

# Practice in resting-state fMRI (rs-fMRI) Analysis: PART III

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## Course Arrangement

### PART I

- REST toolbox <http://restfmri.net/forum/index.php?q=rest>
- ReHo, ALFF, fALFF, Statistics

### PART II

- REST toolbox
- Functional connectivity (seed-based, atlas-based)
- FC strength mapping

## Course Arrangement

### PART III

- Large-scale network analysis
- Graph theory: topological properties (degree, strength, efficiency, clustering...)

### PART IV

- Dynamic functional connectivity

## Employed Software/Package

### 1. SPM preprocessing

- <http://www.fil.ion.ucl.ac.uk/spm/>

### 2. REST functional connectivity, ReHo, ALFF, fALFF, VMHC

- <http://restfmri.net/forum/index.php?q=rest>

### 3. IBASPM 64-bit

- [http://www.ym.edu.tw/~cflu/software/Ibaspm\\_64.zip](http://www.ym.edu.tw/~cflu/software/Ibaspm_64.zip)

### 4. Brain Connectivity Toolbox/Network Based Statistic Toolbox

- <https://sites.google.com/site/bctnet/>

### 5. GraphVar

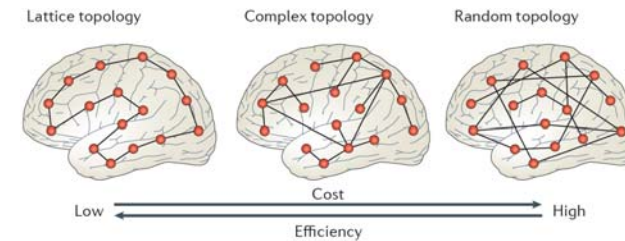
- <http://www.rfmri.org/graphvar>

## Complex Network Analysis

## Human brain networks Wiring costs $\leftrightarrow$ efficiency



- Clusters of lattice-like short-distance connections between spatially neighboring nodes
- Topologically direct interconnections between spatially remote brain regions  $\rightarrow$  increase efficiency of information processing
- Nodes aggregated topologically and anatomically as modules  $\rightarrow$  minimize wiring cost



Bullmore et al., Nature Reviews Neuroscience, 13: 336-349, 2012.

## Network construction

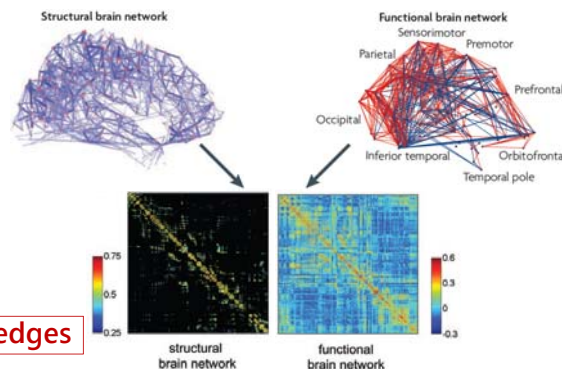


### Nodes

- Cortical regions

### Edges

- Cortical thickness correlations
- Fiber connections
  - DSI, DTI, transneuronal tracers
- Functional connectivity
  - fMRI, EEG, MEG

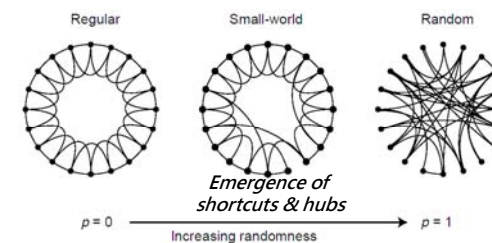


**Network = nodes + edges**

## Complex networks



- Brain have a small-world architecture.

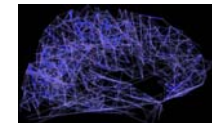


- $\triangleright$  High local clustering  $\triangleright$  Local segregation
- $\triangleright$  Low separation  $\triangleright$  Global integration

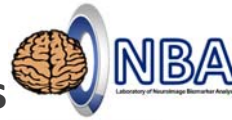
high signal-propagation speed, computational power, and synchronizability

Watts DJ, Strogatz SH, Nature 393:440-442, 1998.

