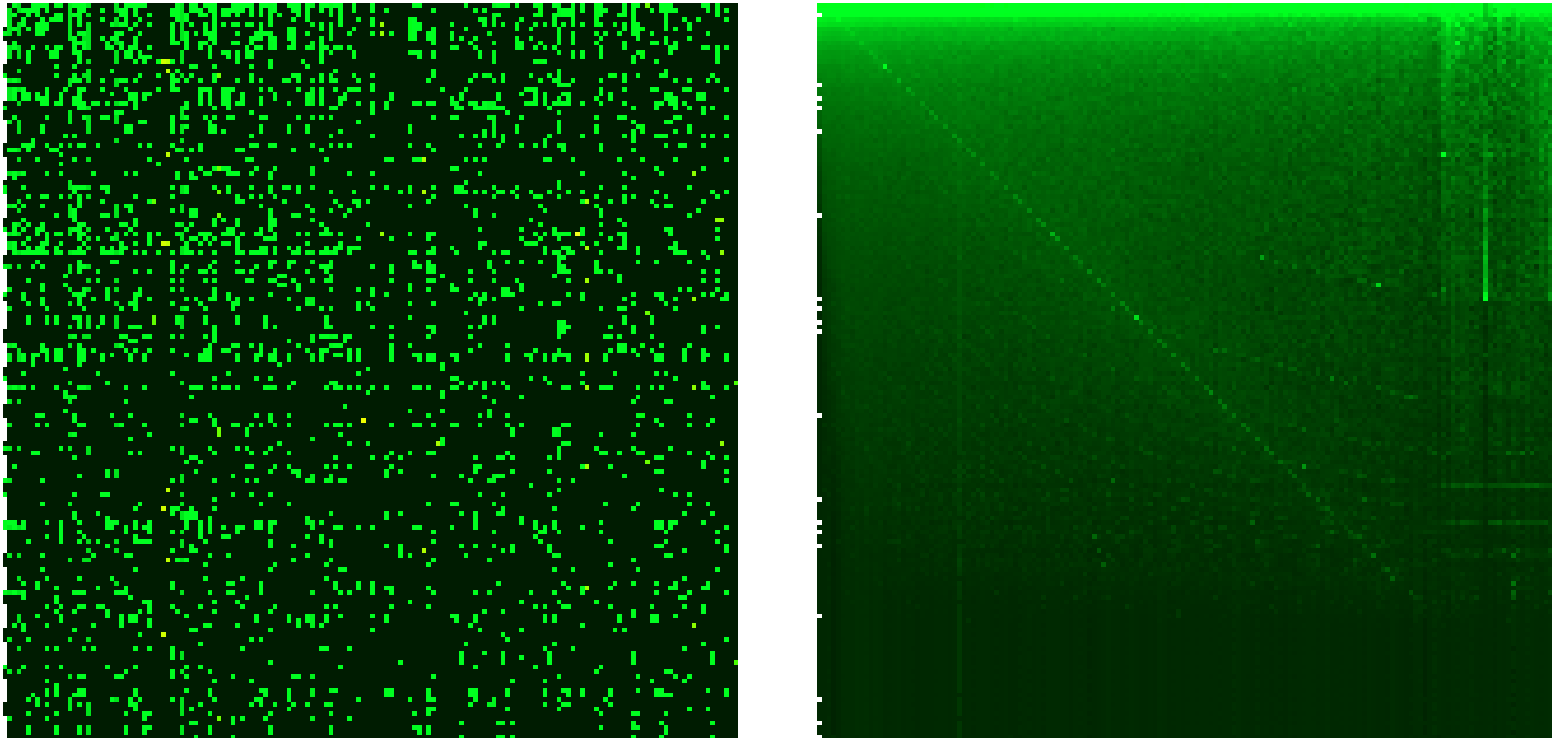


ApliGoogle

Applications of Google matrix to directed networks and Big Data

Défi MASTODONS CNRS
Colloque de restitution 2017



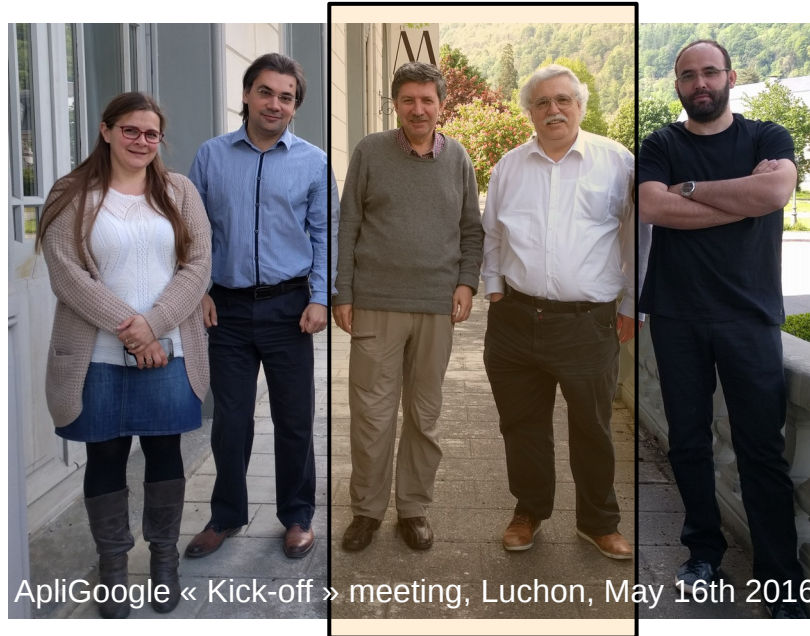
Google matrix G of the English Wikipedia network (Aug. '09) $N=3\ 282\ 257$. Left panel : close up 200x200 first elements.

<http://www.quantware.ups-tlse.fr/APLIGOOGLE/>

ApliGoogle partners



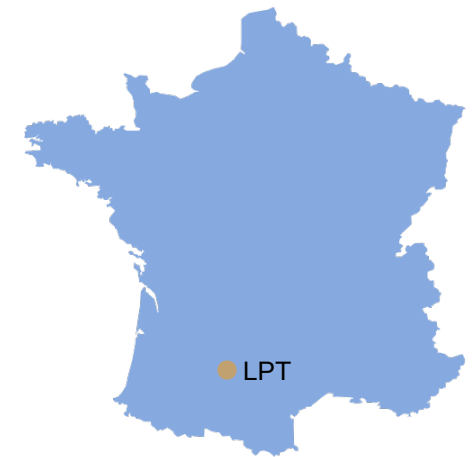
ApliGoogle partners



Laboratoire de Physique Théorique de Toulouse (UMR CNRS 5152)

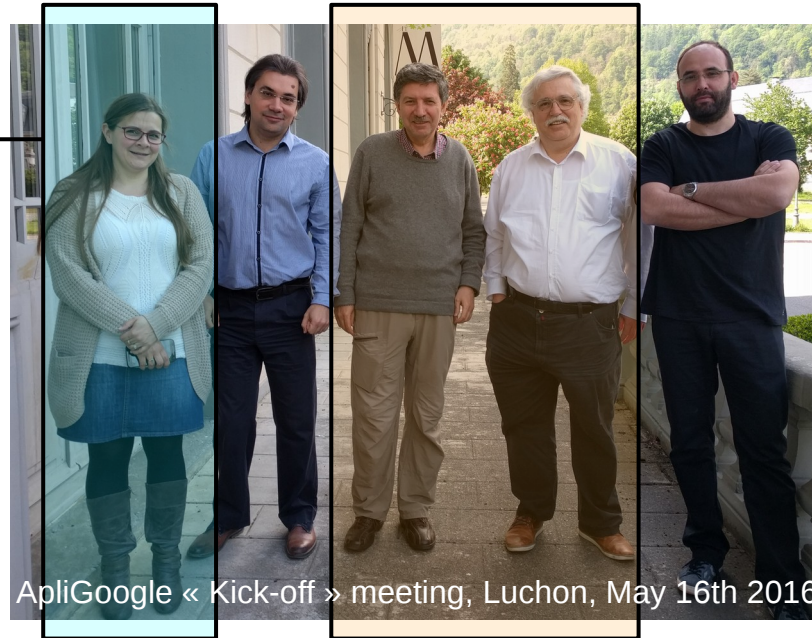
- DR1 D. Shepelyansky (PI)
- Pr. K. Frahm

Expertises: Quantum chaos,
Random Matrix Theory, Complex
directed networks



<http://www.quantware.ups-tlse.fr/dima>

ApliGoogle partners



ApliGoogle « Kick-off » meeting, Luchon, May 16th 2016

Institut de Recherche en Informatique de Toulouse
(UMR CNRS 5152)

- MCF K. Jaffrès-Runser
- S. El Zant (doctorant)
- Dr. T. Peng (post-doctorant)

Expertises: Wireless networks



<http://www.irit.fr/~Katia.Jaffres>

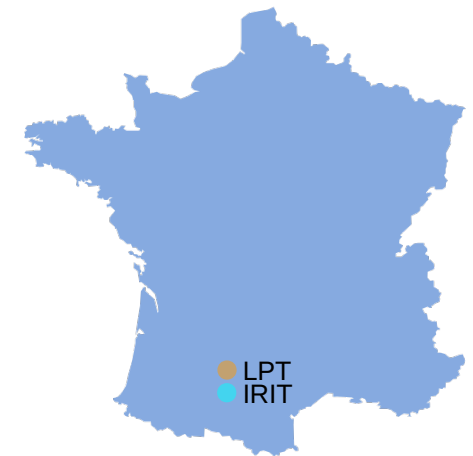
Laboratoire de Physique Théorique de Toulouse (UMR CNRS 5152)

- DR1 D. Shepelyansky (PI)
- Pr. K. Frahm

Expertises: Quantum chaos,
Random Matrix Theory, Complex
directed networks



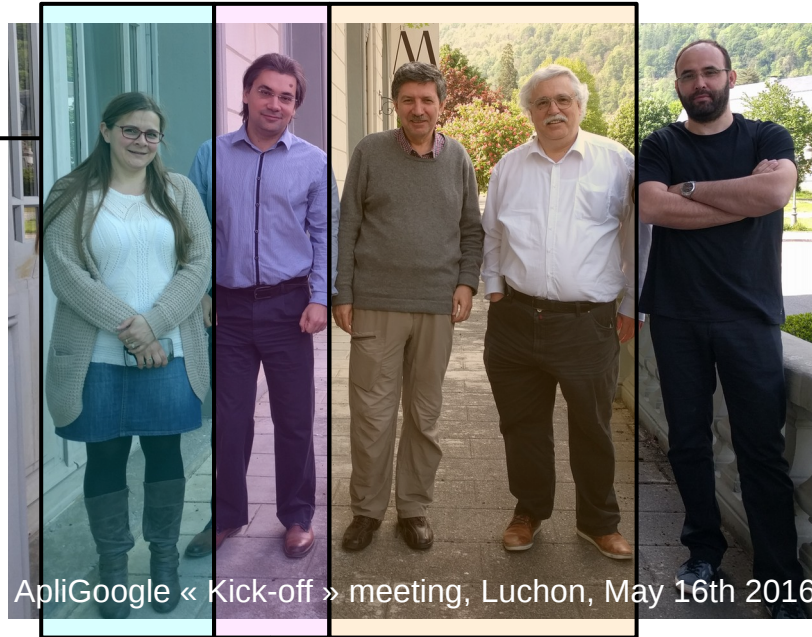
<http://www.quantware.ups-tlse.fr/dima>



ApliGoogle
Applications of Google matrix to directed networks and Big Data

Colloque de restitution 2017 - Défi MASTODONS CNRS – Jeudi 1er février 2018

ApliGoogle partners



ApliGoogle « Kick-off » meeting, Luchon, May 16th 2016

Institut de Recherche en Informatique de Toulouse (UMR CNRS 5152)

- MCF K. Jaffrès-Runser
- S. El Zant (doctorant)
- Dr. T. Peng (post-doctorant)

Expertises: **Wireless networks**



<http://www.irit.fr/~Katia.Jaffres>

Institut Curie (INSERM U900)

- Dr. A. Zinovyev
- Dr. I. Kuperstein
- Dr. L. Calzone
- U. Czerwinska (doctorante)

Expertises: **Computational Biology, Systems Biology of Cancer**



<http://www.ihes.fr/~zinovyev/>

ApliGoogle
Applications of Google matrix to directed networks and Big Data

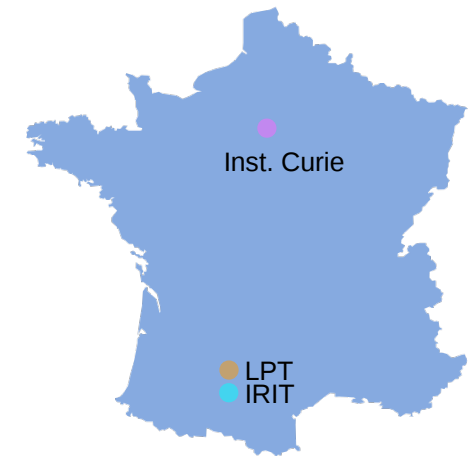
Laboratoire de Physique Théorique de Toulouse (UMR CNRS 5152)

- DR1 D. Shepelyansky (PI)
- Pr. K. Frahm

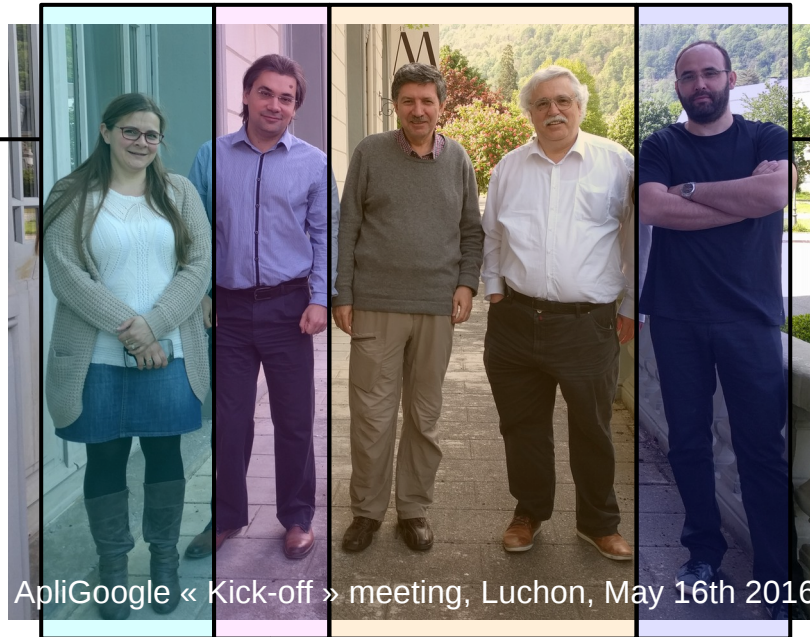
Expertises: **Quantum chaos, Random Matrix Theory, Complex directed networks**



