Spectral graph wavelets: a tool for multiscale community mining in graphs

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Purpose of community detection?



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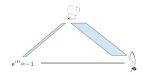
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1) Gives us a sketch:



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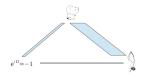
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Purpose of community detection?



1) Gives us a sketch:



2) Gives us intuition:



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Multiscale community structure in a graph

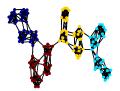
finest scale (16 com.):



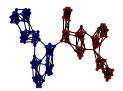
coarser scale (8 com.):



even coarser scale (4 com.):



coarsest scale (2 com.):



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Multiscale community structure in a graph

Classical community detection algorithm do not have this "scale-vision" of a graph. Modularity optimisation finds:



Where the modularity function reads: $Q = \frac{1}{2N} \sum_{ij} \left[A_{ij} - \frac{d_i d_j}{2N} \right] \delta(c_i, c_j)$

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Q=0.74 :



Q=0.83:



Q=0.50 :



All representations are correct but modularity optimisation favours one.

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Related work

- Lambiotte, "Multiscale modularity in complex networks" (2010)
- Schaub et al., "Markov dynamics as a zooming lens for multiscale community detection: non clique-like communities and the field-of-view limit" (2012)
- Arenas et al., "Analysis of the structure of complex networks at different resolution levels" (2008)
- Reichardt et al., "Statistical Mechanics of Community Detection" (2006)
- Mucha et al., "Community Structure in Time-Dependent, Multiscale, and Multiplex Networks" (2010)

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Purpose of this work

Develop a scale dependent community mining tool

General Ideas

- Take advantage of local information encoded in Graph Wavelets
- Cluster together nodes whose local environments are similar

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Notations

$$egin{array}{c|c} \mathcal{G} = (V, E, w) & \mathsf{a} \ N = |V| & \mathsf{nu} \ \mathcal{A} & \mathsf{ac} \ \mathcal{d} & \mathsf{vec} \end{array}$$

a weighted graph number of nodes adjacency matrix vector of strengths

$$\begin{array}{l} A(i,j) = w_{ij} \\ d_i = \sum_{j \in V} w_{ij} \end{array}$$

Laplacian matrix

Objective

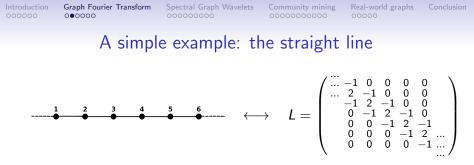
f : signal defined on V \longleftrightarrow \hat{f} : Fourier transform of f

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On the regular line, *L* is the 1-D classical laplacian operator (i.e. double derivative operator): its eigenvectors are the Fourier vectors, and its eigenvalues the associated (squared) frequencies.

Fundamental analogy

On *any* graph, the eigenvectors χ_i of the Laplacian matrix L will be considered as the Fourier vectors, and its eigenvalues λ_i the associated (squared) frequencies.

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The graph Fourier transform

• \hat{f} is obtained from f's decomposition on the eigenvectors χ_i :

$$\hat{f} = egin{pmatrix} <\chi_0,f>\ <\chi_1,f>\ <\chi_2,f>\ \dots\ <\chi_{N-1},f>\end{pmatrix}$$

Define
$$\boldsymbol{\chi} = (\chi_0 | \chi_1 | ... | \chi_{N-1}) : \widehat{f} = \boldsymbol{\chi}^\top f$$

- Reciprocally, the inverse Fourier transform reads: $\left| f = \chi \, \hat{f} \,
 ight|$
- The Parseval theorem is valid: $\forall (g,h) < g,h > = <\hat{g},\hat{h} >$

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Filtering

Definition of graph filtering

We define a filter function g in the Fourier space. It is discrete and defined on the eigenvalues $\lambda_i \rightarrow g(\lambda_i)$.

