

COMMUNITY DETECTION IN BRAIN FUNCTIONAL NETWORKS BEYOND THE RESOLUTION LIMIT

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ANGELO BIFONE

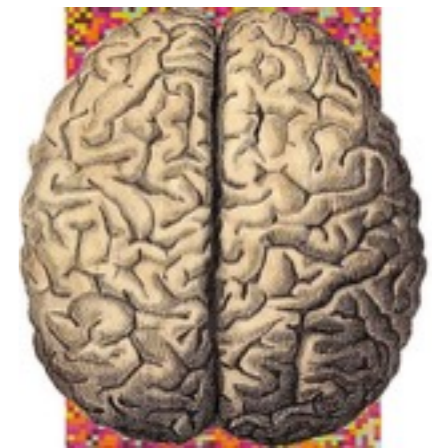
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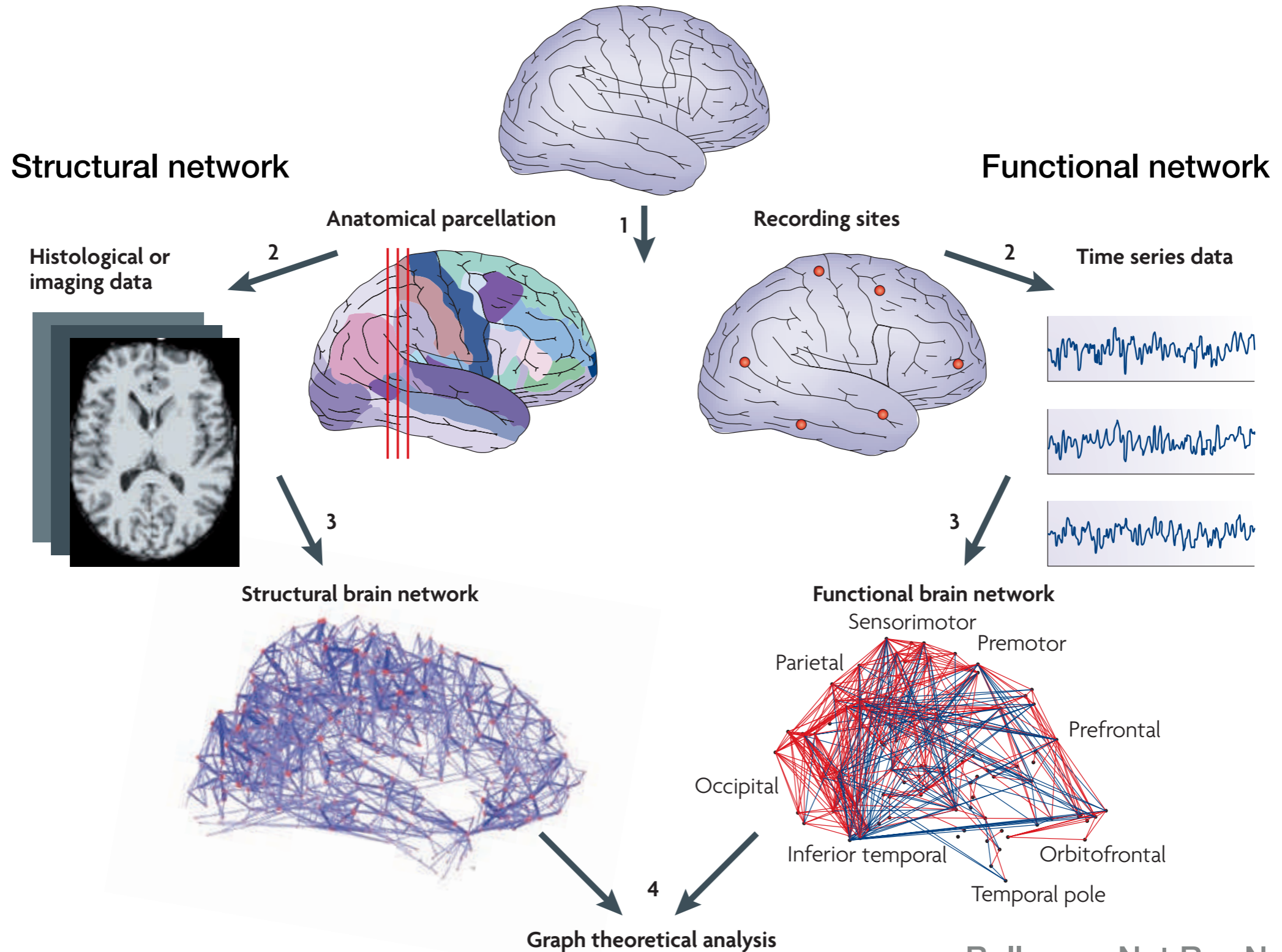
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 [brainetlab.github.io](https://github.com/brainetlab)

