

Outlier detection

EDOUARD PAUWELS

M2-MAT SID

- Data cleaning
- Attack / intrusion detection (IT security)
- Fraud detection (banking, insurance).
- Medical diagnosis and monitoring of unusual symptoms
- Industrial monitoring, damage detection, predictive maintenance
- Image processing, video surveillance
- Text mining (news detection)
- Sensor networks, fault / attack
- etc. . .

What is an outlier?

Often used interchangeably with *anomaly*

Hawkins (1980) :

An observation that deviates so much from other observations as to arouse suspicion that it was generated by a different mechanism.

Johnson (1992) :

An observation in a data set which appears to be inconsistent with the remainder of that set of data.

Barnett and Lewis (1994) :

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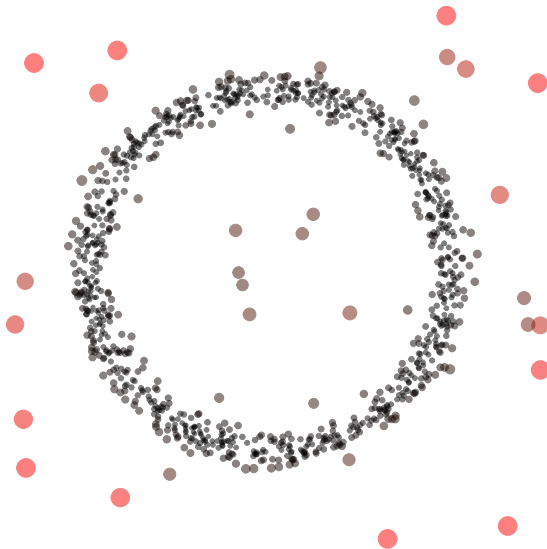
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Main ideas :

- Need a reference distribution, sample.
- Outliers and non outliers are mixed in the same sample.
- The proportion of outliers is small

Illustration



Input space : $\mathcal{X} = \mathbb{R}^p$, $n \in \mathbb{N}$, $S_n = (x_i)_{i=1}^n$, $x_i \in \mathcal{X}$, $i = 1, \dots, n$.

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Binary classification with labels $(y_i)_{i=1}^n$ describing the status (anomaly or not).
Unbalanced classes.

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- ▶ S_n consists only of points which are not anomalies.
- ▶ PU learning : S_n has some point labeled as normal and the rest could be either normal or abnormal.

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We will still have labels : evaluate methods performances, only used for test purposes.

1. What is an outlier ?
2. Score based detection
3. Univariate methods
4. Multivariate approaches
5. Practical

Setting :

Training data : $S_n = (x_i)_{1 \leq i \leq n}$,

Goal : predict $y \in \{0, 1\}$ (anomaly or not).

Scoring : Compute a scoring function $s_n: \mathcal{X} \mapsto \mathbb{R}$. $s_n: x \mapsto h(x, x_1, \dots, x_n)$.

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Out of sample intrusion / change detection : compare score and class on unseen data $s_n(\tilde{x}), \tilde{y}$.

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Anomaly if $s_n(x_i) \geq \bar{s}$.

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Remark : All the methods which we will see can be used for out of sample anomaly detection. The evaluation is then close to what is done in supervised learning settings.

Evaluation metrics : precision and recall

		Reality		Total
		Abnormal	Normal	
Prediction	Abnormal	TP	FP	$TP + FP$
	Normal	FN	TN	$FN + TN$
Total		$TP + FN$	$FP + TN$	n

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$$\text{Precision} \quad \frac{TP}{TP+FP} = \frac{TP}{|\text{predicted anomalies}|} \cdot$$

$$\text{Recall} \quad \frac{TP}{TP+FN} = \frac{TP}{|\text{real anomalies}|} \cdot$$

$$\text{F1 score} \quad F1 = 2 \times \frac{Pr \times Rec}{Pr + Rec}$$

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$$\text{Precision}(\bar{s}) : \frac{TP(\bar{s})}{TP(\bar{s}) + FP(\bar{s})} = \frac{TP(\bar{s})}{|\text{predicted anomalies}(\bar{s})|}.$$

$$\text{Recall}(\bar{s}) : \frac{TP(\bar{s})}{TP(\bar{s}) + FN(\bar{s})} = \frac{TP(\bar{s})}{|\text{real anomalies}(\bar{s})|} \quad (= \text{TPR}(\bar{s})).$$

$$\text{FPR}(\bar{s}) : \frac{FP(\bar{s})}{FP(\bar{s}) + TN(\bar{s})} = \frac{FP(\bar{s})}{|\text{real normal}(\bar{s})|}.$$