$$\hat{f}^{g} = \begin{pmatrix} \hat{f}(0) g(\lambda_{0}) \\ \hat{f}(1) g(\lambda_{1}) \\ \hat{f}(2) g(\lambda_{2}) \\ \vdots \\ \hat{f}(N-1) g(\lambda_{N-1}) \end{pmatrix} = \hat{\mathbf{G}} \hat{f} \text{ with } \hat{\mathbf{G}} = \begin{pmatrix} g(\lambda_{0}) & 0 & 0 & \dots & 0 \\ 0 & g(\lambda_{1}) & 0 & \dots & 0 \\ 0 & 0 & g(\lambda_{2}) & \dots & 0 \\ \vdots & \vdots & \vdots & \vdots \\ 0 & 0 & 0 & \dots & g(\lambda_{N-1}) \end{pmatrix}$$

In the node-space, the filtered signal f^g can be written: $f^g = \chi \hat{\mathbf{G}} \chi^\top f$

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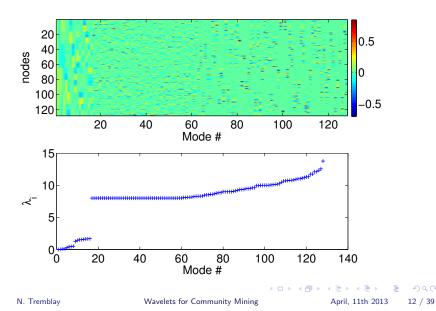
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Spectral analysis: the χ_i and λ_i of the multi scale toy graph



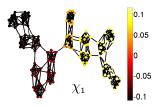
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Some Fourier modes



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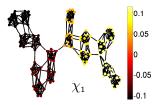
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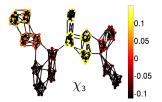
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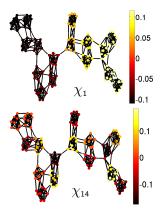
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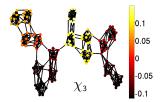
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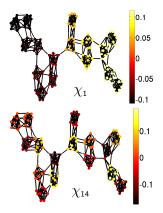
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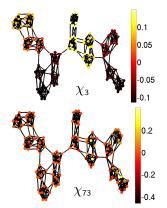
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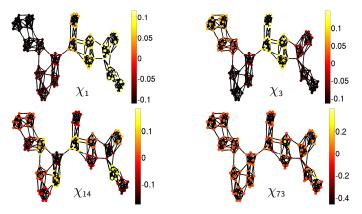
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The first few eigenvectors are very important for community detection

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Graph Wavelets

- Fourier is a global analysis. Fourier modes (eigenvectors of the laplacian) are used in classical spectral clustering, but do not enable a scale dependent analysis: we need wavelets.
- Classical wavelets *by analogy* Graph wavelets

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The classical wavelets

Each wav. $\psi_{\textit{s},\textit{a}}$ is derived by translating and scaling a mother wav. ψ :

$$\psi_{s,a}(x) = \frac{1}{s}\psi\left(\frac{x-a}{s}\right)$$

Equivalently, in the Fourier domain:

$$\hat{\psi}_{s,a}(\omega) = \int_{-\infty}^{\infty} \frac{1}{s} \psi\left(\frac{x-a}{s}\right) \exp^{-i\omega x} dx$$
$$= \exp^{-i\omega a} \int_{-\infty}^{\infty} \frac{1}{s} \psi\left(\frac{X}{s}\right) \exp^{-i\omega X} dX$$
$$= \exp^{-i\omega a} \int_{-\infty}^{\infty} \psi\left(X'\right) \exp^{-i\omega X'} dX'$$
$$= \hat{\delta}_{a}(\omega) \hat{\psi}(s\omega) \quad \text{where} \quad \delta_{a} = \delta(x-a)$$

One possible definition: $\psi_{s,a}(x) = \int_{-\infty}^{\infty} \hat{\delta}_{a}(\omega) \hat{\psi}(s\omega) \exp^{i\omega x} d\omega$

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The classical wavelets

$$\psi_{s,a}(x) \;=\; \int_{-\infty}^{\infty} \hat{\delta_a}(\omega) \hat{\psi}(s\omega) \exp^{i\omega x} \, d\omega$$

