

Stochastic Self-Organizing Map variants with the R package **SOMbrero**

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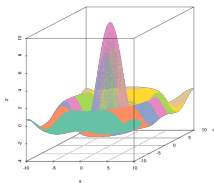
<http://www.nathalievilla.org>



WSOM 2017
Nancy, June 29th 2017

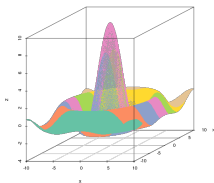
Outline

- 1 SOMbrero: an R package for stochastic SOM
- 2 KORRESP
- 3 Dissimilarity data
- 4 Applications



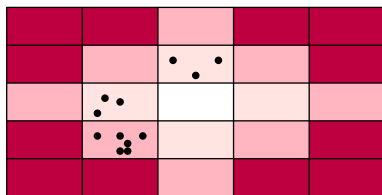
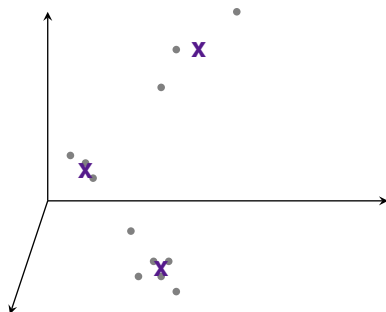
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Basics on stochastic SOM

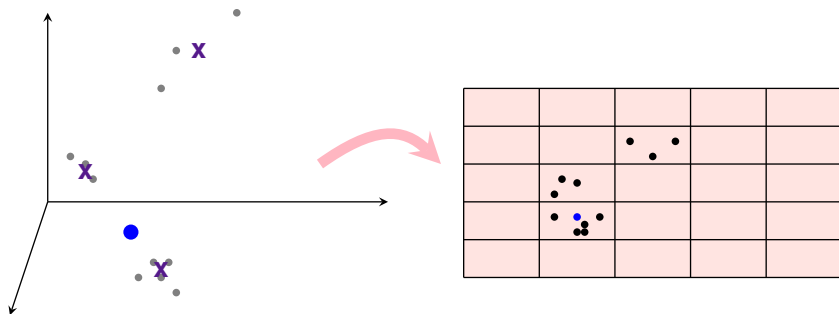
[Kohonen, 2001]



- $(\mathbf{x}_i)_{i=1,\dots,n} \subset \mathbb{R}^d$ are affected to a unit $f(x_i) \in \{1, \dots, U\}$
- the grid is equipped with a “distance” between units: $d(u, u')$ and observations affected to close units are close in \mathbb{R}^d
- every unit u corresponds to a **prototype**, $p_u(\mathbf{x})$ in \mathbb{R}^d

Basics on stochastic SOM

[Kohonen, 2001]

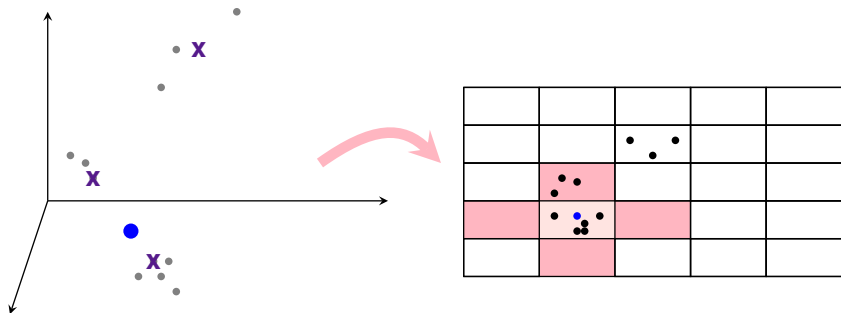


Iterative learning (assignment step): \mathbf{x}_i is picked at random within $(\mathbf{x}_k)_k$ and affected to *best matching unit*:

$$f^t(x_i) = \arg \min_u \|\mathbf{x}_i - p_u^t\|^2$$

Basics on stochastic SOM

[Kohonen, 2001]



Iterative learning (representation step): all prototypes in neighboring units are updated with a gradient descent like step:

$$p_u^{t+1} \leftarrow p_u^t + \mu(t)H^t(d(f(\mathbf{x}_i), u))(\mathbf{x}_i - p_u^t)$$

Implementations of SOM algorithm

- Matlab: SOM Toolbox [[Kohonen, 2014](#)]

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- R (CRAN only)
 - ▶ **class**: batch training, crude implementation
 - ▶ **som** (2016): training (two-step batch), basic plots and quality criteria
 - ▶ **popsom** (2017): vectorized stochastic learning (fortran), stochastic learning (C++), batch (C), several plots and quality criteria
 - ▶ **kohonen** (2017): various variants of SOM (including supervised) and super-clustering

SOMbrero

[Boelaert et al., 2014]

- **SOMbrero** is an R package implementing stochastic variants of SOM for non vectorial data

Specifically well adapted to...

non expert use and teaching

- first release: March 2013; latest release: Sept. 2016 (version 1.2)

- depends on R (version $\geq 3.1.0$) <http://www.r-project.org> and on several packages available on CRAN:



wordcloud, **igraph**, **RColorBrewer**, **scatterplot3d**, **knitr**, **shiny**

- available at <https://cran.r-project.org/package=SOMbrero> (licence GPL) and can be installed from inside R using

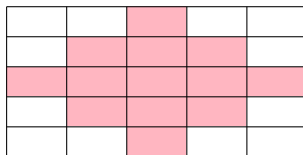
```
install.packages("SOMbrero")
```

Training

```
mysom <- trainSOM(iris[,1:4], ...)
```

Options to train the SOM:

