Audio signal classification

Contents:

- Introduction to pattern classification
- Features
- Feature selection
- Classification methods

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Audio signal classification

- Concrete problems
 - musical genre classification, musical instrument recognition
 - speaker recognition, language recognition
 - audio context recognition (office vs. restaurant vs. street), video segmenting based on audio, sound effects retrieval
- Closely related to sound source recognition in humans
 - source recognition includes segmentation (perceptual sound separation) in polyphonic signals
- Many efficient methods have been developed in the speech / speaker recognition field

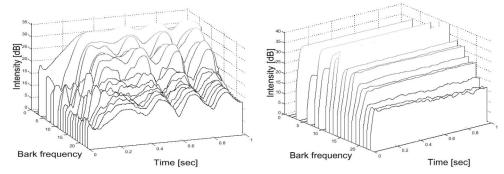
1 Introduction to pattern classification

Refresher of the basic concepts of pattern classification: Introductory slides accompanying with the book "Pattern Classification" by Duda, Hart and Stork, John Wiley & Sons, 2000.

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Example: musical instrument recognition

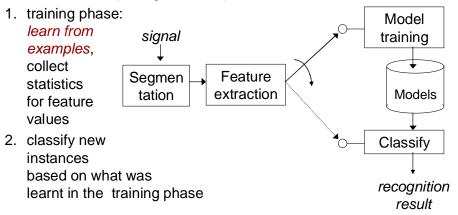
- Different acoustic properties of sound sources make them recognizable
 - properties result from sound production mechanism
- But: even a single source produces varying sounds
 - we must find someting characteristic to the source, not sound events alone: source invariants
- Examples below: flute (left) and clarinet (right) what to measure? [Eronen&Klapuri,2000]



Supervised classification system

Simplified:

- A. Feature extraction
- B. Classification (using models)



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2.1 Spectral features

- = Features that characterize the short-time spectrum
- Usually the most successful features for audio classification
- Also the most general-purpose features for different problems
 - in contrast, temporal features are typically different for musical genre recognition, instrument recognition, or speaker recognition, for example

In extracting spectral features

- Phase spectrum is typically discarded
 - 50% reduction of information
- Spectral fine structure is usually discarded (in most tasks)
 - even more reduction of irrelevant information
- → Retain only the coarse spectral energy distribution
 - effective for general audio classification
 - basis for speech and speaker recognition

2 Feature extraction

- Feature extraction is inevitable
 - time-domain signal as such contains too much irrelevant data to use it directly for classification



- Using appropriate features is crucial for successful classification
- Good features simplify the design of a classifier whereas lousy features (with little discriminating power) can hardly be compensated for by any classifier

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Spectral features

Cepstral coefficients

 Cepstral coefficients c(k) are a very convenient way to model spectral energy distribution

$$c(k) = IDFT\{\log|DFT\{x(n)\}|\}$$

where DFT denotes the Fourier-tranform and IDFT its inverse

In Matlab

```
% see "type rceps"
c = real( ifft( log( abs( fft(x)))));
```

- only real part, real(), is taken because numerical precision produces infinidecimally small imaginary part
- Cepstrum coefficients are calculated in short frames over time
 - can be further modeled by calculating e.g. the mean and variance of each coefficient over time

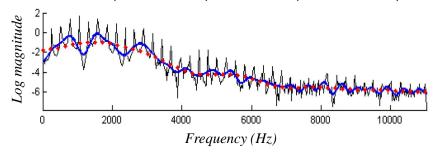
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Spectral features

Cepstral coefficients

- Only the first M cepstrum coefficients are used as features
 - all coefficients model the precise spectrum
 - coarse spectral shape is modeled by the first coefficients
 - precision is selected by the number of coefficients taken
 - the first coefficient (energy) is usually discarded
- Usually $M = f_s / (2000 \text{ Hz})$ is a good first guess for M
- Figure: piano spectrum (thin black line) and spectrum modeled with the first 50 coeffs (thick blue line) or 10 coeffs (red broken line)