<http://www.quantware.ups-tlse.fr/dima>



ApliGoogle partners



Institut de Recherche en Informatique de Toulouse (UMR CNRS 5152)

- MCF K. Jaffrès-Runser
- S. El Zant (doctorant)
- Dr. T. Peng (post-doctorant)

Expertises: **Wireless networks**



<http://www.irit.fr/~Katia.Jaffres>

Institut UTINAM (UMR CNRS 6213)

- MCF J. Lages
- C. Coquidé (doctorant)
- G. Rollin (post-doctorant)

Expertises: **Quantum chaos, Random Matrix Theory, Complex directed networks**



<http://perso.utinam.cnrs.fr/~lages>

Institut Curie (INSERM U900)

- Dr. A. Zinovyev
- Dr. I. Kuperstein
- Dr. L. Calzone
- U. Czerwinska (doctorante)

Expertises: **Computational Biology, Systems Biology of Cancer**



<http://www.ihes.fr/~zinovyev/>

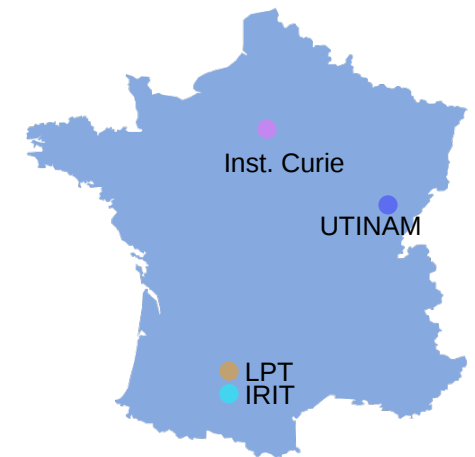
Laboratoire de Physique Théorique de Toulouse (UMR CNRS 5152)

- DR1 D. Shepelyansky (PI)
- Pr. K. Frahm

Expertises: **Quantum chaos, Random Matrix Theory, Complex directed networks**



<http://www.quantware.ups-tlse.fr/dima>



ApliGoogle

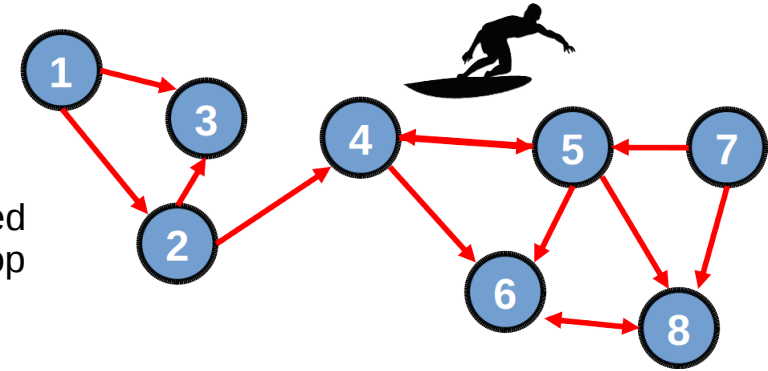
Applications of Google matrix to directed networks and Big Data

Main methods

How Google works

From Markov (1906) to Brin & Page (1998)

Markovian process : a random surfer probe the structure of a directed network. A each step, the surfer choose randomly an adjacent node to hop and continue its journey.



Adjacency matrix

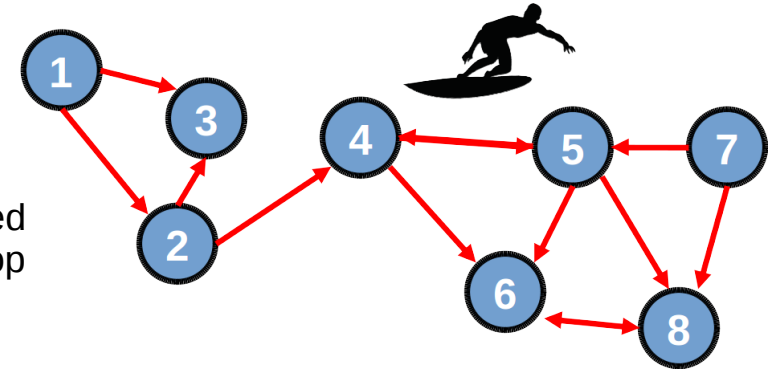
$$A_{ij} = \begin{cases} 1 & \text{si } j \rightarrow i \\ 0 & \text{si } j \not\rightarrow i \end{cases}$$

$$A = \begin{pmatrix} 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 1 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 1 & 1 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 & 1 & 0 & 0 & 0 \\ 0 & 0 & 0 & 1 & 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 1 & 1 & 0 & 0 & 1 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 1 & 1 & 1 & 0 \end{pmatrix}$$

How Google works

From Markov (1906) to Brin & Page (1998)

Markovian process : a random surfer probe the structure of a directed network. At each step, the surfer chooses randomly an adjacent node to hop and continue its journey.



Adjacency matrix

$$A_{ij} = \begin{cases} 1 & \text{si } j \rightarrow i \\ 0 & \text{si } j \nrightarrow i \end{cases}$$

Stochastic matrix

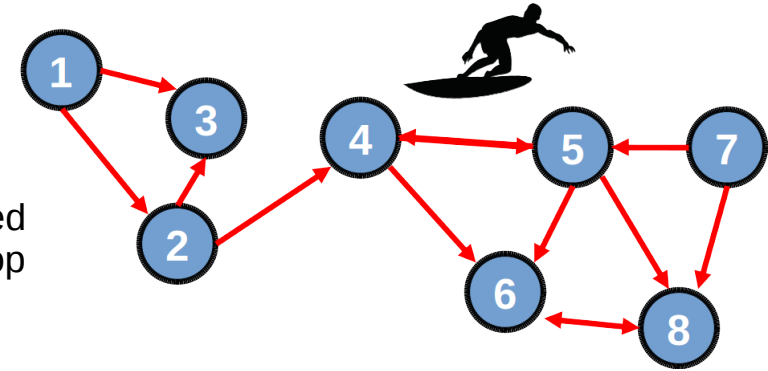
$$S_{ij} = \begin{cases} A_{ij} / \sum_{k=1}^N A_{kj} & \text{si } \sum_{k=1}^N A_{kj} \neq 0 \\ 1/N & \text{sinon} \end{cases}$$

$$S = \begin{pmatrix} 0 & 0 & 1/8 & 0 & 0 & 0 & 0 & 0 \\ 1/2 & 0 & 1/8 & 0 & 0 & 0 & 0 & 0 \\ 1/2 & 1/2 & 1/8 & 0 & 0 & 0 & 0 & 0 \\ 0 & 1/2 & 1/8 & 0 & 1/3 & 0 & 0 & 0 \\ 0 & 0 & 1/8 & 1/2 & 0 & 0 & 1/2 & 0 \\ 0 & 0 & 1/8 & 1/2 & 1/3 & 0 & 0 & 1 \\ 0 & 0 & 1/8 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 1/8 & 0 & 1/3 & 1 & 1/2 & 0 \end{pmatrix}$$

How Google works

From Markov (1906) to Brin & Page (1998)

Markovian process : a random surfer probe the structure of a directed network. A each step, the surfer choose randomly an adjacent node to hop and continue its journey.



Adjacency matrix

$$A_{ij} = \begin{cases} 1 & \text{si } j \rightarrow i \\ 0 & \text{si } j \nrightarrow i \end{cases}$$

Stochastic matrix

$$S_{ij} = \begin{cases} A_{ij} / \sum_{k=1}^N A_{kj} & \text{si } \sum_{k=1}^N A_{kj} \neq 0 \\ 1/N & \text{sinon} \end{cases}$$

Google matrix

$$G_{ij} = \alpha S_{ij} + (1 - \alpha) / N$$

avec $0.5 < \alpha < 1$

Perron-Frobenius operator

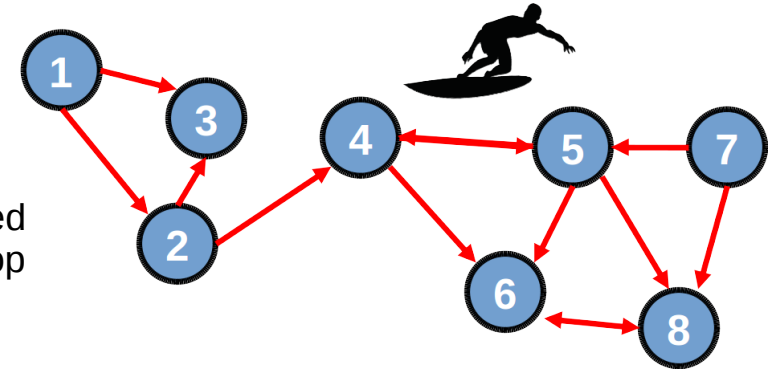
$$G = \begin{pmatrix} 1/40 & 1/40 & 1/8 & 1/40 & 1/40 & 1/40 & 1/40 & 1/40 \\ 17/40 & 1/40 & 1/8 & 1/40 & 1/40 & 1/40 & 1/40 & 1/40 \\ 17/40 & 17/40 & 1/8 & 1/40 & 1/40 & 1/40 & 1/40 & 1/40 \\ 1/40 & 17/40 & 1/8 & 1/40 & 7/24 & 1/40 & 1/40 & 1/40 \\ 1/40 & 1/40 & 1/8 & 17/40 & 1/40 & 1/40 & 17/40 & 1/40 \\ 1/40 & 1/40 & 1/8 & 17/40 & 7/24 & 1/40 & 1/40 & 33/40 \\ 1/40 & 1/40 & 1/8 & 1/40 & 1/40 & 1/40 & 1/40 & 1/40 \\ 1/40 & 1/40 & 1/8 & 1/40 & 7/24 & 33/40 & 17/40 & 1/40 \end{pmatrix}$$

$\alpha = 0.8$

How Google works

From Markov (1906) to Brin & Page (1998)

Markovian process : a random surfer probe the structure of a directed network. A each step, the surfer choose randomly an adjacent node to hop and continue its journey.



Adjacency matrix

$$A_{ij} = \begin{cases} 1 & \text{si } j \rightarrow i \\ 0 & \text{si } j \nrightarrow i \end{cases}$$

Stochastic matrix

$$S_{ij} = \begin{cases} A_{ij} / \sum_{k=1}^N A_{kj} & \text{si } \sum_{k=1}^N A_{kj} \neq 0 \\ 1/N & \text{sinon} \end{cases}$$

Google matrix

$$G_{ij} = \alpha S_{ij} + (1 - \alpha) / N$$

avec $0.5 < \alpha < 1$

Perron-Frobenius operator

PageRank vector

$$\mathbf{P} = \lim_{n \rightarrow \infty} \mathbf{P}^{(n)} = \lim_{n \rightarrow \infty} \mathbf{G}^n \mathbf{P}^{(0)}$$

$P_i^{(n)}$ is the probability that random surfer arrives at node i at the n th step.

\mathbf{P} is the \mathbf{G} matrix eigenvector associated with eigenvalue 1

$$\mathbf{P} = \mathbf{G}\mathbf{P}$$

$$\mathbf{P} = \begin{pmatrix} 0.03109452568730597 \\ 0.04353233614756617 \\ 0.06094527086606558 \\ 0.06729412361797826 \\ 0.07044998599586171 \\ \mathbf{0.35181679356094489} \\ 0.03109452568730597 \\ 0.34377243843697143 \end{pmatrix}$$

Distribution $P(K)$

where K is the rank index:

$$P(1) = \mathbf{0.35181679356094489}$$

$$P(2) = 0.34377243843697143$$

$$P(3) = 0.07044998599586171$$

$$P(4) = 0.06729412361797826$$

$$P(5) = 0.06094527086606558$$

$$P(6) = 0.04353233614756617$$

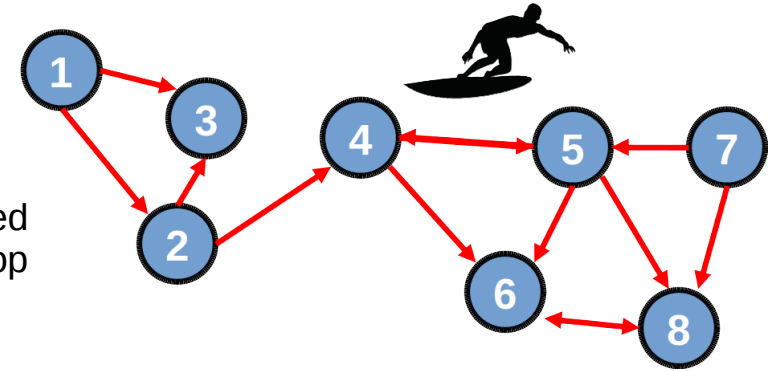
$$P(7) = P(8) = 0.03109452568730597$$



How Google works

From Markov (1906) to Brin & Page (1998)

Markovian process : a random surfer probe the structure of a directed network. At each step, the surfer choose randomly an adjacent node to hop and continue its journey.



Adjacency matrix

$$A_{ij} = \begin{cases} 1 & \text{si } j \rightarrow i \\ 0 & \text{si } j \nrightarrow i \end{cases}$$

Stochastic matrix

$$S_{ij} = \begin{cases} A_{ij} / \sum_{k=1}^N A_{kj} & \text{si } \sum_{k=1}^N A_{kj} \neq 0 \\ 1/N & \text{sinon} \end{cases}$$

Google matrix

$$G_{ij} = \alpha S_{ij} + (1 - \alpha) / N$$

avec $0.5 < \alpha < 1$

Perron-Frobenius operator

PageRank vector

$$\mathbf{P} = \lim_{n \rightarrow \infty} \mathbf{P}^{(n)} = \lim_{n \rightarrow \infty} \mathbf{G}^n \mathbf{P}^{(0)}$$

$P_i^{(n)}$ is the probability that random surfer arrives at node i at the n th step.

\mathbf{P} is the \mathbf{G} matrix eigenvector associated with eigenvalue 1

$$\mathbf{P} = \mathbf{G}\mathbf{P}$$

The most important node is the one with the highest probability.