- **grid**: square grid, with arbitrary width and length
 - ▶ *distance between units*: standard distances as in `dist` or "letremy" (Euclidean then "maximum")



- ▶ *neighborhood relationship*: Gaussian or "letremy"
- **prototypes**: initialized randomly, with a PCA, with random observations from the training sample
- **preprocessing**: centering, scaling to unit variance or nothing
- **training**: number of iterations, standard or Heskes's assignment step

$$f^t(\mathbf{x}_i) \leftarrow \arg \min_{u=1, \dots, U} \sum_{u'=1}^U H^t(d(u, u')) \|\mathbf{x}_i - p_{u'}^{t-1}\|^2$$

Diagnostic tools

quality(mysom)

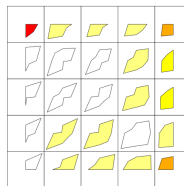
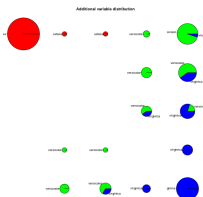
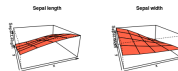
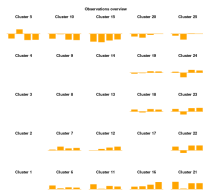
- *topographic error*: average frequency (over the samples) for which the prototypes that comes closest is in the direct neighborhood on the grid of the BMU
- *quantization error*

$$Q = \frac{1}{n} \sum_{i=1}^n \|\mathbf{x}_i - p_{f(\mathbf{x}_i)}\|^2$$



Plots...

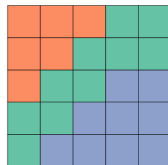
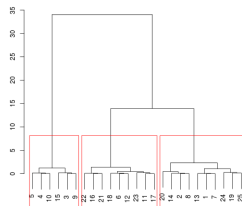
```
plot(mysom,  
     what = c("observations", "prototypes", "add"),  
     type = ..., ...)
```



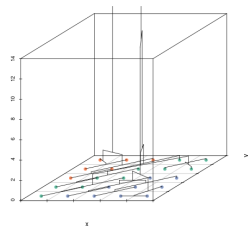
Super-clustering

```
mysom.sc <- superClass(mysom)
```

Super-clusters dendrogram



■ Super-cluster 1
■ Super-cluster 2
■ Super-cluster 3

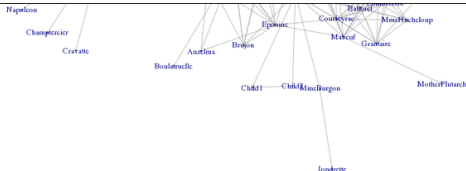


Start with SOMbrero

- 3 datasets corresponding to the three types of data that **SOMbrero** can handle (iris, presidentielles2002 and lesmis, a graph from “Les Misérables”)

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- comprehensive (HTML) vignettes included in the package and available on the website



The `dissim.lesmis` object is a matrix with entries equal to the length of the shortest path between two characters (obtained with the function `shortest_paths` of `pac` characters' names to ease the use of the graphical functions of `SOMbrero`).

Training the SOM

```
set.seed(4031719)
mis.som <- trainSOM(x.data = dissim.lesmis, type = "relational", nb.save = 10,
  _init.proto = "random")
plot(mis.som, what = "energy")
```

Energy evolution



Start with SOMbrero

- 3 datasets corresponding to the three types of data that **SOMbrero** can handle (iris, presidentielles2002 and lesmis, a graph from “Les Misérables”)
- comprehensive (HTML) vignettes included in the package and available on the website
- **Web User Interface** (made with **shiny**) for using the package even if you do not know R programming language (included in the package with `sombreroGUI()` Tested and approved on an historian!

SOMbrero Web User Interface (v0.1)

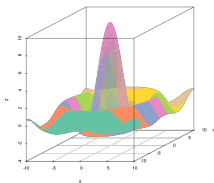
The screenshot displays the SOMbrero Web User Interface (v0.1) with the following elements:

- Navigation tabs:** Import Data, Self-Organize, Plot Map (active), Superclasses, Combine with external information, Help.
- Left Panel:**
 - Header: Select the data type: Numeric
 - 3D surface plot showing a self-organizing map.
 - Text: "Welcome to SOMbrero, the open-source on-line interface for self-organizing maps (SOM). This interface trains SOM for numerical data, contingency tables and dissimilarity data using the R package SOMbrero (v0.4). Train a map on your data and visualize their topology in three simple steps using the panels on the right."
 - Logo: SAMM (Science and Impact)
 - Text: "It is kindly provided by the SAMM team under the GPL 2.0"
- Right Panel:**
 - Section: Third step: plot the self-organizing map
 - Text: "In this panel and the next ones you can visualize the computed self-organizing map. This panel contains the standard plots used to analyze the map."
 - Section: Options
 - Plot what?: Prototypes
 - Type of plot: polygon distances
 - Checkbox: Show cluster names (checked)
 - Visualization grid: A 2x5 grid of plots showing different views of the self-organizing map, including colored polygons and outlines.



