### Anomaly detection Part I: Multivariate data

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Tutorial for the chair DSAIDIS

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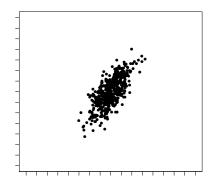
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#### A real task

Regard two measurements during a test in a production process:

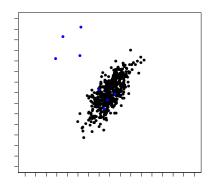


Given training data, polluted or not with anomalies:

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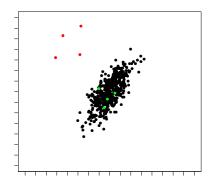
#### For new data, determine:

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$$\mathbf{X} = \{\mathbf{x}_1, ..., \mathbf{x}_n\} \subset \mathbb{R}^d$$

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- Construct a decision function:

$$\mathbb{R}^d \to \{-1,+1\} : \boldsymbol{x} \mapsto g(\boldsymbol{x}),$$

which attributes to any (possible)  $\mathbf{x} \in \mathbb{R}^d$  a label whether it is an anomaly (e.g., +1) or a normal observation (e.g., -1).

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▶ It is more useful to provide an ordering on  $\mathbb{R}^d$ :

$$\mathbb{R}^d \to \mathbb{R} : \mathbf{x} \mapsto g(\mathbf{x}),$$

such that abnormal observations obtain higher anomaly score.



### Practical session (parts I and II)

#### Notebooks:

- anomdet\_simulation1.Rmd,
- anomdet\_hurricanes.Rmd,
- anomdet\_brainimaging.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.
- ▶ 101\_1\_dwi\_fa.nii: Anatomical brain volume data.
- ▶ 101\_1\_dwi.voxelcoordsL.txt: Left brain fiber's bundle.
- ▶ 101\_1\_dwi.voxelcoordsR.txt: Right brain fiber's bundle.

#### 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.
- ▶ DTI.R: Routines for input of brain imaging data: <=> <=> > = > > <=>

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- Generalized portrait is the vector:

$$\psi = rac{arphi}{\min_{oldsymbol{x} \in oldsymbol{X}} \langle oldsymbol{x}, oldsymbol{arphi} 
angle} \quad ext{with } arphi ext{ from} \quad \max_{\|oldsymbol{arphi}\| = 1} \min_{oldsymbol{x} \in oldsymbol{X}} \langle oldsymbol{x}, oldsymbol{arphi} 
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Kernel trick (Boser, Guyon, Vapnik; 1992):

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Soft margin (Cortes, Vapnik; 1995):

- ▶ Allow for a portion of points from **X** to be beyond the margin, label points far from the origin by "1", those close by "-1".
- ▶ Controlled by a parameter  $\nu \in (0,1)$  (Schölkopf, Platt, Shawe-Taylor, Smola, Williamson; 1999).

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This can be formulated as a quadratic programming problem

$$\begin{split} \min_{\boldsymbol{\psi} \in \mathcal{H}, \boldsymbol{\xi} \in \mathbb{R}^n, \rho \in \mathbb{R}} \quad & \frac{1}{2} \|\boldsymbol{\psi}\|^2 + \frac{1}{\nu n} \sum_{i=1}^n \xi_i - \rho \\ \text{subject to} \quad & \langle \boldsymbol{\xi}, \Phi(\boldsymbol{x}_i) \rangle \geq \rho - \xi_i \,, \,\, \xi_i \geq 0 \,\, \text{for} \,\, i = 1, ..., n \,, \end{split}$$
 with  $\boldsymbol{\xi} = (\xi_1, ..., \xi_n)^\top$ .

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One can reformulate the optimization problem to employ the kernel trick.



In dual formulation, using the Lagrangian, one can restate the optimization problem as follows:

$$\begin{split} \min_{\boldsymbol{\alpha}} & \quad \frac{1}{2} \sum_{i=1}^n \sum_{j=1}^n \alpha_i \alpha_j K(\boldsymbol{x}_i, \boldsymbol{x}_j) \\ \text{subject to} & \quad 0 \leq \alpha_i \leq \frac{1}{\nu n} \text{ for } i = 1, ..., n, \sum_{i=1}^n \alpha_i = 1, \\ \text{with } \boldsymbol{\alpha} = (\alpha_1, ..., \alpha_n)^\top. \end{split}$$

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The decision function is then:

$$g_{OCSVM}(\mathbf{x}) = \operatorname{sgn}\left(\sum_{i=1}^{n} \alpha_{i} K(\mathbf{x}_{i}, \mathbf{x}) - \rho\right),$$

where  $\rho$  can be recovered from any  $\mathbf{x}_i$  such that  $0 < \alpha_i < \frac{1}{n}$ :

$$\rho = \langle \boldsymbol{\psi}, \Phi(\boldsymbol{x}_i) \rangle = \sum_{i=1}^{n} \alpha_i K(\boldsymbol{x}_i, \boldsymbol{x}_j).$$



Idea 2: Put points into a small ball.

$$\min_{\substack{R \in \mathbb{R}, \boldsymbol{\xi} \in \mathbb{R}^n, \boldsymbol{c} \in \mathcal{H}, \\ \text{subject to}}} R^2 + \frac{1}{\nu n} \sum_{i=1}^n \xi_i$$

$$\sup_{i=1}^n \{ \boldsymbol{\xi}_i \} \| \boldsymbol{\Phi}(\boldsymbol{x}_i) - \boldsymbol{c} \| \leq R^2 + \xi_i, \ \xi_i \geq 0 \text{ for } i = 1, ..., n.$$

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This leads to the dual:

$$\begin{split} & \min_{\alpha} & & \sum_{i=1}^{n} \sum_{j=1}^{n} \alpha_{i} \alpha_{j} K(\mathbf{x}_{i}, \mathbf{x}_{j}) - \sum_{i=1}^{n} \alpha_{i} K(\mathbf{x}_{i}, \mathbf{x}_{i}) \\ & \text{subject to} & & 0 \leq \alpha_{i} \leq \frac{1}{\nu n}, \text{ for } i = 1, ..., n, \; \sum_{i=1}^{n} \alpha_{i} = 1 \,. \end{split}$$

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$$\|\Phi(\boldsymbol{x}_i) - \boldsymbol{c}\| < R^2 + \xi_i, \ \xi_i > 0 \text{ for } i = 1, ..., n.$$

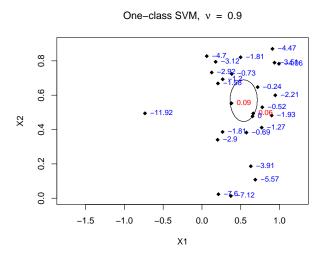
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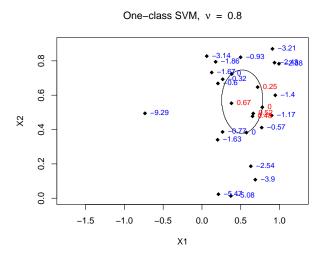
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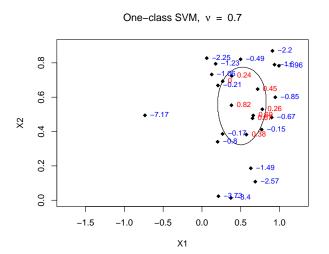
which leads to the decision function:

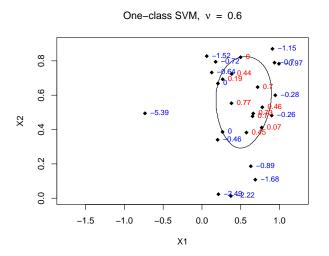
$$g_{OCSVM}(\mathbf{x}) = \left(R^2 - \sum_{i=1}^n \sum_{j=1}^n \alpha_i \alpha_j K(\mathbf{x}_i, \mathbf{x}_j) + 2 \sum_{i=1}^n \alpha_i K(\mathbf{x}_i, \mathbf{x}) - K(\mathbf{x}, \mathbf{x})\right),$$

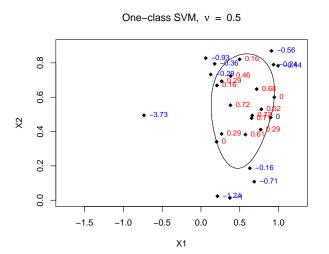
with  $R^2 = \sum_{i,j} \alpha_i \alpha_j K(\mathbf{x}_i, \mathbf{x}_j) - 2 \sum_i \alpha_i K(\mathbf{x}_i, \mathbf{x}_k) + K(\mathbf{x}_k, \mathbf{x}_k)$  for any  $\mathbf{x}_k$  such that  $0 < \alpha_k < 1/(\nu n)$ .

