

A comparison between dissimilarity SOM & kernel SOM for clustering the vertices of a graph

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Modelization through large graphs

Various fields:

- Computer science: World Wide Web, P2P network...
- Social networks
- Biology: Protein interactions, Neuronal network,...
- Business, management: Transportation networks, Industry partnerships...



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- Clustering: building relevant homogeneous groups;
- Graph drawing: giving a global representation of the graph.



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Here: **Self-Organizing Map for nonvectorial data.**



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 - Kernel SOM (on line)
 - Dissimilarity SOM (batch)
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 - A toy example
 - Social network



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The data

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- **Data:** x_1, x_2, \dots, x_n from a (possibly) non vectorial space \mathcal{G} ;
- **Kernel:** $k : \mathcal{G} \times \mathcal{G} \rightarrow \mathbb{R}$ such that k is:
 - **symmetric:** $k(x, x') = k(x', x)$;
 - **positive definite:** $\sum_{i,j=1}^m \alpha_i \alpha_j k(x_i, x_j) \geq 0$ for all $m \in \mathbb{N}$, all $(x_i)_i \in \mathcal{G}$ and all $(\alpha_i)_i \in \mathbb{R}$.



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Main consequence:

\exists Hilbert space $(\mathcal{H}, \langle \cdot, \cdot \rangle)$ and $\phi : \mathcal{G} \rightarrow \mathcal{H}$ such that:

$$k(x, x') = \langle \phi(x), \phi(x') \rangle.$$



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Nonvectorial data are mapped onto a vectorial space where a SOM can be performed by the use of k as a scalar product.



kernel SOM [Lau *et al.*, 2006]: prototypes

Practical computation:

- **Prototype** of neuron i is in \mathcal{H} and is of the form

$$p_i = \sum_{j=1}^n \gamma_{ji} \phi(x_j);$$



kernel SOM [Lau *et al.*, 2006]: prototypes

Practical computation:

- **Prototype** of neuron i is in \mathcal{H} and is of the form

$$p_i = \sum_{j=1}^n \gamma_{ji} \phi(x_j);$$

- ϕ is **implicit** as $\forall i, j = 1, \dots, n,$

$$\|\phi(x_i) - \phi(x_j)\|^2 = k(x_i, x_i) + k(x_j, x_j) - 2k(x_i, x_j);$$



kernel SOM: practical computation (online)

- **assignment step:** for x_l ,

$$\arg \min_{j=1, \dots, M} \left(\sum_{i=1}^n \gamma_{ij} k(x_l, x_i) - \sum_{i, i'=1}^n \gamma_{ij} \gamma_{i'j} k(x_i, x_{i'}) \right)$$



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- **representation step:** $p_i^l = \sum_{j=1}^M \gamma_{ji}^l \phi(x_j)$:

$$\gamma_{ji}^l = \gamma_{ji}^{l-1} + \alpha(l) h(f^l(x_l), j) (\delta_{ij} - \gamma_{ji}^{l-1})$$



Another assumptions for the data set

We are given:

- **Data:** x_1, x_2, \dots, x_n from a (possibly) non vectorial space \mathcal{G} ;
- **A dissimilarity measure:** $\delta : \mathcal{G} \times \mathcal{G} \rightarrow \mathbb{R}$ such that:
 - δ is **symmetric**: $\delta(x, x') = \delta(x', x)$;
 - δ is **positive**: $\delta(x, x') \geq 0$;
 - $\delta(x, x) = 0$.



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Adaptation for SOM:

- The **prototypes** are one of the elements of the data set $(x_i)_{i=1, \dots, n}$;



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Adaptation for SOM:

- The **prototypes** are one of the elements of the data set $(x_i)_{i=1, \dots, n}$;
- δ is used **instead of the usual distance**.



Dissimilarity SOM: practical computation

- **assignment step:** for x_i ,

$$\arg \min_{j=1, \dots, M} \delta(x_i, p_j^{l-1})$$

- **representation step:**

$$p_j^l = \arg \min_{x \in (x_{i'})_{i'=1, \dots, n}} \sum_{i=1}^n h(f^l(x_i), j) \delta(x_i, x)$$



Link with kernel SOM

A **dissimilarity built from a kernel k** :

$$\begin{aligned}\delta_{\text{mean}}(x, x') &= \|\phi(x) - \phi(x')\|^2 \\ &= k(x, x) + k(x', x') - 2k(x, x').\end{aligned}$$



Link with kernel SOM

A **dissimilarity built from a kernel** k :

$$\begin{aligned}\delta_{\text{mean}}(x, x') &= \|\phi(x) - \phi(x')\|^2 \\ &= k(x, x) + k(x', x') - 2k(x, x').\end{aligned}$$

Consequence: By generalizing dissimilarity SOM to the case where the prototype of neuron i is of the form

$$p_j = \sum_{i=1}^n \gamma_{ji} \phi(x_i);$$

we can derive a **batch version** of kernel SOM from dissimilarity SOM.



