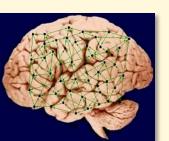


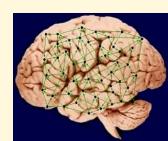
Multivariate Modeling and Inference for Brain Networks: ERGMs and Mixed Models

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Outline

I. Motivation

II. Brain Network Construction and Description

III. Multivariate Modeling and Inference: ERGMs

IV. Multivariate Modeling and Inference: Mixed Models

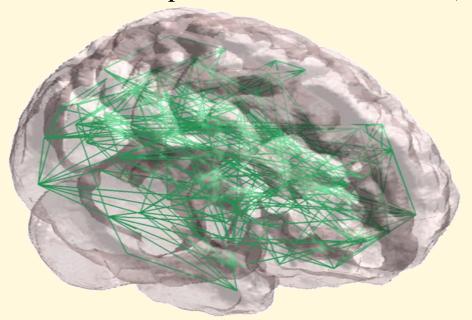
V. Summary

VI. Useful References

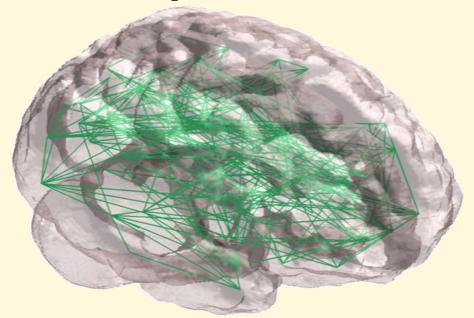
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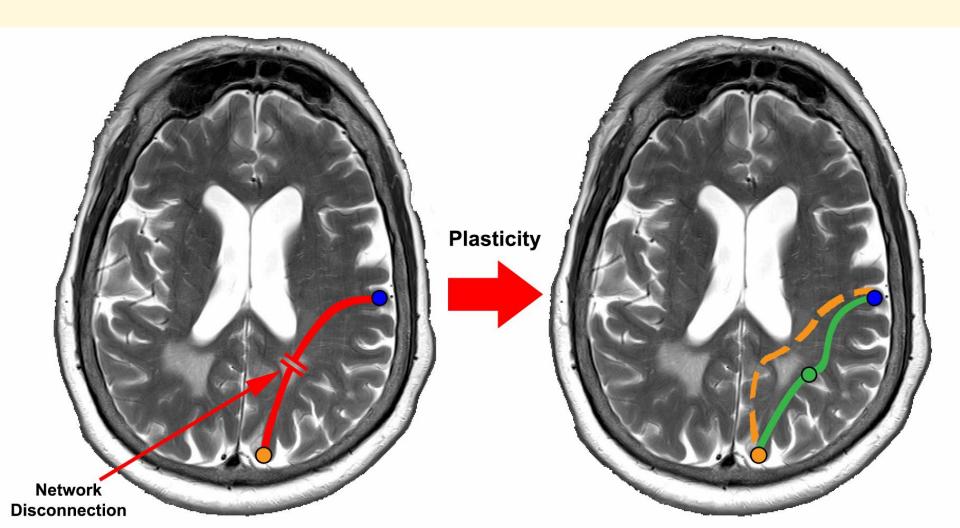


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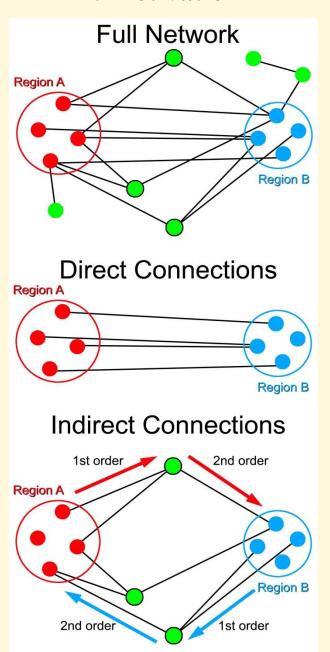


• FC underlie network analyses, subtle distinction overlooked in the literature.

