#### Gaussian Mixture Model - method and application

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# Gaussian Mixture Models – method and applications



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- Method
  - Introduction to Gaussian Mixture Process (GMM)
  - Standard construction of GMM
  - Clustering (Silhouette and Akaike criterion)
- Case studies
  - Monitoring a secondary settler tank
  - Residual and fault detection criteria
- Conclusions



# Gaussian Mixture Model (GMM) - standard construction

A linear superposition of *K*-Gaussians

 $\mu_k$ : mean

 $\sigma_k$ : covariance

$$p(\mathbf{x}_i) = \sum_{k=1}^{K} \underbrace{\pi_k}_{p(k)} \underbrace{\mathcal{N}(\mathbf{x}_i | \mu_k, \sigma_k)}_{p(\mathbf{x}_i | k)}, i = 1, ..., N$$

is called a **Gaussian mixture (GM)**. The mixture coefficient  $\pi_k$  satisfies

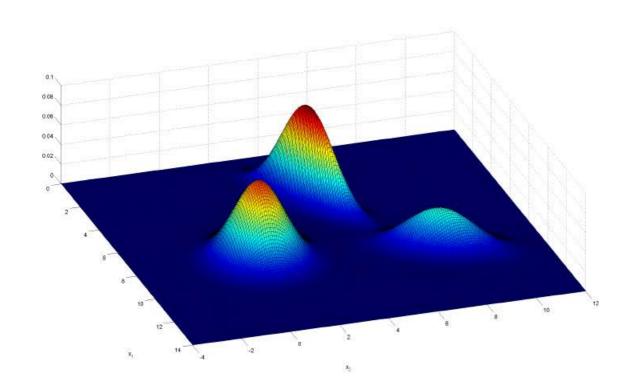
$$\sum_{k=1}^{K} \pi_k = 1, \qquad 0 \le \pi_k \le 1$$

**Interpretation**: The density  $p(\mathbf{x}|k) = \mathcal{N}(\mathbf{x}|\mu_k, \sigma_k)$  is the probability of  $\mathbf{x}$ , given that component k was chosen. The probability of choosing component k is given by the prior probability p(k).



#### For example, consider the following GMM:

$$p(x) = \underbrace{0.3}_{\pi_1} \mathcal{N}\left(x \mid \underbrace{\begin{pmatrix} 4 \\ 4.5 \end{pmatrix}}_{\mu_1}, \underbrace{\begin{pmatrix} 1.2 & 0.6 \\ 0.6 & 0.5 \end{pmatrix}}_{\Sigma_1}\right) + \underbrace{0.5}_{\pi_2} \mathcal{N}\left(x \mid \underbrace{\begin{pmatrix} 8 \\ 1 \end{pmatrix}}_{\mu_2}, \underbrace{\begin{pmatrix} 1 & 0 \\ 0 & 1 \end{pmatrix}}_{\Sigma_2}\right) + \underbrace{0.2}_{\pi_3} \mathcal{N}\left(x \mid \underbrace{\begin{pmatrix} 9 \\ 8 \end{pmatrix}}_{\mu_3}, \underbrace{\begin{pmatrix} 0.6 & 0.5 \\ 0.5 & 1.5 \end{pmatrix}}_{\Sigma_3}\right)$$



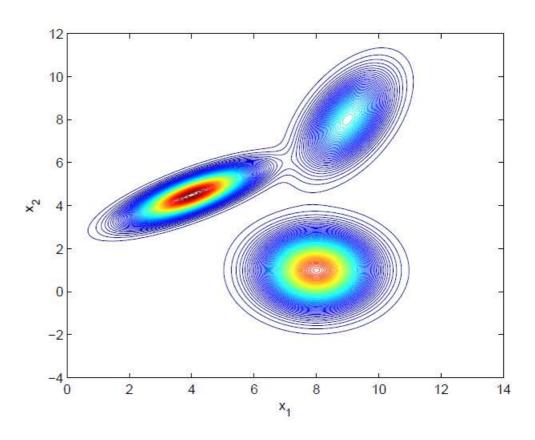


Figure: Probability density function.

Figure: Contour plot.



The form of the GM distribution is governed by the parameters  $\pi$ ,  $\mu$  and  $\sigma$ . One way to get them is by **maximum likelihood**.

Given N observations  $\{x_n\}_{n=1}^N$ , the log-likelihood function is

$$\ln p(X; \pi_{1:K}, \mu_{1:K}, \sigma_{1:K}) = \sum_{n=1}^{N} \ln \left( \sum_{k=1}^{K} \pi_k \mathcal{N}(\mathbf{x}_n | \mu_k, \sigma_k) \right)$$

There is **no closed-form solution** available (due to the sum inside the logarithm).

This problem can be separated into two simple problems using the *expectation-maximization (EM)* algorithm.



#### Conditions to be satisfied at a maximum of the likelihood function

$$\frac{\mathrm{d}}{\mathrm{d}\mu_k} \left[ \ln p(\mathbf{x}|\pi, \mu, \sigma) \right] = 0 \quad \to \quad 0 = -\sum_{n=1}^N \underbrace{\frac{\pi_k \mathcal{N}(\mathbf{x}_n | \mu_k, \sigma_k)}{\sum_j \pi_j \mathcal{N}(\mathbf{x}_n | \mu_j, \sigma_j)}}_{\gamma(z_{nk})} \sigma_k(\mathbf{x}_n - \mu_k)$$

which gives 
$$\rightarrow \mu_k = \frac{1}{N_k} \sum_{n=1}^N \gamma(z_{nk}) \mathbf{x}_n$$

$$\frac{\mathrm{d}}{\mathrm{d}\sigma_k} \left[ \ln p(\mathbf{x}|\pi, \mu, \sigma) \right] = 0 \quad \to \quad \sigma_k = \frac{1}{N_k} \sum_{n=1}^N \gamma(z_{nk}) \left( \mathbf{x}_n - \mu_k \right) \left( \mathbf{x}_n - \mu_k \right)^T$$