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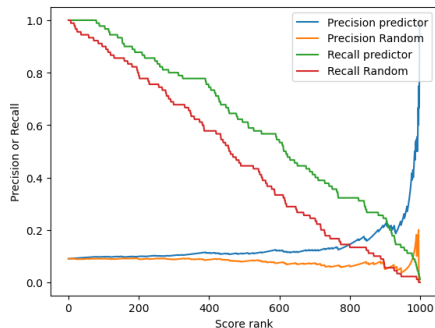
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- Ex : semi-supervised approach.
 - ▶ Choose \bar{s} which has the largest F1-score.
 - ▶ Evaluate using cross validation.

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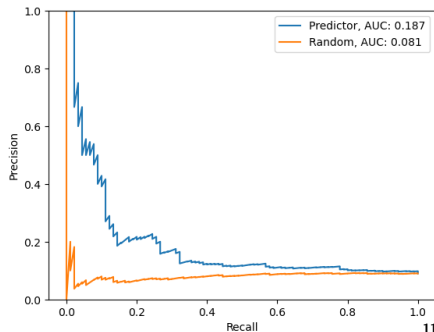
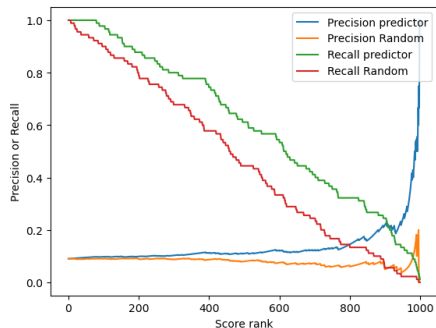
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Evaluation metrics : PR curves

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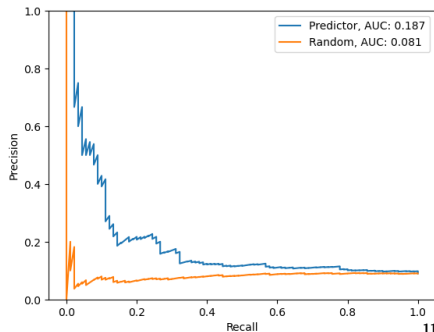
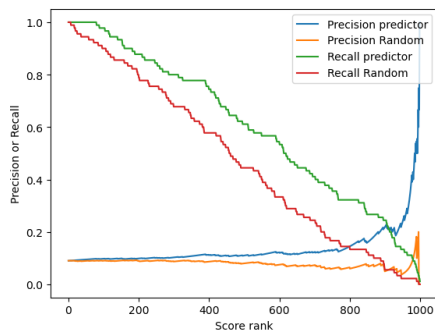
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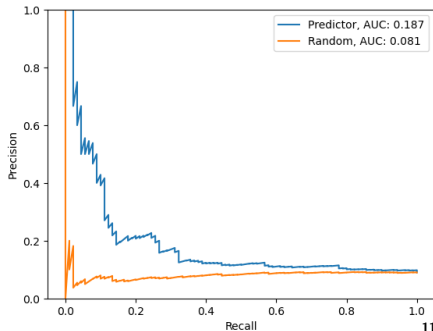
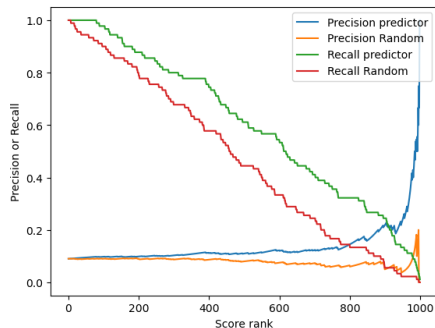
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Comments :

- Compare methods ability to order by degree of outlyingness.
- Allows to compare methods without having to select \bar{s}
- More general but less taylored to certain regimes.
- **In any case** : \bar{s} will be needed in practice.



