Unsupervised learning: Anomaly detection Part II: Functional data

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Anomaly detection in functional framework

Functional isolation forest

The method

FIF parameters

Real data benchmarking

Extension of FIF: Connection to data depth

Data depth: the integrated approach

Depth for curve data

Motivation

Methodology

Computation and properties

Illustrations

Brain imaging

Anomaly detection in functional framework

Functional isolation forest

i ne metnod

FIF parameters

Real data benchmarking

Extension of FIF: Connection to data depth

Data depth: the integrated approach

Depth for curve data

Motivation

Methodology

Computation and properties

Illustrations

Brain imaging

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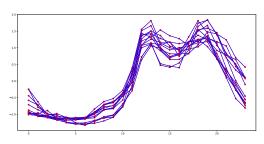
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Functional data framework

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Functional data framework

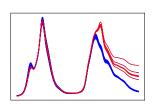
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- ➤ The first step: reconstruct **functional object** from partial observations (time-series) with interpolation or basis decomposition.



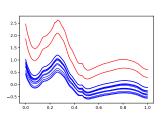
Taxonomy of functional anomalies (Hubert et al., 2015)

A non-complete taxomony of functional abnormalities:

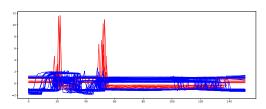
Shape anomalies



Shift anomalies

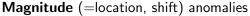


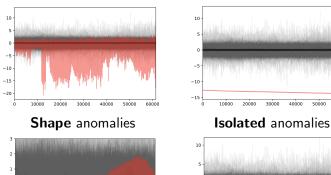
Isolated anomalies

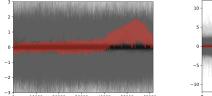


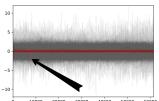
Taxonomy of functional anomalies (Airbus data)

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Anomaly detection in functional framework

Functional isolation forest

The method

FIF parameters

Real data benchmarking

Extension of FIF: Connection to data depth

Data depth: the integrated approach

Depth for curve data

Motivation

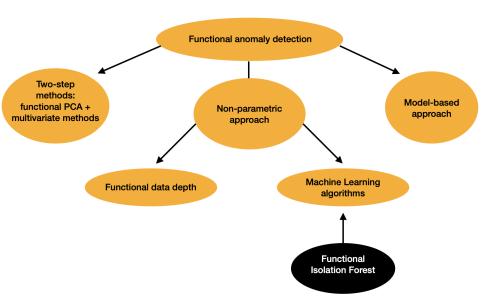
Methodology

Computation and properties

Illustrations

Brain imaging

FIF in the context of FAD contributions



Anomaly detection in functional framework

Functional isolation forest

The method

FIF parameters

Real data benchmarking

Extension of FIF: Connection to data depth

Data depth: the integrated approach

Depth for curve data

Motivation

Methodology

Computation and properties

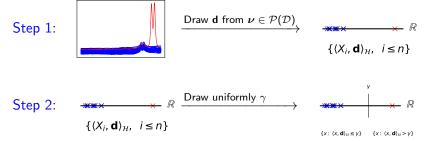
Illustrations

Brain imaging

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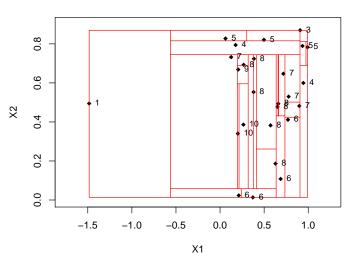
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The trick: an anomaly should be isolated faster than normal data.

Illustration: Isolation tree

Isolation tree, split 25



If a node (j, k) is non terminal, it is split in three steps as follows:

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$$\left[\min_{\mathbf{x}\in\mathcal{S}_{j,k}}\langle\mathbf{x},\mathbf{d}\rangle_{\mathcal{H}},\max_{\mathbf{x}\in\mathcal{S}_{j,k}}\langle\mathbf{x},\mathbf{d}\rangle_{\mathcal{H}}\right],$$

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3. Form the children subsets

$$\begin{array}{rcl} \mathcal{C}_{j+1,2k} & = & \mathcal{C}_{j,k} \cap \{\mathbf{x} \in \mathcal{H} : \ \langle \mathbf{x}, \mathbf{d} \rangle_{\mathcal{H}} \leq \gamma\}, \\ \mathcal{C}_{j+1,2k+1} & = & \mathcal{C}_{j,k} \cap \{\mathbf{x} \in \mathcal{H} : \ \langle \mathbf{x}, \mathbf{d} \rangle_{\mathcal{H}} > \gamma\}. \end{array}$$

as well as the children training datasets

$$S_{j+1,2k} = S_{j,k} \cap C_{j+1,2k}$$
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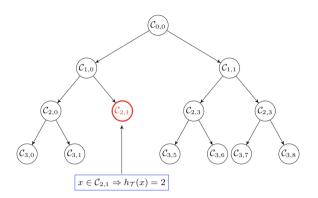
Stop when only one observation is in each node; isolation.



Anomaly score prediction

• One may then define the piecewise constant function $h_{\tau}: \mathcal{H} \to \mathbb{N}$ by: $\forall \mathbf{x} \in \mathcal{H}$,

 $h_{ au}(m{x}) = j$ if and only if $x \in \mathcal{C}_{j,k}$ and $\mathcal{C}_{j,k}$ is associated to a terminal



Anomaly score prediction

Anomaly score calculation for observation x:

- 1. For each isolation tree $i \in \{1, ..., N\}$, locate \mathbf{x} in a terminal node and calculate the depth of this node $h_i(\mathbf{x})$.
- 2. Attribute the anomaly score:

$$s_n(\mathbf{x}) = 2^{-\frac{1}{N \cdot c(n)} \sum_{i=1}^N h_i(\mathbf{x})},$$

with $c(n) = 2H(n-1) - \frac{2(n-1)}{n}$ where H(k) is the harmonic number and can be estimated by $\ln(k) + 0.5772156649$.