“Recursive definition”: the more a node is pointed by important nodes, the more it is important.

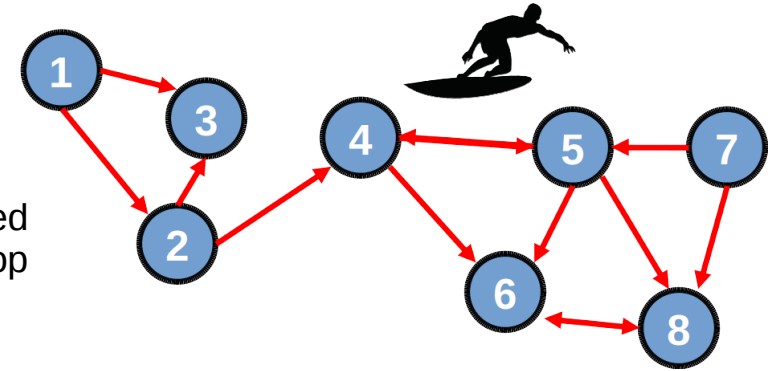
PageRank measures the influence of a node.

PageRank is (was?) at the heart of **Google** search engine (Brin, Page '98).

How Google works

From Markov (1906) to Brin & Page (1998)

Markovian process : a random surfer probe the structure of a directed network. At each step, the surfer choose randomly an adjacent node to hop and continue its journey.



Adjacency matrix

$$A_{ij} = \begin{cases} 1 & \text{si } j \rightarrow i \\ 0 & \text{si } j \nrightarrow i \end{cases}$$

Stochastic matrix

$$S_{ij} = \begin{cases} A_{ij} / \sum_{k=1}^N A_{kj} & \text{si } \sum_{k=1}^N A_{kj} \neq 0 \\ 1/N & \text{sinon} \end{cases}$$

Google matrix

$$G_{ij} = \alpha S_{ij} + (1 - \alpha)/N$$

avec $0.5 < \alpha < 1$

Perron-Frobenius operator

PageRank vector

$$\mathbf{P} = \lim_{n \rightarrow \infty} \mathbf{P}^{(n)} = \lim_{n \rightarrow \infty} \mathbf{G}^n \mathbf{P}^{(0)}$$

$P_i^{(n)}$ is the probability that random surfer arrives at node i at the n th step.

\mathbf{P} is the \mathbf{G} matrix eigenvector associated with eigenvalue 1

$$\mathbf{P} = \mathbf{G}\mathbf{P}$$

The most important node is the one with the highest probability.

“Recursive definition”: the more a node is pointed by important nodes, the more it is important.

PageRank measures the influence of a node.

PageRank is (was?) at the heart of **Google** search engine (Brin, Page ‘98).

CheiRank vector $\mathbf{P}^* = \mathbf{G}^* \mathbf{P}^*$

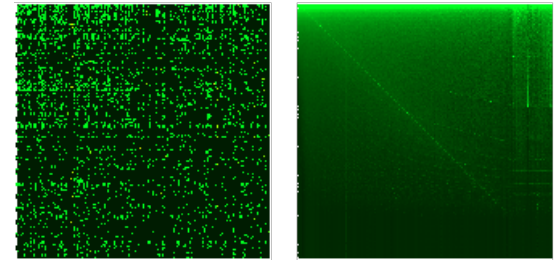
Similar to PageRank of the inverted network. With inverted adjacency matrix elements $A_{ij}^* = A_{ji}$, it is possible to define the stochastic matrix elements $S_{ij}^* \neq S_{ji}$, and the Google matrix elements $G_{ij}^* \neq G_{ji}$ associated to the inverted network (Fogaras ‘03, Chepelianskii ‘10).

“Recursive definition”: the more a node pointed toward important nodes, the more it is important.

CheiRank measures the diffusion/the communication of a node.

Colloque de restitution 2017 - Défi MASTODONS CNRS – Jeudi 1er février 2018

Real directed complex networks



Google matrix G of the English Wikipedia network (Aug. '09)
N=3 282 257

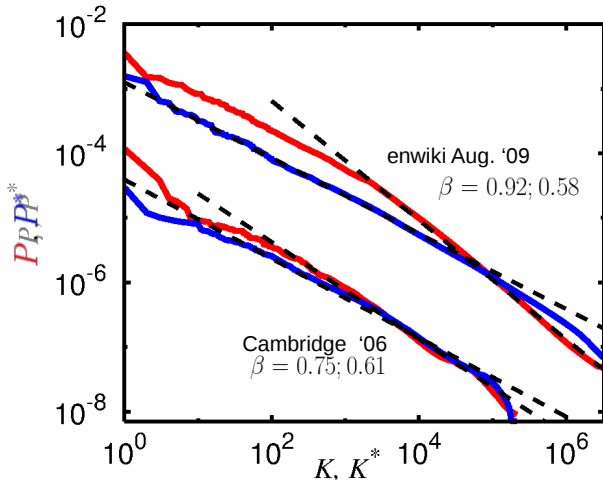
Properties of real networks (WWW, social networks, ...)

- **“Small world” property:** average distance between two nodes $\sim \log N$
- **Scale-free property:** distribution of the number of ingoing or outgoing links is $\rho(k) \sim k^{-\nu}$

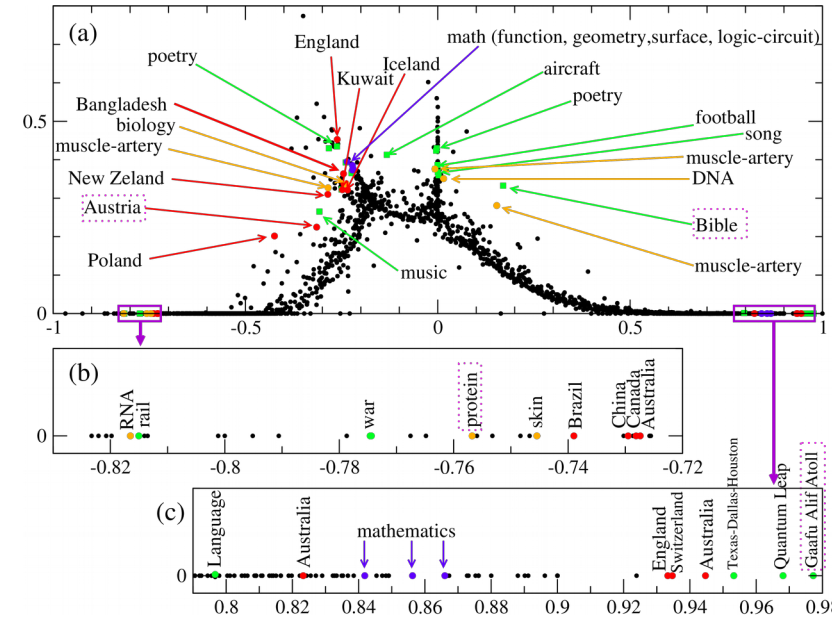
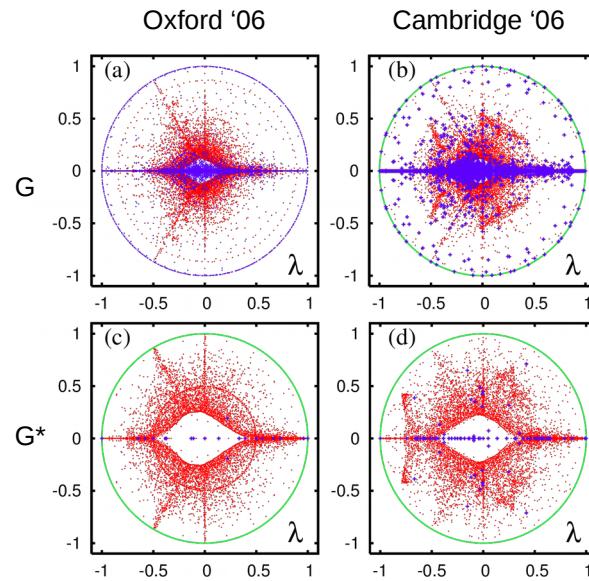
PageRank distribution properties

$$P(K) \sim 1/K^\beta \text{ with } \nu_{\text{in}} = 1 + 1/\beta$$

$$P^*(K^*) \sim 1/K^{*\beta} \text{ with } \nu_{\text{out}} = 1 + 1/\beta$$



Google matrix spectrum



Wikipedia english '13 network

Theory of real directed complex networks

Random Matrix Theory introduced by Wigner '67 describes universal spectral properties shared by complex nuclei/atoms/molecules and also mesoscopic and quantum chaos systems (Hermitean and unitary matrices statistical ensembles).

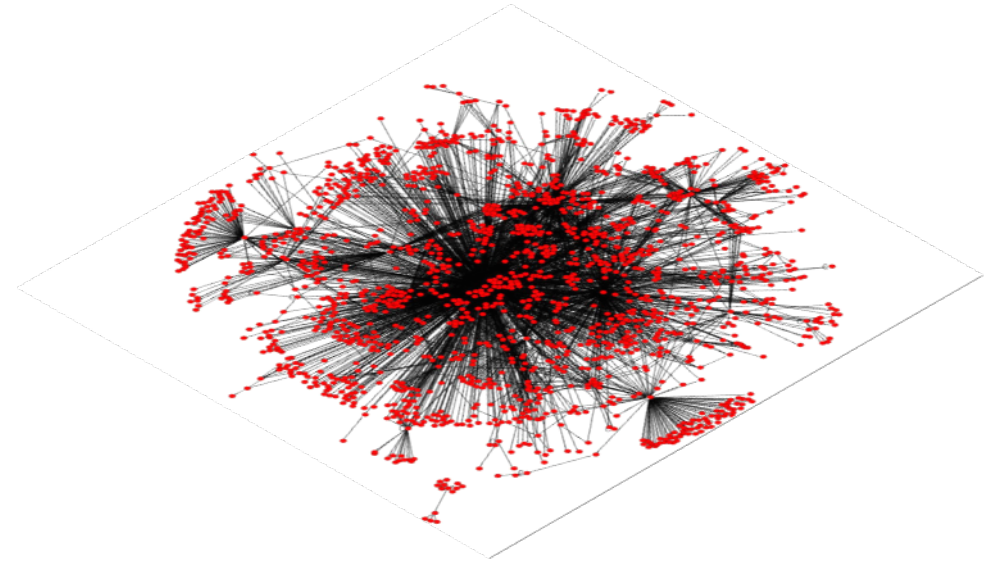
Challenge : A Random Matrix Theory for Markov chains and Google matrix ensembles is still lacking ➔ We need more examples / more applications

Figures from :
Ermann, Frahm, Shepelyansky (2016), Scholarpedia, 11(11):30944

Ermann, Frahm, Shepelyansky (2015), Rev. Mod. Phys. 87, 1261.

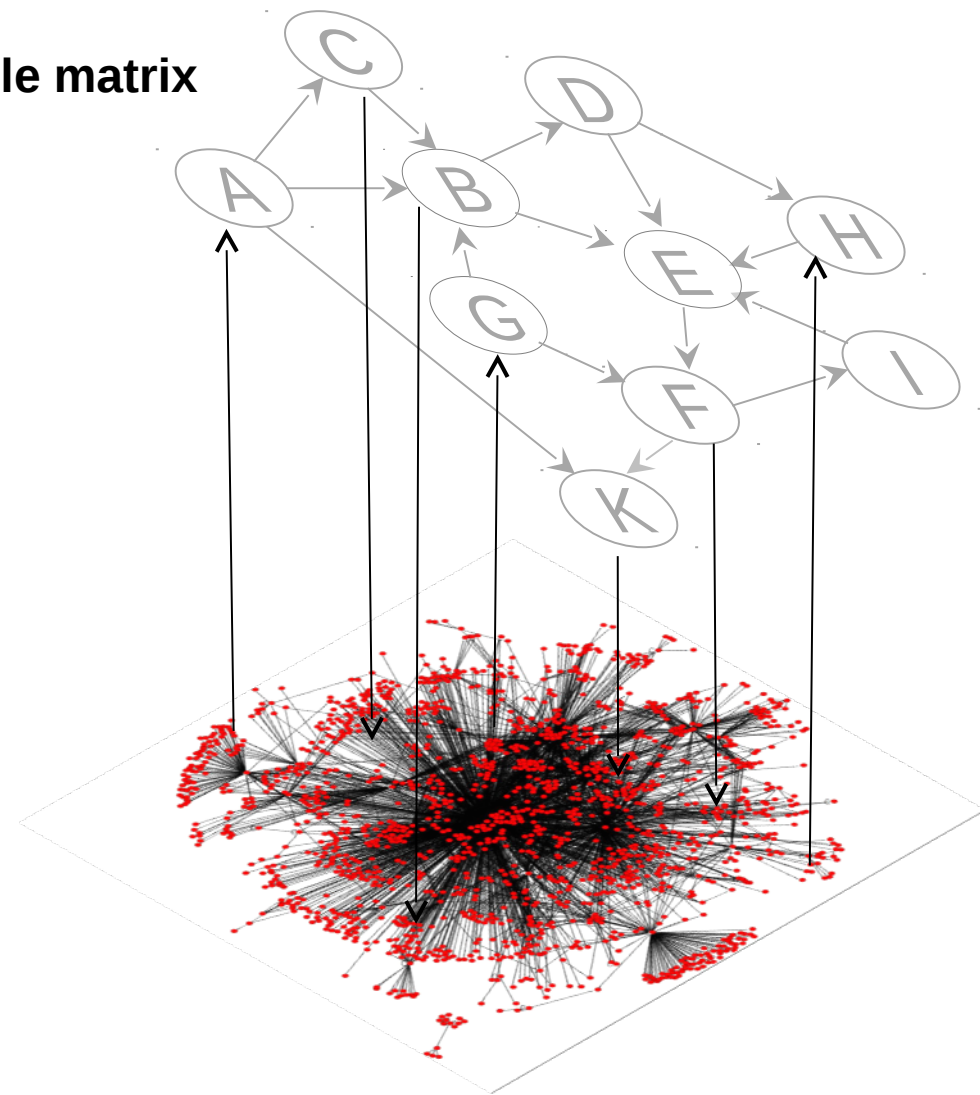
Reduced Google matrix

Consider a network with $N \gg 1$ nodes.



Reduced Google matrix

Consider a network with $N \gg 1$ nodes.
Consider a sub-network (a community) of $N_r \ll N$ nodes.