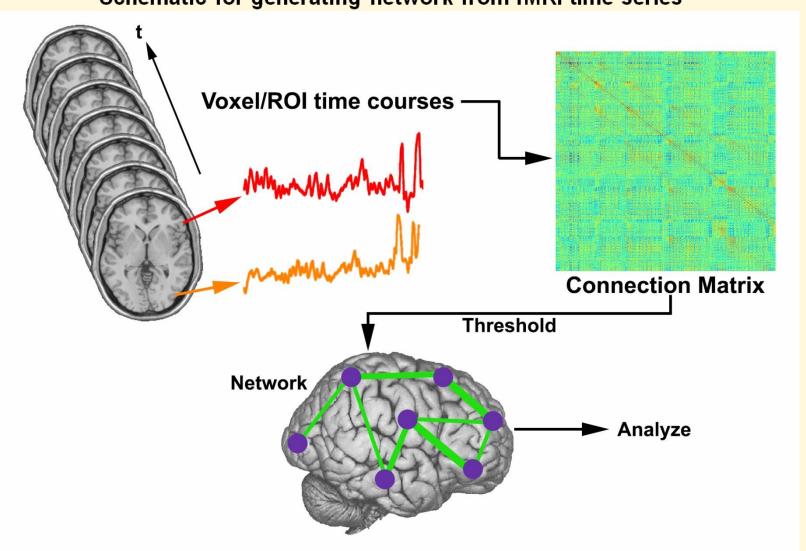
• Systemic organization confers functional abilities as connections may be lost due to adverse health condition, but compensatory connections may develop to maintain organizational consistency and functional performance.



• Also,...



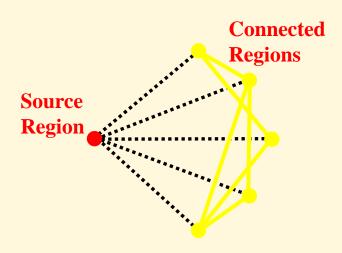
Schematic for generating network from fMRI time series



SMALL-WORLD METRICS

clustering coefficient (C)

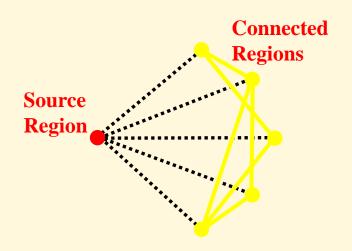
Proportion of a region's connections that are connected to each other

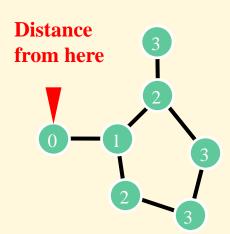


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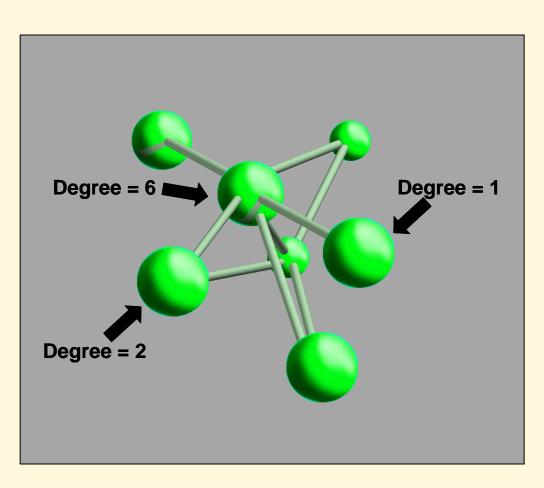




path length (L)

Average shortest distance between region pairs

<u>Degree</u>

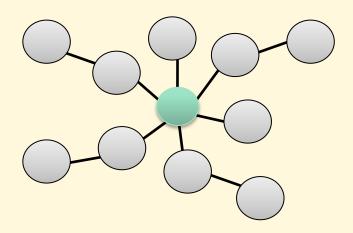


• Degree -K

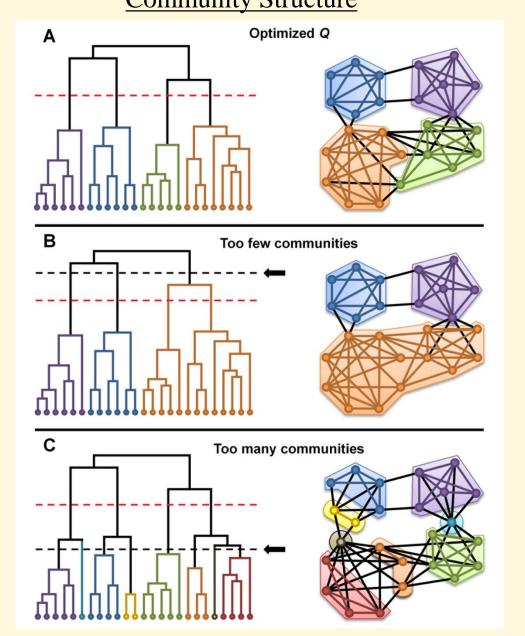
- Number of connections for each node
- Distribution is assessed to evaluate network type/resilience properties
- Assortativity is assessed to evaluate network type/resilience properties