Outline

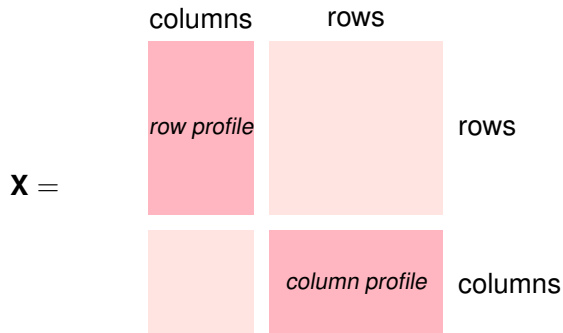
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Extensions to non vectorial data 1

KORRESP [Cottrell and Letrémy, 2005]

Data: contingency table $\mathbf{T} = (n_{ij})_{ij}$ with p rows and q columns transformed into a numeric dataset \mathbf{X} :



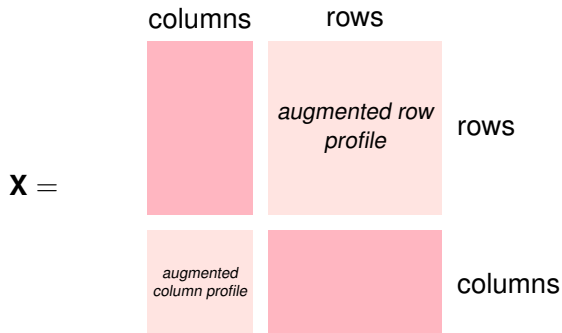
with

- $\forall i = 1, \dots, p$ and $\forall j = 1, \dots, q$, $\mathbf{x}_{ij} = \frac{n_{ij}}{n_i} \times \sqrt{\frac{n}{n_j}}$

Extensions to non vectorial data 1

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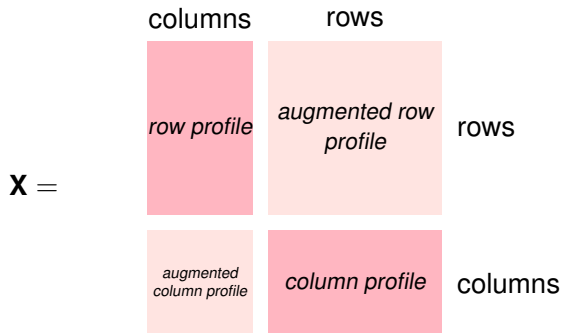
with

- $\forall i = 1, \dots, p$ and $\forall j = q + 1, \dots, q + p$, $\mathbf{x}_{ij} = \mathbf{x}_{k(i)+p,j}$ with $k(i) = \arg \max_{k=1, \dots, q} \mathbf{x}_{ik}$

Extensions to non vectorial data 1

KORRESP [Cottrell and Letrémy, 2005]

Data: contingency table $\mathbf{T} = (n_{ij})_{ij}$ with p rows and q columns transformed into a numeric dataset \mathbf{X} :



- **assignment** uses reduced profile
- **representation** uses augmented profile
- alternatively process row profiles and column profiles

Also available in **SOMbrero**

```
mysom <- trainSOM(presidentialles2002, type = "korresp")  
plot(mysom, what = "obs", type = "names")
```

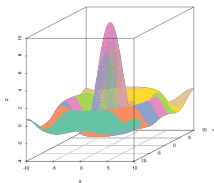
Observations overview

GLUCKSTEIN	LEPAGE #504NF		BAYROU	bas_rhin	yvelines		
	MEGRET	MADELIN		ihone			
martinique guadeloupe #1518A guyane		MAMERE CHEVENEMENT	paris			CHIRAC	
	pas de calais seine_mattime_	7000 #100 LAGUILLER BESANCONOT		hauts_de_seine_			ile et vilaine #100 loire_atlantique
			es.corse				



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In this case, data can frequently be described by a pairwise relationship:

a kernel (similarity measure) $K : \mathcal{X} \times \mathcal{X} \rightarrow \mathbb{R} \dots$

- *symmetric*: $K(z_i, z_j) = K(z_j, z_i)$
- *positive*: $\sum_{i,i'=1}^N \alpha_i \alpha_{i'} K(z_i, z_{i'}) \geq 0$

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$\Rightarrow \exists$ Hilbert space $(\mathcal{H}, \langle \cdot, \cdot \rangle)$ and a feature map $\phi : \mathcal{X} \rightarrow \mathcal{H}$:

$$K(z, z') = \langle \phi(z), \phi(z') \rangle$$

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a dissimilarity $\delta : \mathcal{X} \times \mathcal{X} \rightarrow \mathbb{R}^+ \dots$

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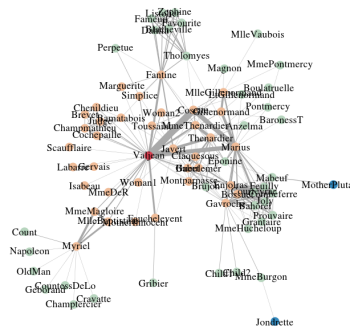
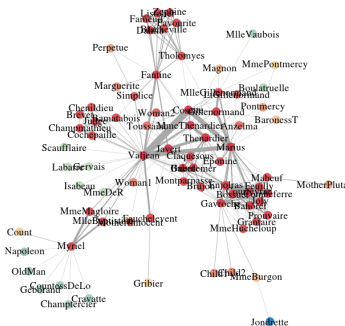
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$\Rightarrow \exists$ Euclidean spaces $(\mathcal{E}_1, \langle \cdot, \cdot \rangle_1)$ and $(\mathcal{E}_2, \langle \cdot, \cdot \rangle_2)$ and feature maps $\phi_1 : \mathcal{X} \rightarrow \mathcal{E}_1$, $\phi_2 : \mathcal{X} \rightarrow \mathcal{E}_2$:

$$\delta(z, z') = \langle \phi_1(z), \phi_1(z') \rangle_1 - \langle \phi_2(z), \phi_2(z') \rangle_2$$

