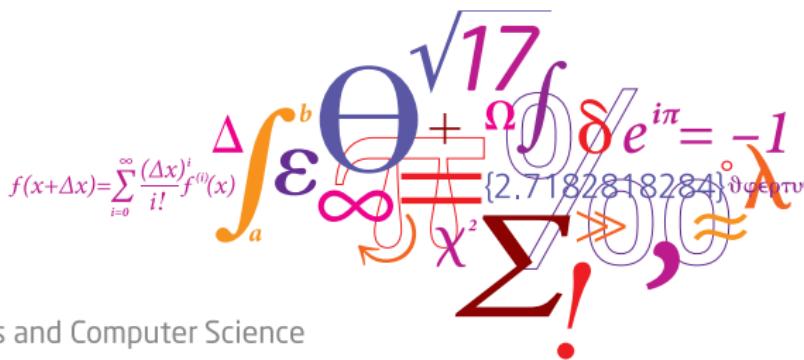


02450: Introduction to Machine Learning and Data Mining

Mixture models and density estimation



# Reading Material

## Reading material:

C17, C18

## Feedback Groups of the day:

- Nathalie Jaure, Andine Havelange, Anne-Line Evenstad Dahlen
- Rebecca Christiansen, Line Karin Mortensen, Simon Eugen Matell
- Andreas Vedel Jantzen, Anne Sofie Talleruphuus
- Mathias Kirkeskov Madsen, Andreas Gramstrup Correia, Iulian Cozma
- Arnor Ingi Sigurdsson, Paul Jacques Connetable, Grigory Solomatov
- David Rosendahl Pedersen, Jacob Bundgaard Knudsen, Sune Mikkel Rasmussen
- Vilde Olerud, Rebecca Mølgaard Sommer,

Tue Herlau, Mikkel N. Schmidt and Morten Mørup

Introduction to Machine Learning and Data Mining

Course notes fall 2016, version 1

August 29, 2016

Technical University of Denmark

# Lecture Schedule

## ① Introduction

30 August: C1

### Supervised learning: Feature extraction, and visualization

## ② Data and feature extraction

6 September: C2, C3

## ③ Measures of similarity and summary statistics

13 September: C4

## ④ Data Visualization and probability

20 September: C5, C6

### Supervised learning: Classification and regression

## ⑤ Decision trees and linear regression

27 September: C7, C8 (Project 1 due before 13:00)

## ⑥ Overfitting and performance evaluation

4 October: C9

## ⑦ Nearest Neighbor, Bayes and Naive Bayes

11 October: C10, C11

## ⑧ Artificial Neural Networks and Bias/Variance

25 October: C12, C13

## ⑨ AUC and ensemble methods

1 November: C14, C15

### Unsupervised learning: Clustering and density estimation

## ⑩ K-means and hierarchical clustering

8 November: C16 (Project 2 due before 13:00)

## ⑪ Mixture models and density estimation

15 November: C17, C18

## ⑫ Association mining

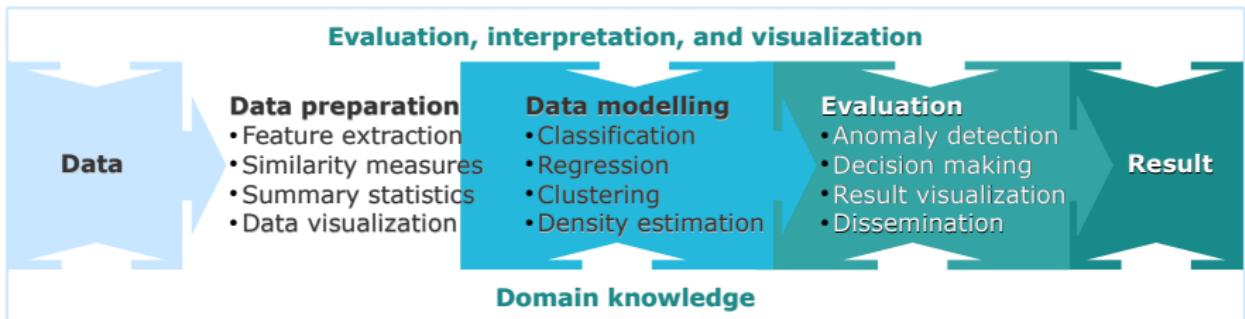
22 November: C19

### Recap

## ⑬ Recap and discussion of the exam

29 November: C1-C19 (Project 3 due before 13:00)

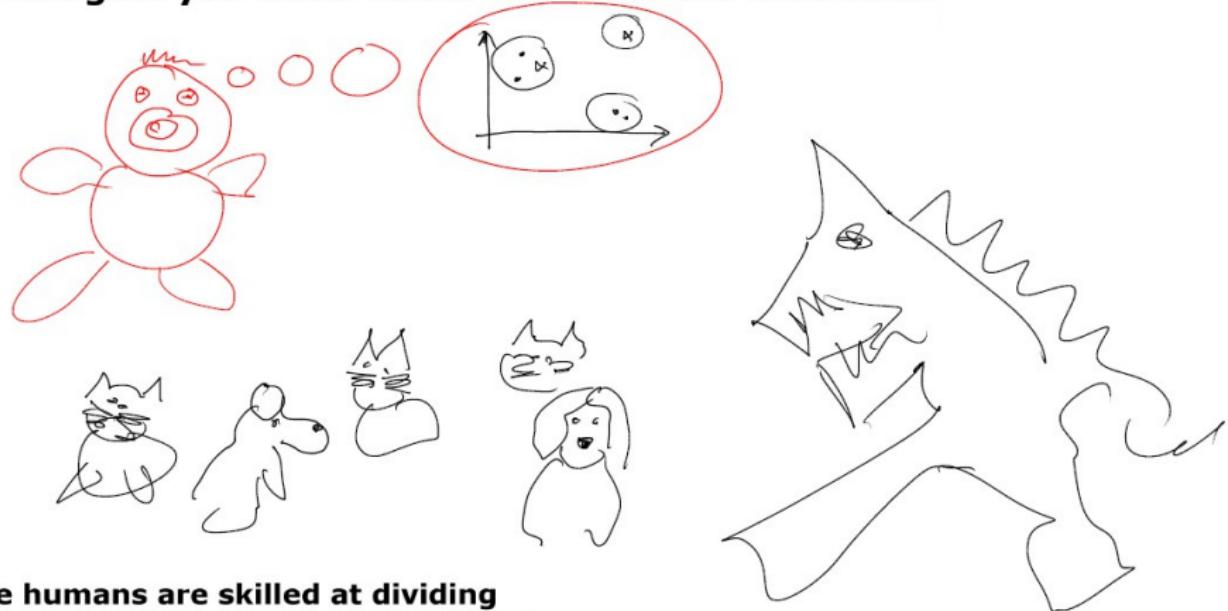
# Data modeling framework



## After today you should be able to:

- Explain the role of the parameters in the Gaussian Mixture Model (GMM)
- and how the parameters are updated using the EM-algorithm
- Explain why cross-validation can be used for GMM
- Understand and apply kernel density, K-nearest neighbour density and average relative density estimation for outlier/anomaly detection.

**Imagine you observe the world for the first time!**

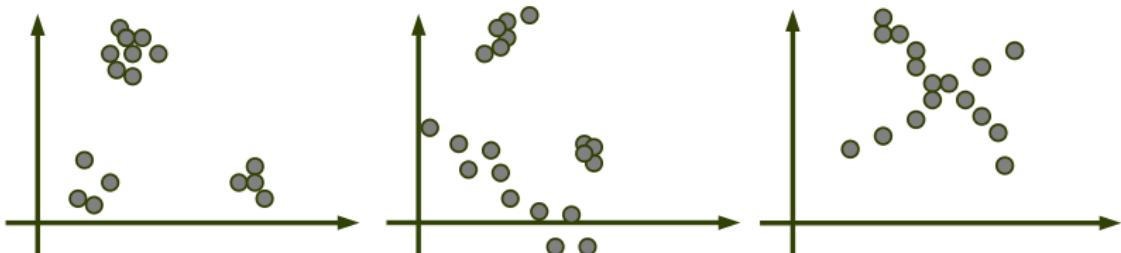


**We humans are skilled at dividing  
objects into groups (clustering), but  
how do we make computers do the  
same?**



## Group exercise

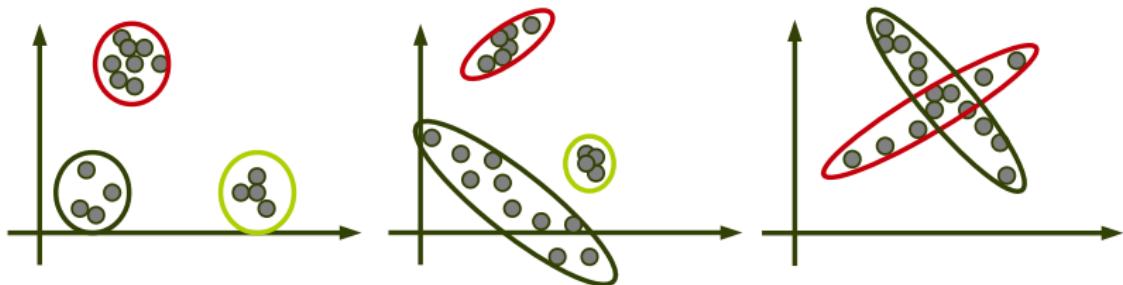
- What are the clusters below and what characterize each cluster?
- Is **k-means** well suited for modeling the clusters below?
  - Will it always find the optimum solution?
  - Can it model the sizes of the clusters?
  - Can it model the shape of the clusters?
  - How can we determine the number of clusters?





## Group exercise

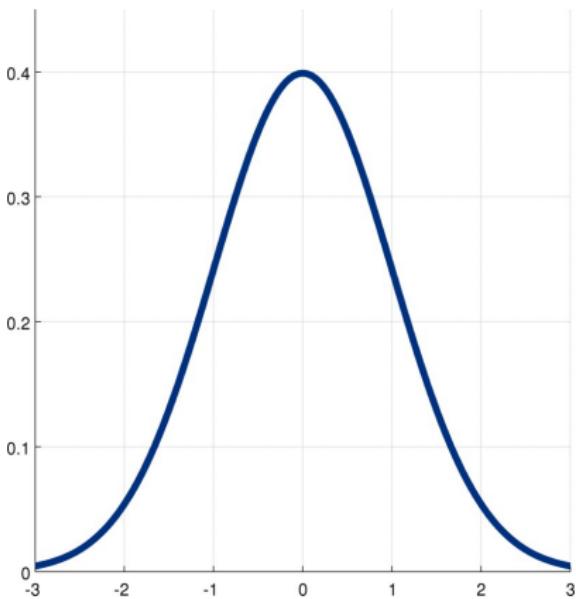
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# Normal distribution

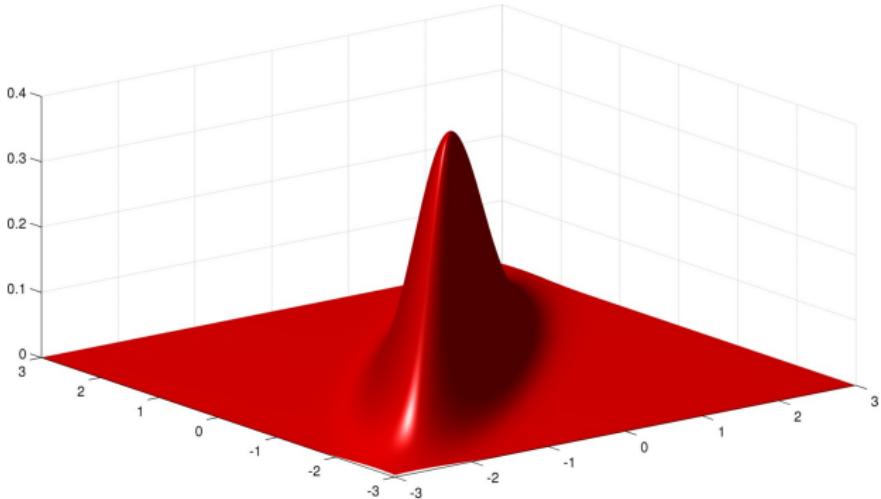
- Probability density function describes the relative chance of a given value to occur
- Normal distribution characterized by
  - Mean
  - Variance

$$p(x) = \frac{1}{\sqrt{2\pi\sigma^2}} \exp\left(-\frac{(x - \mu)^2}{2\sigma^2}\right)$$