Maximize  $\ln p(\mathbf{x}|\pi,\mu,\sigma)$  with respect to  $\pi_k$  (using Lagrange multipliers) gives

$$\pi_k = \frac{N_k}{N}$$
, where  $N_k = \sum_{n=1}^{N} \gamma(z_{nk})$ 

#### Algorithm 1 EM for Gaussian mixtures

- 1: Initialize  $\mu_k^1, \sigma_k^1, \pi_k^1$  and set i = 1.
- 2: while not converged do
- 3: Compute  $\gamma(z_{nk})$ .  $\triangleright$  Expectation step 4: Compute  $\mu_k^{i+1}; \pi_k^{i+1}; N_k; \sigma_k^{i+1}$ .  $\triangleright$  Maximization step
- 5:  $i \leftarrow i + 1$ .
- 6: end while

$$\gamma(z_{nk}) = \frac{\pi_k^i \mathcal{N}(\mathbf{x}_n | \mu_k^i, \sigma_k^i)}{\sum_{j=1}^K \pi_j^i \mathcal{N}(\mathbf{x}_n | \mu_j^i, \sigma_j^i)}, n = 1, ..., N; k = 1, ..., K$$

$$\mu_k^{i+1} = \frac{1}{N_k} \sum_{n=1}^{N} \gamma(z_{nk}) \mathbf{x}_n,$$

$$\pi_k^{i+1} = \frac{N_k}{N}, \quad N_k = \sum_{n=1}^N \gamma(z_{nk}),$$

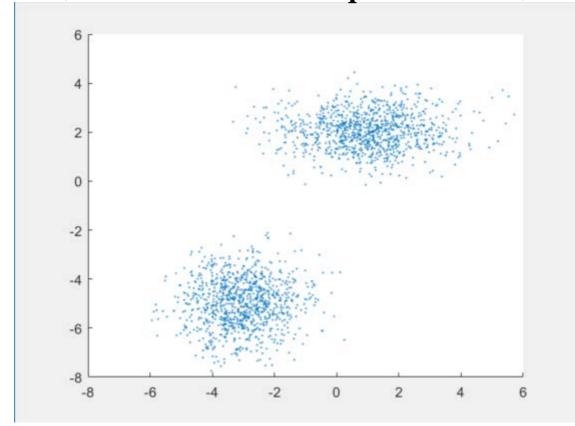
$$\sigma_k^{i+1} = \frac{1}{N_k} \sum_{n=1}^{N} \gamma(z_{nk}) \left( \mathbf{x}_n - \mu_k^{i+1} \right) \left( \mathbf{x}_n - \mu_k^{i+1} \right)^T.$$



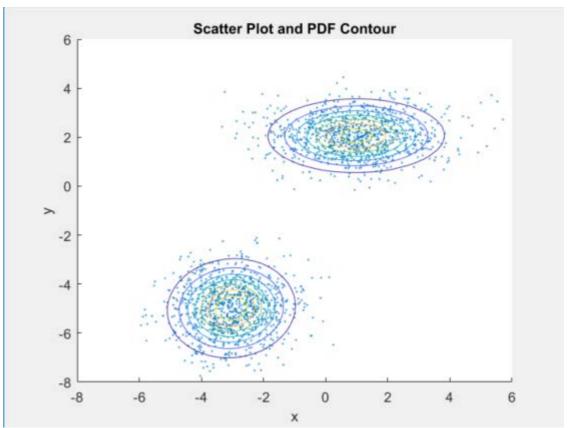
## A simple Matlab example

- Matlab functions:
  - fitgmdist (Fit a Gaussian mixture distribution to data)
  - pdf (Density function of a specific ditribution)

Raw data (2 clusters of 1000 points each)



# Data model with 2 Gaussian Mixture distributions



Run: gmm\_example.m



## A simple Matlab example (cont.)

Silhouette value (S)

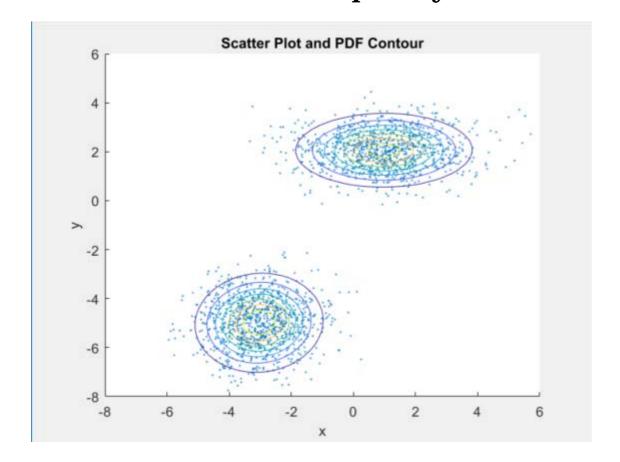
It is a measure of how similar a point is to a point in its own cluster.

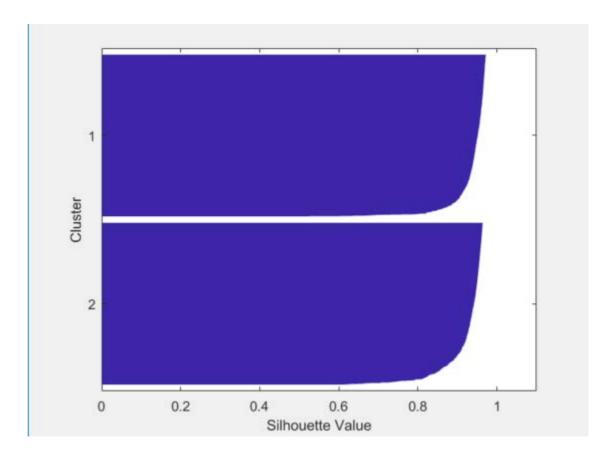
Minimum average distance from the  $i^{th}$  point to points in a different cluster

Average distance from  $i^{th}$  point to other points in the same cluster

For well match of i in its own cluster,  $b_i$  should be large and  $a_i$  small.

 $S_i$  ranges between -1 to +1. High  $S_i$  indicates that i is well-matched to its own cluster, and poorly-matched to neighboring clusters.