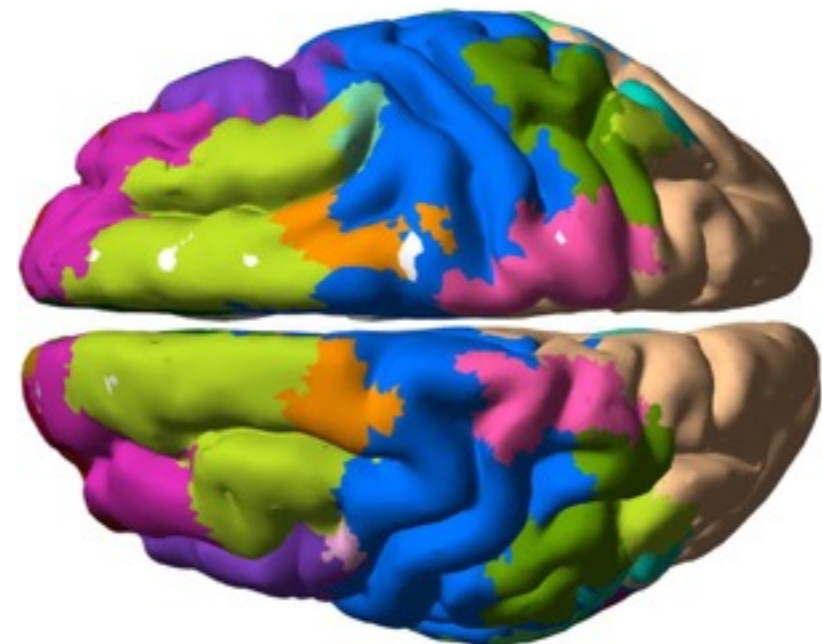
arxiv.org/1609.04316

GRAPH THEORY FOR BRAIN NETWORKS



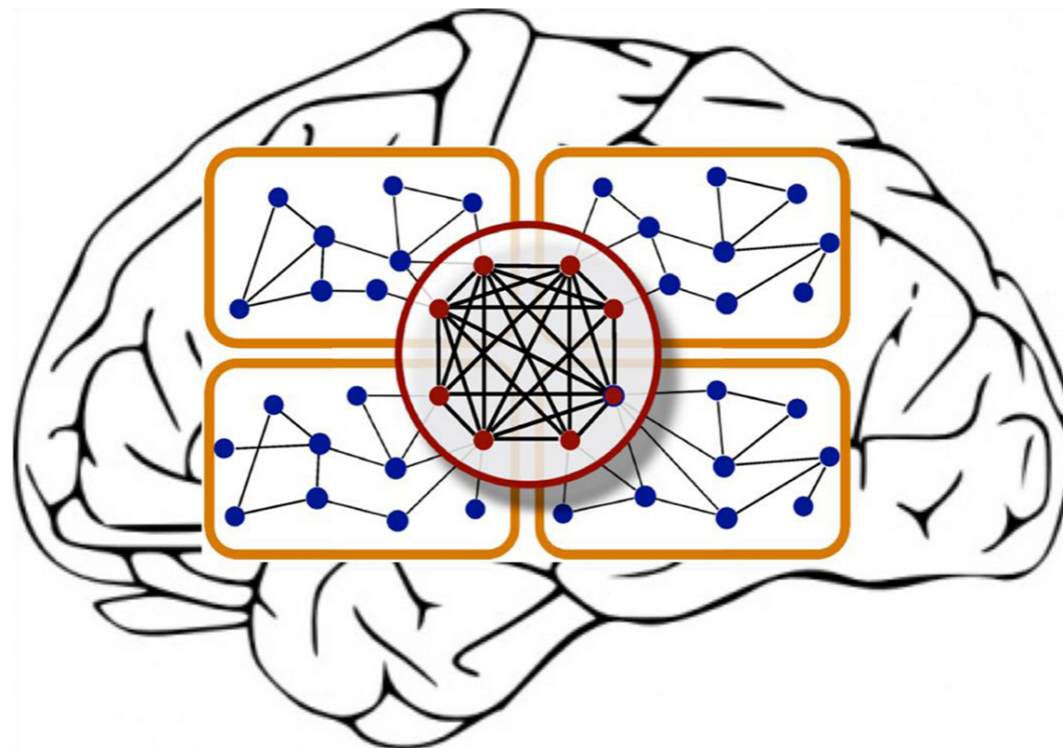
FUNCTIONAL BRAIN NETWORKS

- ▶ A mirror over the living brain.
- ▶ Clinically important biomarker.
- ▶ Aberrant connectivity is observed in many diseases.
- ▶ Modular structure of FC connectivity.
- ▶ Graph theoretical community detection unveils the mesoscopic organization of functional connectivity.



WHY LOOKING FOR MODULES IN THE BRAIN?

- ▶ “Nearly decomposable systems” are faster to adapt and evolve in a changing environment [Simon 1962].
- ▶ Confers stability against abrupt external changes (lesions).
- ▶ Allows for functional segregation and integration.
- ▶ Coevolution of structural and functional connectivity.



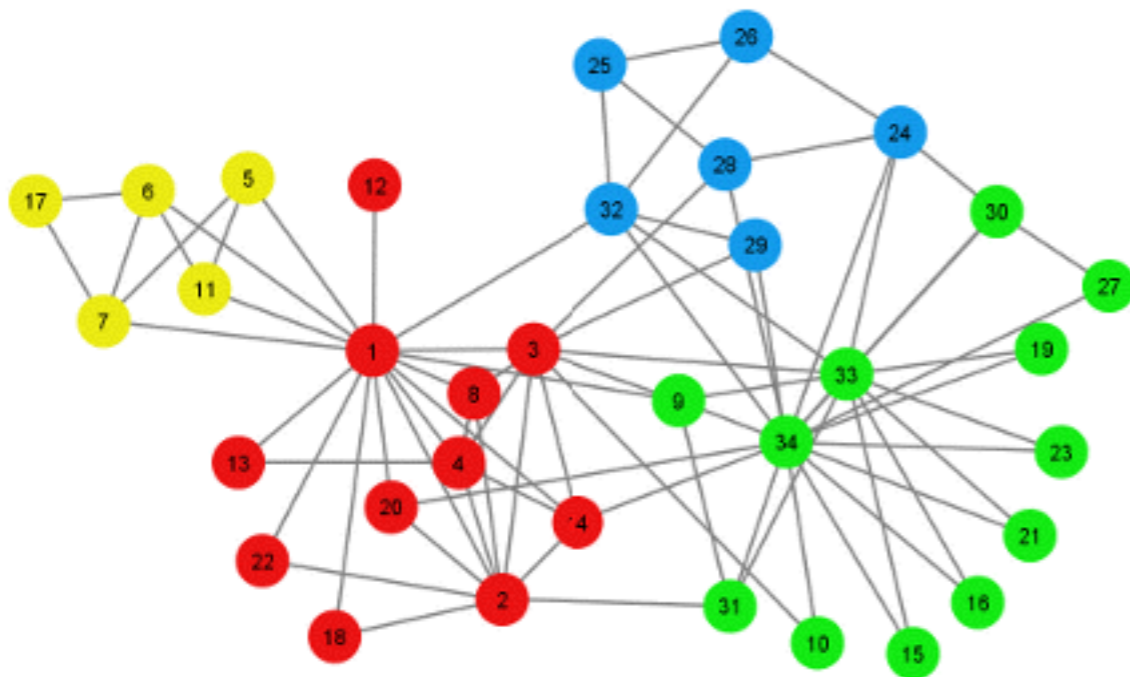
NEWMAN-GIRVAN MODULARITY

$$Q = \frac{1}{2m} \sum_{ij} \left[A_{ij} - \frac{k_i k_j}{2m} \right] \delta(\sigma_i, \sigma_j)$$

1 if node i and node j in the same community

Observed fraction intracluster edges

Expected fraction of intracluster edges



- ▶ Based on a null configuration model
- ▶ Same degree sequence
- ▶ Randomly rewired

Newman, 2006
Zachary, 1977

MOST USED QUALITY FUNCTION FOR COMMUNITY DETECTION

But it has some problems:

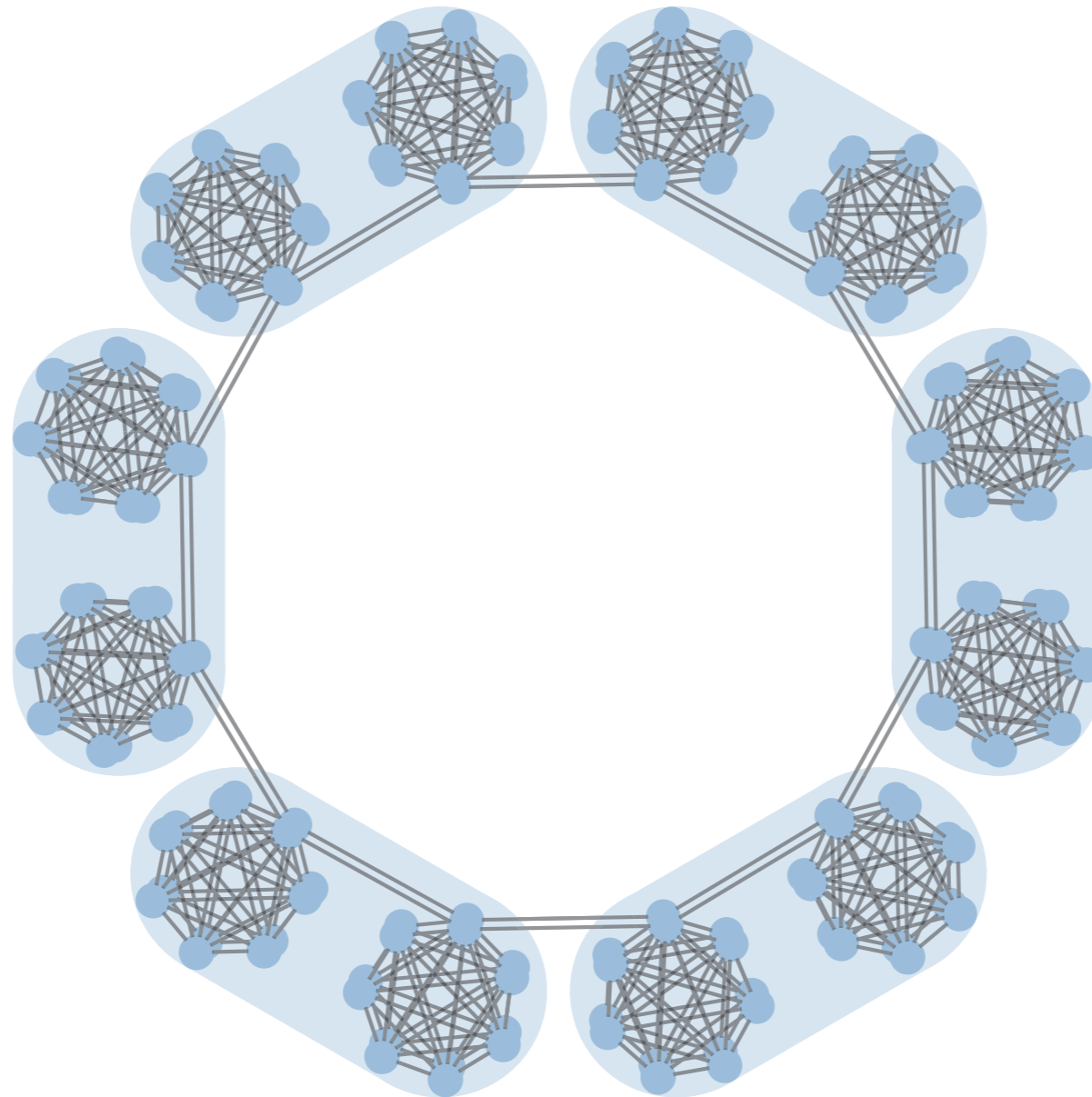
▶ **Resolution limit:**

Inability to detect communities smaller than a certain scale.