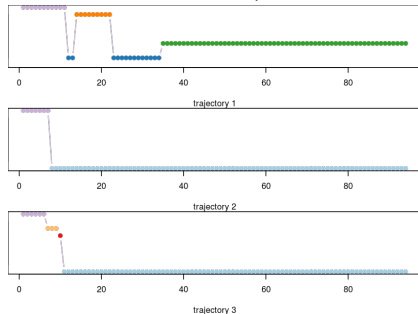
Practical examples

- kernel/dissimilarity for nodes of a graph:
 - ▶ based on the Laplacian (heat kernel / commute time kernel...) [Kondor and Lafferty, 2002, Fouss et al., 2007]
 - ▶ shortest path length



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- kernel/dissimilarity for **nodes of a graph**:
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 - ▶ shortest path length
- kernel/dissimilarity for **strings**
 - ▶ χ^2 dissimilarity emphasizes the contemporary identical situations, whether these identical moments are contiguous or not
 - ▶ Optimal-matching dissimilarities [**Needleman and Wunsch, 1970**] (or “edit distance”, “Levenshtein distance”)

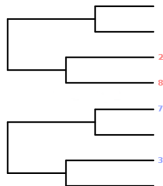


trajectory 1 is closer to trajectory 2 than to trajectory 3 according to OM distance and the opposite according to χ^2 distance



Practical examples

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 - ▶ χ^2 dissimilarity emphasizes the contemporary identical situations, whether these identical moments are contiguous or not
 - ▶ Optimal-matching dissimilarities [Needleman and Wunsch, 1970] (or “edit distance”, “Levenshtein distance”)
- in metagenomics (with phylogeny information)
 - ▶ unifrac, weighted unifrac... [Lozupone et al., 2007]



Extension of SOM to data described by a kernel or a dissimilarity

[Olteanu and Villa-Vialaneix, 2015a]

Data: $(\mathbf{x}_i)_{i=1,\dots,n} \in \mathbb{R}^d$

1: Initialization:

randomly set p_1^0, \dots, p_U^0 in \mathbb{R}^d

2: **for** $t = 1 \rightarrow T$ **do**

3: pick at random $i \in \{1, \dots, n\}$

4: **Assignment**

$$f^t(\mathbf{x}_i) = \arg \min_{u=1,\dots,U} \|\mathbf{x}_i - p_u^t\|^2$$

5: **for all** $u = 1 \rightarrow U$ **do Representation**

6:

$$p_u^{t+1} = p_u^t + \mu(t)H^t(d(f^t(\mathbf{x}_i), u))(\mathbf{x}_i - p_u^t)$$

7: **end for**

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$$f^t(\mathbf{x}_i) = \arg \min_{u=1,\dots,U} \sum_{j,j'=1}^n \beta_{uj}^t \beta_{uj'}^t K(\mathbf{x}_j, \mathbf{x}_{j'}) - 2 \sum_{j=1}^n \beta_{uj}^t K(\mathbf{x}_j, \mathbf{x}_i)$$

5: **for all** $u = 1 \rightarrow U$ **do Representation**

6:

$$\beta_u^{t+1} = \beta_u^t + \mu(t) H^t(d(f^t(\mathbf{x}_i), u)) (\mathbf{1}_i - \beta_u^t)$$

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$$p_u^{t+1} = p_u^t + \mu(t) H^t(d(f^t(\mathbf{x}_i), u)) (\sim \mathbf{x}_i - p_u^t)$$

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4: **Assignment**

$$f^t(\mathbf{x}_i) = \arg \min_{u=1,\dots,U} \sum_{j=1}^n \beta_{uj}^t \delta(\mathbf{x}_i, \mathbf{x}_j) - \frac{1}{2} \sum_{j,j'=1}^n \beta_{uj}^t \beta_{uj'}^t \delta(\mathbf{x}_j, \mathbf{x}_{j'})$$

5: **for all** $u = 1 \rightarrow U$ **do Representation**

6:

$$\beta_u^{t+1} = \beta_u^t + \mu(t) H^t(d(f^t(\mathbf{x}_i), u)) (\mathbf{1}_i - \beta_u^t)$$

7: **end for**

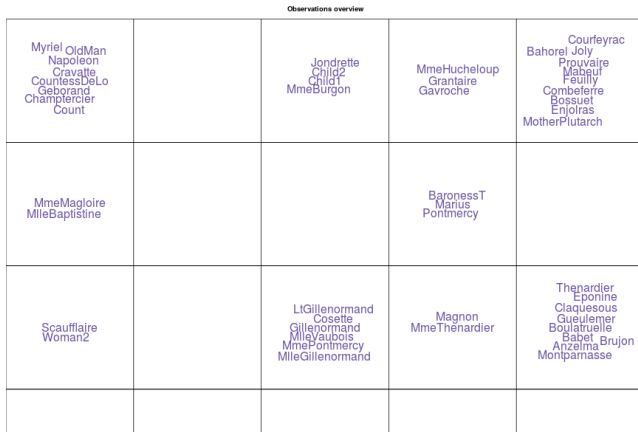
8: **end for**



Relational SOM is available in **SOMbrero**

Kernel SOM equivalent to RSOM for δ square distance induced by the kernel

```
mysom <- trainSOM(dissim.lesmis, type = "relational")  
plot(mysom, what="obs", type="names")
```



Problems with RSOM and KSOM

Data: $(\mathbf{x}_i)_{i=1,\dots,n} \in \mathcal{X}$

1: **Initialization:** $p_u^0 = \sum_{i=1}^n \beta_{ui}^0 \phi(\mathbf{x}_i)$ (convex combination)

2: **for do**

3: pick randomly $i \in \{1, \dots, n\}$

4: **Assignment**

5: **for all** $u = 1 \rightarrow U$ **do Representation**

6:

7: **end for**

8: **end for**

Problems with RSOM and KSOM

Data: $(\mathbf{x}_i)_{i=1,\dots,n} \in \mathcal{X}$

- 1: **Initialization:** $p_u^0 = \sum_{i=1}^n \beta_{ui}^0 \phi(\mathbf{x}_i)$ (convex combination)
- 2: **for** $t = 1 \rightarrow T$ **do**
- 3: pick randomly $i \in \{1, \dots, n\}$
- 4: **Assignment**

$$f^t(\mathbf{x}_i) = \arg \min_{u=1,\dots,U} \|\phi(\mathbf{x}_i) - p_u^t\|_{\mathcal{H}}^2$$

- 5: **for all** $u = 1 \rightarrow U$ **do Representation**

- 6:
$$p_u^{t+1} = p_u^t + \mu(t) (\phi(\mathbf{x}_i) - p_u^t)$$

- 7: **end for**
- 8: **end for**

Problems with RSOM and KSOM

Data: $(\mathbf{x}_i)_{i=1,\dots,n} \in \mathcal{X}$

- 1: **Initialization:** $p_u^0 = \sum_{i=1}^n \beta_{ui}^0 \phi(\mathbf{x}_i)$ (convex combination)
- 2: **for** $t = 1 \rightarrow T$ **do**
- 3: pick randomly $i \in \{1, \dots, n\}$
- 4: **Assignment**

$$f^t(\mathbf{x}_i) = \arg \min_{u=1,\dots,U} \sum_{j,j'=1}^n \beta_{uj}^t \beta_{uj'}^t K(\mathbf{x}_j, \mathbf{x}_{j'}) - 2 \sum_{j=1}^n \beta_{uj}^t K(\mathbf{x}_j, \mathbf{x}_i)$$

- 5: **for all** $u = 1 \rightarrow U$ **do** **Representation**
- 6: $\beta_u^{t+1} = \beta_u^t + \mu(t) H^t(d(f^t(\mathbf{x}_i), u)) (\mathbf{1}_i - \beta_u^t)$
- 7: **end for**
- 8: **end for**

Problems with RSOM and KSOM

Data: $(\mathbf{x}_i)_{i=1,\dots,n} \in \mathcal{X}$

- 1: **Initialization:** $p_u^0 = \sum_{i=1}^n \beta_{ui}^0 \phi(\mathbf{x}_i)$ (convex combination)
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$$f^t(\mathbf{x}_i) = \arg \min_{u=1,\dots,U} \sum_{j,j'=1}^n \beta_{uj}^t \beta_{uj'}^t K(\mathbf{x}_j, \mathbf{x}_{j'}) - 2 \sum_{j=1}^n \beta_{uj}^t K(\mathbf{x}_j, \mathbf{x}_i) \rightarrow O(n^2 U)$$

- 5: **for all** $u = 1 \rightarrow U$ **do Representation**

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- 7: **end for**
- 8: **end for**

Problems with RSOM and KSOM

Data: $(\mathbf{x}_i)_{i=1,\dots,n} \in \mathcal{X}$

- 1: **Initialization:** $p_u^0 = \sum_{i=1}^n \beta_{ui}^0 \phi(\mathbf{x}_i)$ (convex combination)
- 2: **for** $t = 1 \rightarrow \gamma n$ **do**
- 3: pick randomly $i \in \{1, \dots, n\}$
- 4: **Assignment**

$$f^t(\mathbf{x}_i) = \arg \min_{u=1,\dots,U} \sum_{j,j'=1}^n \beta_{uj}^t \beta_{uj'}^t K(\mathbf{x}_j, \mathbf{x}_{j'}) - 2 \sum_{j=1}^n \beta_{uj}^t K(\mathbf{x}_j, \mathbf{x}_i) \rightarrow O(n^2 U)$$

- 5: **for all** $u = 1 \rightarrow U$ **do Representation**

6:

$$\beta_u^{t+1} = \beta_u^t + \mu(t) H^t(d(f^t(\mathbf{x}_i), u)) (\mathbf{1}_i - \beta_u^t) \rightarrow O(nU)$$

- 7: **end for**
- 8: **end for**

→ algorithm complexity: $O(\gamma n^3 U)$ (compared to $O(\gamma U d n)$ for numeric)



Reducing the stochastic K-SOM complexity

[Mariette et al., 2017]

Data: $(\mathbf{x}_i)_{i=1,\dots,n} \in \mathcal{X}$

1: **Initialization:** $p_u^0 = \sum_{i=1}^n \beta_{ui} \phi(\mathbf{x}_i)$ (convex combination)

2: **for** $t = 1 \rightarrow \gamma n$ **do**

3: pick at random $i \in \{1, \dots, n\}$

4: **Assignment** $f^t(\mathbf{x}_i) = \arg \min_{u=1,\dots,U} \sum_{j,j'=1}^n \beta_{uj}^t \beta_{uj'}^t K(\mathbf{x}_j, \mathbf{x}_{j'}) - 2 \sum_{j=1}^n \beta_{uj}^t K(\mathbf{x}_j, \mathbf{x}_i)$

5: **for all** $u = 1 \rightarrow U$ **do Representation**

6: $\beta_u^{t+1} = \beta_u^t + \mu(t) H^t(d(f^t(\mathbf{x}_i), u)) (\mathbf{1}_i - \beta_u^t)$

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8: **end for**

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2: **for** $t = 1 \rightarrow \gamma n$ **do**