# Multivariate Normal distribution

$$p(x) = \frac{1}{(2\pi)^{k/2} |\Sigma|^{1/2}} \exp \left( -\frac{1}{2} (x - \mu)^\top \Sigma^{-1} (x - \mu) \right)$$



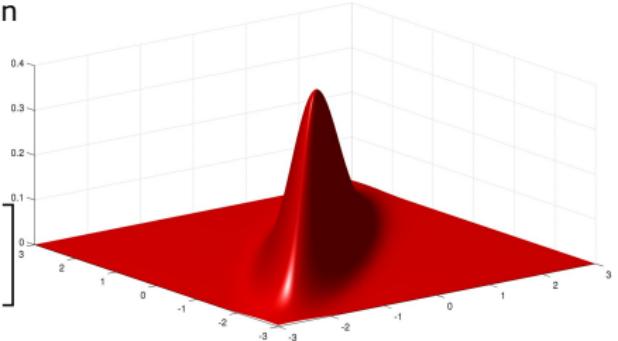
# Multivariate Normal distribution

$$p(x) = \frac{1}{(2\pi)^{k/2} |\Sigma|^{1/2}} \exp \left( -\frac{1}{2} (x - \mu)^\top \Sigma^{-1} (x - \mu) \right)$$

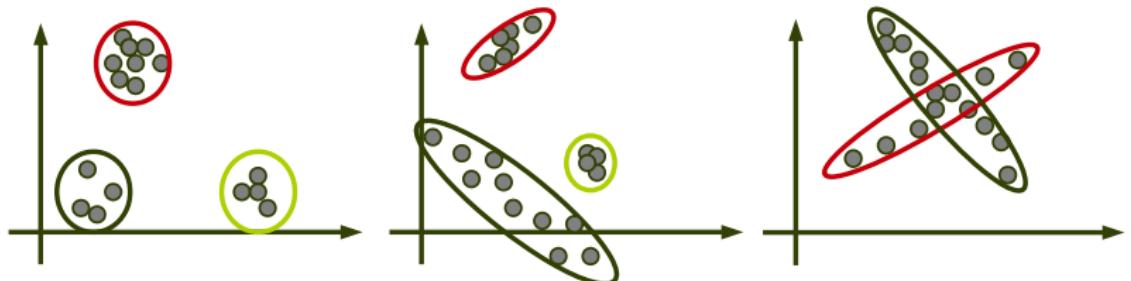
- Example: 2-dimensional Normal distribution

$$\mu = \begin{bmatrix} \bar{x}_1 \\ \bar{x}_2 \end{bmatrix}$$

$$\Sigma = \begin{bmatrix} \text{var}(x_1) & \text{cov}(x_1, x_2) \\ \text{cov}(x_2, x_1) & \text{var}(x_2) \end{bmatrix}$$



# The Gaussian Mixture Model (GMM)



- Different locations
- Different shape
- Different sizes

$$\mu_{(k)}$$

$$\Sigma_{(k)}$$

$$w_k$$

Data density

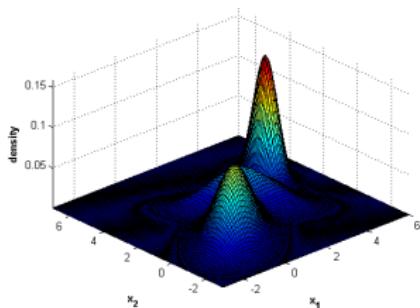
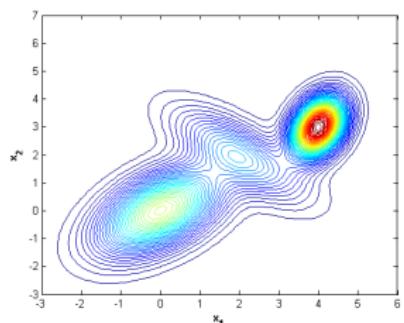
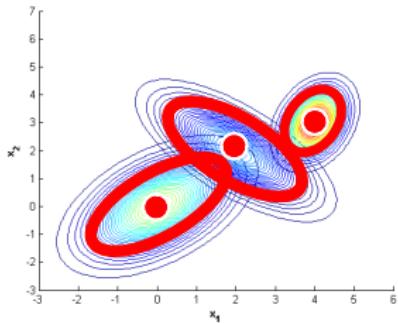
$$p(\mathbf{x}) = \sum_{k=1}^K w_k \mathcal{N}(\mathbf{x}, \mu_{(k)}, \Sigma_{(k)})$$

$$\text{s.t. } \sum_{k=1}^K w_k = 1, \quad w_k \geq 0$$

Sum of cluster specific densities  
assumed normal distributed

## GMM example

$$p(\mathbf{x}) = 0.5\mathcal{N}(\mathbf{x} | \begin{bmatrix} 0 \\ 0 \end{bmatrix}, \begin{bmatrix} 1 & 0 \\ 0 & 0.5 \end{bmatrix}) + 0.2\mathcal{N}(\mathbf{x} | \begin{bmatrix} 2 \\ 2 \end{bmatrix}, \begin{bmatrix} 1 & 0.7 \\ -0.7 & 1 \end{bmatrix}) + 0.3\mathcal{N}(\mathbf{x} | \begin{bmatrix} 4 \\ 3 \end{bmatrix}, \begin{bmatrix} 0.2 & 0.1 \\ 0.1 & 0.5 \end{bmatrix})$$



$\mu_{(k)}$ : Cluster center (prototypical example in cluster)

$\Sigma_{(k)}$ : Shape of the cluster

$w_k$  : Relative size/density of the cluster



## Group exercise

- Consider the Gaussian mixture model (GMM)

$$p(\mathbf{x}) = \sum_{k=1}^K w_k \mathcal{N}(\mathbf{x} | \boldsymbol{\mu}_{(k)}, \boldsymbol{\Sigma}_{(k)}) \quad \text{s.t. } \sum_{k=1}^K w_k = 1, \quad w_k \geq 0$$

- What is the value of the integral?

$$\int p(\mathbf{x}) d\mathbf{x}$$

# Gaussian mixture models, EM algorithm

Select an initial set of model parameters  
(mean and covariance for each cluster)

**Repeat**

- **Expectation**

- For each object, calculate the probability of belonging to each distribution

- **Maximization**

- For each probability distribution, estimate parameters by maximum likelihood

**Until** the parameters do not change

## E-step

$$\boxed{p(z_n = k | \mathbf{x}_n) = \frac{w_k \mathcal{N}(\mathbf{x}_n | \boldsymbol{\mu}_{(k)}, \boldsymbol{\Sigma}_{(k)})}{\sum_{K=1}^K w_k \mathcal{N}(\mathbf{x}_n | \boldsymbol{\mu}_{(k)}, \boldsymbol{\Sigma}_{(k)})}}$$

## M-step

$$N_k = \sum_{n=1}^N p(z_n = k | \mathbf{x}_n)$$

$$\boldsymbol{\mu}_{(k)} = \frac{1}{N_k} \sum_{n=1}^N \mathbf{x}_n p(z_n = k | \mathbf{x}_n)$$

$$w_k = \frac{N_k}{N}$$

$$\boldsymbol{\Sigma}_{(k)} = \frac{1}{N_k} \sum_{n=1}^N (\mathbf{x}_n - \boldsymbol{\mu}_{(k)}) (\mathbf{x}_n - \boldsymbol{\mu}_{(k)})^\top p(z_n = k | \mathbf{x}_n)$$