Graph Centrality and Information Flow

<u>Leverage centrality (LC)</u> identifies nodes that have <u>high</u> degree relative to neighbors



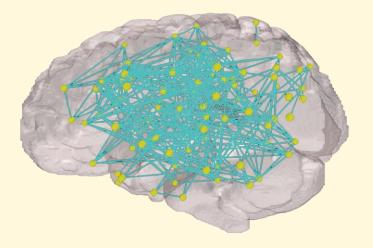
II. Brain Network Construction and Description <u>Community Structure</u>

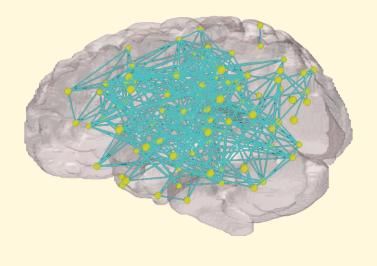


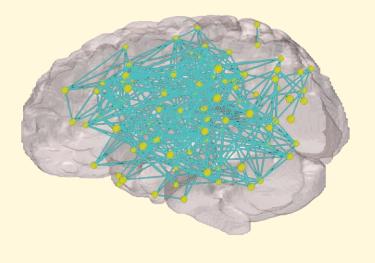
Need a multivariate explanatory and predictive brain network model.

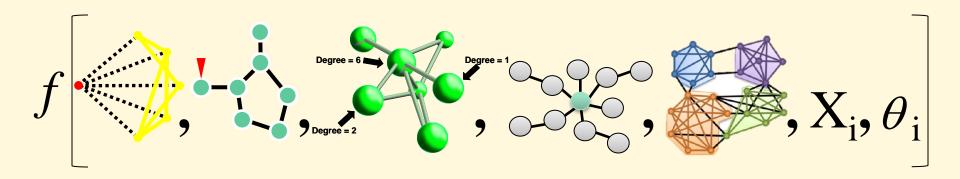
```
Data \begin{cases} Y_i : \text{ network of subject i} \\ X_i : \text{ covariate information (network metrics, demographics, etc.)} \end{cases} \theta_i : \text{ parameters}
```

Want $P(Y_i|X_i,\theta_i)$









Exponential random graph models have the following form:

$$P(\mathbf{Y} = \mathbf{y}) = \kappa(\boldsymbol{\theta})^{-1} \exp\{\boldsymbol{\theta}^{\mathrm{T}} \mathbf{g}(\mathbf{y})\}$$
(1)

where

Y $n \times n$ (n nodes) random symmetric adjacency matrix, $Y_{ij} = 1$ if an edge exists between nodes i and j and $Y_{ij} = 0$ otherwise;

 $\mathbf{g}(y)$ vector of prespecified network statistics (functions of network);

 $\boldsymbol{\theta}$ vector of parameters associated with $\mathbf{g}(\boldsymbol{y})$ (importance, Δ log-odds);

 $\kappa(\boldsymbol{\theta})$ normalizing constant ensuring probabilities sum to one.

Goal: Identify local metrics $\mathbf{g}(y)$ that concisely summarize the global (whole-brain) network structure.

• Once most appropriate statistics established, parameter profiles (θ) can be utilized to classify and compare whole-brain networks.

E.g., Best Model:

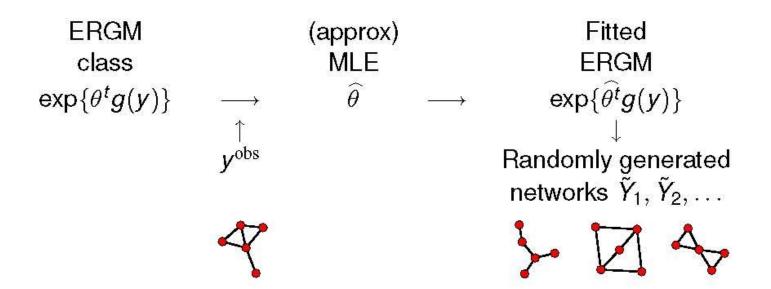
$$P(Y = y) = \frac{1}{\kappa} \exp\left\{\theta_1 + \theta_2 + \theta_3 + \theta_3\right\}$$

- Caveats: comparisons require use of a uniform set of predictors for all networks (due to predictor interdependencies) and balanced networks (same number of nodes for all networks) due to dependence of predictors on network size.

