



Big Data <-> Big Networks

Albert Diaz Guilera

- “Donat el caràcter i la finalitat exclusivament docent i eminentment il·lustrativa de les explicacions a classe d'aquesta presentació, l'autor s'acull a l'article 32 de la Llei de propietat intel·lectual vigent respecte de l'ús parcial d'obres alienes com ara imatges, gràfics o altre material contingudes en les diferents diapositives”
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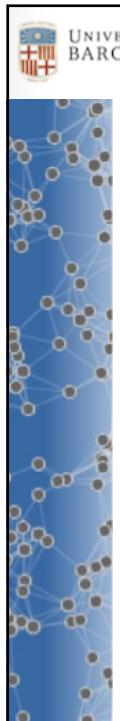
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BARCELONA

Complex Networks

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@anduviera

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Multiplex & Time Dependent Networks

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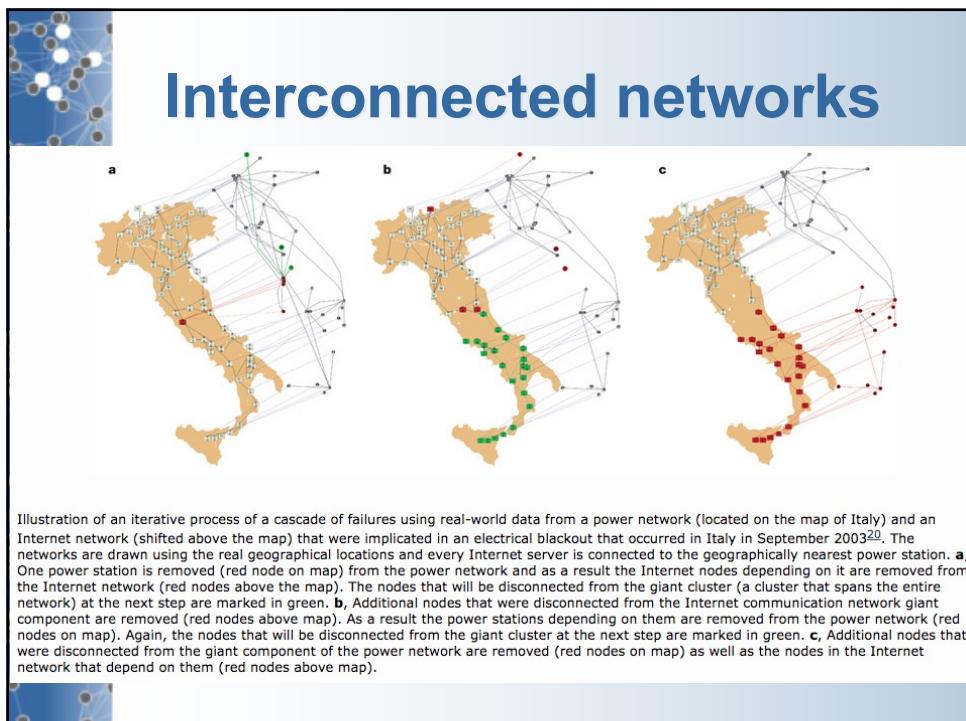
COMPLEXITAT

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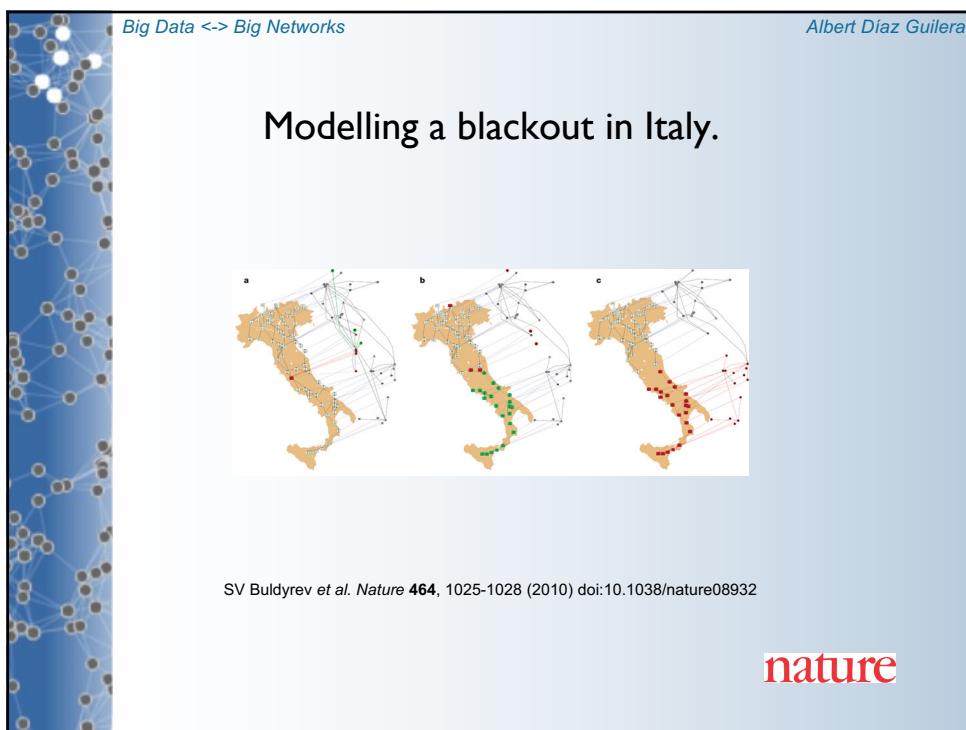
Big data => more complex networks

- New paradigms:
 - Network of networks
 - Time-dependent networks
 - Interconnected networks
 - Multilayer networks. Multiplex networks