# Gaussian mixture models, EM algorithm

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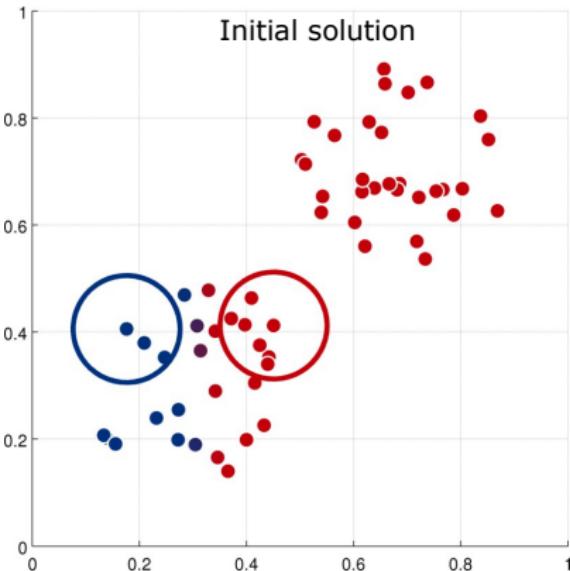
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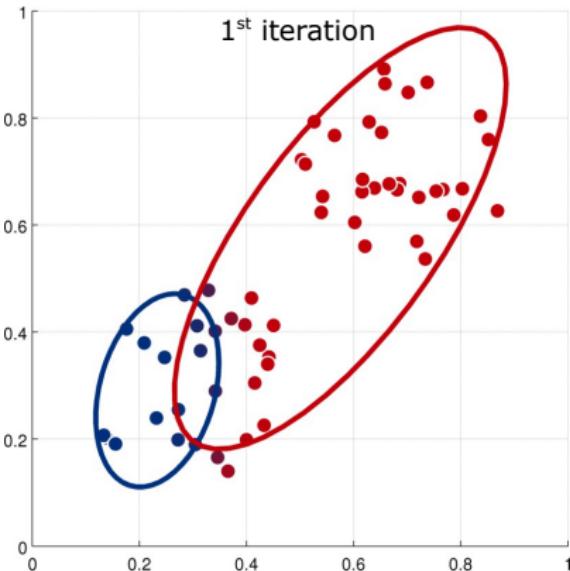
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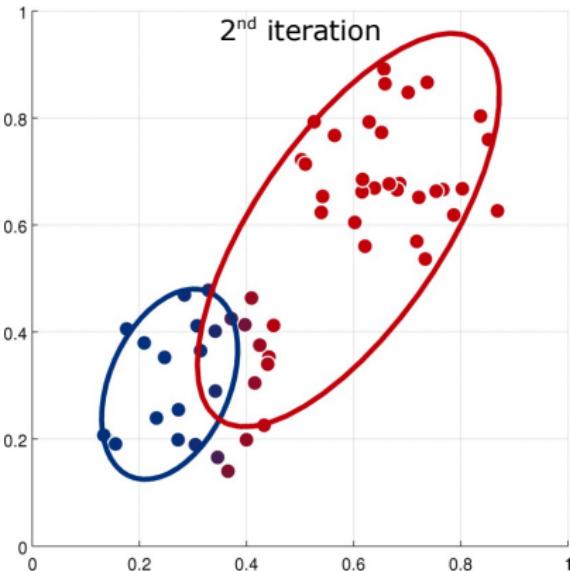
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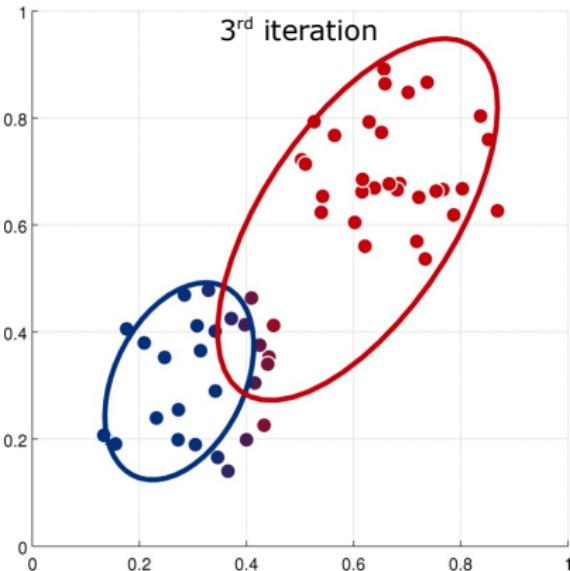
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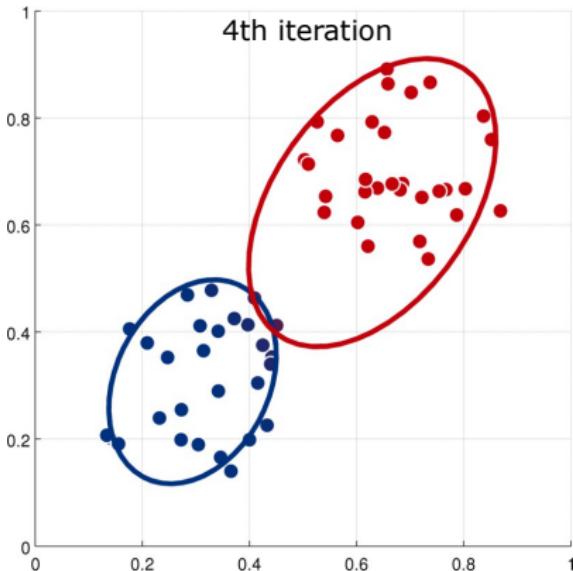
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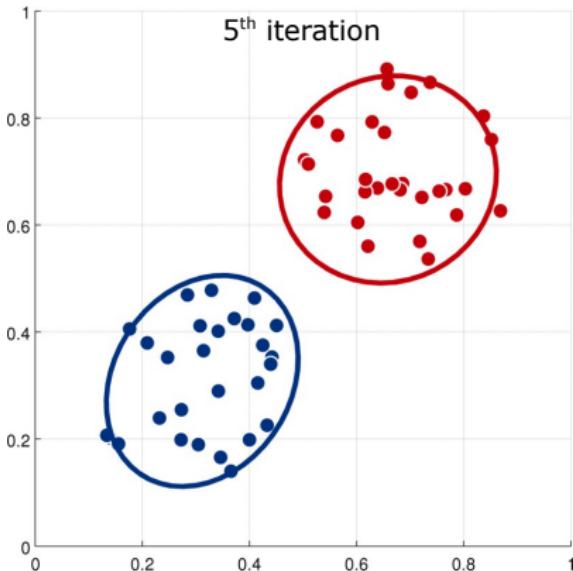
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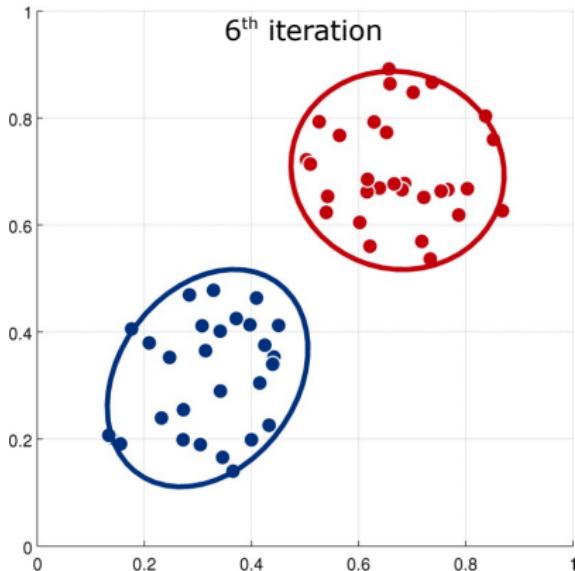
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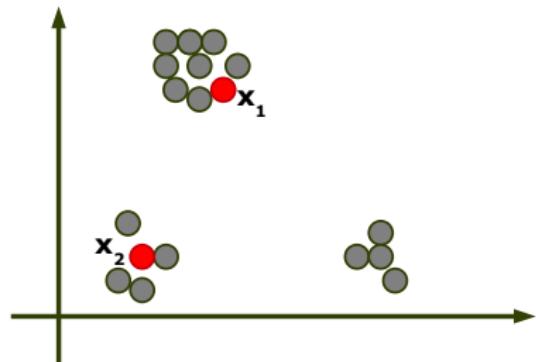
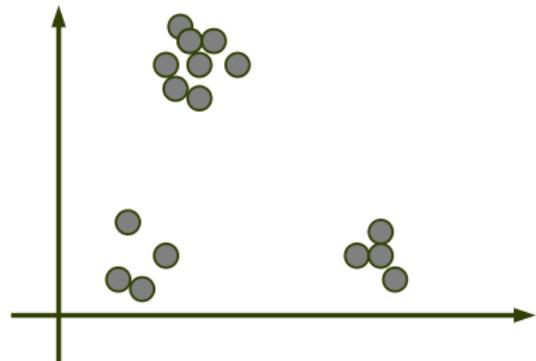
**Until** the parameters do not change





## Group exercise

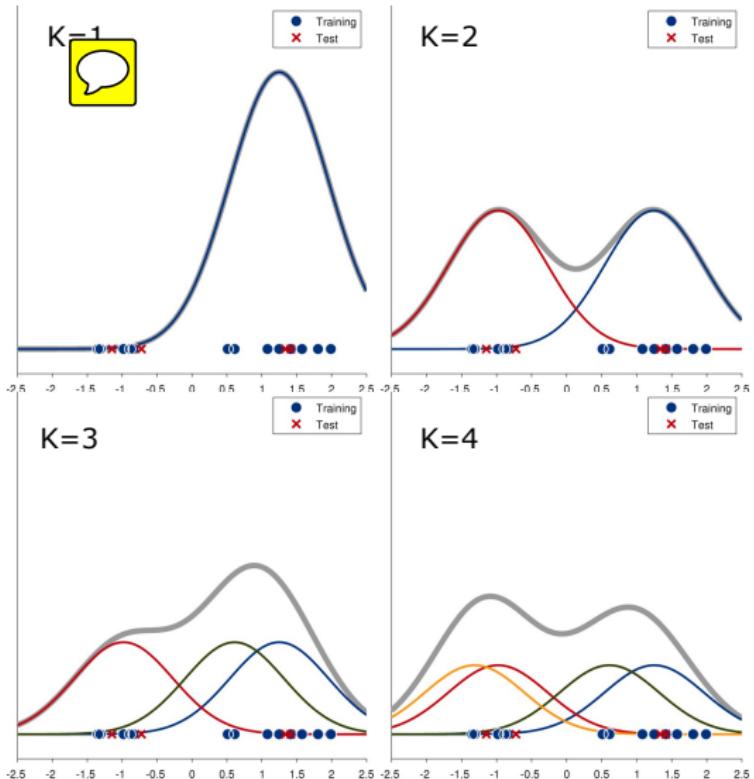
- Consider the data to the right with 16 observations.
  - What would ideally happen if we used a GMM with  $K=16$  clusters to model the data?
- Imagine we have two **test observations** denoted  $\mathbf{x}_1$  and  $\mathbf{x}_2$  (red points) that are not used for training.
  - What happens to  $p(\mathbf{x}_1)$  and  $p(\mathbf{x}_2)$  if we use  $K=3$  and  $K=16$  clusters



## EM Initial solution

# Mixture models

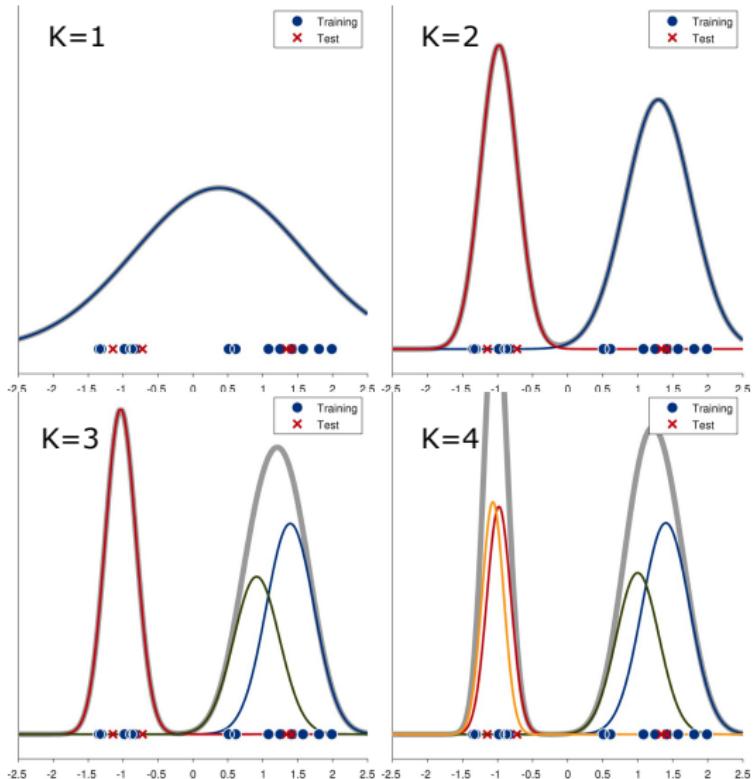
- Selecting complexity using crossvalidation



EM 1<sup>st</sup> iteration

# Mixture models

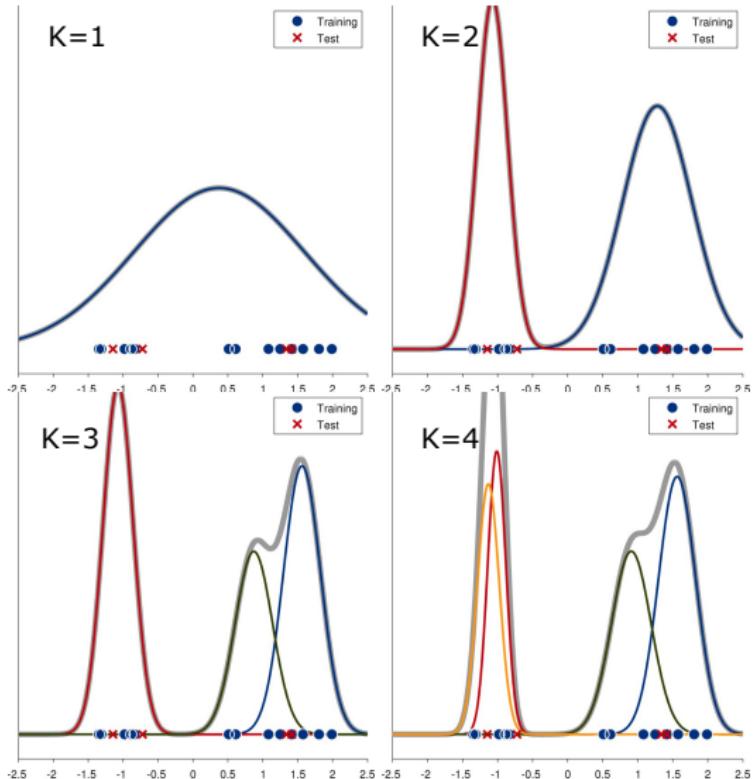
- Selecting complexity using crossvalidation



EM 2<sup>nd</sup> iteration

# Mixture models

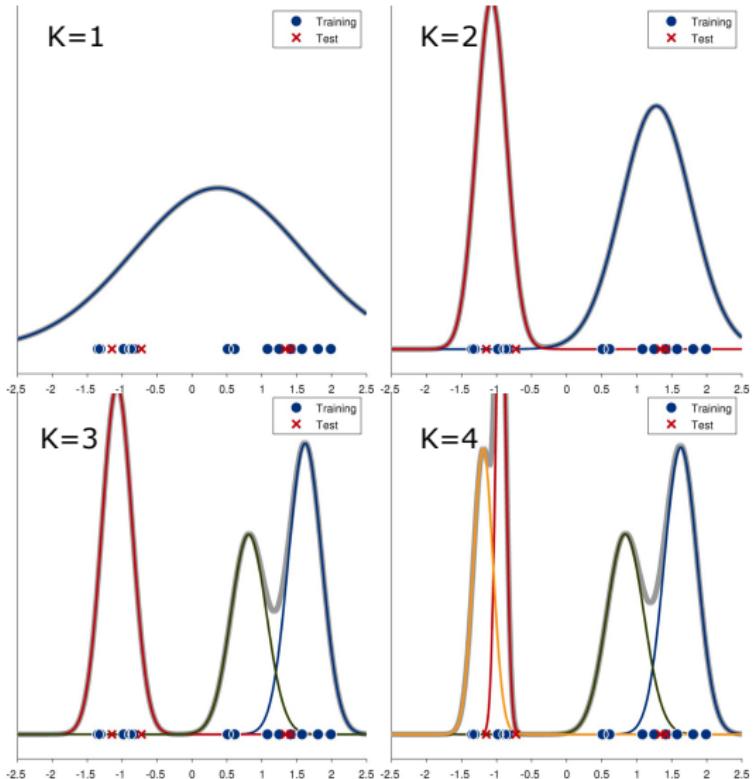
- Selecting complexity using crossvalidation