Spectral features

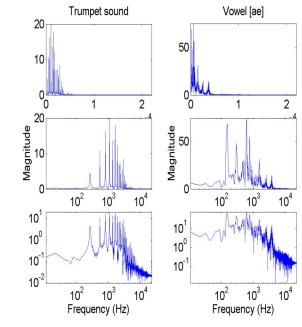
Frequency and magnitude warping

Linear scale

usually hard to "see" anything

Log-frequency

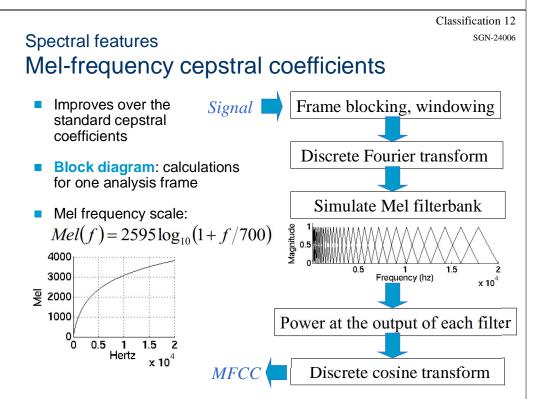
- each octave is approximately equally important perceptually
- Log-magnitude
 - perceived change from 50dB to 60dB about the same as from 60dB to 70dB



Spectral features

Cepstral coefficients

- A drawback of the cepstral coefficients: *linear frequency scale*
- Perceptually, the frequency ranges 100–200Hz and 10kHz 20kHz are approximately equally important
 - the standard cepstral coefficients do not take this into account
 - logarithmic frequency scale would be better
- Why mimic perception?
 - typically we want to classify sounds according to perceptual (dis)similarity
 - perceptually relevant features often lead to robust classification, too
- Desirable for features:
 - small change in feature vector → small perceptual change (and vice versa)
 - → Mel-frequency cepstral coefficients fulfill this criterion



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Spectral features

Mel-frequency cepstral coefficients

- Some reasons why MFCCs are successful:
 - $\begin{array}{c|c} \text{ Mel-frequency scale} \\ \text{ Log of power} \end{array} \end{array} \begin{array}{c} \textit{Large change in MFCC vector} \\ \leftrightarrow \textit{large perceptual change} \end{array}$
 - Discrete cosine transform $\left. \begin{array}{l} \textit{Drop spectral fine structure,} \\ \textit{decorrelate the features} \end{array} \right.$
- - let \mathbf{v}_{t} be the feature vector, then the first-order delta feature is

$$\mathbf{D}\mathbf{v}_{t} = \frac{\mathbf{v}_{t+1} - \mathbf{v}_{t-1}}{2}$$

 the delta features are added after the static features in the feature vector

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2.2 Temporal features

- Characterize the temporal evolution of an audio signal
- Temporal features tend to be more task-specific
- Mid-level representation for feature extraction:
 - power envelope of the signal sampled at 100Hz...1kHz rate
 - or: power envelopes of the signal at 3...40 subbands
- Note: also these drop the phase spectrum and all spectral fine structure!

Spectral features

Other spectral features

Spectral centroid (correlates with brightness)

$$SC = \left(\sum_{k=1}^{K} k \times |X(k)|^{2}\right) / \left(\sum_{k=1}^{K} |X(k)|^{2}\right)$$

where |X(k)| is the magnitude of frequency component k

Bandwidth

$$BW = \sqrt{\sum_{k=1}^{K} (k - SC)^{2} |X(k)|^{2} / \left(\sum_{k=1}^{K} |X(k)|^{2}\right)}$$