- In this definition, $\hat{\psi}(s\omega)$ acts as a filter bank defined by scaling by a factor s a filter kernel function defined in the Fourier space: $\hat{\psi}(\omega)$
- The filter kernel function $\hat{\psi}(\omega)$ is necessarily a bandpass filter with:
 - $\hat{\psi}(0) = 0$: the mean of ψ is by definition null
 - $\lim_{\omega o +\infty} \hat{\psi}(\omega) = 0$: the norm of ψ is by definition finite

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Classical wavelets $\xrightarrow{by \text{ analogy}}$ Graph wavelets (Hammond '11)		
	Classical (continuous) world	Graph world
Real domain variable	х	node <i>a</i>
Fourier domain variable	ω	eigenvalues λ_i
Filter kernel function	$\hat{\psi}(\omega)$	$g(\lambda_i) \Leftrightarrow \hat{\mathbf{G}}$
Filter bank	$\hat{\psi}(m{s}\omega)$	$g(s\lambda_i) \Leftrightarrow \hat{\mathbf{G}}_{\mathbf{s}}$
Fourier modes	$\exp^{-i\omega x}$	eigenvectors χ_i
Fourier transform of f	$\hat{f}(\omega) = \int_{-\infty}^{\infty} f(x) \exp^{-i\omega x} dx$	$\hat{f} = oldsymbol{\chi}^ op f$

The wavelet at scale *s* centered around node *a* is given by:

$$\psi_{s,a}(x) = \int_{-\infty}^{\infty} \hat{\delta}_{a}(\omega) \hat{\psi}(s\omega) \exp^{i\omega x} d\omega \longrightarrow \psi_{s,a} = \chi \, \hat{\mathbf{G}}_{s} \hat{\delta}_{a} = \chi \, \hat{\mathbf{G}}_{s} \, \chi^{\top} \, \delta_{a}$$

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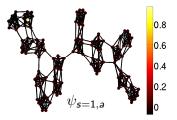
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Examples of wavelets



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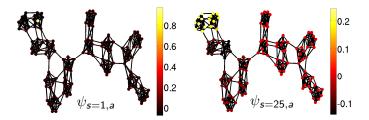
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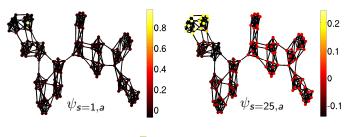
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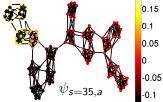
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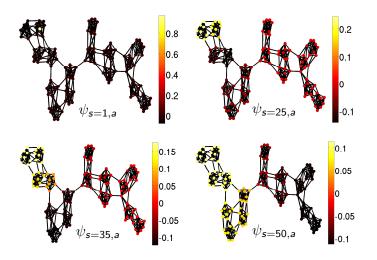
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The graph scaling functions

• Consider the following *lowpass filter kernel h*:

$$h(\omega) = \left(\int_{\omega}^{\infty} rac{|g(\omega')|^2}{\omega'} d\omega'
ight)^{1/2}$$

Classically, if g corresponds to a wavelet filter kernel, this equation defines the associated scaling function filter kernel.

• With the same arguments as previously, we define the graph scaling function at scale *s* centered around *a* as:

$$\phi_{\mathbf{s},\mathbf{a}} = \boldsymbol{\chi}\, \hat{\mathbf{H}}_{\mathbf{s}} \hat{\delta_{\mathbf{a}}} = \boldsymbol{\chi}\, \hat{\mathbf{H}}_{\mathbf{s}}\, \boldsymbol{\chi}^{\top}\, \delta_{\mathbf{a}}$$

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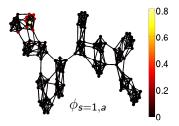
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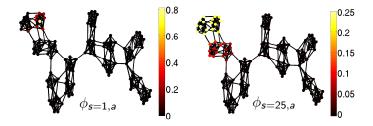
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> 0.25 0.2

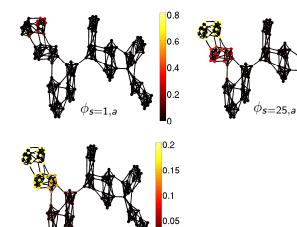
> 0.15

0.1

0.05

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Examples of scaling functions