EM 3<sup>rd</sup> iteration

# Mixture models

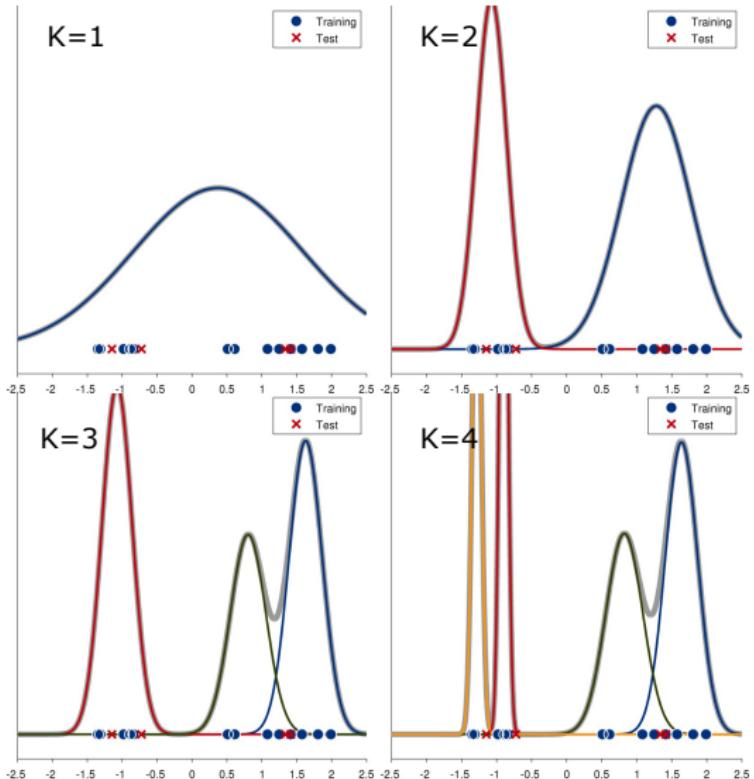
- Selecting complexity using crossvalidation



EM 4<sup>th</sup> iteration

# Mixture models

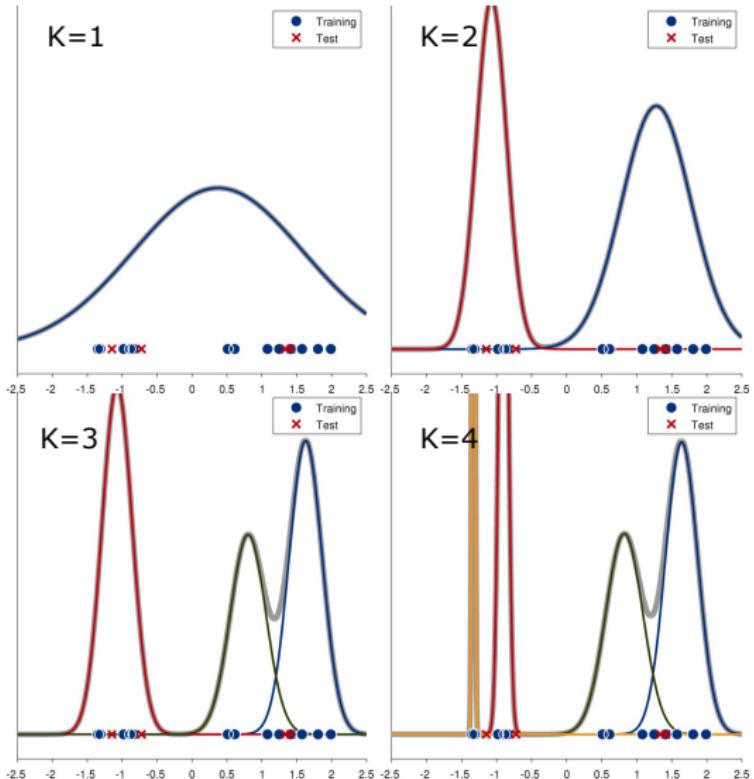
- Selecting complexity using crossvalidation



EM 5<sup>th</sup> iteration

# Mixture models

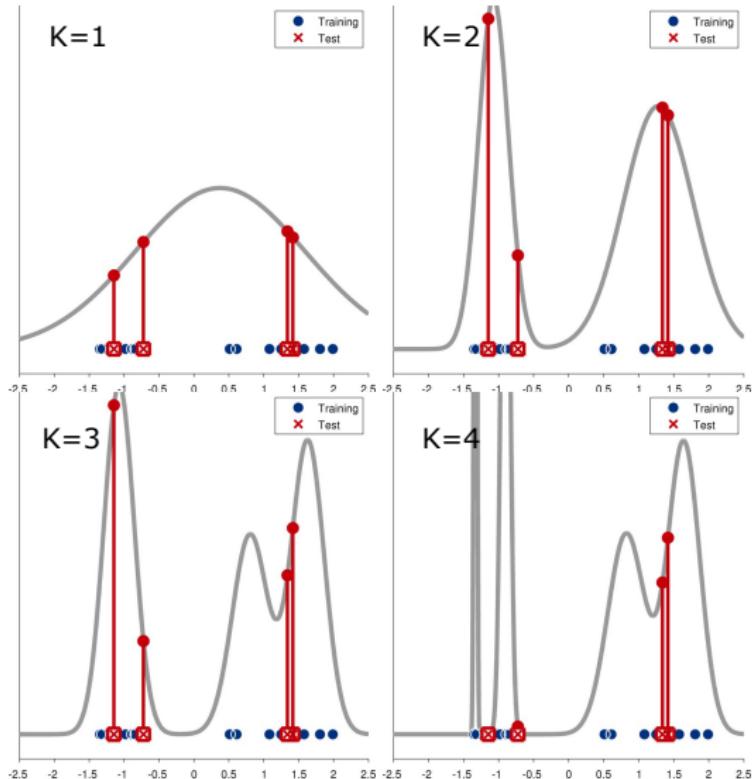
- Selecting complexity using crossvalidation



## Test data evaluation

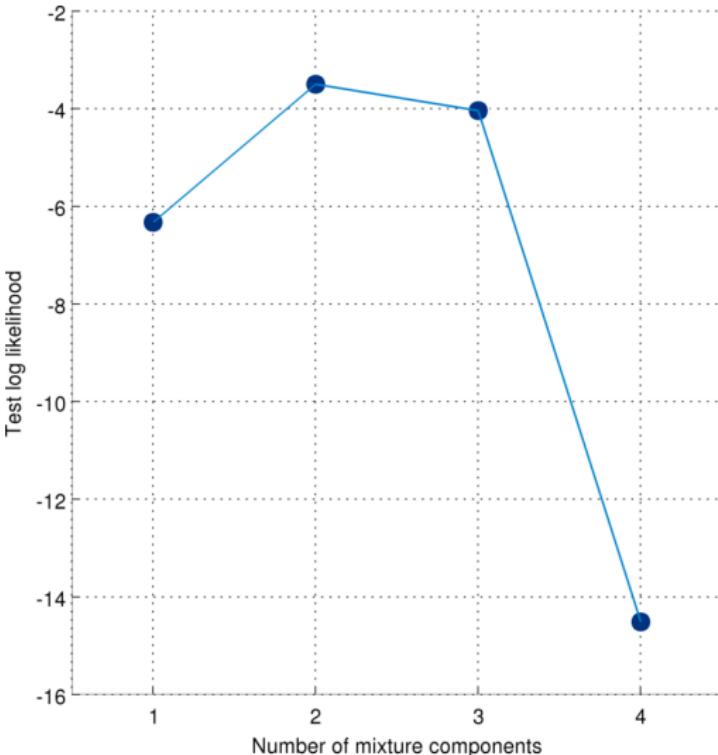
# Mixture models

- Selecting complexity using crossvalidation



## Mixture models

- Selecting complexity using crossvalidation



# K-means versus GMM

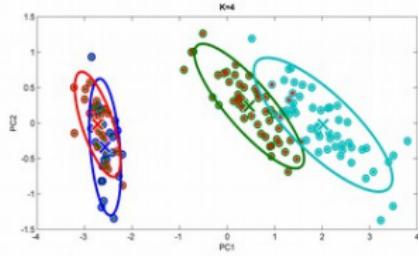
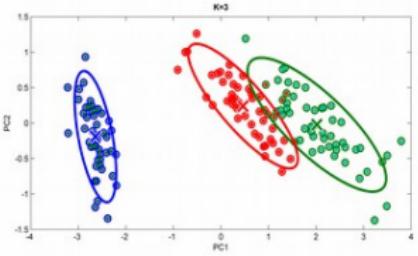
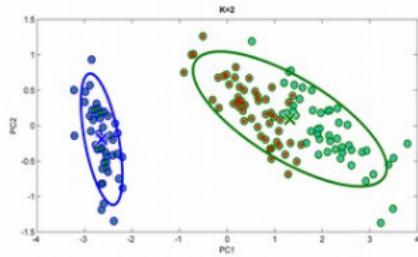
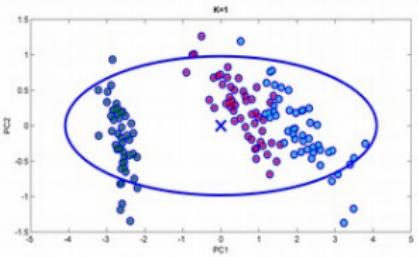
## K-means

- No guarantee of optimal solution
- Does not model shape of clusters
- Does not model the size of clusters
- Difficult to assess the number of clusters to use particularly when there is no ground truth

## Gaussian mixture model (GMM)

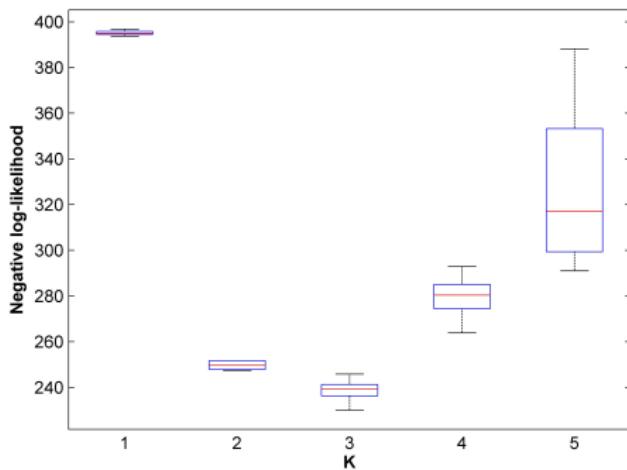
- No guarantee of optimal solution (even more local minima issues due to the additional model parameters)
- Models shape of cluster as ellipsoid
- Models the size of clusters
- Possible to estimate the number of components by cross-validation

# GMM on Iris data using 1,2,3 and 4 components



## Recap of GMM on Iris data

GMM 10 fold cross-validation on Iris data repeated five times where the five runs are plotted using box-plots.