• Use graphical goodness-of-fit (GOF) approach (Hunter et al., 2008) to establish most appropriate set of explanatory metrics for each subject's brain network.

(Hunter, 2007)

Goodness of fit intuition

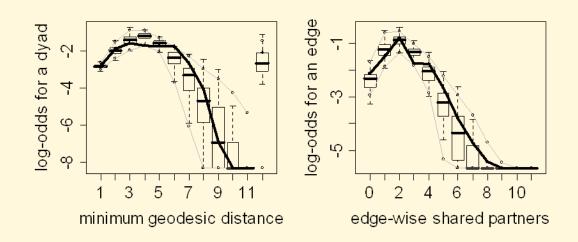


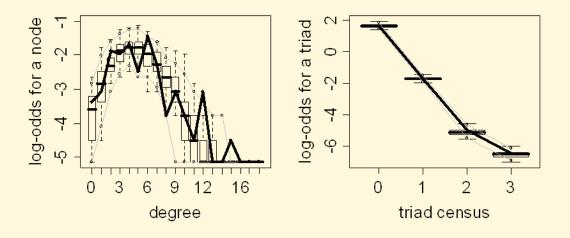
• Question: How does y^{obs} "look" as a representative of the sample $\tilde{Y}_1, \tilde{Y}_2, \ldots$?

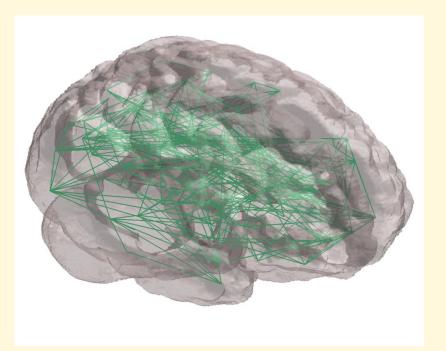
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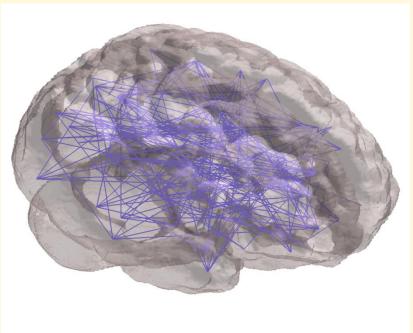
- POC: ERGMs fitted to networks from 10 normal subjects (Simpson et al., 2011)
 - Several R packages available: ergm, ergm.count, GERGM, Bergm, btergm, tergm xergm, xergm.common, blkergm, hergm.

Final ERGMs (composed of most informative explanatory metrics) for each subject provided a good fit to the data as evidenced by graphical GOF plots.







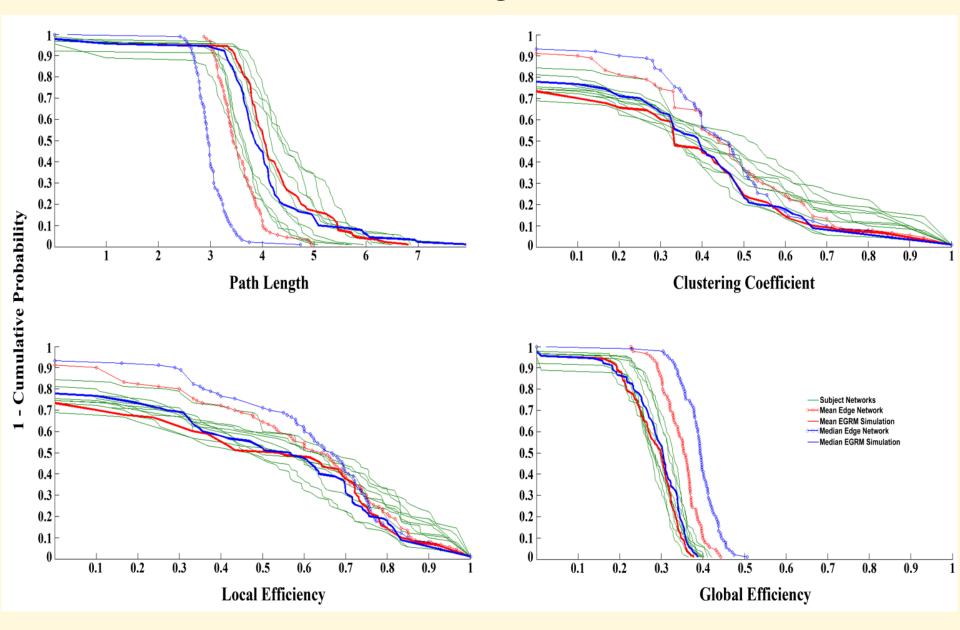


Observed Network

Simulated Network

- Create group "representative" networks via simulation (Simpson et al., 2012).
 - Traditional mean/median networks are edge-based and topologically differ greatly.