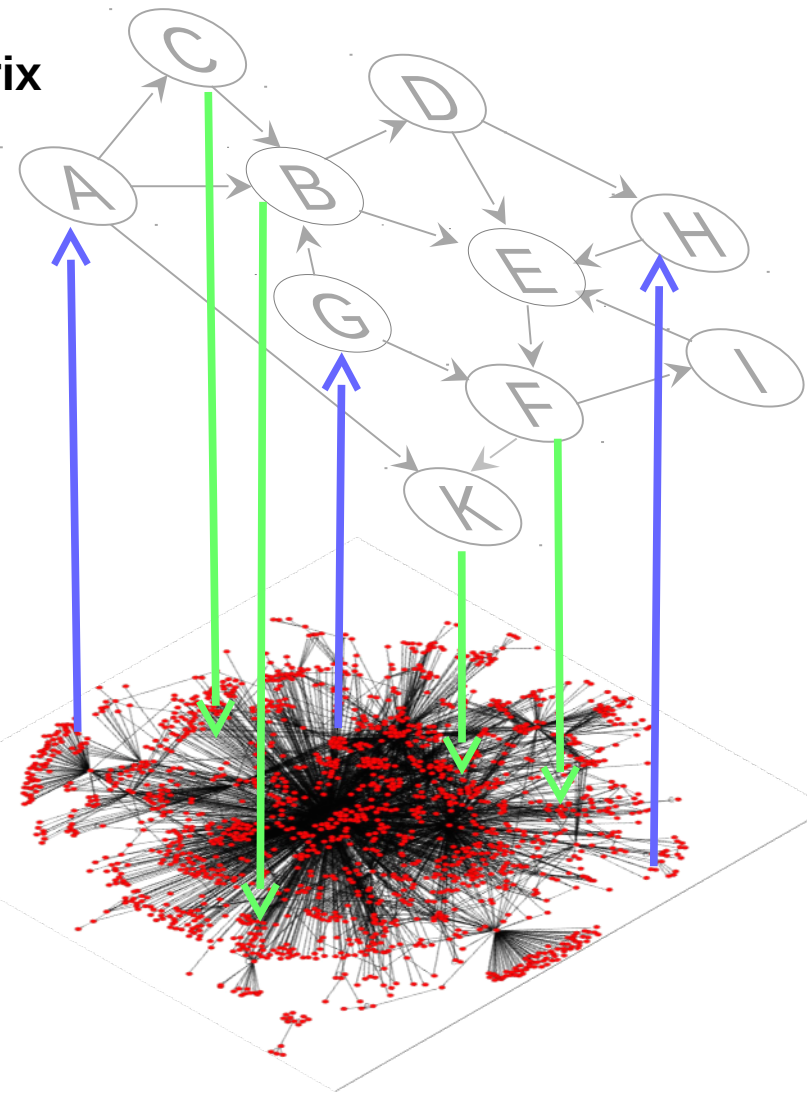
Reduced Google matrix

Consider a network with $N \gg 1$ nodes.
Consider a sub-network (a community) of $N_r \ll N$ nodes.
The Google matrix of the size N network and the associated
PageRank vector can be written as

$$\mathbf{G} = \begin{pmatrix} \mathbf{G}_{rr} & \mathbf{G}_{rs} \\ \mathbf{G}_{sr} & \mathbf{G}_{ss} \end{pmatrix},$$

$$\mathbf{P} = \begin{pmatrix} \mathbf{P}_r \\ \mathbf{P}_s \end{pmatrix}$$

$$\mathbf{G}\mathbf{P} = \mathbf{P}$$



Reduced Google matrix

Consider a network with $N \gg 1$ nodes.

Consider a sub-network (a community) of $N_r \ll N$ nodes.

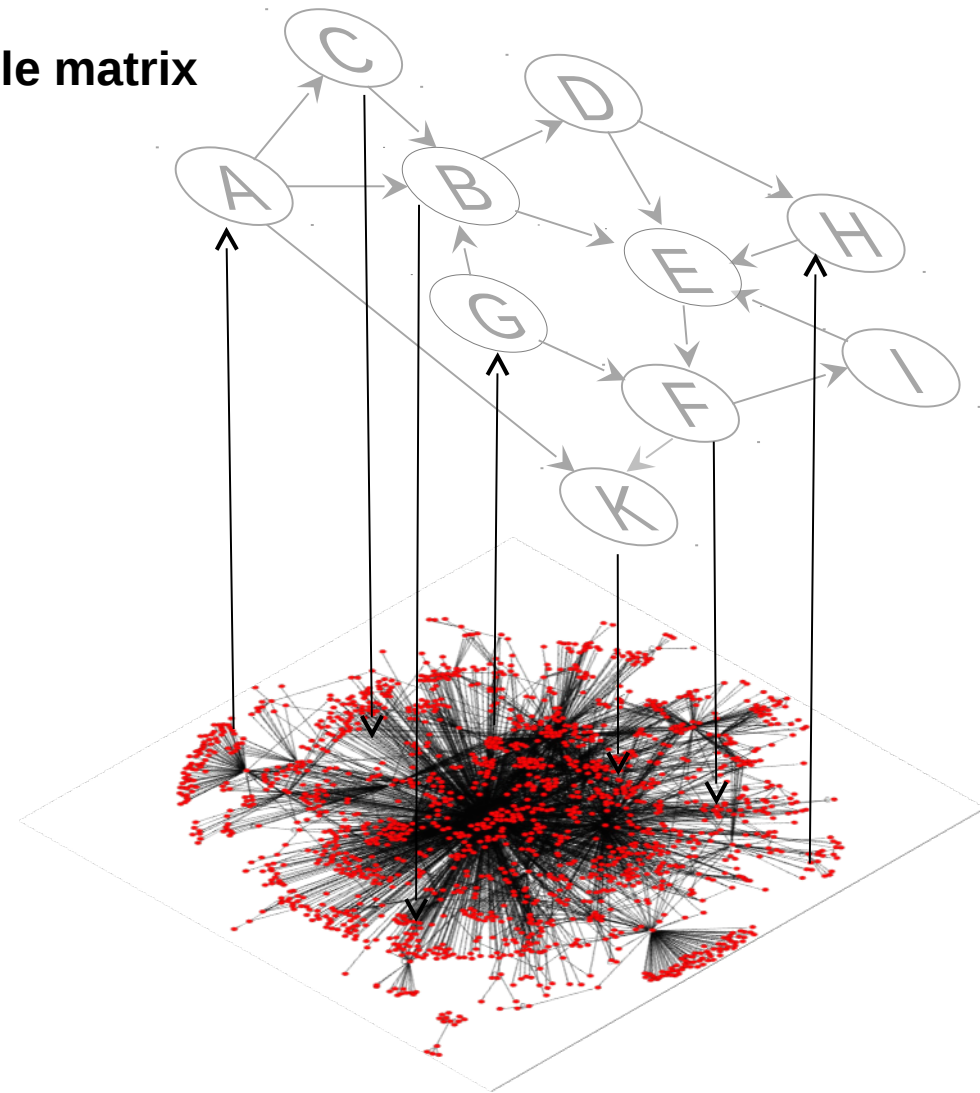
The Google matrix of the size N network and the associated PageRank vector can be written as

$$\mathbf{G} = \begin{pmatrix} \mathbf{G}_{rr} & \mathbf{G}_{rs} \\ \mathbf{G}_{sr} & \mathbf{G}_{ss} \end{pmatrix}, \quad \mathbf{P} = \begin{pmatrix} \mathbf{P}_r \\ \mathbf{P}_s \end{pmatrix}$$

$$\mathbf{G}\mathbf{P} = \mathbf{P}$$

We define the reduced Google matrix \mathbf{G}_R associated to the community of size N_r such as

$$\mathbf{G}_R \mathbf{P}_r = \mathbf{P}_r$$



Reduced Google matrix

Consider a network with $N \gg 1$ nodes.
 Consider a sub-network (a community) of $N_r \ll N$ nodes.
 The Google matrix of the size N network and the associated PageRank vector can be written as

$$\mathbf{G} = \begin{pmatrix} \mathbf{G}_{rr} & \mathbf{G}_{rs} \\ \mathbf{G}_{sr} & \mathbf{G}_{ss} \end{pmatrix}, \quad \mathbf{P} = \begin{pmatrix} \mathbf{P}_r \\ \mathbf{P}_s \end{pmatrix}$$

$$\mathbf{G}\mathbf{P} = \mathbf{P}$$

We define the reduced Google matrix \mathbf{G}_R associated to the community of size N_r such as

$$\mathbf{G}_R \mathbf{P}_r = \mathbf{P}_r$$

The reduced Google matrix can be written

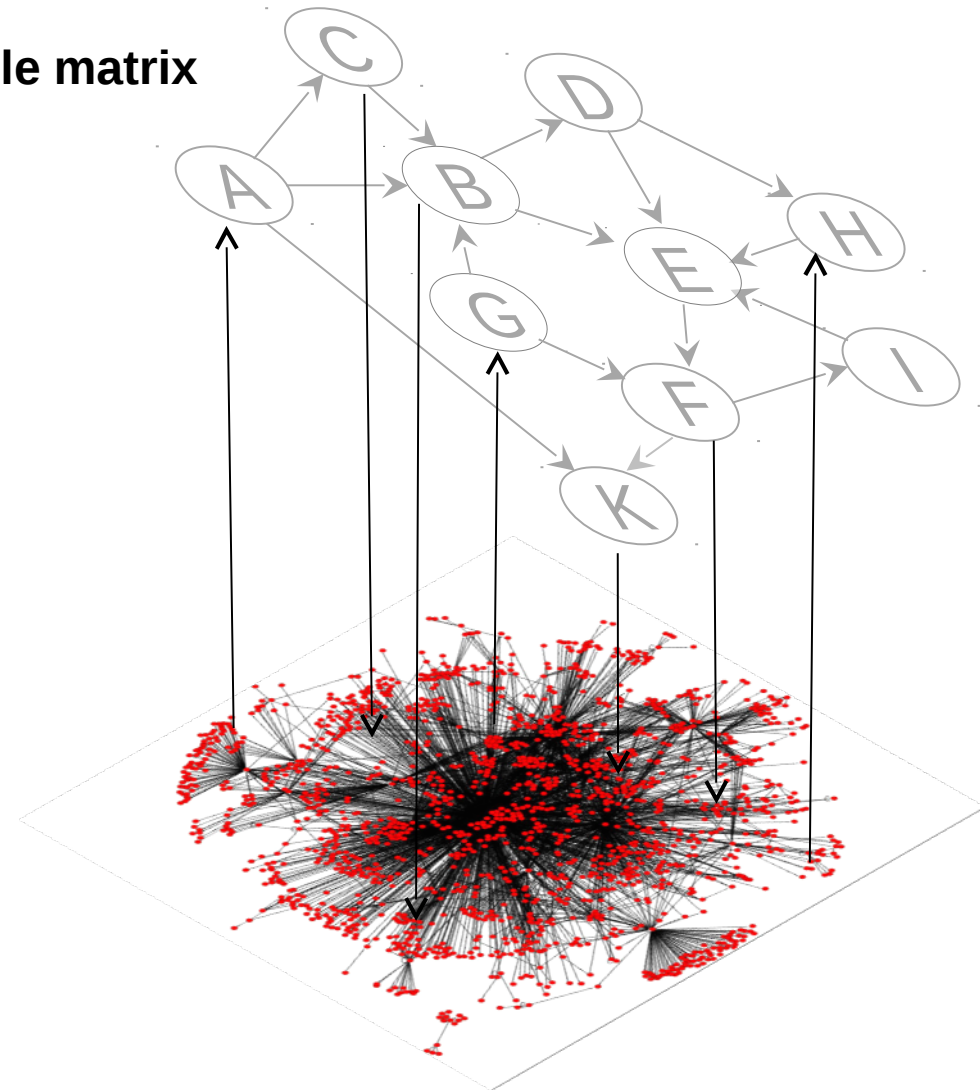
$$\mathbf{G}_R = \mathbf{G}_{rr} + \mathbf{G}_{rs} (\mathbf{1} - \mathbf{G}_{ss})^{-1} \mathbf{G}_{sr}$$

Contribution
from direct
links

Contribution from
indirect links
(scattering term)

Very slow
convergence since
the eigenvalue λ_c
of $\mathbf{G}_{ss} \sim \mathbf{G}$ is
very close to 1.

$$(\mathbf{1} - \mathbf{G}_{ss})^{-1} = \sum_{l=0}^{\infty} \mathbf{G}_{ss}^l$$



J. Lages, D. Shepelyansky, A. Zinovyev, PLoS ONE 13(1): e0190812 (2018)
 K. M. Frahm, and D. L. Shepelyansky, arXiv:1602.02394 [physics.soc-ph]

