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# Introducing a mesoscopic scale with Confluence in Modularity, to improve graph clustering resolution

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#### Abstract

Given a graph G = (V, E) and two vertices  $i, j \in V$ , we introduce Confluence(G, i, j), a vertex mesoscopic closeness measure which brings together vertices from the same link-dense region of the graph G, and separates vertices coming from two distinct dense regions.

Confluence becomes a useful tool to avoid the resolution problems of the standard  $Modularity(G, \Gamma)$  measure for a given clustering  $\Gamma$ , as evidenced by our comparative study between these two measures on toy graphs. We additionally present a heuristic to find a partitional clustering of a graph that tentatively optimizes a clustering quality function derived from *Confluence*, comparing the new heuristic's behaviour to the state of the art Louvain and Infomap methods on real terrain networks, while introducing a way to control the size of the resulting clusters along the way.

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- (2) **DONE** Choisir les graphes de l'état de l'art:
  - https://snap.stanford.edu/data/com-DBLP.html
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  - https://networkrepository.com/actor-movie.php
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- (5) DONE Traduction en bon anglais: Alp et Alexandre
- (6) Relecture: David et Quentin
- (7) Formater le .tex selon les consignes d'édition de Physical-Review(E) : Bruno
- (8) Soumettre l'article à Physical-Review(E) et Déposer l'article sur arxiv : Bruno

Cet article est en cours de rédaction et il est déposé sur https://arxiv.org/user/ pour permetre au logiciel libre Gartgantexte de concourir à un prix du logiciel libre.

Le code Haskell provisoire est déposé ici: https://gitlab.iscpif.fr/gargantext/ gargantext-graph sous la Licence accessible ici: https://gitlab.iscpif.fr/ gargantext/haskell-gargantext/blob/dev/LICENSE

## 1 Introduction

Terrain networks are real world networks that model data gathered by field work, in diverse fields such as sociology, linguistics, biology, or graphs from the internet. Most terrain networks contrast with artificial graphs (deterministic or random) and share four similar properties [Watts and Strogatz, 1998, Albert and Barabasi, 2002, Newman, 2003]. They exhibit:

- **p**<sub>1</sub> : A low density (not many edges);
- $\mathbf{p_2}$ : Short paths (the average number of edges L on the shortest path between two vertices is low);
- $\mathbf{p_3}$ : A high clustering rate  $C = \frac{3 \times number \ of \ triangles}{number \ of \ connected \ triplets}$  (locally densely connected subgraphs can be found whereas the whole graph is globally sparse);
- $\mathbf{p_4}$ : A heavy-tailed degree distribution (the distribution of the degrees of the vertices of the graph can be approximated by a power law).

Clustering a terrain network consists in grouping together in modules vertices that belong to the same densely connected region of the graph (property  $p_3$ ), while keeping separate vertices that do not (property  $p_1$ ). The difference with a classification task is that the number of groups is not known in advance.

## Let $\mathbf{G} = (\mathbf{V}, \mathbf{E})$ be a graph:

-**Module:** A module  $\gamma$  of G is a non-empty subset of the graph's vertices:  $\gamma \neq \emptyset$  and  $\gamma \subset V$ ;

-Clustering: A *clustering*  $\Gamma$  of G is a set of modules of G such that  $\bigcup_{\gamma \in \Gamma} \gamma = V$ ;

-Partitional clustering: If  $\forall \gamma_i, \gamma_j \in \Gamma$ ,  $(i \neq j) \Rightarrow (\gamma_i \cap \gamma_j = \emptyset)$ , then  $\Gamma$  is a *partitional clustering* of G, where modules of G are not allowed to overlap. Given such a  $\Gamma$  we can define an equivalence relation  $\stackrel{\Gamma}{\sim}$  on the set of vertices:  $\forall u, v \in V$ ,  $(u \stackrel{\Gamma}{\sim} v) \Leftrightarrow (\exists \gamma \in \Gamma \text{ such that } u \in \gamma \text{ and } v \in \gamma)$ ;

-Clustering quality function: A clustering quality function  $Q(G, \Gamma)$  is an  $\mathbb{R}$ -valued function whose goal is to measure the adequacy of the modules with the densely connected regions of terrain networks (property  $p_3$ ).

In order to establish a good partitional clustering for a graph G = (V, E), given a clustering quality function Q, it would in theory be sufficient to build all the possible partitionings of the set of vertices V, and to pick a partitioning  $\Gamma$  such that  $Q(G, \Gamma)$  is optimal. This method is however obviously impractical, since the number of partionings of a set of size n = |V| is equal to the  $n^{th}$  Bell number, a sequence known to grow exponentially [Knuth, 1968].

Many graph clustering methods therefore consist in defining a heuristic that can find in a reasonable amount of time a clustering  $\Gamma$  that tentatively optimises  $Q(G, \Gamma)$  for a given clustering quality function Q. Several partitional clustering methods, such as *Louvain* [Blondel et al., 2008], use the *Modularity* quality function suggested in 2004 by Newman and Girvan [Newman and Girvan, 2004].

In section 2 we present the *Modularity* quality function, describing its limits in section 2.1. To avoid these limits, section 3 revisits the definition of *Modularity* to introduce a mesoscopic scale, with a vertices mesoscopic closeness measure *Confluence*, that we define in section 3.1. We then compare the clusterings which maximize this new quality function with the ones that maximize *Modularity*, on a few small toy graphs, in section 3.3.

We then describe, in section 4, a new heuristic to optimize this new quality function on bigger graphs, comparing in section 4.2 the results with those obtained using two of the most used state of the art heuristics: *Louvain* [Blondel et al., 2008] which tries to maximize *Modularity*, and *Infomap*, among the most elegant and efficient heuritics, which tries to maximize the quality function described in 2008 by Rosvall and Bergstrom [Rosvall and Bergstrom, 2008].

Finally, in section 5, we describe a way to control the size of the modules and conclude in section 6.

## 2 Modularity

The modularity of a partitional clustering for a graph G = (V, E) with m = |E| edges is equal to the difference between the proportion of links internal to modules of the clustering, and the same quantity in a null model, where no community structure is expected. The null model is a random graph  $G_{Null}$  with the same number of vertices and edges, as well as the same distribution of degrees as G, where the probability of having an edge between two vertices x and y is equal to  $\frac{d_G(x).d_G(y)}{2m}$ , with  $d_G(i) = |\{v \in V/\{i, v\} \in E\}|$  is the degree of vertex i in G.

Let G = (V, E) be a graph with m edges and  $\Gamma$  a partitioning of V. The modularity of  $\Gamma$  can be defined as follows.

$$Modularity(G,\Gamma) = \frac{1}{2m} \sum_{\gamma \in \Gamma} \sum_{i,j \in \gamma} Pedge(G,i,j) - Pedge(G_{Null},i,j)$$
(1)

Where Pedge(G, x, y) is a symmetrical vertex closeness measure equal to the probability of  $\{x, y\}$  being an edge of G, that is:

$$Pedge(G, i, j) = \begin{cases} 1 \text{ if } \{i, j\} \in E, \\ 0 \text{ otherwise.} \end{cases}$$
(2)

$$Pedge(G_{Null}, i, j) = \frac{d_G(i).d_G(j)}{2m}$$
(3)

In equation 1, the first term  $\frac{1}{2m}$  is purely conventional, so that modularity values all live in the [-1, 1] interval, but plays no role when maximizing modularity, since it is constant for a given graph G.

We then define  $Q_{Pedge}$  as Newman and Girvan's quality function, to be maximized:

$$Q_{Pedge}(G,\Gamma) = \sum_{\gamma \in \Gamma} \sum_{i,j \in \gamma} Pedge(G,i,j) - Pedge(G_{Null},i,j)$$
$$= \sum_{\gamma \in \Gamma} \sum_{i,j \in \gamma} \begin{cases} 1 - \frac{d_G(i).d_G(j)}{2m} & \text{if } \{i,j\} \in E, \\ -\frac{d_G(i).d_G(j)}{2m} & \text{otherwise.} \end{cases}$$
(4)

A good partitional clustering  $\Gamma$  as per 4 is one that groups in the same module vertices that are linked (especially ones with low degrees, but also to a lesser extent ones with high degrees), while avoiding as much as possible the grouping of non-linked vertices (especially ones with high degrees, but to a lesser extent ones with low degrees).

#### 2.1 Limits of Modularity

Several authors [Fortunato and Barthelemy, 2006, Kumpula et al., 2007] showed that optimizing *Modularity* leads to merging small modules into larger ones, even when those small modules are well defined and weakly connected to one another. To address this problem, some authors [Reichardt and Bornholdt, 2006, Arenas et al., 2008] defined multiresolution variants of *Modularity*, adding a resolution parameter to control module sizes.