▶ **Degeneracy:**

Many high Q solutions are different.

RESOLUTION LIMIT: AN EXAMPLE

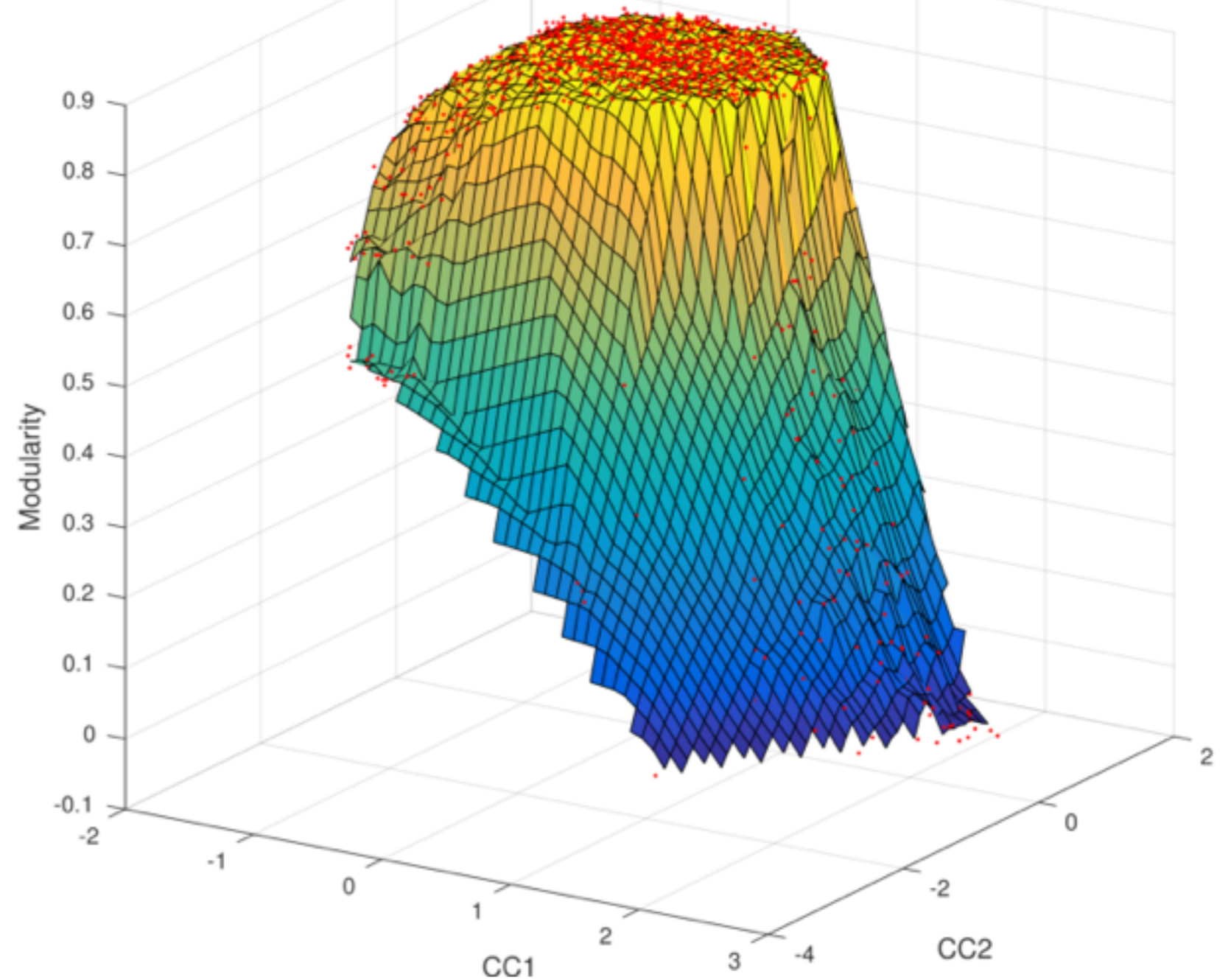
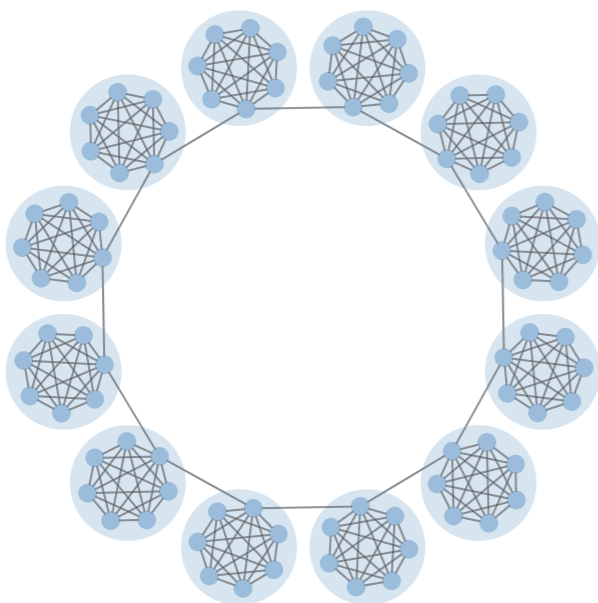


Original
Maximality
partition

Adapted from Traag, 2011

DEGENERACY

- ▶ Degeneracy landscape of a $k=24, n=5$ ring of cliques.
- ▶ Curvilinear components analysis.
- ▶ Red points are solutions.
- ▶ Distance embedding.



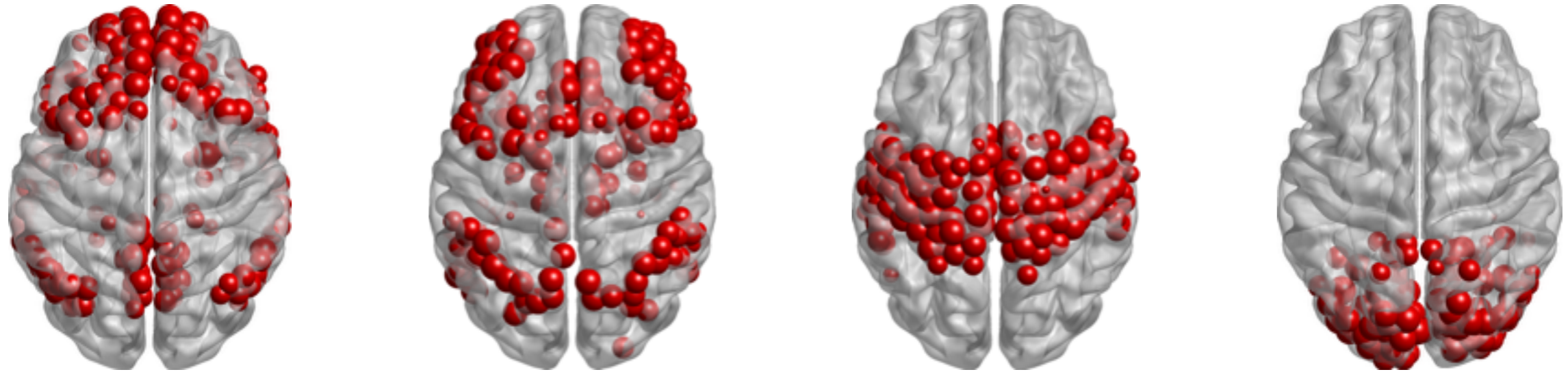
Adapted from Good et al. (2008)

RESOLUTION LIMIT

Resolution limit is an almost ubiquitous phenomenon:

- ▶ Resolution parameter γ [Arenas 2008, Reichardt 2006] only shifts the problem at different scales.
- ▶ It depends on Modularity, not on the heuristic.
- ▶ In Infomap depends on intercluster edges [Kawamoto 2015].
- ▶ Global parameters? Resolution limit kicks in [Fortunato 2016].