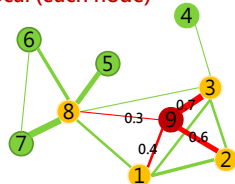
- Complex networks
- Social network
- WWW internet
- Biological system
- Brain network



## Graph theory: topological properties



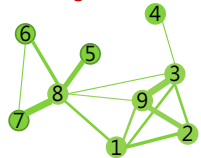
### Local (each node)



- **degree** (the number of neighbors)  
*e.g. degree of node 9 = 4*

- **strength** (the connected correlation coefficient)  
*e.g. strength of node 9*  
 $= (0.3 + 0.4 + 0.6 + 0.7) / 4 = 0.5$

### Global (average over all nodes)



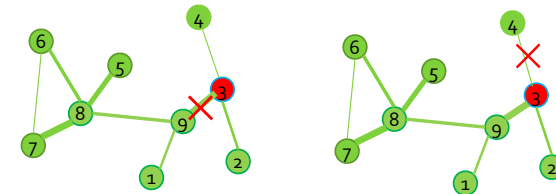
- **clustering coefficient** (the connection between neighbors, [0~1])  
*e.g. clustering coefficient of node 9*  
 $= 5/6 = 0.83$
- **shortest path length (separation)** (the minimal steps for connection)  
*e.g. path length from node 9 to node 6*  
 $= 2 \text{ steps } (9 \rightarrow 8 \rightarrow 6)$

Salvador et al, *Philos Trans R Soc Lond B Biol Sci*, 360, 937-946, 2005

## Network properties



- The topological observations can reveal a "hidden" or "high-level" relations between nodes.



## Brain Connectivity Toolbox



NeuroImage 52 (2010) 1059-1069



Contents lists available at ScienceDirect

NeuroImage

journal homepage: [www.elsevier.com/locate/ynimg](http://www.elsevier.com/locate/ynimg)



### Complex network measures of brain connectivity: Uses and interpretations

Mikhail Rubinov<sup>a,b,c</sup>, Olaf Sporns<sup>d,\*</sup>

<sup>a</sup> Black Dog Institute and School of Psychiatry, University of New South Wales, Sydney, Australia  
<sup>b</sup> Mental Health Research Division, Queensland Institute of Medical Research, Brisbane, Australia  
<sup>c</sup> CSIRO Information and Communication Technologies Centre, Sydney, Australia  
<sup>d</sup> Department of Psychological and Brain Sciences, Indiana University, Bloomington, IN 47405, USA

## Topological Measurements



### Basic Measures

- degree, strength, shortest path length

### Measures of integration

- global efficiency

### Measures of segregation

- Clustering coefficient, local efficiency, modularity

### Measures of centrality

- Betweenness, within-module degree, participation coefficient

### Network motifs

### Measures of resilience

- Degree distribution, neighbor degree, assortativity coefficient

### Network small-worldness

# Network Analysis Using GraphVar

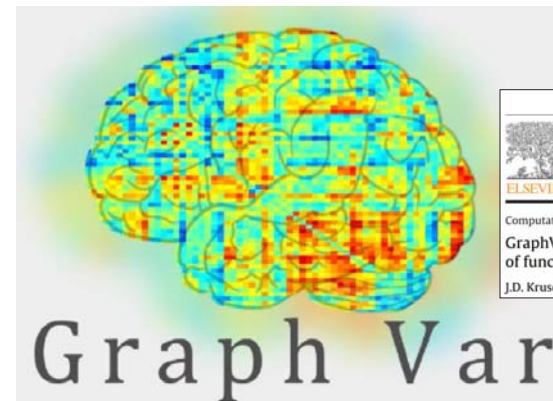
MATLAB R2013b or older version is recommended.