## Anomaly detection: Definition

- Given a collection of data objects
  - Each object has associated a number of features
- Detect which objects **deviate from normal** behaviour

# Anomaly detection: Example

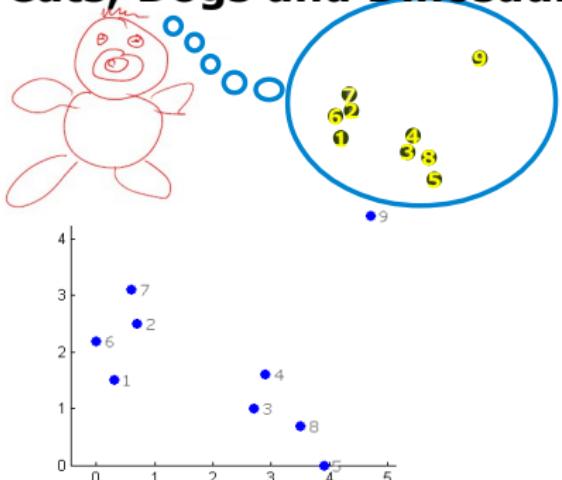
- Credit card **fraud detection**
  - Recognize dubious credit card transactions based on the transaction history of the card holder
- Network **intrusion detection**
  - Detect hacker attacks, web crawlers etc.
- **Ecosystem disturbances**
  - Detect hurricanes, floods droughts, heat waves and fires
- **Health and medicine monitoring**
  - Detect abnormal behaviour in populations and patients
- **Fault detection in industry systems**
  - Detect when a wind turbine performs poorly due to ice coating on blades
- Detection of **outliers** in data measurements
  - Remove erroneous measurements due to misreading from an instrument



## Group exercise

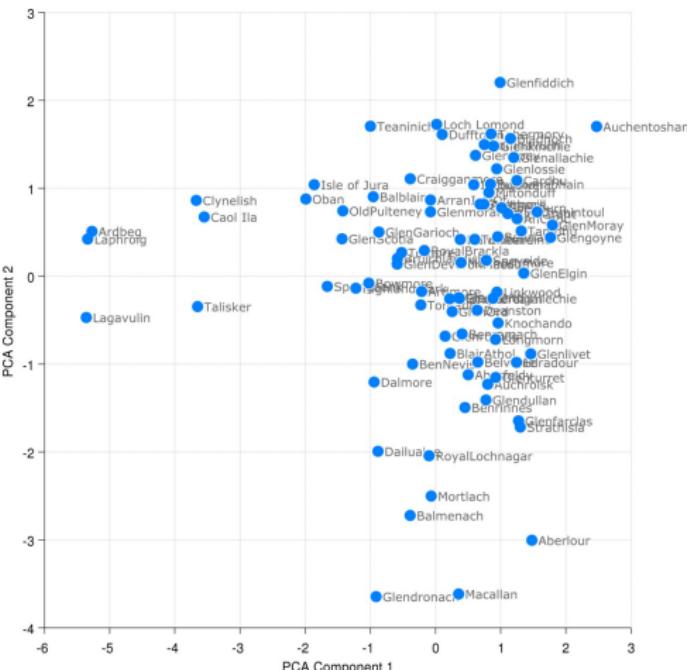
- Come up with **your own definition** of an outlier / anomaly
- How can we detect outliers using some of the methods you have already learned in the course?

## **Data example I: Cats, Dogs and Dinosaurs**

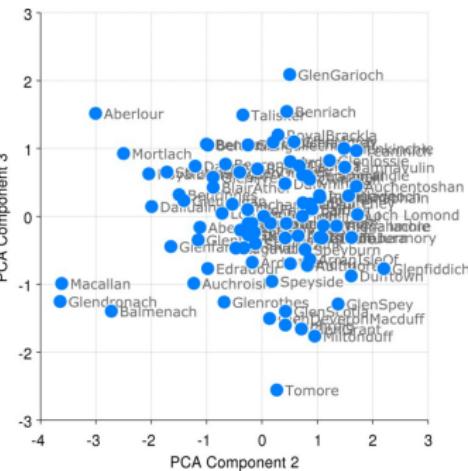
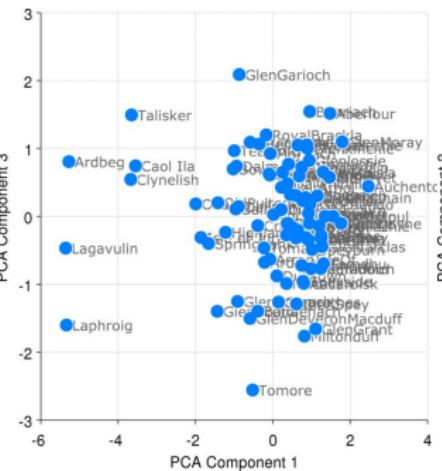
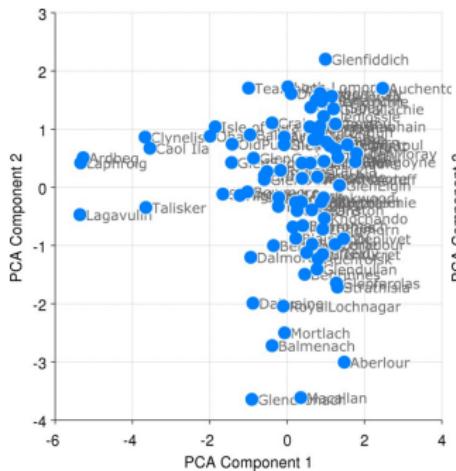


## Data example II: Whisky

- 86 types of Scotch whisky
  - Human ratings 1-5
  - 12 taste categories
    - body, sweetness, smoky, medicinal, tobacco, honey, spicy, winey, nutty, malty, fruity, floral



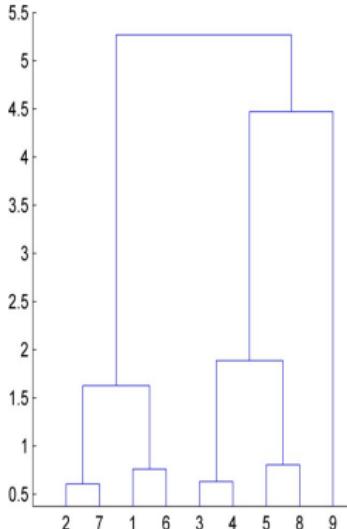
## PCA plot



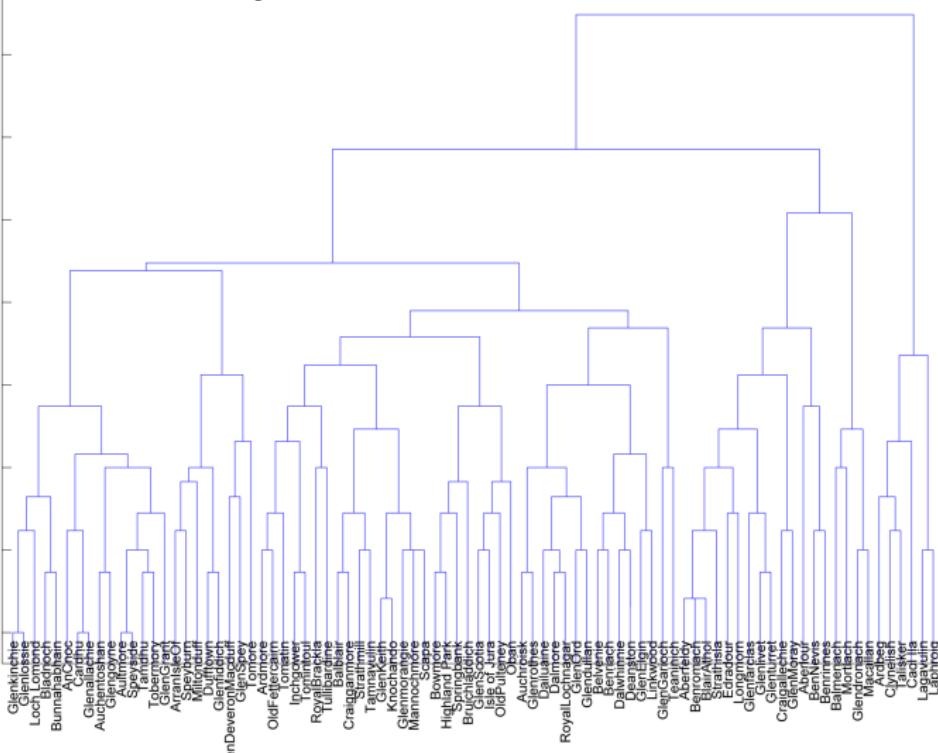
# Dendrogram

- Dendograms can be used to visualize relative distances between the observations

**Data I: Cats, Dogs and Dinosaurs**



**Data II: Whisky**



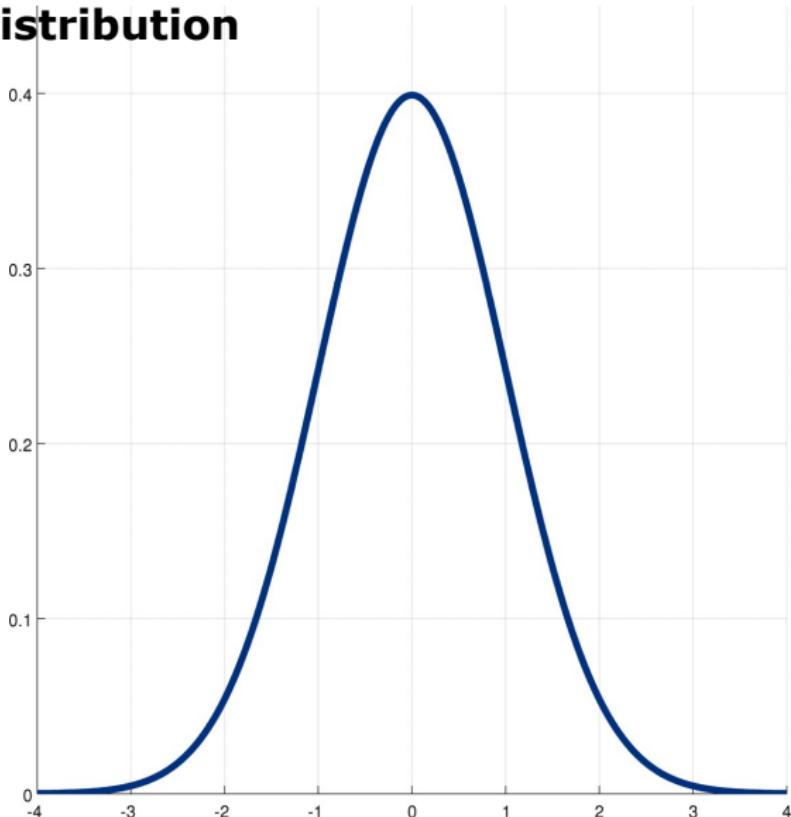
# Density based techniques: Univariate normal distribution

- Map attribute to standard Normal variable

$$z = \frac{x - \mu}{\sigma}$$

- Choose a threshold

c	$p( z >c)$
1.0	0.3173
1.5	0.1336
2.0	0.0455
2.5	0.0124
3.0	0.0027
3.5	0.0005
4.0	0.0001