Subjects (and others)



Advantages

• Statistically principled approach to topologically modeling, analyzing and simulating complex brain networks.

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- Greatest appeal lies in ability to efficiently represent complex network data and allow examining way in which a network's global structure and function depend on its local structure.

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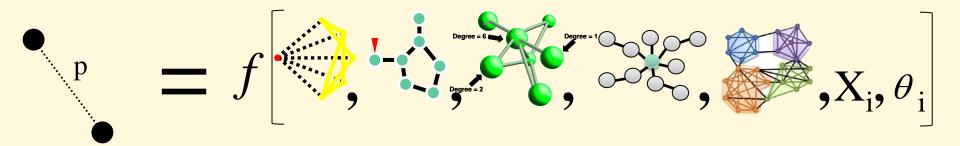
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 - Due to degeneracy issues that may arise.
- Developed for static binary networks.
 - Development for longitudinal and weighted networks in infancy.

IV. Multivariate Modeling and Inference: Mixed Models

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Presence:

$$logit(p_{ijk}) = \boldsymbol{X'_{ijk,1}} \boldsymbol{\beta_{Net}} + \boldsymbol{X'_{ijk,2}} \boldsymbol{\beta_{COI,Con,Int}} + \theta_{ijk}$$

Strength:

$$FZT(S_{ijk}) = \boldsymbol{X'_{ijk,1}} \boldsymbol{\beta_{Net}} + \boldsymbol{X'_{ijk,2}} \boldsymbol{\beta_{COI,Con,Int}} + \theta_{ijk}$$

$$\theta_{pi} = \mathbf{Z}'_{ijk} \boldsymbol{b}_{pi} = \mathbf{Z}'_{ijk} [b_{pi,0} \ \boldsymbol{b}_{pi,net} \ \boldsymbol{b}_{pi,dist} \ \boldsymbol{\delta}_{pi,j} \ \boldsymbol{\delta}_{pi,k}]'$$

$$\theta_{si} = \mathbf{Z}'_{ijk} \boldsymbol{b}_{si} + e_{ijk} = \mathbf{Z}'_{ijk} [b_{si,0} \ \boldsymbol{b}_{si,net} \ \boldsymbol{b}_{si,dist} \ \boldsymbol{\delta}_{si,j} \ \boldsymbol{\delta}_{si,k}]' + e_{ijk}$$

 $b_{i,0}$ deviation of subject-specific intercepts (from population)

 $\boldsymbol{b}_{i,net}$ deviation of subject-specific metric-edge relationships

 $\boldsymbol{b}_{i,dist}$ deviation of subject-specific spatial distance-edge relationships

 $\delta_{i,j/k}$ propensity for node j/k (of given dyad) to be connected and magnitude of its connections

1) Explain: quantifies relationship between Net/COI/Con and probability/strength of connections.

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- 4) Threshold: leverages group-level data to better distinguish between "true" weak connections and noise in individual-level networks.
- 5) Simulate: simulates group- and individual-level networks useful for model GOF assessments, representative network creation, and network variability assessment.

- **Aging Brain:** assess neurological underpinnings of cognitive decline by examining effects of aging on integration of sensory information.
- Young Adults: 27 ± 5.8 y/o (n=20) Older Adults: 73 ± 6.6 y/o (n=19)
- Three separate conditions of fMRI scans:
 - Rest
 - Visual (viewing of a silent movie)
 - Multisensory (MS) (visual and auditory movie with sound)
- 90 node AAL atlas based networks constructed for each participant.