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## GUI for network analysis

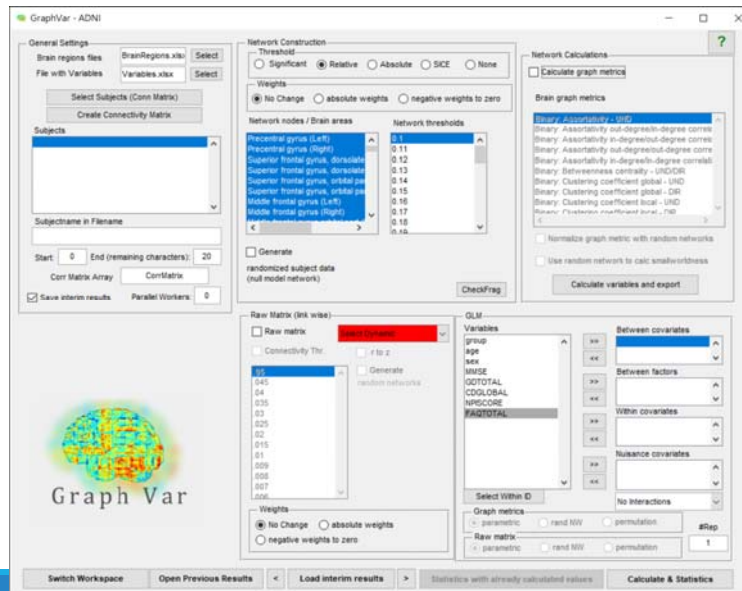


Set Path  
>> start\_GraphVar

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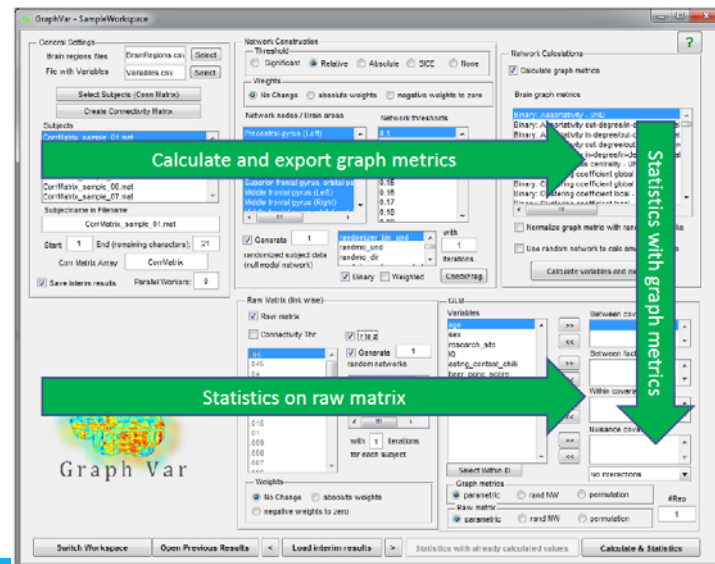


GraphVar  
using BCT toolbox

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Workflow

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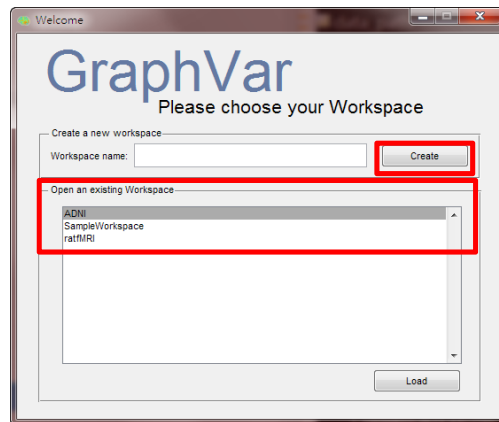
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## Step 1: Preparation of Workspace



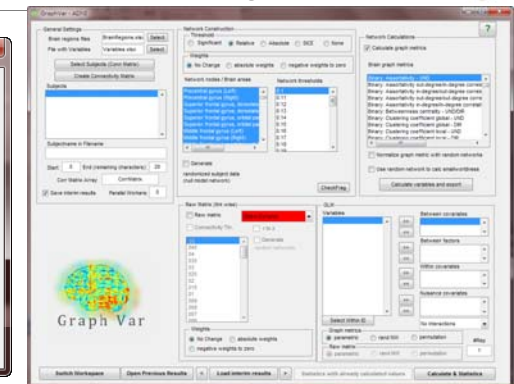
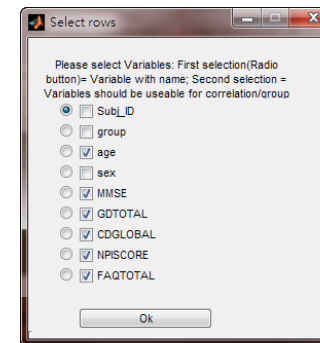
Set Path

>> start\_GraphVar



Use pre-prepared workspace folder may be better!