For instance [Reichardt and Bornholdt, 2006] introduces a parameter  $\lambda \in \mathbb{R}$  in equation 4:

$$Q_{\lambda} = \sum_{\gamma \in \Gamma} \sum_{i,j \in \gamma} \begin{cases} 1 - \lambda . \frac{d_G(i).d_G(j)}{2m} \text{ if } \{i,j\} \in E, \\ -\lambda . \frac{d_G(i).d_G(j)}{2m} \text{ otherwise.} \end{cases}$$
(5)

where  $\lambda$  is a resolution parameter: the higher  $\lambda$  is, the smaller the modules get (high resolution).

However, in [Lancichinetti and Fortunato, 2011], the authors show that "... multiresolution Modularity suffers from two opposite coexisting problems: the tendency to merge small subgraphs, which dominates when the resolution is low; the tendency to split large subgraphs, which dominates when the resolution is high. In benchmark networks with heterogeneous distributions of cluster sizes, the simultaneous elimination of both biases is not possible and multiresolution Modularity is not capable to recover the planted community structure, not even when it is pronounced and easily detectable by other methods, for any value of the resolution parameter. This holds for other multiresolution techniques and it is likely to be a general problem of methods based on global optimization.

... real networks are characterized by the coexistence of clusters of very different sizes, whose distributions are quite well described by power laws [Clauset et al., 2004, Radicchi et al., 2004]. Therefore there is no characteristic cluster size and tuning a resolution parameter may not help."

## 3 Avoiding *Modularity* limits by introducing a mesoscopic scale

In equation 4, with regards to a graph G:

- $Pedge(G, i, j) = \begin{cases} 1 \text{ if } \{i, j\} \in E, \\ 0 \text{ otherwise.} \end{cases}$  is a **local** (microscopic) vertices closeness measure relative to G:
- $Pedge(G_{Null}, i, j) = \frac{d_G(i) \cdot d_G(j)}{2m}$  is a **global** (macroscopic) vertices closeness measure relative to G.

To avoid limits described in section 2.1, we introduce in equation 4 an intermediate **meso**scopic<sup>1</sup> vertices closeness measure relative to G: Confluence(G, i, j) that we define below.

## 3.1 Confluence, a vertices mesoscopic closeness measure

If G = (V, E) is a reflexive<sup>2</sup> and undirected graph, let us imagine a walker wandering on the graph G: at time  $t \in \mathbb{N}$ , the walker is on one vertex  $i \in V$ ; at time t + 1, the walker can reach any neighbouring vertex of i, with uniform probability. This process is called a simple random walk [Bollobas, 2002]. It can be defined by a Markov chain on V with an

 $<sup>^{1}</sup>$ A **meso**scopic scale is an intermediate scale between a local **micro**scopic scale and a global **macro**scopic scale.

 $<sup>^{2}</sup>$  i.e. each vertex is connected to itself. If such self-loops do not exist in the data, they may be added without loss of information.

 $n \times n$  transition matrix [G]:

$$[G] = (g_{i,j})_{i,j \in V} \quad \text{with} \quad g_{i,j} = \begin{cases} \frac{1}{d_G(i)} & \text{if } \{i,j\} \in E, \\ 0 & \text{otherwise.} \end{cases}$$
(6)

Since G is reflexive, each vertex has at least one neighbour (itself) and [G] is therefore well defined. Furthermore, by construction, [G] is a stochastic matrix:  $\forall i \in V, \sum_{j \in V} g_{i,j} =$ 1. The probability  $P_G^t(i \rightsquigarrow j)$  of a walker starting on vertex *i* and reaching vertex *j* after *t* steps is:

$$P_G^t(i \leadsto j) = ([G]^t)_{i,j} \tag{7}$$

**Propsition 1** Let G = (V, E) be a reflexive graph with m edges, and  $G_{null} = (V, E_{null})$  its null model such that the probability of the existence of a link between two vertices i and j is  $e_{i,j} = \frac{d_G(i).d_G(j)}{2m}$ .

$$\forall t \in \mathbb{N}^*, \ \forall i, j \in V, \ P_{G_{null}}^t(i \leadsto j) = \frac{d_G(j)}{2m}$$
(8)

**Proof** by induction on t:

(a) True for t = 1:

$$\forall i, j \in V, P_{G_{null}}^1(i \leadsto j) = e_{i,j} \cdot \frac{1}{d_G(i)} = \frac{d_G(i) \cdot d_G(j)}{2m} \cdot \frac{1}{d_G(i)} = \frac{d_G(j)}{2m}$$

(b) If true for t then true for t + 1:

$$\forall i, j \in V, P_{G_{null}}^{t+1}(i \rightsquigarrow j) = \sum_{k \in V} \left( P_{G_{null}}^t(i \rightsquigarrow k) \cdot P_{G_{null}}^1(k \rightsquigarrow j) \right)$$

$$= \sum_{k \in V} \left( P_{G_{null}}^t(i \rightsquigarrow k) \cdot \frac{d_G(j)}{2m} \right) = \frac{d_G(j)}{2m} \cdot \sum_{k \in V} P_{G_{null}}^t(i \rightsquigarrow k)$$

$$= \frac{d_G(j)}{2m} \cdot \sum_{k \in V} \frac{d_G(k)}{2m} = \frac{d_G(j)}{2m}$$

$$(1) \qquad 2$$

(a) & (b)  $\Rightarrow 8$ 

**On a graph G** = (**V**, **E**) the trajectory of a random walker is completely governed by the topology of the graph in the vicinity of the starting node: after t steps, any vertex j located at a distance of t links or less can be reached. The probability of this event depends on the number of paths between i and j, and on the structure of the graph around the intermediary vertices along those paths. The more short paths exist between vertices i and j, the higher the probability  $P_G^t(i \rightsquigarrow j)$  of reaching j from i. On the graph  $G_{null}$  the trajectory of a random walker is only governed by the degrees of the vertices, and no longer at all by the topology of the graph in the vicinity of the starting node.

We want to consider as "close" each pair of vertices  $\{i, j\}$  having a probability of reaching j from i after a short random walk in G, greater than the probability of reaching j from i in  $G_{null}$ . We therefore define the t-confluence  $Conf^t(G, i, j)$  between two vertices i, j on a graph G as follows:

$$Conf^{t}(G, i, j) = \begin{cases} 0 \text{ if } i = j, \\ \frac{P_{G}^{t}(i \rightsquigarrow j) - P_{G_{null}}^{t}(i \leadsto j)}{P_{G}^{t}(i \leadsto j) + P_{G_{null}}^{t}(i \leadsto j)} = \frac{P_{G}^{t}(i \leadsto j) - \frac{d_{G}(j)}{2m}}{P_{G}^{t}(i \leadsto j) + \frac{d_{G}(j)}{2m}} \text{ otherwise.} \end{cases}$$
(9)

**Propsition 2** Let G = (V, E) be a reflexive graph with m edges, and  $G_{null}$  its null model such that the probability of the existence of a link between two vertices i and j is  $e_{i,j} = \frac{d_G(i) \cdot d_G(j)}{2m}$ .

$$\forall t \in \mathbb{N}^*, \ \forall i, j \in V, \ Conf^t(G_{Null}, i, j) = 0$$
(10)

#### **Proof** :

If 
$$i = j$$
, the result follows directly from definition 9.  
If  $i \neq j$ ,  $Conf^t(G_{Null}, i, j) = \frac{P_{G_{Null}}^t(i \sim j) - \frac{d_{G_{Null}}(j)}{2m}}{P_{G_{Null}}^t(i \sim j) + \frac{d_{G_{Null}}(j)}{2m}}$  (by definition 9)  
 $= \frac{P_{G_{Null}}^t(i \sim j) - \frac{d_G(j)}{2m}}{P_{G_{Null}}^t(i \sim j) + \frac{d_G(j)}{2m}}$  (by definition of  $G_{Null}$ )  
 $= \frac{\frac{d_G(j)}{2m} - \frac{d_G(j)}{2m}}{\frac{d_G(j)}{2m} + \frac{d_G(j)}{2m}}$  (by proposition 1)  
 $= 0$ 

To prove that  $Conf^t(G, \cdot, \cdot)$  is symmetric, we first need to prove proposition 3.

