

Network inference from time-series measurements

Nicolás Rubido^{1,2}, Arturo C. Martí², Ezequiel Bianco-Martínez¹, Celso Grebogi¹, Murilo S. Baptista¹, and Cristina Masoller³

1 – Institute for Complex Systems and Mathematical Biology, University of Aberdeen, UK.

2 – Instituto de Física, Facultad de Ciencias, Universidad de la República, Uruguay

3 – Departament de Física i Enginyeria Nuclear, Universitat Politècnica de Catalunya, Colom 11, E-08222 Terrassa, Barcelona, Spain.



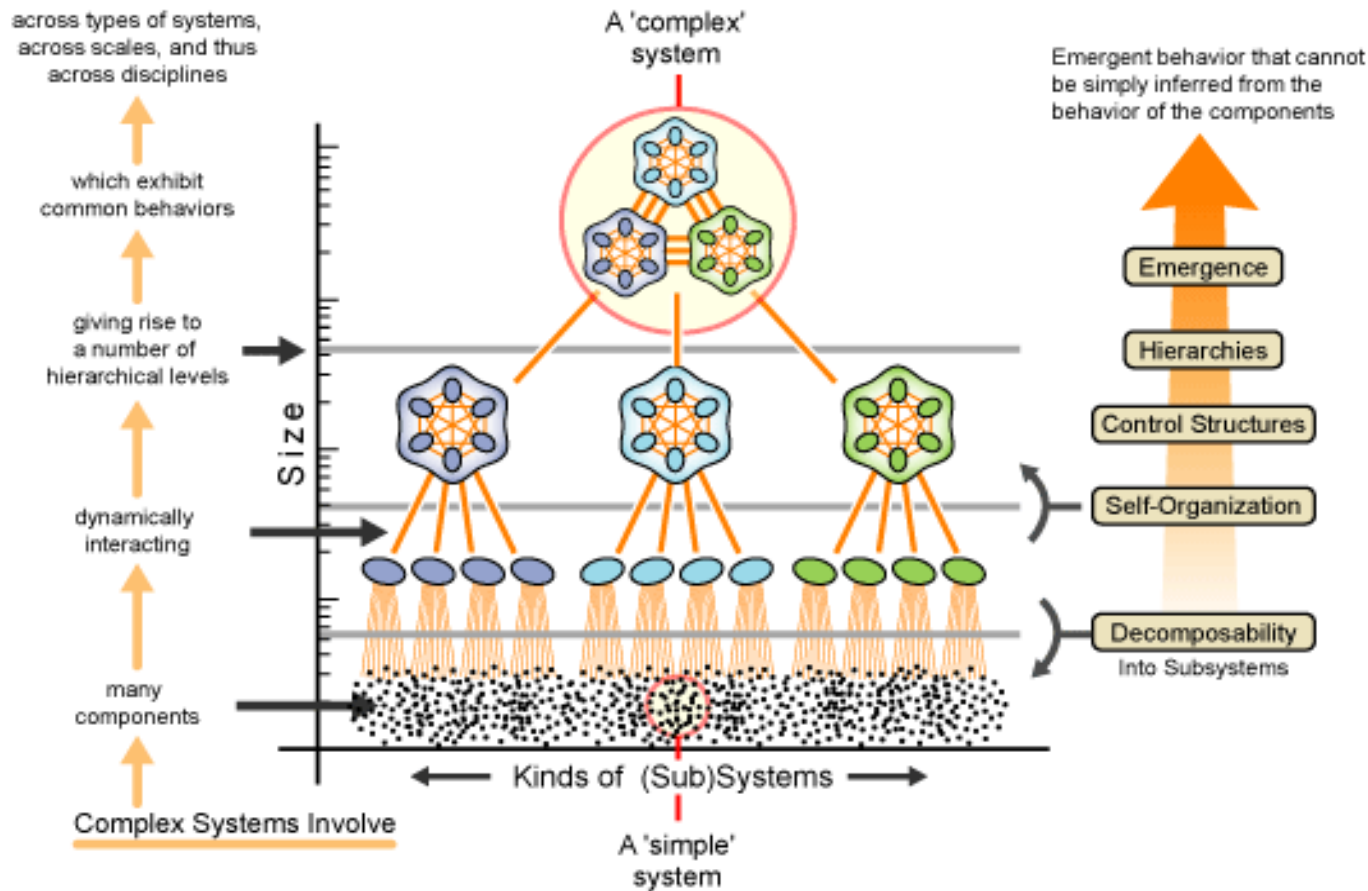
Causality,
Information Transfer and
Dynamical Networks

References:

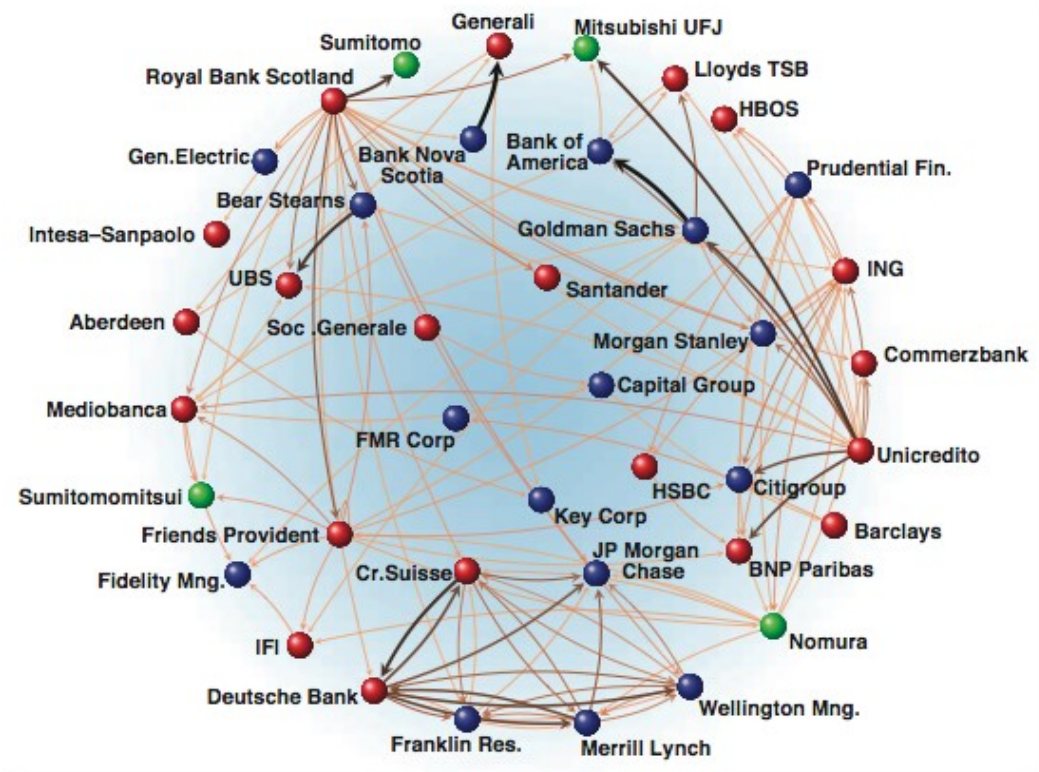
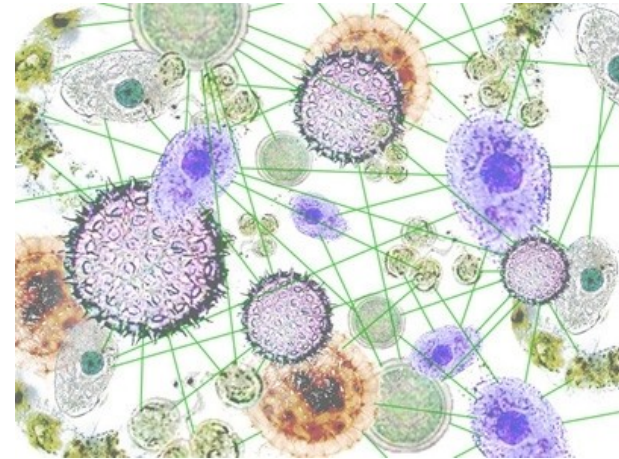
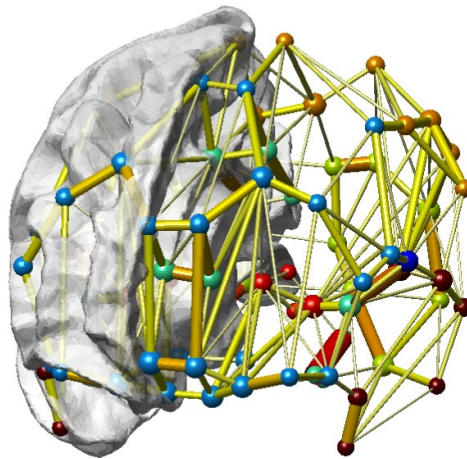
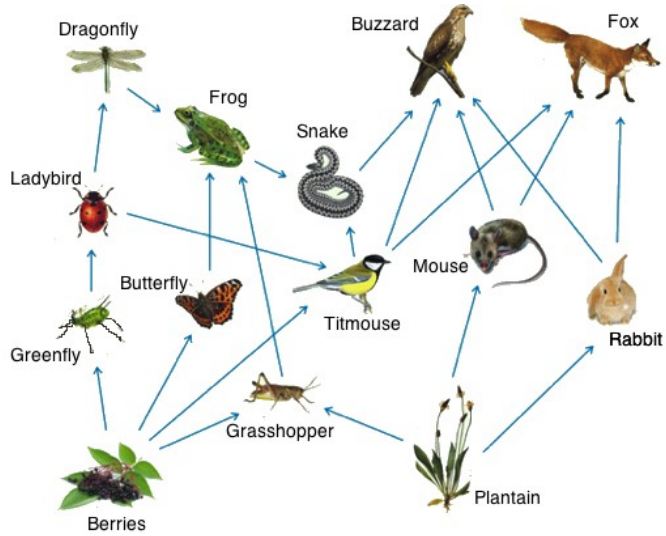
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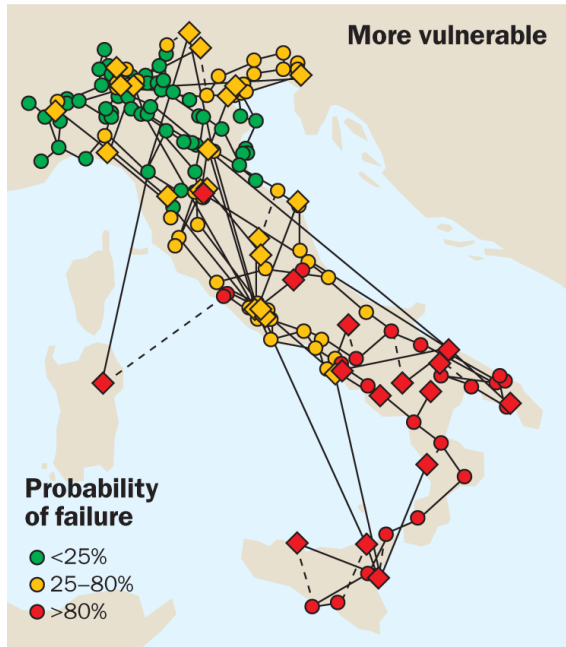
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Complex Systems

