

Clustering: from modeling to visualizing Mapping clusters as spherical Gaussians

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Clustering: from modeling to visualizing Mapping clusters as spherical Gaussians Numerical illustrations for complex data Axes interpretability Discussion

Gaussian Based Visualization of Gaussian and Non-Gaussian Based Clustering

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Take home message

Traditionally: spaces for visualizing clusters are fixed for their user-convenience Natural extension: models for visualizing clusters should follow the same principle!

Outline

1 Clustering: from modeling to visualizing

- 2 Mapping clusters as spherical Gaussians
- **3** Numerical illustrations for complex data
- 4 Axes interpretability
- **5** Discussion

Model-based clustering: pitch¹

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- **Data set:** $\mathbf{x} = (\mathbf{x}_1, \dots, \mathbf{x}_n)$, each $\mathbf{x}_i \in \mathcal{X}$ with d_X variables
- Partition (unknown): $z = (z_1, \ldots, z_n)$ with binary notation $z_i = (z_{i1}, \ldots, z_{iK})$
- Statistical model: couples (x_i, z_i) independently arise from the parametrized pdf

$$
\underbrace{f(\mathbf{x}_i, \mathbf{z}_i)}_{\in \mathcal{F}} = \prod_{k=1}^K \left[\pi_k f_k(\mathbf{x}_i) \right]^{z_{ik}}
$$

- **Example 1** Estimating *f*: implement the MLE principle through an EM-like algorithm
- **Estimating K:** use some information criteria as BIC, ICL,...
- Estimating z: use the MAP principle $\hat{z}_{ik} = 1$ iif $k = \arg \max_{\ell} t_{i\ell}(\hat{f})$ where

$$
t_{ik}(f) = p(z_{ik} = 1 | \mathbf{x}_i; f) = \frac{\pi_k f_k(\mathbf{x}_i)}{\sum_{\ell=1}^K \pi_\ell f_\ell(\mathbf{x}_i)}.
$$

 $1_{4/42}$ ¹See for instance [McLachlan & Peel 2004], [Biernacki 2017]

Model-based clustering: flexibility of *F* for complex *X*

- **Continuous data** $(X = \mathbb{R}^{d_X})$: multivariate Gaussian/t distrib. [McNicholas 2016]
- Gategorical data: product of multinomial distributions [Goodman 1974]
- Mixing cont./cat.: product Gaussian/multinomial [Moustaki & Papageorgiou 2005]
- **Functional data: the discriminative functional mixture [Bouveyron** *et al.* **2015]**
- Network data: the Erdös Rényi mixture [Zanghi *et al.* 2008]
- \blacksquare Other kinds of data, missing data, high dimension,...

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Model-based clustering: poor user-friendly understanding

- \blacksquare *n* or *K* large: poor overview of partition \hat{z}
- d_X large: too many parameters to embrace as a whole in \hat{f}_k
- Complex \mathcal{X} : specific and non trivial parameters involved in \hat{f}_k

Visualization procedures

Aim at proposing user-friendly understanding of the mathematical clustering results

Overview of clustering visualization: mapping *vs.* drawing

Visualization is the achievement of two different successive steps:

- \blacksquare The mapping step:
	- **Performs a transformation, typically space dimension reduction of a data set or of a pdf**
	- It produces no graphical output at all (deliver just a mathematical object)
- **The drawing step:**
	- Provides the final graphical display from the output of the previous mapping step
	- Usually involves classical graphical toolboxes and tunes any graphical parameters

Mathematician is first concerned by the more challenging mapping step

Overview of clustering visualization: individual mapping

- Aims at visualizing simultaneously the data set x and its estimated partition \hat{z}
- **Transforms x**, defined on X, into $\mathbf{y} = (\mathbf{y}_1, \ldots, \mathbf{y}_n)$, defined on a new space \mathcal{Y}

 $M^{\text{ind}} \in \mathcal{M}^{\text{ind}} : \mathbf{x} \in \mathcal{X}^n \mapsto \mathbf{y} = M^{\text{ind}}(\mathbf{x}) \in \mathcal{Y}^n$