## Step 1: Preparation of Workspace



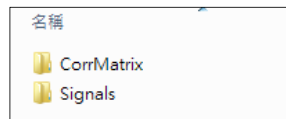
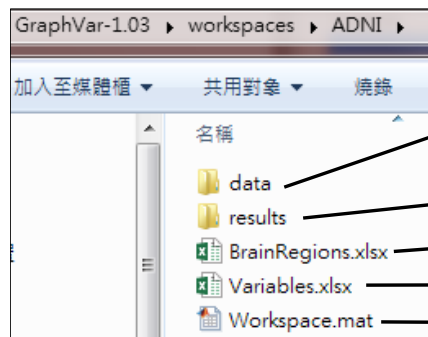
↑ This query window displays only when the **"ImportSettings.mat"** file is absent in the study folder. (delete ImportSettings.mat file and restart GraphVar to reset configuration)

## Step 1: Preparation of Workspace



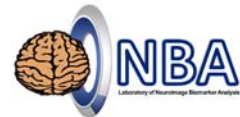
Under **..\GraphVar-1.03\workspaces**

Create a Folder for your study (ex: ADNI, ratfMRI)



- 8 default empty folders and InterimResultsID.mat (Copy to use)
- Node definition for graph analyses
- Subject specific data for statistical analyses
- Default configuration file (Copy to use)

## Step 1: Preparation of Workspace



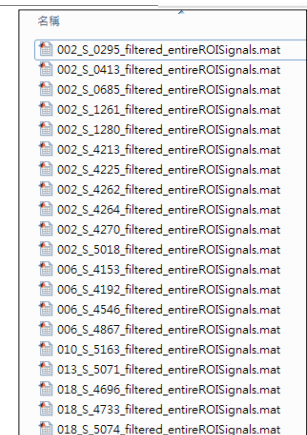
(1) **data** folder

**CorrMatrix**

- One mat-file of connectivity matrix (**Node x Node**) for each subject.
- Should have 20 files for 20 subjects.

**Signals (recommended for GraphVar)**

- One mat-file of preprocessed regional signals (**Time frame x Node**) for each subject.
- Should have 20 files for 20 subjects.





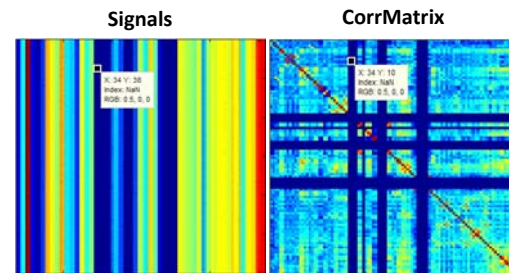
## Step 1: Preparation of Workspace



Nan in Signals/CorrMatrix (caused by empty brain regions) will cause calculation error later!

tlstsq dgelsd error !!

(nan when polyfit) during GLM



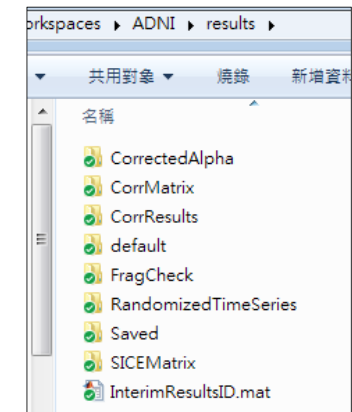
Exclude empty regions for all subjects!!

## Step 1: Preparation of Workspace



(2) **result** folder

8 default empty folders and InterimResultsID.mat



## Step 1: Preparation of Workspace



(3) **BrainRegions.xlsx** file (default: AAL 116)

Category	Node (abbr.)	Node (full name)	Coordinates (X, Y, Z)			
	A	B	C	D	E	F
1		1 Precentral_L	Precentral gyrus (Left)	-40	-6	51
2		1 Precentral_R	Precentral gyrus (Right)	40	-8	52
3		1 Frontal_Sup_L	Superior frontal gyrus, dorsolateral (Left)	-19	35	42
4		1 Frontal_Sup_R	Superior frontal gyrus, dorsolateral (Right)	20	31	44
5		1 Frontal_Sup_Orb_L	Superior frontal gyrus, orbital part (Left)	-18	47	-13
6		1 Frontal_Sup_Orb_R	Superior frontal gyrus, orbital part (Right)	17	48	-14
7		1 Frontal_Mid_L	Middle frontal gyrus (Left)	-34	33	35
8		1 Frontal_Mid_R	Middle frontal gyrus (Right)	37	33	34
9		1 Frontal_Mid_Orb_L	Middle frontal gyrus orbital part (Left)	-32	50	-10
10		1 Frontal_Mid_Orb_R	Middle frontal gyrus orbital part (Right)	32	53	-11
11		1 Frontal_Inf_Oper_L	Inferior frontal gyrus, opercular part (Left)	-49	13	19
12		1 Frontal_Inf_Oper_R	Inferior frontal gyrus, opercular part (Right)	49	15	21

(1 for default selection)

The order must comply with that listed in data!