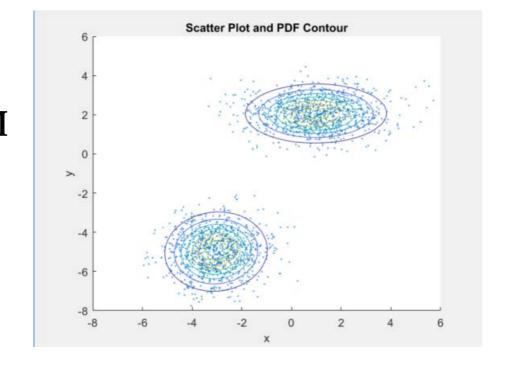


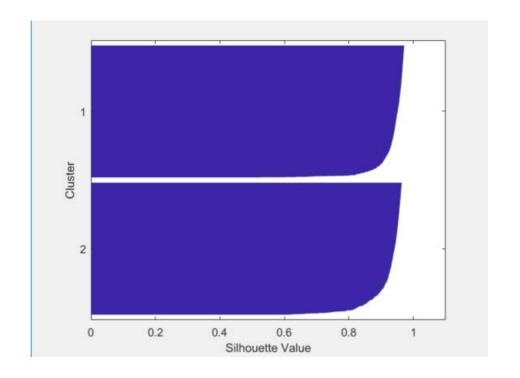


# A simple Matlab example (cont.)

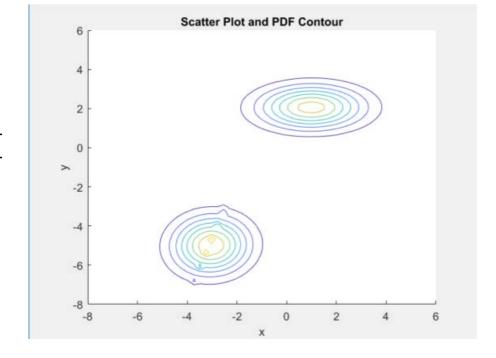
### Silhouette value (S)

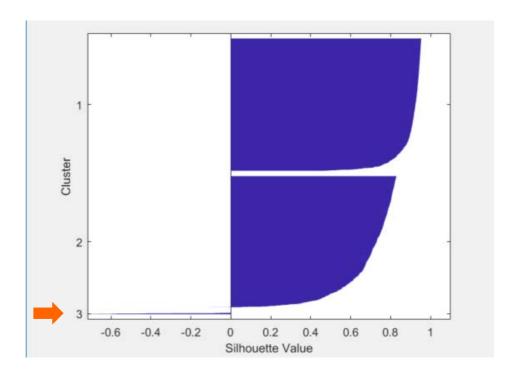
K=2 GM





K=3 GM



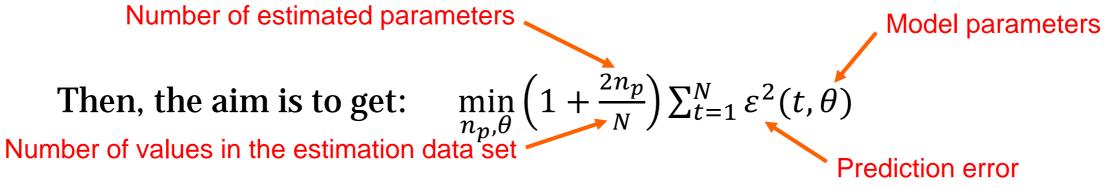




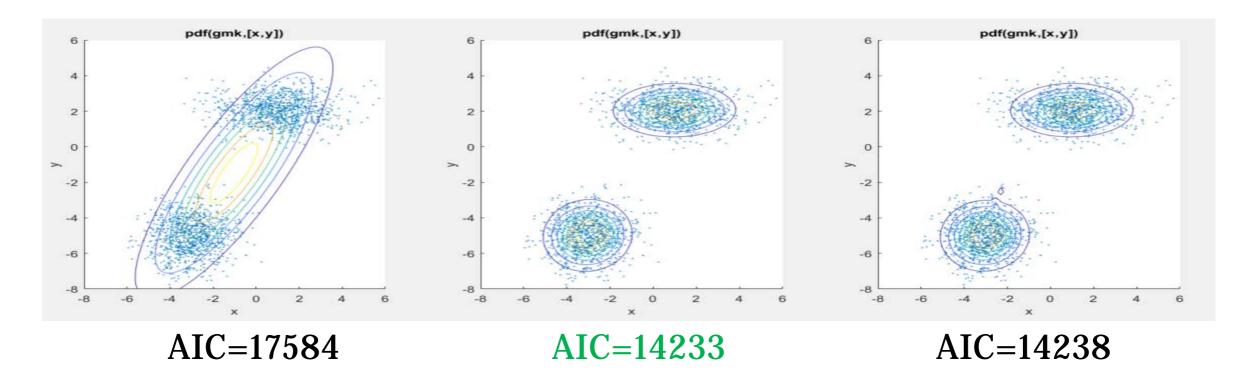
## A simple Matlab example (cont.)

Akaike's Information Criterion (AIC)

Provides a measure of the relative quality of a model for a given set of data.



The most accurate model has the smallest AIC.