REAL WORLD EFFECTS OF RESOLUTION LIMIT



- ▶ Resting state group average over 27 healthy subjects.
- ▶ 4 modules found by modularity maximization.

$$m_c \geq \sqrt{\frac{m}{2}}$$

We need to move this limit away.

SURPRISE

$$S = -\log_{10} \sum_{i=m_\zeta}^m \frac{\binom{p_\zeta}{i} \binom{p-p_\zeta}{m-i}}{\binom{p}{m}}$$

- ▶ p-value of a Fisher exact test based on **urn** model.
- ▶ Measures how surprising is to observe that the intracluster density is the same as graph density.
- ▶ The higher Surprise, the better the clustering.
- ▶ Attention to the statistical significance of the partitioning.

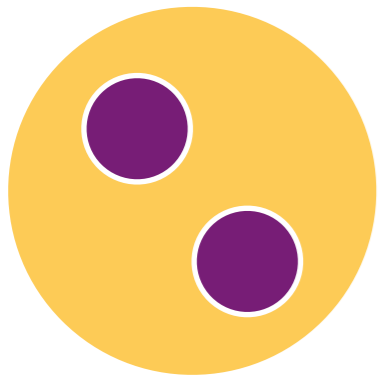
URN MODEL

p total balls, p_ξ yellow and $p - p_\xi$ red.

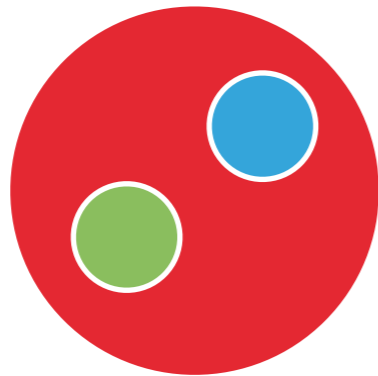
Pick m marbles, randomly, what is the probability of having at least m_ξ yellow balls?

Every marble is a node pair.