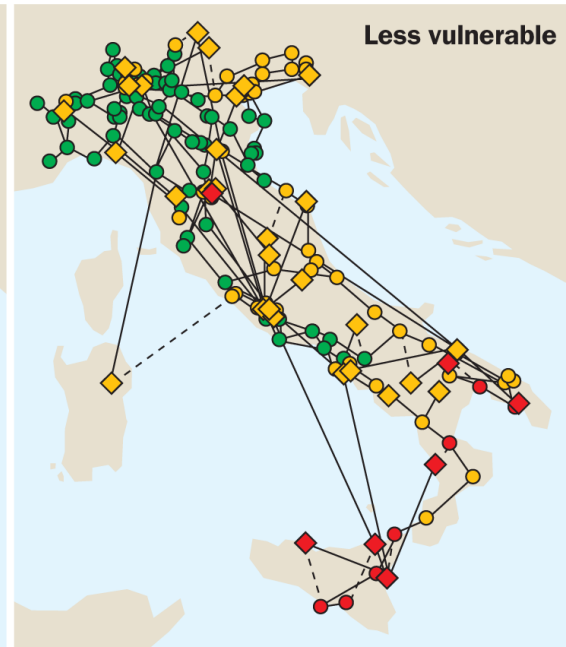


Networks

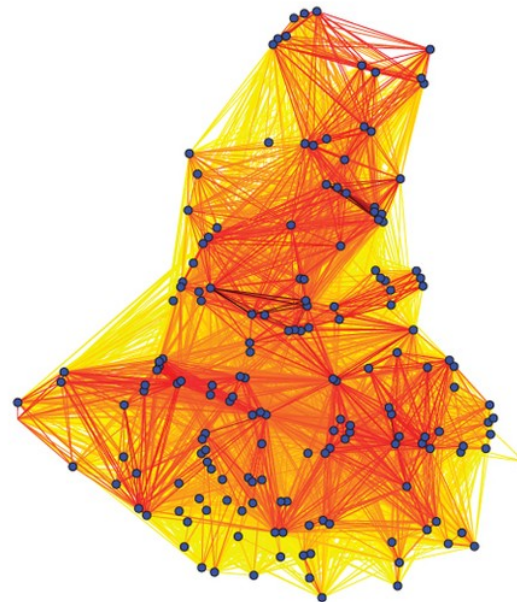
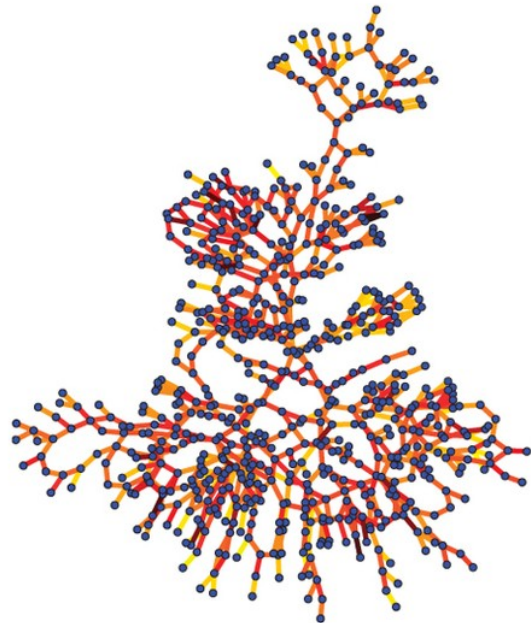