Hyperparameters : number of neighbors, polynomial degree ...

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Tools from supervised learning : cross validation, validation set.

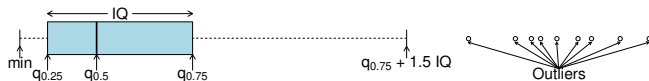
TP_PR_ROC

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Interquartile range :



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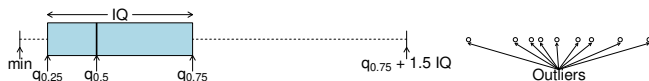


Z-score :

$$s_n : t \mapsto \frac{|t - \bar{x}|}{\sigma_x}$$

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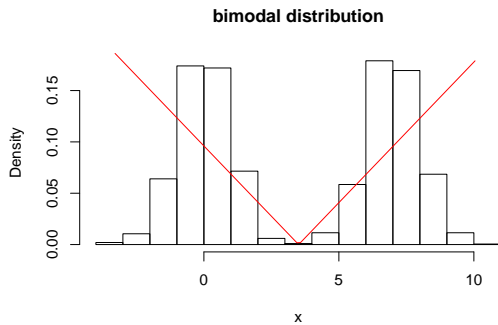


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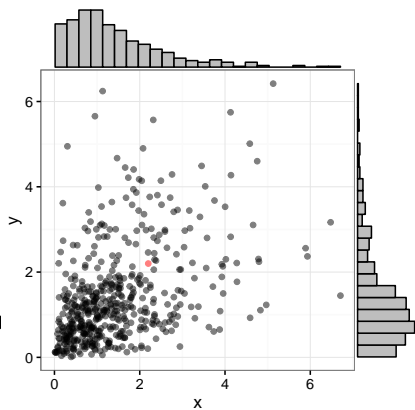
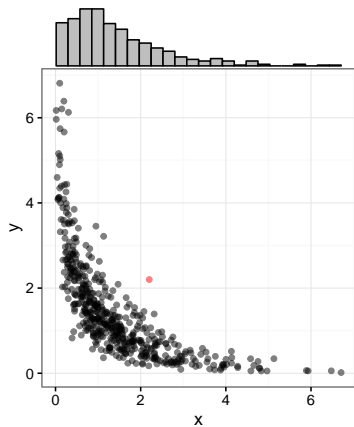
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Shortcomings and limitations?

A bimodal distribution, Z-score in red.



Bi-variate \neq $2 \times$ univariate



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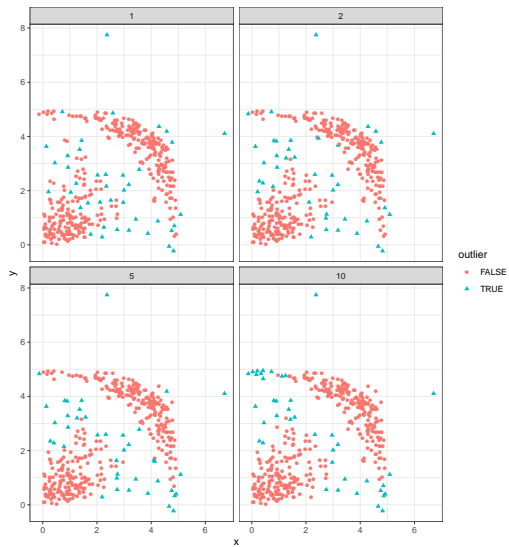
Distance to the k -th neighbor :

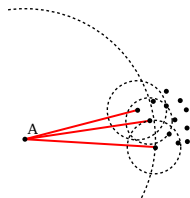
$$s_n : x \mapsto k\text{dist}(x) := \text{dist}(x, x_l)$$

where x_l is the k -th nearest neighbor of x in $S_n = (x_i)_{i=1}^n$.