From dissimilarity SOM to batch kernel SOM

Assignment step

for x_i ,

$$\arg \min_{j=1,\dots,M} \delta(x_i, p_j^{l-1})$$

Representation step

$$p_j^l = \arg \min_{x \in (x_{i'})_{i'=1,\dots,n}} \sum_{i=1}^n h(f^l(x_i), j) \delta(x_i, x)$$



From dissimilarity SOM to batch kernel SOM

Assignment step

for x_i ,

$$\arg \min_{j=1,\dots,M} \left\| x_i - \sum_{i=1}^n \gamma_{ji} \phi(x_i) \right\|$$

Representation step

$$\gamma_j^l = \arg \min_{\gamma \in \mathbb{R}^n} \sum_{i=1}^n h(f^l(x_i), j) \left\| x_i - \sum_{l'=1}^n \gamma_{l'} \phi(x_{l'}) \right\|^2$$



From dissimilarity SOM to batch kernel SOM

Assignment step

for x_i ,

$$\arg \min_{j=1, \dots, M} \sum_{u, u'=1}^n \gamma_{ju} \gamma_{ju'} k(x_u, x_{u'}) - 2 \sum_{u=1}^n \gamma_{ju} k(x_u, x_i)$$

Representation step

$$\gamma_{ji}^l = \frac{h(f^l(x_i), j)}{\sum_{i'=1}^n h(f^l(x_{i'}), j)}$$



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Laplacian and diffusion kernel

Definitions

For a graph with vertices $V = \{x_1, \dots, x_n\}$ having positive weights $(w_{i,j})_{i,j=1,\dots,n}$ such that, for all $i, j = 1, \dots, n$, $w_{i,j} = w_{j,i}$ and $d_i = \sum_{j=1}^n w_{i,j}$,

- **Laplacian:** $L = (L_{i,j})_{i,j=1,\dots,n}$ where

$$L_{i,j} = \begin{cases} -w_{i,j} & \text{if } i \neq j \\ d_i & \text{if } i = j \end{cases} ;$$



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\Rightarrow

$$\begin{aligned} k^\beta : V \times V &\rightarrow \mathbb{R} \\ (x_i, x_j) &\rightarrow K_{i,j}^\beta \end{aligned}$$

is the **diffusion kernel** (or heat kernel).



Properties

- ① **Diffusion on the graph:** $k^\beta(x_i, x_j) \simeq$ quantity of energy accumulated in x_j after a given time if energy is injected in x_i at time 0 and if diffusion is done along the edges.
 $\beta \simeq$ intensity of diffusion;



Properties

- 1 Diffusion on the graph:** $k^\beta(x_i, x_j) \simeq$ quantity of energy accumulated in x_j after a given time if energy is injected in x_i at time 0 and if diffusion is done along the edges.
 $\beta \simeq$ intensity of diffusion;
- 2 Regularization operator:** for $u \in \mathbb{R}^n \sim V$, $u^T K^\beta u$ is higher for vectors u that vary a lot over “close” vertices of the graph.
 $\beta \simeq$ intensity of regularization (for small β , direct neighbors are more important);



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Simulated data

Random generation

50 non-weighted graphs having:

- 5 cliques $(C_i)_{i=1,\dots,5}$ with $(n_i)_{i=1,\dots,5}$ vertices where $n_i \sim \mathcal{P}(50)$;
- for $i = 1, \dots, 5$, l_i links between C_i and the other vertices
 $l_i \sim \mathcal{U}(\{1, \dots, 100n_i\})$.



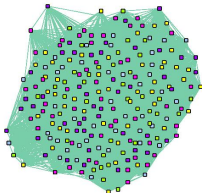
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Example:



Number of vertices: 237

Total of degrees: 12 924

Diameter: 2

Density ≈ 0.46



Compared results ($\beta = 0.1$)

	k-SOM (online)	d-SOM	k-SOM (batch)
Mean energy	0.10	0.13	0.10
Mean nb of classes	8.04	9	6.56
Mean % of good classif.	79.84	77.89	94.34
Time of computation (min)	260	80	20



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	94 %	
	6 %	2 %
		98 %

A medieval social network [Villa *et al.*, 2007]

Graph built from a large corpus of medieval contracts



From a corpus of 1000 agrarian contracts made in South West of France (1250-1350), we built a weighted graph:

- **vertices**: peasants found in the contracts;
- **edges**: number of contracts where two peasants are mentioned together.



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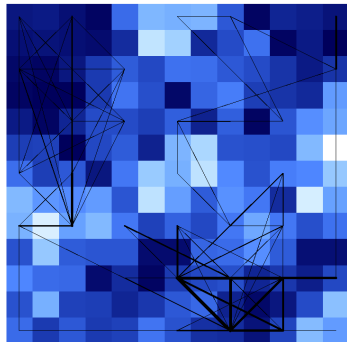
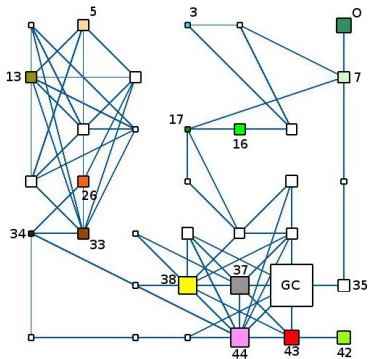
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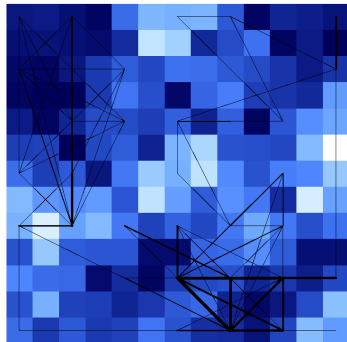
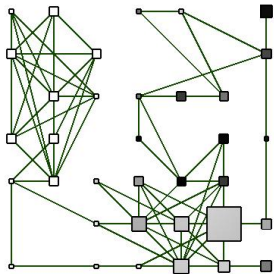
Number of vertices: 515
Number of edges: 4193
Total of weights: 40 329
Diameter: 10
Density: 2,2%



Results on a 7×7 rectangular map



Results on a 7×7 rectangular map



References



Lau, K., Yin, H. & Hubbard, S. (2006).

Kernel self-organising maps for classification.

Neurocomputing, **69**, 2033–2040.



Villa, N., Boulet, R., Rossi, F. & Jouve, B. (2007).

Mining a medieval social network by spectral clustering and kernel self-organizing map.

Submitted.