## Step 1: Preparation of Workspace



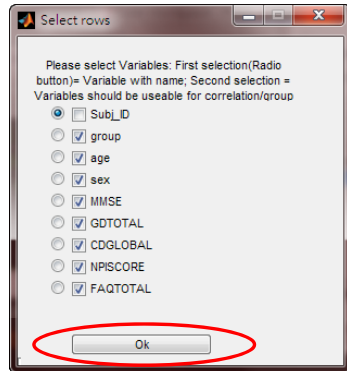
(4) **Variables.xlsx** file

Subject ID		group		Personal/Clinical Data					
	A	B	C	D	E	F	G	H	I
1	Subj_ID	group	age	sex	MMSE	GDTOTAL	CDGLOBAL	NPISCORE	FAQTOTAL
2	002_S_5018	AD	74.4	M	22	2	1	3	17
3	006_S_4153	AD	81.4583333	M	19	2	0.5	0	8
4	006_S_4192	AD	84.3416667	M	17	1	0.5	3	7
5	006_S_4546	AD	71.5777778	M	24	3	1	2	14
6	006_S_4867	AD	75.6527778	M	24	1	2	12	22
7	010_S_5163	AD	67.4111111	M	23	2	0.5	0	8
8	013_S_5071	AD	76.5777778	M	21	4	1	14	20
9	018_S_4696	AD	73.5333333	F	16	0	1	5	20
10	018_S_4733	AD	75.8472222	M	21	1	1	4	26
11	018_S_5074	AD	74.8166667	F	25	1	1	4	26
12	002_S_0295	NC	90.9083333	M	22	0	0	0	0
13	002_S_0413	NC	83.3611111	F	30	0	0	0	0

Subject ID must be included in the filename.

## Step 2: General Settings

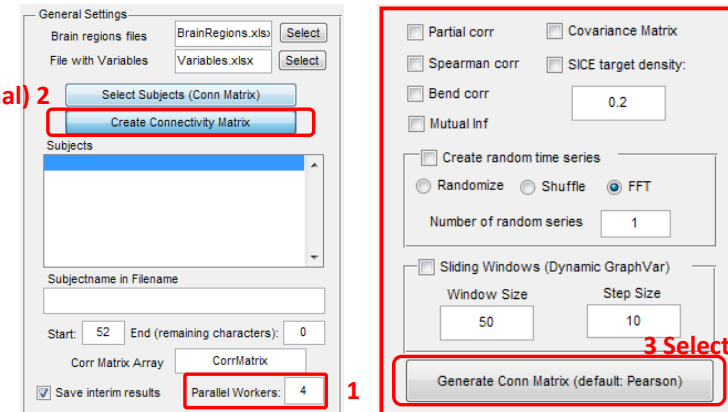
**Select Variables** (First column: Subject ID; Second column: other variables)



The **"ImportSettings.mat"** file will be generated in the study folder (delete ImportSettings.mat file and restart GraphVar to reset configuration)