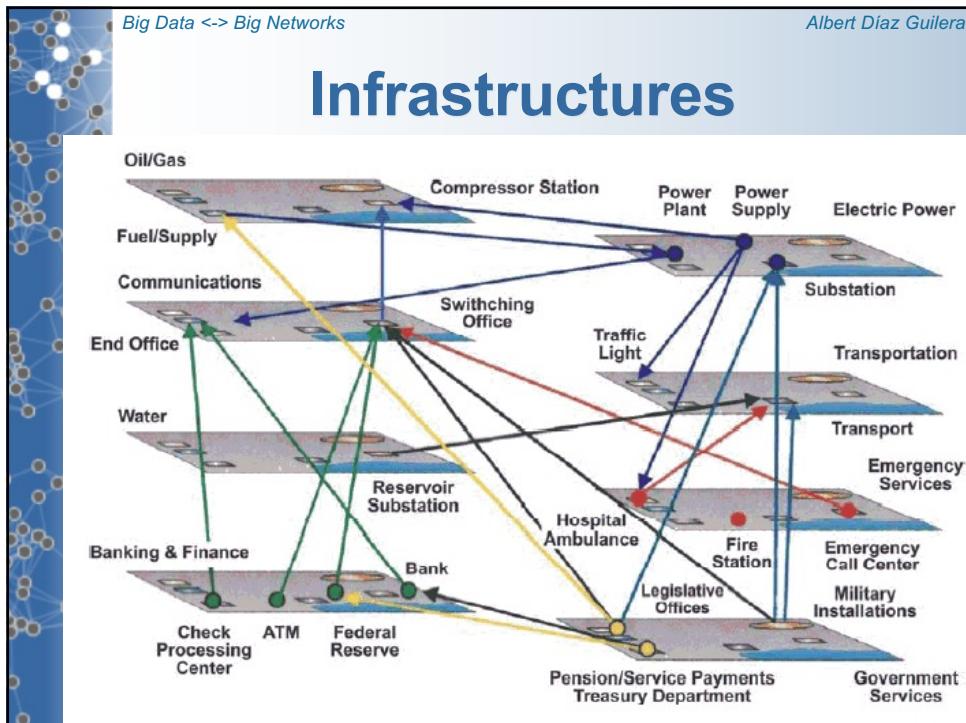
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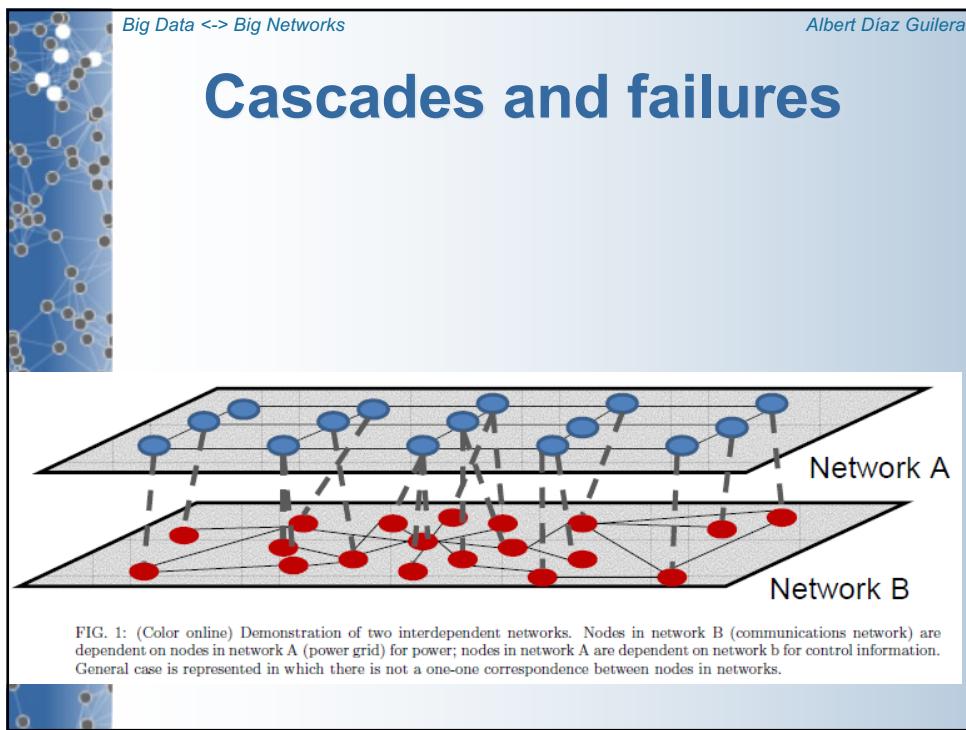
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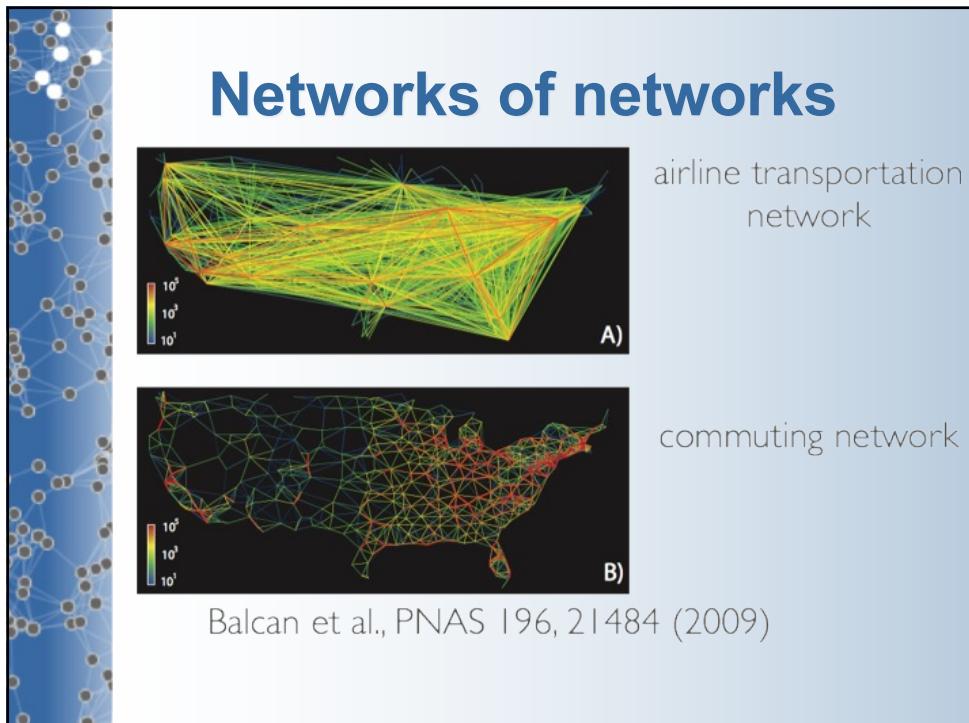
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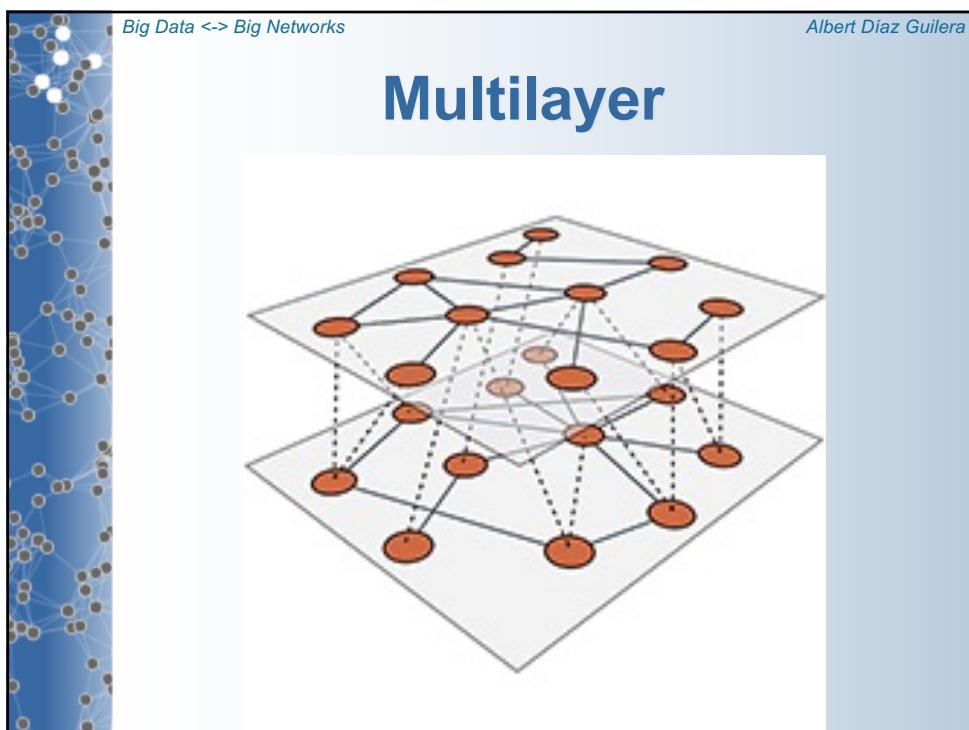