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Temporal features

- Musical instrument classification:
 - rise time: time interval between the onset and instant of maximal amplitude
 - onset asynchrony at different frequencies
 - frequency modulation: amplitude and rate (vibrato: 4-8Hz)
 - amplitude modulation: amplitude and rate
 - tremolo, roughness (fast amplitude modulation at 15-300 Hz rate)
- General audio classification
 - e.g. amplitude modulation
 - also ∆MFCCs can be seen as temporal features

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2.3 Features calculated in the time domain

- Sometimes (computationally) ultra-light features are needed and the Fourier transform is avoided
- Zero-crossing rate

Zero-crossing rate
$$ZCR = \frac{1}{N} \sum_{n=2}^{N} |sign(x(n)) - sign(x(n-1))| \qquad sign(x) = \begin{cases} 1, x > 0 \\ 0, x = 0 \\ -1, x < 0 \end{cases}$$

- Figure: ZCR correlates strongly with spectral centroid (~ brightness)
 - spectral centroid zero-crossing rate [Peltonen, MSc thesis, 2001]
- Short-time energy

$$STE = \frac{1}{N} \sum_{n=1}^{N} x(n)^2$$

lousy feature as such, but different statistics of STE are useful

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Feature selection and transformation¹

1) This section is based on lecture notes of Antti Eronen

- Let us consider methods to select a set of features from a larger set of available features
- The number of features at the disposal of the designer is usually very large (tens or even hundreds)
- Reasons for reducing the number of features to a sufficient minimum:
 - less features → simpler models → less training data needed (remember the "curse of dimensionality")
 - amount of training data: the higher the ratio of the number of training patterns N to the number of free classifier parameters, the better the *generalisation properties* of the resulting classifier
 - computational complexity, correlating (redundant) features

More features...

- **Table**: some features for musical instrument recognition [Eronen, Klapuri, 2000]
- One can measure many things...

	Table 1: Feature descriptions	
1	Rise time, i.e., the duration of attack	
2	Slope of line fitted into rms-energy curve after attack	
3	Mean square error of line fit in 2	
4	Decay time	
5	Time between the end of attack and the maximum of	
	rms-energy	
6	Crest factor, i.e., max / rms of amplitude	
7	Maximum of normalized spectral centroid	
8	Mean of normalized spectral centroid	
9	Mean of spectral centroid	
10	Standard deviation of spectral centroid	
11	Standard deviation of normalized spectral centroid	
12	Frequency of amplitude modulation, range 4-8Hz	
13	Strength of amplitude modulation, range 4-8Hz	
14	Heuristic strength of the amplitude modulation in	
	range 4-8Hz	
15	Frequency of amplitude modulation, range 10-40Hz	
16	Strength of amplitude modulation, range 10-40Hz	
17	Standard deviation of rise times at each Bark band	
18	Mean error of the fit between each of steady state	
	intensities and mean steady state intensity	
19	Mean error of fit between each of onset intensities	
	and mean onset intensity	
20	Overall variation of intensities at each band	
21	Fundamental frequency	
22	Standard deviation of fundamental frequency	
23-33	Average cepstral coefficients during onset	
34-44	Average cepstral coefficients after onset	

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Feature selection

- Feature selection (or, reduction) problem:
 - reduce the dimension of the feature vectors while preserving as much class discriminatory information as possible
- Good features result in large between-class distance and small within-class variance
- Approaches for finding good sets of features:
 - examine features individually and discard those with little discrimination ability
 - a better way is to examine the features in combinations
 - linear (or nonlinear) transformation of the feature vector

3.1 Data normalization

- Features with large values may have larger influence on the cost function than features with small values (e.g. Euclidean distance)
- For *N* available data points of the *d*th feature (d = 1, 2, ..., D);

mean:
$$m_d = \frac{1}{N} \sum_{n=1}^{N} x_{nd}$$

variance:
$$\sigma_d^2 = \frac{1}{N-1} \sum_{n=1}^{N} (x_{nd} - m_d)^2$$

normalized feature:
$$\hat{x}_{nd} = \frac{x_{nd} - m_d}{\sigma_d}$$

- The resulting normalized features have zero mean and unit variance
 - if desired, feature weighting can be performed separately after normalizat.