**Propsition 3** Let G = (V, E) be a reflexive graph.

$$\forall t \in \mathbb{N}^*, \ \forall i, j \in V, P_G^t(i \leadsto j) = \frac{d_G(j)}{d_G(i)} \cdot P_G^t(j \leadsto i)$$
(11)

**Proof** by induction on t:

## (a) True for t = 1:

 $\begin{aligned} \forall i, j \in V, \\ \text{if } \{i, j\} \notin E, \text{ then } P_G^1(i \rightsquigarrow j) &= 0 \text{ and } P_G^1(j \rightsquigarrow i) = 0, \text{ therefore } P_G^1(i \rightsquigarrow j) = \frac{d_G(j)}{d_G(i)}.P_G^1(j \rightsquigarrow i) = 0 \\ \text{otherwise } P_G^1(i \leadsto j) &= \frac{1}{d_G(i)} = \frac{d_G(j)}{d_G(i)}.\frac{1}{d_G(j)} = \frac{d_G(j)}{d_G(i)}.P_G^1(j \rightsquigarrow i) \end{aligned}$ 

(b) If true for t then true for t + 1:

$$\begin{aligned} \forall i, j \in V, P_G^{t+1}(i \rightsquigarrow j) &= \sum_{k \in V} \left( P_G^t(i \rightsquigarrow k) . P_G^1(k \rightsquigarrow j) \right) \\ &= \sum_{k \in V} \left( P_G^t(k \rightsquigarrow i) . \frac{d_G(k)}{d_G(i)} . P_G^1(k \rightsquigarrow j) \right) = \sum_{k \in V} \left( P_G^t(k \rightsquigarrow i) . \frac{d_G(k)}{d_G(i)} . P_G^1(j \rightsquigarrow k) . \frac{d_G(j)}{d_G(k)} \right) \\ &= \sum_{k \in V} \left( P_G^t(k \rightsquigarrow i) . P_G^1(j \rightsquigarrow k) . \frac{d_G(j)}{d_G(i)} \right) = \frac{d_G(j)}{d_G(i)} \sum_{k \in V} \left( P_G^1(j \rightsquigarrow k) . P_G^t(k \rightsquigarrow i) \right) \\ &= \frac{d_G(j)}{d_G(i)} . P_G^{t+1}(j \rightsquigarrow i) \end{aligned}$$
(a) & (b)  $\Rightarrow 11$ 

**Propsition 4** Let G = (V, E) be a reflexive graph.

$$\forall t \in \mathbb{N}^*, \ \forall i, j \in V, Conf^t(G, i, j) = Conf^t(G, j, i)$$
(12)

**Proof** :

$$\begin{aligned} \text{If } i=j: \text{ it follows directly from definition 9.} \\ \text{If } i\neq j: Conf^t(G,i,j) &= \frac{P_G^t(i \rightsquigarrow j) - P_{G_{null}}^t(i \rightsquigarrow j)}{P_G^t(i \rightsquigarrow j) + P_{G_{null}}^t(i \rightsquigarrow j)} = \frac{P_G^t(i \rightsquigarrow j) - \frac{d_G(j)}{2m}}{P_G^t(i \rightsquigarrow j) + \frac{d_G(j)}{2m}} = \frac{\frac{d_G(j)}{d_G(i)} \cdot P_G^t(j \rightsquigarrow i) - \frac{d_G(j)}{2m}}{\frac{d_G(j)}{d_G(i)} \cdot P_G^t(j \rightsquigarrow i) + \frac{d_G(j)}{2m}} \\ &= \frac{(\frac{d_G(j)}{d_G(i)} \cdot P_G^t(j \rightsquigarrow i) - \frac{d_G(j)}{2m}) \cdot \frac{d_G(i)}{d_G(j)}}{(\frac{d_G(j)}{d_G(i)} \cdot P_G^t(j \rightsquigarrow i) + \frac{d_G(j)}{2m}) \cdot \frac{d_G(j)}{d_G(j)}} = \frac{P_G^t(j \rightsquigarrow i) - \frac{d_G(j)}{2m}}{P_G^t(j \rightsquigarrow i) + \frac{d_G(j)}{2m}} = \frac{P_G^t(j \rightsquigarrow i) - P_{G_{null}}^t(j \rightsquigarrow i)}{P_G^t(j \rightsquigarrow i) + P_{G_{null}}^t(j \rightsquigarrow i)} \\ &= Conf^t(G, j, i) \end{aligned}$$

Confluence actually defines an infinity of symmetrical vertex closeness measures, one for each random walk length t. For clarity, in the rest of this paper, we set t = 3 and define  $Conf(G, i, j) = Conf^3(G, i, j)$ .

Most terrain networks exhibit the properties  $p_2$  (short paths) and  $p_3$  (high clustering rate). With a classic distance such as the shortest path between two vertices, all vertices would be close to each other in a terrain network (because of property  $p_2$ ). On the contrary, Confluence allows us to identify vertices living in the same higher density zones of G (property  $p_3$ ):

If i, j are in the same high local density region:

$$P_G^3(i \leadsto j) > P_{G_{null}}^3(i \leadsto j), \text{ thus } Conf(G, i, j) > 0$$
<sup>(13)</sup>

If i, j are in two distinct high local density regions:

$$P_G^3(i \leadsto j) < P_{G_{null}}^3(i \leadsto j), \text{ thus } Conf(G, i, j) < 0$$
(14)

## 3.2 Introducing a mesoscopic scale in Modularity

To avoid the limits of *Modularity* described in section 2.1, we propose  $Q_{Conf}$ , a new clustering quality function, which introduces a mesoscopic scale into *Modularity* through *Confluence*:

$$Q_{Conf}(G,\Gamma) = \sum_{\gamma \in \Gamma} \sum_{i \neq j \in \gamma} Conf(G,i,j) + \left(Pedge(G,i,j) - Pedge(G_{null},i,j)\right)$$

By the definitions of Conf (equation 9) and Pedge (equations 2 and 3) :

$$Q_{Conf}(G,\Gamma) = \sum_{\gamma \in \Gamma} \sum_{i \neq j \in \gamma} \frac{P_G^3(i \rightsquigarrow j) - \frac{d_G(j)}{2m}}{P_G^3(i \leadsto j) + \frac{d_G(j)}{2m}} + \begin{cases} 1 - \frac{d_G(i).d_G(j)}{2m} & \text{if } \{i, j\} \in E, \\ -\frac{d_G(i).d_G(j)}{2m} & \text{otherwise.} \end{cases}$$

Leading us to the following definition of  $Q_{Conf}$ :

$$Q_{Conf}(G,\Gamma) = \sum_{\gamma \in \Gamma} \sum_{i \neq j \in \gamma} \begin{cases} \left(\frac{P_G^3(i \rightarrow j) - \frac{d_G(j)}{2m}}{P_G^3(i \rightarrow j) + \frac{d_G(j)}{2m}}\right) + \left(1 - \frac{d_G(i).d_G(j)}{2m}\right) \text{ if } \{i, j\} \in E, \\ \left(\frac{P_G^3(i \rightarrow j) - \frac{d_G(j)}{2m}}{P_G^3(i \rightarrow j) + \frac{d_G(j)}{2m}}\right) - \left(\frac{d_G(i).d_G(j)}{2m}\right) \text{ otherwise.} \end{cases}$$
(15)

To make it easier to interpret  $Q_{Conf}$ , we define the GooDness of a clustering  $\Gamma$  for a graph G = (V, E):

$$GooDness(G,\Gamma) = \frac{1}{2|E|} Q_{Conf}(G,\Gamma)$$
(16)

and the ConFness of a set of vertices  $\gamma \in \Gamma$  for a graph G = (V, E):

$$ConFness(G,\gamma) = \frac{1}{|\gamma| \cdot (|\gamma| - 1)} \sum_{i \neq j \in \gamma} Confluence(G,i,j)$$
(17)

#### 3.3 Optimality for $Q_{Pedge}$ versus optimality for $Q_{Conf}$

A partitional clustering  $\Delta$  is optimal for a quality function Q iff: For all partitioning  $\Gamma$  of  $V, Q(G, \Delta) \geq Q(G, \Gamma)$ . Computing a  $\Delta$  that maximizes  $Q_{Pedge}(G, \Delta)$  is  $\mathcal{N}P$  - complete [Brandes et al., 2008], and the same holds for computing a clustering that maximizes  $Q_{Conf}$ .