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Score behavior:

- when $\frac{1}{N}\sum_{i=1}^{N}h_i(\mathbf{x}) \rightarrow c(n)$, $s_n(\mathbf{x}) \rightarrow 0.5$,
- when $\frac{1}{N} \sum_{i=1}^{N} h_i(\mathbf{x}) \to 0$, $s_n(\mathbf{x}) \to 1$,
- when $\frac{1}{N}\sum_{i=1}^{N}h_i(\mathbf{x}) \rightarrow n-1$, $s_n(\mathbf{x}) \rightarrow 0$.

Anomaly detection in functional framework

Functional isolation forest

- ine method

FIF parameters

Real data benchmarking

Extension of FIF: Connection to data depth

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Depth for curve data

Motivation

Methodology

Computation and properties

Illustrations

Brain imaging

Parameters of FIF

- ► Classical parameters of ISOLATION FOREST :
 - number of trees,
 - size of the subsample,
 - height limit.

- New parameters due to the functional setup :
 - 1. The dictionary \mathcal{D} .
 - 2. The probability measure ν .
 - 3. The scalar product $\langle .,. \rangle_{\mathcal{H}}$.

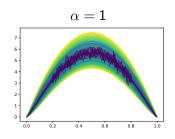
The role of the scalar product

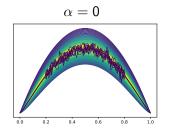
Compromise between both location and shape :

$$\langle \mathbf{f}, \mathbf{g} \rangle := \alpha \times \frac{\langle \mathbf{f}, \mathbf{g} \rangle_{L_2}}{||\mathbf{f}|| \, ||\mathbf{g}||} + (1 - \alpha) \times \frac{\langle \mathbf{f}', \mathbf{g}' \rangle_{L_2}}{||\mathbf{f}'|| \, ||\mathbf{g}'||}, \quad \alpha \in [0, 1],$$

Example on a toy dataset:

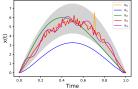
- ▶ 90 curves defined by $\mathbf{x}(t) = 30(1-t)^q t^q$ with q equispaced in [1, 1.4],
- ▶ 10 abnormal curves defined by $\mathbf{x}(t) = 30(1-t)^{1.2}t^{1.2}$ noised by $\varepsilon \sim \mathcal{N}(0, 0.3^2)$ on the interval [0.2, 0.8].

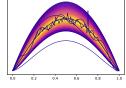


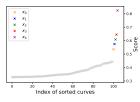


Ability to detect a variety of anomalies

- ► Sobolev inner product: $\langle .,. \rangle_{W_{1,2}}$.
- Gaussian wavelets dictionary $\mathbf{d}_{\theta,\sigma}(t) = \frac{2}{\sqrt{3\sigma}\pi^{1/4}} \left(1 \left(\frac{t-\theta}{\sigma}\right)^2\right) \exp\left(\frac{-(t-\theta)^2}{2\sigma^2}\right).$
- ightharpoonup Uniform measure ν .







Anomaly detection in functional framework

Functional isolation forest

FIF parameters

Real data benchmarking

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Motivation

Methodology

Computation and properties

Illustrations

Brain imaging

Performance on real datasets (1)

- ► FIF with 4 setups (Dictionary+scalar product):
 - ▶ Dyadic indicator (DI)+ L_2
 - ightharpoonup Cosine (Cos)+ L_2
 - Cosine (Cos)+Sobolev
 - ▶ Dataset itself (Self)+ L_2

Competitors:

- ► Isolation Forest, Local Outlier Factor, One-class SVM after dimension reduction by FPCA.
- fHD_{RP}: Random projection method with functional Halspace depth.
- ► fSDO : Functional Stahel-Donoho Outlyingness.

Performance on real datasets (2)

Methods :	DI_{L_2}	Cos _{Sob}	Cos_{L_2}	$Self_{L_2}$	IF	LOF	OCSVM	fHD_{RP}	fSDO
Chinatown	0.93	0.82	0.74	0.77	0.69	0.68	0.70	0.76	0.98
Coffee	0.76	0.87	0.73	0.77	0.60	0.51	0.59	0.74	0.67
ECGFiveDays	0.78	0.75	0.81	0.56	0.81	0.89	0.90	0.60	0.81
ECG200	0.86	0.88	0.88	0.87	0.80	0.80	0.79	0.85	0.86
Handoutlines	0.73	0.76	0.73	0.72	0.68	0.61	0.71	0.73	0.76
SonyRobotAI1	0.89	0.80	0.85	0.83	0.79	0.69	0.74	0.83	0.94
SonyRobotAI2	0.77	0.75	0.79	0.92	0.86	0.78	0.80	0.86	0.81
StarLightCurves	0.82	0.81	0.76	0.86	0.76	0.72	0.77	0.77	0.85
TwoLeadECG	0.71	0.61	0.61	0.56	0.71	0.63	0.71	0.65	0.69
Yoga	0.62	0.54	0.60	0.58	0.57	0.52	0.59	0.55	0.55
EOGHorizontal	0.72	0.76	0.81	0.74	0.70	0.69	0.74	0.73	0.75
CinECGTorso	0.70	0.92	0.86	0.43	0.51	0.46	0.41	0.64	0.80
ECG5000	0.93	0.98	0.98	0.95	0.96	0.93	0.95	0.91	0.93

Table: AUC of different anomaly detection methods calculated on the test set. Bold numbers correspond to the best result.