a



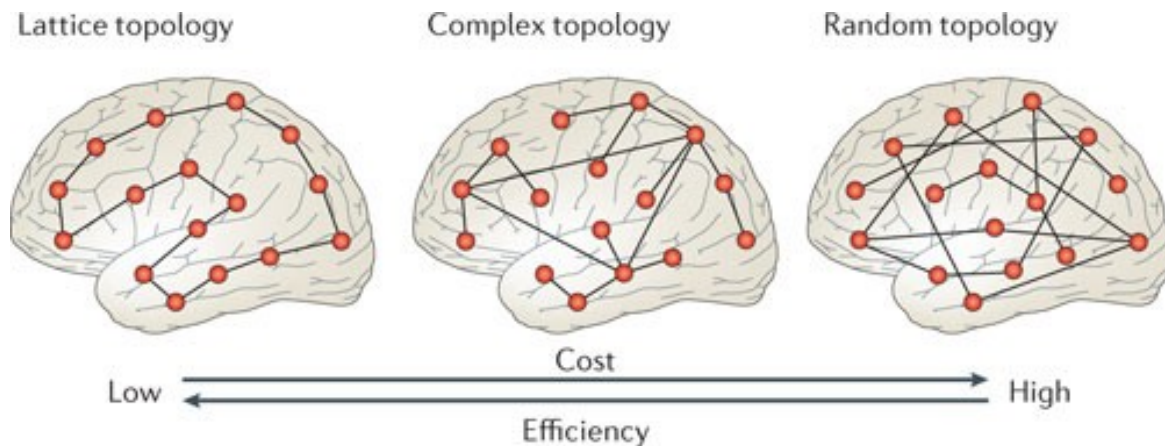
b



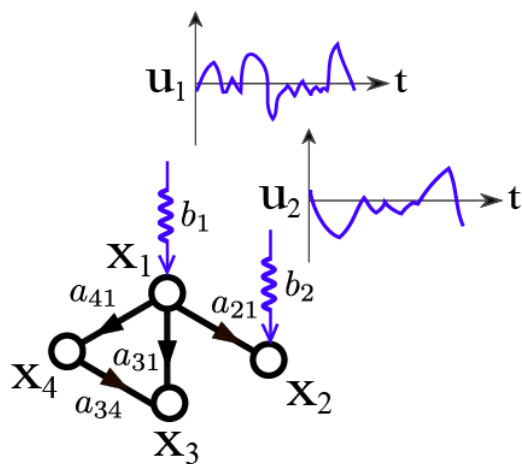
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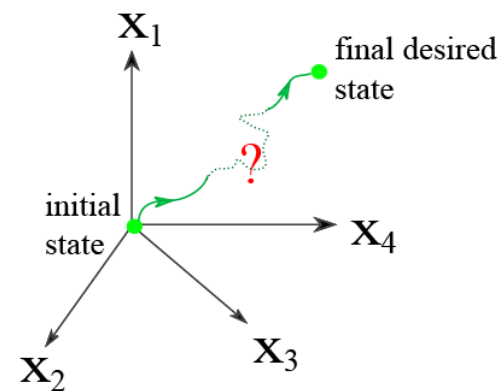
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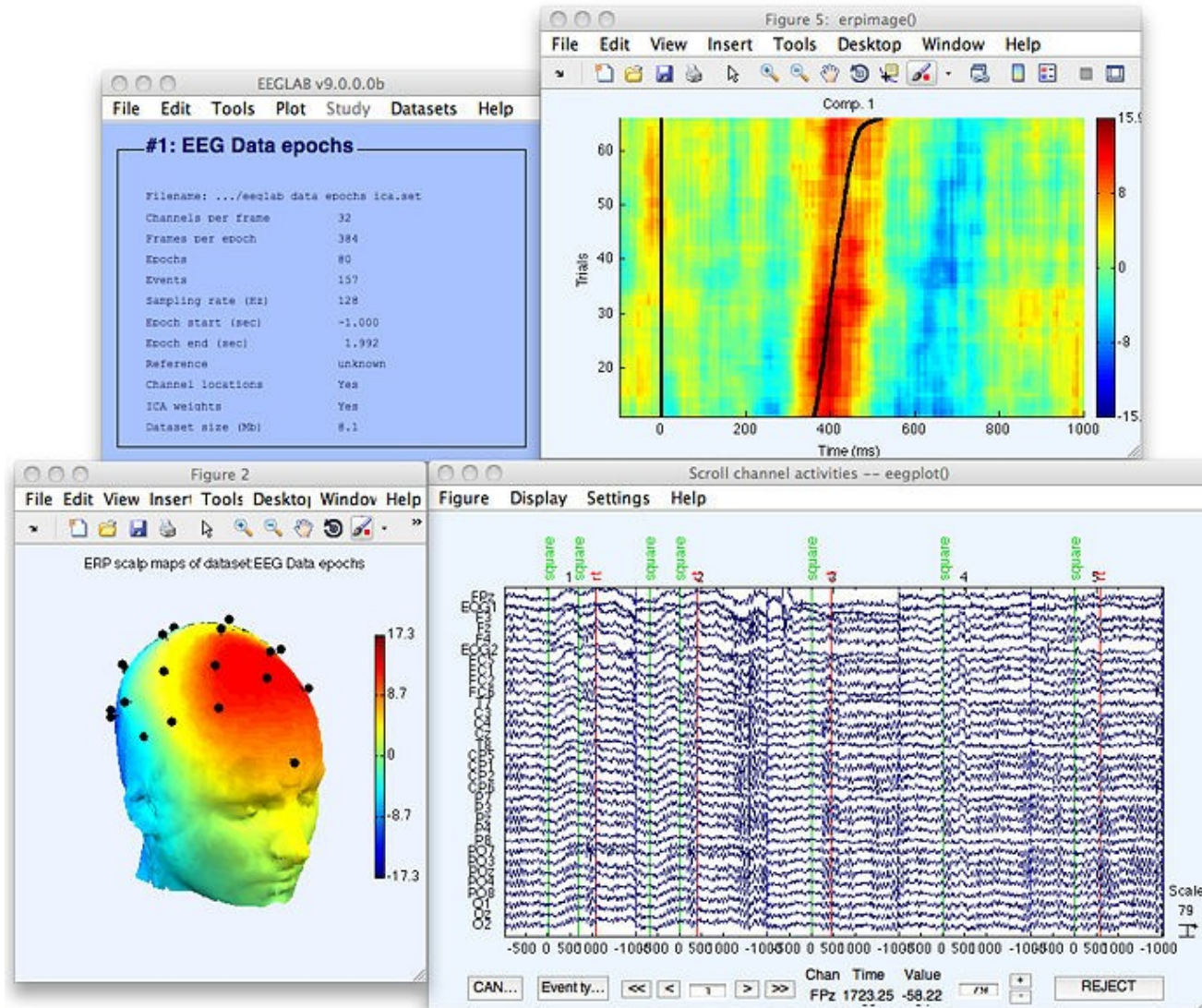
Nature Reviews | Neuroscience

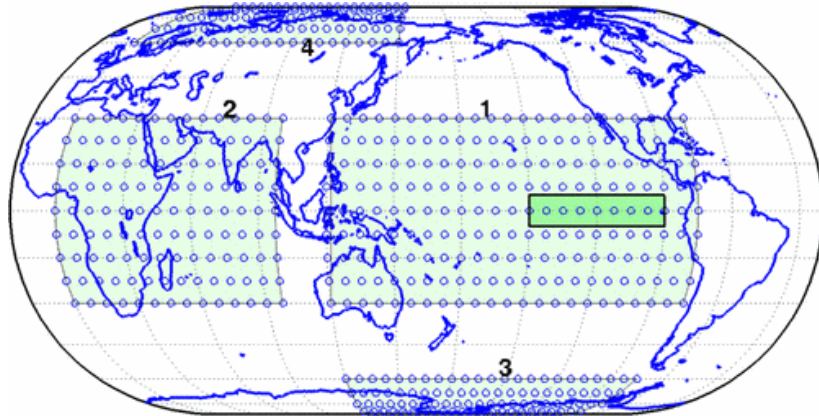


$$\mathbf{A} = \begin{pmatrix} 0 & 0 & 0 & 0 \\ a_{21} & 0 & 0 & 0 \\ a_{31} & 0 & 0 & a_{34} \\ a_{41} & 0 & 0 & 0 \end{pmatrix}; \quad \mathbf{B} = \begin{pmatrix} b_1 & 0 \\ 0 & b_2 \\ 0 & 0 \\ 0 & 0 \end{pmatrix}; \quad \mathbf{C} = \begin{pmatrix} b_1 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & b_2 & a_{21}b_1 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & a_{31}b_1 & 0 & a_{34}a_{41}b_1 & 0 & 0 & 0 \\ 0 & 0 & a_{41}b_1 & 0 & 0 & 0 & 0 & 0 \end{pmatrix}$$

$$N = 4, \quad M = 2, \quad \text{rank}(\mathbf{C}) = 4 = N$$

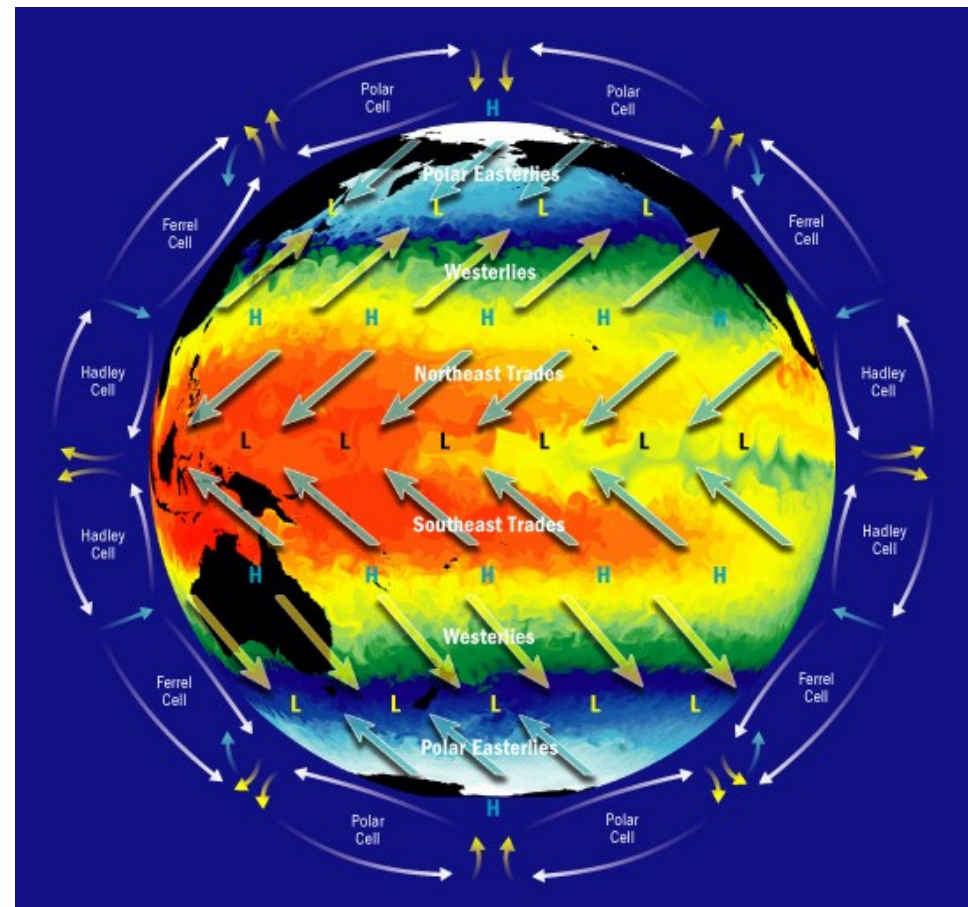
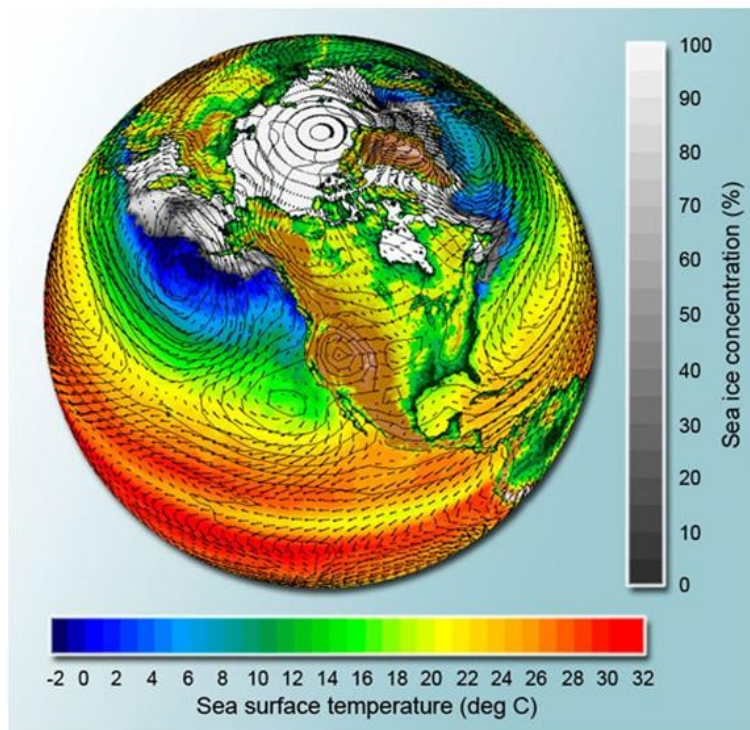
Time-series measurements