 $\phi_{s=35,a}$

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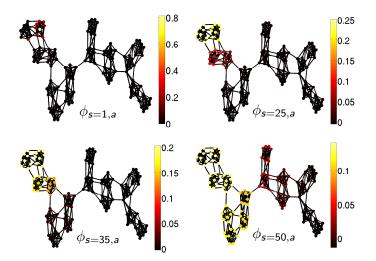
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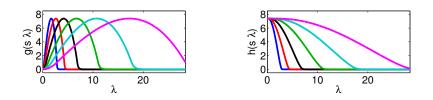
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Example of filters

For each graph under study, we automatically find the good filter parameters for g by imposing:

- The coarsest scale needs to be focused on the first mode χ_1 .
- All scales need at least to keep some information from χ_1 .
- The finest scale needs to use the information from all modes.



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Important note

In the following, we will not *actually* perform a Wavelet Transform of any signal: we will rather focus on the wavelets ψ_i and take advantage of the topological information encoded in them

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Application to detection of communities

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The three key points of clustering

Any clustering technique is based on the choice of:

- 1. feature vectors for each node
- 2. a distance to measure two given vectors' closeness
- 3. a clustering algorithm to separate nodes in clusters

We choose to use:

- 1. wavelets (resp. scaling functions) as feature vectors
- 2. the correlation distance
- 3. the complete linkage clustering algorithm

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Complete linkage clustering

- It is a bottom to top hierarchical algorithm: it starts with as many clusters as nodes and works its way up to fewer clusters (by linking subclusters together) until it reaches one global cluster.
- To compute the distance between two subclusters under examination : all possible pairs of nodes, taking one from each cluster, are considered. The *maximum* possible node-to-node distance is declared to be the cluster-to-cluster closeness.
- Outputs a dendrogram (from Greek dendron "tree" and gramma "drawing").

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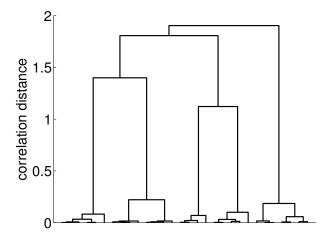
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Example of a dendrogram at a given scale s



The big question: where should we cut the dendrogram?

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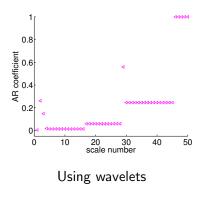
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With prior knowledge

Let us cheat by using prior knowledge on the number of communities we are looking for. If we cut each dendrogram in two clusters



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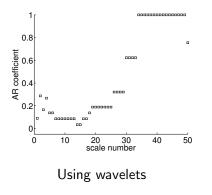
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With prior knowledge

Let us cheat by using prior knowledge on the number of communities we are looking for. If we cut each dendrogram in four clusters



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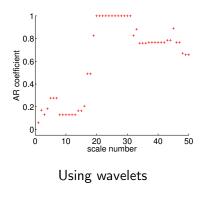
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With prior knowledge

Let us cheat by using prior knowledge on the number of communities we are looking for. If we cut each dendrogram in eight clusters



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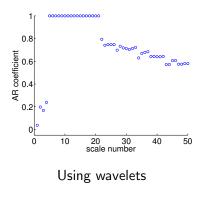
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With prior knowledge

Let us cheat by using prior knowledge on the number of communities we are looking for. If we cut each dendrogram in sixteen clusters



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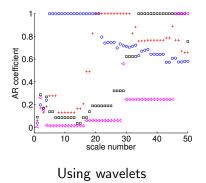
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With prior knowledge

Let us cheat by using prior knowledge on the number of communities we are looking for. The four levels of communities.



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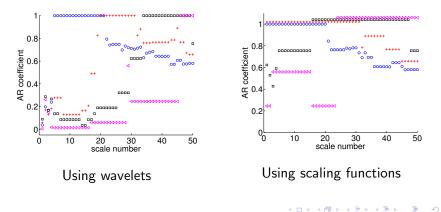
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Let us cheat by using prior knowledge on the number of communities we are looking for. The four levels of communities.