# Case study

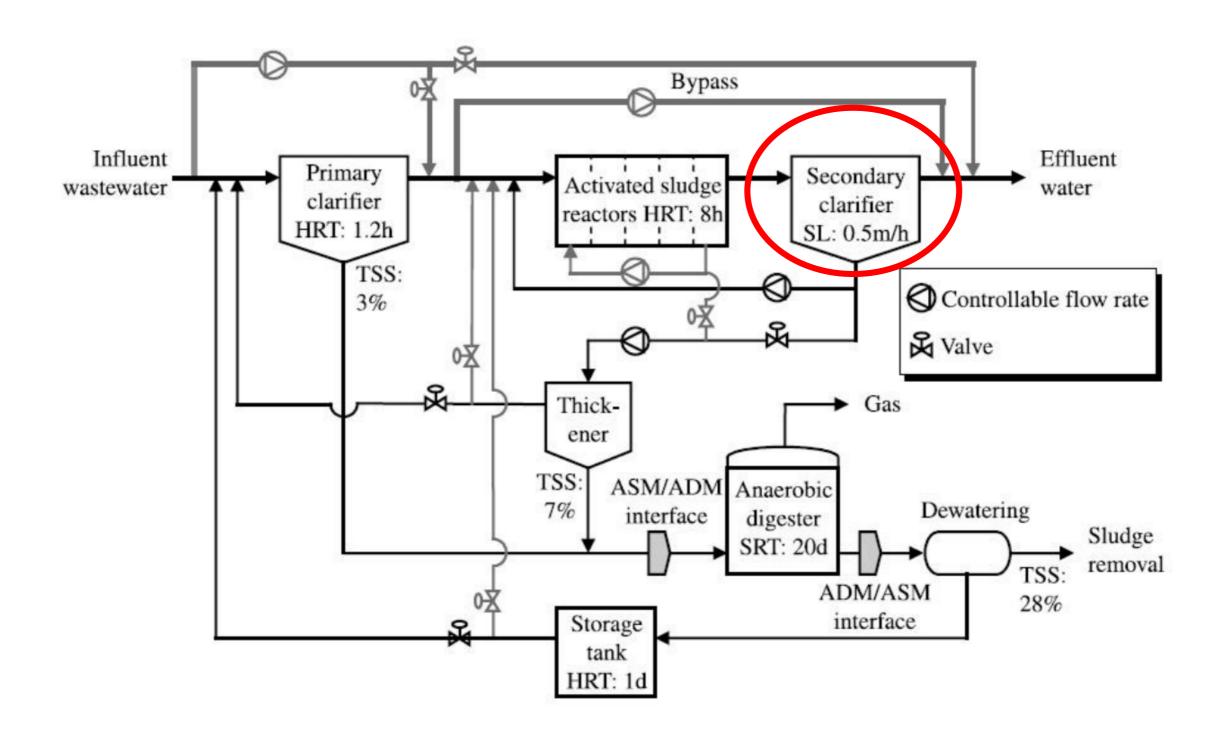


# A wastewater treatment plant





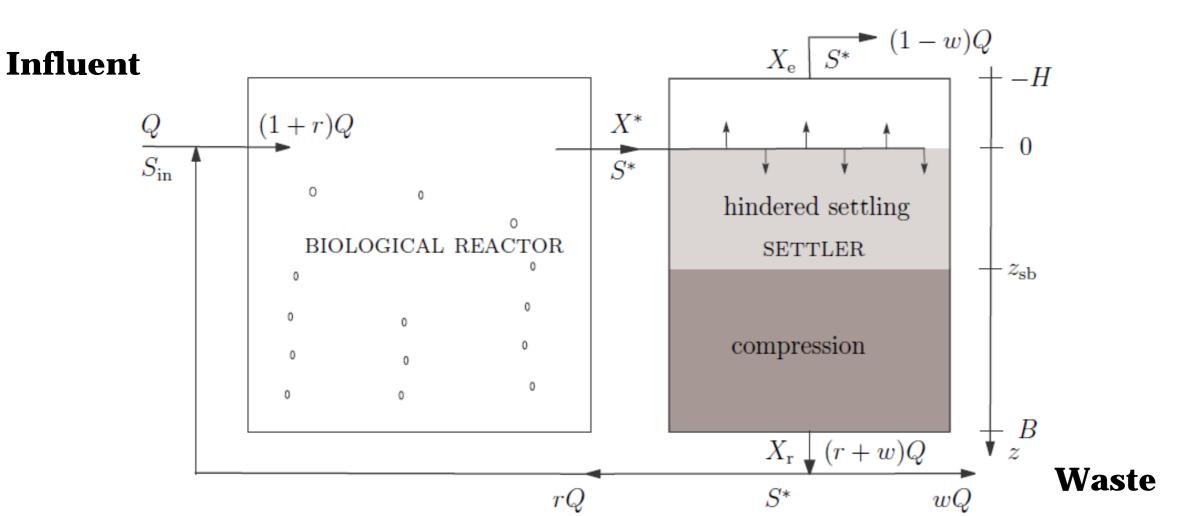
## A wastewater treatment plant (cont.)