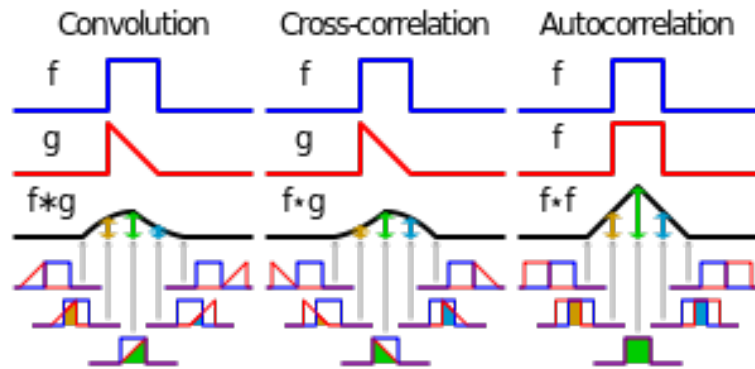


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"The backbone of the climate network",
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C. Tominski, J.F. Donges, and T. Nocke,
"Information Visualization in Climate Research",
IEEE 15th Int. Conf. Inf. Vis. **4**, 298-305 (2011).

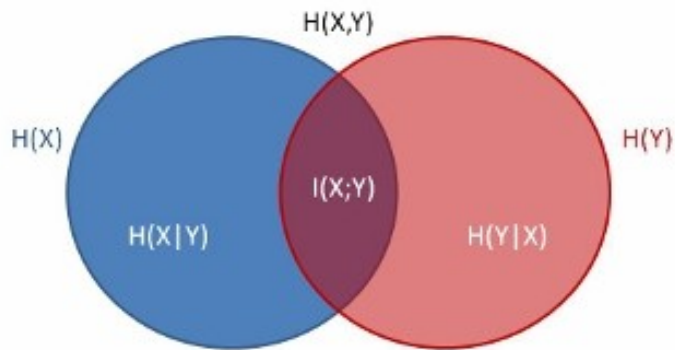


Similarity measures



Cross-Correlation

Mutual Information



$$I(X;Y) = H(X \cap Y)$$

$$H(X;Y) = H(X) + H(Y) - H(X \cap Y)$$

Large networks



Unified Causal Model

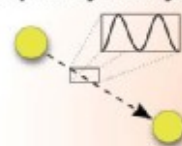


vs Bayesian Networks

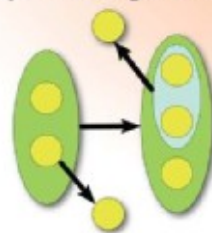


Granger Causality

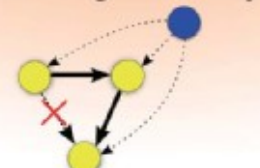
Frequency Analysis



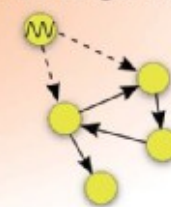
