

# Face Processing

from face detection to face recognition

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# Overview

Part 1 Introduction

Part 2 Pre-requisites

Part 3 Face Detection

Part 4 **Face Recognition**

# Outline

## 1 Face Recognition

- Face Recognition
- Challenges
- Normalization
- Feature Extraction
- Classification
- Statistical Model based Approaches
- Evaluation
- Research Directions































# Feature Extraction

## Local

- Local PCA, Padgett and Cottrell [1997]
- 2D Gabor Wavelet, Daugman [1985], Lades [1993]: Gabor filters are known as good feature detectors and such filters remove most of the variability in images that is due to variations in lighting.
- 2D Discrete Cosine Transform: Face images are analyzed on a block by block basis. Each block is decomposed in terms of 2D Discrete Cosine Transform (DCT) basis functions. A feature vector for each block is then constructed with the DCT coefficients.
- Modification of the 2D DCT: Sanderson [2002] proposed the DCTmod2, where the first three DCT coefficients are replaced by their respective horizontal and vertical deltas in order to reduce the effects of illumination direction changes.
- Local Binary Patterns: Ahonen [2005] proposed to use LBP histograms computed in face regions

# Principal Component Analysis (PCA)

## Eigenfaces

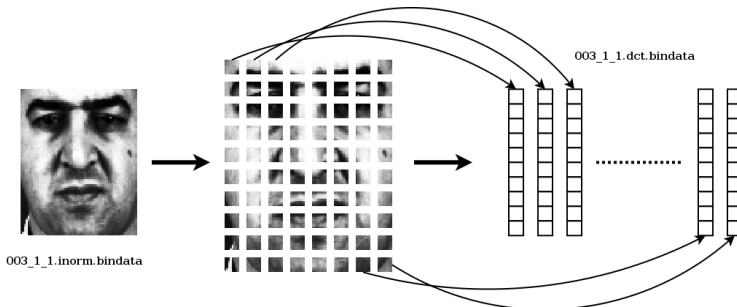
- $x_i \in \mathbb{R}^n$  is a face image which  $n = w \times h$ ,
- $\mathbf{W} = [\mathbf{e}_1 \dots \mathbf{e}_m]^T$ :



# Discrete Cosine Transform (DCT)

## Eigenfaces

- the face image is decomposed into blocks,
- the DCT is computed for each block





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# Classification

## Definitions

- Classification consists of attributing a label to the input data and differs according to the specific task (closed or open set identification, verification)
- All system provides a score  $\Lambda_l(X)$  corresponding to an opinion on the probe face pattern  $X$  to be the identity  $l$ .
  - verification: the label is true (client) or false (impostor)
  - closed set identification: the label is the identity
  - open set identification: the label is the identity or *unknown*

# Classification

## Definitions

- verification: given a threshold  $\tau$ , the claim is accepted when  $\Lambda_I(X) \geq \tau$  and rejected when  $\Lambda_I(X) < \tau$
- closed set identification: we can recognize identity  $I^*$  corresponding to the probe face pattern  $X$  as follows

$$I^* = \arg \max_I \Lambda_I(X) \quad (1)$$

- open set identification: the recognized identity  $I^*$  corresponding to the probe face is found as follows

$$I^* = \begin{cases} \text{unknown} & \text{if } \Lambda_I(X) < \tau \quad \forall I \\ \arg \max_I \Lambda_I(X) & \text{otherwise} \end{cases} \quad (2)$$





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# Statistical Model based Approaches

## Discriminant vs Generative

- **Discriminant models** such as Multi-Layer Perceptrons or Support Vector Machines:
  - training dataset of  $l$  pairs  $(X_i, y_i)$  where  $X_i$  is a vector containing the pattern, while  $y_i$  is the class of the corresponding pattern,
  - we train one model per identity,  $y_i$  being coded as  $+1$  for patterns corresponding to this identity and as  $-1$  for patterns corresponding to an other identity,
  - Drawback: difficulty to train them with a small training dataset.
- **Generative models** estimate the likelihood of the face image being a specific identity using models representing identities.

# Generative Models

Simple to complex models to compute  $\Lambda_C(X) = P(X|\lambda_C)$

- Gaussian Mixture Models (GMM) [Sanderson:2003],
- 1D Hidden Markov Models (1D-HMM) [Eickeler:2000],
- Pseudo-2D Hidden Markov Models (P2D-HMM) [Nefian:1999],
- Bayesian Networks (BNface) [Heusch:2009].

# Generative Models

## Training

- using the Maximum Likelihood (ML) criterion via the Expectation Maximization (EM),

A lot of data is required to properly estimate model parameters.

- using a well trained generic (non-person specific) model as the starting point for ML training,

ML training still produces poor models.

- using Maximum *a Posteriori* (MAP) training [Gauvain:1994] (also called *MAP adaptation*).

This approach derives a client specific model from a generic model and circumvents the lack of data problem.

