

MACHINE LEARNING: CLUSTERING, AND CLASSIFICATION

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- Cross Validation?

Supervised vs. Unsupervised

- Unsupervised "clustering"
- Supervised binary classifiers (2 classes)
- Multiclass is derived from binary

Clustering

- Unsupervised learning find pockets of data to group together
- Statistical analysis techniques

Clustering

K = # of clusters

 Choosing the number of clusters – note that choosing the "best" number of clusters according to minimizing total squared distance will always result in same # of clusters as data points.

Clustering

The basic goal of clustering is to divide the data into groups such that the points within a group are close to each other, but far from items in other groups.

Hard clustering – each point is assigned to one and only one cluster.



The key points relating to k-means clustering are:

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- Clustering algorithm chooses a set of clusters with the minimum within-cluster variance



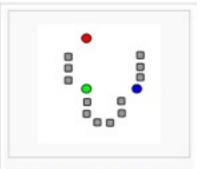
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- it requires a pre-specified number of clusters;
- Clustering algorithm chooses a set of clusters with the minimum within-cluster variance
- Guaranteed to converge (eventually)



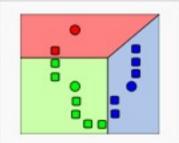
Demo

http://home.dei.polimi.it/matteucc/Clustering/tutorial_html/ http://home.dei.polimi.it/matteucc/Clustering/tutorial_html/ http://home.dei.polimi.it/matteucc/Clustering/tutorial_html/

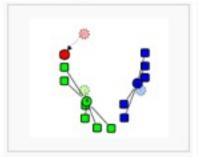
Demonstration of the standard algorithm



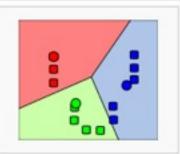
 k initial "means" (in this case k=3) are randomly selected from the data set (shown in color).



 k clusters are created by associating every observation with the nearest mean. The partitions here represent the Voronoi diagram generated by the means.



 The centroid of each of the k clusters becomes the new means.



 Steps 2 and 3 are repeated until convergence has been reached.

The initialization method needs to be further specified. There are several possible ways to initialize the cluster centers:

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- Find the mean for the whole data set then perturb into k means



ANALYSIS AND DECISION MAKING: GMMS

Mixture Models (GMM)

- K-means = hard clusters.
- GMM = soft clusters.

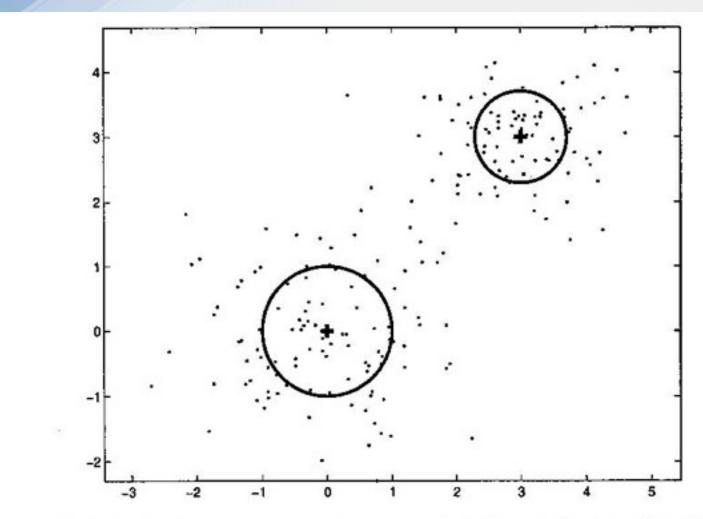


Fig. 3.1. Spherical covariance mixture model. Sampled data (dots), centres (crosses) and one standard deviation error bars (lines).

Mixture Models (GMM)

GMM is good because:

- 1. Can approximate any pdf with enough components
- 2. EM makes it easy to find components parameters
 - EM the means and variances adapt to fit the data as well as possible
- 3. Compresses data considerably
- Can make softer decisions (decide further downstream given additional information)





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 - "A prior is often the purely subjective assessment of an experienced expert."