## Step 2: General Settings

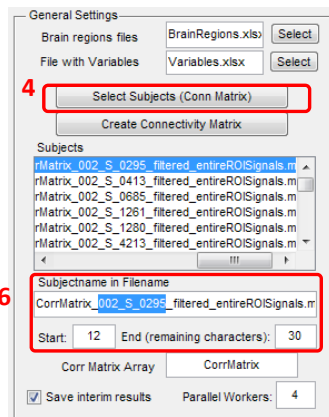
(optional) 2



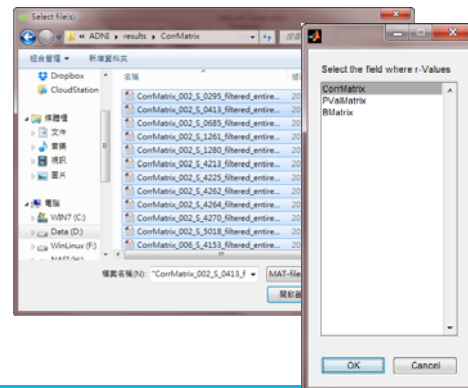
Mouse-over help

3 Select Signals

## Step 2: General Settings

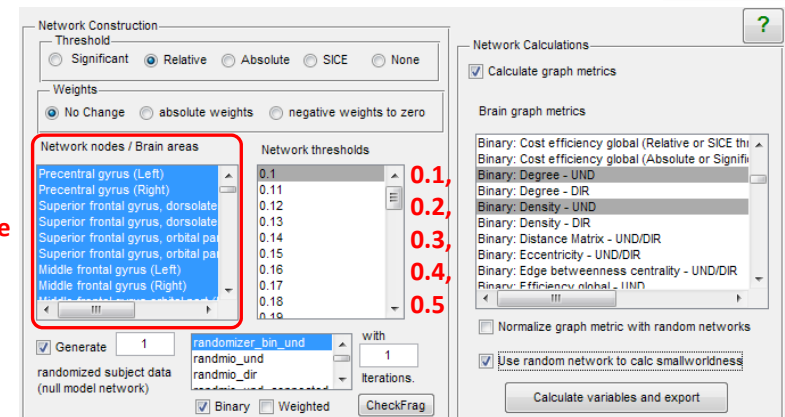


5 Select results/CorrMatrix



## Step 3: Network Construction

Can select a subset for the subnetwork analysis



## Step 3: Network Construction



### (1) **Threshold** method

#### Significant

- Thresholded based on the p-values.

#### Relative

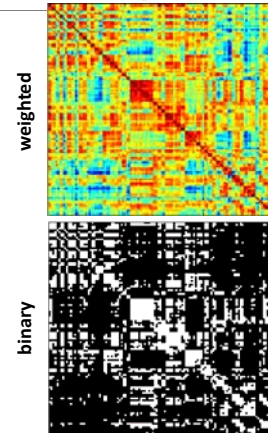
- Thresholded in a proportional way.

#### Absolute

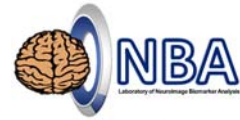
- Thresholded based on the correlation coefficients.

#### None

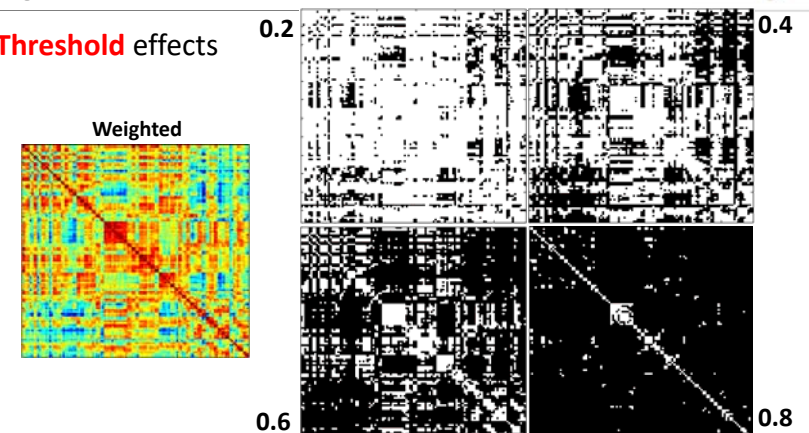
- No thresholding. Preserve full graph.



## Step 3: Network Construction



### (1) **Threshold** effects



## Step 3: Network Construction



### Common Abbreviations in Network Analysis

- bin/BIN: binary graph
- und/UND: undirected graph
- dir/DIR: directed graph
- w/W: weighted graph
- wu/WU: weighted and undirected graph
- bu/BU: binary and undirected graph

## Step 3: Network Construction



### (2) Generate **null-model network**

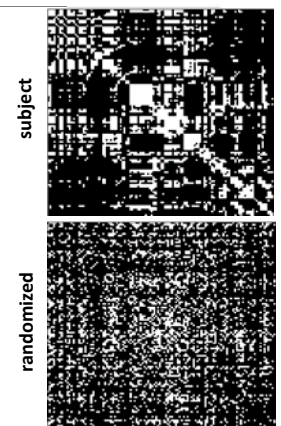
#### Usage

- Used for normalization of the network topological measures.
- Non-parametric testing of the significance against random networks.