Here,

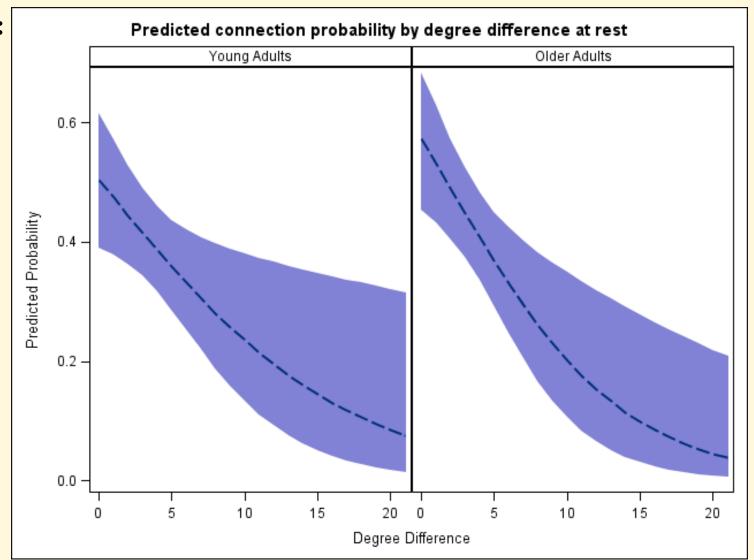
$$oldsymbol{eta}_{Net} = \left[eta_{C_{ ext{avg}}} \; eta_{Eglob_{ ext{avg}}} \; eta_{K_{ ext{diff}}} \; eta_{LC_{ ext{avg}}} \; eta_{Q}
ight]'.$$

$$\beta_{COI} = \beta_{age}$$
.

$$oldsymbol{eta}_{Con} = [eta_{sex} \ eta_{educ} \ eta_{dist} \ eta_{dist^2}]'.$$

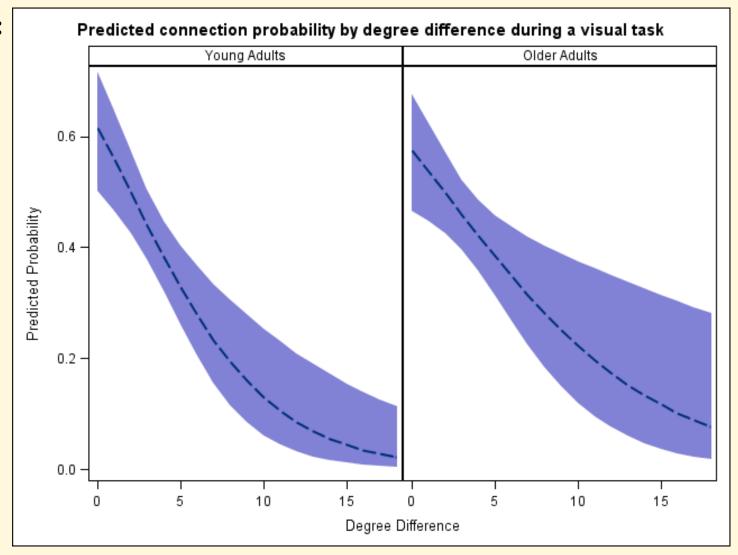
$$\boldsymbol{\beta}_{Int} = [\beta_{age \times C} \ \beta_{age \times Eglob} \ \beta_{age \times K} \ \beta_{age \times LC} \ \beta_{age \times Q} \ \beta_{age \times sex}]'.$$

Predict:



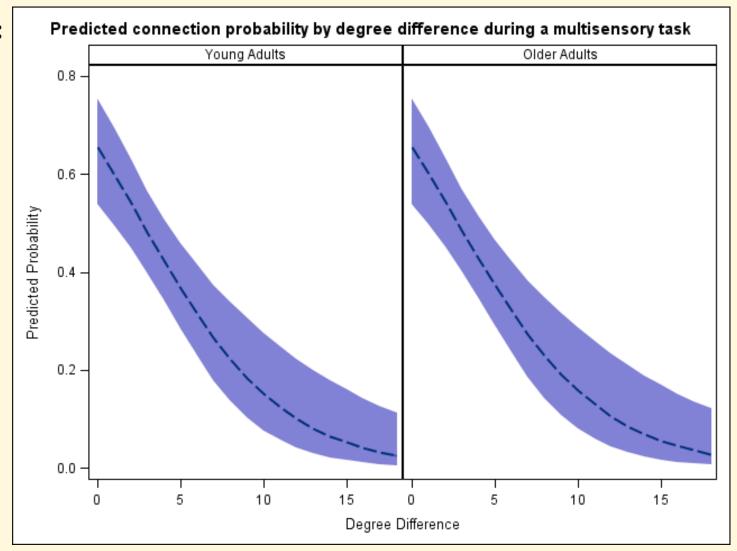
Prediction intervals for connection probability as a function of degree difference in young and older participants at rest.