3: pick at random $i \in \{1, \dots, n\}$

4: **Assignment** $f^t(\mathbf{x}_i) = \arg \min_{u=1,\dots,U} \underbrace{\sum_{j,j'=1}^n \beta_{uj}^t \beta_{uj'}^t K(\mathbf{x}_j, \mathbf{x}_{j'})}_{A_u^t} - 2 \underbrace{\sum_{j=1}^n \beta_{uj}^t K(\mathbf{x}_j, \mathbf{x}_i)}_{B_{ui}^t}$

5: **for all** $u = 1 \rightarrow U$ **do Representation**

6: $\beta_u^{t+1} = \beta_u^t + \mu(t) H^t(d(f^t(\mathbf{x}_i), u)) (\mathbf{1}_i - \beta_u^t)$

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- 2: **for** $t = 1 \rightarrow \gamma n$ **do**
- 3: pick at random $i \in \{1, \dots, n\}$
- 4: **Assignment** $f^t(\mathbf{x}_i) = \arg \min_{u=1,\dots,U} A_u^t - 2B_{ui}^t$
- 5: **for all** $u = 1 \rightarrow U$ **do Representation**
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6: $\beta_u^{t+1} = \beta_u^t + \underbrace{\mu(t) H^t(d(f^t(\mathbf{x}_i), u))}_{\lambda_u(t)} (\mathbf{1}_i - \beta_u^t)$

7: **end for**

8: **end for**

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[Mariette et al., 2017]

Data: $(\mathbf{x}_i)_{i=1,\dots,n} \in \mathcal{X}$

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- 6: $\beta_u^{t+1} = (1 - \lambda_u(t))\beta_u^t + \lambda_u(t)\mathbf{1}_i$
- 7: **end for**
- 8: **end for**

Reducing the stochastic K-SOM complexity

[Mariette et al., 2017]

Data: $(\mathbf{x}_i)_{i=1,\dots,n} \in \mathcal{X}$

- 1: **Initialization:** $p_u^0 = \sum_{i=1}^n \beta_{ui} \phi(\mathbf{x}_i)$ (convex combination)
- 2: $A_u^0 = \sum_{j,j'=1}^n \beta_{uj}^0 \beta_{uj'}^0 K(\mathbf{x}_j, \mathbf{x}_{j'})$
- 3: $B_{ui}^0 = \sum_{j=1}^n \beta_{uj}^0 K(\mathbf{x}_j, \mathbf{x}_i)$
- 4: **for** $t = 1 \rightarrow \gamma n$ **do**
- 5: pick at random $i \in \{1, \dots, n\}$
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- 10: **end for**

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2: $A_u^0 = \sum_{j,j'=1}^n \beta_{uj'} \beta_{uj}^0 K(\mathbf{x}_j, \mathbf{x}_{j'})$

3: $B_{ui}^0 = \sum_{j=1}^n \beta_{uj}^0 K(\mathbf{x}_j, \mathbf{x}_i)$

4: **for** $t = 1 \rightarrow \gamma n$ **do**

5: pick at random $i \in \{1, \dots, n\}$

6: **Assignment** $f^t(\mathbf{x}_i) = \arg \min_{u=1,\dots,U} A_u^t - 2B_{ui}^t$

7: **for all** $u = 1 \rightarrow U$ **do Representation**

8: $\beta_u^{t+1} = (1 - \lambda_u(t))\beta_u^t + \lambda_u(t)\mathbf{1}_i$

$B_{ui'}^{t+1} = \sum_{j=1}^n \beta_{uj'}^{t+1} K(\mathbf{x}_{j'}, \mathbf{x}_j) = (1 - \lambda_u(t))B_{ui'}^t + \lambda_u(t)K(\mathbf{x}_i, \mathbf{x}_{i'})$

9: **end for**

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Data: $(\mathbf{x}_j)_{j=1,\dots,n} \in \mathcal{X}$

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$$B_{uj'}^{t+1} = \sum_{j=1}^n \beta_{uj}^{t+1} K(\mathbf{x}_j, \mathbf{x}_{j'}) = (1 - \lambda_u(t))B_{uj'}^t + \lambda_u(t)K(\mathbf{x}_i, \mathbf{x}_{j'})$$

$$A_u^{t+1} = \sum_{j,j'=1}^n \beta_{uj}^{t+1} \beta_{uj'}^{t+1} K(\mathbf{x}_j, \mathbf{x}_{j'}) = \\ (1 - \lambda_u(t))^2 A_u^t + \lambda_u(t)^2 K(\mathbf{x}_i, \mathbf{x}_i) + 2\lambda_u(t)(1 - \lambda_u(t))B_{ui}^t$$

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Reducing the stochastic K-SOM complexity

[Mariette et al., 2017]

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2: $A_u^0 = \sum_{j,j'=1}^n \beta_{uj}^0 \beta_{uj'}^0 K(\mathbf{x}_j, \mathbf{x}_{j'}) \rightarrow \mathcal{O}(n^2 U)$

3: $B_{ui}^0 = \sum_{j=1}^n \beta_{uj}^0 K(\mathbf{x}_j, \mathbf{x}_i) \rightarrow \mathcal{O}(nU)$

4: **for** $t = 1 \rightarrow \gamma n$ **do**

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4: **for** $t = 1 \rightarrow \gamma n$ **do**

5: pick at random $i \in \{1, \dots, n\}$

6: **Assignment** $f^t(\mathbf{x}_i) = \arg \min_{u=1,\dots,U} A_u^t - 2B_{ui}^t \rightarrow$ does not depend on n