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Functional isolation forest

The method

FIF parameters

Real data benchmarking

Extension of FIF: Connection to data depth

Data depth: the integrated approach

Depth for curve data

Motivation

Methodology

Computation and properties

Illustrations

Brain imaging

Extension to multivariate functional data

FIF can be easily extended to the multivariate functional data, *i.e.* when the quantity of interest lies in \mathbb{R}^d for each moment of time:

$$x:[0,1] \longrightarrow \mathbb{R}^d$$

$$t \longmapsto \left((x^1(t), \ldots, x^d(t)) \right)$$

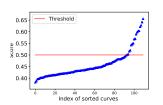
Coordinate-wise sum of the d corresponding scalar products:

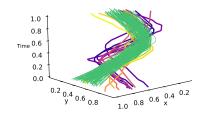
$$\langle \mathbf{f}, \mathbf{g} \rangle_{L_2^{\otimes d}} := \sum_{i=1}^d \langle f^{(i)}, g^{(i)} \rangle_{L_2}$$

▶ Dictionaries : Composed by univariate function on each axis, multivariate wavelets, multivariate Brownian motion ...

Example with MNIST dataset

We extract the digits' contours and obtain bivariate functional curves from MNIST dataset. Each digit is transformed into a curve in $(L_2([0,1]) \times L_2([0,1]))$ using length parametrization on [0,1].







Connection to data depth and supervised classification

One may define a functional depth by $D_{FIF}(x; S) = 1 - s_n(x; S)$.

Assume that we have a training classification dataset of q classes $S = S^1 \cup ... \cup S^q$.

 Low dimensional representation based on depth-based map can be defined by

$$\mathbf{x} \mapsto \phi(\mathbf{x}) = (D_{FIF}(\mathbf{x}; \mathcal{S}^1), ..., D_{FIF}(\mathbf{x}; \mathcal{S}^q)) \in [0, 1]^q.$$

▶ One may define a DD-plot classifier by using a classifier on the low dimension representation of the functional dataset.

Example of depth map on MNIST dataset

 ${\cal S}$ is constructed by taking 100 digits from class 1, 100 from class 5 and 100 from class 7.

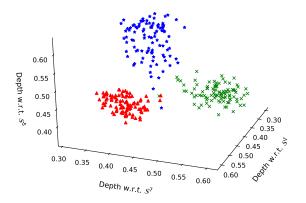


Figure: Depth space embedding of the three digits (1, 5 and 7) of the MNIST dataset.

Some remarks on FIF

- New anomaly detection algorithm for functional data:
 - Great flexibility but dictionaries (and scalar product) are tricky to choose in an unsupervised setting.
 - Low complexity and memory requierement.
- Lack of theoretical garanties!

STAERMAN, G., MOZHAROVSKYI, P., CLÉMENÇON, S., AND D'ALCHÉ-BUC, F. Functional Isolation Forest. ACML 2019.

All codes are available at: https://github.com/guillaumestaermanML/FIF.

Anomaly detection in functional framework

Functional isolation forest

The method

FIF parameters

Real data benchmarking

Extension of FIF: Connection to data depth

Data depth: the integrated approach

Depth for curve data

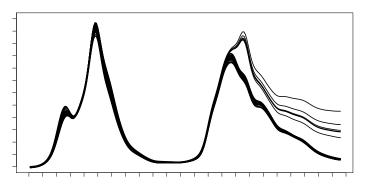
Motivation

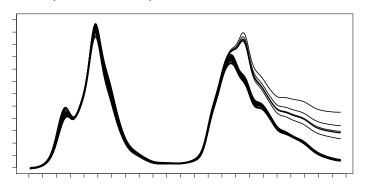
Methodology

Computation and properties

Illustrations

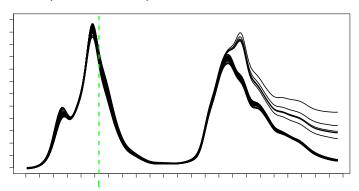
Brain imaging





▶ Functional depth of f w.r.t. $\mathcal{F} = \{f_i\}_{i=1}^n$:

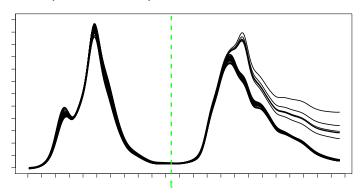
$$D(\boldsymbol{f}|\mathcal{F}) = \int_{t_{\min}}^{t^{\max}} D^1(\boldsymbol{f}(t)|\mathcal{F}(t)) dt$$
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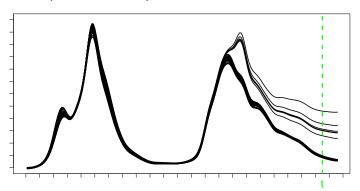
where $D^d(\cdot|\cdot)$ is a multivariate data depth, as defined above.



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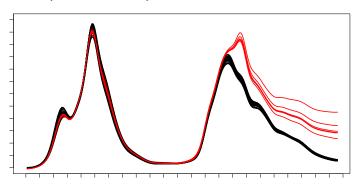
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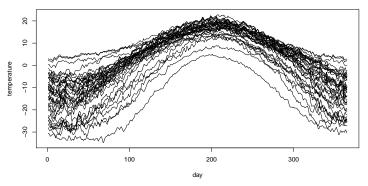
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▶ Label f as anomaly if $D(f|\mathcal{F}) < \min(D)$.

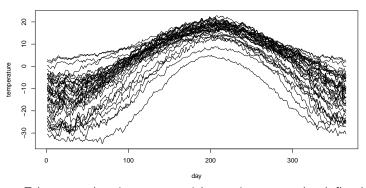


Integrated depth for functional data



Let ${\bf F}$ be a stochastic process with continuous paths defined on [0,1], and ${\bf f}$ its realization.

Integrated depth for functional data



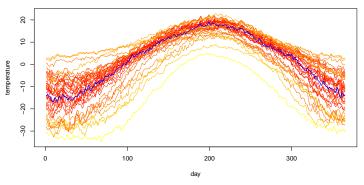
Let \mathbf{F} be a stochastic process with continuous paths defined on [0,1], and \mathbf{f} its realization. Then:

$$D(\mathbf{f}|\mathbf{F}) = \int_0^1 D(\mathbf{f}(t)|\mathbf{F}(t)) dt.$$

see Fraiman, Muniz, 2001; also López-Pintado, Romo, 2011.