#### BCT functions

- randomizer\_bin\_und*, *randmio\_und*, *randmio\_und\_connected*, *null\_model\_und\_signed*.

<https://sites.google.com/site/bctnet/Home/functions>





## Step 3: Network Construction

**Progress monitoring window**

Network Calculations

☒ Calculate graph metrics

Brain graph metrics

Binary: Assortativity - UND

Binary: Assortativity out-degree/in-degree correl

Binary: Assortativity in-degree/out-degree correl

Binary: Assortativity out-degree/out-degree correl

Binary: Assortativity in-degree/in-degree correl

Binary: Betweenness centrality - UND/DIR

Binary: Clustering coefficient global - UND

Binary: Clustering coefficient global - DIR

Binary: Clustering coefficient local - UND

Binary: Clustering coefficient local - DIR

☒ Normalize graph metric with random networks

☒ Use random network to calc smallworldness

**Calculate variables and export**

Progress

All Tasks 11479 of 11480 Operations (100%)

Thresholding 41 of 41

Randomize Subject 20 of 20

Graph Function 6 of 6

Subject 40 of 40

Exporting

Cancel (or MainWin.Ctrl+C)

**Calculate & export to Excel files**

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## Step 4: Statistics (GLM)

GLM

Variables

Between covariates

MMSE

CDGTOTAL

CDGLOBAL

Between factors

group

Within covariates

Nuisance covariates

age

sex

No interactions

Select Within ID

Graph metrics

☐ parametric ☐ rand V/V ☒ permutation

Raw matrix

☒ parametric ☐ rand NW ☐ permutation

#Rep

1000

Statistics with already calculated values

**Calculate & Statistics**

Progress

All Tasks 690 of 690 Operations (88%)

Thresholding 5 of 5

Randomize Subject

Graph Function 3 of 3

Subject 40 of 40

GLM

Cancel (or MainWin.Ctrl+C)

**Relevant clinical data (multivariate regression)**

**Group labeling**

**Repeated measurement**

**Confounding effects to remove**

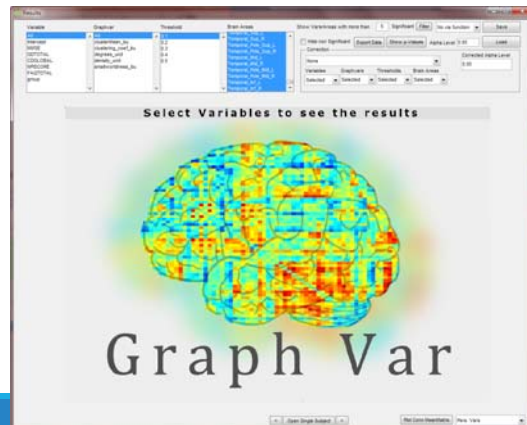
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## Step 5: Results viewer

Automatically pop-out once the [Calculate & Statistics] is done.



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## Step 5: Results viewer

Variable	Graphvar	Threshold	Brain Areas
All	All	0.1	Temporal_Sup_L
Intercept	cluster/mean_bu	0.2	Temporal_Sup_R
MMSE	clustering_coef_bu	0.3	Temporal_Pole_Sup_L
CDGTOTAL	degrees_und	0.4	Temporal_Pole_Sup_R
CDGLOBAL	density_und	0.5	Temporal_Mid_L
NPISCORE	smallworldness_bu		Temporal_Mid_R
FAQTOTAL			Temporal_Pole_Mid_L
group			Temporal_Pole_Mid_R
			Temporal_Inf_L
			Temporal_Inf_R