7: **for all** $u = 1 \rightarrow U$ **do Representation**

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$$B_{uj'}^{t+1} = \sum_{j=1}^n \beta_{uj}^{t+1} K(\mathbf{x}_j, \mathbf{x}_{j'}) = (1 - \lambda_u(t))B_{uj'}^t + \lambda_u(t)K(\mathbf{x}_i, \mathbf{x}_{j'})$$

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[Mariette et al., 2017]

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4: **for** $t = 1 \rightarrow \gamma n$ **do**

5: pick at random $i \in \{1, \dots, n\}$

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$B_{uj'}^{t+1} = \sum_{j=1}^n \beta_{uj}^{t+1} K(\mathbf{x}_j, \mathbf{x}_{j'}) = (1 - \lambda_u(t))B_{uj'}^t + \lambda_u(t)K(\mathbf{x}_i, \mathbf{x}_{j'}) \rightarrow O(nU)$

$A_u^{t+1} = \sum_{j,j'=1}^n \beta_{uj}^{t+1} \beta_{uj'}^{t+1} K(\mathbf{x}_j, \mathbf{x}_{j'}) =$
 $(1 - \lambda_u(t))^2 A_u^t + \lambda_u(t)^2 K(\mathbf{x}_i, \mathbf{x}_i) + 2\lambda_u(t)(1 - \lambda_u(t))B_{ui}^t \rightarrow O(U)$

9: **end for**

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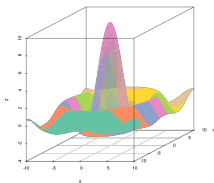
 $B_{uj'}^{t+1} = \sum_{j=1}^n \beta_{uj'}^{t+1} K(\mathbf{x}_j, \mathbf{x}_i) = (1 - \lambda_u(t))B_{uj'}^t + \lambda_u(t)K(\mathbf{x}_i, \mathbf{x}_i) \rightarrow O(nU)$
 $A_u^{t+1} = \sum_{j,j'=1}^n \beta_{uj}^{t+1} \beta_{uj'}^{t+1} K(\mathbf{x}_j, \mathbf{x}_{j'}) =$
 $(1 - \lambda_u(t))^2 A_u^t + \lambda_u(t)^2 K(\mathbf{x}_i, \mathbf{x}_i) + 2\lambda_u(t)(1 - \lambda_u(t))B_{ui}^t \rightarrow O(U)$
- 9: **end for**
- 10: **end for**

Final complexity: $O(\gamma n^2 U)$ with additional storage memory of $O(U)$ and $O(Un)$.



Outline

- 1 SOMbrero: an R package for stochastic SOM
- 2 KORRESP
- 3 Dissimilarity data
- 4 Applications

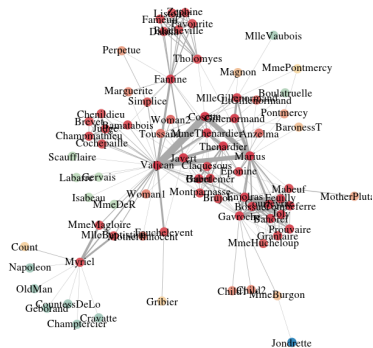


SOMbrero is well designed to handle graphs...

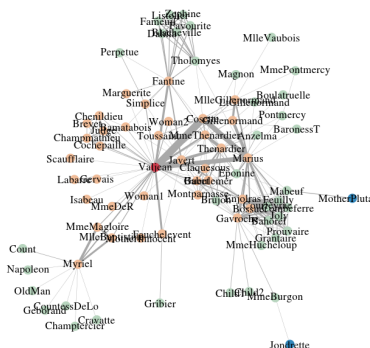
[Olteanu and Villa-Vialaneix, 2015b]

Graphs are kernel/dissimilarity data

Commute time kernel



Shortest path length



SOMbrero is well designed to handle graphs...

[Olteanu and Villa-Vialaneix, 2015b]

Graphs are kernel/dissimilarity data

Analysis of SOM results

SOM produces a map *st*:

- each node of the graph is associated to a unit of the map
- “close” nodes (according to the similarity/dissimilarity measure)

SOMbrero is well designed to handle graphs...

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Analysis of SOM results

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- each node of the graph is associated to a unit of the map
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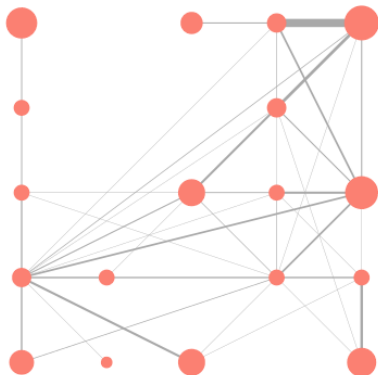
Using the result to produce a **simplified representation of the graph** (function `projectIGraph`), which is the representation of a meta-graph:

- meta-nodes are positionned on (non empty) clusters of the map;
- size of a meta-node is proportionnal to the number of nodes classified in this unit;
- edges between meta-nodes have width proportional to the number of edges between nodes of the two corresponding units.

SOMbrero is well designed to handle graphs...

[Olteanu and Villa-Vialaneix, 2015b]

Graphs are kernel/dissimilarity data

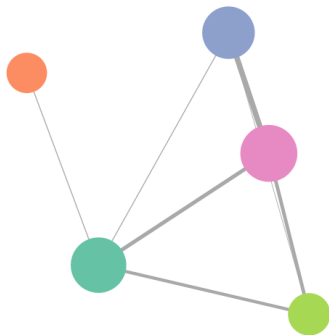


SOMbrero is well designed to handle graphs...

