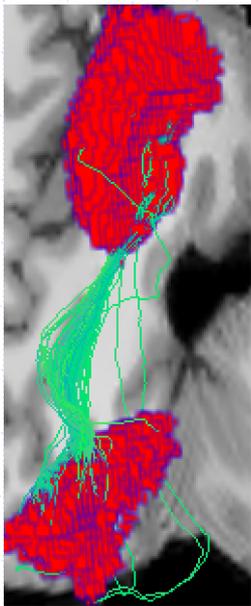
1. What is a good measure of the connectivity strength between two ROIs?

Right-Putamen (16)

ctx\_rh\_S\_parieto\_occipital(160)

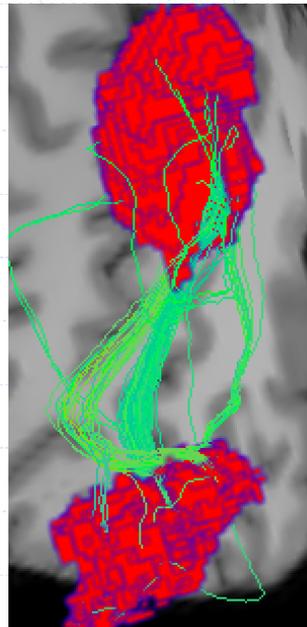
ID: 104820

|  $CM(16,115)$  | = 89



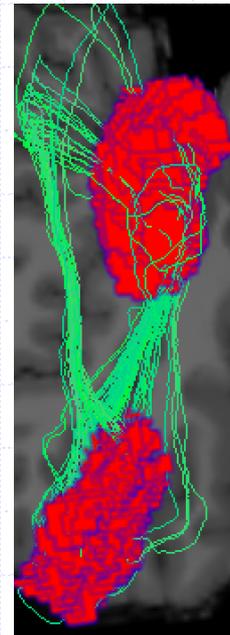
ID: 101006

|  $CM(16,115)$  | = 226



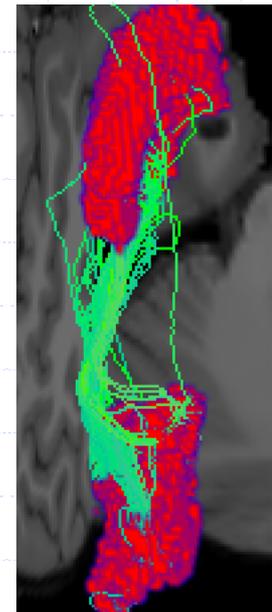
ID: 102311

|  $CM(16,160)$  | = 76



ID: 145836

|  $CM(16,160)$  | = 249





## Robust Coupling Strength Measures

### 1. What is a good measure of the connectivity strength between two ROIs?

- Current people use only counts
- Should include:
  - Diffusion properties: FA values, AFD values along fibers
  - Geometry properties: Shapes, Loops, Clusters
  - Nodes information: Volume of nodes, Connected surface areas
  - ...
- Working on extracting them now.
- Question: how to verify the measures?



HUMAN  
**Connectome**  
PROJECT

Mapping structural and functional connections in the human brain

**Thank You**