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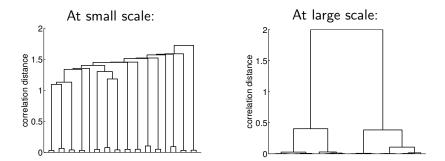
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Without prior knowledge

We cut the dendrogram at its maximal gap.



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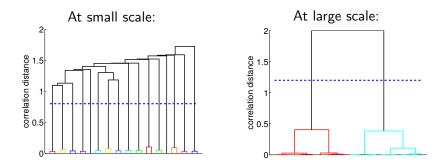
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Without prior knowledge

We cut the dendrogram at its maximal gap.



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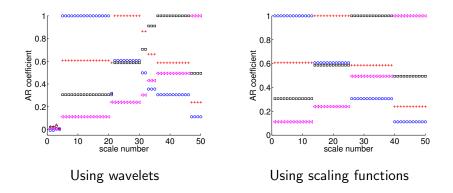
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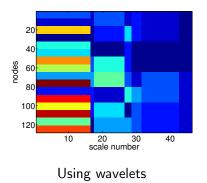
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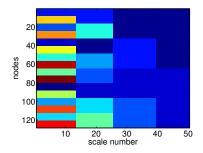
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Using scaling functions

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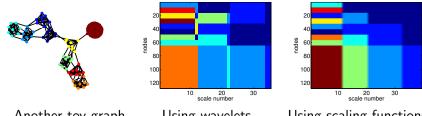
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Another toy graph



Another toy graph

Using wavelets

Using scaling functions

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The filtered modularity

We define the filtered adjacency matrices at scale s:

• recall that $A = D^{\frac{1}{2}}\chi(I - \Lambda)\chi^{\top}D^{\frac{1}{2}}$

•
$$A_{s}^{g} = A + D^{\frac{1}{2}} \chi \hat{G}_{s} \chi^{\top} D^{-\frac{1}{2}} A$$

•
$$A_s^h = D^{\frac{1}{2}} \chi \hat{H}_s \chi^\top D^{-\frac{1}{2}} A$$

The classical modularity reads: $B(A) = \frac{1}{2m}(A + \frac{dd^{\top}}{2m})$ where *d* is the strength vector and $2m = \sum d(i)$

We define the filtered modularity matrices at scale s:

$$B_s^g = B(A_s^g)$$
 and $B_s^h = B(A_s^h)$

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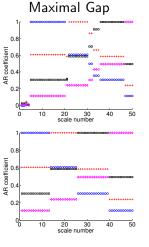
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Maximize filtered modularity



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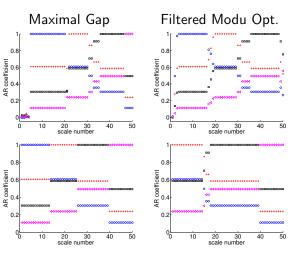
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Maximize filtered modularity



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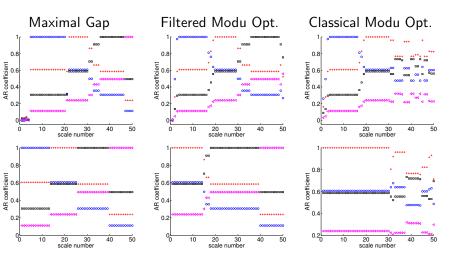
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Maximize filtered modularity



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Two real-world graphs

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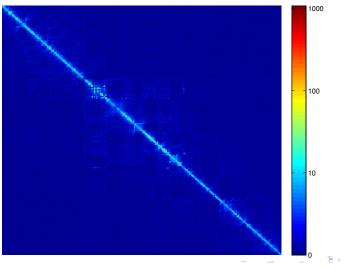
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Intra-chromosomic interaction data

Collaboration with R. Boulos, B. Audit (ENS Lyon)



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Evolution of the correlation matrix of the wavelets with respect to scale

Collaboration with R. Boulos, B. Audit (ENS Lyon)

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The dynamic social network of a primary school

Collaboration with A. Barrat (CPT Marseille)