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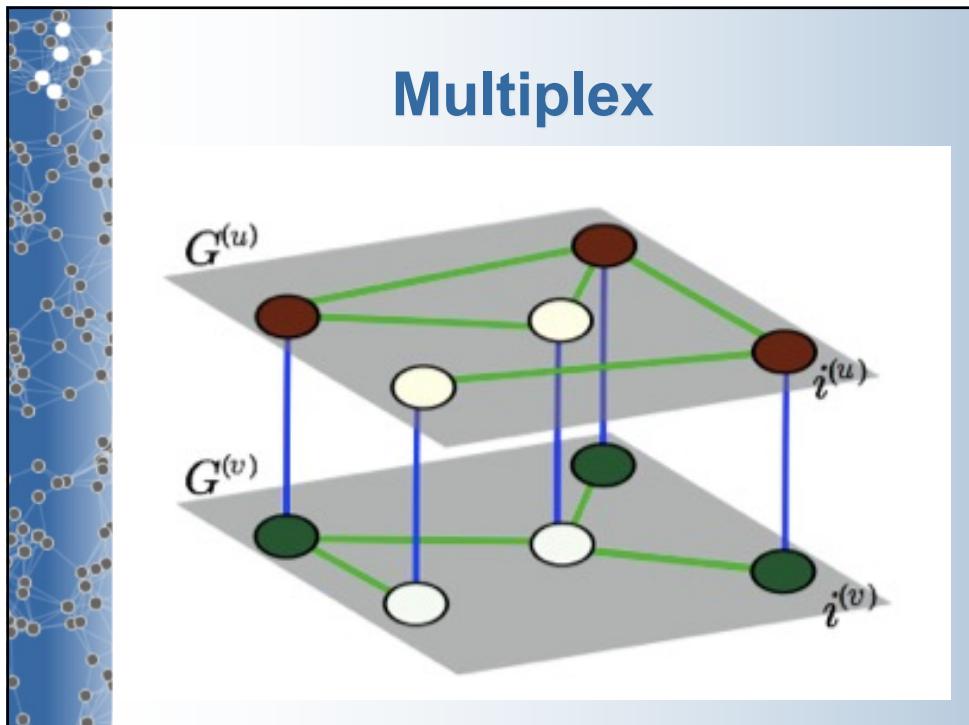
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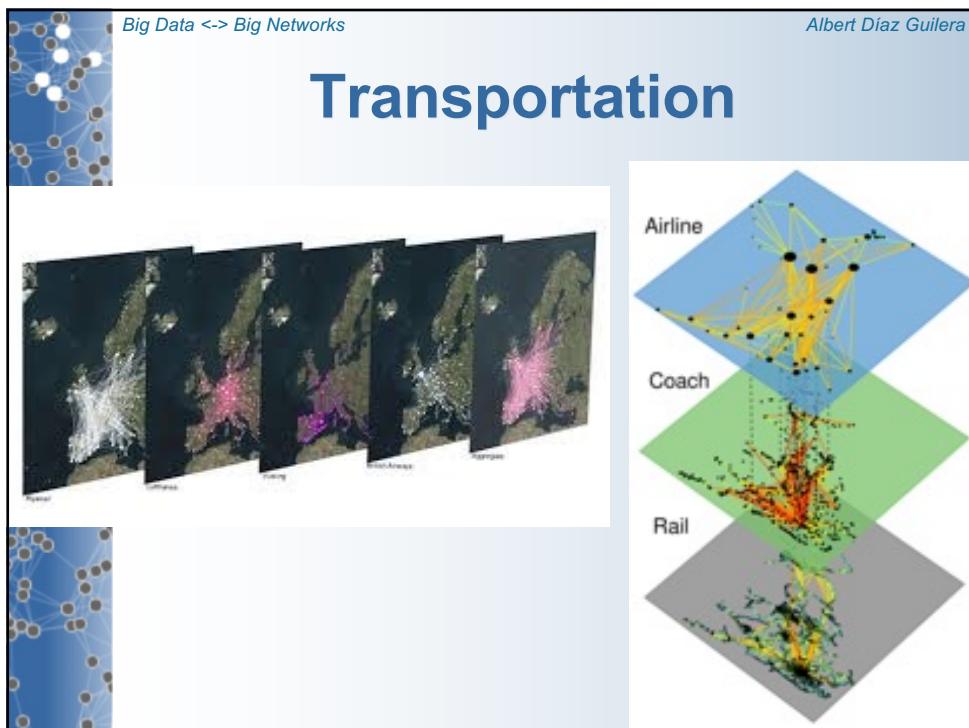
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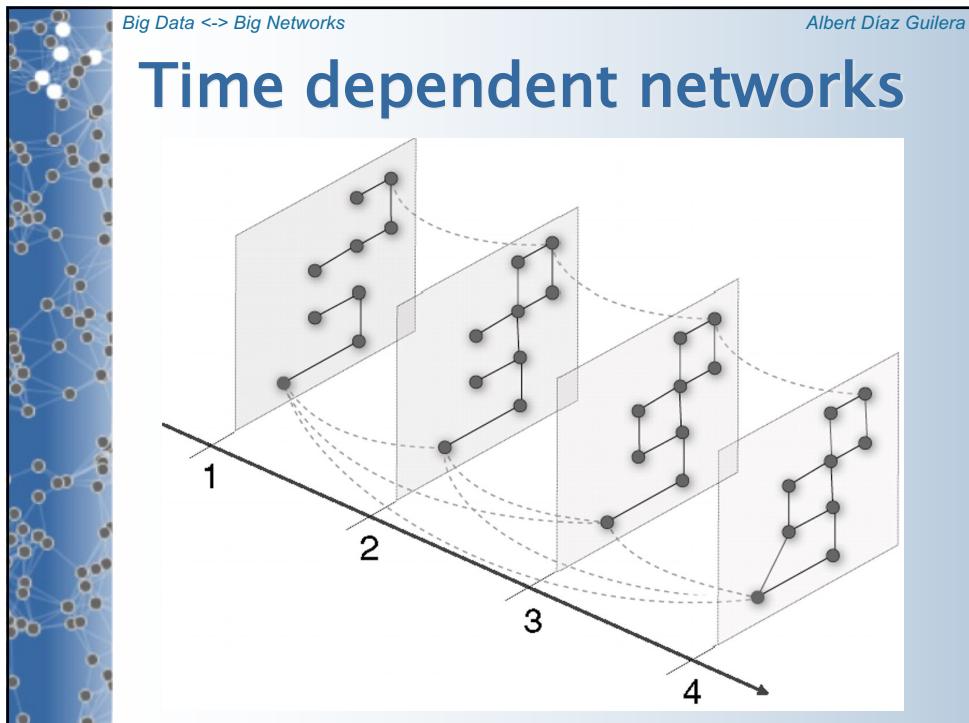
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Enzo Nicosia's