Complex Granger Causality

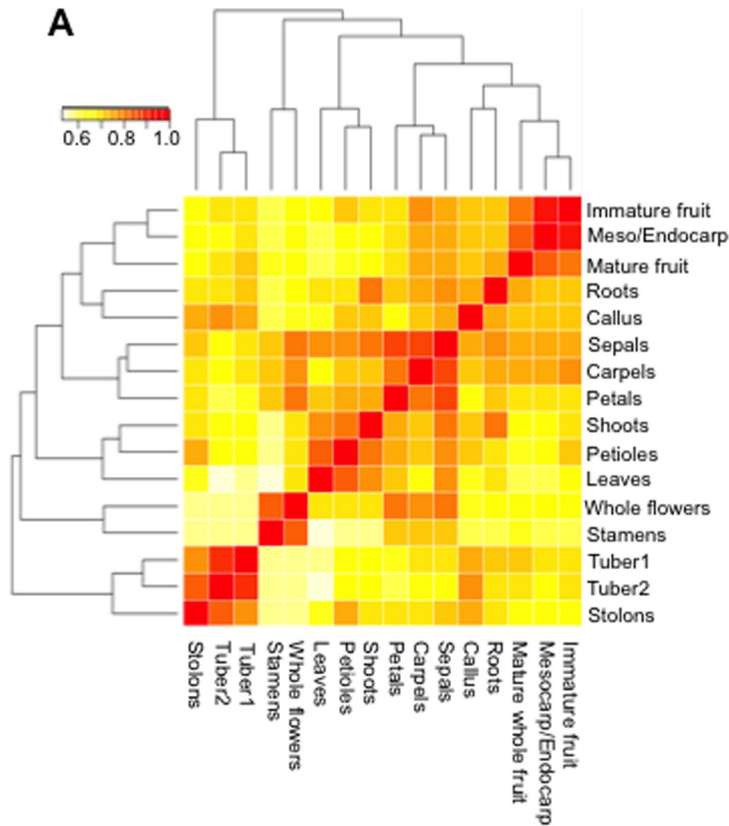


Partial Granger Causality



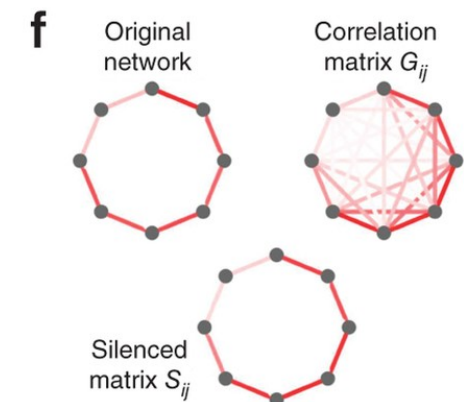
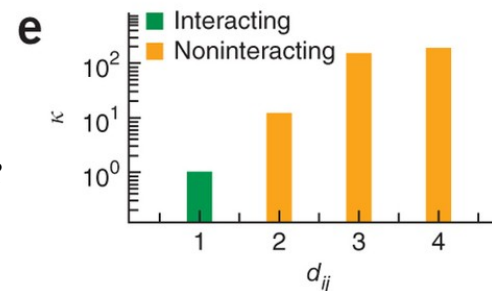
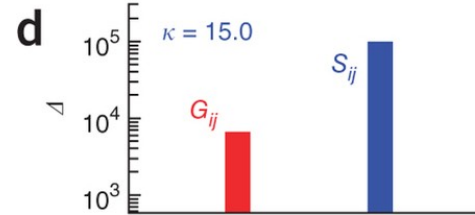
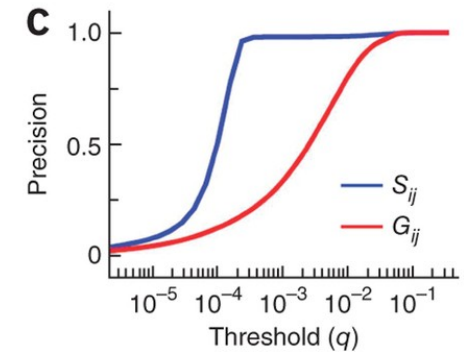
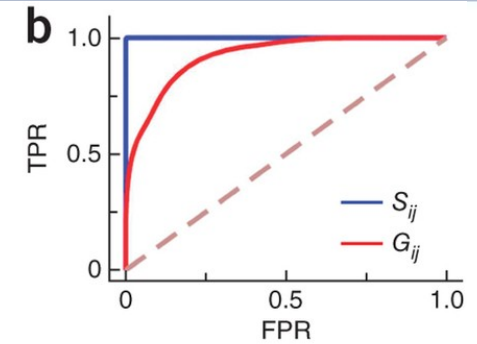
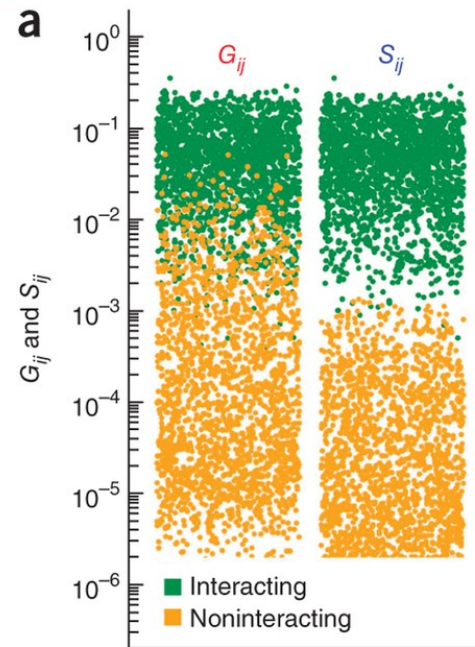
Harmonic Granger Causality



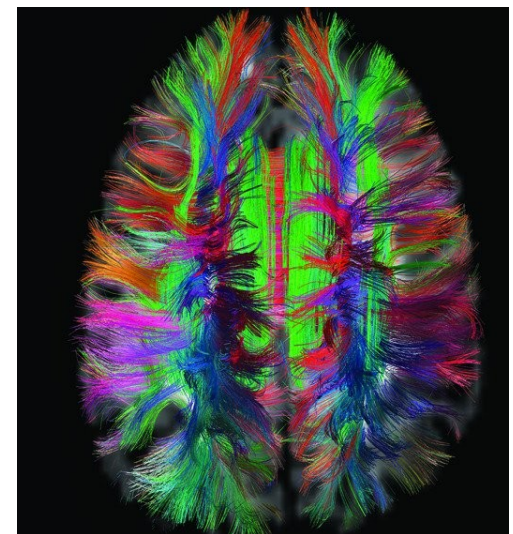
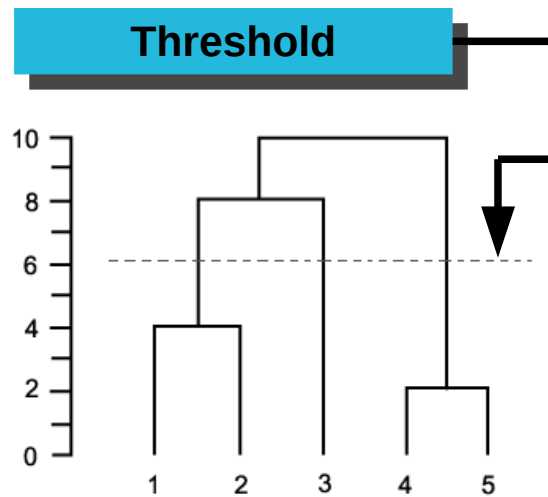
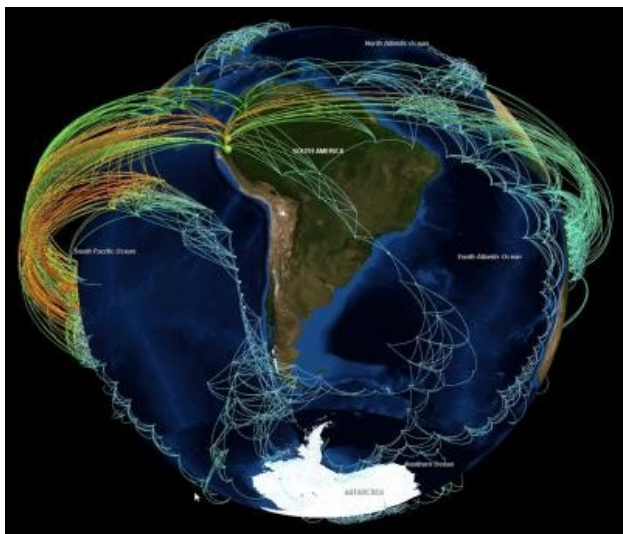
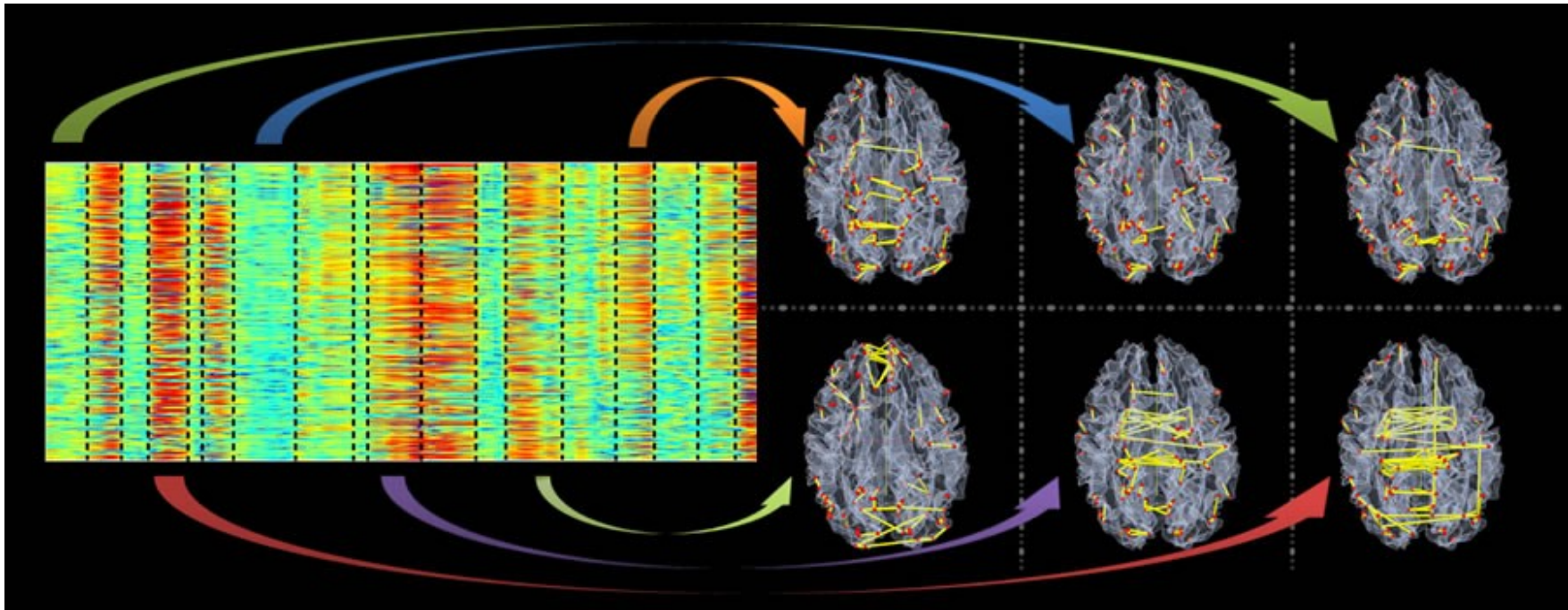


Cluster heat map of gene expression data

A.N. Massa, K.L. Childs, H. Lin, G.J. Bryan, G. Giuliano, and C.R. Buell, “*The Transcriptome of the Reference Potato Genome Solanum tuberosum Group Phureja Clone DM1-3 516R44*”, PloS ONE **6**(10), e26801 (2011).