Reduced Google matrix

Consider a network with $N \gg 1$ nodes.
 Consider a sub-network (a community) of $N_r \ll N$ nodes.
 The Google matrix of the size N network and the associated PageRank vector can be written as

$$\mathbf{G} = \begin{pmatrix} \mathbf{G}_{rr} & \mathbf{G}_{rs} \\ \mathbf{G}_{sr} & \mathbf{G}_{ss} \end{pmatrix}, \quad \mathbf{P} = \begin{pmatrix} \mathbf{P}_r \\ \mathbf{P}_s \end{pmatrix}$$

$$\mathbf{G}\mathbf{P} = \mathbf{P}$$

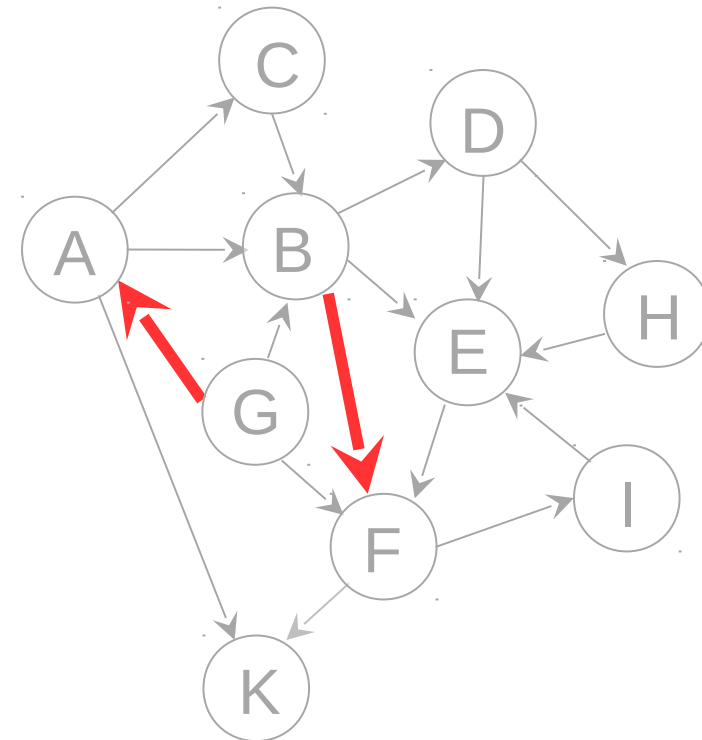
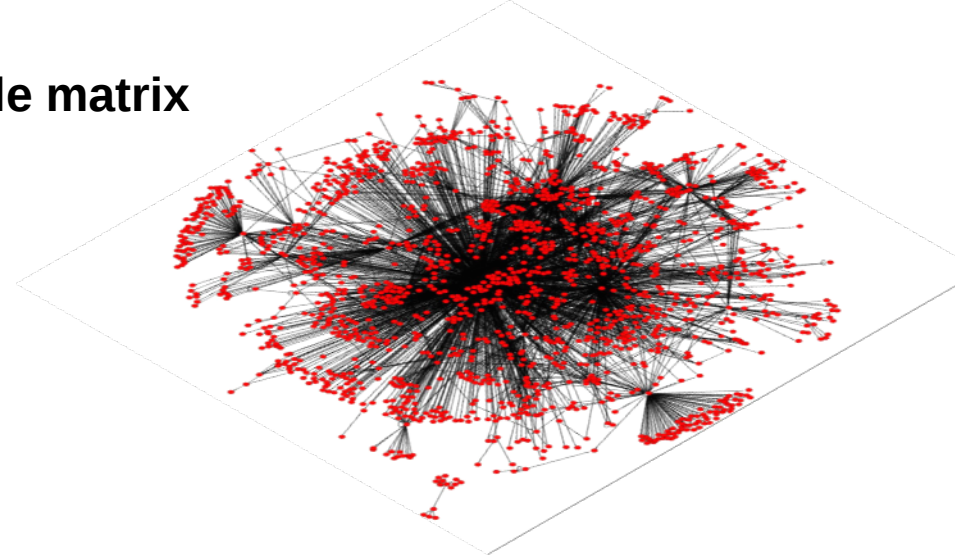
We define the reduced Google matrix \mathbf{G}_R associated to the community of size N_r such as

$$\mathbf{G}_R \mathbf{P}_r = \mathbf{P}_r$$

The reduced Google matrix can be written

$$\mathbf{G}_R = \mathbf{G}_{rr} + \mathbf{G}_{pr} + \mathbf{G}_{qr}$$

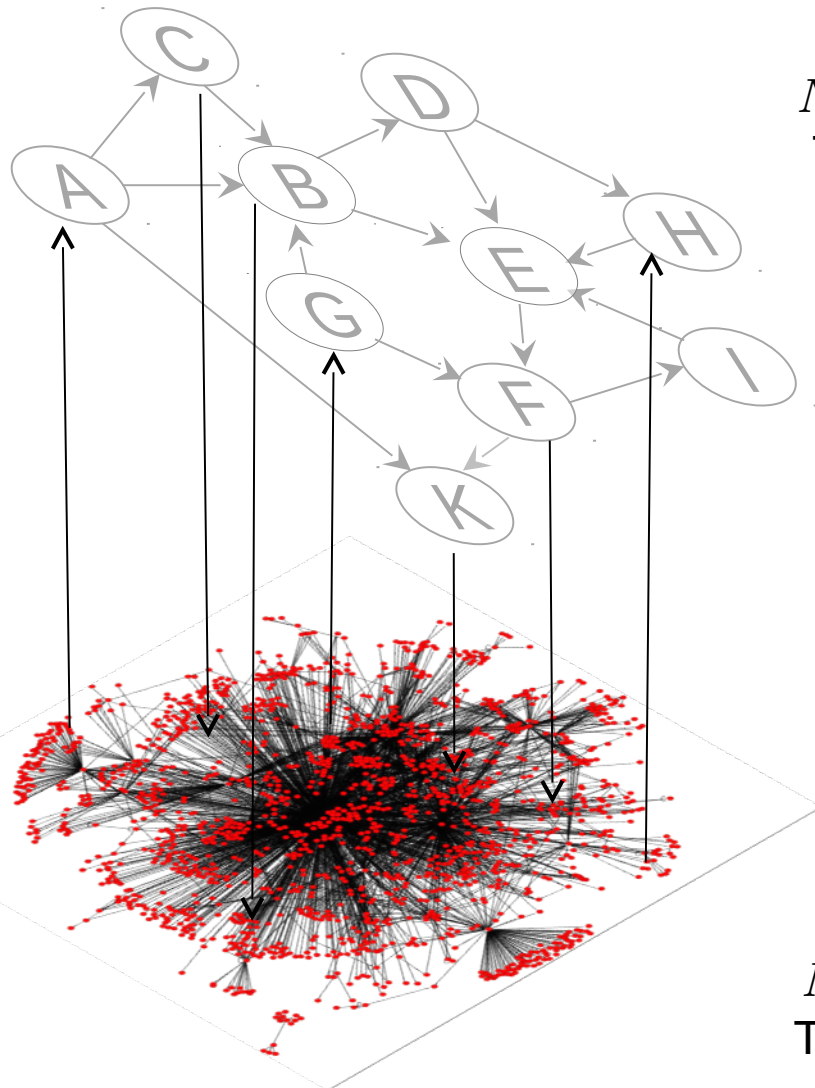
↓ Contribution from direct links
 ↓ Contribution from « PageRank »
 ↓ Contribution from hidden links