- C. Elegans (synapses and gap junctions)
- BioGrid (Physical and Genetic interactions among all proteins)
- OpenFlights continental airport networks (each continent, layers represent airlines)
- APS scientific collaboration network (layers are PACS 0-9)

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Manlio de Domenico



- Twitter (different events)
- Genetic interactions for different organisms
- FAO Multiplex Trade Network (layers are for specific products)

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Resources

SOFTWARE FOR MULTIPLEX METRICS AND MODELS (back to top)

MAMMUL (Metrics And Models for MULTiplex networks) is a collection of tools for the analysis and modelling of multiplex networks.

MAMMUL can be downloaded from GitHub: <https://github.com/KatolaZ/mammult>

MultiRed is a Python module for the structural reduction of multiplex networks, based on the paper:

M. De Domenico*, V. Nicosia*, A. Arenas, V. Latora, "Structural Reducibility of Multilayer Networks", *Nat. Commun.* 6, 6864 (2015).

MultiRed can be downloaded from GitHub: <https://github.com/KatolaZ/multired>

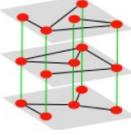
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MultinetX

<https://github.com/nkoub/multinetx>



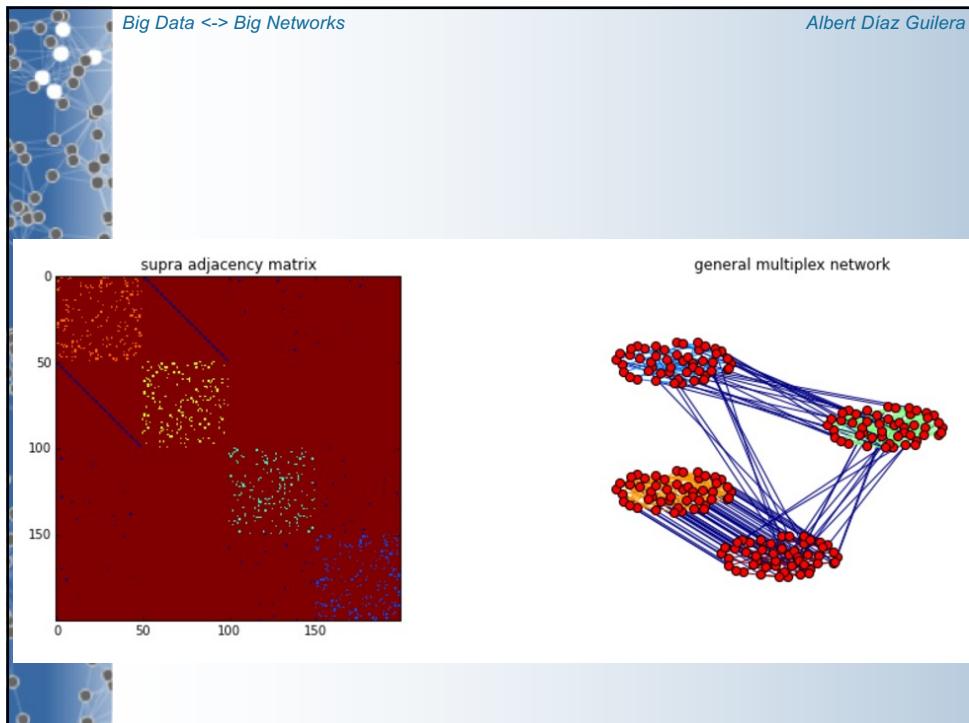
multiNetX v1.0

multiNetX is a python package for the manipulation and visualization of multilayer networks. The core of this package is a MultilayerGraph, a class that inherits all properties from networkx.Graph().

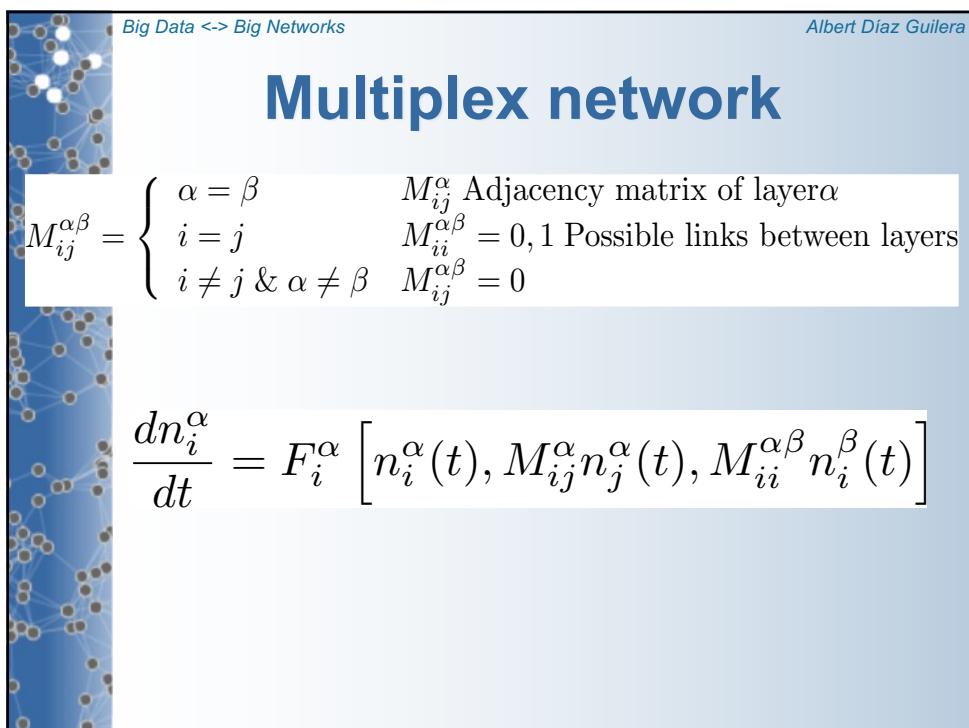
This allows for:

- Creating networks with weighted or unweighted links (only undirected networks are supported in this version)
- Analysing the spectral properties of adjacency or Laplacian matrices
- Visualizing dynamical processes by coloring the nodes and links accordingly

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One-to-one

$$M_{ii}^{\alpha\beta} = M^{\alpha\beta} = 0, 1 \text{ the same } \forall i$$

Matrix of layers

$$\frac{dn_i^\alpha}{dt} = F_i^\alpha \left[n_i^\alpha(t), M_{ij}^\alpha n_j^\alpha(t), M^{\alpha\beta} n_i^\beta(t) \right]$$

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CHARACTERIZATION

- Degree
- Clustering
- Centrality
- Communities
- Layer comparison

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Degree. Activity

$$\mathbf{k}_i = \{k_i^{[1]}, k_i^{[2]}, \dots, k_i^{[M]}\}$$

We say that node i , with $i = 1, 2, \dots, N$, is *active* at layer α if $k_i^{[\alpha]} > 0$. We can then associate to each node i a *node-activity vector*

$$\mathbf{b}_i = \{b_i^{[1]}, b_i^{[2]}, \dots, b_i^{[M]}\} \quad (2)$$

where

$$b_i^{[\alpha]} = 1 - \delta_{0, k_i^{[\alpha]}}$$

i.e., $b_i^{[\alpha]} = 1$ if node i has at least one edge at layer α , and is 0 otherwise. We call *node-activity* B_i of node i the number of layers on which node i is active:

$$B_i = \sum_{\alpha} b_i^{[\alpha]} \quad (3)$$

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Degree. Participation ratio

$$P_i = \frac{M}{M-1} \left[1 - \sum_{\alpha=1}^M \left(\frac{k_i^{[\alpha]}}{o_i} \right)^2 \right]$$

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Layers comparison

- Hamming distance

$$H^{[\alpha, \beta]} = \frac{\sum_i b_i^{[\alpha]} (1 - b_i^{[\beta]}) + b_i^{[\beta]} (1 - b_i^{[\alpha]})}{\min(N^{[\alpha]} + N^{[\beta]}, N)}.$$

