

# Structure of and Inquiries about Healthy Connectivity Matrix Data

Nooreen S Dabbish and Kaijun Wang

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# Outline

## Data Structure

- Introduction to connectivity matrix data

- UMCD: USC Multimodal Connectivity Database

- Our particular dataset

## Summary Plots and Numbers:

- Overview

- Correlation Plots

- qgraph, spring layout

- Degree Distribution

## Literature Research

- Raw datapreparation and processing

- Classification:

- Classification2

## Possible Topics

## Methods

- Graph Kernel

- Two-sample Test

# Basics

How it is organized. What types; what measurements are selected?

# Connectivity Matrices

- ▶ symmetric/undirected  $n \times n$  matrix
- ▶  $n \times 3$  list of Montreal Neurological Institute (MNI) coordinate spaces for each node
- ▶  $n \times 1$  lists of node names and abbreviations
- ▶ [data types] fMRI, Diffusion Tensor Images (DTI), Diffusion Spectrum Images (DSI), structural MRI, EEG, MEG

<http://umcd.humanconnectomeproject.org>

# Description in (Brown 2016)

- ▶ central repository for connectivity matrices
- ▶ click-of-the-mouse analyses
- ▶ (as of 1/17/16) 2254 brain networks (CMs), 21 studies
- ▶ all ages fetus to 89 yo.
- ▶ healthy, ADHD, autism, OCD, APOE-4 carrier status (risk for AD)

# Beijing Zang

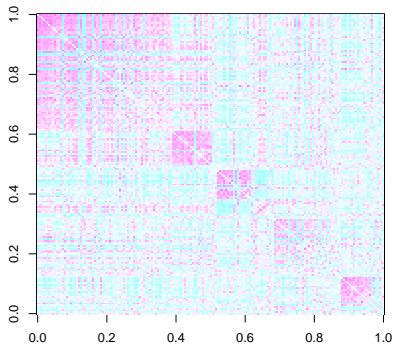
The dataset to be used: R-fMRI BOLD data from 148 subjects (74 female and 74 male; matched by age (21 years old) recruited as part of larger studies conducted in Beijing China. See the MATERIALS AND METHODOLOGY, Resting State Data of <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC3271304/pdf/TONIJ-6-1.pdf>

# Overview

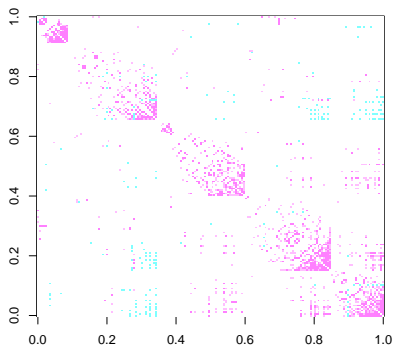
number of samples/variables/..histogram plot of some measurement to give a “feel” for the data. List of interesting modeling questions and why they are scientifically interesting?



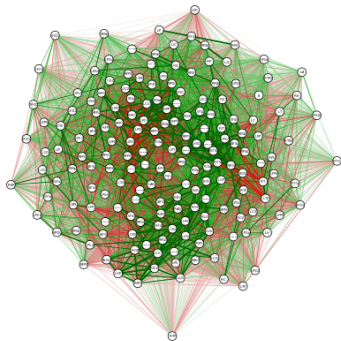
Before Thresholding



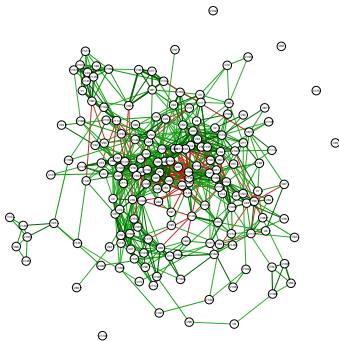
After thresholding (with 0.5)