Highlights: major results obtained in 2017

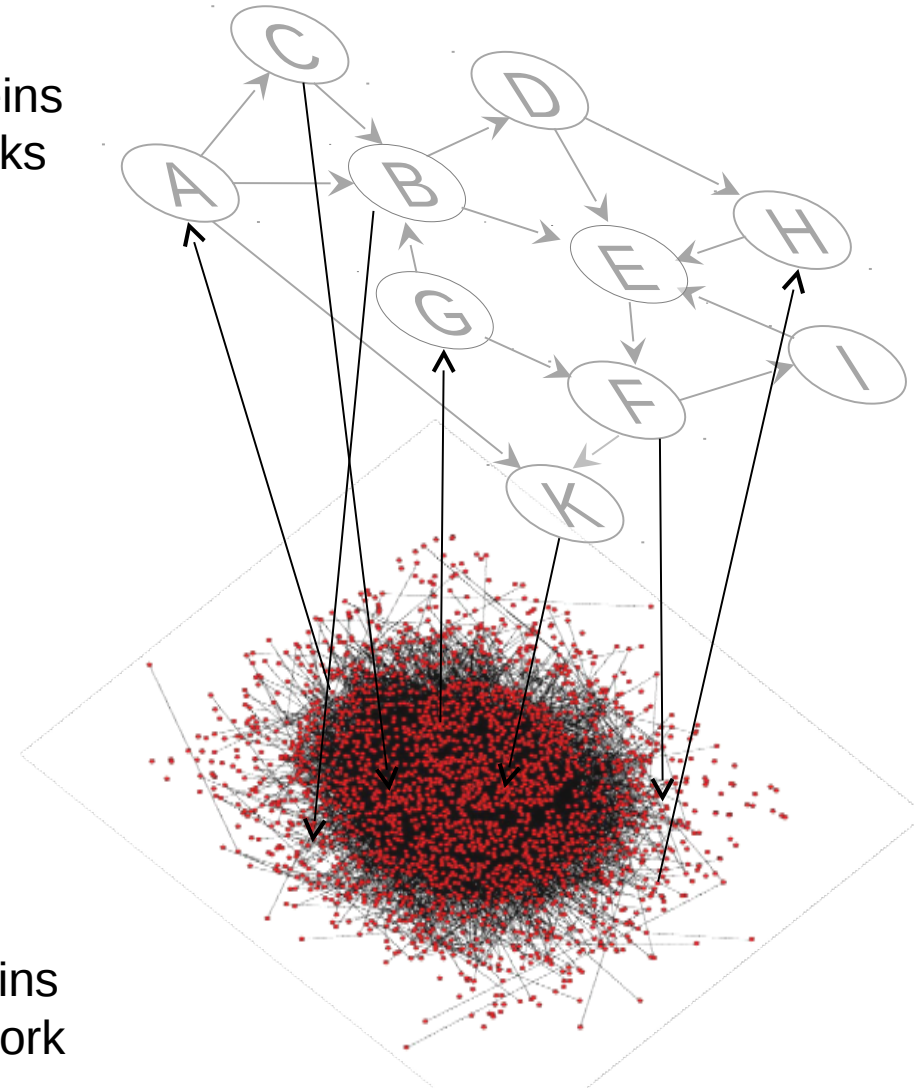
Googlomics

Inferring hidden causal relations between proteins



TRN1 ("normal")
Normal B-Lymphocytes

$N_r < 100$ proteins
The subnetworks



TRN2 ("cancer")
Leukemia cell line

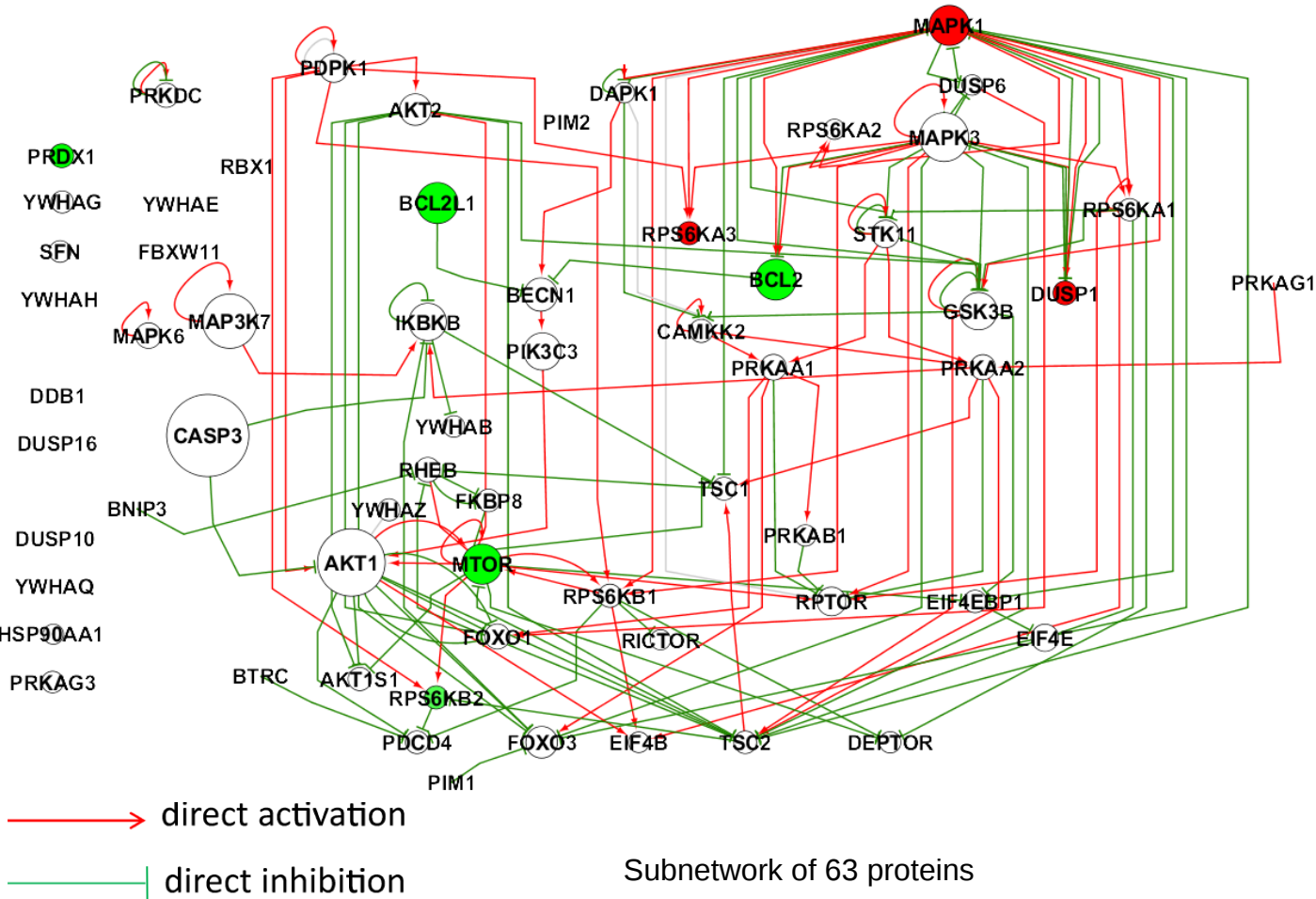
$N = 2432$ proteins
The whole network

J. Lages, D. Shepelyansky, A. Zinovyev, PLoS ONE 13(1): e0190812 (2018)

Googlomics

Inferring hidden causal relations between proteins

Inferring indirect (hidden) causal connections between AKT-mTOR pathway members

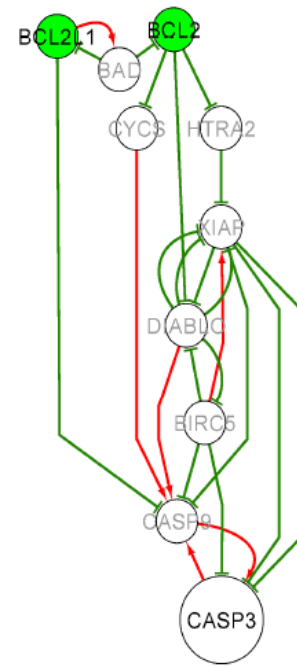
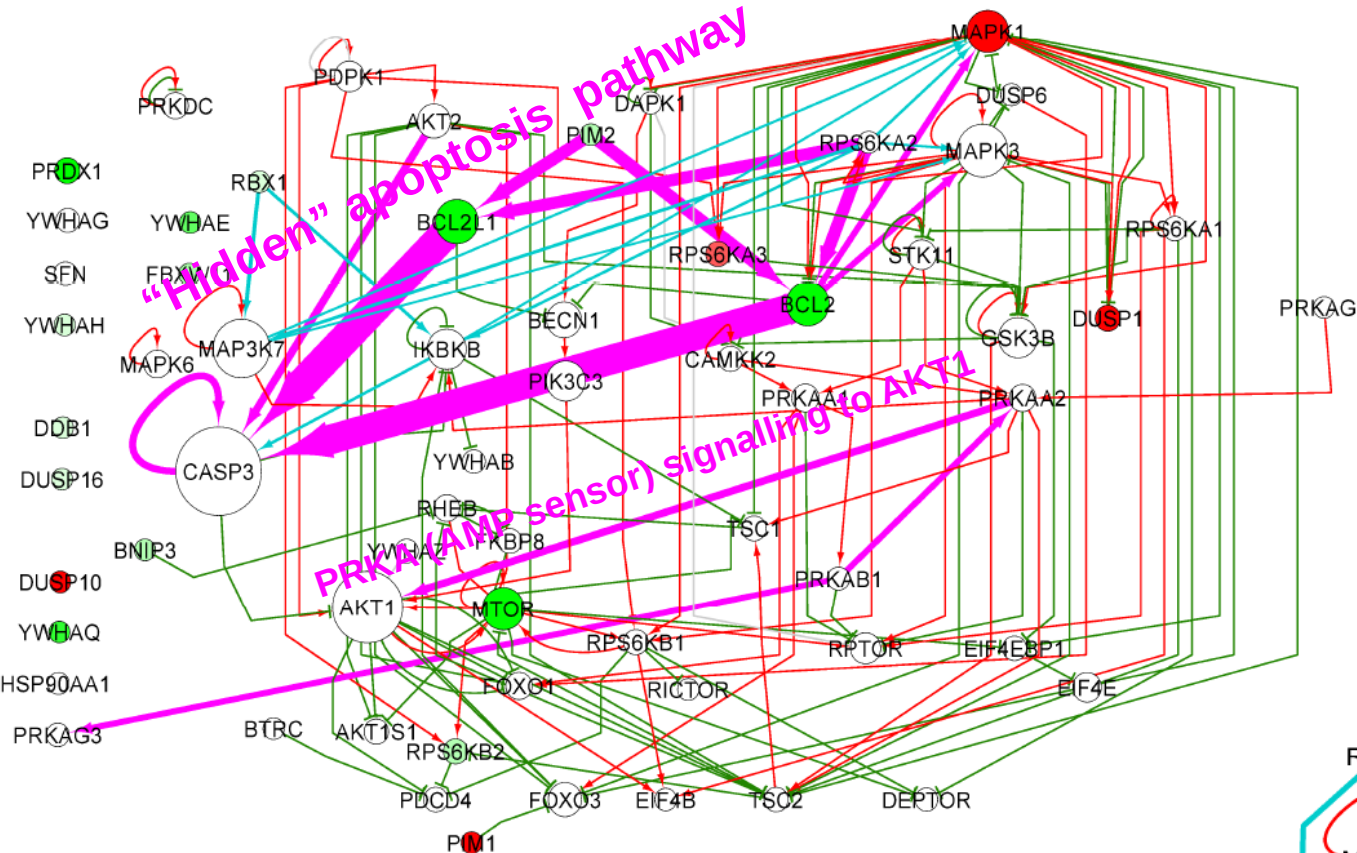


J. Lages, D. Shepelyansky, A. Zinovyev, PLoS ONE 13(1): e0190812 (2018)

Googlomics

Inferring hidden causal relations between proteins

Inferring indirect (hidden) causal connections between AKT-mTOR pathway members

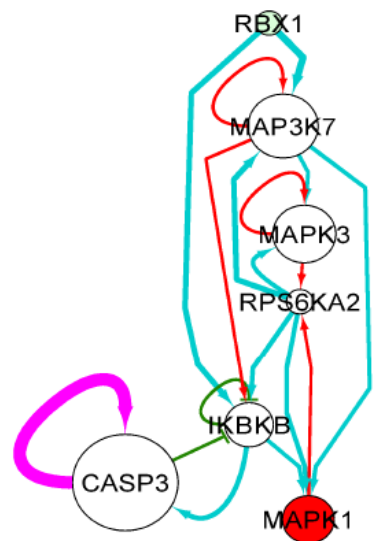


- direct activation
- | direct inhibition
- hidden (indirect)

Subnetwork of 63 proteins

hidden emergent oncogenic

Emergent oncogenic signaling between RBX1 (cell cycle protein degradation proteasome) and MAPK1

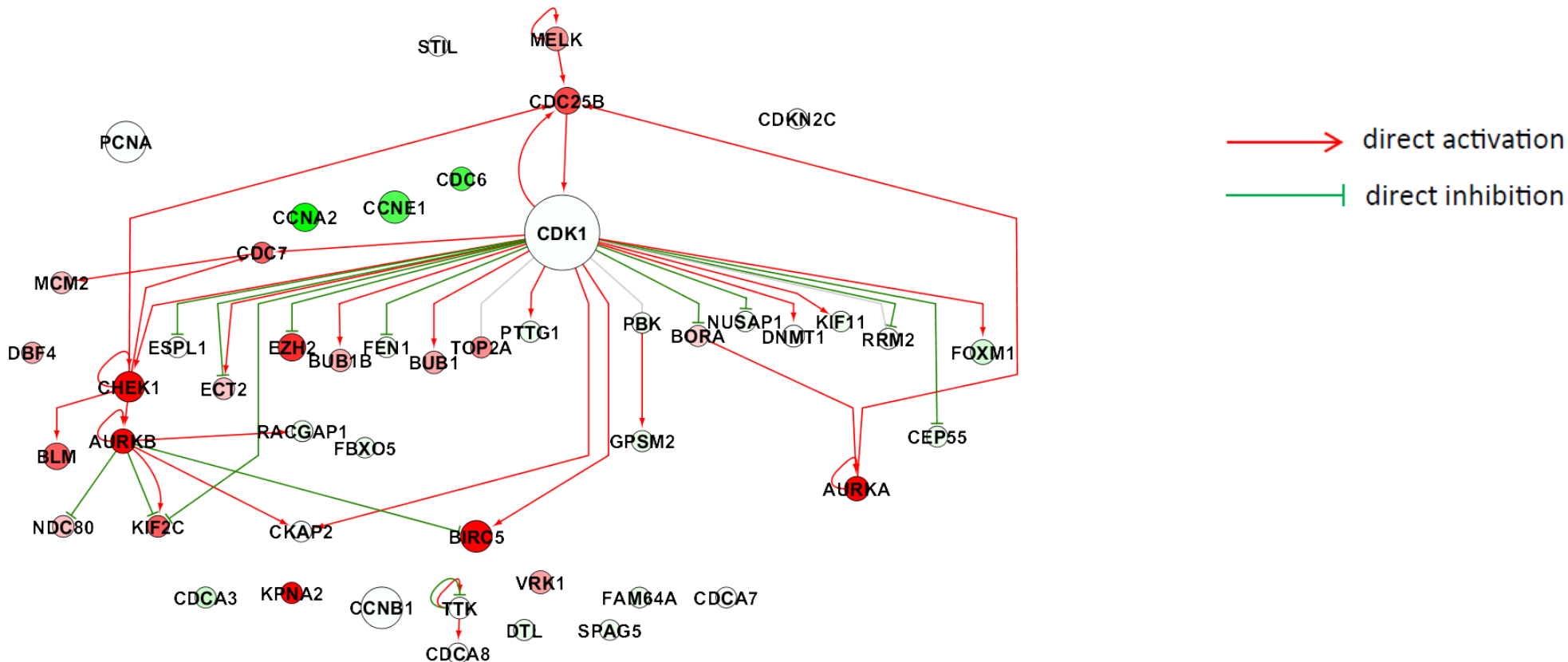


J. Lages, D. Shepelyansky, A. Zinovyev, PLoS ONE 13(1): e0190812 (2018)

Googlomics

Inferring hidden causal relations between proteins

Genes of a proliferative signature resulted from pancancer transcriptomic analysis



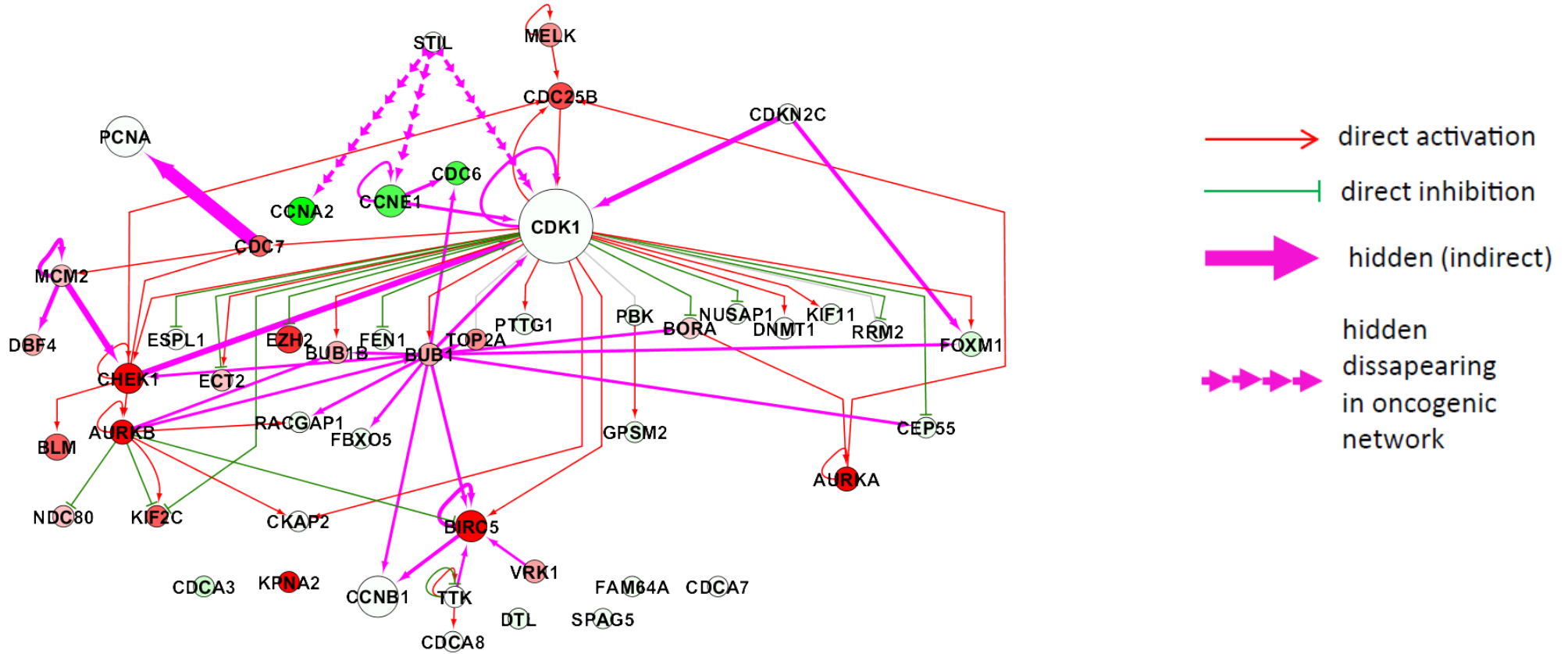
Subnetwork of 49 proteins

J. Lages, D. Shepelyansky, A. Zinovyev, PLoS ONE 13(1): e0190812 (2018)

Googlomics

Inferring hidden causal relations between proteins

Genes of a proliferative signature resulted from pancancer transcriptomic analysis



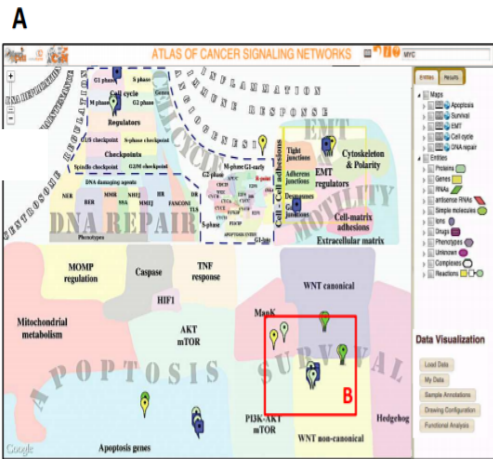
Subnetwork of 49 proteins

- More genes are connected into the network
- Emergence of a new “hidden” hub BUB1
- Connection to PCNA (DNA replication and DNA repair)
- Many cell cycle proteins improves in PageRank (AURK)
- Connection between STIL (mitotic spindle checkpoint regulator) and CCNA2, CCNE1

J. Lages, D. Shepelyansky, A. Zinovyev, PLoS ONE 13(1): e0190812 (2018)

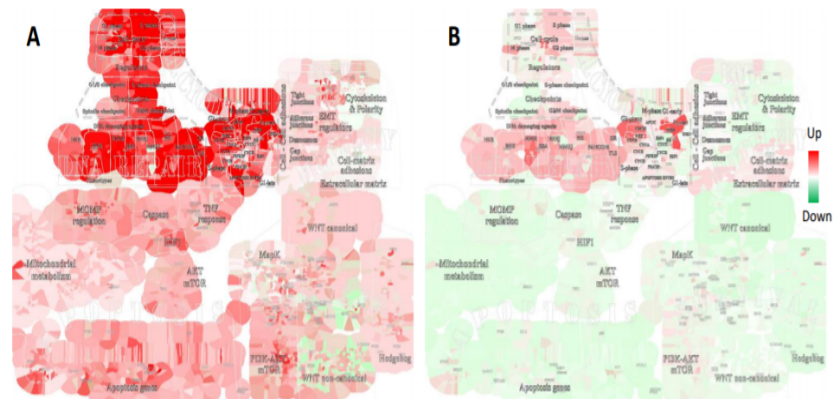
Atlas of Cancer Signaling Network

Atlas of Cancer Signaling Network pathway database and its application in pre-clinical studies (Monraz Gomez et al, <http://dx.doi.org/10.1101/234823> (2017), submitted to Briefings in Bioinformatics)



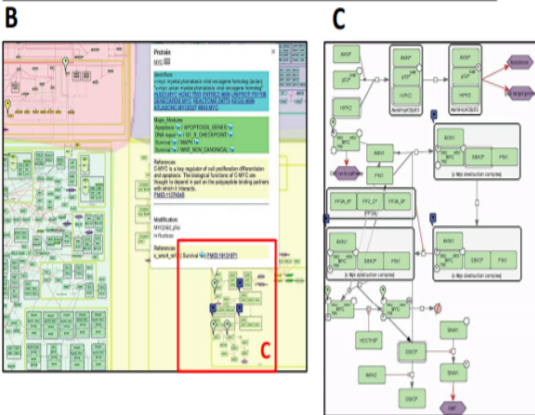
Network propagation methods applied to ACSN allows :

Visualizing molecular omics data for different cancer subtypes (A vs B)



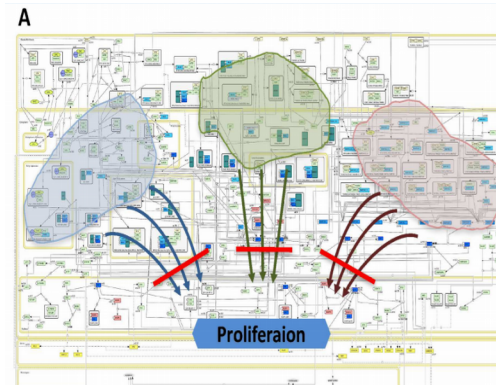
Red -upregulated modules, green - downregulated

Designing therapeutic interventions to disrupt malignant cancer signaling



<http://acsn.curie.fr>

ACSN - pathway database created by Institut Curie, large reaction network describing processes deregulated in cancer



WikiProteins project

Inferring directed network of proteins from Wikipedia using reduced Google Matrix

(ongoing work...)