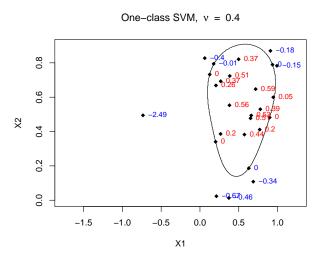












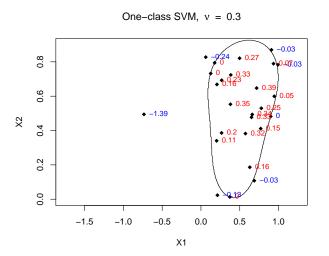


Illustration: Case 1

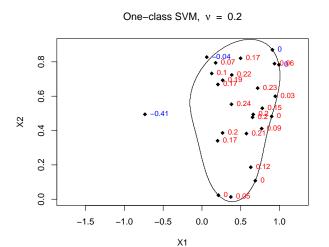
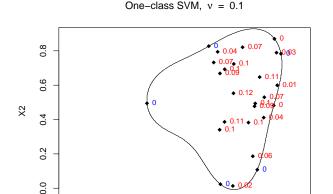


Illustration: Case 1



-0.5

X1

0.0

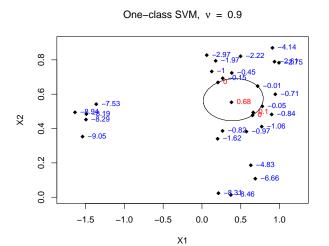
0.5

1.0

-1.5

-1.0

Illustration: Case 2



One-class SVM, v = 0.8

Illustration: Case 2

9.0

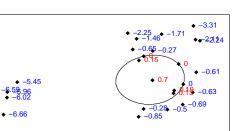
0.0

-1.5

-1.0

-0.5

X1



0.0

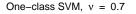
→ -3.42→ -4.88

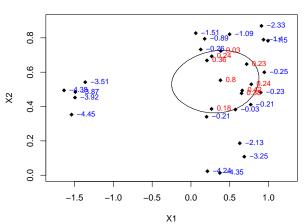
1.0

♦ -6.176.31

0.5

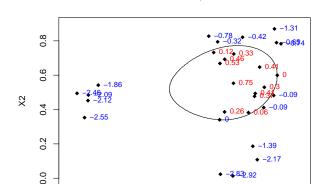
Illustration: Case 2





One-class SVM, v = 0.6

Illustration: Case 2



-0.5

X1

0.0

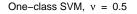
0.5

1.0

-1.5

-1.0

Illustration: Case 2



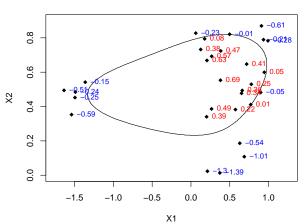
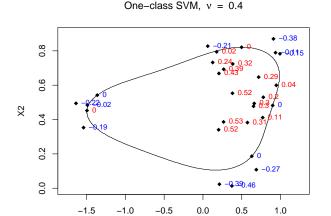


Illustration: Case 2



X1

Illustration: Case 2

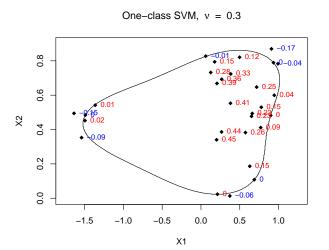


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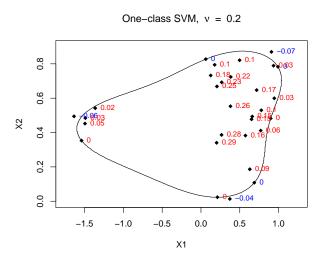
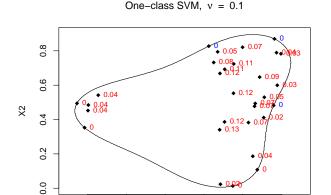


Illustration: Case 2



-0.5

X1

0.0

0.5

1.0

-1.5

-1.0

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#### k-distance of a point x:

For any integer k > 0, the k-distance of point  $\boldsymbol{x}$ , denoted as k- $dist(\boldsymbol{x})$ , is defined as the distance  $d(\boldsymbol{x}, \boldsymbol{o})$  between  $\boldsymbol{x}$  and a point  $\boldsymbol{o} \in \boldsymbol{X}$  such that:

- ▶ for at least k points  $o' \in X \setminus \{x\}$  it holds that  $d(x, o') \le d(x, o)$ , and
- for at most k-1 points  $\boldsymbol{o}' \in \boldsymbol{X} \setminus \{\boldsymbol{x}\}$  it holds that  $d(\boldsymbol{x}, \boldsymbol{o}') < d(\boldsymbol{x}, \boldsymbol{o})$ .

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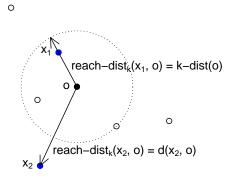
Given the k-dist(x), the k-neighborhood of x, denoted  $N_k(x)$ , contains every point whose distance from x is not greater than the k-dist(x), i.e.:

$$N_k(\mathbf{x}) = \{\mathbf{q} \in \mathbf{X} \setminus \{\mathbf{x}\} \mid d(\mathbf{x}, \mathbf{q}) \leq k \text{-}dist(\mathbf{x})\}.$$



**Reachability distance** of order k of point x w.r.t. point o: For  $k \in \mathbb{N}$ , the reachability distance of order k of point x with respect to point o is defined as:

$$reach-dist_k(\mathbf{x}, \mathbf{o}) = \max\{k-dist(\mathbf{o}), d(\mathbf{x}, \mathbf{o})\}.$$



Local reachability density of a point x:

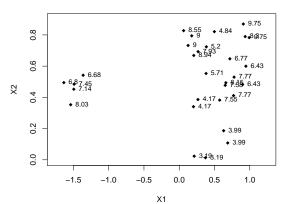
The local reachability density of x is defined as:

$$Ird_k(\mathbf{x}) = \frac{|N_k(\mathbf{x})|}{\sum_{\mathbf{o} \in N_k(\mathbf{x})} reach-dist_k(\mathbf{x}, \mathbf{o})}.$$

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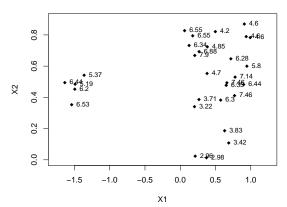
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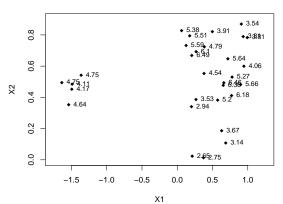
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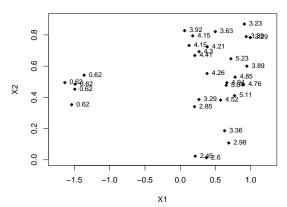
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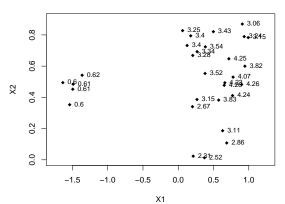
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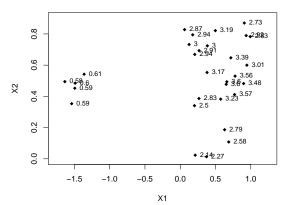
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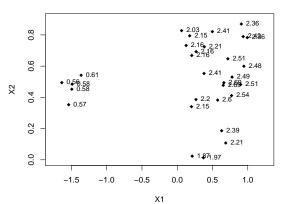
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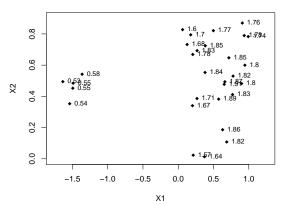
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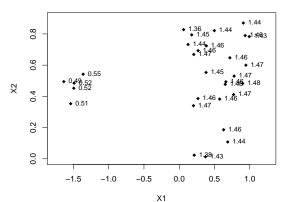
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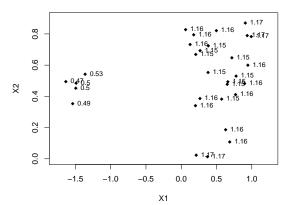
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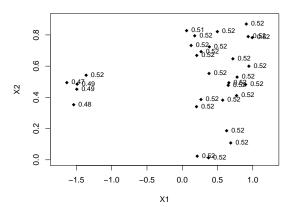
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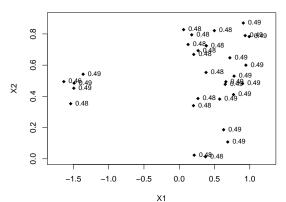
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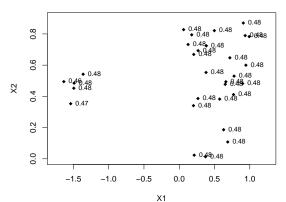
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**Local outlier factor** of a point x:

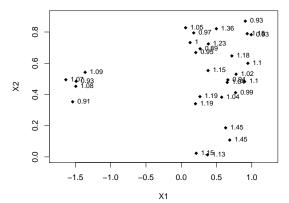
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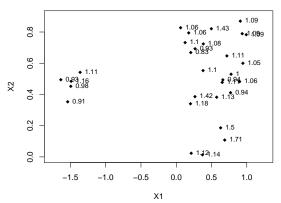
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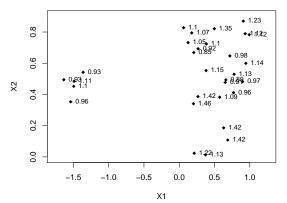
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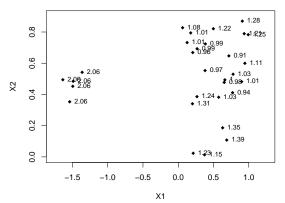
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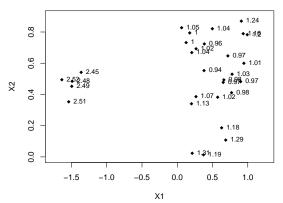
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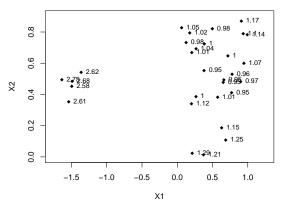
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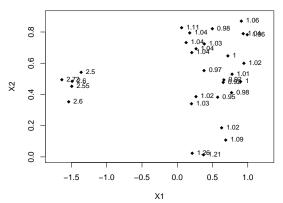
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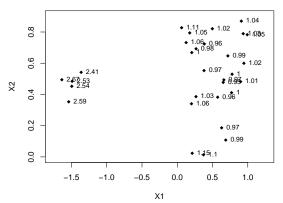
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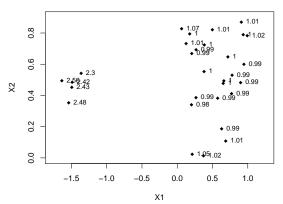
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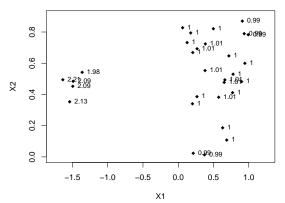
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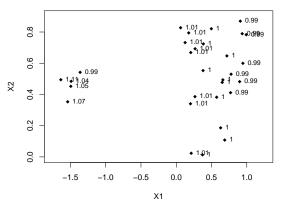
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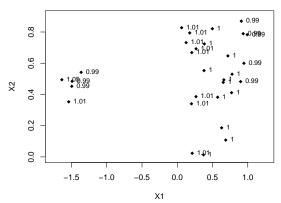
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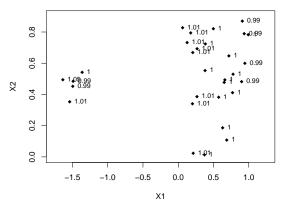
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#### Practical session

- ▶ Isolation forest (Liu, Ting, Zhou; 2008) is an anomaly detection method inherited from the famous random forest algorithm (Breiman, 2001).
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- ► Since no supervised feedback is given, isolation forest is based on purely random (uniform) variable-based partitioning.
- ▶ Main idea: Outlying observations are isolated faster.
- Tree-kind partitioning is done until "full isolation": outlying observations will have smaller depth (on an average) in the isolation tree.
- ► A monotone transform is usually applied to the aggregated estimate.
- ► To reduce both masking effect and computation cost, small-size sub-sampling is used instead of bootstrap.

► Each isolation tree is grown **recursively** using the described below node-construction procedure

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1. Choose a split variable I uniformly from  $\{1, ..., d\}$ .

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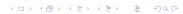
$$\left[\min_{\boldsymbol{x}\in\mathcal{S}_{j,k}}\langle\boldsymbol{x},\boldsymbol{e}_{l}\rangle,\max_{\boldsymbol{x}\in\mathcal{S}_{j,k}}\langle\boldsymbol{x},\boldsymbol{e}_{j}\rangle\right].$$

3. Form the children subsets

$$C_{j+1,2k} = C_{j,k} \cap \{ \boldsymbol{x} \in \mathbb{R}^d : \langle \boldsymbol{x}, \boldsymbol{e}_l \rangle \leq \kappa \},$$
  
$$C_{j+1,2k+1} = C_{j,k} \cap \{ \boldsymbol{x} \in \mathbb{R}^d : \langle \boldsymbol{x}, \boldsymbol{e}_l \rangle > \kappa \}.$$

as well as the children training datasets

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Stop when only one observation is in each node: isolation.

Illustration: Isolation tree

#### **Isolation forest**

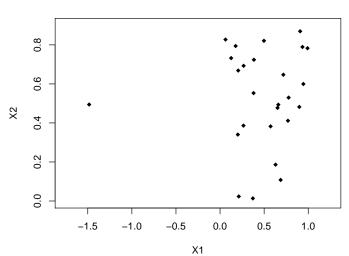


Illustration: Isolation tree

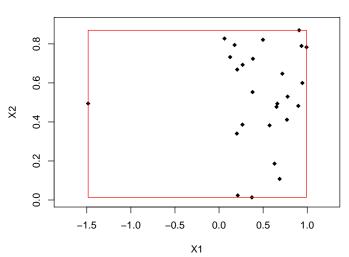


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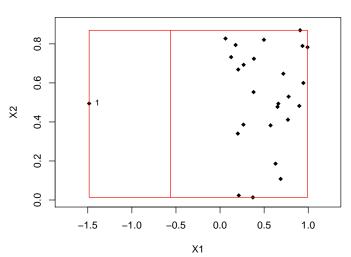


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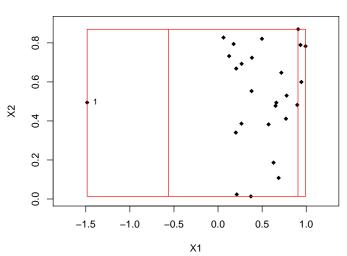
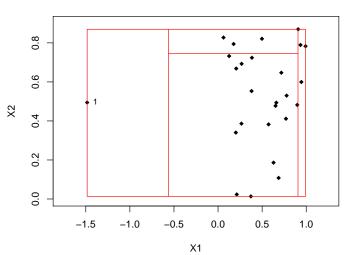


Illustration: Isolation tree



Isolation tree, split 4

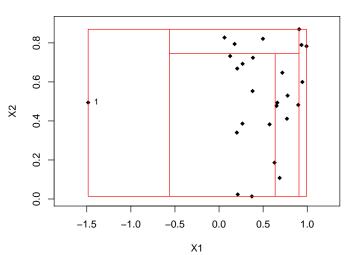


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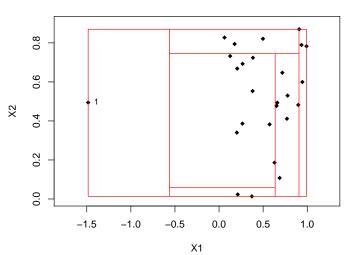


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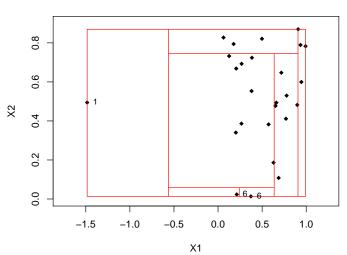


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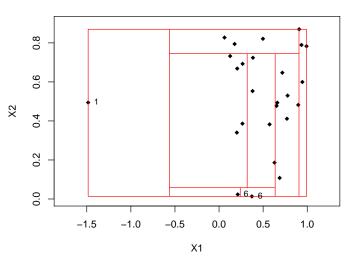


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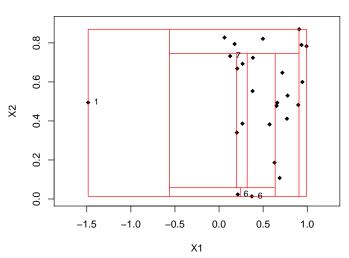
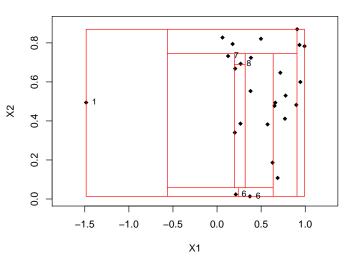
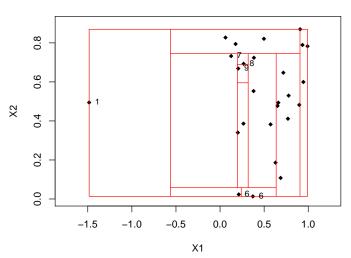