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3.2 Principal component analysis

- Idea: find a linear transform A such that when applied to the feature vectors x, the resulting new features y are uncorrelated.
 - \rightarrow the covariance matrix of the feature vectors y is diagonal.
- Define the covariance matrix of x by (here we use the training data!)

$$C_x = E\{(x - m)(x - m)^T\}.$$

Because C_x is real and symmetric, it is always possible to find a set of D orthonormal *eigenvectors*, and hence it can be diagonalised:

$$A C_{r} A^{T} = \Lambda$$

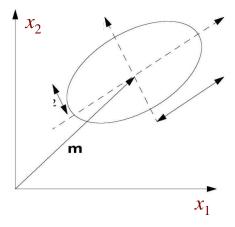
Above, A is an D by D matrix whose *rows* are formed from the eigenvectors of C_x and Λ is a diagonal matrix with the corresponding eigenvalues on its diagonal

Now the transform can be written as

$$y = A(x - m)$$

where m is the mean of the feature vectors x.

Mean, variance, correlation



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Principal component analysis

Eigenvectors

Recall the eigenvector equations:

$$Ce_d = \lambda_d e_d$$
$$(C - \lambda_d I)e_d = 0$$

where e_d are the eigenvectors and λ_d are the eigenvalues and C is square matrix

- In Matlab: [A, lambdas] = eig(C_x)
- The transform matrix A in y = A(x m) consists of the eigenvectors of C_x : $A = [e_1, e_2, ..., e_D]^T$
 - Since C_x is real and symmetric, A is orthonormal $(AA^T = I, A^T = A^{-1})$

Transformed features

■ The transformed feature vectors y = A(x - m) have zero mean and their covariance matrix (ignoring the mean terms):

$$C_y = E\{ A(\boldsymbol{x} - \boldsymbol{m}) (A(\boldsymbol{x} - \boldsymbol{m}))^T \} = AC_x A^T = \Lambda$$

i.e. C_{v} is diagonal and the features are thus *uncorrelated*.

■ Furthermore, let us denote by $\Lambda^{1/2}$ the diagonal matrix whose elements are the square roots of the eigenvalues of C_x . Then the transformed vectors

$$y = \Lambda^{-1/2} A(x - m)$$

have uncorrelated elements with unit variance:

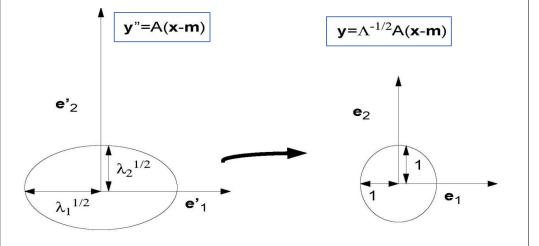
$$C_y = (\Lambda^{-1/2}A) C_x (\Lambda^{-1/2}A)^T = \dots = I.$$

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Principal component analysis

Transformed features

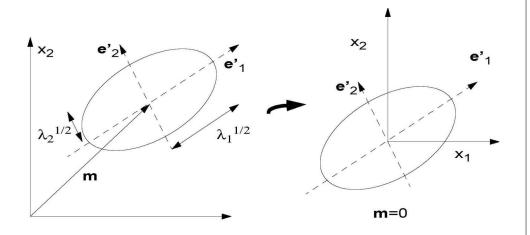
- Left: decorrelated features \rightarrow basis vectors orthogonal $e_1^T e_2 = 0$,
- Right: variance scaling \rightarrow basis vectors orthonormal $e_1^T e_1 = e_2^T e_2 = 1$



Principal component analysis

Transformed features

■ Mean removal: y' = x - m



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Principal component analysis