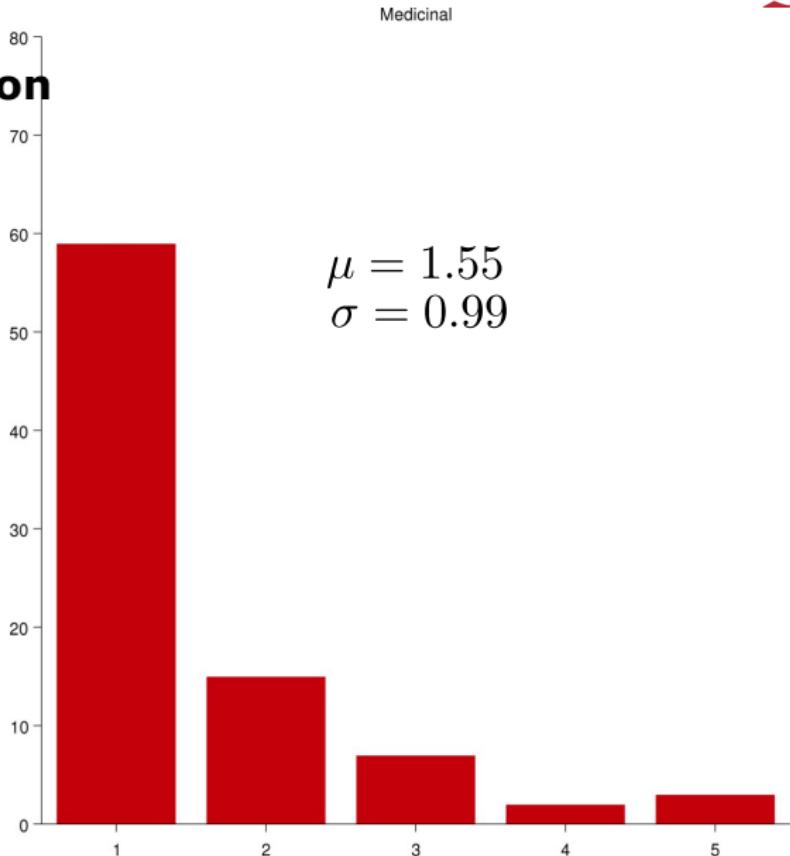
# Normal distribution

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Medicinal: z-score

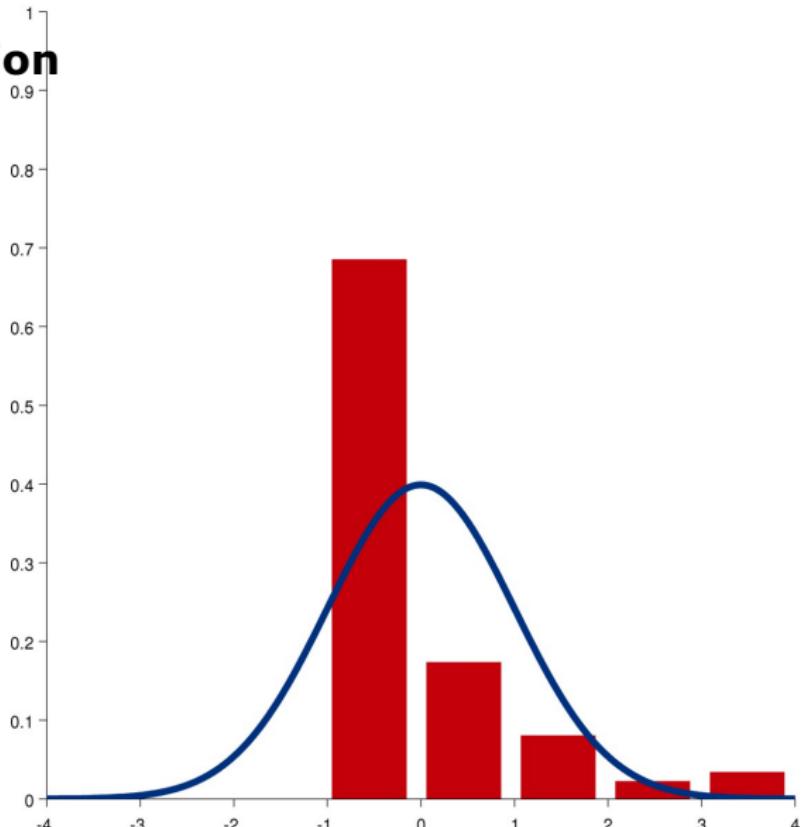
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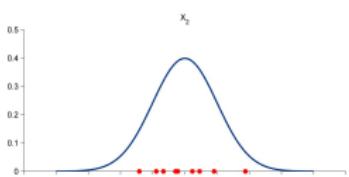
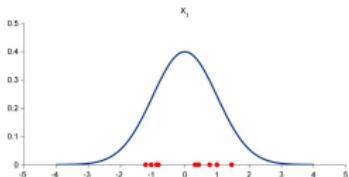
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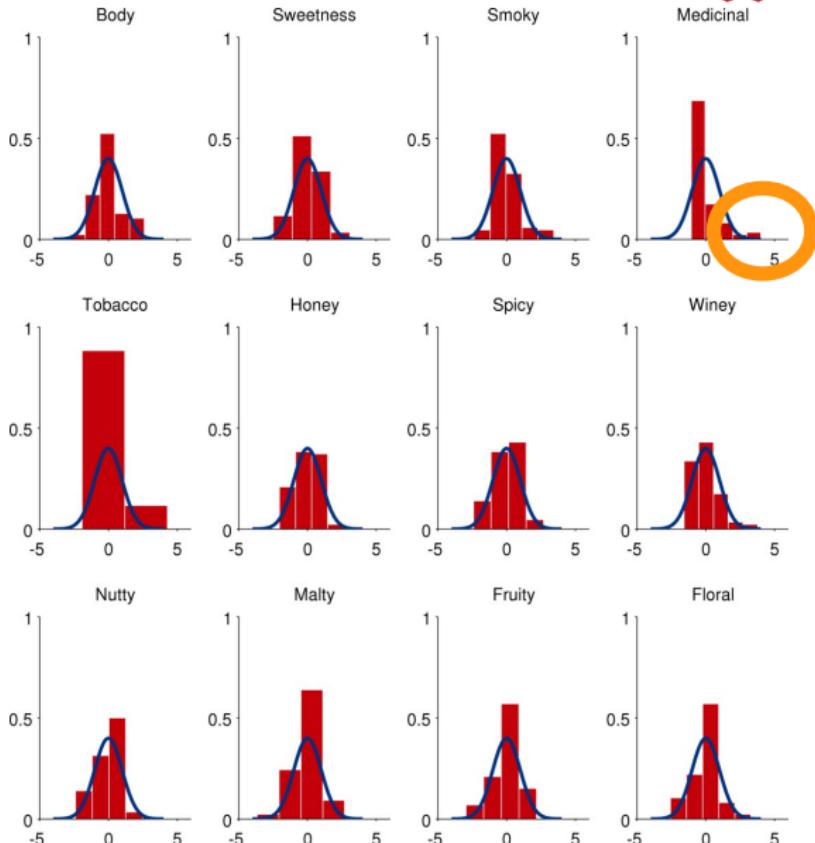
$$z = \frac{x - \mu}{\sigma}$$

$$p(|z| > c) = 0.001$$
$$c = 3.2905$$

## Data I: Cats, Dogs and Dinosaurs



## Data II: Whisky



Medicinal: z-score

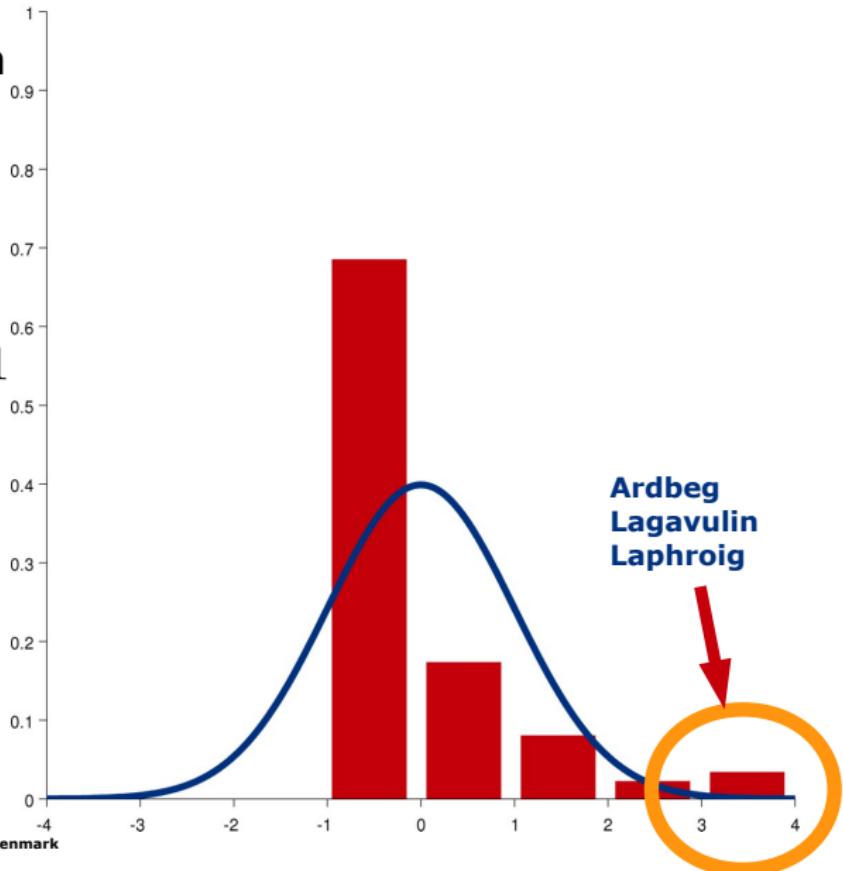
# Normal distribution

- Map attribute to standard Normal variable

$$z = \frac{x - \mu}{\sigma}$$

- Choose a threshold

$$p(|z| > c) = 0.001$$
$$c = 3.2905$$



# Approaches to anomaly detection

- **Density-based techniques**
  - Estimate the density of data objects
  - Outliers are:
    - Data objects in low density area
- **We can of course use the GMM to evaluate the density of test data.**
  - why not on the training data?**
- **Approaches we will presently also consider:**
  - Kernel density estimation
  - Inverse average distance to K nearest neighbours (KNN density)
  - Average relative KNN density

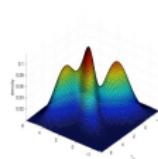
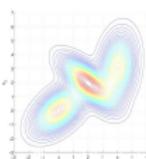
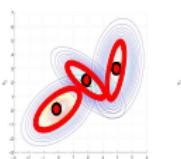
# Density based techniques: Kernel Density Estimator

Remember from Last week:  
Gaussian Mixture Model (GMM)

Data density      Sum of cluster specific densities assumed normal distributed

$$p(\mathbf{x}) = \sum_{k=1}^K w_k \mathcal{N}(\mathbf{x} | \boldsymbol{\mu}_{(k)}, \boldsymbol{\Sigma}_{(k)})$$

$$(s.t. \sum_{k=1}^K w_k = 1, \quad w_k \geq 0)$$



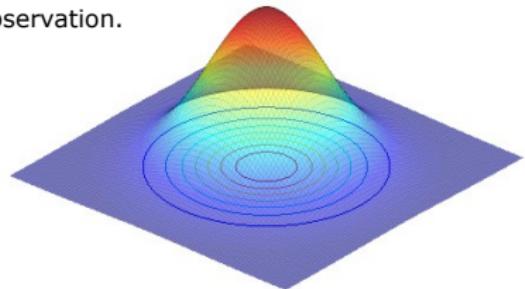
$\boldsymbol{\mu}_{(k)}$  : Cluster center (prototypical example in cluster)



$\boldsymbol{\Sigma}_{(k)}$  : Shape of the cluster

$w_k$  : Relative density of the cluster

Kernel Density estimation based on Gaussian Kernel:  
Consider the GMM and define a Gaussian with mean  $\mathbf{x}$  and co-variance  $\sigma^2 \mathbf{I}$  around each Observation.



Let all observation weight the same, i.e.  $w_n = 1/N$

$$p(\mathbf{x}) = \sum_{n=1}^N \frac{1}{N} \mathcal{N}(\mathbf{x} | \mathbf{x}_n, \sigma^2 \mathbf{I})$$

Only free parameter  $\sigma^2$ !



There is nothing special about the normal distribution. For a general mixture distribution  $p$  the general form of kernel density estimator is:

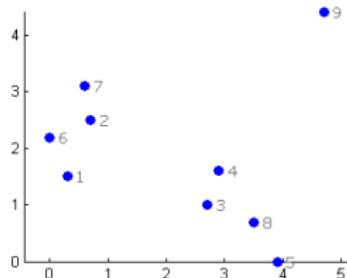
$$p(\mathbf{x}) = \sum_{n=1}^N \frac{1}{N} p(\mathbf{x} | \mathbf{x}_n, \theta)$$

This may be useful if  $\mathbf{x}$  is discrete or non-negative.