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Query

Obtain similarity via Likelihood

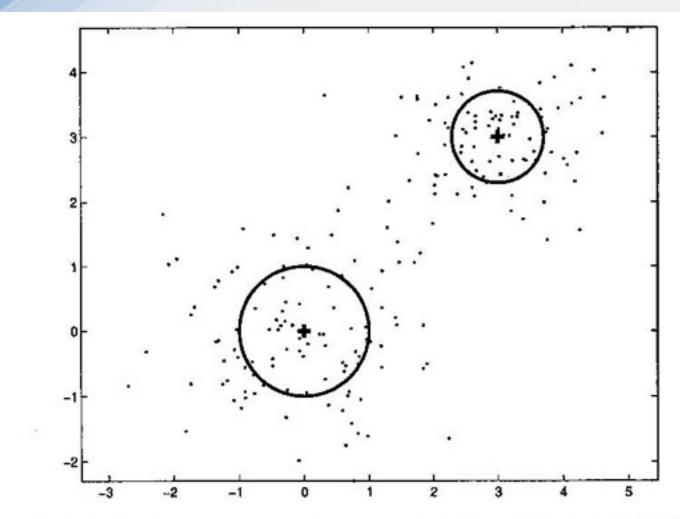
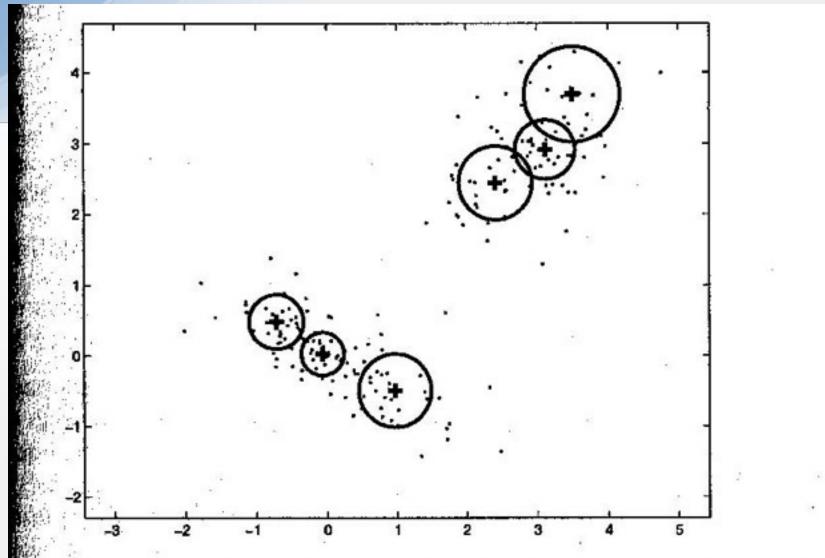


Fig. 3.1. Spherical covariance mixture model. Sampled data (dots), centres (crosses) and one standard deviation error bars (lines).



2. Spherical covariance mixture model with six components fitted to the **mpled** from the full covariance two-component model in Fig. 3.3. Sampled **tots**), centres (*crosses*) and one standard deviation error bars (*lines*).

From Netlab (p82-83)

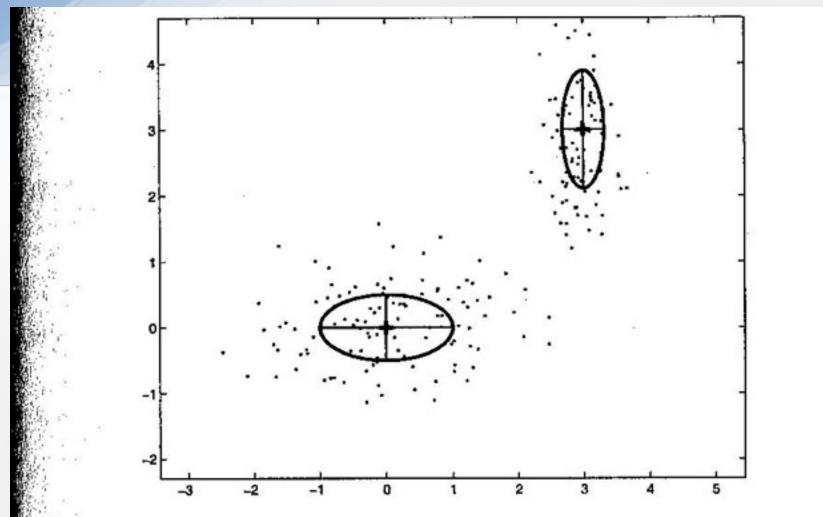
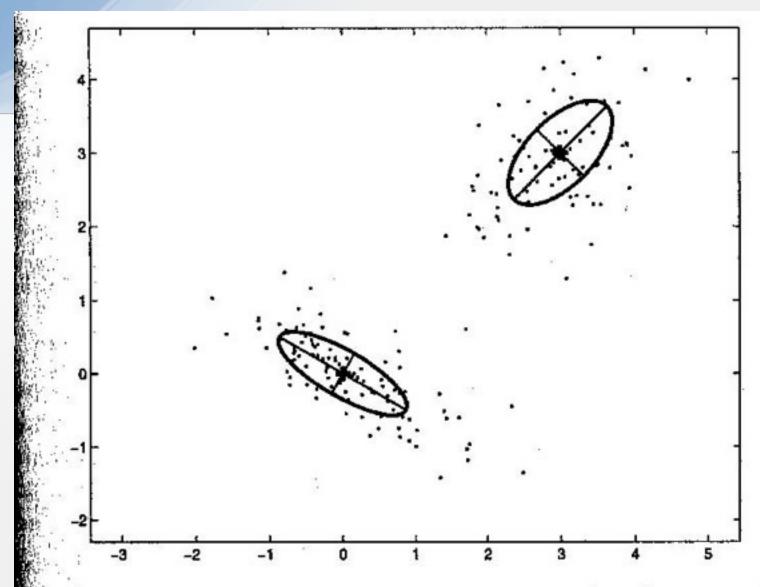