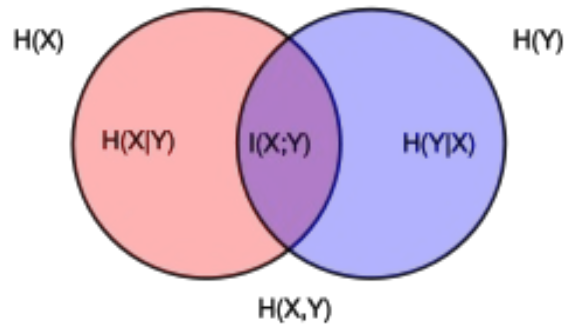


B. Barzel and A.-L. Barabási, “*Network link prediction by global silencing of indirect correlations*”, Nat. Biotech. **31**, 720-725 (2013).



Network Inference Problems

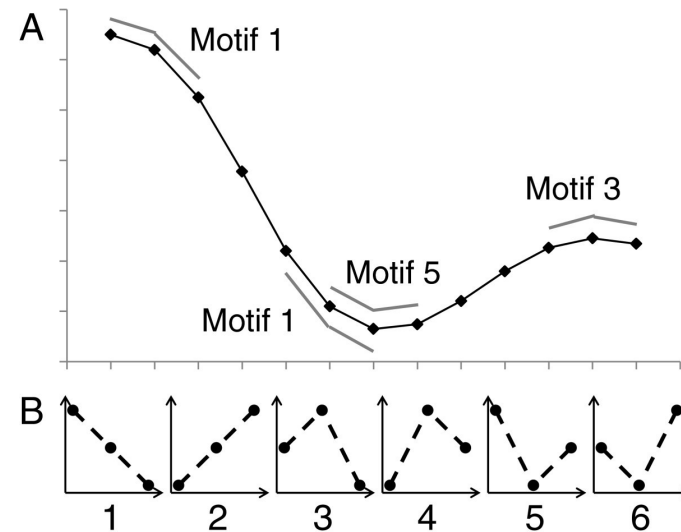
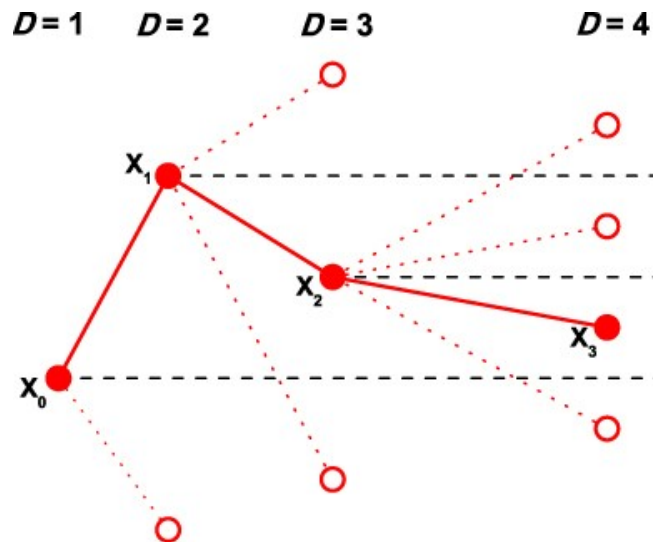
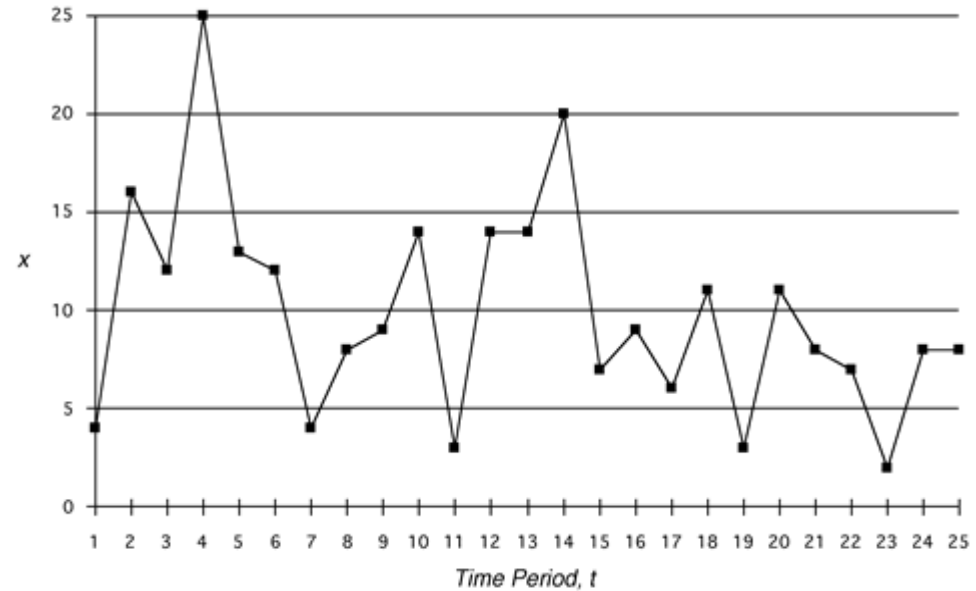
- Which similarity measure to use
- How to choose a threshold
- How much data is available
- How to avoid the (usual) noise in the data
- How to recover coupling strengths
- Which are the directions in the interactions
- How many “units” observed
- ...



$$I(X;Y) = H(X) - H(X|Y)$$

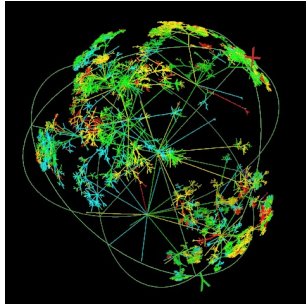
$$= H(X) + H(Y) - H(X,Y),$$

$$I(X;Y) = \sum_x \sum_y p(x,y) \log \frac{p(x,y)}{p(x)p(y)},$$

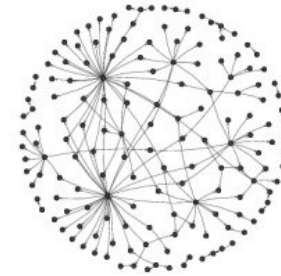


C. Bandt and B. Pompe, "Permutation Entropy: A Natural Complexity Measure for Time Series", Phys. Rev. Lett. **88**(17), 174102(4) (2002).