However, when the number of vertices of a graph G = (V, E) is small, the problem of maximizing the modularity can be turned into a reasonably tractable Integer Linear Program (see [Brandes et al., 2008]): We define  $n^2$  decision variables  $X_{ij} \in \{0, 1\}$ , one for each pair of vertices  $\{i, j\} \in V$ . The key idea is that we can build an equivalence relation on V ( $i \sim j$  iff  $X_{ij} = 1$ ) and therefore a partitioning of V. To guarantee that the decision variables give rise to an equivalence relation, they must satisfy the following constraints:

**Reflexivity:**  $\forall i \in V, X_{ii} = 1;$ 

Symmetry:  $\forall i, j \in V : X_{ij} = X_{ji};$ 

$$\begin{array}{l} \textbf{Transitivity: } \forall i,j,k \in V : \begin{cases} \forall i,j,k \in V : X_{ij} + X_{jk} - 2.X_{ik} \leq 1; \\ \forall i,j,k \in V : X_{ik} + X_{ij} - 2.X_{jk} \leq 1; \\ \forall i,j,k \in V : X_{jk} + X_{ik} - 2.X_{ij} \leq 1. \end{cases} \end{array}$$

With the following objective functions to maximize:

For 
$$\mathbf{Q}_{\mathbf{Pedge}}$$
:  $\sum_{i,j\in V} X_{ij} \cdot \left( Pedge(G, i, j) - Pedge(G_{Null}, i, j) \right)$   
For  $\mathbf{Q}_{\mathbf{Conf}}$ :  $\sum_{i\neq j\in V} X_{ij} \cdot \left( Conf(G, i, j) + Pedge(G, i, j) - Pedge(G_{null}, i, j) \right)$ 

$$(18)$$

On four small atificial graphs,  $G_{toy}^1$ , ...,  $G_{toy}^4$ , we compare optimal clusterings  $\Delta_{Pedge}^G$ and  $\Delta_{Conf}^G$  respectively computed for  $Q_{Pedge}$  and  $Q_{Conf}$ , with results illustrated in Figure 1.

The optimal clusterings for  $Q_{Conf}$  do not necessarily have a higher resolution than with  $Q_{Pedge}$ , they can even have a lower resolution<sup>3</sup>:

• Fig. 6(b):  $\Delta_{Conf}^{G_{toy}^1} \langle 0.20, 0.40 \rangle$  versus  $\Delta_{Pedge}^{G_{toy}^1} \langle 0.21, 0.38 \rangle$ :

$$\Delta_{\text{Conf}}^{\mathbf{G}_{\text{toy}}^{1}} \text{ has a higer resolution than } \Delta_{\text{Pedge}}^{\mathbf{G}_{\text{toy}}^{1}}, \text{ with } \delta_{Pedge}^{1} = \delta_{Conf}^{2} \cup \delta_{Conf}^{3};$$

$$\circ \Delta_{Conf}^{G_{toy}^{1}} = \left\{ \delta_{Conf}^{1} = \{0, 4, 5, 6\} \ \langle 0.10 \rangle, \ \delta_{Conf}^{2} = \{1, 2, 3\} \ \langle 0.18 \rangle, \ \delta_{Conf}^{3} = \{7, 8\} \ \langle 0.18 \rangle \right\};$$

$$\circ \Delta_{Pedge}^{G_{toy}^{1}} = \left\{ \delta_{Pedge}^{1} = \{1, 2, 3, 7, 8\} \ \langle 0.04 \rangle, \ \delta_{Pedge}^{2} = \{0, 4, 5, 6\} \ \langle 0.10 \rangle \right\}.$$

<sup>3</sup>Where  $\langle \_, \_ \rangle$  for the  $\langle Modularity, GooDness \rangle$  of the clustering, and  $\langle \_ \rangle$  for the  $\langle ConFness \rangle$  of the modules

• Fig 1(b):  $\Delta_{Conf}^{G_{toy}^2} \langle 0.37, 0.75 \rangle$  versus  $\Delta_{Pedge}^{G_{toy}^2} \langle 0.38, 0.70 \rangle$ :  $\Delta_{Conf}^{G_{toy}^2}$  has a lower resolution than  $\Delta_{Pedge}^{G_{toy}^2}$ , with  $\delta_{Conf}^1 = \delta_{Pedge}^1 \cup \delta_{Pedge}^2$ :

$$\begin{split} &\circ \ \Delta^{G^2_{toy}}_{Conf} = \Big\{ \delta^1_{Conf} = \{0, 1, 2, 3, 4, 5, 6, 7, 8, 9\} \ \langle 0.13 \rangle, & \delta^2_{Conf} = \{13, 14, 15, 16\} \ \langle 0.59 \rangle, \\ &\delta^3_{Conf} = \{10, 11, 12\} \ \langle 0.67 \rangle \Big\}; \\ &\circ \ \Delta^{G^2_{toy}}_{Pedge} = \Big\{ \delta^1_{Pedge} = \{0, 1, 3, 4, 8\} \ \langle 0.17 \rangle, \ \delta^2_{Pedge} = \{2, 5, 6, 7, 9\} \ \langle 0.22 \rangle, \\ & \delta^3_{Pedge} = \{13, 14, 15, 16\} \ \langle 0.59 \rangle, \ \delta^4_{Pedge} = \{10, 11, 12\} \ \langle 0.67 \rangle \Big\}. \end{split}$$

• Fig 1(c):  $\Delta_{Conf}^{G_{toy}^3} \langle 0.30, 0.62 \rangle$  versus  $\Delta_{Pedge}^{G_{toy}^3} \langle 0.33, 0.61 \rangle$ :

 $\Delta^{\mathbf{G}^{3}_{\text{toy}}}_{\mathbf{Conf}} \text{ is different from } \Delta^{\mathbf{G}^{3}_{\text{toy}}}_{\mathbf{Pedge}}, \text{ with } \delta^{1}_{Conf} = \delta^{2}_{Pedge} \cup \delta^{3}_{Pedge} \text{ and } \delta^{1}_{Pedge} = \delta^{2}_{Conf} \cup \delta^{3}_{Conf}$ 

$$\begin{split} \circ \ \Delta^{G^3_{toy}}_{Conf} = & \Big\{ \delta^1_{Conf} {=} \{ 0, 1, 2, 3, 4, 5, 6, 7, 8, 9 \} \ \langle 0.10 \rangle, \ \delta^2_{Conf} {=} \{ 10, 11, 12 \} \ \langle 0.65 \rangle, \\ & \delta^3_{Conf} {=} \{ 13, 14, 15 \} \ \langle 0.66 \rangle \Big\}; \\ \circ \ \Delta^{G^3_{toy}}_{Pedge} = & \Big\{ \delta^1_{Pedge} {=} \{ 10, 11, 12, 13, 14, 15 \} \ \langle 0.26 \rangle, \ \delta^2_{Pedge} {=} \{ 0, 2, 6, 7, 8 \} \ \langle 0.17 \rangle, \\ & \delta^3_{Pedge} {=} \{ 1, 3, 4, 5, 9 \} \ \langle 0.15 \rangle \Big\}. \end{split}$$

• Fig 1(d):  $\Delta_{Conf}^{G_{toy}^4} \langle 0.32, 0.53 \rangle$  versus  $\Delta_{Pedge}^{G_{toy}^4} \langle 0.32, 0.53 \rangle$ :  $\Delta_{Conf}^{G_{toy}^4}$  is equal to  $\Delta_{Pedge}^{G_{toy}^4}$ :

$$\circ \ \Delta_{Conf}^{G_{toy}^{4}} = \Delta_{Pedge}^{G_{toy}^{4}} = \Big\{ \delta^{1} = \{0, 1, 2, 3, 4, 5, 6, 7\} \ \langle 0.19 \rangle, \ \delta^{2} = \{8, 9, 10, 11, 12, 13, 14, 15\} \ \langle 0.12 \rangle, \\ \delta^{3} = \{16, 17, 18, 19, 20, 21, 22, 23\} \ \langle 0.19 \rangle \Big\}.$$



Figure 1: Optimality with respect to  $\mathbf{Q}_{Pedge}$  versus optimality with respect to  $\mathbf{Q}_{Conf}$ . Shapes describe an optimal clustering for  $Q_{Conf}$  (if two vertices have same shape, then they are in a same module for  $Q_{Conf}$ ). Colors describe an optimal clustering for  $Q_{Pedge}$  (if two vertices have same color, then they are in a same module for  $Q_{Pedge}$ ).

## 4 A heuristic by edges confluence

We describe in this section a heuristic for tentatively maximizing  $Q_{Conf}$ . To this end, we start with a variant on Conf, named Confr, where the confluence between two adjacent vertices is computed by removing the edge between them<sup>4</sup>:

$$Confr(G, i, j) = \begin{cases} Conf(G, i, j) \text{ if } \{i, j\} \notin E, \\ Conf(G = (V, E - \{i, j\}), i, j) \text{ otherwise.} \end{cases}$$
(19)

Confr gives us an ordering on the edges  $\{i, j\} \in E$  of the graph. In particular, sorting the edges by decreasing Confr(G, i, j) forms the basis of a new strategy for constructing the cluster's modules, Hbec (Heuristic by edges confluence), described<sup>5</sup> in algorithm 1. On  $G_{toy}^5$ , the graph depicted in Figure 2, Hbec yields the following clusters:  $Hbec(G_{toys}^5) = \{\{0, 1, 3, 4, 5\}, \{6, 7, 8, 9, 10\}, \{2\}\}$ . It is interesting to note how vertex 2 is left isolated in its own cluster by Hbec, whereas it joins the  $\{6, 7, 8, 9, 10\}$  module in the optimal clustering for  $Q_{Conf}$ :  $\Delta_{Conf}^{G_{toy}^5} = \{\{0, 1, 3, 4, 5\}, \{2, 6, 7, 8, 9, 10\}\}$ .

