# Two provably consistent divide and conquer clustering algorithms for large networks

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Based on joint work with

Peter Bickel (UC Berkeley) and Purnamrita Sarkar (UT Austin)

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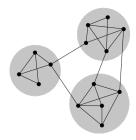


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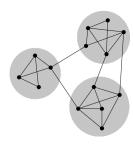


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**Communities**: Groups of nodes that are more densely connected within than across (or the other way round).

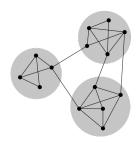


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- political orientation of blogs,
- functionally similar genes,
- facebook groups,
- groups according to feeding mode in ecological networks (who eats who) etc.

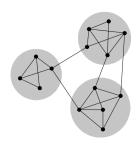


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Stochastic Block Model (SBM)

K: number of communities/blocks  $\sigma$ : latent block membership vector B: block connection probabilities

$$\bullet \ \mathbb{P}_{\sigma(i)=a,\sigma(j)=b}(A_{ij}=1)=B_{ab}.$$

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 Likelihood based methods, modularity optimization, semi-definite programming (SDP), spectral clustering etc.

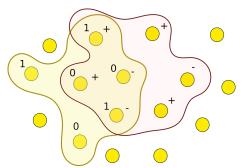
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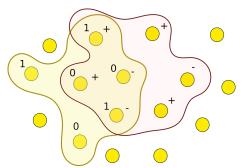
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- Added advantage: parallelizable.

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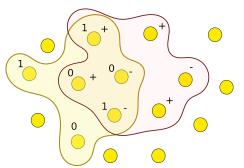


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- Will talk about PACE only.

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Our main idea is to concentrate on estimating whether two individuals belong to the same community or not, instead of estimating Z directly, i.e. we focus on

$$C = ZZ^{\top}$$
,

the so-called clustering matrix. However, this is equivalent to estimating the community structure.

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Let  $A_{S_\ell}$ : the adjacency matrix of the network induced by  $S_\ell$ .

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- $② \ \mathsf{Set} \ \hat{C}_{S_\ell} = \hat{Z}_{S_\ell} \hat{Z}_{S_\ell}^\top.$
- 3 Extend  $\hat{C}_{S_{\ell}}$  to an  $n \times n$  matrix  $\hat{C}^{(\ell)}$ :

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Can use any standard algorithm in the first step, e.g. profile likelihood (PL), mean field method (MF), spectral clustering (SC), semi-definite programming (SDP) etc.

#### PACE: patching up

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Here  $1 \le \tau \le T$  is an integer tuning parameter. We will call  $\hat{C}_{\tau}$  as Piecewise Averaged Community Estimator (PACE).

#### Simulations

Algorithm	ME(%)	Time (seconds)
SDP	0	1588
SDP + PACE + spectral	0	288
SDP + PACE + RPKmeans	9.1	281

Table: PACE with SDP. n = 5000,  $d_n = 128$ , m = 500, 4 equal sized clusters, T = 100, 20 parallel workers in Matlab.

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Algorithm	ME(%)	Time (seconds)
RSC	39.6	87
RSC + PACE + spectral	3.4	21
RSC + PACE + RPKmeans	34.2	14

Table: PACE with RSC. n = 5000,  $d_n = 7$ , cluster proportions  $\pi = (0.2, 0.8)$ , T = 100, 20 parallel workers in Matlab, 3-hop neighbourhoods were used as subgraphs

#### Consistency

Measure of distance between two clustering matrices  $C = ZZ^T$  and  $C' = Z'Z'^T$ :

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#### Theorem (informal)

Suppose  $\tilde{\delta}_S$  denotes the misclustering error on a randomly selected subgraph S. Then

$$\mathbb{E}\tilde{\delta}_{\mathsf{PACE}} \leq \mathbb{E}\tilde{\delta}_{\mathsf{S}} + \textit{coverage error}.$$

Coverage error is  $O(\exp(-O(Tm^2/n^2)))$  for random *m*-subgraphs.

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Using results of Lei and Rinaldo (2015), Guédon and Vershynin (2015) can show

- ① one needs subgraphs of size  $m\gg nd_n^{-1}$ , and  $T\gg \frac{n^2}{m^2}$  for consistency.
- ② Complexity becomes  $O_{\star}(\frac{n^3}{d_n})$ , from  $O(n^3)$ .
- Also a significant boost comes from parallelizability.

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# Thank You!

Note that

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- Thus we expect that any distance based clustering algorithm working on rows of  $\hat{C}$  will do well.
- Computing distance for n-dimensional vectors is expensive, so we do a random projection of the rows to  $\Omega(\log n)$  dimensions (**Johnson-Lindenstrauss**) first, and then do clustering of those projected rows.