## **The Process**

#### **Effluent**



*Q*: flowrate

*S*: conc. soluble substrate

*X*: conc. biomass

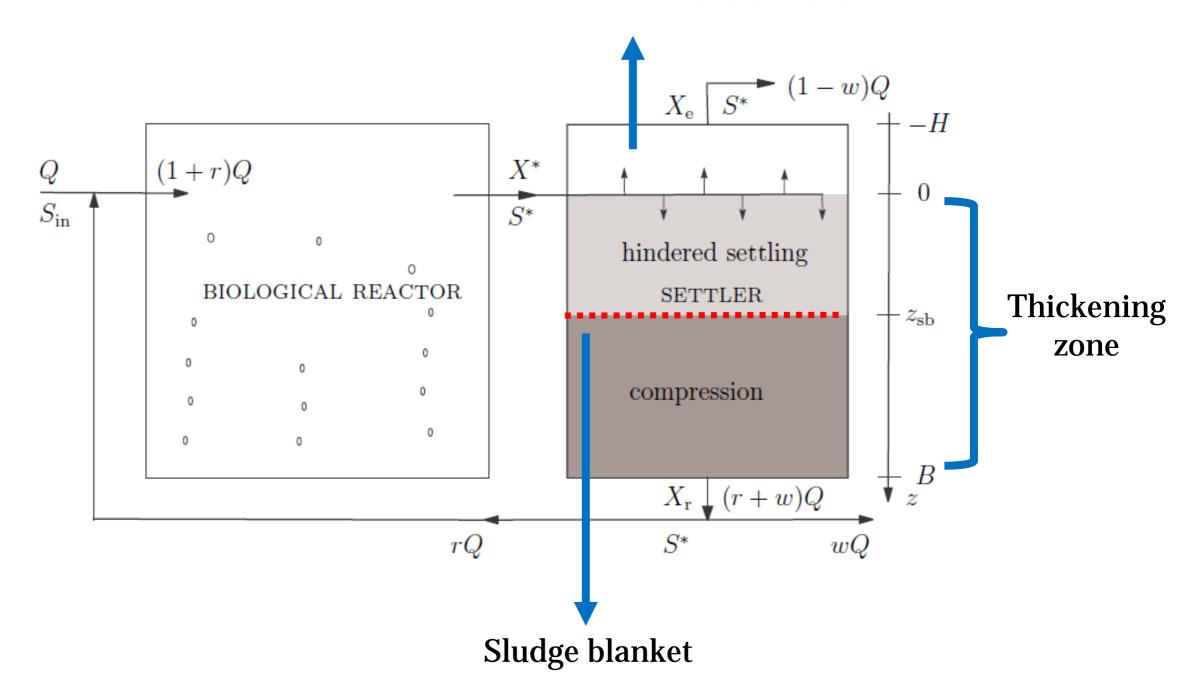
*r*: recycle ratio

w: wastage ratio



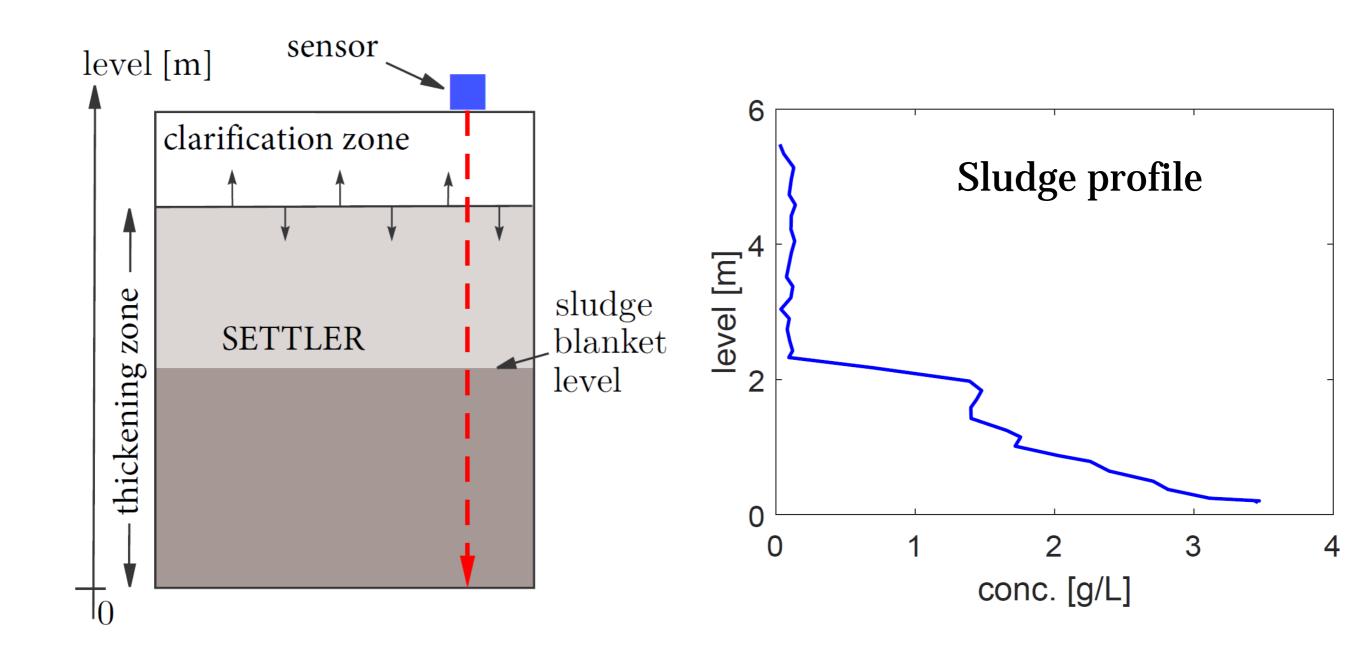
## The Process (cont.)