<sup>&</sup>lt;sup>4</sup>A comparative study of Confr with over 80 similarity measures between vertices of a graph has been done by Emmanuel Navarro in his thesis, looking into the sensitivity of various methods to the density of graphs, paths of length 1 and the degrees of vertices, among other aspects. This thesis is available at https://oatao.univ-toulouse.fr/12024/1/navarro.pdf. The study shows that, among the 80 similarity measures considered, Confr(G, i, j) is the only one that, even though it does not systematically regroup pairs of vertices  $\{i, j\} \in E$ , is correlated to the local edge density around vertices i and j, while being independent of the global density of the graphs.

<sup>&</sup>lt;sup>5</sup>Different edges might happen to have the exact same Confr value, making the process non-deterministic in general, because of its sensitivity on the order in which the edges with identical Confr values are processed. A simple solution to this problem is to sort edges by first comparing their Confr values and then using the lexicographic order on (i, j) when Confr values are strictly identical.



Figure 2: **G**<sup>5</sup><sub>toys</sub>

Indeed, when a vertex is linked to two or more distinct dense regions of a graph, Hbec tends to leave that vertex isolated due to its Confr-centered strategy, which does not push the said vertex towards one of those regions over the other. It is therefore desirable to add one last step to the process, to get even closer to the optimal clustering by trying to merge more modules with a greedy algorithm:

**FastGreedy**(**Hbec**(**G**)) : we iteratively merge pairs of modules resulting from Hbec(G) when the union of modules is locally optimal with respect to  $Q_{Conf}$ , stopping when merging any pair of the remaining modules would not result in an increase of  $Q_{Conf}$ .

We call  $\mathbf{Kodex}(\mathbf{G})$  the complete algorithm, including the final  $\mathbf{FastGreedy}$  step. We now get:  $Kodex(G_{toys}^5) = \Delta_{Conf}^{G_{toy}^5}$ .

### 4.1 Implementation

Our implementation of *Kodex* reflects the high-level description from the previous sections and is therefore split into three main parts: pre-computing the confluence of all the edges of the graph (and sorting the said list of edges in decreasing confluence order), doing a first clustering pass (*Hbec*), and the final greedy step. Let G = (V, E), and n = |V|.

## Edges confluence:

- First, construct the reflexive sparse adjacency matrix of the input graph,  $Adj(G) \in \mathcal{M}_n(\mathbb{R})$ , represented using the Compressed Sparse Column format, supported by the vast majority of sparse linear algebra libraries, by supplying the appropriate routine with all the (i, j, 1) triplets, where  $\{i, j\} \in E$  or i = j. That is,  $Adj(G)_{i,j} = 1$  if  $i, j \in E$  or i = j, 0 otherwise.
- Then compute the corresponding transition matrix, T(G), assuming a uniform probability of reaching any neighbor from a given vertex. This amounts to setting all non-zero values of a column j to  $\frac{1}{deg_G j}$ . That is,  $T(G)_{i,j} = \frac{1}{deg_G(j)}$  if  $i, j \in E$ , 0 otherwise.

Algorithm 1 Hbec(G) Heuristic by edges confluence for optimizing  $Q_{Conf}$ 

```
Input: G = (V, E) an undirected graph
Output: C a partitional clustering of G
for i \in V do
  mod_i \leftarrow \{i\}
  v_i \leftarrow i
q \leftarrow 0
\Upsilon \hookleftarrow \varnothing
While \Upsilon \neq E do
  \{i, j\} \leftarrow \underset{\{x, y\} \in E - \Upsilon}{\operatorname{arg\,max}} Confr(G, x, y) \triangleright (\mathbf{a})
  \Upsilon \leftarrow \Upsilon \cup \{\{i,j\}\}
  if v_i \neq v_j then \blacktriangleright i and j are not in the same module
    q_{candidate} \leftarrow q + 2. \sum_{x \in m_i} \sum_{y \in m_i} Conf(G, x, y) + Pedge(G, x, y) - Pedge(G_{null}, x, y) \triangleright (b)
    if q \leq q_{candidate} then
          q \longleftarrow q_{candidate}
         mod_i \leftarrow mod_i \cup mod_i
         for y \in mod_j do
              v_y \leftarrow v_i
         m_j \leftarrow \emptyset
C \hookleftarrow \varnothing
for i \in V do
  if mod_i \neq \emptyset
    C \leftarrow C \cup \{mod_i\}
return C
```

- Then compute the proxemy matrix,  $P(G) = T(G)^3$ .
- Next, we iterate over the list of edges  $\{a, b\} \in E$ , constructing another list:
  - Define  $T(G \{a, b\})$  to be the same as T(G) except for the entries at (a, b) and (b, a) that should be set to 0, to emulate the removal of the link between vertices a and b, and an update to the other entries of the two relevant columns to reflect a decrease by 1 in the degree of vertices a and b, to account for the freshly removed link.

That is,

$$T(G - \{a, b\})_{i,j} = \begin{cases} 0 \text{ if } \{i, j\} = \{a, b\} \\ \frac{1}{\deg_G(j)} \text{ if } j = a \text{ or } j = b \text{ but } i \notin \{a, b\} \\ T(G)_{i,j} \text{ otherwise} \end{cases}$$

- Define  $e_b \in \mathbb{R}^n$  as  $(e_b)_i = 1$  if i = b, 0 otherwise, ideally using a sparse representation.
- Multiply  $T(G \{a, b\})$  and  $e_b$ , and multiply the result by  $T(G \{a, b\})$ , and again, as many times as the length of the random walk is, which has been 3 in most of this paper. That is, for t = 3, we compute:  $T(G - \{a, b\})T(G - \{a, b\})T(G - \{a, b\})e_b$ . We call the resulting vector  $w \in \mathbb{R}^n$
- Finally, compute  $conf_{a,b} = \frac{w_a \frac{deg_G(a) 1}{nnz(Adj(G)) 2}}{w_a + \frac{deg_G(a) 1}{nnz(Adj(G)) 2}}$  where nnz is a function returning the number of non-zero entries in a sparse matrix, typically provided by sparse linear algebra libraries. Return  $(a, b, conf_{a,b})$ .
- Sort the list of  $(a, b, conf_{a,b})$  according to  $conf_{a,b}$ , in decreasing order, breaking equalities by discriminating on the lexicographic ordering on a and then b.

## Hbec

- We create a clustering with all vertices isolated in their own 1-point cluster. Clusters are represented with associative maps from cluster identifiers to sets of vertex identifiers, i.e  $clust[i] = \{a, b, c\}$  means that vertices a, b, c are in cluster number i, while maintaining another associative map from vertex identifiers to cluster identifier, i.e index[i] = c means that vertex i belongs to cluster c. We additionally maintain a mapping from cluster identifiers to scores, score[i], to avoid save some computations that we will keep reusing, instead of performing them from scratch every time.
- We set the score of our clustering,  $clust_score$  to 0 initially.
- We iterate over the sorted list of edges from the previous phase. When processing entry  $(a, b, conf_{a,b})$ :

We compute the clustering quality improvement we would get by merging clusters index[a] and index[b]:

$$score(index[a] \text{ and } index[b]) = \sum_{i \in clust[index[a]], j \in clust[index[b]]} f(i,j)$$
 (20)

where f is defined as:

$$f(i,j) = \frac{P(G)_{i,j} - \frac{\deg_G(i)}{nnz(Adj(G))}}{P(G)_{i,j} + \frac{\deg_G(i)}{nnz(Adj(G))}} + Adj(G)_{i,j} - \frac{(\deg_G(i) - 1)(\deg_G(j) - 1)}{nnz(Adj(G)) - n}$$

- If score(index[a] and index[b]) score[index[a]] score[index[b]] > 0, we merge index[b] into index[a]:
  - \*  $clust_score := clust_score + score(index[a] \text{ and } index[b]) score[index[a]] score[index[b]]$
  - $* \ clust[index[a]] := clust[index[a]] \cup clust[index[b]]$
  - \* score[index[a]] := score(index[a] and index[b])
  - \* For all  $i \in clust[index[b]]$ , we set index[i] := index[a].

Otherwise, we move on to the next edge.

• Once all the edges have been processed, clust describes a partitional clustering of G.