# Generative Models

## Decision

- Let us denote the parameter set for client  $C$  as  $\lambda_C$  and the parameter set describing a generic face (non-client specific) as  $\lambda_{\bar{C}}$ .
- Given a claim for client  $C$ 's identity and a set of  $T$  feature vectors  $X = \{\vec{x}_t\}_{t=1}^T$  supporting the claim (extracted from the given face).
- We find an opinion on the claim using
$$\Lambda(X) = \log P(X|\lambda_C) - \log P(X|\lambda_{\bar{C}})$$

# Generative Models

## Decision

- We find an opinion on the claim using
$$\Lambda(X) = \log P(X|\lambda_C) - \log P(X|\lambda_{\bar{C}})$$
where:
  - $P(X|\lambda_C)$  is the likelihood of the claim coming from the true claimant
  - $P(X|\lambda_{\bar{C}})$  is the likelihood of the claim coming from an impostor.
- The generic face model (also called *world model* or *Universal Background Model*) is trained with data from many people.
- The decision is then reached as follows: given a threshold  $\tau$ , the claim is accepted when  $\Lambda(X) \geq \tau$  and rejected when  $\Lambda(X) < \tau$ .

# Gaussian Mixture Model

## GMM

- The likelihood of a set of feature vectors is given by

$$P(X|\lambda) = \prod_{t=1}^T P(\vec{x}_t|\lambda) \quad (3)$$

where

$$P(\vec{x}|\lambda) = \sum_{k=1}^{N_G} m_k \mathcal{N}(\vec{x}|\vec{\mu}_k, \Sigma_k) \quad (4)$$

$$\lambda = \{m_k, \vec{\mu}_k, \Sigma_k\}_{k=1}^{N_G} \quad (5)$$

# Gaussian Mixture Model

## GMM

- The likelihood of a set of feature vectors is given by

$$P(X|\lambda) = \prod_{t=1}^T \sum_{k=1}^{N_G} m_k \mathcal{N}(\vec{x}^t | \vec{\mu}_k, \Sigma_k) \quad (6)$$

- $\mathcal{N}(\vec{x} | \vec{\mu}, \Sigma)$  is a  $D$ -dimensional Gaussian density function with mean  $\vec{\mu}$  and diagonal covariance matrix  $\Sigma$ .
- $N_G$  is the number of gaussians and  $m_k$  is the weight for gaussian  $k$  (with constraints  $\sum_{k=1}^{N_G} m_k = 1$  and  $\forall k : m_k \geq 0$ ).



# Gaussian Mixture Model

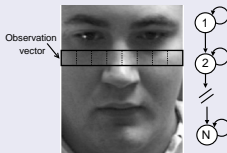
## GMM

- Generally, each feature vector  $X$  describes a different part of the face (a local approach).
- We note that the spatial relations between face parts are lost (the position of each part does not matter in the likelihood estimation).
  - Advantage: this lead to a robustness to imperfect localization of the face,
  - Drawback: discriminatory information carried by spatial relations is lost. Fortunately, there is a simple way to restore a degree of spatial relations.

# 1D-Hidden Markov Model

## 1D HMM

- The face is represented as a sequence of overlapping *rectangular* blocks from top to bottom of the face:



- The model is characterized by the following:
  - $N$ , the number of states in the model,
  - The state transition matrix  $A = \{a_{ij}\}$ ,
  - The state probability distribution  $B = \{b_j(\vec{x}_t)\}$ .

# 1D-Hidden Markov Model

$N$  the number of states in the model

each state corresponds to a region of the face;

$S = \{S_1, S_2, \dots, S_N\}$  is the set of states. The state of the model at row  $t$  is given by  $q_t \in S$ ,  $1 \leq t \leq T$ , where  $T$  is the length of the observation sequence (number of rectangular blocks).

The state transition matrix  $A = \{a_{ij}\}$

The topology of the 1D-HMM allows only self transitions or transitions to the next state:

$$a_{ij} = \begin{cases} P(q_t = S_j | q_{t-1} = S_i) & \text{for } j = i, j = i + 1 \\ 0 & \text{otherwise} \end{cases} \quad (7)$$

The state probability distribution  $B = \{b_j(\vec{x}_t)\}$

$$b_j(\vec{x}_t) = P(\vec{x}_t | q_t = S_j) \quad (8)$$

# 1D Hidden Markov Model

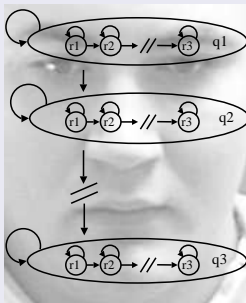
## 1D HMM

- Compared to the GMM approach the spatial constraints are much more strict, mainly due to the rigid preservation of horizontal spatial relations (e.g. distance between the eyes).
- The vertical constraints are more relaxed, though they still enforce the top-to-bottom segmentation (e.g. the eyes have to be above the mouth).
- The relaxation of constraints allows for a degree of vertical translation and some vertical stretching (caused, for example, by an imperfect face localization).