#### Clarification zone





## Scanning a secondary settler

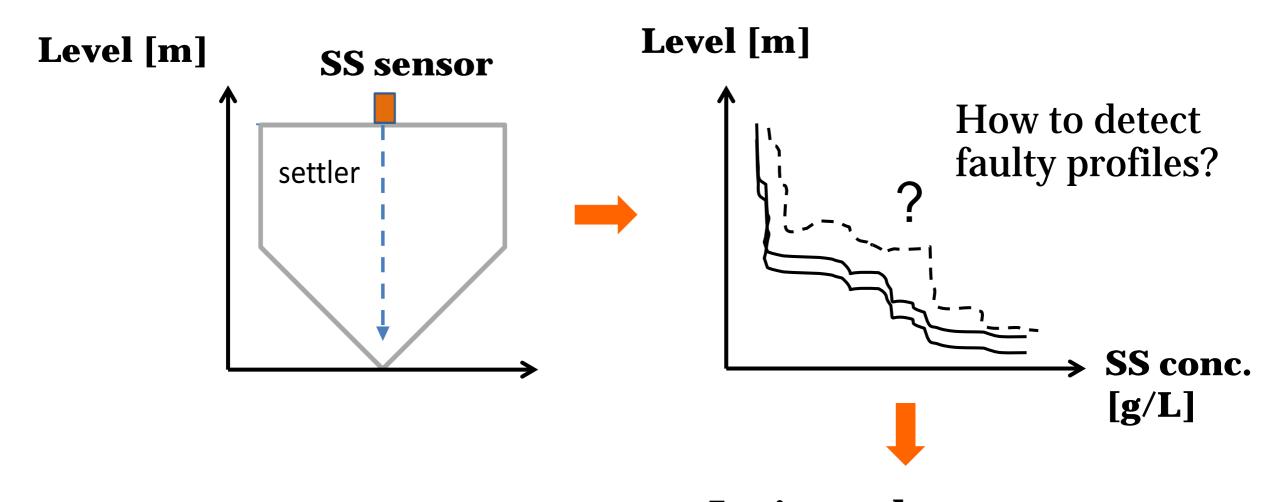




## The Problem

Scanning

Sludge profiles

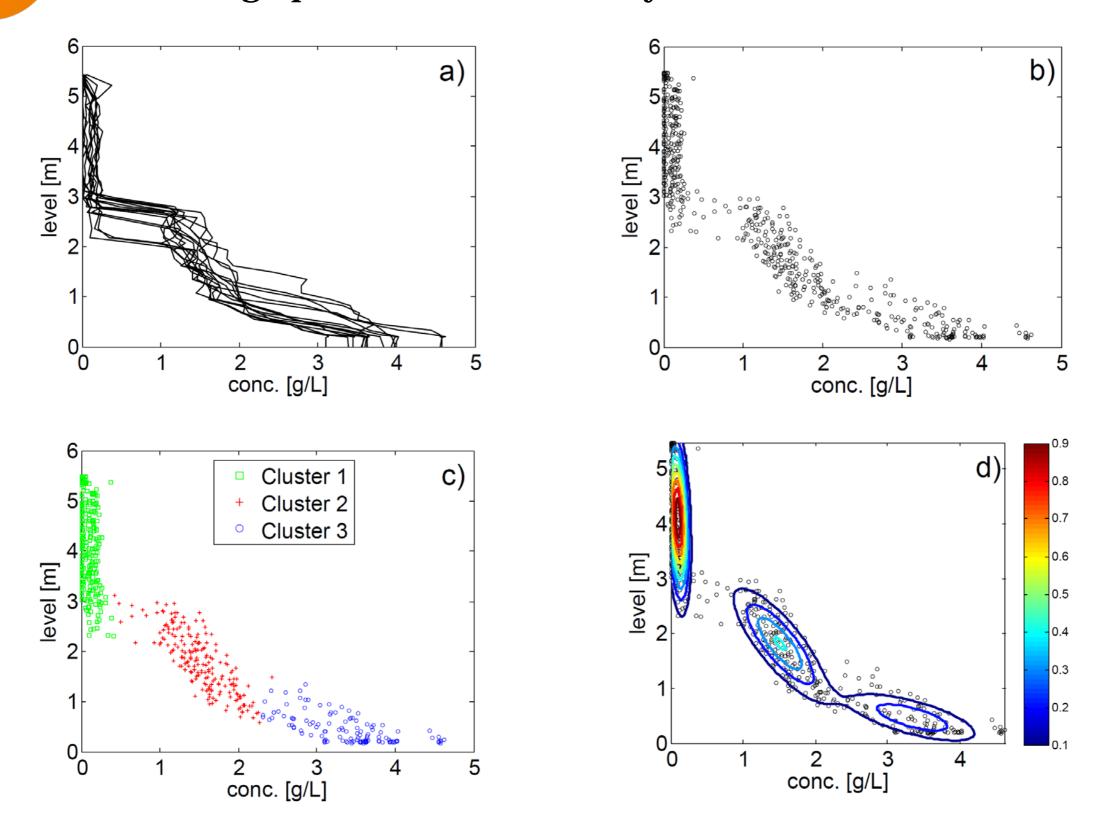


Let's apply Gaussian Mixture Models!



## **GMM** for the settler

### 15 sludge profiles in non-faulty conditions





## GMM for the settler (cont.)

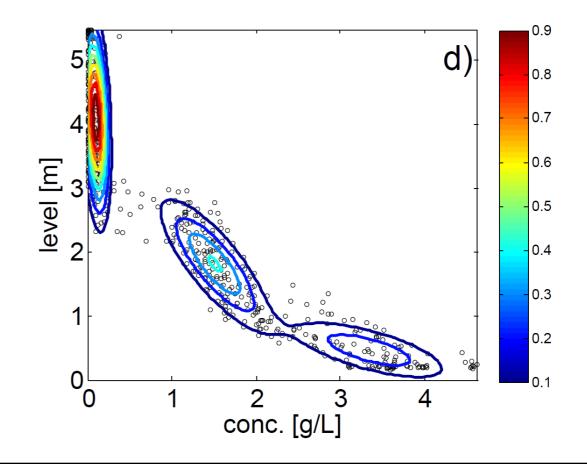
### GMM parameters $\pi_k$ , $\mu_k$ , $\sigma_k$ :

#### We denote

$$x_1 = \{SS \text{ conc.}\}\ \text{and}\ x_2 = \{level\}$$

$$\mu_k = \begin{bmatrix} \operatorname{mean}(x_1) \\ \operatorname{mean}(x_2) \end{bmatrix},$$

$$\sigma_k = \begin{bmatrix} \operatorname{cov}(x_1, x_1) & \operatorname{cov}(x_1, x_2) \\ \operatorname{cov}(x_2, x_1) & \operatorname{cov}(x_2, x_2) \end{bmatrix},$$

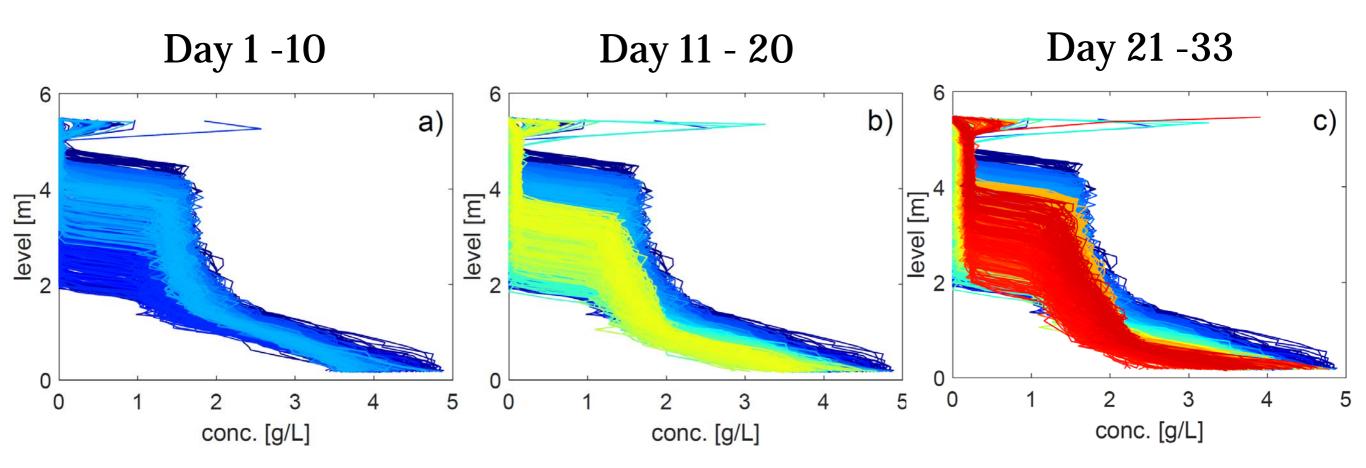


Weight	Mean	Covariance
$\pi_1 = 0.4329$	$\mu_1 = \begin{bmatrix} 0.0958 \\ 4.1102 \end{bmatrix}$	$\sigma_1 = \begin{bmatrix} 0.0074 & -0.0223 \\ -0.0223 & 0.7084 \end{bmatrix}$
$\pi_2 = 0.3405$	$\mu_2 = \begin{bmatrix} 1.5065 \\ 1.8203 \end{bmatrix}$	$\sigma_2 = \begin{bmatrix} 0.1446 & -0.1840 \\ -0.1840 & 0.3550 \end{bmatrix}$
$\pi_3 = 0.2265$	$\mu_3 = \begin{bmatrix} 3.3421 \\ 0.4691 \end{bmatrix}$	$\sigma_3 = \begin{bmatrix} 0.3612 & -0.1208 \\ -0.1208 & 0.0866 \end{bmatrix}$



## Settler monitoring

- Sludge profiles from day 1 (blue) to day 33 (red).
- New profile every 15 minutes = 3168 profiles.