## How do we determine $\sigma^2$ ?

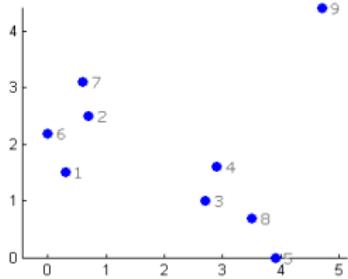
### Data I: Cats, Dogs and Dinosaurs





## How do we determine $\sigma^2$

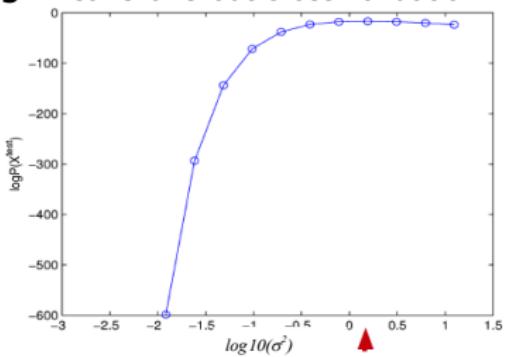
### Data I: Cats, Dogs and Dinosaurs



$$\sigma^2 = 0.01$$

$$\sigma^2 = 0.1$$

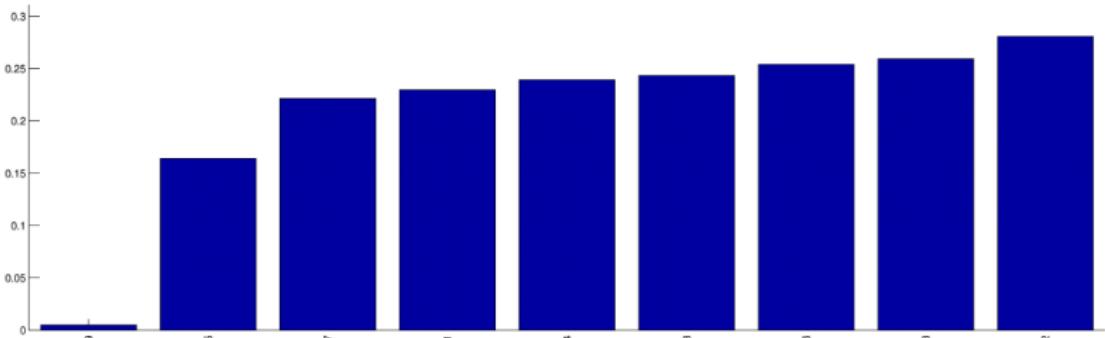
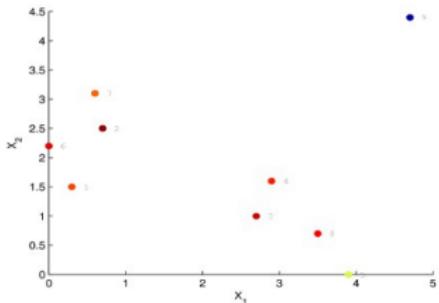
Density of test set based on leave-one-out cross validation



$$\sigma^2 = 1$$

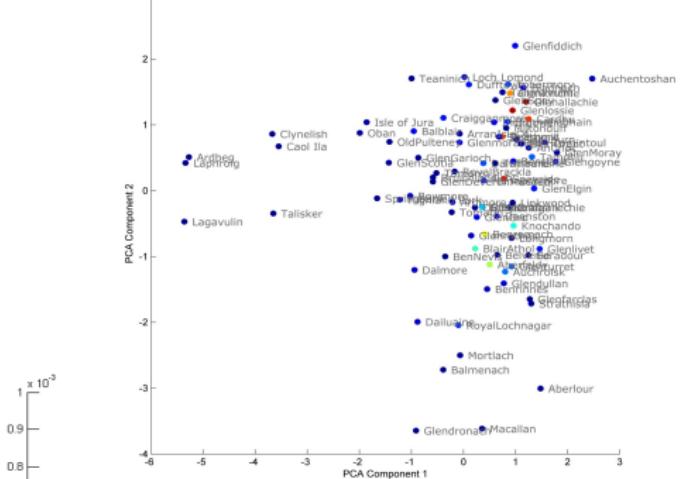
Optimal  $\sigma^2 = 1.55$   
 $\sigma^2 = 5$

# Estimated leave-one-out density evaluated at each observation

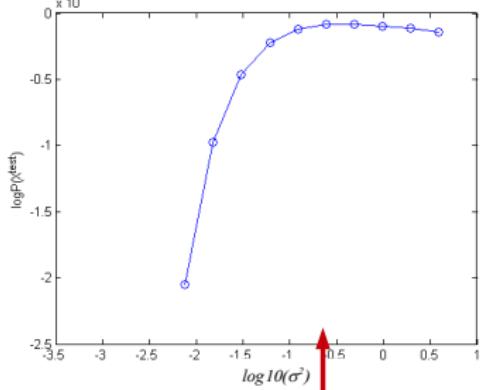


# Estimated density evaluated at each observation

## Data II: Whisky

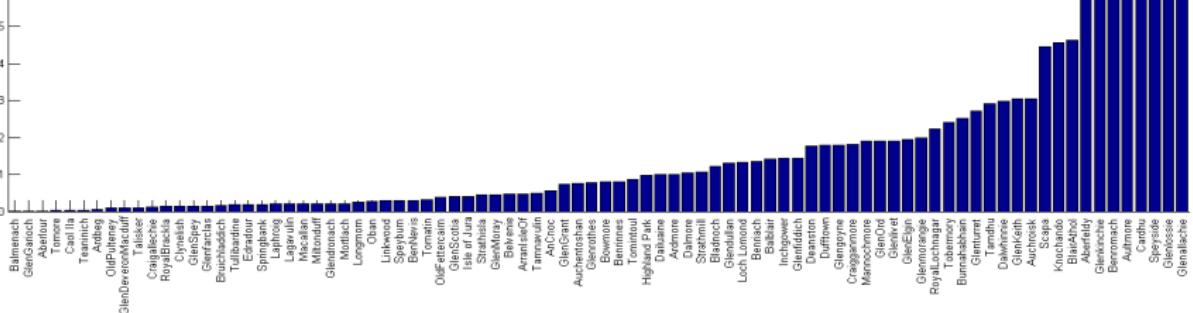


Density of test set based on leave-one-out cross validation



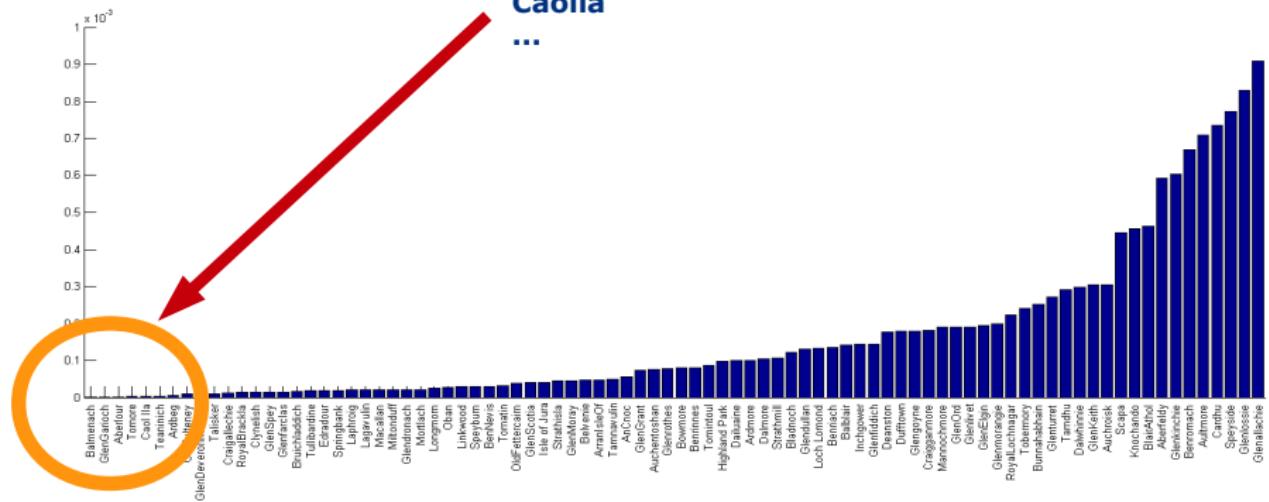
Optimal  $\sigma^2 = 0.49$

## leave-one-out densities



## Data II: Whisky

Balmenach  
Glen Garioch  
Aberlour  
Tomore  
Caolla  
...



# Inverse distance density estimation

- **Distance based measure of density**

- Density is inverse proportional to average distance to k nearest neighbors
- Density is low if nearest neighbors are far away

$$\text{density}_{\mathbf{X}_{\setminus i}}(\mathbf{x}_i, K) = \frac{1}{\frac{1}{K} \sum_{\mathbf{x}' \in N_{\mathbf{X}_{\setminus i}}(\mathbf{x}_i, K)} d(\mathbf{x}_i, \mathbf{x}')}$$

- **Relative density**

- Density compared to density at nearest neighbors


$$\text{relative density}_{\mathbf{X}}(\mathbf{x}_i, K) = \frac{\text{density}_{\mathbf{X}_{\setminus i}}(\mathbf{x}_i, K)}{\frac{1}{K} \sum_{\mathbf{x}_j \in N_{\mathbf{X}_{\setminus i}}(\mathbf{x}_i, K)} \text{density}_{\mathbf{X}_{\setminus j}}(\mathbf{x}_j, K)}$$

$N_{\mathbf{X}}(\mathbf{x}, K) = \{\text{The } K \text{ observations in } \mathbf{X} \text{ which are nearest to } \mathbf{x}\}$

$$\mathbf{X}_{\setminus i}^T = [x_1 \ x_2 \ \cdots \ x_{i-2} \ x_{i-1} \ \mathbf{x}_{i+1} \ x_{i+2} \ \cdots \ x_N]$$



Consider the pairwise distance matrix given to the left. What is the density and average relative density of the first observation for  $k=2$ ? DTU

$d(\mathbf{x}_i, \mathbf{x}_j)$	$\mathbf{x}_1$	$\mathbf{x}_2$	$\mathbf{x}_3$	$\mathbf{x}_4$	$\mathbf{x}_5$	$\mathbf{x}_6$	$\mathbf{x}_7$
$\mathbf{x}_1$	0	2.5	2.4	4.0	0.8	0.6	3.3
$\mathbf{x}_2$	2.5	0	0.6	1.6	2.9	3.0	1.1
$\mathbf{x}_3$	2.4	0.6	0	1.9	3.0	2.7	1.0
$\mathbf{x}_4$	4.0	1.6	1.9	0	4.5	4.6	3.8
$\mathbf{x}_5$	0.8	2.9	3.0	4.5	0	1.1	3.9
$\mathbf{x}_6$	0.6	3.0	2.7	4.6	1.1	0	3.8
$\mathbf{x}_7$	3.3	1.1	1.0	3.8	3.9	3.8	0

$$\text{density}_{\mathbf{X} \setminus i}(\mathbf{x}_i, K) = \frac{1}{\sum_{\mathbf{x}' \in N_{\mathbf{X} \setminus i}(\mathbf{x}_i, K)} d(\mathbf{x}_i, \mathbf{x}')}$$

$$\text{ard}_{\mathbf{X}}(\mathbf{x}_i, K) = \frac{\text{density}_{\mathbf{X} \setminus i}(\mathbf{x}_i, K)}{\sum_{\mathbf{x}_j \in N_{\mathbf{X} \setminus i}(\mathbf{x}_i, K)} \text{density}_{\mathbf{X} \setminus j}(\mathbf{x}_j, K)}$$



Consider the pairwise distance matrix given to the left. What is the density and average relative density of the first observation for  $k=2$ ? DTU