# Pseudo-2D Hidden Markov Model

## 2D HMM

- Emission probabilities of the HMM (now referred to as the “main HMM”) are estimated through a secondary HMM (referred to as an “embedded HMM”):



- The states of the embedded HMMs are in turn modeled by a mixture of gaussians.

# Pseudo-2D Hidden Markov Model

## 2D HMM

- The degree of spatial constraints present in the P2D-HMM approach can be thought of as being somewhere in between the GMM and the 1D-HMM approaches. While the GMM approach has no spatial constraints and the 1D-HMM has rigid horizontal constraints, the P2D-HMM approach has relaxed constraints in both directions.
- However, the constraints still enforce the left-to-right segmentation of the embedded HMMs (e.g. the left eye has to be before the right eye), and top-to-bottom segmentation (e.g. like in the 1D-HMM approach, the eyes have to be above the mouth). The relaxed constraints allow for a degree of both vertical and horizontal translations, as well as some vertical and horizontal stretching of the face.

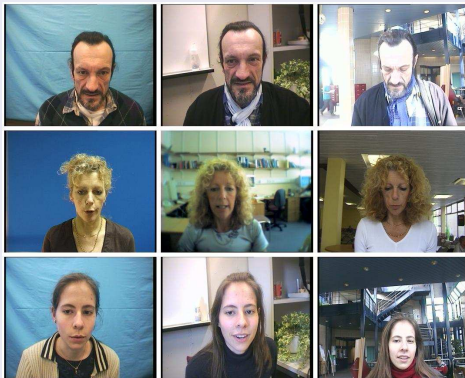
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# Database

## BANCA

- BANCA (English) database with realistic conditions:  
*controlled, degraded and adverse*

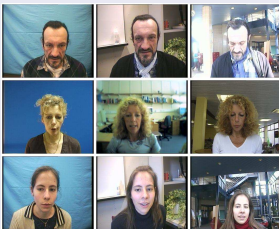




# Database

## BANCA

- 12 recording sessions over several months, in different conditions and with different cameras,
- high variability in illumination, pose, resolution, background and quality of the camera.



# Protocols

## BANCA Protocols

- 7 distinct configurations that specify which images can be used for training and testing:

Test Sessions	Train Sessions			
	1	5	9	1,5,9
C: 2-4 I: 1-4	Mc			
C: 6-8 I: 5-8	Ud	Md		
C: 10-12 I: 9-12	Ua		Ma	
C: 2-4,6-8,10-12 I: 1-12	P			G

Matched Controlled (Mc), Matched Degraded (Md), Matched Adverse (Ma), Unmatched Degraded (Ud), Unmatched Adverse (Ua), Pooled test (P) and Grand test (G).



# Experiment Results (manual)

System	Protocol			
	Mc	Ud	Ua	P
PCA	9.5	20.9	20.8	18.4
LDA/NC	4.9	16.0	20.2	14.8
SVM	5.4	25.4	30.1	20.3
GMM <i>ML</i>	12.9	28.9	26.0	22.9
GMM <i>init</i>	12.8	29.7	28.3	23.8
GMM <i>MAP</i>	<b>8.9</b>	<b>17.3</b>	<b>20.9</b>	<b>17.0</b>
1D-HMM <i>ML</i>	9.1	17.8	17.1	15.9
1D-HMM <i>init</i>	9.1	<b>15.6</b>	17.4	<b>14.7</b>
1D-HMM <i>MAP</i>	<b>6.9</b>	16.3	<b>17.0</b>	<b>14.7</b>
P2D-HMM <i>ML</i>	9.0	19.0	18.0	17.5
P2D-HMM <i>init</i>	8.6	16.5	19.2	17.0
P2D-HMM <i>MAP</i>	* <b>4.6</b>	* <b>15.3</b>	* <b>13.1</b>	* <b>13.5</b>

## Experiment Results (auto)

System	Protocol			
	Mc	Ud	Ua	P
PCA	22.4	29.7	33.7	29.0
LDA/NC	22.6	25.4	27.1	25.2
SVM	19.7	30.4	33.2	27.8
GMM <i>ML</i>	16.7	33.3	33.3	27.7
GMM <i>init</i>	19.8	35.0	35.1	29.7
GMM <i>MAP</i>	<b>9.5</b>	<b>21.0</b>	<b>24.8</b>	<b>19.5</b>
1D-HMM <i>ML</i>	21.0	28.8	29.5	27.0
1D-HMM <i>init</i>	21.3	30.1	31.4	28.1
1D-HMM <i>MAP</i>	<b>13.8</b>	<b>25.9</b>	<b>23.4</b>	<b>21.7</b>
P2D-HMM <i>ML</i>	12.1	25.2	26.9	22.3
P2D-HMM <i>init</i>	13.5	24.6	26.5	22.5
P2D-HMM <i>MAP</i>	* <b>6.5</b>	* <b>15.9</b>	* <b>14.7</b>	* <b>14.7</b>



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