Integrated depth for functional data



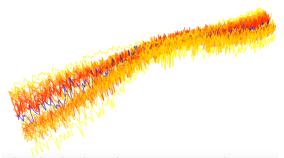
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$$D(\boldsymbol{f}|\boldsymbol{F}) = \int_0^1 \min\{F_{\boldsymbol{F}(t)}(\boldsymbol{f}(t)), 1 - F_{\boldsymbol{F}(t)}(\boldsymbol{f}(t)^-)\}dt.$$

see Fraiman, Muniz, 2001; also López-Pintado, Romo, 2011.



Multivariate functional halfspace depth



Let \mathbf{F} be a d-real-valued stochastic process with continuous paths defined on [0,1], and \mathbf{f} its realization. Then:

$$MFD(\mathbf{f}|\mathbf{F}) = \int_0^1 D(\mathbf{f}(t)|\mathbf{F}(t)) \cdot w(t)dt,$$
 $w(t) = w_{\alpha}(t, \mathbf{F}(t)) = rac{\operatorname{vol}\{D_{\alpha}(\mathbf{F}(t))\}}{\int_0^1 \operatorname{vol}\{D_{\alpha}(\mathbf{F}(u))\}du}.$

see Claeskens, Hubert, Slaets, Vakili, 2014.



Contents

Anomaly detection in functional framework

Functional isolation forest

i ne metnod

FIF parameters

Real data benchmarking

Extension of FIF: Connection to data depth

Data depth: the integrated approach

Depth for curve data

Motivation

Methodology

Computation and properties

Illustrations

Brain imaging

Practical session



Contents

Anomaly detection in functional framework

Functional isolation forest

The method

FIF parameters

Real data benchmarking

Extension of FIF: Connection to data depth

Data depth: the integrated approach

Depth for curve data

Motivation

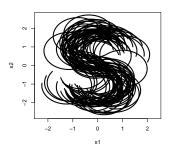
Methodology

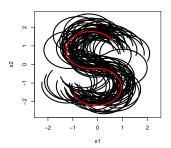
Computation and properties

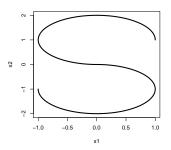
Illustrations

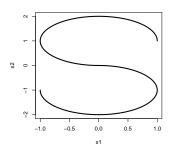
Brain imaging

Practical session







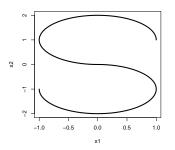


Regard the following different parametrizations of a curve:

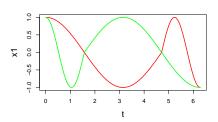
Parametrization A:

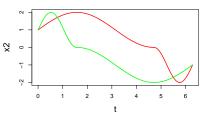
$$\begin{array}{l} x_1(t) = -\big(\cos(t) + 1\big)\mathbb{1}\{t < \frac{3\pi}{2}\} - \big(\cos(3t - 3\pi) + 1\big)\mathbb{1}\{t \geq \frac{3\pi}{2}\} + 1 \\ x_2(t) = \big(\sin(t) + 1\big)\mathbb{1}\{t < \frac{3\pi}{2}\} - \big(\sin(3t - 3\pi) + 1\big)\mathbb{1}\{t \geq \frac{3\pi}{2}\} \\ \text{Parametrization B:} \end{array}$$

$$\begin{array}{ll} x_1(t) = -\big(\cos(3t) + 1\big)\mathbb{1}\{t < \frac{\pi}{2}\} - \big(\cos(t+\pi) + 1\big)\mathbb{1}\{t \ge \frac{\pi}{2}\} + 1 \\ x_2(t) = & \big(\sin(3t) + 1\big)\mathbb{1}\{t < \frac{\pi}{2}\} - \big(\sin(t+\pi) + 1\big)\mathbb{1}\{t \ge \frac{\pi}{2}\} \end{array}$$

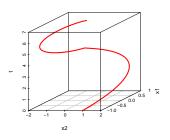


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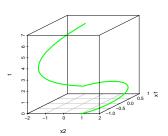




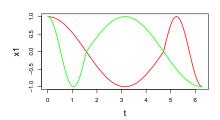
Parametrization A

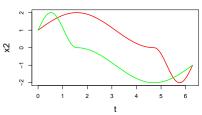


Parametrization B

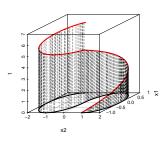


Parametrization:

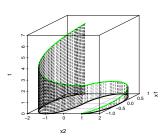




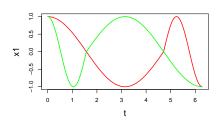
Parametrization A

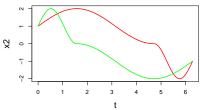


Parametrization B

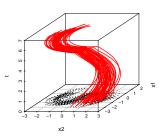


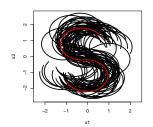
Parametrization:



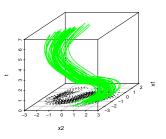


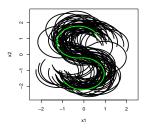
Parametrization A



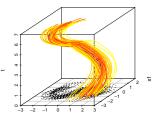


Parametrization B

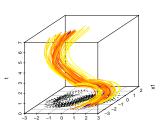




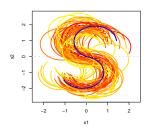
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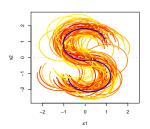


Parametrization B

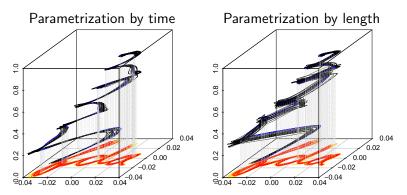


The depth-induced orders differ! x2





Functional halfspace depth for the FDA-data

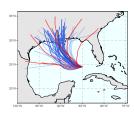


Depth-induced ranking for parametrizations by time and by length:

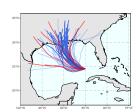
Time	2	3	13	12	4	8	1	17	11	9	7	19	15	20	18	16	14	5	6	10
Length	6	3	16	14	5	7	13	11	1	17	2	19	8	20	12	18	15	4	9	10

Simulated hurricane tracks: curve boxplot

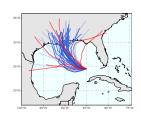
MFH depth - par. time



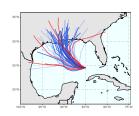
mSB depth - par. time



MFH depth - par. length



mSB depth - par. length



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▶ We endow 𝔻 with the Fréchet *metric*:

$$d_{\mathfrak{B}}\left(\mathcal{C}_{1},\mathcal{C}_{2}\right)=\inf\left\{\|\beta_{1}-\beta_{2}\|_{\infty},\beta_{1}\in\mathcal{C}_{1},\;\beta_{2}\in\mathcal{C}_{2}\right\},\quad\mathcal{C}_{1},\mathcal{C}_{2}\in\mathfrak{B}.$$



► n

- r
- ▶ Let C be an unparameterized curve. The *length of* C:

$$L(\mathcal{C}) = \sup_{\tau} \left\{ \sum_{i=1}^{N} |\beta(\tau_i) - \beta(\tau_{i-1})|_2 : \tau \text{ is a partition of } [0,1] \right\},$$
 for all $\beta \in \mathcal{C}$.

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An unparametrized curve \mathcal{C} is called *rectifiable* if $L(\mathcal{C})$ is finite. The length $L: \mathfrak{B} \to \mathbb{R} + \cup \{\infty\}$ is measurable:

$$\mathcal{P} = \Big\{ P \text{ prob. measure on } (\mathfrak{B}, d_{\mathfrak{B}}) \ : \ P(\{\mathcal{C} \in \mathfrak{B}; 0 < \mathit{L}(\mathcal{C}) < \infty\}) = 1 \Big\}.$$

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- We derive the probability distribution Q_P on \mathbb{R}^d as follows: if $X \sim Q_P$, then distribution of $X \mid \mathcal{X} = \mathcal{C}$ is the (uniform on \mathcal{C}) probability distribution $\mu_{\mathcal{C}}$:

$$\mu_{\mathcal{C}}(A) = \int_{\mathcal{C}} \mathbb{1}_{A}(x) dx.$$

The statistical model:

$$\mathcal{X}_1,\ldots,\mathcal{X}_n$$
 i.i.d. from P .

For Monte-Carlo estimation, we can consider the following **sampling scheme**:

$$\left\{ \begin{array}{l} \mathcal{X}_1, \dots, \mathcal{X}_n \text{ i.i.d. from } P, \\ \text{for all } i = 1, \dots, n \\ X_{i,1}, \dots, X_{i,m} \text{ i.i.d. from } \mu_{\mathcal{X}_i}. \end{array} \right.$$

Data depth for an unparametrized curve

Definition

The **Tukey curve depth** of $C \in \mathfrak{B}$ w.r.t. Q_P is defined as:

$$D(C|Q_P) = \int_C D(\mathbf{x}|Q_P, \mu_C) d\mu_C(\mathbf{x}),$$

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The sample Tukey curve depth of $C \in \mathfrak{B}$ w.r.t. $\mathcal{X}_1, ..., \mathcal{X}_n$ is:

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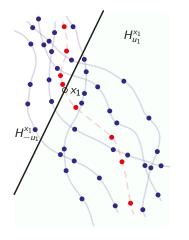
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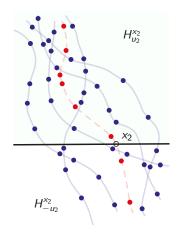
$$D(C|\mathcal{X}_1,\ldots,\mathcal{X}_n)=\int_{C}D(\boldsymbol{x}|Q_n,\mu_C)d\mu_C(\boldsymbol{x}),$$

where
$$Q_n = (\mu_{\chi_1} + \cdots + \mu_{\chi_n})/n$$
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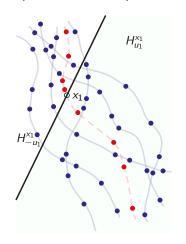


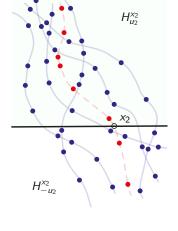
Data depth for an unparametrized curve: intuition





Data depth for an unparametrized curve: intuition





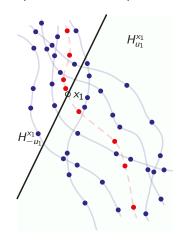
Traditional reasoning:

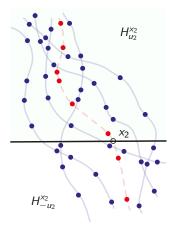
$$\widehat{Q}_{P}(H_{u_{1}}^{x_{1}}) = \frac{25}{40}, \ \widehat{\mu}_{C}(H_{u_{1}}^{x_{1}}) = \frac{4}{8}
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Curve-based reasoning:

$$\widehat{Q}_{P}(H_{u_{2}}^{x_{2}}) = \frac{25}{40}, \ \widehat{\mu}_{C}(H_{u_{2}}^{x_{2}}) = \frac{6}{8}
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Data depth for an unparametrized curve: intuition





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Let a chosen curve consist of two (independently drawn on \mathcal{C}) parts $\mathbb{Y}_{1,m} = (Y_{1,1}, \ldots, Y_{1,m})$ and $\mathbb{Y}_{2,m} = (Y_{2,1}, \ldots, Y_{2,m})$ with empirical distribution

$$\widehat{\mu}_m = \frac{1}{m} \sum_{i=1}^m \delta_{Y_{1,i}},$$

where $\delta_{\mathbf{x}}$ is the Dirac measure in $\mathbf{x} \in \mathbb{R}^d$.