#### Final greedy step

- For all pairs of non-empty clusters resulting from *Hbec*, compute the sum described in equation 20, using an associative container of some sort to record the mapping between the pair of cluster identifiers and the resulting score.
- If there is a pair of clusters for which the sum yields a strictly positive number (i.e a pair that would improve the quality of the clustering, if merged), merge them, updating the scores computed in the first step to account for the merge, wherever one of those clusters is involved.
- Repeat until no such pair of clusters exists anymore.

## 4.2 Tests

We compare three heuristics for computing a partitional clustering:

• Louvain : The Louvain method [Blondel et al., 2008] is the state of the art heuristic that tentatively maximizes Modularity;

- Infomap : The Infomap method is a heuristic for tentatively maximizing the quality function described in 2008 by Rosvall and Bergstrom [Rosvall and Bergstrom, 2008]. This quality function is based on the minimum description length principle [Grünwald, 2007]. It consists in measuring the compression ratio that a given partitioning  $\Gamma$  provides for describing the trajectory of a random walk on a graph. The trajectory description happens on two levels. When the walker enters a module, we write down its name. We then write the vertices that the walker visits, with a notation local to the module, so that an identical short name may be used for different vertices from different modules. A concise description of the trajectory, with a good compression ratio, is therefore possible when the modules of  $\Gamma$  are such that the walker tends to stay in them, which corresponds to the idea that the walker is "captured" when it enters a good module, which is supposed to be a densely linked region that is only weakly connected to other modules. The quality<sup>6</sup> of a partitional clustering is the compression ratio that  $\Gamma$  provides for describing the trajectory of a random walker on G;
- Kodex : The heuristic described in section 4 that tentatively maximizes  $Q_{Conf}$ .

#### 4.2.1 Louvain, Infomap and Kodex on four G<sub>toys</sub> graphs

We start by testing Louvain, Infomap and Kodex on the four  $G_{toys}$  graphs of section 3.3. The results are summarized in Figure 3.

On these four  $G_{toys}$  graphs, Louvain finds the optimal modules with respect to  $Q_{Pedge}$ while Kodex finds the optimal modules with respect to  $Q_{Conf}$ . On  $G_{toy}^2$  and  $G_{toy}^3$ , Infomap and Kodex agree on the clustering, both finding the optimal modules with respect to  $Q_{Conf}$ . However, on  $G_{toy}^1$  and  $G_{toy}^4$ , Infomap collects all the vertices into a single module, because there is no way to compress the description of the path of a random walker on either of those graphs.

Indeed,  $G_{toy}^4$  has been artificially constructed as follow:  $G_{toy}^4 = (V, E)$  where V is the union of 3 sets of 8 vertices,  $\gamma_1 = \{0, ..., 7\}$ ,  $\gamma_2 = \{8, ..., 15\}$ ,  $\gamma_3 = \{16, ..., 23\}$ , and edges are randomly drawn using two different probabilities: we use a probability of 0.5 for drawing an edge between two vertices from the same  $\gamma_i$ , and a probability of 0.25 when the two vertices come from different sets.

Vertices from a given  $\gamma_i$  are expected to have as many internal links (within  $\gamma_i$ ) as they have links to vertices from the two other sets. At every single step, the random walker is equally likely to leave a  $\gamma_i$  or stay within it, which effectively prevents any path compression to take place in  $G_{toy}^4$ .

<sup>&</sup>lt;sup>6</sup>Note that this quality function cannot be expressed as  $\sum_{\gamma \in \Gamma} \sum_{i,j \in \gamma} sim(G,i,j)$ , with sim(G,.,.) an  $\mathbb{R}$ -valued symmetric similarity measure between vertices of G. We therefore left out this quality function in our study of optimality in section 3.3, not having the ability to define the corresponding objective function to maximize in a similar fashion to what was done for  $Q_{Pedge}$  and  $Q_{Conf}$  with the formulas 18.



Figure 3: Clusterings with Louvain, Infomap and Kodex: shapes describe Kodex(G), colors describe Louvain(G), and blocks describe Infomap(G).

- On  $\mathbf{G}_{toy}^1$ :  $Louvain(G_{toy}^1) = \Delta_{Pedge}^{G_{toy}^1}$ ,  $Infomap(G_{toy}^1) = \{V\}$ ,  $Kodex(G_{toy}^1) = \Delta_{Conf}^{G_{toy}^1}$
- On  $\mathbf{G}_{\mathbf{toy}}^2$ :  $Louvain(G_{toy}^2) = \Delta_{Pedge}^{G_{toy}^2}$ ,  $Infomap(G_{toy}^2) = Kodex(G_{toy}^2) = \Delta_{Conf}^{G_{toy}^2}$
- On  $\mathbf{G}^{\mathbf{3}}_{\mathbf{toy}}$ :  $Louvain(G^3_{toy}) = \Delta^{G^3_{toy}}_{Pedge}$ ,  $Infomap(G^3_{toy}) = Kodex(G^3_{toy}) = \Delta^{G^3_{toy}}_{Conf}$
- On  $\mathbf{G}_{\mathbf{toy}}^4$ :  $Louvain(G_{toy}^4) = Kodex(G_{toy}^4) = \Delta_{Conf}^{G_{toy}^4} = \Delta_{Pedge}^{G_{toy}^4}$ ,  $Infomap(G_{toy}^4) = \{V\}$

## 4.2.2 Louvain, Infomap and Kodex on terrain graphs

We now compare Louvain, Infomap and Kodex on four state-of-the-art terrain graphs.

- $G_{FrSynV}$ : The *Dicosyn* resource comes from a collaboration between IBM and the CNRS laboratory ATILF http://www.atilf.fr/. Starting with seven classical French dictionaries (Bailly, Benac, Du Chazaud, Guizot, Lafaye, Larousse and Robert), they extracted synonymic relationships, and the resulting graph  $G_{FrSyn}$  was then symmetrized and split by PoS into three graphs:  $G_{FrSynV}$  for verbs,  $G_{FrSynN}$  for nouns and  $G_{FrSynA}$  for adjectives.
- **G**<sub>Actors</sub>: The well known network of actors [Watts and Strogatz, 1998] has been built from the Internet Movie Database (April 1997) http://us.imdb.com; nodes are actors, and two actors are linked if they have played in a film together [Rossi and Ahmed, 2015] https://networkrepository.com/actor-collaboration.php.
- **G**<sub>DBLP</sub>: The DBLP computer science bibliography provides a comprehensive list of research papers in computer science [Leskovec and Krevl, 2014]. Two authors are connected if they have published at least one paper together https://snap.stanford.edu/data/com-DBLP.html.

• **G**<sub>Amazon</sub>: A network was collected by crawling the Amazon website. It is based on the *Customers Who Bought This Item Also Bought* feature of the Amazon website [Leskovec and Krevl, 2014]. If a product *i* is frequently co-purchased with product *j*, the graph contains an undirected edge from *i* to *j* https://snap.stanford. edu/data/com-Amazon.html.

Table 1 illustrates the pedigrees<sup>7</sup> of these terrain graphs, and Table 2 illustrates the execution times, number of modules, and the length of the biggest module found by the different methods..

Table 1: **Pedigrees:** n and m are the number of vertices and edges,  $\langle k \rangle$  is the mean degree of vertices, C is the clustering coefficient of the graph,  $L_{lcc}$  is the average shortest path length between any two nodes of the largest connected component (largest subgraph in which there exist at least one path between any two nodes) and  $n_{lcc}$  the number of vertices of this component,  $\lambda$ is the coefficient of the best fitting power law of the degree distribution and  $r^2$  is the correlation coefficient of the fit, measuring how well the data is modelled by the power law.