[Olteanu and Villa-Vialaneix, 2015b]

Graphs are kernel/dissimilarity data

Combined with super-clustering, positioning meta-nodes at the centers of gravity of the clusters on the map (superClass + projectIGraph)

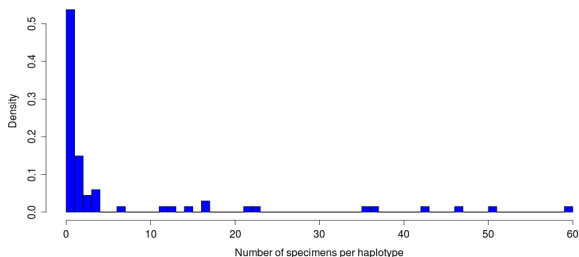


SC1: main characters (including Valjean), SC2: Bishop Myriel and related characters, SC3: Gavroche and related characters, SC4: rest of the

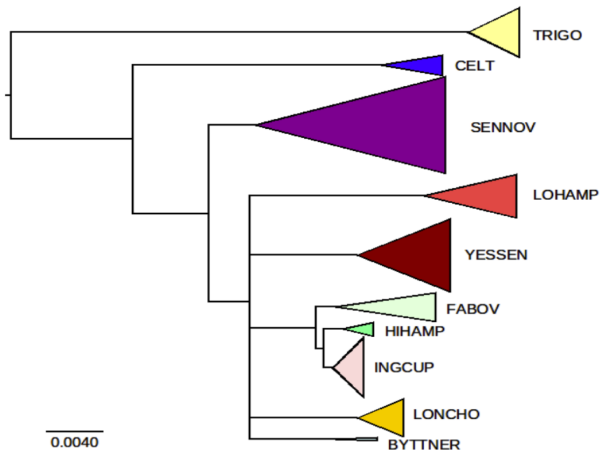


Astrartes Fulgurator data

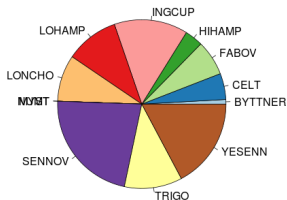
- 465 samples (67 haplotypes)
- 663 sites CO1
- Low separation level for the species (1 to 19 haplotypes within each species)
- 10 described species [[Hebert et al., 2004](#)]



Phylogenetic tree for the *Astraptes* data



Relative distribution of the ten species and MDS projection of the samples

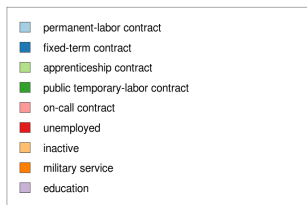
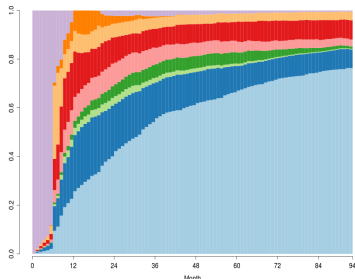


Topographic error: 0.0022



Relational SOM applied to career paths

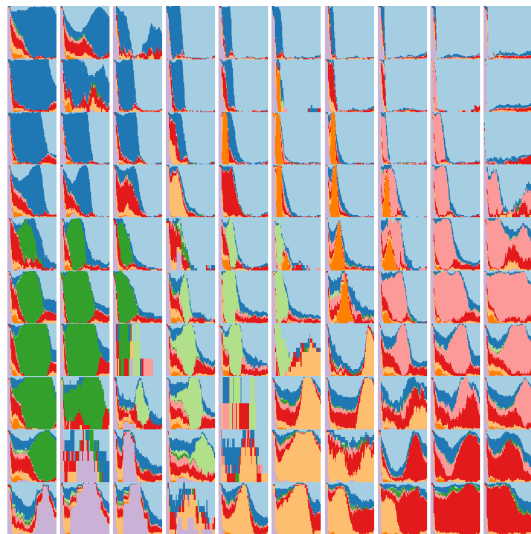
[Massoni et al., 2013]



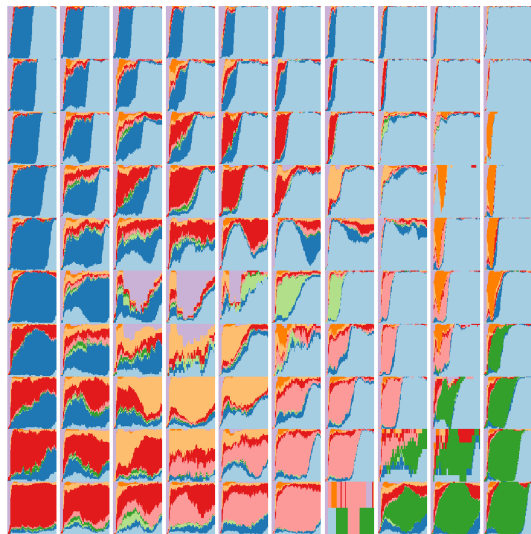
Two 10x10 maps trained on the 16040 sequences with :

- The χ^2 distance - emphasizes the contemporaneity of the situations
- The LCS distance - emphasizes the transitions, the order of the situations and the common subsequences

Results with the χ^2 distance



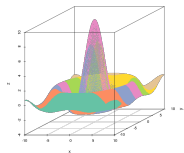
Results with the LCS distance



Perspectives

SOMbrero would benefit from:

- faster implementation using C/C++
- nicer graphics with *e.g.*, **ggplot2**
- more flexible way to handle topology of the map
- (currently under implementation) kernel computation
- better coverage with **testthat**
- public **shiny** interface
- ...



Do not hesitate to **use it** and send us your feedback!



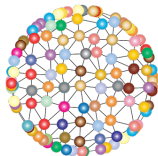
Special thanks to



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<http://user2019.fr>







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