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Dissimilarity

$D_r(p, q) = 0.8146$
 $D(p, q) = 0.5837$

$D_r(p, q) = 0.8905$
 $D(p, q) = 0.5723$

$D_r(p, q) = 1.0000$
 $D(p, q) = 0.8316$

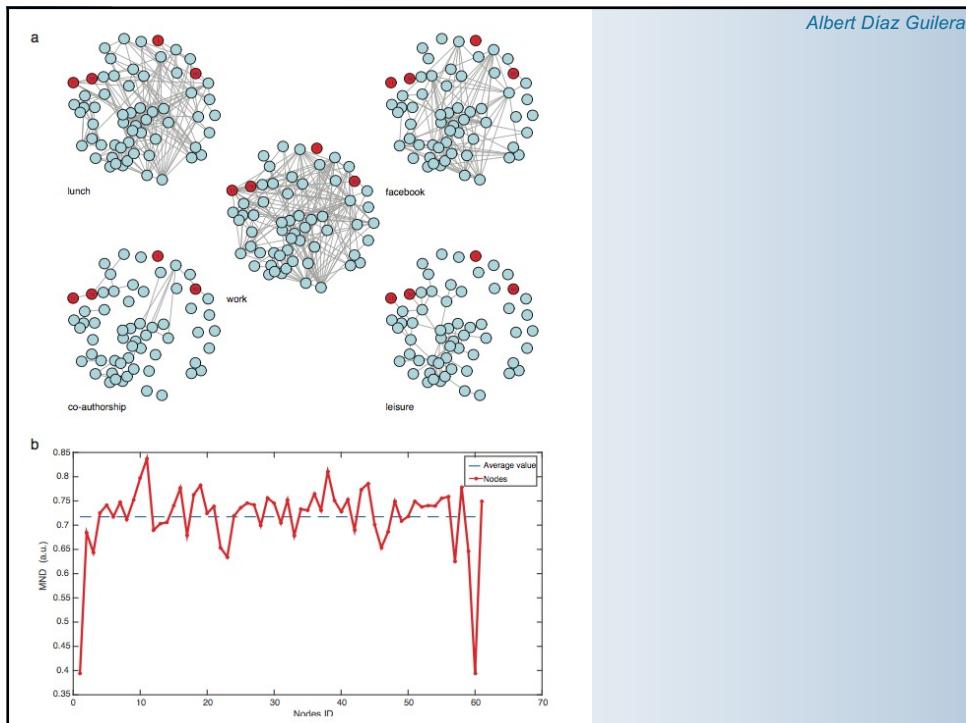
Quantification of network structural dissimilarities

Tiago A. Schieber, Laura Carpi, Albert Díaz-Guilera, Panos M. Pardalos, Cristina Masoller & Martín G. Ravetti

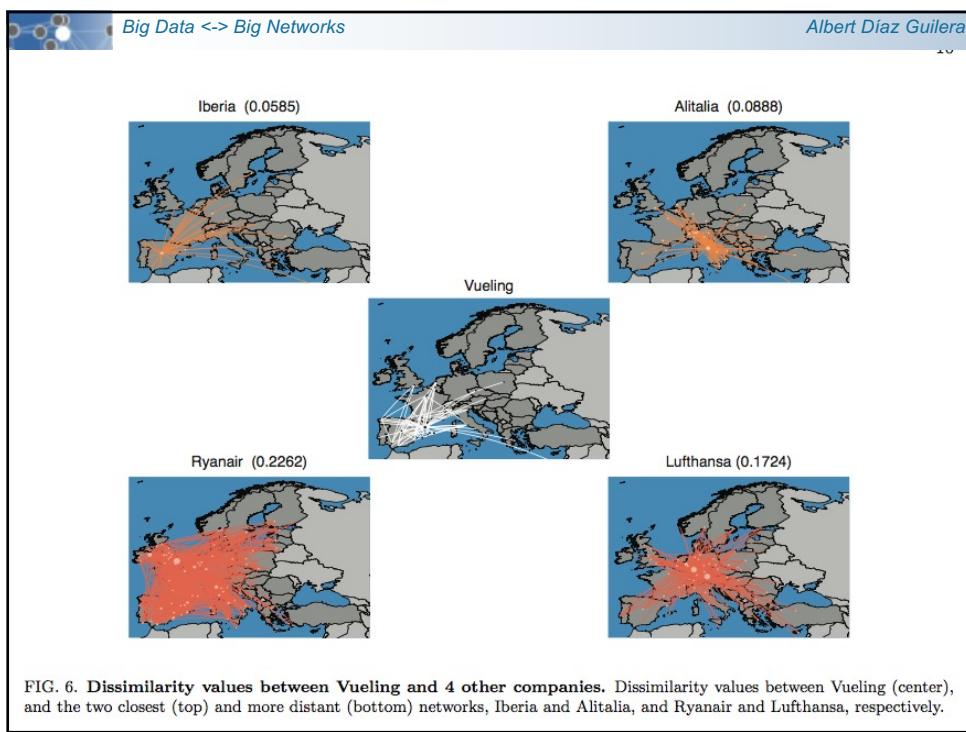
Nature Communications 8,
Article number: 13928 (2017)
doi:10.1038/ncomms13928

Received: 26 May 2016
Accepted: 15 November 2016
Published online: 09 January 2017

