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Lecture "Data Mining in Bioinformatics"

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Recapitulation	Soft Clustering	Research	Conclusion
Overview			









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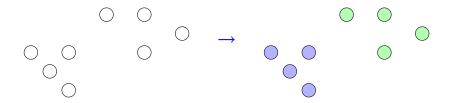






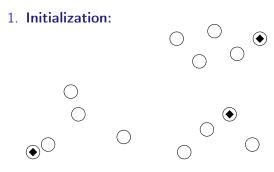
Recapitulation	Soft Clustering	Research	Conclusion
Clustering			

- Discovery of classes in a set of objects
- Unsupervised learning



# K-Means Clustering in a Nutshell

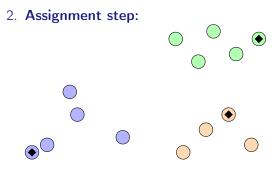
- Build k clusters
- Minimize intra-cluster variance



Pick k random points as means

Research

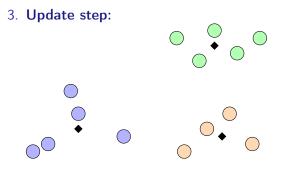
# K-Means Clustering in a Nutshell (2)



Assign each point to the nearest mean

Research

# K-Means Clustering in a Nutshell (3)



Recalculate means from corresponding points

4. Go to step 2. if at least for one point the cluster was changed

Recapitulation	Soft Clustering	Research	Conclusion
Overview			



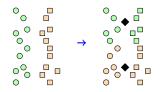




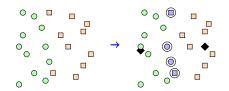




- Sometimes classes **can not** be defined by the least distance to a central point
- Examples:



• Elongated clusters often are not detected correctly



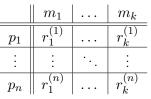
• Points on the border between two means should not be assigned to one cluster

# Taking Care of Uncertainty

- Points can belong to more than one cluster
- Clustering should reflect the **degree of association** between points and means
- Replace hard (absolute) decisions in algorithms with soft (relative) ones
- The final clustering allows interpretation of uncertain points



• New representation of associations as responsibility matrix:



•  $r_c^{(i)}$  describes the **responsibility** of cluster c for point i:

$$r_{c}^{(p)} = \frac{\exp\left(-\beta d\left(m_{c}, p_{i}\right)\right)}{\sum_{k'} \exp\left(-\beta d\left(m_{k'}, p_{i}\right)\right)}$$

 $\Rightarrow$  New parameter  $\beta$ , which describes the "stiffness" of the clustering

# Soft K-Means Clustering (2)

### • New update step:

Refine all clusters c via:

$$m_c = \frac{\sum_{i=1}^n r_c^{(i)} p_i}{\sum_{i=1}^n r_c^{(i)}} = \frac{\text{Weighted points sum}}{\text{Total responsibility}}$$

• New association step:

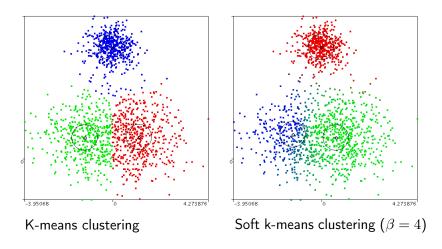
Update the responsibility matrix

Recapitulation	Soft Clustering	Research	Conclusion
C			

Stiffness

- $\bullet\,$  The stiffness  $\beta$  influences the difference to the hard k-means clustering
- Soft k-means clustering with  $\beta \to \infty$  would yield the same result as hard k-means clustering
- Figuring out the right value for  $\beta$  is non-trivial even with "try and error"

## Example



- In general soft clustering can reduce information loss due to discarding all clusters except one
- Document clustering e.g. for web search engines:
  - $\Rightarrow$  Soft clustering allows for documents to occur in several topics
- Analysis of gene expression data (microarray experiments):
  - $\Rightarrow\,$  Soft clustering decreases the sensitivity towards noise
- Prediction of molecule functions from protein-protein-interaction networks
  - ⇒ Soft clustering can assign several relevant functions to a protein

Recapitulation	Soft Clustering	Research	Conclusion
Overview			



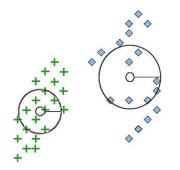


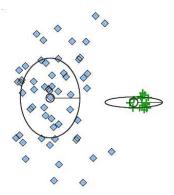


## Further Improvements of Soft K-Means Clustering

- $\bullet\,$  Choice of  $\beta\,$  decides about usefulness of the results
  - Modification using Gaussian maximum likelihood function
  - Assumption: Each cluster is **Gaussian sphere** with its own width
  - During the update step the algorithm recalculates  $\beta$  itself
  - $\Rightarrow$  Clusters with **different sizes** can be detected
- Similar enhancement using axis-aligned Gaussians is possible
  - $\Rightarrow$  Clusters with elongated shapes can be detected

## Examples





Source: MacKay, 2003

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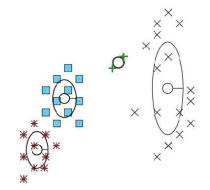






## The Last Slide

- Soft decisions can improve the chances for finding clusters with special shapes
- This can go terribly wrong and make the results worse
- Association of points to clusters is the crucial step in k-means clustering
- ⇒ Many more improvements and modifications possible
  - Applications for soft clustering are innumerable



Source: MacKay, 2003

#### Thank you for your attention.

#### Questions? Remarks?

#### **References:**



#### K. Borgwardt.

Data mining in bioinformatics. Lecture, March 2010.



D.J.C. MacKay.

Information theory, inference, and learning algorithms. Cambridge Univ Press, 2003.