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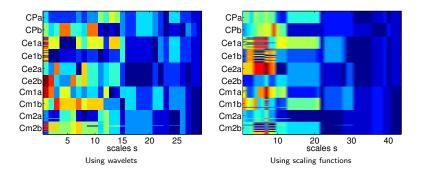
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Multi-scale Communities in Primary School

Collaboration with A. Barrat (CPT Marseille)



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Conclusion

- Wavelet $\psi_{s,a}$ gives an "egocentered view" of the network seen from node *a* at scale *s*
- Correlation between these different views gives us a distance between nodes at scale *s*
- This enables multi-scale clustering of nodes in communities

I did not mention:

- the design of the filters
- the scale boundaries of the parameter "s"
- how we choose the relevant scales (we use a notion of stability of each partition)

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Thank you for your attention!

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The Adjusted Rand Index

Let:

- \mathcal{C} and \mathcal{C}' be two partitions we want to compare.
- a be the # of pairs of nodes that are in the same community in C and in the same community in C'
- b be the # of pairs of nodes that are in different communities in C and in different communities in C'
- c be the # of pairs of nodes that are in the same community in C and in different communities in C'
- d be the # of pairs of nodes that are in different communities in C and in the same community in C'

a + b is the number of "agreements" between C and C'. c + d is the number of "disagreements" between C and C'.

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The Adjusted Rand Index

The Rand index, R, is:

$$R = \frac{a+b}{a+b+c+d} = \frac{a+b}{\binom{n}{2}}$$

The Adjusted Rand index AR is the corrected-for-chance version of the Rand index:

$$AR = \frac{R - ExpectedIndex}{MaxIndex - ExpectedIndex}$$

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Conclusion

The filtered modularity

$$A_{s}^{g} = A + D^{\frac{1}{2}}\chi \hat{G}_{s}\chi^{\top}D^{-\frac{1}{2}}A$$

Consider d the vector of strengths of A and 2m the sum of the strengths. The classical modularity reads:

$$\mathsf{B} = rac{A}{2m} - rac{dd^ op}{(2m)^2}$$

Consider d' the vector of strengths of A_s^g and 2m' the sum of the strengths. We can show that:

$$rac{dd^{ op}}{(2m)^2} = rac{d'd'^{ op}}{(2m')^2}$$

Moreover, if $g_s(1) = 0$ (which is the case), the filtered modularity reads:

$$B_{s}^{g} = \frac{A + D^{\frac{1}{2}}\chi\hat{G}_{s}\chi^{\top}D^{-\frac{1}{2}}A}{2m} - \frac{dd^{\top}}{(2m)^{2}}$$
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The filtered modularity

$$B_{s}^{g} = \frac{A + D^{\frac{1}{2}}\chi \hat{G}_{s}\chi^{\top} D^{-\frac{1}{2}}A}{2m} - \frac{dd^{\top}}{(2m)^{2}}$$

Recall that modularity compares the actual normalised weight $\frac{A_{ij}}{2m}$ to the expected weight if the graph was a random Chung-Lu graph: $\frac{d_i d_j}{(2m)^2}$. The filtered modularity does not change the expected weight but rather changes the actual normalised weight: it adds or retrieve value to $\frac{A_{ij}}{2m}$. At small scale, it will increase the weights important

for small scale structures and decrease the weights important for superstructures.

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The filtered modularity

It can be written:

$$B_{s}^{g} = rac{1}{2m} \sum_{i=2}^{N} (1 + g_{s}(i))(1 - \lambda_{i})D^{rac{1}{2}}\chi_{i}\chi_{i}^{ op}D^{rac{1}{2}}$$

To compare to Delvenne's filtered modularity:

$$B_t = rac{1}{2m} \sum_{i=2}^N (1-\lambda_i)^t D^{rac{1}{2}} \chi_i \chi_i^\top D^{rac{1}{2}}$$

And Arenas' version: (here for regular networks)

$$B_{\alpha} = \frac{1}{2m} \sum_{i=2}^{N} (1 - \frac{\lambda_i}{\alpha}) D^{\frac{1}{2}} \chi_i \chi_i^{\top} D^{\frac{1}{2}}$$

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