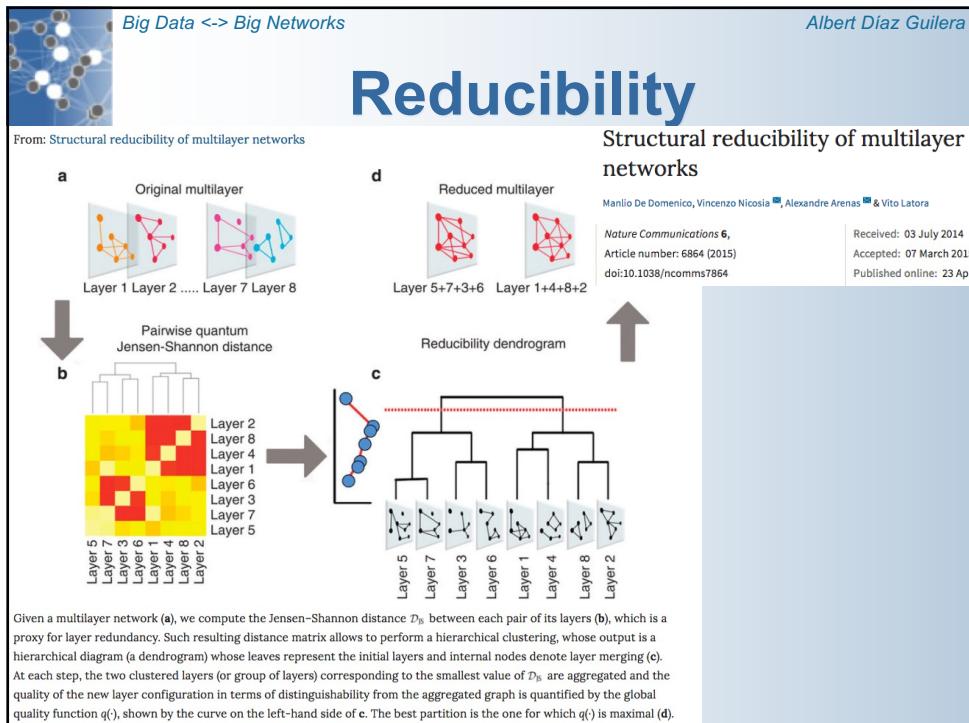
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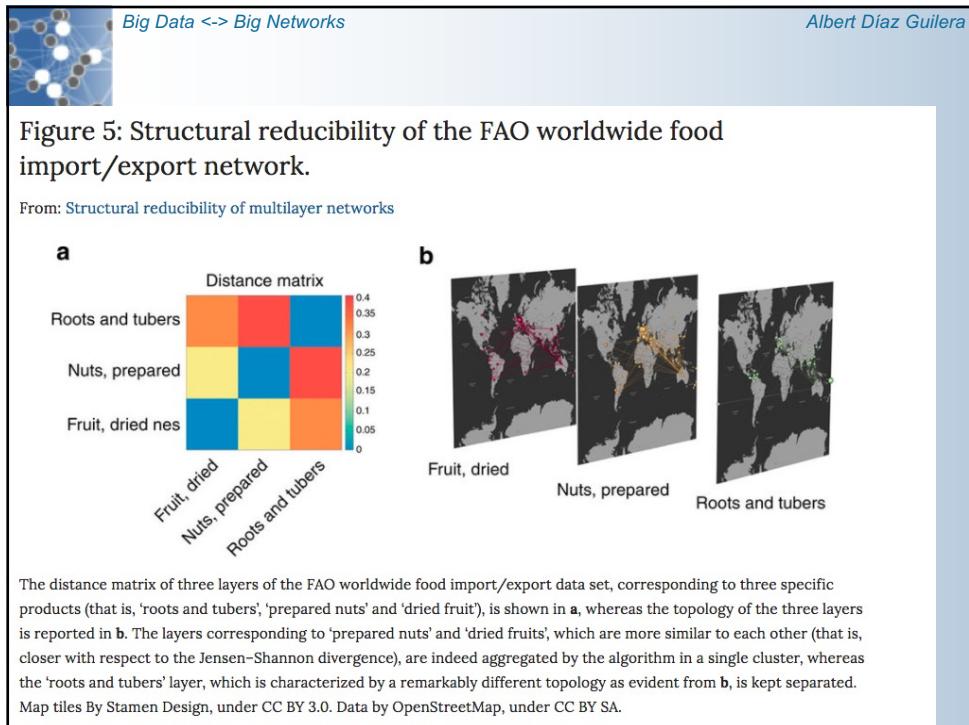
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Clustering

II. STATISTICS OF CYCLES

The diagram shows two cycles. The left cycle has nodes connected by edges of type 1 (solid red), type 2 (dashed green), and type 3 (dash-dot blue). The counts are: $s_{11} = 2$, $s_{12} = 2$, $s_{22} = 1$. The right cycle has nodes connected by edges of type 2 (dashed green) and type 3 (dash-dot blue). The counts are: $s_{11} = 0$, $s_{22} = 0$, $s_{33} = 0$, $s_{12} = 2$, $s_{13} = 2$, $s_{23} = 2$.

FIG. 1. Notation of multiplex cycles.
We characterize a given cycle in a multiplex network, by the matrix $\mathbf{s} = \{s_{ab}\}_{a,b=1,\dots,M}$, where each element s_{ab} defines the number of nodes in the cycle which connect an edge of type a with an edge of type b . When

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Centralities

Ranking in interconnected multilayer networks reveals versatile nodes

Manlio De Domenico , Albert Solé-Ribalta, Elisa Omodei, Sergio Gómez & Alex Arenas

<i>Nature Communications</i> 6 , Article number: 6868 (2015) doi:10.1038/ncomms7868	Received: 17 October 2014 Accepted: 07 March 2015 Published online: 23 April 2015
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Communities

Detection of gene communities in multi-networks reveals cancer drivers

Laura Cantini , Enzo Medico , Santo Fortunato & Michele Caselle

Scientific Reports 5, Article number: 17386
(2015)
doi:10.1038/srep17386

Received: 08 September 2015
Accepted: 29 October 2015
Published online: 07 December 2015



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- To address this problem, we combined, in a single multi-network, four different gene networks:
 - (i) Transcription Factor (TF) co-targeting network,
 - (ii) microRNA co-targeting network,
 - (iii) Protein-Protein Interaction (PPI) network
 - (iv) gene co-expression network.

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Community detection in the multi-network

After filtering, all the layers of the multi-network were sparse enough to perform community detection. The design of community detection algorithms on multi-networks is still an open problem²⁸. We propose here a possible solution based on the use of the consensus clustering procedure described in²⁹. We used five widely adopted algorithms: Infomap¹¹, OSLOM¹², Label propagation¹³, Louvain¹⁴ and Modularity optimization via simulated annealing¹⁵. We integrated in our software all five algorithms, leaving the choice of the preferred one to the user. We

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MODELS

- Individual layers
- Correlations between layers

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Time dependent networks

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Static network

- Topology is fixed
- The state of the nodes evolve according to:
 - Their own rules
 - The interaction with their neighbours

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Dynamic network

- Links change!!!
- What determines the rules of change of the links?

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Classification

- Adaptive systems
- Time dependence of the links

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Adaptive networks

© 2009

Adaptive Networks

Theory, Models and Applications

Editors: Gross, Thilo, Sayama, Hiroki (Eds.)

Free Preview

This volume is a state-of-the-art introduction to and survey on adaptive networks, combining aspects of the evolution of networks and dynamics on networks

Dynamics of epidemic diseases on a growing adaptive network

Güven Demirel, Edmund Barter & Thilo Gross

Scientific Reports 7, Article number: 42352
 (2017)
 doi:10.1038/srep42352

Received: 12 October 2016
 Accepted: 08 January 2017
 Published online: 10 February 2017

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- Nodes look for improving their outcome.
- “Better” neighbours, depending on node states
- Complex systems are usually self-organized, not engineered, not centralized

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- Social agent: looking for more connected
- Economic agent: looking for more wealthy
- Access to limited information

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Dynamics of the system

- Importance of the two time scales in the final outcome of the complex system

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• **Adaptive coevolutionary networks: a review**

Thilo Gross, Bernd Blasius

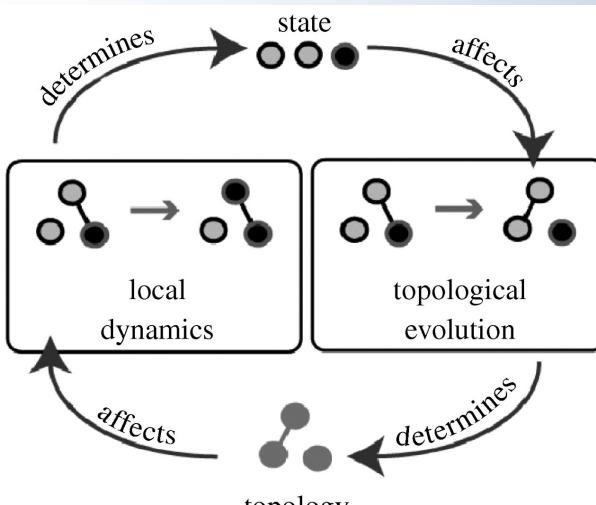
Published 6 March 2008. DOI: 10.1098/rsif.2007.1229



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In an adaptive network, the evolution of the topology depends on the dynamics of the nodes.