- The transformation can be used to reduce the amount of needed data
 - order the eigenvectors and values so that the first row of A corresponds to the largest eigenvalue and the last row to the smallest eigenvalue
 - take only the first M principal components to create an M by D matrix A for the data projection

$$\hat{\boldsymbol{x}} = A_{\scriptscriptstyle M} \left(\boldsymbol{x} - \boldsymbol{m} \right)$$

- We obtain an approximation for x, which essentially is the projection of x onto the subspace spanned by the M orthonormal eigenvectors.
- This projection is optimal in the sense that it minimises the mean square error (MSE)

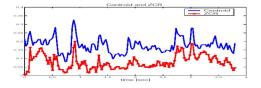
$$E\left\|\hat{\boldsymbol{x}}-\boldsymbol{x}\right\|^2$$

for any approximation with M components

Principal component analysis

Practical use of PCA

- PCA is useful for preprocessing features before classification
 - Note: PCA does not take different classes into account it only considers the properties of different features
- If two features x_i and x_j are redundant, then one eigenvalue in A is very small and one dimension can be dropped
 - we do not need to choose between two correlating features!: it is better to do the linear transform and then drop the least significant dimension
 - → both of the correlating features are utilized
- You need to scale the feature variances before eigenanalysis, since eigenvalues are proportional to the numerical range of the features
 - procedure: 1. normalize → PCA → are there unnecessary dimensions?
- Example of correlating features:
 - ZCR and spectral centroid



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Class separability measures

Between-class scatter matrix

$$S_b = \sum_{k=1}^K p(\omega_k) (\boldsymbol{m}_k - \boldsymbol{m}_0) (\boldsymbol{m}_k - \boldsymbol{m}_0)^T$$

where m_k is the mean of class k and m_0 is the global mean vector

$$\boldsymbol{m}_0 = \sum_{k=1}^K p(\boldsymbol{\omega}_k) \boldsymbol{m}_k$$

trace{S_b} is a measure of the average distance of the mean of each class from the global mean value over all classes

3.3 Class separability measures

- PCA does not take different classes into accout
 - features remaining after PCA are efficient in characterising sounds but do not necessarily discriminate between different classes
- Now we will look at a set of simple criteria that measure the discriminating properties of feature vectors
- Within-class scatter matrix (K different classes)

$$S_{w} = \sum_{k=1}^{K} p(\omega_{k}) C_{k}$$

where $C_k = E((x - m_k)(x - m_k)^T)$ is the covariance matrix for class k and $p(\omega_k)$ the prior probability of the class ω_k , i.e., $p(\omega_k) = N_k / N$, where N_k is the number of samples from class ω_k (of total N samples).

trace {S_w} is the sum of the diagonal elements of S_w and here quantifies the average within-class variance of the features over all classes

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Class separability measures

Now we can define a criterion

$$J_{1} = \frac{trace\{S_{b}\}}{trace\{S_{w}\}}$$

- It obtains large values when
 - samples in the $D\text{-}{\rm dimensional}$ space are well clustered around their mean within each class (small $\mathit{trace}\{S_w\}$)
 - the clusters of the different classes are well separated (large $trace{S_b}$)

3.4 Feature subset selection

- Problem: how to select a subset of M features from the D originally available so that
 - 1. we reduce the dimensionality of the feature vector (M < D)
 - 2. we optimize the desired class separability criterion
- To find the "optimal" subset of features, we should form all possible combinations of M features out of the D originally available
 - the best combination is then selected according to any desired class separability measure J
- In practice, it is not possible to evaluate all the possible feature combinations!

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Feature vector selection

Sequential forward selection (SFS)

- Particularly suitable for finding a few "golden" features:
- Compute criterion J value for all individual features.
 Select the best.
- 2. Form all possible two-dimensional vectors that contain the winner from the previous step. Calculate the criterion for each vector and select the best.
- 3. Continue adding features one at time, taking always the one that results in the largest value of the criterion J.
- 4. Stop when the desired vector dimension *M* is reached.
- Both SBS and SFS suffer from the nesting effect: once a feature is discarded in SBS (selected in SFS), it cannot be reconsidered again (discarded in SFS).