Predict:



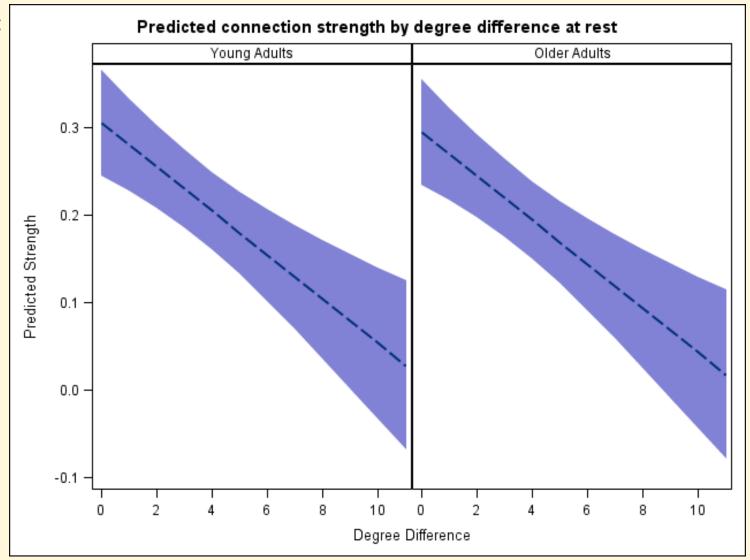
Prediction intervals for connection probability as a function of degree difference in young and older participants during a visual task.

Predict:



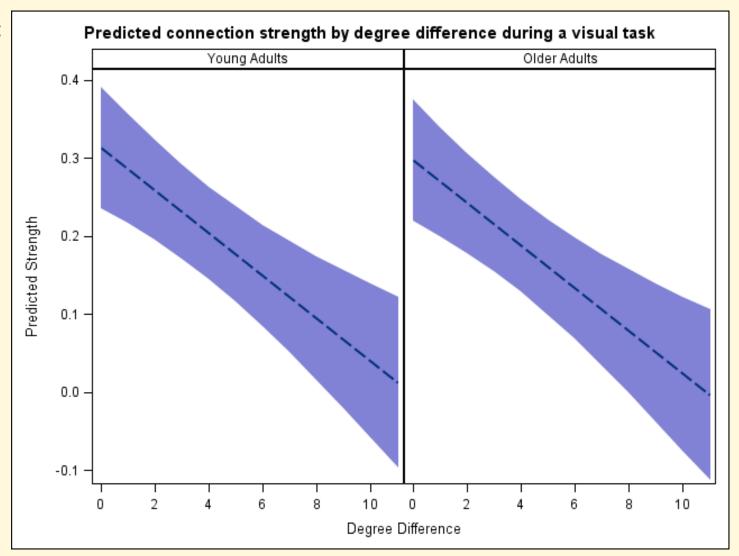
Prediction intervals for connection probability as a function of degree difference in young and older participants during a multisensory task.

Predict:



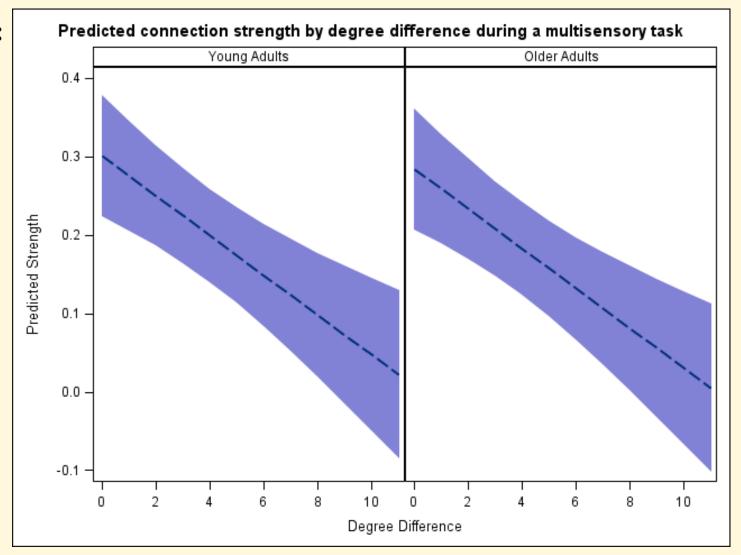
Prediction intervals for connection strength as a function of degree difference in young and older participants at rest.