The diagram illustrates the relationship between local dynamics and topological evolution in an adaptive network. It shows two boxes: 'local dynamics' and 'topological evolution'. Arrows indicate a 'determines' relationship from 'local dynamics' to 'state' (represented by three nodes) and from 'state' to 'topological evolution'. Arrows also indicate an 'affects' relationship from 'topological evolution' back to 'local dynamics' and from 'topology' (represented by three nodes) to 'local dynamics'.

Thilo Gross, and Bernd Blasius J. R. Soc. Interface
2008;5:259-271

© 2007 The Royal Society



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Time-dependent

- Links change because some external (not coupled to states) rules
 - Random
 - Due to mobility → contact networks

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Physics Reports

Volume 519, Issue 3, October 2012, Pages 97–125



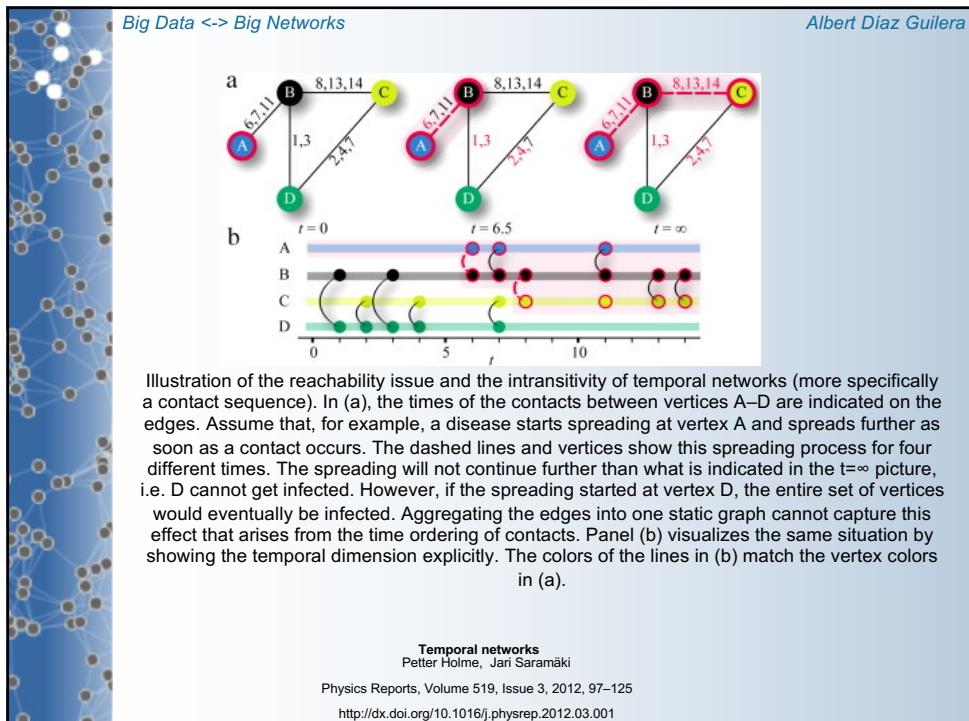
Temporal networks

Petter Holme^{a, b, c},    Jari Saramäki^d

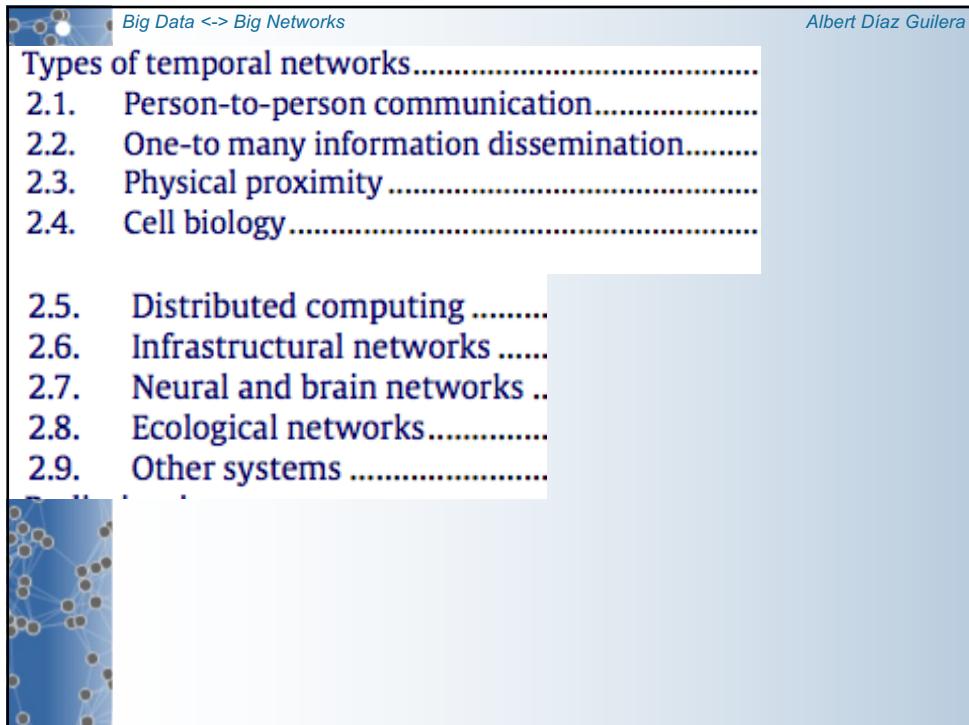
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<https://doi.org.sire.ub.edu/10.1016/j.physrep.2012.03.001> [Get rights and content](#)

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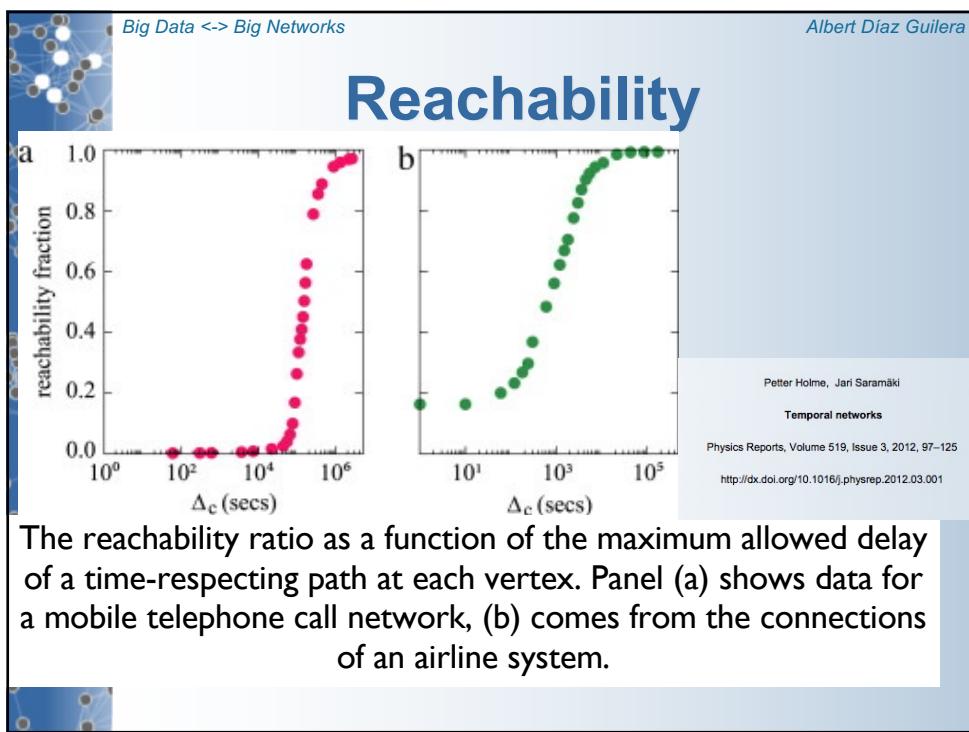
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Reachability graph



Fig. 10.