Fig. 3.2. Diagonal covariance mixture model. Sampled data (dots), centres (crosses), covariance axes (thin lines) and one standard deviation error bars (thick lines).



3. Full covariance mixture model. Sampled data (dots), centres (crosses), ace axes (thin lines) and one standard deviation error bars (thick lines).

GMM

"Pooled covariance" – using a single covariance to describe all clusters (saves on parameter computation)



Evaluate the probability of that mixture modeling your point.
 likelihoodgm1 = gmmprob
 (gm1,testing_features)



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 Log-function is "order-preserving" – maximizing a function vs. maximizing its log



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- EM is gradient-based it does not find the global maximum in the general case, unless properly initialized in the general region of interest.
- Error wants to be -inf, which occurs when Gaussian is fit for each data point. (mean = data point and variance = 0)
- "There are often a large number of local minima which correspond to poor models. Solution is to build models from many different initialization

GMM

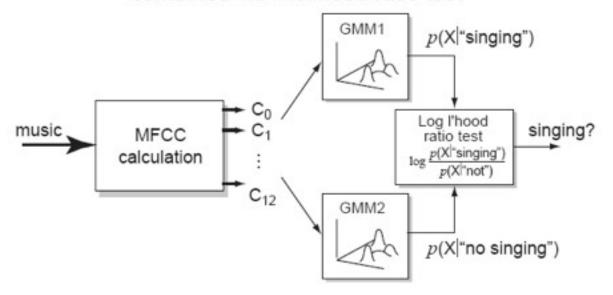
Application:

- State-of-the-art speech recognition systems
- estimate up to 30,000 separate GMMs, each with about 32 components. This means that these systems can have up to a million Gaussian components!! All the parameters are estimated from (a lot of) data by the EM algorithm.

Application: Speaker Recognition

GMM System

- Separate models for p(x|sing), p(x|no sing)
 - combined via likelihood ratio test



- How many Gaussians for each?
 - say 20; depends on data & complexity
- What kind of covariance?
 - diagonal (spherical?)





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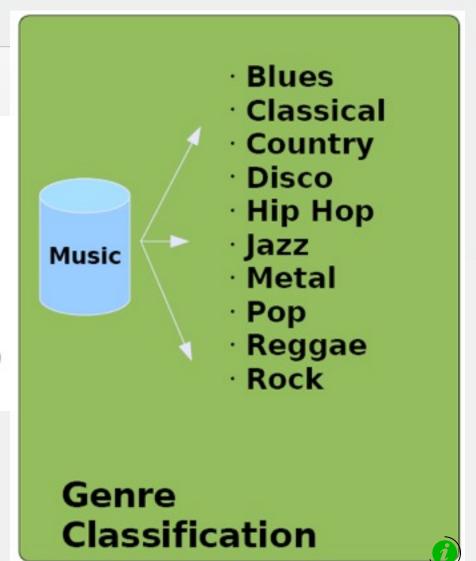
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Genre Classification:

- Manual: 72% (Perrot/Gjerdigen)
- Automated (2002) 60% (Tzanetakis)
- Automated (2005) 82% (Bergstra/Casagrande/Eck)
- Automated (2007) 76%

From ISMIR 2007 Music Recommender Tutorial (Lamere & Celma)





How?