Intracluster pairs



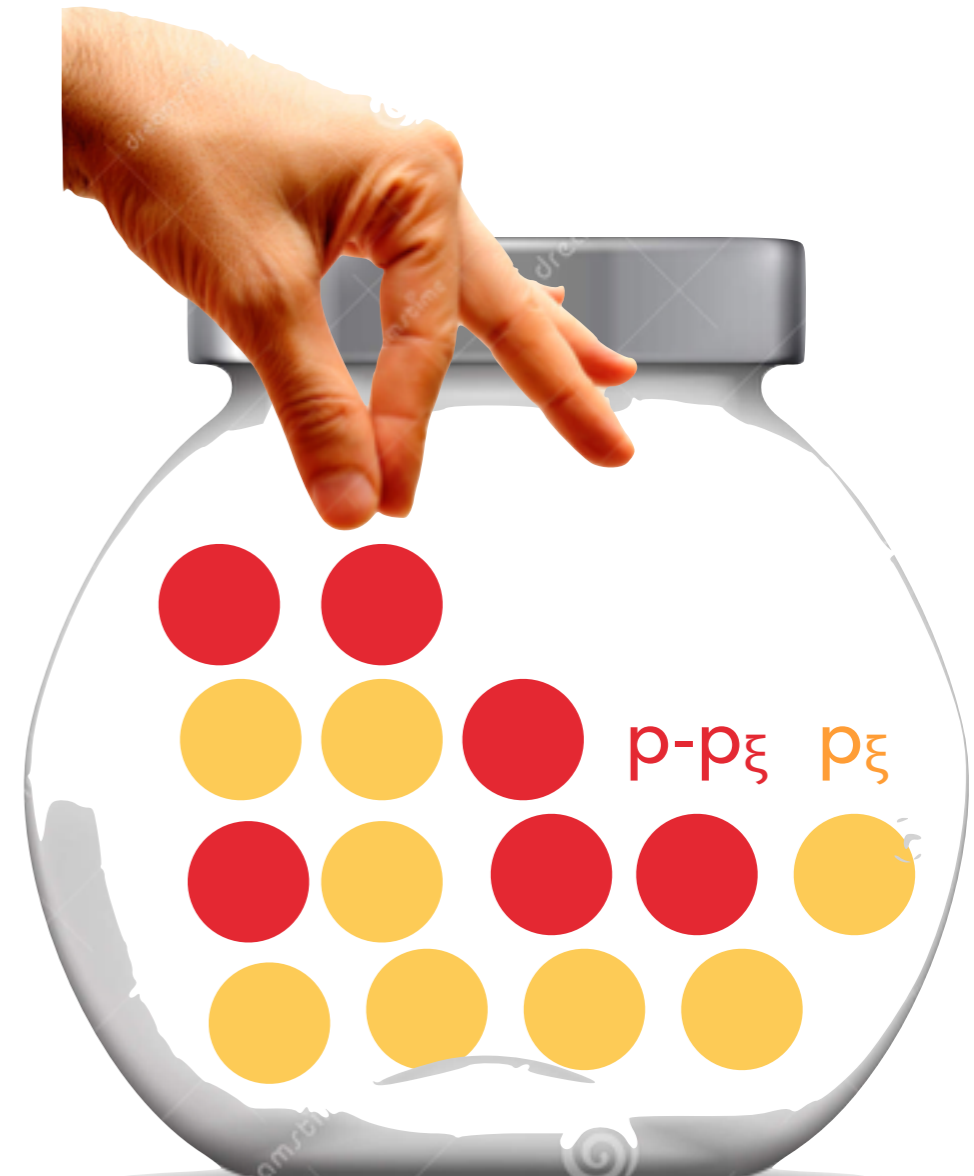
Intercluster pairs



$m - m_\xi$ m_ξ

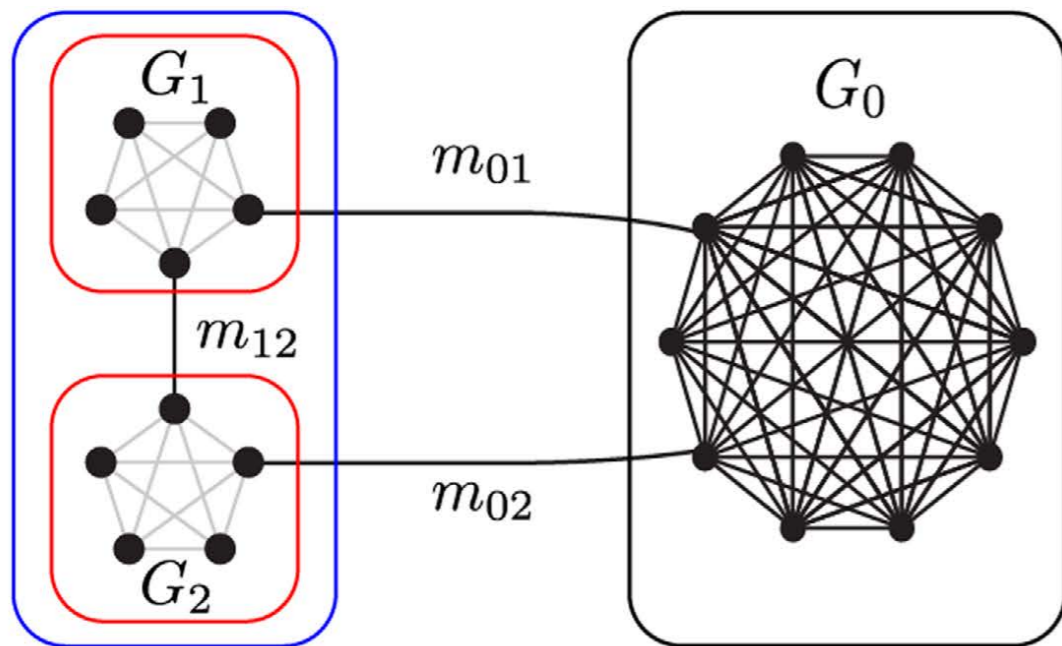


m balls



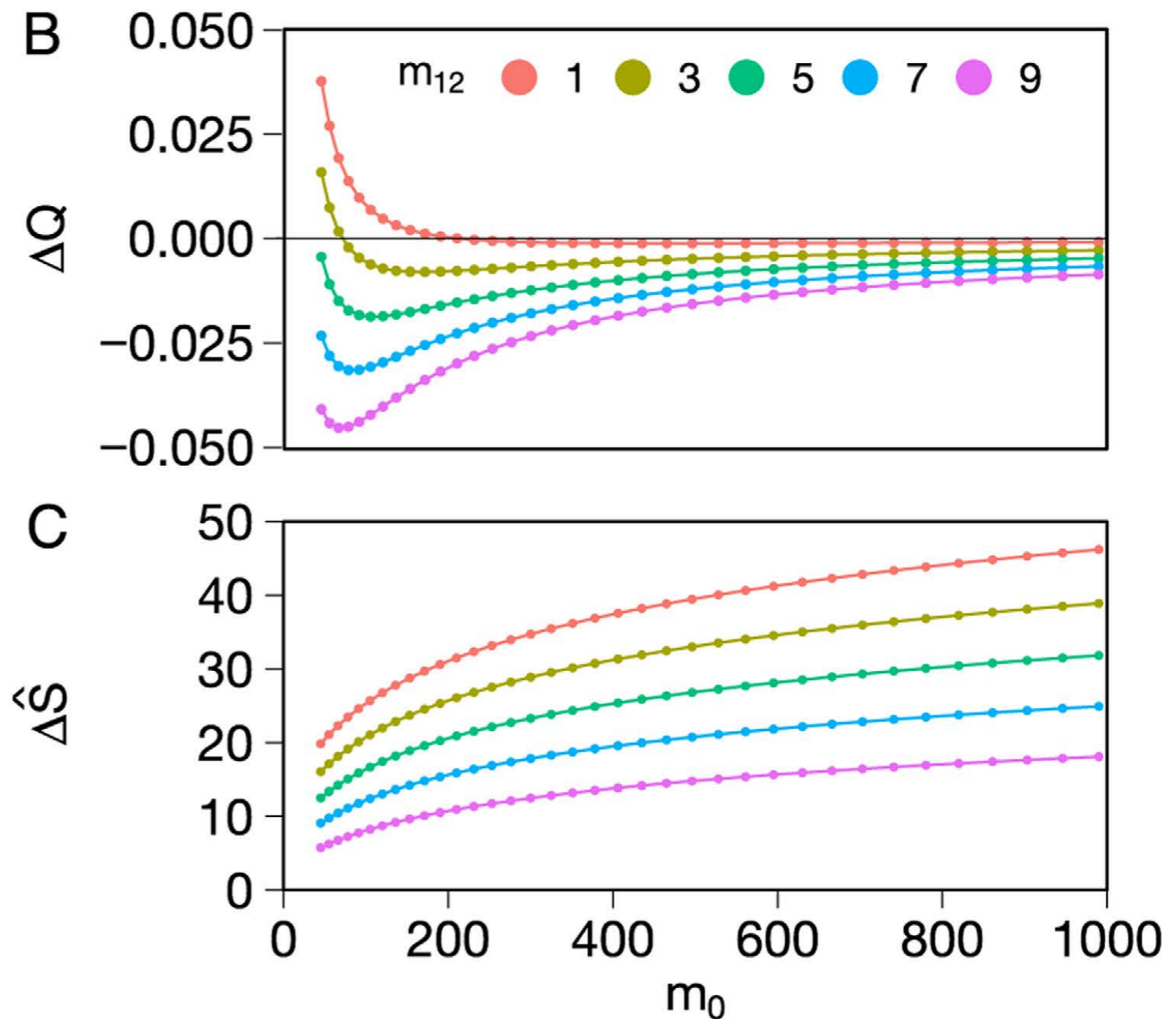
p balls

RESOLUTION LIMIT AND SURPRISE



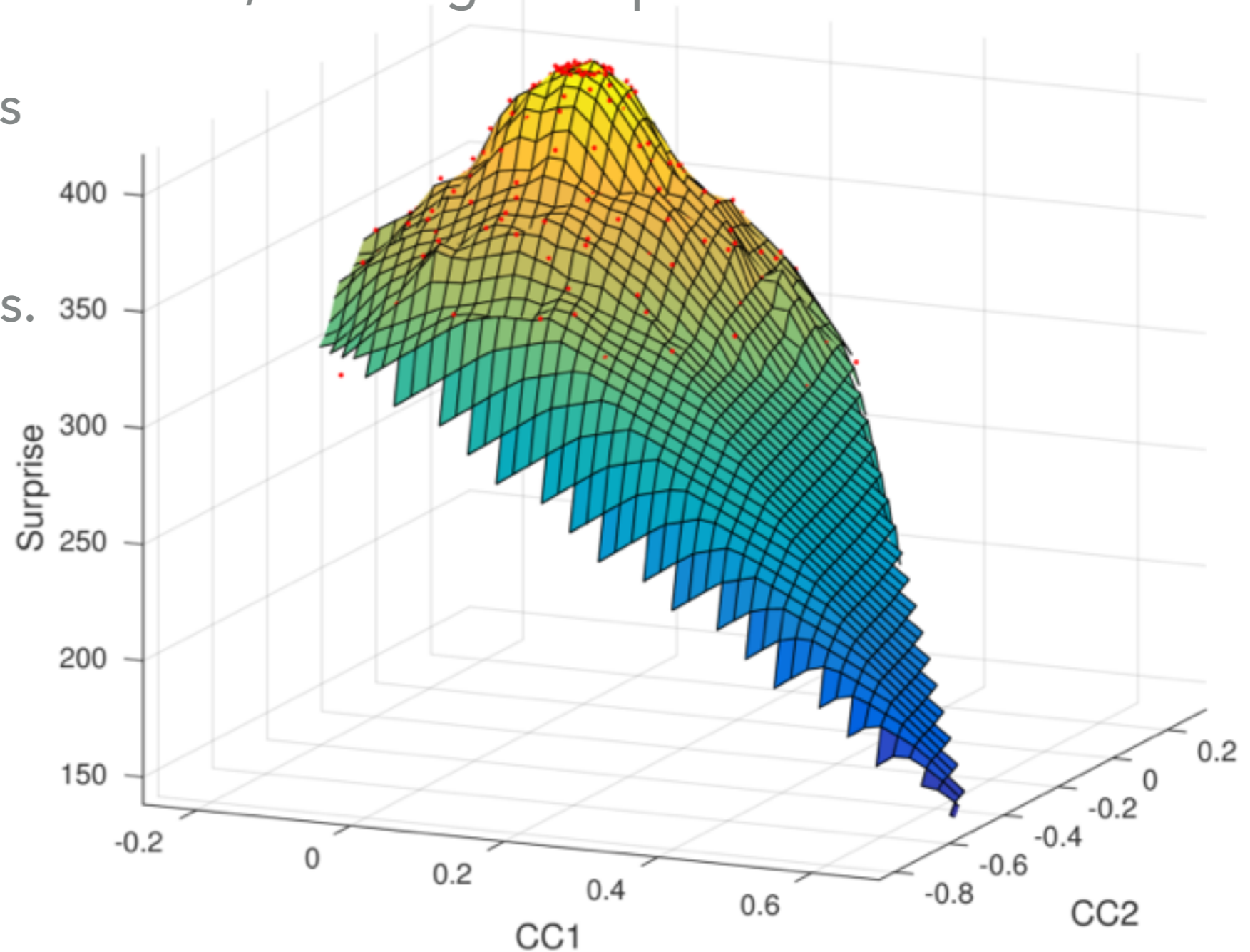
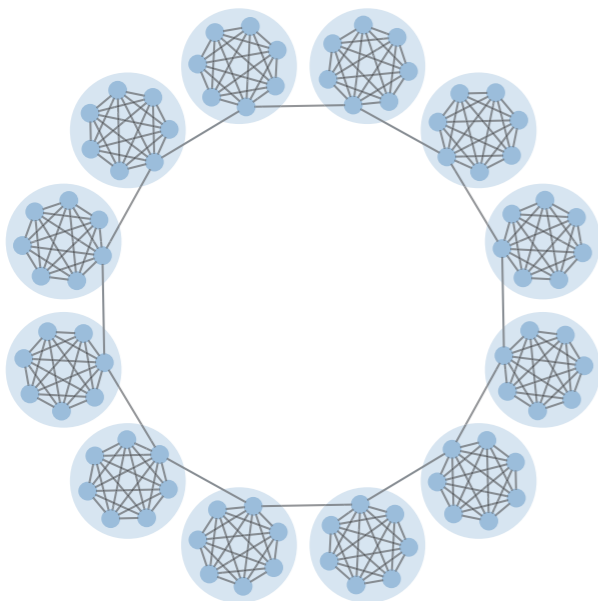
$$\Delta Q = Q_A - Q_B$$