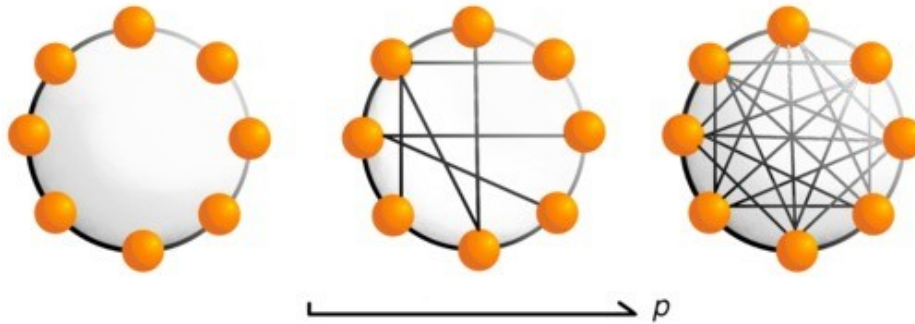
Inferred



Comparison

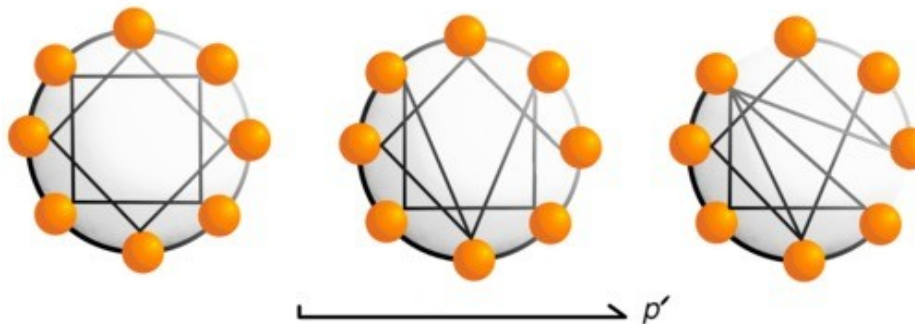


Underlying



$$E[M_N] = p \frac{N(N-3)}{2} + N$$

Expected number of edges



$$E[M_N] = \frac{N k}{2} = \frac{N(N/4)}{2}$$

Expected number of edges

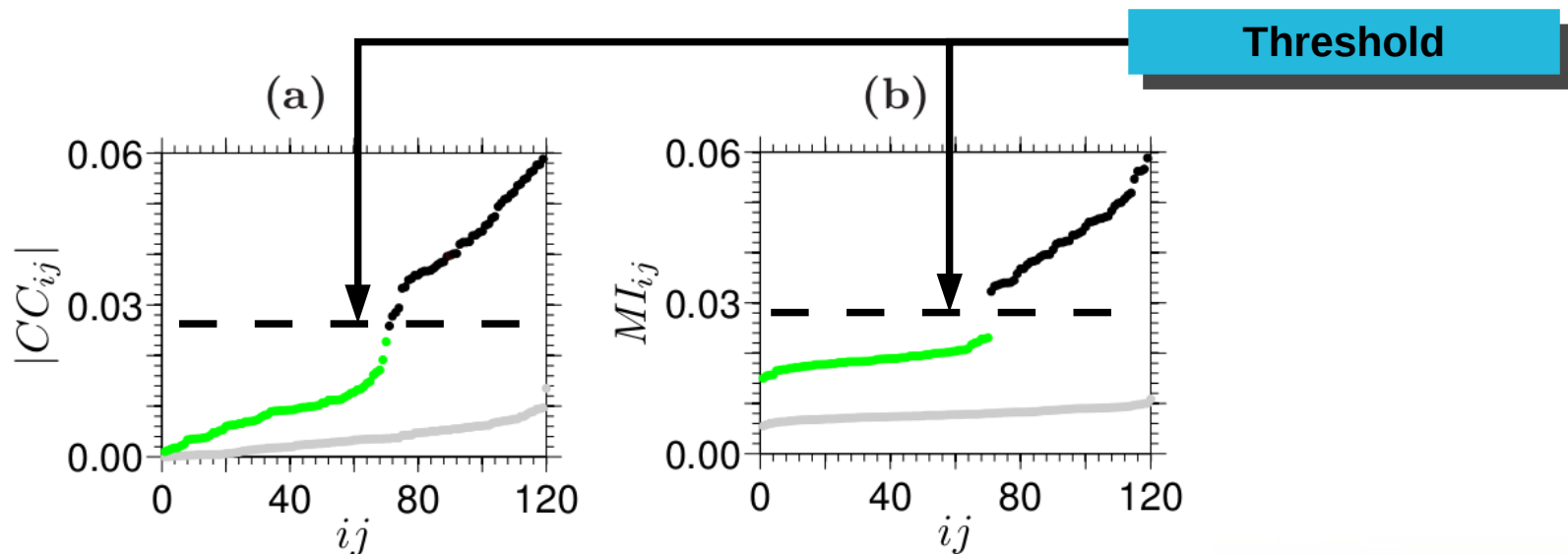
Poster: N. Rubido, et al., “Exact detection of direct links in networks of interacting dynamical units”.

Results

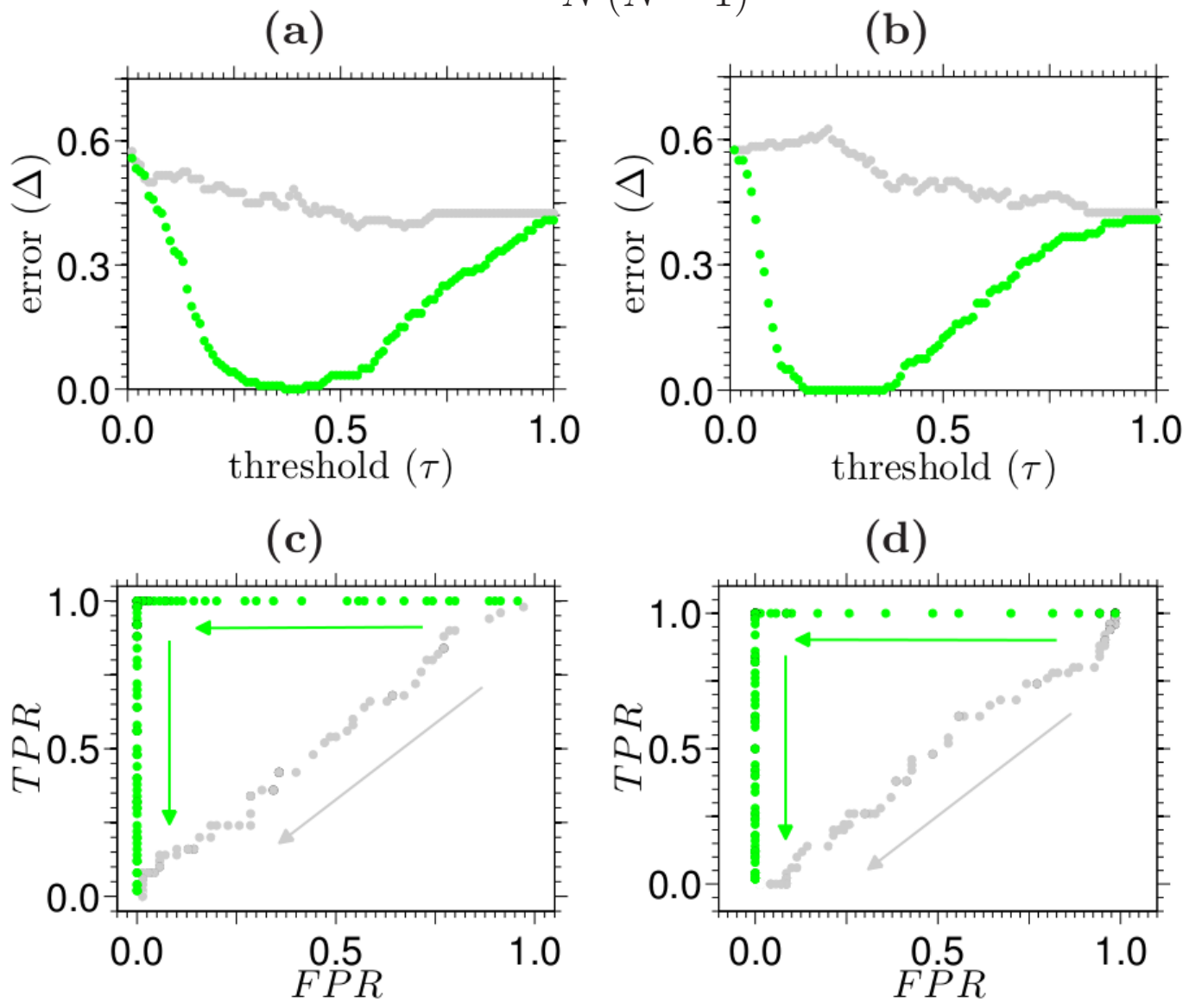
$$x_{n+1}^{(i)} = (1 - \epsilon) f_i(x_n^{(i)}) + \frac{\epsilon}{d_i} \sum_{j=1}^N W_{ij} f_j(x_n^{(j)}) \quad W_{ij} = A_{ij} (1 + g \xi_{ij})$$