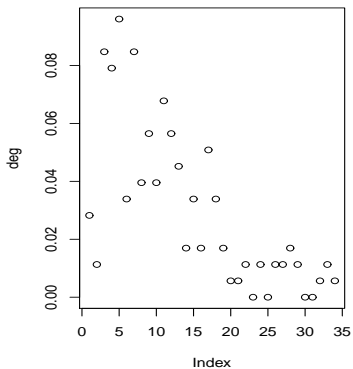
Before Thresholding



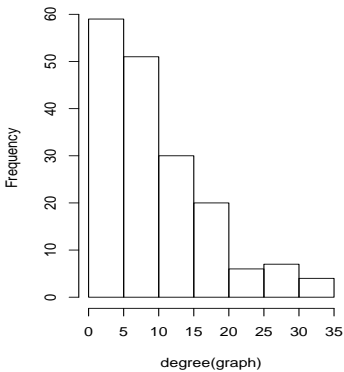
After Thresholding



**Degree Distribution of Nodes**



**Histogram of degree for Corr> .5**



# Literatures: On fMRI Data

- ▶ Network Centrality in the Human Functional Connectome  
Xi-Nian Zuo, et al. (2011)
- ▶ Toward reliable characterization of functional homogeneity in the human brain: Preprocessing, scan duration, imaging resolution and computational space  
Xi-Nian Zuo, et al. (2013)
- ▶ Fully exploratory network independent component analysis of the 1000 functional connectomes database  
Klaudius Kalcher, et al.(2013)

# Literatures: Classification Problems

- ▶ Combining Graph and Machine Learning Methods to Analyze Differences in Functional Connectivity Across Sex (2012) Casanova R., et al.  
Response: Gender  
Random forest and Lasso
- ▶ The Kernel Two-Sample Test for Brain Networks, Emanuele Olivetti et al. (2015)  
Response: Gender  
Classification based method vs Kernel Two-sample

# Literatures: Classification Problems

- ▶ Distinct neural signatures detected for ADHD subtypes after controlling for micro-movements in resting state functional connectivity MRI data Damien A. Fair, et al. (2013)  
Response: ADHD
- ▶ The Autism Brain Imaging Data Exchange: Towards Large-Scale Evaluation of the Intrinsic Brain Architecture in Autism Di Martino, et al.(2014)  
Response: Autism Spectrum Disorders  
Intrinsic Functional Connectivity Analyses between Autism and Control Group

# Possible Topics

- ▶ Classification according to the connectivity matrices, given exogenous data.
- ▶ Signal Path Clustering, normal pattern in brain activities
- ▶ Evaluate different classification methods on this data (e.g. random forest and lasso in [Cassanova 2012])

# Graph Kernel

Graph pairs  $G_i, G_j \rightarrow$  Kernel  $k_{ij} = \kappa(G_i, G_j) \rightarrow$  Classifier



# KTST

Functional connectomes, 148 graphs, 74 from group A and 74 from group B

Kernel matrix: 148-by-148, of pairwise kernels

graphs	$A_1$	$A_2$	$\dots$	$A_m$	$B_1$	$B_2$	$\dots$	$B_n$
$A_1$	$k_{A1,A1}$	$k_{A1,A2}$	$\dots$		$k_{A1,B1}$	$\dots$		
$A_2$	$k_{A2,A1}$							
$\vdots$	$\vdots$		$\ddots$					
$A_m$	$k_{Am,A1}$	$\dots$						
$B_1$	$k_{B1,A1}$				$k_{B1,B1}$	$k_{B1,B2}$	$\dots$	
$B_2$		$\ddots$			$k_{B2,B1}$			
$\vdots$	$\vdots$				$\vdots$		$\ddots$	
$B_n$	$k_{Bn,A1}$							

maximum mean discrepancy (MMD):

$$MMD^2 = E[k(x_A, x'_A)] + E[k(x_B, x'_B)] - 2E[k(x_A, x'_B)]$$

$$\widehat{MMD}^2 = \frac{1}{m(m-1)} \sum_{i \neq j} k(x_i^A, x_j^A) + \frac{1}{n(n-1)} \sum_{i \neq j} k(x_i^B, x_j^B) - 2 \frac{1}{nm}$$

Test hypothesis:  $H_0 : MMD = 0$

# KTST

- ▶ MMD changes with different matrix permutations
- ▶ in practice we calculate different MMDs according to various permutations, to see its distribution
- ▶ Total calculation complexity:  $M \cdot N^2$ , where, M is the number of permutations, and N is the number of graphs.

# Factor Analysis

- ▶ input factor number  $k$ , covariance matrix  $A$
- ▶ find  $k$  factors  $F_1, \dots, F_k$  according to  $A$
- ▶ find projection from factors to the original variables in  $A$

$$X_1 = \lambda_{11}F_1 + \lambda_{12}F_2 + \dots + \lambda_{1k}F_k$$

...

$$X_n = \lambda_{n1}F_1 + \lambda_{n2}F_2 + \dots + \lambda_{nk}F_k$$

# Factor Analysis

- ▶ The coefficients  $\lambda_{ij}$  are called loadings, group  $X_i$ s according to their highest loading into  $k$  groups
- ▶ permute the original matrix

Example:

if we have two groups with  $\{X_1, X_2, X_4\}$ ,  $\{X_3, X_5\}$  the original matrix will exchange the third and fourth row and column.