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Let $\widehat{Q}_{n,m}$ be the empirical distribution (observed sample) $\mathbb{X}_{n,m} = \{X_{i,i}, i = 1, ..., n, j = 1, ..., m\}$

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► To compute the sample Tukey curve depth, a Monte Carlo approximation is used.



Let H be a closed halfspace in \mathbb{R}^d and $\mathcal{H}^{n,m}_{\Delta}$ denote a collection of such halfspaces such that for all $H \in \mathcal{H}^{n,m}_{\Delta}$ either $\widehat{Q}_{n,m}(H) = 0$ or $\widehat{\mu}_m(H) > \Delta$, almost surely, for $\Delta \in (0,\frac{1}{2})$.

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Definition

The Monte Carlo approximation of the Tukey curve depth of C w.r.t. $\mathcal{X}_1, ..., \mathcal{X}_n$ is defined as:

$$\widehat{D}_{n,m,\Delta}(\mathcal{C}|\mathcal{X}_1,...,\mathcal{X}_n) = \frac{1}{m} \sum_{i=1}^m \widehat{D}(Y_{2,i}|\widehat{Q}_{n,m},\widehat{\mu}_m,\mathcal{H}^{n,m}_{\Delta}),$$

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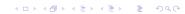
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with the depth of an arbitrary point $\pmb{x} \in \mathbb{R}^d$ w.r.t. $\widehat{Q}_{n,m}$ being:

$$\widehat{D}(\mathbf{x}|\widehat{Q}_{n,m},\widehat{\mu}_m,\mathcal{H}^{n,m}_{\Delta}) = \inf\{\frac{\widehat{Q}_{n,m}(H)}{\widehat{\mu}_m(H)} : H \in \mathcal{H}^{n,m}_{\Delta}, \mathbf{x} \in \partial H\}$$

and $\frac{0}{0} = 0$ as before.



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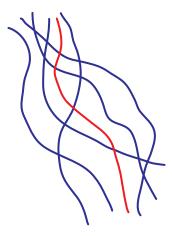
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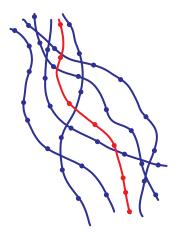
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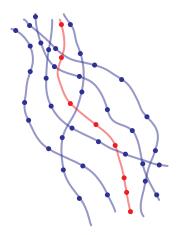
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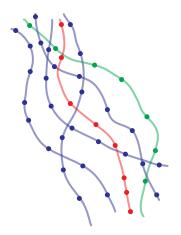
Brain imaging

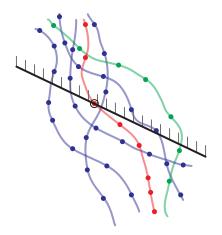
Practical session

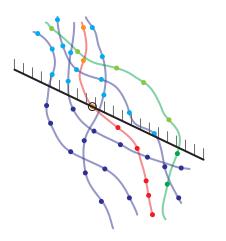




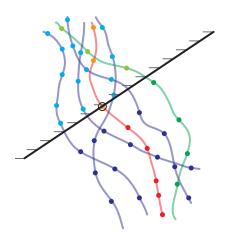




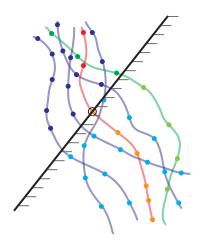




$$D(\mathbb{Y}_{2,c}|Q_m,\mathcal{H}_{m,b}) = \frac{\frac{1}{5}\left(\frac{5}{7} + \frac{3}{8} + \frac{6}{8} + \frac{2}{7} + \frac{3}{6}\right)}{\frac{2}{8}} = 2.1$$



$$D(\mathbb{Y}_{2,c}|Q_m,\mathcal{H}_{m,b}) = \frac{\frac{1}{5}\left(\frac{3}{7} + \frac{5}{8} + \frac{4}{8} + \frac{3}{7} + \frac{3}{6}\right)}{\frac{2}{8}} = 1.9857$$



$$D(\mathbb{Y}_{2,c}|Q_m,\mathcal{H}_{m,b}) = \frac{\frac{1}{5}\left(\frac{4}{7} + \frac{3}{8} + \frac{4}{8} + \frac{4}{7} + \frac{4}{6}\right)}{\frac{5}{8}} = 0.7159$$

Theorem

Let $\mathcal{C} \in \mathfrak{B}$ be a rectifiable curve, and let P be a probability measure in the space of curves such that $P \in \mathcal{P}$. Let (Δ_m) be a decreasing sequence of positive numbers such that (Δ_m) and $(\sqrt{\frac{\log(m)}{m}}/\Delta_m^2)$ converges to zero when $m \to \infty$.

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Then:

▶ the Monte Carlo approximation $\widehat{D}_{n,m,\Delta_m}(\mathcal{C}|\mathcal{X}_1,...,\mathcal{X}_n)$ converges in probability to $D(\mathcal{C}|\mathcal{X}_1,...,\mathcal{X}_n)$ as $m \to \infty$;

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- ▶ the sample Tukey curve depth $D(C|X_1,...,X_n)$ converges in probability to D(C|P) as $n \to \infty$.

Data depth for an unparametrized curve: properties

Restrict to \mathfrak{B}_{ℓ} , the subset of unparametrized curves of positive length bounded by $\ell>0$. Then the Tukey curve depth satisfies the following properties:

► Nonnegativity and boundedness by one:

$$D(C|Q_P) \in [0,1]$$
.