Reachability graphs. Panel (a) shows a contact sequence (same as in Fig. 1) and (b) shows its reachability graph.

Panel (a) shows a contact sequence graph with nodes A, B, C, and D. Edges are labeled with contact IDs: A-B (6,7,11), A-D (1,3), B-C (8,13,14), and B-D (2,4,7). Panel (b) shows the corresponding reachability graph with nodes A, B, C, and P. Directed edges exist from A to B, A to C, B to C, B to P, C to P, and C to A.

```

graph LR
    subgraph a [ ]
        A((A)) --- B((B))
        A --- D((D))
        B --- C((C))
        B --- D
        A --- B
    end
    subgraph b [ ]
        A((A)) --> B((B))
        A --> C((C))
        B --> C
        B --> P((P))
        C --> P
        C --> A
    end
    
```

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Generalizations



- Distances
- Components
- Centrality measures
- Communities

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 **Physics Reports**
Volume 486, Issues 3–5, February 2010, Pages 75–174 

Community detection in graphs

Santo Fortunato  · 
 Show more

<https://doi.org.sire.ub.edu/10.1016/j.physrep.2009.11.002> Get rights and content

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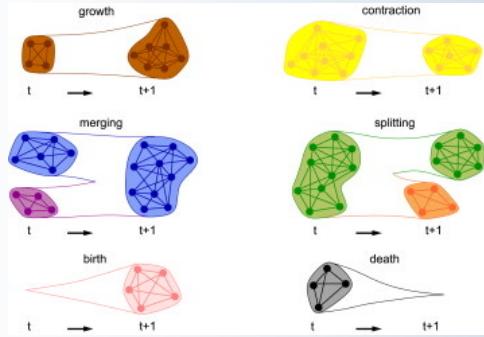


Fig. 27. Possible scenarios in the evolution of communities. Reprinted figure with permission from Ref. [380].

Santo Fortunato
Community detection in graphs
 Physics Reports, Volume 486, Issues 3–5, 2010, 75–174
<http://dx.doi.org/10.1016/j.physrep.2009.11.002>

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Contact networks

- Agents (humans) move
- Creating a time-dependent network of connectivities

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SocioPatterns

<http://www.sociopatterns.org/>

SocioPatterns

ABOUT | GALLERY | PUBLICATIONS

WELCOME

SocioPatterns is an interdisciplinary research collaboration formed in 2008 that adopts a data-driven methodology to study social dynamics and human activity.

Since 2008, we have collected longitudinal data on the physical proximity and face-to-face contacts of individuals in numerous real-world environments, covering widely varying contexts across several countries: schools, museums, hospitals, etc. We use the data to study human behaviour and to develop agent-based models for the transmission of infectious diseases.

We make most of the collected data freely available to the scientific community.

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Data

DATASETS

This page provides a collection of datasets obtained through the SocioPatterns sensing platform.

- Household
- Schools
- Hospitals
- Conferences

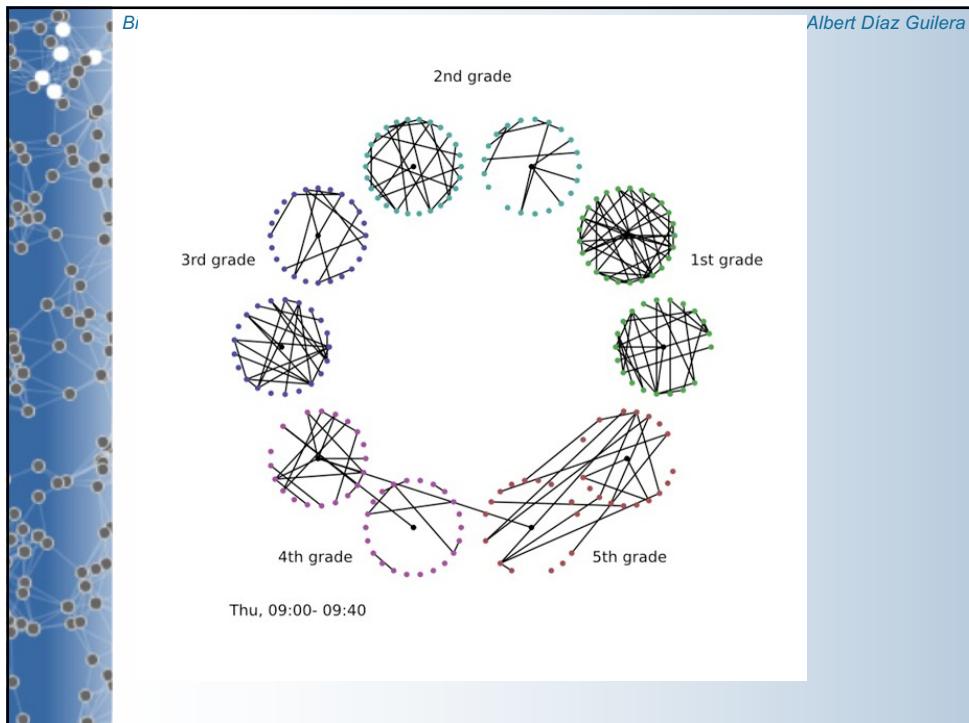
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Conference

What's in a crowd? Analysis of face-to-face behavioral networks. L. Isella et al.

J. Theor. Bio. 271 (2011) 166

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Modeling Human Dynamics of Face-to-Face Interaction Networks

Michele Starnini, Andrea Baronchelli, and Romualdo Pastor-Satorras
 Phys. Rev. Lett. **110**, 168701 – Published 15 April 2013

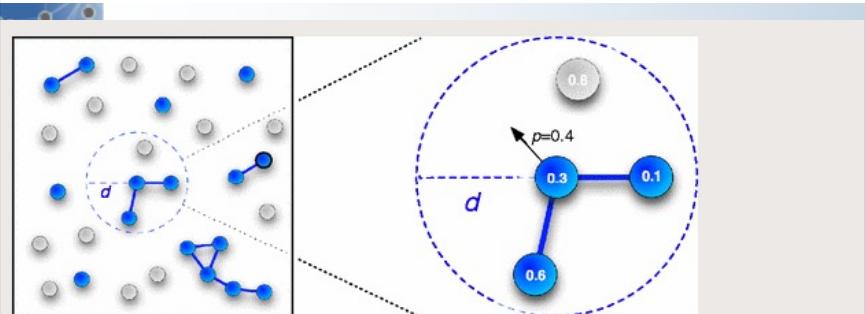


FIG. 1

[Open in new window](#)

Left: Blue (dark) colored agents are active, gray (light) agents do not move nor interact. Interacting agents, within a distance d , are connected by a link. Right: Each individual is characterized by a number representing her attractiveness. The probability for the central individual to move is $p = 1.0 - 0.6 = 0.4$, since the attractiveness of the inactive agent is not taken into account.

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