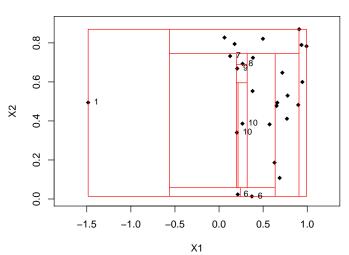
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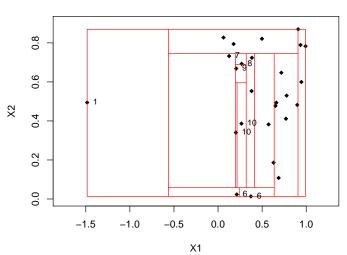
Isolation tree, split 10



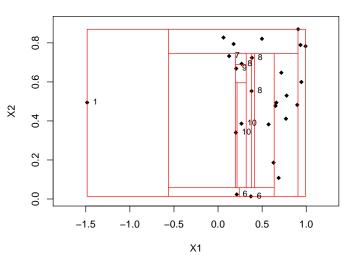
Isolation tree, split 11



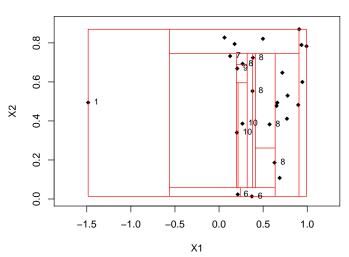
Isolation tree, split 12



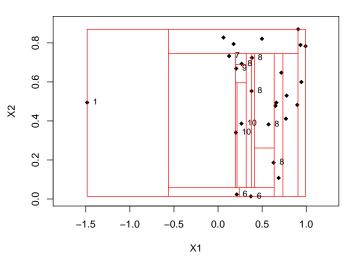
Isolation tree, split 13



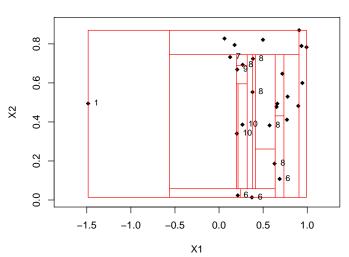
Isolation tree, split 14



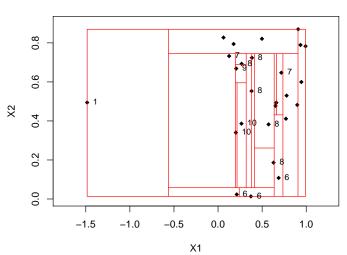
Isolation tree, split 15



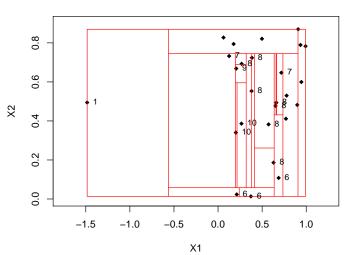
Isolation tree, split 16



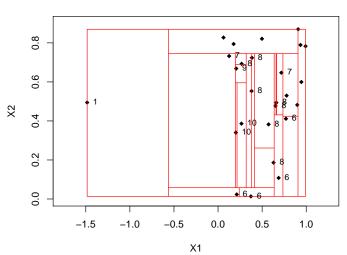
Isolation tree, split 17



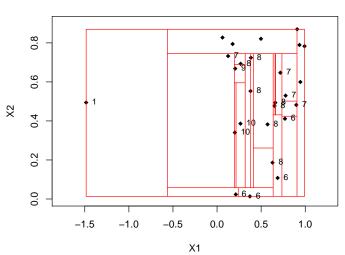
Isolation tree, split 18



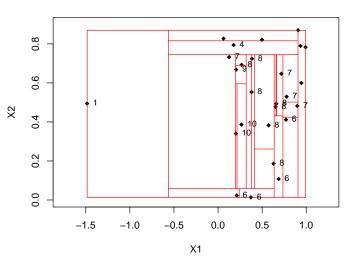
Isolation tree, split 19



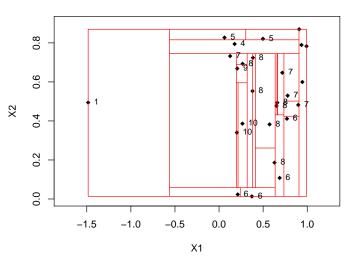
Isolation tree, split 20



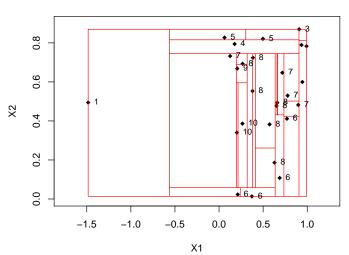
Isolation tree, split 21



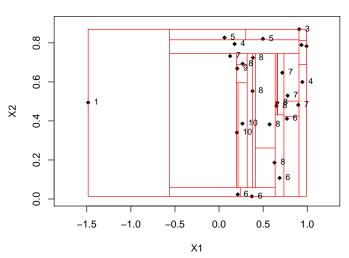
Isolation tree, split 22



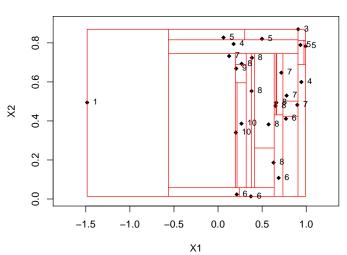
Isolation tree, split 23



Isolation tree, split 24



Isolation tree, split 25



#### Anomaly score calculation for observation x:

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

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

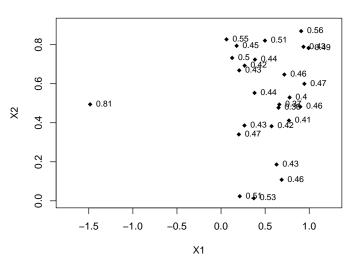
with  $c(n) = 2H(n-1) - \frac{2(n-1)}{n}$  where H(k) is the harmonic number and can be estimated by  $\ln(k) + 0.5772156649$ .

#### Score behavior:

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

Illustration: Anomaly score

#### Isolation forest score, 100 trees



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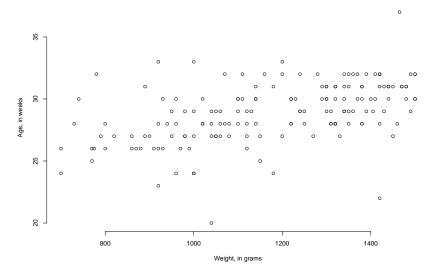
# Systematic orderings: data depth The notion of data depth

The Tukey depth function Central regions

#### Practical session

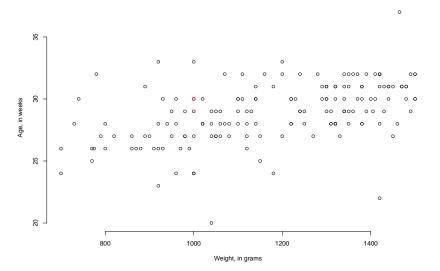
# Data depth

#### Babies with low birth weight



# Data depth

#### Babies with low birth weight



A **data depth** measures how close a given point is located to the center of a distribution. For  $x \in \mathbb{R}^p$  and a p-variate random vector X distributed as  $P \in \mathcal{P}$ , a data depth is a function

$$D: \mathbb{R}^p \times \mathcal{P} \to [0,1], (\boldsymbol{x},P) \mapsto D(\boldsymbol{x}|P)$$

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that is:

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Depth notions: Mahalanobis ('36), projection (Stahel, '81; Donoho, '82), simplicial volume (Oja, '83), simplicial (Liu, '90), zonoid (Koshevoy, Mosler, '97), spatial (Vardi, Zhang, '00; Serfling, '02), lens (Liu, Modarres, '11), ... depth.



### Applications of data depth:

- Multivariate data analysis (Liu, Parelius, Singh '99);
- Statistical quality control (Liu, Singh '93);
- ► Cluster analysis and classification (Mosler, Hoberg '06; Li, Cuesta-Albertos, Liu '12; M., Mosler, Lange '15);
- ► Tests for multivariate location, scale, symmetry (Liu '92; Dyckerhoff '02; Dyckerhoff, Ley, Paindaveine '15);
- Outlier detection (Hubert, Rousseeuw, Segaert '15);
- Multivariate risk measurement (Cascos, Mochalov '07);
- Robust linear programming (Bazovkin, Mosler '15);
- Missing data imputation (M., Josse, Husson '20);
- etc.