Feature vector selection

Sequential backward selection (SBS)

- Particularly suitable for discarding a few worst features:
- 1. Choose a class separability criterion J, and calculate its value for the feature vector which consists of *all* available features (\rightarrow length D)
- 2. Eliminate one feature, and for each possible resulting combinations (of length *D*–1) compute *J*. Select the best.
- 3. Continue this for the remaining features, and stop when you have obtained the desired dimension M.
- This is a *suboptimal* search procedure, since we cannot guarantee that the optimal *r*–1 dimensional vector has to originate from the optimal *r* dimensional one!

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3.5 Linear transforms

- Feature "generation" by combining all the D features to obtain M (with M < D) features</p>
- Define a mapping A that transforms the original feature vectors x to M-dimensional feature vectors such that

$$y = Ax$$

- Objectives:
 - 1. reduce the dimensionality of the feature vector
 - 2. optimize the desired class separability criterion
- Note the similarity with PCA the difference is that here we consider the class separability

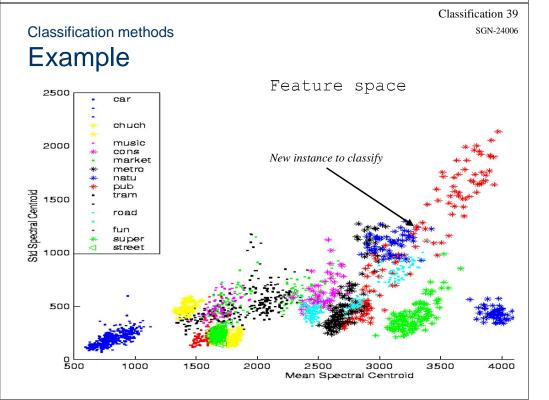
Linear transforms

Linear discriminant analysis (LDA)

Choose A to optimize

$$\boldsymbol{J}_{2} = trace \left\{ \boldsymbol{S}_{w}^{-1} \boldsymbol{S}_{b} \right\}$$

- Very similar to PCA. The difference is that here the eigenanalysis is performed for the matrix $S_w^{-1}S_b$ instead of the global covariance.
- The rows of the transform matrix are obtained by choosing the M largest eigenvectors of $S_w^{-1}S_h$
- Leads to a feature vector dimension M ≤ K-1 where K is the number of classes
 - this is because the rank of $S_w^{-1} S_b$ is K-1



4 Classification methods

- Goal is to classify previously unseen instances after the classifier has learned the training data
- Supervised vs. unsupervised classification
 - supervised: the classes of training instances are told during learning
 - unsupervised: *clustering* into hitherto unknown classes
- Supervised classification is the focus here
- Example on the next page
 - data is represented as M-dimensional feature vectors (here M=2)
 - 18 different classes
 - several training samples from each class

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4.1 Classification by distance functions

- Minimum distance classification
 - calculate the distance $D(x, y_i)$ between the unknown sample x and all the training samples y_i from all classes
 - for example the Euclidean distance $D_{E}(x,y) = \sqrt{(x-y)^{T}(x-y)}$
 - choose the class according to the closest training sample
- k-nearest neighbour (k-NN) classifier
 - pick k nearest neighbours to x and then choose the class which was most often picked
- These are a *lazy classifiers*
 - training is trivial: just store the training samples y,
 - classification gets complex with a lot of training data
 - → must measure distance to all training samples
- Computational efficiency can be improved by storing only a sufficient number of class prototypes for each class
 - or by using efficient indexing techniques (locality sensitive hashing)