- \blacksquare Many methods, depending on $\mathcal X$ definition: PCA, MCA, MFA, FPCA, MDS...
- Some of them use \hat{z} in M^{ind}: LDA, mixture entropy preservation [Scrucca 2010]
- Nearly always, $\mathcal{Y} = \mathbb{R}^2$

Model \hat{f} is is not taken into account through this approach which is focused on **x**

Overview of clustering visualization: pdf mapping

- Aims at displaying information relative to the mapping of the *f* distribution
- Transforms $f = \sum_{k} \pi_{k} f_{k} \in \mathcal{F}$, into a new mixture $g = \sum_{k} \pi_{k} g_{k} \in \mathcal{G}$

 $M^{pdf} \in M^{pdf} : f \in \mathcal{F} \mapsto g = M^{pdf}(f) \in G$

- *G* is a pdf family defined on the space $\mathcal Y$
- *M*^{pdf} is often obtained as a by product of M^{ind} (tedious outside linear mappings)
- For large *n*, *M*ind finally displays *M*pdf
- Often, both **y** and *g* are overlaid

Summary of traditional visualization strategies²

- Nature of G can dramatically depend on the choice of M^{pdf}
- \blacksquare It can potentially lead to very different cluster shapes!
- **Arguments for traditional** M^{pdf} **: user-friendly, easy-to-compute**
- Examples: linear mappings in all PCA-like methods

2Similar thinking with *^M*ind

New visualization strategy

- **If** It is the reversed situation where G is defined instead of M^{pdf}
- **O**ffer opportunity to impose directly G to be a user-friendly mixture family
- Strategy M and Strategy_G are both valid but Strategy_G is rarely explored!

This work: explore Strategy*^G*

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Spherical Gaussians as candidates

- **Users are usually familiar with multivariate spherical Gaussians on** $\mathcal{V} = \mathbb{R}^{d_Y}$
- **Thus a simple and "user-friendly" candidate** g **is a mixture of spherical Gaussians**

$$
g(\mathbf{y}; \boldsymbol{\mu}) = \sum_{k=1}^K \underset{\text{from } f}{\pi_k} \phi_{d_Y}(\mathbf{y}; \underbrace{\boldsymbol{\mu}_k}_{?}, \boldsymbol{l})
$$

where $\mu = (\mu_1, \ldots, \mu_K)$ and $\phi_{d_v}(\cdot; \mu_k, I)$ the pdf of the Gaussian distribution ■ with mean $\boldsymbol{\mu}_k = (\mu_{k1}, \ldots, \mu_{kd_{\mathcal{V}}}) \in \mathbb{R}^{d_{\mathcal{Y}}}$ ■ with covariance matrix equal to identity *I*

 $g(\cdot;\mu)$ should be then linked with *f* in order to define a sensible G

 $G = \{g : g(\cdot; \mu), \mu \in \text{arg min } \delta(f, g(\cdot; \mu)), f \in \mathcal{F}\}$

g as the "clustering twin" of *f*

Question: how to choose δ since generally $\mathcal{X} \neq \mathcal{Y}$? Answer: in our clustering context, δ should measure the clustering ability difference

Kullback-Leibler divergence of clustering ability between both *f* and $g(\cdot;\mu)^3$

$$
\delta_{\text{KL}}(f,g(\cdot;\boldsymbol{\mu})) = \int_{\mathcal{T}} \mathsf{p}_f(\boldsymbol{t}) \ln \frac{\mathsf{p}_f(\boldsymbol{t})}{\mathsf{p}_g(\boldsymbol{t};\boldsymbol{\mu})} d\boldsymbol{t}
$$

where

 \mathbf{p}_f : pdf of proba. of classification $\mathbf{t}(f) = (\mathbf{t}_i(f))_{i=1}^n$, with $\mathbf{t}_i(f) = (t_{ik}(f))_{k=1}^{K-1}$ $p_g(\cdot;\mu)$: pdf of proba. of classif. $\mathbf{t}(g)=(\mathbf{t}_i(g))_{i=1}^n$, with $\mathbf{t}_i(g)=(t_{ik}(g))_{k=1}^{K-1}$ $\mathcal{T} = \{ \bm{t} : \bm{t} = (t_1, \ldots, t_{K-1}), t_k > 0, \sum_k t_k < 1 \}$