Graph	n	m	$\langle {f k}  angle$	$\mathbf{C}$	$\mathbf{L_{lcc}}(\mathbf{n_{lcc}})$	$\lambda(\mathbf{r^2})$
$G_{FrSynV}$	$9,\!147$	$51,\!423$	11.24	0.14	$4.20(8,\!993)$	-1.88(0.91)
G <sub>Actors</sub>	$68,\!684$	$3,\!007,\!298$	87.57	0.20	$3.51(68,\!019)$	-1.49(0.90)
$G_{DBLP}$	$317,\!080$	1,049,866	6.62	0.31	$6.79(317,\!080)$	-2.71(0.95)
$\mathbf{G}_{\mathbf{Amazon}}$	$334,\!863$	$925,\!872$	5.53	0.21	$11.95(334,\!863)$	-2.81(0.93)

(TODO: Commenter les resultats de la table 2: Bruno)

Table 2: Clusterings of four Terrain Graphs with Louvain, Kodex, Infomap: (T) Time computation in seconds (Louvain and Infomap in C++, Kodex in Haskell (see section 4.1), (N) Number of modules, (M) Length of the biggest module and  $\langle -; -; - \rangle$  for  $\langle Louvain; Kodex; Infomap \rangle$ 

	$G_{FrSynV}$	$G_{Actors}$	$G_{DBLP}$	$G_{Amazon}$
$(\mathbf{T})$	$\langle 0 \ ; \ ?? \ ; \ 13  angle$	$\langle ?? \ ; \ ?? \ ; \ ??  angle$	$\langle 11 \ ; \ 9, 639 \ ; \ 1, 933  angle$	$\langle 6 \ ; \ 7,830 \ ; \ 1,795  angle$
$(\mathbf{N})$	$\langle 90 \ ; \ 671 \ ; \ 575  angle$	$\langle ?? \ ; \ ?? \ ; \ ??  angle$	$\langle 238 \ ; \ 17,437 \ ; \ 16,960 \rangle$	$\langle 234 \ ; \ 20,703 \ ; \ 17,292 \rangle$
$(\mathbf{M})$	$\langle 1, 189 \ ; \ 155 \ ; \ 173 \rangle$	$\langle ????; ; ???; ; ??? \rangle$	$\langle 25, 893 \ ; \ 249 \ ; \ 599 \rangle$	$\langle 12,783\ ;\ 535\ ;\ 347  angle$

<sup>&</sup>lt;sup>7</sup>The power law estimation we give,  $\lambda(r^2)$  is not very accurate (see for instance [Newman, 2005]). However, giving a correct estimation of the odds that a given discrete distribution is heavy-tailed is a difficult issue ([Goldstein et al., 2004, Clauset et al., 2009]), and refining the power-law estimates is beyond the scope of this paper.

## 5 Modulating the size of modules

In many domains such as linguistics, sociology or biology, it can be useful to look into sets of objects (documents, sentences, words, individuals, cells, ...) at different levels of granularity. In order to gain some control over the size of modules, we introduce a  $\beta$  parameter into  $Q_{Conf}$ , resulting in the following  $Q_{Conf}^{\beta}(G, \Gamma)$  quality function:

$$Q_{Conf}^{\beta}(G,\Gamma) = \sum_{\gamma \in \Gamma} \sum_{i \neq j \in \gamma} \begin{cases} Conf(G, i, j) + \left(1 - \frac{d_G(i).d_G(j)}{2m}\right) \text{ if } \{i, j\} \in E, \\ Conf(G, i, j) - \left(\frac{d_G(i).d_G(j)}{2m}\right) - \beta * \left(1 - Conf(G, i, j)\right) \text{ otherwise.} \end{cases}$$

$$= \sum_{\gamma \in \Gamma} \sum_{i \neq j \in \gamma} \begin{cases} \frac{P_{G}^{3}(i \sim j) - \frac{d_{G}(j)}{2m}}{P_{G}^{3}(i \sim j) + \frac{d_{G}(j)}{2m}} + \left(1 - \frac{d_{G}(i).d_{G}(j)}{2m}\right) \text{ if } \{i, j\} \in E, \\ \frac{P_{G}^{3}(i \sim j) - \frac{d_{G}(j)}{2m}}{P_{G}^{3}(i \sim j) + \frac{d_{G}(j)}{2m}} - \left(\frac{d_{G}(i).d_{G}(j)}{2m}\right) - \beta * \left(1 - \frac{P_{G}^{3}(i \sim j) - \frac{d_{G}(j)}{2m}}{P_{G}^{3}(i \sim j) + \frac{d_{G}(j)}{2m}}\right) \text{ otherwise.} \end{cases}$$
(21)

Since  $P_G^3(i \rightsquigarrow j) \in [0, 1]$ , it follows that  $\frac{P_G^3(i \rightsquigarrow j) - \frac{d_G(j)}{2m}}{P_G^3(i \leadsto j) + \frac{d_G(j)}{2m}} \in [-1, 1]$ , and therefore: if  $\beta > 0$ , then  $-\beta * \left(1 - \frac{P_G^3(i \rightsquigarrow j) - \frac{d_G(j)}{2m}}{P_G^3(i \leadsto j) + \frac{d_G(j)}{2m}}\right) \leqslant 0$ . That is, with  $\beta > 0$ :

- If  $\{i, j\} \in E$ , then noting changes ;
- If {i, j} ∉ E and i, j are in the same high density region, then we slightly penalize the {i, j} pair (by 13);
- If {i, j} ∉ E and i, j are in two distinct high density regions, then we strongly penalize the {i, j} pair (by 14).

Because we must avoid as much as possible that the modules contain pairs  $\{i, j\} \notin E$  (with increasing efficiency as  $\beta$  grows), this forces modules to be smaller, as illustrated by Figure 4 on graph  $G_{toy}^3$ . Increasing  $\beta$  does not simply split modules (resulting in a hierarchical approach). This can be observed when going from  $\beta = 0.5$  to  $\beta = 1.0$ , noticing that there is no  $\delta \in \Delta_{Q_{conf}}^{G_{toy}^3}$  such that  $\delta_{Q_{conf}^{1.0}}^2 = \{0, 2, 4\} \subset \delta$ .

Revisiting the previous algorithm to account for the  $\beta$  parameter, yielding  $Kodex(G, \beta)$ , simply amounts to replacing line  $\blacktriangleright$  (b) in algorithm 1 by:

$$q_{candidate} \leftarrow q+2. \sum_{x \in m_i} \sum_{y \in m_j} \begin{cases} Conf(G, x, y) + Pedge(G, x, y) - Pedge(G_{null}, x, y) \text{ if } \{x, y\} \in E, \\ (1+\beta).Conf(G, x, y) + Pedge(G, x, y) - Pedge(G_{null}, x, y) - \beta \text{ otherwise.} \end{cases}$$



Figure 4: **Optimal clusterings of**  $G_{toy}^3$  **for**  $\mathbf{Q}_{Conf}^{\beta=0.0}$ ,  $\mathbf{Q}_{Conf}^{\beta=0.5}$ ,  $\mathbf{Q}_{Conf}^{\beta=1.0}$ : In Fig. 4(a), blocks describe  $\Delta_{Q_{Conf}^{0.0}}^{G_{toy}^3}$  an optimal clustering of  $G_{toy}^3$  for  $Q_{Conf}^{\beta=0.0}$ , shapes describe  $\Delta_{Q_{Conf}^{0.5}}^{G_{toy}^3}$  for  $Q_{Conf}^{\beta=0.5}$ , and colors describe  $\Delta_{Q_{Conf}^{1.0}}^{G_{toy}^3}$  for  $Q_{Conf}^{\beta=1.0}$ .

$$\circ \ \Delta_{Q_{Conf}^{0}}^{G_{toy}^{3}} = \left\{ \delta_{Q_{Conf}^{0}}^{1} = \{0, 1, 2, 3, 4, 5, 6, 7, 8, 9\} \ \langle 0.10 \rangle, \ \delta_{Q_{Conf}^{0}}^{2} = \{10, 11, 12\} \ \langle 0.65 \rangle, \\ \delta_{Q_{Conf}^{0}}^{3} = \{13, 14, 15\} \ \langle 0.66 \rangle \right\};$$

$$\circ \ \Delta_{Q_{Conf}^{0,5}}^{G_{toy}^{3}} = \left\{ \delta_{Q_{Conf}^{0,5}}^{1} = \{0, 2, 6, 7, 8\} \ \langle 0.17 \rangle, \ \delta_{Q_{Conf}^{0,5}}^{2} = \{1, 3, 4, 5, 9\} \ \langle 0.15 \rangle, \ \delta_{Q_{Conf}^{0,5}}^{3} = \{10, 11, 12\} \ \langle 0.65 \rangle, \\ \delta_{Q_{Conf}^{0,5}}^{4} = \{13, 14, 15\} \ \langle 0.66 \rangle \right\};$$

$$\circ \ \Delta_{Q_{Conf}^{0,5}}^{G_{toy}^{3}} = \left\{ \delta_{Q_{Conf}^{1,0}}^{1} = \{1, 3, 5, 9\} \ \langle 0.17 \rangle, \ \delta_{Q_{Conf}^{1,0}}^{2} = \{0, 2, 4\} \ \langle 0.14 \rangle, \ \delta_{Q_{Conf}^{1,0}}^{3} = \{6, 7, 8\} \ \langle 0.27 \rangle, \\ \delta_{Q_{Conf}^{1,0}}^{4} = \{10, 11, 12\} \ \langle 0.65 \rangle, \ \delta_{Q_{Conf}^{1,0}}^{5} = \{13, 14, 15\} \ \langle 0.66 \rangle \right\}.$$

Table 3 shows the results for  $Kodex(G,\beta)$  with  $\beta = 0.0$ ,  $\beta = 0.5$ ,  $\beta = 1.0$  on four real terrain graphs. To compare two clusterings  $\Gamma_1$  and  $\Gamma_2$ , of a graph G = (V, E) we define:

Figure 5 shows  $Fscore(\Gamma_1, \Gamma_2)$  between clusterings Kodex(G, 0.0), Kodex(G, 0.5), Kodex(G, 1.0), Louvain(G) et Infomap(G) on the graphs  $G_{FrSynV}$ ,  $G_{Actors}$ ,  $G_{DBLP}$  and  $G_{Amazon}$ .