(Red does not mean alarm!)



## Residual and Fault detection criteria

#### Algorithm 2 GMM-based residual calculation

- 1: Collect a group of M-profiles in non-faulty conditions.
- 2: Set K and compute the iterative EM algorithm (see Algorithm 1) to get  $\pi_k, \mu_k, \sigma_k$ .
- 3: while monitoring a new profile do
- for every profile do

5:

threshold

$$r = \frac{1}{p(\mathbf{x}; \pi_{1:K}, \mu_{1:K}, \sigma_{1:K})},$$
 (7)

where

$$p(\mathbf{x}; \pi_{1:K}, \mu_{1:K}, \sigma_{1:K}) = \sum_{n=1}^{N} \sum_{k=1}^{K} \pi_k \mathcal{N}(\mathbf{x}_n | \mu_k, \sigma_k).$$
(8)

- end for 6:
- 7: end while



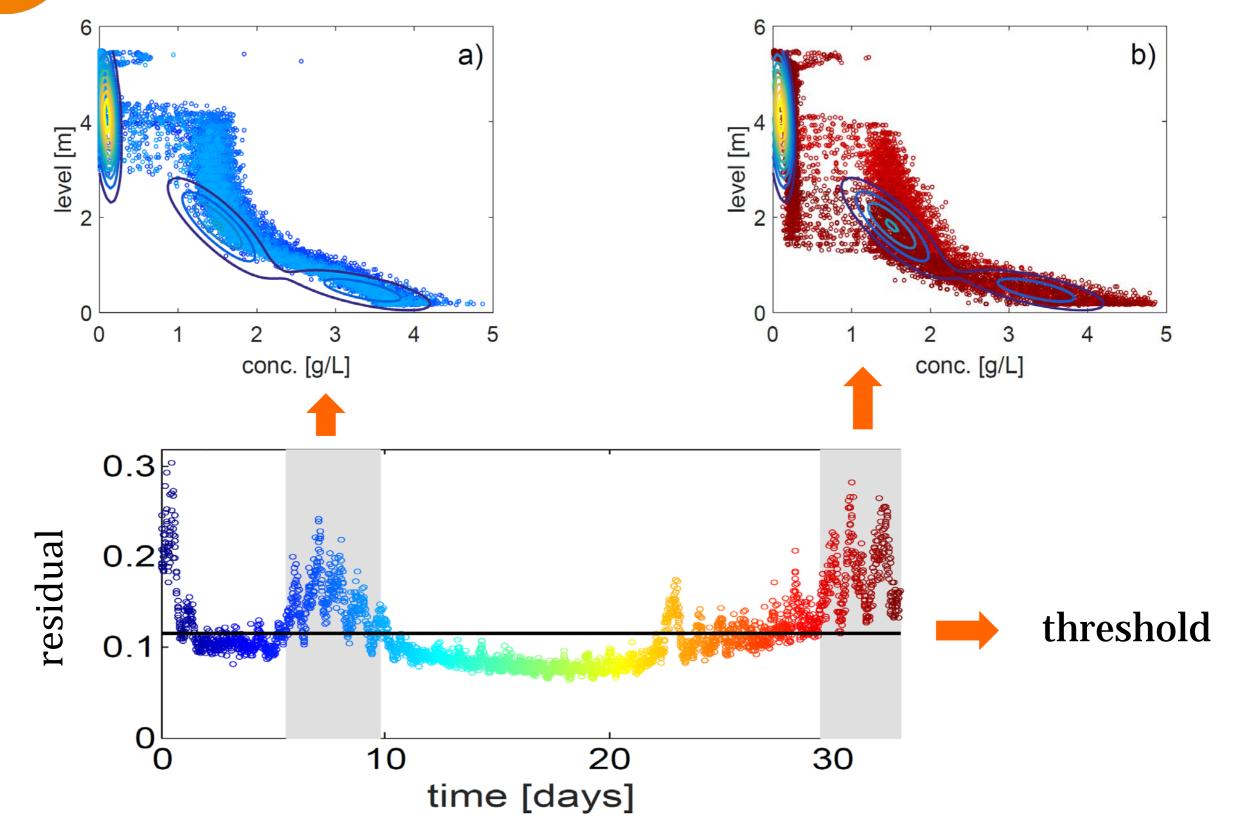
 $H_0: r \le h$  normal where  $H_1: r > h$  faulty!  $h = \max\{r\}$ 

$$h = \max\{r\} \Big|_{t \in H_0}$$

Classical binary hypothesis testing problem



# Settler monitoring (cont.)





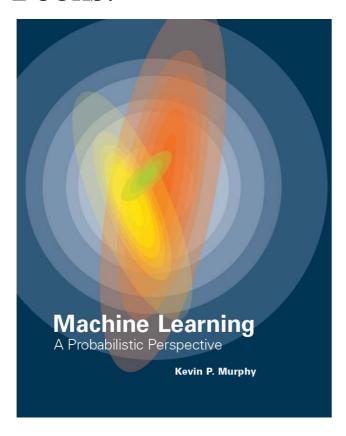
## Conclusions

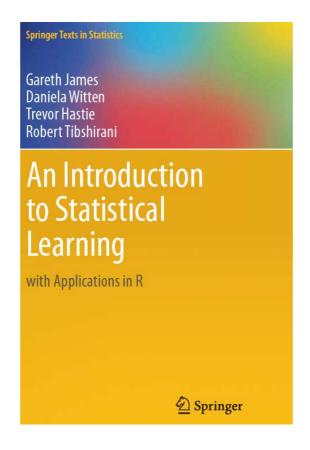
- Valuable information can be obtained by monitoring a Secondary Settler in a wastewater treatment plant.
- Gaussian Mixture Models provide a novel tool for fault detection in this process.
- The proposed method is general and could be implemented in settlers with different geometries and sludge profiles.
- The method is also suitable for monitoring deviations in a process with repetitive data profiles.