Distance to the k-th neighbor

Variation of k





$$N_k(x)$$

$$\text{REACH}_k(x, y) = \max \{ k\text{dist}(y), \text{dist}(x, y) \}$$

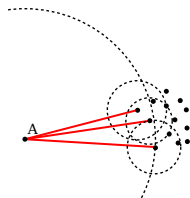
k nearest neighbors of x
reachability

$$\text{LRD}_k(x) = \left(\frac{1}{k} \sum_{y \in N_k(x)} \text{REACH}_k(x, y) \right)^{-1}$$

Local Reachability Density

$$\text{LOF}_k(x) = \frac{1}{k} \sum_{y \in N_k(x)} \frac{\text{LRD}_k(y)}{\text{LRD}_k(x)}$$

Local Outlier Factor



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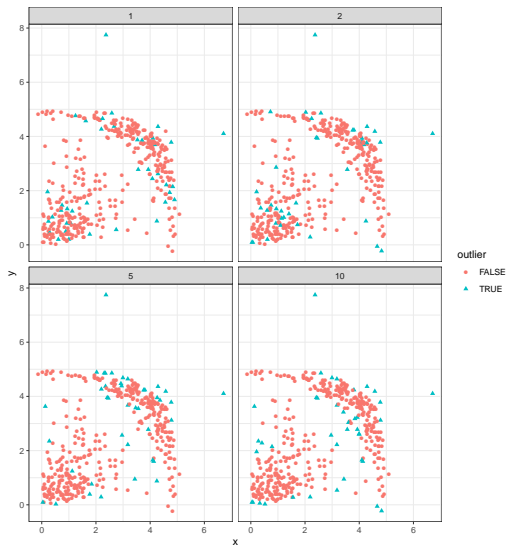
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Local Outlier Factor

$\text{LOF} > 1$ implies smaller density as neighbors. The LOF is used as a score s_n .

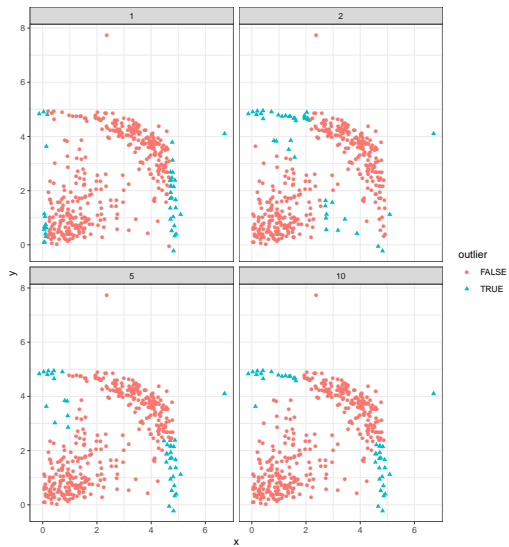
Variation of k



k clusters

- Perform clustering using
- $s_n: x \mapsto \text{dist}(x, c)$, where c is the centroid the closest to x .

Variation of the number of clusters



Family of parametrized models density functions p_θ :

- Maximum likelihood : $\hat{\theta} \in \arg \max_{\theta} \sum_{i=1}^n \log(p_\theta(x_i))$.
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Mahalanobis distance and Gaussian model :

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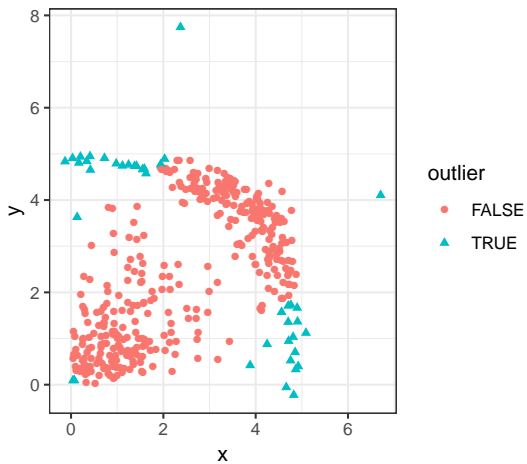
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Mahalanobis distance and Gaussian model :

$$s_n : x \mapsto \exp \left(- (x - \bar{x}_n) \Sigma_n^{-1} (x - \bar{x}_n) \right)$$

where \bar{x}_n is the empirical mean and Σ_n is the empirical covariance matrix.

No tuning parameter



Density of the form

$$p_{\theta} : x \mapsto \sum_{i=1}^K \tau_i p(x|\mu_i, \Sigma_i)$$

where $\tau_i > 0$, $\sum_i \tau_i = 1$,

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Maximum likelihood : using EM algorithm (\sim extension of k-means).