Show Vars/Areas with more than 5 Significant Filter No vis function Save

☐ Hide non Significant Export Data Show p-Values Alpha Level 0.05 Load

Correction

None

Corrected Alpha Level 0.05

Variables Graphvars Thresholds Brain Areas

Selected Selected Selected Selected

Open Single Subject

**Left upper panel**

**Right upper panel**

**Bottom**

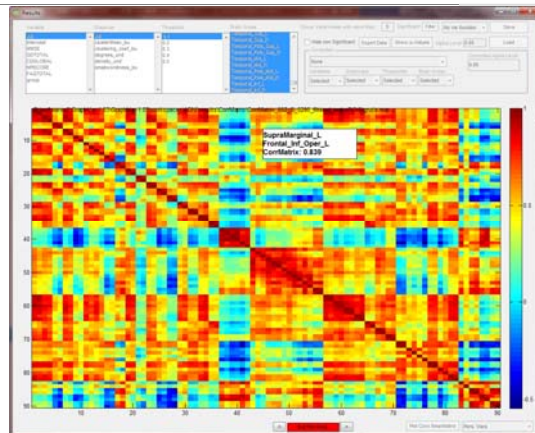
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## Step 5: Results viewer

Display of single subject's CorrMatrix



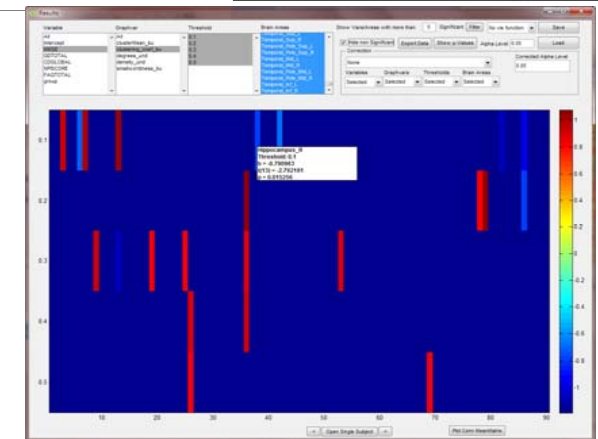
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## Step 5: Results viewer

Display of regression results



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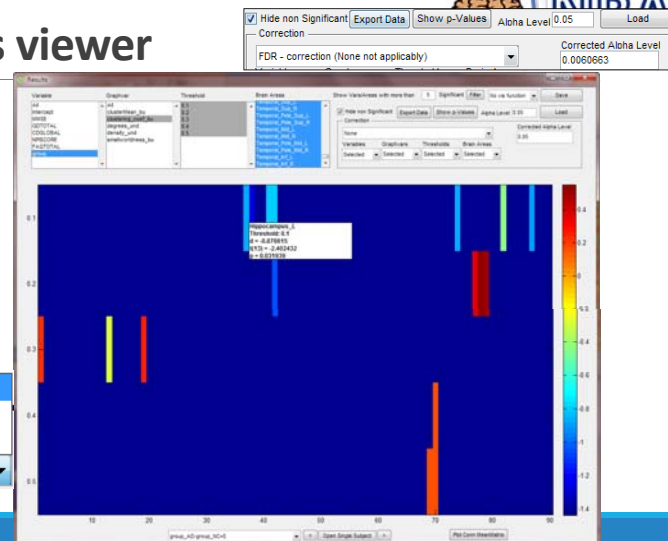
## Step 5: Results viewer

Display mode for local measurements:

Threshold x Brain regions

Group Comparisons:

group\_AD-group\_NC=0  
group\_AD=0  
group\_NC=0  
group\_AD-group\_NC=0



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## Step 5: Results viewer

Display mode for global measurements:

Measures under different thresholds

group\_AD-group\_NC=0  
group\_AD=0  
group\_NC=0  
group\_AD=0



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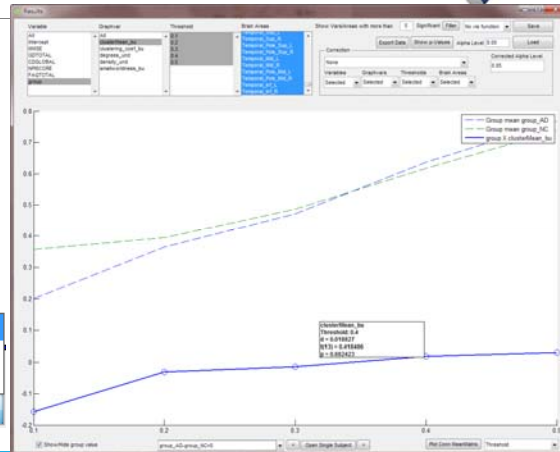
40

## Step 5: Results viewer

Display mode for global measurements:

Measures under different thresholds

group\_AD-group\_NC=0  
group\_AD=0  
group\_NC=0  
group\_AD-group\_NC=0



# Q & A