~10000 wiki pages devoted to proteins

5009 proteins with described interactions,
16468 direct connections

The general properties of extracted network
is similar to manually curated protein-
protein interaction networks

This network is embedded in the global
Wikipedia network

The rest of the Wikipedia defines a
context of hyperlinks (model of external
world)

Reduced Google matrix allows finding
“hidden” functional interactions
between proteins

FLNA

From Wikipedia, the free encyclopedia

For the armed group in northern Mali, see National Liberation Front of Azawad.

Filamin A, alpha (FLNA) is a protein that in humans is encoded by the *FLNA* gene.^{[R][C]}

Contents [hide]

- Function
- Structure
- Interactions
- RNA editing
 - Type
 - Site
 - Structure
 - Function
- References
- Further reading
- External links

Function [edit]

Actin-binding protein, or filamin, is a 280-kD protein that crosslinks actin filaments into orthogonal networks in cortical cytoplasm and participates in the anchoring of membrane proteins for the actin cytoskeleton. Remodeling of the cytoskeleton is central to the modulation of cell shape and migration. Filamin A, encoded by the *FLNA* gene, is a widely expressed protein that regulates reorganization of the actin cytoskeleton by interacting with integrins, transmembrane receptor complexes, and second messengers [supplied by OMIM]^[?]

Structure [edit]

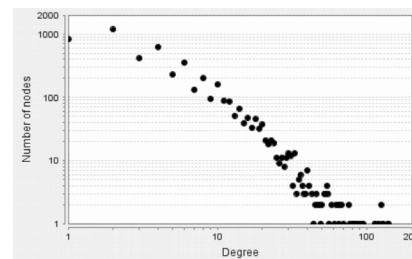
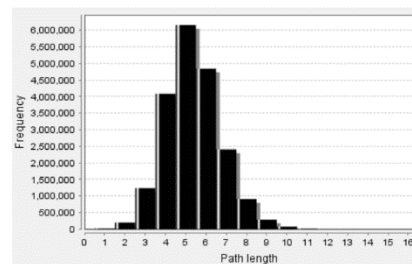
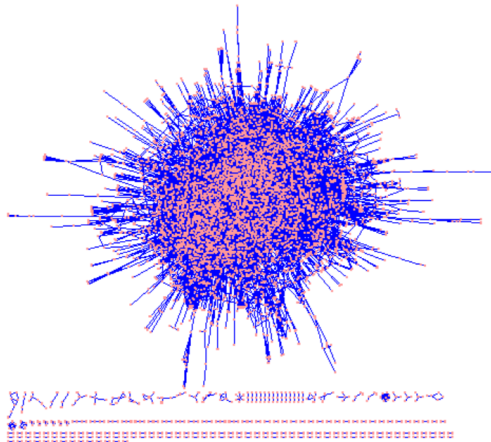
The protein structure includes an actin binding N terminal domain, 24 internal repeats and 2 hinge regions.^{[R][C]}

Interactions [edit]

Filamin has been shown to interact with:

- BRCA2^[18]
- CD29^{[13][4]}
- CASR^{[13][4]}
- FBLN1^[15]
- FILIP1^[16]
- FLNB^[17]
- TRIO^[21] and
- RALA^[19]
- VHL^{[22][23]}
- NHP1^[14]
- RALGAPB^[19]
- SH2B3^[20]

RNA editing [edit]



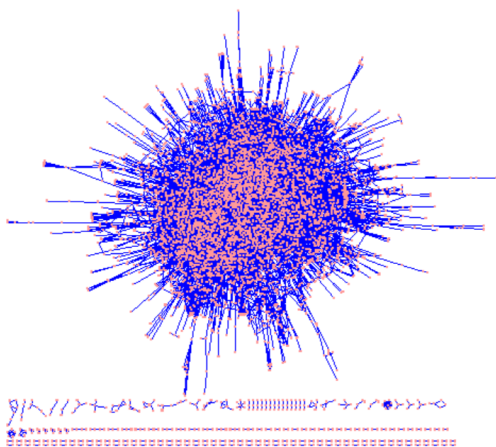
WikiProteins project

Inferring directed network of proteins from Wikipedia using reduced Google Matrix

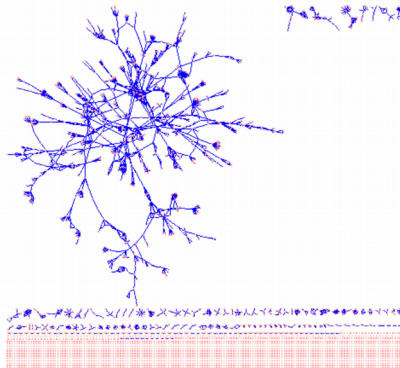
(ongoing work...)

Comparing direct and hidden networks of protein connections

Direct connections

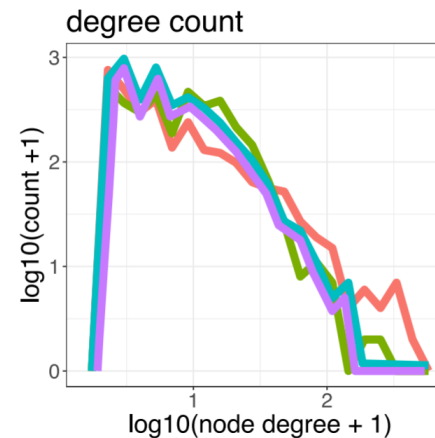
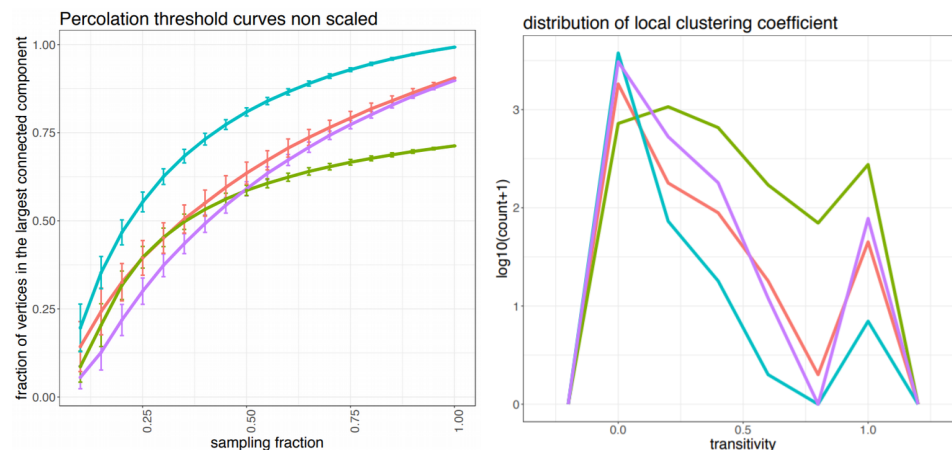


Hidden connections (Gqr>0.005)



Hidden connections are characterized by existence of tight network communities

Comparative topological analysis



- Manually curated network SIGNOR
- Network of hidden interactions
- Randomized direct WikiProtein network
- Direct WikiProtein network

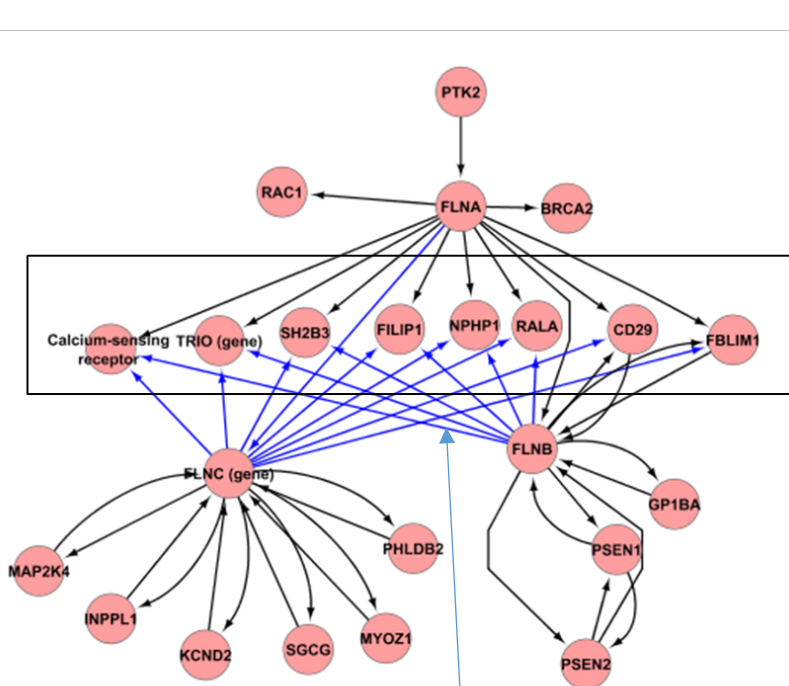
WikiProteins project

Inferring directed network of proteins from Wikipedia using reduced Google Matrix

(ongoing work...)

Studying “hidden” communities of proteins in Wikipedia is informative

Example: Discovering common functional partners of filamins

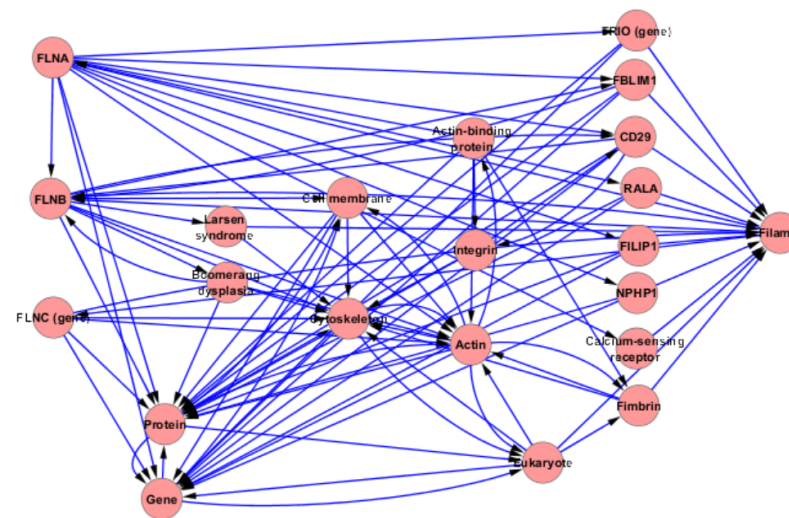


Direct links

Hidden links

“Hidden” connection community

Direct shortest/second shortest paths from filamins FLN* to targets



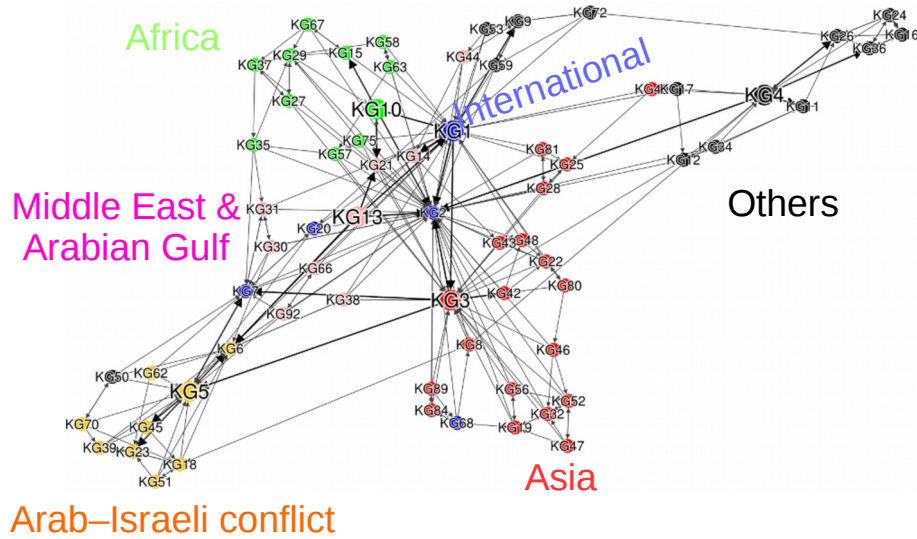
Direct links

Automatically extracted annotation: Filamins are actin-binding proteins, related to cytoskeleton and involved in Larsen syndrome and Boomerang dysplasia