R-package **ddalpha** (Pokotylo, M., Dyckerhoff, Nagy): calculates a number of depths; performs depth-based classification of multivariate and functional data; contains 50 multivariate and 5 functional data sets.

Python library data-depth: to be released soon.

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Tukey (1975) — "Mathematics and the picturing of data"

Tukey depth of  $\mathbf{x} \in \mathbb{R}^p$  w.r.t. a d-variate random vector X distributed as P is defined as the smallest probability mass of a closed halfspace containing  $\mathbf{x}$ :

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$$D^{T(n)}(m{x}|m{X}) = \frac{1}{n} \min_{m{u} \in \mathbb{S}^{p-1}} \sharp\{i: m{u}'m{x}_i \geq m{u}'m{x}\}.$$

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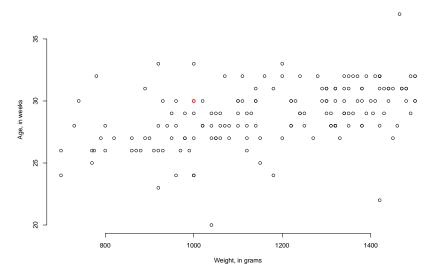
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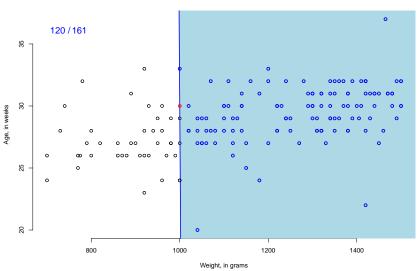
- satisfies all the above postulates,
- ▶ is purely non-parametric and robust,
- has direct connection to quantiles and many applications.



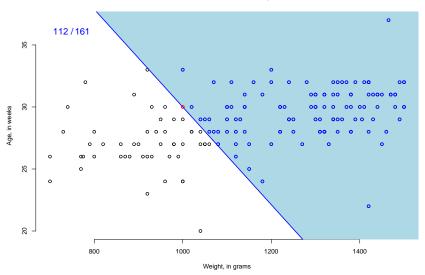
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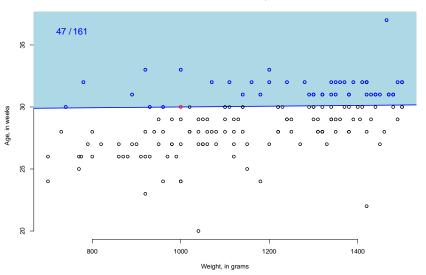


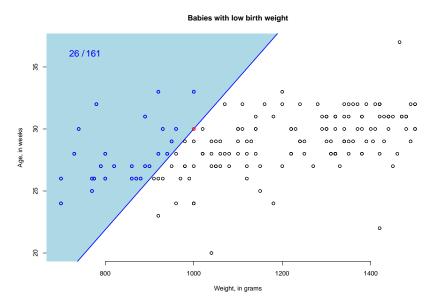




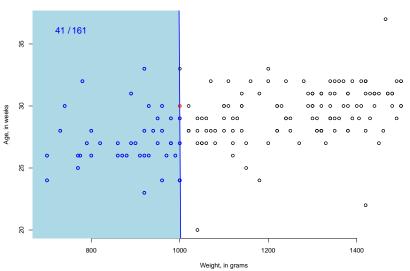


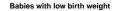
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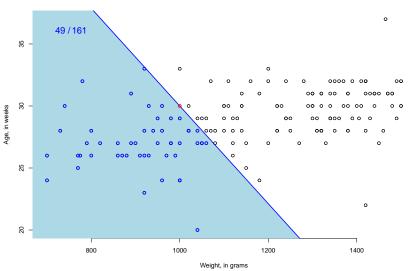