Version 1 – One feature vector per song
- High-level features extracted from data
• Timbral (MFCCs, etc), Rhythmic content (beat

histogram, autocor, tempos), Pitch info

Sampling of the frames in the song
Statistics of features extracted from a piece (includes means, weights, etc)

- Representative of MFCC spectral shape

- Could further use "Anchor space" where classifiers are

training to represent musically meaningful classifiers. (Euclidean distance between anchor space)

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- Version 2 Cloud of points
 Extract audio every N frames
 K-Means or GMM representing a "cloud of points" for song
 Clusters: mean, covariance and weight of each cluster

= signature for song/artist/genre

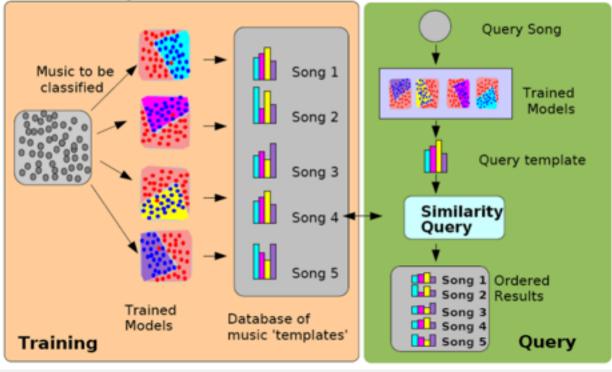
Music Recommendation

Cloud of points from frames of song

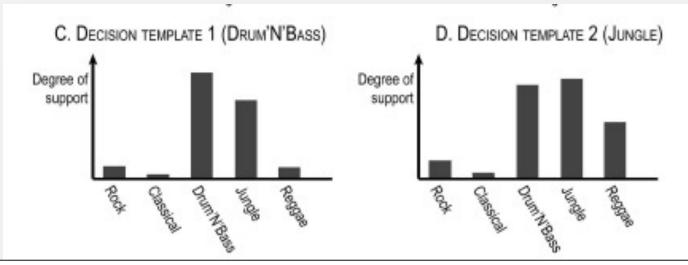
- High-level features extracted from data
- Classifier: Weighted attribute nearest neighbors or fast distance measures.
- k-Means or GMM used to create clusters.
- The mean, covariance and weight of each cluster
 signature for the song.
- Compare distance between other songs
 (signature) using various techniques to measure
 distance between probability distributions. (Most
 similar = closest distance)

Automatic annotation

Similarity based on classification



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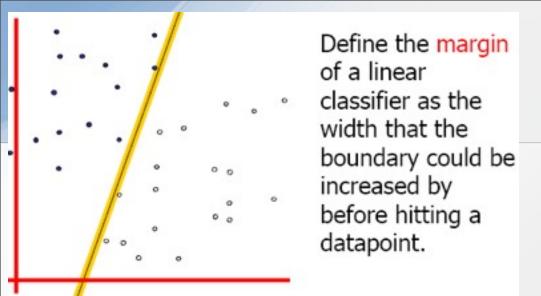




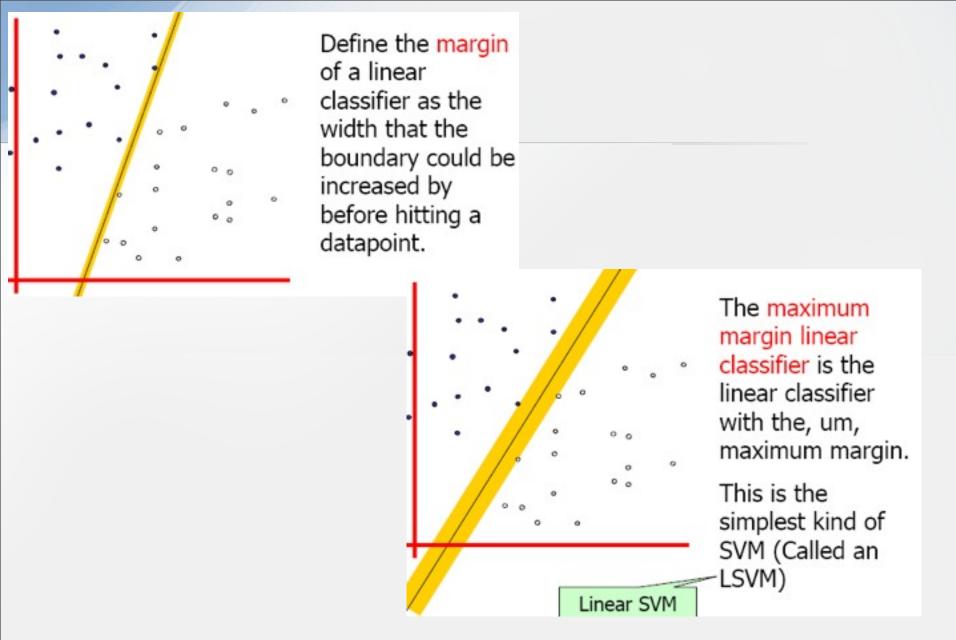
SUPPORT VECTOR MACHINES (SVM)