$$\Delta S = S_A - S_B$$



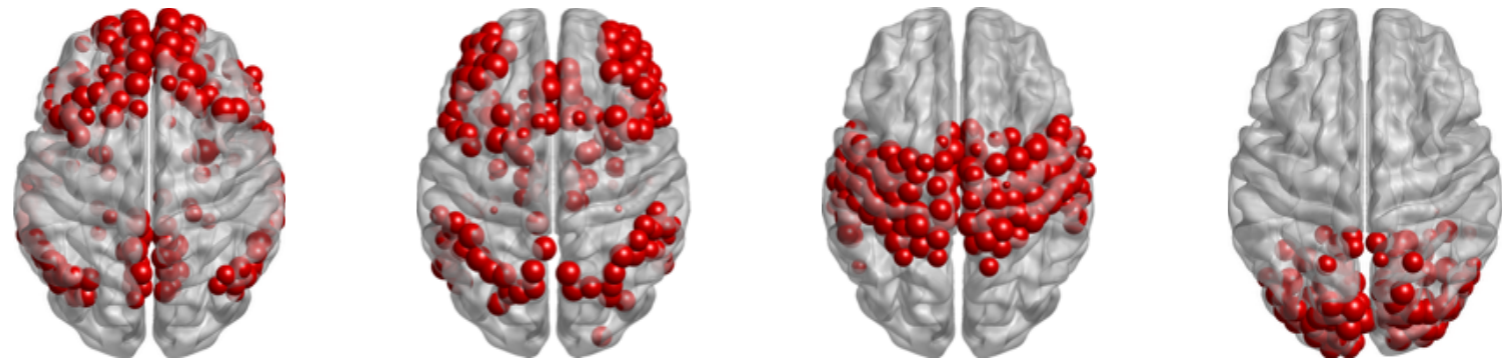
NO DEGENERACY

- ▶ Degeneracy landscape of a $k=24, n=5$ ring of cliques.
- ▶ Curvilinear components analysis.
- ▶ Red points are solutions.
- ▶ Distance embedding.