Classification by distance functions

Distance metrics

- Choice of the distance metric is very important
 - Euclidean distance metric:

$$D_E(\mathbf{x}, \mathbf{y}) = \sqrt{(\mathbf{x} - \mathbf{y})^T (\mathbf{x} - \mathbf{y})} = \sqrt{\sum_d (x_d - y_d)^2}$$

sqrt is order-preserving, thus it is equivalent to minimize

$$D_E^2(x,y) = (x-y)^T(x-y)$$

Mahalanobis distance between x and y

$$D_{M}(\boldsymbol{x},\boldsymbol{y}) = (\boldsymbol{x}-\boldsymbol{y})^{T} C^{-1}(\boldsymbol{x}-\boldsymbol{y})$$

where *C* is the covariance matrix of training data.

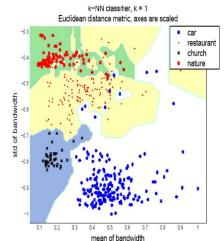
- Mahalanobis distance D_M is generally a good choice
- Comment: k-NN is a decent classifier yet very easy to implement

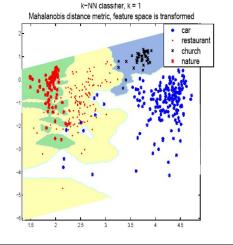
Classification by distance functions

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Example: decision boundaries for 1-NN classifier

- Left: Euclidean distance, axes scaled
- Right: Mahalanobis distance scales the axes and rotates the feature space to decorrelates the features [Peltonen, MSc thesis, 2001]



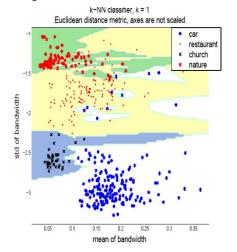


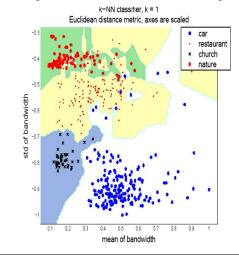
Classification by distance functions

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Example: decision boundaries for 1-NN classifier

- Left: Euclidean distance, axes not scaled
 → undesired weighting (elongated horizontal contours)
- Right: Euclidean distance, axes scaled [Peltonen, MSc thesis, 2001]





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4.2 Statistical classification

- Idea: interpret the feature vector x as a random variable whose distribution depends on the class
- Bayes' formula a *very* important tool:

$$P(\omega_i|\mathbf{x}) = \frac{P(\mathbf{x}|\omega_i)P(\omega_i)}{P(\mathbf{x})}$$

where $P(\omega_i|x)$ denotes the probability of class ω_i given an observed feature vector x

- Note: we do not know $P(\omega_i|\mathbf{x})$, but $P(\mathbf{x}/\omega_i)$ can be estimated from the training data
- Moreover, since P(x) is the same for all classes, we can perform classification based on (MAP classifier):

$$\omega_i = \operatorname{arg\,max}_i \{ P(\mathbf{x} | \omega_i) P(\omega_i) \}$$

A remaining task is to parametrize and learn $P(x/\omega_i)$ and to define (or learn) the prior probabilities $P(\omega_i)$

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Statistical classification

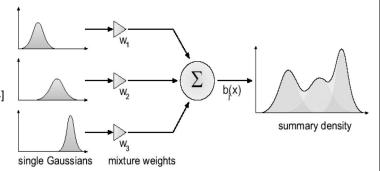
Gaussian mixture model (GMM)

- GMMs are very convenient for representing $P(x/\omega_i)$, i.e., the multidimensional distribution of features for the class ω_i
- Weighted sum of multidimensional Gaussian distributions

$$P(\boldsymbol{x}|\omega_i) = \sum_{q=1}^{Q} w_{i,q} N(\boldsymbol{x}; \boldsymbol{\mu}_{i,q}, \Sigma_{i,q})$$

where $w_{i,q}$ are weights, $N(x,\mu,\Sigma)$ is a Gaussian distribution, and $\Sigma_q w_{i,q} = 1$