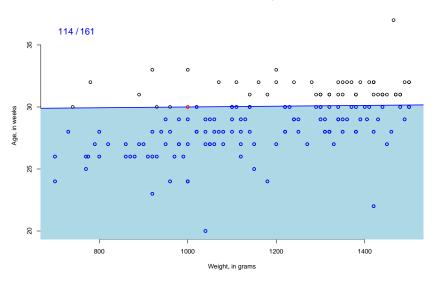


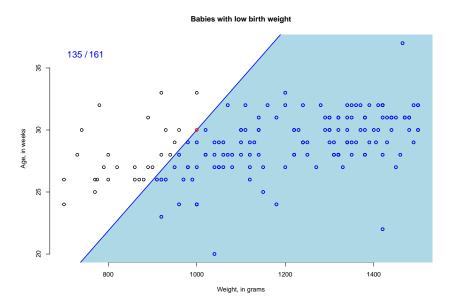


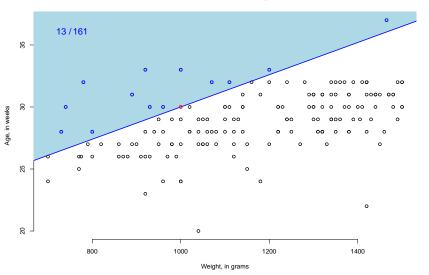


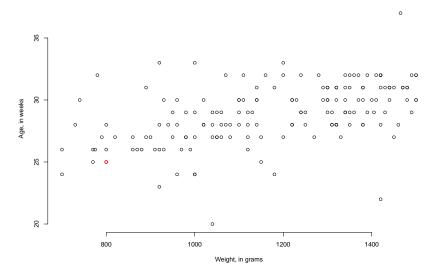


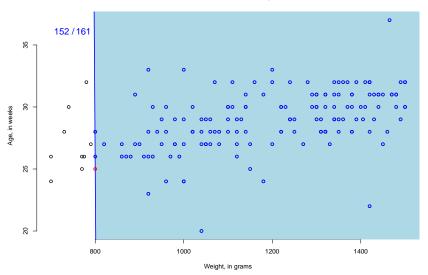


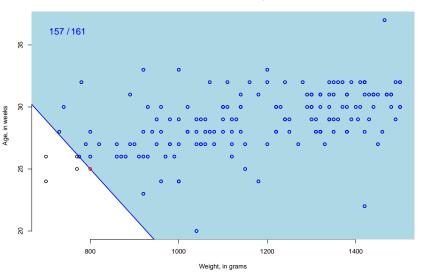


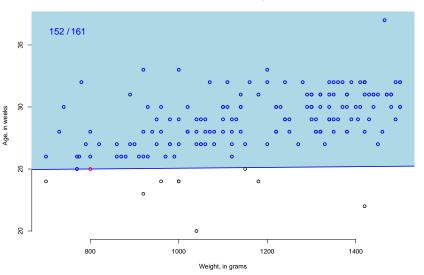


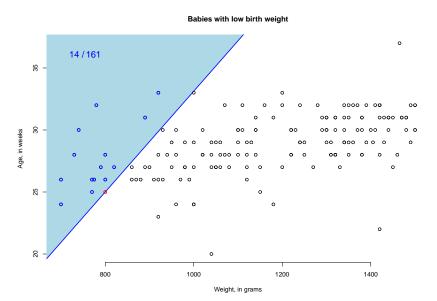


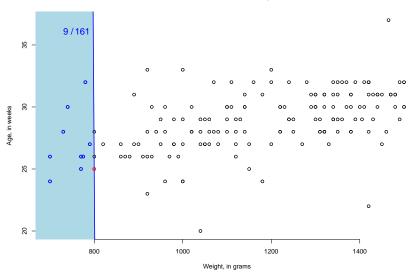


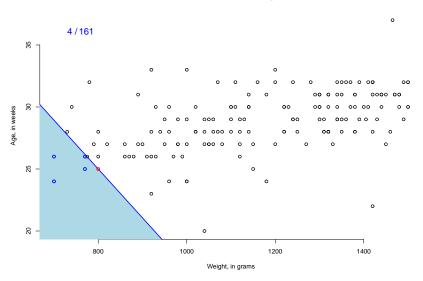


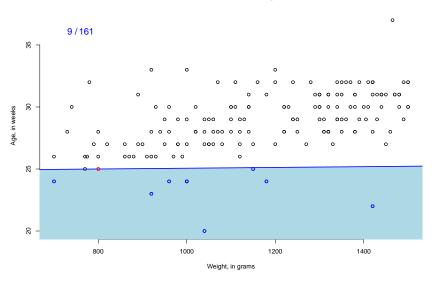


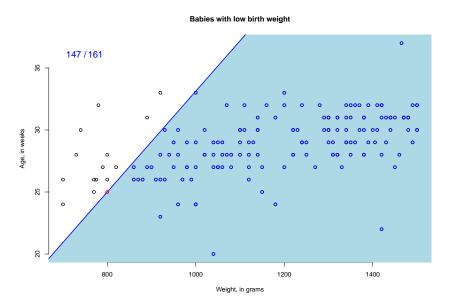


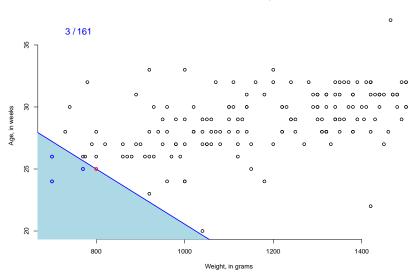


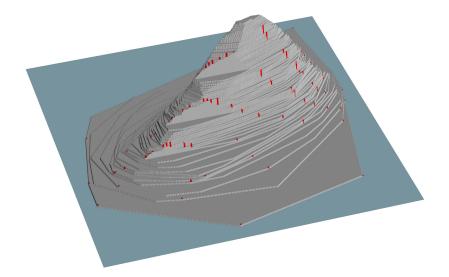












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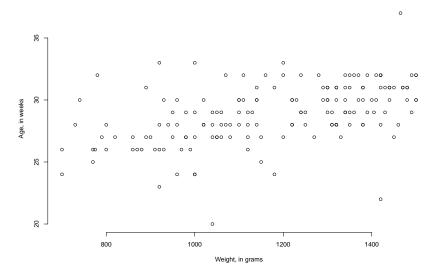
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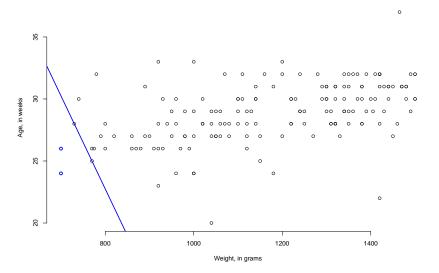
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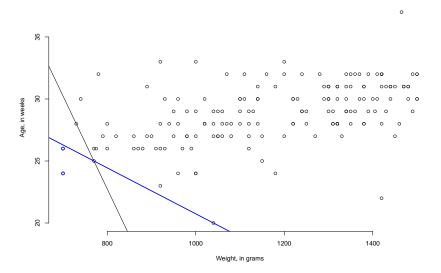
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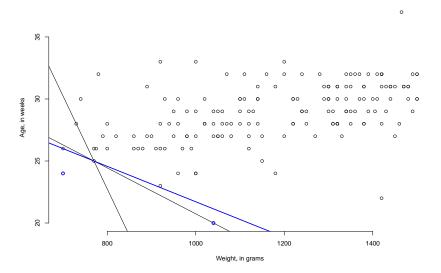
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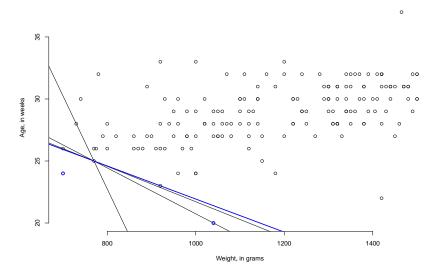
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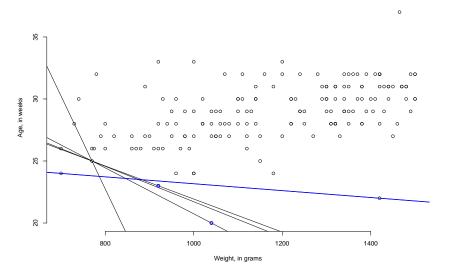