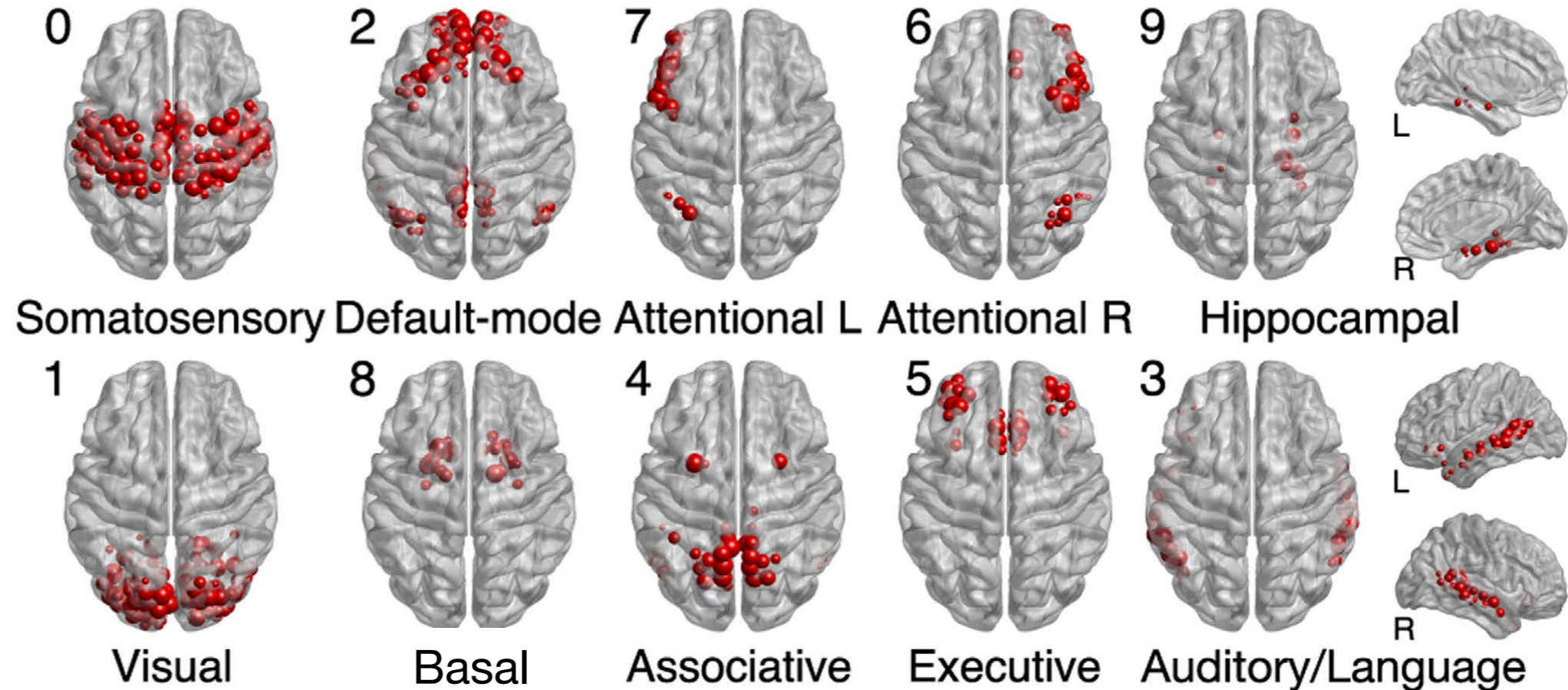


APPLICATION OF SURPRISE OPTIMIZATION

Modularity



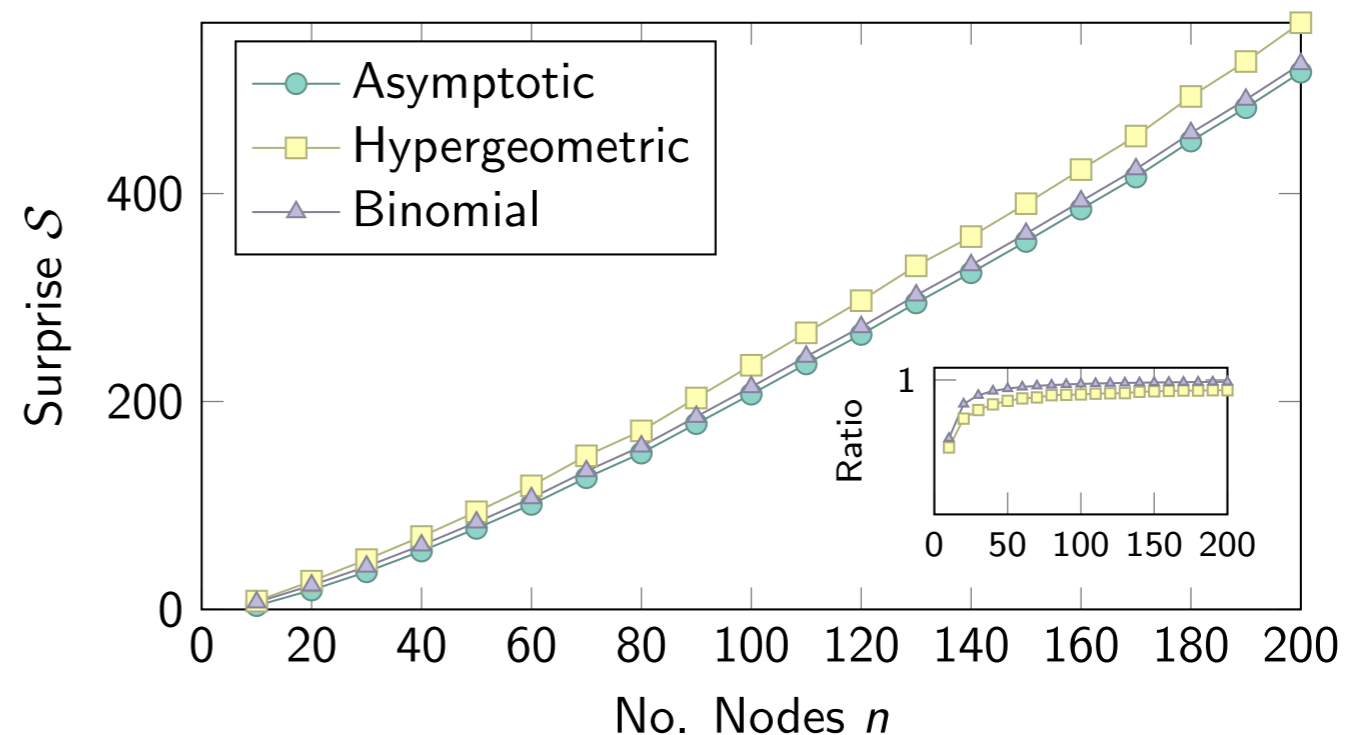
Surprise



ASYMPTOTICAL SURPRISE

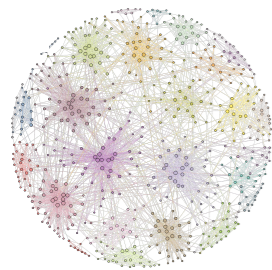
$$S_a = m D_{\text{KL}} \left(\frac{m_\zeta}{m} \parallel \frac{p_\zeta}{p} \right)$$

- ▶ Asymptotical approximation valid for large n .
- ▶ Information theoretic distributions pseudo-distance.
- ▶ Information gained.
- ▶ Supports weighted graphs.



COMPARING ASYMPTOTICAL SURPRISE WITH OTHER METHODS

How to make fair comparison on brain networks if we don't have the brain networks community structure?



**GENERATE A NETWORK
WITH GIVEN COMMUNITY
STRUCTURE**

**COMPUTE THE NEAREST
POSITIVE DEFINITE
MATRIX**

LFR Model

neuroSim R package

Rician distribution

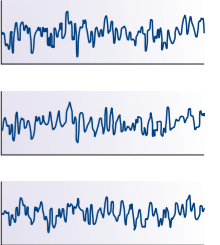
Fisher Transformation

Surprise, Infomap, Modularity

NMI

GRAPH CREATION

**SIMULATE RS BOLD
SIGNALS FOR MANY
VIRTUAL SUBJECTS**



**INJECT CORRELATION
INTO SYNTHETIC TIME
SERIES**

**ADD REALISTIC NOISE TO
TIME SERIES**

**GET PARTITION
SIMILARITY WITH THE
PLANTED ONE**

**RUN COMMUNITY
DETECTION TO ASSESS
EFFECTS OF NOISE**

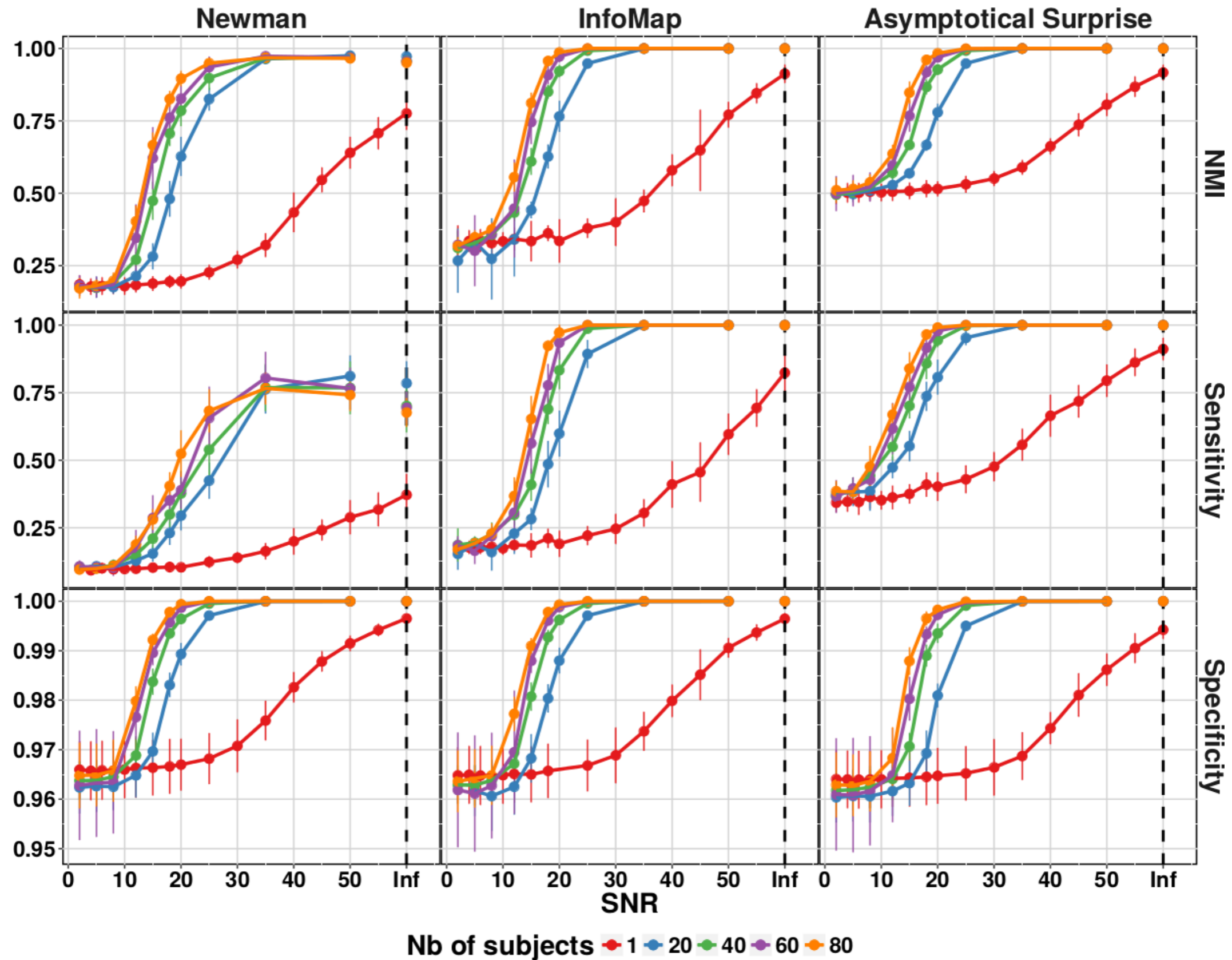
COMPARING COMMUNITY DETECTION ON BRAIN NETWORKS

- ▶ Varied $SNR = \langle S \rangle / \sigma_n$ and number of subjects.
- ▶ Normalized Mutual Information (NMI)
- ▶ Matrix C_{ij} is the number of nodes in the planted community- i appearing in the detected community- j .
- ▶ Sensitivity (Recall) = $TP / (TP + FN)$
- ▶ Specificity = $TN / (TN + FP)$