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$$D(f \circ C|Q_{P_f}) = D(C|Q_P).$$

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Vanishing at infinity:

$$\lim_{d_{\mathbb{D}}(\mathcal{C},\mathbf{0})\to\infty,\mathcal{C}\in\mathfrak{B}_{\ell}}D(\mathcal{C},Q_{P})=\inf_{\mathcal{C}\in\mathfrak{B}_{\ell}}D(\mathcal{C},Q_{P})=0\,.$$

Contents

Anomaly detection in functional framework

Functional isolation forest

FIF parameters

Real data benchmarking

Extension of FIF: Connection to data depth

Data depth: the integrated approach

Depth for curve data

Motivation

Methodology

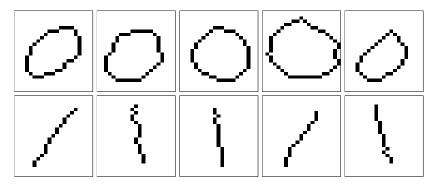
Computation and properties

Illustrations

Brain imaging

Practical session

Some examples:



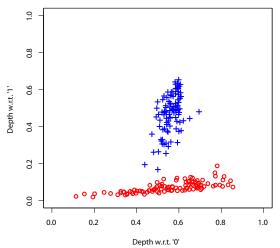
Given: training sample $S_0 = \{C_1, ..., C_m\}$ stemming from P_0 and $S_1 = \{C_{m+1}, ..., C_{m+n}\}$ stemming from P_1 in \mathfrak{B} .

Find: classifier $g: \mathfrak{B} \to \{0,1\}$ best separating P_0 and P_1 .

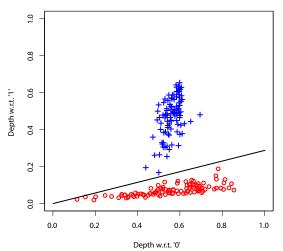


$$\mathbf{Z} = \{\mathbf{z}_i \, : \, \mathbf{z}_i = (D(C_i|Q_{P_0}), D(C_i|Q_{P_1})), \, i = 1, ..., m+n\}.$$

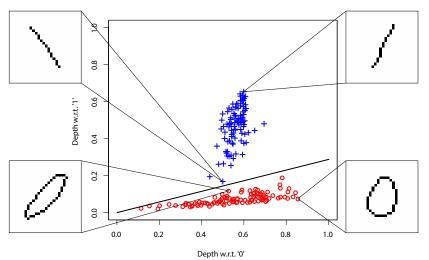
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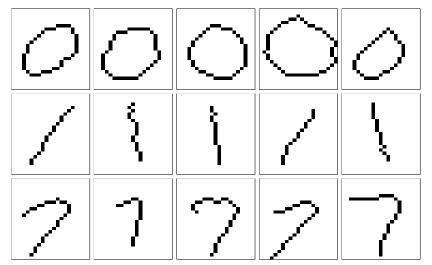


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Unsupervised classification: MNIST ("0", "1", and "7")

Some examples:



Task: Find reasonable grouping with data depth (Jörnsten '04).

Unsupervised classification: MNIST ("0", "1", and "7")

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Let $\{\mathcal{C}_1,...,\mathcal{C}_{\sum_j n_j}\}$ be the observed sample and let I_j , j=1,...,J denote the corresponding partitioning into J clusters (indices of observations belonging to each cluster j) with $\bigcup_j I_j = \{1,...,\sum_j n_j\}$ and $I_{j_1} \cap I_{j_2} = \emptyset$ for all $j_1 \neq j_2$.

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- ▶ Define the silhouette width of an observation i belonging to cluster j as
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$$Sil_i^j = \frac{\bar{d}_i^{-J} - \bar{d}_i^J}{\max\{\bar{d}_i^{-j}, \bar{d}_i^j\}},$$

where $\bar{d}_i^J = \frac{1}{\#I_j-1} \sum_{i' \in I_j, \, i' \neq i} d_{\mathfrak{B}}(\mathcal{C}_i, \mathcal{C}_{i'})$ and $\bar{d}_i^{-j} \in \operatorname{argmin}_{j' \neq j} \frac{1}{\#I_{j'}} \sum_{i' \in I_{j'}} d_{\mathfrak{B}}(\mathcal{C}_i, \mathcal{C}_{i'})$ are average distances to the observations in its own cluster and in the closest among foreign clusters respectively.

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► The **relative depth** is defined as

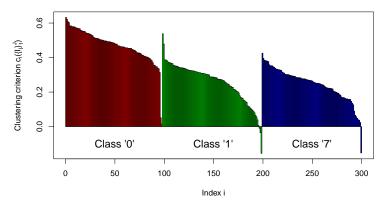
$$ReD_i^j = D(\mathcal{C}_i|\{\mathcal{C}_{i'}\}_{i' \in I_j}) - \max_{i' \neq i} D(\mathcal{C}_i|\{\mathcal{C}_{i'}\}_{i' \in I_{j'}}).$$

Clustering criterion:

$$C(\{I_j\}_1^J) = \frac{1}{\sum_j n_j} \sum_{j=1}^J \sum_{i \in I_j} c_i(\{I_j\}_1^J),$$

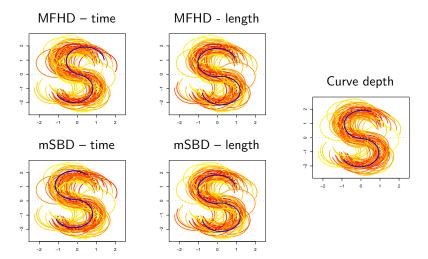
with the observation-wise clustering criterion:

$$c_i(\{I_j\}_1^J) = (1-\lambda)Sil_i^j + \lambda ReD_i^j$$
.