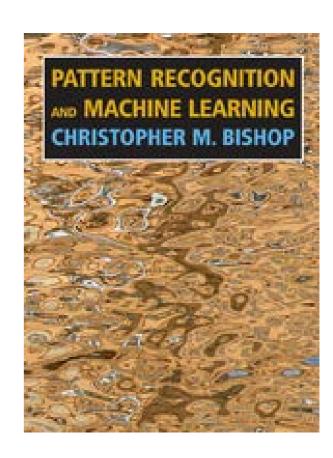


## Sources of information

Books:



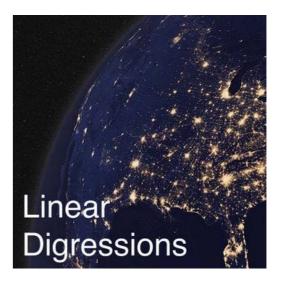




Podcasts:



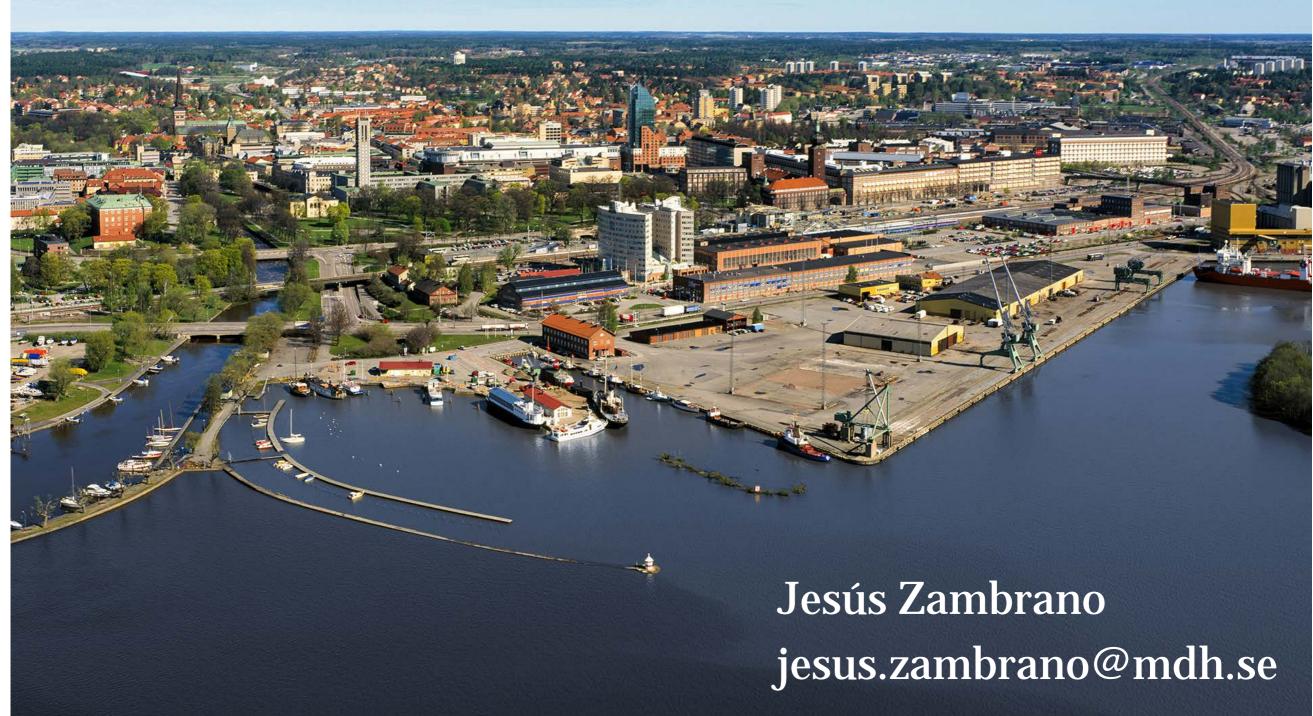


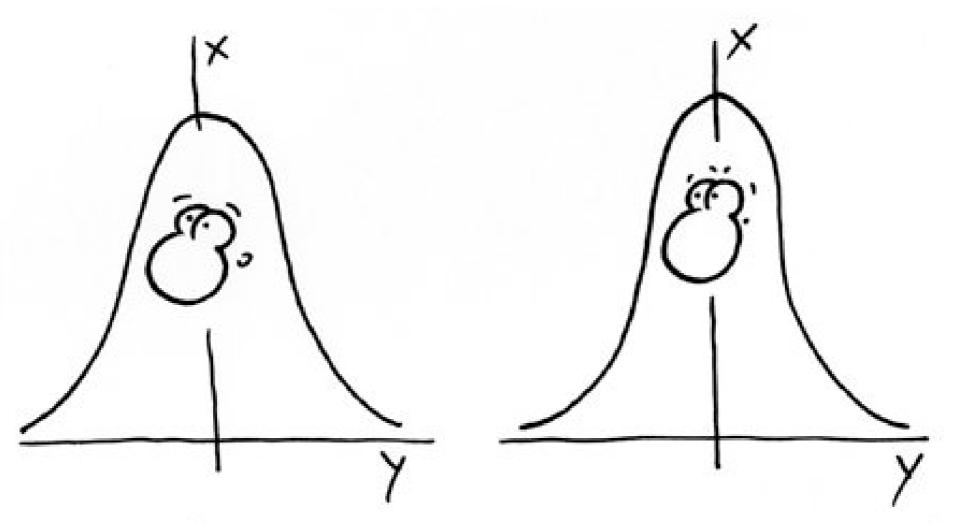




# Thanks for your attention!







"I always feel so normal, so bored, you know. Sometimes I would like to do something... you know... something... mmm... Poissonian."