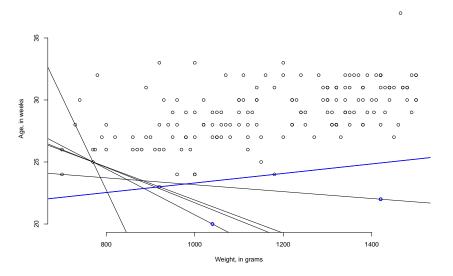


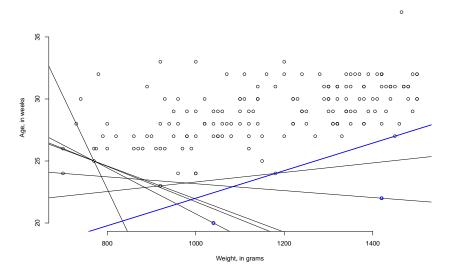


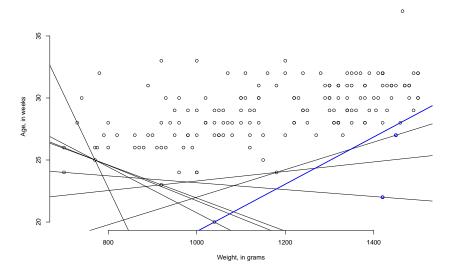


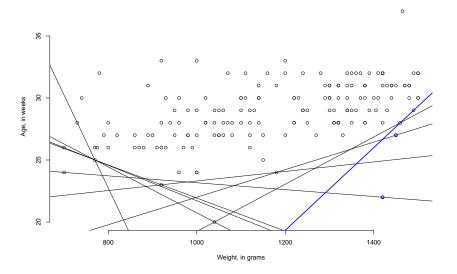


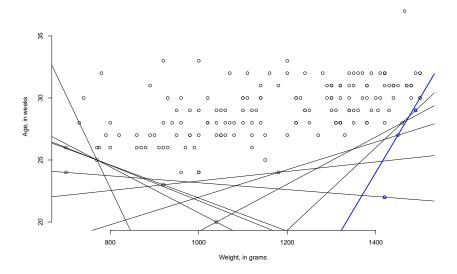


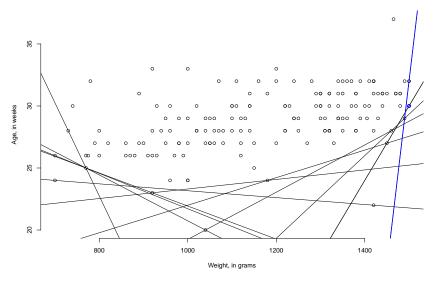


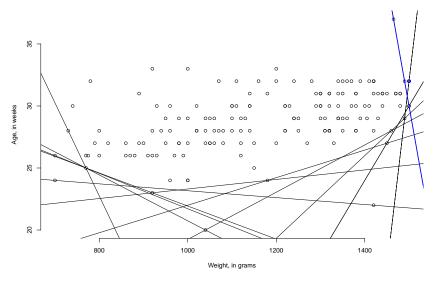




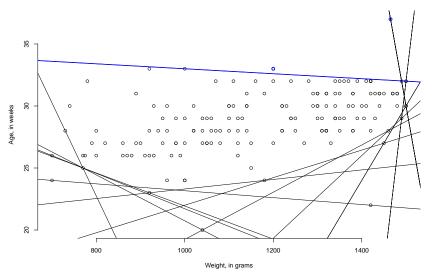




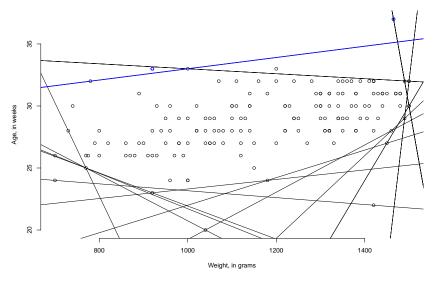




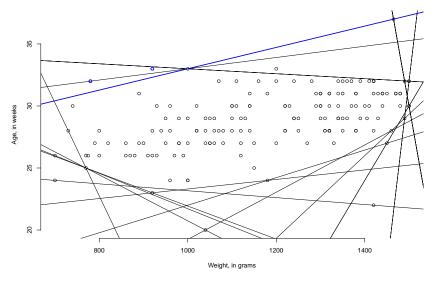


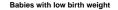


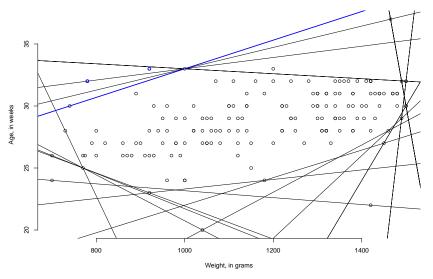


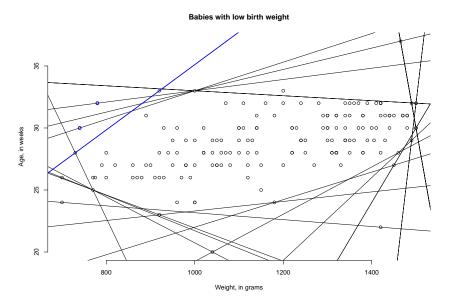


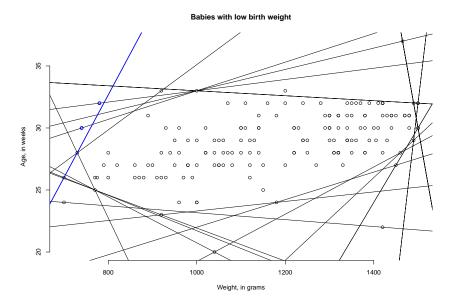


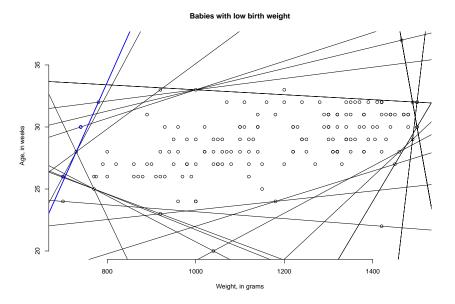


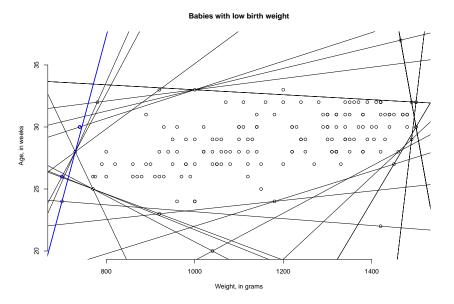


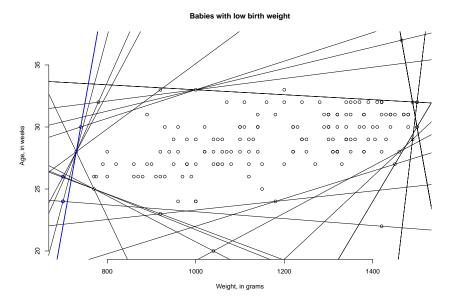


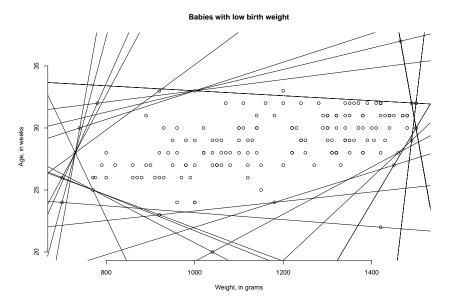


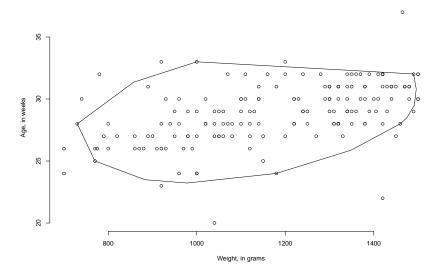




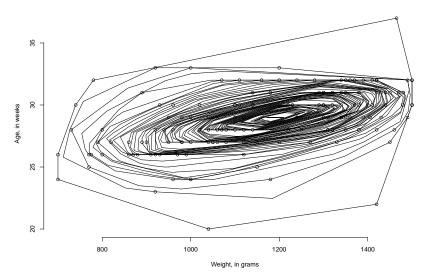




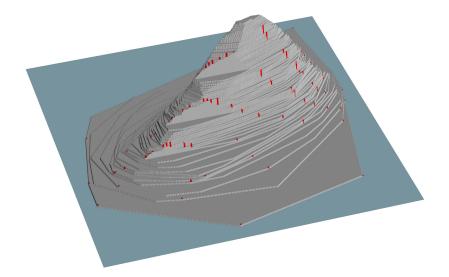






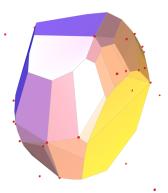


# Tukey (=halfspace, location) data depth





#### Tukey (=halfspace, location) depth region: $\tau = 2/161$



### Tukey (=halfspace, location) depth region: $\tau = 5/161$



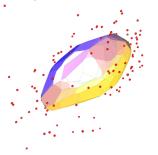
### Tukey (=halfspace, location) depth region: $\tau = 9/161$



#### Tukey (=halfspace, location) depth region: $\tau = 13/161$



#### Tukey (=halfspace, location) depth region: $\tau = 17/161$



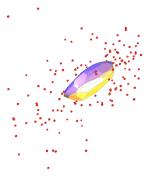
#### Tukey (=halfspace, location) depth region: $\tau = 25/161$



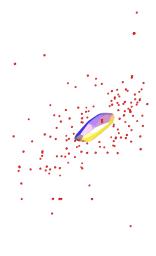
#### Tukey (=halfspace, location) depth region: $\tau = 33/161$



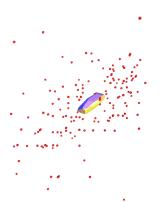
#### Tukey (=halfspace, location) depth region: $\tau = 41/161$



#### Tukey (=halfspace, location) depth region: $\tau = 49/161$



#### Tukey (=halfspace, location) depth region: $\tau = 57/161$



#### Tukey (=halfspace, location) depth region: $\tau = 65/161$



#### Tukey (=halfspace, location) depth region: $\tau = 68/161$



#### Contents

#### Introduction

#### Non-parametric approaches

One-class support vector machines Local outlier factor Isolation forest

#### Systematic orderings: data depth

The notion of data depth
The Tukey depth function
Central regions

Further depth notions

#### Practical session

### Mahalanobis depth (Mahalanobis, 1936)

▶ Mahalanobis depth is defined as:

$$D^{Mah}(\mathbf{x}|X) = \frac{1}{1 + (\delta^{Mah})^2(\mathbf{x}|X)},$$

based on Mahalanobis distance:

$$(\delta^{\mathit{Mah}})^2(\mathbf{x}|X) = (\mathbf{x} - \boldsymbol{\mu}_X)^T \mathbf{\Sigma}_X^{-1} (\mathbf{x} - \boldsymbol{\mu}_X)$$
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  - moment estimates;
  - robust estimates such as minimum volume ellipsoid or minimum covariance determinant (MCD).

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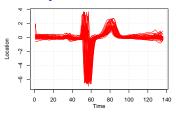
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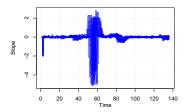
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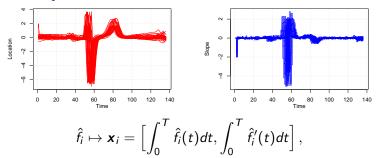
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  - by a single elliptical contour characterizes a multivariate normal distribution or one within an affine family of non-degenerate elliptical distributions.

### ECG five days data



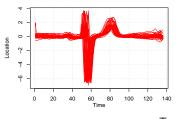


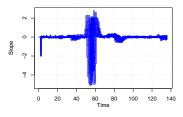
#### ECG five days data



with  $\hat{f}_i(t)$  being the function obtained by connecting the points  $(t_{ij}, f_i(t_{ij})), j = 1, \ldots, N_i$  with line segments,  $\hat{f}'_i(t)$  its derivative.

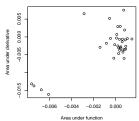
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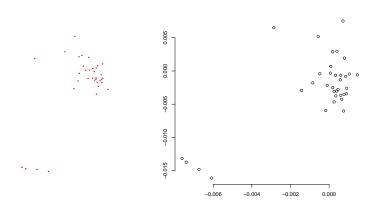


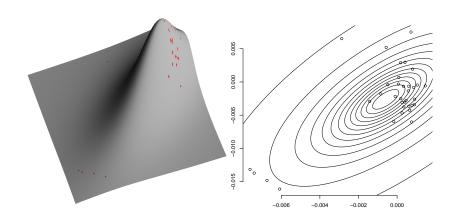


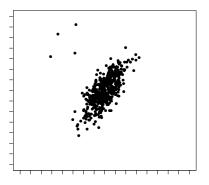
$$\hat{f}_i \mapsto \boldsymbol{x}_i = \left[\int_0^T \hat{f}_i(t)dt, \int_0^T \hat{f}_i'(t)dt\right],$$

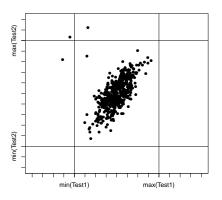
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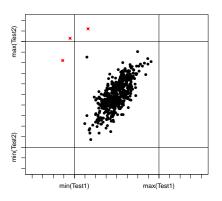






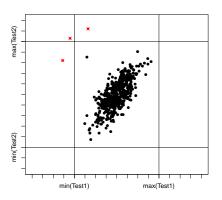


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- ► Label observation **x** as anomaly if:

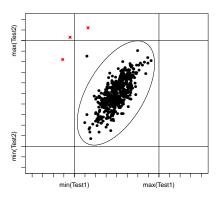
$$\textit{\textbf{x}} \notin [\mathsf{min}(\mathsf{Test1}), \mathsf{max}(\mathsf{Test1})] \times [\mathsf{min}(\mathsf{Test2}), \mathsf{max}(\mathsf{Test2})] \,.$$



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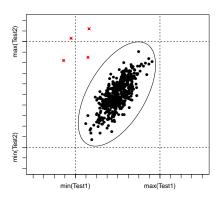
Not all anomalies can be detected.