 $3p_f$ is the reference measure

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G reduced to a unique distribution

- A natural requirement: $p_g(\cdot;\mu)$ and *g* should be linked by a one-to-one mapping
- \blacksquare Currently not true since rotations and/or translations are possible
- It means: for one distribution *f*, there is a unique optimal distribution $g(\cdot;\mu)$
- Additional constraints on $g(\cdot;\mu): d_Y = K 1$, $\mu_K = 0$, $\mu_{kh} = 0$ ($h > k$), $\mu_{kk} \ge 0$

Clustering: from modeling to visualizing Mapping clusters as spherical Gaussians Numerical illustrations for complex data Axes interpretability Discussion Estimating the Gaussian centers (pitch)

- **The Kullback-Leibler divergence** δ_{KL} has generally no closed-form
- Estimate it by the following consistent (in S) Monte-Carlo expression

$$
\hat{\delta}_{\text{KL}}(f, g(\cdot; \boldsymbol{\mu})) = \underbrace{\frac{1}{S} \sum_{s=1}^{S} \ln p_g(t^{(s)}; \boldsymbol{\mu})}_{L(\boldsymbol{\mu}; t)} + \text{cst}
$$

with *S* independent draws of conditional proba. $\mathbf{t} = (\mathbf{t}^{(1)}, \dots, \mathbf{t}^{(S)})$ from p_f

- It is the normalized (observed-data) log-likelihood function of a mixture model
- \blacksquare But, by construction, all the conditional probabilities are fixed in this mixture
- **Thus, just maximize the normalized complete-data log-likelihood** $L_{\text{comp}}(\mu; t)$ **:**
	- $K = 2$: this maximization is straightforward
	- $K > 2$: use a standard Quasi-Newton algorithm with different random initializations, for avoiding possible local optima

From a multivariate to a bivariate Gaussian mixture

- *g* is defined on \mathbb{R}^{K-1} but it is more convenient to be on \mathbb{R}^2
- **Just apply LDA** on *g* to display this distribution on its most discriminative map
- It leads to the bivariate spherical Gaussian mixture \tilde{g}

$$
\tilde{g}(\tilde{\mathbf{y}};\tilde{\boldsymbol{\mu}})=\sum_{k=1}^K \pi_k \phi_2(\tilde{\mathbf{y}};\tilde{\boldsymbol{\mu}}_k,\boldsymbol{I}),
$$

where
$$
\tilde{\mathbf{y}} \in \mathbb{R}^2
$$
, $\tilde{\boldsymbol{\mu}} = (\tilde{\boldsymbol{\mu}}_1, \dots, \tilde{\boldsymbol{\mu}}_K)$ and $\tilde{\boldsymbol{\mu}}_k \in \mathbb{R}^2$

Use the % of inertia of LDA to measure the quality of the mapping from g **to** \tilde{g}

Remark

If $\mathcal{X} = \mathbb{R}^d$ and *f* is a Gaussian mixture with isotropic covariance matrices, then the proposed mapping is equivalent to applying a LDA to the centers of *f*

Overall accuracy of the mapping between *f* and ˜*g*

Use the following difference between the normalized entropies of *f* and \tilde{g}

$$
\delta_{\mathsf{E}}(f, \tilde{\mathbf{g}}) = -\frac{1}{\ln K} \sum_{k=1}^K \left\{ \int_{\mathcal{X}} t_k(\mathbf{x}; f) \ln t_k(\mathbf{x}; f) d\mathbf{x} - \int_{\mathbb{R}^2} t_k(\tilde{\mathbf{y}}; \tilde{\mathbf{g}}) \ln t_k(\tilde{\mathbf{y}}; \tilde{\mathbf{g}}) d\tilde{\mathbf{y}} \right\}
$$