Gaussian mixture model

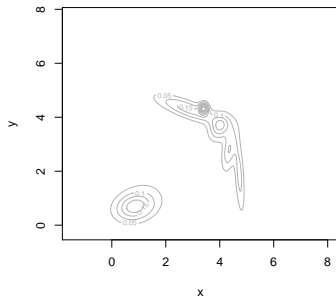
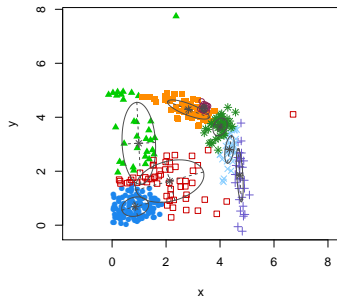
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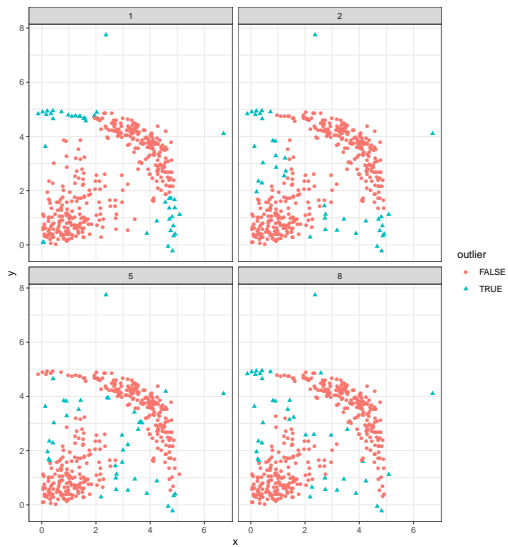
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Variation of the number of clusters



Gaussian kernel with bandwidth σ

$$k(x, y) = \frac{1}{\sqrt{2\pi}\sigma} e^{-\frac{\|y-x\|^2}{2\sigma^2}}$$

Kernel density estimator :

$$p_\sigma : x \mapsto \frac{1}{n} \sum_{i=1}^n k(x, x_i)$$

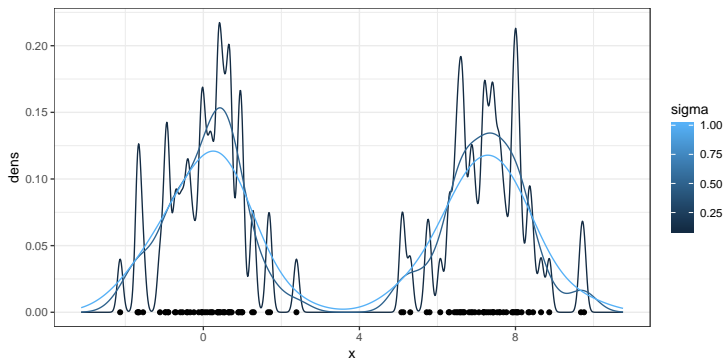
Density based

Gaussian kernel with bandwidth σ

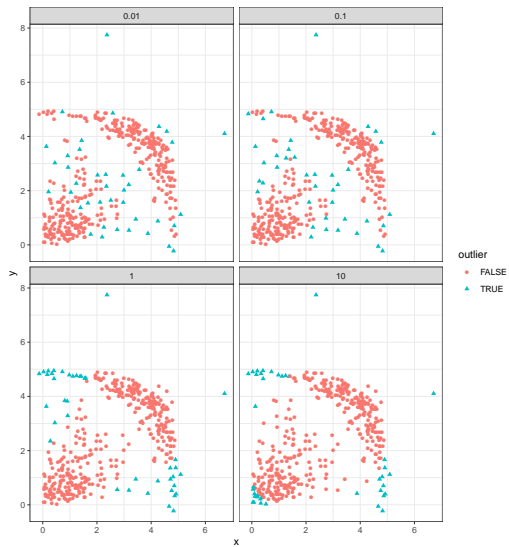
$$k(x, y) = \frac{1}{\sqrt{2\pi}\sigma} e^{-\frac{\|y-x\|^2}{\sigma^2}}$$

Kernel density estimator :