Table 3: Kodex clusterings of four terrain graphs with  $\beta = 0.0$ ,  $\beta = 0.5$ ,  $\beta = 1.0$  $\langle -; - \rangle$  for  $\langle$  Number of modules ; Length of the biggest modules  $\rangle$ 

	${ m G}_{{ m FrSynV}}$	$\mathbf{G}_{\mathbf{Actors}}$	$G_{DBLP}$	$\mathbf{G}_{\mathbf{Amazon}}$
$\beta = 0.0$	$\langle 671 \ ; \ 155 \rangle$	$\langle ?? \ ; \ ??  angle$	$\langle 17,437 \ ; \ 249 \rangle$	$\langle 20,703 \ ; \ 535 \rangle$
$\beta = 0.5$	$\langle 785 \ ; \ 126 \rangle$	$\langle ?? \ ; \ ??  angle$	$\langle ?? \ ; \ ?? \rangle$	$\langle ?? \ ; \ ?? \rangle$
$\beta = 1.0$	$\langle 861 \ ; \ 89 \rangle$	$\langle ?? \ ; \ ??  angle$	$\langle ?? \ ; \ ?? \rangle$	$\langle ?? \ ; \ ?? \rangle$



Figure 5: **Fscore**( $\Gamma_1$ ,  $\Gamma_2$ ): Comparison of clusterings Kodex(G, 0.0), Kodex(G, 0.5), Kodex(G, 1.0), Louvain(G) and Infomap(G) on  $G_{FrSynV}$ ,  $G_{Actors}$ ,  $G_{DBLP}$  et  $G_{Amazon}$ .

(TODO: Commenter les resultats de la table 3 et de la figure 5: Bruno)

## 6 Conclusions and perspectives

In this paper, we defined *Confluence*, a mesoscopic vertex closeness measure based on short random walks, that we introduced into *Modularity* to define  $Q_{Conf}$ , a new clustering quality function. With four small toy graphs, we showed that optimal clusterings for  $Q_{Conf}$  improve the resolution of optimal clusterings for *Modularity*.

We then introduced Kodex(G), a heuristic based on the *Confluence* of edges to optimize  $Q_{Conf}$  on a graph G. On the same four little toy graphs, we showed that Kodexfinds an optimal clustering for  $Q_{Conf}$  and that for two of those toy graphs, Kodex(G) =Infomap(G), while on the other two, Kodex(G) identifies the modules we expected while Infomap(G) regroups all vertices into a single module.

We showed that *Kodex* avoids the resolution problem of *Louvain* on four real terrain graphs, and that the modules computed by *Infomap* are closer to those computed by

Kodex than the ones produced by Louvain.

Finally, to get a handle on the size of modules, we introduced the  $\beta \in \mathbb{R}$  parameter, resulting in the  $Q_{Conf}^{\beta}$  quality function and the  $Kodex(G,\beta)$  variant of the algorithm. We then showed that this approach is not hierarchical<sup>8</sup>.

(TODO: Alp et Alexandre: Avantages de l'implémentation Haskel: Monotache/Parallelisme, Complexité/Efficacité, Maintenance, Portabilité ...)

## Length of Random Walks:

For clarity and simplicity, we restricted the random walks of  $P_G^t(i \rightarrow j)$  to a length of t = 3. A first study of the impact of the length of those random walks was done in [Gaume et al., 2010], but a deeper one should be carried to understand how the length influences the **meso**scopicity of *Confluence* and its effect on  $Q_{Conf}$  and *Kodex*. Indeed, if G is connected and  $i \neq j$ :

• When  $\mathbf{t} = \mathbf{1}$ :  $Conf^t(G, i, j) = \begin{cases} \frac{2m - d_G(i).d_G(j)}{2m + d_G(i).d_G(j)} & \text{if } \{i, j\} \in E, \\ -1 & \text{otherwise.} \end{cases}$  Confluence is inde-

pendent of the intermediate structures between the two vertices i and j ;

- When 1 < t < ∞: Conf<sup>t</sup>(G, i, j) = P<sup>t</sup><sub>G</sub>(i→j) dG(j)/2m/2m/2m/2m}, Confluence is dependent of the t-intermediate structures (t-mesoscopicity) between the two vertices i and j (see13 and 14);
- When  $\mathbf{t} \to \infty$ :  $\lim_{t\to\infty} Conf^t(G, i, j) = 0$ , Confluence is constant<sup>9</sup>.

For example we can build the graph  $G_{toy}^{1\star}$  from  $G_{toy}^{1}$  by inserting a new vertex in the middle of each edge. Figure 6 illustrates the optimal clusterings on  $G_{toy}^{1}$  and on  $G_{toy}^{1\star_{self.memory}}$  for  $Q_{Conf}^{0.0}$  with t = 3 and with t = 6, allowing us to see that:

$$\begin{aligned} \mathbf{G}_{toy}^{1 \bigstar} \text{ with } \mathbf{t} &= \mathbf{6}: \text{ we get the modules } \Delta_{Conf}^{G_{toy}^{*}} = \left\{ \delta_{Conf}^{1} = \{0, 4, 5, 6\}, \ \delta_{Conf}^{2} = \{1, 2, 3, \}, \ \delta_{Conf}^{3} = \{7, 8\} \right\} \text{ i.e the optimal clusters of } G_{toy}^{1} \text{ for } Q_{Conf}^{0.0} \text{ with } t = 3: \\ &- \delta_{Conf}^{1} = \{0, 4, 5, 6\} \subset \{0, 4, 5, 6, cut(0/4), cut(0/5), cut(0/6), cut(4/5), cut(4/6), cut(5/6)\}; \\ &- \delta_{Conf}^{2} = \{1, 2, 3, \} \subset \{1, 2, 3, cut(0/1), cut(0/2), cut(0/3), cut(1/2), cut(1/3), cut(2/3)\}; \end{aligned}$$

<sup>&</sup>lt;sup>8</sup>If  $\beta_1 < \beta_2$  then the optimal modules for  $Q_{Conf}^{\beta_2}$  are not necessarily subsets of the optimal modules for  $Q_{Conf}^{\beta_1}$ .

 $- \ \delta^{3}_{Conf} = \{7, 8\} \subset \{7, 8, cut(7/8), cut(3/7), cut(4/7), cut(2/8), cut(5/8)\}.$ On  $G^{1\bigstar}_{toy}$  with  $\mathbf{t} = 3$ : we do not get the modules of  $\Delta^{G^{1}_{toy}}_{Conf}$ .



Figure 6: **Optimal clusterings for**  $Q_{Conf}^{0.0}$  **with** t = 3 **and with** t = 6: shapes describe an optimal clustering for  $Q_{Conf}^{0.0}$  with t = 3, colors describe an optimal clustering for  $Q_{Conf}^{0.0}$  with t = 6.

## **Directed graphs:**

If G is a positively weighted graph by  $W = \{w_{i,j} \text{ such } \{i, j\} \in E\}$ , then we can apply  $Q_{Conf}$  and Kodex by replacing equations 6 and 9 by 22 and 23 respectively:

$$[G] = (g_{i,j})_{i,j\in V} \quad \text{with} \quad g_{i,j} = \begin{cases} \frac{w_{i,j}}{\sum_{k\in V} w_{i,k}} & \text{if } \{i,j\}\in E, \\ 0 & \text{otherwise.} \end{cases}$$
(22)

$$Conf^{t}(G, i, j) = \begin{cases} 0 \text{ if } i = j, \\ \frac{P_{G}^{t}(i \leadsto j) - \frac{\sum_{k \in V} w_{k,j}}{\sum_{w \in W} w}}{P_{G}^{t}(i \leadsto j) + \frac{\sum_{k \in V} w_{k,j}}{\sum_{w \in W} w}} \text{ otherwise.} \end{cases}$$
(23)

If G is a directed graph, one can also consider using a variant of page rank [Gaume and Mathieu, 2016] in place of equation 7.

### **Overlaps:**

In order to define a heuristic for computing clusters with potential overlaps between modules, one could consider using *Hbec* to isolate vertices between several dense regions of the graph (see section 4) and allowing them to belong to the modules corresponding to those dense regions, identified with the help of *ConFness* (see 17).

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