- Such a quantity can be easily estimated by empirical values
- \blacksquare Its meaning is particularly relevant:
	- $\delta_F(f, \tilde{g}) \approx 0$: the component overlap conveyed by \tilde{g} (over f) is accurate
	- $\delta_F(f, \tilde{g}) \approx 1$: \tilde{g} strongly underestimates the component overlap of *f*
	- $\delta_{\rm E}(f, \tilde{g}) \approx -1$: \tilde{g} strongly overestimates the component overlap of *f*

 $\delta_F(f, \tilde{g})$ permits to evaluate the bias of the visualization

Drawing \tilde{g}

- Cluster centers: the locations of $\tilde{\mu}_1, \ldots, \tilde{\mu}_K$ are materialized by vectors
- Cluster spread: the 95% confidence level displayed by a black border
- Cluster overlap: iso-probability curves of the MAP classification for different levels
- **Mapping accuracy:** $\delta_E(f, \tilde{g})$ and also % of inertia by axis

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20/42

House of Representatives Congressmen: data⁴ and model

- Votes of the $n = 435$ U.S. Congressmen on the $d_X = 16$ key votes
- Gategorical data: for each vote, three levels are considered (yea, nay, ?)
- Data clustered by a mixture of product of multinomial distributions [Goodman 1974]
- $K = 4$ selected by BIC [Schwarz 1974]
- Use the R package Rmixmod [Lebret *et al.* 2015]
- Gomplex output: 435 individual memberships, $192 = 16 \times 3 \times 4$ parameters

House of Representatives Congressmen: standard MCA visualization

First map of the MCA (R package FactoMineR [Lê *et al.* 2008]): difficult to interpret

House of Representatives Congressmen: Gaussian visualization

Difference between entropies: - 0.08

Mapping of *f* on this graph is accurate because $\delta_E(f, \tilde{g}) = 0.01$

Contraceptive method choice: data⁵ and model

- Subset of the 1987 National Indonesia Contraceptive Prevalence Survey
- Mixed data: 1473 Indian women with two numerical variables (age and number of children) and eight categorical variables (education level, education level of the husband, religion, occupation, occupation of the husband, standard-of-living index and media exposure)
- Clustered by a mixture *f* assuming that variables are independent within components
- Model selection is done by the BIC criterion which detects six components
- Use the R package Rmixmod [Lebret *et al.* 2015]

Contraceptive method choice: estimated parameters

Table: Parameters of the continuous variables for the Contraceptive method choice.

Table: Modes of the categorical variables for the Contraceptive method choice.

Contraceptive method choice: Gaussian visualization

Difference between entropies: -0.34

Mapping of *f* on this graph is accurate because $\delta_E(f, \tilde{g}) = 0.04$

Bike sharing system: $data^6$ and model

- Station occupancy data collected over the course of one month on the bike sharing system in Paris
- Data collected over 5 weeks, between February, 24 and March, 30, 2014, on 1189 bike stations
- Functional data: station status information (available bikes/docks) downloaded every hour from the open-data APIs of JCDecaux company
- The final data set contains 1 189 loading profiles, one per station, sampled at 1 448 time points
- Model: profiles of the stations were projected on a basis of 25 Fourier functions
- Model-based clustering of these functional data [Bouveyron *et al.* 2015] with the R package FunFEM [Bouveyron 2015]
- Retain 10 clusters

Bike sharing system: cluster of curves visualization

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Difference between entropies: - 0.05

Mapping of *f* on this graph is accurate because $\delta_E(f, \tilde{g}) = -0.03$

French political blogosphere: $data^7$ and model

- Not oriented network data: a single day snapshot of over 1 100 political blogs automatically extracted the October, 14th, 2006 and manually classified by the "Observatoire Présidentielle" project.
- \blacksquare Nodes represent hostnames (= a set of pages) and edges represent hyperlinks between different hostnames
- Gather different communities organization due to the existence of several political parties and commentators
- Assumption: authors of these blogs tend to link, by political affinities, blogs with similar political positions
- Use the graph clustering via Erdös–Rényi mixture proposed by [Zanghi et al. 2008]
- \blacksquare Use the R package mixer
- As proposed by these authors, we consider $K = 6$ components

French political blogosphere: confusion matrix

Table: Confusion matrix between the component memberships and the political party memberships.