Predict:



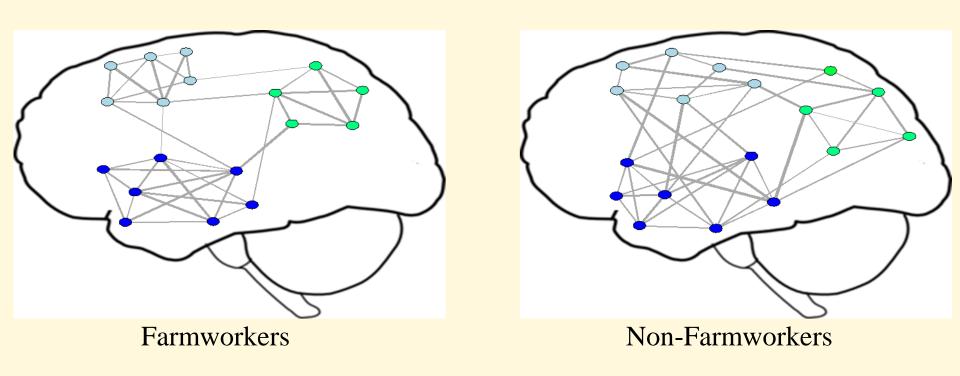
Prediction intervals for connection strength as a function of degree difference in young and older participants during a visual task.

Predict:



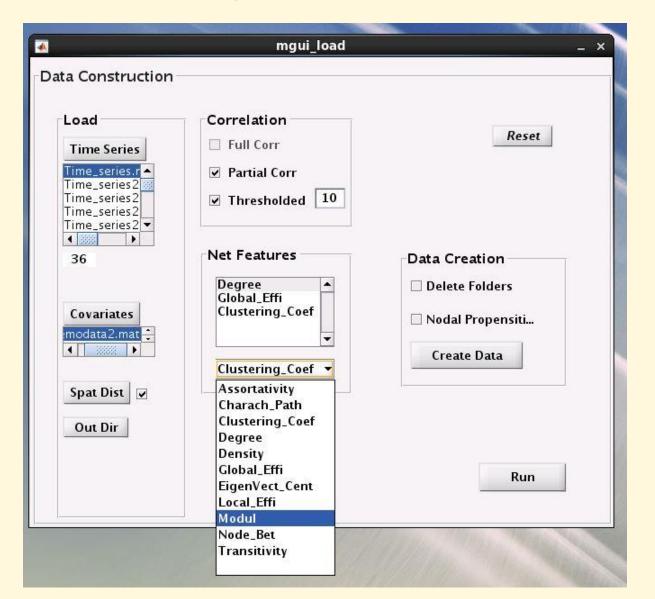
Prediction intervals for connection strength as a function of degree difference in young and older participants during a multisensory task.

• Another example: Used to examine the impacts of pesticide and nicotine exposures on farmworkers' functional brain networks.



• FW: More modularly organized with higher functional specificity and lower inter-modular integrity

Matlab GUI interface coming soon!



ERGMs vs. Mixed Models

- Provide complementary multivariate approaches for analyzing at network level.
 - I.e., assessing systemic infrastructural properties of networks as opposed to properties of specific nodes or connections

ERGMs vs. Mixed Models

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 - I.e., assessing systemic infrastructural properties of network as opposed to properties of specific nodes or connections

ERGMs

- Efficiently represent network data by modeling global structure as function of local substructural (network) properties.
- Not well-suited for examining specific connections, comparing groups, or assessing network-phenotype relationships.

ERGMs vs. Mixed Models

Mixed Models

- Well-suited for examining specific connections, group comparisons, and network-phenotype relationship assessment.
- Limited in ability to capture inherent complex dependence structure of networks.
 - Simpson and Laurienti (2015) adapt to brain network context and account for dependence structure.

ERGMs vs. Mixed Models

Mixed Models

- Well-suited for examining specific connections, group comparisons, and network-phenotype relationship assessment.
- Limited in ability to capture inherent complex dependence structure of networks.
 - Simpson & Laurienti (2015) adapt to brain network context and account for dependence structure.
- Rudimentary connectivity/network hybrid method (Simpson & Laurienti, 2016).
- May provide machinery to develop needed advanced hybrid methods.
- Will at least be beneficial in joint network/connectivity analyses in conjunction with an appropriate connectivity method.

ACKNOWLEDGEMENTS

Collaborators

Paul J. Laurienti

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Mohsen Bahrami

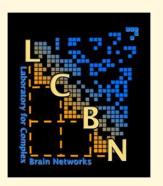
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<u>Others</u>

Members of the Laboratory for Complex Brain Networks



VI. Useful References

ERGMs

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Mixed Models

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