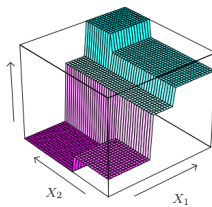
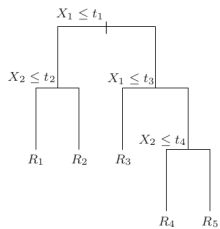
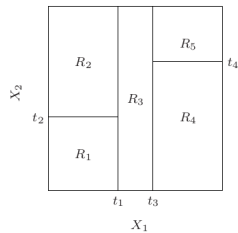
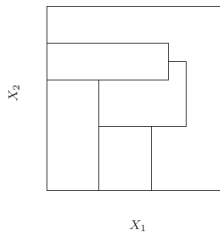
$$\rho_\sigma : x \mapsto \frac{1}{n} \sum_{i=1}^n k(x, x_i)$$



Variation of the bandwidth



Isolation forest



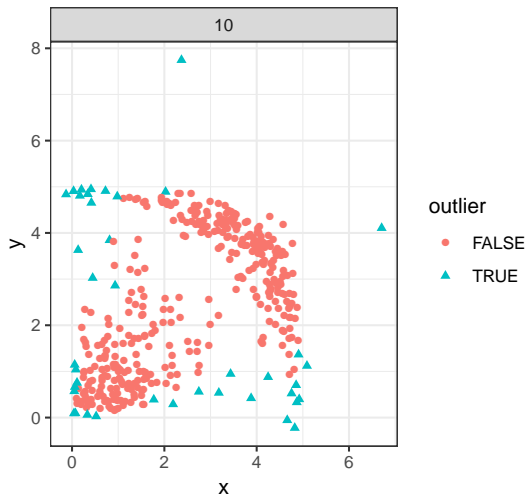
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- A tree is grown randomly by induction.
- Given a rectangle which contains more than 1 point, we split it in two by choosing one variable and one threshold randomly.
- Stop when points are alone in their rectangle.

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A tree provides a notion of depth which can be used as a score to measure abnormality. An isolation forest consists of several such trees, s_n is the average depth across trees.

Isolation forest

No parameter (number of trees in the forest)



Main idea, find a ball of minimal radius which encloses all the points :

$$\begin{aligned} \min_{r \in \mathbb{R}, c \in \mathbb{R}^p} \quad & r^2 \\ \text{s.t.} \quad & \|x_i - c\|^2 \leq r^2, \quad i = 1, \dots, n. \end{aligned}$$

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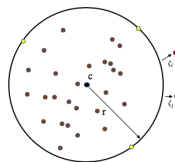
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s_n is roughly the distance to the center.



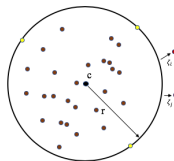
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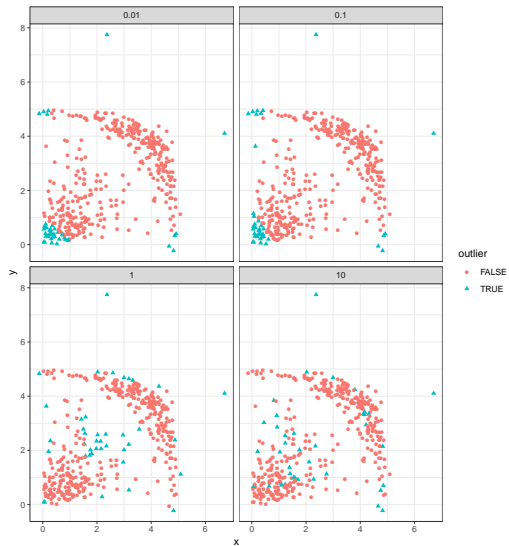


Kernel trick : $\phi: x \mapsto X \in \mathbb{R}^P$ sends x to a high (infinite) dimensional feature space.

Implicitly : $x_i \rightarrow \phi(x_i), \quad i = 1, \dots, n.$

Positive definite kernel (ex : Gaussian) implicitly encodes ϕ .

Gaussian kernel with varying bandwidth



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Exercise : For each method that we have seen describe

- How many parameters ?
- Is the computed score random ?
 - ▶ if you run the algorithm twice, do you get the same result ?
- Do anomaly correspond to large or small values of the score ?

1. What is an outlier ?
2. Score based detection
3. Univariate methods
4. Multivariate approaches
5. Practical