French political blogosphere: Gaussian visualization

Difference between entropies: - 0.17

The graph slightly over-represents the component overlaps: $\delta_E(f, \tilde{g}) = -0.216$

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Correspondence between new axis $1⁸$ and initial features

We measure the overlap strength of each feature *j* by the following entropy ratio:

$$
\bar{E}_j = \frac{E_j}{\sum_{j=1}^d E_j} \qquad \in [0,1]
$$

where E_i corresponds to the marginal entropy of axis 1 re-estimated without feature j

The larger is \bar{E}_j , the more discriminant is feature *j* on axis 1

Since axis 1 is the most discriminant axis, \bar{E}_j measures the contribution of feature *j* to the mixture overlap on axis 1

 $\frac{8}{34/42}$ ⁸The same principle could be applied to all other new axes.

Prostate cancer data¹⁰: definition

- Individuals: 506 patients with prostatic cancer grouped on clinical criteria into two Stages 3 and 4 of the disease
- \blacksquare Variables: $d = 12$ pre-trial variates were measured on each patient, composed by eight continuous variables (age, weight, systolic blood pressure, diastolic blood pressure, serum haemoglobin, size of primary tumour, index of tumour stage and histolic grade, serum prostatic acid phosphatase) and four categorical variables with various numbers of levels (performance rating, cardiovascular disease history, electrocardiogram code, bone metastases)
- Some missing data: 62 missing values ($\approx 1\%$)

Use the Rmixtcomp package⁹ to estimate the mixture...

⁹https://cran.r-project.org/web/packages/RMixtComp/index.html

 10 Byar DP, Green SB (1980): Bulletin Cancer, Paris 67:477-488

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Prostate cancer data: visualization 11

Difference between entropies: 0.13

¹¹ Biernacki, C., Marbac, M., Vandewalle, V. Gaussian-Based Visualization of Gaussian and Non-Gaussian-Based Sh₍₄₂ Clustering. J Classif (2020).

Prostate cancer data: axis 1 interpretation

Features SG (and AP) are the most discriminant features

■ Thus axis 1 can be interpreted as a "SG/AP axis"

Prostate cancer data: validation of axis 1 interpretation

In the Rmixcomp package, there exists an estimation of the feature importance (which is independent of the Gaussian-based visualization)

We remark that Rmixcomp feature importance and visualization interpretation are in accordance

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Conclusion

- Generic method for visualizing the results of any model-based clustering result
- Very easy to understand output since "Gaussian-like"
- Permit visualization for any type of data, because only based on proba. of classif.
- Can be used after any existing package of model-based clustering
- The overall accuracy of the visualization is also provided

R package on the CRAN: ClusVis

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Extensions

- **Possibility to explore other pdf visualizations than Gaussians**
- However, should keep in mind simple visualizations are targeted
- **Possibility to compare/select pdf candidates through** $\delta \kappa$ **or** $\delta \epsilon$

About individual visualization

Theoretically, impossible to obtain individual visualization from pdf visualization

However, we can propose a pseudo scatter plot of x as follows

$$
\mathbf{x}_i \longmapsto \mathbf{t}_i(f) = \mathbf{t}_i(g) \stackrel{\text{bijection}}{\longmapsto} \mathbf{y}_i \in \mathbb{R}^{K-1} \stackrel{\text{LDA}}{\longmapsto} \tilde{\mathbf{y}}_i \in \mathbb{R}^2
$$

 \bullet $\ddot{\mathbf{v}}$ allows only to visualize the classification position of **x**

Example for the congressmen data set

Difference between entropies: 0.01

Gaution: do not overlay pdf and individual plots since $\tilde{\mathbf{y}} = (\tilde{y}_1, \ldots, \tilde{y}_n)$ is not necessarily drawn from a Gaussian mixture