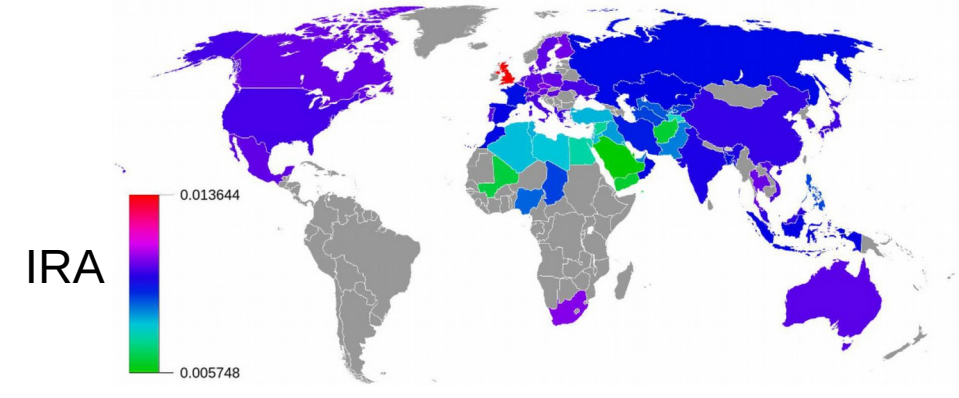
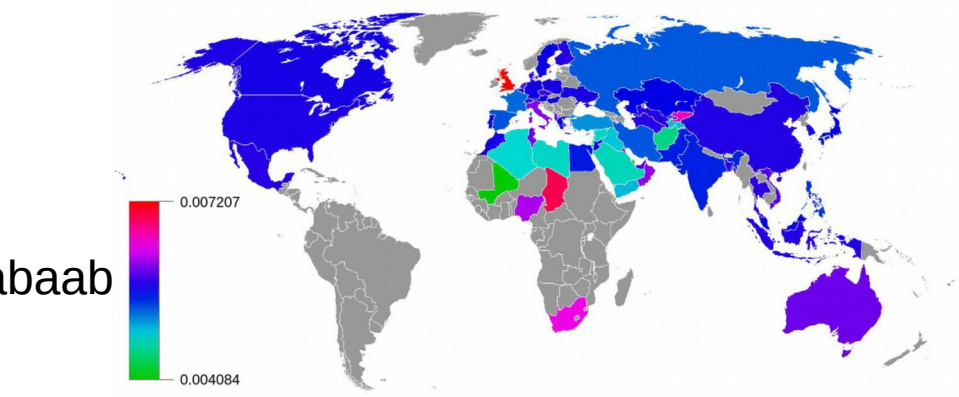
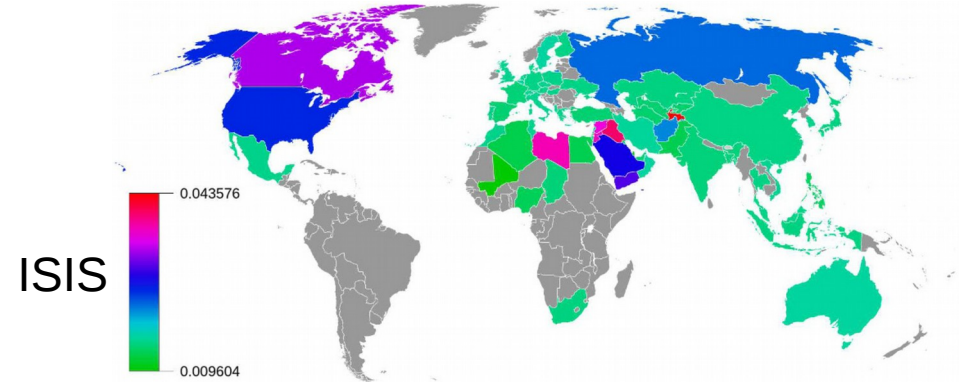
World Terror Networks from the reduced Google matrix of Wikipedia

Reduced Google matrix of 95 terrorist group and 64 world countries

Interactions between terrorist groups

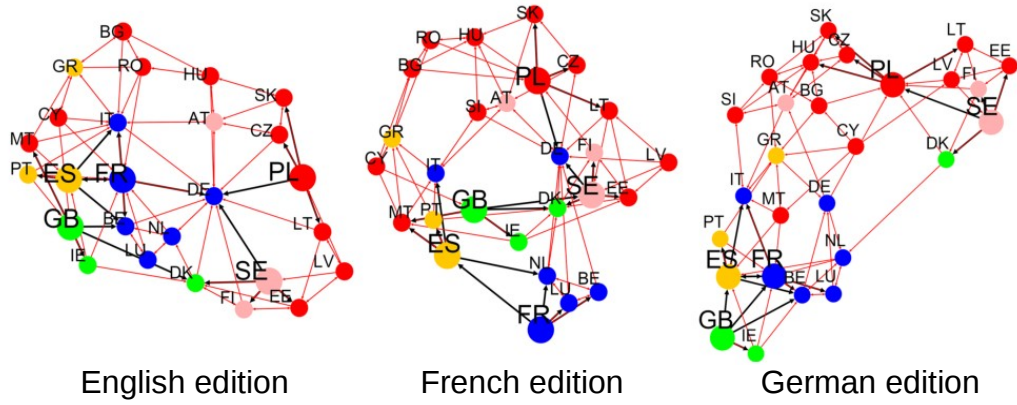


World map of the influence of terrorist groups on countries



Samer El Zant, Klaus M. Frahm, Katia Jaffres-Runser, Dima L. Shepelyansky, Eur. Phys. J. B v.91, p,7 (2018)

EU countries relationships according to Wikipedia



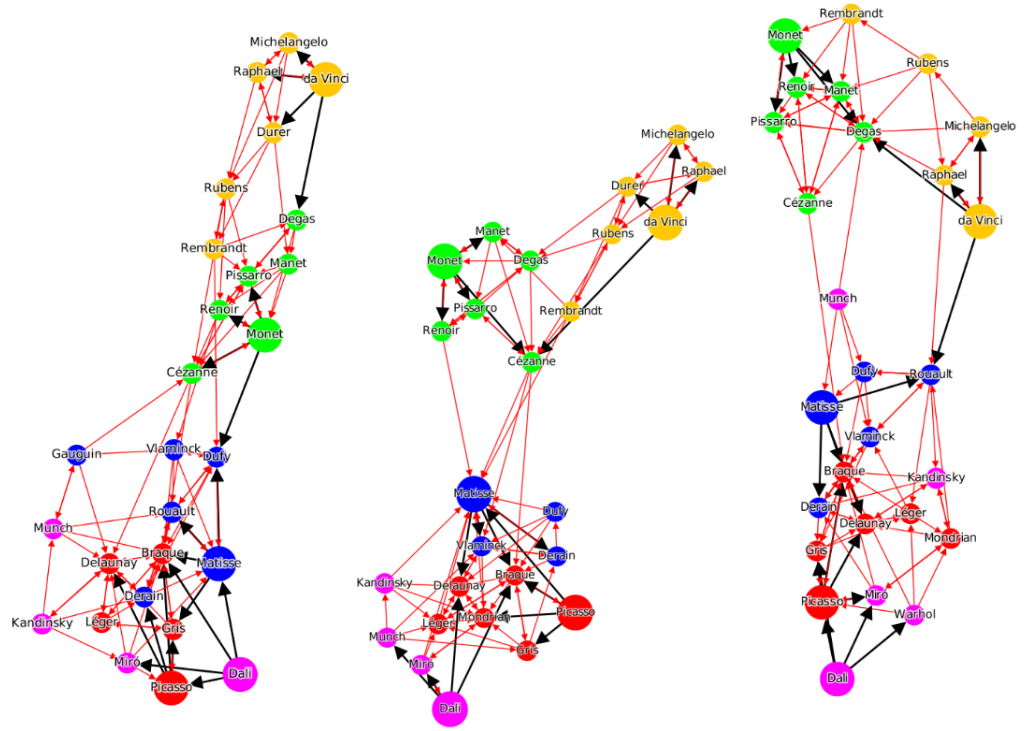
English edition

French edition

German edition

S. El Zant, K.Jaffres-Runser, and D.L.Shepelyansky, "Geopolitical interactions from reduced Google matrix analysis of Wikipedia", submitted.

Painters relationships according to Wikipedia



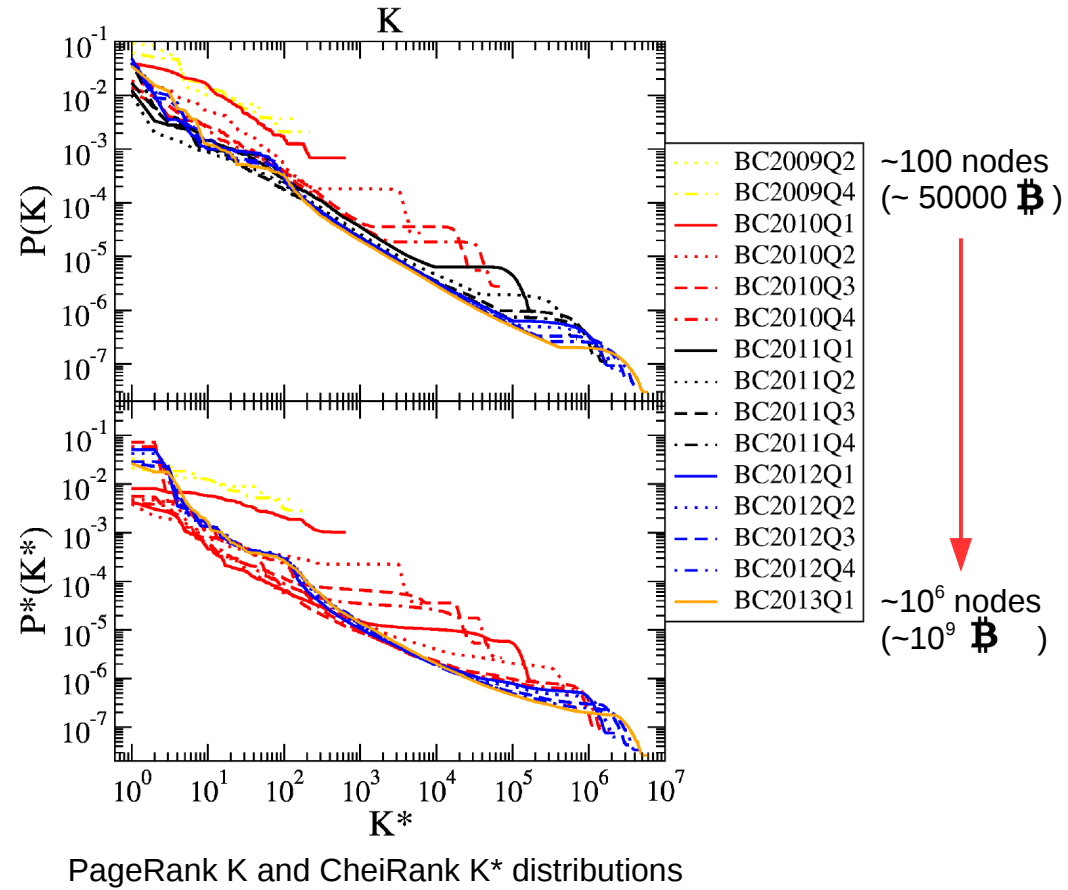
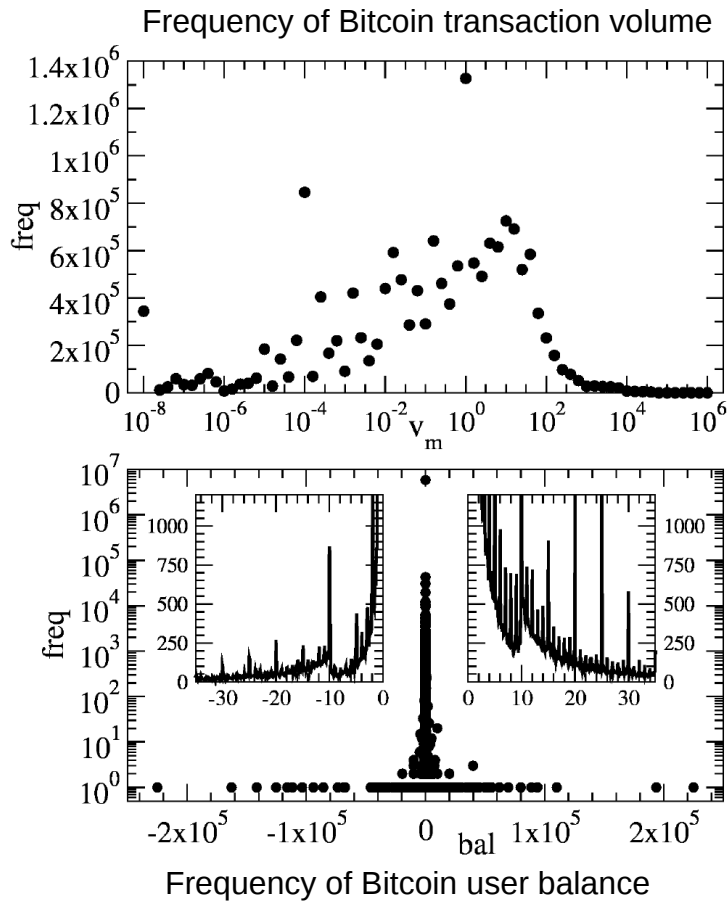
English edition

French edition

German edition

S. El Zant, K.Jaffres-Runser, K.M.Frahm, and D.L.Shepelyansky, "Reduced Google matrix analysis of Art: Painters", submitted.

Google matrix of Bitcoin network

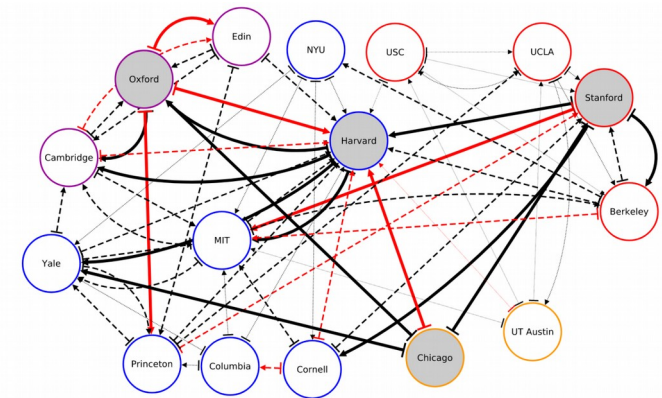


L.Ermann, K.Frahm and D. Shepelyansky "Google matrix of Bitcoin networks", submitted to Eur. Phys. J. B (2017)

Planned further works

Planned further works

- **Global multiproduct world trade network**
Merging UN, WTO and OECD data to construct a global network of about 2 million nodes.
Analysis of global world trade along decades.
Use of reduced Google matrix to find hidden links between activity sectors or products.
- **Global multilingual Wikipedia network**
Build a global Wikipedia network (size about 20 million nodes/articles) using the links between articles from different language editions.
Use of reduced Google matrix to find hidden links between concepts (many possible directions of investigation).
- **The reduced Google matrix method as a Cytoscape plug-in**
The reduced Google matrix method will be available as Cytoscape plugin.
- **Googlomics : application of Google matrix to multifunctional network of omics (proteins, genes, ...)**
taking into account the nature of protein-protein interactions.



Hidden relationships between top 20 universities in Wikipedia [preliminary result]

Thank You !