$d(\mathbf{x}_i, \mathbf{x}_j)$	$\mathbf{x}_1$	$\mathbf{x}_2$	$\mathbf{x}_3$	$\mathbf{x}_4$	$\mathbf{x}_5$	$\mathbf{x}_6$	$\mathbf{x}_7$
$\mathbf{x}_1$	0	2.5	2.4	4.0	0.8	0.6	3.3
$\mathbf{x}_2$	2.5	0	0.6	1.6	2.9	3.0	1.1
$\mathbf{x}_3$	2.4	0.6	0	1.9	3.0	2.7	1.0
$\mathbf{x}_4$	4.0	1.6	1.9	0	4.5	4.6	3.8
$\mathbf{x}_5$	0.8	2.9	3.0	4.5	0	1.1	3.9
$\mathbf{x}_6$	0.6	3.0	2.7	4.6	1.1	0	3.8
$\mathbf{x}_7$	3.3	1.1	1.0	3.8	3.9	3.8	0

$$\text{density}_{\mathbf{X} \setminus i}(\mathbf{x}_i, K) = \frac{1}{\frac{1}{K} \sum_{\mathbf{x}' \in N_{\mathbf{X} \setminus i}(\mathbf{x}_i, K)} d(\mathbf{x}_i, \mathbf{x}')}}$$

$$\text{ard}_{\mathbf{X}}(\mathbf{x}_i, K) = \frac{\text{density}_{\mathbf{X} \setminus i}(\mathbf{x}_i, K)}{\frac{1}{K} \sum_{\mathbf{x}_j \in N_{\mathbf{X} \setminus i}(\mathbf{x}_i, K)} \text{density}_{\mathbf{X} \setminus j}(\mathbf{x}_j, K)}$$

$$\text{density}(\mathbf{x}_1, 2) = [1/2(0.6+0.8)]^{-1} = 2/1.4$$

$$\text{density}(\mathbf{x}_6, 2) = [1/2(0.6+1.1)]^{-1} = 2/1.7$$

$$\text{density}(\mathbf{x}_5, 2) = [1/2(0.8+1.1)]^{-1} = 2/1.9$$

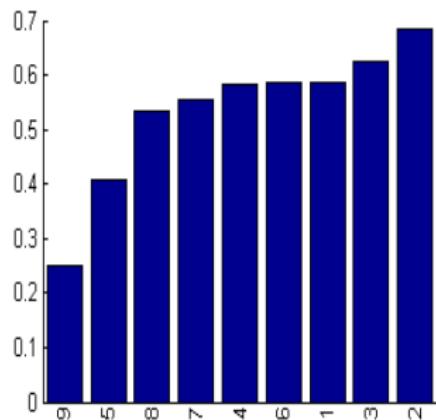
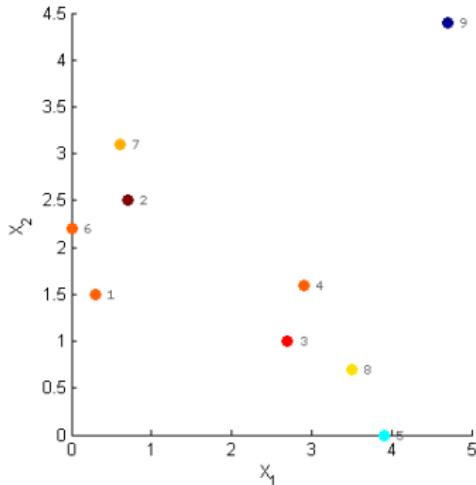


$$\text{Av. Rel. Density}(\mathbf{x}_1, 2) = [2/1.4]/[1/2 (2/1.7 + 2/1.9)] = 1.2817$$

# Inverse distance density estimation

- KNN density (5 nearest neighbors)

## Data I: Cats , dogs and dinosaurs

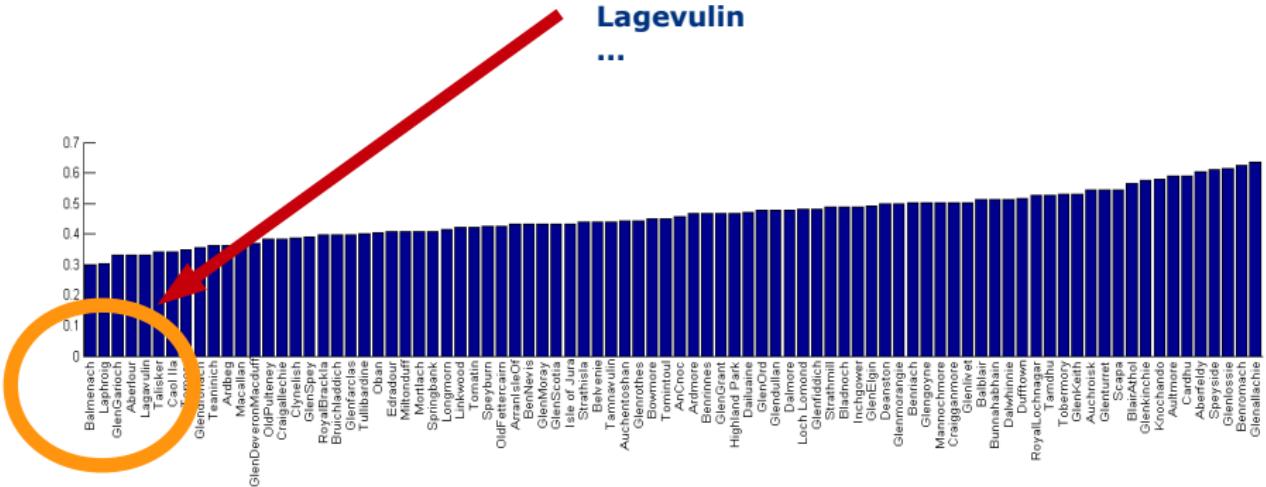


# Inverse distance density estimation

- KNN density (5 nearest neighbors)

Balmenach  
Laphroig  
GlenGarioch  
Aberlour  
Lagevulin  
...

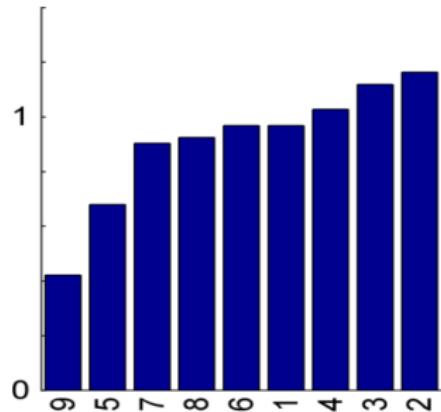
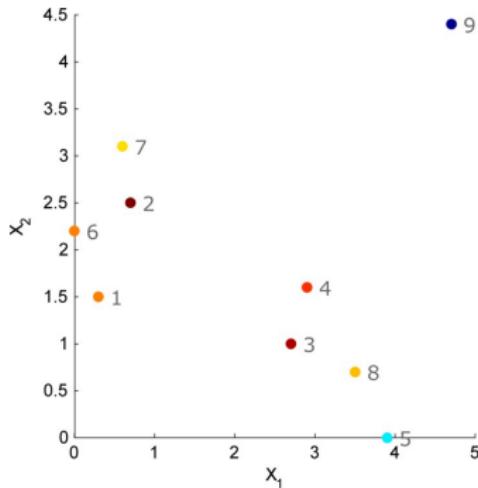
...



# Average Relative density

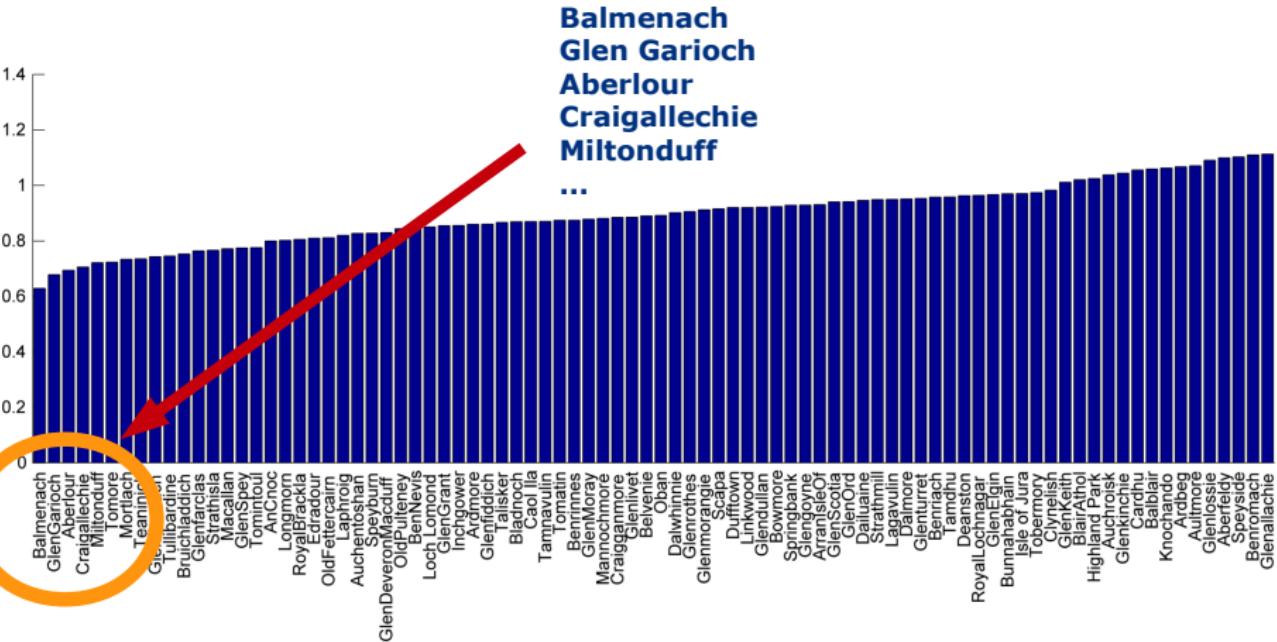
- Average Relative KNN density (5 nearest neighbors)

## Data I: Cats , dogs and dinosaurs



# Average relative density

- Average relative KNN density (5 nearest neighbors)



# Results using different methods

- **Kernel Density Estimation**

- Balmenach
- Glen Garioch
- Aberlour
- Tomore
- Caolla

- **KNN density**

- Balmenach
- Laphroig
- Glen Garioch
- Aberlour
- Lagavulin

- **KNN average relative density**

- Balmenach
- Glen Garioch
- Aberlour
- Craigallechie
- Miltonduff

**Common:** Balmenach, Glen Garioch,  
Aberlour

# Example of exam questions



Q1: What is the average relative density for observation 2 (i.e.  $\mathbf{x}_2$ ) for k=2 nearest neighbours?

- A: 1/5
- B: 3/10
- C: 7/10
- D: 1

$d(\mathbf{x}_i, \mathbf{x}_j)$	$\mathbf{x}_1$	$\mathbf{x}_2$	$\mathbf{x}_3$	$\mathbf{x}_4$	$\mathbf{x}_5$
$\mathbf{x}_1$	0	2.0	0.2	0.9	0.2
$\mathbf{x}_2$	2.0	0	1.5	0.5	2.0
$\mathbf{x}_3$	0.2	1.5	0	1.2	1.4
$\mathbf{x}_4$	0.9	0.5	1.2	0	1.0
$\mathbf{x}_5$	0.2	2.0	1.4	1.0	0

$$\text{density}_{\mathbf{X}_{\setminus i}}(\mathbf{x}_i, K) = \frac{1}{\sum_{\mathbf{x}' \in N_{\mathbf{X}_{\setminus i}}(\mathbf{x}_i, K)} d(\mathbf{x}_i, \mathbf{x}')}}$$

$$\text{ard}_{\mathbf{X}}(\mathbf{x}_i, K) = \frac{\text{density}_{\mathbf{X}_{\setminus i}}(\mathbf{x}_i, K)}{\sum_{\mathbf{x}_j \in N_{\mathbf{X}_{\setminus i}}(\mathbf{x}_i, K)} \text{density}_{\mathbf{X}_{\setminus j}}(\mathbf{x}_j, K)}$$

# Example of exam questions

QI: What is the average relative density for observation 2 (i.e.  $\mathbf{x}_2$ ) for k=2 nearest neighbours?

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$\mathbf{x}_1$	0	2.0	0.2	0.9	0.2
$\mathbf{x}_2$	2.0	0	1.5	0.5	2.0
$\mathbf{x}_3$	0.2	1.5	0	1.2	1.4
$\mathbf{x}_4$	0.9	0.5	1.2	0	1.0
$\mathbf{x}_5$	0.2	2.0	1.4	1.0	0

$$\text{density}_{\mathbf{X}_{\setminus i}}(\mathbf{x}_i, K) = \frac{1}{\frac{1}{K} \sum_{\mathbf{x}' \in N_{\mathbf{X}_{\setminus i}}(\mathbf{x}_i, K)} d(\mathbf{x}_i, \mathbf{x}')}}$$

$$\text{ard}_{\mathbf{X}}(\mathbf{x}_i, K) = \frac{\text{density}_{\mathbf{X}_{\setminus i}}(\mathbf{x}_i, K)}{\frac{1}{K} \sum_{\mathbf{x}_j \in N_{\mathbf{X}_{\setminus i}}(\mathbf{x}_i, K)} \text{density}_{\mathbf{X}_{\setminus j}}(\mathbf{x}_j, K)}$$