▶ Mahalanobis distance of an observation  $\mathbf{x} \in \mathbb{R}^2$  (from the mean) is defined as follows:

$$d_{\mathsf{Mah}}(\pmb{x}|\pmb{X}) = (\pmb{x}-\pmb{\mu})^{ op} \pmb{\Sigma}^{-1}(\pmb{x}-\pmb{\mu})\,,$$

where  $\mu$  is the **mean** and  $\Sigma$  is the **covariance** matrix.



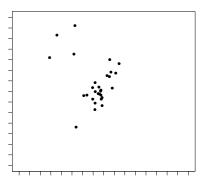
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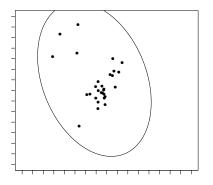
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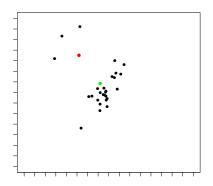
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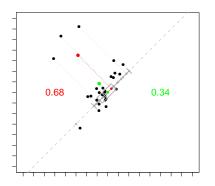


► Mahalanobis distance (moment estimators) **not robust**.



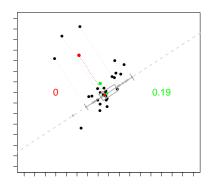
- Mahalanobis distance (moment estimators) not robust.
- ▶ Stahel-Donoho outlyingness of x w.r.t.  $X = \{x_i\}_{i=1}^n$ :

$$O_{SD}(\boldsymbol{x}|\boldsymbol{X}) = \max_{\boldsymbol{u} \in \mathcal{S}^{d-1}} \frac{|\boldsymbol{x}^{\top}\boldsymbol{u} - \mathsf{med}(\boldsymbol{X}\boldsymbol{u})|}{\mathsf{MAD}(\boldsymbol{X}\boldsymbol{u})}.$$



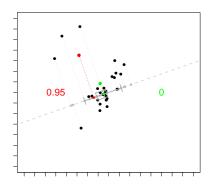
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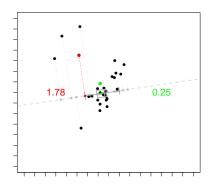
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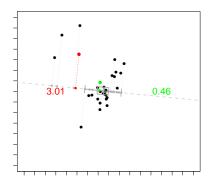
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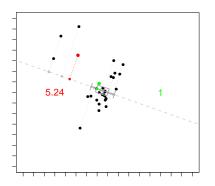
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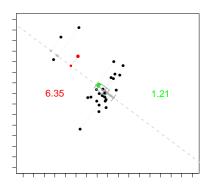
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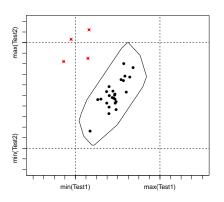
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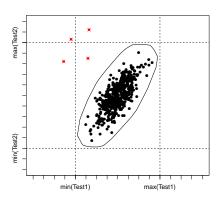


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is the **projected outlyingness** (Stahel, 1981; Donoho, 1982),  $\operatorname{med}(Y)$  and  $\operatorname{MAD}(Y) = \operatorname{med}(\big|Y - \operatorname{med}(Y)\big|)$  are the univariate median and median absolute deviation from the median, respectively.

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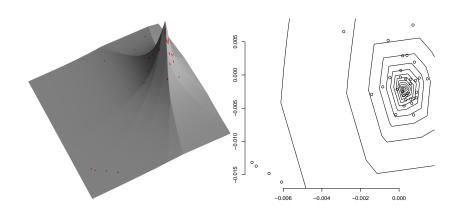
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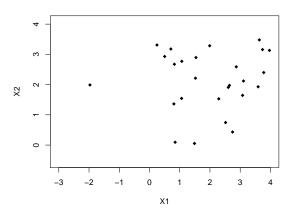
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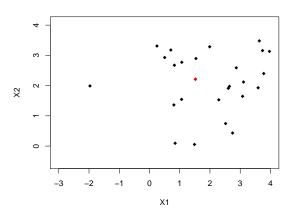


$$D^{spt}(\boldsymbol{x}|X) = 1 - \left\| \mathbb{E} \big[ \frac{\boldsymbol{x} - \boldsymbol{X}}{\|\boldsymbol{x} - \boldsymbol{X}\|} \big] \right\| \quad \text{with} \quad \frac{\boldsymbol{x} - \boldsymbol{X}}{\|\boldsymbol{x} - \boldsymbol{X}\|} = 0 \quad \text{if} \quad \boldsymbol{x} - \boldsymbol{X} = \boldsymbol{0} \,.$$

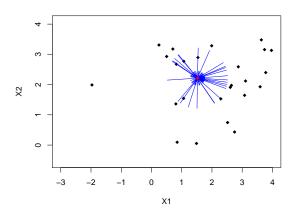
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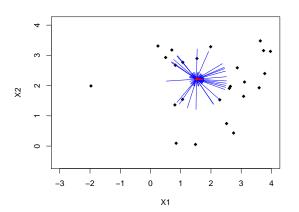
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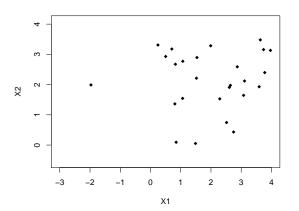
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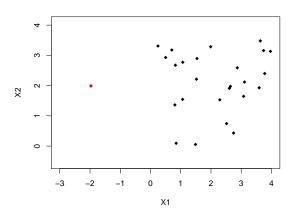
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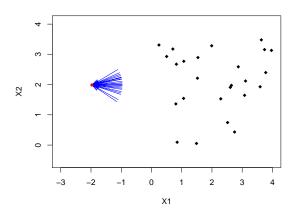
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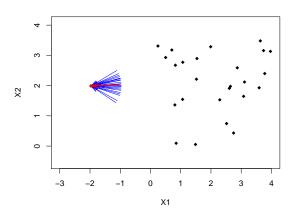
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Exploiting the idea of spatial quantiles of Chaudhuri (1996) and Koltchinskii (1997), Vardi & Zhang (2000) and Serflig (2002) formulate the **spatial depth** (also  $L_1$ -depth) as:

$$D^{spt}(\mathbf{x}|X) = 1 - \left\| \mathbb{E}\left[v(\mathbf{\Sigma}^{-\frac{1}{2}}(\mathbf{x} - X))\right] \right\|,$$

with

$$u(\mathbf{y}) = \begin{cases} \frac{\mathbf{y}}{\|\mathbf{y}\|} & \text{if } \mathbf{y} \neq \mathbf{0}, \\ \mathbf{0} & \text{if } \mathbf{y} = \mathbf{0}. \end{cases}$$

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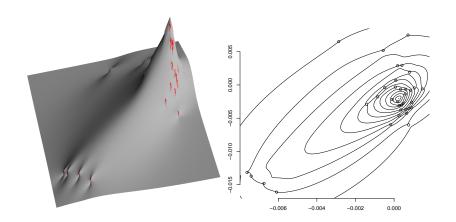
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- ▶ satisfies D1 D5, but not D4con, is continuous;
- if Σ is orthogonal, satisfies D2iso only;
- with D2iso its maximum (say x\*) is referred to as spatial median, a multivariate location estimator having asymptotic breakdown point of 0.5.



### Contents

#### Introduction

### Non-parametric approaches

One-class support vector machines Local outlier factor Isolation forest

### Systematic orderings: data depth

The notion of data depth
The Tukey depth function
Central regions
Further depth notions

#### 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.
- Breunig, M.M., Kriegel, H.-P., Ng, R.T., and Sander, J. (2000). LOF: Identifying density-based local outliers. In: Proceedings of the 2000 ACM SIGMOD International Conference on Management of Data, 29, 93–104.
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- ▶ Mosler, K. (2013). Depth statistics. In: Robustness and Complex Data Structures: Festschrift in Honour of Ursula Gather, 17-34.

# Practical session (part I)

#### Notebooks:

- anomdet\_simulation1.Rmd,
- ▶ anomdet\_hurricanes.Rmd,
- ▶ anomdet\_brainimaging.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
- ▶ 101\_1\_dwi\_fa.nii: Anatomical brain volume data.
- ► 101\_1\_dwi.voxelcoordsL.txt: Left brain fiber's bundle.
- ▶ 101\_1\_dwi.voxelcoordsR.txt: Right brain fiber's bundle.

### 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.

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## Literature (mentioned in the tutorial) (2)

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