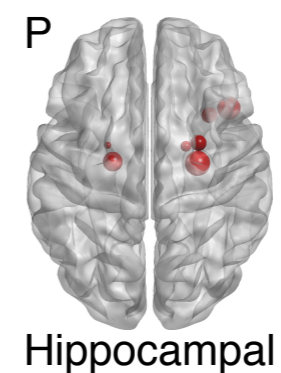
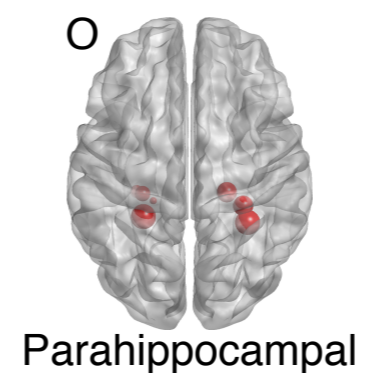
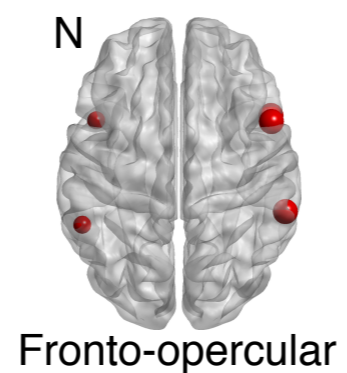
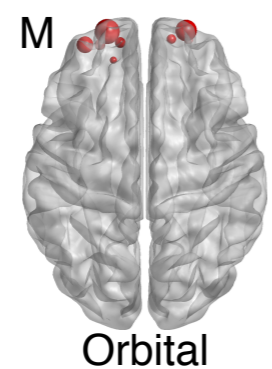
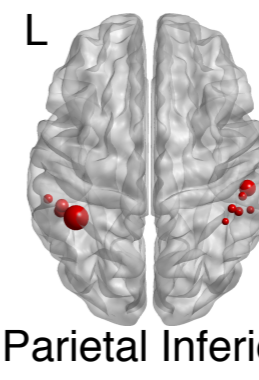
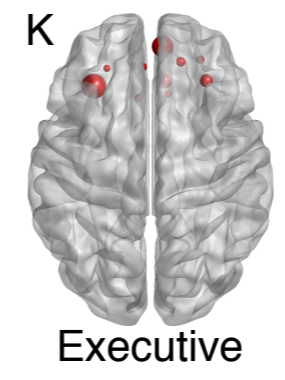
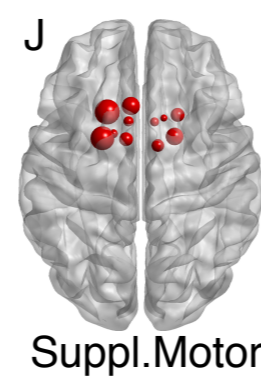
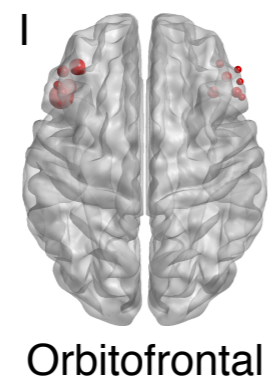
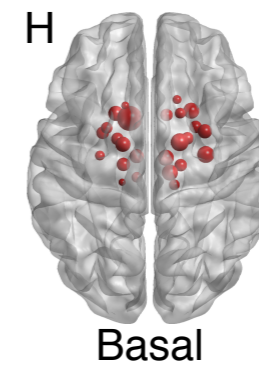
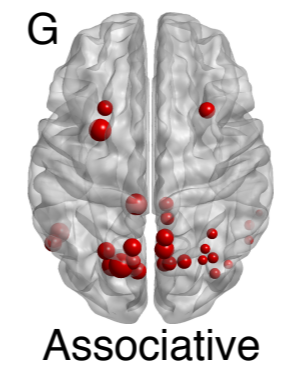
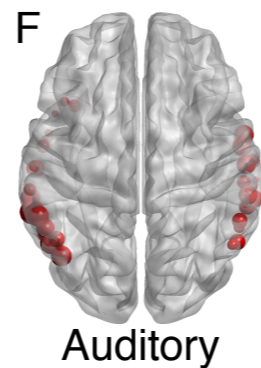
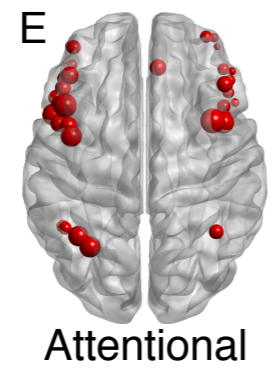
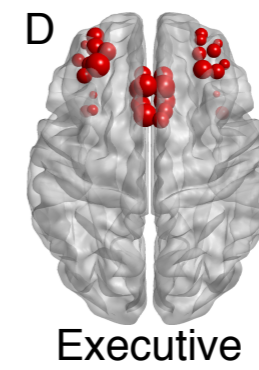
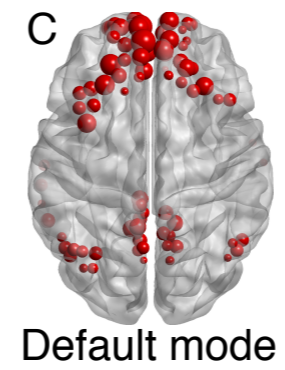
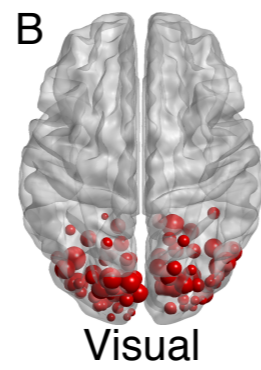
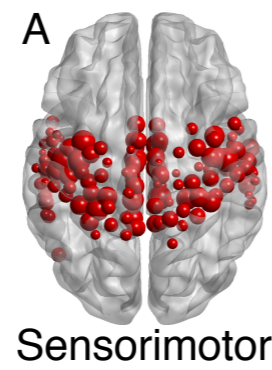
TP	FN
FP	TN

		Planted communities		
		1	2	3
Detected communities	A	4	2	
	B		1	1
	C			2

RESULTS



HUMAN RESTING STATE DATE



CONCLUSIONS

- ▶ Functional connectivity can be studied with graph-theoretical approaches.
- ▶ Resolution limit hindered detection of functional modules.
- ▶ Coarse resolution hides small details and differences between groups.
- ▶ Asymptotical Surprise can identify neurofunctionally plausible and anatomically well-defined substructures.

But ...

- ▶ It may overfit the community structure due to its improved sensitivity.

THANK YOU!



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github.com/carlonicolini

brainetlab.github.io



Cecile Bordier

REFERENCES

Nicolini C., Bifone A. Scientific Reports 6, 19250, (2016)

Nicolini C., Bordier C., Bifone A. Arvix 1609.04316 arxiv.org/1609.04316 