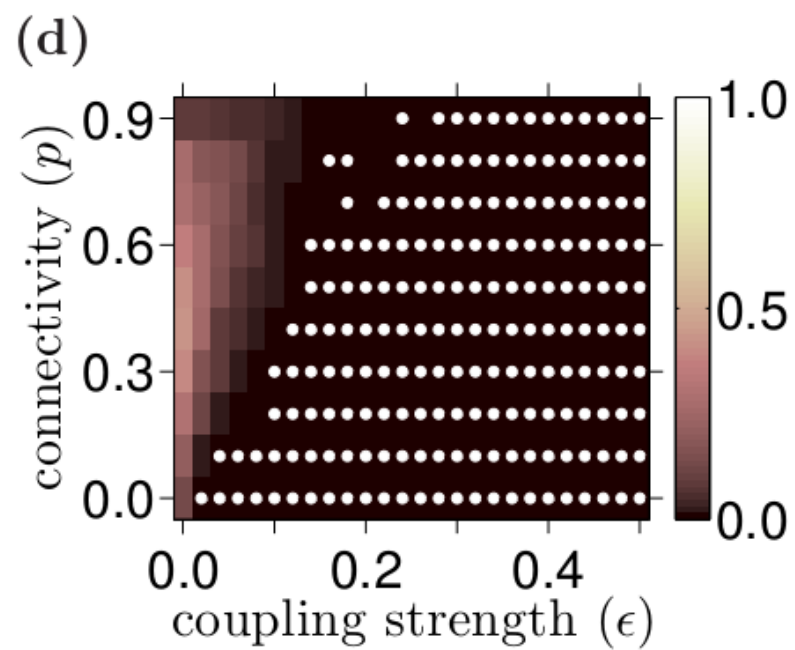
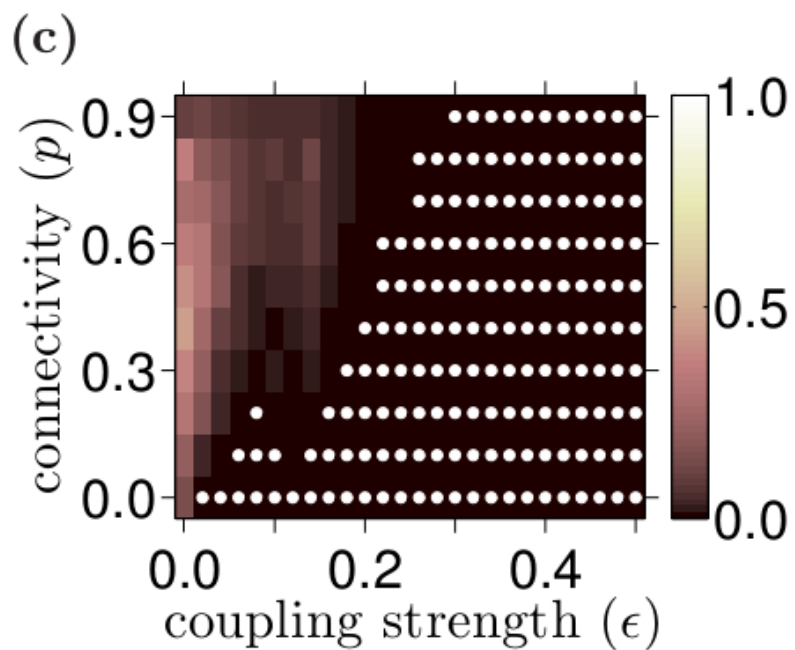
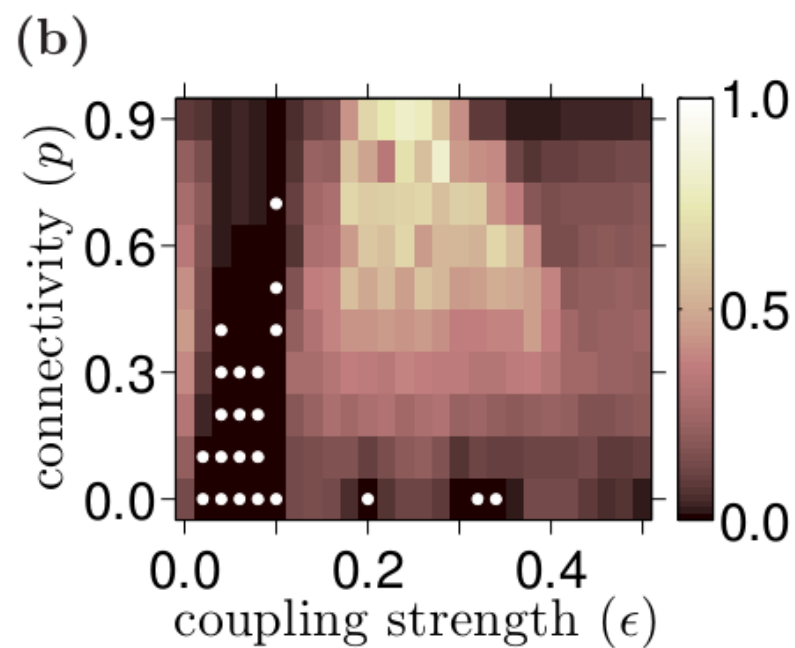
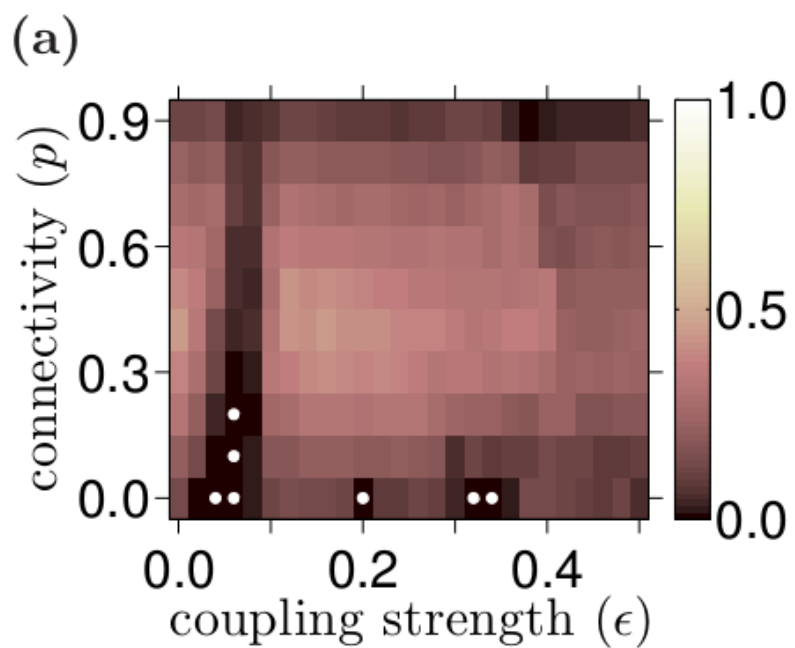
Kunihiko Kaneko, "Overview of coupled map lattices", Chaos 2(3), 279 (1992).

- Logistic maps
- Circle maps
- Optical maps
- Tent maps



$$\Delta = \frac{\sum_{i,j=1}^N |A_{ij} - A_{\tau,ij}|}{N(N-1)}$$





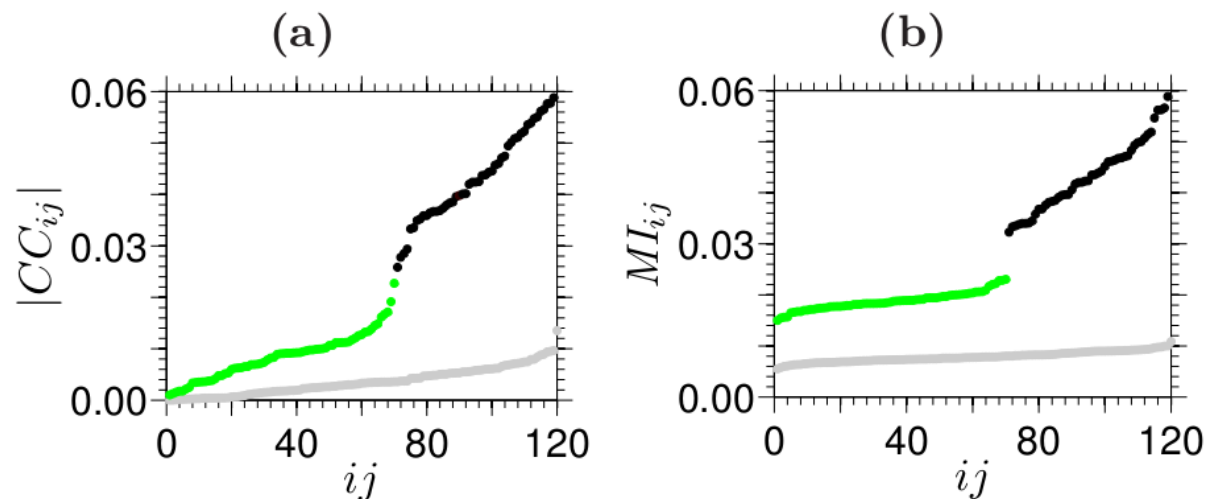
Articles:

N. Rubido, A.C. Martí, E. Bianco-Martínez, C. Grebogi, M.S. Baptista, and C. Masoller, “*Exact detection of direct links in networks of interacting dynamical units*”, submitted (2014) [available at: <http://arxiv.org/abs/1403.4839>].

CONCLUSIONS (take home messages):

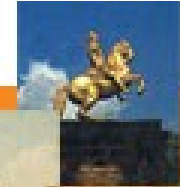
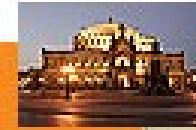
CC and MI allow to infer the underlying networks of coupled dynamical systems, without errors, from finite-size time-series measurements.
The correct detection of links depends on the existence of a gap in the ordered values of the similarity measures between pairs of nodes.

E. Bianco-Martínez, N. Rubido, C.G. Antonopoulos, and M.S. Baptista, “*Network Inference by Mutual Information Rates from Complex Time-series*”, in preparation (2014).





Workshops,
seminars,
conferences



THANK YOU

Ongoing projects:

A. L'Her, P. Amil, R. García, F. Abellá, M. S. Baptista, A. C. Martí, C. Cabeza, and N. Rubido, “*Electronic circuit implementation of a network of Logistic maps*”.
Universidad de la República (UdelaR), Montevideo, Uruguay.

N. Rubido and A.J. Pons, “*Neural circuits and transfer functions*”.
Universidad Politécnica de Barcelona (UPC), Terrassa, Spain.



Institute for Complex Systems
and Mathematical Biology (ICSMB)

“To understand biology at the system level, we must examine the structure and dynamics of cellular and organismal function, rather than the characteristics of isolated parts of a cell or organism.”

Hiroaki Kitano