From: http://www.autonlab.org/tutorials/



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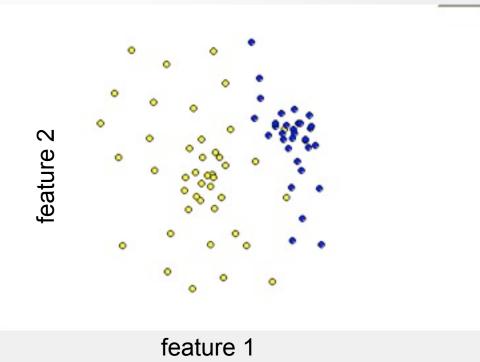
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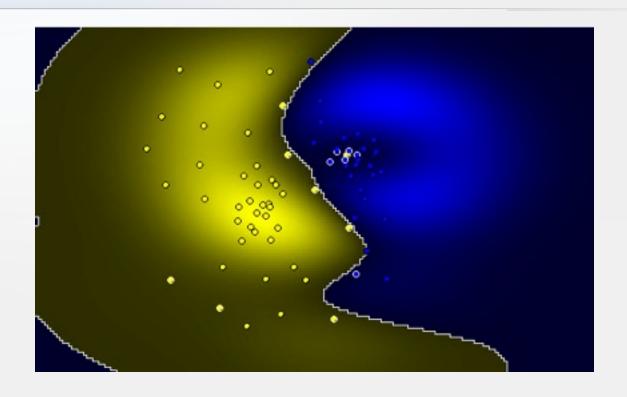
SVM

- Hyperplane separates the data from the two classes with a "maximum margin".
- Support Vectors are those data points that the margin pushes up against
- SVM training is guaranteed to find the global minimum of the cost function.
- Less experience needed fewer parameters to tune

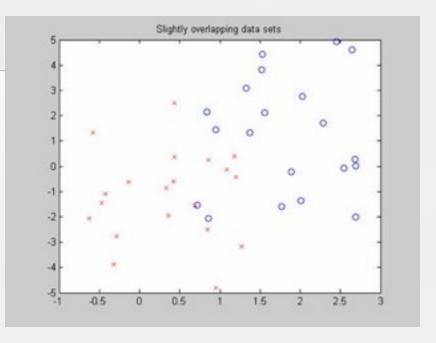
SVM with polynomial kernel visualization by Udi Aharoni

http://www.youtube.com/watch?
 v=3liCbRZPrZA





SVM Parameters



What effect do the parameters of an radial-basis-function SVM have on the separating the two data sets?

Using the RBF kernel, we have to choose values of :

gamma = degree of curviness of the hyperplane / complexity of the contour

C = allowance for points to overlap into each other's class

Video 1 Video 2

RBF Parameters: C and gamma

- Grid search using cross-validation to find the best one. Coarse then fine grid search.
- e.g., 2-5, 2-3, ... 2+15, gamma = 2-15, 2-13, 2+3

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- Grid search using cross-validation to find the best one. Coarse then fine grid search.
- e.g., 2-5, 2-3, ... 2+15, gamma = 2-15, 2-13, 2+3
- Why grid search
 - Psychological (If you have time for brute force...
 why chance it on approximations or heuristics)
 - Since there are only 2 params, grid search isn't all the different from advanced estimation techniques
 - Easily parallelized (C and gamma are

Practical Guide to SVM: The Lab

- Feature selection?
- Scale feature data
 - Save scaling stats so we can scale the test data to be in the same range
- Feature format
- Class labels {1,-1} or {0,1}
- Kernels (linear, polynomial, RBF, sigmoid)
- Find best C and gamma (cross-validation)
- Train with entire training set
- Test with validation or test set
- easy.py or grid.py