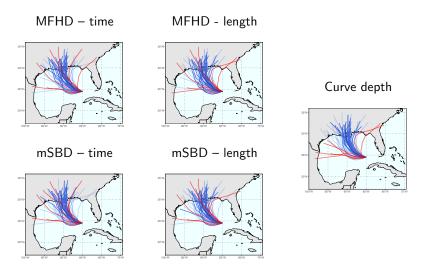
Comparison with functional depth: Example 1

Simulated S letters: depth-induced ranking



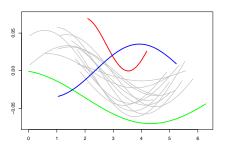
Comparison with functional depth: Example 2

Simulated hurricane tracks: curve boxplot

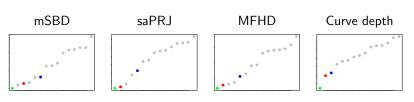


Comparison with functional depth: Anomaly detection 1

Data set 1 with introduced anomalies:

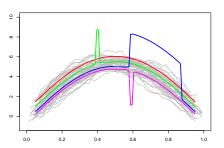


Ordered depth values:

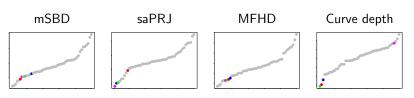


Comparison with functional depth: Anomaly detection 2

Data set 2 with introduced anomalies:



Ordered depth values:



Contents

Anomaly detection in functional framework

Functional isolation forest

The method

FIF parameters

Real data benchmarking

Extension of FIF: Connection to data depth

Data depth: the integrated approach

Depth for curve data

Motivation

Methodology

Computation and properties

Illustrations

Brain imaging

Practical session

Application: brain imaging - OATS data

► The **Older Australian Twins Study** (OATS) includes diffusion tensor magnetic resonance images (DTI) of 34 twin pairs: 11 dizygotic (DZ) and 23 monozygotic (MZ).

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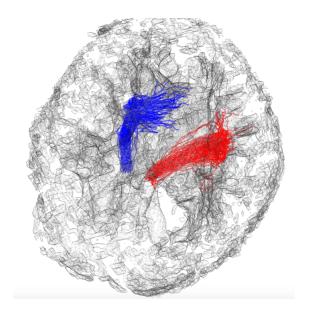
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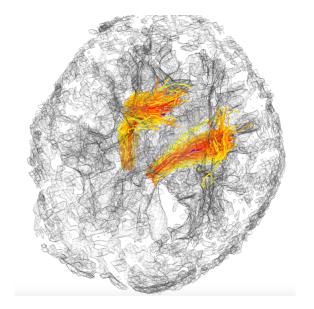
- ► **Information compression** for better understanding of brain functioning.
- Outlier detection for indication of wrongly tracked fibers.
- ► Curve registration for aligning data from different individuals before further analysis.
- Studying genetic dependency (DZ vs. MZ) for identifying disease causes.



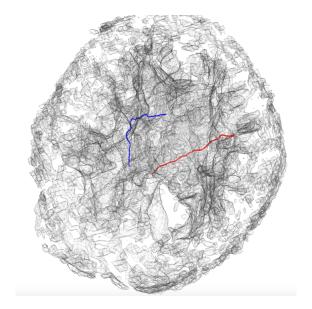
Application: brain imaging



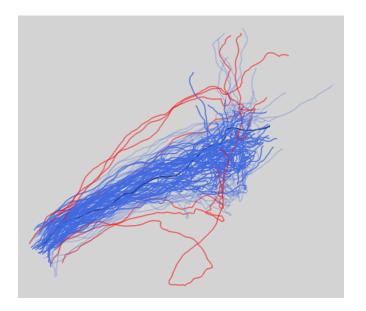
Application: brain imaging – depth-based ordering



Application: brain imaging – information compression



Application: brain imaging, right stem – outlier detection

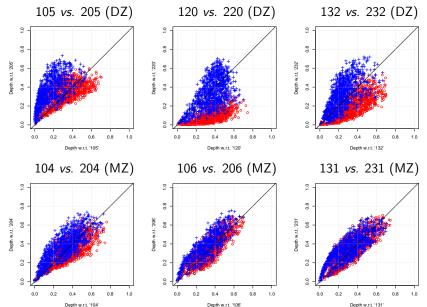


Application: brain imaging, right stem - registration



- ► The red and the dark blue curves are respectively the **deepest** curves before registration of the respective subject and subject **235**, the subject whose deepest curve is the **deepest of all**.
- ▶ We **bring** the red curve **as close as possible** (in terms of the *distance*) to the black curve. The transformed curve (after registration) is the light blue curve.
- ▶ **Distances** from each curve to the deepest one (dark blue) before (red) and after (light blue) registration are 10.271 and 3.245 (for subject 104), 4.539 and 3.395 (for subject 110), 3.329 and 2.084 (for subject 131), respectively.

Application: brain imaging, right stem - twins comparison



Contents

Anomaly detection in functional framework

Functional isolation forest

The method

FIF parameters

Real data benchmarking

Extension of FIF: Connection to data depth

Data depth: the integrated approach

Depth for curve data

Motivation

Methodology

Computation and properties

Illustrations

Brain imaging

Practical session

Thank you for attention! (and a short list of literature)

- ► Chandola, V., Banerjee, A., and Kumar, V. (2009). Anomaly detection: A survey. ACM Computing Surveys (CSUR), 41(3):15, 1–58.
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- ▶ Mosler, K. (2013). Depth statistics. In: Robustness and Complex Data Structures: Festschrift in Honour of Ursula Gather, 17—34.
- Hubert, M., Rousseeuw, P.J., and Segaert, P. (2015). Multivariate functional outlier detection. Statistical Methods & Applications, 24(2), 177—202.



Practical session (part II)

Notebooks:

- ▶ anomdet_simulation1.Rmd,
- anomdet_hurricanes.Rmd,
- anomdet_cars.ipynb,
- anomdet_airbus.ipynb.

Data sets:

- carsanom.csv: Data set on anomaly detection for cars.
- airbus_data.csv: Data set from Airbus.
- hurdat2-1851-2019-052520.txt: Historical hurricane data.

Supplementary scripts:

- ▶ depth_routines.py: Routines for data depth calculation.
- ► FIF.py: Implementation of the functional isolation forest.
- depth_routines.R: Routines for curves' parametrization.

Literature (mentioned in the tutorial) (1)

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Literature (mentioned in the tutorial) (4)

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