Figure: GMM in one dimension [Heittola. MSc thesis, 2004]



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Statistical classification

MAP classification

• After having learned $P(x/\omega_i)$ and $P(\omega_i)$ for each class, maximum a posteriori classification of a new instance x:

$$\omega_i = \arg\max_i \{ P(\mathbf{x}|\omega_i) P(\omega_i) \}$$

■ Usually we have a sequence of feature vectors $x_{1:T}$ and

$$\omega_i = \arg\max_i \left\{ \prod_{t=1}^T P(\mathbf{x}_t | \omega_i) P(\omega_i) \right\}$$

(assuming that feature vectors are i.i.d. given the class)

since logarithm is order-preserving we can maximize

$$\omega_{i} = \arg\max_{i} \left\{ \log \left(\prod_{t=1}^{T} P(\mathbf{x}_{t} | \omega_{i}) P(\omega_{i}) \right) \right\}$$

$$= \arg\max_{i} \left\{ \sum_{t=1}^{T} \log \left[P(\mathbf{x}_{t} | \omega_{i}) P(\omega_{i}) \right] \right\}$$

Gaussian mixture model (GMM)

GMMs can fit any distribution given a sufficient number of Gaussians

- successful generalization requires limiting the number of components

$$P(\mathbf{x}|\omega_i) = \sum_{q=1}^{Q} w_{i,q} N(\mathbf{x}; \boldsymbol{\mu}_{i,q}, \boldsymbol{\Sigma}_{i,q})$$

$$N(\mathbf{x}; \boldsymbol{\mu}, \boldsymbol{\Sigma}) = \frac{1}{\sqrt{(2\pi)^D |\boldsymbol{\Sigma}|}} \exp\left(-\frac{1}{2}(\mathbf{x} - \boldsymbol{\mu})^T \boldsymbol{\Sigma}^{-1} (\mathbf{x} - \boldsymbol{\mu})\right)$$

- GMM is a parametric model
 - parameters: weights $w_{i,q}$, means $\pmb{\mu}_{i,q}$, covariance matrices $\Sigma_{i,q}$
 - diagonal covariance matrices can be used if features are decorrelated → *much fewer* parameters
- Parameters of the GMM of each class are estimated to maximize $P(x/\omega_i)$, i.e., probability of the training data for each class ω_i
 - iterative EM algorithm
- Toolboxes: insert training data → get the model → use to classify new

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Statistical classification

Hidden Markov model (HMM)

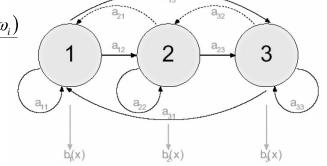
- HMMs account for the time-varying nature of sounds
 - probability of observation x in state j: typically GMM model $b_i(x)$
 - transition probabilities btw states: matrix A with $a_{ii} = P(q_i = i | q_{t-1} = j)$
 - the state variable is hidden, and usually not of interest in classification
- Train own HMM for each class
- computing

 $P(\omega_i|\mathbf{x}_{1:T}) = \frac{P(\mathbf{x}_{1:T}|\omega_i)P(\omega_i)}{P(\mathbf{x}_{1:T})}$

where $x_{1:T}$ is an observation sequence

HMMs allow

Figure: [Heittola. MSc thesis, 2004]



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Statistical classification

Hidden Markov model (HMM)

- Hidden Markov models are a generalization over GMMs
 - GMM = HMM with only one state (when using GMMs as stateconditional observation distributions)
- HMMs do not necessarily perform better than GMMs
 - → only if some temporal information is truly learned to the state transition probability matrix, then HMMs are useful → e.g. speech
 - Example: genre classification, general audio classification
- In speech recogntion, HMMs are a key technology (sequential order of phonemes is crucial)
- More about HMMs on speech recognition lectures